

## MeSH Tree Changes for 2017

Type	Tree - heading
-	A01 Body Regions
-	A01.111 Anatomic Landmarks
-	A01.236 Breast
-	A01.236.249 Mammary Glands, Human
-	A01.236.500 Nipples
-	A01.378 Extremities
-	A01.378.100 Amputation Stumps
-	A01.378.610 Lower Extremity
-	A01.378.610.100 Buttocks
-	A01.378.610.250 Foot
-	A01.378.610.250.149 Ankle
-	A01.378.610.250.300 Forefoot, Human
-	A01.378.610.250.300.480 Metatarsus
-	A01.378.610.250.300.792 Toes
-	A01.378.610.250.300.792.380 Hallux
-	A01.378.610.250.510 Heel
-	A01.378.610.400 Hip
-	A01.378.610.450 Knee
-	A01.378.610.500 Leg
-	A01.378.610.750 Thigh
-	A01.378.800 Upper Extremity
-	A01.378.800.075 Arm
-	A01.378.800.090 Axilla
-	A01.378.800.420 Elbow
-	A01.378.800.585 Forearm
-	A01.378.800.667 Hand
-	A01.378.800.667.430 Fingers
-	A01.378.800.667.430.705 Thumb
-	A01.378.800.667.572 Metacarpus
-	A01.378.800.667.715 Wrist
-	A01.378.800.750 Shoulder
-	A01.456 Head
-	A01.456.313 Ear
-	A01.456.505 Face
-	A01.456.505.173 Cheek
-	A01.456.505.259 Chin

## MeSH Tree Changes for 2017

Type	Tree - heading
-	A01.456.505.420 Eye
-	A01.456.505.420.338 Eyebrows
-	A01.456.505.420.504 Eyelids
-	A01.456.505.420.504.421 Eyelashes
-	A01.456.505.580 Forehead
-	A01.456.505.631 Mouth
-	A01.456.505.631.515 Lip
-	A01.456.505.682 Nasolabial Fold
-	A01.456.505.733 Nose
-	A01.456.505.750 Parotid Region
-	A01.456.505.875 Superficial Musculoaponeurotic System
-	A01.456.810 Scalp
-	A01.456.830 Skull Base
-	A01.456.830.150 Cranial Fossa, Anterior
-	A01.456.830.165 Cranial Fossa, Middle
-	A01.456.830.200 Cranial Fossa, Posterior
-	A01.598 Neck
-	A01.598.500 Superficial Musculoaponeurotic System
-	A01.635 Organs at Risk
-	A01.719 Perineum
-	A01.923 Torso
-	A01.923.047 Abdomen
-	A01.923.047.025 Abdominal Cavity
-	A01.923.047.025.600 Peritoneum
-	A01.923.047.025.600.225 Douglas' Pouch
-	A01.923.047.025.600.451 Mesentery
-	A01.923.047.025.600.451.535 Mesocolon
-	A01.923.047.025.600.573 Omentum
-	A01.923.047.025.600.678 Peritoneal Cavity
-	A01.923.047.025.600.700 Peritoneal Stomata
-	A01.923.047.025.750 Retroperitoneal Space
-	A01.923.047.050 Abdominal Wall
-	A01.923.047.365 Groin
-	A01.923.047.412 Inguinal Canal
-	A01.923.047.849 Umbilicus
-	A01.923.176 Back

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	A01.923.176.519	Lumbosacral Region
-	A01.923.176.780	Sacrococcygeal Region
-	A01.923.600	Pelvis
-	A01.923.600.500	Lesser Pelvis
-	A01.923.600.600	Pelvic Floor
-	A01.923.761	Thorax
-	A01.923.761.800	Thoracic Cavity
-	A01.923.761.800.500	Mediastinum
-	A01.923.761.800.650	Pleural Cavity
-	A01.923.761.850	Thoracic Wall
-	A01.935	Transplant Donor Site
-	A01.941	Transplants
-	A01.941.500	Allografts
-	A01.941.750	Autografts
-	A01.941.812	Bone-Patellar Tendon-Bone Grafts
-	A01.941.843	Composite Tissue Allografts
-	A01.941.875	Heterografts
-	A01.941.937	Isografts
-	A01.947	Trigger Points
-	A01.960	Viscera
-	A02	Musculoskeletal System
New Heading	<b>A02.083</b>	<b>Aponeurosis</b>
-	A02.165	Cartilage
-	A02.165.257	Elastic Cartilage
-	A02.165.257.250	Ear Cartilage
-	A02.165.257.625	Laryngeal Cartilages
-	A02.165.257.625.083	Arytenoid Cartilage
-	A02.165.257.625.211	Cricoid Cartilage
-	A02.165.257.625.411	Epiglottis
-	A02.165.257.625.870	Thyroid Cartilage
-	A02.165.308	Fibrocartilage
-	A02.165.308.410	Intervertebral Disc
New Heading	<b>A02.165.308.410.250</b>	<b>Annulus Fibrosus</b>
New Heading	<b>A02.165.308.410.500</b>	<b>Nucleus Pulposus</b>

## MeSH Tree Changes for 2017

Type	Tree - heading	
Old Tree	A02.165.308.500	Menisci, Tibial
New Heading	A02.165.308.538	Meniscus
New Tree	A02.165.308.538.500	Menisci, Tibial
-	A02.165.308.575	Palmar Plate
-	A02.165.308.650	Plantar Plate
-	A02.165.308.800	Triangular Fibrocartilage
-	A02.165.407	Hyaline Cartilage
-	A02.165.407.150	Cartilage, Articular
-	A02.165.407.325	Costal Cartilage
-	A02.165.407.500	Laryngeal Cartilages
-	A02.165.407.500.083	Arytenoid Cartilage
-	A02.165.407.500.211	Cricoid Cartilage
-	A02.165.407.500.411	Epiglottis
-	A02.165.407.500.870	Thyroid Cartilage
-	A02.165.407.550	Nasal Cartilages
-	A02.340	Fascia
-	A02.340.424	Fascia Lata
-	A02.340.800	Tenon Capsule
-	A02.513	Ligaments
-	A02.513.170	Broad Ligament
-	A02.513.514	Ligaments, Articular
-	A02.513.514.100	Anterior Cruciate Ligament
-	A02.513.514.162	Collateral Ligaments
New Heading	A02.513.514.162.250	Collateral Ligament, Ulnar
-	A02.513.514.162.500	Lateral Ligament, Ankle
-	A02.513.514.162.600	Medial Collateral Ligament, Knee
-	A02.513.514.287	Ligamentum Flavum
-	A02.513.514.350	Longitudinal Ligaments
-	A02.513.514.475	Patellar Ligament
-	A02.513.514.538	Plantar Plate
-	A02.513.514.600	Posterior Cruciate Ligament
-	A02.513.514.800	Round Ligament of Femur
-	A02.513.901	Round Ligaments
-	A02.513.901.500	Round Ligament of Femur

## MeSH Tree Changes for 2017

Type	Tree - heading
-	A02.513.901.750 Round Ligament of Liver
-	A02.513.901.875 Round Ligament of Uterus
-	A02.633 Muscles
-	A02.633.567 Muscle, Skeletal
-	A02.633.567.050 Abdominal Muscles
New Heading	<b>A02.633.567.050.375 Abdominal Oblique Muscles</b>
-	A02.633.567.050.750 Pelvic Floor
-	A02.633.567.050.800 Rectus Abdominis
-	A02.633.567.175 Back Muscles
-	A02.633.567.175.500 Intermediate Back Muscles
-	A02.633.567.175.750 Paraspinal Muscles
-	A02.633.567.175.875 Superficial Back Muscles
-	A02.633.567.300 Deltoid Muscle
-	A02.633.567.400 Facial Muscles
New Heading	<b>A02.633.567.425 Gracilis Muscle</b>
New Heading	<b>A02.633.567.450 Hamstring Muscles</b>
-	A02.633.567.500 Laryngeal Muscles
-	A02.633.567.600 Masticatory Muscles
-	A02.633.567.600.500 Masseter Muscle
-	A02.633.567.600.700 Pterygoid Muscles
-	A02.633.567.600.850 Temporal Muscle
-	A02.633.567.650 Neck Muscles
-	A02.633.567.700 Oculomotor Muscles
-	A02.633.567.750 Palatal Muscles
-	A02.633.567.775 Pectoralis Muscles
-	A02.633.567.800 Pharyngeal Muscles
-	A02.633.567.800.360 Esophageal Sphincter, Upper
-	A02.633.567.800.680 Velopharyngeal Sphincter
-	A02.633.567.825 Psoas Muscles
-	A02.633.567.850 Quadriceps Muscle
-	A02.633.567.900 Respiratory Muscles
-	A02.633.567.900.300 Diaphragm
-	A02.633.567.900.500 Intercostal Muscles
-	A02.633.567.912 Rotator Cuff

## MeSH Tree Changes for 2017

Type	Tree - heading
-	A02.633.567.950                      Stapedius
-	A02.633.567.975                      Tensor Tympani
-	A02.633.570                            Muscle, Smooth
-	A02.633.570.210                      Esophageal Sphincter, Lower
-	A02.633.570.491                      Muscle, Smooth, Vascular
-	A02.633.570.500                      Myometrium
-	A02.633.580                            Myocardium
-	A02.633.580.680                      Papillary Muscles
-	A02.835                                 Skeleton
-	A02.835.232                            Bone and Bones
-	A02.835.232.022                      Bone-Implant Interface
-	A02.835.232.043                      Bones of Lower Extremity
-	A02.835.232.043.300                      Foot Bones
-	A02.835.232.043.300.492                      Metatarsal Bones
-	A02.835.232.043.300.710                      Tarsal Bones
-	A02.835.232.043.300.710.300                      Calcaneus
-	A02.835.232.043.300.710.780                      Talus
-	A02.835.232.043.300.800                      Toe Phalanges
-	A02.835.232.043.650                      Leg Bones
-	A02.835.232.043.650.247                      Femur
-	A02.835.232.043.650.247.343                      Femur Head
-	A02.835.232.043.650.247.510                      Femur Neck
-	A02.835.232.043.650.321                      Fibula
-	A02.835.232.043.650.624                      Patella
-	A02.835.232.043.650.883                      Tibia
-	A02.835.232.043.825                      Pelvic Bones
-	A02.835.232.043.825.108                      Acetabulum
-	A02.835.232.043.825.434                      Ilium
-	A02.835.232.043.825.548                      Ischium
-	A02.835.232.043.825.781                      Pubic Bone
-	A02.835.232.087                      Bones of Upper Extremity
-	A02.835.232.087.090                      Arm Bones
-	A02.835.232.087.090.400                      Humerus
-	A02.835.232.087.090.400.400                      Humeral Head
-	A02.835.232.087.090.700                      Radius
-	A02.835.232.087.090.850                      Ulna

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	A02.835.232.087.090.850.600	Olecranon Process
-	A02.835.232.087.227	Clavicle
-	A02.835.232.087.319	Hand Bones
-	A02.835.232.087.319.150	Carpal Bones
-	A02.835.232.087.319.150.150	Capitate Bone
-	A02.835.232.087.319.150.400	Hamate Bone
-	A02.835.232.087.319.150.500	Lunate Bone
-	A02.835.232.087.319.150.600	Pisiform Bone
-	A02.835.232.087.319.150.750	Scaphoid Bone
-	A02.835.232.087.319.150.800	Trapezium Bone
-	A02.835.232.087.319.150.805	Trapezoid Bone
-	A02.835.232.087.319.150.831	Triquetrum Bone
-	A02.835.232.087.319.350	Finger Phalanges
-	A02.835.232.087.319.550	Metacarpal Bones
-	A02.835.232.087.783	Scapula
-	A02.835.232.087.783.261	Acromion
New Heading	<b>A02.835.232.087.783.356</b>	<b>Coracoid Process</b>
-	A02.835.232.087.783.450	Glenoid Cavity
-	A02.835.232.169	Diaphyses
-	A02.835.232.251	Epiphyses
-	A02.835.232.251.352	Growth Plate
-	A02.835.232.409	Hyoid Bone
New Heading	<b>A02.835.232.570</b>	<b>Rib Cage</b>
New Tree	<a href="#">A02.835.232.570.500</a>	Ribs
New Tree	<a href="#">A02.835.232.570.500.150</a>	Cervical Rib
New Tree	<a href="#">A02.835.232.570.750</a>	Sternum
New Tree	<a href="#">A02.835.232.570.750.442</a>	Manubrium
New Tree	<a href="#">A02.835.232.570.750.825</a>	Xiphoid Bone
-	A02.835.232.730	Sesamoid Bones
New Tree	<a href="#">A02.835.232.730.500</a>	Patella
-	A02.835.232.781	Skull

## MeSH Tree Changes for 2017

Type	Tree - heading
-	A02.835.232.781.175      Cranial Fontanelles
-	A02.835.232.781.200      Cranial Sutures
-	A02.835.232.781.292      Ethmoid Bone
-	A02.835.232.781.324      Facial Bones
-	A02.835.232.781.324.502      Jaw
-	A02.835.232.781.324.502.125      Alveolar Process
-	A02.835.232.781.324.502.125.800      Tooth Socket
-	A02.835.232.781.324.502.320      Dental Arch
-	A02.835.232.781.324.502.632      Mandible
-	A02.835.232.781.324.502.632.130      Chin
-	A02.835.232.781.324.502.632.600      Mandibular Condyle
-	A02.835.232.781.324.502.645      Maxilla
-	A02.835.232.781.324.502.660      Palate, Hard
-	A02.835.232.781.324.665      Nasal Bone
-	A02.835.232.781.324.690      Orbit
-	A02.835.232.781.324.948      Turbinates
-	A02.835.232.781.324.971      Vomer
-	A02.835.232.781.324.995      Zygoma
-	A02.835.232.781.375      Frontal Bone
-	A02.835.232.781.572      Occipital Bone
-	A02.835.232.781.572.434      Foramen Magnum
-	A02.835.232.781.651      Parietal Bone
-	A02.835.232.781.670      Pterygopalatine Fossa
-	A02.835.232.781.750      Skull Base
-	A02.835.232.781.750.150      Cranial Fossa, Anterior
-	A02.835.232.781.750.165      Cranial Fossa, Middle
-	A02.835.232.781.750.400      Cranial Fossa, Posterior
-	A02.835.232.781.802      Sphenoid Bone
-	A02.835.232.781.802.662      Sella Turcica
-	A02.835.232.781.885      Temporal Bone
-	A02.835.232.781.885.444      Mastoid
-	A02.835.232.781.885.681      Petrous Bone
-	A02.835.232.834      Spine
-	A02.835.232.834.151      Cervical Vertebrae
-	A02.835.232.834.151.383      Axis, Cervical Vertebra
-	A02.835.232.834.151.383.668      Odontoid Process



## MeSH Tree Changes for 2017

Type	Tree - heading
-	A02.835.232.834.151.500                      Cervical Atlas
-	A02.835.232.834.229                              Coccyx
-	A02.835.232.834.432                              Intervertebral Disc
New Heading	<b>A02.835.232.834.432.250</b> <b>Annulus Fibrosus</b>
New Heading	<b>A02.835.232.834.432.500</b> <b>Nucleus Pulposus</b>
-	A02.835.232.834.519                              Lumbar Vertebrae
-	A02.835.232.834.717                              Sacrum
-	A02.835.232.834.803                              Spinal Canal
-	A02.835.232.834.803.350                              Epidural Space
-	A02.835.232.834.892                              Thoracic Vertebrae
Old Tree	<b>A02.835.232.904</b> <b>Thorax</b>
Old Tree	<b>A02.835.232.904.567</b> <b>Ribs</b>
Old Tree	<b>A02.835.232.904.567.150</b> <b>Cervical Rib</b>
Old Tree	<b>A02.835.232.904.766</b> <b>Sternum</b>
Old Tree	<b>A02.835.232.904.766.442</b> <b>Manubrium</b>
Old Tree	<b>A02.835.232.904.766.825</b> <b>Xiphoid Bone</b>
-	A02.835.583    Joints
-	A02.835.583.032    Acromioclavicular Joint
-	A02.835.583.097    Atlanto-Axial Joint
-	A02.835.583.101    Atlanto-Occipital Joint
-	A02.835.583.156    Bursa, Synovial
-	A02.835.583.192    Cartilage, Articular
-	A02.835.583.290    Elbow Joint
-	A02.835.583.378    Foot Joints
-	A02.835.583.378.062    Ankle Joint
-	A02.835.583.378.531    Metatarsophalangeal Joint
-	A02.835.583.378.531.500    Plantar Plate
-	A02.835.583.378.831    Tarsal Joints
-	A02.835.583.378.831.780    Subtalar Joint
-	A02.835.583.378.900    Toe Joint
-	A02.835.583.378.900.500    Plantar Plate
-	A02.835.583.405    Hand Joints
-	A02.835.583.405.174    Carpal Joints
-	A02.835.583.405.200    Carpometacarpal Joints

## MeSH Tree Changes for 2017

Type	Tree - heading
-	A02.835.583.405.350                      Finger Joint
-	A02.835.583.405.500                      Metacarpophalangeal Joint
-	A02.835.583.405.500.900                      Palmar Plate
-	A02.835.583.405.930                      Wrist Joint
-	A02.835.583.405.930.800                      Triangular Fibrocartilage
-	A02.835.583.411                      Hip Joint
-	A02.835.583.411.500                      Round Ligament of Femur
-	A02.835.583.443                      Joint Capsule
-	A02.835.583.443.800                      Synovial Membrane
-	A02.835.583.443.800.800                      Synovial Fluid
-	A02.835.583.475                      Knee Joint
-	A02.835.583.475.590                      Menisci, Tibial
-	A02.835.583.475.650                      Patellofemoral Joint
-	A02.835.583.512                      Ligaments, Articular
-	A02.835.583.512.100                      Anterior Cruciate Ligament
-	A02.835.583.512.162                      Collateral Ligaments
New Heading	<b>A02.835.583.512.162.250                      Collateral Ligament, Ulnar</b>
-	A02.835.583.512.162.500                      Lateral Ligament, Ankle
-	A02.835.583.512.162.600                      Medial Collateral Ligament, Knee
-	A02.835.583.512.287                      Ligamentum Flavum
-	A02.835.583.512.350                      Longitudinal Ligaments
-	A02.835.583.512.475                      Patellar Ligament
-	A02.835.583.512.538                      Plantar Plate
-	A02.835.583.512.600                      Posterior Cruciate Ligament
-	A02.835.583.512.900                      Palmar Plate
-	A02.835.583.512.950                      Round Ligament of Femur
-	A02.835.583.656                      Pubic Symphysis
-	A02.835.583.707                      Sacroiliac Joint
-	A02.835.583.748                      Shoulder Joint
-	A02.835.583.781                      Sternoclavicular Joint
-	A02.835.583.790                      Sternocostal Joints
-	A02.835.583.861                      Temporomandibular Joint
-	A02.835.583.861.900                      Temporomandibular Joint Disc
-	A02.835.583.979                      Zygapophyseal Joint
-	A02.880                      Tendons

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	A02.880.176	Achilles Tendon
New Heading	<b>A02.880.307</b>	<b>Hamstring Tendons</b>
-	A02.880.438	Patellar Ligament
-	A02.880.700	Rotator Cuff
-	A03	Digestive System
-	A03.159	Biliary Tract
-	A03.159.183	Bile Ducts
-	A03.159.183.079	Bile Ducts, Extrahepatic
-	A03.159.183.079.300	Common Bile Duct
-	A03.159.183.079.300.950	Ampulla of Vater
-	A03.159.183.079.300.950.600	Sphincter of Oddi
-	A03.159.183.079.450	Cystic Duct
-	A03.159.183.079.600	Hepatic Duct, Common
-	A03.159.183.158	Bile Ducts, Intrahepatic
-	A03.159.183.158.125	Bile Canaliculi
-	A03.159.439	Gallbladder
-	A03.556	Gastrointestinal Tract
-	A03.556.124	Intestines
-	A03.556.124.369	Intestinal Mucosa
-	A03.556.124.369.290	Enterocytes
-	A03.556.124.369.320	Goblet Cells
-	A03.556.124.369.700	Paneth Cells
-	A03.556.124.526	Intestine, Large
-	A03.556.124.526.070	Anal Canal
-	A03.556.124.526.209	Cecum
-	A03.556.124.526.209.290	Appendix
-	A03.556.124.526.356	Colon
-	A03.556.124.526.356.333	Colon, Ascending
-	A03.556.124.526.356.500	Colon, Descending
-	A03.556.124.526.356.668	Colon, Sigmoid
-	A03.556.124.526.356.834	Colon, Transverse
-	A03.556.124.526.767	Rectum
-	A03.556.124.684	Intestine, Small
-	A03.556.124.684.124	Duodenum
-	A03.556.124.684.124.236	Ampulla of Vater

## MeSH Tree Changes for 2017

Type	Tree - heading
-	A03.556.124.684.124.236.572 Sphincter of Oddi
-	A03.556.124.684.124.322 Brunner Glands
-	A03.556.124.684.249 Ileum
-	A03.556.124.684.249.400 Ileocecal Valve
-	A03.556.124.684.249.612 Meckel Diverticulum
-	A03.556.124.684.500 Jejunum
-	A03.556.249 Lower Gastrointestinal Tract
-	A03.556.249.124 Ileum
-	A03.556.249.124.400 Ileocecal Valve
-	A03.556.249.124.612 Meckel Diverticulum
-	A03.556.249.249 Intestine, Large
-	A03.556.249.249.070 Anal Canal
-	A03.556.249.249.209 Cecum
-	A03.556.249.249.209.290 Appendix
-	A03.556.249.249.356 Colon
-	A03.556.249.249.356.333 Colon, Ascending
-	A03.556.249.249.356.500 Colon, Descending
-	A03.556.249.249.356.668 Colon, Sigmoid
-	A03.556.249.249.356.834 Colon, Transverse
-	A03.556.249.249.767 Rectum
-	A03.556.249.750 Jejunum
-	A03.556.500 Mouth
-	A03.556.500.379 Dentition
-	A03.556.500.760 Salivary Glands
-	A03.556.500.760.464 Parotid Gland
-	A03.556.500.760.640 Salivary Ducts
-	A03.556.500.760.650 Salivary Glands, Minor
-	A03.556.500.760.687 Sublingual Gland
-	A03.556.500.760.812 Submandibular Gland
-	A03.556.500.760.906 von Ebner Glands
-	A03.556.500.885 Tongue
-	A03.556.500.885.431 Lingual Frenum
-	A03.556.500.885.779 Taste Buds
-	A03.556.750 Pharynx
-	A03.556.875 Upper Gastrointestinal Tract
-	A03.556.875.249 Duodenum

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	A03.556.875.249.160	Ampulla of Vater
-	A03.556.875.249.160.572	Sphincter of Oddi
-	A03.556.875.249.322	Brunner Glands
-	A03.556.875.500	Esophagus
New Heading	<b>A03.556.875.500.180</b>	<b>Esophageal Mucosa</b>
-	A03.556.875.500.360	Esophageal Sphincter, Upper
-	A03.556.875.500.414	Esophagogastric Junction
-	A03.556.875.500.414.350	Esophageal Sphincter, Lower
-	A03.556.875.875	Stomach
-	A03.556.875.875.163	Cardia
-	A03.556.875.875.330	Esophagogastric Junction
-	A03.556.875.875.330.350	Esophageal Sphincter, Lower
-	A03.556.875.875.419	Gastric Fundus
-	A03.556.875.875.440	Gastric Mucosa
-	A03.556.875.875.440.150	Chief Cells, Gastric
-	A03.556.875.875.440.250	Enterochromaffin Cells
-	A03.556.875.875.440.300	Gastrin-Secreting Cells
-	A03.556.875.875.440.708	Parietal Cells, Gastric
-	A03.556.875.875.440.854	Somatostatin-Secreting Cells
-	A03.556.875.875.578	Gastric Stump
-	A03.556.875.875.716	Pyloric Antrum
-	A03.556.875.875.799	Pylorus
-	A03.620	Liver
-	A03.620.150	Bile Ducts, Intrahepatic
-	A03.620.150.125	Bile Canaliculi
-	A03.620.575	Round Ligament of Liver
-	A03.734	Pancreas
-	A03.734.414	Islets of Langerhans
-	A03.734.414.065	Glucagon-Secreting Cells
-	A03.734.414.131	Insulin-Secreting Cells
-	A03.734.414.587	Pancreatic Polypeptide-Secreting Cells
-	A03.734.414.793	Somatostatin-Secreting Cells
-	A03.734.540	Pancreas, Exocrine
-	A03.734.667	Pancreatic Ducts
-	A04	Respiratory System



## MeSH Tree Changes for 2017

Type	Tree - heading
-	A04.623.490.600                      Pyriform Sinus
-	A04.623.557                              Nasopharynx
-	A04.623.557.500                      Adenoids
-	A04.623.603                              Oropharynx
-	A04.623.603.925                      Palatine Tonsil
-	A04.623.617                              Pharyngeal Muscles
-	A04.623.617.360                      Esophageal Sphincter, Upper
-	A04.623.617.680                      Velopharyngeal Sphincter
-	A04.716                                      Pleura
-	A04.760                                      Respiratory Mucosa
-	A04.760.259                              Goblet Cells
-	A04.760.520                              Laryngeal Mucosa
-	A04.760.600                              Nasal Mucosa
-	A04.760.600.640                      Olfactory Mucosa
-	A04.760.600.640.640                      Olfactory Receptor Neurons
-	A04.889                                      Trachea
-	A05    Urogenital System
-	A05.360                                      Genitalia
-	A05.360.319                              Genitalia, Female
-	A05.360.319.114                      Adnexa Uteri
-	A05.360.319.114.170                      Broad Ligament
-	A05.360.319.114.373                      Fallopian Tubes
-	A05.360.319.114.630                      Ovary
-	A05.360.319.114.630.278                      Corpus Luteum
-	A05.360.319.114.630.278.400                      Luteal Cells
-	A05.360.319.114.630.535                      Ovarian Follicle
-	A05.360.319.114.630.535.150                      Follicular Fluid
-	A05.360.319.114.630.535.200                      Granulosa Cells
-	A05.360.319.114.630.535.200.500                      Cumulus Cells
-	A05.360.319.114.630.535.400                      Theca Cells
-	A05.360.319.114.815                      Round Ligament of Uterus
-	A05.360.319.679                              Uterus
-	A05.360.319.679.256                      Cervix Uteri
-	A05.360.319.679.490                      Endometrium
-	A05.360.319.679.490.373                      Decidua
-	A05.360.319.679.490.373.500                      Deciduoma

## MeSH Tree Changes for 2017

Type	Tree - heading
-	A05.360.319.679.690 Myometrium
-	A05.360.319.779 Vagina
-	A05.360.319.779.479 Hymen
-	A05.360.319.887 Vulva
-	A05.360.319.887.220 Bartholin's Glands
-	A05.360.319.887.436 Clitoris
-	A05.360.444 Genitalia, Male
-	A05.360.444.123 Bulbourethral Glands
-	A05.360.444.251 Ejaculatory Ducts
-	A05.360.444.371 Epididymis
-	A05.360.444.492 Penis
-	A05.360.444.492.362 Foreskin
-	A05.360.444.492.726 Urethra
-	A05.360.444.575 Prostate
-	A05.360.444.661 Scrotum
-	A05.360.444.713 Seminal Vesicles
-	A05.360.444.777 Spermatic Cord
-	A05.360.444.849 Testis
-	A05.360.444.849.513 Leydig Cells
-	A05.360.444.849.600 Rete Testis
-	A05.360.444.849.700 Seminiferous Tubules
-	A05.360.444.849.700.349 Blood-Testis Barrier
-	A05.360.444.849.700.700 Seminiferous Epithelium
-	A05.360.444.849.789 Sertoli Cells
-	A05.360.444.930 Vas Deferens
-	A05.360.490 Germ Cells
-	A05.360.490.690 Ovum
-	A05.360.490.690.680 Oocytes
-	A05.360.490.690.680.500 Polar Bodies
-	A05.360.490.690.700 Oogonia
-	A05.360.490.690.950 Zona Pellucida
-	A05.360.490.690.970 Zygote
-	A05.360.490.890 Spermatozoa
-	A05.360.490.890.820 Sperm Head
-	A05.360.490.890.820.100 Acrosome
-	A05.360.490.890.830 Sperm Midpiece



## MeSH Tree Changes for 2017

Type	Tree - heading
-	A05.360.490.890.840 Sperm Tail
-	A05.360.490.890.860 Spermatids
-	A05.360.490.890.880 Spermatocytes
-	A05.360.490.890.900 Spermatogonia
-	A05.360.576 Gonads
-	A05.360.576.497 Ovary
-	A05.360.576.782 Testis
-	A05.810 Urinary Tract
-	A05.810.453 Kidney
-	A05.810.453.324 Kidney Cortex
-	A05.810.453.324.359 Kidney Glomerulus
-	A05.810.453.324.359.372 Glomerular Filtration Barrier
-	A05.810.453.324.359.372.400 Glomerular Basement Membrane
-	A05.810.453.324.359.372.650 Podocytes
-	A05.810.453.324.359.520 Juxtaglomerular Apparatus
-	A05.810.453.324.359.620 Mesangial Cells
-	A05.810.453.324.359.620.500 Glomerular Mesangium
-	A05.810.453.466 Kidney Medulla
-	A05.810.453.537 Kidney Pelvis
-	A05.810.453.537.503 Kidney Calices
-	A05.810.453.736 Nephrons
-	A05.810.453.736.520 Kidney Glomerulus
-	A05.810.453.736.520.224 Glomerular Basement Membrane
-	A05.810.453.736.520.372 Glomerular Filtration Barrier
-	A05.810.453.736.520.520 Juxtaglomerular Apparatus
-	A05.810.453.736.520.620 Mesangial Cells
-	A05.810.453.736.520.620.500 Glomerular Mesangium
-	A05.810.453.736.520.720 Podocytes
-	A05.810.453.736.560 Kidney Tubules
-	A05.810.453.736.560.254 Bowman Capsule
-	A05.810.453.736.560.510 Kidney Tubules, Collecting
-	A05.810.453.736.560.540 Kidney Tubules, Distal
-	A05.810.453.736.560.570 Kidney Tubules, Proximal
-	A05.810.453.736.560.610 Loop of Henle
-	A05.810.776 Ureter
-	A05.810.876 Urethra

## MeSH Tree Changes for 2017

Type	Tree - heading
-	A05.810.890                      Urinary Bladder
-	A06                                      Endocrine System
-	A06.224                              Chromaffin System
-	A06.224.161                      Chromaffin Cells
-	A06.224.161.500                      Chromaffin Granules
-	A06.224.358                      Enterochromaffin Cells
-	A06.224.365                      Enterochromaffin-like Cells
-	A06.224.636                      Para-Aortic Bodies
-	A06.224.736                      Paraganglia, Chromaffin
-	A06.390                              Enteroendocrine Cells
-	A06.390.021                      Enterochromaffin Cells
-	A06.390.043                      Enterochromaffin-like Cells
-	A06.390.065                      Gastrin-Secreting Cells
-	A06.390.087                      Glucagon-Secreting Cells
-	A06.390.131                      Insulin-Secreting Cells
-	A06.390.650                      Pancreatic Polypeptide-Secreting Cells
-	A06.390.825                      Somatostatin-Secreting Cells
-	A06.407                              Endocrine Glands
-	A06.407.071                      Adrenal Glands
-	A06.407.071.140                      Adrenal Cortex
-	A06.407.071.140.950                      Zona Fasciculata
-	A06.407.071.140.960                      Zona Glomerulosa
-	A06.407.071.140.970                      Zona Reticularis
-	A06.407.071.265                      Adrenal Medulla
-	A06.407.312                      Gonads
-	A06.407.312.497                      Ovary
-	A06.407.312.497.278                      Corpus Luteum
-	A06.407.312.497.278.400                      Luteal Cells
-	A06.407.312.497.535                      Ovarian Follicle
-	A06.407.312.497.535.150                      Follicular Fluid
-	A06.407.312.497.535.300                      Granulosa Cells
-	A06.407.312.497.535.300.500                      Cumulus Cells
-	A06.407.312.497.535.600                      Theca Cells
-	A06.407.312.782                      Testis
-	A06.407.312.782.513                      Leydig Cells
-	A06.407.414                      Islets of Langerhans

## MeSH Tree Changes for 2017

Type	Tree - heading
-	A06.407.414.043 Glucagon-Secreting Cells
-	A06.407.414.087 Insulin-Secreting Cells
-	A06.407.414.587 Pancreatic Polypeptide-Secreting Cells
-	A06.407.414.793 Somatostatin-Secreting Cells
-	A06.407.560 Parathyroid Glands
-	A06.407.635 Pineal Gland
-	A06.407.691 Pituitary-Adrenal System
-	A06.407.747 Pituitary Gland
-	A06.407.747.500 Pituitary Gland, Anterior
-	A06.407.747.500.500 Corticotrophs
-	A06.407.747.500.750 Gonadotrophs
-	A06.407.747.500.812 Lactotrophs
-	A06.407.747.500.937 Somatotrophs
-	A06.407.747.500.968 Thyrotrophs
-	A06.407.747.750 Pituitary Gland, Intermediate
-	A06.407.747.750.500 Melanotrophs
-	A06.407.747.875 Pituitary Gland, Posterior
-	A06.407.900 Thyroid Gland
-	A06.688 Neurosecretory Systems
-	A06.688.178 Circumventricular Organs
-	A06.688.178.500 Area Postrema
-	A06.688.178.750 Median Eminence
-	A06.688.178.875 Pituitary Gland, Posterior
-	A06.688.178.937 Subcommissural Organ
-	A06.688.178.968 Subfornical Organ
-	A06.688.267 Glomus Tympanicum
-	A06.688.357 Hypothalamo-Hypophyseal System
-	A06.688.357.500 Median Eminence
-	A06.688.357.750 Pituitary Gland
-	A06.688.357.750.500 Pituitary Gland, Anterior
-	A06.688.357.750.500.500 Corticotrophs
-	A06.688.357.750.500.750 Gonadotrophs
-	A06.688.357.750.500.812 Lactotrophs
-	A06.688.357.750.500.937 Somatotrophs
-	A06.688.357.750.500.968 Thyrotrophs
-	A06.688.357.750.750 Pituitary Gland, Intermediate



## MeSH Tree Changes for 2017

Type	Tree - heading
-	A07.231.114.248 Ciliary Arteries
-	A07.231.114.269 Coronary Vessels
-	A07.231.114.330 Epigastric Arteries
-	A07.231.114.351 Femoral Artery
-	A07.231.114.379 Gastroepiploic Artery
-	A07.231.114.407 Hepatic Artery
-	A07.231.114.444 Iliac Artery
-	A07.231.114.523 Maxillary Artery
-	A07.231.114.545 Meningeal Arteries
-	A07.231.114.565 Mesenteric Arteries
-	A07.231.114.565.510 Mesenteric Artery, Inferior
-	A07.231.114.565.755 Mesenteric Artery, Superior
-	A07.231.114.622 Ophthalmic Artery
-	A07.231.114.681 Popliteal Artery
-	A07.231.114.715 Pulmonary Artery
-	A07.231.114.740 Radial Artery
-	A07.231.114.745 Renal Artery
-	A07.231.114.765 Retinal Artery
-	A07.231.114.814 Splenic Artery
-	A07.231.114.839 Subclavian Artery
-	A07.231.114.891 Thoracic Arteries
-	A07.231.114.891.525 Mammary Arteries
-	A07.231.114.895 Tibial Arteries
-	A07.231.114.920 Ulnar Artery
-	A07.231.114.929 Umbilical Arteries
-	A07.231.114.940 Uterine Artery
-	A07.231.114.955 Vertebral Artery
-	A07.231.461 Microvessels
-	A07.231.461.080 Arterioles
-	A07.231.461.085 Arteriovenous Anastomosis
-	A07.231.461.165 Capillaries
-	A07.231.461.920 Venules
-	A07.231.611 Retinal Vessels
-	A07.231.611.647 Retinal Artery
-	A07.231.611.773 Retinal Vein
-	A07.231.700 Tunica Intima

## MeSH Tree Changes for 2017

Type	Tree - heading
-	A07.231.700.500 Endothelium, Vascular
-	A07.231.700.750 Pericytes
-	A07.231.733 Tunica Media
-	A07.231.733.500 Muscle, Smooth, Vascular
-	A07.231.765 Vasa Nervorum
-	A07.231.836 Vasa Vasorum
-	A07.231.908 Veins
-	A07.231.908.077 Axillary Vein
-	A07.231.908.106 Azygos Vein
-	A07.231.908.130 Brachiocephalic Veins
-	A07.231.908.155 Cerebral Veins
-	A07.231.908.194 Coronary Vessels
-	A07.231.908.194.500 Coronary Sinus
-	A07.231.908.224 Cranial Sinuses
-	A07.231.908.224.334 Cavernous Sinus
-	A07.231.908.224.667 Superior Sagittal Sinus
-	A07.231.908.224.833 Transverse Sinuses
-	A07.231.908.314 Femoral Vein
-	A07.231.908.380 Hepatic Veins
-	A07.231.908.427 Iliac Vein
-	A07.231.908.498 Jugular Veins
-	A07.231.908.641 Popliteal Vein
-	A07.231.908.670 Portal System
-	A07.231.908.670.385 Mesenteric Veins
-	A07.231.908.670.567 Portal Vein
-	A07.231.908.670.730 Splenic Vein
-	A07.231.908.670.874 Umbilical Veins
-	A07.231.908.713 Pulmonary Veins
-	A07.231.908.752 Renal Veins
-	A07.231.908.783 Retinal Vein
-	A07.231.908.819 Saphenous Vein
-	A07.231.908.877 Subclavian Vein
-	A07.231.908.949 Venae Cavae
-	A07.231.908.949.648 Vena Cava, Inferior
-	A07.231.908.949.815 Vena Cava, Superior
-	A07.231.908.950 Venous Valves

## MeSH Tree Changes for 2017

Type	Tree - heading
-	A07.231.908.952 Venules
-	A07.500 Glomerular Filtration Barrier
-	A07.541 Heart
-	A07.541.207 Endocardium
-	A07.541.278 Fetal Heart
-	A07.541.278.395 Ductus Arteriosus
-	A07.541.278.930 Truncus Arteriosus
-	A07.541.358 Heart Atria
-	A07.541.358.100 Atrial Appendage
-	A07.541.409 Heart Conduction System
-	A07.541.409.147 Atrioventricular Node
-	A07.541.409.273 Bundle of His
-	A07.541.409.683 Purkinje Fibers
-	A07.541.409.819 Sinoatrial Node
-	A07.541.459 Heart Septum
-	A07.541.459.249 Atrial Septum
-	A07.541.459.374 Endocardial Cushions
-	A07.541.459.500 Foramen Ovale
-	A07.541.459.750 Ventricular Septum
-	A07.541.510 Heart Valves
-	A07.541.510.110 Aortic Valve
-	A07.541.510.240 Chordae Tendineae
-	A07.541.510.507 Mitral Valve
-	A07.541.510.619 Papillary Muscles
-	A07.541.510.738 Pulmonary Valve
-	A07.541.510.893 Tricuspid Valve
-	A07.541.560 Heart Ventricles
-	A07.541.704 Myocardium
-	A07.541.704.500 Myoblasts, Cardiac
-	A07.541.704.570 Myocytes, Cardiac
-	A07.541.704.750 Papillary Muscles
-	A07.541.795 Pericardium
-	A07.541.795.500 Pericardial Fluid
-	A08 Nervous System
-	A08.186 Central Nervous System
-	A08.186.211 Brain

## MeSH Tree Changes for 2017

Type	Tree - heading
-	A08.186.211.035 Blood-Brain Barrier
-	A08.186.211.132 Brain Stem
-	A08.186.211.132.659 Mesencephalon
-	A08.186.211.132.659.413 Cerebral Peduncle
-	A08.186.211.132.659.413.437 Cerebral Crus
-	A08.186.211.132.659.413.656 Substantia Nigra
-	A08.186.211.132.659.413.656.249 Pars Compacta
-	A08.186.211.132.659.413.656.500 Pars Reticulata
-	A08.186.211.132.659.413.875 Tegmentum Mesencephali
-	A08.186.211.132.659.413.875.187 Cerebral Aqueduct
-	A08.186.211.132.659.413.875.284 Midbrain Reticular Formation
-	A08.186.211.132.659.413.875.284.500 Pedunculopontine Tegmental Nucleus
-	A08.186.211.132.659.413.875.381 Oculomotor Nuclear Complex
-	A08.186.211.132.659.413.875.381.500 Edinger-Westphal Nucleus
-	A08.186.211.132.659.413.875.595 Periaqueductal Gray
-	A08.186.211.132.659.413.875.618 Raphe Nuclei
-	A08.186.211.132.659.413.875.618.249 Dorsal Raphe Nucleus
-	A08.186.211.132.659.413.875.618.436 Interpeduncular Nucleus
-	A08.186.211.132.659.413.875.618.624 Midbrain Raphe Nuclei
-	A08.186.211.132.659.413.875.642 Red Nucleus
-	A08.186.211.132.659.413.875.820 Ventral Tegmental Area
-	A08.186.211.132.659.473 Locus Coeruleus
-	A08.186.211.132.659.800 Tectum Mesencephali
-	A08.186.211.132.659.800.407 Inferior Colliculi
-	A08.186.211.132.659.800.611 Posterior Cerebellar Commissure
-	A08.186.211.132.659.800.662 Pretectal Region
-	A08.186.211.132.659.800.713 Subcommissural Organ
-	A08.186.211.132.659.800.816 Superior Colliculi
-	A08.186.211.132.772 Reticular Formation
-	A08.186.211.132.772.322 Midbrain Reticular Formation
-	A08.186.211.132.772.322.500 Pedunculopontine Tegmental Nucleus
-	A08.186.211.132.772.646 Respiratory Center
-	A08.186.211.132.810 Rhombencephalon
-	A08.186.211.132.810.428 Metencephalon
-	A08.186.211.132.810.428.200 Cerebellum
-	A08.186.211.132.810.428.200.212 Cerebellar Cortex



## MeSH Tree Changes for 2017

Type	Tree - heading
-	A08.186.211.132.810.428.200.212.299 Cerebellar Vermis
-	A08.186.211.132.810.428.200.212.600 Purkinje Cells
-	A08.186.211.132.810.428.200.337 Cerebellar Nuclei
-	A08.186.211.132.810.428.200.462 Cerebellopontine Angle
-	A08.186.211.132.810.428.600 Pons
-	A08.186.211.132.810.428.600.067 Barrington's Nucleus
-	A08.186.211.132.810.428.600.135 Cochlear Nucleus
-	A08.186.211.132.810.428.600.392 Kolliker-Fuse Nucleus
-	A08.186.211.132.810.428.600.521 Middle Cerebellar Peduncle
-	A08.186.211.132.810.428.600.650 Pontine Tegmentum
-	A08.186.211.132.810.428.600.650.249 Abducens Nucleus
-	A08.186.211.132.810.428.600.650.374 Facial Nucleus
-	A08.186.211.132.810.428.600.650.437 Locus Coeruleus
-	A08.186.211.132.810.428.600.650.500 Parabrachial Nucleus
-	A08.186.211.132.810.428.600.650.562 Raphe Nuclei
-	A08.186.211.132.810.428.600.650.562.500 Nucleus Raphe Magnus
-	A08.186.211.132.810.428.600.650.625 Superior Olivary Complex
-	A08.186.211.132.810.428.600.650.687 Trapezoid Body
-	A08.186.211.132.810.428.600.650.718 Trigeminal Motor Nucleus
-	A08.186.211.132.810.428.600.650.750 Trigeminal Nucleus, Spinal
-	A08.186.211.132.810.428.600.800 Vestibular Nuclei
-	A08.186.211.132.810.428.600.800.800 Vestibular Nucleus, Lateral
-	A08.186.211.132.810.591 Myelencephalon
-	A08.186.211.132.810.591.500 Medulla Oblongata
-	A08.186.211.132.810.591.500.286 Area Postrema
-	A08.186.211.132.810.591.500.574 Olivary Nucleus
-	A08.186.211.132.810.591.500.662 Raphe Nuclei
-	A08.186.211.132.810.591.500.662.436 Nucleus Raphe Obscurus
-	A08.186.211.132.810.591.500.662.624 Nucleus Raphe Pallidus
-	A08.186.211.132.810.591.500.750 Solitary Nucleus
-	A08.186.211.132.810.591.500.875 Trigeminal Nucleus, Spinal
-	A08.186.211.132.810.591.500.875.500 Trigeminal Caudal Nucleus
-	A08.186.211.132.810.753 Tectospinal Fibers
-	A08.186.211.132.931 Trigeminal Nuclei
-	A08.186.211.132.931.459 Trigeminal Motor Nucleus
-	A08.186.211.132.931.920 Trigeminal Nucleus, Spinal

## MeSH Tree Changes for 2017

Type	Tree - heading
-	A08.186.211.132.931.920.927 Trigeminal Caudal Nucleus
-	A08.186.211.168 Gray Matter
-	A08.186.211.204 White Matter
-	A08.186.211.276 Cerebral Ventricles
-	A08.186.211.276.187 Cerebral Aqueduct
-	A08.186.211.276.298 Choroid Plexus
-	A08.186.211.276.460 Ependyma
-	A08.186.211.276.500 Fourth Ventricle
-	A08.186.211.276.650 Lateral Ventricles
-	A08.186.211.276.814 Septum Pellucidum
-	A08.186.211.276.840 Third Ventricle
-	A08.186.211.464 Limbic System
-	A08.186.211.464.090 Amygdala
-	A08.186.211.464.090.500 Basolateral Nuclear Complex
-	A08.186.211.464.090.750 Central Amygdaloid Nucleus
-	A08.186.211.464.090.875 Corticomедial Nuclear Complex
-	A08.186.211.464.090.875.500 Periamygdaloid Cortex
-	A08.186.211.464.200 Epithalamus
-	A08.186.211.464.200.360 Habenula
-	A08.186.211.464.200.680 Pineal Gland
-	A08.186.211.464.405 Hippocampus
-	A08.186.211.464.405.099 CA1 Region, Hippocampal
-	A08.186.211.464.405.149 CA2 Region, Hippocampal
-	A08.186.211.464.405.174 CA3 Region, Hippocampal
-	A08.186.211.464.405.200 Dentate Gyrus
-	A08.186.211.464.405.200.500 Mossy Fibers, Hippocampal
-	A08.186.211.464.405.600 Fornix, Brain
-	A08.186.211.464.497 Hypothalamus
-	A08.186.211.464.497.300 Hypothalamic Area, Lateral
-	A08.186.211.464.497.342 Hypothalamus, Anterior
-	A08.186.211.464.497.342.063 Anterior Hypothalamic Nucleus
-	A08.186.211.464.497.342.231 Organum Vasculosum
-	A08.186.211.464.497.342.400 Paraventricular Hypothalamic Nucleus
-	A08.186.211.464.497.342.450 Preoptic Area
-	A08.186.211.464.497.342.625 Suprachiasmatic Nucleus
-	A08.186.211.464.497.342.650 Supraoptic Nucleus

## MeSH Tree Changes for 2017

Type	Tree - heading
-	A08.186.211.464.497.352 Hypothalamus, Middle
-	A08.186.211.464.497.352.081 Arcuate Nucleus of Hypothalamus
-	A08.186.211.464.497.352.270 Dorsomedial Hypothalamic Nucleus
-	A08.186.211.464.497.352.435 Hypothalamo-Hypophyseal System
-	A08.186.211.464.497.352.435.249 Median Eminence
-	A08.186.211.464.497.352.435.500 Pituitary Gland
-	A08.186.211.464.497.352.435.500.500 Pituitary Gland, Anterior
-	A08.186.211.464.497.352.435.500.500.500 Corticotrophs
-	A08.186.211.464.497.352.435.500.500.750 Gonadotrophs
-	A08.186.211.464.497.352.435.500.500.812 Lactotrophs
-	A08.186.211.464.497.352.435.500.500.937 Somatotrophs
-	A08.186.211.464.497.352.435.500.500.968 Thyrotrophs
-	A08.186.211.464.497.352.435.500.750 Pituitary Gland, Intermediate
-	A08.186.211.464.497.352.435.500.750.500 Melanotrophs
-	A08.186.211.464.497.352.435.500.875 Pituitary Gland, Posterior
-	A08.186.211.464.497.352.870 Tuber Cinereum
-	A08.186.211.464.497.352.953 Ventromedial Hypothalamic Nucleus
-	A08.186.211.464.497.362 Hypothalamus, Posterior
-	A08.186.211.464.497.362.500 Mammillary Bodies
-	A08.186.211.464.590 Limbic Lobe
-	A08.186.211.464.590.500 Gyrus Cinguli
-	A08.186.211.464.590.750 Parahippocampal Gyrus
-	A08.186.211.464.590.750.225 Entorhinal Cortex
-	A08.186.211.464.590.750.612 Periamygdaloid Cortex
-	A08.186.211.464.699 Olfactory Pathways
-	A08.186.211.464.710 Parahippocampal Gyrus
-	A08.186.211.464.710.225 Entorhinal Cortex
-	A08.186.211.464.710.612 Periamygdaloid Cortex
-	A08.186.211.464.730 Perforant Pathway
-	A08.186.211.464.750 Septum of Brain
-	A08.186.211.464.750.800 Septal Nuclei
-	A08.186.211.464.750.900 Septum Pellucidum
-	A08.186.211.464.820 Substantia Innominata
-	A08.186.211.730 Prosencephalon
-	A08.186.211.730.317 Diencephalon
-	A08.186.211.730.317.200 Epithalamus

## MeSH Tree Changes for 2017

Type	Tree - heading
-	A08.186.211.730.317.200.360                      Habenula
-	A08.186.211.730.317.200.620                      Pineal Gland
-	A08.186.211.730.317.357                              Hypothalamus
-	A08.186.211.730.317.357.300                      Hypothalamic Area, Lateral
-	A08.186.211.730.317.357.342                      Hypothalamus, Anterior
-	A08.186.211.730.317.357.342.063                      Anterior Hypothalamic Nucleus
-	A08.186.211.730.317.357.342.315                      Organum Vasculosum
-	A08.186.211.730.317.357.342.400                      Paraventricular Hypothalamic Nucleus
-	A08.186.211.730.317.357.342.450                      Preoptic Area
-	A08.186.211.730.317.357.342.625                      Suprachiasmatic Nucleus
-	A08.186.211.730.317.357.342.650                      Supraoptic Nucleus
-	A08.186.211.730.317.357.352                              Hypothalamus, Middle
-	A08.186.211.730.317.357.352.081                      Arcuate Nucleus of Hypothalamus
-	A08.186.211.730.317.357.352.270                      Dorsomedial Hypothalamic Nucleus
-	A08.186.211.730.317.357.352.435                      Hypothalamo-Hypophyseal System
-	A08.186.211.730.317.357.352.435.249                      Median Eminence
-	A08.186.211.730.317.357.352.435.500                      Pituitary Gland
-	A08.186.211.730.317.357.352.435.500.500                      Pituitary Gland, Anterior
-	A08.186.211.730.317.357.352.435.500.500.500                      Corticotrophs
-	A08.186.211.730.317.357.352.435.500.500.750                      Gonadotrophs
-	A08.186.211.730.317.357.352.435.500.500.812                      Lactotrophs
-	A08.186.211.730.317.357.352.435.500.500.937                      Somatotrophs
-	A08.186.211.730.317.357.352.435.500.500.968                      Thyrotrophs
-	A08.186.211.730.317.357.352.435.500.750                      Pituitary Gland, Intermediate
-	A08.186.211.730.317.357.352.435.500.750.500                      Melanotrophs
-	A08.186.211.730.317.357.352.435.500.875                      Pituitary Gland, Posterior
-	A08.186.211.730.317.357.352.870                      Tuber Cinereum
-	A08.186.211.730.317.357.352.953                      Ventromedial Hypothalamic Nucleus
-	A08.186.211.730.317.357.362                              Hypothalamus, Posterior
-	A08.186.211.730.317.357.362.500                      Mammillary Bodies
-	A08.186.211.730.317.578                              Optic Chiasm
-	A08.186.211.730.317.689                              Optic Tract
-	A08.186.211.730.317.800                              Subthalamus
-	A08.186.211.730.317.800.240                      Entopeduncular Nucleus
-	A08.186.211.730.317.800.800                      Subthalamic Nucleus
-	A08.186.211.730.317.800.900                      Zona Incerta



## MeSH Tree Changes for 2017

Type	Tree - heading
-	A08.186.211.730.885.287.500.345 Hippocampus
-	A08.186.211.730.885.287.500.345.099 CA1 Region, Hippocampal
-	A08.186.211.730.885.287.500.345.149 CA2 Region, Hippocampal
-	A08.186.211.730.885.287.500.345.174 CA3 Region, Hippocampal
-	A08.186.211.730.885.287.500.345.200 Dentate Gyrus
-	A08.186.211.730.885.287.500.345.200.500 Mossy Fibers, Hippocampal
-	A08.186.211.730.885.287.500.345.600 Fornix, Brain
-	A08.186.211.730.885.287.500.382 Limbic Lobe
-	A08.186.211.730.885.287.500.382.500 Gyrus Cinguli
-	A08.186.211.730.885.287.500.382.750 Parahippocampal Gyrus
-	A08.186.211.730.885.287.500.382.750.225 Entorhinal Cortex
-	A08.186.211.730.885.287.500.382.750.612 Periamygdaloid Cortex
-	A08.186.211.730.885.287.500.420 Neocortex
-	A08.186.211.730.885.287.500.571 Occipital Lobe
-	A08.186.211.730.885.287.500.571.735 Visual Cortex
-	A08.186.211.730.885.287.500.620 Olfactory Cortex
-	A08.186.211.730.885.287.500.620.374 Basal Forebrain
-	A08.186.211.730.885.287.500.620.374.500 Organum Vasculosum
-	A08.186.211.730.885.287.500.620.562 Entorhinal Cortex
-	A08.186.211.730.885.287.500.620.656 Olfactory Tubercle
-	A08.186.211.730.885.287.500.620.656.500 Islands of Calleja
-	A08.186.211.730.885.287.500.620.750 Piriform Cortex
-	A08.186.211.730.885.287.500.670 Parietal Lobe
-	A08.186.211.730.885.287.500.670.675 Somatosensory Cortex
-	A08.186.211.730.885.287.500.670.837 Wernicke Area
-	A08.186.211.730.885.287.500.814 Sensorimotor Cortex
-	A08.186.211.730.885.287.500.814.249 Auditory Cortex
-	A08.186.211.730.885.287.500.814.624 Motor Cortex
-	A08.186.211.730.885.287.500.814.695 Olfactory Cortex
-	A08.186.211.730.885.287.500.814.695.249 Basal Forebrain
-	A08.186.211.730.885.287.500.814.695.374 Entorhinal Cortex
-	A08.186.211.730.885.287.500.814.695.500 Olfactory Tubercle
-	A08.186.211.730.885.287.500.814.695.500.500 Islands of Calleja
-	A08.186.211.730.885.287.500.814.906 Somatosensory Cortex
-	A08.186.211.730.885.287.500.814.953 Visual Cortex
-	A08.186.211.730.885.287.500.863 Temporal Lobe

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	A08.186.211.730.885.287.500.863.297	Auditory Cortex
New Heading	<b>A08.186.211.730.885.287.500.863.574</b>	<b>Perirhinal Cortex</b>
-	A08.186.211.730.885.287.500.863.850	Wernicke Area
-	A08.186.211.730.885.287.750	Olfactory Cortex
-	A08.186.211.730.885.287.750.374	Basal Forebrain
-	A08.186.211.730.885.287.750.562	Entorhinal Cortex
-	A08.186.211.730.885.287.750.656	Olfactory Tubercle
-	A08.186.211.730.885.287.750.656.500	Islands of Calleja
-	A08.186.211.730.885.287.750.750	Piriform Cortex
-	A08.186.211.730.885.380	Diagonal Band of Broca
-	A08.186.211.730.885.385	External Capsule
-	A08.186.211.730.885.388	Olfactory Bulb
-	A08.186.211.730.885.390	Telencephalic Commissures
-	A08.186.211.730.885.390.500	Anterior Cerebellar Commissure
-	A08.186.211.730.885.390.750	Corpus Callosum
-	A08.186.211.730.885.435	Internal Capsule
-	A08.186.211.730.885.750	Septum of Brain
-	A08.186.211.730.885.750.800	Septal Nuclei
-	A08.186.211.730.885.750.814	Septum Pellucidum
-	A08.186.566	Meninges
-	A08.186.566.166	Arachnoid
-	A08.186.566.166.686	Subarachnoid Space
-	A08.186.566.166.686.351	Cisterna Magna
-	A08.186.566.395	Dura Mater
-	A08.186.566.395.687	Subdural Space
-	A08.186.566.731	Pia Mater
-	A08.186.854	Spinal Cord
-	A08.186.854.126	Cervical Cord
-	A08.186.854.253	Extrapyramidal Tracts
-	A08.186.854.348	Gray Matter
-	A08.186.854.443	Spinal Cord Lateral Horn
-	A08.186.854.633	Pyramidal Tracts
-	A08.186.854.697	Spinal Cord Dorsal Horn
-	A08.186.854.697.500	Posterior Horn Cells
-	A08.186.854.697.500.500	Substantia Gelatinosa

## MeSH Tree Changes for 2017

Type	Tree - heading
-	A08.186.854.729 Spinal Cord Ventral Horn
-	A08.186.854.729.500 Anterior Horn Cells
-	A08.186.854.761 Spinothalamic Tracts
-	A08.186.854.880 White Matter
-	A08.340 Ganglia
-	A08.340.315 Ganglia, Autonomic
-	A08.340.315.300 Ganglia, Parasympathetic
-	A08.340.315.350 Ganglia, Sympathetic
-	A08.340.315.350.800 Stellate Ganglion
-	A08.340.315.350.850 Superior Cervical Ganglion
-	A08.340.352 Ganglia, Invertebrate
-	A08.340.390 Ganglia, Sensory
-	A08.340.390.340 Ganglia, Spinal
-	A08.340.390.380 Genuiculate Ganglion
-	A08.340.390.550 Nodose Ganglion
-	A08.340.390.800 Spiral Ganglion
-	A08.340.390.850 Trigeminal Ganglion
-	A08.340.685 Satellite Cells, Perineuronal
-	A08.511 Nerve Net
-	A08.511.500 Central Pattern Generators
-	A08.612 Neural Pathways
-	A08.612.220 Afferent Pathways
-	A08.612.220.110 Auditory Pathways
-	A08.612.220.640 Olfactory Pathways
-	A08.612.220.725 Spinocerebellar Tracts
-	A08.612.220.735 Spinothalamic Tracts
-	A08.612.220.830 Visceral Afferents
-	A08.612.220.860 Visual Pathways
-	A08.612.380 Efferent Pathways
-	A08.612.380.239 Extrapyramidal Tracts
-	A08.612.380.730 Pyramidal Tracts
-	A08.612.435 Internal Capsule
-	A08.612.492 Medial Forebrain Bundle
-	A08.612.600 Perforant Pathway
-	A08.637 Neuroglia
-	A08.637.200 Astrocytes



## MeSH Tree Changes for 2017

Type	Tree - heading	
-	A08.637.300	Ependymoglia Cells
-	A08.637.400	Microglia
-	A08.637.500	Neuropil
-	A08.637.500.550	Neuropil Threads
-	A08.637.600	Oligodendroglia
-	A08.637.600.500	Myelin Sheath
-	A08.637.685	Satellite Cells, Perineuronal
-	A08.637.800	Schwann Cells
-	A08.637.800.500	Myelin Sheath
-	A08.637.800.500.550	Neurilemma
-	A08.637.800.500.700	Ranvier's Nodes
-	A08.675	Neurons
-	A08.675.100	Adrenergic Neurons
-	A08.675.100.500	Adrenergic Fibers
-	A08.675.127	Cholinergic Neurons
New Tree	<a href="#">A08.675.127.500</a>	<a href="#">Cholinergic Fibers</a>
New Tree	<a href="#">A08.675.127.500.060</a>	<a href="#">Autonomic Fibers, Preganglionic</a>
New Tree	<a href="#">A08.675.127.500.700</a>	<a href="#">Parasympathetic Fibers, Postganglionic</a>
-	A08.675.256	Dendrites
-	A08.675.256.200	Dendritic Spines
-	A08.675.256.500	Neurites
-	A08.675.278	Dopaminergic Neurons
-	A08.675.289	GABAergic Neurons
New Heading	<b>A08.675.324</b>	<b>Grid Cells</b>
-	A08.675.358	Interneurons
-	A08.675.358.050	Amacrine Cells
-	A08.675.358.350	Commissural Interneurons
-	A08.675.358.650	Renshaw Cells
-	A08.675.358.700	Retinal Bipolar Cells
-	A08.675.460	Lewy Bodies
-	A08.675.500	Mirror Neurons
-	A08.675.542	Nerve Fibers
-	A08.675.542.075	Adrenergic Fibers

## MeSH Tree Changes for 2017

Type	Tree - heading
-	A08.675.542.075.800                      Sympathetic Fibers, Postganglionic
-	A08.675.542.100                              Autonomic Fibers, Postganglionic
-	A08.675.542.100.700                      Parasympathetic Fibers, Postganglionic
-	A08.675.542.100.800                      Sympathetic Fibers, Postganglionic
-	A08.675.542.145                              Axons
New Heading	<b>A08.675.542.145.250                      Axon Initial Segment</b>
-	A08.675.542.145.500                      Neurites
-	A08.675.542.145.750                      Presynaptic Terminals
-	A08.675.542.145.750.500                      Mossy Fibers, Hippocampal
-	A08.675.542.234                              Cholinergic Fibers
-	A08.675.542.234.060                      Autonomic Fibers, Preganglionic
-	A08.675.542.234.700                      Parasympathetic Fibers, Postganglionic
-	A08.675.542.512                              Nerve Fibers, Myelinated
-	A08.675.542.512.560                      Myelin Sheath
-	A08.675.542.512.560.550                      Neurilemma
-	A08.675.542.512.560.700                      Ranvier's Nodes
-	A08.675.542.756                              Nerve Fibers, Unmyelinated
-	A08.675.575                                      Neural Analyzers
-	A08.675.609                                      Neurofibrils
-	A08.675.609.520                              Neurofibrillary Tangles
-	A08.675.650                                      Neurons, Afferent
-	A08.675.650.250                              Hair Cells, Auditory
-	A08.675.650.250.250                      Hair Cells, Auditory, Inner
-	A08.675.650.250.315                      Hair Cells, Auditory, Outer
-	A08.675.650.395                              Hair Cells, Vestibular
-	A08.675.650.675                              Posterior Horn Cells
-	A08.675.650.675.800                      Substantia Gelatinosa
-	A08.675.650.850                              Retinal Neurons
-	A08.675.650.850.500                      Amacrine Cells
-	A08.675.650.850.625                      Photoreceptor Cells
-	A08.675.650.850.625.660                      Photoreceptor Cells, Invertebrate
-	A08.675.650.850.625.670                      Photoreceptor Cells, Vertebrate
-	A08.675.650.850.625.670.049                      Photoreceptor Connecting Cilium
-	A08.675.650.850.625.670.100                      Retinal Cone Photoreceptor Cells
-	A08.675.650.850.625.670.237                      Retinal Photoreceptor Cell Inner Segment

## MeSH Tree Changes for 2017

Type	Tree - heading
-	A08.675.650.850.625.670.375 Retinal Photoreceptor Cell Outer Segment
-	A08.675.650.850.625.670.375.500 Rod Cell Outer Segment
-	A08.675.650.850.625.670.650 Retinal Rod Photoreceptor Cells
-	A08.675.650.850.625.670.650.650 Rod Cell Outer Segment
-	A08.675.650.850.750 Retinal Bipolar Cells
-	A08.675.650.850.875 Retinal Ganglion Cells
-	A08.675.650.850.937 Retinal Horizontal Cells
-	A08.675.650.915 Sensory Receptor Cells
-	A08.675.650.915.500 Chemoreceptor Cells
-	A08.675.650.915.500.530 Neuroepithelial Cells
-	A08.675.650.915.500.530.550 Neuroepithelial Bodies
-	A08.675.650.915.500.540 Olfactory Receptor Neurons
-	A08.675.650.915.500.600 Paraganglia, Nonchromaffin
-	A08.675.650.915.500.600.050 Aortic Bodies
-	A08.675.650.915.500.600.150 Carotid Body
-	A08.675.650.915.500.600.350 Glomus Jugulare
-	A08.675.650.915.500.600.360 Glomus Tympanicum
-	A08.675.650.915.500.800 Taste Buds
-	A08.675.650.915.750 Mechanoreceptors
-	A08.675.650.915.750.300 Golgi-Mazzoni Corpuscles
-	A08.675.650.915.750.425 Merkel Cells
-	A08.675.650.915.750.500 Muscle Spindles
-	A08.675.650.915.750.600 Neuroepithelial Cells
-	A08.675.650.915.750.600.350 Hair Cells, Auditory
-	A08.675.650.915.750.600.350.350 Hair Cells, Auditory, Inner
-	A08.675.650.915.750.600.350.365 Hair Cells, Auditory, Outer
-	A08.675.650.915.750.600.675 Hair Cells, Vestibular
-	A08.675.650.915.750.700 Pacinian Corpuscles
-	A08.675.650.915.750.750 Pressoreceptors
-	A08.675.650.915.750.780 Pulmonary Stretch Receptors
-	A08.675.650.915.875 Nociceptors
-	A08.675.650.915.937 Photoreceptor Cells
-	A08.675.650.915.937.650 Photoreceptor Cells, Invertebrate
-	A08.675.650.915.937.670 Photoreceptor Cells, Vertebrate
-	A08.675.650.915.937.670.049 Photoreceptor Connecting Cilium
-	A08.675.650.915.937.670.100 Retinal Cone Photoreceptor Cells

## MeSH Tree Changes for 2017

Type	Tree - heading
-	A08.675.650.915.937.670.237      Retinal Photoreceptor Cell Inner Segment
-	A08.675.650.915.937.670.375      Retinal Photoreceptor Cell Outer Segment
-	A08.675.650.915.937.670.375.500      Rod Cell Outer Segment
-	A08.675.650.915.937.670.650      Retinal Rod Photoreceptor Cells
-	A08.675.650.915.937.670.650.650      Rod Cell Outer Segment
-	A08.675.650.915.968      Thermoreceptors
-	A08.675.655      Neurons, Efferent
-	A08.675.655.500      Motor Neurons
-	A08.675.655.500.050      Anterior Horn Cells
-	A08.675.655.500.525      Motor Neurons, Gamma
-	A08.675.703      Neuropil
-	A08.675.703.550      Neuropil Threads
-	A08.675.712      Nissl Bodies
-	A08.675.748      Nitregic Neurons
-	A08.675.784      Purkinje Cells
-	A08.675.790      Pyramidal Cells
New Heading	<b>A08.675.790.500      Place Cells</b>
-	A08.675.895      Serotonergic Neurons
-	A08.713      Neurosecretory Systems
-	A08.713.049      Circumventricular Organs
-	A08.713.049.500      Area Postrema
-	A08.713.049.750      Median Eminence
-	A08.713.049.875      Pituitary Gland, Posterior
-	A08.713.049.937      Subcommissural Organ
-	A08.713.049.968      Subfornical Organ
-	A08.713.100      Corpora Allata
-	A08.713.228      Glomus Tympanicum
-	A08.713.357      Hypothalamo-Hypophyseal System
-	A08.713.357.500      Median Eminence
-	A08.713.357.750      Pituitary Gland
-	A08.713.357.750.500      Pituitary Gland, Anterior
-	A08.713.357.750.500.500      Corticotrophs
-	A08.713.357.750.500.750      Gonadotrophs
-	A08.713.357.750.500.812      Lactotrophs
-	A08.713.357.750.500.937      Somatotrophs

## MeSH Tree Changes for 2017

Type	Tree - heading
-	A08.713.357.750.500.968 Thyrotrophs
-	A08.713.357.750.750 Pituitary Gland, Intermediate
-	A08.713.357.750.750.500 Melanotrophs
-	A08.713.357.750.875 Pituitary Gland, Posterior
-	A08.713.733 Pineal Gland
-	A08.800 Peripheral Nervous System
-	A08.800.050 Autonomic Nervous System
-	A08.800.050.050 Autonomic Pathways
-	A08.800.050.050.050 Autonomic Fibers, Postganglionic
-	A08.800.050.050.050.700 Parasympathetic Fibers, Postganglionic
-	A08.800.050.050.050.800 Sympathetic Fibers, Postganglionic
-	A08.800.050.050.060 Autonomic Fibers, Preganglionic
-	A08.800.050.050.150 Celiac Plexus
-	A08.800.050.050.275 Facial Nerve
-	A08.800.050.050.337 Glossopharyngeal Nerve
-	A08.800.050.050.400 Hypogastric Plexus
-	A08.800.050.050.500 Myenteric Plexus
-	A08.800.050.050.650 Oculomotor Nerve
-	A08.800.050.050.800 Splanchnic Nerves
-	A08.800.050.050.850 Submucous Plexus
-	A08.800.050.050.925 Vagus Nerve
-	A08.800.050.050.925.450 Laryngeal Nerves
-	A08.800.050.050.925.450.700 Recurrent Laryngeal Nerve
-	A08.800.050.050.925.550 Nodose Ganglion
-	A08.800.050.150 Enteric Nervous System
-	A08.800.050.150.500 Myenteric Plexus
-	A08.800.050.150.750 Submucous Plexus
-	A08.800.050.300 Ganglia, Autonomic
-	A08.800.050.300.250 Ganglia, Parasympathetic
-	A08.800.050.300.300 Ganglia, Sympathetic
-	A08.800.050.300.300.800 Stellate Ganglion
-	A08.800.050.300.300.850 Superior Cervical Ganglion
-	A08.800.050.600 Parasympathetic Nervous System
-	A08.800.050.600.149 Facial Nerve
-	A08.800.050.600.300 Ganglia, Parasympathetic
-	A08.800.050.600.387 Glossopharyngeal Nerve

## MeSH Tree Changes for 2017

Type	Tree - heading
-	A08.800.050.600.475 Oculomotor Nerve
-	A08.800.050.600.650 Parasympathetic Fibers, Postganglionic
-	A08.800.050.600.825 Vagus Nerve
-	A08.800.050.600.825.450 Laryngeal Nerves
-	A08.800.050.600.825.450.700 Recurrent Laryngeal Nerve
-	A08.800.050.600.825.550 Nodose Ganglion
-	A08.800.050.800 Sympathetic Nervous System
-	A08.800.050.800.300 Ganglia, Sympathetic
-	A08.800.050.800.300.800 Stellate Ganglion
-	A08.800.050.800.300.850 Superior Cervical Ganglion
-	A08.800.050.800.800 Splanchnic Nerves
-	A08.800.050.800.900 Vasomotor System
-	A08.800.050.800.900.700 Pressoreceptors
-	A08.800.350 Ganglia, Sensory
-	A08.800.350.340 Ganglia, Spinal
-	A08.800.350.380 Genuiculate Ganglion
-	A08.800.350.550 Nodose Ganglion
-	A08.800.350.800 Spiral Ganglion
-	A08.800.350.850 Trigeminal Ganglion
-	A08.800.550 Nerve Endings
-	A08.800.550.550 Neuroeffector Junction
-	A08.800.550.550.550 Neuromuscular Junction
-	A08.800.550.550.550.500 Motor Endplate
-	A08.800.800 Peripheral Nerves
-	A08.800.800.060 Autonomic Pathways
-	A08.800.800.060.050 Autonomic Fibers, Postganglionic
-	A08.800.800.060.050.700 Parasympathetic Fibers, Postganglionic
-	A08.800.800.060.050.800 Sympathetic Fibers, Postganglionic
-	A08.800.800.060.060 Autonomic Fibers, Preganglionic
-	A08.800.800.060.150 Celiac Plexus
-	A08.800.800.060.275 Facial Nerve
-	A08.800.800.060.337 Glossopharyngeal Nerve
-	A08.800.800.060.400 Hypogastric Plexus
-	A08.800.800.060.500 Myenteric Plexus
-	A08.800.800.060.650 Oculomotor Nerve
-	A08.800.800.060.800 Splanchnic Nerves

## MeSH Tree Changes for 2017

Type	Tree - heading
-	A08.800.800.060.840                      Submucous Plexus
-	A08.800.800.060.920                      Vagus Nerve
-	A08.800.800.060.920.450                      Laryngeal Nerves
-	A08.800.800.060.920.450.700                      Recurrent Laryngeal Nerve
-	A08.800.800.060.920.550                      Nodose Ganglion
-	A08.800.800.090                      Blood-Nerve Barrier
-	A08.800.800.120                      Cranial Nerves
-	A08.800.800.120.030                      Abducens Nerve
-	A08.800.800.120.060                      Accessory Nerve
-	A08.800.800.120.250                      Facial Nerve
-	A08.800.800.120.250.120                      Chorda Tympani Nerve
-	A08.800.800.120.250.280                      Geniculate Ganglion
-	A08.800.800.120.290                      Glossopharyngeal Nerve
-	A08.800.800.120.330                      Hypoglossal Nerve
-	A08.800.800.120.600                      Oculomotor Nerve
-	A08.800.800.120.640                      Olfactory Nerve
-	A08.800.800.120.680                      Optic Nerve
-	A08.800.800.120.680.600                      Optic Chiasm
-	A08.800.800.120.680.660                      Optic Disk
-	A08.800.800.120.760                      Trigeminal Nerve
-	A08.800.800.120.760.500                      Mandibular Nerve
-	A08.800.800.120.760.500.450                      Lingual Nerve
-	A08.800.800.120.760.550                      Maxillary Nerve
-	A08.800.800.120.760.650                      Ophthalmic Nerve
-	A08.800.800.120.760.825                      Trigeminal Ganglion
-	A08.800.800.120.800                      Trochlear Nerve
-	A08.800.800.120.900                      Vagus Nerve
-	A08.800.800.120.900.450                      Laryngeal Nerves
-	A08.800.800.120.900.450.700                      Recurrent Laryngeal Nerve
-	A08.800.800.120.900.550                      Nodose Ganglion
-	A08.800.800.120.910                      Vestibulocochlear Nerve
-	A08.800.800.120.910.120                      Cochlear Nerve
-	A08.800.800.120.910.120.800                      Spiral Ganglion
-	A08.800.800.120.910.900                      Vestibular Nerve
-	A08.800.800.690                      Schwann Cells
-	A08.800.800.690.500                      Myelin Sheath

## MeSH Tree Changes for 2017

Type	Tree - heading
-	A08.800.800.690.500.550                      Neurilemma
-	A08.800.800.690.500.700                      Ranvier's Nodes
-	A08.800.800.720                                  Spinal Nerves
-	A08.800.800.720.050                          Brachial Plexus
-	A08.800.800.720.050.500                      Median Nerve
-	A08.800.800.720.050.540                      Musculocutaneous Nerve
-	A08.800.800.720.050.700                      Radial Nerve
-	A08.800.800.720.050.850                      Ulnar Nerve
-	A08.800.800.720.150                          Cervical Plexus
-	A08.800.800.720.150.700                      Phrenic Nerve
-	A08.800.800.720.450                          Lumbosacral Plexus
-	A08.800.800.720.450.250                      Femoral Nerve
-	A08.800.800.720.450.600                      Obturator Nerve
-	A08.800.800.720.450.680                      Pudendal Nerve
-	A08.800.800.720.450.760                      Sciatic Nerve
-	A08.800.800.720.450.760.640                      Peroneal Nerve
-	A08.800.800.720.450.760.820                      Tibial Nerve
-	A08.800.800.720.450.760.820.820                      Sural Nerve
-	A08.800.800.720.725                          Spinal Nerve Roots
-	A08.800.800.720.725.150                      Cauda Equina
-	A08.800.800.720.725.350                      Ganglia, Spinal
-	A08.800.800.720.800                          Thoracic Nerves
-	A08.800.800.720.800.350                      Intercostal Nerves
-	A08.800.950                                      Sensory Receptor Cells
-	A08.800.950.500                                  Chemoreceptor Cells
-	A08.800.950.500.530                          Neuroepithelial Cells
-	A08.800.950.500.530.550                      Neuroepithelial Bodies
-	A08.800.950.500.540                          Olfactory Receptor Neurons
-	A08.800.950.500.600                          Paraganglia, Nonchromaffin
-	A08.800.950.500.600.050                      Aortic Bodies
-	A08.800.950.500.600.150                      Carotid Body
-	A08.800.950.500.600.350                      Glomus Jugulare
-	A08.800.950.500.600.360                      Glomus Tympanicum
-	A08.800.950.500.800                          Taste Buds
-	A08.800.950.750                                  Mechanoreceptors
-	A08.800.950.750.300                          Golgi-Mazzoni Corpuscles



## MeSH Tree Changes for 2017

Type	Tree - heading
-	A08.800.950.750.425 Merkel Cells
-	A08.800.950.750.500 Muscle Spindles
-	A08.800.950.750.600 Neuroepithelial Cells
-	A08.800.950.750.600.350 Hair Cells, Auditory
-	A08.800.950.750.600.350.350 Hair Cells, Auditory, Inner
-	A08.800.950.750.600.350.365 Hair Cells, Auditory, Outer
-	A08.800.950.750.600.675 Hair Cells, Vestibular
-	A08.800.950.750.700 Pacinian Corpuscles
-	A08.800.950.750.750 Pressoreceptors
-	A08.800.950.750.780 Pulmonary Stretch Receptors
-	A08.800.950.875 Nociceptors
-	A08.800.950.937 Photoreceptor Cells
-	A08.800.950.937.650 Photoreceptor Cells, Invertebrate
-	A08.800.950.937.670 Photoreceptor Cells, Vertebrate
-	A08.800.950.937.670.049 Photoreceptor Connecting Cilium
-	A08.800.950.937.670.100 Retinal Cone Photoreceptor Cells
-	A08.800.950.937.670.237 Retinal Photoreceptor Cell Inner Segment
-	A08.800.950.937.670.375 Retinal Photoreceptor Cell Outer Segment
-	A08.800.950.937.670.375.500 Rod Cell Outer Segment
-	A08.800.950.937.670.650 Retinal Rod Photoreceptor Cells
-	A08.800.950.937.670.650.650 Rod Cell Outer Segment
-	A08.800.950.968 Thermoreceptors
-	A08.850 Synapses
-	A08.850.180 Electrical Synapses
-	A08.850.550 Neuroeffector Junction
-	A08.850.550.550 Neuromuscular Junction
-	A08.850.550.550.500 Motor Endplate
-	A08.850.700 Presynaptic Terminals
-	A08.850.700.500 Mossy Fibers, Hippocampal
-	A08.850.800 Synaptic Membranes
-	A08.850.800.500 Post-Synaptic Density
-	A08.850.840 Synaptic Vesicles
-	A09 Sense Organs
-	A09.246 Ear
-	A09.246.272 Ear, External
-	A09.246.272.197 Ear Auricle

## MeSH Tree Changes for 2017

Type	Tree - heading
-	A09.246.272.396 Ear Canal
-	A09.246.272.549 Ear Cartilage
-	A09.246.272.702 Tympanic Membrane
-	A09.246.397 Ear, Middle
-	A09.246.397.247 Ear Ossicles
-	A09.246.397.247.362 Incus
-	A09.246.397.247.524 Malleus
-	A09.246.397.247.806 Stapes
-	A09.246.397.369 Eustachian Tube
-	A09.246.397.400 Glomus Tympanicum
-	A09.246.397.727 Stapedius
-	A09.246.397.749 Tensor Tympani
-	A09.246.631 Ear, Inner
-	A09.246.631.246 Cochlea
-	A09.246.631.246.125 Basilar Membrane
-	A09.246.631.246.280 Cochlear Aqueduct
-	A09.246.631.246.292 Cochlear Duct
-	A09.246.631.246.292.876 Stria Vascularis
-	A09.246.631.246.292.938 Tectorial Membrane
-	A09.246.631.246.577 Organ of Corti
-	A09.246.631.246.577.325 Hair Cells, Auditory
-	A09.246.631.246.577.325.315 Hair Cells, Auditory, Inner
-	A09.246.631.246.577.325.380 Hair Cells, Auditory, Outer
-	A09.246.631.246.577.543 Labyrinth Supporting Cells
-	A09.246.631.246.814 Round Window, Ear
-	A09.246.631.246.848 Scala Tympani
-	A09.246.631.246.874 Scala Vestibuli
-	A09.246.631.246.900 Spiral Ganglion
-	A09.246.631.246.930 Spiral Lamina
-	A09.246.631.246.965 Spiral Ligament of Cochlea
-	A09.246.631.455 Labyrinthine Fluids
-	A09.246.631.663 Semicircular Canals
-	A09.246.631.663.500 Semicircular Ducts
-	A09.246.631.663.500.500 Hair Cells, Ampulla
-	A09.246.631.909 Vestibule, Labyrinth
-	A09.246.631.909.551 Oval Window, Ear

## MeSH Tree Changes for 2017

Type	Tree - heading
-	A09.246.631.909.625                      Sacculle and Utricle
-	A09.246.631.909.625.125                      Acoustic Maculae
-	A09.246.631.909.625.125.340                      Hair Cells, Vestibular
-	A09.246.631.909.625.125.680                      Otolithic Membrane
-	A09.246.631.909.957                      Vestibular Aqueduct
-	A09.246.631.909.957.360                      Endolymphatic Duct
-	A09.246.631.909.957.360.701                      Endolymphatic Sac
-	A09.371                      Eye
-	A09.371.060                      Anterior Eye Segment
-	A09.371.060.067                      Anterior Chamber
-	A09.371.060.067.070                      Aqueous Humor
-	A09.371.060.067.318                      Endothelium, Corneal
-	A09.371.060.160                      Ciliary Body
-	A09.371.060.200                      Conjunctiva
-	A09.371.060.217                      Cornea
-	A09.371.060.217.113                      Bowman Membrane
-	A09.371.060.217.228                      Corneal Stroma
-	A09.371.060.217.271                      Descemet Membrane
-	A09.371.060.217.318                      Endothelium, Corneal
-	A09.371.060.217.325                      Epithelium, Corneal
-	A09.371.060.217.659                      Limbus Corneae
-	A09.371.060.450                      Iris
-	A09.371.060.450.780                      Pupil
-	A09.371.060.500                      Lens, Crystalline
-	A09.371.060.500.155                      Lens Capsule, Crystalline
-	A09.371.060.500.155.500                      Posterior Capsule of the Lens
-	A09.371.060.500.225                      Lens Cortex, Crystalline
-	A09.371.060.500.670                      Lens Nucleus, Crystalline
-	A09.371.060.932                      Trabecular Meshwork
-	A09.371.061                      Anterior Capsule of the Lens
-	A09.371.199                      Axial Length, Eye
-	A09.371.337                      Eyelids
-	A09.371.337.168                      Conjunctiva
-	A09.371.337.338                      Eyelashes
-	A09.371.337.614                      Meibomian Glands
-	A09.371.463                      Lacrimal Apparatus

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	A09.371.463.640	Nasolacrimal Duct
Old Tree	A09.371.613	Oculomotor Muscles
-	A09.371.670	Pigment Epithelium of Eye
-	A09.371.670.500	Retinal Pigment Epithelium
-	A09.371.714	Posterior Eye Segment
-	A09.371.714.500	Vitreous Body
-	A09.371.729	Retina
-	A09.371.729.055	Blood-Retinal Barrier
-	A09.371.729.313	Fundus Oculi
-	A09.371.729.522	Macula Lutea
-	A09.371.729.522.436	Fovea Centralis
-	A09.371.729.690	Optic Disk
-	A09.371.729.831	Retinal Neurons
-	A09.371.729.831.500	Amacrine Cells
-	A09.371.729.831.625	Photoreceptor Cells
-	A09.371.729.831.625.660	Photoreceptor Cells, Invertebrate
-	A09.371.729.831.625.670	Photoreceptor Cells, Vertebrate
-	A09.371.729.831.625.670.049	Photoreceptor Connecting Cilium
-	A09.371.729.831.625.670.100	Retinal Cone Photoreceptor Cells
-	A09.371.729.831.625.670.237	Retinal Photoreceptor Cell Inner Segment
-	A09.371.729.831.625.670.375	Retinal Photoreceptor Cell Outer Segment
-	A09.371.729.831.625.670.375.500	Rod Cell Outer Segment
-	A09.371.729.831.625.670.650	Retinal Rod Photoreceptor Cells
-	A09.371.729.831.625.670.650.650	Rod Cell Outer Segment
-	A09.371.729.831.750	Retinal Bipolar Cells
-	A09.371.729.831.875	Retinal Ganglion Cells
-	A09.371.729.831.937	Retinal Horizontal Cells
-	A09.371.729.887	Retinal Pigment Epithelium
-	A09.371.784	Sclera
-	A09.371.839	Tenon Capsule
-	A09.371.894	Uvea
-	A09.371.894.030	Blood-Aqueous Barrier
-	A09.371.894.223	Choroid
-	A09.371.894.223.250	Bruch Membrane
-	A09.371.894.280	Ciliary Body
-	A09.371.894.513	Iris

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	A09.371.894.513.780	Pupil
-	A09.531	Nose
-	A09.531.623	Olfactory Mucosa
-	A09.531.623.580	Olfactory Receptor Neurons
-	A09.531.940	Vomeronasal Organ
-	A09.846	Taste Buds
-	A10	Tissues
-	A10.082	Body Fluid Compartments
-	A10.082.500	Extracellular Space
-	A10.082.750	Intracellular Space
-	A10.165	Connective Tissue
-	A10.165.114	Adipose Tissue
New Heading	<b>A10.165.114.161</b>	<b>Adipose Tissue, Beige</b>
-	A10.165.114.322	Adipose Tissue, Brown
-	A10.165.114.830	Adipose Tissue, White
-	A10.165.114.830.500	Abdominal Fat
-	A10.165.114.830.500.500	Intra-Abdominal Fat
-	A10.165.114.830.500.750	Subcutaneous Fat, Abdominal
-	A10.165.114.830.750	Subcutaneous Fat
-	A10.165.114.830.750.500	Subcutaneous Fat, Abdominal
-	A10.165.189	Adventitia
-	A10.165.265	Bone and Bones
-	A10.165.265.166	Bone Matrix
-	A10.165.265.183	Bone-Implant Interface
-	A10.165.265.200	Bony Callus
New Heading	<b>A10.165.265.414</b>	<b>Cancellous Bone</b>
Old Tree	<b>A10.165.265.507</b>	<b>Haversian System</b>
New Heading	<b>A10.165.265.521</b>	<b>Cortical Bone</b>
New Tree	<b>A10.165.265.521.500</b>	<b>Haversian System</b>
-	A10.165.265.746	Periosteum
-	A10.165.382	Cartilage
-	A10.165.382.300	Elastic Cartilage
-	A10.165.382.350	Fibrocartilage

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	<a href="#">A10.165.382.350.050</a>	<a href="#">Intervertebral Disc</a>
New Heading	<b>A10.165.382.350.050.250</b>	<b>Annulus Fibrosus</b>
New Heading	<b>A10.165.382.350.050.500</b>	<b>Nucleus Pulposus</b>
New Heading	<b>A10.165.382.350.163</b>	<b>Meniscus</b>
New Tree	<a href="#">A10.165.382.350.163.500</a>	<a href="#">Menisci, Tibial</a>
-	A10.165.382.350.200	Palmar Plate
-	A10.165.382.350.400	Plantar Plate
-	A10.165.382.350.800	Triangular Fibrocartilage
-	A10.165.382.400	Hyaline Cartilage
-	A10.165.400	Elastic Tissue
New Tree	<a href="#">A10.165.425</a>	<a href="#">Fascia</a>
-	A10.165.450	Granulation Tissue
-	A10.165.450.300	Cicatrix
-	A10.165.450.300.125	Cicatrix, Hypertrophic
-	A10.165.450.300.425	Keloid
-	A10.165.450.300.425.125	Acne Keloid
-	A10.165.887	Subcutaneous Tissue
-	A10.165.970	Wharton Jelly
-	A10.272	Epithelium
-	A10.272.220	Basement Membrane
-	A10.272.220.250	Bruch Membrane
-	A10.272.491	Endothelium
-	A10.272.491.318	Endothelium, Corneal
-	A10.272.491.332	Endothelium, Lymphatic
-	A10.272.491.355	Endothelium, Vascular
-	A10.272.491.677	Pericytes
-	A10.272.497	Epidermis
-	A10.272.497.500	Hair Follicle
-	A10.272.510	Epithelium, Corneal
-	A10.272.640	Pigment Epithelium of Eye
-	A10.272.700	Seminiferous Epithelium
-	A10.272.850	Urothelium

## MeSH Tree Changes for 2017

Type	Tree - heading
-	A10.336 Exocrine Glands
-	A10.336.095 Bartholin's Glands
-	A10.336.197 Bulbourethral Glands
-	A10.336.422 Lacrimal Apparatus
-	A10.336.482 Mammary Glands, Animal
-	A10.336.532 Mammary Glands, Human
-	A10.336.645 Pancreas, Exocrine
-	A10.336.707 Prostate
-	A10.336.779 Salivary Glands
-	A10.336.779.464 Parotid Gland
-	A10.336.779.640 Salivary Ducts
-	A10.336.779.650 Salivary Glands, Minor
-	A10.336.779.687 Sublingual Gland
-	A10.336.779.812 Submandibular Gland
-	A10.336.779.906 von Ebner Glands
-	A10.336.827 Sebaceous Glands
-	A10.336.827.600 Meibomian Glands
-	A10.336.899 Sweat Glands
-	A10.336.899.206 Apocrine Glands
-	A10.336.899.480 Eccrine Glands
-	A10.549 Lymphoid Tissue
-	A10.549.100 Adenoids
-	A10.549.175 Bursa of Fabricius
-	A10.549.400 Lymph Nodes
-	A10.549.400.500 Germinal Center
-	A10.549.400.500.200 Dendritic Cells, Follicular
New Heading	<b>A10.549.400.750 Sentinel Lymph Node</b>
-	A10.549.580 Palatine Tonsil
-	A10.549.600 Peyer's Patches
-	A10.549.700 Spleen
New Heading	<b>A10.549.725 Tertiary Lymphoid Structures</b>
-	A10.549.750 Thymus Gland
-	A10.615 Membranes
-	A10.615.179 Basement Membrane

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	A10.615.179.124	Basilar Membrane
-	A10.615.179.250	Bruch Membrane
-	A10.615.179.437	Descemet Membrane
-	A10.615.179.625	Glomerular Basement Membrane
-	A10.615.284	Extraembryonic Membranes
-	A10.615.284.147	Allantois
-	A10.615.284.277	Amnion
-	A10.615.284.375	Chorioallantoic Membrane
-	A10.615.284.473	Chorion
-	A10.615.284.473.200	Chorionic Villi
-	A10.615.284.981	Yolk Sac
-	A10.615.550	Mucous Membrane
New Heading	<b>A10.615.550.146</b>	<b>Esophageal Mucosa</b>
-	A10.615.550.291	Gastric Mucosa
-	A10.615.550.291.150	Chief Cells, Gastric
-	A10.615.550.291.162	Enterochromaffin Cells
-	A10.615.550.291.300	Gastrin-Secreting Cells
-	A10.615.550.291.650	Parietal Cells, Gastric
-	A10.615.550.291.825	Somatostatin-Secreting Cells
-	A10.615.550.368	Hymen
-	A10.615.550.444	Intestinal Mucosa
-	A10.615.550.444.290	Enterocytes
-	A10.615.550.444.321	Goblet Cells
-	A10.615.550.444.700	Paneth Cells
-	A10.615.550.599	Mouth Mucosa
-	A10.615.550.760	Respiratory Mucosa
-	A10.615.550.760.520	Laryngeal Mucosa
-	A10.615.550.760.520.320	Goblet Cells
-	A10.615.550.760.600	Nasal Mucosa
-	A10.615.550.760.600.320	Goblet Cells
-	A10.615.550.760.600.640	Olfactory Mucosa
-	A10.615.550.760.600.640.640	Olfactory Receptor Neurons
-	A10.615.789	Serous Membrane
-	A10.615.789.470	Pericardium
-	A10.615.789.470.500	Pericardial Fluid



## MeSH Tree Changes for 2017

Type	Tree - heading	
-	A10.615.789.596	Peritoneum
-	A10.615.789.736	Pleura
-	A10.690	Muscles
-	A10.690.467	Muscle, Smooth
-	A10.690.467.491	Muscle, Smooth, Vascular
-	A10.690.467.500	Myometrium
-	A10.690.552	Muscle, Striated
-	A10.690.552.500	Muscle, Skeletal
New Heading	<b>A10.690.552.500.250</b>	<b>Hamstring Muscles</b>
-	A10.690.552.500.500	Muscle Fibers, Skeletal
-	A10.690.552.500.500.600	Muscle Fibers, Fast-Twitch
-	A10.690.552.500.500.700	Muscle Fibers, Slow-Twitch
-	A10.690.552.500.500.850	Sarcoplasmic Reticulum
-	A10.690.552.500.750	Muscle Spindles
-	A10.690.552.750	Myocardium
-	A10.690.552.750.500	Myoblasts, Cardiac
-	A10.690.552.750.570	Myocytes, Cardiac
-	A10.690.552.875	Myofibrils
-	A10.690.552.875.700	Sarcomeres
-	A10.755	Nerve Tissue
-	A10.755.260	Ependyma
-	A10.755.503	Myelin Sheath
-	A10.755.503.550	Neurilemma
-	A10.755.503.700	Ranvier's Nodes
-	A10.802	Organoids
New Heading	<b>A10.806</b>	<b>Parenchymal Tissue</b>
-	A10.810	Peritoneal Stomata
New Heading	<b>A10.830</b>	<b>Margins of Excision</b>
-	A10.850	Surgically-Created Structures
-	A10.850.200	Colonic Pouches
-	A10.850.700	Skeletal Muscle Ventricle
-	A10.850.710	Surgical Flaps
-	A10.850.710.500	Free Tissue Flaps
-	A10.850.710.625	Myocutaneous Flap

## MeSH Tree Changes for 2017

Type	Tree - heading
-	A10.850.710.750                      Perforator Flap
-	A10.850.720                              Surgical Stomas
-	A10.850.750                              Urinary Reservoirs, Continent
-	A11    Cells
-	A11.031                                    Acinar Cells
-	A11.063                                    Antibody-Producing Cells
-	A11.063.438                              B-Lymphocytes
-	A11.063.438.450                        B-Lymphocyte Subsets
-	A11.063.438.450.300                    B-Lymphocytes, Regulatory
-	A11.063.438.725                        Plasma Cells
-	A11.066                                    Antigen-Presenting Cells
-	A11.066.270                              Dendritic Cells
-	A11.066.270.500                        Langerhans Cells
-	A11.066.275                              Dendritic Cells, Follicular
-	A11.118                                    Blood Cells
-	A11.118.188                              Blood Platelets
-	A11.118.290                              Erythrocytes
-	A11.118.290.270                        Erythrocyte Membrane
-	A11.118.290.330                        Erythrocytes, Abnormal
-	A11.118.290.330.100                    Acanthocytes
-	A11.118.290.330.315                    Erythrocyte Inclusions
-	A11.118.290.330.315.335              Heinz Bodies
-	A11.118.290.330.531                    Megaloblasts
-	A11.118.290.330.802                    Spherocytes
-	A11.118.290.760                        Reticulocytes
-	A11.118.480                              Hemocytes
-	A11.118.637                              Leukocytes
-	A11.118.637.415                        Granulocytes
-	A11.118.637.415.120                    Basophils
-	A11.118.637.415.345                    Eosinophils
-	A11.118.637.415.583                    Neutrophils
-	A11.118.637.555                        Leukocytes, Mononuclear
-	A11.118.637.555.283                    Cytokine-Induced Killer Cells
-	A11.118.637.555.283.500              Killer Cells, Lymphokine-Activated
-	A11.118.637.555.283.750              Monocytes, Activated Killer
-	A11.118.637.555.283.875              T-Lymphocytes, Cytotoxic

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	A11.118.637.555.567	Lymphocytes
-	A11.118.637.555.567.537	Killer Cells, Natural
-	A11.118.637.555.567.537.500	Killer Cells, Lymphokine-Activated
-	A11.118.637.555.567.550	Lymphocyte Subsets
-	A11.118.637.555.567.550.450	B-Lymphocyte Subsets
-	A11.118.637.555.567.550.450.300	B-Lymphocytes, Regulatory
-	A11.118.637.555.567.550.500	T-Lymphocyte Subsets
-	A11.118.637.555.567.550.500.200	T-Lymphocytes, Cytotoxic
-	A11.118.637.555.567.550.500.400	T-Lymphocytes, Helper-Inducer
-	A11.118.637.555.567.550.500.400.900	Th1 Cells
-	A11.118.637.555.567.550.500.400.905	Th2 Cells
-	A11.118.637.555.567.550.500.400.915	Th17 Cells
-	A11.118.637.555.567.550.500.700	T-Lymphocytes, Regulatory
-	A11.118.637.555.567.562	B-Lymphocytes
-	A11.118.637.555.567.562.200	B-Lymphocyte Subsets
-	A11.118.637.555.567.562.200.300	B-Lymphocytes, Regulatory
-	A11.118.637.555.567.562.725	Plasma Cells
-	A11.118.637.555.567.562.800	Precursor Cells, B-Lymphoid
-	A11.118.637.555.567.569	T-Lymphocytes
-	A11.118.637.555.567.569.200	CD4-Positive T-Lymphocytes
-	A11.118.637.555.567.569.200.400	T-Lymphocytes, Helper-Inducer
-	A11.118.637.555.567.569.200.400.900	Th1 Cells
-	A11.118.637.555.567.569.200.400.905	Th2 Cells
-	A11.118.637.555.567.569.200.400.915	Th17 Cells
-	A11.118.637.555.567.569.200.700	T-Lymphocytes, Regulatory
-	A11.118.637.555.567.569.220	CD8-Positive T-Lymphocytes
-	A11.118.637.555.567.569.220.200	T-Lymphocytes, Cytotoxic
-	A11.118.637.555.567.569.290	Natural Killer T-Cells
-	A11.118.637.555.567.569.360	Precursor Cells, T-Lymphoid
-	A11.118.637.555.567.569.360.800	Thymocytes
-	A11.118.637.555.567.569.500	T-Lymphocyte Subsets
New Heading	<b>A11.118.637.555.567.569.500.100</b>	<b>Mucosal-Associated Invariant T Cells</b>
-	A11.118.637.555.567.569.500.200	T-Lymphocytes, Cytotoxic
-	A11.118.637.555.567.569.500.400	T-Lymphocytes, Helper-Inducer
-	A11.118.637.555.567.569.500.400.900	Th1 Cells

## MeSH Tree Changes for 2017

Type	Tree - heading
-	A11.118.637.555.567.569.500.400.905 Th2 Cells
-	A11.118.637.555.567.569.500.400.915 Th17 Cells
-	A11.118.637.555.567.569.500.700 T-Lymphocytes, Regulatory
-	A11.118.637.555.567.622 Lymphocytes, Null
-	A11.118.637.555.567.650 Lymphocytes, Tumor-Infiltrating
-	A11.118.637.555.652 Monocytes
-	A11.118.637.555.652.500 Monocytes, Activated Killer
-	A11.148 Bone Marrow Cells
-	A11.148.174 Endothelial Progenitor Cells
-	A11.148.350 Granulocytes
-	A11.148.350.350 Granulocyte Precursor Cells
-	A11.148.378 Hematopoietic Stem Cells
-	A11.148.378.294 Lymphoid Progenitor Cells
-	A11.148.378.294.374 Precursor Cells, B-Lymphoid
-	A11.148.378.294.750 Precursor Cells, T-Lymphoid
-	A11.148.378.294.750.800 Thymocytes
-	A11.148.378.590 Myeloid Progenitor Cells
-	A11.148.378.590.675 Granulocyte-Macrophage Progenitor Cells
-	A11.148.378.590.675.500 Granulocyte Precursor Cells
-	A11.148.378.590.675.750 Monocyte-Macrophage Precursor Cells
-	A11.148.378.590.837 Megakaryocyte-Erythroid Progenitor Cells
-	A11.148.378.590.837.250 Erythroid Precursor Cells
-	A11.148.378.590.837.250.200 Erythroblasts
-	A11.148.378.590.837.250.200.500 Megaloblasts
-	A11.148.378.590.837.625 Megakaryocyte Progenitor Cells
-	A11.148.479 Megakaryocytes
-	A11.148.580 Monocytes
-	A11.148.580.500 Monocyte-Macrophage Precursor Cells
-	A11.148.790 Reticulocytes
-	A11.251 Cells, Cultured
-	A11.251.210 Cell Line
-	A11.251.210.100 3T3 Cells
-	A11.251.210.100.080 BALB 3T3 Cells
-	A11.251.210.100.550 NIH 3T3 Cells
-	A11.251.210.100.775 Swiss 3T3 Cells
-	A11.251.210.100.775.800 3T3-L1 Cells

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	A11.251.210.172	Cell Line, Transformed
-	A11.251.210.172.500	COS Cells
-	A11.251.210.172.750	HEK293 Cells
-	A11.251.210.172.875	RAW 264.7 Cells
-	A11.251.210.190	Cell Line, Tumor
New Heading	<b>A11.251.210.190.080</b>	<b>A549 Cells</b>
-	A11.251.210.190.160	Caco-2 Cells
-	A11.251.210.190.380	HCT116 Cells
-	A11.251.210.190.400	HeLa Cells
-	A11.251.210.190.400.500	KB Cells
-	A11.251.210.190.465	HL-60 Cells
-	A11.251.210.190.475	HT29 Cells
-	A11.251.210.190.495	Jurkat Cells
-	A11.251.210.190.510	K562 Cells
-	A11.251.210.190.630	MCF-7 Cells
-	A11.251.210.190.750	PC12 Cells
-	A11.251.210.190.880	U937 Cells
-	A11.251.210.200	CHO Cells
-	A11.251.210.505	L Cells (Cell Line)
-	A11.251.210.520	LLC-PK1 Cells
-	A11.251.210.827	Madin Darby Canine Kidney Cells
-	A11.251.210.891	Sf9 Cells
-	A11.251.210.955	Vero Cells
-	A11.251.353	Clone Cells
-	A11.251.353.485	Hybridomas
-	A11.251.476	Cytokine-Induced Killer Cells
-	A11.251.538	Feeder Cells
-	A11.251.600	Hybrid Cells
-	A11.251.600.485	Hybridomas
-	A11.251.800	Spheroids, Cellular
-	A11.251.860	Tumor Cells, Cultured
-	A11.251.860.180	Cell Line, Tumor
New Heading	<b>A11.251.860.180.080</b>	<b>A549 Cells</b>
-	A11.251.860.180.160	Caco-2 Cells

## MeSH Tree Changes for 2017

Type	Tree - heading
-	A11.251.860.180.380 HCT116 Cells
-	A11.251.860.180.400 HeLa Cells
-	A11.251.860.180.400.500 KB Cells
-	A11.251.860.180.432 Hep G2 Cells
-	A11.251.860.180.465 HL-60 Cells
-	A11.251.860.180.475 HT29 Cells
-	A11.251.860.180.495 Jurkat Cells
-	A11.251.860.180.510 K562 Cells
-	A11.251.860.180.750 PC12 Cells
-	A11.251.860.180.880 U937 Cells
-	A11.251.860.590 Embryonal Carcinoma Stem Cells
-	A11.270 Cells, Immobilized
-	A11.284 Cellular Structures
-	A11.284.074 Cell Body
-	A11.284.149 Cell Membrane
-	A11.284.149.165 Cell Membrane Structures
-	A11.284.149.165.165 Cell-Matrix Junctions
-	A11.284.149.165.165.142 Costameres
-	A11.284.149.165.165.285 Focal Adhesions
-	A11.284.149.165.165.360 Hemidesmosomes
-	A11.284.149.165.175 Coated Pits, Cell-Membrane
-	A11.284.149.165.175.160 Caveolae
-	A11.284.149.165.355 Glycocalyx
-	A11.284.149.165.420 Intercellular Junctions
-	A11.284.149.165.420.020 Adherens Junctions
-	A11.284.149.165.420.297 Desmosomes
-	A11.284.149.165.420.471 Gap Junctions
-	A11.284.149.165.420.471.180 Electrical Synapses
-	A11.284.149.165.420.548 Immunological Synapses
-	A11.284.149.165.420.625 Plasmodesmata
-	A11.284.149.165.420.780 Synapses
-	A11.284.149.165.420.780.180 Electrical Synapses
-	A11.284.149.165.420.780.550 Neuroeffector Junction
-	A11.284.149.165.420.780.550.550 Neuromuscular Junction
-	A11.284.149.165.420.780.550.550.500 Motor Endplate
-	A11.284.149.165.420.780.700 Presynaptic Terminals

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	A11.284.149.165.420.780.800	Synaptic Membranes
-	A11.284.149.165.420.780.800.500	Post-Synaptic Density
-	A11.284.149.165.420.820	Tight Junctions
-	A11.284.149.165.570	Membrane Microdomains
-	A11.284.149.165.570.160	Caveolae
-	A11.284.149.165.600	Myelin Sheath
-	A11.284.149.165.630	Nuclear Envelope
-	A11.284.149.165.630.500	Nuclear Lamina
-	A11.284.149.165.695	Phycobilisomes
-	A11.284.149.165.760	Ranvier's Nodes
-	A11.284.149.356	Erythrocyte Membrane
-	A11.284.149.450	Intracellular Membranes
-	A11.284.149.450.349	Mitochondrial Membranes
-	A11.284.149.450.700	Nuclear Envelope
-	A11.284.149.648	Purple Membrane
-	A11.284.149.707	Sarcolemma
-	A11.284.149.844	Synaptic Membranes
-	A11.284.180	Cell Surface Extensions
-	A11.284.180.075	Axons
New Heading	<b>A11.284.180.075.125</b>	<b>Axon Initial Segment</b>
-	A11.284.180.075.249	Growth Cones
-	A11.284.180.120	Cellulosomes
-	A11.284.180.165	Cilia
-	A11.284.180.225	Dendrites
-	A11.284.180.225.169	Dendritic Spines
-	A11.284.180.225.340	Growth Cones
-	A11.284.180.285	Fimbriae, Bacterial
-	A11.284.180.290	Flagella
-	A11.284.180.290.835	Sperm Tail
-	A11.284.180.565	Microvilli
-	A11.284.180.610	Neurites
-	A11.284.180.610.345	Growth Cones
-	A11.284.180.690	Pili, Sex
-	A11.284.180.695	Podosomes
-	A11.284.180.700	Pseudopodia

## MeSH Tree Changes for 2017

Type	Tree - heading
-	A11.284.180.800 Stereocilia
-	A11.284.180.900 Telopodes
-	A11.284.183 Cell Wall
-	A11.284.183.200 Cell Wall Skeleton
-	A11.284.187 Chromosomes
-	A11.284.187.167 Chromosomes, Archaeal
-	A11.284.187.178 Chromosomes, Artificial
-	A11.284.187.178.170 Chromosomes, Artificial, Bacterial
-	A11.284.187.178.190 Chromosomes, Artificial, Mammalian
-	A11.284.187.178.190.117 Chromosomes, Artificial, Human
-	A11.284.187.178.195 Chromosomes, Artificial, P1 Bacteriophage
-	A11.284.187.178.200 Chromosomes, Artificial, Yeast
-	A11.284.187.190 Chromosomes, Bacterial
-	A11.284.187.190.170 Chromosomes, Artificial, Bacterial
-	A11.284.187.360 Chromosomes, Fungal
-	A11.284.187.360.800 Chromosomes, Artificial, Yeast
-	A11.284.187.440 Chromosomes, Insect
-	A11.284.187.440.500 Polytene Chromosomes
-	A11.284.187.520 Chromosomes, Mammalian
-	A11.284.187.520.190 Chromosomes, Artificial, Mammalian
-	A11.284.187.520.190.117 Chromosomes, Artificial, Human
-	A11.284.187.520.300 Chromosomes, Human
-	A11.284.187.520.300.117 Chromosomes, Artificial, Human
-	A11.284.187.520.300.235 Chromosomes, Human, 1-3
-	A11.284.187.520.300.235.240 Chromosomes, Human, Pair 1
-	A11.284.187.520.300.235.245 Chromosomes, Human, Pair 2
-	A11.284.187.520.300.235.250 Chromosomes, Human, Pair 3
-	A11.284.187.520.300.280 Chromosomes, Human, 4-5
-	A11.284.187.520.300.280.285 Chromosomes, Human, Pair 4
-	A11.284.187.520.300.280.290 Chromosomes, Human, Pair 5
-	A11.284.187.520.300.325 Chromosomes, Human, 6-12 and X
-	A11.284.187.520.300.325.330 Chromosomes, Human, Pair 6
-	A11.284.187.520.300.325.335 Chromosomes, Human, Pair 7
-	A11.284.187.520.300.325.340 Chromosomes, Human, Pair 8
-	A11.284.187.520.300.325.345 Chromosomes, Human, Pair 9
-	A11.284.187.520.300.325.345.500 Philadelphia Chromosome



## MeSH Tree Changes for 2017

Type	Tree - heading	
-	A11.284.187.520.300.325.350	Chromosomes, Human, Pair 10
-	A11.284.187.520.300.325.355	Chromosomes, Human, Pair 11
-	A11.284.187.520.300.325.360	Chromosomes, Human, Pair 12
-	A11.284.187.520.300.325.680	Chromosomes, Human, X
-	A11.284.187.520.300.370	Chromosomes, Human, 13-15
-	A11.284.187.520.300.370.375	Chromosomes, Human, Pair 13
-	A11.284.187.520.300.370.380	Chromosomes, Human, Pair 14
-	A11.284.187.520.300.370.385	Chromosomes, Human, Pair 15
-	A11.284.187.520.300.415	Chromosomes, Human, 16-18
-	A11.284.187.520.300.415.420	Chromosomes, Human, Pair 16
-	A11.284.187.520.300.415.425	Chromosomes, Human, Pair 17
-	A11.284.187.520.300.415.430	Chromosomes, Human, Pair 18
-	A11.284.187.520.300.460	Chromosomes, Human, 19-20
-	A11.284.187.520.300.460.465	Chromosomes, Human, Pair 19
-	A11.284.187.520.300.460.470	Chromosomes, Human, Pair 20
-	A11.284.187.520.300.505	Chromosomes, Human, 21-22 and Y
-	A11.284.187.520.300.505.510	Chromosomes, Human, Pair 21
-	A11.284.187.520.300.505.515	Chromosomes, Human, Pair 22
-	A11.284.187.520.300.505.515.500	Philadelphia Chromosome
-	A11.284.187.520.300.505.757	Chromosomes, Human, Y
-	A11.284.187.560	Chromosomes, Plant
-	A11.284.187.570	Isochromosomes
-	A11.284.187.788	Ring Chromosomes
-	A11.284.187.865	Sex Chromosomes
New Heading	<b>A11.284.187.865.400</b>	<b>Pseudoautosomal Regions</b>
-	A11.284.187.865.800	Sex Chromatin
-	A11.284.187.865.982	X Chromosome
-	A11.284.187.865.982.500	Chromosomes, Human, X
-	A11.284.187.865.983	Y Chromosome
-	A11.284.187.865.983.500	Chromosomes, Human, Y
-	A11.284.295	Extracellular Space
-	A11.284.295.260	Extracellular Fluid
-	A11.284.295.310	Extracellular Matrix
-	A11.284.295.310.279	Glomerular Basement Membrane
-	A11.284.295.310.560	Microfibrils

## MeSH Tree Changes for 2017

Type	Tree - heading
-	A11.284.295.310.990                      Zona Pellucida
-	A11.284.295.495                            Extracellular Traps
-	A11.284.295.588                           Extracellular Vesicles
-	A11.284.295.588.500                    Cell-Derived Microparticles
-	A11.284.295.588.750                    Exosomes
-	A11.284.295.680                           Periplasm
-	A11.284.420                                Inclusion Bodies
-	A11.284.420.204                         Erythrocyte Inclusions
-	A11.284.420.204.335                    Heinz Bodies
-	A11.284.420.390                         Inclusion Bodies, Viral
-	A11.284.420.400                         Intranuclear Inclusion Bodies
-	A11.284.420.460                         Lewy Bodies
-	A11.284.420.500                         Mallory Bodies
-	A11.284.430                                Intracellular Space
-	A11.284.430.106                         Cell Nucleus
-	A11.284.430.106.279                    Cell Nucleus Structures
-	A11.284.430.106.279.345                Intranuclear Space
-	A11.284.430.106.279.345.175           Cell Nucleolus
-	A11.284.430.106.279.345.190           Chromosomes
-	A11.284.430.106.279.345.190.160            Chromosome Structures
-	A11.284.430.106.279.345.190.160.165            Centromere
-	A11.284.430.106.279.345.190.160.165.500            Kinetochores
-	A11.284.430.106.279.345.190.160.175            Chromatids
-	A11.284.430.106.279.345.190.160.180            Chromatin
-	A11.284.430.106.279.345.190.160.180.270            Euchromatin
-	A11.284.430.106.279.345.190.160.180.270.500            Chromosomal Puffs
-	A11.284.430.106.279.345.190.160.180.383            Heterochromatin
-	A11.284.430.106.279.345.190.160.180.383.800            Sex Chromatin
-	A11.284.430.106.279.345.190.160.180.625            Nucleosomes
-	A11.284.430.106.279.345.190.160.650            Nucleolus Organizer Region
-	A11.284.430.106.279.345.190.160.830            Synaptonemal Complex
-	A11.284.430.106.279.345.190.160.845            Telomere
-	A11.284.430.106.279.345.190.580            Polytene Chromosomes
-	A11.284.430.106.279.345.190.580.500            Chromosomal Puffs
-	A11.284.430.106.279.345.195            Coiled Bodies
-	A11.284.430.106.279.345.447            Gemini of Coiled Bodies

## MeSH Tree Changes for 2017

Type	Tree - heading
-	A11.284.430.106.279.345.700 Nuclear Matrix
-	A11.284.430.106.279.345.700.700 Nuclear Lamina
-	A11.284.430.106.279.345.850 Spliceosomes
-	A11.284.430.106.279.692 Nuclear Envelope
-	A11.284.430.106.279.692.314 Nuclear Lamina
-	A11.284.430.106.279.692.630 Nuclear Pore
-	A11.284.430.106.550 Macronucleus
-	A11.284.430.106.570 Micronuclei, Chromosome-Defective
-	A11.284.430.106.575 Micronucleus, Germline
-	A11.284.430.214 Cytoplasm
-	A11.284.430.214.190 Cytoplasmic Structures
-	A11.284.430.214.190.500 Cytoplasmic Granules
-	A11.284.430.214.190.500.207 Chromaffin Granules
-	A11.284.430.214.190.500.560 Melanosomes
-	A11.284.430.214.190.500.585 Microbodies
-	A11.284.430.214.190.500.585.250 Glyoxysomes
-	A11.284.430.214.190.500.585.600 Peroxisomes
-	A11.284.430.214.190.500.610 Nissl Bodies
-	A11.284.430.214.190.500.950 Weibel-Palade Bodies
-	A11.284.430.214.190.750 Cytoskeleton
-	A11.284.430.214.190.750.050 Actin Cytoskeleton
-	A11.284.430.214.190.750.050.414 Filamins
-	A11.284.430.214.190.750.050.830 Stress Fibers
-	A11.284.430.214.190.750.410 Intermediate Filaments
-	A11.284.430.214.190.750.585 Microtubule-Organizing Center
-	A11.284.430.214.190.750.585.079 Basal Bodies
-	A11.284.430.214.190.750.585.160 Centrosome
-	A11.284.430.214.190.750.585.160.130 Centrioles
-	A11.284.430.214.190.750.585.580 Spindle Pole Bodies
-	A11.284.430.214.190.750.602 Microtubules
-	A11.284.430.214.190.750.602.309 Axoneme
-	A11.284.430.214.190.750.602.620 Neuropil Threads
-	A11.284.430.214.190.750.620 Myofibrils
-	A11.284.430.214.190.750.640 Neurofibrils
-	A11.284.430.214.190.750.640.520 Neurofibrillary Tangles
-	A11.284.430.214.190.750.820 Spindle Apparatus

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	A11.284.430.214.190.750.820.500	Spindle Poles
-	A11.284.430.214.190.750.820.500.500	Centrosome
-	A11.284.430.214.190.750.820.500.500.130	Centrioles
-	A11.284.430.214.190.750.820.500.750	Spindle Pole Bodies
-	A11.284.430.214.190.875	Organelles
-	A11.284.430.214.190.875.080	Bacterial Chromatophores
-	A11.284.430.214.190.875.117	Cell Nucleus
-	A11.284.430.214.190.875.117.550	Macronucleus
-	A11.284.430.214.190.875.117.570	Micronuclei, Chromosome-Defective
-	A11.284.430.214.190.875.117.575	Micronucleus, Germline
-	A11.284.430.214.190.875.190	Cytoplasmic Vesicles
-	A11.284.430.214.190.875.190.190	Cytoplasmic Granules
-	A11.284.430.214.190.875.190.190.207	Chromaffin Granules
-	A11.284.430.214.190.875.190.190.560	Melanosomes
-	A11.284.430.214.190.875.190.190.755	Microbodies
-	A11.284.430.214.190.875.190.190.755.250	Glyoxysomes
-	A11.284.430.214.190.875.190.190.755.600	Peroxisomes
-	A11.284.430.214.190.875.190.190.950	Weibel-Palade Bodies
-	A11.284.430.214.190.875.190.550	Lysosomes
-	A11.284.430.214.190.875.190.550.040	Acrosome
-	A11.284.430.214.190.875.190.700	Phagosomes
New Heading	<b>A11.284.430.214.190.875.190.700.500</b>	<b>Autophagosomes</b>
-	A11.284.430.214.190.875.190.880	Transport Vesicles
-	A11.284.430.214.190.875.190.880.180	Coated Vesicles
-	A11.284.430.214.190.875.190.880.180.160	Caveolae
-	A11.284.430.214.190.875.190.880.180.170	Clathrin-Coated Vesicles
-	A11.284.430.214.190.875.190.880.180.180	COP-Coated Vesicles
-	A11.284.430.214.190.875.190.880.337	Endosomes
-	A11.284.430.214.190.875.190.880.337.500	Multivesicular Bodies
-	A11.284.430.214.190.875.190.880.495	Exosomes
-	A11.284.430.214.190.875.190.880.810	Secretory Vesicles
-	A11.284.430.214.190.875.190.880.830	Synaptic Vesicles
-	A11.284.430.214.190.875.190.920	Vacuoles
-	A11.284.430.214.190.875.248	Endoplasmic Reticulum
-	A11.284.430.214.190.875.248.300	Endoplasmic Reticulum, Rough

## MeSH Tree Changes for 2017

Type	Tree - heading
-	A11.284.430.214.190.875.248.300.610 Nissl Bodies
-	A11.284.430.214.190.875.248.310 Endoplasmic Reticulum, Smooth
-	A11.284.430.214.190.875.248.310.800 Sarcoplasmic Reticulum
-	A11.284.430.214.190.875.336 Golgi Apparatus
-	A11.284.430.214.190.875.336.850 trans-Golgi Network
-	A11.284.430.214.190.875.393 Lipid Droplets
-	A11.284.430.214.190.875.450 Magnetosomes
-	A11.284.430.214.190.875.564 Mitochondria
-	A11.284.430.214.190.875.564.461 Mitochondria, Liver
-	A11.284.430.214.190.875.564.627 Mitochondria, Muscle
-	A11.284.430.214.190.875.564.627.603 Mitochondria, Heart
-	A11.284.430.214.190.875.564.755 Mitochondrial Ribosomes
-	A11.284.430.214.190.875.564.882 Submitochondrial Particles
-	A11.284.430.214.190.875.700 Plastids
-	A11.284.430.214.190.875.700.069 Apicoplasts
-	A11.284.430.214.190.875.700.140 Chloroplasts
-	A11.284.430.214.190.875.700.140.800 Thylakoids
-	A11.284.430.214.190.875.811 Ribosomes
-	A11.284.430.214.190.875.811.370 Mitochondrial Ribosomes
-	A11.284.430.214.190.875.811.740 Polyribosomes
-	A11.284.430.214.190.875.811.740.300 Endoplasmic Reticulum, Rough
-	A11.284.430.214.190.875.811.740.300.610 Nissl Bodies
-	A11.284.430.214.190.875.811.870 Ribosome Subunits
-	A11.284.430.214.190.875.811.870.700 Ribosome Subunits, Large
-	A11.284.430.214.190.875.811.870.700.349 Ribosome Subunits, Large, Archaeal
-	A11.284.430.214.190.875.811.870.700.700 Ribosome Subunits, Large, Bacterial
-	A11.284.430.214.190.875.811.870.700.750 Ribosome Subunits, Large, Eukaryotic
-	A11.284.430.214.190.875.811.870.750 Ribosome Subunits, Small
-	A11.284.430.214.190.875.811.870.750.349 Ribosome Subunits, Small, Archaeal
-	A11.284.430.214.190.875.811.870.750.700 Ribosome Subunits, Small, Bacterial
-	A11.284.430.214.190.875.811.870.750.750 Ribosome Subunits, Small, Eukaryotic
-	A11.284.430.214.190.875.820 Sarcomeres

## MeSH Tree Changes for 2017

Type	Tree - heading
-	A11.284.430.214.190.875.930 Vault Ribonucleoprotein Particles
-	A11.284.430.214.200 Cytosol
-	A11.284.430.429 Intracellular Fluid
-	A11.284.430.429.200 Cytosol
-	A11.284.835 Subcellular Fractions
-	A11.284.835.168 Cell-Free System
-	A11.284.835.450 Intracellular Fluid
-	A11.284.835.450.200 Cytosol
-	A11.284.835.514 Intracellular Membranes
-	A11.284.835.514.349 Mitochondrial Membranes
-	A11.284.835.514.700 Nuclear Envelope
-	A11.284.835.514.700.500 Nuclear Lamina
-	A11.284.835.540 Microsomes
-	A11.284.835.540.541 Microsomes, Liver
-	A11.284.835.626 Mitochondria
-	A11.284.835.626.461 Mitochondria, Liver
-	A11.284.835.626.627 Mitochondria, Muscle
-	A11.284.835.626.627.603 Mitochondria, Heart
-	A11.284.835.626.882 Submitochondrial Particles
-	A11.284.835.859 Synaptosomes
-	A11.284.917 Trichomes
-	A11.299 Chromaffin Cells
-	A11.299.500 PC12 Cells
-	A11.329 Connective Tissue Cells
-	A11.329.114 Adipocytes
New Heading	<b>A11.329.114.125 Adipocytes, Beige</b>
-	A11.329.114.249 Adipocytes, Brown
-	A11.329.114.500 Adipocytes, White
-	A11.329.171 Chondrocytes
-	A11.329.228 Fibroblasts
-	A11.329.228.100 3T3 Cells
-	A11.329.228.100.080 BALB 3T3 Cells
-	A11.329.228.100.550 NIH 3T3 Cells
-	A11.329.228.100.775 Swiss 3T3 Cells
-	A11.329.228.100.775.800 3T3-L1 Cells

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Heading	<b>A11.329.228.105</b>	<b>Cancer-Associated Fibroblasts</b>
-	A11.329.228.109	Corneal Keratocytes
-	A11.329.228.220	COS Cells
-	A11.329.228.505	L Cells (Cell Line)
-	A11.329.228.950	Mesangial Cells
-	A11.329.228.975	Myofibroblasts
-	A11.329.372	Macrophages
-	A11.329.372.300	Epithelioid Cells
-	A11.329.372.368	Foam Cells
-	A11.329.372.376	Giant Cells, Foreign-Body
-	A11.329.372.380	Giant Cells, Langhans
-	A11.329.372.385	Histiocytes
-	A11.329.372.588	Kupffer Cells
-	A11.329.372.600	Macrophages, Alveolar
-	A11.329.372.630	Macrophages, Peritoneal
-	A11.329.372.700	Osteoclasts
-	A11.329.427	Mast Cells
-	A11.329.522	Odontoblasts
-	A11.329.629	Osteoblasts
-	A11.329.629.500	Osteocytes
-	A11.329.830	Stromal Cells
-	A11.329.830.500	Mesenchymal Stromal Cells
-	A11.329.830.750	Telocytes
-	A11.329.830.750.500	Telopodes
New Heading	<b>A11.329.835</b>	<b>Synoviocytes</b>
New Heading	<b>A11.329.840</b>	<b>Tenocytes</b>
-	A11.329.850	Theca Cells
-	A11.382	Endocrine Cells
-	A11.382.249	APUD Cells
-	A11.382.500	Corticotrophs
-	A11.382.625	Enteroendocrine Cells
-	A11.382.625.031	Enterochromaffin Cells
-	A11.382.625.047	Enterochromaffin-like Cells
-	A11.382.625.055	Gastrin-Secreting Cells

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	A11.382.625.064	Glucagon-Secreting Cells
-	A11.382.625.092	Insulin-Secreting Cells
-	A11.382.625.900	Pancreatic Polypeptide-Secreting Cells
-	A11.382.625.950	Somatostatin-Secreting Cells
-	A11.382.750	Gonadotrophs
-	A11.382.812	Granulosa Cells
-	A11.382.812.500	Cumulus Cells
-	A11.382.875	Lactotrophs
-	A11.382.906	Leydig Cells
-	A11.382.921	Luteal Cells
-	A11.382.937	Melanotrophs
-	A11.382.944	Neuroendocrine Cells
-	A11.382.952	Sertoli Cells
-	A11.382.968	Somatotrophs
-	A11.382.984	Thyrotrophs
-	A11.382.992	Trophoblasts
-	A11.436	Epithelial Cells
New Heading	<b>A11.436.054</b>	<b>A549 Cells</b>
New Tree	<a href="#">A11.436.081</a>	<a href="#">Alveolar Epithelial Cells</a>
New Tree	<a href="#">A11.436.081</a>	<a href="#">Pneumocytes</a>
-	A11.436.107	Ameloblasts
-	A11.436.123	Atypical Squamous Cells of the Cervix
-	A11.436.140	Caco-2 Cells
-	A11.436.150	Chief Cells, Gastric
-	A11.436.155	CHO Cells
-	A11.436.265	Chromatophores
-	A11.436.265.531	Melanophores
-	A11.436.265.531.560	Melanosomes
-	A11.436.267	Corticotrophs
-	A11.436.270	Dendritic Cells
-	A11.436.270.545	Langerhans Cells
-	A11.436.275	Endothelial Cells
-	A11.436.275.182	Endothelial Progenitor Cells
-	A11.436.275.365	Hemangioblasts



## MeSH Tree Changes for 2017

Type	Tree - heading
-	A11.436.275.682 Human Umbilical Vein Endothelial Cells
-	A11.436.275.841 Podosomes
-	A11.436.290 Enterocytes
-	A11.436.294 Enteroendocrine Cells
-	A11.436.294.031 Enterochromaffin Cells
-	A11.436.294.047 Enterochromaffin-like Cells
-	A11.436.294.055 Gastrin-Secreting Cells
-	A11.436.294.064 Glucagon-Secreting Cells
-	A11.436.294.092 Insulin-Secreting Cells
-	A11.436.294.900 Pancreatic Polypeptide-Secreting Cells
-	A11.436.294.950 Somatostatin-Secreting Cells
-	A11.436.298 Goblet Cells
-	A11.436.319 Gonadotrophs
-	A11.436.329 Granulosa Cells
-	A11.436.329.500 Cumulus Cells
-	A11.436.334 HEK293 Cells
-	A11.436.340 HeLa Cells
-	A11.436.340.500 KB Cells
-	A11.436.348 Hepatocytes
-	A11.436.348.500 Hep G2 Cells
-	A11.436.365 HT29 Cells
-	A11.436.397 Keratinocytes
-	A11.436.495 Labyrinth Supporting Cells
-	A11.436.507 Lactotrophs
-	A11.436.513 Leydig Cells
-	A11.436.520 LLC-PK1 Cells
-	A11.436.566 Luteal Cells
-	A11.436.589 Madin Darby Canine Kidney Cells
-	A11.436.613 Melanocytes
-	A11.436.613.560 Melanosomes
-	A11.436.636 Melanotrophs
-	A11.436.660 Merkel Cells
-	A11.436.690 Neuroepithelial Cells
-	A11.436.690.550 Neuroepithelial Bodies
-	A11.436.700 Paneth Cells
-	A11.436.708 Parietal Cells, Gastric

## MeSH Tree Changes for 2017

Type	Tree - heading	
Old Tree	A11.436.714	Alveolar Epithelial Cells
Old Tree	A11.436.714	Pneumocytes
-	A11.436.720	Podocytes
-	A11.436.837	Sertoli Cells
-	A11.436.896	Somatotrophs
New Heading	<b>A11.436.911</b>	<b>Thyroid Epithelial Cells</b>
-	A11.436.925	Thyrotrophs
-	A11.436.955	Vero Cells
-	A11.443	Erythroid Cells
-	A11.443.240	Erythrocytes
-	A11.443.240.330	Erythrocytes, Abnormal
-	A11.443.240.330.100	Acanthocytes
-	A11.443.240.330.531	Megaloblasts
-	A11.443.240.330.802	Spherocytes
-	A11.443.240.497	Erythroid Precursor Cells
-	A11.443.240.497.200	Erythroblasts
-	A11.443.240.497.200.500	Megaloblasts
-	A11.443.240.497.480	K562 Cells
-	A11.443.240.665	Reticulocytes
-	A11.450	Eukaryotic Cells
-	A11.497	Germ Cells
-	A11.497.124	Embryonic Germ Cells
-	A11.497.248	Germ Cells, Plant
-	A11.497.497	Ovum
-	A11.497.497.600	Oocytes
-	A11.497.497.600.500	Polar Bodies
-	A11.497.497.650	Oogonia
-	A11.497.497.900	Zona Pellucida
-	A11.497.497.950	Zygote
-	A11.497.760	Spermatozoa
-	A11.497.760.400	Sperm Head
-	A11.497.760.400.100	Acrosome
-	A11.497.760.450	Sperm Midpiece
-	A11.497.760.500	Sperm Tail
-	A11.497.760.600	Spermatids

## MeSH Tree Changes for 2017

Type	Tree - heading
-	A11.497.760.700                      Spermatocytes
-	A11.497.760.800                      Spermatogonia
-	A11.500                                      Giant Cells
-	A11.500.376                              Giant Cells, Foreign-Body
-	A11.500.380                              Giant Cells, Langhans
-	A11.561                                      Hepatic Stellate Cells
-	A11.600                                      Mesophyll Cells
-	A11.620                                      Muscle Cells
-	A11.620.124                              Interstitial Cells of Cajal
-	A11.620.249                              Muscle Fibers, Skeletal
-	A11.620.249.400                        Muscle Fibers, Fast-Twitch
-	A11.620.249.700                        Muscle Fibers, Slow-Twitch
-	A11.620.249.850                        Myofibrils
-	A11.620.249.850.700                    Sarcomeres
-	A11.620.500                              Myocytes, Cardiac
-	A11.620.500.500                        Myofibrils
-	A11.620.500.500.700                    Sarcomeres
-	A11.620.520                              Myocytes, Smooth Muscle
-	A11.620.520.500                        Myofibroblasts
-	A11.627                                      Myeloid Cells
-	A11.627.340                              Granulocytes
-	A11.627.340.120                        Basophils
-	A11.627.340.345                        Eosinophils
-	A11.627.340.360                        Granulocyte Precursor Cells
-	A11.627.340.360.500                    HL-60 Cells
-	A11.627.340.583                        Neutrophils
-	A11.627.482                              Macrophages
-	A11.627.482.300                        Epithelioid Cells
-	A11.627.482.368                        Foam Cells
-	A11.627.482.376                        Giant Cells, Foreign-Body
-	A11.627.482.380                        Giant Cells, Langhans
-	A11.627.482.385                        Histiocytes
-	A11.627.482.588                        Kupffer Cells
-	A11.627.482.600                        Macrophages, Alveolar
-	A11.627.482.630                        Macrophages, Peritoneal
-	A11.627.482.665                        Monocyte-Macrophage Precursor Cells

## MeSH Tree Changes for 2017

Type	Tree - heading
-	A11.627.482.665.500 U937 Cells
-	A11.627.482.700 Osteoclasts
-	A11.627.624 Monocytes
-	A11.627.624.249 Monocyte-Macrophage Precursor Cells
-	A11.627.624.249.500 U937 Cells
-	A11.627.624.500 Monocytes, Activated Killer
-	A11.627.635 Myeloid Progenitor Cells
-	A11.627.635.675 Granulocyte-Macrophage Progenitor Cells
-	A11.627.635.675.500 Granulocyte Precursor Cells
-	A11.627.635.675.750 Monocyte-Macrophage Precursor Cells
-	A11.627.635.675.750.500 U937 Cells
New Heading	<b>A11.627.817 Myeloid-Derived Suppressor Cells</b>
-	A11.642 Neoplastic Cells, Circulating
-	A11.650 Neuroglia
-	A11.650.200 Astrocytes
-	A11.650.300 Ependymogial Cells
-	A11.650.400 Microglia
-	A11.650.500 Neuropil
-	A11.650.500.550 Neuropil Threads
-	A11.650.600 Oligodendroglia
-	A11.650.600.500 Myelin Sheath
-	A11.650.775 Satellite Cells, Perineuronal
-	A11.650.800 Schwann Cells
-	A11.650.800.500 Myelin Sheath
-	A11.650.800.500.550 Neurilemma
-	A11.650.800.500.700 Ranvier's Nodes
-	A11.671 Neurons
-	A11.671.050 Adrenergic Neurons
-	A11.671.137 Axons
New Heading	<b>A11.671.137.170 Axon Initial Segment</b>
-	A11.671.137.340 Growth Cones
-	A11.671.137.560 Mossy Fibers, Hippocampal
-	A11.671.137.750 Presynaptic Terminals
-	A11.671.188 Cholinergic Neurons

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	<a href="#">A11.671.188.500</a>	<a href="#">Cholinergic Fibers</a>
New Tree	<a href="#">A11.671.188.500.060</a>	<a href="#">Autonomic Fibers, Preganglionic</a>
New Tree	<a href="#">A11.671.188.500.700</a>	<a href="#">Parasympathetic Fibers, Postganglionic</a>
-	A11.671.240	Dendrites
-	A11.671.240.169	Dendritic Spines
-	A11.671.240.340	Growth Cones
-	A11.671.270	Dopaminergic Neurons
-	A11.671.285	GABAergic Neurons
New Heading	<b>A11.671.322</b>	<b>Grid Cells</b>
-	A11.671.358	Interneurons
-	A11.671.358.050	Amacrine Cells
-	A11.671.358.212	Commissural Interneurons
-	A11.671.358.375	Renshaw Cells
-	A11.671.358.700	Retinal Bipolar Cells
-	A11.671.460	Lewy Bodies
-	A11.671.487	Mirror Neurons
-	A11.671.501	Nerve Fibers
-	A11.671.501.075	Adrenergic Fibers
-	A11.671.501.075.800	Sympathetic Fibers, Postganglionic
-	A11.671.501.100	Autonomic Fibers, Postganglionic
-	A11.671.501.100.700	Parasympathetic Fibers, Postganglionic
-	A11.671.501.100.800	Sympathetic Fibers, Postganglionic
-	A11.671.501.145	Axons
New Heading	<b>A11.671.501.145.250</b>	<b>Axon Initial Segment</b>
-	A11.671.501.145.500	Neurites
-	A11.671.501.145.750	Presynaptic Terminals
-	A11.671.501.145.750.500	Mossy Fibers, Hippocampal
-	A11.671.501.234	Cholinergic Fibers
-	A11.671.501.234.060	Autonomic Fibers, Preganglionic
-	A11.671.501.234.700	Parasympathetic Fibers, Postganglionic
-	A11.671.501.512	Nerve Fibers, Myelinated
-	A11.671.501.512.560	Myelin Sheath

## MeSH Tree Changes for 2017

Type	Tree - heading
-	A11.671.501.512.560.550                      Neurilemma
-	A11.671.501.512.560.700                      Ranvier's Nodes
-	A11.671.501.756                                  Nerve Fibers, Unmyelinated
-	A11.671.514                                      Nerve Fibers, Myelinated
-	A11.671.514.553                              Myelin Sheath
-	A11.671.514.553.640                      Ranvier's Nodes
-	A11.671.543                                      Neurites
-	A11.671.573                                      Neurofibrils
-	A11.671.573.520                              Neurofibrillary Tangles
-	A11.671.650                                      Neurons, Afferent
-	A11.671.650.250                              Hair Cells, Auditory
-	A11.671.650.250.250                      Hair Cells, Auditory, Inner
-	A11.671.650.250.315                      Hair Cells, Auditory, Outer
-	A11.671.650.395                              Hair Cells, Vestibular
-	A11.671.650.675                              Posterior Horn Cells
-	A11.671.650.675.800                      Substantia Gelatinosa
-	A11.671.650.850                              Retinal Neurons
-	A11.671.650.850.500                      Amacrine Cells
-	A11.671.650.850.625                      Photoreceptor Cells
-	A11.671.650.850.625.660                      Photoreceptor Cells, Invertebrate
-	A11.671.650.850.625.670                      Photoreceptor Cells, Vertebrate
-	A11.671.650.850.625.670.049                      Photoreceptor Connecting Cilium
-	A11.671.650.850.625.670.100                      Retinal Cone Photoreceptor Cells
-	A11.671.650.850.625.670.237                      Retinal Photoreceptor Cell Inner Segment
-	A11.671.650.850.625.670.375                      Retinal Photoreceptor Cell Outer Segment
-	A11.671.650.850.625.670.375.500                      Rod Cell Outer Segment
-	A11.671.650.850.625.670.650                      Retinal Rod Photoreceptor Cells
-	A11.671.650.850.625.670.650.650                      Rod Cell Outer Segment
-	A11.671.650.850.750                              Retinal Bipolar Cells
-	A11.671.650.850.875                              Retinal Ganglion Cells
-	A11.671.650.850.937                              Retinal Horizontal Cells
-	A11.671.650.915                              Sensory Receptor Cells
-	A11.671.650.915.500                              Chemoreceptor Cells
-	A11.671.650.915.500.530                      Neuroepithelial Cells
-	A11.671.650.915.500.530.550                      Neuroepithelial Bodies
-	A11.671.650.915.500.540                      Olfactory Receptor Neurons

## MeSH Tree Changes for 2017

Type	Tree - heading
-	A11.671.650.915.500.600 Paraganglia, Nonchromaffin
-	A11.671.650.915.500.600.050 Aortic Bodies
-	A11.671.650.915.500.600.150 Carotid Body
-	A11.671.650.915.500.600.350 Glomus Jugulare
-	A11.671.650.915.500.600.360 Glomus Tympanicum
-	A11.671.650.915.500.800 Taste Buds
-	A11.671.650.915.750 Mechanoreceptors
-	A11.671.650.915.750.300 Golgi-Mazzoni Corpuscles
-	A11.671.650.915.750.425 Merkel Cells
-	A11.671.650.915.750.500 Muscle Spindles
-	A11.671.650.915.750.600 Neuroepithelial Cells
-	A11.671.650.915.750.600.350 Hair Cells, Auditory
-	A11.671.650.915.750.600.350.350 Hair Cells, Auditory, Inner
-	A11.671.650.915.750.600.350.365 Hair Cells, Auditory, Outer
-	A11.671.650.915.750.600.675 Hair Cells, Vestibular
-	A11.671.650.915.750.700 Pacinian Corpuscles
-	A11.671.650.915.750.750 Pressoreceptors
-	A11.671.650.915.750.780 Pulmonary Stretch Receptors
-	A11.671.650.915.875 Nociceptors
-	A11.671.650.915.937 Photoreceptor Cells
-	A11.671.650.915.937.650 Photoreceptor Cells, Invertebrate
-	A11.671.650.915.937.670 Photoreceptor Cells, Vertebrate
-	A11.671.650.915.937.670.049 Photoreceptor Connecting Cilium
-	A11.671.650.915.937.670.100 Retinal Cone Photoreceptor Cells
-	A11.671.650.915.937.670.237 Retinal Photoreceptor Cell Inner Segment
-	A11.671.650.915.937.670.375 Retinal Photoreceptor Cell Outer Segment
-	A11.671.650.915.937.670.375.500 Rod Cell Outer Segment
-	A11.671.650.915.937.670.650 Retinal Rod Photoreceptor Cells
-	A11.671.650.915.937.670.650.650 Rod Cell Outer Segment
-	A11.671.650.915.968 Thermoreceptors
-	A11.671.655 Neurons, Efferent
-	A11.671.655.500 Motor Neurons
-	A11.671.655.500.050 Anterior Horn Cells
-	A11.671.655.500.525 Motor Neurons, Gamma
-	A11.671.685 Neuropil
-	A11.671.685.550 Neuropil Threads

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	A11.671.693	Nissl Bodies
-	A11.671.738	Nitregic Neurons
-	A11.671.784	Purkinje Cells
-	A11.671.790	Pyramidal Cells
New Heading	<b>A11.671.790.500</b>	<b>Place Cells</b>
-	A11.671.895	Serotonergic Neurons
-	A11.690	Oxyphil Cells
-	A11.700	Pancreatic Stellate Cells
-	A11.710	Pericytes
-	A11.733	Phagocytes
-	A11.733.397	Macrophages
-	A11.733.397.300	Epithelioid Cells
-	A11.733.397.368	Foam Cells
-	A11.733.397.376	Giant Cells, Foreign-Body
-	A11.733.397.380	Giant Cells, Langhans
-	A11.733.397.385	Histiocytes
-	A11.733.397.588	Kupffer Cells
-	A11.733.397.600	Macrophages, Alveolar
-	A11.733.397.630	Macrophages, Peritoneal
-	A11.733.397.815	RAW 264.7 Cells
-	A11.733.547	Monocytes
-	A11.733.547.500	Monocytes, Activated Killer
-	A11.733.689	Neutrophils
-	A11.750	Plant Cells
-	A11.760	Prokaryotic Cells
-	A11.789	Protoplasts
-	A11.828	Reed-Sternberg Cells
-	A11.868	Spheroplasts
-	A11.870	Spores
-	A11.870.700	Spores, Bacterial
-	A11.870.710	Spores, Fungal
-	A11.870.740	Spores, Protozoan
-	A11.870.740.600	Oocysts
-	A11.870.740.600.800	Sporozoites
-	A11.870.740.800	Schizonts



## MeSH Tree Changes for 2017

Type	Tree - heading
-	A11.870.740.800.500 Merozoites
-	A11.870.740.900 Trophozoites
-	A11.872 Stem Cells
-	A11.872.040 Adult Stem Cells
New Heading	<b>A11.872.040.250 Adult Germline Stem Cells</b>
-	A11.872.040.500 Induced Pluripotent Stem Cells
Old Tree	<b>A11.872.205 Endothelial Progenitor Cells</b>
-	A11.872.220 Fetal Stem Cells
-	A11.872.378 Hematopoietic Stem Cells
-	A11.872.378.270 Hemangioblasts
-	A11.872.378.294 Lymphoid Progenitor Cells
-	A11.872.378.294.500 Precursor Cells, B-Lymphoid
-	A11.872.378.294.750 Precursor Cells, T-Lymphoid
-	A11.872.378.294.750.800 Thymocytes
-	A11.872.378.590 Myeloid Progenitor Cells
-	A11.872.378.590.635 Granulocyte-Macrophage Progenitor Cells
-	A11.872.378.590.635.500 Granulocyte Precursor Cells
-	A11.872.378.590.635.875 Monocyte-Macrophage Precursor Cells
-	A11.872.378.590.817 Megakaryocyte-Erythroid Progenitor Cells
-	A11.872.378.590.817.250 Erythroid Precursor Cells
-	A11.872.378.590.817.625 Megakaryocyte Progenitor Cells
New Heading	<b>A11.872.378.795 Peripheral Blood Stem Cells</b>
-	A11.872.590 Multipotent Stem Cells
-	A11.872.590.500 Mesenchymal Stromal Cells
New Tree	<b>A11.872.620 Myoblasts</b>
New Tree	<b>A11.872.620.470 Myoblasts, Cardiac</b>
New Tree	<b>A11.872.620.500 Myoblasts, Skeletal</b>
New Tree	<b>A11.872.620.500.700 Satellite Cells, Skeletal Muscle</b>
New Tree	<b>A11.872.620.510 Myoblasts, Smooth Muscle</b>
Old Tree	<b>A11.872.645 Myoblasts</b>
Old Tree	<b>A11.872.645.470 Myoblasts, Cardiac</b>

## MeSH Tree Changes for 2017

Type	Tree - heading	
Old Tree	A11.872.645.500	Myoblasts, Skeletal
Old Tree	A11.872.645.500.700	Satellite Cells, Skeletal Muscle
Old Tree	A11.872.645.510	Myoblasts, Smooth Muscle
-	A11.872.650	Neoplastic Stem Cells
-	A11.872.650.500	Embryonal Carcinoma Stem Cells
-	A11.872.653	Neural Stem Cells
New Heading	<b>A11.872.677</b>	<b>Oogonial Stem Cells</b>
-	A11.872.700	Pluripotent Stem Cells
-	A11.872.700.250	Embryonic Stem Cells
-	A11.872.700.250.130	Blastomeres
-	A11.872.700.250.260	Embryoid Bodies
-	A11.872.700.250.500	Embryonal Carcinoma Stem Cells
-	A11.872.700.250.625	Embryonic Germ Cells
-	A11.872.700.250.750	Human Embryonic Stem Cells
-	A11.872.700.250.875	Mouse Embryonic Stem Cells
-	A11.872.700.500	Induced Pluripotent Stem Cells
-	A11.872.785	Side-Population Cells
-	A11.872.870	Totipotent Stem Cells
-	A11.900	Thymocytes
-	A12	Fluids and Secretions
-	A12.098	Amniotic Fluid
-	A12.200	Bodily Secretions
-	A12.200.087	Bile
-	A12.200.147	Cerumen
-	A12.200.194	Colostrum
-	A12.200.250	Dander
-	A12.200.307	Gastric Juice
-	A12.200.307.603	Gastric Acid
-	A12.200.390	Intestinal Secretions
-	A12.200.455	Milk
-	A12.200.467	Milk, Human
-	A12.200.503	Mucus
-	A12.200.503.339	Cervix Mucus
-	A12.200.567	Pancreatic Juice
-	A12.200.666	Saliva

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	A12.200.702	Sebum
-	A12.200.732	Semen
-	A12.200.769	Smegma
-	A12.200.808	Sputum
-	A12.200.849	Sweat
-	A12.200.882	Tears
-	A12.200.935	Venoms
-	A12.200.946	Vernix Caseosa
-	A12.207	Body Fluids
-	A12.207.119	Ascitic Fluid
-	A12.207.152	Blood
-	A12.207.152.200	Fetal Blood
-	A12.207.152.693	Plasma
-	A12.207.152.693.600	Platelet-Rich Plasma
-	A12.207.152.846	Serum
-	A12.207.152.846.500	Immune Sera
-	A12.207.152.846.500.203	Antilymphocyte Serum
-	A12.207.180	Body Fluid Compartments
-	A12.207.200	Body Water
-	A12.207.270	Extracellular Fluid
-	A12.207.270.040	Aqueous Humor
-	A12.207.270.210	Cerebrospinal Fluid
-	A12.207.270.300	Extravascular Lung Water
-	A12.207.270.340	Follicular Fluid
-	A12.207.270.517	Labyrinthine Fluids
-	A12.207.270.517.324	Endolymph
-	A12.207.270.517.678	Perilymph
-	A12.207.270.606	Lymph
-	A12.207.270.606.350	Chyle
-	A12.207.270.651	Pericardial Fluid
-	A12.207.270.695	Plasma
-	A12.207.270.695.600	Platelet-Rich Plasma
-	A12.207.270.847	Synovial Fluid
-	A12.207.515	Intracellular Fluid
Old Tree	A12.207.721	Nipple Aspirate Fluid
New	A12.207.824	Nipple Discharge

## MeSH Tree Changes for 2017

Type	Tree - heading	
Heading		
New Tree	<a href="#">A12.207.824.500</a>	<a href="#">Nipple Aspirate Fluid</a>
-	A12.207.927	Urine
-	A12.383	Exudates and Transudates
-	A12.383.200	Cyst Fluid
-	A12.383.250	Dentinal Fluid
-	A12.383.500	Gingival Crevicular Fluid
-	A12.383.800	Subretinal Fluid
-	A12.459	Feces
-	A12.459.529	Meconium
-	A12.459.764	Melena
-	A12.519	Gastrointestinal Contents
-	A12.580	Hyalin
-	A13	Animal Structures
-	A13.048	Air Sacs
-	A13.073	Anal Sacs
-	A13.077	Animal Fins
-	A13.079	Animal Shells
-	A13.093	Arthropod Antennae
New Tree	<a href="#">A13.093.500</a>	<a href="#">Sensilla</a>
-	A13.114	Beak
-	A13.163	Bursa of Fabricius
-	A13.223	Cloaca
-	A13.242	Comb and Wattles
-	A13.246	Compound Eye, Arthropod
-	A13.250	Corpora Allata
-	A13.265	Crop, Avian
-	A13.316	Egg Shell
-	A13.332	Electric Organ
-	A13.350	Embryo, Nonmammalian
-	A13.350.150	Chick Embryo
-	A13.350.575	Chorioallantoic Membrane
-	A13.365	Fat Body
-	A13.370	Feathers

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	A13.395	Forelimb
-	A13.395.248	Carpus, Animal
-	A13.395.823	Wings, Animal
-	A13.408	Ganglia, Invertebrate
-	A13.421	Gills
New Heading	<b>A13.433</b>	<b>Gizzard, Non-avian</b>
-	A13.445	Harderian Gland
-	A13.449	Head Kidney
-	A13.453	Hemolymph
-	A13.463	Hepatopancreas
-	A13.468	High Vocal Center
-	A13.473	Hindlimb
-	A13.473.683	Stifle
-	A13.473.821	Tarsus, Animal
-	A13.491	Hoof and Claw
-	A13.507	Horns
-	A13.507.288	Antlers
-	A13.518	Imaginal Discs
-	A13.530	Interrenal Gland
-	A13.564	Lateral Line System
-	A13.574	Malpighian Tubules
-	A13.589	Mammary Glands, Animal
-	A13.600	Metrial Gland
-	A13.641	Mushroom Bodies
-	A13.655	Nematocyst
-	A13.660	Nictitating Membrane
-	A13.686	Optic Lobe, Nonmammalian
-	A13.706	Oviducts
-	A13.706.500	Fallopian Tubes
-	A13.734	Perianal Glands
-	A13.750	Photoreceptor Cells, Invertebrate
-	A13.811	Salt Gland
-	A13.820	Scent Glands
Old Tree	<b>A13.825</b>	<b>Sensilla</b>
-	A13.853	Stomach, Avian

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	<a href="#">A13.853.355</a>	<a href="#">Gizzard</a>
New Tree	<a href="#">A13.853.355</a>	<a href="#">Gizzard, Avian</a>
Old Tree	<a href="#">A13.853.363</a>	<a href="#">Gizzard</a>
Old Tree	<a href="#">A13.853.363</a>	<a href="#">Gizzard, Avian</a>
-	A13.853.710	Proventriculus
-	A13.869	Stomach, Ruminant
-	A13.869.106	Abomasum
-	A13.869.524	Omasum
-	A13.869.697	Reticulum
-	A13.869.804	Rumen
-	A13.895	Tail
-	A13.939	Ultimobranchial Body
-	A13.950	Vibrissae
-	A13.970	Wool
-	A14	Stomatognathic System
-	A14.194	Cheek
-	A14.363	Facial Muscles
-	A14.521	Jaw
-	A14.521.125	Alveolar Process
-	A14.521.125.800	Tooth Socket
-	A14.521.320	Dental Arch
-	A14.521.632	Mandible
-	A14.521.632.300	Chin
-	A14.521.632.600	Mandibular Condyle
-	A14.521.645	Maxilla
-	A14.521.658	Palate
-	A14.521.658.660	Palate, Hard
-	A14.530	Masticatory Muscles
-	A14.530.630	Masseter Muscle
-	A14.530.790	Pterygoid Muscles
-	A14.530.940	Temporal Muscle
-	A14.549	Mouth
-	A14.549.167	Dentition
-	A14.549.167.229	Dentition, Mixed

## MeSH Tree Changes for 2017

Type	Tree - heading
-	A14.549.167.237 Dentition, Permanent
-	A14.549.167.646 Periodontium
-	A14.549.167.646.094 Alveolar Process
-	A14.549.167.646.094.800 Tooth Socket
-	A14.549.167.646.267 Dental Cementum
-	A14.549.167.646.374 Epithelial Attachment
-	A14.549.167.646.480 Gingiva
-	A14.549.167.646.700 Periapical Tissue
-	A14.549.167.646.771 Periodontal Ligament
-	A14.549.167.860 Tooth
-	A14.549.167.860.150 Bicuspid
-	A14.549.167.860.200 Cuspid
-	A14.549.167.860.425 Incisor
-	A14.549.167.860.525 Molar
-	A14.549.167.860.525.500 Molar, Third
-	A14.549.167.860.700 Tooth, Deciduous
-	A14.549.167.860.700.500 Natal Teeth
-	A14.549.167.860.715 Tooth, Unerupted
-	A14.549.167.900 Tooth Components
-	A14.549.167.900.250 Dental Cementum
-	A14.549.167.900.255 Dental Enamel
-	A14.549.167.900.255.500 Dental Pellicle
-	A14.549.167.900.260 Dental Pulp
-	A14.549.167.900.265 Dental Pulp Cavity
-	A14.549.167.900.280 Dentin
-	A14.549.167.900.280.280 Dentin, Secondary
-	A14.549.167.900.700 Tooth Cervix
-	A14.549.167.900.710 Tooth Crown
-	A14.549.167.900.720 Tooth Germ
-	A14.549.167.900.720.250 Dental Papilla
-	A14.549.167.900.720.255 Dental Sac
-	A14.549.167.900.720.265 Enamel Organ
-	A14.549.167.900.750 Tooth Root
-	A14.549.167.900.750.700 Tooth Apex
-	A14.549.336 Lip
-	A14.549.336.505 Labial Frenum

## MeSH Tree Changes for 2017

Type	Tree - heading
-	A14.549.441 Mouth Floor
-	A14.549.512 Mouth Mucosa
-	A14.549.617 Palate
-	A14.549.617.623 Palatal Muscles
-	A14.549.617.660 Palate, Hard
-	A14.549.617.780 Palate, Soft
-	A14.549.617.780.729 Uvula
-	A14.549.760 Salivary Glands
-	A14.549.760.464 Parotid Gland
-	A14.549.760.640 Salivary Ducts
-	A14.549.760.650 Salivary Glands, Minor
-	A14.549.760.687 Sublingual Gland
-	A14.549.760.812 Submandibular Gland
-	A14.549.760.906 von Ebner Glands
-	A14.549.885 Tongue
-	A14.549.885.431 Lingual Frenum
-	A14.549.885.779 Taste Buds
-	A14.724 Pharynx
-	A14.724.490 Hypopharynx
-	A14.724.557 Nasopharynx
-	A14.724.557.500 Adenoids
-	A14.724.603 Oropharynx
-	A14.724.603.925 Palatine Tonsil
-	A14.724.617 Pharyngeal Muscles
-	A14.724.617.360 Esophageal Sphincter, Upper
-	A14.724.617.680 Velopharyngeal Sphincter
-	A14.907 Temporomandibular Joint
-	A14.907.900 Temporomandibular Joint Disc
-	A15 Hemic and Immune Systems
-	A15.145 Blood
-	A15.145.229 Blood Cells
-	A15.145.229.093 Blood Buffy Coat
-	A15.145.229.188 Blood Platelets
-	A15.145.229.334 Erythrocytes
-	A15.145.229.334.270 Erythrocyte Membrane
-	A15.145.229.334.330 Erythrocytes, Abnormal



## MeSH Tree Changes for 2017

Type	Tree - heading
-	A15.145.229.334.330.100 Acanthocytes
-	A15.145.229.334.330.340 Erythrocyte Inclusions
-	A15.145.229.334.330.340.335 Heinz Bodies
-	A15.145.229.334.330.531 Megaloblasts
-	A15.145.229.334.330.802 Spherocytes
-	A15.145.229.334.760 Reticulocytes
-	A15.145.229.480 Hemocytes
-	A15.145.229.637 Leukocytes
-	A15.145.229.637.415 Granulocytes
-	A15.145.229.637.415.120 Basophils
-	A15.145.229.637.415.345 Eosinophils
-	A15.145.229.637.415.583 Neutrophils
-	A15.145.229.637.555 Leukocytes, Mononuclear
-	A15.145.229.637.555.283 Cytokine-Induced Killer Cells
-	A15.145.229.637.555.283.500 Killer Cells, Lymphokine-Activated
-	A15.145.229.637.555.283.750 Monocytes, Activated Killer
-	A15.145.229.637.555.283.875 T-Lymphocytes, Cytotoxic
-	A15.145.229.637.555.567 Lymphocytes
-	A15.145.229.637.555.567.537 Killer Cells, Natural
-	A15.145.229.637.555.567.537.500 Killer Cells, Lymphokine-Activated
-	A15.145.229.637.555.567.550 Lymphocyte Subsets
-	A15.145.229.637.555.567.550.450 B-Lymphocyte Subsets
-	A15.145.229.637.555.567.550.450.300 B-Lymphocytes, Regulatory
-	A15.145.229.637.555.567.550.500 T-Lymphocyte Subsets
-	A15.145.229.637.555.567.550.500.200 T-Lymphocytes, Cytotoxic
-	A15.145.229.637.555.567.550.500.400 T-Lymphocytes, Helper-Inducer
-	A15.145.229.637.555.567.550.500.400.500 Th1 Cells
-	A15.145.229.637.555.567.550.500.400.750 Th2 Cells
-	A15.145.229.637.555.567.550.500.400.770 Th17 Cells
-	A15.145.229.637.555.567.550.500.700 T-Lymphocytes, Regulatory
-	A15.145.229.637.555.567.562 B-Lymphocytes
-	A15.145.229.637.555.567.562.200 B-Lymphocyte Subsets
-	A15.145.229.637.555.567.562.200.300 B-Lymphocytes, Regulatory
-	A15.145.229.637.555.567.562.725 Plasma Cells
-	A15.145.229.637.555.567.569 T-Lymphocytes
-	A15.145.229.637.555.567.569.200 CD4-Positive T-Lymphocytes

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	A15.145.229.637.555.567.569.200.400	T-Lymphocytes, Helper-Inducer
-	A15.145.229.637.555.567.569.200.400.500	Th1 Cells
-	A15.145.229.637.555.567.569.200.400.750	Th2 Cells
-	A15.145.229.637.555.567.569.200.400.770	Th17 Cells
-	A15.145.229.637.555.567.569.200.700	T-Lymphocytes, Regulatory
-	A15.145.229.637.555.567.569.220	CD8-Positive T-Lymphocytes
-	A15.145.229.637.555.567.569.220.200	T-Lymphocytes, Cytotoxic
-	A15.145.229.637.555.567.569.360	Natural Killer T-Cells
-	A15.145.229.637.555.567.569.500	T-Lymphocyte Subsets
New Heading	<b>A15.145.229.637.555.567.569.500.100 Cells</b>	<b>Mucosal-Associated Invariant T</b>
-	A15.145.229.637.555.567.569.500.200	T-Lymphocytes, Cytotoxic
-	A15.145.229.637.555.567.569.500.400	T-Lymphocytes, Helper-Inducer
-	A15.145.229.637.555.567.569.500.400.500	Th1 Cells
-	A15.145.229.637.555.567.569.500.400.750	Th2 Cells
-	A15.145.229.637.555.567.569.500.400.770	Th17 Cells
-	A15.145.229.637.555.567.569.500.700	T-Lymphocytes, Regulatory
-	A15.145.229.637.555.567.584	Lymphocytes, Null
-	A15.145.229.637.555.567.650	Lymphocytes, Tumor-Infiltrating
-	A15.145.229.637.555.652	Monocytes
-	A15.145.229.637.555.652.500	Monocytes, Activated Killer
-	A15.145.300	Fetal Blood
-	A15.145.693	Plasma
-	A15.145.693.600	Platelet-Rich Plasma
-	A15.145.846	Serum
-	A15.378	Hematopoietic System
-	A15.378.316	Bone Marrow Cells
-	A15.378.316.340	Granulocytes
-	A15.378.316.340.350	Granulocyte Precursor Cells
-	A15.378.316.378	Hematopoietic Stem Cells
-	A15.378.316.378.550	Lymphoid Progenitor Cells
-	A15.378.316.378.590	Myeloid Progenitor Cells
-	A15.378.316.378.590.675	Granulocyte-Macrophage Progenitor Cells
-	A15.378.316.378.590.675.500	Granulocyte Precursor Cells
-	A15.378.316.378.590.675.750	Monocyte-Macrophage Precursor Cells
-	A15.378.316.378.590.837	Megakaryocyte-Erythroid Progenitor Cells

## MeSH Tree Changes for 2017

Type	Tree - heading
-	A15.378.316.378.590.837.250 Erythroid Precursor Cells
-	A15.378.316.378.590.837.250.200 Erythroblasts
-	A15.378.316.378.590.837.250.200.500 Megaloblasts
-	A15.378.316.378.590.837.625 Megakaryocyte Progenitor Cells
-	A15.378.316.479 Megakaryocytes
-	A15.378.316.580 Monocytes
-	A15.378.316.790 Reticulocytes
-	A15.382 Immune System
-	A15.382.032 Antibody-Producing Cells
-	A15.382.032.438 B-Lymphocytes
-	A15.382.032.438.450 B-Lymphocyte Subsets
-	A15.382.032.438.450.300 B-Lymphocytes, Regulatory
-	A15.382.032.438.725 Plasma Cells
-	A15.382.066 Antigen-Presenting Cells
-	A15.382.066.270 Dendritic Cells
-	A15.382.066.270.500 Langerhans Cells
-	A15.382.066.275 Dendritic Cells, Follicular
-	A15.382.216 Bone Marrow
-	A15.382.250 Enterochromaffin Cells
-	A15.382.370 Immunological Synapses
-	A15.382.490 Leukocytes
-	A15.382.490.315 Granulocytes
-	A15.382.490.315.120 Basophils
-	A15.382.490.315.251 Eosinophils
-	A15.382.490.315.583 Neutrophils
-	A15.382.490.555 Leukocytes, Mononuclear
-	A15.382.490.555.283 Cytokine-Induced Killer Cells
-	A15.382.490.555.283.500 Killer Cells, Lymphokine-Activated
-	A15.382.490.555.283.750 Monocytes, Activated Killer
-	A15.382.490.555.283.875 T-Lymphocytes, Cytotoxic
-	A15.382.490.555.567 Lymphocytes
-	A15.382.490.555.567.537 Killer Cells, Natural
-	A15.382.490.555.567.537.500 Killer Cells, Lymphokine-Activated
-	A15.382.490.555.567.550 Lymphocyte Subsets
-	A15.382.490.555.567.550.300 B-Lymphocyte Subsets
-	A15.382.490.555.567.550.500 T-Lymphocyte Subsets

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	A15.382.490.555.567.550.500.200	T-Lymphocytes, Cytotoxic
-	A15.382.490.555.567.550.500.400	T-Lymphocytes, Helper-Inducer
-	A15.382.490.555.567.550.500.400.900	Th1 Cells
-	A15.382.490.555.567.550.500.400.905	Th2 Cells
-	A15.382.490.555.567.550.500.400.915	Th17 Cells
-	A15.382.490.555.567.550.500.700	T-Lymphocytes, Regulatory
-	A15.382.490.555.567.562	B-Lymphocytes
-	A15.382.490.555.567.562.450	B-Lymphocyte Subsets
-	A15.382.490.555.567.562.450.300	B-Lymphocytes, Regulatory
-	A15.382.490.555.567.562.725	Plasma Cells
-	A15.382.490.555.567.569	T-Lymphocytes
-	A15.382.490.555.567.569.200	CD4-Positive T-Lymphocytes
-	A15.382.490.555.567.569.200.400	T-Lymphocytes, Helper-Inducer
-	A15.382.490.555.567.569.200.400.900	Th1 Cells
-	A15.382.490.555.567.569.200.400.905	Th2 Cells
-	A15.382.490.555.567.569.200.400.915	Th17 Cells
-	A15.382.490.555.567.569.200.700	T-Lymphocytes, Regulatory
-	A15.382.490.555.567.569.220	CD8-Positive T-Lymphocytes
-	A15.382.490.555.567.569.220.200	T-Lymphocytes, Cytotoxic
-	A15.382.490.555.567.569.440	Jurkat Cells
-	A15.382.490.555.567.569.470	Natural Killer T-Cells
-	A15.382.490.555.567.569.500	T-Lymphocyte Subsets
New Heading	<b>A15.382.490.555.567.569.500.100</b>	<b>Mucosal-Associated Invariant T Cells</b>
-	A15.382.490.555.567.569.500.200	T-Lymphocytes, Cytotoxic
-	A15.382.490.555.567.569.500.400	T-Lymphocytes, Helper-Inducer
-	A15.382.490.555.567.569.500.400.900	Th1 Cells
-	A15.382.490.555.567.569.500.400.905	Th2 Cells
-	A15.382.490.555.567.569.500.400.915	Th17 Cells
-	A15.382.490.555.567.569.500.700	T-Lymphocytes, Regulatory
-	A15.382.490.555.567.622	Lymphocytes, Null
-	A15.382.490.555.567.650	Lymphocytes, Tumor-Infiltrating
-	A15.382.490.555.652	Monocytes
-	A15.382.490.555.652.500	Monocytes, Activated Killer
-	A15.382.520	Lymphatic System
-	A15.382.520.150	Lymph

## MeSH Tree Changes for 2017

Type	Tree - heading
-	A15.382.520.150.350 Chyle
-	A15.382.520.301 Lymphatic Vessels
-	A15.382.520.301.500 Endothelium, Lymphatic
-	A15.382.520.301.750 Thoracic Duct
-	A15.382.520.604 Lymphoid Tissue
-	A15.382.520.604.100 Adenoids
-	A15.382.520.604.114 Bursa of Fabricius
-	A15.382.520.604.412 Lymph Nodes
-	A15.382.520.604.412.500 Germinal Center
-	A15.382.520.604.412.500.200 Dendritic Cells, Follicular
New Heading	<b>A15.382.520.604.412.750</b> <b>Sentinel Lymph Node</b>
-	A15.382.520.604.580 Palatine Tonsil
-	A15.382.520.604.600 Peyer's Patches
-	A15.382.520.604.700 Spleen
New Heading	<b>A15.382.520.604.725</b> <b>Tertiary Lymphoid Structures</b>
-	A15.382.520.604.750 Thymus Gland
-	A15.382.652 Mast Cells
-	A15.382.680 Phagocytes
-	A15.382.680.397 Macrophages
-	A15.382.680.397.300 Epithelioid Cells
-	A15.382.680.397.368 Foam Cells
-	A15.382.680.397.376 Giant Cells, Foreign-Body
-	A15.382.680.397.380 Giant Cells, Langhans
-	A15.382.680.397.385 Histiocytes
-	A15.382.680.397.588 Kupffer Cells
-	A15.382.680.397.600 Macrophages, Alveolar
-	A15.382.680.397.630 Macrophages, Peritoneal
-	A15.382.680.547 Monocytes
-	A15.382.680.547.500 Monocytes, Activated Killer
-	A15.382.680.689 Neutrophils
-	A15.382.812 Mononuclear Phagocyte System
-	A15.382.812.260 Dendritic Cells
-	A15.382.812.260.500 Langerhans Cells
-	A15.382.812.522 Macrophages

## MeSH Tree Changes for 2017

Type	Tree - heading
-	A15.382.812.522.300 Epithelioid Cells
-	A15.382.812.522.368 Foam Cells
-	A15.382.812.522.376 Giant Cells, Foreign-Body
-	A15.382.812.522.380 Giant Cells, Langhans
-	A15.382.812.522.385 Histiocytes
-	A15.382.812.522.588 Kupffer Cells
-	A15.382.812.522.600 Macrophages, Alveolar
-	A15.382.812.522.630 Macrophages, Peritoneal
-	A15.382.812.547 Monocytes
-	A15.382.812.547.500 Monocytes, Activated Killer
-	A16 Embryonic Structures
-	A16.094 Blastomeres
-	A16.142 Branchial Region
-	A16.166 Cleavage Stage, Ovum
-	A16.178 Cloaca
-	A16.254 Embryo, Mammalian
-	A16.254.500 Blastocyst
-	A16.254.500.533 Blastocyst Inner Cell Mass
-	A16.254.500.766 Trophoblasts
-	A16.254.750 Extraembryonic Membranes
-	A16.254.750.147 Allantois
-	A16.254.750.277 Amnion
-	A16.254.750.473 Chorion
-	A16.254.750.473.200 Chorionic Villi
-	A16.254.750.981 Yolk Sac
-	A16.331 Embryo, Nonmammalian
-	A16.331.024 Blastoderm
-	A16.331.042 Blastodisc
-	A16.331.099 Blastula
-	A16.331.200 Chick Embryo
-	A16.331.400 Chorioallantoic Membrane
-	A16.331.800 Yolk Sac
-	A16.378 Fetus
-	A16.378.099 Aborted Fetus
-	A16.378.149 Amniotic Fluid
-	A16.378.200 Fetal Blood

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	A16.378.303	Fetal Heart
-	A16.378.303.395	Ductus Arteriosus
-	A16.378.303.662	Endocardial Cushions
-	A16.378.303.930	Truncus Arteriosus
-	A16.378.529	Meconium
-	A16.378.693	Umbilical Cord
-	A16.378.693.641	Umbilical Arteries
-	A16.378.693.807	Umbilical Veins
-	A16.378.857	Vernix Caseosa
-	A16.441	Gastrula
-	A16.504	Germ Layers
-	A16.504.273	Ectoderm
-	A16.504.407	Endoderm
-	A16.504.660	Mesoderm
-	A16.504.660.600	Pericytes
-	A16.504.660.750	Somites
-	A16.535	Gestational Sac
New Heading	<b>A16.551</b>	<b>Gubernaculum</b>
-	A16.567	Limb Buds
-	A16.599	Mesonephros
-	A16.615	Morula
-	A16.623	Mullerian Ducts
-	A16.627	Neural Crest
-	A16.629	Neural Plate
-	A16.630	Neural Tube
-	A16.660	Notochord
-	A16.675	Organizers, Embryonic
-	A16.690	Ovum
-	A16.690.325	Egg Yolk
-	A16.690.886	Vitelline Membrane
-	A16.690.900	Zona Pellucida
-	A16.710	Placenta
-	A16.710.189	Chorionic Villi
-	A16.710.289	Decidua
-	A16.710.289.500	Deciduoma

## MeSH Tree Changes for 2017

Type	Tree - heading
-	A16.710.802 Trophoblasts
-	A16.830 Primitive Streak
-	A16.835 Pronephros
-	A16.835.400 Head Kidney
-	A16.890 Urachus
-	A16.920 Vitelline Duct
-	A16.935 Wolffian Ducts
-	A16.950 Zygote
-	A17 Integumentary System
-	A17.360 Hair
-	A17.360.296 Eyebrows
-	A17.360.421 Eyelashes
-	A17.360.710 Hair Follicle
-	A17.360.855 Wool
-	A17.600 Nails
-	A17.815 Skin
-	A17.815.180 Dermis
-	A17.815.180.040 Acellular Dermis
-	A17.815.250 Epidermis
-	A17.815.250.500 Hair Follicle
-	A17.815.805 Sebaceous Glands
-	A17.815.830 Sweat Glands
-	A17.815.830.206 Apocrine Glands
-	A17.815.830.480 Eccrine Glands
-	A18 Plant Structures
-	A18.005 Chromosomes, Plant
-	A18.024 Plant Components, Aerial
-	A18.024.249 Flowering Tops
-	A18.024.249.500 Flowers
-	A18.024.249.500.249 Germ Cells, Plant
-	A18.024.249.500.249.249 Ovule
-	A18.024.249.500.249.500 Pollen
-	A18.024.249.500.249.500.500 Pollen Tube
-	A18.024.249.500.624 Inflorescence
-	A18.024.500 Fruit
-	A18.024.500.500 Nuts



## MeSH Tree Changes for 2017

Type	Tree - heading
-	A18.024.500.750                      Seeds
-	A18.024.500.750.333                      Cotyledon
-	A18.024.500.750.500                      Edible Grain
-	A18.024.500.750.666                      Endosperm
-	A18.024.750                      Plant Epidermis
-	A18.024.750.200                      Plant Bark
-	A18.024.750.650                      Plant Stomata
-	A18.024.750.825                      Trichomes
-	A18.024.812                      Plant Leaves
-	A18.024.812.324                      Mesophyll Cells
-	A18.024.812.650                      Plant Stomata
-	A18.024.812.700                      Pulvinus
-	A18.024.875                      Plant Shoots
-	A18.024.875.500                      Cotyledon
-	A18.024.875.750                      Hypocotyl
-	A18.024.875.875                      Meristem
-	A18.024.875.875.500                      Cambium
-	A18.024.937                      Plant Stems
-	A18.024.937.249                      Hypocotyl
-	A18.024.937.500                      Meristem
-	A18.024.937.750                      Rhizome
-	A18.400                      Plant Roots
-	A18.400.500                      Meristem
-	A18.400.500.299                      Cambium
-	A18.400.500.600                      Plant Root Cap
-	A18.400.525                      Mycorrhizae
-	A18.400.625                      Plant Tubers
-	A18.400.750                      Rhizome
-	A18.400.875                      Root Nodules, Plant
-	A18.450                      Plant Vascular Bundle
-	A18.450.250                      Phloem
-	A18.450.500                      Xylem
-	A18.450.500.500                      Wood
-	A18.550                      Seedlings
-	A18.550.500                      Hypocotyl
-	A18.560                      Sporangia

## MeSH Tree Changes for 2017

Type	Tree - heading
-	A19 Fungal Structures
-	A19.311 Chromosomes, Fungal
-	A19.311.800 Chromosomes, Artificial, Yeast
-	A19.374 Fruiting Bodies, Fungal
-	A19.374.500 Spores, Fungal
-	A19.530 Fungal Capsules
-	A19.687 Mycelium
-	A19.687.400 Hyphae
-	A19.690 Mycorrhizae
-	A19.800 Sporangia
-	A20 Bacterial Structures
-	A20.186 Bacterial Capsules
-	A20.374 Bacterial Chromatophores
-	A20.812 Chromosomes, Bacterial
-	A20.812.170 Chromosomes, Artificial, Bacterial
-	A20.843 Fimbriae, Bacterial
-	A20.859 Magnetosomes
-	A20.867 Pili, Sex
-	A20.871 Purple Membrane
-	A20.873 Ribosome Subunits, Large, Bacterial
-	A20.874 Ribosome Subunits, Small, Bacterial
-	A21 Viral Structures
-	A21.249 Virion
-	A21.249.500 Nucleocapsid
-	A21.249.500.250 Capsid
-	B01 Eukaryota
-	B01.043 Alveolata
-	B01.043.075 Apicomplexa
-	B01.043.075.189 Coccidia
-	B01.043.075.189.250 Eimeriida
-	B01.043.075.189.250.150 Cryptosporidiidae
-	B01.043.075.189.250.150.160 Cryptosporidium
-	B01.043.075.189.250.150.160.170 Cryptosporidium parvum
-	B01.043.075.189.250.250 Eimeriidae
-	B01.043.075.189.250.250.200 Cyclospora
-	B01.043.075.189.250.250.250 Eimeria

## MeSH Tree Changes for 2017

Type	Tree - heading
-	B01.043.075.189.250.250.250.250 Eimeria tenella
-	B01.043.075.189.250.250.400 Isospora
-	B01.043.075.189.250.750 Sarcocystidae
-	B01.043.075.189.250.750.550 Neospora
-	B01.043.075.189.250.750.750 Sarcocystis
-	B01.043.075.189.250.750.800 Toxoplasma
-	B01.043.075.189.275 Eucoccidiida
-	B01.043.075.380 Haemosporida
-	B01.043.075.380.611 Plasmodium
-	B01.043.075.380.611.461 Plasmodium berghei
-	B01.043.075.380.611.490 Plasmodium chabaudi
-	B01.043.075.380.611.505 Plasmodium cynomolgi
-	B01.043.075.380.611.561 Plasmodium falciparum
-	B01.043.075.380.611.576 Plasmodium gallinaceum
-	B01.043.075.380.611.610 Plasmodium knowlesi
-	B01.043.075.380.611.661 Plasmodium malariae
-	B01.043.075.380.611.700 Plasmodium ovale
-	B01.043.075.380.611.761 Plasmodium vivax
-	B01.043.075.380.611.780 Plasmodium yoelii
-	B01.043.075.600 Piroplasmia
-	B01.043.075.600.580 Piroplasmida
-	B01.043.075.600.580.070 Babesia
-	B01.043.075.600.580.070.100 Babesia bovis
-	B01.043.075.600.580.070.550 Babesia microti
-	B01.043.075.600.580.700 Theileria
-	B01.043.075.600.580.700.100 Theileria annulata
-	B01.043.075.600.580.700.500 Theileria parva
-	B01.043.185 Ciliophora
-	B01.043.185.375 Kinetofragminophorea
-	B01.043.185.375.650 Trichostomatida
-	B01.043.185.375.650.650 Trichostomatina
-	B01.043.185.375.650.650.153 Balantidium
-	B01.043.185.650 Oligohymenophorea
-	B01.043.185.650.375 Hymenostomatida
-	B01.043.185.650.375.550 Peniculina
-	B01.043.185.650.375.550.637 Paramecium

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	B01.043.185.650.375.550.637.074	Paramecium aurelia
-	B01.043.185.650.375.550.637.150	Paramecium caudatum
-	B01.043.185.650.375.550.637.700	Paramecium tetraurelia
-	B01.043.185.650.375.750	Tetrahymenina
-	B01.043.185.650.375.750.850	Tetrahymena
-	B01.043.185.650.375.750.850.700	Tetrahymena pyriformis
-	B01.043.185.650.375.750.850.875	Tetrahymena thermophila
-	B01.043.185.700	Polymenophorea
-	B01.043.185.700.400	Hypotrichida
-	B01.043.185.700.400.750	Sporadotrichina
-	B01.043.185.700.400.750.375	Euplotes
-	B01.043.185.700.400.750.600	Oxytricha
-	B01.043.214	Dinoflagellida
-	B01.043.214.600	Pfiesteria piscicida
-	B01.046	Amoebozoa
-	B01.046.060	Archamoebae
-	B01.046.090	Balamuthia mandrillaris
-	B01.046.500	Lobosea
-	B01.046.500.100	Amoebida
-	B01.046.500.100.075	Acanthopodina
-	B01.046.500.100.075.080	Acanthamoeba
-	B01.046.500.100.075.080.150	Acanthamoeba castellanii
-	B01.046.500.100.200	Blastocystina
-	B01.046.500.100.200.200	Blastocystis
-	B01.046.500.100.200.200.375	Blastocystis hominis
-	B01.046.500.100.700	Tubulina
-	B01.046.500.100.700.089	Amoeba
New Heading	<b>B01.046.500.100.700.089.500</b>	<b>Endamoeba histolytica</b>
-	B01.046.500.100.700.325	Endolimax
-	B01.046.500.100.700.335	Entamoeba
-	B01.046.500.100.700.335.330	Entamoeba histolytica
-	B01.046.500.100.700.430	Hartmannella
-	B01.046.500.700	Schizopyrenida
-	B01.046.500.700.710	Naegleria
-	B01.046.500.700.710.375	Naegleria fowleri

## MeSH Tree Changes for 2017

Type	Tree - heading
-	B01.046.550 Mycetozoa
-	B01.046.550.200 Dictyosteliida
-	B01.046.550.200.300 Dictyostelium
-	B01.046.550.550 Myxomycetes
-	B01.046.550.550.600 Physarida
-	B01.046.550.550.600.700 Physarum
-	B01.046.550.550.600.700.550 Physarum polycephalum
-	B01.050 Animals
-	B01.050.050 Animal Population Groups
-	B01.050.050.116 Animals, Domestic
-	B01.050.050.116.500 Livestock
-	B01.050.050.116.600 Pets
-	B01.050.050.116.625 Poultry
-	B01.050.050.126 Animals, Exotic
-	B01.050.050.136 Animals, Genetically Modified
-	B01.050.050.136.500 Mice, Transgenic
-	B01.050.050.136.500.500 Mice, Knockout
-	B01.050.050.136.700 Rats, Transgenic
-	B01.050.050.199 Animals, Laboratory
-	B01.050.050.199.520 Animals, Inbred Strains
-	B01.050.050.199.520.040 Animals, Congenic
-	B01.050.050.199.520.040.500 Mice, Congenic
-	B01.050.050.199.520.520 Mice, Inbred Strains
-	B01.050.050.199.520.520.200 Mice, Hairless
-	B01.050.050.199.520.520.300 Mice, Inbred A
-	B01.050.050.199.520.520.318 Mice, Inbred AKR
-	B01.050.050.199.520.520.338 Mice, Inbred BALB C
-	B01.050.050.199.520.520.388 Mice, Inbred C3H
-	B01.050.050.199.520.520.420 Mice, Inbred C57BL
-	B01.050.050.199.520.520.420.500 Mice, Inbred mdx
-	B01.050.050.199.520.520.440 Mice, Inbred CBA
-	B01.050.050.199.520.520.445 Mice, Inbred CFTR
-	B01.050.050.199.520.520.500 Mice, Inbred DBA
-	B01.050.050.199.520.520.510 Mice, Inbred ICR
-	B01.050.050.199.520.520.555 Mice, Inbred MRL lpr
-	B01.050.050.199.520.520.565 Mice, Inbred NOD

## MeSH Tree Changes for 2017

Type	Tree - heading
-	B01.050.050.199.520.520.580 Mice, Inbred NZB
-	B01.050.050.199.520.520.600 Mice, Inbred SENCAR
-	B01.050.050.199.520.760 Rats, Inbred Strains
-	B01.050.050.199.520.760.080 Rats, Inbred ACI
-	B01.050.050.199.520.760.090 Rats, Inbred BB
-	B01.050.050.199.520.760.110 Rats, Inbred BN
-	B01.050.050.199.520.760.130 Rats, Inbred BUF
-	B01.050.050.199.520.760.165 Rats, Inbred Dahl
-	B01.050.050.199.520.760.200 Rats, Inbred F344
-	B01.050.050.199.520.760.275 Rats, Inbred LEC
-	B01.050.050.199.520.760.280 Rats, Inbred Lew
-	B01.050.050.199.520.760.290 Rats, Inbred OLETF
-	B01.050.050.199.520.760.300 Rats, Inbred SHR
-	B01.050.050.199.520.760.360 Rats, Inbred WF
-	B01.050.050.199.520.760.390 Rats, Inbred WKY
-	B01.050.050.282 Animals, Newborn
-	B01.050.050.284 Animals, Outbred Strains
-	B01.050.050.286 Animals, Poisonous
-	B01.050.050.286.500 Fishes, Poisonous
-	B01.050.050.293 Animals, Suckling
-	B01.050.050.300 Animals, Wild
-	B01.050.050.448 Animals, Zoo
-	B01.050.050.565 Endangered Species
-	B01.050.050.580 Introduced Species
-	B01.050.150 Chordata
-	B01.050.150.200 Chordata, Nonvertebrate
-	B01.050.150.200.199 Cephalochordata
-	B01.050.150.200.199.500 Lancelets
-	B01.050.150.200.400 Hyperotreti
-	B01.050.150.200.400.380 Hagfishes
-	B01.050.150.200.727 Urochordata
-	B01.050.150.200.727.300 Ciona intestinalis
-	B01.050.150.900 Vertebrates
-	B01.050.150.900.090 Amphibians
-	B01.050.150.900.090.180 Anura
-	B01.050.150.900.090.180.210 Bufonidae

## MeSH Tree Changes for 2017

Type	Tree - heading
-	B01.050.150.900.090.180.210.080 Bufo arenarum
-	B01.050.150.900.090.180.210.108 Bufo bufo
-	B01.050.150.900.090.180.210.580 Bufo marinus
-	B01.050.150.900.090.180.610 Pipidae
-	B01.050.150.900.090.180.610.500 Xenopus
-	B01.050.150.900.090.180.610.500.562 Xenopus laevis
-	B01.050.150.900.090.180.708 Ranidae
-	B01.050.150.900.090.180.708.180 Rana catesbeiana
-	B01.050.150.900.090.180.708.210 Rana clamitans
-	B01.050.150.900.090.180.708.240 Rana esculenta
-	B01.050.150.900.090.180.708.310 Rana pipiens
-	B01.050.150.900.090.180.708.360 Rana ridibunda
-	B01.050.150.900.090.180.708.420 Rana temporaria
-	B01.050.150.900.090.608 Urodela
-	B01.050.150.900.090.608.080 Ambystomatidae
-	B01.050.150.900.090.608.080.068 Ambystoma
-	B01.050.150.900.090.608.080.068.525 Ambystoma mexicanum
-	B01.050.150.900.090.608.630 Proteidae
-	B01.050.150.900.090.608.630.510 Necturus
-	B01.050.150.900.090.608.630.510.508 Necturus maculosus
-	B01.050.150.900.090.608.700 Salamandridae
-	B01.050.150.900.090.608.700.500 Notophthalmus
-	B01.050.150.900.090.608.700.500.480 Notophthalmus viridescens
-	B01.050.150.900.090.608.700.540 Pleurodeles
-	B01.050.150.900.090.608.700.610 Salamandra
-	B01.050.150.900.090.608.700.670 Triturus
-	B01.050.150.900.248 Birds
-	B01.050.150.900.248.050 Anseriformes
-	B01.050.150.900.248.050.200 Ducks
-	B01.050.150.900.248.050.350 Geese
-	B01.050.150.900.248.150 Charadriiformes
-	B01.050.150.900.248.165 Columbiformes
-	B01.050.150.900.248.165.150 Columbidae
-	B01.050.150.900.248.350 Galliformes
-	B01.050.150.900.248.350.150 Chickens
-	B01.050.150.900.248.350.650 Quail

## MeSH Tree Changes for 2017

Type	Tree - heading
-	B01.050.150.900.248.350.650.300 Colinus
-	B01.050.150.900.248.350.650.350 Coturnix
-	B01.050.150.900.248.350.800 Turkeys
-	B01.050.150.900.248.600 Palaeognathae
-	B01.050.150.900.248.600.200 Dromaiidae
-	B01.050.150.900.248.600.750 Rheiformes
-	B01.050.150.900.248.600.800 Struthioniformes
-	B01.050.150.900.248.617 Poultry
-	B01.050.150.900.248.617.192 Chickens
-	B01.050.150.900.248.617.345 Ducks
-	B01.050.150.900.248.617.492 Geese
-	B01.050.150.900.248.617.800 Turkeys
-	B01.050.150.900.248.620 Passeriformes
-	B01.050.150.900.248.620.750 Songbirds
-	B01.050.150.900.248.620.750.150 Crows
-	B01.050.150.900.248.620.750.250 Finches
-	B01.050.150.900.248.620.750.250.150 Canaries
-	B01.050.150.900.248.620.750.700 Sparrows
-	B01.050.150.900.248.620.750.712 Starlings
-	B01.050.150.900.248.620.750.725 Swallows
-	B01.050.150.900.248.710 Psittaciformes
-	B01.050.150.900.248.710.251 Cockatoos
-	B01.050.150.900.248.710.672 Parrots
-	B01.050.150.900.248.710.672.040 Agapornis
-	B01.050.150.900.248.710.672.050 Amazona
-	B01.050.150.900.248.710.672.650 Parakeets
-	B01.050.150.900.248.710.672.650.500 Melopsittacus
-	B01.050.150.900.248.710.672.650.650 Psittacula
-	B01.050.150.900.248.815 Raptors
-	B01.050.150.900.248.815.350 Falconiformes
-	B01.050.150.900.248.815.350.200 Eagles
-	B01.050.150.900.248.815.350.400 Hawks
-	B01.050.150.900.248.815.550 Strigiformes
-	B01.050.150.900.248.860 Spheniscidae
-	B01.050.150.900.493 Fishes
-	B01.050.150.900.493.039 Batrachoidiformes



## MeSH Tree Changes for 2017

Type	Tree - heading
-	B01.050.150.900.493.080                      Catfishes
-	B01.050.150.900.493.080.148                      Ictaluridae
-	B01.050.150.900.493.130                      Characiformes
-	B01.050.150.900.493.130.150                      Characidae
-	B01.050.150.900.493.200                      Cypriniformes
-	B01.050.150.900.493.200.244                      Cyprinidae
-	B01.050.150.900.493.200.244.248                      Carps
-	B01.050.150.900.493.200.244.248.480                      Goldfish
-	B01.050.150.900.493.200.244.828                      Zebrafish
-	B01.050.150.900.493.338                      Eels
-	B01.050.150.900.493.338.282                      Anguilla
-	B01.050.150.900.493.370                      Elasmobranchii
-	B01.050.150.900.493.370.853                      Sharks
-	B01.050.150.900.493.370.853.392                      Dogfish
-	B01.050.150.900.493.370.853.392.800                      Squalus
-	B01.050.150.900.493.370.853.392.800.500                      Squalus acanthias
-	B01.050.150.900.493.370.870                      Skates (Fish)
-	B01.050.150.900.493.370.935                      Torpedo
-	B01.050.150.900.493.378                      Electric Fish
-	B01.050.150.900.493.378.430                      Gymnotiformes
-	B01.050.150.900.493.378.430.250                      Electrophorus
-	B01.050.150.900.493.378.682                      Skates (Fish)
-	B01.050.150.900.493.378.722                      Torpedo
-	B01.050.150.900.493.385                      Esociformes
-	B01.050.150.900.493.385.300                      Esocidae
-	B01.050.150.900.493.385.850                      Umbridae
-	B01.050.150.900.493.392                      Fishes, Poisonous
-	B01.050.150.900.493.418                      Flatfishes
-	B01.050.150.900.493.418.438                      Flounder
-	B01.050.150.900.493.467                      Gadiformes
-	B01.050.150.900.493.467.400                      Gadus morhua
-	B01.050.150.900.493.516                      Hagfishes
-	B01.050.150.900.493.559                      Lampreys
-	B01.050.150.900.493.559.600                      Petromyzon
-	B01.050.150.900.493.595                      Osmeriformes
-	B01.050.150.900.493.602                      Perciformes

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	B01.050.150.900.493.602.105	Bass
-	B01.050.150.900.493.602.200	Cichlids
-	B01.050.150.900.493.602.200.800	Tilapia
-	B01.050.150.900.493.602.600	Perches
-	B01.050.150.900.493.602.750	Sea Bream
-	B01.050.150.900.493.602.825	Tuna
-	B01.050.150.900.493.817	Salmoniformes
-	B01.050.150.900.493.817.750	Salmonidae
-	B01.050.150.900.493.817.750.705	Salmon
-	B01.050.150.900.493.817.750.705.580	Oncorhynchus
-	B01.050.150.900.493.817.750.705.580.400	Oncorhynchus keta
-	B01.050.150.900.493.817.750.705.580.410	Oncorhynchus kisutch
-	B01.050.150.900.493.817.750.705.790	Salmo salar
-	B01.050.150.900.493.817.750.825	Trout
-	B01.050.150.900.493.817.750.825.580	Oncorhynchus
-	B01.050.150.900.493.817.750.825.580.600	Oncorhynchus mykiss
-	B01.050.150.900.493.850	Smegmamorpha
-	B01.050.150.900.493.850.139	Beloniformes
-	B01.050.150.900.493.850.139.650	Oryzias
-	B01.050.150.900.493.850.280	Cyprinodontiformes
-	B01.050.150.900.493.850.280.214	Fundulidae
-	B01.050.150.900.493.850.280.500	Killifishes
-	B01.050.150.900.493.850.280.680	Poecilia
-	B01.050.150.900.493.875	Tetraodontiformes
-	B01.050.150.900.493.875.800	Takifugu
-	B01.050.150.900.649	Mammals
-	B01.050.150.900.649.077	Artiodactyla
New Heading	<b>B01.050.150.900.649.077.190</b>	<b>Camelidae</b>
New Tree	<a href="#">B01.050.150.900.649.077.190.090</a>	<a href="#">Camelids, New World</a>
New Tree	<a href="#">B01.050.150.900.649.077.190.180</a>	<a href="#">Camels</a>
New Tree	<a href="#">B01.050.150.900.649.077.190.180</a>	<a href="#">Camelus</a>
-	B01.050.150.900.649.077.380	Ruminants
-	B01.050.150.900.649.077.380.100	Antelopes

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	B01.050.150.900.649.077.380.120	Bison
-	B01.050.150.900.649.077.380.135	Buffaloes
Old Tree	<b>B01.050.150.900.649.077.380.175</b>	<b>Camelids, New World</b>
Old Tree	<b>B01.050.150.900.649.077.380.201</b>	<b>Camels</b>
Old Tree	<b>B01.050.150.900.649.077.380.201</b>	<b>Camelus</b>
-	B01.050.150.900.649.077.380.271	Cattle
-	B01.050.150.900.649.077.380.373	Deer
-	B01.050.150.900.649.077.380.373.322	Muntjacs
-	B01.050.150.900.649.077.380.373.644	Reindeer
-	B01.050.150.900.649.077.380.443	Giraffes
-	B01.050.150.900.649.077.380.513	Goats
-	B01.050.150.900.649.077.380.513.500	Rupicapra
-	B01.050.150.900.649.077.380.791	Sheep
-	B01.050.150.900.649.077.380.791.100	Sheep, Bighorn
-	B01.050.150.900.649.077.380.791.150	Sheep, Domestic
-	B01.050.150.900.649.077.880	Swine
-	B01.050.150.900.649.077.880.399	Sus scrofa
-	B01.050.150.900.649.077.880.399.800	Swine, Miniature
-	B01.050.150.900.649.147	Carnivora
-	B01.050.150.900.649.147.050	Ailuridae
-	B01.050.150.900.649.147.153	Canidae
-	B01.050.150.900.649.147.153.150	Coyotes
-	B01.050.150.900.649.147.153.200	Dogs
-	B01.050.150.900.649.147.153.250	Foxes
-	B01.050.150.900.649.147.153.400	Jackals
-	B01.050.150.900.649.147.153.700	Raccoon Dogs
-	B01.050.150.900.649.147.153.900	Wolves
-	B01.050.150.900.649.147.320	Eupleridae
-	B01.050.150.900.649.147.354	Felidae
-	B01.050.150.900.649.147.354.030	Acinonyx
-	B01.050.150.900.649.147.354.250	Felis
-	B01.050.150.900.649.147.354.250.125	Cats
-	B01.050.150.900.649.147.354.500	Lynx
-	B01.050.150.900.649.147.354.600	Panthera
-	B01.050.150.900.649.147.354.600.500	Lions
-	B01.050.150.900.649.147.354.600.800	Tigers

## MeSH Tree Changes for 2017

Type	Tree - heading
-	B01.050.150.900.649.147.354.630 Puma
-	B01.050.150.900.649.147.400 Herpestidae
-	B01.050.150.900.649.147.410 Hyaenidae
-	B01.050.150.900.649.147.500 Mephitidae
-	B01.050.150.900.649.147.550 Mustelidae
-	B01.050.150.900.649.147.550.350 Ferrets
-	B01.050.150.900.649.147.550.500 Mink
-	B01.050.150.900.649.147.550.600 Otters
-	B01.050.150.900.649.147.575 Nandiniidae
-	B01.050.150.900.649.147.600 Pinnipedia
-	B01.050.150.900.649.147.600.288 Fur Seals
-	B01.050.150.900.649.147.600.670 Sea Lions
-	B01.050.150.900.649.147.600.700 Seals, Earless
-	B01.050.150.900.649.147.600.700.600 Phoca
-	B01.050.150.900.649.147.600.821 Walruses
-	B01.050.150.900.649.147.650 Procyonidae
-	B01.050.150.900.649.147.650.700 Raccoons
-	B01.050.150.900.649.147.890 Ursidae
-	B01.050.150.900.649.147.900 Viverridae
-	B01.050.150.900.649.189 Cetacea
-	B01.050.150.900.649.189.267 Dolphins
-	B01.050.150.900.649.189.267.100 Bottle-Nosed Dolphin
-	B01.050.150.900.649.189.267.150 Common Dolphins
-	B01.050.150.900.649.189.267.750 Stenella
-	B01.050.150.900.649.189.267.900 Whale, Killer
-	B01.050.150.900.649.189.267.920 Whales, Pilot
-	B01.050.150.900.649.189.566 Porpoises
-	B01.050.150.900.649.189.566.600 Phocoena
-	B01.050.150.900.649.189.865 Whales
-	B01.050.150.900.649.189.865.100 Balaenoptera
-	B01.050.150.900.649.189.865.100.350 Fin Whale
-	B01.050.150.900.649.189.865.100.500 Minke Whale
-	B01.050.150.900.649.189.865.110 Beluga Whale
-	B01.050.150.900.649.189.865.130 Bowhead Whale
-	B01.050.150.900.649.189.865.400 Humpback Whale
-	B01.050.150.900.649.189.865.750 Sperm Whale

## MeSH Tree Changes for 2017

Type	Tree - heading
-	B01.050.150.900.649.225 Chiroptera
-	B01.050.150.900.649.400 Hyraxes
-	B01.050.150.900.649.473 Insectivora
-	B01.050.150.900.649.473.376 Hedgehogs
-	B01.050.150.900.649.473.450 Moles
-	B01.050.150.900.649.473.751 Shrews
-	B01.050.150.900.649.521 Lagomorpha
-	B01.050.150.900.649.521.400 Hares
-	B01.050.150.900.649.521.700 Rabbits
-	B01.050.150.900.649.573 Marsupialia
-	B01.050.150.900.649.573.500 Macropodidae
-	B01.050.150.900.649.573.575 Opossums
-	B01.050.150.900.649.573.575.150 Didelphis
-	B01.050.150.900.649.573.575.500 Monodelphis
-	B01.050.150.900.649.573.587 Phalangeridae
-	B01.050.150.900.649.573.587.800 Trichosurus
-	B01.050.150.900.649.573.590 Phascolarctidae
-	B01.050.150.900.649.573.600 Potoroidae
-	B01.050.150.900.649.627 Monotremata
-	B01.050.150.900.649.627.300 Echidna
-	B01.050.150.900.649.627.600 Platypus
-	B01.050.150.900.649.709 Perissodactyla
-	B01.050.150.900.649.709.235 Equidae
-	B01.050.150.900.649.709.235.472 Horses
-	B01.050.150.900.649.801 Primates
-	B01.050.150.900.649.801.400 Haplorhini
-	B01.050.150.900.649.801.400.112 Catarrhini
-	B01.050.150.900.649.801.400.112.199 Cercopithecidae
-	B01.050.150.900.649.801.400.112.199.120 Cercopithecinae
-	B01.050.150.900.649.801.400.112.199.120.120 Cercocebus
-	B01.050.150.900.649.801.400.112.199.120.120.110 Cercocebus atys
-	B01.050.150.900.649.801.400.112.199.120.126 Cercopithecus
-	B01.050.150.900.649.801.400.112.199.120.126.110 Cercopithecus aethiops
-	B01.050.150.900.649.801.400.112.199.120.260 Erythrocebus
-	B01.050.150.900.649.801.400.112.199.120.260.260 Erythrocebus patas
-	B01.050.150.900.649.801.400.112.199.120.510 Macaca

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	B01.050.150.900.649.801.400.112.199.120.510.520	Macaca fascicularis
-	B01.050.150.900.649.801.400.112.199.120.510.550	Macaca mulatta
-	B01.050.150.900.649.801.400.112.199.120.510.560	Macaca nemestrina
-	B01.050.150.900.649.801.400.112.199.120.510.570	Macaca radiata
-	B01.050.150.900.649.801.400.112.199.120.530	Mandrillus
-	B01.050.150.900.649.801.400.112.199.120.610	Papio
-	B01.050.150.900.649.801.400.112.199.120.610.050	Papio anubis
-	B01.050.150.900.649.801.400.112.199.120.610.150	Papio cynocephalus
-	B01.050.150.900.649.801.400.112.199.120.610.400	Papio hamadryas
-	B01.050.150.900.649.801.400.112.199.120.610.600	Papio papio
-	B01.050.150.900.649.801.400.112.199.120.610.800	Papio ursinus
-	B01.050.150.900.649.801.400.112.199.120.730	Theropithecus
-	B01.050.150.900.649.801.400.112.199.150	Colobinae
-	B01.050.150.900.649.801.400.112.199.150.150	Colobus
-	B01.050.150.900.649.801.400.112.400	Hominidae
-	B01.050.150.900.649.801.400.112.400.375	Gorilla gorilla
-	B01.050.150.900.649.801.400.112.400.400	Humans
-	B01.050.150.900.649.801.400.112.400.550	Neanderthals
-	B01.050.150.900.649.801.400.112.400.600	Pan paniscus
-	B01.050.150.900.649.801.400.112.400.620	Pan troglodytes
-	B01.050.150.900.649.801.400.112.400.635	Pongo
-	B01.050.150.900.649.801.400.112.400.635.050	Pongo abelii
-	B01.050.150.900.649.801.400.112.400.635.650	Pongo pygmaeus
-	B01.050.150.900.649.801.400.112.420	Hylobatidae
-	B01.050.150.900.649.801.400.112.420.390	Hylobates
-	B01.050.150.900.649.801.400.600	Platyrrhini
-	B01.050.150.900.649.801.400.600.037	Aotidae
-	B01.050.150.900.649.801.400.600.037.092	Aotus trivirgatus
-	B01.050.150.900.649.801.400.600.075	Atelidae
-	B01.050.150.900.649.801.400.600.075.050	Alouattinae
-	B01.050.150.900.649.801.400.600.075.050.075	Alouatta
-	B01.050.150.900.649.801.400.600.075.075	Atelinae
-	B01.050.150.900.649.801.400.600.150	Cebidae
-	B01.050.150.900.649.801.400.600.150.120	Cebinae
-	B01.050.150.900.649.801.400.600.150.120.120	Cebus
-	B01.050.150.900.649.801.400.600.150.150	Callitrichinae

## MeSH Tree Changes for 2017

Type	Tree - heading
-	B01.050.150.900.649.801.400.600.150.150.105 Callimico
-	B01.050.150.900.649.801.400.600.150.150.114 Callithrix
-	B01.050.150.900.649.801.400.600.150.150.450 Leontopithecus
-	B01.050.150.900.649.801.400.600.150.150.710 Saguinus
-	B01.050.150.900.649.801.400.600.150.710 Saimirinae
-	B01.050.150.900.649.801.400.600.150.710.710 Saimiri
-	B01.050.150.900.649.801.400.600.575 Pitheciidae
-	B01.050.150.900.649.801.400.800 Tarsii
-	B01.050.150.900.649.801.400.800.500 Tarsiidae
-	B01.050.150.900.649.801.700 Strepsirhini
-	B01.050.150.900.649.801.700.200 Cheirogaleidae
-	B01.050.150.900.649.801.700.508 Lemuridae
-	B01.050.150.900.649.801.700.508.490 Lemur
-	B01.050.150.900.649.801.700.572 Lorisidae
-	B01.050.150.900.649.801.700.572.408 Galago
-	B01.050.150.900.649.833 Proboscidea Mammal
-	B01.050.150.900.649.833.249 Elephants
-	B01.050.150.900.649.833.500 Mammoths
-	B01.050.150.900.649.833.525 Mastodons
-	B01.050.150.900.649.865 Rodentia
-	B01.050.150.900.649.865.328 Chinchilla
-	B01.050.150.900.649.865.373 Cuniculidae
-	B01.050.150.900.649.865.419 Dasyproctidae
-	B01.050.150.900.649.865.510 Dipodomys
-	B01.050.150.900.649.865.545 Gophers
-	B01.050.150.900.649.865.550 Guinea Pigs
-	B01.050.150.900.649.865.600 Mole Rats
-	B01.050.150.900.649.865.600.700 Spalax
-	B01.050.150.900.649.865.635 Muridae
-	B01.050.150.900.649.865.635.070 Arvicolinae
-	B01.050.150.900.649.865.635.150 Cricetinae
-	B01.050.150.900.649.865.635.150.250 Cricetulus
-	B01.050.150.900.649.865.635.150.500 Mesocricetus
-	B01.050.150.900.649.865.635.150.630 Phodopus
-	B01.050.150.900.649.865.635.300 Gerbillinae
-	B01.050.150.900.649.865.635.505 Murinae

## MeSH Tree Changes for 2017

Type	Tree - heading
-	B01.050.150.900.649.865.635.505.500 Mice
-	B01.050.150.900.649.865.635.505.500.150 Mice, Congenic
-	B01.050.150.900.649.865.635.505.500.400 Mice, Inbred Strains
-	B01.050.150.900.649.865.635.505.500.400.025 Mice, 129 Strain
-	B01.050.150.900.649.865.635.505.500.400.200 Mice, Hairless
-	B01.050.150.900.649.865.635.505.500.400.300 Mice, Inbred A
-	B01.050.150.900.649.865.635.505.500.400.318 Mice, Inbred AKR
-	B01.050.150.900.649.865.635.505.500.400.338 Mice, Inbred BALB C
-	B01.050.150.900.649.865.635.505.500.400.388 Mice, Inbred C3H
-	B01.050.150.900.649.865.635.505.500.400.420 Mice, Inbred C57BL
-	B01.050.150.900.649.865.635.505.500.400.420.500 Mice, Inbred mdx
-	B01.050.150.900.649.865.635.505.500.400.440 Mice, Inbred CBA
-	B01.050.150.900.649.865.635.505.500.400.445 Mice, Inbred CFTR
-	B01.050.150.900.649.865.635.505.500.400.500 Mice, Inbred DBA
-	B01.050.150.900.649.865.635.505.500.400.510 Mice, Inbred ICR
-	B01.050.150.900.649.865.635.505.500.400.555 Mice, Inbred MRL lpr
-	B01.050.150.900.649.865.635.505.500.400.565 Mice, Inbred NOD
-	B01.050.150.900.649.865.635.505.500.400.580 Mice, Inbred NZB
-	B01.050.150.900.649.865.635.505.500.400.600 Mice, Inbred SENCAR
-	B01.050.150.900.649.865.635.505.500.550 Mice, Mutant Strains
-	B01.050.150.900.649.865.635.505.500.550.025 Mice, 129 Strain
-	B01.050.150.900.649.865.635.505.500.550.100 Mice, Biozzi
-	B01.050.150.900.649.865.635.505.500.550.230 Mice, Hairless
-	B01.050.150.900.649.865.635.505.500.550.265 Mice, Inbred mdx
-	B01.050.150.900.649.865.635.505.500.550.430 Mice, Jimpy
-	B01.050.150.900.649.865.635.505.500.550.455 Mice, Knockout
-	B01.050.150.900.649.865.635.505.500.550.480 Mice, Neurologic Mutants
-	B01.050.150.900.649.865.635.505.500.550.500 Mice, Nude
-	B01.050.150.900.649.865.635.505.500.550.530 Mice, Obese
-	B01.050.150.900.649.865.635.505.500.550.560 Mice, Quaking
-	B01.050.150.900.649.865.635.505.500.550.780 Mice, SCID
-	B01.050.150.900.649.865.635.505.500.800 Mice, Transgenic
-	B01.050.150.900.649.865.635.505.500.800.500 Mice, Knockout
-	B01.050.150.900.649.865.635.505.500.800.500.025 Mice, 129 Strain
-	B01.050.150.900.649.865.635.505.700 Rats
-	B01.050.150.900.649.865.635.505.700.400 Rats, Inbred Strains



## MeSH Tree Changes for 2017

Type	Tree - heading	
-	B01.050.150.900.649.865.635.505.700.400.080	Rats, Inbred ACI
-	B01.050.150.900.649.865.635.505.700.400.090	Rats, Inbred BB
-	B01.050.150.900.649.865.635.505.700.400.110	Rats, Inbred BN
-	B01.050.150.900.649.865.635.505.700.400.130	Rats, Inbred BUF
-	B01.050.150.900.649.865.635.505.700.400.165	Rats, Inbred Dahl
-	B01.050.150.900.649.865.635.505.700.400.200	Rats, Inbred F344
-	B01.050.150.900.649.865.635.505.700.400.275	Rats, Inbred LEC
-	B01.050.150.900.649.865.635.505.700.400.280	Rats, Inbred Lew
-	B01.050.150.900.649.865.635.505.700.400.290	Rats, Inbred OLETF
-	B01.050.150.900.649.865.635.505.700.400.300	Rats, Inbred SHR
-	B01.050.150.900.649.865.635.505.700.400.360	Rats, Inbred WF
-	B01.050.150.900.649.865.635.505.700.400.390	Rats, Inbred WKY
-	B01.050.150.900.649.865.635.505.700.500	Rats, Long-Evans
-	B01.050.150.900.649.865.635.505.700.550	Rats, Mutant Strains
-	B01.050.150.900.649.865.635.505.700.550.110	Rats, Brattleboro
-	B01.050.150.900.649.865.635.505.700.550.308	Rats, Gunn
-	B01.050.150.900.649.865.635.505.700.550.408	Rats, Hairless
-	B01.050.150.900.649.865.635.505.700.550.508	Rats, Nude
-	B01.050.150.900.649.865.635.505.700.550.700	Rats, Zucker
-	B01.050.150.900.649.865.635.505.700.750	Rats, Sprague-Dawley
-	B01.050.150.900.649.865.635.505.700.825	Rats, Transgenic
-	B01.050.150.900.649.865.635.505.700.900	Rats, Wistar
-	B01.050.150.900.649.865.635.680	Sigmodontinae
-	B01.050.150.900.649.865.635.680.510	Peromyscus
-	B01.050.150.900.649.865.635.700	Spalax
-	B01.050.150.900.649.865.640	Myoxidae
-	B01.050.150.900.649.865.663	Octodon
-	B01.050.150.900.649.865.692	Porcupines
-	B01.050.150.900.649.865.750	Sciuridae
-	B01.050.150.900.649.865.750.460	Marmota
-	B01.050.150.900.649.875	Scandentia
-	B01.050.150.900.649.875.770	Tupaiaidae
-	B01.050.150.900.649.875.770.770	Tupaia
-	B01.050.150.900.649.879	Sirenia
-	B01.050.150.900.649.879.125	Dugong
-	B01.050.150.900.649.879.750	Trichechus

## MeSH Tree Changes for 2017

Type	Tree - heading
-	B01.050.150.900.649.879.750.350 Trichechus inunguis
-	B01.050.150.900.649.879.750.500 Trichechus manatus
-	B01.050.150.900.649.900 Xenarthra
-	B01.050.150.900.649.900.200 Armadillos
-	B01.050.150.900.649.900.750 Sloths
-	B01.050.150.900.833 Reptiles
-	B01.050.150.900.833.100 Alligators and Crocodiles
-	B01.050.150.900.833.150 Dinosaurs
-	B01.050.150.900.833.393 Lizards
-	B01.050.150.900.833.393.500 Iguanas
-	B01.050.150.900.833.672 Snakes
-	B01.050.150.900.833.672.250 Boidae
-	B01.050.150.900.833.672.280 Colubridae
-	B01.050.150.900.833.672.350 Elapidae
-	B01.050.150.900.833.672.350.225 Bungarus
-	B01.050.150.900.833.672.875 Viperidae
-	B01.050.150.900.833.672.875.120 Agkistrodon
-	B01.050.150.900.833.672.875.200 Bothrops
-	B01.050.150.900.833.672.875.280 Crotalus
-	B01.050.150.900.833.672.875.750 Russell's Viper
-	B01.050.150.900.833.672.875.875 Trimeresurus
-	B01.050.150.900.833.848 Turtles
-	B01.050.500 Invertebrates
-	B01.050.500.091 Annelida
-	B01.050.500.091.426 Leeches
-	B01.050.500.091.426.400 Hirudo medicinalis
-	B01.050.500.091.657 Oligochaeta
-	B01.050.500.091.700 Polychaeta
-	B01.050.500.131 Arthropods
-	B01.050.500.131.166 Arachnida
-	B01.050.500.131.166.132 Acari
-	B01.050.500.131.166.132.419 Mites
-	B01.050.500.131.166.132.419.050 Acaridae
-	B01.050.500.131.166.132.419.575 Psoroptidae
-	B01.050.500.131.166.132.419.600 Pyroglyphidae
-	B01.050.500.131.166.132.419.600.200 Dermatophagoides farinae

## MeSH Tree Changes for 2017

Type	Tree - heading
-	B01.050.500.131.166.132.419.600.215 Dermatophagoides pteronyssinus
-	B01.050.500.131.166.132.419.750 Sarcoptidae
-	B01.050.500.131.166.132.419.750.800 Sarcoptes scabiei
-	B01.050.500.131.166.132.419.850 Tetranychidae
-	B01.050.500.131.166.132.419.875 Trombiculidae
-	B01.050.500.131.166.132.419.900 Varroidae
-	B01.050.500.131.166.132.832 Ticks
-	B01.050.500.131.166.132.832.100 Argasidae
-	B01.050.500.131.166.132.832.100.100 Argas
-	B01.050.500.131.166.132.832.100.700 Ornithodoros
-	B01.050.500.131.166.132.832.400 Ixodidae
-	B01.050.500.131.166.132.832.400.200 Dermacentor
-	B01.050.500.131.166.132.832.400.425 Ixodes
-	B01.050.500.131.166.132.832.400.712 Rhipicephalus
-	B01.050.500.131.166.132.832.400.712.700 Rhipicephalus sanguineus
-	B01.050.500.131.166.661 Scorpions
-	B01.050.500.131.166.803 Spiders
-	B01.050.500.131.166.803.200 Black Widow Spider
-	B01.050.500.131.166.803.600 Brown Recluse Spider
-	B01.050.500.131.365 Crustacea
-	B01.050.500.131.365.055 Amphipoda
-	B01.050.500.131.365.060 Anostraca
-	B01.050.500.131.365.060.050 Artemia
-	B01.050.500.131.365.070 Arguloida
-	B01.050.500.131.365.150 Cladocera
-	B01.050.500.131.365.150.200 Daphnia
-	B01.050.500.131.365.160 Copepoda
-	B01.050.500.131.365.190 Decapoda (Crustacea)
-	B01.050.500.131.365.190.050 Anomura
-	B01.050.500.131.365.190.070 Astacoidea
-	B01.050.500.131.365.190.110 Brachyura
-	B01.050.500.131.365.190.150 Crangonidae
-	B01.050.500.131.365.190.550 Nephropidae
-	B01.050.500.131.365.190.615 Palaemonidae
-	B01.050.500.131.365.190.625 Palinuridae
-	B01.050.500.131.365.190.635 Pandalidae

## MeSH Tree Changes for 2017

Type	Tree - heading
-	B01.050.500.131.365.190.660 Penaeidae
-	B01.050.500.131.365.250 Euphausiacea
-	B01.050.500.131.365.400 Isopoda
-	B01.050.500.131.365.640 Pentastomida
-	B01.050.500.131.365.880 Thoracica
-	B01.050.500.131.450 Horseshoe Crabs
-	B01.050.500.131.617 Insects
-	B01.050.500.131.617.069 Beetles
-	B01.050.500.131.617.069.307 Fireflies
-	B01.050.500.131.617.069.615 Tenebrio
-	B01.050.500.131.617.069.799 Tribolium
-	B01.050.500.131.617.069.900 Weevils
-	B01.050.500.131.617.104 Ephemeroptera
-	B01.050.500.131.617.140 Cockroaches
-	B01.050.500.131.617.140.100 Blattellidae
-	B01.050.500.131.617.140.580 Periplaneta
-	B01.050.500.131.617.289 Diptera
-	B01.050.500.131.617.289.208 Ceratopogonidae
-	B01.050.500.131.617.289.260 Chironomidae
-	B01.050.500.131.617.289.275 Culicidae
-	B01.050.500.131.617.289.275.100 Aedes
-	B01.050.500.131.617.289.275.120 Anopheles
-	B01.050.500.131.617.289.275.120.400 Anopheles gambiae
-	B01.050.500.131.617.289.275.225 Culex
-	B01.050.500.131.617.289.275.612 Ochlerotatus
-	B01.050.500.131.617.289.310 Drosophilidae
-	B01.050.500.131.617.289.310.250 Drosophila
-	B01.050.500.131.617.289.310.250.500 Drosophila melanogaster
-	B01.050.500.131.617.289.310.250.750 Drosophila simulans
-	B01.050.500.131.617.289.400 Glossinidae
-	B01.050.500.131.617.289.400.700 Tsetse Flies
-	B01.050.500.131.617.289.642 Muscidae
-	B01.050.500.131.617.289.642.470 Houseflies
-	B01.050.500.131.617.289.781 Psychodidae
-	B01.050.500.131.617.289.781.602 Phlebotomus
-	B01.050.500.131.617.289.797 Sarcophagidae

## MeSH Tree Changes for 2017

Type	Tree - heading
-	B01.050.500.131.617.289.800 Simuliidae
-	B01.050.500.131.617.289.850 Tephritidae
-	B01.050.500.131.617.289.850.150 Ceratitis capitata
-	B01.050.500.131.617.412 Hemiptera
-	B01.050.500.131.617.412.165 Aphids
-	B01.050.500.131.617.412.420 Heteroptera
-	B01.050.500.131.617.412.420.150 Cimicidae
-	B01.050.500.131.617.412.420.150.100 Bedbugs
-	B01.050.500.131.617.412.420.700 Reduviidae
-	B01.050.500.131.617.412.420.700.850 Triatominae
-	B01.050.500.131.617.412.420.700.850.600 Panstrongylus
-	B01.050.500.131.617.412.420.700.850.700 Rhodnius
-	B01.050.500.131.617.412.420.700.850.800 Triatoma
-	B01.050.500.131.617.412.600 Planococcus Insect
-	B01.050.500.131.617.479 Hymenoptera
-	B01.050.500.131.617.479.205 Ants
-	B01.050.500.131.617.479.387 Bees
-	B01.050.500.131.617.479.900 Wasps
-	B01.050.500.131.617.485 Isoptera
-	B01.050.500.131.617.561 Lepidoptera
-	B01.050.500.131.617.561.200 Butterflies
-	B01.050.500.131.617.561.650 Moths
-	B01.050.500.131.617.561.650.100 Bombyx
-	B01.050.500.131.617.561.650.525 Manduca
-	B01.050.500.131.617.561.650.700 Spodoptera
-	B01.050.500.131.617.590 Mantodea
-	B01.050.500.131.617.634 Odonata
-	B01.050.500.131.617.678 Orthoptera
-	B01.050.500.131.617.678.369 Grasshoppers
-	B01.050.500.131.617.678.369.500 Locusta migratoria
-	B01.050.500.131.617.678.410 Gryllidae
-	B01.050.500.131.617.690 Phthiraptera
-	B01.050.500.131.617.690.050 Amblycera
-	B01.050.500.131.617.690.060 Anoplura
-	B01.050.500.131.617.690.060.716 Pediculus
-	B01.050.500.131.617.690.060.730 Phthirus

## MeSH Tree Changes for 2017

Type	Tree - heading
-	B01.050.500.131.617.690.450 Ischnocera
-	B01.050.500.131.617.750 Siphonaptera
-	B01.050.500.131.617.750.200 Ctenocephalides
-	B01.050.500.131.617.750.800 Tunga
-	B01.050.500.131.617.750.950 Xenopsylla
-	B01.050.500.131.617.800 Thysanoptera
-	B01.050.500.217 Bryozoa
-	B01.050.500.272 Chordata, Nonvertebrate
-	B01.050.500.272.199 Cephalochordata
-	B01.050.500.272.199.500 Lancelets
-	B01.050.500.272.400 Hyperotreti
-	B01.050.500.272.400.380 Hagfishes
-	B01.050.500.272.727 Urochordata
-	B01.050.500.272.727.300 Ciona intestinalis
-	B01.050.500.308 Cnidaria
-	B01.050.500.308.237 Anthozoa
-	B01.050.500.308.237.700 Renilla
-	B01.050.500.308.237.750 Sea Anemones
-	B01.050.500.308.361 Cubozoa
-	B01.050.500.308.485 Hydrozoa
-	B01.050.500.308.485.400 Hydra
-	B01.050.500.308.550 Myxozoa
-	B01.050.500.308.550.500 Myxobolus
-	B01.050.500.308.690 Scyphozoa
-	B01.050.500.308.690.750 Sea Nettle, East Coast
-	B01.050.500.325 Ctenophora
-	B01.050.500.408 Echinodermata
-	B01.050.500.408.560 Sea Cucumbers
-	B01.050.500.408.560.150 Cucumaria
-	B01.050.500.408.560.400 Holothuria
-	B01.050.500.408.560.700 Stichopus
-	B01.050.500.408.578 Sea Urchins
-	B01.050.500.408.578.050 Anthocidaris
-	B01.050.500.408.578.070 Arbacia
-	B01.050.500.408.578.400 Hemicentrotus
-	B01.050.500.408.578.500 Lytechinus

## MeSH Tree Changes for 2017

Type	Tree - heading
-	B01.050.500.408.578.600 Paracentrotus
-	B01.050.500.408.578.800 Strongylocentrotus
-	B01.050.500.408.578.800.600 Strongylocentrotus purpuratus
-	B01.050.500.408.765 Starfish
-	B01.050.500.408.765.070 Asterias
-	B01.050.500.408.765.075 Asterina
-	B01.050.500.500 Helminths
-	B01.050.500.500.132 Acanthocephala
-	B01.050.500.500.132.546 Moniliformis
-	B01.050.500.500.294 Nematoda
-	B01.050.500.500.294.100 Adenophorea
-	B01.050.500.500.294.100.275 Enoplida
-	B01.050.500.500.294.100.275.200 Diocetophymatoidea
-	B01.050.500.500.294.100.275.570 Mermithoidea
-	B01.050.500.500.294.100.275.780 Trichuroidea
-	B01.050.500.500.294.100.275.780.180 Capillaria
-	B01.050.500.500.294.100.275.780.608 Trichinella
-	B01.050.500.500.294.100.275.780.608.700 Trichinella spiralis
-	B01.050.500.500.294.100.275.780.628 Trichuris
-	B01.050.500.500.294.700 Secernentea
-	B01.050.500.500.294.700.100 Ascaridida
-	B01.050.500.500.294.700.100.090 Ascaridia
-	B01.050.500.500.294.700.100.100 Ascaridoidea
-	B01.050.500.500.294.700.100.100.075 Anisakis
-	B01.050.500.500.294.700.100.100.108 Ascaris
-	B01.050.500.500.294.700.100.100.108.425 Ascaris lumbricoides
-	B01.050.500.500.294.700.100.100.108.700 Ascaris suum
-	B01.050.500.500.294.700.100.100.700 Toxascaris
-	B01.050.500.500.294.700.100.100.780 Toxocara
-	B01.050.500.500.294.700.100.100.780.225 Toxocara canis
-	B01.050.500.500.294.700.500 Oxyurida
-	B01.050.500.500.294.700.500.500 Oxyuroidea
-	B01.050.500.500.294.700.500.500.308 Enterobius
-	B01.050.500.500.294.700.700 Rhabditida
-	B01.050.500.500.294.700.700.640 Rhabdiasoidea
-	B01.050.500.500.294.700.700.640.680 Strongyloides

## MeSH Tree Changes for 2017

Type	Tree - heading
-	B01.050.500.500.294.700.700.640.680.600 Strongyloides ratti
-	B01.050.500.500.294.700.700.640.680.700 Strongyloides stercoralis
-	B01.050.500.500.294.700.700.660 Rhabditoidea
-	B01.050.500.500.294.700.700.660.250 Caenorhabditis
-	B01.050.500.500.294.700.700.660.250.250 Caenorhabditis elegans
-	B01.050.500.500.294.700.750 Spirurida
-	B01.050.500.500.294.700.750.225 Camallanina
-	B01.050.500.500.294.700.750.225.250 Dracunculoidea
-	B01.050.500.500.294.700.750.225.250.250 Dracunculus Nematode
-	B01.050.500.500.294.700.750.700 Spirurina
-	B01.050.500.500.294.700.750.700.300 Filarioidea
-	B01.050.500.500.294.700.750.700.300.043 Acanthocheilonema
-	B01.050.500.500.294.700.750.700.300.088 Brugia
-	B01.050.500.500.294.700.750.700.300.088.500 Brugia malayi
-	B01.050.500.500.294.700.750.700.300.088.550 Brugia pahangi
-	B01.050.500.500.294.700.750.700.300.208 Dipetalonema
-	B01.050.500.500.294.700.750.700.300.230 Dirofilaria
-	B01.050.500.500.294.700.750.700.300.230.300 Dirofilaria immitis
-	B01.050.500.500.294.700.750.700.300.230.650 Dirofilaria repens
-	B01.050.500.500.294.700.750.700.300.410 Loa
-	B01.050.500.500.294.700.750.700.300.438 Mansonella
-	B01.050.500.500.294.700.750.700.300.470 Microfilaria
-	B01.050.500.500.294.700.750.700.300.470 Microfilariae
-	B01.050.500.500.294.700.750.700.300.510 Onchocerca
-	B01.050.500.500.294.700.750.700.300.510.850 Onchocerca volvulus
-	B01.050.500.500.294.700.750.700.300.700 Setaria Nematode
-	B01.050.500.500.294.700.750.700.300.708 Wuchereria
-	B01.050.500.500.294.700.750.700.300.708.150 Wuchereria bancrofti
-	B01.050.500.500.294.700.750.700.680 Spiruroidea
-	B01.050.500.500.294.700.750.700.722 Thelazioidea
-	B01.050.500.500.294.700.750.700.722.375 Gnathostoma
-	B01.050.500.500.294.700.775 Strongylida
-	B01.050.500.500.294.700.775.100 Ancylostomatoidea
-	B01.050.500.500.294.700.775.100.100 Ancylostoma
-	B01.050.500.500.294.700.775.100.550 Necator
-	B01.050.500.500.294.700.775.100.550.100 Necator americanus



## MeSH Tree Changes for 2017

Type	Tree - heading
-	B01.050.500.500.294.700.775.400 Heligmosomatoidea
-	B01.050.500.500.294.700.775.400.550 Nematospiroides
-	B01.050.500.500.294.700.775.400.550.275 Nematospiroides dubius
-	B01.050.500.500.294.700.775.400.575 Nippostrongylus
-	B01.050.500.500.294.700.775.588 Metastrongyloidea
-	B01.050.500.500.294.700.775.588.080 Angiostrongylus
-	B01.050.500.500.294.700.775.588.080.225 Angiostrongylus cantonensis
-	B01.050.500.500.294.700.775.600 Molineoidea
-	B01.050.500.500.294.700.775.600.550 Nematodirus
-	B01.050.500.500.294.700.775.710 Strongyloidea
-	B01.050.500.500.294.700.775.710.600 Oesophagostomum
-	B01.050.500.500.294.700.775.710.700 Strongylus
-	B01.050.500.500.294.700.775.746 Trichostrongyloidea
-	B01.050.500.500.294.700.775.746.300 Dictyocaulus
-	B01.050.500.500.294.700.775.746.410 Haemonchus
-	B01.050.500.500.294.700.775.746.432 Ostertagia
-	B01.050.500.500.294.700.775.746.610 Trichostrongylus
-	B01.050.500.500.294.700.825 Tylenchida
-	B01.050.500.500.294.700.825.825 Tylenchoidea
-	B01.050.500.500.736 Platyhelminths
-	B01.050.500.500.736.215 Cestoda
-	B01.050.500.500.736.215.228 Diphyllbothrium
-	B01.050.500.500.736.215.228.831 Sparganum
-	B01.050.500.500.736.215.327 Echinococcus
-	B01.050.500.500.736.215.327.400 Echinococcus granulosus
-	B01.050.500.500.736.215.327.500 Echinococcus multilocularis
-	B01.050.500.500.736.215.454 Hymenolepis
-	B01.050.500.500.736.215.454.400 Hymenolepis diminuta
-	B01.050.500.500.736.215.454.410 Hymenolepis nana
-	B01.050.500.500.736.215.520 Mesocestoides
-	B01.050.500.500.736.215.800 Spirometra
-	B01.050.500.500.736.215.800.810 Sparganum
-	B01.050.500.500.736.215.895 Taenia
-	B01.050.500.500.736.215.895.286 Cysticercus
-	B01.050.500.500.736.215.895.643 Taenia saginata
-	B01.050.500.500.736.215.895.775 Taenia solium

## MeSH Tree Changes for 2017

Type	Tree - heading
-	B01.050.500.500.736.715 Trematoda
-	B01.050.500.500.736.715.300 Dicrocoeliidae
-	B01.050.500.500.736.715.300.308 Dicrocoelium
-	B01.050.500.500.736.715.360 Echinostomatidae
-	B01.050.500.500.736.715.360.310 Echinostoma
-	B01.050.500.500.736.715.408 Fasciolidae
-	B01.050.500.500.736.715.408.380 Fasciola
-	B01.050.500.500.736.715.408.380.420 Fasciola hepatica
-	B01.050.500.500.736.715.465 Heterophyidae
-	B01.050.500.500.736.715.520 Opisthorchidae
-	B01.050.500.500.736.715.520.210 Clonorchis sinensis
-	B01.050.500.500.736.715.520.700 Opisthorchis
-	B01.050.500.500.736.715.600 Paramphistomatidae
-	B01.050.500.500.736.715.770 Schistosomatidae
-	B01.050.500.500.736.715.770.680 Schistosoma
-	B01.050.500.500.736.715.770.680.510 Schistosoma haematobium
-	B01.050.500.500.736.715.770.680.570 Schistosoma japonicum
-	B01.050.500.500.736.715.770.680.700 Schistosoma mansoni
-	B01.050.500.500.736.715.800 Troglotrematidae
-	B01.050.500.500.736.715.800.610 Paragonimus
-	B01.050.500.500.736.715.800.610.900 Paragonimus westermani
-	B01.050.500.500.736.847 Turbellaria
-	B01.050.500.500.736.847.610 Planarians
-	B01.050.500.500.770 Rotifera
-	B01.050.500.644 Mollusca
-	B01.050.500.644.080 Bivalvia
-	B01.050.500.644.080.050 Arcidae
-	B01.050.500.644.080.050.700 Scapharca
-	B01.050.500.644.080.150 Cardiidae
-	B01.050.500.644.080.175 Corbicula
-	B01.050.500.644.080.225 Dreissena
-	B01.050.500.644.080.500 Mercenaria
-	B01.050.500.644.080.535 Mya
-	B01.050.500.644.080.537 Mytilidae
-	B01.050.500.644.080.537.500 Mytilus
-	B01.050.500.644.080.537.500.500 Mytilus edulis

## MeSH Tree Changes for 2017

Type	Tree - heading
-	B01.050.500.644.080.537.600 Perna
-	B01.050.500.644.080.643 Ostreidae
-	B01.050.500.644.080.643.150 Crassostrea
-	B01.050.500.644.080.643.600 Ostrea
-	B01.050.500.644.080.660 Pectinidae
-	B01.050.500.644.080.660.600 Pecten
-	B01.050.500.644.080.670 Pinctada
-	B01.050.500.644.080.750 Spisula
-	B01.050.500.644.080.850 Unionidae
-	B01.050.500.644.080.850.050 Anodonta
-	B01.050.500.644.080.850.850 Unio
-	B01.050.500.644.116 Cephalopoda
-	B01.050.500.644.116.150 Decapodiformes
-	B01.050.500.644.116.150.500 Loligo
-	B01.050.500.644.116.150.700 Sepia
-	B01.050.500.644.116.550 Nautilus
-	B01.050.500.644.116.600 Octopodiformes
-	B01.050.500.644.400 Gastropoda
-	B01.050.500.644.400.060 Aplysia
-	B01.050.500.644.400.150 Clione
-	B01.050.500.644.400.275 Conus Snail
-	B01.050.500.644.400.400 Hermisenda
-	B01.050.500.644.400.600 Pleurobranchaea
-	B01.050.500.644.400.750 Snails
-	B01.050.500.644.400.750.185 Biomphalaria
-	B01.050.500.644.400.750.323 Bulinus
-	B01.050.500.644.400.750.450 Helix (Snails)
-	B01.050.500.644.400.750.645 Lymnaea
-	B01.050.500.644.400.800 Tritonia Sea Slug
-	B01.050.500.644.600 Polyplacophora
-	B01.050.500.714 Parasites
-	B01.050.500.740 Placozoa
-	B01.050.500.802 Porifera
-	B01.050.500.802.039 Agelas
-	B01.050.500.802.080 Axinella
-	B01.050.500.802.160 Callyspongia

## MeSH Tree Changes for 2017

Type	Tree - heading
-	B01.050.500.802.175 Crambe Sponge
-	B01.050.500.802.200 Dysidea
-	B01.050.500.802.360 Geodia
-	B01.050.500.802.380 Haliclona
-	B01.050.500.802.650 Petrosia
-	B01.050.500.802.670 Plakortis
-	B01.050.500.802.775 Suberites
-	B01.050.500.802.785 Theonella
-	B01.050.500.802.900 Xestospongia
-	B01.050.500.901 Tardigrada
-	B01.175 Choanoflagellata
-	B01.206 Cryptophyta
-	B01.237 Diplomonadida
-	B01.237.385 Giardia
-	B01.237.385.400 Giardia lamblia
-	B01.268 Euglenozoa
-	B01.268.250 Euglenida
-	B01.268.250.320 Euglena
-	B01.268.250.320.418 Euglena gracilis
-	B01.268.250.320.500 Euglena longa
-	B01.268.475 Kinetoplastida
-	B01.268.475.868 Trypanosomatina
-	B01.268.475.868.110 Crithidia
-	B01.268.475.868.110.350 Crithidia fasciculata
-	B01.268.475.868.488 Leishmania
-	B01.268.475.868.488.080 Leishmania braziliensis
-	B01.268.475.868.488.230 Leishmania donovani
-	B01.268.475.868.488.240 Leishmania enriettii
-	B01.268.475.868.488.285 Leishmania guyanensis
-	B01.268.475.868.488.325 Leishmania infantum
-	B01.268.475.868.488.405 Leishmania major
-	B01.268.475.868.488.410 Leishmania mexicana
-	B01.268.475.868.488.680 Leishmania tropica
-	B01.268.475.868.887 Trypanosoma
-	B01.268.475.868.887.080 Trypanosoma brucei brucei
-	B01.268.475.868.887.110 Trypanosoma brucei gambiense

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	B01.268.475.868.887.115	Trypanosoma brucei rhodesiense
-	B01.268.475.868.887.128	Trypanosoma congolense
-	B01.268.475.868.887.140	Trypanosoma cruzi
-	B01.268.475.868.887.410	Trypanosoma lewisi
-	B01.268.475.868.887.680	Trypanosoma rangeli
-	B01.268.475.868.887.850	Trypanosoma vivax
-	B01.300	Fungi
-	B01.300.053	Agricultural Inoculants
-	B01.300.107	Ascomycota
-	B01.300.107.320	Eurotiales
-	B01.300.107.320.125	Byssoschlamys
-	B01.300.107.320.200	Emericella
-	B01.300.107.320.215	Eupenicillium
-	B01.300.107.320.218	Eurotium
-	B01.300.107.320.500	Monascus
-	B01.300.107.320.550	Neosartorya
-	B01.300.107.320.800	Talaromyces
-	B01.300.107.320.820	Thermoascus
-	B01.300.107.501	Hypocreales
-	B01.300.107.501.200	Claviceps
-	B01.300.107.501.220	Cordyceps
-	B01.300.107.501.310	Epichloe
-	B01.300.107.501.400	Gibberella
-	B01.300.107.501.430	Hypocrea
-	B01.300.107.501.550	Nectria
-	B01.300.107.575	Magnaporthe
-	B01.300.107.650	Onygenales
-	B01.300.107.650.080	Arthrodermataceae
-	B01.300.107.655	Ophiostomatales
-	B01.300.107.655.600	Ophiostoma
-	B01.300.107.685	Parmeliaceae
-	B01.300.107.685.500	Usnea
-	B01.300.107.715	Phyllachorales
-	B01.300.107.730	Pneumocystis
-	B01.300.107.730.650	Pneumocystis carinii
Old Tree	<b>B01.300.107.730.700</b>	<b>Pneumocystis jirovecii</b>

## MeSH Tree Changes for 2017

Type	Tree - heading
-	B01.300.107.780 Pseudallescheria
-	B01.300.107.795 Saccharomycetales
-	B01.300.107.795.190 Debaromyces
-	B01.300.107.795.192 Dekkera
-	B01.300.107.795.200 Dipodascus
-	B01.300.107.795.230 Endomyces
-	B01.300.107.795.250 Eremothecium
-	B01.300.107.795.393 Hanseniaspora
-	B01.300.107.795.536 Kluyveromyces
-	B01.300.107.795.550 Lipomyces
-	B01.300.107.795.575 Metschnikowia
-	B01.300.107.795.700 Pichia
-	B01.300.107.795.785 Saccharomyces
New Heading	<b>B01.300.107.795.785.400 Saccharomyces boulardii</b>
-	B01.300.107.795.785.800 Saccharomyces cerevisiae
-	B01.300.107.795.815 Saccharomycopsis
-	B01.300.107.795.830 Torulaspora
-	B01.300.107.795.905 Williopsis
-	B01.300.107.795.980 Yarrowia
-	B01.300.107.795.990 Zygosaccharomyces
-	B01.300.107.797 Schizosaccharomyces
-	B01.300.107.800 Sordariales
-	B01.300.107.800.150 Chaetomium
-	B01.300.107.800.629 Neurospora
-	B01.300.107.800.629.564 Neurospora crassa
-	B01.300.107.800.700 Podospora
-	B01.300.107.950 Xylariales
-	B01.300.179 Basidiomycota
-	B01.300.179.100 Agaricales
-	B01.300.179.100.105 Agaricus
-	B01.300.179.100.107 Agrocybe
-	B01.300.179.100.110 Amanita
-	B01.300.179.100.125 Armillaria
-	B01.300.179.100.200 Coprinus
-	B01.300.179.100.205 Cortinarius

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	B01.300.179.100.215	Cyathus
-	B01.300.179.100.320	Flammulina
-	B01.300.179.100.350	Grifola
-	B01.300.179.100.360	Hebeloma
-	B01.300.179.100.502	Laccaria
-	B01.300.179.100.515	Lentinula
-	B01.300.179.100.515.810	Shiitake Mushrooms
-	B01.300.179.100.550	Marasmius
-	B01.300.179.100.645	Pholiota
-	B01.300.179.100.650	Pleurotus
-	B01.300.179.100.660	Psilocybe
-	B01.300.179.100.750	Schizophyllum
-	B01.300.179.100.780	Termitomyces
-	B01.300.179.100.790	Tricholoma
-	B01.300.179.100.875	Volvariella
-	B01.300.179.110	Phakopsora pachyrhizi
-	B01.300.179.120	Polyporales
-	B01.300.179.120.174	Coriolaceae
-	B01.300.179.120.174.070	Antrodia
-	B01.300.179.120.174.600	Poria
-	B01.300.179.120.174.850	Trametes
-	B01.300.179.120.379	Ganoderma
-	B01.300.179.120.379.700	Reishi
-	B01.300.179.120.760	Polyporaceae
-	B01.300.179.120.760.649	Phanerochaete
-	B01.300.179.120.760.675	Polyporus
-	B01.300.179.120.760.690	Pycnoporus
New Heading	<b>B01.300.179.120.760.845</b>	<b>Wolfiporia</b>
-	B01.300.179.853	Ustilaginales
-	B01.300.179.853.800	Ustilago
-	B01.300.230	Blastocladiomycota
-	B01.300.230.074	Allomyces
-	B01.300.230.150	Blastocladiella
-	B01.300.230.200	Coelomomyces
Old Tree	<b>B01.300.248</b>	<b>Blood-Borne Pathogens</b>

## MeSH Tree Changes for 2017

Type	Tree - heading
-	B01.300.283 Chytridiomycota
-	B01.300.300 Fungi, Unclassified
-	B01.300.300.300 Entomophthorales
-	B01.300.300.300.150 Conidiobolus
-	B01.300.300.300.300 Entomophthora
-	B01.300.300.500 Mucorales
-	B01.300.300.500.030 Absidia
-	B01.300.300.500.150 Cunninghamella
-	B01.300.300.500.480 Mortierella
-	B01.300.300.500.500 Mucor
-	B01.300.300.500.750 Phycomyces
-	B01.300.300.500.795 Rhizomucor
-	B01.300.300.500.800 Rhizopus
-	B01.300.320 Glomeromycota
-	B01.300.340 Lichens
-	B01.300.340.458 Parmeliaceae
-	B01.300.340.458.500 Usnea
-	B01.300.360 Microsporidia
-	B01.300.360.500 Microsporea
-	B01.300.360.500.500 Microsporida
-	B01.300.360.500.500.100 Apansporoblastina
-	B01.300.360.500.500.100.175 Encephalitozoon
-	B01.300.360.500.500.100.175.175 Encephalitozoon cuniculi
-	B01.300.360.500.500.100.200 Enterocytozoon
-	B01.300.360.500.500.100.400 Nosema
-	B01.300.360.500.500.100.900 Vittiforma
-	B01.300.360.500.500.600 Pansporablastina
-	B01.300.360.500.500.600.050 Amblyospora
-	B01.300.360.500.500.600.400 Glugea
-	B01.300.360.500.500.600.500 Loma
-	B01.300.360.500.500.600.650 Pleistophora
-	B01.300.360.500.500.600.800 Thelohania
-	B01.300.360.520 Microsporidia, Unclassified
-	B01.300.381 Mitosporic Fungi
-	B01.300.381.025 Acremonium
-	B01.300.381.075 Alternaria



## MeSH Tree Changes for 2017

Type	Tree - heading
-	B01.300.381.081 Aspergillus
-	B01.300.381.081.170 Aspergillus flavus
-	B01.300.381.081.295 Aspergillus fumigatus
-	B01.300.381.081.420 Aspergillus nidulans
-	B01.300.381.081.450 Aspergillus niger
-	B01.300.381.081.480 Aspergillus ochraceus
-	B01.300.381.081.500 Aspergillus oryzae
-	B01.300.381.103 Beauveria
-	B01.300.381.125 Blastomyces
-	B01.300.381.128 Botrytis
-	B01.300.381.137 Brettanomyces
-	B01.300.381.147 Candida
-	B01.300.381.147.326 Candida albicans
-	B01.300.381.147.400 Candida glabrata
-	B01.300.381.147.800 Candida tropicalis
-	B01.300.381.170 Chrysosporium
-	B01.300.381.200 Cladosporium
-	B01.300.381.230 Coccidioides
-	B01.300.381.235 Colletotrichum
-	B01.300.381.258 Cryptococcus
-	B01.300.381.258.300 Cryptococcus gattii
-	B01.300.381.258.366 Cryptococcus neoformans
-	B01.300.381.299 Duddingtonia
-	B01.300.381.340 Epidermophyton
-	B01.300.381.355 Exophiala
-	B01.300.381.366 Fusarium
-	B01.300.381.380 Geotrichum
-	B01.300.381.385 Gliocladium
-	B01.300.381.420 Helminthosporium
-	B01.300.381.440 Histoplasma
-	B01.300.381.490 Kloeckera
-	B01.300.381.502 Lacazia
-	B01.300.381.515 Madurella
-	B01.300.381.522 Malassezia
-	B01.300.381.541 Metarhizium
-	B01.300.381.560 Microsporum

## MeSH Tree Changes for 2017

Type	Tree - heading
-	B01.300.381.570 Neotyphodium
-	B01.300.381.640 Paecilomyces
-	B01.300.381.648 Paracoccidioides
-	B01.300.381.662 Penicillium
-	B01.300.381.662.592 Penicillium chrysogenum
-	B01.300.381.706 Phialophora
-	B01.300.381.723 Pyricularia grisea
-	B01.300.381.740 Rhizoctonia
-	B01.300.381.750 Rhodotorula
-	B01.300.381.785 Scedosporium
-	B01.300.381.802 Scopulariopsis
-	B01.300.381.820 Sporothrix
-	B01.300.381.840 Stachybotrys
-	B01.300.381.910 Trichoderma
-	B01.300.381.928 Trichophyton
-	B01.300.381.940 Trichosporon
-	B01.300.381.950 Verticillium
-	B01.300.655 Mycorrhizae
-	B01.300.665 Neocallimastigomycota
-	B01.300.665.600 Neocallimastigales
-	B01.300.665.600.600 Neocallimastix
-	B01.300.665.600.675 Piromyces
-	B01.300.930 Yeasts
-	B01.300.930.150 Brettanomyces
-	B01.300.930.176 Candida
-	B01.300.930.176.326 Candida albicans
-	B01.300.930.176.400 Candida glabrata
-	B01.300.930.176.800 Candida tropicalis
-	B01.300.930.316 Cryptococcus
-	B01.300.930.316.300 Cryptococcus gattii
-	B01.300.930.316.366 Cryptococcus neoformans
-	B01.300.930.323 Debaromyces
-	B01.300.930.326 Dekkera
-	B01.300.930.330 Dipodascus
-	B01.300.930.340 Endomyces
-	B01.300.930.350 Eremothecium

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	B01.300.930.443	Hanseniaspora
-	B01.300.930.530	Kloeckera
-	B01.300.930.536	Kluyveromyces
-	B01.300.930.555	Lipomyces
-	B01.300.930.574	Malassezia
-	B01.300.930.587	Metschnikowia
-	B01.300.930.600	Pichia
-	B01.300.930.650	Rhodotorula
-	B01.300.930.705	Saccharomyces
New Heading	<b>B01.300.930.705.328</b>	<b>Saccharomyces boulardii</b>
-	B01.300.930.705.655	Saccharomyces cerevisiae
-	B01.300.930.710	Saccharomycopsis
-	B01.300.930.720	Schizosaccharomyces
-	B01.300.930.910	Torulaspora
-	B01.300.930.930	Trichosporon
-	B01.300.930.945	Williopsis
-	B01.300.930.980	Yarrowia
-	B01.400	Haptophyta
-	B01.500	Mesomycetozoea
-	B01.500.700	Rhinosporidium
-	B01.625	Oxymonadida
-	B01.630	Parabasalidea
-	B01.630.399	Hypermastigia
-	B01.630.800	Trichomonadida
-	B01.630.800.180	Dientamoeba
-	B01.630.800.808	Trichomonas
-	B01.630.800.808.717	Trichomonas vaginalis
-	B01.630.800.849	Tritrichomonas
-	B01.630.800.849.325	Tritrichomonas foetus
-	B01.650	Plants
New Tree	<b>B01.650.160</b>	<b>Crops, Agricultural</b>
New Tree	<b>B01.650.160.250</b>	<b>Edible Grain</b>
New Tree	<b>B01.650.160.956</b>	<b>Vegetables</b>

## MeSH Tree Changes for 2017

Type	Tree - heading
-	B01.650.232      Glaucophyta
-	B01.650.232.200      Cyanophora
-	B01.650.449      Plant Weeds
-	B01.650.510      Plants, Edible
New Tree	<a href="#">B01.650.510.250</a> <a href="#">Edible Grain</a>
Old Tree	<a href="#">B01.650.510.344</a> <a href="#">Crops, Agricultural</a>
Old Tree	<a href="#">B01.650.510.344.250</a> <a href="#">Edible Grain</a>
-	B01.650.510.956      Vegetables
-	B01.650.520      Plants, Genetically Modified
-	B01.650.560      Plants, Medicinal
-	B01.650.660      Plants, Toxic
-	B01.650.700      Rhodophyta
-	B01.650.700.150      Chondrus
-	B01.650.700.325      Gracilaria
-	B01.650.700.500      Laurencia
-	B01.650.700.585      Plocamium
-	B01.650.700.600      Porphyra
-	B01.650.700.610      Porphyridium
-	B01.650.723      Salt-Tolerant Plants
-	B01.650.819      Seedlings
-	B01.650.915      Trees
-	B01.650.940      Viridiplantae
-	B01.650.940.150      Chlorophyta
-	B01.650.940.150.133      Acetabularia
-	B01.650.940.150.300      Caulerpa
-	B01.650.940.150.469      Chlorella
-	B01.650.940.150.469.400      Chlorella vulgaris
-	B01.650.940.150.634      Prototheca
-	B01.650.940.150.800      Scenedesmus
-	B01.650.940.150.900      Ulva
-	B01.650.940.150.925      Volvocida
-	B01.650.940.150.925.344      Chlamydomonas
-	B01.650.940.150.925.344.650      Chlamydomonas reinhardtii
-	B01.650.940.150.950      Volvox
-	B01.650.940.800      Streptophyta

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	B01.650.940.800.150	Charophyceae
-	B01.650.940.800.150.150	Characeae
-	B01.650.940.800.150.150.150	Chara
-	B01.650.940.800.150.150.575	Nitella
-	B01.650.940.800.150.200	Desmidiiales
-	B01.650.940.800.150.200.150	Closterium
-	B01.650.940.800.150.200.500	Micrasterias
-	B01.650.940.800.150.990	Zygnematales
-	B01.650.940.800.150.990.550	Mougeotia
-	B01.650.940.800.150.990.800	Spirogyra
-	B01.650.940.800.575	Embryophyta
-	B01.650.940.800.575.100	Angiosperms
-	B01.650.940.800.575.100.009	Aceraceae
-	B01.650.940.800.575.100.009.500	Acer
-	B01.650.940.800.575.100.010	Acoraceae
-	B01.650.940.800.575.100.010.500	Acorus
-	B01.650.940.800.575.100.014	Adoxaceae
Old Tree	B01.650.940.800.575.100.018	Agavaceae
Old Tree	B01.650.940.800.575.100.018.249	Agave
Old Tree	B01.650.940.800.575.100.018.500	Yucca
-	B01.650.940.800.575.100.020	Aizoaceae
-	B01.650.940.800.575.100.020.500	Mesembryanthemum
-	B01.650.940.800.575.100.022	Alangiaceae
-	B01.650.940.800.575.100.025	Alismatidae
-	B01.650.940.800.575.100.025.033	Alismataceae
-	B01.650.940.800.575.100.025.033.044	Alisma
-	B01.650.940.800.575.100.025.033.777	Sagittaria
-	B01.650.940.800.575.100.025.399	Hydrocharitaceae
-	B01.650.940.800.575.100.025.699	Potamogetonaceae
-	B01.650.940.800.575.100.025.977	Zosteraceae
Old Tree	B01.650.940.800.575.100.027	Aloe
-	B01.650.940.800.575.100.037	Amaranthaceae
-	B01.650.940.800.575.100.037.100	Achyranthes
-	B01.650.940.800.575.100.037.130	Amaranthus
-	B01.650.940.800.575.100.037.333	Celosia
-	B01.650.940.800.575.100.044	Anacardiaceae

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	B01.650.940.800.575.100.044.277	Anacardium
-	B01.650.940.800.575.100.044.416	Mangifera
-	B01.650.940.800.575.100.044.555	Pistacia
-	B01.650.940.800.575.100.044.666	Rhus
-	B01.650.940.800.575.100.044.705	Semecarpus
-	B01.650.940.800.575.100.044.744	Toxicodendron
-	B01.650.940.800.575.100.065	Annonaceae
-	B01.650.940.800.575.100.065.500	Annona
-	B01.650.940.800.575.100.065.750	Asimina
-	B01.650.940.800.575.100.065.781	Cananga
-	B01.650.940.800.575.100.065.812	Goniothalamus
-	B01.650.940.800.575.100.065.875	Guatteria
-	B01.650.940.800.575.100.065.937	Polyalthia
-	B01.650.940.800.575.100.065.968	Rollinia
-	B01.650.940.800.575.100.065.984	Uvaria
-	B01.650.940.800.575.100.065.992	Xylopia
-	B01.650.940.800.575.100.075	Apiaceae
-	B01.650.940.800.575.100.075.077	Anethum graveolens
-	B01.650.940.800.575.100.075.088	Angelica
-	B01.650.940.800.575.100.075.088.077	Angelica archangelica
-	B01.650.940.800.575.100.075.088.538	Angelica sinensis
-	B01.650.940.800.575.100.075.099	Apium graveolens
-	B01.650.940.800.575.100.075.144	Bupleurum
-	B01.650.940.800.575.100.075.166	Carum
-	B01.650.940.800.575.100.075.180	Centella
-	B01.650.940.800.575.100.075.199	Cnidium
-	B01.650.940.800.575.100.075.211	Coriandrum
-	B01.650.940.800.575.100.075.233	Cuminum
-	B01.650.940.800.575.100.075.288	Daucus carota
-	B01.650.940.800.575.100.075.294	Eryngium
-	B01.650.940.800.575.100.075.300	Ferula
-	B01.650.940.800.575.100.075.333	Foeniculum
-	B01.650.940.800.575.100.075.388	Hemlock
-	B01.650.940.800.575.100.075.388.155	Cicuta
-	B01.650.940.800.575.100.075.388.188	Conium
-	B01.650.940.800.575.100.075.411	Heracleum

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	B01.650.940.800.575.100.075.488	Levisticum
-	B01.650.940.800.575.100.075.511	Ligusticum
-	B01.650.940.800.575.100.075.568	Oenanthe
-	B01.650.940.800.575.100.075.625	Pastinaca
-	B01.650.940.800.575.100.075.750	Petroselinum
-	B01.650.940.800.575.100.075.798	Pimpinella
-	B01.650.940.800.575.100.075.875	Sanicula
-	B01.650.940.800.575.100.075.937	Thapsia
Old Tree	B01.650.940.800.575.100.081	Apocynaceae
Old Tree	B01.650.940.800.575.100.081.043	Alstonia
Old Tree	B01.650.940.800.575.100.081.088	Amsonia
Old Tree	B01.650.940.800.575.100.081.131	Apocynum
Old Tree	B01.650.940.800.575.100.081.153	Asclepias
Old Tree	B01.650.940.800.575.100.081.174	Aspidosperma
Old Tree	B01.650.940.800.575.100.081.217	Calotropis
Old Tree	B01.650.940.800.575.100.081.260	Catharanthus
Old Tree	B01.650.940.800.575.100.081.303	Cryptolepis
Old Tree	B01.650.940.800.575.100.081.325	Cynanchum
Old Tree	B01.650.940.800.575.100.081.336	Gymnema
Old Tree	B01.650.940.800.575.100.081.336.500	Gymnema sylvestre
Old Tree	B01.650.940.800.575.100.081.341	Hemidesmus
Old Tree	B01.650.940.800.575.100.081.346	Holarrhena
Old Tree	B01.650.940.800.575.100.081.389	Hoodia
Old Tree	B01.650.940.800.575.100.081.411	Marsdenia
Old Tree	B01.650.940.800.575.100.081.432	Nerium
Old Tree	B01.650.940.800.575.100.081.604	Ochrosia
Old Tree	B01.650.940.800.575.100.081.691	Periploca
Old Tree	B01.650.940.800.575.100.081.777	Rauwolfia
Old Tree	B01.650.940.800.575.100.081.833	Strophanthus
Old Tree	B01.650.940.800.575.100.081.888	Tabernaemontana
Old Tree	B01.650.940.800.575.100.081.944	Thevetia
Old Tree	B01.650.940.800.575.100.081.958	Tylophora
Old Tree	B01.650.940.800.575.100.081.972	Vinca
Old Tree	B01.650.940.800.575.100.081.979	Vincetoxicum
Old Tree	B01.650.940.800.575.100.081.986	Voacanga
-	B01.650.940.800.575.100.084	Aquifoliaceae

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	B01.650.940.800.575.100.084.500	Ilex
-	B01.650.940.800.575.100.084.500.500	Ilex guayusa
-	B01.650.940.800.575.100.084.500.750	Ilex paraguariensis
-	B01.650.940.800.575.100.084.500.875	Ilex vomitoria
-	B01.650.940.800.575.100.085	Araceae
-	B01.650.940.800.575.100.085.066	Alocasia
-	B01.650.940.800.575.100.085.077	Amorphophallus
-	B01.650.940.800.575.100.085.088	Arisaema
-	B01.650.940.800.575.100.085.109	Arum
-	B01.650.940.800.575.100.085.177	Calla Plant
-	B01.650.940.800.575.100.085.199	Colocasia
-	B01.650.940.800.575.100.085.211	Cyrtosperma
-	B01.650.940.800.575.100.085.488	Philodendron
-	B01.650.940.800.575.100.085.500	Pinellia
-	B01.650.940.800.575.100.085.944	Xanthosoma
-	B01.650.940.800.575.100.085.972	Zantedeschia
-	B01.650.940.800.575.100.087	Araliaceae
-	B01.650.940.800.575.100.087.249	Aralia
-	B01.650.940.800.575.100.087.374	Eleutherococcus
-	B01.650.940.800.575.100.087.412	Hedera
-	B01.650.940.800.575.100.087.431	Kalopanax
-	B01.650.940.800.575.100.087.450	Oplopanax
-	B01.650.940.800.575.100.087.500	Panax
-	B01.650.940.800.575.100.087.500.500	Panax notoginseng
-	B01.650.940.800.575.100.093	Arecaceae
-	B01.650.940.800.575.100.093.088	Areca
-	B01.650.940.800.575.100.093.188	Calamus
-	B01.650.940.800.575.100.093.211	Cocos
-	B01.650.940.800.575.100.093.480	Euterpe
-	B01.650.940.800.575.100.093.615	Phoenixaceae
-	B01.650.940.800.575.100.093.750	Serenoa
-	B01.650.940.800.575.100.096	Aristolochiaceae
-	B01.650.940.800.575.100.096.211	Aristolochia
-	B01.650.940.800.575.100.096.233	Asarum
New Heading	<b>B01.650.940.800.575.100.099</b>	<b>Asparagales</b>



## MeSH Tree Changes for 2017

Type	Tree - heading	
New Heading	<b>B01.650.940.800.575.100.099.050</b>	<b>Amaryllidaceae</b>
New Tree	<a href="#">B01.650.940.800.575.100.099.050.060</a>	Allium
New Tree	<a href="#">B01.650.940.800.575.100.099.050.060.149</a>	Chive
New Tree	<a href="#">B01.650.940.800.575.100.099.050.060.300</a>	Garlic
New Tree	<a href="#">B01.650.940.800.575.100.099.050.060.600</a>	Onions
New Tree	<a href="#">B01.650.940.800.575.100.099.050.060.800</a>	Shallots
New Tree	<a href="#">B01.650.940.800.575.100.099.050.178</a>	Crinum
New Tree	<a href="#">B01.650.940.800.575.100.099.050.295</a>	Galanthus
New Tree	<a href="#">B01.650.940.800.575.100.099.050.530</a>	Lycoris
New Tree	<a href="#">B01.650.940.800.575.100.099.050.600</a>	Narcissus
New Heading	<b>B01.650.940.800.575.100.099.060</b>	<b>Asparagaceae</b>
New Tree	<a href="#">B01.650.940.800.575.100.099.060.032</a>	Agave
New Tree	<a href="#">B01.650.940.800.575.100.099.060.048</a>	Anemarrhena
New Tree	<a href="#">B01.650.940.800.575.100.099.060.063</a>	Asparagus Plant
New Tree	<a href="#">B01.650.940.800.575.100.099.060.125</a>	Camassia
New Tree	<a href="#">B01.650.940.800.575.100.099.060.188</a>	Convallaria
New Tree	<a href="#">B01.650.940.800.575.100.099.060.195</a>	Cordyline
New Tree	<a href="#">B01.650.940.800.575.100.099.060.250</a>	Dracaena
New Tree	<a href="#">B01.650.940.800.575.100.099.060.313</a>	Drimia
New Tree	<a href="#">B01.650.940.800.575.100.099.060.313</a>	Urginea
New Tree	<a href="#">B01.650.940.800.575.100.099.060.375</a>	Hosta
New Tree	<a href="#">B01.650.940.800.575.100.099.060.400</a>	Hyacinthus

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	<a href="#">B01.650.940.800.575.100.099.060.500</a>	Liriope Plant
New Tree	<a href="#">B01.650.940.800.575.100.099.060.560</a>	Maianthemum
New Tree	<a href="#">B01.650.940.800.575.100.099.060.560</a>	Smilacina
New Tree	<a href="#">B01.650.940.800.575.100.099.060.600</a>	Ophiopogon
New Tree	<a href="#">B01.650.940.800.575.100.099.060.650</a>	Polygonatum
New Tree	<a href="#">B01.650.940.800.575.100.099.060.725</a>	Ruscus
New Tree	<a href="#">B01.650.940.800.575.100.099.060.800</a>	Sansevieria
New Tree	<a href="#">B01.650.940.800.575.100.099.060.810</a>	Scilla
New Tree	<a href="#">B01.650.940.800.575.100.099.060.980</a>	Yucca
New Heading	<b><a href="#">B01.650.940.800.575.100.099.230</a></b>	<b>Hypoxidaceae</b>
New Tree	<a href="#">B01.650.940.800.575.100.099.230.190</a>	Curculigo
New Tree	<a href="#">B01.650.940.800.575.100.099.230.395</a>	Hypoxis
New Tree	<a href="#">B01.650.940.800.575.100.099.400</a>	Iridaceae
New Tree	<a href="#">B01.650.940.800.575.100.099.400.500</a>	Crocus
New Tree	<a href="#">B01.650.940.800.575.100.099.400.750</a>	Iris Plant
New Tree	<a href="#">B01.650.940.800.575.100.099.640</a>	Orchidaceae
New Tree	<a href="#">B01.650.940.800.575.100.099.640.222</a>	Dendrobium
New Tree	<a href="#">B01.650.940.800.575.100.099.640.500</a>	Gastrodia
New Tree	<a href="#">B01.650.940.800.575.100.099.640.888</a>	Vanilla
New Heading	<b><a href="#">B01.650.940.800.575.100.099.970</a></b>	<b>Xanthorrhoeaceae</b>
New Tree	<a href="#">B01.650.940.800.575.100.099.970.500</a>	Aloe
New Tree	<a href="#">B01.650.940.800.575.100.099.970.750</a>	Hemerocallis

## MeSH Tree Changes for 2017

Type	Tree - heading
-	B01.650.940.800.575.100.100 Asteraceae
-	B01.650.940.800.575.100.100.037 Achillea
-	B01.650.940.800.575.100.100.041 Achyrocline
-	B01.650.940.800.575.100.100.046 Ageratina
-	B01.650.940.800.575.100.100.056 Ageratum
-	B01.650.940.800.575.100.100.065 Ambrosia
-	B01.650.940.800.575.100.100.070 Ammi
-	B01.650.940.800.575.100.100.072 Arctium
-	B01.650.940.800.575.100.100.075 Arnica
-	B01.650.940.800.575.100.100.100 Artemisia
-	B01.650.940.800.575.100.100.100.099 Artemisia absinthium
-	B01.650.940.800.575.100.100.100.144 Artemisia annua
-	B01.650.940.800.575.100.100.111 Aster Plant
-	B01.650.940.800.575.100.100.118 Atractylis
-	B01.650.940.800.575.100.100.126 Atractylodes
-	B01.650.940.800.575.100.100.133 Baccharis
-	B01.650.940.800.575.100.100.137 Bidens
-	B01.650.940.800.575.100.100.141 Calendula
-	B01.650.940.800.575.100.100.154 Callilepis
-	B01.650.940.800.575.100.100.168 Carduus
-	B01.650.940.800.575.100.100.181 Carthamus
-	B01.650.940.800.575.100.100.181.500 Carthamus tinctorius
-	B01.650.940.800.575.100.100.188 Centaurea
-	B01.650.940.800.575.100.100.195 Chamomile
-	B01.650.940.800.575.100.100.195.099 Anthemis
-	B01.650.940.800.575.100.100.195.288 Chamaemelum
-	B01.650.940.800.575.100.100.195.488 Matricaria
-	B01.650.940.800.575.100.100.195.744 Tripleurospermum
-	B01.650.940.800.575.100.100.200 Chicory
-	B01.650.940.800.575.100.100.203 Chromolaena
-	B01.650.940.800.575.100.100.206 Chrysanthemum
-	B01.650.940.800.575.100.100.206.222 Chrysanthemum cinerariifolium
-	B01.650.940.800.575.100.100.227 Cirsium
-	B01.650.940.800.575.100.100.241 Cnicus
-	B01.650.940.800.575.100.100.244 Conyza
-	B01.650.940.800.575.100.100.248 Coreopsis

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	B01.650.940.800.575.100.100.255	Crepis
-	B01.650.940.800.575.100.100.269	Cynara
-	B01.650.940.800.575.100.100.269.500	Cynara scolymus
-	B01.650.940.800.575.100.100.289	Dahlia
-	B01.650.940.800.575.100.100.310	Echinacea
-	B01.650.940.800.575.100.100.317	Echinops Plant
-	B01.650.940.800.575.100.100.337	Eclipta
-	B01.650.940.800.575.100.100.342	Erigeron
-	B01.650.940.800.575.100.100.347	Eupatorium
-	B01.650.940.800.575.100.100.349	Flaveria
-	B01.650.940.800.575.100.100.352	Geigeria
-	B01.650.940.800.575.100.100.355	Gnaphalium
-	B01.650.940.800.575.100.100.356	Grindelia
-	B01.650.940.800.575.100.100.358	Haplopappus
-	B01.650.940.800.575.100.100.400	Helianthus
-	B01.650.940.800.575.100.100.425	Helichrysum
-	B01.650.940.800.575.100.100.471	Inula
-	B01.650.940.800.575.100.100.500	Lettuce
-	B01.650.940.800.575.100.100.506	Leuzea
-	B01.650.940.800.575.100.100.512	Mikania
-	B01.650.940.800.575.100.100.525	Milk Thistle
-	B01.650.940.800.575.100.100.543	Montanoa
-	B01.650.940.800.575.100.100.562	Onopordum
-	B01.650.940.800.575.100.100.596	Petasites
New Heading	<b>B01.650.940.800.575.100.100.605</b>	<b>Psacalium</b>
-	B01.650.940.800.575.100.100.613	Pulicaria
-	B01.650.940.800.575.100.100.631	Ratibida
-	B01.650.940.800.575.100.100.665	Rudbeckia
-	B01.650.940.800.575.100.100.700	Saussurea
-	B01.650.940.800.575.100.100.750	Scolymus
-	B01.650.940.800.575.100.100.775	Scorzonera
-	B01.650.940.800.575.100.100.800	Senecio
-	B01.650.940.800.575.100.100.816	Solidago
-	B01.650.940.800.575.100.100.833	Sonchus
-	B01.650.940.800.575.100.100.843	Stevia

## MeSH Tree Changes for 2017

Type	Tree - heading
-	B01.650.940.800.575.100.100.853 Tagetes
-	B01.650.940.800.575.100.100.874 Tanacetum
-	B01.650.940.800.575.100.100.874.500 Tanacetum parthenium
-	B01.650.940.800.575.100.100.895 Taraxacum
-	B01.650.940.800.575.100.100.916 Tragopogon
-	B01.650.940.800.575.100.100.958 Tussilago
-	B01.650.940.800.575.100.100.979 Verbesina
-	B01.650.940.800.575.100.100.984 Vernonia
-	B01.650.940.800.575.100.100.989 Wedelia
-	B01.650.940.800.575.100.100.994 Xanthium
-	B01.650.940.800.575.100.106 Balanophoraceae
-	B01.650.940.800.575.100.106.500 Cynomorium
-	B01.650.940.800.575.100.112 Balsaminaceae
-	B01.650.940.800.575.100.112.500 Impatiens
-	B01.650.940.800.575.100.118 Begoniaceae
-	B01.650.940.800.575.100.125 Berberidaceae
-	B01.650.940.800.575.100.125.249 Berberis
-	B01.650.940.800.575.100.125.311 Caulophyllum
-	B01.650.940.800.575.100.125.342 Epimedium
-	B01.650.940.800.575.100.125.374 Mahonia
-	B01.650.940.800.575.100.125.500 Podophyllum
-	B01.650.940.800.575.100.125.500.500 Podophyllum peltatum
-	B01.650.940.800.575.100.137 Betulaceae
-	B01.650.940.800.575.100.137.500 Alnus
-	B01.650.940.800.575.100.137.750 Betula
-	B01.650.940.800.575.100.137.875 Corylus
-	B01.650.940.800.575.100.146 Bixaceae
-	B01.650.940.800.575.100.148 Bombacaceae
-	B01.650.940.800.575.100.148.500 Adansonia
-	B01.650.940.800.575.100.148.750 Bombax
-	B01.650.940.800.575.100.148.875 Ceiba
-	B01.650.940.800.575.100.150 Boraginaceae
-	B01.650.940.800.575.100.150.122 Amsinckia
-	B01.650.940.800.575.100.150.188 Borago
-	B01.650.940.800.575.100.150.210 Comfrey
-	B01.650.940.800.575.100.150.327 Cordia

## MeSH Tree Changes for 2017

Type	Tree - heading
-	B01.650.940.800.575.100.150.385 Echium
-	B01.650.940.800.575.100.150.444 Heliotropium
-	B01.650.940.800.575.100.150.501 Lithospermum
-	B01.650.940.800.575.100.150.750 Pulmonaria
-	B01.650.940.800.575.100.157 Brassicaceae
-	B01.650.940.800.575.100.157.100 Arabidopsis
-	B01.650.940.800.575.100.157.150 Arabis
-	B01.650.940.800.575.100.157.175 Armoracia
-	B01.650.940.800.575.100.157.187 Barbarea
-	B01.650.940.800.575.100.157.200 Brassica
-	B01.650.940.800.575.100.157.200.249 Brassica napus
-	B01.650.940.800.575.100.157.200.277 Brassica rapa
-	B01.650.940.800.575.100.157.200.500 Mustard Plant
-	B01.650.940.800.575.100.157.244 Capsella
-	B01.650.940.800.575.100.157.255 Cardamine
-	B01.650.940.800.575.100.157.277 Crambe Plant
-	B01.650.940.800.575.100.157.357 Erysimum
-	B01.650.940.800.575.100.157.438 Isatis
-	B01.650.940.800.575.100.157.600 Lepidium
-	B01.650.940.800.575.100.157.600.500 Lepidium sativum
-	B01.650.940.800.575.100.157.644 Nasturtium
-	B01.650.940.800.575.100.157.711 Raphanus
-	B01.650.940.800.575.100.157.744 Rorippa
-	B01.650.940.800.575.100.157.811 Sinapis
-	B01.650.940.800.575.100.157.905 Thlaspi
-	B01.650.940.800.575.100.157.952 Wasabia
-	B01.650.940.800.575.100.159 Bromeliaceae
-	B01.650.940.800.575.100.159.099 Ananas
-	B01.650.940.800.575.100.159.244 Bromelia
-	B01.650.940.800.575.100.159.622 Tillandsia
-	B01.650.940.800.575.100.162 Burseraceae
-	B01.650.940.800.575.100.162.111 Boswellia
-	B01.650.940.800.575.100.162.138 Bursera
-	B01.650.940.800.575.100.162.166 Commiphora
-	B01.650.940.800.575.100.165 Buxaceae
-	B01.650.940.800.575.100.165.500 Buxus

## MeSH Tree Changes for 2017

Type	Tree - heading
-	B01.650.940.800.575.100.165.750 Pachysandra
-	B01.650.940.800.575.100.168 Cactaceae
-	B01.650.940.800.575.100.168.500 Opuntia
-	B01.650.940.800.575.100.169 Calycanthaceae
-	B01.650.940.800.575.100.171 Campanulaceae
-	B01.650.940.800.575.100.171.200 Codonopsis
-	B01.650.940.800.575.100.171.600 Lobelia
-	B01.650.940.800.575.100.171.800 Platycodon
-	B01.650.940.800.575.100.175 Cannabaceae
-	B01.650.940.800.575.100.175.500 Cannabis
-	B01.650.940.800.575.100.175.750 Humulus
-	B01.650.940.800.575.100.177 Capparaceae
-	B01.650.940.800.575.100.177.500 Capparis
-	B01.650.940.800.575.100.177.750 Cleome
-	B01.650.940.800.575.100.180 Caprifoliaceae
-	B01.650.940.800.575.100.180.388 Lonicera
-	B01.650.940.800.575.100.180.500 Sambucus
-	B01.650.940.800.575.100.180.500.500 Sambucus nigra
-	B01.650.940.800.575.100.180.625 Symphoricarpos
-	B01.650.940.800.575.100.180.750 Viburnum
-	B01.650.940.800.575.100.192 Caricaceae
-	B01.650.940.800.575.100.192.500 Carica
-	B01.650.940.800.575.100.195 Caryophyllaceae
-	B01.650.940.800.575.100.195.055 Agrostemma
-	B01.650.940.800.575.100.195.077 Arenaria Plant
-	B01.650.940.800.575.100.195.222 Dianthus
-	B01.650.940.800.575.100.195.416 Lychnis
-	B01.650.940.800.575.100.195.611 Saponaria
-	B01.650.940.800.575.100.195.655 Silene
-	B01.650.940.800.575.100.195.827 Stellaria
-	B01.650.940.800.575.100.195.913 Vaccaria
-	B01.650.940.800.575.100.196 Cecropiaceae
-	B01.650.940.800.575.100.196.188 Cecropia Plant
-	B01.650.940.800.575.100.197 Celastraceae
-	B01.650.940.800.575.100.197.155 Catha
-	B01.650.940.800.575.100.197.199 Celastrus

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	B01.650.940.800.575.100.197.366	Euonymus
-	B01.650.940.800.575.100.197.577	Maytenus
-	B01.650.940.800.575.100.197.682	Salacia
-	B01.650.940.800.575.100.197.788	Tripterygium
-	B01.650.940.800.575.100.200	Chenopodiaceae
-	B01.650.940.800.575.100.200.199	Atriplex
-	B01.650.940.800.575.100.200.299	Bassia scoparia
-	B01.650.940.800.575.100.200.399	Beta vulgaris
-	B01.650.940.800.575.100.200.599	Chenopodium
-	B01.650.940.800.575.100.200.599.500	Chenopodium album
-	B01.650.940.800.575.100.200.599.750	Chenopodium ambrosioides
-	B01.650.940.800.575.100.200.599.875	Chenopodium quinoa
-	B01.650.940.800.575.100.200.699	Salsola
-	B01.650.940.800.575.100.200.800	Spinacia oleracea
-	B01.650.940.800.575.100.204	Cistaceae
-	B01.650.940.800.575.100.204.177	Cistus
-	B01.650.940.800.575.100.216	Clethraceae
-	B01.650.940.800.575.100.228	Combretaceae
-	B01.650.940.800.575.100.228.166	Combretum
-	B01.650.940.800.575.100.228.583	Terminalia
-	B01.650.940.800.575.100.233	Commelinaceae
-	B01.650.940.800.575.100.233.222	Commelina
-	B01.650.940.800.575.100.233.611	Tradescantia
-	B01.650.940.800.575.100.238	Convolvulaceae
-	B01.650.940.800.575.100.238.124	Bonamia Plant
-	B01.650.940.800.575.100.238.249	Calystegia
-	B01.650.940.800.575.100.238.374	Convolvulus
-	B01.650.940.800.575.100.238.500	Ipomoea
-	B01.650.940.800.575.100.238.500.500	Ipomoea batatas
-	B01.650.940.800.575.100.238.500.750	Ipomoea nil
-	B01.650.940.800.575.100.241	Cornaceae
-	B01.650.940.800.575.100.241.299	Cornus
Old Tree	B01.650.940.800.575.100.270	Corsiaceae
-	B01.650.940.800.575.100.300	Cucurbitaceae
-	B01.650.940.800.575.100.300.093	Bryonia
-	B01.650.940.800.575.100.300.140	Citrullus



## MeSH Tree Changes for 2017

Type	Tree - heading
-	B01.650.940.800.575.100.300.140.500 Citrullus colocynthis
-	B01.650.940.800.575.100.300.188 Cucumis
-	B01.650.940.800.575.100.300.188.444 Cucumis melo
-	B01.650.940.800.575.100.300.188.666 Cucumis sativus
-	B01.650.940.800.575.100.300.222 Cucurbita
-	B01.650.940.800.575.100.300.444 Gynostemma
-	B01.650.940.800.575.100.300.522 Luffa
-	B01.650.940.800.575.100.300.655 Momordica
-	B01.650.940.800.575.100.300.655.249 Momordica charantia
-	B01.650.940.800.575.100.300.827 Trichosanthes
-	B01.650.940.800.575.100.306 Cuscuta
-	B01.650.940.800.575.100.312 Cyperaceae
-	B01.650.940.800.575.100.312.099 Carex Plant
-	B01.650.940.800.575.100.312.199 Cyperus
-	B01.650.940.800.575.100.312.299 Eleocharis
-	B01.650.940.800.575.100.318 Dilleniaceae
-	B01.650.940.800.575.100.321 Dioncophyllaceae
-	B01.650.940.800.575.100.325 Dioscoreaceae
-	B01.650.940.800.575.100.325.500 Dioscorea
-	B01.650.940.800.575.100.325.750 Tamus
-	B01.650.940.800.575.100.327 Dipsacaceae
-	B01.650.940.800.575.100.329 Droseraceae
-	B01.650.940.800.575.100.329.500 Drosera
-	B01.650.940.800.575.100.333 Ebenaceae
-	B01.650.940.800.575.100.333.500 Diospyros
-	B01.650.940.800.575.100.336 Elaeagnaceae
-	B01.650.940.800.575.100.336.500 Hippophae
-	B01.650.940.800.575.100.339 Elaeocarpaceae
-	B01.650.940.800.575.100.343 Ericaceae
-	B01.650.940.800.575.100.343.099 Arctostaphylos
-	B01.650.940.800.575.100.343.211 Calluna
-	B01.650.940.800.575.100.343.267 Gaultheria
-	B01.650.940.800.575.100.343.324 Ledum
-	B01.650.940.800.575.100.343.549 Rhododendron
-	B01.650.940.800.575.100.343.774 Vaccinium
-	B01.650.940.800.575.100.343.774.249 Blueberry Plant

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	B01.650.940.800.575.100.343.774.249	Blueberry Plants
-	B01.650.940.800.575.100.343.774.500	Huckleberry Plant
-	B01.650.940.800.575.100.343.774.750	Vaccinium macrocarpon
-	B01.650.940.800.575.100.343.774.875	Vaccinium myrtillus
-	B01.650.940.800.575.100.343.774.937	Vaccinium vitis-idaea
-	B01.650.940.800.575.100.344	Eriocaulaceae
-	B01.650.940.800.575.100.346	Erythroxylaceae
-	B01.650.940.800.575.100.346.500	Coca
-	B01.650.940.800.575.100.348	Eucommiaceae
-	B01.650.940.800.575.100.350	Euphorbiaceae
-	B01.650.940.800.575.100.350.038	Acalypha
-	B01.650.940.800.575.100.350.077	Aleurites
-	B01.650.940.800.575.100.350.199	Croton
-	B01.650.940.800.575.100.350.333	Euphorbia
-	B01.650.940.800.575.100.350.399	Hevea
-	B01.650.940.800.575.100.350.461	Hippomane
-	B01.650.940.800.575.100.350.488	Jatropha
-	B01.650.940.800.575.100.350.505	Mallotus Plant
-	B01.650.940.800.575.100.350.535	Manihot
-	B01.650.940.800.575.100.350.622	Phyllanthus
-	B01.650.940.800.575.100.350.622.500	Phyllanthus emblica
-	B01.650.940.800.575.100.350.650	Ricinus
-	B01.650.940.800.575.100.350.650.300	Castor Bean
-	B01.650.940.800.575.100.350.737	Sapium
-	B01.650.940.800.575.100.350.825	Suregada
-	B01.650.940.800.575.100.401	Fabaceae
-	B01.650.940.800.575.100.401.012	Abrus
-	B01.650.940.800.575.100.401.025	Acacia
-	B01.650.940.800.575.100.401.051	Albizzia
-	B01.650.940.800.575.100.401.077	Arachis
-	B01.650.940.800.575.100.401.082	Aspalathus
-	B01.650.940.800.575.100.401.087	Astragalus Plant
-	B01.650.940.800.575.100.401.087.500	Astragalus gummifer
-	B01.650.940.800.575.100.401.087.750	Astragalus membranaceus
-	B01.650.940.800.575.100.401.088	Bauhinia
-	B01.650.940.800.575.100.401.090	Butea

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	B01.650.940.800.575.100.401.093	Caesalpinia
-	B01.650.940.800.575.100.401.094	Cajanus
-	B01.650.940.800.575.100.401.096	Canavalia
-	B01.650.940.800.575.100.401.098	Caragana
-	B01.650.940.800.575.100.401.100	Cassia
-	B01.650.940.800.575.100.401.112	Castanospermum
-	B01.650.940.800.575.100.401.125	Chamaecrista
-	B01.650.940.800.575.100.401.150	Cicer
-	B01.650.940.800.575.100.401.162	Clitoria
-	B01.650.940.800.575.100.401.175	Crotalaria
-	B01.650.940.800.575.100.401.181	Cyamopsis
-	B01.650.940.800.575.100.401.184	Cyclopia Plant
-	B01.650.940.800.575.100.401.187	Cytisus
-	B01.650.940.800.575.100.401.190	Dalbergia
-	B01.650.940.800.575.100.401.191	Derris
-	B01.650.940.800.575.100.401.192	Dioclea
-	B01.650.940.800.575.100.401.194	Dipteryx
-	B01.650.940.800.575.100.401.196	Dolichos
-	B01.650.940.800.575.100.401.200	Erythrina
-	B01.650.940.800.575.100.401.250	Galega
-	B01.650.940.800.575.100.401.262	Genista
-	B01.650.940.800.575.100.401.275	Gleditsia
-	B01.650.940.800.575.100.401.300	Glycyrrhiza
-	B01.650.940.800.575.100.401.300.500	Glycyrrhiza uralensis
-	B01.650.940.800.575.100.401.318	Griffonia
-	B01.650.940.800.575.100.401.327	Hymenaea
-	B01.650.940.800.575.100.401.337	Indigofera
-	B01.650.940.800.575.100.401.356	Laburnum
-	B01.650.940.800.575.100.401.375	Lathyrus
-	B01.650.940.800.575.100.401.450	Lens Plant
-	B01.650.940.800.575.100.401.525	Lespedeza
New Heading	<b>B01.650.940.800.575.100.401.544</b>	<b>Loteae</b>
Old Tree	<b>B01.650.940.800.575.100.401.562</b>	<b>Lotus</b>
-	B01.650.940.800.575.100.401.571	Lupinus
-	B01.650.940.800.575.100.401.581	Maackia

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	B01.650.940.800.575.100.401.590	Medicago
-	B01.650.940.800.575.100.401.590.500	Medicago sativa
-	B01.650.940.800.575.100.401.590.750	Medicago truncatula
-	B01.650.940.800.575.100.401.592	Melilotus
-	B01.650.940.800.575.100.401.594	Millettia
-	B01.650.940.800.575.100.401.596	Mimosa
-	B01.650.940.800.575.100.401.598	Mucuna
-	B01.650.940.800.575.100.401.601	Myroxylon
-	B01.650.940.800.575.100.401.604	Ononis
-	B01.650.940.800.575.100.401.606	Oxytropis
-	B01.650.940.800.575.100.401.611	Pachyrhizus
-	B01.650.940.800.575.100.401.625	Peas
-	B01.650.940.800.575.100.401.649	Phaseolus
-	B01.650.940.800.575.100.401.694	Physostigma
-	B01.650.940.800.575.100.401.698	Pongamia
-	B01.650.940.800.575.100.401.702	Prosopis
-	B01.650.940.800.575.100.401.710	Psoralea
-	B01.650.940.800.575.100.401.712	Pterocarpus
-	B01.650.940.800.575.100.401.714	Pueraria
-	B01.650.940.800.575.100.401.716	Robinia
-	B01.650.940.800.575.100.401.718	Senna Plant
-	B01.650.940.800.575.100.401.726	Sesbania
-	B01.650.940.800.575.100.401.734	Sophora
-	B01.650.940.800.575.100.401.750	Soybeans
-	B01.650.940.800.575.100.401.812	Spartium
-	B01.650.940.800.575.100.401.843	Sphenostylis
-	B01.650.940.800.575.100.401.875	Tamarindus
-	B01.650.940.800.575.100.401.890	Tephrosia
-	B01.650.940.800.575.100.401.898	Tetrapleura
-	B01.650.940.800.575.100.401.906	Trifolium
-	B01.650.940.800.575.100.401.937	Trigonella
-	B01.650.940.800.575.100.401.952	Ulex
-	B01.650.940.800.575.100.401.968	Vicia
-	B01.650.940.800.575.100.401.968.500	Vicia faba
-	B01.650.940.800.575.100.401.968.750	Vicia sativa
New	<b>B01.650.940.800.575.100.401.976</b>	<b>Vigna</b>

## MeSH Tree Changes for 2017

Type	Tree - heading	
Heading		
-	B01.650.940.800.575.100.401.984	Wisteria
-	B01.650.940.800.575.100.425	Fagaceae
-	B01.650.940.800.575.100.425.249	Fagus
-	B01.650.940.800.575.100.425.500	Quercus
-	B01.650.940.800.575.100.437	Flacourtiaceae
-	B01.650.940.800.575.100.437.249	Casearia
-	B01.650.940.800.575.100.437.500	Ryania
-	B01.650.940.800.575.100.443	Fumariaceae
-	B01.650.940.800.575.100.443.500	Corydalis
-	B01.650.940.800.575.100.443.750	Fumaria
Old Tree	<b>B01.650.940.800.575.100.450</b>	<b>Gentianaceae</b>
Old Tree	<b>B01.650.940.800.575.100.450.188</b>	<b>Centaurium</b>
Old Tree	<b>B01.650.940.800.575.100.450.391</b>	<b>Gentiana</b>
Old Tree	<b>B01.650.940.800.575.100.450.492</b>	<b>Gentianella</b>
Old Tree	<b>B01.650.940.800.575.100.450.594</b>	<b>Swertia</b>
New Heading	<b>B01.650.940.800.575.100.456</b>	<b>Gentianales</b>
New Tree	<b>B01.650.940.800.575.100.456.500</b>	<b>Apocynaceae</b>
New Tree	<b>B01.650.940.800.575.100.456.500.043</b>	<b>Alstonia</b>
New Tree	<b>B01.650.940.800.575.100.456.500.088</b>	<b>Amsonia</b>
New Tree	<b>B01.650.940.800.575.100.456.500.131</b>	<b>Apocynum</b>
New Tree	<b>B01.650.940.800.575.100.456.500.153</b>	<b>Asclepias</b>
New Tree	<b>B01.650.940.800.575.100.456.500.174</b>	<b>Aspidosperma</b>
New Tree	<b>B01.650.940.800.575.100.456.500.217</b>	<b>Calotropis</b>
New Tree	<b>B01.650.940.800.575.100.456.500.260</b>	<b>Catharanthus</b>
New Tree	<b>B01.650.940.800.575.100.456.500.303</b>	<b>Cryptolepis</b>
New Tree	<b>B01.650.940.800.575.100.456.500.325</b>	<b>Cynanchum</b>
New Tree	<b>B01.650.940.800.575.100.456.500.336</b>	<b>Gymnema</b>

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	B01.650.940.800.575.100.456.500.336.500 <span style="float: right;">Gymnema sylvestre</span>
New Tree	B01.650.940.800.575.100.456.500.341 <span style="float: right;">Hemidesmus</span>
New Tree	B01.650.940.800.575.100.456.500.346 <span style="float: right;">Holarrhena</span>
New Tree	B01.650.940.800.575.100.456.500.389 <span style="float: right;">Hoodia</span>
New Tree	B01.650.940.800.575.100.456.500.411 <span style="float: right;">Marsdenia</span>
New Tree	B01.650.940.800.575.100.456.500.432 <span style="float: right;">Nerium</span>
New Tree	B01.650.940.800.575.100.456.500.604 <span style="float: right;">Ochrosia</span>
New Tree	B01.650.940.800.575.100.456.500.691 <span style="float: right;">Periploca</span>
New Tree	B01.650.940.800.575.100.456.500.777 <span style="float: right;">Rauwolfia</span>
New Tree	B01.650.940.800.575.100.456.500.833 <span style="float: right;">Strophanthus</span>
New Tree	B01.650.940.800.575.100.456.500.888 <span style="float: right;">Tabernaemontana</span>
New Tree	B01.650.940.800.575.100.456.500.944 <span style="float: right;">Thevetia</span>
New Tree	B01.650.940.800.575.100.456.500.958 <span style="float: right;">Tylophora</span>
New Tree	B01.650.940.800.575.100.456.500.972 <span style="float: right;">Vinca</span>
New Tree	B01.650.940.800.575.100.456.500.979 <span style="float: right;">Vincetoxicum</span>
New Tree	B01.650.940.800.575.100.456.500.986 <span style="float: right;">Voacanga</span>
New Tree	B01.650.940.800.575.100.456.750 <span style="float: right;">Gentianaceae</span>
New Tree	B01.650.940.800.575.100.456.750.188 <span style="float: right;">Centaurium</span>
New Tree	B01.650.940.800.575.100.456.750.391 <span style="float: right;">Gentiana</span>
New Tree	B01.650.940.800.575.100.456.750.492 <span style="float: right;">Gentianella</span>
New Tree	B01.650.940.800.575.100.456.750.594 <span style="float: right;">Swertia</span>
New Tree	B01.650.940.800.575.100.456.875 <span style="float: right;">Loganiaceae</span>

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	B01.650.940.800.575.100.456.875.500	Gelsemium
New Tree	B01.650.940.800.575.100.456.875.750	Strychnos
New Tree	B01.650.940.800.575.100.456.875.750.500	Strychnos nux-vomica
New Tree	B01.650.940.800.575.100.456.937	Rubiaceae
New Tree	B01.650.940.800.575.100.456.937.200	Cephaelis
New Tree	B01.650.940.800.575.100.456.937.250	Cinchona
New Tree	B01.650.940.800.575.100.456.937.275	Coffea
New Tree	B01.650.940.800.575.100.456.937.377	Galium
New Tree	B01.650.940.800.575.100.456.937.388	Gardenia
New Tree	B01.650.940.800.575.100.456.937.404	Hamelia
New Tree	B01.650.940.800.575.100.456.937.422	Hedyotis
New Tree	B01.650.940.800.575.100.456.937.555	Mitragyna
New Tree	B01.650.940.800.575.100.456.937.566	Morinda
New Tree	B01.650.940.800.575.100.456.937.611	Oldenlandia
New Tree	B01.650.940.800.575.100.456.937.805	Pausinystalia
New Tree	B01.650.940.800.575.100.456.937.844	Psychotria
New Tree	B01.650.940.800.575.100.456.937.876	Rubia
New Tree	B01.650.940.800.575.100.456.937.909	Uncaria
New Tree	B01.650.940.800.575.100.456.937.909.500	Cat's Claw
-	B01.650.940.800.575.100.462	Geraniaceae
-	B01.650.940.800.575.100.462.500	Geranium
-	B01.650.940.800.575.100.462.750	Pelargonium
-	B01.650.940.800.575.100.475	Hamamelidaceae
-	B01.650.940.800.575.100.475.500	Hamamelis

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	B01.650.940.800.575.100.475.750	Liquidambar
-	B01.650.940.800.575.100.489	Hernandiaceae
-	B01.650.940.800.575.100.503	Hippocastanaceae
-	B01.650.940.800.575.100.503.500	Aesculus
-	B01.650.940.800.575.100.517	Hippocrateaceae
-	B01.650.940.800.575.100.531	Hydrophyllaceae
-	B01.650.940.800.575.100.531.500	Eriodictyon
-	B01.650.940.800.575.100.539	Illicium
Old Tree	B01.650.940.800.575.100.549	Iridaceae
Old Tree	B01.650.940.800.575.100.549.500	Crocus
Old Tree	B01.650.940.800.575.100.549.750	Iris Plant
-	B01.650.940.800.575.100.569	Juglandaceae
-	B01.650.940.800.575.100.569.249	Carya
-	B01.650.940.800.575.100.569.500	Juglans
-	B01.650.940.800.575.100.572	Krameriaceae
-	B01.650.940.800.575.100.583	Lamiales
-	B01.650.940.800.575.100.583.040	Acanthaceae
-	B01.650.940.800.575.100.583.040.022	Justicia
-	B01.650.940.800.575.100.583.040.044	Andrographis
-	B01.650.940.800.575.100.583.040.522	Avicennia
-	B01.650.940.800.575.100.583.080	Bignoniaceae
-	B01.650.940.800.575.100.583.080.500	Tabebuia
-	B01.650.940.800.575.100.583.190	Calceolariaceae
-	B01.650.940.800.575.100.583.300	Craterostigma
-	B01.650.940.800.575.100.583.520	Lamiaceae
-	B01.650.940.800.575.100.583.520.049	Agastache
-	B01.650.940.800.575.100.583.520.051	Ajuga
-	B01.650.940.800.575.100.583.520.075	Ballota
-	B01.650.940.800.575.100.583.520.098	Callicarpa
-	B01.650.940.800.575.100.583.520.121	Clerodendrum
-	B01.650.940.800.575.100.583.520.166	Coleus
-	B01.650.940.800.575.100.583.520.231	Hedeoma
-	B01.650.940.800.575.100.583.520.297	Hyptis
-	B01.650.940.800.575.100.583.520.330	Hyssopus Plant
-	B01.650.940.800.575.100.583.520.362	Isodon
-	B01.650.940.800.575.100.583.520.428	Lavandula



## MeSH Tree Changes for 2017

Type	Tree - heading	
-	B01.650.940.800.575.100.583.520.446	Leonurus
-	B01.650.940.800.575.100.583.520.455	Lycopus
-	B01.650.940.800.575.100.583.520.465	Marrubium
-	B01.650.940.800.575.100.583.520.474	Melissa
-	B01.650.940.800.575.100.583.520.483	Mentha
-	B01.650.940.800.575.100.583.520.483.444	Mentha piperita
-	B01.650.940.800.575.100.583.520.483.500	Mentha pulegium
-	B01.650.940.800.575.100.583.520.483.750	Mentha spicata
-	B01.650.940.800.575.100.583.520.510	Monarda
-	B01.650.940.800.575.100.583.520.538	Nepeta
-	B01.650.940.800.575.100.583.520.647	Ocimum
-	B01.650.940.800.575.100.583.520.647.500	Ocimum basilicum
New Heading	<b>B01.650.940.800.575.100.583.520.647.750</b>	<b>Ocimum sanctum</b>
-	B01.650.940.800.575.100.583.520.702	Origanum
-	B01.650.940.800.575.100.583.520.729	Orthosiphon
-	B01.650.940.800.575.100.583.520.743	Perilla
-	B01.650.940.800.575.100.583.520.743.500	Perilla frutescens
-	B01.650.940.800.575.100.583.520.750	Phlomis
-	B01.650.940.800.575.100.583.520.753	Plectranthus
New Heading	<b>B01.650.940.800.575.100.583.520.754</b>	<b>Pogostemon</b>
-	B01.650.940.800.575.100.583.520.755	Prunella
-	B01.650.940.800.575.100.583.520.757	Rosmarinus
-	B01.650.940.800.575.100.583.520.922	Salvia
-	B01.650.940.800.575.100.583.520.922.750	Salvia miltiorrhiza
-	B01.650.940.800.575.100.583.520.922.770	Salvia officinalis
-	B01.650.940.800.575.100.583.520.935	Satureja
-	B01.650.940.800.575.100.583.520.942	Scutellaria
-	B01.650.940.800.575.100.583.520.942.500	Scutellaria baicalensis
-	B01.650.940.800.575.100.583.520.945	Sideritis
-	B01.650.940.800.575.100.583.520.947	Stachys
-	B01.650.940.800.575.100.583.520.948	Teucrium
-	B01.650.940.800.575.100.583.520.974	Thymus Plant
-	B01.650.940.800.575.100.583.520.987	Vitex
-	B01.650.940.800.575.100.583.580	Mimulus

## MeSH Tree Changes for 2017

Type	Tree - heading
-	B01.650.940.800.575.100.583.640 Oleaceae
-	B01.650.940.800.575.100.583.640.333 Forsythia
-	B01.650.940.800.575.100.583.640.374 Fraxinus
-	B01.650.940.800.575.100.583.640.416 Jasminum
-	B01.650.940.800.575.100.583.640.499 Ligustrum
-	B01.650.940.800.575.100.583.640.666 Olea
-	B01.650.940.800.575.100.583.640.833 Syringa
-	B01.650.940.800.575.100.583.670 Orobanchaceae
-	B01.650.940.800.575.100.583.670.175 Cistanche
-	B01.650.940.800.575.100.583.670.270 Euphrasia
-	B01.650.940.800.575.100.583.670.670 Orobanche
-	B01.650.940.800.575.100.583.670.690 Pedicularis
-	B01.650.940.800.575.100.583.670.875 Striga
-	B01.650.940.800.575.100.583.690 Pedaliaceae
-	B01.650.940.800.575.100.583.690.500 Harpagophytum
-	B01.650.940.800.575.100.583.690.750 Sesamum
-	B01.650.940.800.575.100.583.700 Plantaginaceae
-	B01.650.940.800.575.100.583.700.067 Antirrhinum
-	B01.650.940.800.575.100.583.700.080 Bacopa
-	B01.650.940.800.575.100.583.700.265 Digitalis
-	B01.650.940.800.575.100.583.700.530 Linaria
-	B01.650.940.800.575.100.583.700.680 Penstemon
-	B01.650.940.800.575.100.583.700.690 Picrorhiza
-	B01.650.940.800.575.100.583.700.700 Plantago
-	B01.650.940.800.575.100.583.700.850 Scoparia
-	B01.650.940.800.575.100.583.700.925 Veronica
-	B01.650.940.800.575.100.583.750 Rehmannia
-	B01.650.940.800.575.100.583.800 Scrophulariaceae
-	B01.650.940.800.575.100.583.800.124 Buddleja
-	B01.650.940.800.575.100.583.800.855 Scrophularia
-	B01.650.940.800.575.100.583.800.899 Verbascum
-	B01.650.940.800.575.100.583.990 Verbenaceae
-	B01.650.940.800.575.100.583.990.411 Lantana
-	B01.650.940.800.575.100.583.990.433 Lippia
-	B01.650.940.800.575.100.583.990.855 Verbena
-	B01.650.940.800.575.100.590 Lauraceae

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	B01.650.940.800.575.100.590.149	Cinnamomum
-	B01.650.940.800.575.100.590.149.124	Cinnamomum aromaticum
-	B01.650.940.800.575.100.590.149.249	Cinnamomum camphora
-	B01.650.940.800.575.100.590.149.500	Cinnamomum zeylanicum
-	B01.650.940.800.575.100.590.177	Cryptocarya
-	B01.650.940.800.575.100.590.399	Laurus
-	B01.650.940.800.575.100.590.524	Lindera
-	B01.650.940.800.575.100.590.555	Litsea
-	B01.650.940.800.575.100.590.587	Ocotea
-	B01.650.940.800.575.100.590.650	Persea
-	B01.650.940.800.575.100.590.737	Sassafras
-	B01.650.940.800.575.100.590.825	Umbellularia
-	B01.650.940.800.575.100.600	Lecythidaceae
-	B01.650.940.800.575.100.600.099	Barringtonia
-	B01.650.940.800.575.100.600.133	Bertholletia
Old Tree	B01.650.940.800.575.100.610	Liliaceae
Old Tree	B01.650.940.800.575.100.610.100	Allium
Old Tree	B01.650.940.800.575.100.610.100.149	Chive
Old Tree	B01.650.940.800.575.100.610.100.300	Garlic
Old Tree	B01.650.940.800.575.100.610.100.600	Onions
Old Tree	B01.650.940.800.575.100.610.100.800	Shallots
Old Tree	B01.650.940.800.575.100.610.111	Alstroemeria
Old Tree	B01.650.940.800.575.100.610.117	Anemarrhena
Old Tree	B01.650.940.800.575.100.610.162	Asparagus Plant
Old Tree	B01.650.940.800.575.100.610.181	Camassia
Old Tree	B01.650.940.800.575.100.610.200	Colchicum
Old Tree	B01.650.940.800.575.100.610.211	Convallaria
Old Tree	B01.650.940.800.575.100.610.222	Cordyline
Old Tree	B01.650.940.800.575.100.610.233	Crinum
Old Tree	B01.650.940.800.575.100.610.255	Curculigo
Old Tree	B01.650.940.800.575.100.610.277	Dracaena
Old Tree	B01.650.940.800.575.100.610.313	Fritillaria
Old Tree	B01.650.940.800.575.100.610.350	Galanthus
Old Tree	B01.650.940.800.575.100.610.387	Hemerocallis
Old Tree	B01.650.940.800.575.100.610.406	Hosta
Old Tree	B01.650.940.800.575.100.610.415	Hyacinthus

## MeSH Tree Changes for 2017

Type	Tree - heading	
Old Tree	<a href="#">B01.650.940.800.575.100.610.425</a>	Hypoxis
Old Tree	<a href="#">B01.650.940.800.575.100.610.500</a>	Lilium
Old Tree	<a href="#">B01.650.940.800.575.100.610.509</a>	Liriope Plant
Old Tree	<a href="#">B01.650.940.800.575.100.610.533</a>	Lycoris
Old Tree	<a href="#">B01.650.940.800.575.100.610.555</a>	Narcissus
Old Tree	<a href="#">B01.650.940.800.575.100.610.588</a>	Ophiopogon
Old Tree	<a href="#">B01.650.940.800.575.100.610.593</a>	Ornithogalum
Old Tree	<a href="#">B01.650.940.800.575.100.610.688</a>	Polygonatum
Old Tree	<a href="#">B01.650.940.800.575.100.610.716</a>	Ruscus
Old Tree	<a href="#">B01.650.940.800.575.100.610.730</a>	Sansevieria
Old Tree	<a href="#">B01.650.940.800.575.100.610.737</a>	Scilla
Old Tree	<a href="#">B01.650.940.800.575.100.610.744</a>	Maianthemum
Old Tree	<a href="#">B01.650.940.800.575.100.610.744</a>	Smilacina
Old Tree	<a href="#">B01.650.940.800.575.100.610.772</a>	Trillium
Old Tree	<a href="#">B01.650.940.800.575.100.610.786</a>	Tulipa
Old Tree	<a href="#">B01.650.940.800.575.100.610.793</a>	Drimia
Old Tree	<a href="#">B01.650.940.800.575.100.610.793</a>	Urginea
Old Tree	<a href="#">B01.650.940.800.575.100.610.800</a>	Veratrum
Old Tree	<a href="#">B01.650.940.800.575.100.610.900</a>	Zigadenus
New Heading	<b><a href="#">B01.650.940.800.575.100.615</a></b>	<b>Liliales</b>
New Tree	<a href="#">B01.650.940.800.575.100.615.088</a>	Alstroemeria
New Heading	<b><a href="#">B01.650.940.800.575.100.615.175</a></b>	<b>Colchicaceae</b>
New Tree	<a href="#">B01.650.940.800.575.100.615.175.170</a>	Colchicum
New Tree	<a href="#">B01.650.940.800.575.100.615.180</a>	Corsiaceae
New Tree	<a href="#">B01.650.940.800.575.100.615.500</a>	Liliaceae
New Tree	<a href="#">B01.650.940.800.575.100.615.500.313</a>	Fritillaria
New Tree	<a href="#">B01.650.940.800.575.100.615.500.500</a>	Lilium
New Tree	<a href="#">B01.650.940.800.575.100.615.500.593</a>	Ornithogalum
New Tree	<a href="#">B01.650.940.800.575.100.615.500.786</a>	Tulipa

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Heading	<b>B01.650.940.800.575.100.615.625</b>	<b>Melanthiaceae</b>
New Tree	B01.650.940.800.575.100.615.625.860	Trillium
New Tree	B01.650.940.800.575.100.615.625.920	Veratrum
New Tree	B01.650.940.800.575.100.615.625.960	Zigadenus
New Tree	B01.650.940.800.575.100.615.750	Smilacaceae
New Tree	B01.650.940.800.575.100.615.750.500	Smilax
-	B01.650.940.800.575.100.620	Linaceae
-	B01.650.940.800.575.100.620.500	Flax
Old Tree	<b>B01.650.940.800.575.100.638</b>	<b>Loganiaceae</b>
Old Tree	<b>B01.650.940.800.575.100.638.500</b>	<b>Gelsemium</b>
Old Tree	<b>B01.650.940.800.575.100.638.750</b>	<b>Strychnos</b>
Old Tree	<b>B01.650.940.800.575.100.638.750.500</b>	<b>Strychnos nux-vomica</b>
-	B01.650.940.800.575.100.665	Lythraceae
-	B01.650.940.800.575.100.665.249	Cuphea
-	B01.650.940.800.575.100.665.374	Lagerstroemia
-	B01.650.940.800.575.100.665.437	Lawsonia Plant
-	B01.650.940.800.575.100.665.500	Lythrum
-	B01.650.940.800.575.100.665.750	Woodfordia
-	B01.650.940.800.575.100.680	Magnoliaceae
-	B01.650.940.800.575.100.680.249	Liriodendron
-	B01.650.940.800.575.100.680.500	Magnolia
-	B01.650.940.800.575.100.690	Malpighiaceae
-	B01.650.940.800.575.100.690.099	Banisteriopsis
-	B01.650.940.800.575.100.690.355	Galphimia
-	B01.650.940.800.575.100.700	Malvaceae
-	B01.650.940.800.575.100.700.100	Abelmoschus
-	B01.650.940.800.575.100.700.115	Althaea
-	B01.650.940.800.575.100.700.244	Gossypium
-	B01.650.940.800.575.100.700.288	Hibiscus
-	B01.650.940.800.575.100.700.555	Malva
-	B01.650.940.800.575.100.707	Melastomataceae
-	B01.650.940.800.575.100.715	Meliaceae

## MeSH Tree Changes for 2017

Type	Tree - heading
-	B01.650.940.800.575.100.715.061 Aglaia
-	B01.650.940.800.575.100.715.124 Azadirachta
-	B01.650.940.800.575.100.715.249 Cedrela
-	B01.650.940.800.575.100.715.500 Melia
-	B01.650.940.800.575.100.715.500.500 Melia azedarach
-	B01.650.940.800.575.100.725 Menispermaceae
-	B01.650.940.800.575.100.725.111 Cissampelos
-	B01.650.940.800.575.100.725.124 Cocculus
-	B01.650.940.800.575.100.725.249 Cyclea
-	B01.650.940.800.575.100.725.500 Menispermum
-	B01.650.940.800.575.100.725.625 Sinomenium
-	B01.650.940.800.575.100.725.750 Stephania
-	B01.650.940.800.575.100.725.750.500 Stephania tetrandra
-	B01.650.940.800.575.100.725.875 Tinospora
-	B01.650.940.800.575.100.735 Mistletoe
-	B01.650.940.800.575.100.735.249 Loranthaceae
-	B01.650.940.800.575.100.735.888 Viscaceae
-	B01.650.940.800.575.100.735.888.699 Phoradendron
-	B01.650.940.800.575.100.735.888.750 Viscum
-	B01.650.940.800.575.100.735.888.750.500 Viscum album
-	B01.650.940.800.575.100.740 Molluginaceae
-	B01.650.940.800.575.100.745 Monimiaceae
-	B01.650.940.800.575.100.745.500 Peumus
-	B01.650.940.800.575.100.750 Moraceae
-	B01.650.940.800.575.100.750.055 Antiaris
-	B01.650.940.800.575.100.750.088 Artocarpus
-	B01.650.940.800.575.100.750.144 Broussonetia
-	B01.650.940.800.575.100.750.500 Ficus
-	B01.650.940.800.575.100.750.599 Maclura
-	B01.650.940.800.575.100.750.633 Morus
-	B01.650.940.800.575.100.752 Moringa
-	B01.650.940.800.575.100.752.500 Moringa oleifera
-	B01.650.940.800.575.100.756 Myoporaceae
-	B01.650.940.800.575.100.756.249 Eremophila Plant
-	B01.650.940.800.575.100.756.500 Myoporum
-	B01.650.940.800.575.100.760 Myricaceae

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	B01.650.940.800.575.100.760.500	Myrica
-	B01.650.940.800.575.100.765	Myristicaceae
-	B01.650.940.800.575.100.765.500	Myristica fragrans
-	B01.650.940.800.575.100.770	Myrsinaceae
-	B01.650.940.800.575.100.770.066	Ardisia
-	B01.650.940.800.575.100.770.533	Embelia
-	B01.650.940.800.575.100.773	Myrtaceae
-	B01.650.940.800.575.100.773.366	Eucalyptus
-	B01.650.940.800.575.100.773.374	Eugenia
-	B01.650.940.800.575.100.773.522	Feijoa
-	B01.650.940.800.575.100.773.649	Kunzea
-	B01.650.940.800.575.100.773.713	Leptospermum
-	B01.650.940.800.575.100.773.745	Melaleuca
-	B01.650.940.800.575.100.773.761	Myrtus
-	B01.650.940.800.575.100.773.777	Pimenta
-	B01.650.940.800.575.100.773.888	Psidium
-	B01.650.940.800.575.100.773.900	Syzygium
-	B01.650.940.800.575.100.774	Nelumbonaceae
-	B01.650.940.800.575.100.774.500	Nelumbo
-	B01.650.940.800.575.100.775	Nyctaginaceae
-	B01.650.940.800.575.100.775.500	Mirabilis
-	B01.650.940.800.575.100.776	Nymphaeaceae
-	B01.650.940.800.575.100.776.488	Nuphar
-	B01.650.940.800.575.100.776.500	Nymphaea
-	B01.650.940.800.575.100.778	Nyssaceae
-	B01.650.940.800.575.100.778.188	Camptotheca
-	B01.650.940.800.575.100.778.594	Nyssa
-	B01.650.940.800.575.100.779	Ochnaceae
-	B01.650.940.800.575.100.780	Olacaceae
-	B01.650.940.800.575.100.786	Onagraceae
-	B01.650.940.800.575.100.786.222	Clarkia
-	B01.650.940.800.575.100.786.416	Epilobium
-	B01.650.940.800.575.100.786.611	Oenothera
-	B01.650.940.800.575.100.786.611.500	Oenothera biennis
Old Tree	B01.650.940.800.575.100.787	Orchidaceae
Old Tree	B01.650.940.800.575.100.787.222	Dendrobium

## MeSH Tree Changes for 2017

Type	Tree - heading	
Old Tree	B01.650.940.800.575.100.787.500	Gastrodia
Old Tree	B01.650.940.800.575.100.787.888	Vanilla
-	B01.650.940.800.575.100.789	Oxalidaceae
-	B01.650.940.800.575.100.789.500	Averrhoa
-	B01.650.940.800.575.100.791	Paeonia
-	B01.650.940.800.575.100.792	Pandanaceae
-	B01.650.940.800.575.100.795	Papaveraceae
-	B01.650.940.800.575.100.795.124	Argemone
-	B01.650.940.800.575.100.795.249	Chelidonium
-	B01.650.940.800.575.100.795.374	Eschscholzia
-	B01.650.940.800.575.100.795.500	Papaver
-	B01.650.940.800.575.100.795.750	Sanguinaria
-	B01.650.940.800.575.100.803	Passifloraceae
-	B01.650.940.800.575.100.803.500	Passiflora
New Tree	B01.650.940.800.575.100.803.750	Turnera
-	B01.650.940.800.575.100.809	Phytolaccaceae
-	B01.650.940.800.575.100.809.500	Phytolacca
-	B01.650.940.800.575.100.809.500.500	Phytolacca americana
-	B01.650.940.800.575.100.809.500.750	Phytolacca dodecandra
-	B01.650.940.800.575.100.812	Piperaceae
-	B01.650.940.800.575.100.812.332	Peperomia
-	B01.650.940.800.575.100.812.666	Piper
-	B01.650.940.800.575.100.812.666.249	Kava
-	B01.650.940.800.575.100.812.666.374	Piper betle
-	B01.650.940.800.575.100.812.666.500	Piper nigrum
-	B01.650.940.800.575.100.821	Plumbaginaceae
-	B01.650.940.800.575.100.822	Poaceae
-	B01.650.940.800.575.100.822.033	Agropyron
-	B01.650.940.800.575.100.822.044	Agrostis
-	B01.650.940.800.575.100.822.055	Andropogon
-	B01.650.940.800.575.100.822.060	Avena
-	B01.650.940.800.575.100.822.066	Bambusa
-	B01.650.940.800.575.100.822.071	Brachiaria
-	B01.650.940.800.575.100.822.072	Brachypodium
-	B01.650.940.800.575.100.822.077	Bromus



## MeSH Tree Changes for 2017

Type	Tree - heading
-	B01.650.940.800.575.100.822.151 Cenchrus
-	B01.650.940.800.575.100.822.227 Coix
-	B01.650.940.800.575.100.822.266 Cymbopogon
-	B01.650.940.800.575.100.822.288 Cynodon
-	B01.650.940.800.575.100.822.299 Dactylis
-	B01.650.940.800.575.100.822.311 Digitaria
-	B01.650.940.800.575.100.822.322 Echinochloa
-	B01.650.940.800.575.100.822.333 Eleusine
-	B01.650.940.800.575.100.822.344 Elymus
-	B01.650.940.800.575.100.822.355 Eragrostis
-	B01.650.940.800.575.100.822.427 Festuca
-	B01.650.940.800.575.100.822.463 Holcus
-	B01.650.940.800.575.100.822.481 Hordeum
-	B01.650.940.800.575.100.822.500 Lolium
-	B01.650.940.800.575.100.822.558 Melinis repens
-	B01.650.940.800.575.100.822.587 Millets
-	B01.650.940.800.575.100.822.616 Oryza
-	B01.650.940.800.575.100.822.680 Panicum
-	B01.650.940.800.575.100.822.744 Paspalum
-	B01.650.940.800.575.100.822.755 Pennisetum
-	B01.650.940.800.575.100.822.777 Phalaris
-	B01.650.940.800.575.100.822.788 Phleum
-	B01.650.940.800.575.100.822.793 Poa
-	B01.650.940.800.575.100.822.835 Saccharum
-	B01.650.940.800.575.100.822.844 Sasa
-	B01.650.940.800.575.100.822.857 Secale
-	B01.650.940.800.575.100.822.871 Setaria Plant
-	B01.650.940.800.575.100.822.894 Sorghum
-	B01.650.940.800.575.100.822.906 Triticale
-	B01.650.940.800.575.100.822.918 Triticum
-	B01.650.940.800.575.100.822.942 Vetiveria
-	B01.650.940.800.575.100.822.966 Zea mays
-	B01.650.940.800.575.100.823 Polygalaceae
-	B01.650.940.800.575.100.823.500 Polygala
-	B01.650.940.800.575.100.823.750 Securidaca
-	B01.650.940.800.575.100.825 Polygonaceae

## MeSH Tree Changes for 2017

Type	Tree - heading
-	B01.650.940.800.575.100.825.306 Eriogonum
-	B01.650.940.800.575.100.825.333 Fagopyrum
-	B01.650.940.800.575.100.825.359 Fallopia
-	B01.650.940.800.575.100.825.359.500 Fallopia japonica
-	B01.650.940.800.575.100.825.359.590 Fallopia multiflora
-	B01.650.940.800.575.100.825.437 Polygonum
-	B01.650.940.800.575.100.825.700 Rheum
-	B01.650.940.800.575.100.825.850 Rumex
-	B01.650.940.800.575.100.826 Pontederiaceae
-	B01.650.940.800.575.100.826.500 Eichhornia
-	B01.650.940.800.575.100.827 Portulacaceae
-	B01.650.940.800.575.100.827.500 Portulaca
-	B01.650.940.800.575.100.829 Primulaceae
-	B01.650.940.800.575.100.829.099 Anagallis
-	B01.650.940.800.575.100.829.549 Cyclamen
-	B01.650.940.800.575.100.829.774 Primula
-	B01.650.940.800.575.100.831 Proteaceae
-	B01.650.940.800.575.100.831.500 Macadamia
-	B01.650.940.800.575.100.833 Punicaceae
-	B01.650.940.800.575.100.835 Pyrolaceae
-	B01.650.940.800.575.100.835.500 Pyrola
-	B01.650.940.800.575.100.838 Ranunculaceae
-	B01.650.940.800.575.100.838.022 Aconitum
-	B01.650.940.800.575.100.838.033 Actaea
-	B01.650.940.800.575.100.838.036 Adonis
-	B01.650.940.800.575.100.838.077 Anemone
-	B01.650.940.800.575.100.838.088 Aquilegia
-	B01.650.940.800.575.100.838.182 Cimicifuga
-	B01.650.940.800.575.100.838.205 Clematis
-	B01.650.940.800.575.100.838.229 Coptis
-	B01.650.940.800.575.100.838.277 Delphinium
-	B01.650.940.800.575.100.838.337 Helleborus
-	B01.650.940.800.575.100.838.397 Hydrastis
-	B01.650.940.800.575.100.838.518 Nigella
-	B01.650.940.800.575.100.838.518.500 Nigella damascena
-	B01.650.940.800.575.100.838.518.750 Nigella sativa

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	B01.650.940.800.575.100.838.666	Pulsatilla
-	B01.650.940.800.575.100.838.733	Ranunculus
-	B01.650.940.800.575.100.838.766	Semiaquilegia
-	B01.650.940.800.575.100.838.883	Thalictrum
-	B01.650.940.800.575.100.838.941	Xanthorhiza
-	B01.650.940.800.575.100.839	Resedaceae
Old Tree	B01.650.940.800.575.100.840	Rhamnaceae
Old Tree	B01.650.940.800.575.100.840.111	Ceanothus
Old Tree	B01.650.940.800.575.100.840.133	Colubrina
Old Tree	B01.650.940.800.575.100.840.316	Frangula
Old Tree	B01.650.940.800.575.100.840.408	Karwinskia
Old Tree	B01.650.940.800.575.100.840.666	Rhamnus
Old Tree	B01.650.940.800.575.100.840.833	Ziziphus
-	B01.650.940.800.575.100.841	Rhizophoraceae
-	B01.650.940.800.575.100.842	Rosales
-	B01.650.940.800.575.100.842.110	Chrysobalanaceae
-	B01.650.940.800.575.100.842.222	Connaraceae
-	B01.650.940.800.575.100.842.249	Crassulaceae
-	B01.650.940.800.575.100.842.249.249	Kalanchoe
-	B01.650.940.800.575.100.842.249.500	Rhodiola
-	B01.650.940.800.575.100.842.249.750	Sedum
-	B01.650.940.800.575.100.842.311	Grossulariaceae
-	B01.650.940.800.575.100.842.311.500	Ribes
-	B01.650.940.800.575.100.842.374	Hydrangeaceae
-	B01.650.940.800.575.100.842.374.500	Hydrangea
New Tree	B01.650.940.800.575.100.842.437	Rhamnaceae
New Tree	B01.650.940.800.575.100.842.437.111	Ceanothus
New Tree	B01.650.940.800.575.100.842.437.133	Colubrina
New Tree	B01.650.940.800.575.100.842.437.316	Frangula
New Tree	B01.650.940.800.575.100.842.437.408	Karwinskia
New Tree	B01.650.940.800.575.100.842.437.666	Rhamnus
New	B01.650.940.800.575.100.842.437.833	Ziziphus

## MeSH Tree Changes for 2017

Type	Tree - heading	
Tree		
-	B01.650.940.800.575.100.842.500	Rosaceae
-	B01.650.940.800.575.100.842.500.033	Agrimonia
-	B01.650.940.800.575.100.842.500.055	Alchemilla
-	B01.650.940.800.575.100.842.500.177	Crataegus
-	B01.650.940.800.575.100.842.500.222	Eriobotrya
-	B01.650.940.800.575.100.842.500.244	Filipendula
-	B01.650.940.800.575.100.842.500.266	Fragaria
-	B01.650.940.800.575.100.842.500.288	Geum
-	B01.650.940.800.575.100.842.500.444	Malus
-	B01.650.940.800.575.100.842.500.522	Photinia
-	B01.650.940.800.575.100.842.500.601	Potentilla
-	B01.650.940.800.575.100.842.500.625	Prunus
-	B01.650.940.800.575.100.842.500.625.250	Prunus africana
-	B01.650.940.800.575.100.842.500.625.375	Prunus armeniaca
-	B01.650.940.800.575.100.842.500.625.400	Prunus avium
-	B01.650.940.800.575.100.842.500.625.450	Prunus domestica
-	B01.650.940.800.575.100.842.500.625.500	Prunus dulcis
-	B01.650.940.800.575.100.842.500.625.750	Prunus persica
-	B01.650.940.800.575.100.842.500.693	Pyracantha
-	B01.650.940.800.575.100.842.500.699	Pyrus
-	B01.650.940.800.575.100.842.500.724	Quillaja
-	B01.650.940.800.575.100.842.500.750	Rosa
-	B01.650.940.800.575.100.842.500.764	Rubus
-	B01.650.940.800.575.100.842.500.779	Sanguisorba
-	B01.650.940.800.575.100.842.500.793	Sorbus
-	B01.650.940.800.575.100.842.500.808	Spiraea
-	B01.650.940.800.575.100.842.750	Saxifragaceae
-	B01.650.940.800.575.100.842.750.500	Heuchera
Old Tree	<b>B01.650.940.800.575.100.850</b>	<b>Rubiaceae</b>
Old Tree	<b>B01.650.940.800.575.100.850.200</b>	<b>Cephaelis</b>
Old Tree	<b>B01.650.940.800.575.100.850.250</b>	<b>Cinchona</b>
Old Tree	<b>B01.650.940.800.575.100.850.275</b>	<b>Coffea</b>
Old Tree	<b>B01.650.940.800.575.100.850.377</b>	<b>Galium</b>
Old Tree	<b>B01.650.940.800.575.100.850.388</b>	<b>Gardenia</b>
Old Tree	<b>B01.650.940.800.575.100.850.404</b>	<b>Hamelia</b>

## MeSH Tree Changes for 2017

Type	Tree - heading	
Old Tree	B01.650.940.800.575.100.850.422	Hedyotis
Old Tree	B01.650.940.800.575.100.850.555	Mitragyna
Old Tree	B01.650.940.800.575.100.850.566	Morinda
Old Tree	B01.650.940.800.575.100.850.611	Oldenlandia
Old Tree	B01.650.940.800.575.100.850.805	Pausinystalia
Old Tree	B01.650.940.800.575.100.850.844	Psychotria
Old Tree	B01.650.940.800.575.100.850.876	Rubia
Old Tree	B01.650.940.800.575.100.850.909	Uncaria
Old Tree	B01.650.940.800.575.100.850.909.500	Cat's Claw
-	B01.650.940.800.575.100.875	Rutaceae
-	B01.650.940.800.575.100.875.077	Aegle
-	B01.650.940.800.575.100.875.155	Casimiroa
-	B01.650.940.800.575.100.875.177	Citrus
-	B01.650.940.800.575.100.875.177.077	Citrus aurantiifolia
-	B01.650.940.800.575.100.875.177.538	Citrus paradisi
-	B01.650.940.800.575.100.875.177.769	Citrus sinensis
-	B01.650.940.800.575.100.875.216	Clausena
-	B01.650.940.800.575.100.875.255	Dictamnus
-	B01.650.940.800.575.100.875.333	Evodia
-	B01.650.940.800.575.100.875.666	Murraya
-	B01.650.940.800.575.100.875.717	Phellodendron
-	B01.650.940.800.575.100.875.733	Pilocarpus
-	B01.650.940.800.575.100.875.755	Poncirus
-	B01.650.940.800.575.100.875.788	Ruta
-	B01.650.940.800.575.100.875.922	Zanthoxylum
-	B01.650.940.800.575.100.881	Salicaceae
-	B01.650.940.800.575.100.881.666	Populus
-	B01.650.940.800.575.100.881.833	Salix
-	B01.650.940.800.575.100.882	Salvadoraceae
-	B01.650.940.800.575.100.884	Santalaceae
-	B01.650.940.800.575.100.884.666	Pyrularia
-	B01.650.940.800.575.100.884.833	Santalum
-	B01.650.940.800.575.100.885	Sapindaceae
-	B01.650.940.800.575.100.885.111	Blighia
-	B01.650.940.800.575.100.885.444	Litchi
-	B01.650.940.800.575.100.885.722	Paullinia

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	B01.650.940.800.575.100.885.861	Sapindus
-	B01.650.940.800.575.100.886	Sapotaceae
-	B01.650.940.800.575.100.886.500	Madhuca
-	B01.650.940.800.575.100.886.555	Manilkara
-	B01.650.940.800.575.100.886.577	Mimusops
-	B01.650.940.800.575.100.886.666	Palaquium
-	B01.650.940.800.575.100.886.699	Pouteria
-	B01.650.940.800.575.100.886.849	Synsepalum
-	B01.650.940.800.575.100.887	Sarraceniaceae
-	B01.650.940.800.575.100.888	Saururaceae
-	B01.650.940.800.575.100.888.500	Houttuynia
-	B01.650.940.800.575.100.889	Schisandraceae
-	B01.650.940.800.575.100.889.500	Kadsura
-	B01.650.940.800.575.100.889.750	Schisandra
-	B01.650.940.800.575.100.898	Simaroubaceae
-	B01.650.940.800.575.100.898.044	Ailanthus
-	B01.650.940.800.575.100.898.111	Brucea
-	B01.650.940.800.575.100.898.333	Eurycoma
-	B01.650.940.800.575.100.898.555	Picrasma
-	B01.650.940.800.575.100.898.599	Quassia
-	B01.650.940.800.575.100.898.799	Simarouba
Old Tree	B01.650.940.800.575.100.901	Smilacaceae
Old Tree	B01.650.940.800.575.100.901.500	Smilax
-	B01.650.940.800.575.100.905	Solanaceae
-	B01.650.940.800.575.100.905.049	Atropa
-	B01.650.940.800.575.100.905.049.222	Atropa belladonna
-	B01.650.940.800.575.100.905.145	Capsicum
-	B01.650.940.800.575.100.905.166	Cestrum
-	B01.650.940.800.575.100.905.188	Datura
-	B01.650.940.800.575.100.905.188.550	Datura metel
-	B01.650.940.800.575.100.905.188.800	Datura stramonium
-	B01.650.940.800.575.100.905.202	Duboisia
-	B01.650.940.800.575.100.905.274	Hyoscyamus
-	B01.650.940.800.575.100.905.310	Lycium
-	B01.650.940.800.575.100.905.322	Lycopersicon esculentum
-	B01.650.940.800.575.100.905.347	Mandragora

## MeSH Tree Changes for 2017

Type	Tree - heading
-	B01.650.940.800.575.100.905.448 Petunia
-	B01.650.940.800.575.100.905.499 Physalis
-	B01.650.940.800.575.100.905.612 Scopolia
-	B01.650.940.800.575.100.905.725 Solanum
-	B01.650.940.800.575.100.905.725.610 Solanum glaucophyllum
-	B01.650.940.800.575.100.905.725.666 Solanum melongena
-	B01.650.940.800.575.100.905.725.699 Solanum nigrum
-	B01.650.940.800.575.100.905.725.777 Solanum tuberosum
-	B01.650.940.800.575.100.905.900 Tobacco
-	B01.650.940.800.575.100.905.950 Withania
-	B01.650.940.800.575.100.910 Stemonaceae
-	B01.650.940.800.575.100.915 Sterculiaceae
-	B01.650.940.800.575.100.915.144 Cacao
-	B01.650.940.800.575.100.915.166 Cola
-	B01.650.940.800.575.100.915.583 Sterculia
-	B01.650.940.800.575.100.920 Styracaceae
-	B01.650.940.800.575.100.920.777 Styrax
-	B01.650.940.800.575.100.924 Tamaricaceae
-	B01.650.940.800.575.100.928 Theales
-	B01.650.940.800.575.100.928.124 Actinidiaceae
-	B01.650.940.800.575.100.928.124.500 Actinidia
-	B01.650.940.800.575.100.928.249 Clusiaceae
-	B01.650.940.800.575.100.928.249.166 Calophyllum
-	B01.650.940.800.575.100.928.249.249 Clusia
-	B01.650.940.800.575.100.928.249.333 Garcinia
-	B01.650.940.800.575.100.928.249.333.200 Garcinia cambogia
-	B01.650.940.800.575.100.928.249.333.400 Garcinia kola
-	B01.650.940.800.575.100.928.249.333.600 Garcinia mangostana
-	B01.650.940.800.575.100.928.249.500 Hypericum
-	B01.650.940.800.575.100.928.249.750 Mammea
-	B01.650.940.800.575.100.928.374 Dipterocarpaceae
-	B01.650.940.800.575.100.928.500 Theaceae
-	B01.650.940.800.575.100.928.500.500 Camellia
-	B01.650.940.800.575.100.928.500.500.500 Camellia sinensis
-	B01.650.940.800.575.100.932 Thymelaeaceae
-	B01.650.940.800.575.100.932.500 Daphne

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	B01.650.940.800.575.100.932.909	Wikstroemia
-	B01.650.940.800.575.100.938	Tiliaceae
-	B01.650.940.800.575.100.938.144	Corchorus
-	B01.650.940.800.575.100.938.333	Grewia
-	B01.650.940.800.575.100.938.777	Tilia
-	B01.650.940.800.575.100.938.888	Triumfetta
-	B01.650.940.800.575.100.942	Tropaeolaceae
-	B01.650.940.800.575.100.942.500	Tropaeolum
Old Tree	B01.650.940.800.575.100.947	Turnera
-	B01.650.940.800.575.100.948	Typhaceae
-	B01.650.940.800.575.100.949	Ulmaceae
-	B01.650.940.800.575.100.949.777	Trema
-	B01.650.940.800.575.100.949.888	Ulmus
-	B01.650.940.800.575.100.950	Urticaceae
-	B01.650.940.800.575.100.950.077	Boehmeria
-	B01.650.940.800.575.100.950.777	Parietaria
-	B01.650.940.800.575.100.950.888	Urtica dioica
-	B01.650.940.800.575.100.955	Valerianaceae
-	B01.650.940.800.575.100.955.443	Nardostachys
-	B01.650.940.800.575.100.955.665	Patrinia
-	B01.650.940.800.575.100.955.888	Valerian
-	B01.650.940.800.575.100.955.944	Valerianella
-	B01.650.940.800.575.100.962	Violaceae
-	B01.650.940.800.575.100.962.888	Viola
-	B01.650.940.800.575.100.965	Vitaceae
-	B01.650.940.800.575.100.965.044	Ampelopsis
-	B01.650.940.800.575.100.965.272	Cissus
-	B01.650.940.800.575.100.965.500	Vitis
-	B01.650.940.800.575.100.970	Winteraceae
-	B01.650.940.800.575.100.970.199	Drimys
-	B01.650.940.800.575.100.970.599	Pseudowintera
-	B01.650.940.800.575.100.975	Zingiberales
-	B01.650.940.800.575.100.975.124	Costus
-	B01.650.940.800.575.100.975.323	Heliconiaceae
-	B01.650.940.800.575.100.975.522	Marantaceae
-	B01.650.940.800.575.100.975.555	Musaceae



## MeSH Tree Changes for 2017

Type	Tree - heading
-	B01.650.940.800.575.100.975.555.500 Musa
-	B01.650.940.800.575.100.975.727 Strelitziaceae
-	B01.650.940.800.575.100.975.900 Zingiberaceae
-	B01.650.940.800.575.100.975.900.033 Alpinia
-	B01.650.940.800.575.100.975.900.099 Amomum
-	B01.650.940.800.575.100.975.900.166 Curcuma
-	B01.650.940.800.575.100.975.900.333 Elettaria
-	B01.650.940.800.575.100.975.900.444 Ginger
-	B01.650.940.800.575.100.987 Zygophyllaceae
-	B01.650.940.800.575.100.987.088 Balanites
-	B01.650.940.800.575.100.987.277 Guaiacum
-	B01.650.940.800.575.100.987.399 Larrea
-	B01.650.940.800.575.100.987.699 Peganum
-	B01.650.940.800.575.100.987.849 Tribulus
-	B01.650.940.800.575.100.987.924 Zygophyllum
-	B01.650.940.800.575.118 Anthocerotophyta
-	B01.650.940.800.575.137 Bryophyta
-	B01.650.940.800.575.137.500 Bryopsida
-	B01.650.940.800.575.137.750 Sphagnopsida
-	B01.650.940.800.575.175 Equisetum
-	B01.650.940.800.575.400 Gymnosperms
-	B01.650.940.800.575.400.122 Coniferophyta
-	B01.650.940.800.575.400.122.166 Cephalotaxus
-	B01.650.940.800.575.400.122.188 Cupressaceae
-	B01.650.940.800.575.400.122.188.199 Chamaecyparis
-	B01.650.940.800.575.400.122.188.266 Cupressus
-	B01.650.940.800.575.400.122.188.500 Juniperus
-	B01.650.940.800.575.400.122.188.599 Libocedrus
-	B01.650.940.800.575.400.122.188.799 Thuja
-	B01.650.940.800.575.400.122.666 Pinaceae
-	B01.650.940.800.575.400.122.666.033 Abies
-	B01.650.940.800.575.400.122.666.222 Cedrus
-	B01.650.940.800.575.400.122.666.444 Larix
-	B01.650.940.800.575.400.122.666.755 Picea
-	B01.650.940.800.575.400.122.666.777 Pinus
-	B01.650.940.800.575.400.122.666.777.600 Pinus ponderosa

## MeSH Tree Changes for 2017

Type	Tree - heading
-	B01.650.940.800.575.400.122.666.777.744 <span style="float: right;">Pinus sylvestris</span>
-	B01.650.940.800.575.400.122.666.777.888 <span style="float: right;">Pinus taeda</span>
-	B01.650.940.800.575.400.122.666.984 <span style="float: right;">Pseudotsuga</span>
-	B01.650.940.800.575.400.122.666.992 <span style="float: right;">Tsuga</span>
-	B01.650.940.800.575.400.122.833 <span style="float: right;">Taxaceae</span>
-	B01.650.940.800.575.400.122.833.500 <span style="float: right;">Taxus</span>
-	B01.650.940.800.575.400.122.855 <span style="float: right;">Taxodiaceae</span>
-	B01.650.940.800.575.400.122.855.166 <span style="float: right;">Cryptomeria</span>
-	B01.650.940.800.575.400.122.855.177 <span style="float: right;">Cunninghamia</span>
-	B01.650.940.800.575.400.122.855.777 <span style="float: right;">Sequoia</span>
-	B01.650.940.800.575.400.122.855.788 <span style="float: right;">Sequoiadendron</span>
-	B01.650.940.800.575.400.122.855.833 <span style="float: right;">Taxodium</span>
-	B01.650.940.800.575.400.186 <span style="float: right;">Cycadophyta</span>
-	B01.650.940.800.575.400.186.199 <span style="float: right;">Cycas</span>
-	B01.650.940.800.575.400.186.955 <span style="float: right;">Zamiaceae</span>
-	B01.650.940.800.575.400.300 <span style="float: right;">Ginkgo biloba</span>
-	B01.650.940.800.575.400.344 <span style="float: right;">Gnetophyta</span>
-	B01.650.940.800.575.400.344.222 <span style="float: right;">Ephedra</span>
-	B01.650.940.800.575.400.344.222.500 <span style="float: right;">Ephedra sinica</span>
-	B01.650.940.800.575.400.344.333 <span style="float: right;">Gnetum</span>
-	B01.650.940.800.575.462 <span style="float: right;">Hepatophyta</span>
-	B01.650.940.800.575.462.500 <span style="float: right;">Frullania</span>
-	B01.650.940.800.575.462.750 <span style="float: right;">Marchantia</span>
-	B01.650.940.800.575.525 <span style="float: right;">Lycopodiaceae</span>
-	B01.650.940.800.575.525.333 <span style="float: right;">Huperzia</span>
-	B01.650.940.800.575.525.666 <span style="float: right;">Lycopodium</span>
-	B01.650.940.800.575.825 <span style="float: right;">Selaginellaceae</span>
-	B01.650.940.900 <span style="float: right;">Tracheobionta</span>
-	B01.650.940.900.500 <span style="float: right;">Pteridophyta</span>
-	B01.650.940.900.500.750 <span style="float: right;">Ferns</span>
-	B01.650.940.900.500.750.199 <span style="float: right;">Dennstaedtiaceae</span>
-	B01.650.940.900.500.750.199.666 <span style="float: right;">Pteridium</span>
-	B01.650.940.900.500.750.222 <span style="float: right;">Dryopteridaceae</span>
-	B01.650.940.900.500.750.222.222 <span style="float: right;">Dryopteris</span>
-	B01.650.940.900.500.750.222.611 <span style="float: right;">Polystichum</span>
-	B01.650.940.900.500.750.461 <span style="float: right;">Marsileaceae</span>

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	B01.650.940.900.500.750.700	Polypodiaceae
-	B01.650.940.900.500.750.700.666	Polypodium
-	B01.650.940.900.500.750.750	Pteridaceae
-	B01.650.940.900.500.750.750.044	Adiantum
-	B01.650.940.900.500.750.750.666	Pteris
-	B01.675	Retortamonadidae
-	B01.680	Rhizaria
-	B01.680.150	Cercozoa
-	B01.680.150.600	Plasmodiophorida
-	B01.680.275	Foraminifera
-	B01.680.400	Haplosporida
-	B01.750	Stramenopiles
-	B01.750.150	Chrysophyta
-	B01.750.150.600	Ochromonas
-	B01.750.600	Phaeophyta
-	B01.750.600.040	Ascophyllum
-	B01.750.600.212	Fucus
-	B01.750.600.425	Kelp
-	B01.750.600.450	Laminaria
-	B01.750.600.480	Macrocystis
-	B01.750.600.725	Sargassum
-	B01.750.600.800	Undaria
-	B01.750.875	Diatoms
-	B01.750.937	Oomycetes
-	B01.750.937.050	Achlya
-	B01.750.937.075	Aphanomyces
-	B01.750.937.500	Lagenidium
-	B01.750.937.710	Peronospora
-	B01.750.937.715	Phytophthora
-	B01.750.937.715.600	Phytophthora infestans
-	B01.750.937.750	Pythium
-	B01.750.937.825	Saprolegnia
-	B02	Archaea
Old Tree	B02.050	Blood-Borne Pathogens
-	B02.075	Crenarchaeota
-	B02.075.200	Desulfurococcales

## MeSH Tree Changes for 2017

Type	Tree - heading
-	B02.075.200.500 Desulfurococcaceae
-	B02.075.200.500.050 Aeropyrum
-	B02.075.200.650 Pyrodictiaceae
-	B02.075.725 Sulfolobales
-	B02.075.725.725 Sulfolobaceae
-	B02.075.725.725.030 Acidianus
-	B02.075.725.725.725 Sulfolobus
-	B02.075.725.725.725.070 Sulfolobus acidocaldarius
-	B02.075.725.725.725.800 Sulfolobus solfataricus
-	B02.075.800 Thermoproteales
-	B02.075.800.780 Thermofilaceae
-	B02.075.800.800 Thermoproteaceae
-	B02.075.800.800.650 Pyrobaculum
-	B02.075.800.800.800 Thermoproteus
-	B02.200 Euryarchaeota
-	B02.200.080 Archaeoglobales
-	B02.200.080.080 Archaeoglobus
-	B02.200.080.080.100 Archaeoglobus fulgidus
-	B02.200.400 Halobacteriales
-	B02.200.400.400 Halobacteriaceae
-	B02.200.400.400.400 Haloarcula
-	B02.200.400.400.400.410 Haloarcula marismortui
-	B02.200.400.400.410 Halobacterium
-	B02.200.400.400.410.450 Halobacterium salinarum
-	B02.200.400.400.420 Halococcus
-	B02.200.400.400.440 Haloferax
-	B02.200.400.400.440.500 Haloferax mediterranei
-	B02.200.400.400.440.900 Haloferax volcanii
-	B02.200.400.400.445 Halorubrum
-	B02.200.400.400.550 Natronobacterium
-	B02.200.400.400.560 Natronococcus
-	B02.200.492 Methanobacteriales
-	B02.200.492.500 Methanobacteriaceae
-	B02.200.492.500.500 Methanobacterium
-	B02.200.492.500.515 Methanobrevibacter
-	B02.200.538 Methanococcales

## MeSH Tree Changes for 2017

Type	Tree - heading
-	B02.200.538.262 Methanocaldococcaceae
-	B02.200.538.262.500 Methanocaldococcus
-	B02.200.538.525 Methanococcaceae
-	B02.200.538.525.500 Methanococcus
-	B02.200.705 Methanomicrobiales
-	B02.200.705.500 Methanomicrobiaceae
-	B02.200.705.750 Methanospirillum
-	B02.200.765 Methanosarcinales
-	B02.200.765.550 Methanosarcinaceae
-	B02.200.765.550.550 Methanosarcina
-	B02.200.765.550.550.200 Methanosarcina barkeri
-	B02.200.825 Thermococcales
-	B02.200.825.800 Thermococcaceae
-	B02.200.825.800.650 Pyrococcus
-	B02.200.825.800.650.020 Pyrococcus abyssi
-	B02.200.825.800.650.300 Pyrococcus furiosus
-	B02.200.825.800.650.400 Pyrococcus horikoshii
-	B02.200.825.800.800 Thermococcus
-	B02.200.850 Thermoplasmatales
-	B02.200.850.800 Thermoplasma
-	B02.500 Korarchaeota
-	B02.600 Nanoarchaeota
-	B03 Bacteria
-	B03.026 Acidobacteria
-	B03.054 Agricultural Inoculants
-	B03.110 Atypical Bacterial Forms
-	B03.110.422 L Forms
-	B03.110.761 Spheroplasts
-	B03.120 Bacteria, Aerobic
-	B03.130 Bacteria, Anaerobic
New Heading	<b>B03.135 Bacteria, Thermotolerant</b>
-	B03.140 Bacteroidetes
-	B03.140.094 Bacteroidaceae
-	B03.140.094.152 Bacteroides
-	B03.140.094.152.400 Bacteroides fragilis

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Heading	<b>B03.140.094.152.700</b>	<b>Bacteroides thetaiotaomicron</b>
-	B03.140.094.625	Porphyromonas
-	B03.140.094.625.350	Porphyromonas endodontalis
-	B03.140.094.625.515	Porphyromonas gingivalis
-	B03.140.094.640	Prevotella
-	B03.140.094.640.375	Prevotella intermedia
-	B03.140.094.640.500	Prevotella melaninogenica
-	B03.140.094.640.550	Prevotella nigrescens
-	B03.140.094.640.700	Prevotella ruminicola
-	B03.140.130	Cytophagaceae
-	B03.140.130.150	Cytophaga
-	B03.140.130.200	Flexibacter
-	B03.140.190	Flavobacteriaceae
-	B03.140.190.100	Capnocytophaga
-	B03.140.190.120	Chryseobacterium
-	B03.140.190.250	Flavobacterium
-	B03.140.190.600	Ornithobacterium
-	B03.140.190.700	Riemerella
-	B03.140.190.800	Tenacibaculum
-	B03.140.600	Pedobacter
-	B03.140.750	Rhodothermus
-	B03.140.800	Sphingobacterium
New Heading	<b>B03.140.900</b>	<b>Tannerella forsythia</b>
Old Tree	<b>B03.165</b>	<b>Blood-Borne Pathogens</b>
-	B03.250	Chlorobi
-	B03.250.140	Chlorobium
-	B03.275	Chloroflexi
-	B03.275.150	Chloroflexus
-	B03.280	Cyanobacteria
-	B03.280.100	Anabaena
-	B03.280.100.150	Anabaena cylindrica
-	B03.280.100.300	Anabaena flos-aquae
-	B03.280.100.900	Anabaena variabilis
-	B03.280.110	Aphanizomenon

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	B03.280.200	Cyanothece
-	B03.280.210	Cylindrospermopsis
-	B03.280.500	Microcystis
-	B03.280.550	Nodularia
-	B03.280.575	Nostoc
-	B03.280.575.150	Nostoc commune
-	B03.280.575.500	Nostoc muscorum
-	B03.280.612	Oscillatoria
-	B03.280.650	Plectonema
-	B03.280.697	Spirulina
-	B03.280.745	Synechococcus
-	B03.280.750	Synechocystis
New Heading	<b>B03.280.875</b>	<b>Trichodesmium</b>
-	B03.300	Endospore-Forming Bacteria
-	B03.300.390	Gram-Positive Endospore-Forming Bacteria
-	B03.300.390.400	Gram-Positive Endospore-Forming Rods
-	B03.300.390.400.158	Bacillaceae
-	B03.300.390.400.158.108	Anoxybacillus
-	B03.300.390.400.158.218	Bacillus
New Heading	<b>B03.300.390.400.158.218.076</b>	<b>Bacillus amyloliquefaciens</b>
-	B03.300.390.400.158.218.151	Bacillus anthracis
-	B03.300.390.400.158.218.252	Bacillus cereus
New Heading	<b>B03.300.390.400.158.218.283</b>	<b>Bacillus clausii</b>
New Heading	<b>B03.300.390.400.158.218.314</b>	<b>Bacillus coagulans</b>
New Heading	<b>B03.300.390.400.158.218.345</b>	<b>Bacillus firmus</b>
New Heading	<b>B03.300.390.400.158.218.376</b>	<b>Bacillus licheniformis</b>
-	B03.300.390.400.158.218.500	Bacillus megaterium
New Heading	<b>B03.300.390.400.158.218.613</b>	<b>Bacillus pumilus</b>
-	B03.300.390.400.158.218.725	Bacillus subtilis
-	B03.300.390.400.158.218.800	Bacillus thuringiensis
-	B03.300.390.400.158.400	Geobacillus

## MeSH Tree Changes for 2017

Type	Tree - heading
-	B03.300.390.400.158.400.400 <span style="float: right;">Geobacillus stearothermophilus</span>
-	B03.300.390.400.158.600 <span style="float: right;">Halobacillus</span>
-	B03.300.390.400.158.900 <span style="float: right;">Virgibacillus</span>
-	B03.300.390.400.160 <span style="float: right;">Brevibacillus</span>
-	B03.300.390.400.200 <span style="float: right;">Clostridium</span>
-	B03.300.390.400.200.025 <span style="float: right;">Clostridium acetobutylicum</span>
-	B03.300.390.400.200.100 <span style="float: right;">Clostridium beijerinckii</span>
-	B03.300.390.400.200.130 <span style="float: right;">Clostridium bifermentans</span>
-	B03.300.390.400.200.160 <span style="float: right;">Clostridium botulinum</span>
-	B03.300.390.400.200.160.050 <span style="float: right;">Clostridium botulinum type A</span>
-	B03.300.390.400.200.160.100 <span style="float: right;">Clostridium botulinum type B</span>
-	B03.300.390.400.200.160.150 <span style="float: right;">Clostridium botulinum type C</span>
-	B03.300.390.400.200.160.200 <span style="float: right;">Clostridium botulinum type D</span>
-	B03.300.390.400.200.160.250 <span style="float: right;">Clostridium botulinum type E</span>
-	B03.300.390.400.200.160.300 <span style="float: right;">Clostridium botulinum type F</span>
-	B03.300.390.400.200.160.350 <span style="float: right;">Clostridium botulinum type G</span>
-	B03.300.390.400.200.180 <span style="float: right;">Clostridium butyricum</span>
-	B03.300.390.400.200.200 <span style="float: right;">Clostridium cellulolyticum</span>
-	B03.300.390.400.200.205 <span style="float: right;">Clostridium cellulovorans</span>
-	B03.300.390.400.200.215 <span style="float: right;">Clostridium chauvoei</span>
-	B03.300.390.400.200.250 <span style="float: right;">Clostridium difficile</span>
-	B03.300.390.400.200.412 <span style="float: right;">Clostridium histolyticum</span>
-	B03.300.390.400.200.493 <span style="float: right;">Clostridium kluveri</span>
-	B03.300.390.400.200.575 <span style="float: right;">Clostridium perfringens</span>
-	B03.300.390.400.200.690 <span style="float: right;">Clostridium septicum</span>
-	B03.300.390.400.200.700 <span style="float: right;">Clostridium sordellii</span>
-	B03.300.390.400.200.710 <span style="float: right;">Clostridium sticklandii</span>
-	B03.300.390.400.200.713 <span style="float: right;">Clostridium symbiosum</span>
-	B03.300.390.400.200.722 <span style="float: right;">Clostridium tertium</span>
-	B03.300.390.400.200.725 <span style="float: right;">Clostridium tetani</span>
-	B03.300.390.400.200.740 <span style="float: right;">Clostridium tetanomorphum</span>
-	B03.300.390.400.200.770 <span style="float: right;">Clostridium thermocellum</span>
-	B03.300.390.400.200.800 <span style="float: right;">Clostridium tyrobutyricum</span>
-	B03.300.390.400.500 <span style="float: right;">Micromonosporaceae</span>
-	B03.300.390.400.500.500 <span style="float: right;">Micromonospora</span>
-	B03.300.390.400.525 <span style="float: right;">Moorella</span>



## MeSH Tree Changes for 2017

Type	Tree - heading	
-	B03.300.390.400.645	Paenibacillus
New Heading	<b>B03.300.390.400.645.500</b>	<b>Paenibacillus larvae</b>
New Heading	<b>B03.300.390.400.645.750</b>	<b>Paenibacillus polymyxa</b>
-	B03.300.390.400.790	Saccharopolyspora
-	B03.300.390.400.800	Staphylococcaceae
-	B03.300.390.400.800.750	Staphylococcus
-	B03.300.390.400.800.750.100	Staphylococcus aureus
-	B03.300.390.400.800.750.100.500 aureus	Methicillin-Resistant Staphylococcus aureus
-	B03.300.390.400.800.750.343	Staphylococcus epidermidis
-	B03.300.390.400.800.750.400	Staphylococcus haemolyticus
-	B03.300.390.400.800.750.425	Staphylococcus hominis
-	B03.300.390.400.800.750.435	Staphylococcus hyicus
-	B03.300.390.400.800.750.445	Staphylococcus intermedius
-	B03.300.390.400.800.750.500	Staphylococcus lugdunensis
-	B03.300.390.400.800.750.750	Staphylococcus saprophyticus
-	B03.300.390.400.810	Streptomycetaceae
-	B03.300.390.400.810.768	Streptomyces
-	B03.300.390.400.810.768.100	Streptomyces antibioticus
-	B03.300.390.400.810.768.125	Streptomyces aureofaciens
-	B03.300.390.400.810.768.200	Streptomyces coelicolor
-	B03.300.390.400.810.768.375	Streptomyces griseus
-	B03.300.390.400.810.768.500	Streptomyces lividans
-	B03.300.390.400.810.768.750	Streptomyces rimosus
-	B03.300.390.650	Pasteuria
-	B03.300.390.825	Sporosarcina
-	B03.353	Firmicutes
-	B03.353.250	Acidaminococcus
-	B03.353.500	Bacillales
-	B03.353.500.050	Alicyclobacillus
-	B03.353.500.100	Bacillaceae
-	B03.353.500.100.108	Anoxybacillus
-	B03.353.500.100.218	Bacillus
New Heading	<b>B03.353.500.100.218.076</b>	<b>Bacillus amyloliquefaciens</b>

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	B03.353.500.100.218.151	Bacillus anthracis
-	B03.353.500.100.218.252	Bacillus cereus
New Heading	<b>B03.353.500.100.218.283</b>	<b>Bacillus clausii</b>
New Heading	<b>B03.353.500.100.218.314</b>	<b>Bacillus coagulans</b>
New Heading	<b>B03.353.500.100.218.345</b>	<b>Bacillus firmus</b>
New Heading	<b>B03.353.500.100.218.376</b>	<b>Bacillus licheniformis</b>
-	B03.353.500.100.218.500	Bacillus megaterium
New Heading	<b>B03.353.500.100.218.613</b>	<b>Bacillus pumilus</b>
-	B03.353.500.100.218.725	Bacillus subtilis
-	B03.353.500.100.218.800	Bacillus thuringiensis
-	B03.353.500.100.400	Geobacillus
-	B03.353.500.100.400.400	Geobacillus stearothermophilus
-	B03.353.500.100.425	Halobacillus
-	B03.353.500.117	Brevibacillus
-	B03.353.500.120	Brochothrix
-	B03.353.500.310	Gemella
-	B03.353.500.500	Listeria
-	B03.353.500.500.500	Listeria monocytogenes
-	B03.353.500.645	Paenibacillus
New Heading	<b>B03.353.500.645.500</b>	<b>Paenibacillus larvae</b>
New Heading	<b>B03.353.500.645.750</b>	<b>Paenibacillus polymyxa</b>
-	B03.353.500.650	Pasteuria
-	B03.353.500.700	Planococcaceae
-	B03.353.500.700.700	Sporosarcina
-	B03.353.500.750	Staphylococcaceae
-	B03.353.500.750.750	Staphylococcus
-	B03.353.500.750.750.100	Staphylococcus aureus
-	B03.353.500.750.750.100.500	Methicillin-Resistant Staphylococcus aureus
-	B03.353.500.750.750.343	Staphylococcus epidermidis
-	B03.353.500.750.750.400	Staphylococcus haemolyticus
-	B03.353.500.750.750.425	Staphylococcus hominis

## MeSH Tree Changes for 2017

Type	Tree - heading
-	B03.353.500.750.750.435      Staphylococcus hyicus
-	B03.353.500.750.750.445      Staphylococcus intermedius
-	B03.353.500.750.750.500      Staphylococcus lugdunensis
-	B03.353.500.750.750.750      Staphylococcus saprophyticus
-	B03.353.500.875      Thermoactinomyces
-	B03.353.625      Clostridiales
-	B03.353.625.250      Butyrivibrio
New Heading	<b>B03.353.625.250.500      Butyrivibrio fibrisolvens</b>
-	B03.353.625.500      Clostridium
-	B03.353.625.500.025      Clostridium acetobutylicum
-	B03.353.625.500.100      Clostridium beijerinckii
-	B03.353.625.500.130      Clostridium bifermentans
-	B03.353.625.500.160      Clostridium botulinum
-	B03.353.625.500.160.050      Clostridium botulinum type A
-	B03.353.625.500.160.100      Clostridium botulinum type B
-	B03.353.625.500.160.150      Clostridium botulinum type C
-	B03.353.625.500.160.200      Clostridium botulinum type D
-	B03.353.625.500.160.250      Clostridium botulinum type E
-	B03.353.625.500.160.300      Clostridium botulinum type F
-	B03.353.625.500.160.350      Clostridium botulinum type G
-	B03.353.625.500.180      Clostridium butyricum
-	B03.353.625.500.200      Clostridium cellulolyticum
-	B03.353.625.500.205      Clostridium cellulovorans
-	B03.353.625.500.215      Clostridium chauvoei
-	B03.353.625.500.250      Clostridium difficile
-	B03.353.625.500.412      Clostridium histolyticum
-	B03.353.625.500.493      Clostridium kluyveri
-	B03.353.625.500.575      Clostridium perfringens
-	B03.353.625.500.690      Clostridium septicum
-	B03.353.625.500.700      Clostridium sordellii
-	B03.353.625.500.710      Clostridium sticklandii
-	B03.353.625.500.713      Clostridium symbiosum
-	B03.353.625.500.722      Clostridium tertium
-	B03.353.625.500.725      Clostridium tetani
-	B03.353.625.500.740      Clostridium tetanomorphum

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	B03.353.625.500.770	Clostridium thermocellum
-	B03.353.625.500.800	Clostridium tyrobutyricum
-	B03.353.625.750	Eubacterium
New Heading	<b>B03.353.625.766</b>	<b>Faecalibacterium</b>
New Heading	<b>B03.353.625.766.500</b>	<b>Faecalibacterium prausnitzii</b>
-	B03.353.625.782	Peptococcaceae
-	B03.353.625.782.149	Desulfitobacterium
-	B03.353.625.782.297	Desulfotomaculum
-	B03.353.625.782.593	Peptococcus
-	B03.353.625.798	Peptostreptococcus
-	B03.353.625.806	Ruminococcus
-	B03.353.625.813	Sarcina
-	B03.353.688	Erysipelothrix
-	B03.353.750	Lactobacillales
-	B03.353.750.030	Aerococcaceae
-	B03.353.750.030.014	Abiotrophia
-	B03.353.750.030.030	Aerococcus
-	B03.353.750.130	Carnobacteriaceae
-	B03.353.750.130.150	Carnobacterium
-	B03.353.750.250	Enterococcaceae
-	B03.353.750.250.250	Enterococcus
-	B03.353.750.250.250.280	Enterococcus faecalis
-	B03.353.750.250.250.300	Enterococcus faecium
New Heading	<b>B03.353.750.250.250.475</b>	<b>Enterococcus hirae</b>
-	B03.353.750.250.250.650	Vancomycin-Resistant Enterococci
-	B03.353.750.450	Lactobacillaceae
-	B03.353.750.450.475	Lactobacillus
-	B03.353.750.450.475.100	Lactobacillus acidophilus
-	B03.353.750.450.475.180	Lactobacillus brevis
-	B03.353.750.450.475.225	Lactobacillus casei
New Heading	<b>B03.353.750.450.475.238</b>	<b>Lactobacillus crispatus</b>
-	B03.353.750.450.475.250	Lactobacillus delbrueckii
-	B03.353.750.450.475.325	Lactobacillus fermentum

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Heading	<b>B03.353.750.450.475.363</b>	<b>Lactobacillus gasseri</b>
-	B03.353.750.450.475.400	Lactobacillus helveticus
New Heading	<b>B03.353.750.450.475.453</b>	<b>Lactobacillus johnsonii</b>
-	B03.353.750.450.475.506	Lactobacillus leichmannii
New Heading	<b>B03.353.750.450.475.559</b>	<b>Lactobacillus paracasei</b>
New Heading	<b>B03.353.750.450.475.586</b>	<b>Lactobacillus pentosus</b>
-	B03.353.750.450.475.612	Lactobacillus plantarum
-	B03.353.750.450.475.680	Lactobacillus reuteri
-	B03.353.750.450.475.700	Lactobacillus rhamnosus
New Heading	<b>B03.353.750.450.475.775</b>	<b>Lactobacillus sakei</b>
New Heading	<b>B03.353.750.450.475.850</b>	<b>Lactobacillus salivarius</b>
-	B03.353.750.450.737	Pediococcus
New Heading	<b>B03.353.750.450.737.500</b>	<b>Pediococcus acidilactici</b>
New Heading	<b>B03.353.750.450.737.750</b>	<b>Pediococcus pentosaceus</b>
-	B03.353.750.475	Leuconostocaceae
-	B03.353.750.475.450	Leuconostoc
New Heading	<b>B03.353.750.475.450.500</b>	<b>Leuconostoc mesenteroides</b>
-	B03.353.750.475.600	Oenococcus
-	B03.353.750.475.900	Weissella
-	B03.353.750.737	Streptococcaceae
-	B03.353.750.737.500	Lactococcus
-	B03.353.750.737.500.400	Lactococcus lactis
-	B03.353.750.737.872	Streptococcus
-	B03.353.750.737.872.100	Streptococcus agalactiae
-	B03.353.750.737.872.150	Streptococcus bovis
-	B03.353.750.737.872.225	Streptococcus equi
New Heading	<b>B03.353.750.737.872.243</b>	<b>Streptococcus gallolyticus</b>
New Heading	<b>B03.353.750.737.872.243.500</b> <b>gallolyticus</b>	<b>Streptococcus gallolyticus subspecies</b> <b>gallolyticus</b>
-	B03.353.750.737.872.260	Streptococcus gordonii

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Heading	<b>B03.353.750.737.872.405</b>	<b>Streptococcus iniae</b>
-	B03.353.750.737.872.550	Streptococcus pneumoniae
-	B03.353.750.737.872.575	Streptococcus pyogenes
New Heading	<b>B03.353.750.737.872.663</b>	<b>Streptococcus salivarius</b>
-	B03.353.750.737.872.750	Streptococcus suis
-	B03.353.750.737.872.800	Streptococcus thermophilus
-	B03.353.750.737.872.875	Viridans Streptococci
-	B03.353.750.737.872.875.475	Streptococcus milleri Group
-	B03.353.750.737.872.875.475.080	Streptococcus anginosus
-	B03.353.750.737.872.875.475.150	Streptococcus constellatus
-	B03.353.750.737.872.875.475.400	Streptococcus intermedius
-	B03.353.750.737.872.875.500	Streptococcus mitis
-	B03.353.750.737.872.875.520	Streptococcus mutans
-	B03.353.750.737.872.875.600	Streptococcus oralis
-	B03.353.750.737.872.875.700	Streptococcus sanguis
-	B03.353.750.737.872.875.750	Streptococcus sobrinus
-	B03.353.875	Moorella
-	B03.353.937	Thermoanaerobacter
-	B03.353.968	Thermoanaerobacterium
-	B03.353.984	Veillonellaceae
-	B03.370	Fusobacteria
-	B03.370.250	Fusobacterium
-	B03.370.250.400	Fusobacterium necrophorum
-	B03.370.250.500	Fusobacterium nucleatum
-	B03.370.437	Leptotrichia
-	B03.370.600	Propionigenium
-	B03.370.700	Streptobacillus
-	B03.440	Gram-Negative Bacteria
-	B03.440.040	Anaplasmataceae
-	B03.440.040.050	Anaplasma
-	B03.440.040.050.100	Anaplasma centrale
-	B03.440.040.050.500	Anaplasma marginale
-	B03.440.040.050.575	Anaplasma ovis
-	B03.440.040.050.600	Anaplasma phagocytophilum

## MeSH Tree Changes for 2017

Type	Tree - heading
-	B03.440.040.287 Ehrlichia
-	B03.440.040.287.090 Ehrlichia canis
-	B03.440.040.287.150 Ehrlichia chaffeensis
-	B03.440.040.287.700 Ehrlichia ruminantium
-	B03.440.040.525 Neorickettsia
-	B03.440.040.525.700 Neorickettsia risticii
-	B03.440.040.525.720 Neorickettsia sennetsu
-	B03.440.050 Arcobacter
-	B03.440.090 Bartonellaceae
-	B03.440.090.100 Bartonella
-	B03.440.090.100.080 Bartonella bacilliformis
-	B03.440.090.100.350 Bartonella henselae
-	B03.440.090.100.650 Bartonella quintana
-	B03.440.097 Brachyspira
-	B03.440.097.100 Brachyspira hyodysenteriae
-	B03.440.100 Buchnera
-	B03.440.180 Campylobacter
-	B03.440.180.200 Campylobacter coli
-	B03.440.180.325 Campylobacter fetus
-	B03.440.180.375 Campylobacter hyointestinalis
-	B03.440.180.425 Campylobacter jejuni
-	B03.440.180.500 Campylobacter lari
-	B03.440.180.650 Campylobacter rectus
-	B03.440.180.700 Campylobacter sputorum
-	B03.440.180.800 Campylobacter upsaliensis
-	B03.440.190 Chlamydiales
-	B03.440.190.190 Chlamydiaceae
-	B03.440.190.190.190 Chlamydia
-	B03.440.190.190.190.500 Chlamydia muridarum
-	B03.440.190.190.190.750 Chlamydia trachomatis
-	B03.440.190.190.230 Chlamydophila
-	B03.440.190.190.230.249 Chlamydophila pneumoniae
-	B03.440.190.190.230.500 Chlamydophila psittaci
-	B03.440.210 Chloroflexus
-	B03.440.305 Fibrobacteres
-	B03.440.305.500 Fibrobacter

## MeSH Tree Changes for 2017

Type	Tree - heading
-	B03.440.400 Gram-Negative Aerobic Bacteria
-	B03.440.400.050 Acidobacteria
-	B03.440.400.280 Caulobacter
-	B03.440.400.280.280 Caulobacter crescentus
-	B03.440.400.400 Gallionellaceae
-	B03.440.400.425 Gram-Negative Aerobic Rods and Cocci
-	B03.440.400.425.100 Acetobacteraceae
-	B03.440.400.425.100.100 Acetobacter
-	B03.440.400.425.100.110 Acidiphilium
-	B03.440.400.425.100.550 Gluconobacter
-	B03.440.400.425.100.550.550 Gluconobacter oxydans
-	B03.440.400.425.103 Acidithiobacillus
-	B03.440.400.425.103.800 Acidithiobacillus thiooxidans
-	B03.440.400.425.117 Alcaligenaceae
-	B03.440.400.425.117.024 Achromobacter
-	B03.440.400.425.117.024.100 Achromobacter cycloclastes
-	B03.440.400.425.117.024.120 Achromobacter denitrificans
-	B03.440.400.425.117.050 Alcaligenes
-	B03.440.400.425.117.050.200 Alcaligenes faecalis
-	B03.440.400.425.117.075 Bordetella
-	B03.440.400.425.117.075.099 Bordetella avium
-	B03.440.400.425.117.075.200 Bordetella bronchiseptica
-	B03.440.400.425.117.075.540 Bordetella parapertussis
-	B03.440.400.425.117.075.550 Bordetella pertussis
-	B03.440.400.425.117.800 Taylorella
-	B03.440.400.425.117.800.200 Taylorella equigenitalis
-	B03.440.400.425.126 Alteromonas
-	B03.440.400.425.127 Azorhizobium
-	B03.440.400.425.127.100 Azorhizobium caulinodans
-	B03.440.400.425.180 Bdellovibrio
New Heading	<b>B03.440.400.425.180.500 Bdellovibrio bacteriovorus</b>
-	B03.440.400.425.200 Bradyrhizobiaceae
-	B03.440.400.425.200.040 Afipia
-	B03.440.400.425.200.090 Bradyrhizobium
-	B03.440.400.425.200.520 Nitrobacter



## MeSH Tree Changes for 2017

Type	Tree - heading
-	B03.440.400.425.200.700 Rhodopseudomonas
-	B03.440.400.425.215 Brucellaceae
-	B03.440.400.425.215.500 Brucella
-	B03.440.400.425.215.500.100 Brucella abortus
-	B03.440.400.425.215.500.150 Brucella canis
-	B03.440.400.425.215.500.500 Brucella melitensis
-	B03.440.400.425.215.500.625 Brucella ovis
-	B03.440.400.425.215.500.750 Brucella suis
-	B03.440.400.425.251 Burkholderiaceae
-	B03.440.400.425.251.100 Burkholderia
-	B03.440.400.425.251.100.110 Burkholderia cepacia complex
-	B03.440.400.425.251.100.110.490 Burkholderia cenocepacia
-	B03.440.400.425.251.100.110.500 Burkholderia cepacia
-	B03.440.400.425.251.100.355 Burkholderia gladioli
-	B03.440.400.425.251.100.477 Burkholderia mallei
-	B03.440.400.425.251.100.600 Burkholderia pseudomallei
-	B03.440.400.425.251.200 Cupriavidus
-	B03.440.400.425.251.200.200 Cupriavidus necator
-	B03.440.400.425.251.650 Ralstonia
-	B03.440.400.425.251.650.600 Ralstonia pickettii
-	B03.440.400.425.251.650.750 Ralstonia solanacearum
-	B03.440.400.425.288 Caulobacteraceae
-	B03.440.400.425.288.100 Caulobacter
-	B03.440.400.425.288.100.100 Caulobacter crescentus
-	B03.440.400.425.292 Comamonadaceae
-	B03.440.400.425.292.150 Comamonas
-	B03.440.400.425.292.150.750 Comamonas testosteroni
-	B03.440.400.425.292.170 Delftia
-	B03.440.400.425.292.170.030 Delftia acidovorans
-	B03.440.400.425.292.500 Leptothrix
-	B03.440.400.425.292.750 Sphaerotilus
-	B03.440.400.425.297 Coxiellaceae
-	B03.440.400.425.297.150 Coxiella
-	B03.440.400.425.297.150.100 Coxiella burnetii
-	B03.440.400.425.300 Cytophagaceae
-	B03.440.400.425.300.150 Cytophaga

## MeSH Tree Changes for 2017

Type	Tree - heading
-	B03.440.400.425.300.200 Flexibacter
-	B03.440.400.425.310 Flavobacteriaceae
-	B03.440.400.425.310.120 Chryseobacterium
-	B03.440.400.425.310.250 Flavobacterium
-	B03.440.400.425.310.600 Ornithobacterium
-	B03.440.400.425.310.700 Riemerella
-	B03.440.400.425.310.800 Tenacibaculum
-	B03.440.400.425.340 Francisella
-	B03.440.400.425.340.590 Francisella tularensis
-	B03.440.400.425.365 Gluconacetobacter
-	B03.440.400.425.365.500 Gluconacetobacter xylinus
-	B03.440.400.425.377 Halomonadaceae
-	B03.440.400.425.377.249 Chromohalobacter
-	B03.440.400.425.377.500 Halomonas
-	B03.440.400.425.395 Halothiobacillus
-	B03.440.400.425.450 Legionellaceae
-	B03.440.400.425.450.450 Legionella
-	B03.440.400.425.450.450.400 Legionella longbeachae
-	B03.440.400.425.450.450.500 Legionella pneumophila
-	B03.440.400.425.475 Leptospiraceae
-	B03.440.400.425.475.475 Leptospira
-	B03.440.400.425.475.475.400 Leptospira interrogans
-	B03.440.400.425.475.475.400.050 Leptospira interrogans serovar australis
-	B03.440.400.425.475.475.400.055 Leptospira interrogans serovar autumnalis
-	B03.440.400.425.475.475.400.150 Leptospira interrogans serovar canicola
-	B03.440.400.425.475.475.400.400 Leptospira interrogans serovar hebdomadis
-	B03.440.400.425.475.475.400.420 Leptospira interrogans serovar icterohaemorrhagiae
-	B03.440.400.425.475.475.400.650 Leptospira interrogans serovar pomona
-	B03.440.400.425.487 Methylobacteriaceae
-	B03.440.400.425.487.500 Methylobacterium
-	B03.440.400.425.487.500.150 Methylobacterium extorquens
-	B03.440.400.425.500 Methylococcaceae
-	B03.440.400.425.500.500 Methylococcus
-	B03.440.400.425.500.500.100 Methylococcus capsulatus
-	B03.440.400.425.500.505 Methylomonas

## MeSH Tree Changes for 2017

Type	Tree - heading
-	B03.440.400.425.525                      Methylophilaceae
-	B03.440.400.425.525.500                      Methylobacillus
-	B03.440.400.425.525.520                      Methylophilus
-	B03.440.400.425.525.520.500                      Methylophilus methylotrophus
-	B03.440.400.425.537                      Moraxellaceae
-	B03.440.400.425.537.050                      Acinetobacter
-	B03.440.400.425.537.050.099                      Acinetobacter baumannii
-	B03.440.400.425.537.050.200                      Acinetobacter calcoaceticus
-	B03.440.400.425.537.525                      Moraxella
-	B03.440.400.425.537.525.200                      Moraxella (Branhamella) catarrhalis
-	B03.440.400.425.537.525.500                      Moraxella (Moraxella) bovis
-	B03.440.400.425.537.650                      Psychrobacter
-	B03.440.400.425.550                      Neisseriaceae
-	B03.440.400.425.550.400                      Kingella
-	B03.440.400.425.550.400.375                      Kingella kingae
-	B03.440.400.425.550.550                      Neisseria
-	B03.440.400.425.550.550.125                      Neisseria cinerea
-	B03.440.400.425.550.550.299                      Neisseria elongata
-	B03.440.400.425.550.550.474                      Neisseria gonorrhoeae
-	B03.440.400.425.550.550.600                      Neisseria lactamica
-	B03.440.400.425.550.550.641                      Neisseria meningitidis
-	B03.440.400.425.550.550.641.700                      Neisseria meningitidis, Serogroup A
-	B03.440.400.425.550.550.641.710                      Neisseria meningitidis, Serogroup B
-	B03.440.400.425.550.550.641.720                      Neisseria meningitidis, Serogroup C
-	B03.440.400.425.550.550.641.800                      Neisseria meningitidis, Serogroup W-135
-	B03.440.400.425.550.550.641.820                      Neisseria meningitidis, Serogroup Y
-	B03.440.400.425.550.550.660                      Neisseria mucosa
-	B03.440.400.425.550.550.800                      Neisseria sicca
-	B03.440.400.425.562                      Nitrosomonadaceae
-	B03.440.400.425.562.500                      Nitrosomonas
-	B03.440.400.425.562.500.200                      Nitrosomonas europaea
-	B03.440.400.425.575                      Ochrobactrum
-	B03.440.400.425.575.650                      Ochrobactrum anthropi
-	B03.440.400.425.587                      Oxalobacteraceae
-	B03.440.400.425.587.400                      Herbaspirillum
-	B03.440.400.425.600                      Paracoccus

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	B03.440.400.425.600.380	Paracoccus denitrificans
-	B03.440.400.425.600.690	Paracoccus pantotrophus
-	B03.440.400.425.611	Pedobacter
-	B03.440.400.425.622	Pseudoalteromonas
-	B03.440.400.425.625	Pseudomonadaceae
-	B03.440.400.425.625.050	Azotobacter
-	B03.440.400.425.625.050.850	Azotobacter vinelandii
-	B03.440.400.425.625.150	Cellvibrio
-	B03.440.400.425.625.625	Pseudomonas
-	B03.440.400.425.625.625.100	Pseudomonas aeruginosa
-	B03.440.400.425.625.625.120	Pseudomonas alcaligenes
New Heading	<b>B03.440.400.425.625.625.223</b>	<b>Pseudomonas chlororaphis</b>
-	B03.440.400.425.625.625.325	Pseudomonas fluorescens
-	B03.440.400.425.625.625.350	Pseudomonas fragi
-	B03.440.400.425.625.625.500	Pseudomonas mendocina
-	B03.440.400.425.625.625.650	Pseudomonas oleovorans
-	B03.440.400.425.625.625.662	Pseudomonas pseudoalcaligenes
-	B03.440.400.425.625.625.675	Pseudomonas putida
-	B03.440.400.425.625.625.750	Pseudomonas stutzeri
-	B03.440.400.425.625.625.770	Pseudomonas syringae
-	B03.440.400.425.700	Rhizobiaceae
-	B03.440.400.425.700.024	Agrobacterium
-	B03.440.400.425.700.024.050	Agrobacterium tumefaciens
-	B03.440.400.425.700.800	Rhizobium
-	B03.440.400.425.700.800.337	Rhizobium etli
-	B03.440.400.425.700.800.450	Rhizobium leguminosarum
-	B03.440.400.425.700.800.600	Rhizobium phaseoli
-	B03.440.400.425.700.800.800	Rhizobium tropici
-	B03.440.400.425.700.887	Sinorhizobium
-	B03.440.400.425.700.887.249	Sinorhizobium fredii
-	B03.440.400.425.700.887.500	Sinorhizobium meliloti
-	B03.440.400.425.708	Rhodospirillaceae
-	B03.440.400.425.708.100	Azospirillum
-	B03.440.400.425.708.100.020	Azospirillum brasilense
-	B03.440.400.425.708.100.510	Azospirillum lipoferum

## MeSH Tree Changes for 2017

Type	Tree - heading
-	B03.440.400.425.708.500                      Magnetospirillum
-	B03.440.400.425.708.733                      Rhodospirillum
-	B03.440.400.425.708.733.150                      Rhodospirillum centenum
-	B03.440.400.425.708.733.650                      Rhodospirillum rubrum
-	B03.440.400.425.710                      Rhodothermus
-	B03.440.400.425.745                      Sphingobacterium
-	B03.440.400.425.750                      Sphingomonas
-	B03.440.400.425.875                      Thermus
-	B03.440.400.425.875.875                      Thermus thermophilus
-	B03.440.400.425.935                      Xanthobacter
-	B03.440.400.425.967                      Xanthomonadaceae
-	B03.440.400.425.967.500                      Lysobacter
-	B03.440.400.425.967.750                      Stenotrophomonas
-	B03.440.400.425.967.750.500                      Stenotrophomonas maltophilia
-	B03.440.400.425.967.930                      Xanthomonas
-	B03.440.400.425.967.930.100                      Xanthomonas campestris
-	B03.440.400.425.967.930.850                      Xanthomonas vesicatoria
-	B03.440.400.425.967.950                      Xylella
-	B03.440.400.425.983                      Zoogloea
-	B03.440.400.450                      Gram-Negative Chemolithotrophic Bacteria
-	B03.440.400.450.800                      Thiobacillus
-	B03.440.400.645                      Thiotrichaceae
-	B03.440.400.645.100                      Beggiatoa
-	B03.440.400.645.800                      Thiothrix
-	B03.440.400.840                      Vitreoscilla
-	B03.440.425                      Gram-Negative Anaerobic Bacteria
-	B03.440.425.400                      Gram-Negative Anaerobic Cocci
-	B03.440.425.400.500                      Megasphaera
New Heading	<b>B03.440.425.400.500.500                      Megasphaera elsdenii</b>
-	B03.440.425.400.750                      Thiocapsa
-	B03.440.425.400.750.660                      Thiocapsa roseopersicina
-	B03.440.425.410                      Gram-Negative Anaerobic Straight, Curved, and Helical Rods
-	B03.440.425.410.073                      Acidaminococcus
-	B03.440.425.410.145                      Anaerobiospirillum
-	B03.440.425.410.194                      Bacteroidaceae

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	B03.440.425.410.194.152	Bacteroides
-	B03.440.425.410.194.152.400	Bacteroides fragilis
New Heading	<b>B03.440.425.410.194.152.700</b>	<b>Bacteroides thetaiotaomicron</b>
-	B03.440.425.410.194.625	Porphyromonas
-	B03.440.425.410.194.625.350	Porphyromonas endodontalis
-	B03.440.425.410.194.625.515	Porphyromonas gingivalis
-	B03.440.425.410.194.640	Prevotella
-	B03.440.425.410.194.640.375	Prevotella intermedia
-	B03.440.425.410.194.640.500	Prevotella melaninogenica
-	B03.440.425.410.194.640.550	Prevotella nigrescens
-	B03.440.425.410.194.640.700	Prevotella ruminicola
-	B03.440.425.410.200	Bilophila
-	B03.440.425.410.220	Brachyspira
-	B03.440.425.410.220.100	Brachyspira hyodysenteriae
-	B03.440.425.410.275	Chlorobium
-	B03.440.425.410.290	Chromatium
-	B03.440.425.410.350	Desulfovibrio
-	B03.440.425.410.350.040	Desulfovibrio africanus
-	B03.440.425.410.350.150	Desulfovibrio desulfuricans
-	B03.440.425.410.350.350	Desulfovibrio gigas
-	B03.440.425.410.350.875	Desulfovibrio vulgaris
-	B03.440.425.410.360	Desulfuromonas
-	B03.440.425.410.400	Dichelobacter nodosus
-	B03.440.425.410.405	Ectothiorhodospiraceae
-	B03.440.425.410.405.500	Ectothiorhodospira
-	B03.440.425.410.405.500.720	Ectothiorhodospira shaposhnikovii
-	B03.440.425.410.405.750	Halorhodospira halophila
New Heading	<b>B03.440.425.410.413</b>	<b>Faecalibacterium</b>
New Heading	<b>B03.440.425.410.413.500</b>	<b>Faecalibacterium prausnitzii</b>
-	B03.440.425.410.420	Fusobacterium
-	B03.440.425.410.420.400	Fusobacterium necrophorum
-	B03.440.425.410.420.500	Fusobacterium nucleatum
-	B03.440.425.410.430	Geobacter
-	B03.440.425.410.535	Leptotrichia

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	B03.440.425.410.580	Oxalobacter formigenes
-	B03.440.425.410.600	Propionigenium
-	B03.440.425.410.711	Spirochaetaceae
-	B03.440.425.410.711.193	Borrelia
-	B03.440.425.410.711.193.150	Borrelia burgdorferi Group
-	B03.440.425.410.711.193.150.125	Borrelia burgdorferi
-	B03.440.425.410.711.679	Spirochaeta
-	B03.440.425.410.711.795	Treponema
-	B03.440.425.410.711.795.700	Treponema denticola
-	B03.440.425.410.711.795.840	Treponema pallidum
-	B03.440.425.410.722	Succinivibrionaceae
-	B03.440.425.410.745	Thauera
-	B03.440.425.410.750	Thermotoga maritima
-	B03.440.425.410.760	Thermotoga neapolitana
-	B03.440.425.410.800	Veillonellaceae
-	B03.440.425.410.800.755	Pectinatus
-	B03.440.425.410.800.877	Selenomonas
-	B03.440.425.410.800.938	Veillonella
-	B03.440.425.410.875	Wolinella
-	B03.440.450	Gram-Negative Facultatively Anaerobic Rods
-	B03.440.450.009	Actinobacillus
-	B03.440.450.009.200	Actinobacillus equuli
-	B03.440.450.009.580	Actinobacillus pleuropneumoniae
-	B03.440.450.009.675	Actinobacillus seminis
-	B03.440.450.009.700	Actinobacillus suis
-	B03.440.450.019	Aeromonadaceae
-	B03.440.450.019.025	Aeromonas
-	B03.440.450.019.025.150	Aeromonas caviae
-	B03.440.450.019.025.380	Aeromonas hydrophila
-	B03.440.450.019.025.690	Aeromonas salmonicida
New Heading	<b>B03.440.450.019.025.845</b>	<b>Aeromonas veronii</b>
-	B03.440.450.040	Azoarcus
-	B03.440.450.340	Capnocytophaga
-	B03.440.450.342	Cardiobacteriaceae
-	B03.440.450.342.100	Cardiobacterium

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	B03.440.450.342.550	Dichelobacter nodosus
-	B03.440.450.360	Chromobacterium
-	B03.440.450.400	Eikenella
-	B03.440.450.400.275	Eikenella corrodens
-	B03.440.450.425	Enterobacteriaceae
-	B03.440.450.425.180	Calymmatobacterium
-	B03.440.450.425.198	Cronobacter
-	B03.440.450.425.198.150	Cronobacter sakazakii
-	B03.440.450.425.200	Citrobacter
-	B03.440.450.425.200.275	Citrobacter freundii
-	B03.440.450.425.200.475	Citrobacter koseri
-	B03.440.450.425.200.737	Citrobacter rodentium
-	B03.440.450.425.260	Edwardsiella
-	B03.440.450.425.260.340	Edwardsiella ictaluri
-	B03.440.450.425.260.750	Edwardsiella tarda
-	B03.440.450.425.275	Enterobacter
-	B03.440.450.425.275.099	Enterobacter aerogenes
-	B03.440.450.425.275.200	Enterobacter cloacae
-	B03.440.450.425.300	Erwinia
-	B03.440.450.425.300.050	Erwinia amylovora
-	B03.440.450.425.325	Escherichia
-	B03.440.450.425.325.300	Escherichia coli
-	B03.440.450.425.325.300.330	Enteropathogenic Escherichia coli
-	B03.440.450.425.325.300.340	Enterotoxigenic Escherichia coli
-	B03.440.450.425.325.300.360	Escherichia coli K12
New Heading	<b>B03.440.450.425.325.300.580</b>	<b>Extraintestinal Pathogenic Escherichia coli</b>
New Tree	<a href="#">B03.440.450.425.325.300.580.500</a>	<a href="#">Uropathogenic Escherichia coli</a>
-	B03.440.450.425.325.300.800	Shiga-Toxigenic Escherichia coli
-	B03.440.450.425.325.300.800.250	Enterohemorrhagic Escherichia coli
New Heading	<b>B03.440.450.425.325.300.800.250.250</b>	<b>Escherichia coli O104</b>
-	B03.440.450.425.325.300.800.250.500	Escherichia coli O157
Old Tree	<a href="#">B03.440.450.425.325.300.900</a>	<a href="#">Uropathogenic Escherichia coli</a>
-	B03.440.450.425.375	Hafnia
-	B03.440.450.425.375.050	Hafnia alvei



## MeSH Tree Changes for 2017

Type	Tree - heading
-	B03.440.450.425.425 Klebsiella
-	B03.440.450.425.425.580 Klebsiella oxytoca
-	B03.440.450.425.425.600 Klebsiella pneumoniae
-	B03.440.450.425.435 Kluyvera
-	B03.440.450.425.500 Morganella
-	B03.440.450.425.500.500 Morganella morgani
-	B03.440.450.425.580 Pantoea
-	B03.440.450.425.585 Pectobacterium
-	B03.440.450.425.585.120 Pectobacterium carotovorum
-	B03.440.450.425.585.150 Pectobacterium chrysanthemi
-	B03.440.450.425.590 Photorhabdus
-	B03.440.450.425.595 Plesiomonas
-	B03.440.450.425.600 Proteus
-	B03.440.450.425.600.501 Proteus mirabilis
-	B03.440.450.425.600.650 Proteus penneri
-	B03.440.450.425.600.800 Proteus vulgaris
-	B03.440.450.425.670 Providencia
-	B03.440.450.425.800 Salmonella
-	B03.440.450.425.800.100 Salmonella arizonae
-	B03.440.450.425.800.200 Salmonella enterica
-	B03.440.450.425.800.200.300 Salmonella enteritidis
-	B03.440.450.425.800.200.600 Salmonella paratyphi A
-	B03.440.450.425.800.200.700 Salmonella paratyphi B
-	B03.440.450.425.800.200.750 Salmonella paratyphi C
-	B03.440.450.425.800.200.800 Salmonella typhi
-	B03.440.450.425.800.200.825 Salmonella typhimurium
-	B03.440.450.425.814 Serratia
-	B03.440.450.425.814.620 Serratia liquefaciens
-	B03.440.450.425.814.664 Serratia marcescens
-	B03.440.450.425.850 Shigella
-	B03.440.450.425.850.175 Shigella boydii
-	B03.440.450.425.850.350 Shigella dysenteriae
-	B03.440.450.425.850.450 Shigella flexneri
-	B03.440.450.425.850.800 Shigella sonnei
-	B03.440.450.425.887 Wigglesworthia
-	B03.440.450.425.890 Xenorhabdus

## MeSH Tree Changes for 2017

Type	Tree - heading
-	B03.440.450.425.900 Yersinia
-	B03.440.450.425.900.300 Yersinia enterocolitica
-	B03.440.450.425.900.600 Yersinia pestis
-	B03.440.450.425.900.615 Yersinia pseudotuberculosis
-	B03.440.450.425.900.650 Yersinia ruckeri
-	B03.440.450.510 Moritella
-	B03.440.450.600 Pasteurellaceae
-	B03.440.450.600.224 Aggregatibacter
-	B03.440.450.600.224.500 Aggregatibacter actinomycetemcomitans
-	B03.440.450.600.224.750 Aggregatibacter aphrophilus
-	B03.440.450.600.224.875 Aggregatibacter segnis
-	B03.440.450.600.450 Haemophilus
-	B03.440.450.600.450.125 Haemophilus ducreyi
-	B03.440.450.600.450.330 Haemophilus influenzae
-	B03.440.450.600.450.330.150 Haemophilus influenzae type b
-	B03.440.450.600.450.665 Haemophilus paragallinarum
-	B03.440.450.600.450.675 Haemophilus parainfluenzae
-	B03.440.450.600.450.690 Haemophilus paraphrophilus
-	B03.440.450.600.450.700 Haemophilus parasuis
-	B03.440.450.600.450.750 Haemophilus somnus
-	B03.440.450.600.500 Mannheimia
-	B03.440.450.600.500.500 Mannheimia haemolytica
-	B03.440.450.600.600 Pasteurella
-	B03.440.450.600.600.500 Pasteurella multocida
-	B03.440.450.600.600.650 Pasteurella pneumotropica
-	B03.440.450.650 Rahnella
-	B03.440.450.690 Shewanella
-	B03.440.450.690.590 Shewanella putrefaciens
-	B03.440.450.700 Streptobacillus
-	B03.440.450.900 Vibrionaceae
-	B03.440.450.900.050 Aliivibrio
-	B03.440.450.900.050.860 Aliivibrio fischeri
-	B03.440.450.900.050.930 Aliivibrio salmonicida
-	B03.440.450.900.500 Listonella
-	B03.440.450.900.604 Photobacterium
-	B03.440.450.900.859 Vibrio

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	B03.440.450.900.859.030	Vibrio alginolyticus
-	B03.440.450.900.859.225	Vibrio cholerae
-	B03.440.450.900.859.225.075	Vibrio cholerae non-O1
-	B03.440.450.900.859.225.151	Vibrio cholerae O1
-	B03.440.450.900.859.225.200	Vibrio cholerae O139
-	B03.440.450.900.859.500	Vibrio mimicus
-	B03.440.450.900.859.550	Vibrio parahaemolyticus
-	B03.440.450.900.859.900	Vibrio vulnificus
-	B03.440.450.980	Zymomonas
-	B03.440.475	Gram-Negative Oxygenic Photosynthetic Bacteria
-	B03.440.475.100	Cyanobacteria
-	B03.440.475.100.100	Anabaena
-	B03.440.475.100.100.150	Anabaena cylindrica
-	B03.440.475.100.100.300	Anabaena flos-aquae
-	B03.440.475.100.100.900	Anabaena variabilis
-	B03.440.475.100.110	Aphanizomenon
-	B03.440.475.100.200	Cyanothece
-	B03.440.475.100.210	Cylindrospermopsis
-	B03.440.475.100.500	Microcystis
-	B03.440.475.100.550	Nodularia
-	B03.440.475.100.575	Nostoc
-	B03.440.475.100.575.150	Nostoc commune
-	B03.440.475.100.575.500	Nostoc muscorum
-	B03.440.475.100.612	Oscillatoria
-	B03.440.475.100.650	Plectonema
-	B03.440.475.100.655	Prochlorophytes
-	B03.440.475.100.655.600	Prochlorococcus
-	B03.440.475.100.655.620	Prochloron
-	B03.440.475.100.655.640	Prochlorothrix
-	B03.440.475.100.745	Synechococcus
-	B03.440.475.100.750	Synechocystis
New Heading	<b>B03.440.475.100.875</b>	<b>Trichodesmium</b>
-	B03.440.500	Helicobacter
-	B03.440.500.250	Helicobacter felis
-	B03.440.500.350	Helicobacter heilmannii

## MeSH Tree Changes for 2017

Type	Tree - heading
-	B03.440.500.362 Helicobacter hepaticus
-	B03.440.500.450 Helicobacter mustelae
-	B03.440.500.550 Helicobacter pylori
-	B03.440.520 Lawsonia Bacteria
-	B03.440.540 Methylosinus
-	B03.440.540.500 Methylosinus trichosporium
-	B03.440.595 Oceanospirillaceae
-	B03.440.602 Ornithobacterium
-	B03.440.612 Piscirickettsiaceae
-	B03.440.614 Planctomycetales
-	B03.440.623 Rhodobacter
-	B03.440.623.225 Rhodobacter capsulatus
-	B03.440.623.700 Rhodobacter sphaeroides
-	B03.440.645 Rhodomicrobium
-	B03.440.646 Rhodovulum
-	B03.440.647 Rickettsiaceae
-	B03.440.647.650 Rickettsiae
-	B03.440.647.650.550 Orientia tsutsugamushi
-	B03.440.647.650.650 Rickettsia
-	B03.440.647.650.650.040 Rickettsia akari
-	B03.440.647.650.650.125 Rickettsia conorii
-	B03.440.647.650.650.362 Rickettsia felis
-	B03.440.647.650.650.600 Rickettsia prowazekii
-	B03.440.647.650.650.650 Rickettsia rickettsii
-	B03.440.647.650.650.810 Rickettsia typhi
-	B03.440.647.825 Wolbachia
-	B03.440.680 Roseobacter
-	B03.440.840 Spirillaceae
-	B03.440.840.750 Spirillum
-	B03.440.860 Tenericutes
-	B03.440.860.074 Acholeplasmataceae
-	B03.440.860.074.150 Acholeplasma
-	B03.440.860.074.150.500 Acholeplasma laidlawii
-	B03.440.860.074.575 Phytoplasma
-	B03.440.860.300 Entomoplasmatales
-	B03.440.860.300.250 Entomoplasmataceae

## MeSH Tree Changes for 2017

Type	Tree - heading
-	B03.440.860.300.750 Spiroplasmataceae
-	B03.440.860.300.750.625 Spiroplasma
-	B03.440.860.300.750.625.500 Spiroplasma citri
-	B03.440.860.580 Mycoplasmatales
-	B03.440.860.580.553 Mycoplasmataceae
-	B03.440.860.580.553.553 Mycoplasma
-	B03.440.860.580.553.553.050 Mycoplasma agalactiae
-	B03.440.860.580.553.553.075 Mycoplasma arthritis
-	B03.440.860.580.553.553.195 Mycoplasma bovigenitalium
-	B03.440.860.580.553.553.200 Mycoplasma bovis
-	B03.440.860.580.553.553.210 Mycoplasma capricolum
-	B03.440.860.580.553.553.220 Mycoplasma conjunctivae
-	B03.440.860.580.553.553.267 Mycoplasma dispar
-	B03.440.860.580.553.553.315 Mycoplasma fermentans
-	B03.440.860.580.553.553.345 Mycoplasma gallisepticum
-	B03.440.860.580.553.553.355 Mycoplasma genitalium
-	B03.440.860.580.553.553.400 Mycoplasma hominis
-	B03.440.860.580.553.553.420 Mycoplasma hyopneumoniae
-	B03.440.860.580.553.553.425 Mycoplasma hyorhinis
-	B03.440.860.580.553.553.430 Mycoplasma hyosynoviae
-	B03.440.860.580.553.553.450 Mycoplasma iowae
-	B03.440.860.580.553.553.570 Mycoplasma meleagridis
-	B03.440.860.580.553.553.585 Mycoplasma mycoides
-	B03.440.860.580.553.553.630 Mycoplasma orale
-	B03.440.860.580.553.553.635 Mycoplasma ovipneumoniae
-	B03.440.860.580.553.553.645 Mycoplasma penetrans
-	B03.440.860.580.553.553.650 Mycoplasma pneumoniae
-	B03.440.860.580.553.553.670 Mycoplasma pulmonis
-	B03.440.860.580.553.553.720 Mycoplasma salivarium
-	B03.440.860.580.553.553.770 Mycoplasma synoviae
-	B03.440.860.580.553.900 Ureaplasma
-	B03.440.860.580.553.900.800 Ureaplasma urealyticum
-	B03.440.930 Verrucomicrobia
-	B03.510 Gram-Positive Bacteria
-	B03.510.024 Actinobacteria
-	B03.510.024.049 Actinomycetales

## MeSH Tree Changes for 2017

Type	Tree - heading
-	B03.510.024.049.050 Actinomycetaceae
-	B03.510.024.049.050.050 Actinomyces
-	B03.510.024.049.050.050.050 Actinomyces viscosus
-	B03.510.024.049.050.287 Arcanobacterium
-	B03.510.024.049.050.525 Mobiluncus
-	B03.510.024.049.100 Brevibacterium
-	B03.510.024.049.150 Cellulomonas
-	B03.510.024.049.180 Corynebacterium
-	B03.510.024.049.180.100 Brevibacterium flavum
-	B03.510.024.049.180.150 Corynebacterium diphtheriae
-	B03.510.024.049.180.300 Corynebacterium glutamicum
-	B03.510.024.049.180.600 Corynebacterium pseudotuberculosis
-	B03.510.024.049.180.625 Corynebacterium pyogenes
-	B03.510.024.049.250 Frankia
-	B03.510.024.049.300 Gordonia Bacterium
-	B03.510.024.049.475 Micrococcaceae
-	B03.510.024.049.475.050 Arthrobacter
-	B03.510.024.049.475.500 Micrococcus
-	B03.510.024.049.475.500.500 Micrococcus luteus
-	B03.510.024.049.500 Micromonosporaceae
-	B03.510.024.049.500.500 Micromonospora
-	B03.510.024.049.525 Mycobacteriaceae
-	B03.510.024.049.525.500 Mycobacterium
-	B03.510.024.049.525.500.300 Mycobacterium avium
-	B03.510.024.049.525.500.300.600 Mycobacterium avium subsp. paratuberculosis
-	B03.510.024.049.525.500.402 Mycobacterium bovis
-	B03.510.024.049.525.500.480 Mycobacterium haemophilum
-	B03.510.024.049.525.500.502 Mycobacterium leprae
-	B03.510.024.049.525.500.602 Mycobacterium lepraemurium
-	B03.510.024.049.525.500.640 Mycobacterium phlei
-	B03.510.024.049.525.500.702 Mycobacterium tuberculosis
-	B03.510.024.049.525.500.720 Nontuberculous Mycobacteria
-	B03.510.024.049.525.500.720.100 Mycobacterium avium Complex
-	B03.510.024.049.525.500.720.225 Mycobacterium chelonae
-	B03.510.024.049.525.500.720.325 Mycobacterium fortuitum

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	B03.510.024.049.525.500.720.400	Mycobacterium kansasii
-	B03.510.024.049.525.500.720.500	Mycobacterium marinum
-	B03.510.024.049.525.500.720.625	Mycobacterium scrofulaceum
-	B03.510.024.049.525.500.720.662	Mycobacterium smegmatis
-	B03.510.024.049.525.500.720.700	Mycobacterium ulcerans
-	B03.510.024.049.525.500.720.950	Mycobacterium xenopi
-	B03.510.024.049.537	Nocardiaceae
-	B03.510.024.049.537.550	Nocardia
-	B03.510.024.049.537.550.550	Nocardia asteroides
-	B03.510.024.049.537.775	Rhodococcus
-	B03.510.024.049.537.775.700	Rhodococcus equi
-	B03.510.024.049.600	Propionibacteriaceae
-	B03.510.024.049.600.600	Propionibacterium
-	B03.510.024.049.600.600.600	Propionibacterium acnes
New Heading	<b>B03.510.024.049.600.600.800</b>	<b>Propionibacterium freudenreichii</b>
-	B03.510.024.049.750	Saccharopolyspora
-	B03.510.024.049.775	Streptomycetaceae
-	B03.510.024.049.775.775	Streptomyces
-	B03.510.024.049.775.775.100	Streptomyces antibioticus
-	B03.510.024.049.775.775.125	Streptomyces aureofaciens
-	B03.510.024.049.775.775.200	Streptomyces coelicolor
-	B03.510.024.049.775.775.375	Streptomyces griseus
-	B03.510.024.049.775.775.500	Streptomyces lividans
-	B03.510.024.049.775.775.750	Streptomyces rimosus
-	B03.510.024.049.887	Tropheryma
-	B03.510.024.100	Bifidobacterium
New Heading	<b>B03.510.024.100.125</b>	<b>Bifidobacterium adolescentis</b>
New Heading	<b>B03.510.024.100.250</b>	<b>Bifidobacterium animalis</b>
New Heading	<b>B03.510.024.100.313</b>	<b>Bifidobacterium bifidum</b>
New Heading	<b>B03.510.024.100.375</b>	<b>Bifidobacterium breve</b>
New Heading	<b>B03.510.024.100.500</b>	<b>Bifidobacterium longum</b>
New	<b>B03.510.024.100.500.500</b>	<b>Bifidobacterium longum subspecies infantis</b>

## MeSH Tree Changes for 2017

Type	Tree - heading	
Heading		
New Heading	<b>B03.510.024.100.750</b>	<b>Bifidobacterium pseudocatenulatum</b>
-	B03.510.024.400	Gardnerella
-	B03.510.024.400.800	Gardnerella vaginalis
-	B03.510.100	Bacillales
-	B03.510.100.050	Alicyclobacillus
-	B03.510.100.100	Bacillaceae
-	B03.510.100.100.108	Anoxybacillus
-	B03.510.100.100.218	Bacillus
New Heading	<b>B03.510.100.100.218.076</b>	<b>Bacillus amyloliquefaciens</b>
-	B03.510.100.100.218.151	Bacillus anthracis
-	B03.510.100.100.218.252	Bacillus cereus
New Heading	<b>B03.510.100.100.218.283</b>	<b>Bacillus clausii</b>
New Heading	<b>B03.510.100.100.218.314</b>	<b>Bacillus coagulans</b>
New Heading	<b>B03.510.100.100.218.345</b>	<b>Bacillus firmus</b>
New Heading	<b>B03.510.100.100.218.376</b>	<b>Bacillus licheniformis</b>
-	B03.510.100.100.218.500	Bacillus megaterium
New Heading	<b>B03.510.100.100.218.613</b>	<b>Bacillus pumilus</b>
-	B03.510.100.100.218.725	Bacillus subtilis
-	B03.510.100.100.218.800	Bacillus thuringiensis
-	B03.510.100.100.400	Geobacillus
-	B03.510.100.100.400.400	Geobacillus stearothermophilus
-	B03.510.100.100.425	Halobacillus
-	B03.510.100.100.600	Planococcaceae
-	B03.510.100.100.600.700	Sporosarcina
-	B03.510.100.117	Brevibacillus
-	B03.510.100.120	Brochothrix
-	B03.510.100.310	Gemella
-	B03.510.100.500	Listeria
-	B03.510.100.500.500	Listeria monocytogenes
-	B03.510.100.645	Paenibacillus



## MeSH Tree Changes for 2017

Type	Tree - heading	
New Heading	<b>B03.510.100.645.500</b>	<b>Paenibacillus larvae</b>
New Heading	<b>B03.510.100.645.750</b>	<b>Paenibacillus polymyxa</b>
-	B03.510.100.650	Pasteuria
-	B03.510.100.750	Staphylococcaceae
-	B03.510.100.750.750	Staphylococcus
-	B03.510.100.750.750.100	Staphylococcus aureus
-	B03.510.100.750.750.100.500	Methicillin-Resistant Staphylococcus aureus
-	B03.510.100.750.750.343	Staphylococcus epidermidis
-	B03.510.100.750.750.400	Staphylococcus haemolyticus
-	B03.510.100.750.750.425	Staphylococcus hominis
-	B03.510.100.750.750.435	Staphylococcus hyicus
-	B03.510.100.750.750.445	Staphylococcus intermedius
-	B03.510.100.750.750.500	Staphylococcus lugdunensis
-	B03.510.100.750.750.750	Staphylococcus saprophyticus
-	B03.510.100.800	Thermoactinomyces
-	B03.510.400	Gram-Positive Cocci
-	B03.510.400.030	Aerococcus
-	B03.510.400.200	Deinococcus
-	B03.510.400.500	Micrococcaceae
-	B03.510.400.500.500	Micrococcus
-	B03.510.400.500.500.400	Micrococcus luteus
-	B03.510.400.531	Oenococcus
-	B03.510.400.562	Peptococcaceae
-	B03.510.400.562.593	Peptococcus
-	B03.510.400.625	Peptostreptococcus
-	B03.510.400.630	Planococcaceae
-	B03.510.400.630.600	Planococcus Bacteria
-	B03.510.400.630.700	Sporosarcina
-	B03.510.400.687	Ruminococcus
-	B03.510.400.750	Sarcina
-	B03.510.400.790	Staphylococcaceae
-	B03.510.400.790.750	Staphylococcus
-	B03.510.400.790.750.100	Staphylococcus aureus
-	B03.510.400.790.750.100.500	Methicillin-Resistant Staphylococcus aureus

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	B03.510.400.790.750.343	Staphylococcus epidermidis
-	B03.510.400.790.750.400	Staphylococcus haemolyticus
-	B03.510.400.790.750.425	Staphylococcus hominis
-	B03.510.400.790.750.435	Staphylococcus hyicus
-	B03.510.400.790.750.445	Staphylococcus intermedius
-	B03.510.400.790.750.500	Staphylococcus lugdunensis
-	B03.510.400.790.750.750	Staphylococcus saprophyticus
-	B03.510.400.800	Streptococcaceae
-	B03.510.400.800.500	Lactococcus
-	B03.510.400.800.500.400	Lactococcus lactis
-	B03.510.400.800.872	Streptococcus
-	B03.510.400.800.872.100	Streptococcus agalactiae
-	B03.510.400.800.872.150	Streptococcus bovis
-	B03.510.400.800.872.225	Streptococcus equi
New Heading	<b>B03.510.400.800.872.243</b>	<b>Streptococcus gallolyticus</b>
New Heading	<b>B03.510.400.800.872.243.500 gallolyticus</b>	<b>Streptococcus gallolyticus subspecies gallolyticus</b>
-	B03.510.400.800.872.260	Streptococcus gordonii
New Heading	<b>B03.510.400.800.872.405</b>	<b>Streptococcus iniae</b>
-	B03.510.400.800.872.550	Streptococcus pneumoniae
-	B03.510.400.800.872.575	Streptococcus pyogenes
New Heading	<b>B03.510.400.800.872.663</b>	<b>Streptococcus salivarius</b>
-	B03.510.400.800.872.750	Streptococcus suis
-	B03.510.400.800.872.800	Streptococcus thermophilus
-	B03.510.400.800.872.875	Viridans Streptococci
-	B03.510.400.800.872.875.475	Streptococcus milleri Group
-	B03.510.400.800.872.875.475.080	Streptococcus anginosus
-	B03.510.400.800.872.875.475.150	Streptococcus constellatus
-	B03.510.400.800.872.875.475.400	Streptococcus intermedius
-	B03.510.400.800.872.875.500	Streptococcus mitis
-	B03.510.400.800.872.875.520	Streptococcus mutans
-	B03.510.400.800.872.875.600	Streptococcus oralis
-	B03.510.400.800.872.875.700	Streptococcus sanguis
-	B03.510.400.800.872.875.750	Streptococcus sobrinus

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	B03.510.415	Gram-Positive Endospore-Forming Bacteria
-	B03.510.415.400	Gram-Positive Endospore-Forming Rods
-	B03.510.415.400.078	Alicyclobacillus
-	B03.510.415.400.158	Bacillaceae
-	B03.510.415.400.158.108	Anoxybacillus
-	B03.510.415.400.158.218	Bacillus
New Heading	<b>B03.510.415.400.158.218.076</b>	<b>Bacillus amyloliquefaciens</b>
-	B03.510.415.400.158.218.151	Bacillus anthracis
-	B03.510.415.400.158.218.252	Bacillus cereus
New Heading	<b>B03.510.415.400.158.218.283</b>	<b>Bacillus clausii</b>
New Heading	<b>B03.510.415.400.158.218.314</b>	<b>Bacillus coagulans</b>
New Heading	<b>B03.510.415.400.158.218.345</b>	<b>Bacillus firmus</b>
New Heading	<b>B03.510.415.400.158.218.376</b>	<b>Bacillus licheniformis</b>
-	B03.510.415.400.158.218.500	Bacillus megaterium
New Heading	<b>B03.510.415.400.158.218.613</b>	<b>Bacillus pumilus</b>
-	B03.510.415.400.158.218.725	Bacillus subtilis
-	B03.510.415.400.158.218.800	Bacillus thuringiensis
-	B03.510.415.400.158.400	Geobacillus
-	B03.510.415.400.158.400.400	Geobacillus stearothermophilus
-	B03.510.415.400.158.425	Halobacillus
-	B03.510.415.400.158.900	Virgibacillus
-	B03.510.415.400.179	Brevibacillus
-	B03.510.415.400.200	Clostridium
-	B03.510.415.400.200.025	Clostridium acetobutylicum
-	B03.510.415.400.200.100	Clostridium beijerinckii
-	B03.510.415.400.200.130	Clostridium bifermentans
-	B03.510.415.400.200.160	Clostridium botulinum
-	B03.510.415.400.200.160.050	Clostridium botulinum type A
-	B03.510.415.400.200.160.100	Clostridium botulinum type B
-	B03.510.415.400.200.160.150	Clostridium botulinum type C
-	B03.510.415.400.200.160.200	Clostridium botulinum type D
-	B03.510.415.400.200.160.250	Clostridium botulinum type E

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	B03.510.415.400.200.160.300	Clostridium botulinum type F
-	B03.510.415.400.200.160.350	Clostridium botulinum type G
-	B03.510.415.400.200.180	Clostridium butyricum
-	B03.510.415.400.200.200	Clostridium cellulolyticum
-	B03.510.415.400.200.205	Clostridium cellulovorans
-	B03.510.415.400.200.215	Clostridium chauvoei
-	B03.510.415.400.200.250	Clostridium difficile
-	B03.510.415.400.200.412	Clostridium histolyticum
-	B03.510.415.400.200.493	Clostridium kluyveri
-	B03.510.415.400.200.575	Clostridium perfringens
-	B03.510.415.400.200.690	Clostridium septicum
-	B03.510.415.400.200.700	Clostridium sordellii
-	B03.510.415.400.200.710	Clostridium sticklandii
-	B03.510.415.400.200.713	Clostridium symbiosum
-	B03.510.415.400.200.722	Clostridium tertium
-	B03.510.415.400.200.725	Clostridium tetani
-	B03.510.415.400.200.740	Clostridium tetanomorphum
-	B03.510.415.400.200.770	Clostridium thermocellum
-	B03.510.415.400.200.800	Clostridium tyrobutyricum
-	B03.510.415.400.230	Desulfotomaculum
-	B03.510.415.400.500	Micromonosporaceae
-	B03.510.415.400.500.500	Micromonospora
-	B03.510.415.400.525	Moorella
-	B03.510.415.400.645	Paenibacillus
New Heading	<b>B03.510.415.400.645.500</b>	<b>Paenibacillus larvae</b>
New Heading	<b>B03.510.415.400.645.750</b>	<b>Paenibacillus polymyxa</b>
-	B03.510.415.400.790	Saccharopolyspora
-	B03.510.415.400.810	Streptomycetaceae
-	B03.510.415.400.810.768	Streptomyces
-	B03.510.415.400.810.768.100	Streptomyces antibioticus
-	B03.510.415.400.810.768.125	Streptomyces aureofaciens
-	B03.510.415.400.810.768.200	Streptomyces coelicolor
-	B03.510.415.400.810.768.375	Streptomyces griseus
-	B03.510.415.400.810.768.500	Streptomyces lividans

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	B03.510.415.400.810.768.750	Streptomyces rimosus
-	B03.510.415.650	Pasteuria
-	B03.510.415.700	Sporosarcina
-	B03.510.460	Gram-Positive Rods
-	B03.510.460.400	Gram-Positive Asporogenous Rods
-	B03.510.460.400.400	Gram-Positive Asporogenous Rods, Irregular
-	B03.510.460.400.400.024	Acetobacterium
-	B03.510.460.400.400.049	Actinobacteria
-	B03.510.460.400.400.049.049	Actinomycetaceae
-	B03.510.460.400.400.049.049.178	Actinomyces
-	B03.510.460.400.400.049.049.178.800	Actinomyces viscosus
-	B03.510.460.400.400.049.049.200	Arcanobacterium
-	B03.510.460.400.400.049.049.589	Mobiluncus
-	B03.510.460.400.400.049.074	Arthrobacter
-	B03.510.460.400.400.049.100	Bifidobacterium
New Heading	<b>B03.510.460.400.400.049.100.125</b>	<b>Bifidobacterium adolescentis</b>
New Heading	<b>B03.510.460.400.400.049.100.250</b>	<b>Bifidobacterium animalis</b>
New Heading	<b>B03.510.460.400.400.049.100.313</b>	<b>Bifidobacterium bifidum</b>
New Heading	<b>B03.510.460.400.400.049.100.375</b>	<b>Bifidobacterium breve</b>
New Heading	<b>B03.510.460.400.400.049.100.500</b>	<b>Bifidobacterium longum</b>
New Heading	<b>B03.510.460.400.400.049.100.500.500 infantis</b>	<b>Bifidobacterium longum subspecies infantis</b>
New Heading	<b>B03.510.460.400.400.049.100.750</b>	<b>Bifidobacterium pseudocatenulatum</b>
-	B03.510.460.400.400.049.550	Brevibacterium
-	B03.510.460.400.400.200	Corynebacterium
-	B03.510.460.400.400.200.150	Corynebacterium diphtheriae
-	B03.510.460.400.400.200.300	Corynebacterium glutamicum
-	B03.510.460.400.400.200.600	Corynebacterium pseudotuberculosis
-	B03.510.460.400.400.200.625	Corynebacterium pyogenes
-	B03.510.460.400.400.250	Eubacterium
-	B03.510.460.400.400.600	Propionibacteriaceae
-	B03.510.460.400.400.600.600	Propionibacterium

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	B03.510.460.400.400.600.600.600	Propionibacterium acnes
New Heading	<b>B03.510.460.400.400.600.600.800</b>	<b>Propionibacterium freudenreichii</b>
-	B03.510.460.400.400.790	Thermoanaerobacter
-	B03.510.460.400.400.800	Thermoanaerobacterium
-	B03.510.460.400.410	Gram-Positive Asporogenous Rods, Regular
-	B03.510.460.400.410.120	Brochothrix
-	B03.510.460.400.410.350	Erysipelothrix
-	B03.510.460.400.410.475	Lactobacillaceae
-	B03.510.460.400.410.475.475	Lactobacillus
-	B03.510.460.400.410.475.475.100	Lactobacillus acidophilus
-	B03.510.460.400.410.475.475.180	Lactobacillus brevis
-	B03.510.460.400.410.475.475.225	Lactobacillus casei
New Heading	<b>B03.510.460.400.410.475.475.238</b>	<b>Lactobacillus crispatus</b>
-	B03.510.460.400.410.475.475.250	Lactobacillus delbrueckii
-	B03.510.460.400.410.475.475.325	Lactobacillus fermentum
New Heading	<b>B03.510.460.400.410.475.475.363</b>	<b>Lactobacillus gasseri</b>
-	B03.510.460.400.410.475.475.400	Lactobacillus helveticus
New Heading	<b>B03.510.460.400.410.475.475.453</b>	<b>Lactobacillus johnsonii</b>
-	B03.510.460.400.410.475.475.506	Lactobacillus leichmannii
New Heading	<b>B03.510.460.400.410.475.475.559</b>	<b>Lactobacillus paracasei</b>
New Heading	<b>B03.510.460.400.410.475.475.586</b>	<b>Lactobacillus pentosus</b>
-	B03.510.460.400.410.475.475.612	Lactobacillus plantarum
-	B03.510.460.400.410.475.475.680	Lactobacillus reuteri
-	B03.510.460.400.410.475.475.700	Lactobacillus rhamnosus
New Heading	<b>B03.510.460.400.410.475.475.775</b>	<b>Lactobacillus sakei</b>
New Heading	<b>B03.510.460.400.410.475.475.850</b>	<b>Lactobacillus salivarius</b>
-	B03.510.460.400.410.485	Listeria
-	B03.510.460.400.410.485.500	Listeria monocytogenes
-	B03.510.460.400.410.552	Mycobacteriaceae
-	B03.510.460.400.410.552.552	Mycobacterium
-	B03.510.460.400.410.552.552.300	Mycobacterium avium

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	B03.510.460.400.410.552.552.300.600 paratuberculosis	Mycobacterium avium subsp.
-	B03.510.460.400.410.552.552.402	Mycobacterium bovis
-	B03.510.460.400.410.552.552.480	Mycobacterium haemophilum
-	B03.510.460.400.410.552.552.502	Mycobacterium leprae
-	B03.510.460.400.410.552.552.602	Mycobacterium lepraemurium
-	B03.510.460.400.410.552.552.640	Mycobacterium phlei
-	B03.510.460.400.410.552.552.702	Mycobacterium tuberculosis
-	B03.510.460.400.410.552.552.720	Nontuberculous Mycobacteria
-	B03.510.460.400.410.552.552.720.100	Mycobacterium avium Complex
-	B03.510.460.400.410.552.552.720.225	Mycobacterium chelonae
-	B03.510.460.400.410.552.552.720.325	Mycobacterium fortuitum
-	B03.510.460.400.410.552.552.720.400	Mycobacterium kansasii
-	B03.510.460.400.410.552.552.720.500	Mycobacterium marinum
-	B03.510.460.400.410.552.552.720.625	Mycobacterium scrofulaceum
-	B03.510.460.400.410.552.552.720.662	Mycobacterium smegmatis
-	B03.510.460.400.410.552.552.720.700	Mycobacterium ulcerans
-	B03.510.460.400.410.552.552.720.950	Mycobacterium xenopi
-	B03.510.460.410	Gram-Positive Endospore-Forming Rods
-	B03.510.460.410.158	Bacillaceae
-	B03.510.460.410.158.108	Anoxybacillus
-	B03.510.460.410.158.218	Bacillus
New Heading	<b>B03.510.460.410.158.218.076</b>	<b>Bacillus amyloliquefaciens</b>
-	B03.510.460.410.158.218.151	Bacillus anthracis
-	B03.510.460.410.158.218.252	Bacillus cereus
New Heading	<b>B03.510.460.410.158.218.283</b>	<b>Bacillus clausii</b>
New Heading	<b>B03.510.460.410.158.218.314</b>	<b>Bacillus coagulans</b>
New Heading	<b>B03.510.460.410.158.218.345</b>	<b>Bacillus firmus</b>
New Heading	<b>B03.510.460.410.158.218.376</b>	<b>Bacillus licheniformis</b>
-	B03.510.460.410.158.218.500	Bacillus megaterium
New Heading	<b>B03.510.460.410.158.218.613</b>	<b>Bacillus pumilus</b>
-	B03.510.460.410.158.218.725	Bacillus subtilis

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	B03.510.460.410.158.218.800	Bacillus thuringiensis
-	B03.510.460.410.158.400	Geobacillus
-	B03.510.460.410.158.400.400	Geobacillus stearothermophilus
-	B03.510.460.410.158.425	Halobacillus
-	B03.510.460.410.158.900	Virgibacillus
-	B03.510.460.410.165	Brevibacillus
-	B03.510.460.410.500	Micromonosporaceae
-	B03.510.460.410.500.500	Micromonospora
-	B03.510.460.410.525	Moorella
-	B03.510.460.410.645	Paenibacillus
New Heading	<b>B03.510.460.410.645.500</b>	<b>Paenibacillus larvae</b>
New Heading	<b>B03.510.460.410.645.750</b>	<b>Paenibacillus polymyxa</b>
-	B03.510.460.410.790	Saccharopolyspora
-	B03.510.460.410.810	Streptomycetaceae
-	B03.510.460.410.810.768	Streptomyces
-	B03.510.460.410.810.768.100	Streptomyces antibioticus
-	B03.510.460.410.810.768.125	Streptomyces aureofaciens
-	B03.510.460.410.810.768.200	Streptomyces coelicolor
-	B03.510.460.410.810.768.375	Streptomyces griseus
-	B03.510.460.410.810.768.500	Streptomyces lividans
-	B03.510.460.410.810.768.750	Streptomyces rimosus
-	B03.510.550	Lactobacillales
-	B03.510.550.030	Aerococcaceae
-	B03.510.550.030.014	Abiotrophia
-	B03.510.550.030.030	Aerococcus
-	B03.510.550.130	Carnobacteriaceae
-	B03.510.550.130.150	Carnobacterium
-	B03.510.550.250	Enterococcaceae
-	B03.510.550.250.250	Enterococcus
-	B03.510.550.250.250.280	Enterococcus faecalis
-	B03.510.550.250.250.300	Enterococcus faecium
New Heading	<b>B03.510.550.250.250.475</b>	<b>Enterococcus hirae</b>
-	B03.510.550.250.250.650	Vancomycin-Resistant Enterococci
-	B03.510.550.450	Lactobacillaceae



## MeSH Tree Changes for 2017

Type	Tree - heading	
-	B03.510.550.450.475	Lactobacillus
-	B03.510.550.450.475.100	Lactobacillus acidophilus
-	B03.510.550.450.475.180	Lactobacillus brevis
-	B03.510.550.450.475.225	Lactobacillus casei
New Heading	<b>B03.510.550.450.475.238</b>	<b>Lactobacillus crispatus</b>
-	B03.510.550.450.475.250	Lactobacillus delbrueckii
-	B03.510.550.450.475.325	Lactobacillus fermentum
New Heading	<b>B03.510.550.450.475.363</b>	<b>Lactobacillus gasseri</b>
-	B03.510.550.450.475.400	Lactobacillus helveticus
New Heading	<b>B03.510.550.450.475.453</b>	<b>Lactobacillus johnsonii</b>
-	B03.510.550.450.475.506	Lactobacillus leichmannii
New Heading	<b>B03.510.550.450.475.559</b>	<b>Lactobacillus paracasei</b>
New Heading	<b>B03.510.550.450.475.586</b>	<b>Lactobacillus pentosus</b>
-	B03.510.550.450.475.612	Lactobacillus plantarum
-	B03.510.550.450.475.680	Lactobacillus reuteri
-	B03.510.550.450.475.700	Lactobacillus rhamnosus
New Heading	<b>B03.510.550.450.475.775</b>	<b>Lactobacillus sakei</b>
New Heading	<b>B03.510.550.450.475.850</b>	<b>Lactobacillus salivarius</b>
-	B03.510.550.450.737	Pediococcus
New Heading	<b>B03.510.550.450.737.500</b>	<b>Pediococcus acidilactici</b>
New Heading	<b>B03.510.550.450.737.750</b>	<b>Pediococcus pentosaceus</b>
-	B03.510.550.475	Leuconostocaceae
-	B03.510.550.475.450	Leuconostoc
New Heading	<b>B03.510.550.475.450.500</b>	<b>Leuconostoc mesenteroides</b>
-	B03.510.550.475.600	Oenococcus
-	B03.510.550.475.900	Weissella
-	B03.510.550.737	Streptococcaceae
-	B03.510.550.737.500	Lactococcus
-	B03.510.550.737.500.400	Lactococcus lactis

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	B03.510.550.737.872	Streptococcus
-	B03.510.550.737.872.100	Streptococcus agalactiae
-	B03.510.550.737.872.150	Streptococcus bovis
-	B03.510.550.737.872.225	Streptococcus equi
New Heading	<b>B03.510.550.737.872.243</b>	<b>Streptococcus gallolyticus</b>
New Heading	<b>B03.510.550.737.872.243.500 gallolyticus</b>	<b>Streptococcus gallolyticus subspecies</b>
-	B03.510.550.737.872.260	Streptococcus gordonii
New Heading	<b>B03.510.550.737.872.405</b>	<b>Streptococcus iniae</b>
-	B03.510.550.737.872.550	Streptococcus pneumoniae
-	B03.510.550.737.872.575	Streptococcus pyogenes
New Heading	<b>B03.510.550.737.872.663</b>	<b>Streptococcus salivarius</b>
-	B03.510.550.737.872.750	Streptococcus suis
-	B03.510.550.737.872.800	Streptococcus thermophilus
-	B03.510.550.737.872.875	Viridans Streptococci
-	B03.510.550.737.872.875.475	Streptococcus milleri Group
-	B03.510.550.737.872.875.475.080	Streptococcus anginosus
-	B03.510.550.737.872.875.475.150	Streptococcus constellatus
-	B03.510.550.737.872.875.475.400	Streptococcus intermedius
-	B03.510.550.737.872.875.500	Streptococcus mitis
-	B03.510.550.737.872.875.520	Streptococcus mutans
-	B03.510.550.737.872.875.600	Streptococcus oralis
-	B03.510.550.737.872.875.700	Streptococcus sanguis
-	B03.510.550.737.872.875.750	Streptococcus sobrinus
New Heading	<b>B03.585</b>	<b>Nitrogen-Fixing Bacteria</b>
New Tree	<a href="#">B03.585.051</a>	<a href="#">Anabaena</a>
New Tree	<a href="#">B03.585.051.150</a>	<a href="#">Anabaena cylindrica</a>
New Tree	<a href="#">B03.585.051.300</a>	<a href="#">Anabaena flos-aquae</a>
New Tree	<a href="#">B03.585.051.900</a>	<a href="#">Anabaena variabilis</a>
New Tree	<a href="#">B03.585.101</a>	<a href="#">Azoarcus</a>

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	B03.585.110	Azorhizobium
New Tree	B03.585.110.100	Azorhizobium caulinodans
New Tree	B03.585.120	Azospirillum brasilense
New Tree	B03.585.130	Azospirillum lipoferum
New Tree	B03.585.200	Beijerinckiaceae
New Tree	B03.585.420	Frankia
New Tree	B03.585.500	Herbaspirillum
New Tree	B03.585.625	Nodularia
New Tree	B03.585.750	Nostoc commune
New Tree	B03.585.760	Nostoc muscorum
New Heading	<b>B03.585.860</b>	<b>Paenibacillus polymyxa</b>
New Tree	B03.585.880	Plectonema
New Tree	B03.585.900	Rhizobium
New Tree	B03.585.900.337	Rhizobium etli
New Tree	B03.585.900.450	Rhizobium leguminosarum
New Tree	B03.585.900.600	Rhizobium phaseoli
New Tree	B03.585.900.800	Rhizobium tropici
New Heading	<b>B03.585.950</b>	<b>Trichodesmium</b>
-	B03.660	Proteobacteria
-	B03.660.050	Alphaproteobacteria
-	B03.660.050.020	Anaplasmataceae
-	B03.660.050.020.050	Anaplasma
-	B03.660.050.020.050.100	Anaplasma centrale
-	B03.660.050.020.050.500	Anaplasma marginale

## MeSH Tree Changes for 2017

Type	Tree - heading
-	B03.660.050.020.050.575                      Anaplasma ovis
-	B03.660.050.020.050.600                      Anaplasma phagocytophilum
-	B03.660.050.020.287                           Ehrlichia
-	B03.660.050.020.287.090                    Ehrlichia canis
-	B03.660.050.020.287.100                   Ehrlichia chaffeensis
-	B03.660.050.020.287.700                   Ehrlichia ruminantium
-	B03.660.050.020.525                        Neorickettsia
-	B03.660.050.020.525.700                   Neorickettsia risticii
-	B03.660.050.020.525.720                   Neorickettsia sennetsu
-	B03.660.050.030                              Bartonellaceae
-	B03.660.050.030.040                        Bartonella
-	B03.660.050.030.040.080                   Bartonella bacilliformis
-	B03.660.050.030.040.350                   Bartonella henselae
-	B03.660.050.030.040.650                   Bartonella quintana
-	B03.660.050.032                              Beijerinckiaceae
-	B03.660.050.035                              Bradyrhizobiaceae
-	B03.660.050.035.040                        Afipia
-	B03.660.050.035.090                        Bradyrhizobium
-	B03.660.050.035.520                        Nitrobacter
-	B03.660.050.035.700                        Rhodopseudomonas
-	B03.660.050.070                              Brucellaceae
-	B03.660.050.070.100                        Brucella
-	B03.660.050.070.100.100                   Brucella abortus
-	B03.660.050.070.100.150                   Brucella canis
-	B03.660.050.070.100.500                   Brucella melitensis
-	B03.660.050.070.100.625                   Brucella ovis
-	B03.660.050.070.100.750                   Brucella suis
-	B03.660.050.070.700                        Ochrobactrum
-	B03.660.050.070.700.650                   Ochrobactrum anthropi
-	B03.660.050.090                              Caulobacteraceae
-	B03.660.050.090.100                        Caulobacter
-	B03.660.050.090.100.100                   Caulobacter crescentus
-	B03.660.050.340                              Holosporaceae
-	B03.660.050.350                              Hyphomicrobiaceae
-	B03.660.050.350.080                        Azorhizobium
-	B03.660.050.350.080.100                   Azorhizobium caulinodans

## MeSH Tree Changes for 2017

Type	Tree - heading
-	B03.660.050.350.400 Hyphomicrobium
-	B03.660.050.350.675 Rhodomicrobium
-	B03.660.050.350.950 Xanthobacter
-	B03.660.050.500 Methylobacteriaceae
-	B03.660.050.500.500 Methylobacterium
-	B03.660.050.500.500.150 Methylobacterium extorquens
-	B03.660.050.512 Methylocystaceae
-	B03.660.050.512.500 Methylosinus
-	B03.660.050.512.500.750 Methylosinus trichosporium
-	B03.660.050.600 Phyllobacteriaceae
-	B03.660.050.600.500 Mesorhizobium
-	B03.660.050.663 Rhodospirillales
-	B03.660.050.663.050 Acetobacteraceae
-	B03.660.050.663.050.010 Acetobacter
-	B03.660.050.663.050.020 Acidiphilium
-	B03.660.050.663.050.400 Gluconacetobacter
-	B03.660.050.663.050.400.500 Gluconacetobacter xylinus
-	B03.660.050.663.050.415 Gluconobacter
-	B03.660.050.663.050.415.550 Gluconobacter oxydans
-	B03.660.050.663.750 Rhodospirillaceae
-	B03.660.050.663.750.100 Azospirillum
-	B03.660.050.663.750.100.020 Azospirillum brasilense
-	B03.660.050.663.750.100.510 Azospirillum lipoferum
-	B03.660.050.663.750.500 Magnetospirillum
-	B03.660.050.663.750.733 Rhodospirillum
-	B03.660.050.663.750.733.150 Rhodospirillum centenum
-	B03.660.050.663.750.733.650 Rhodospirillum rubrum
-	B03.660.050.730 Rhizobiaceae
-	B03.660.050.730.024 Agrobacterium
-	B03.660.050.730.024.500 Agrobacterium tumefaciens
-	B03.660.050.730.670 Rhizobium
-	B03.660.050.730.670.337 Rhizobium etli
-	B03.660.050.730.670.450 Rhizobium leguminosarum
-	B03.660.050.730.670.600 Rhizobium phaseoli
-	B03.660.050.730.670.800 Rhizobium tropici
-	B03.660.050.730.835 Sinorhizobium

## MeSH Tree Changes for 2017

Type	Tree - heading
-	B03.660.050.730.835.399 Sinorhizobium fredii
-	B03.660.050.730.835.800 Sinorhizobium meliloti
-	B03.660.050.750 Rhodobacteraceae
-	B03.660.050.750.600 Paracoccus
-	B03.660.050.750.600.125 Paracoccus denitrificans
-	B03.660.050.750.600.600 Paracoccus pantotrophus
-	B03.660.050.750.700 Rhodobacter
-	B03.660.050.750.700.225 Rhodobacter capsulatus
-	B03.660.050.750.700.700 Rhodobacter sphaeroides
-	B03.660.050.750.725 Rhodovulum
-	B03.660.050.750.750 Roseobacter
-	B03.660.050.765 Rickettsiaceae
-	B03.660.050.765.650 Rickettsieae
-	B03.660.050.765.650.600 Orientia tsutsugamushi
-	B03.660.050.765.650.650 Rickettsia
-	B03.660.050.765.650.650.040 Rickettsia akari
-	B03.660.050.765.650.650.180 Rickettsia conorii
-	B03.660.050.765.650.650.320 Rickettsia felis
-	B03.660.050.765.650.650.600 Rickettsia prowazekii
-	B03.660.050.765.650.650.650 Rickettsia rickettsii
-	B03.660.050.765.650.650.810 Rickettsia typhi
-	B03.660.050.765.825 Wolbachia
-	B03.660.050.800 Sphingomonadaceae
-	B03.660.050.800.750 Sphingomonas
-	B03.660.050.800.990 Zymomonas
-	B03.660.075 Betaproteobacteria
-	B03.660.075.027 Alcaligenaceae
-	B03.660.075.027.030 Achromobacter
-	B03.660.075.027.030.100 Achromobacter cycloclastes
-	B03.660.075.027.030.120 Achromobacter denitrificans
-	B03.660.075.027.050 Alcaligenes
-	B03.660.075.027.050.200 Alcaligenes faecalis
-	B03.660.075.027.075 Bordetella
-	B03.660.075.027.075.024 Bordetella avium
-	B03.660.075.027.075.050 Bordetella bronchiseptica
-	B03.660.075.027.075.590 Bordetella parapertussis

## MeSH Tree Changes for 2017

Type	Tree - heading
-	B03.660.075.027.075.600 Bordetella pertussis
-	B03.660.075.027.800 Taylorella
-	B03.660.075.027.800.200 Taylorella equigenitalis
-	B03.660.075.077 Burkholderiaceae
-	B03.660.075.077.100 Burkholderia
-	B03.660.075.077.100.110 Burkholderia cepacia complex
-	B03.660.075.077.100.110.490 Burkholderia cenocepacia
-	B03.660.075.077.100.110.500 Burkholderia cepacia
-	B03.660.075.077.100.355 Burkholderia gladioli
-	B03.660.075.077.100.477 Burkholderia mallei
-	B03.660.075.077.100.600 Burkholderia pseudomallei
-	B03.660.075.077.200 Cupriavidus
-	B03.660.075.077.200.200 Cupriavidus necator
-	B03.660.075.077.650 Ralstonia
-	B03.660.075.077.650.600 Ralstonia pickettii
-	B03.660.075.077.650.750 Ralstonia solanacearum
-	B03.660.075.102 Comamonadaceae
-	B03.660.075.102.150 Comamonas
-	B03.660.075.102.150.750 Comamonas testosteroni
-	B03.660.075.102.170 Delftia
-	B03.660.075.102.170.030 Delftia acidovorans
-	B03.660.075.102.500 Leptothrix
-	B03.660.075.102.750 Sphaerotilus
-	B03.660.075.375 Gallionellaceae
-	B03.660.075.400 Hydrogenophilaceae
-	B03.660.075.400.800 Thiobacillus
-	B03.660.075.495 Methylophilaceae
-	B03.660.075.495.500 Methylobacillus
-	B03.660.075.495.520 Methylophilus
-	B03.660.075.495.520.500 Methylophilus methylotrophus
-	B03.660.075.525 Neisseriaceae
-	B03.660.075.525.100 Chromobacterium
-	B03.660.075.525.170 Eikenella
-	B03.660.075.525.170.100 Eikenella corrodens
-	B03.660.075.525.410 Kingella
-	B03.660.075.525.410.410 Kingella kingae

## MeSH Tree Changes for 2017

Type	Tree - heading
-	B03.660.075.525.520                      Neisseria
-	B03.660.075.525.520.124                      Neisseria cinerea
-	B03.660.075.525.520.262                      Neisseria elongata
-	B03.660.075.525.520.400                      Neisseria gonorrhoeae
-	B03.660.075.525.520.475                      Neisseria lactamica
-	B03.660.075.525.520.500                      Neisseria meningitidis
-	B03.660.075.525.520.500.700                      Neisseria meningitidis, Serogroup A
-	B03.660.075.525.520.500.710                      Neisseria meningitidis, Serogroup B
-	B03.660.075.525.520.500.720                      Neisseria meningitidis, Serogroup C
-	B03.660.075.525.520.500.800                      Neisseria meningitidis, Serogroup W-135
-	B03.660.075.525.520.500.820                      Neisseria meningitidis, Serogroup Y
-	B03.660.075.525.520.520                      Neisseria mucosa
-	B03.660.075.525.520.760                      Neisseria sicca
-	B03.660.075.525.900                      Vitreoscilla
-	B03.660.075.550                      Nitrosomonadaceae
-	B03.660.075.550.500                      Nitrosomonas
-	B03.660.075.550.500.200                      Nitrosomonas europaea
-	B03.660.075.600                      Oxalobacteraceae
-	B03.660.075.600.400                      Herbaspirillum
-	B03.660.075.600.600                      Oxalobacter formigenes
-	B03.660.075.655                      Rhodocyclaceae
-	B03.660.075.655.070                      Azoarcus
-	B03.660.075.655.800                      Thauera
-	B03.660.075.655.975                      Zoogloea
-	B03.660.075.710                      Spirillaceae
-	B03.660.075.710.750                      Spirillum
-	B03.660.125                      Deltaproteobacteria
-	B03.660.125.050                      Bdellovibrio
New Heading	<b>B03.660.125.050.500                      Bdellovibrio bacteriovorus</b>
-	B03.660.125.087                      Bilophila
-	B03.660.125.125                      Desulfovibrio
-	B03.660.125.125.040                      Desulfovibrio africanus
-	B03.660.125.125.150                      Desulfovibrio desulfuricans
-	B03.660.125.125.350                      Desulfovibrio gigas
-	B03.660.125.125.875                      Desulfovibrio vulgaris



## MeSH Tree Changes for 2017

Type	Tree - heading	
-	B03.660.125.150	Desulfuromonas
-	B03.660.125.305	Geobacter
-	B03.660.125.460	Lawsonia Bacteria
-	B03.660.125.500	Myxococcales
-	B03.660.125.500.605	Myxococcus
-	B03.660.125.500.605.900	Myxococcus xanthus
-	B03.660.125.500.710	Stigmatella
-	B03.660.125.500.710.050	Stigmatella aurantiaca
-	B03.660.150	Epsilonproteobacteria
-	B03.660.150.050	Arcobacter
-	B03.660.150.100	Campylobacter
-	B03.660.150.100.100	Campylobacter coli
-	B03.660.150.100.220	Campylobacter fetus
-	B03.660.150.100.297	Campylobacter hyointestinalis
-	B03.660.150.100.375	Campylobacter jejuni
-	B03.660.150.100.450	Campylobacter lari
-	B03.660.150.100.700	Campylobacter rectus
-	B03.660.150.100.740	Campylobacter sputorum
-	B03.660.150.100.850	Campylobacter upsaliensis
-	B03.660.150.280	Helicobacter
-	B03.660.150.280.350	Helicobacter felis
-	B03.660.150.280.400	Helicobacter heilmannii
-	B03.660.150.280.410	Helicobacter hepaticus
-	B03.660.150.280.480	Helicobacter mustelae
-	B03.660.150.280.550	Helicobacter pylori
-	B03.660.150.920	Wolinella
-	B03.660.250	Gammaproteobacteria
-	B03.660.250.015	Acidithiobacillus
-	B03.660.250.015.800	Acidithiobacillus thiooxidans
-	B03.660.250.017	Aeromonadaceae
-	B03.660.250.017.025	Aeromonas
-	B03.660.250.017.025.150	Aeromonas caviae
-	B03.660.250.017.025.380	Aeromonas hydrophila
-	B03.660.250.017.025.690	Aeromonas salmonicida
New Heading	<b>B03.660.250.017.025.845</b>	<b>Aeromonas veronii</b>

## MeSH Tree Changes for 2017

Type	Tree - heading
-	B03.660.250.019 Alcanivoraceae
-	B03.660.250.021 Alteromonadaceae
-	B03.660.250.021.020 Alteromonas
-	B03.660.250.021.480 Marinobacter
-	B03.660.250.021.500 Moritella
-	B03.660.250.021.600 Pseudoalteromonas
-	B03.660.250.021.755 Shewanella
-	B03.660.250.021.755.750 Shewanella putrefaciens
-	B03.660.250.080 Cardiobacteriaceae
-	B03.660.250.080.100 Cardiobacterium
-	B03.660.250.080.550 Dichelobacter nodosus
-	B03.660.250.110 Chromatiaceae
-	B03.660.250.110.150 Chromatium
-	B03.660.250.110.400 Halothiobacillus
-	B03.660.250.110.750 Thiocapsa
-	B03.660.250.110.750.660 Thiocapsa roseopersicina
-	B03.660.250.132 Coxiellaceae
-	B03.660.250.132.150 Coxiella
-	B03.660.250.132.150.100 Coxiella burnetii
-	B03.660.250.145 Ectothiorhodospiraceae
-	B03.660.250.145.500 Ectothiorhodospira
-	B03.660.250.145.500.720 Ectothiorhodospira shaposhnikovii
-	B03.660.250.145.750 Halorhodospira halophila
-	B03.660.250.150 Enterobacteriaceae
-	B03.660.250.150.047 Buchnera
-	B03.660.250.150.095 Calymmatobacterium
-	B03.660.250.150.098 Cronobacter
-	B03.660.250.150.098.150 Cronobacter sakazakii
-	B03.660.250.150.100 Citrobacter
-	B03.660.250.150.100.210 Citrobacter freundii
-	B03.660.250.150.100.475 Citrobacter koseri
-	B03.660.250.150.100.737 Citrobacter rodentium
-	B03.660.250.150.160 Edwardsiella
-	B03.660.250.150.160.340 Edwardsiella ictaluri
-	B03.660.250.150.160.750 Edwardsiella tarda
-	B03.660.250.150.170 Enterobacter

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	B03.660.250.150.170.049	Enterobacter aerogenes
-	B03.660.250.150.170.100	Enterobacter cloacae
-	B03.660.250.150.175	Erwinia
-	B03.660.250.150.175.050	Erwinia amylovora
-	B03.660.250.150.180	Escherichia
-	B03.660.250.150.180.100	Escherichia coli
-	B03.660.250.150.180.100.330	Enteropathogenic Escherichia coli
-	B03.660.250.150.180.100.340	Enterotoxigenic Escherichia coli
-	B03.660.250.150.180.100.360	Escherichia coli K12
New Heading	<b>B03.660.250.150.180.100.580</b>	<b>Extraintestinal Pathogenic Escherichia coli</b>
New Tree	<b>B03.660.250.150.180.100.580.500</b>	<b>Uropathogenic Escherichia coli</b>
-	B03.660.250.150.180.100.800	Shiga-Toxigenic Escherichia coli
-	B03.660.250.150.180.100.800.250	Enterohemorrhagic Escherichia coli
New Heading	<b>B03.660.250.150.180.100.800.250.250</b>	<b>Escherichia coli O104</b>
-	B03.660.250.150.180.100.800.250.500	Escherichia coli O157
Old Tree	<b>B03.660.250.150.180.100.900</b>	<b>Uropathogenic Escherichia coli</b>
-	B03.660.250.150.280	Hafnia
-	B03.660.250.150.280.050	Hafnia alvei
-	B03.660.250.150.400	Klebsiella
-	B03.660.250.150.400.580	Klebsiella oxytoca
-	B03.660.250.150.400.590	Klebsiella pneumoniae
-	B03.660.250.150.410	Kluyvera
-	B03.660.250.150.500	Morganella
-	B03.660.250.150.500.500	Morganella morganii
-	B03.660.250.150.540	Pantoea
-	B03.660.250.150.542	Pectobacterium
-	B03.660.250.150.542.120	Pectobacterium carotovorum
-	B03.660.250.150.542.150	Pectobacterium chrysanthemi
-	B03.660.250.150.545	Photobacterium
-	B03.660.250.150.550	Plesiomonas
-	B03.660.250.150.590	Proteus
-	B03.660.250.150.590.500	Proteus mirabilis
-	B03.660.250.150.590.670	Proteus penneri
-	B03.660.250.150.590.840	Proteus vulgaris

## MeSH Tree Changes for 2017

Type	Tree - heading
-	B03.660.250.150.600 Providencia
-	B03.660.250.150.655 Rahnella
-	B03.660.250.150.710 Salmonella
-	B03.660.250.150.710.050 Salmonella arizonae
-	B03.660.250.150.710.160 Salmonella enterica
-	B03.660.250.150.710.160.160 Salmonella enteritidis
-	B03.660.250.150.710.160.580 Salmonella paratyphi A
-	B03.660.250.150.710.160.710 Salmonella paratyphi B
-	B03.660.250.150.710.160.711 Salmonella paratyphi C
-	B03.660.250.150.710.160.750 Salmonella typhi
-	B03.660.250.150.710.160.760 Salmonella typhimurium
-	B03.660.250.150.720 Serratia
-	B03.660.250.150.720.480 Serratia liquefaciens
-	B03.660.250.150.720.500 Serratia marcescens
-	B03.660.250.150.730 Shigella
-	B03.660.250.150.730.050 Shigella boydii
-	B03.660.250.150.730.125 Shigella dysenteriae
-	B03.660.250.150.730.210 Shigella flexneri
-	B03.660.250.150.730.710 Shigella sonnei
-	B03.660.250.150.880 Wigglesworthia
-	B03.660.250.150.910 Xenorhabdus
-	B03.660.250.150.950 Yersinia
-	B03.660.250.150.950.160 Yersinia enterocolitica
-	B03.660.250.150.950.580 Yersinia pestis
-	B03.660.250.150.950.590 Yersinia pseudotuberculosis
-	B03.660.250.150.950.650 Yersinia ruckeri
-	B03.660.250.200 Francisella
-	B03.660.250.200.750 Francisella tularensis
-	B03.660.250.350 Halomonadaceae
-	B03.660.250.350.249 Chromohalobacter
-	B03.660.250.350.500 Halomonas
-	B03.660.250.460 Legionellaceae
-	B03.660.250.460.460 Legionella
-	B03.660.250.460.460.480 Legionella longbeachae
-	B03.660.250.460.460.580 Legionella pneumophila
-	B03.660.250.500 Methylococcaceae

## MeSH Tree Changes for 2017

Type	Tree - heading
-	B03.660.250.500.500                      Methylococcus
-	B03.660.250.500.500.100                      Methylococcus capsulatus
-	B03.660.250.500.505                      Methylomonas
-	B03.660.250.530                      Moraxellaceae
-	B03.660.250.530.050                      Acinetobacter
-	B03.660.250.530.050.099                      Acinetobacter baumannii
-	B03.660.250.530.050.200                      Acinetobacter calcoaceticus
-	B03.660.250.530.525                      Moraxella
-	B03.660.250.530.525.200                      Moraxella (Branhamella) catarrhalis
-	B03.660.250.530.525.500                      Moraxella (Moraxella) bovis
-	B03.660.250.530.650                      Psychrobacter
-	B03.660.250.540                      Oceanospirillaceae
-	B03.660.250.540.500                      Marinomonas
-	B03.660.250.550                      Pasteurellaceae
-	B03.660.250.550.050                      Actinobacillus
-	B03.660.250.550.050.200                      Actinobacillus equuli
-	B03.660.250.550.050.580                      Actinobacillus pleuropneumoniae
-	B03.660.250.550.050.675                      Actinobacillus seminis
-	B03.660.250.550.050.700                      Actinobacillus suis
-	B03.660.250.550.170                      Aggregatibacter
-	B03.660.250.550.170.500                      Aggregatibacter actinomycetemcomitans
-	B03.660.250.550.170.750                      Aggregatibacter aphrophilus
-	B03.660.250.550.170.875                      Aggregatibacter segnis
-	B03.660.250.550.290                      Haemophilus
-	B03.660.250.550.290.125                      Haemophilus ducreyi
-	B03.660.250.550.290.330                      Haemophilus influenzae
-	B03.660.250.550.290.330.150                      Haemophilus influenzae type b
-	B03.660.250.550.290.665                      Haemophilus paragallinarum
-	B03.660.250.550.290.675                      Haemophilus parainfluenzae
-	B03.660.250.550.290.690                      Haemophilus paraphrophilus
-	B03.660.250.550.290.700                      Haemophilus parasuis
-	B03.660.250.550.290.750                      Haemophilus somnus
-	B03.660.250.550.500                      Mannheimia
-	B03.660.250.550.500.500                      Mannheimia haemolytica
-	B03.660.250.550.590                      Pasteurella
-	B03.660.250.550.590.500                      Pasteurella multocida

## MeSH Tree Changes for 2017

Type	Tree - heading
-	B03.660.250.550.590.650 <span style="float: right;">Pasteurella pneumotropica</span>
-	B03.660.250.565 <span style="float: right;">Piscirickettsiaceae</span>
-	B03.660.250.565.600 <span style="float: right;">Piscirickettsia</span>
-	B03.660.250.580 <span style="float: right;">Pseudomonadaceae</span>
-	B03.660.250.580.022 <span style="float: right;">Azotobacter</span>
-	B03.660.250.580.022.820 <span style="float: right;">Azotobacter vinelandii</span>
-	B03.660.250.580.100 <span style="float: right;">Cellvibrio</span>
-	B03.660.250.580.590 <span style="float: right;">Pseudomonas</span>
-	B03.660.250.580.590.050 <span style="float: right;">Pseudomonas aeruginosa</span>
-	B03.660.250.580.590.070 <span style="float: right;">Pseudomonas alcaligenes</span>
New Heading	<b>B03.660.250.580.590.140</b> <span style="float: right;"><b>Pseudomonas chlororaphis</b></span>
-	B03.660.250.580.590.210 <span style="float: right;">Pseudomonas fluorescens</span>
-	B03.660.250.580.590.230 <span style="float: right;">Pseudomonas fragi</span>
-	B03.660.250.580.590.500 <span style="float: right;">Pseudomonas mendocina</span>
-	B03.660.250.580.590.570 <span style="float: right;">Pseudomonas oleovorans</span>
-	B03.660.250.580.590.575 <span style="float: right;">Pseudomonas pseudoalcaligenes</span>
-	B03.660.250.580.590.580 <span style="float: right;">Pseudomonas putida</span>
-	B03.660.250.580.590.750 <span style="float: right;">Pseudomonas stutzeri</span>
-	B03.660.250.580.590.770 <span style="float: right;">Pseudomonas syringae</span>
-	B03.660.250.740 <span style="float: right;">Succinivibrionaceae</span>
-	B03.660.250.740.050 <span style="float: right;">Anaerobiospirillum</span>
-	B03.660.250.770 <span style="float: right;">Thiotrichaceae</span>
-	B03.660.250.770.100 <span style="float: right;">Beggiatoa</span>
-	B03.660.250.770.800 <span style="float: right;">Thiothrix</span>
-	B03.660.250.830 <span style="float: right;">Vibrionaceae</span>
-	B03.660.250.830.050 <span style="float: right;">Aliivibrio</span>
-	B03.660.250.830.050.860 <span style="float: right;">Aliivibrio fischeri</span>
-	B03.660.250.830.050.930 <span style="float: right;">Aliivibrio salmonicida</span>
-	B03.660.250.830.500 <span style="float: right;">Listonella</span>
-	B03.660.250.830.590 <span style="float: right;">Photobacterium</span>
-	B03.660.250.830.830 <span style="float: right;">Vibrio</span>
-	B03.660.250.830.830.030 <span style="float: right;">Vibrio alginolyticus</span>
-	B03.660.250.830.830.100 <span style="float: right;">Vibrio cholerae</span>
-	B03.660.250.830.830.100.075 <span style="float: right;">Vibrio cholerae non-O1</span>
-	B03.660.250.830.830.100.151 <span style="float: right;">Vibrio cholerae O1</span>



## MeSH Tree Changes for 2017

Type	Tree - heading
-	B03.900.350.150 Desulfovibrio desulfuricans
-	B03.900.350.350 Desulfovibrio gigas
-	B03.900.350.875 Desulfovibrio vulgaris
-	B03.900.370 Desulfuromonas
-	B04 Viruses
-	B04.080 Arboviruses
-	B04.100 Archaeal Viruses
-	B04.100.224 Fuselloviridae
-	B04.100.250 Guttaviridae
-	B04.100.450 Lipothrixviridae
-	B04.100.500 Myoviridae
-	B04.100.700 Rudiviridae
-	B04.100.750 Siphoviridae
-	B04.123 Bacteriophages
-	B04.123.100 Bacillus Phages
-	B04.123.150 Caudovirales
-	B04.123.150.500 Myoviridae
-	B04.123.150.500.260 Bacteriophage mu
-	B04.123.150.500.300 Bacteriophage P1
-	B04.123.150.500.305 Bacteriophage P2
-	B04.123.150.500.350 Bacteriophage T4
-	B04.123.150.700 Podoviridae
-	B04.123.150.700.065 Bacteriophage N4
-	B04.123.150.700.070 Bacteriophage P22
-	B04.123.150.700.100 Bacteriophage T3
-	B04.123.150.700.230 Bacteriophage T7
-	B04.123.150.800 Siphoviridae
-	B04.123.150.800.200 Bacteriophage HK022
-	B04.123.150.800.230 Bacteriophage lambda
-	B04.123.205 Coliphages
-	B04.123.205.200 Bacteriophage HK022
-	B04.123.205.230 Bacteriophage lambda
-	B04.123.205.250 Bacteriophage M13
-	B04.123.205.260 Bacteriophage mu
-	B04.123.205.280 Bacteriophage N4
-	B04.123.205.300 Bacteriophage P1



## MeSH Tree Changes for 2017

Type	Tree - heading
-	B04.123.205.305 Bacteriophage P2
-	B04.123.205.320 Bacteriophage phi X 174
-	B04.123.205.350 Bacteriophage PRD1
-	B04.123.205.600 Leviviridae
-	B04.123.205.600.050 Allolevivirus
-	B04.123.205.600.500 Levivirus
-	B04.123.205.891 T-Phages
-	B04.123.205.891.100 Bacteriophage T3
-	B04.123.205.891.200 Bacteriophage T4
-	B04.123.205.891.230 Bacteriophage T7
-	B04.123.210 Corticoviridae
-	B04.123.370 Inoviridae
-	B04.123.370.400 Inovirus
-	B04.123.370.400.240 Bacteriophage IKe
-	B04.123.370.400.250 Bacteriophage M13
-	B04.123.370.400.300 Bacteriophage Pf1
-	B04.123.370.600 Plectrovirus
-	B04.123.470 Microviridae
-	B04.123.470.500 Microvirus
-	B04.123.470.500.320 Bacteriophage phi X 174
-	B04.123.502 Mycobacteriophages
-	B04.123.655 Prophages
-	B04.123.660 Pseudomonas Phages
-	B04.123.660.535 Bacteriophage Pf1
-	B04.123.660.550 Bacteriophage phi 6
-	B04.123.691 RNA Phages
-	B04.123.691.230 Cystoviridae
-	B04.123.691.230.070 Bacteriophage phi 6
-	B04.123.691.600 Leviviridae
-	B04.123.691.600.050 Allolevivirus
-	B04.123.691.600.500 Levivirus
-	B04.123.706 Salmonella Phages
-	B04.123.706.070 Bacteriophage P22
-	B04.123.831 Staphylococcus Phages
-	B04.123.850 Streptococcus Phages
-	B04.123.900 Tectiviridae

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	B04.123.900.150	Bacteriophage PRD1
Old Tree	B04.194	Blood-Borne Pathogens
-	B04.265	Defective Viruses
-	B04.265.270	Hepatitis Delta Virus
-	B04.265.590	Sarcoma Viruses, Feline
-	B04.265.600	Sarcoma Viruses, Murine
-	B04.265.600.400	Harvey murine sarcoma virus
-	B04.265.600.500	Kirsten murine sarcoma virus
-	B04.265.600.600	Moloney murine sarcoma virus
-	B04.265.658	Satellite Viruses
-	B04.265.658.850	Tobacco mosaic satellite virus
-	B04.265.658.860	Tobacco necrosis satellite virus
-	B04.265.700	SSPE Virus
-	B04.280	DNA Viruses
-	B04.280.030	Adenoviridae
-	B04.280.030.080	Atadenovirus
-	B04.280.030.100	Aviadenovirus
-	B04.280.030.100.250	Fowl adenovirus A
-	B04.280.030.500	Mastadenovirus
-	B04.280.030.500.200	Adenoviruses, Canine
-	B04.280.030.500.350	Adenoviruses, Human
-	B04.280.030.500.675	Adenoviruses, Porcine
-	B04.280.030.500.700	Adenoviruses, Simian
-	B04.280.030.750	Siadenovirus
-	B04.280.038	Anelloviridae
-	B04.280.038.500	Torque teno virus
-	B04.280.045	Ascoviridae
-	B04.280.049	Asfarviridae
-	B04.280.049.035	African Swine Fever Virus
-	B04.280.065	Baculoviridae
-	B04.280.065.249	Granulovirus
-	B04.280.065.500	Nucleopolyhedrovirus
-	B04.280.090	Caudovirales
-	B04.280.090.500	Myoviridae
-	B04.280.090.500.260	Bacteriophage mu
-	B04.280.090.500.300	Bacteriophage P1

## MeSH Tree Changes for 2017

Type	Tree - heading
-	B04.280.090.500.305 Bacteriophage P2
-	B04.280.090.500.350 Bacteriophage T4
-	B04.280.090.700 Podoviridae
-	B04.280.090.700.065 Bacteriophage N4
-	B04.280.090.700.070 Bacteriophage P22
-	B04.280.090.700.100 Bacteriophage T3
-	B04.280.090.700.230 Bacteriophage T7
-	B04.280.090.800 Siphoviridae
-	B04.280.090.800.200 Bacteriophage HK022
-	B04.280.090.800.230 Bacteriophage lambda
-	B04.280.092 Caulimoviridae
-	B04.280.092.074 Badnavirus
-	B04.280.092.150 Caulimovirus
-	B04.280.092.850 Tungrovirus
-	B04.280.120 Circoviridae
-	B04.280.120.150 Circovirus
-	B04.280.120.400 Gyrovirus
-	B04.280.120.400.150 Chicken anemia virus
-	B04.280.210 DNA Tumor Viruses
-	B04.280.210.400 Gammaherpesvirinae
-	B04.280.210.400.500 Lymphocryptovirus
-	B04.280.210.400.500.450 Herpesvirus 4, Human
-	B04.280.210.400.700 Rhadinovirus
-	B04.280.210.400.700.300 Herpesvirus 4, Bovine
-	B04.280.210.400.700.330 Herpesvirus 8, Human
-	B04.280.210.400.700.400 Herpesvirus 2, Saimiriine
-	B04.280.210.430 Herpesvirus 1, Ranid
-	B04.280.210.600 Leporipoxvirus
-	B04.280.210.600.250 Fibroma Virus, Rabbit
-	B04.280.210.600.550 Myxoma virus
-	B04.280.210.620 Polyomaviridae
-	B04.280.210.620.615 Polyomavirus
-	B04.280.210.620.615.100 BK Virus
-	B04.280.210.620.615.400 JC Virus
-	B04.280.210.620.615.550 Merkel cell polyomavirus
-	B04.280.210.620.615.700 Simian virus 40

## MeSH Tree Changes for 2017

Type	Tree - heading
-	B04.280.210.655 Papillomaviridae
-	B04.280.210.655.050 Alphapapillomavirus
-	B04.280.210.655.050.606 Human papillomavirus 6
-	B04.280.210.655.050.606.611 Human papillomavirus 11
-	B04.280.210.655.050.616 Human papillomavirus 16
-	B04.280.210.655.050.616.500 Human papillomavirus 31
-	B04.280.210.655.050.618 Human papillomavirus 18
-	B04.280.210.655.100 Betapapillomavirus
-	B04.280.210.655.200 Deltapapillomavirus
-	B04.280.210.655.200.100 Bovine papillomavirus 1
-	B04.280.210.655.400 Gammapapillomavirus
-	B04.280.210.655.500 Kappapapillomavirus
-	B04.280.210.655.500.150 Cottontail rabbit papillomavirus
-	B04.280.210.655.520 Lambdapapillomavirus
-	B04.280.210.655.540 Mupapillomavirus
-	B04.280.210.655.950 Xipapillomavirus
-	B04.280.210.655.950.150 Bovine papillomavirus 4
-	B04.280.210.950 Yatapoxvirus
-	B04.280.210.950.950 Yaba monkey tumor virus
-	B04.280.224 Fuselloviridae
-	B04.280.350 Geminiviridae
-	B04.280.350.100 Begomovirus
-	B04.280.350.500 Maize streak virus
New Heading	<b>B04.280.356 Giant Viruses</b>
-	B04.280.362 Guttaviridae
-	B04.280.375 Hepadnaviridae
-	B04.280.375.100 Avihepadnavirus
-	B04.280.375.100.450 Hepatitis B Virus, Duck
-	B04.280.375.650 Orthohepadnavirus
-	B04.280.375.650.425 Hepatitis B virus
-	B04.280.375.650.460 Hepatitis B Virus, Woodchuck
-	B04.280.382 Herpesviridae
-	B04.280.382.100 Alphaherpesvirinae
-	B04.280.382.100.374 Iltovirus
-	B04.280.382.100.374.450 Herpesvirus 1, Gallid

## MeSH Tree Changes for 2017

Type	Tree - heading
-	B04.280.382.100.562 Mardivirus
-	B04.280.382.100.562.400 Herpesvirus 2, Gallid
-	B04.280.382.100.562.405 Herpesvirus 3, Gallid
-	B04.280.382.100.562.550 Herpesvirus 1, Meleagrid
-	B04.280.382.100.750 Simplexvirus
-	B04.280.382.100.750.200 Herpesvirus 2, Bovine
-	B04.280.382.100.750.350 Herpesvirus 1, Cercopithecine
-	B04.280.382.100.750.390 Herpesvirus 1, Human
-	B04.280.382.100.750.440 Herpesvirus 2, Human
-	B04.280.382.100.900 Varicellovirus
-	B04.280.382.100.900.400 Herpesvirus 1, Bovine
-	B04.280.382.100.900.410 Herpesvirus 5, Bovine
-	B04.280.382.100.900.420 Herpesvirus 1, Canid
-	B04.280.382.100.900.430 Herpesvirus 1, Equid
-	B04.280.382.100.900.435 Herpesvirus 3, Equid
-	B04.280.382.100.900.440 Herpesvirus 4, Equid
-	B04.280.382.100.900.460 Herpesvirus 3, Human
-	B04.280.382.100.900.510 Herpesvirus 1, Suid
-	B04.280.382.150 Betaherpesvirinae
-	B04.280.382.150.150 Cytomegalovirus
-	B04.280.382.150.500 Muromegalovirus
-	B04.280.382.150.750 Roseolovirus
-	B04.280.382.150.750.400 Herpesvirus 6, Human
-	B04.280.382.150.750.420 Herpesvirus 7, Human
-	B04.280.382.400 Gammaherpesvirinae
-	B04.280.382.400.500 Lymphocryptovirus
-	B04.280.382.400.500.400 Herpesvirus 4, Human
-	B04.280.382.400.700 Rhadinovirus
-	B04.280.382.400.700.300 Herpesvirus 4, Bovine
-	B04.280.382.400.700.330 Herpesvirus 8, Human
-	B04.280.382.400.700.400 Herpesvirus 2, Saimiriine
-	B04.280.382.500 Herpesvirus 1, Ranid
-	B04.280.382.530 Ictalurivirus
-	B04.280.400 Inoviridae
-	B04.280.400.400 Inovirus
-	B04.280.400.400.240 Bacteriophage IKE

## MeSH Tree Changes for 2017

Type	Tree - heading
-	B04.280.400.400.250 Bacteriophage M13
-	B04.280.400.400.300 Bacteriophage Pf1
-	B04.280.400.600 Plectrovirus
-	B04.280.410 Iridoviridae
-	B04.280.410.400 Iridovirus
-	B04.280.410.700 Ranavirus
-	B04.280.450 Lipothrixviridae
-	B04.280.470 Microviridae
-	B04.280.470.500 Microvirus
-	B04.280.470.500.320 Bacteriophage phi X 174
-	B04.280.475 Mimiviridae
-	B04.280.498 Nanoviridae
-	B04.280.498.500 Nanovirus
-	B04.280.505 Nimaviridae
-	B04.280.505.900 White spot syndrome virus 1
-	B04.280.580 Parvoviridae
-	B04.280.580.325 Densovirinae
-	B04.280.580.325.200 Densovirus
-	B04.280.580.650 Parvovirinae
-	B04.280.580.650.040 Amdovirus
-	B04.280.580.650.040.050 Aleutian Mink Disease Virus
-	B04.280.580.650.085 Bocavirus
-	B04.280.580.650.085.500 Human bocavirus
-	B04.280.580.650.170 Dependovirus
-	B04.280.580.650.200 Erythrovirus
-	B04.280.580.650.200.650 Parvovirus B19, Human
-	B04.280.580.650.600 Parvovirus
-	B04.280.580.650.600.325 Feline Panleukopenia Virus
-	B04.280.580.650.600.325.330 Mink enteritis virus
-	B04.280.580.650.600.325.660 Parvovirus, Canine
-	B04.280.580.650.600.437 H-1 parvovirus
-	B04.280.580.650.600.550 Minute Virus of Mice
-	B04.280.580.650.600.690 Parvovirus, Porcine
-	B04.280.600 Phycodnaviridae
-	B04.280.630 Polydnaviridae
-	B04.280.650 Poxviridae

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	B04.280.650.160	Chordopoxvirinae
-	B04.280.650.160.100	Avipoxvirus
-	B04.280.650.160.100.149	Canarypox virus
-	B04.280.650.160.100.300	Fowlpox virus
-	B04.280.650.160.150	Capripoxvirus
-	B04.280.650.160.150.500	Lumpy skin disease virus
-	B04.280.650.160.500	Leporipoxvirus
-	B04.280.650.160.500.250	Fibroma Virus, Rabbit
-	B04.280.650.160.500.550	Myxoma virus
-	B04.280.650.160.550	Molluscipoxvirus
-	B04.280.650.160.550.500	Molluscum contagiosum virus
-	B04.280.650.160.650	Orthopoxvirus
-	B04.280.650.160.650.150	Cowpox virus
-	B04.280.650.160.650.200	Ectromelia virus
-	B04.280.650.160.650.500	Monkeypox virus
-	B04.280.650.160.650.900	Vaccinia virus
-	B04.280.650.160.650.930	Variola virus
-	B04.280.650.160.670	Parapoxvirus
-	B04.280.650.160.670.600	Orf virus
-	B04.280.650.160.670.650	Pseudocowpox Virus
-	B04.280.650.160.800	Suipoxvirus
-	B04.280.650.160.980	Yatapoxvirus
-	B04.280.650.160.980.950	Yaba monkey tumor virus
-	B04.280.650.250	Entomopoxvirinae
-	B04.280.775	Rudiviridae
-	B04.280.900	Tectiviridae
New Heading	<b>B04.280.950</b>	<b>Virophages</b>
-	B04.352	Fungal Viruses
-	B04.423	Helper Viruses
-	B04.450	Hepatitis Viruses
-	B04.450.100	Adenoviruses, Canine
-	B04.450.355	GB virus B
-	B04.450.360	GB virus C
-	B04.450.390	Hepadnaviridae
-	B04.450.390.100	Avihepadnavirus

## MeSH Tree Changes for 2017

Type	Tree - heading
-	B04.450.390.100.450 Hepatitis B Virus, Duck
-	B04.450.390.650 Orthohepadnavirus
-	B04.450.390.650.425 Hepatitis B virus
-	B04.450.390.650.460 Hepatitis B Virus, Woodchuck
-	B04.450.410 Hepacivirus
-	B04.450.411 Hepatitis Delta Virus
-	B04.450.412 Hepatitis E virus
-	B04.450.416 Hepatitis Virus, Duck
-	B04.450.420 Hepatovirus
-	B04.450.420.410 Hepatitis A virus
-	B04.450.420.410.500 Hepatitis A Virus, Human
-	B04.450.580 Murine hepatitis virus
-	B04.525 Insect Viruses
-	B04.525.045 Ascoviridae
-	B04.525.100 Baculoviridae
-	B04.525.100.249 Granulovirus
-	B04.525.100.500 Nucleopolyhedrovirus
-	B04.525.150 Densovirinae
-	B04.525.150.150 Densovirus
-	B04.525.165 Dicistroviridae
-	B04.525.250 Entomopoxvirinae
-	B04.525.400 Iridovirus
-	B04.525.515 Nodaviridae
-	B04.525.630 Polydnviridae
-	B04.613 Oncogenic Viruses
-	B04.613.204 DNA Tumor Viruses
-	B04.613.204.500 Gammaherpesvirinae
-	B04.613.204.500.500 Lymphocryptovirus
-	B04.613.204.500.500.400 Herpesvirus 4, Human
-	B04.613.204.500.700 Rhadinovirus
-	B04.613.204.500.700.300 Herpesvirus 4, Bovine
-	B04.613.204.500.700.330 Herpesvirus 8, Human
-	B04.613.204.500.700.400 Herpesvirus 2, Saimiriine
-	B04.613.204.550 Herpesvirus 1, Ranid
-	B04.613.204.600 Leporipoxvirus
-	B04.613.204.600.250 Fibroma Virus, Rabbit



## MeSH Tree Changes for 2017

Type	Tree - heading
-	B04.613.204.600.550 Myxoma virus
-	B04.613.204.655 Papillomaviridae
-	B04.613.204.655.050 Alphapapillomavirus
-	B04.613.204.655.050.606 Human papillomavirus 6
-	B04.613.204.655.050.606.611 Human papillomavirus 11
-	B04.613.204.655.050.616 Human papillomavirus 16
-	B04.613.204.655.050.616.500 Human papillomavirus 31
-	B04.613.204.655.050.618 Human papillomavirus 18
-	B04.613.204.655.100 Betapapillomavirus
-	B04.613.204.655.200 Deltapapillomavirus
-	B04.613.204.655.200.100 Bovine papillomavirus 1
-	B04.613.204.655.400 Gammapapillomavirus
-	B04.613.204.655.500 Kappapapillomavirus
-	B04.613.204.655.500.150 Cottontail rabbit papillomavirus
-	B04.613.204.655.520 Lambdapapillomavirus
-	B04.613.204.655.540 Mupapillomavirus
-	B04.613.204.655.950 Xipapillomavirus
-	B04.613.204.655.950.150 Bovine papillomavirus 4
-	B04.613.204.670 Polyomaviridae
-	B04.613.204.670.615 Polyomavirus
-	B04.613.204.670.615.100 BK Virus
-	B04.613.204.670.615.400 JC Virus
-	B04.613.204.670.615.550 Merkel cell polyomavirus
-	B04.613.204.670.615.700 Simian virus 40
-	B04.613.204.950 Yatapoxvirus
-	B04.613.204.950.950 Yaba monkey tumor virus
-	B04.613.807 Retroviridae
-	B04.613.807.070 Alpharetrovirus
-	B04.613.807.070.100 Avian Leukosis Virus
-	B04.613.807.070.110 Avian Myeloblastosis Virus
-	B04.613.807.070.120 Avian Sarcoma Viruses
-	B04.613.807.070.120.700 Rous sarcoma virus
-	B04.613.807.124 Betaretrovirus
-	B04.613.807.124.400 Jaagsiekte sheep retrovirus
-	B04.613.807.124.500 Mammary Tumor Virus, Mouse
-	B04.613.807.124.520 Mason-Pfizer monkey virus

## MeSH Tree Changes for 2017

Type	Tree - heading
-	B04.613.807.200 Deltaretrovirus
-	B04.613.807.200.600 Leukemia Virus, Bovine
-	B04.613.807.200.725 Primate T-lymphotropic virus 1
-	B04.613.807.200.725.400 Human T-lymphotropic virus 1
-	B04.613.807.200.725.750 Simian T-lymphotropic virus 1
-	B04.613.807.200.730 Primate T-lymphotropic virus 2
-	B04.613.807.200.730.399 Human T-lymphotropic virus 2
-	B04.613.807.200.730.810 Simian T-lymphotropic virus 2
-	B04.613.807.200.735 Primate T-lymphotropic virus 3
-	B04.613.807.200.735.374 Human T-lymphotropic virus 3
-	B04.613.807.200.735.750 Simian T-lymphotropic virus 3
-	B04.613.807.250 Endogenous Retroviruses
-	B04.613.807.260 Epsilonretrovirus
-	B04.613.807.375 Gammaretrovirus
-	B04.613.807.375.500 Leukemia Virus, Feline
-	B04.613.807.375.510 Leukemia Virus, Gibbon Ape
-	B04.613.807.375.525 Leukemia Virus, Murine
-	B04.613.807.375.525.020 Abelson murine leukemia virus
-	B04.613.807.375.525.050 AKR murine leukemia virus
-	B04.613.807.375.525.225 Friend murine leukemia virus
-	B04.613.807.375.525.580 Mink Cell Focus-Inducing Viruses
-	B04.613.807.375.525.596 Moloney murine leukemia virus
-	B04.613.807.375.525.750 Radiation Leukemia Virus
-	B04.613.807.375.525.770 Rauscher Virus
-	B04.613.807.375.525.850 Spleen Focus-Forming Viruses
-	B04.613.807.375.700 Reticuloendotheliosis Viruses, Avian
-	B04.613.807.375.700.700 Reticuloendotheliosis virus
-	B04.613.807.375.770 Sarcoma Virus, Woolly Monkey
-	B04.613.807.375.800 Sarcoma Viruses, Feline
-	B04.613.807.375.860 Sarcoma Viruses, Murine
-	B04.613.807.375.860.400 Harvey murine sarcoma virus
-	B04.613.807.375.860.500 Kirsten murine sarcoma virus
-	B04.613.807.375.860.600 Moloney murine sarcoma virus
-	B04.613.807.805 Retroviruses, Simian
-	B04.613.807.805.510 Leukemia Virus, Gibbon Ape
-	B04.613.807.805.550 Mason-Pfizer monkey virus

## MeSH Tree Changes for 2017

Type	Tree - heading
-	B04.613.807.805.601 Sarcoma Virus, Woolly Monkey
-	B04.613.807.805.851 Simian T-lymphotropic virus 1
-	B04.613.807.805.855 Simian T-lymphotropic virus 2
-	B04.613.807.805.858 Simian T-lymphotropic virus 3
-	B04.700 Oncolytic Viruses
-	B04.715 Plant Viruses
-	B04.715.081 Bromoviridae
-	B04.715.081.050 Alfamovirus
-	B04.715.081.050.050 Alfalfa mosaic virus
-	B04.715.081.080 Bromovirus
-	B04.715.081.180 Cucumovirus
-	B04.715.081.400 Ilarvirus
-	B04.715.081.700 Oleavirus
-	B04.715.102 Caulimoviridae
-	B04.715.102.074 Badnavirus
-	B04.715.102.150 Caulimovirus
-	B04.715.102.850 Tungrovirus
-	B04.715.110 Closteroviridae
-	B04.715.110.150 Closterovirus
-	B04.715.110.175 Crinivirus
-	B04.715.150 Comoviridae
-	B04.715.150.150 Comovirus
-	B04.715.150.250 Fabavirus
-	B04.715.150.500 Nepovirus
-	B04.715.260 Flexiviridae
-	B04.715.260.150 Carlavirus
-	B04.715.260.575 Potexvirus
-	B04.715.270 Geminiviridae
-	B04.715.270.100 Begomovirus
-	B04.715.270.500 Maize streak virus
-	B04.715.435 Luteoviridae
-	B04.715.435.500 Luteovirus
-	B04.715.464 Mosaic Viruses
-	B04.715.464.040 Alfamovirus
-	B04.715.464.040.050 Alfalfa mosaic virus
-	B04.715.464.080 Bromovirus

## MeSH Tree Changes for 2017

Type	Tree - heading
-	B04.715.464.100 Caulimovirus
-	B04.715.464.150 Comovirus
-	B04.715.464.180 Cucumovirus
-	B04.715.464.600 Potyvirus
-	B04.715.464.600.600 Plum Pox Virus
-	B04.715.464.725 Tobamovirus
-	B04.715.464.725.800 Tobacco Mosaic Virus
-	B04.715.464.750 Tymovirus
-	B04.715.499 Nanoviridae
-	B04.715.499.100 Babuvirus
-	B04.715.499.500 Nanovirus
-	B04.715.600 Phycodnaviridae
-	B04.715.635 Potyviridae
-	B04.715.635.600 Potyvirus
-	B04.715.635.600.600 Plum Pox Virus
-	B04.715.750 Sequiviridae
-	B04.715.750.750 Sequivirus
-	B04.715.750.900 Waikavirus
-	B04.715.780 Tenuivirus
-	B04.715.810 Tombusviridae
-	B04.715.810.150 Carmovirus
-	B04.715.810.870 Tombusvirus
-	B04.715.820 Tospovirus
-	B04.715.850 Tymoviridae
-	B04.715.850.860 Tymovirus
-	B04.725 Proviruses
-	B04.800 Reassortant Viruses
-	B04.820 RNA Viruses
-	B04.820.057 Arenaviridae
-	B04.820.057.070 Arenavirus
-	B04.820.057.070.100 Arenaviruses, Old World
-	B04.820.057.070.100.500 Lassa virus
-	B04.820.057.070.100.525 Lujo virus
-	B04.820.057.070.100.550 Lymphocytic choriomeningitis virus
-	B04.820.057.070.800 Arenaviruses, New World
-	B04.820.057.070.800.450 Junin virus

## MeSH Tree Changes for 2017

Type	Tree - heading
-	B04.820.057.070.800.650 Pichinde virus
-	B04.820.069 Astroviridae
-	B04.820.069.080 Avastrovirus
-	B04.820.069.500 Mamastrovirus
-	B04.820.075 Birnaviridae
-	B04.820.075.050 Aquabirnavirus
-	B04.820.075.050.430 Infectious pancreatic necrosis virus
-	B04.820.075.060 Avibirnavirus
-	B04.820.075.060.400 Infectious bursal disease virus
-	B04.820.075.200 Entomobirnavirus
-	B04.820.081 Bromoviridae
-	B04.820.081.050 Alfamovirus
-	B04.820.081.050.050 Alfalfa mosaic virus
-	B04.820.081.080 Bromovirus
-	B04.820.081.180 Cucumovirus
-	B04.820.081.400 Ilarvirus
-	B04.820.081.700 Oleavirus
-	B04.820.087 Bunyaviridae
-	B04.820.087.440 Hantavirus
-	B04.820.087.440.400 Hantaan virus
-	B04.820.087.440.600 Puumala virus
-	B04.820.087.440.690 Seoul virus
-	B04.820.087.440.700 Sin Nombre virus
-	B04.820.087.565 Nairovirus
-	B04.820.087.565.400 Hemorrhagic Fever Virus, Crimean-Congo
-	B04.820.087.565.600 Nairobi sheep disease virus
-	B04.820.087.640 Orthobunyavirus
-	B04.820.087.640.147 Bunyamwera virus
-	B04.820.087.640.300 Encephalitis Virus, California
-	B04.820.087.640.300.500 La Crosse virus
-	B04.820.087.640.855 Simbu virus
-	B04.820.087.710 Phlebovirus
-	B04.820.087.710.700 Rift Valley fever virus
-	B04.820.087.710.750 Sandfly fever Naples virus
-	B04.820.087.710.900 Uukuniemi virus
-	B04.820.087.855 Tospovirus

## MeSH Tree Changes for 2017

Type	Tree - heading
-	B04.820.095                      Caliciviridae
-	B04.820.095.500                  Lagovirus
-	B04.820.095.500.380              Hemorrhagic Disease Virus, Rabbit
-	B04.820.095.550                  Norovirus
-	B04.820.095.550.500              Norwalk virus
-	B04.820.095.775                  Sapovirus
-	B04.820.095.887                  Vesivirus
-	B04.820.095.887.150              Calicivirus, Feline
-	B04.820.095.887.900              Vesicular exanthema of swine virus
-	B04.820.110                      Closteroviridae
-	B04.820.110.150                  Closterovirus
-	B04.820.110.175                  Crinivirus
-	B04.820.150                      Comoviridae
-	B04.820.150.150                  Comovirus
-	B04.820.150.250                  Fabavirus
-	B04.820.150.500                  Nepovirus
-	B04.820.180                      Cystoviridae
-	B04.820.180.070                  Bacteriophage phi 6
-	B04.820.215                      Dicistroviridae
-	B04.820.230                      Encephalitis Viruses
-	B04.820.230.100                  Encephalitis Virus, California
-	B04.820.230.100.500              La Crosse virus
-	B04.820.230.150                  Encephalitis Virus, Eastern Equine
-	B04.820.230.369                  Encephalitis Virus, Venezuelan Equine
-	B04.820.230.440                  Encephalitis Virus, Western Equine
-	B04.820.230.475                  Encephalitis Viruses, Japanese
-	B04.820.230.475.227              Encephalitis Virus, Japanese
-	B04.820.230.475.300              Encephalitis Virus, Murray Valley
-	B04.820.230.475.330              Encephalitis Virus, St. Louis
-	B04.820.230.475.950              West Nile virus
-	B04.820.230.511                  Encephalitis Viruses, Tick-Borne
-	B04.820.250                      Flaviviridae
-	B04.820.250.350                  Flavivirus
-	B04.820.250.350.270              Dengue Virus
-	B04.820.250.350.300              Encephalitis Viruses, Japanese
-	B04.820.250.350.300.227              Encephalitis Virus, Japanese

## MeSH Tree Changes for 2017

Type	Tree - heading
-	B04.820.250.350.300.300                      Encephalitis Virus, Murray Valley
-	B04.820.250.350.300.330                      Encephalitis Virus, St. Louis
-	B04.820.250.350.300.950                      West Nile virus
-	B04.820.250.350.350                          Encephalitis Viruses, Tick-Borne
-	B04.820.250.350.990                          Yellow fever virus
-	B04.820.250.350.995                          Zika Virus
-	B04.820.250.400                              GB virus A
-	B04.820.250.410                              GB virus C
-	B04.820.250.475                              Hepacivirus
-	B04.820.250.475.405                          GB virus B
-	B04.820.250.700                              Pestivirus
-	B04.820.250.700.100                          Border disease virus
-	B04.820.250.700.150                          Diarrhea Viruses, Bovine Viral
-	B04.820.250.700.150.100                      Diarrhea Virus 1, Bovine Viral
-	B04.820.250.700.150.120                      Diarrhea Virus 2, Bovine Viral
-	B04.820.250.700.400                          Classical swine fever virus
-	B04.820.260                                  Flexiviridae
-	B04.820.260.150                              Carlavirus
-	B04.820.260.575                              Potexvirus
-	B04.820.300                                  Hepatitis Delta Virus
-	B04.820.340                                  Hepevirus
-	B04.820.340.400                              Hepatitis E virus
-	B04.820.410                                  Leviviridae
-	B04.820.410.050                              Allolevivirus
-	B04.820.410.500                              Levivirus
-	B04.820.420                                  Luteoviridae
-	B04.820.420.500                              Luteovirus
-	B04.820.455                                  Mononegavirales
-	B04.820.455.149                              Bornaviridae
-	B04.820.455.149.135                          Borna disease virus
-	B04.820.455.300                              Filoviridae
-	B04.820.455.300.200                          Ebolavirus
-	B04.820.455.300.650                          Marburgvirus
-	B04.820.455.600                              Paramyxoviridae
-	B04.820.455.600.650                          Paramyxovirinae
-	B04.820.455.600.650.070                      Avulavirus

## MeSH Tree Changes for 2017

Type	Tree - heading
-	B04.820.455.600.650.070.600 Newcastle disease virus
-	B04.820.455.600.650.400 Henipavirus
-	B04.820.455.600.650.400.400 Hendra Virus
-	B04.820.455.600.650.400.550 Nipah Virus
-	B04.820.455.600.650.500 Morbillivirus
-	B04.820.455.600.650.500.280 Distemper Virus, Canine
-	B04.820.455.600.650.500.320 Distemper Virus, Phocine
-	B04.820.455.600.650.500.500 Measles virus
-	B04.820.455.600.650.500.500.890 SSPE Virus
-	B04.820.455.600.650.500.650 Peste-des-petits-ruminants virus
-	B04.820.455.600.650.500.675 Rinderpest virus
-	B04.820.455.600.650.700 Respirovirus
-	B04.820.455.600.650.700.700 Parainfluenza Virus 3, Bovine
-	B04.820.455.600.650.700.715 Parainfluenza Virus 1, Human
-	B04.820.455.600.650.700.735 Parainfluenza Virus 3, Human
-	B04.820.455.600.650.700.800 Sendai virus
-	B04.820.455.600.650.750 Rubulavirus
-	B04.820.455.600.650.750.550 Mumps virus
-	B04.820.455.600.650.750.715 Parainfluenza Virus 2, Human
-	B04.820.455.600.650.750.717 Parainfluenza Virus 4, Human
-	B04.820.455.600.650.750.800 Parainfluenza Virus 5
-	B04.820.455.600.670 Pneumovirinae
-	B04.820.455.600.670.500 Metapneumovirus
-	B04.820.455.600.670.600 Pneumovirus
-	B04.820.455.600.670.600.550 Murine pneumonia virus
-	B04.820.455.600.670.600.750 Respiratory Syncytial Viruses
-	B04.820.455.600.670.600.750.700 Respiratory Syncytial Virus, Bovine
-	B04.820.455.600.670.600.750.730 Respiratory Syncytial Virus, Human
-	B04.820.455.750 Rhabdoviridae
-	B04.820.455.750.250 Ephemerovirus
-	B04.820.455.750.250.200 Ephemeral Fever Virus, Bovine
-	B04.820.455.750.500 Lyssavirus
-	B04.820.455.750.500.700 Rabies virus
-	B04.820.455.750.600 Novirhabdovirus
-	B04.820.455.750.600.400 Infectious hematopoietic necrosis virus
-	B04.820.455.750.900 Vesiculovirus



## MeSH Tree Changes for 2017

Type	Tree - heading
-	B04.820.455.750.900.900 Vesicular stomatitis Indiana virus
-	B04.820.455.750.900.910 Vesicular stomatitis New Jersey virus
-	B04.820.504 Nidovirales
-	B04.820.504.080 Arteriviridae
-	B04.820.504.080.500 Arterivirus
-	B04.820.504.080.500.100 Arteritis Virus, Equine
-	B04.820.504.080.500.500 Lactate dehydrogenase-elevating virus
-	B04.820.504.080.500.800 Porcine respiratory and reproductive syndrome virus
-	B04.820.504.540 Coronaviridae
-	B04.820.504.540.150 Coronavirus
-	B04.820.504.540.150.150 Coronavirus, Bovine
-	B04.820.504.540.150.160 Coronavirus, Canine
-	B04.820.504.540.150.190 Coronavirus, Feline
-	B04.820.504.540.150.210 Coronavirus 229E, Human
-	B04.820.504.540.150.215 Coronavirus NL63, Human
-	B04.820.504.540.150.220 Coronavirus OC43, Human
-	B04.820.504.540.150.260 Coronavirus, Rat
-	B04.820.504.540.150.300 Coronavirus, Turkey
-	B04.820.504.540.150.500 Infectious bronchitis virus
-	B04.820.504.540.150.540 Middle East Respiratory Syndrome Coronavirus
-	B04.820.504.540.150.580 Murine hepatitis virus
-	B04.820.504.540.150.665 Porcine epidemic diarrhea virus
-	B04.820.504.540.150.750 SARS Virus
-	B04.820.504.540.150.800 Transmissible gastroenteritis virus
-	B04.820.504.540.150.800.600 Porcine Respiratory Coronavirus
-	B04.820.504.540.800 Torovirus
-	B04.820.504.770 Roniviridae
-	B04.820.515 Nodaviridae
-	B04.820.545 Orthomyxoviridae
-	B04.820.545.405 Influenzavirus A
-	B04.820.545.405.400 Influenza A virus
-	B04.820.545.405.400.106 Influenza A Virus, H10N8 Subtype
-	B04.820.545.405.400.214 Influenza A Virus, H1N1 Subtype
-	B04.820.545.405.400.216 Influenza A Virus, H1N2 Subtype
-	B04.820.545.405.400.250 Influenza A Virus, H2N2 Subtype
-	B04.820.545.405.400.300 Influenza A Virus, H3N2 Subtype

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	B04.820.545.405.400.350	Influenza A Virus, H3N8 Subtype
-	B04.820.545.405.400.500	Influenza A Virus, H5N1 Subtype
-	B04.820.545.405.400.505	Influenza A Virus, H5N2 Subtype
New Heading	<b>B04.820.545.405.400.598</b>	<b>Influenza A Virus, H5N8 Subtype</b>
-	B04.820.545.405.400.690	Influenza A Virus, H7N1 Subtype
-	B04.820.545.405.400.692	Influenza A Virus, H7N2 Subtype
-	B04.820.545.405.400.694	Influenza A Virus, H7N3 Subtype
-	B04.820.545.405.400.700	Influenza A Virus, H7N7 Subtype
-	B04.820.545.405.400.800	Influenza A Virus, H7N9 Subtype
-	B04.820.545.405.400.900	Influenza A Virus, H9N2 Subtype
-	B04.820.545.405.400.920	Influenza A Virus, H10N7 Subtype
-	B04.820.545.407	Influenzavirus B
-	B04.820.545.407.410	Influenza B virus
-	B04.820.545.410	Influenzavirus C
-	B04.820.545.450	Isavirus
-	B04.820.545.850	Thogotovirus
-	B04.820.562	Picobirnavirus
-	B04.820.565	Picornaviridae
-	B04.820.565.070	Aphthovirus
-	B04.820.565.070.250	Foot-and-Mouth Disease Virus
-	B04.820.565.170	Cardiovirus
-	B04.820.565.170.200	Encephalomyocarditis virus
-	B04.820.565.170.200.150	Columbia SK virus
-	B04.820.565.170.200.480	Maus Elberfeld virus
-	B04.820.565.170.200.500	Mengovirus
-	B04.820.565.170.800	Theilovirus
-	B04.820.565.284	Enterovirus
-	B04.820.565.284.180	Enterovirus A, Human
-	B04.820.565.284.182	Enterovirus B, Human
-	B04.820.565.284.182.225	Echovirus 6, Human
-	B04.820.565.284.182.250	Echovirus 9
-	B04.820.565.284.184	Enterovirus C, Human
-	B04.820.565.284.188	Enterovirus D, Human
-	B04.820.565.284.200	Enterovirus, Bovine
-	B04.820.565.284.600	Enteroviruses, Porcine

## MeSH Tree Changes for 2017

Type	Tree - heading
-	B04.820.565.284.689 Poliovirus
-	B04.820.565.290 Erbovirus
-	B04.820.565.342 Hepatitis Virus, Duck
-	B04.820.565.400 Hepatovirus
-	B04.820.565.400.250 Encephalomyelitis Virus, Avian
-	B04.820.565.400.410 Hepatitis A virus
-	B04.820.565.400.410.500 Hepatitis A Virus, Human
-	B04.820.565.500 Kobuvirus
-	B04.820.565.700 Parechovirus
-	B04.820.565.775 Rhinovirus
-	B04.820.565.830 Teschovirus
-	B04.820.590 Potyviridae
-	B04.820.590.600 Potyvirus
-	B04.820.590.600.600 Plum Pox Virus
-	B04.820.630 Reoviridae
-	B04.820.630.160 Coltivirus
-	B04.820.630.160.200 Colorado tick fever virus
-	B04.820.630.550 Orbivirus
-	B04.820.630.550.050 African Horse Sickness Virus
-	B04.820.630.550.100 Bluetongue virus
-	B04.820.630.550.400 Hemorrhagic Disease Virus, Epizootic
-	B04.820.630.550.750 Palyam Virus
-	B04.820.630.590 Orthoreovirus
-	B04.820.630.590.500 Orthoreovirus, Avian
-	B04.820.630.590.550 Orthoreovirus, Mammalian
-	B04.820.630.590.550.700 Mammalian orthoreovirus 3
-	B04.820.630.790 Rotavirus
-	B04.820.650 Retroviridae
-	B04.820.650.070 Alpharetrovirus
-	B04.820.650.070.100 Avian Leukosis Virus
-	B04.820.650.070.110 Avian Myeloblastosis Virus
-	B04.820.650.070.120 Avian Sarcoma Viruses
-	B04.820.650.070.120.700 Rous sarcoma virus
-	B04.820.650.124 Betaretrovirus
-	B04.820.650.124.400 Jaagsiekte sheep retrovirus
-	B04.820.650.124.500 Mammary Tumor Virus, Mouse

## MeSH Tree Changes for 2017

Type	Tree - heading
-	B04.820.650.124.520 Mason-Pfizer monkey virus
-	B04.820.650.200 Deltaretrovirus
-	B04.820.650.200.600 Leukemia Virus, Bovine
-	B04.820.650.200.725 Primate T-lymphotropic virus 1
-	B04.820.650.200.725.400 Human T-lymphotropic virus 1
-	B04.820.650.200.725.755 Simian T-lymphotropic virus 1
-	B04.820.650.200.730 Primate T-lymphotropic virus 2
-	B04.820.650.200.730.404 Human T-lymphotropic virus 2
-	B04.820.650.200.730.810 Simian T-lymphotropic virus 2
-	B04.820.650.200.735 Primate T-lymphotropic virus 3
-	B04.820.650.200.735.374 Human T-lymphotropic virus 3
-	B04.820.650.200.735.750 Simian T-lymphotropic virus 3
-	B04.820.650.250 Endogenous Retroviruses
-	B04.820.650.260 Epsilonretrovirus
-	B04.820.650.375 Gammaretrovirus
-	B04.820.650.375.500 Leukemia Virus, Feline
-	B04.820.650.375.510 Leukemia Virus, Gibbon Ape
-	B04.820.650.375.525 Leukemia Virus, Murine
-	B04.820.650.375.525.020 Abelson murine leukemia virus
-	B04.820.650.375.525.050 AKR murine leukemia virus
-	B04.820.650.375.525.225 Friend murine leukemia virus
-	B04.820.650.375.525.580 Mink Cell Focus-Inducing Viruses
-	B04.820.650.375.525.596 Moloney murine leukemia virus
-	B04.820.650.375.525.750 Radiation Leukemia Virus
-	B04.820.650.375.525.770 Rauscher Virus
-	B04.820.650.375.525.850 Spleen Focus-Forming Viruses
-	B04.820.650.375.700 Reticuloendotheliosis Viruses, Avian
-	B04.820.650.375.700.700 Reticuloendotheliosis virus
-	B04.820.650.375.700.850 Trager duck spleen necrosis virus
-	B04.820.650.375.770 Sarcoma Virus, Woolly Monkey
-	B04.820.650.375.800 Sarcoma Viruses, Feline
-	B04.820.650.375.860 Sarcoma Viruses, Murine
-	B04.820.650.375.860.400 Harvey murine sarcoma virus
-	B04.820.650.375.860.500 Kirsten murine sarcoma virus
-	B04.820.650.375.860.600 Moloney murine sarcoma virus
-	B04.820.650.375.930 Xenotropic murine leukemia virus-related virus

## MeSH Tree Changes for 2017

Type	Tree - heading
-	B04.820.650.589                      Lentivirus
-	B04.820.650.589.500                      Lentiviruses, Bovine
-	B04.820.650.589.500.400                      Immunodeficiency Virus, Bovine
-	B04.820.650.589.520                      Lentiviruses, Equine
-	B04.820.650.589.520.400                      Infectious Anemia Virus, Equine
-	B04.820.650.589.530                      Lentiviruses, Feline
-	B04.820.650.589.530.400                      Immunodeficiency Virus, Feline
-	B04.820.650.589.600                      Lentiviruses, Ovine-Caprine
-	B04.820.650.589.600.070                      Arthritis-Encephalitis Virus, Caprine
-	B04.820.650.589.600.900                      Visna-maedi virus
-	B04.820.650.589.650                      Lentiviruses, Primate
-	B04.820.650.589.650.350                      HIV
-	B04.820.650.589.650.350.400                      HIV-1
-	B04.820.650.589.650.350.410                      HIV-2
-	B04.820.650.589.650.800                      Simian Immunodeficiency Virus
-	B04.820.650.805                      Retroviruses, Simian
-	B04.820.650.805.510                      Leukemia Virus, Gibbon Ape
-	B04.820.650.805.550                      Mason-Pfizer monkey virus
-	B04.820.650.805.601                      Sarcoma Virus, Woolly Monkey
-	B04.820.650.805.700                      Simian Immunodeficiency Virus
-	B04.820.650.805.851                      Simian T-lymphotropic virus 1
-	B04.820.650.805.855                      Simian T-lymphotropic virus 2
-	B04.820.650.850                      Spumavirus
-	B04.820.650.850.750                      Simian foamy virus
-	B04.820.750                      Sequiviridae
-	B04.820.750.750                      Sequivirus
-	B04.820.750.900                      Waikavirus
-	B04.820.800                      Tenuivirus
-	B04.820.850                      Togaviridae
-	B04.820.850.054                      Alphavirus
-	B04.820.850.054.198                      Chikungunya virus
-	B04.820.850.054.320                      Encephalitis Virus, Eastern Equine
-	B04.820.850.054.340                      Encephalitis Virus, Venezuelan Equine
-	B04.820.850.054.360                      Encephalitis Virus, Western Equine
-	B04.820.850.054.586                      O'nyong-nyong Virus
-	B04.820.850.054.813                      Ross River virus

## MeSH Tree Changes for 2017

Type	Tree - heading
-	B04.820.850.054.840 Semliki forest virus
-	B04.820.850.054.860 Sindbis Virus
-	B04.820.850.700 Rubivirus
-	B04.820.850.700.700 Rubella virus
-	B04.820.860 Tombusviridae
-	B04.820.860.150 Carmovirus
-	B04.820.860.870 Tombusvirus
-	B04.820.880 Totiviridae
-	B04.820.880.400 Giardiavirus
-	B04.820.880.500 Leishmanivirus
-	B04.820.880.800 Totivirus
-	B04.820.900 Tymoviridae
-	B04.820.900.850 Tymovirus
-	B04.950 Virion
-	B04.950.500 Nucleocapsid
-	B04.950.500.250 Capsid
-	B04.956 Viroids
-	B04.970 Viruses, Unclassified
-	B04.970.270 Anelloviridae
-	B04.970.540 Nanovirus
-	B04.970.610 Picobirnavirus
-	B05 Organism Forms
-	B05.080 Aquatic Organisms
-	B05.080.500 Plankton
-	B05.080.500.600 Phytoplankton
-	B05.080.500.600.500 Microalgae
-	B05.080.500.980 Zooplankton
-	B05.080.750 Seaweed
-	B05.080.750.450 Kelp
-	B05.110 Atypical Bacterial Forms
-	B05.110.422 L Forms
-	B05.110.761 Spheroplasts
New Tree	<a href="#">B05.155</a> <a href="#">Blood-Borne Pathogens</a>
-	B05.200 Chimera
-	B05.200.750 Transplantation Chimera

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	B05.200.750.760	Radiation Chimera
-	B05.237	Endophytes
New Heading	<b>B05.256</b>	<b>Extremophiles</b>
-	B05.275	Hermaphroditic Organisms
-	B05.350	Lichens
-	B05.500	Life Cycle Stages
-	B05.500.500	Larva
-	B05.500.500.150	Cercaria
-	B05.500.500.150.500	Metacercariae
-	B05.500.650	Nymph
-	B05.500.675	Oocysts
-	B05.500.675.800	Sporozoites
-	B05.500.700	Pupa
-	B05.500.800	Schizonts
-	B05.500.800.500	Merozoites
-	B05.500.950	Trophozoites
New Tree	<b>B05.525</b>	<b>Microfilaria</b>
New Tree	<b>B05.525</b>	<b>Microfilariae</b>
-	B05.550	Mycorrhizae
-	B05.620	Organisms, Genetically Modified
-	B05.620.136	Animals, Genetically Modified
-	B05.620.600	Plants, Genetically Modified
New Heading	<b>B05.698</b>	<b>Sentinel Species</b>
-	B05.775	Spores
-	B05.775.700	Spores, Bacterial
-	B05.775.710	Spores, Fungal
-	B05.775.740	Spores, Protozoan
-	B05.775.740.600	Oocysts
-	B05.775.740.600.800	Sporozoites
-	B05.775.740.800	Schizonts
-	B05.775.740.800.500	Merozoites
-	B05.775.740.900	Trophozoites
-	C01	Bacterial Infections and Mycoses

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C01.252 Bacterial Infections
-	C01.252.100 Bacteremia
-	C01.252.100.375 Hemorrhagic Septicemia
-	C01.252.200 Central Nervous System Bacterial Infections
-	C01.252.200.450 Lyme Neuroborreliosis
-	C01.252.200.500 Meningitis, Bacterial
-	C01.252.200.500.400 Meningitis, Escherichia coli
-	C01.252.200.500.450 Meningitis, Haemophilus
-	C01.252.200.500.500 Meningitis, Listeria
-	C01.252.200.500.550 Meningitis, Meningococcal
-	C01.252.200.500.550.800 Waterhouse-Friderichsen Syndrome
-	C01.252.200.500.600 Meningitis, Pneumococcal
-	C01.252.200.500.800 Tuberculosis, Meningeal
-	C01.252.200.750 Neurosyphilis
-	C01.252.300 Endocarditis, Bacterial
-	C01.252.300.407 Endocarditis, Subacute Bacterial
-	C01.252.354 Eye Infections, Bacterial
-	C01.252.354.225 Conjunctivitis, Bacterial
-	C01.252.354.225.250 Conjunctivitis, Inclusion
-	C01.252.354.225.625 Ophthalmia Neonatorum
-	C01.252.354.225.800 Trachoma
-	C01.252.354.400 Hordeolum
-	C01.252.354.450 Keratoconjunctivitis, Infectious
-	C01.252.354.800 Tuberculosis, Ocular
-	C01.252.354.900 Uveitis, Suppurative
-	C01.252.354.900.675 Panophthalmitis
-	C01.252.377 Fournier Gangrene
-	C01.252.400 Gram-Negative Bacterial Infections
-	C01.252.400.050 Aliivibrio Infections
-	C01.252.400.054 Anaplasmataceae Infections
-	C01.252.400.054.050 Anaplasmosis
-	C01.252.400.054.160 Ehrlichiosis
-	C01.252.400.054.160.400 Heartwater Disease
-	C01.252.400.110 Bacteroidaceae Infections
-	C01.252.400.110.109 Bacteroides Infections
-	C01.252.400.126 Bartonellaceae Infections



## MeSH Tree Changes for 2017

Type	Tree - heading
-	C01.252.400.126.100                      Bartonella Infections
-	C01.252.400.126.100.050                      Angiomatosis, Bacillary
-	C01.252.400.126.100.150                      Cat-Scratch Disease
-	C01.252.400.126.100.800                      Trench Fever
-	C01.252.400.143                      Bordetella Infections
-	C01.252.400.143.740                      Whooping Cough
-	C01.252.400.155                      Borrelia Infections
-	C01.252.400.155.569                      Lyme Disease
-	C01.252.400.155.569.200                      Erythema Chronicum Migrans
-	C01.252.400.155.569.600                      Lyme Neuroborreliosis
-	C01.252.400.155.644                      Relapsing Fever
-	C01.252.400.167                      Brucellosis
-	C01.252.400.167.322                      Brucellosis, Bovine
-	C01.252.400.170                      Burkholderia Infections
-	C01.252.400.170.400                      Glanders
-	C01.252.400.170.531                      Melioidosis
-	C01.252.400.177                      Campylobacter Infections
-	C01.252.400.200                      Cat-Scratch Disease
-	C01.252.400.210                      Chlamydiaceae Infections
-	C01.252.400.210.210                      Chlamydia Infections
-	C01.252.400.210.210.124                      Chlamydial Pneumonia
-	C01.252.400.210.210.250                      Conjunctivitis, Inclusion
-	C01.252.400.210.210.490                      Lymphogranuloma Venereum
-	C01.252.400.210.210.800                      Trachoma
-	C01.252.400.210.250                      Chlamydophila Infections
-	C01.252.400.210.250.299                      Chlamydial Pneumonia
-	C01.252.400.210.250.600                      Psittacosis
-	C01.252.400.245                      Cytophagaceae Infections
-	C01.252.400.260                      Desulfovibrionaceae Infections
-	C01.252.400.310                      Enterobacteriaceae Infections
-	C01.252.400.310.229                      Dysentery, Bacillary
-	C01.252.400.310.330                      Escherichia coli Infections
-	C01.252.400.310.330.500                      Meningitis, Escherichia coli
-	C01.252.400.310.416                      Granuloma Inguinale
-	C01.252.400.310.503                      Klebsiella Infections
-	C01.252.400.310.503.650                      Rhinoscleroma

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C01.252.400.310.708 Proteus Infections
-	C01.252.400.310.821 Salmonella Infections
-	C01.252.400.310.821.438 Paratyphoid Fever
-	C01.252.400.310.821.606 Salmonella Food Poisoning
-	C01.252.400.310.821.706 Salmonella Infections, Animal
-	C01.252.400.310.821.873 Typhoid Fever
-	C01.252.400.310.850 Serratia Infections
-	C01.252.400.310.980 Yersinia Infections
-	C01.252.400.310.980.745 Plague
-	C01.252.400.310.980.780 Yersinia pseudotuberculosis Infections
-	C01.252.400.349 Flavobacteriaceae Infections
-	C01.252.400.388 Fusobacteriaceae Infections
-	C01.252.400.388.350 Fusobacterium Infections
-	C01.252.400.388.350.400 Gingivitis, Necrotizing Ulcerative
-	C01.252.400.388.350.700 Lemierre Syndrome
-	C01.252.400.466 Helicobacter Infections
-	C01.252.400.500 Legionellosis
-	C01.252.400.500.501 Legionnaires' Disease
-	C01.252.400.511 Leptospirosis
-	C01.252.400.511.739 Weil Disease
-	C01.252.400.560 Moraxellaceae Infections
-	C01.252.400.560.022 Acinetobacter Infections
-	C01.252.400.610 Mycoplasmatales Infections
-	C01.252.400.610.610 Mycoplasma Infections
-	C01.252.400.610.610.717 Pleuropneumonia, Contagious
-	C01.252.400.610.610.760 Pneumonia, Mycoplasma
-	C01.252.400.610.850 Ureaplasma Infections
-	C01.252.400.625 Neisseriaceae Infections
-	C01.252.400.625.391 Gonorrhea
-	C01.252.400.625.391.658 Ophthalmia Neonatorum
-	C01.252.400.625.549 Meningococcal Infections
-	C01.252.400.625.549.449 Meningitis, Meningococcal
-	C01.252.400.625.549.449.800 Waterhouse-Friderichsen Syndrome
-	C01.252.400.700 Pasteurellaceae Infections
-	C01.252.400.700.030 Actinobacillus Infections
-	C01.252.400.700.030.178 Actinobacillosis

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C01.252.400.700.433 Haemophilus Infections
-	C01.252.400.700.433.257 Chancroid
-	C01.252.400.700.433.615 Meningitis, Haemophilus
-	C01.252.400.700.662 Pasteurella Infections
-	C01.252.400.700.662.297 Hemorrhagic Septicemia
-	C01.252.400.700.675 Pasteurellosis, Pneumonic
-	C01.252.400.719 Piscirickettsiaceae Infections
-	C01.252.400.739 Pseudomonas Infections
-	C01.252.400.755 Q Fever
-	C01.252.400.771 Rat-Bite Fever
-	C01.252.400.780 Rickettsiaceae Infections
-	C01.252.400.780.600 Pneumonia, Rickettsial
-	C01.252.400.780.790 Rickettsia Infections
-	C01.252.400.780.790.125 Boutonneuse Fever
-	C01.252.400.780.790.700 Rocky Mountain Spotted Fever
-	C01.252.400.780.790.800 Typhus, Endemic Flea-Borne
-	C01.252.400.780.790.805 Typhus, Epidemic Louse-Borne
-	C01.252.400.780.850 Scrub Typhus
-	C01.252.400.825 Tick-Borne Diseases
-	C01.252.400.825.050 Anaplasmosis
-	C01.252.400.825.125 Boutonneuse Fever
-	C01.252.400.825.200 Ehrlichiosis
-	C01.252.400.825.300 Heartwater Disease
-	C01.252.400.825.480 Lyme Disease
-	C01.252.400.825.480.400 Erythema Chronicum Migrans
-	C01.252.400.825.480.700 Lyme Neuroborreliosis
-	C01.252.400.825.750 Relapsing Fever
-	C01.252.400.825.775 Rocky Mountain Spotted Fever
-	C01.252.400.840 Treponemal Infections
-	C01.252.400.840.558 Pinta
-	C01.252.400.840.744 Syphilis
-	C01.252.400.840.744.161 Chancre
-	C01.252.400.840.744.456 Neurosyphilis
-	C01.252.400.840.744.456.778 Tabes Dorsalis
-	C01.252.400.840.744.657 Syphilis, Cardiovascular
-	C01.252.400.840.744.725 Syphilis, Congenital

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C01.252.400.840.744.800 Syphilis, Cutaneous
-	C01.252.400.840.744.871 Syphilis, Latent
-	C01.252.400.840.892 Yaws
-	C01.252.400.939 Tularemia
-	C01.252.400.959 Vibrio Infections
-	C01.252.400.959.347 Cholera
-	C01.252.410 Gram-Positive Bacterial Infections
-	C01.252.410.040 Actinomycetales Infections
-	C01.252.410.040.137 Actinomycosis
-	C01.252.410.040.137.262 Actinomycosis, Cervicofacial
-	C01.252.410.040.137.631 Whipple Disease
-	C01.252.410.040.246 Corynebacterium Infections
-	C01.252.410.040.246.388 Diphtheria
-	C01.252.410.040.246.430 Erythrasma
-	C01.252.410.040.552 Mycobacterium Infections
-	C01.252.410.040.552.386 Leprosy
-	C01.252.410.040.552.386.775 Leprosy, Multibacillary
-	C01.252.410.040.552.386.775.500 Leprosy, Lepromatous
-	C01.252.410.040.552.386.850 Leprosy, Paucibacillary
-	C01.252.410.040.552.386.850.249 Leprosy, Borderline
-	C01.252.410.040.552.386.850.500 Leprosy, Tuberculoid
-	C01.252.410.040.552.475 Mycobacterium Infections, Nontuberculous
-	C01.252.410.040.552.475.247 Buruli Ulcer
-	C01.252.410.040.552.475.495 Mycobacterium avium-intracellulare Infection
-	C01.252.410.040.552.588 Paratuberculosis
-	C01.252.410.040.552.846 Tuberculosis
-	C01.252.410.040.552.846.122 Latent Tuberculosis
-	C01.252.410.040.552.846.246 Peritonitis, Tuberculous
-	C01.252.410.040.552.846.493 Tuberculoma
-	C01.252.410.040.552.846.493.400 Tuberculoma, Intracranial
-	C01.252.410.040.552.846.516 Tuberculosis, Avian
-	C01.252.410.040.552.846.538 Tuberculosis, Bovine
-	C01.252.410.040.552.846.561 Tuberculosis, Cardiovascular
-	C01.252.410.040.552.846.561.595 Pericarditis, Tuberculous
-	C01.252.410.040.552.846.570 Tuberculosis, Central Nervous System
-	C01.252.410.040.552.846.570.300 Tuberculoma, Intracranial

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C01.252.410.040.552.846.570.600 Tuberculosis, Meningeal
-	C01.252.410.040.552.846.583 Tuberculosis, Cutaneous
-	C01.252.410.040.552.846.583.260 Erythema Induratum
-	C01.252.410.040.552.846.583.470 Lupus Vulgaris
-	C01.252.410.040.552.846.606 Tuberculosis, Endocrine
-	C01.252.410.040.552.846.628 Tuberculosis, Gastrointestinal
-	C01.252.410.040.552.846.651 Tuberculosis, Hepatic
-	C01.252.410.040.552.846.696 Tuberculosis, Laryngeal
-	C01.252.410.040.552.846.719 Tuberculosis, Lymph Node
-	C01.252.410.040.552.846.719.500 King's Evil
-	C01.252.410.040.552.846.764 Tuberculosis, Miliary
-	C01.252.410.040.552.846.775 Tuberculosis, Multidrug-Resistant
-	C01.252.410.040.552.846.775.500 Extensively Drug-Resistant Tuberculosis
-	C01.252.410.040.552.846.786 Tuberculosis, Ocular
-	C01.252.410.040.552.846.809 Tuberculosis, Oral
-	C01.252.410.040.552.846.831 Tuberculosis, Osteoarticular
-	C01.252.410.040.552.846.831.722 Tuberculosis, Spinal
-	C01.252.410.040.552.846.877 Tuberculosis, Pleural
-	C01.252.410.040.552.846.877.405 Empyema, Tuberculous
-	C01.252.410.040.552.846.899 Tuberculosis, Pulmonary
-	C01.252.410.040.552.846.899.669 Silicotuberculosis
-	C01.252.410.040.552.846.922 Tuberculosis, Splenic
-	C01.252.410.040.552.846.944 Tuberculosis, Urogenital
-	C01.252.410.040.552.846.944.596 Tuberculosis, Female Genital
-	C01.252.410.040.552.846.944.721 Tuberculosis, Male Genital
-	C01.252.410.040.552.846.944.847 Tuberculosis, Renal
-	C01.252.410.040.692 Nocardia Infections
-	C01.252.410.040.692.606 Mycetoma
-	C01.252.410.090 Bacillaceae Infections
-	C01.252.410.090.072 Anthrax
-	C01.252.410.110 Bifidobacteriales Infections
-	C01.252.410.222 Clostridium Infections
-	C01.252.410.222.151 Botulism
-	C01.252.410.222.310 Enterocolitis, Pseudomembranous
-	C01.252.410.222.325 Enterotoxemia
-	C01.252.410.222.440 Gas Gangrene

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C01.252.410.222.864 Tetanus
-	C01.252.410.334 Erysipelothrix Infections
-	C01.252.410.334.329 Erysipeloid
-	C01.252.410.334.776 Swine Erysipelas
-	C01.252.410.514 Listeriosis
-	C01.252.410.514.533 Meningitis, Listeria
-	C01.252.410.868 Staphylococcal Infections
-	C01.252.410.868.675 Pneumonia, Staphylococcal
-	C01.252.410.868.806 Staphylococcal Food Poisoning
-	C01.252.410.868.820 Staphylococcal Skin Infections
-	C01.252.410.868.820.270 Furunculosis
-	C01.252.410.868.820.270.200 Carbuncle
-	C01.252.410.868.820.504 Impetigo
-	C01.252.410.868.820.770 Staphylococcal Scalded Skin Syndrome
-	C01.252.410.890 Streptococcal Infections
-	C01.252.410.890.210 Ecthyma
-	C01.252.410.890.240 Endocarditis, Subacute Bacterial
-	C01.252.410.890.328 Erysipelas
-	C01.252.410.890.485 Impetigo
-	C01.252.410.890.670 Pneumococcal Infections
-	C01.252.410.890.670.595 Meningitis, Pneumococcal
-	C01.252.410.890.670.750 Pneumonia, Pneumococcal
-	C01.252.410.890.731 Rheumatic Fever
-	C01.252.410.890.731.649 Rheumatic Heart Disease
-	C01.252.410.890.823 Scarlet Fever
-	C01.252.620 Pneumonia, Bacterial
-	C01.252.620.249 Chlamydial Pneumonia
-	C01.252.620.500 Pneumonia, Mycoplasma
-	C01.252.620.530 Pneumonia of Calves, Enzootic
-	C01.252.620.540 Pneumonia of Swine, Mycoplasmal
-	C01.252.620.550 Pneumonia, Pneumococcal
-	C01.252.620.600 Pneumonia, Rickettsial
-	C01.252.620.620 Pneumonia, Staphylococcal
-	C01.252.810 Sexually Transmitted Diseases, Bacterial
-	C01.252.810.201 Chancroid
-	C01.252.810.301 Chlamydia Infections

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C01.252.810.301.490 Lymphogranuloma Venereum
-	C01.252.810.401 Gonorrhea
-	C01.252.810.451 Granuloma Inguinale
-	C01.252.810.859 Syphilis
-	C01.252.825 Skin Diseases, Bacterial
-	C01.252.825.110 Actinomycosis, Cervicofacial
-	C01.252.825.150 Angiomatosis, Bacillary
-	C01.252.825.180 Digital Dermatitis
-	C01.252.825.210 Ecthyma
-	C01.252.825.260 Erysipelas
-	C01.252.825.310 Erythema Chronicum Migrans
-	C01.252.825.320 Erythrasma
-	C01.252.825.340 Fasciitis, Necrotizing
-	C01.252.825.360 Granuloma Inguinale
-	C01.252.825.420 Hidradenitis Suppurativa
-	C01.252.825.557 Mycetoma
-	C01.252.825.630 Pinta
-	C01.252.825.705 Rhinoscleroma
-	C01.252.825.770 Staphylococcal Skin Infections
-	C01.252.825.770.270 Furunculosis
-	C01.252.825.770.270.200 Carbuncle
-	C01.252.825.770.360 Impetigo
-	C01.252.825.770.770 Staphylococcal Scalded Skin Syndrome
-	C01.252.825.790 Syphilis, Cutaneous
-	C01.252.825.820 Tuberculosis, Cutaneous
-	C01.252.825.820.260 Erythema Induratum
-	C01.252.825.820.470 Lupus Vulgaris
-	C01.252.825.910 Yaws
-	C01.252.847 Spirochaetales Infections
-	C01.252.847.193 Borrelia Infections
-	C01.252.847.193.569 Lyme Disease
-	C01.252.847.193.569.200 Erythema Chronicum Migrans
-	C01.252.847.193.569.600 Lyme Neuroborreliosis
-	C01.252.847.193.644 Relapsing Fever
-	C01.252.847.511 Leptospirosis
-	C01.252.847.511.739 Weil Disease

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C01.252.847.840 Treponemal Infections
-	C01.252.847.840.558 Pinta
-	C01.252.847.840.744 Syphilis
-	C01.252.847.840.744.161 Chancre
-	C01.252.847.840.744.456 Neurosyphilis
-	C01.252.847.840.744.456.778 Tabes Dorsalis
-	C01.252.847.840.744.657 Syphilis, Cardiovascular
-	C01.252.847.840.744.725 Syphilis, Congenital
-	C01.252.847.840.744.800 Syphilis, Cutaneous
-	C01.252.847.840.744.871 Syphilis, Latent
-	C01.252.847.840.892 Yaws
-	C01.252.954 Vaginosis, Bacterial
-	C01.395 Central Nervous System Infections
-	C01.395.250 Brain Abscess
-	C01.395.250.500 Toxoplasmosis, Cerebral
-	C01.395.500 Central Nervous System Bacterial Infections
-	C01.539 Infection
-	C01.539.069 Aneurysm, Infected
-	C01.539.100 Arthritis, Infectious
-	C01.539.100.500 Arthritis, Reactive
-	C01.539.160 Bone Diseases, Infectious
-	C01.539.160.495 Osteomyelitis
-	C01.539.160.495.500 Mastoiditis
-	C01.539.160.495.750 Petrositis
-	C01.539.160.495.875 Pott Puffy Tumor
-	C01.539.160.595 Periostitis
-	C01.539.160.762 Spondylitis
-	C01.539.160.762.301 Discitis
-	C01.539.160.886 Tuberculosis, Osteoarticular
-	C01.539.160.886.722 Tuberculosis, Spinal
-	C01.539.190 Cardiovascular Infections
-	C01.539.190.249 Endocarditis, Bacterial
-	C01.539.190.249.407 Endocarditis, Subacute Bacterial
-	C01.539.190.500 Syphilis, Cardiovascular
-	C01.539.190.750 Tuberculosis, Cardiovascular
-	C01.539.190.750.595 Pericarditis, Tuberculous



## MeSH Tree Changes for 2017

Type	Tree - heading
-	C01.539.195 Catheter-Related Infections
-	C01.539.218 Coinfection
-	C01.539.221 Communicable Diseases
-	C01.539.221.500 Communicable Diseases, Emerging
-	C01.539.221.750 Waterborne Diseases
-	C01.539.234 Community-Acquired Infections
-	C01.539.248 Cross Infection
-	C01.539.248.500 Pneumonia, Ventilator-Associated
-	C01.539.375 Eye Infections
-	C01.539.375.177 Corneal Ulcer
-	C01.539.375.265 Endophthalmitis
-	C01.539.375.265.500 Uveitis, Suppurative
-	C01.539.375.354 Eye Infections, Bacterial
-	C01.539.375.354.220 Conjunctivitis, Bacterial
-	C01.539.375.354.220.250 Conjunctivitis, Inclusion
-	C01.539.375.354.220.625 Ophthalmia Neonatorum
-	C01.539.375.354.220.800 Trachoma
-	C01.539.375.354.400 Hordeolum
-	C01.539.375.354.450 Keratoconjunctivitis, Infectious
-	C01.539.375.354.800 Tuberculosis, Ocular
-	C01.539.375.354.900 Uveitis, Suppurative
-	C01.539.375.354.900.675 Panophthalmitis
-	C01.539.375.450 Eye Infections, Fungal
-	C01.539.375.450.900 Uveitis, Suppurative
-	C01.539.375.450.900.675 Panophthalmitis
-	C01.539.392 Focal Infection
-	C01.539.392.433 Focal Infection, Dental
-	C01.539.424 Gingivitis, Necrotizing Ulcerative
-	C01.539.463 Intraabdominal Infections
-	C01.539.463.099 Appendicitis
-	C01.539.463.298 Diverticulitis
-	C01.539.463.600 Peritonitis
-	C01.539.463.600.249 Peritonitis, Tuberculous
-	C01.539.463.600.500 Subphrenic Abscess
-	C01.539.463.850 Typhlitis
-	C01.539.503 Laboratory Infection

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	C01.539.535	Ludwig's Angina
-	C01.539.597	Opportunistic Infections
-	C01.539.597.050	AIDS-Related Opportunistic Infections
-	C01.539.597.880	Superinfection
-	C01.539.635	Pelvic Infection
-	C01.539.635.500	Pelvic Inflammatory Disease
-	C01.539.674	Pregnancy Complications, Infectious
-	C01.539.674.173	Abortion, Septic
-	C01.539.674.715	Puerperal Infection
-	C01.539.685	Prosthesis-Related Infections
-	C01.539.730	Reproductive Tract Infections
-	C01.539.739	Respiratory Tract Infections
-	C01.539.739.484	Empyema, Pleural
-	C01.539.739.484.320	Empyema, Tuberculous
-	C01.539.739.969	Whooping Cough
-	C01.539.757	Sepsis
-	C01.539.757.100	Bacteremia
-	C01.539.757.100.275	Endotoxemia
-	C01.539.757.100.375	Hemorrhagic Septicemia
-	C01.539.757.360	Fungemia
-	C01.539.757.360.150	Candidemia
New Heading	<b>C01.539.757.580</b>	<b>Neonatal Sepsis</b>
-	C01.539.757.800	Shock, Septic
-	C01.539.778	Sexually Transmitted Diseases
-	C01.539.778.281	Sexually Transmitted Diseases, Bacterial
-	C01.539.778.281.201	Chancroid
-	C01.539.778.281.301	Chlamydia Infections
-	C01.539.778.281.301.490	Lymphogranuloma Venereum
-	C01.539.778.281.401	Gonorrhea
-	C01.539.778.281.451	Granuloma Inguinale
-	C01.539.778.281.859	Syphilis
-	C01.539.800	Skin Diseases, Infectious
-	C01.539.800.130	Cellulitis
-	C01.539.800.200	Dermatomycoses
-	C01.539.800.200.055	Blastomycosis

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	C01.539.800.200.100	Candidiasis, Chronic Mucocutaneous
-	C01.539.800.200.105	Candidiasis, Cutaneous
-	C01.539.800.200.110	Chromoblastomycosis
-	C01.539.800.200.383	Hyalohyphomycosis
-	C01.539.800.200.383.125	Alternariosis
Old Tree	C01.539.800.200.383.249	Aspergillosis
Old Tree	C01.539.800.200.383.249.074	Aspergillosis, Allergic Bronchopulmonary
Old Tree	C01.539.800.200.383.249.537	Neuroaspergillosis
-	C01.539.800.200.383.375	Cerebral Phaeohyphomycosis
-	C01.539.800.200.383.500	Fusariosis
-	C01.539.800.200.475	Lobomycosis
-	C01.539.800.200.500	Mycetoma
-	C01.539.800.200.675	Sporotrichosis
-	C01.539.800.200.720	Tinea
-	C01.539.800.200.720.550	Onychomycosis
-	C01.539.800.200.720.730	Tinea Capitis
-	C01.539.800.200.720.730.740	Tinea Favosa
-	C01.539.800.200.720.760	Tinea Pedis
-	C01.539.800.200.860	Tinea Versicolor
-	C01.539.800.460	Paronychia
-	C01.539.800.720	Skin Diseases, Bacterial
-	C01.539.800.720.110	Actinomycosis, Cervicofacial
-	C01.539.800.720.150	Angiomatosis, Bacillary
-	C01.539.800.720.180	Digital Dermatitis
-	C01.539.800.720.210	Ecthyma
-	C01.539.800.720.260	Erysipelas
-	C01.539.800.720.310	Erythema Chronicum Migrans
-	C01.539.800.720.320	Erythrasma
-	C01.539.800.720.360	Granuloma Inguinale
-	C01.539.800.720.420	Hidradenitis Suppurativa
-	C01.539.800.720.557	Mycetoma
-	C01.539.800.720.630	Pinta
-	C01.539.800.720.705	Rhinoscleroma
-	C01.539.800.720.770	Staphylococcal Skin Infections
-	C01.539.800.720.770.270	Furunculosis
-	C01.539.800.720.770.270.200	Carbuncle

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C01.539.800.720.770.360                      Impetigo
-	C01.539.800.720.770.770                      Staphylococcal Scalded Skin Syndrome
-	C01.539.800.720.790                      Syphilis, Cutaneous
-	C01.539.800.720.820                      Tuberculosis, Cutaneous
-	C01.539.800.720.820.260                      Erythema Induratum
-	C01.539.800.720.820.470                      Lupus Vulgaris
-	C01.539.800.720.910                      Yaws
-	C01.539.820                      Soft Tissue Infections
-	C01.539.830                      Suppuration
-	C01.539.830.025                      Abscess
-	C01.539.830.025.020                      Abdominal Abscess
-	C01.539.830.025.020.455                      Liver Abscess
-	C01.539.830.025.020.455.460                      Liver Abscess, Amebic
-	C01.539.830.025.020.455.730                      Liver Abscess, Pyogenic
-	C01.539.830.025.020.810                      Subphrenic Abscess
-	C01.539.830.025.160                      Brain Abscess
-	C01.539.830.025.160.800                      Toxoplasmosis, Cerebral
-	C01.539.830.025.325                      Epidural Abscess
-	C01.539.830.025.490                      Lung Abscess
-	C01.539.830.025.650                      Periapical Abscess
-	C01.539.830.025.665                      Periodontal Abscess
-	C01.539.830.025.675                      Peritonsillar Abscess
-	C01.539.830.025.700                      Psoas Abscess
-	C01.539.830.025.780                      Retropharyngeal Abscess
-	C01.539.830.200                      Cellulitis
-	C01.539.830.305                      Empyema
-	C01.539.830.305.310                      Empyema, Pleural
-	C01.539.830.305.310.320                      Empyema, Tuberculous
-	C01.539.830.305.330                      Empyema, Subdural
-	C01.539.830.499                      Hidradenitis Suppurativa
-	C01.539.830.694                      Otitis Media, Suppurative
-	C01.539.830.767                      Pyomyositis
-	C01.539.830.840                      Thyroiditis, Suppurative
-	C01.539.830.900                      Uveitis, Suppurative
-	C01.539.861                      Toxemia
-	C01.539.861.375                      Endotoxemia

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C01.539.895 Urinary Tract Infections
-	C01.539.895.219 Bacteriuria
-	C01.539.895.719 Pyuria
-	C01.539.895.775 Schistosomiasis haematobia
-	C01.539.947 Wound Infection
-	C01.539.947.692 Surgical Wound Infection
-	C01.703 Mycoses
New Tree	<a href="#">C01.703.080</a> <a href="#">Aspergillosis</a>
New Tree	<a href="#">C01.703.080.074</a> <a href="#">Aspergillosis, Allergic Bronchopulmonary</a>
New Tree	<a href="#">C01.703.080.537</a> <a href="#">Neuroaspergillosis</a>
New Tree	<a href="#">C01.703.080.768</a> <a href="#">Pulmonary Aspergillosis</a>
New Tree	<a href="#">C01.703.080.768.500</a> <a href="#">Aspergillosis, Allergic Bronchopulmonary</a>
New Tree	<a href="#">C01.703.080.768.750</a> <a href="#">Invasive Pulmonary Aspergillosis</a>
-	C01.703.160 Candidiasis
-	C01.703.160.165 Candidiasis, Chronic Mucocutaneous
-	C01.703.160.170 Candidiasis, Cutaneous
-	C01.703.160.175 Candidiasis, Invasive
-	C01.703.160.175.150 Candidemia
-	C01.703.160.180 Candidiasis, Oral
-	C01.703.160.190 Candidiasis, Vulvovaginal
-	C01.703.181 Central Nervous System Fungal Infections
-	C01.703.181.500 Meningitis, Fungal
-	C01.703.181.500.500 Meningitis, Cryptococcal
-	C01.703.203 Coccidioidomycosis
-	C01.703.248 Cryptococcosis
-	C01.703.248.290 Meningitis, Cryptococcal
-	C01.703.295 Dermatomycoses
-	C01.703.295.055 Blastomycosis
-	C01.703.295.165 Candidiasis, Chronic Mucocutaneous
-	C01.703.295.170 Candidiasis, Cutaneous
-	C01.703.295.182 Chromoblastomycosis
-	C01.703.295.328 Hyalohyphomycosis

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	C01.703.295.328.125	Alternariosis
-	C01.703.295.328.249	Aspergillosis
-	C01.703.295.328.249.074	Aspergillosis, Allergic Bronchopulmonary
-	C01.703.295.328.249.537	Neuroaspergillosis
-	C01.703.295.328.375	Cerebral Phaeohyphomycosis
-	C01.703.295.328.500	Fusariosis
-	C01.703.295.475	Lobomycosis
-	C01.703.295.522	Mycetoma
-	C01.703.295.675	Sporotrichosis
-	C01.703.295.872	Tinea
-	C01.703.295.872.458	Onychomycosis
-	C01.703.295.872.541	Tinea Capitis
-	C01.703.295.872.541.500	Tinea Favosa
-	C01.703.295.872.707	Tinea Pedis
-	C01.703.295.936	Tinea Versicolor
-	C01.703.343	Eye Infections, Fungal
-	C01.703.343.900	Uveitis, Suppurative
-	C01.703.343.900.675	Panophthalmitis
Old Tree	C01.703.360	Fungemia
Old Tree	C01.703.360.150	Candidemia
-	C01.703.392	Geotrichosis
-	C01.703.450	Histoplasmosis
New Heading	<b>C01.703.492</b>	<b>Invasive Fungal Infections</b>
New Tree	C01.703.492.500	Candidiasis, Invasive
New Tree	C01.703.492.500.150	Candidemia
New Tree	C01.703.492.625	Fungemia
New Tree	C01.703.492.625.150	Candidemia
New Tree	C01.703.492.688	Invasive Pulmonary Aspergillosis
New Tree	C01.703.492.750	Neuroaspergillosis
-	C01.703.534	Lung Diseases, Fungal
New Tree	C01.703.534.045	Aspergillosis

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	<a href="#">C01.703.534.045.074</a>	<a href="#">Aspergillosis, Allergic Bronchopulmonary</a>
New Tree	<a href="#">C01.703.534.045.537</a>	<a href="#">Neuroaspergillosis</a>
Old Tree	<del>C01.703.534.090</del>	<del>Aspergillosis, Allergic Bronchopulmonary</del>
-	C01.703.534.395	Blastomycosis
-	C01.703.534.700	Pneumonia, Pneumocystis
-	C01.703.617	Microsporidiosis
-	C01.703.617.300	Encephalitozoonosis
-	C01.703.658	Otomycosis
-	C01.703.700	Paracoccidioidomycosis
-	C01.703.726	Phaeohyphomycosis
-	C01.703.726.500	Cerebral Phaeohyphomycosis
-	C01.703.753	Piedra
-	C01.703.770	Pneumocystis Infections
-	C01.703.770.700	Pneumonia, Pneumocystis
-	C01.703.875	Trichosporonosis
-	C01.703.980	Zygomycosis
-	C01.703.980.600	Mucormycosis
-	C01.908	Zoonoses
-	C02	Virus Diseases
-	C02.081	Arbovirus Infections
-	C02.081.030	African Horse Sickness
-	C02.081.125	Bluetongue
-	C02.081.270	Dengue
-	C02.081.270.200	Severe Dengue
-	C02.081.343	Encephalitis, Arbovirus
-	C02.081.343.340	Encephalitis, California
-	C02.081.343.345	Encephalitis, Japanese
-	C02.081.343.350	Encephalitis, St. Louis
-	C02.081.343.360	Encephalitis, Tick-Borne
-	C02.081.343.950	West Nile Fever
-	C02.081.355	Encephalomyelitis, Equine
-	C02.081.355.177	Encephalomyelitis, Eastern Equine
-	C02.081.355.355	Encephalomyelitis, Venezuelan Equine
-	C02.081.355.677	Encephalomyelitis, Western Equine

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C02.081.700 Phlebotomus Fever
-	C02.081.810 Rift Valley Fever
-	C02.081.885 Tick-Borne Diseases
-	C02.081.885.125 African Swine Fever
-	C02.081.885.200 Colorado Tick Fever
-	C02.081.885.400 Encephalitis, Tick-Borne
-	C02.081.885.430 Hemorrhagic Fever, Crimean
-	C02.081.885.440 Hemorrhagic Fever, Omsk
-	C02.081.885.475 Kyasanur Forest Disease
-	C02.081.885.550 Nairobi Sheep Disease
-	C02.081.980 Yellow Fever
-	C02.081.990 Zika Virus Infection
-	C02.109 Bronchiolitis, Viral
-	C02.182 Central Nervous System Viral Diseases
-	C02.182.525 Encephalitis, Viral
-	C02.182.525.300 Encephalitis, Arbovirus
-	C02.182.525.300.200 Encephalitis, California
-	C02.182.525.300.250 Encephalitis, Japanese
-	C02.182.525.300.300 Encephalitis, St. Louis
-	C02.182.525.300.350 Encephalitis, Tick-Borne
-	C02.182.525.300.850 West Nile Fever
-	C02.182.525.350 Encephalitis, Herpes Simplex
-	C02.182.525.400 Encephalitis, Varicella Zoster
-	C02.182.525.450 Encephalomyelitis, Equine
-	C02.182.525.450.250 Encephalomyelitis, Venezuelan Equine
-	C02.182.525.450.300 Encephalomyelitis, Western Equine
-	C02.182.525.500 Leukoencephalopathy, Progressive Multifocal
-	C02.182.525.600 Subacute Sclerosing Panencephalitis
-	C02.182.550 Meningitis, Viral
-	C02.182.550.500 Lymphocytic Choriomeningitis
-	C02.182.710 Pseudorabies
-	C02.219 Coinfection
-	C02.256 DNA Virus Infections
-	C02.256.076 Adenoviridae Infections
-	C02.256.076.045 Adenovirus Infections, Human
-	C02.256.076.381 Hepatitis, Infectious Canine



## MeSH Tree Changes for 2017

Type	Tree - heading
-	C02.256.142 African Swine Fever
-	C02.256.200 Circoviridae Infections
-	C02.256.200.500 Porcine Postweaning Multisystemic Wasting Syndrome
-	C02.256.430 Hepadnaviridae Infections
-	C02.256.430.400 Hepatitis B
-	C02.256.430.400.100 Hepatitis B, Chronic
-	C02.256.466 Herpesviridae Infections
-	C02.256.466.087 Bell Palsy
-	C02.256.466.175 Chickenpox
-	C02.256.466.245 Cytomegalovirus Infections
-	C02.256.466.245.150 Cytomegalovirus Retinitis
-	C02.256.466.262 Encephalitis, Herpes Simplex
-	C02.256.466.279 Encephalitis, Varicella Zoster
-	C02.256.466.313 Epstein-Barr Virus Infections
-	C02.256.466.313.165 Burkitt Lymphoma
-	C02.256.466.313.400 Infectious Mononucleosis
-	C02.256.466.313.500 Leukoplakia, Hairy
-	C02.256.466.382 Herpes Simplex
-	C02.256.466.382.290 Herpes Genitalis
-	C02.256.466.382.316 Herpes Labialis
-	C02.256.466.382.410 Kaposi Varicelliform Eruption
-	C02.256.466.382.465 Keratitis, Herpetic
-	C02.256.466.382.465.450 Keratitis, Dendritic
-	C02.256.466.382.834 Stomatitis, Herpetic
-	C02.256.466.423 Herpes Zoster
-	C02.256.466.423.466 Herpes Zoster Ophthalmicus
-	C02.256.466.423.733 Herpes Zoster Oticus
-	C02.256.466.423.970 Zoster Sine Herpete
-	C02.256.466.488 Infectious Bovine Rhinotracheitis
-	C02.256.466.606 Malignant Catarrh
-	C02.256.466.650 Marek Disease
-	C02.256.466.793 Pseudorabies
-	C02.256.466.850 Roseolovirus Infections
-	C02.256.466.850.290 Exanthema Subitum
-	C02.256.466.860 Sarcoma, Kaposi
-	C02.256.650 Papillomavirus Infections

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C02.256.650.810 Warts
-	C02.256.650.810.217 Condylomata Acuminata
-	C02.256.650.810.217.500 Buschke-Lowenstein Tumor
-	C02.256.650.810.345 Epidermodysplasia Verruciformis
-	C02.256.700 Parvoviridae Infections
-	C02.256.700.091 Aleutian Mink Disease
-	C02.256.700.300 Erythema Infectiosum
-	C02.256.700.363 Feline Panleukopenia
-	C02.256.700.550 Mink Viral Enteritis
-	C02.256.721 Polyomavirus Infections
-	C02.256.721.150 Carcinoma, Merkel Cell
-	C02.256.721.500 Leukoencephalopathy, Progressive Multifocal
-	C02.256.743 Poxviridae Infections
-	C02.256.743.175 Cowpox
-	C02.256.743.193 Ecthyma, Contagious
-	C02.256.743.239 Ectromelia, Infectious
-	C02.256.743.366 Fowlpox
-	C02.256.743.494 Lumpy Skin Disease
-	C02.256.743.611 Molluscum Contagiosum
-	C02.256.743.615 Monkeypox
-	C02.256.743.665 Myxomatosis, Infectious
-	C02.256.743.826 Smallpox
-	C02.256.743.929 Vaccinia
-	C02.290 Encephalitis, Viral
-	C02.290.310 Encephalitis, Arbovirus
-	C02.290.310.140 Encephalitis, California
-	C02.290.310.280 Encephalitis, Japanese
-	C02.290.310.406 Encephalitis, St. Louis
-	C02.290.310.530 Encephalitis, Tick-Borne
-	C02.290.310.950 West Nile Fever
-	C02.290.325 Encephalitis, Herpes Simplex
-	C02.290.355 Encephalitis, Varicella Zoster
-	C02.290.450 Encephalomyelitis, Equine
-	C02.290.450.200 Encephalomyelitis, Eastern Equine
-	C02.290.450.225 Encephalomyelitis, Venezuelan Equine
-	C02.290.450.250 Encephalomyelitis, Western Equine

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C02.290.575 Leukoencephalopathy, Progressive Multifocal
-	C02.290.700 Subacute Sclerosing Panencephalitis
-	C02.325 Eye Infections, Viral
-	C02.325.250 Conjunctivitis, Viral
-	C02.325.250.250 Conjunctivitis, Acute Hemorrhagic
-	C02.325.270 Cytomegalovirus Retinitis
-	C02.325.450 Herpes Zoster Ophthalmicus
-	C02.325.465 Keratitis, Herpetic
-	C02.325.465.450 Keratitis, Dendritic
-	C02.330 Fatigue Syndrome, Chronic
-	C02.407 Hepatitis, Viral, Animal
-	C02.407.432 Hepatitis, Infectious Canine
-	C02.407.810 Rift Valley Fever
-	C02.440 Hepatitis, Viral, Human
-	C02.440.420 Hepatitis A
-	C02.440.435 Hepatitis B
-	C02.440.435.100 Hepatitis B, Chronic
-	C02.440.440 Hepatitis C
-	C02.440.440.120 Hepatitis C, Chronic
-	C02.440.450 Hepatitis D
-	C02.440.450.100 Hepatitis D, Chronic
-	C02.440.470 Hepatitis E
-	C02.597 Opportunistic Infections
-	C02.597.050 AIDS-Related Opportunistic Infections
-	C02.597.880 Superinfection
-	C02.705 Pneumonia, Viral
-	C02.782 RNA Virus Infections
-	C02.782.082 Arenaviridae Infections
-	C02.782.082.440 Hemorrhagic Fever, American
-	C02.782.082.545 Lassa Fever
-	C02.782.082.580 Lymphocytic Choriomeningitis
-	C02.782.105 Astroviridae Infections
-	C02.782.123 Birnaviridae Infections
-	C02.782.147 Bunyaviridae Infections
-	C02.782.147.340 Encephalitis, California
-	C02.782.147.420 Hantavirus Infections

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C02.782.147.420.380 Hantavirus Pulmonary Syndrome
-	C02.782.147.420.400 Hemorrhagic Fever with Renal Syndrome
-	C02.782.147.444 Hemorrhagic Fever, Crimean
-	C02.782.147.633 Nairobi Sheep Disease
-	C02.782.147.700 Phlebotomus Fever
-	C02.782.147.810 Rift Valley Fever
-	C02.782.160 Caliciviridae Infections
-	C02.782.160.927 Vesicular Exanthema of Swine
-	C02.782.310 Encephalitis, Arbovirus
-	C02.782.310.340 Encephalitis, California
-	C02.782.310.345 Encephalitis, Japanese
-	C02.782.310.350 Encephalitis, St. Louis
-	C02.782.310.360 Encephalitis, Tick-Borne
-	C02.782.310.950 West Nile Fever
-	C02.782.350 Flaviviridae Infections
-	C02.782.350.250 Flavivirus Infections
-	C02.782.350.250.214 Dengue
-	C02.782.350.250.214.200 Severe Dengue
-	C02.782.350.250.300 Encephalitis, Japanese
-	C02.782.350.250.450 Encephalitis, St. Louis
-	C02.782.350.250.500 Encephalitis, Tick-Borne
-	C02.782.350.250.560 Hemorrhagic Fever, Omsk
-	C02.782.350.250.635 Kyasanur Forest Disease
-	C02.782.350.250.650 Louping Ill
-	C02.782.350.250.900 West Nile Fever
-	C02.782.350.250.980 Yellow Fever
-	C02.782.350.250.990 Zika Virus Infection
-	C02.782.350.350 Hepatitis C
-	C02.782.350.350.120 Hepatitis C, Chronic
-	C02.782.350.675 Pestivirus Infections
-	C02.782.350.675.100 Border Disease
-	C02.782.350.675.106 Bovine Virus Diarrhea-Mucosal Disease
-	C02.782.350.675.200 Classical Swine Fever
-	C02.782.350.675.400 Hemorrhagic Syndrome, Bovine
-	C02.782.417 Hemorrhagic Fevers, Viral
-	C02.782.417.214 Dengue

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C02.782.417.214.200 Severe Dengue
-	C02.782.417.400 Hemorrhagic Fever, American
-	C02.782.417.412 Hemorrhagic Fever, Crimean
-	C02.782.417.415 Hemorrhagic Fever, Ebola
-	C02.782.417.435 Hemorrhagic Fever, Omsk
-	C02.782.417.450 Hemorrhagic Fever with Renal Syndrome
-	C02.782.417.475 Kyasanur Forest Disease
-	C02.782.417.505 Lassa Fever
-	C02.782.417.560 Marburg Virus Disease
-	C02.782.417.762 Rift Valley Fever
-	C02.782.417.881 Yellow Fever
-	C02.782.450 Hepatitis D
-	C02.782.450.100 Hepatitis D, Chronic
-	C02.782.455 Hepatitis E
-	C02.782.580 Mononegavirales Infections
-	C02.782.580.124 Borna Disease
-	C02.782.580.250 Filoviridae Infections
-	C02.782.580.250.400 Hemorrhagic Fever, Ebola
-	C02.782.580.250.500 Marburg Virus Disease
-	C02.782.580.600 Paramyxoviridae Infections
-	C02.782.580.600.080 Avulavirus Infections
-	C02.782.580.600.080.600 Newcastle Disease
-	C02.782.580.600.400 Henipavirus Infections
-	C02.782.580.600.500 Morbillivirus Infections
-	C02.782.580.600.500.285 Distemper
-	C02.782.580.600.500.500 Measles
-	C02.782.580.600.500.500.800 Subacute Sclerosing Panencephalitis
-	C02.782.580.600.500.600 Peste-des-Petits-Ruminants
-	C02.782.580.600.500.700 Rinderpest
-	C02.782.580.600.600 Respirovirus Infections
-	C02.782.580.600.600.648 Pasteurellosis, Pneumonic
-	C02.782.580.600.620 Pneumovirus Infections
-	C02.782.580.600.620.750 Respiratory Syncytial Virus Infections
-	C02.782.580.600.680 Rubulavirus Infections
-	C02.782.580.600.680.500 Mumps
-	C02.782.580.830 Rhabdoviridae Infections

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C02.782.580.830.375 Ephemeral Fever
-	C02.782.580.830.450 Hemorrhagic Septicemia, Viral
-	C02.782.580.830.750 Rabies
-	C02.782.580.830.825 Vesicular Stomatitis
-	C02.782.600 Nidovirales Infections
-	C02.782.600.100 Arterivirus Infections
-	C02.782.600.100.700 Porcine Reproductive and Respiratory Syndrome
-	C02.782.600.550 Coronaviridae Infections
-	C02.782.600.550.200 Coronavirus Infections
-	C02.782.600.550.200.325 Enteritis, Transmissible, of Turkeys
-	C02.782.600.550.200.360 Feline Infectious Peritonitis
-	C02.782.600.550.200.400 Gastroenteritis, Transmissible, of Swine
-	C02.782.600.550.200.750 Severe Acute Respiratory Syndrome
-	C02.782.600.550.800 Torovirus Infections
-	C02.782.620 Orthomyxoviridae Infections
-	C02.782.620.365 Influenza, Human
-	C02.782.620.375 Influenza in Birds
-	C02.782.687 Picornaviridae Infections
-	C02.782.687.150 Cardiovirus Infections
-	C02.782.687.207 Common Cold
-	C02.782.687.359 Enterovirus Infections
-	C02.782.687.359.201 Conjunctivitis, Acute Hemorrhagic
-	C02.782.687.359.213 Coxsackievirus Infections
-	C02.782.687.359.213.331 Hand, Foot and Mouth Disease
-	C02.782.687.359.213.466 Herpangina
-	C02.782.687.359.213.737 Pleurodynia, Epidemic
-	C02.782.687.359.347 Echovirus Infections
New Tree	<a href="#">C02.782.687.359.347.500</a> <a href="#">Herpangina</a>
-	C02.782.687.359.456 Encephalomyelitis, Enzootic Porcine
-	C02.782.687.359.500 Hepatitis A
-	C02.782.687.359.764 Poliomyelitis
-	C02.782.687.359.764.614 Poliomyelitis, Bulbar
-	C02.782.687.359.764.650 Postpoliomyelitis Syndrome
-	C02.782.687.359.855 Swine Vesicular Disease
-	C02.782.687.484 Foot-and-Mouth Disease

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C02.782.791 Reoviridae Infections
-	C02.782.791.142 African Horse Sickness
-	C02.782.791.315 Bluetongue
-	C02.782.791.482 Colorado Tick Fever
-	C02.782.791.814 Rotavirus Infections
-	C02.782.815 Retroviridae Infections
-	C02.782.815.096 Avian Leukosis
-	C02.782.815.200 Deltaretrovirus Infections
-	C02.782.815.200.260 Enzootic Bovine Leukosis
-	C02.782.815.200.470 HTLV-I Infections
-	C02.782.815.200.470.710 Paraparesis, Tropical Spastic
-	C02.782.815.200.480 HTLV-II Infections
-	C02.782.815.616 Lentivirus Infections
-	C02.782.815.616.300 Equine Infectious Anemia
-	C02.782.815.616.350 Feline Acquired Immunodeficiency Syndrome
-	C02.782.815.616.400 HIV Infections
-	C02.782.815.616.400.040 Acquired Immunodeficiency Syndrome
New Heading	<b>C02.782.815.616.400.044 Acute Retroviral Syndrome</b>
-	C02.782.815.616.400.048 AIDS Arteritis, Central Nervous System
-	C02.782.815.616.400.050 AIDS-Associated Nephropathy
-	C02.782.815.616.400.070 AIDS Dementia Complex
-	C02.782.815.616.400.080 AIDS-Related Complex
-	C02.782.815.616.400.100 AIDS-Related Opportunistic Infections
-	C02.782.815.616.400.400 HIV-Associated Lipodystrophy Syndrome
-	C02.782.815.616.400.480 HIV Enteropathy
-	C02.782.815.616.400.500 HIV Seropositivity
-	C02.782.815.616.400.520 HIV Wasting Syndrome
-	C02.782.815.616.660 Pneumonia, Progressive Interstitial, of Sheep
-	C02.782.815.616.850 Simian Acquired Immunodeficiency Syndrome
-	C02.782.815.616.900 Visna
-	C02.782.815.622 Leukemia, Feline
-	C02.782.815.650 Murine Acquired Immunodeficiency Syndrome
-	C02.782.815.725 Pulmonary Adenomatosis, Ovine
-	C02.782.815.800 Sarcoma, Avian
-	C02.782.930 Togaviridae Infections

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C02.782.930.100                      Alphavirus Infections
-	C02.782.930.100.184                      Chikungunya Fever
-	C02.782.930.100.370                      Encephalomyelitis, Equine
-	C02.782.930.100.370.162                      Encephalomyelitis, Eastern Equine
-	C02.782.930.100.370.325                      Encephalomyelitis, Venezuelan Equine
-	C02.782.930.100.370.662                      Encephalomyelitis, Western Equine
-	C02.782.930.700                      Rubivirus Infections
-	C02.782.930.700.700                      Rubella
-	C02.782.930.700.700.700                      Rubella Syndrome, Congenital
-	C02.800                      Sexually Transmitted Diseases
-	C02.800.801                      Sexually Transmitted Diseases, Viral
-	C02.800.801.220                      Condylomata Acuminata
-	C02.800.801.220.500                      Buschke-Lowenstein Tumor
-	C02.800.801.350                      Herpes Genitalis
-	C02.800.801.400                      HIV Infections
-	C02.800.801.400.040                      Acquired Immunodeficiency Syndrome
New Heading	<b>C02.800.801.400.044                      Acute Retroviral Syndrome</b>
-	C02.800.801.400.048                      AIDS Arteritis, Central Nervous System
-	C02.800.801.400.050                      AIDS-Associated Nephropathy
-	C02.800.801.400.070                      AIDS Dementia Complex
-	C02.800.801.400.080                      AIDS-Related Complex
-	C02.800.801.400.400                      HIV-Associated Lipodystrophy Syndrome
-	C02.800.801.400.480                      HIV Enteropathy
-	C02.800.801.400.500                      HIV Seropositivity
-	C02.800.801.400.520                      HIV Wasting Syndrome
-	C02.825                      Skin Diseases, Viral
-	C02.825.260                      Erythema Infectiosum
-	C02.825.290                      Exanthema Subitum
-	C02.825.320                      Herpes Simplex
-	C02.825.320.320                      Herpes Labialis
-	C02.825.320.410                      Kaposi Varicelliform Eruption
-	C02.825.550                      Molluscum Contagiosum
-	C02.825.810                      Warts
-	C02.825.810.110                      Condylomata Acuminata
-	C02.825.810.110.500                      Buschke-Lowenstein Tumor



## MeSH Tree Changes for 2017

Type	Tree - heading
-	C02.825.810.260                      Epidermodysplasia Verruciformis
-	C02.839                                      Slow Virus Diseases
-	C02.839.040                              Acquired Immunodeficiency Syndrome
-	C02.839.080                              AIDS-Related Complex
-	C02.839.091                              Aleutian Mink Disease
-	C02.839.375                              Equine Infectious Anemia
-	C02.839.400                              Feline Acquired Immunodeficiency Syndrome
-	C02.839.550                              Leukoencephalopathy, Progressive Multifocal
-	C02.839.660                              Pneumonia, Progressive Interstitial, of Sheep
-	C02.839.850                              Simian Acquired Immunodeficiency Syndrome
-	C02.839.862                              Subacute Sclerosing Panencephalitis
-	C02.839.900                              Visna
-	C02.928                                      Tumor Virus Infections
-	C02.928.120                              Avian Leukosis
-	C02.928.216                              Carcinoma, Merkel Cell
-	C02.928.313                              Epstein-Barr Virus Infections
-	C02.928.313.165                              Burkitt Lymphoma
-	C02.928.489                              Marek Disease
-	C02.928.650                              Murine Acquired Immunodeficiency Syndrome
-	C02.928.725                              Papillomavirus Infections
-	C02.928.740                              Pulmonary Adenomatosis, Ovine
-	C02.928.800                              Sarcoma, Avian
-	C02.928.914                              Warts
-	C02.928.914.217                              Condylomata Acuminata
-	C02.928.914.217.500                              Buschke-Lowenstein Tumor
-	C02.928.914.345                              Epidermodysplasia Verruciformis
-	C02.937                                      Viremia
-	C02.968                                      Zoonoses
-	C03    Parasitic Diseases
-	C03.105                                      Central Nervous System Parasitic Infections
-	C03.105.250                              Central Nervous System Helminthiasis
-	C03.105.250.550                              Neurocysticercosis
-	C03.105.250.600                              Neuroschistosomiasis
-	C03.105.300                              Central Nervous System Protozoal Infections
-	C03.105.300.500                              Malaria, Cerebral
-	C03.105.300.800                              Toxoplasmosis, Cerebral

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	C03.202	Coinfection
-	C03.300	Eye Infections, Parasitic
-	C03.300.125	Acanthamoeba Keratitis
-	C03.300.562	Onchocerciasis, Ocular
-	C03.300.781	Toxoplasmosis, Ocular
-	C03.335	Helminthiasis
-	C03.335.190	Cestode Infections
-	C03.335.190.304	Diphyllobothriasis
-	C03.335.190.304.780	Sparganosis
-	C03.335.190.396	Echinococcosis
-	C03.335.190.396.314	Echinococcosis, Hepatic
-	C03.335.190.396.480	Echinococcosis, Pulmonary
-	C03.335.190.519	Hymenolepiasis
-	C03.335.190.628	Monieziasis
-	C03.335.190.902	Taeniasis
-	C03.335.190.902.185	Cysticercosis
-	C03.335.190.902.185.550	Neurocysticercosis
-	C03.335.349	Helminthiasis, Animal
-	C03.335.349.315	Dictyocaulus Infections
-	C03.335.349.320	Dirofilariasis
-	C03.335.349.444	Fascioloidiasis
-	C03.335.349.656	Monieziasis
-	C03.335.349.820	Setariasis
-	C03.335.349.840	Strongyle Infections, Equine
-	C03.335.349.868	Toxocariasis
-	C03.335.508	Nematode Infections
-	C03.335.508.100	Adenophorea Infections
-	C03.335.508.100.275	Enoplida Infections
-	C03.335.508.100.275.882	Trichinellosis
-	C03.335.508.100.275.895	Trichuriasis
-	C03.335.508.523	Larva Migrans
-	C03.335.508.523.780	Larva Migrans, Visceral
-	C03.335.508.700	Secernentea Infections
-	C03.335.508.700.100	Ascaridida Infections
-	C03.335.508.700.100.060	Anisakiasis
-	C03.335.508.700.100.070	Ascariasis

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C03.335.508.700.100.080 Ascaridiasis
-	C03.335.508.700.100.850 Toxascariasis
-	C03.335.508.700.100.868 Toxocariasis
-	C03.335.508.700.100.868.420 Larva Migrans, Visceral
-	C03.335.508.700.550 Oxyurida Infections
-	C03.335.508.700.550.550 Oxyuriasis
-	C03.335.508.700.550.550.375 Enterobiasis
-	C03.335.508.700.700 Rhabditida Infections
-	C03.335.508.700.700.799 Strongyloidiasis
-	C03.335.508.700.750 Spirurida Infections
-	C03.335.508.700.750.299 Dracunculiasis
-	C03.335.508.700.750.361 Filariasis
-	C03.335.508.700.750.361.137 Acanthocheilonemiasis
-	C03.335.508.700.750.361.275 Dipetalonema Infections
-	C03.335.508.700.750.361.290 Dirofilariasis
-	C03.335.508.700.750.361.350 Elephantiasis, Filarial
-	C03.335.508.700.750.361.518 Loiasis
-	C03.335.508.700.750.361.588 Mansonelliasis
-	C03.335.508.700.750.361.699 Onchocerciasis
-	C03.335.508.700.750.361.699.500 Onchocerciasis, Ocular
-	C03.335.508.700.750.361.852 Setariasis
-	C03.335.508.700.750.380 Gnathostomiasis
-	C03.335.508.700.775 Strongylida Infections
-	C03.335.508.700.775.455 Hookworm Infections
-	C03.335.508.700.775.455.154 Ancylostomiasis
-	C03.335.508.700.775.455.683 Necatoriasis
-	C03.335.508.700.775.583 Oesophagostomiasis
-	C03.335.508.700.775.773 Strongyle Infections, Equine
-	C03.335.508.700.775.825 Trichostrongyloidiasis
-	C03.335.508.700.775.825.315 Dictyocaulus Infections
-	C03.335.508.700.775.825.400 Haemonchiasis
-	C03.335.508.700.775.825.580 Ostertagiasis
-	C03.335.508.700.775.825.842 Trichostrongylosis
-	C03.335.865 Trematode Infections
-	C03.335.865.148 Clonorchiasis
-	C03.335.865.224 Dicrocoeliasis

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C03.335.865.282 Echinostomiasis
-	C03.335.865.354 Fascioliasis
-	C03.335.865.399 Fascioloidiasis
-	C03.335.865.685 Opisthorchiasis
-	C03.335.865.741 Paragonimiasis
-	C03.335.865.859 Schistosomiasis
-	C03.335.865.859.213 Neuroschistosomiasis
-	C03.335.865.859.427 Schistosomiasis haematobia
-	C03.335.865.859.521 Schistosomiasis japonica
-	C03.335.865.859.576 Schistosomiasis mansoni
-	C03.432 Intestinal Diseases, Parasitic
-	C03.432.060 Anisakiasis
-	C03.432.146 Balantidiasis
-	C03.432.250 Blastocystis Infections
-	C03.432.269 Cryptosporidiosis
-	C03.432.275 Dientamoebiasis
-	C03.432.396 Dysentery, Amebic
-	C03.432.481 Giardiasis
-	C03.518 Liver Diseases, Parasitic
-	C03.518.314 Echinococcosis, Hepatic
-	C03.518.424 Fascioliasis
-	C03.518.600 Liver Abscess, Amebic
-	C03.582 Lung Diseases, Parasitic
-	C03.582.314 Echinococcosis, Pulmonary
-	C03.600 Mesomycetozoea Infections
-	C03.600.700 Rhinosporidiosis
-	C03.684 Opportunistic Infections
-	C03.684.050 AIDS-Related Opportunistic Infections
-	C03.684.880 Superinfection
-	C03.695 Parasitemia
-	C03.701 Parasitic Diseases, Animal
-	C03.701.377 Helminthiasis, Animal
-	C03.701.377.315 Dictyocaulus Infections
-	C03.701.377.320 Dirofilariasis
-	C03.701.377.444 Fascioloidiasis
-	C03.701.377.656 Monieziasis

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C03.701.377.820                      Setariasis
-	C03.701.377.840                      Strongyle Infections, Equine
-	C03.701.377.868                      Toxocariasis
-	C03.701.688                            Protozoan Infections, Animal
-	C03.701.688.122                      Babesiosis
-	C03.701.688.235                      Cryptosporidiosis
-	C03.701.688.367                      Dourine
-	C03.701.688.735                      Theileriasis
-	C03.701.688.817                      Toxoplasmosis, Animal
-	C03.701.688.896                      Trypanosomiasis, Bovine
-	C03.718                                 Pregnancy Complications, Parasitic
-	C03.752                                 Protozoan Infections
-	C03.752.049                            Amebiasis
-	C03.752.049.203                      Acanthamoeba Keratitis
-	C03.752.049.250                      Blastocystis Infections
-	C03.752.049.328                      Dysentery, Amebic
-	C03.752.049.407                      Entamoebiasis
-	C03.752.049.703                      Liver Abscess, Amebic
-	C03.752.100                            Central Nervous System Protozoal Infections
-	C03.752.200                            Ciliophora Infections
-	C03.752.200.146                      Balantidiasis
-	C03.752.250                            Coccidiosis
-	C03.752.250.269                      Cryptosporidiosis
-	C03.752.250.280                      Cyclosporiasis
-	C03.752.250.410                      Isosporiasis
-	C03.752.250.634                      Sarcocystosis
-	C03.752.250.800                      Toxoplasmosis
-	C03.752.250.800.110                 Toxoplasmosis, Animal
-	C03.752.250.800.250                 Toxoplasmosis, Cerebral
-	C03.752.250.800.445                 Toxoplasmosis, Congenital
-	C03.752.250.800.640                 Toxoplasmosis, Ocular
-	C03.752.275                            Dientamoebiasis
-	C03.752.300                            Euglenozoa Infections
-	C03.752.300.500                      Leishmaniasis
-	C03.752.300.500.400                 Leishmaniasis, Cutaneous
-	C03.752.300.500.400.350            Leishmaniasis, Diffuse Cutaneous

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C03.752.300.500.400.385 Leishmaniasis, Mucocutaneous
-	C03.752.300.500.510 Leishmaniasis, Visceral
-	C03.752.300.900 Trypanosomiasis
-	C03.752.300.900.200 Chagas Disease
-	C03.752.300.900.200.190 Chagas Cardiomyopathy
-	C03.752.300.900.226 Dourine
-	C03.752.300.900.719 Trypanosomiasis, African
-	C03.752.300.900.802 Trypanosomiasis, Bovine
-	C03.752.400 Giardiasis
-	C03.752.530 Malaria
-	C03.752.530.606 Malaria, Avian
-	C03.752.530.620 Malaria, Cerebral
-	C03.752.530.650 Malaria, Falciparum
-	C03.752.530.650.228 Blackwater Fever
-	C03.752.530.700 Malaria, Vivax
-	C03.752.625 Protozoan Infections, Animal
-	C03.752.625.122 Babesiosis
-	C03.752.625.235 Cryptosporidiosis
-	C03.752.625.367 Dourine
-	C03.752.625.735 Theileriasis
-	C03.752.625.817 Toxoplasmosis, Animal
-	C03.752.625.896 Trypanosomiasis, Bovine
-	C03.752.875 Tick-Borne Diseases
-	C03.752.875.175 Babesiosis
-	C03.752.875.850 Theileriasis
-	C03.752.890 Trichomonas Infections
-	C03.752.890.633 Trichomonas Vaginitis
-	C03.858 Skin Diseases, Parasitic
-	C03.858.211 Ectoparasitic Infestations
-	C03.858.211.250 Flea Infestations
-	C03.858.211.250.800 Tungiasis
-	C03.858.211.465 Lice Infestations
-	C03.858.211.480 Mite Infestations
-	C03.858.211.480.708 Scabies
-	C03.858.211.480.869 Trombiculiasis
-	C03.858.211.503 Myiasis

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C03.858.211.503.398 Hypodermyiasis
-	C03.858.211.503.744 Screw Worm Infection
-	C03.858.211.857 Tick Infestations
-	C03.858.424 Larva Migrans
-	C03.858.560 Leishmaniasis
-	C03.858.560.400 Leishmaniasis, Cutaneous
-	C03.858.560.400.350 Leishmaniasis, Diffuse Cutaneous
-	C03.858.560.400.385 Leishmaniasis, Mucocutaneous
-	C03.858.650 Onchocerciasis
-	C03.908 Zoonoses
-	C03.908.650 Pythiosis
-	C04 Neoplasms
-	C04.182 Cysts
-	C04.182.044 Arachnoid Cysts
-	C04.182.089 Bone Cysts
-	C04.182.089.265 Bone Cysts, Aneurysmal
-	C04.182.089.530 Jaw Cysts
-	C04.182.089.530.660 Nonodontogenic Cysts
-	C04.182.089.530.690 Odontogenic Cysts
-	C04.182.089.530.690.150 Basal Cell Nevus Syndrome
-	C04.182.089.530.690.310 Dentigerous Cyst
-	C04.182.089.530.690.605 Odontogenic Cyst, Calcifying
-	C04.182.089.530.690.790 Periodontal Cyst
-	C04.182.089.530.690.790.820 Radicular Cyst
-	C04.182.117 Branchioma
-	C04.182.156 Breast Cyst
-	C04.182.195 Bronchogenic Cyst
-	C04.182.197 Chalazion
-	C04.182.198 Choledochal Cyst
-	C04.182.199 Colloid Cysts
-	C04.182.201 Dermoid Cyst
-	C04.182.254 Epidermal Cyst
-	C04.182.281 Esophageal Cyst
-	C04.182.300 Follicular Cyst
-	C04.182.347 Ganglion Cysts
-	C04.182.430 Lymphocele

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C04.182.444                    Mediastinal Cyst
-	C04.182.473                    Mesenteric Cyst
-	C04.182.511                    Mucocele
-	C04.182.612                    Ovarian Cysts
-	C04.182.612.765                Polycystic Ovary Syndrome
-	C04.182.640                    Pancreatic Cyst
-	C04.182.640.692                Pancreatic Pseudocyst
-	C04.182.668                    Parovarian Cyst
-	C04.182.710                    Pilonidal Sinus
-	C04.182.766                    Ranula
-	C04.182.867                    Synovial Cyst
-	C04.182.867.500                Popliteal Cyst
-	C04.182.872                    Tarlov Cysts
-	C04.182.902                    Thyroglossal Cyst
-	C04.182.946                    Urachal Cyst
-	C04.445                        Hamartoma
-	C04.445.435                    Hamartoma Syndrome, Multiple
-	C04.445.435.500                Proteus Syndrome
-	C04.445.622                    Pallister-Hall Syndrome
-	C04.445.810                    Tuberous Sclerosis
-	C04.557                        Neoplasms by Histologic Type
-	C04.557.227                    Histiocytic Disorders, Malignant
-	C04.557.227.190                Dendritic Cell Sarcoma, Follicular
-	C04.557.227.199                Dendritic Cell Sarcoma, Interdigitating
-	C04.557.227.380                Histiocytic Sarcoma
-	C04.557.227.500                Langerhans Cell Sarcoma
-	C04.557.337                    Leukemia
-	C04.557.337.100                Enzootic Bovine Leukosis
-	C04.557.337.372                Leukemia, Experimental
-	C04.557.337.372.216            Avian Leukosis
-	C04.557.337.372.594            Leukemia L1210
-	C04.557.337.372.602            Leukemia L5178
-	C04.557.337.372.782            Leukemia P388
-	C04.557.337.385                Leukemia, Feline
-	C04.557.337.415                Leukemia, Hairy Cell
-	C04.557.337.428                Leukemia, Lymphoid



## MeSH Tree Changes for 2017

Type	Tree - heading
-	C04.557.337.428.080                      Leukemia, B-Cell
-	C04.557.337.428.080.125                      Leukemia, Lymphocytic, Chronic, B-Cell
-	C04.557.337.428.080.562                      Leukemia, Prolymphocytic, B-Cell
-	C04.557.337.428.100                      Leukemia, Biphenotypic, Acute
-	C04.557.337.428.565                      Leukemia, Prolymphocytic
-	C04.557.337.428.565.745                      Leukemia, Prolymphocytic, B-Cell
-	C04.557.337.428.565.750                      Leukemia, Prolymphocytic, T-Cell
-	C04.557.337.428.580                      Leukemia, T-Cell
-	C04.557.337.428.580.049                      Leukemia, Large Granular Lymphocytic
-	C04.557.337.428.580.100                      Leukemia-Lymphoma, Adult T-Cell
-	C04.557.337.428.580.125                      Leukemia, Prolymphocytic, T-Cell
-	C04.557.337.428.600                      Precursor Cell Lymphoblastic Leukemia-Lymphoma
-	C04.557.337.428.600.600                      Precursor B-Cell Lymphoblastic Leukemia-Lymphoma
-	C04.557.337.428.600.620                      Precursor T-Cell Lymphoblastic Leukemia-Lymphoma
-	C04.557.337.440                      Leukemia, Mast-Cell
-	C04.557.337.539                      Leukemia, Myeloid
-	C04.557.337.539.250                      Leukemia, Myelogenous, Chronic, BCR-ABL Positive
-	C04.557.337.539.250.100                      Blast Crisis
-	C04.557.337.539.250.300                      Leukemia, Myeloid, Accelerated Phase
-	C04.557.337.539.250.400                      Leukemia, Myeloid, Chronic-Phase
-	C04.557.337.539.275                      Leukemia, Myeloid, Acute
-	C04.557.337.539.275.125                      Leukemia, Basophilic, Acute
-	C04.557.337.539.275.300                      Leukemia, Eosinophilic, Acute
-	C04.557.337.539.275.325                      Leukemia, Erythroblastic, Acute
-	C04.557.337.539.275.440                      Leukemia, Mast-Cell
-	C04.557.337.539.275.450                      Leukemia, Megakaryoblastic, Acute
-	C04.557.337.539.275.484                      Leukemia, Monocytic, Acute
-	C04.557.337.539.275.700                      Leukemia, Promyelocytic, Acute
-	C04.557.337.539.300                      Leukemia, Myeloid, Chronic, Atypical, BCR-ABL Negative
-	C04.557.337.539.520                      Leukemia, Myelomonocytic, Acute
-	C04.557.337.539.522                      Leukemia, Myelomonocytic, Chronic
-	C04.557.337.539.525                      Leukemia, Myelomonocytic, Juvenile
-	C04.557.337.539.775                      Sarcoma, Myeloid
-	C04.557.337.595                      Leukemia, Plasma Cell

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C04.557.337.650                      Leukemia, Radiation-Induced
-	C04.557.375                              Lymphatic Vessel Tumors
-	C04.557.375.450                        Lymphangioma
-	C04.557.375.450.450                    Lymphangioma, Cystic
-	C04.557.375.460                        Lymphangiomyoma
-	C04.557.375.460.465                    Lymphangiomyomatosis
-	C04.557.375.480                        Lymphangiosarcoma
-	C04.557.386                              Lymphoma
-	C04.557.386.150                        Composite Lymphoma
-	C04.557.386.355                        Hodgkin Disease
-	C04.557.386.390                        Immunoproliferative Small Intestinal Disease
-	C04.557.386.435                        Intraocular Lymphoma
-	C04.557.386.480                        Lymphoma, Non-Hodgkin
-	C04.557.386.480.150                    Lymphoma, B-Cell
-	C04.557.386.480.150.165                Burkitt Lymphoma
-	C04.557.386.480.150.450                Lymphoma, AIDS-Related
-	C04.557.386.480.150.570                Lymphoma, B-Cell, Marginal Zone
-	C04.557.386.480.150.585                Lymphoma, Large B-Cell, Diffuse
-	C04.557.386.480.150.585.500            Plasmablastic Lymphoma
-	C04.557.386.480.150.592                Lymphoma, Primary Effusion
-	C04.557.386.480.150.600                Lymphomatoid Granulomatosis
-	C04.557.386.480.350                    Lymphoma, Follicular
-	C04.557.386.480.493                    Lymphoma, Large-Cell, Immunoblastic
-	C04.557.386.480.525                    Lymphoma, Mantle-Cell
-	C04.557.386.480.750                    Lymphoma, T-Cell
-	C04.557.386.480.750.099                Enteropathy-Associated T-Cell Lymphoma
-	C04.557.386.480.750.199                Lymphoma, Extranodal NK-T-Cell
-	C04.557.386.480.750.399                Lymphoma, Large-Cell, Anaplastic
-	C04.557.386.480.750.800                Lymphoma, T-Cell, Cutaneous
-	C04.557.386.480.750.800.507 Large Cell                              Lymphoma, Primary Cutaneous Anaplastic
-	C04.557.386.480.750.800.528            Lymphomatoid Papulosis
-	C04.557.386.480.750.800.550            Mycosis Fungoides
-	C04.557.386.480.750.800.550.600      Pagetoid Reticulosis
-	C04.557.386.480.750.800.775            Sezary Syndrome
-	C04.557.386.480.750.825                Lymphoma, T-Cell, Peripheral

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C04.557.435 Neoplasms, Complex and Mixed
-	C04.557.435.075 Adenolymphoma
-	C04.557.435.090 Adenoma, Pleomorphic
-	C04.557.435.108 Adenomyoepithelioma
-	C04.557.435.110 Adenomyoma
-	C04.557.435.135 Adenosarcoma
-	C04.557.435.250 Carcinoma, Adenosquamous
-	C04.557.435.290 Carcinosarcoma
-	C04.557.435.290.210 Carcinoma 256, Walker
-	C04.557.435.295 Composite Lymphoma
-	C04.557.435.380 Hepatoblastoma
-	C04.557.435.500 Mesenchymoma
-	C04.557.435.525 Mixed Tumor, Malignant
-	C04.557.435.530 Mixed Tumor, Mesodermal
-	C04.557.435.540 Mixed Tumor, Mullerian
-	C04.557.435.585 Myoepithelioma
-	C04.557.435.595 Wilms Tumor
-	C04.557.435.595.220 Denys-Drash Syndrome
-	C04.557.435.595.950 WAGR Syndrome
-	C04.557.435.600 Nephroma, Mesoblastic
-	C04.557.435.675 Pulmonary Blastoma
-	C04.557.435.710 Rhabdoid Tumor
-	C04.557.435.775 Sarcoma, Endometrial Stromal
-	C04.557.435.850 Thymoma
-	C04.557.450 Neoplasms, Connective and Soft Tissue
-	C04.557.450.550 Neoplasms, Adipose Tissue
-	C04.557.450.550.100 Angiolipoma
-	C04.557.450.550.125 Angiomyolipoma
-	C04.557.450.550.400 Lipoma
-	C04.557.450.550.400.500 Lipoblastoma
-	C04.557.450.550.420 Liposarcoma
-	C04.557.450.550.420.425 Liposarcoma, Myxoid
-	C04.557.450.550.710 Myelolipoma
-	C04.557.450.565 Neoplasms, Connective Tissue
-	C04.557.450.565.250 Chondroblastoma
-	C04.557.450.565.265 Chondroma

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C04.557.450.565.265.270 Chondromatosis
-	C04.557.450.565.280 Chondrosarcoma
-	C04.557.450.565.280.280 Chondrosarcoma, Mesenchymal
-	C04.557.450.565.325 Endometrial Stromal Tumors
-	C04.557.450.565.370 Gastrointestinal Stromal Tumors
-	C04.557.450.565.380 Giant Cell Tumors
-	C04.557.450.565.380.380 Giant Cell Tumor of Bone
New Heading	<b>C04.557.450.565.380.690 Giant Cell Tumor of Tendon Sheath</b>
New Tree	<b>C04.557.450.565.380.690.500 Synovitis, Pigmented Villonodular</b>
-	C04.557.450.565.465 Mastocytosis
-	C04.557.450.565.465.124 Mast-Cell Sarcoma
-	C04.557.450.565.465.249 Mastocytoma
-	C04.557.450.565.465.249.500 Mastocytoma, Skin
-	C04.557.450.565.465.500 Mastocytosis, Cutaneous
-	C04.557.450.565.465.500.500 Mastocytoma, Skin
-	C04.557.450.565.465.500.850 Urticaria Pigmentosa
-	C04.557.450.565.465.750 Mastocytosis, Systemic
-	C04.557.450.565.465.750.500 Leukemia, Mast-Cell
-	C04.557.450.565.540 Myofibroma
-	C04.557.450.565.550 Myxoma
-	C04.557.450.565.550.312 Carney Complex
-	C04.557.450.565.550.625 Neurothekeoma
-	C04.557.450.565.560 Myxosarcoma
-	C04.557.450.565.575 Neoplasms, Bone Tissue
-	C04.557.450.565.575.400 Fibroma, Ossifying
-	C04.557.450.565.575.420 Giant Cell Tumor of Bone
-	C04.557.450.565.575.600 Osteoblastoma
-	C04.557.450.565.575.610 Osteochondroma
-	C04.557.450.565.575.610.615 Osteochondromatosis
-	C04.557.450.565.575.610.615.325 Exostoses, Multiple Hereditary
-	C04.557.450.565.575.625 Osteoma
-	C04.557.450.565.575.625.625 Osteoma, Osteoid
-	C04.557.450.565.575.650 Osteosarcoma
-	C04.557.450.565.575.650.655 Osteosarcoma, Juxtacortical

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C04.557.450.565.575.650.800 Sarcoma, Ewing
-	C04.557.450.565.590 Neoplasms, Fibrous Tissue
-	C04.557.450.565.590.340 Fibroma
New Tree	<a href="#">C04.557.450.565.590.340.173</a> Dupuytren Contracture
-	C04.557.450.565.590.340.345 Fibroma, Desmoplastic
-	C04.557.450.565.590.340.360 Fibroma, Ossifying
-	C04.557.450.565.590.340.400 Fibromatosis, Abdominal
-	C04.557.450.565.590.340.410 Fibromatosis, Aggressive
New Heading	<b>C04.557.450.565.590.340.705</b> <b>Fibromatosis, Plantar</b>
-	C04.557.450.565.590.350 Fibrosarcoma
-	C04.557.450.565.590.350.320 Dermatofibrosarcoma
-	C04.557.450.565.590.350.590 Neurofibrosarcoma
-	C04.557.450.565.590.425 Histiocytoma
-	C04.557.450.565.590.425.350 Histiocytoma, Benign Fibrous
-	C04.557.450.565.590.425.360 Histiocytoma, Malignant Fibrous
-	C04.557.450.565.590.550 Myofibromatosis
-	C04.557.450.565.590.595 Neoplasms, Fibroepithelial
-	C04.557.450.565.590.595.050 Adenofibroma
-	C04.557.450.565.590.595.050.500 Cystadenofibroma
-	C04.557.450.565.590.595.150 Brenner Tumor
-	C04.557.450.565.590.595.350 Fibroadenoma
-	C04.557.450.565.590.797 Solitary Fibrous Tumors
-	C04.557.450.565.590.797.750 Solitary Fibrous Tumor, Pleural
-	C04.557.450.565.800 Sarcoma, Clear Cell
-	C04.557.450.565.825 Sarcoma, Small Cell
-	C04.557.450.565.835 Sarcoma, Synovial
-	C04.557.450.590 Neoplasms, Muscle Tissue
-	C04.557.450.590.350 Granular Cell Tumor
-	C04.557.450.590.450 Leiomyoma
-	C04.557.450.590.450.125 Angiomyoma
-	C04.557.450.590.450.455 Leiomyoma, Epithelioid
-	C04.557.450.590.450.465 Leiomyomatosis
-	C04.557.450.590.455 Leiomyosarcoma
-	C04.557.450.590.540 Myoma

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C04.557.450.590.540.700 Rhabdomyoma
-	C04.557.450.590.550 Myosarcoma
-	C04.557.450.590.550.660 Rhabdomyosarcoma
-	C04.557.450.590.550.660.665 Rhabdomyosarcoma, Alveolar
-	C04.557.450.590.550.660.675 Rhabdomyosarcoma, Embryonal
-	C04.557.450.590.775 Sarcoma, Alveolar Soft Part
-	C04.557.450.590.800 Smooth Muscle Tumor
-	C04.557.450.692 Perivascular Epithelioid Cell Neoplasms
-	C04.557.450.692.249 Angiomyolipoma
-	C04.557.450.692.500 Lymphangioliomyomatosis
-	C04.557.450.795 Sarcoma
-	C04.557.450.795.135 Adenosarcoma
-	C04.557.450.795.290 Carcinosarcoma
-	C04.557.450.795.290.210 Carcinoma 256, Walker
-	C04.557.450.795.300 Chondrosarcoma
-	C04.557.450.795.300.280 Chondrosarcoma, Mesenchymal
-	C04.557.450.795.315 Desmoplastic Small Round Cell Tumor
-	C04.557.450.795.332 Endometrial Stromal Tumors
-	C04.557.450.795.332.500 Sarcoma, Endometrial Stromal
-	C04.557.450.795.350 Fibrosarcoma
-	C04.557.450.795.350.320 Dermatofibrosarcoma
-	C04.557.450.795.350.590 Neurofibrosarcoma
-	C04.557.450.795.390 Hemangiosarcoma
-	C04.557.450.795.400 Histiocytoma, Malignant Fibrous
-	C04.557.450.795.455 Leiomyosarcoma
-	C04.557.450.795.465 Liposarcoma
-	C04.557.450.795.465.425 Liposarcoma, Myxoid
-	C04.557.450.795.480 Lymphangiosarcoma
-	C04.557.450.795.530 Mixed Tumor, Mesodermal
-	C04.557.450.795.550 Myosarcoma
-	C04.557.450.795.550.660 Rhabdomyosarcoma
-	C04.557.450.795.550.660.665 Rhabdomyosarcoma, Alveolar
-	C04.557.450.795.550.660.675 Rhabdomyosarcoma, Embryonal
-	C04.557.450.795.560 Myxosarcoma
-	C04.557.450.795.620 Osteosarcoma
-	C04.557.450.795.620.655 Osteosarcoma, Juxtacortical

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C04.557.450.795.620.800 Sarcoma, Ewing
-	C04.557.450.795.650 Phyllodes Tumor
-	C04.557.450.795.775 Sarcoma, Alveolar Soft Part
-	C04.557.450.795.800 Sarcoma, Clear Cell
-	C04.557.450.795.830 Sarcoma, Experimental
-	C04.557.450.795.830.760 Sarcoma 37
-	C04.557.450.795.830.780 Sarcoma 180
-	C04.557.450.795.830.800 Sarcoma, Avian
-	C04.557.450.795.830.850 Sarcoma, Yoshida
-	C04.557.450.795.850 Sarcoma, Kaposi
-	C04.557.450.795.853 Sarcoma, Myeloid
-	C04.557.450.795.870 Sarcoma, Small Cell
-	C04.557.450.795.875 Sarcoma, Synovial
-	C04.557.465 Neoplasms, Germ Cell and Embryonal
-	C04.557.465.200 Carcinoma, Embryonal
-	C04.557.465.220 Chordoma
-	C04.557.465.330 Germinoma
-	C04.557.465.330.300 Dysgerminoma
-	C04.557.465.330.800 Seminoma
-	C04.557.465.420 Gonadoblastoma
-	C04.557.465.510 Mesonephroma
-	C04.557.465.510.350 Endodermal Sinus Tumor
-	C04.557.465.625 Neuroectodermal Tumors
-	C04.557.465.625.200 Craniopharyngioma
-	C04.557.465.625.600 Neoplasms, Neuroepithelial
-	C04.557.465.625.600.355 Ganglioneuroma
-	C04.557.465.625.600.380 Glioma
-	C04.557.465.625.600.380.080 Astrocytoma
-	C04.557.465.625.600.380.080.335 Glioblastoma
-	C04.557.465.625.600.380.290 Ependymoma
-	C04.557.465.625.600.380.290.390 Glioma, Subependymal
-	C04.557.465.625.600.380.350 Ganglioglioma
-	C04.557.465.625.600.380.400 Gliosarcoma
-	C04.557.465.625.600.380.515 Medulloblastoma
-	C04.557.465.625.600.380.590 Oligodendroglioma
-	C04.557.465.625.600.380.795 Optic Nerve Glioma

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C04.557.465.625.600.580                      Neurocytoma
-	C04.557.465.625.600.590                      Neuroectodermal Tumors, Primitive
-	C04.557.465.625.600.590.500                      Medulloblastoma
-	C04.557.465.625.600.590.650                      Neuroectodermal Tumors, Primitive, Peripheral
-	C04.557.465.625.600.590.650.550                      Neuroblastoma
-	C04.557.465.625.600.590.650.550.150                      Esthesioneuroblastoma, Olfactory
-	C04.557.465.625.600.590.650.550.300                      Ganglioneuroblastoma
-	C04.557.465.625.600.657                      Pinealoma
-	C04.557.465.625.600.725                      Retinoblastoma
-	C04.557.465.625.630                      Neuroectodermal Tumor, Melanotic
-	C04.557.465.625.650                      Neuroendocrine Tumors
-	C04.557.465.625.650.025                      Adenoma, Acidophil
-	C04.557.465.625.650.075                      Adenoma, Basophil
-	C04.557.465.625.650.095                      Adenoma, Chromophobe
-	C04.557.465.625.650.135                      Apudoma
-	C04.557.465.625.650.200                      Carcinoid Tumor
-	C04.557.465.625.650.200.500                      Malignant Carcinoid Syndrome
-	C04.557.465.625.650.200.500.205                      Carcinoid Heart Disease
-	C04.557.465.625.650.240                      Carcinoma, Neuroendocrine
-	C04.557.465.625.650.240.315                      Carcinoma, Medullary
-	C04.557.465.625.650.240.325                      Carcinoma, Merkel Cell
-	C04.557.465.625.650.240.695                      Somatostatinoma
-	C04.557.465.625.650.240.847                      Vipoma
-	C04.557.465.625.650.510                      Melanoma
-	C04.557.465.625.650.510.385                      Hutchinson's Melanotic Freckle
-	C04.557.465.625.650.510.515                      Melanoma, Amelanotic
-	C04.557.465.625.650.510.525                      Melanoma, Experimental
-	C04.557.465.625.650.595                      Neurilemmoma
-	C04.557.465.625.650.595.610                      Neuroma, Acoustic
-	C04.557.465.625.650.595.610.500                      Neurofibromatosis 2
-	C04.557.465.625.650.700                      Paraganglioma
-	C04.557.465.625.650.700.705                      Paraganglioma, Extra-Adrenal
-	C04.557.465.625.650.700.705.220                      Carotid Body Tumor
-	C04.557.465.625.650.700.705.340                      Glomus Jugulare Tumor
-	C04.557.465.625.650.700.705.360                      Glomus Tympanicum Tumor
-	C04.557.465.625.650.700.725                      Pheochromocytoma



## MeSH Tree Changes for 2017

Type	Tree - heading
-	C04.557.465.900 Teratocarcinoma
-	C04.557.465.910 Teratoma
-	C04.557.465.910.250 Dermoid Cyst
-	C04.557.465.910.850 Struma Ovarii
-	C04.557.465.955 Trophoblastic Neoplasms
-	C04.557.465.955.207 Choriocarcinoma
-	C04.557.465.955.207.438 Choriocarcinoma, Non-gestational
-	C04.557.465.955.207.875 Trophoblastic Tumor, Placental Site
-	C04.557.465.955.416 Gestational Trophoblastic Disease
-	C04.557.465.955.416.812 Hydatidiform Mole
-	C04.557.465.955.416.812.500 Hydatidiform Mole, Invasive
-	C04.557.470 Neoplasms, Glandular and Epithelial
-	C04.557.470.035 Adenoma
-	C04.557.470.035.012 ACTH-Secreting Pituitary Adenoma
-	C04.557.470.035.025 Adenoma, Acidophil
-	C04.557.470.035.075 Adenoma, Basophil
-	C04.557.470.035.085 Adenoma, Bile Duct
-	C04.557.470.035.095 Adenoma, Chromophobe
-	C04.557.470.035.100 Adenoma, Islet Cell
-	C04.557.470.035.100.852 Insulinoma
-	C04.557.470.035.120 Adenoma, Liver Cell
-	C04.557.470.035.140 Adenoma, Oxyphilic
-	C04.557.470.035.155 Adenoma, Pleomorphic
-	C04.557.470.035.175 Adenoma, Sweat Gland
-	C04.557.470.035.175.125 Acrospiroma
-	C04.557.470.035.175.125.600 Poroma
-	C04.557.470.035.175.375 Hidrocystoma
-	C04.557.470.035.175.800 Syringoma
-	C04.557.470.035.185 Adenoma, Villous
-	C04.557.470.035.200 Adenomatoid Tumor
-	C04.557.470.035.210 Adenomatosis, Pulmonary
-	C04.557.470.035.215 Adenomatous Polyps
-	C04.557.470.035.215.100 Adenomatous Polyposis Coli
-	C04.557.470.035.215.100.500 Gardner Syndrome
-	C04.557.470.035.232 Adrenal Rest Tumor
-	C04.557.470.035.250 Apudoma

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C04.557.470.035.320 Cystadenoma
-	C04.557.470.035.320.225 Cystadenoma, Mucinous
-	C04.557.470.035.320.230 Cystadenoma, Papillary
-	C04.557.470.035.320.240 Cystadenoma, Serous
-	C04.557.470.035.415 Growth Hormone-Secreting Pituitary Adenoma
-	C04.557.470.035.510 Mesothelioma
-	C04.557.470.035.510.515 Mesothelioma, Cystic
-	C04.557.470.035.625 Prolactinoma
-	C04.557.470.200 Carcinoma
-	C04.557.470.200.025 Adenocarcinoma
-	C04.557.470.200.025.014 Adenocarcinoma in Situ
-	C04.557.470.200.025.030 Adenocarcinoma, Bronchiolo-Alveolar
-	C04.557.470.200.025.045 Adenocarcinoma, Clear Cell
-	C04.557.470.200.025.060 Adenocarcinoma, Follicular
-	C04.557.470.200.025.060.225 Carcinoma, Papillary, Follicular
-	C04.557.470.200.025.075 Adenocarcinoma, Mucinous
-	C04.557.470.200.025.085 Adenocarcinoma, Papillary
-	C04.557.470.200.025.085.225 Carcinoma, Papillary, Follicular
-	C04.557.470.200.025.095 Adenocarcinoma, Scirrhus
-	C04.557.470.200.025.095.410 Linitis Plastica
-	C04.557.470.200.025.105 Adenocarcinoma, Sebaceous
-	C04.557.470.200.025.152 Adrenocortical Carcinoma
-	C04.557.470.200.025.200 Carcinoid Tumor
-	C04.557.470.200.025.200.500 Malignant Carcinoid Syndrome
-	C04.557.470.200.025.200.500.205 Carcinoid Heart Disease
-	C04.557.470.200.025.215 Carcinoma, Acinar Cell
-	C04.557.470.200.025.220 Carcinoma, Adenoid Cystic
-	C04.557.470.200.025.232 Carcinoma, Ductal
-	C04.557.470.200.025.232.500 Carcinoma, Ductal, Breast
-	C04.557.470.200.025.232.750 Carcinoma, Pancreatic Ductal
-	C04.557.470.200.025.240 Carcinoma, Endometrioid
-	C04.557.470.200.025.255 Carcinoma, Hepatocellular
-	C04.557.470.200.025.275 Carcinoma, Intraductal, Noninfiltrating
Old Tree	C04.557.470.200.025.275.625 Paget's Disease, Mammary
-	C04.557.470.200.025.290 Carcinoma, Islet Cell
-	C04.557.470.200.025.290.500 Gastrinoma

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	C04.557.470.200.025.290.750	Glucagonoma
-	C04.557.470.200.025.305	Carcinoma, Lobular
-	C04.557.470.200.025.340	Carcinoma, Mucoepidermoid
-	C04.557.470.200.025.370	Carcinoma, Neuroendocrine
-	C04.557.470.200.025.370.315	Carcinoma, Medullary
-	C04.557.470.200.025.370.325	Carcinoma, Merkel Cell
-	C04.557.470.200.025.370.695	Somatostatinoma
-	C04.557.470.200.025.370.847	Vipoma
-	C04.557.470.200.025.390	Carcinoma, Renal Cell
-	C04.557.470.200.025.415	Carcinoma, Signet Ring Cell
-	C04.557.470.200.025.415.410	Krukenberg Tumor
-	C04.557.470.200.025.420	Carcinoma, Skin Appendage
-	C04.557.470.200.025.450	Cholangiocarcinoma
-	C04.557.470.200.025.450.500	Klatskin Tumor
-	C04.557.470.200.025.455	Choriocarcinoma
-	C04.557.470.200.025.455.750	Choriocarcinoma, Non-gestational
-	C04.557.470.200.025.455.875	Trophoblastic Tumor, Placental Site
-	C04.557.470.200.025.480	Cystadenocarcinoma
-	C04.557.470.200.025.480.225	Cystadenocarcinoma, Mucinous
-	C04.557.470.200.025.480.230	Cystadenocarcinoma, Papillary
-	C04.557.470.200.025.480.240	Cystadenocarcinoma, Serous
-	C04.557.470.200.025.500	Eccrine Porocarcinoma
-	C04.557.470.200.025.660	Paget Disease, Extramammary
-	C04.557.470.200.025.715	Pulmonary Adenomatosis, Ovine
-	C04.557.470.200.150	Carcinoma, Adenosquamous
-	C04.557.470.200.165	Carcinoma, Basal Cell
-	C04.557.470.200.165.150	Basal Cell Nevus Syndrome
-	C04.557.470.200.170	Carcinoma, Basosquamous
-	C04.557.470.200.200	Carcinoma, Ehrlich Tumor
-	C04.557.470.200.220	Carcinoma, Giant Cell
-	C04.557.470.200.240	Carcinoma in Situ
-	C04.557.470.200.240.124	Adenocarcinoma in Situ
New Heading	<b>C04.557.470.200.240.187</b>	<b>Breast Carcinoma In Situ</b>
New Tree	<a href="#">C04.557.470.200.240.187.250</a>	<a href="#">Carcinoma, Intraductal, Noninfiltrating</a>

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	C04.557.470.200.240.187.500 <span style="float: right;">Paget's Disease, Mammary</span>
-	C04.557.470.200.240.250 <span style="float: right;">Cervical Intraepithelial Neoplasia</span>
-	C04.557.470.200.240.500 <span style="float: right;">Prostatic Intraepithelial Neoplasia</span>
-	C04.557.470.200.255 <span style="float: right;">Carcinoma, Krebs 2</span>
-	C04.557.470.200.260 <span style="float: right;">Carcinoma, Large Cell</span>
-	C04.557.470.200.280 <span style="float: right;">Carcinoma, Lewis Lung</span>
-	C04.557.470.200.360 <span style="float: right;">Carcinoma, Papillary</span>
-	C04.557.470.200.380 <span style="float: right;">Carcinoma, Small Cell</span>
-	C04.557.470.200.400 <span style="float: right;">Carcinoma, Squamous Cell</span>
-	C04.557.470.200.400.130 <span style="float: right;">Bowen's Disease</span>
-	C04.557.470.200.430 <span style="float: right;">Carcinoma, Transitional Cell</span>
-	C04.557.470.200.450 <span style="float: right;">Carcinoma, Verrucous</span>
-	C04.557.470.200.450.500 <span style="float: right;">Buschke-Lowenstein Tumor</span>
-	C04.557.470.200.588 <span style="float: right;">Mammary Analogue Secretory Carcinoma</span>
-	C04.557.470.200.725 <span style="float: right;">Thyroid Carcinoma, Anaplastic</span>
-	C04.557.470.550 <span style="float: right;">Neoplasms, Adnexal and Skin Appendage</span>
-	C04.557.470.550.105 <span style="float: right;">Adenocarcinoma, Sebaceous</span>
-	C04.557.470.550.175 <span style="float: right;">Adenoma, Sweat Gland</span>
-	C04.557.470.550.175.125 <span style="float: right;">Acrospiroma</span>
-	C04.557.470.550.175.125.600 <span style="float: right;">Poroma</span>
-	C04.557.470.550.175.375 <span style="float: right;">Hidrocystoma</span>
-	C04.557.470.550.175.800 <span style="float: right;">Syringoma</span>
-	C04.557.470.550.420 <span style="float: right;">Carcinoma, Skin Appendage</span>
-	C04.557.470.565 <span style="float: right;">Neoplasms, Basal Cell</span>
-	C04.557.470.565.165 <span style="float: right;">Carcinoma, Basal Cell</span>
-	C04.557.470.565.165.150 <span style="float: right;">Basal Cell Nevus Syndrome</span>
-	C04.557.470.565.170 <span style="float: right;">Carcinoma, Basosquamous</span>
-	C04.557.470.565.625 <span style="float: right;">Pilomatrixoma</span>
-	C04.557.470.590 <span style="float: right;">Neoplasms, Cystic, Mucinous, and Serous</span>
-	C04.557.470.590.075 <span style="float: right;">Adenocarcinoma, Mucinous</span>
-	C04.557.470.590.340 <span style="float: right;">Carcinoma, Mucoepidermoid</span>
-	C04.557.470.590.415 <span style="float: right;">Carcinoma, Signet Ring Cell</span>
-	C04.557.470.590.415.410 <span style="float: right;">Krukenberg Tumor</span>
-	C04.557.470.590.480 <span style="float: right;">Cystadenocarcinoma</span>
-	C04.557.470.590.480.225 <span style="float: right;">Cystadenocarcinoma, Mucinous</span>

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C04.557.470.590.480.230 Cystadenocarcinoma, Papillary
-	C04.557.470.590.480.240 Cystadenocarcinoma, Serous
-	C04.557.470.590.482 Cystadenofibroma
-	C04.557.470.590.485 Cystadenoma
-	C04.557.470.590.485.225 Cystadenoma, Mucinous
-	C04.557.470.590.485.230 Cystadenoma, Papillary
-	C04.557.470.590.485.240 Cystadenoma, Serous
-	C04.557.470.590.580 Mucoepidermoid Tumor
-	C04.557.470.590.782 Pseudomyxoma Peritonei
-	C04.557.470.615 Neoplasms, Ductal, Lobular, and Medullary
-	C04.557.470.615.132 Carcinoma, Ductal
-	C04.557.470.615.132.500 Carcinoma, Ductal, Breast
-	C04.557.470.615.132.750 Carcinoma, Pancreatic Ductal
-	C04.557.470.615.275 Carcinoma, Intraductal, Noninfiltrating
-	C04.557.470.615.275.625 Paget's Disease, Mammary
-	C04.557.470.615.305 Carcinoma, Lobular
-	C04.557.470.615.315 Carcinoma, Medullary
-	C04.557.470.615.660 Paget Disease, Extramammary
-	C04.557.470.615.670 Papilloma, Intraductal
-	C04.557.470.625 Neoplasms, Fibroepithelial
-	C04.557.470.625.050 Adenofibroma
-	C04.557.470.625.050.500 Cystadenofibroma
-	C04.557.470.625.150 Brenner Tumor
-	C04.557.470.625.350 Fibroadenoma
-	C04.557.470.660 Neoplasms, Mesothelial
-	C04.557.470.660.200 Adenomatoid Tumor
-	C04.557.470.660.510 Mesothelioma
-	C04.557.470.660.510.515 Mesothelioma, Cystic
-	C04.557.470.670 Neoplasms, Neuroepithelial
-	C04.557.470.670.355 Ganglioneuroma
-	C04.557.470.670.380 Glioma
-	C04.557.470.670.380.080 Astrocytoma
-	C04.557.470.670.380.080.335 Glioblastoma
-	C04.557.470.670.380.290 Ependymoma
-	C04.557.470.670.380.290.390 Glioma, Subependymal
-	C04.557.470.670.380.350 Ganglioglioma

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C04.557.470.670.380.400                      Gliosarcoma
-	C04.557.470.670.380.515                      Medulloblastoma
-	C04.557.470.670.380.590                      Oligodendroglioma
-	C04.557.470.670.380.795                      Optic Nerve Glioma
-	C04.557.470.670.580                          Neurocytoma
-	C04.557.470.670.590                          Neuroectodermal Tumors, Primitive
-	C04.557.470.670.590.500                      Medulloblastoma
-	C04.557.470.670.590.650                      Neuroectodermal Tumors, Primitive, Peripheral
-	C04.557.470.670.590.650.550                      Neuroblastoma
-	C04.557.470.670.590.650.550.150                      Esthesioneuroblastoma, Olfactory
-	C04.557.470.670.590.650.550.300                      Ganglioneuroblastoma
-	C04.557.470.670.657                          Pinealoma
-	C04.557.470.670.725                          Retinoblastoma
-	C04.557.470.700                              Neoplasms, Squamous Cell
-	C04.557.470.700.040                          Acanthoma
-	C04.557.470.700.360                          Carcinoma, Papillary
-	C04.557.470.700.400                          Carcinoma, Squamous Cell
-	C04.557.470.700.400.130                      Bowen's Disease
-	C04.557.470.700.450                          Carcinoma, Verrucous
-	C04.557.470.700.450.500                      Buschke-Lowenstein Tumor
-	C04.557.470.700.600                          Papilloma
-	C04.557.470.700.600.610                      Papilloma, Inverted
-	C04.557.475                                  Neoplasms, Gonadal Tissue
-	C04.557.475.395                              Gonadoblastoma
-	C04.557.475.750                              Sex Cord-Gonadal Stromal Tumors
-	C04.557.475.750.656                          Granulosa Cell Tumor
-	C04.557.475.750.751                          Luteoma
-	C04.557.475.750.847                          Sertoli-Leydig Cell Tumor
-	C04.557.475.750.847.249                      Leydig Cell Tumor
-	C04.557.475.750.847.500                      Sertoli Cell Tumor
-	C04.557.475.750.875                          Thecoma
-	C04.557.580                                  Neoplasms, Nerve Tissue
-	C04.557.580.520                              Meningioma
-	C04.557.580.600                              Nerve Sheath Neoplasms
-	C04.557.580.600.580                          Neurofibroma
-	C04.557.580.600.580.585                      Neurofibroma, Plexiform

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C04.557.580.600.580.590 Neurofibromatoses
-	C04.557.580.600.580.590.650 Neurofibromatosis 1
-	C04.557.580.600.580.590.655 Neurofibromatosis 2
-	C04.557.580.600.580.795 Neurofibrosarcoma
-	C04.557.580.600.610 Neuroma
-	C04.557.580.600.610.595 Neurilemmoma
-	C04.557.580.600.610.595.610 Neuroma, Acoustic
-	C04.557.580.600.610.595.610.500 Neurofibromatosis 2
-	C04.557.580.600.625 Neurothekeoma
-	C04.557.580.625 Neuroectodermal Tumors
-	C04.557.580.625.200 Craniopharyngioma
-	C04.557.580.625.600 Neoplasms, Neuroepithelial
-	C04.557.580.625.600.355 Ganglioneuroma
-	C04.557.580.625.600.380 Glioma
-	C04.557.580.625.600.380.080 Astrocytoma
-	C04.557.580.625.600.380.080.335 Glioblastoma
-	C04.557.580.625.600.380.290 Ependymoma
-	C04.557.580.625.600.380.290.390 Glioma, Subependymal
-	C04.557.580.625.600.380.350 Ganglioglioma
-	C04.557.580.625.600.380.400 Gliosarcoma
-	C04.557.580.625.600.380.515 Medulloblastoma
-	C04.557.580.625.600.380.590 Oligodendroglioma
-	C04.557.580.625.600.380.795 Optic Nerve Glioma
-	C04.557.580.625.600.580 Neurocytoma
-	C04.557.580.625.600.590 Neuroectodermal Tumors, Primitive
-	C04.557.580.625.600.590.500 Medulloblastoma
-	C04.557.580.625.600.590.650 Neuroectodermal Tumors, Primitive, Peripheral
-	C04.557.580.625.600.590.650.550 Neuroblastoma
-	C04.557.580.625.600.590.650.550.150 Esthesioneuroblastoma, Olfactory
-	C04.557.580.625.600.590.650.550.300 Ganglioneuroblastoma
-	C04.557.580.625.600.657 Pinealoma
-	C04.557.580.625.600.725 Retinoblastoma
-	C04.557.580.625.630 Neuroectodermal Tumor, Melanotic
-	C04.557.580.625.650 Neuroendocrine Tumors
-	C04.557.580.625.650.025 Adenoma, Acidophil
-	C04.557.580.625.650.075 Adenoma, Basophil

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C04.557.580.625.650.095      Adenoma, Chromophobe
-	C04.557.580.625.650.135      Apudoma
-	C04.557.580.625.650.200      Carcinoid Tumor
-	C04.557.580.625.650.200.500      Malignant Carcinoid Syndrome
-	C04.557.580.625.650.200.500.205      Carcinoid Heart Disease
-	C04.557.580.625.650.240      Carcinoma, Neuroendocrine
-	C04.557.580.625.650.240.315      Carcinoma, Medullary
-	C04.557.580.625.650.240.325      Carcinoma, Merkel Cell
-	C04.557.580.625.650.510      Melanoma
-	C04.557.580.625.650.510.385      Hutchinson's Melanotic Freckle
-	C04.557.580.625.650.510.515      Melanoma, Amelanotic
-	C04.557.580.625.650.510.525      Melanoma, Experimental
-	C04.557.580.625.650.595      Neurilemmoma
-	C04.557.580.625.650.595.610      Neuroma, Acoustic
-	C04.557.580.625.650.700      Paraganglioma
-	C04.557.580.625.650.700.705      Paraganglioma, Extra-Adrenal
-	C04.557.580.625.650.700.705.220      Carotid Body Tumor
-	C04.557.580.625.650.700.705.340      Glomus Jugulare Tumor
-	C04.557.580.625.650.700.705.360      Glomus Tympanicum Tumor
-	C04.557.580.625.650.700.725      Pheochromocytoma
-	C04.557.595      Neoplasms, Plasma Cell
-	C04.557.595.500      Multiple Myeloma
-	C04.557.595.500.500      Leukemia, Plasma Cell
-	C04.557.595.600      Plasmacytoma
-	C04.557.595.925      Waldenstrom Macroglobulinemia
-	C04.557.645      Neoplasms, Vascular Tissue
-	C04.557.645.100      Angiofibroma
-	C04.557.645.115      Angiokeratoma
-	C04.557.645.350      Glomus Tumor
-	C04.557.645.375      Hemangioma
-	C04.557.645.375.185      Central Nervous System Venous Angioma
-	C04.557.645.375.370      Hemangioendothelioma
-	C04.557.645.375.370.380      Hemangioendothelioma, Epithelioid
-	C04.557.645.375.380      Hemangioma, Capillary
-	C04.557.645.375.380.370      Hemangioblastoma
-	C04.557.645.375.385      Hemangioma, Cavernous



## MeSH Tree Changes for 2017

Type	Tree - heading
-	C04.557.645.375.385.500 Hemangioma, Cavernous, Central Nervous System
-	C04.557.645.375.617 Kasabach-Merritt Syndrome
-	C04.557.645.375.850 Sturge-Weber Syndrome
-	C04.557.645.380 Hemangiopericytoma
-	C04.557.645.390 Hemangiosarcoma
-	C04.557.645.520 Meningioma
-	C04.557.645.750 Sarcoma, Kaposi
-	C04.557.665 Nevi and Melanomas
-	C04.557.665.510 Melanoma
-	C04.557.665.510.385 Hutchinson's Melanotic Freckle
-	C04.557.665.510.515 Melanoma, Amelanotic
-	C04.557.665.510.525 Melanoma, Experimental
-	C04.557.665.560 Nevus
-	C04.557.665.560.260 Dysplastic Nevus Syndrome
-	C04.557.665.560.580 Nevus, Halo
-	C04.557.665.560.590 Nevus, Intradermal
-	C04.557.665.560.615 Nevus, Pigmented
-	C04.557.665.560.615.530 Mongolian Spot
-	C04.557.665.560.615.550 Nevus, Blue
-	C04.557.665.560.615.585 Nevus of Ota
-	C04.557.665.560.615.625 Nevus, Spindle Cell
-	C04.557.665.560.615.625.585 Nevus, Epithelioid and Spindle Cell
-	C04.557.665.560.700 Nevus, Sebaceous of Jadassohn
-	C04.557.695 Odontogenic Tumors
-	C04.557.695.065 Ameloblastoma
-	C04.557.695.210 Cementoma
-	C04.557.695.605 Odontogenic Cyst, Calcifying
-	C04.557.695.607 Odontogenic Tumor, Squamous
-	C04.557.695.610 Odontoma
-	C04.588 Neoplasms by Site
-	C04.588.033 Abdominal Neoplasms
-	C04.588.033.513 Peritoneal Neoplasms
-	C04.588.033.731 Retroperitoneal Neoplasms
-	C04.588.033.740 Sister Mary Joseph's Nodule
-	C04.588.083 Anal Gland Neoplasms
-	C04.588.149 Bone Neoplasms

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	C04.588.149.030	Adamantinoma
-	C04.588.149.276	Femoral Neoplasms
-	C04.588.149.721	Skull Neoplasms
-	C04.588.149.721.450	Jaw Neoplasms
-	C04.588.149.721.450.583	Mandibular Neoplasms
-	C04.588.149.721.450.601	Maxillary Neoplasms
-	C04.588.149.721.450.692	Palatal Neoplasms
-	C04.588.149.721.600	Nose Neoplasms
-	C04.588.149.721.656	Orbital Neoplasms
-	C04.588.149.721.828	Skull Base Neoplasms
-	C04.588.149.828	Spinal Neoplasms
-	C04.588.180	Breast Neoplasms
New Heading	<b>C04.588.180.130</b>	<b>Breast Carcinoma In Situ</b>
-	C04.588.180.260	Breast Neoplasms, Male
-	C04.588.180.390	Carcinoma, Ductal, Breast
New Tree	<b>C04.588.180.437</b>	<b>Carcinoma, Lobular</b>
-	C04.588.180.483	Hereditary Breast and Ovarian Cancer Syndrome
-	C04.588.180.576	Inflammatory Breast Neoplasms
-	C04.588.180.682	Unilateral Breast Neoplasms
-	C04.588.180.788	Triple Negative Breast Neoplasms
-	C04.588.274	Digestive System Neoplasms
-	C04.588.274.120	Biliary Tract Neoplasms
-	C04.588.274.120.250	Bile Duct Neoplasms
-	C04.588.274.120.250.250	Common Bile Duct Neoplasms
-	C04.588.274.120.401	Gallbladder Neoplasms
-	C04.588.274.476	Gastrointestinal Neoplasms
-	C04.588.274.476.205	Esophageal Neoplasms
-	C04.588.274.476.411	Intestinal Neoplasms
-	C04.588.274.476.411.184	Cecal Neoplasms
-	C04.588.274.476.411.184.290	Appendiceal Neoplasms
-	C04.588.274.476.411.307	Colorectal Neoplasms
-	C04.588.274.476.411.307.089	Adenomatous Polyposis Coli
-	C04.588.274.476.411.307.089.393	Gardner Syndrome
-	C04.588.274.476.411.307.180	Colonic Neoplasms

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C04.588.274.476.411.307.180.800 Sigmoid Neoplasms
-	C04.588.274.476.411.307.190 Colorectal Neoplasms, Hereditary Nonpolyposis
-	C04.588.274.476.411.307.790 Rectal Neoplasms
-	C04.588.274.476.411.307.790.040 Anus Neoplasms
-	C04.588.274.476.411.307.790.040.040 Anal Gland Neoplasms
-	C04.588.274.476.411.445 Duodenal Neoplasms
-	C04.588.274.476.411.501 Ileal Neoplasms
-	C04.588.274.476.411.523 Jejunal Neoplasms
-	C04.588.274.476.767 Stomach Neoplasms
-	C04.588.274.623 Liver Neoplasms
-	C04.588.274.623.040 Adenoma, Liver Cell
-	C04.588.274.623.160 Carcinoma, Hepatocellular
-	C04.588.274.623.460 Liver Neoplasms, Experimental
-	C04.588.274.761 Pancreatic Neoplasms
-	C04.588.274.761.249 Adenoma, Islet Cell
-	C04.588.274.761.249.500 Insulinoma
-	C04.588.274.761.500 Carcinoma, Islet Cell
-	C04.588.274.761.500.124 Gastrinoma
-	C04.588.274.761.500.249 Glucagonoma
-	C04.588.274.761.500.500 Somatostatinoma
-	C04.588.274.761.500.750 Vipoma
-	C04.588.274.761.750 Carcinoma, Pancreatic Ductal
-	C04.588.274.780 Peritoneal Neoplasms
-	C04.588.322 Endocrine Gland Neoplasms
-	C04.588.322.078 Adrenal Gland Neoplasms
-	C04.588.322.078.265 Adrenal Cortex Neoplasms
-	C04.588.322.078.265.500 Adrenocortical Adenoma
-	C04.588.322.078.265.750 Adrenocortical Carcinoma
-	C04.588.322.400 Multiple Endocrine Neoplasia
-	C04.588.322.400.500 Multiple Endocrine Neoplasia Type 1
-	C04.588.322.400.505 Multiple Endocrine Neoplasia Type 2a
-	C04.588.322.400.510 Multiple Endocrine Neoplasia Type 2b
-	C04.588.322.455 Ovarian Neoplasms
-	C04.588.322.455.398 Granulosa Cell Tumor
-	C04.588.322.455.431 Hereditary Breast and Ovarian Cancer Syndrome
-	C04.588.322.455.464 Luteoma

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C04.588.322.455.531 Meigs Syndrome
-	C04.588.322.455.648 Sertoli-Leydig Cell Tumor
-	C04.588.322.455.765 Thecoma
-	C04.588.322.475 Pancreatic Neoplasms
-	C04.588.322.475.249 Adenoma, Islet Cell
-	C04.588.322.475.249.500 Insulinoma
-	C04.588.322.475.500 Carcinoma, Islet Cell
-	C04.588.322.475.500.124 Gastrinoma
-	C04.588.322.475.500.249 Glucagonoma
-	C04.588.322.475.500.500 Somatostatinoma
-	C04.588.322.475.500.750 Vipoma
-	C04.588.322.475.750 Carcinoma, Pancreatic Ductal
-	C04.588.322.490 Paraneoplastic Endocrine Syndromes
-	C04.588.322.525 Parathyroid Neoplasms
-	C04.588.322.609 Pituitary Neoplasms
-	C04.588.322.609.145 ACTH-Secreting Pituitary Adenoma
-	C04.588.322.609.145.500 Nelson Syndrome
-	C04.588.322.609.292 Growth Hormone-Secreting Pituitary Adenoma
-	C04.588.322.609.792 Prolactinoma
-	C04.588.322.762 Testicular Neoplasms
-	C04.588.322.762.500 Sertoli-Leydig Cell Tumor
-	C04.588.322.762.500.249 Leydig Cell Tumor
-	C04.588.322.762.500.500 Sertoli Cell Tumor
-	C04.588.322.894 Thyroid Neoplasms
-	C04.588.322.894.800 Thyroid Nodule
-	C04.588.364 Eye Neoplasms
-	C04.588.364.235 Conjunctival Neoplasms
-	C04.588.364.447 Intraocular Lymphoma
-	C04.588.364.659 Orbital Neoplasms
-	C04.588.364.738 Paraneoplastic Syndromes, Ocular
-	C04.588.364.818 Retinal Neoplasms
-	C04.588.364.818.760 Retinoblastoma
-	C04.588.364.978 Uveal Neoplasms
-	C04.588.364.978.223 Choroid Neoplasms
-	C04.588.364.978.400 Iris Neoplasms
-	C04.588.443 Head and Neck Neoplasms

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C04.588.443.353 Esophageal Neoplasms
-	C04.588.443.392 Facial Neoplasms
-	C04.588.443.392.500 Eyelid Neoplasms
-	C04.588.443.591 Mouth Neoplasms
-	C04.588.443.591.402 Gingival Neoplasms
-	C04.588.443.591.545 Leukoplakia, Oral
-	C04.588.443.591.545.500 Leukoplakia, Hairy
-	C04.588.443.591.550 Lip Neoplasms
-	C04.588.443.591.692 Palatal Neoplasms
-	C04.588.443.591.824 Salivary Gland Neoplasms
-	C04.588.443.591.824.695 Parotid Neoplasms
-	C04.588.443.591.824.882 Sublingual Gland Neoplasms
-	C04.588.443.591.824.885 Submandibular Gland Neoplasms
-	C04.588.443.591.925 Tongue Neoplasms
-	C04.588.443.665 Otorhinolaryngologic Neoplasms
-	C04.588.443.665.312 Ear Neoplasms
-	C04.588.443.665.481 Laryngeal Neoplasms
-	C04.588.443.665.650 Nose Neoplasms
-	C04.588.443.665.650.693 Paranasal Sinus Neoplasms
-	C04.588.443.665.650.693.575 Maxillary Sinus Neoplasms
-	C04.588.443.665.710 Pharyngeal Neoplasms
-	C04.588.443.665.710.485 Hypopharyngeal Neoplasms
-	C04.588.443.665.710.650 Nasopharyngeal Neoplasms
-	C04.588.443.665.710.684 Oropharyngeal Neoplasms
-	C04.588.443.665.710.684.800 Tonsillar Neoplasms
-	C04.588.443.680 Parathyroid Neoplasms
-	C04.588.443.915 Thyroid Neoplasms
-	C04.588.443.915.800 Thyroid Nodule
-	C04.588.443.925 Tracheal Neoplasms
-	C04.588.448 Hematologic Neoplasms
-	C04.588.448.200 Bone Marrow Neoplasms
New Tree	<a href="#">C04.588.448.200.500</a> <a href="#">Polycythemia Vera</a>
-	C04.588.531 Mammary Neoplasms, Animal
-	C04.588.531.500 Mammary Neoplasms, Experimental
-	C04.588.614 Nervous System Neoplasms

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C04.588.614.250 Central Nervous System Neoplasms
-	C04.588.614.250.195 Brain Neoplasms
-	C04.588.614.250.195.205 Cerebral Ventricle Neoplasms
-	C04.588.614.250.195.205.200 Choroid Plexus Neoplasms
-	C04.588.614.250.195.205.200.500 Papilloma, Choroid Plexus
-	C04.588.614.250.195.411 Infratentorial Neoplasms
-	C04.588.614.250.195.411.100 Brain Stem Neoplasms
-	C04.588.614.250.195.411.211 Cerebellar Neoplasms
-	C04.588.614.250.195.648 Neurocytoma
-	C04.588.614.250.195.766 Pinealoma
-	C04.588.614.250.195.885 Supratentorial Neoplasms
-	C04.588.614.250.195.885.500 Hypothalamic Neoplasms
-	C04.588.614.250.195.885.500.299 Pallister-Hall Syndrome
-	C04.588.614.250.195.885.500.600 Pituitary Neoplasms
-	C04.588.614.250.387 Central Nervous System Cysts
-	C04.588.614.250.387.100 Arachnoid Cysts
-	C04.588.614.250.387.200 Colloid Cysts
-	C04.588.614.250.580 Meningeal Neoplasms
-	C04.588.614.250.580.150 Meningeal Carcinomatosis
-	C04.588.614.250.580.500 Meningioma
-	C04.588.614.250.803 Spinal Cord Neoplasms
-	C04.588.614.250.803.342 Epidural Neoplasms
-	C04.588.614.300 Cranial Nerve Neoplasms
-	C04.588.614.300.015 Neuroma, Acoustic
-	C04.588.614.300.600 Optic Nerve Neoplasms
-	C04.588.614.300.600.600 Optic Nerve Glioma
-	C04.588.614.550 Paraneoplastic Syndromes, Nervous System
-	C04.588.614.550.112 Anti-N-Methyl-D-Aspartate Receptor Encephalitis
-	C04.588.614.550.225 Lambert-Eaton Myasthenic Syndrome
-	C04.588.614.550.450 Limbic Encephalitis
-	C04.588.614.550.550 Myelitis, Transverse
-	C04.588.614.550.600 Opsoclonus-Myoclonus Syndrome
-	C04.588.614.550.650 Paraneoplastic Cerebellar Degeneration
-	C04.588.614.550.700 Paraneoplastic Polyneuropathy
-	C04.588.614.596 Peripheral Nervous System Neoplasms
-	C04.588.614.596.240 Cranial Nerve Neoplasms

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C04.588.614.596.240.015 Neuroma, Acoustic
-	C04.588.614.596.240.240 Optic Nerve Neoplasms
-	C04.588.614.596.240.240.500 Optic Nerve Glioma
-	C04.588.699 Pelvic Neoplasms
-	C04.588.805 Skin Neoplasms
-	C04.588.805.040 Acanthoma
-	C04.588.805.578 Sebaceous Gland Neoplasms
-	C04.588.805.578.500 Muir-Torre Syndrome
-	C04.588.805.776 Sweat Gland Neoplasms
-	C04.588.839 Soft Tissue Neoplasms
-	C04.588.839.500 Muscle Neoplasms
-	C04.588.839.750 Vascular Neoplasms
-	C04.588.842 Splenic Neoplasms
-	C04.588.894 Thoracic Neoplasms
-	C04.588.894.309 Heart Neoplasms
-	C04.588.894.309.500 Carney Complex
-	C04.588.894.479 Mediastinal Neoplasms
-	C04.588.894.797 Respiratory Tract Neoplasms
-	C04.588.894.797.520 Lung Neoplasms
-	C04.588.894.797.520.109 Bronchial Neoplasms
-	C04.588.894.797.520.109.220 Carcinoma, Bronchogenic
-	C04.588.894.797.520.109.220.249 Carcinoma, Non-Small-Cell Lung
-	C04.588.894.797.520.109.220.624 Small Cell Lung Carcinoma
-	C04.588.894.797.520.237 Multiple Pulmonary Nodules
-	C04.588.894.797.520.734 Pancoast Syndrome
-	C04.588.894.797.520.867 Pulmonary Blastoma
-	C04.588.894.797.520.933 Pulmonary Sclerosing Hemangioma
-	C04.588.894.797.640 Pleural Neoplasms
-	C04.588.894.797.640.700 Pleural Effusion, Malignant
-	C04.588.894.797.640.800 Solitary Fibrous Tumor, Pleural
-	C04.588.894.797.760 Tracheal Neoplasms
-	C04.588.894.949 Thymus Neoplasms
-	C04.588.894.949.500 Thymoma
-	C04.588.945 Urogenital Neoplasms
-	C04.588.945.418 Genital Neoplasms, Female
-	C04.588.945.418.365 Fallopian Tube Neoplasms

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C04.588.945.418.948 Uterine Neoplasms
-	C04.588.945.418.948.585 Endometrial Neoplasms
-	C04.588.945.418.948.585.124 Carcinoma, Endometrioid
-	C04.588.945.418.948.850 Uterine Cervical Neoplasms
-	C04.588.945.418.955 Vaginal Neoplasms
-	C04.588.945.418.968 Vulvar Neoplasms
-	C04.588.945.440 Genital Neoplasms, Male
-	C04.588.945.440.715 Penile Neoplasms
-	C04.588.945.440.770 Prostatic Neoplasms
-	C04.588.945.440.770.500 Prostatic Neoplasms, Castration-Resistant
-	C04.588.945.440.915 Testicular Neoplasms
-	C04.588.945.440.915.500 Sertoli-Leydig Cell Tumor
-	C04.588.945.440.915.500.249 Leydig Cell Tumor
-	C04.588.945.440.915.500.500 Sertoli Cell Tumor
-	C04.588.945.947 Urologic Neoplasms
-	C04.588.945.947.535 Kidney Neoplasms
-	C04.588.945.947.535.160 Carcinoma, Renal Cell
-	C04.588.945.947.535.585 Wilms Tumor
-	C04.588.945.947.535.585.220 Denys-Drash Syndrome
-	C04.588.945.947.535.585.950 WAGR Syndrome
-	C04.588.945.947.535.790 Nephroma, Mesoblastic
-	C04.588.945.947.940 Ureteral Neoplasms
-	C04.588.945.947.945 Urethral Neoplasms
-	C04.588.945.947.960 Urinary Bladder Neoplasms
-	C04.588.945.956 Venereal Tumors, Veterinary
-	C04.619 Neoplasms, Experimental
-	C04.619.045 Carcinoma 256, Walker
-	C04.619.124 Carcinoma, Brown-Pearce
-	C04.619.169 Carcinoma, Ehrlich Tumor
-	C04.619.214 Carcinoma, Krebs 2
-	C04.619.230 Carcinoma, Lewis Lung
-	C04.619.531 Leukemia, Experimental
-	C04.619.531.216 Avian Leukosis
-	C04.619.531.594 Leukemia L1210
-	C04.619.531.602 Leukemia L5178
-	C04.619.531.782 Leukemia P388



## MeSH Tree Changes for 2017

Type	Tree - heading
-	C04.619.540      Liver Neoplasms, Experimental
-	C04.619.590      Mammary Neoplasms, Experimental
-	C04.619.600      Melanoma, Experimental
-	C04.619.857      Sarcoma, Experimental
-	C04.619.857.573      Sarcoma 37
-	C04.619.857.656      Sarcoma 180
-	C04.619.857.800      Sarcoma, Avian
-	C04.619.857.822      Sarcoma, Yoshida
-	C04.626      Neoplasms, Hormone-Dependent
-	C04.651      Neoplasms, Multiple Primary
-	C04.651.435      Hamartoma Syndrome, Multiple
-	C04.651.435.500      Proteus Syndrome
-	C04.651.600      Multiple Endocrine Neoplasia
-	C04.651.600.500      Multiple Endocrine Neoplasia Type 1
-	C04.651.600.505      Multiple Endocrine Neoplasia Type 2a
-	C04.651.600.510      Multiple Endocrine Neoplasia Type 2b
-	C04.651.800      Tuberous Sclerosis
-	C04.666      Neoplasms, Post-Traumatic
-	C04.682      Neoplasms, Radiation-Induced
-	C04.682.512      Leukemia, Radiation-Induced
-	C04.692      Neoplasms, Second Primary
-	C04.697      Neoplastic Processes
-	C04.697.045      Anaplasia
-	C04.697.098      Carcinogenesis
-	C04.697.098.500      Cell Transformation, Neoplastic
-	C04.697.098.500.110      Blast Crisis
-	C04.697.098.500.160      Cell Transformation, Viral
-	C04.697.098.875      Cocarcinogenesis
-	C04.697.645      Neoplasm Invasiveness
-	C04.697.645.500      Leukemic Infiltration
-	C04.697.650      Neoplasm Metastasis
-	C04.697.650.560      Lymphatic Metastasis
-	C04.697.650.695      Neoplasm Micrometastasis
-	C04.697.650.830      Neoplasm Seeding
-	C04.697.650.895      Neoplasms, Unknown Primary
-	C04.697.650.900      Neoplastic Cells, Circulating

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C04.697.655 Neoplasm Recurrence, Local
-	C04.697.670 Neoplasm Regression, Spontaneous
-	C04.697.700 Neoplasm, Residual
-	C04.700 Neoplastic Syndromes, Hereditary
-	C04.700.100 Adenomatous Polyposis Coli
-	C04.700.100.392 Gardner Syndrome
-	C04.700.175 Basal Cell Nevus Syndrome
-	C04.700.212 Birt-Hogg-Dube Syndrome
-	C04.700.250 Colorectal Neoplasms, Hereditary Nonpolyposis
-	C04.700.250.500 Lynch Syndrome II
-	C04.700.250.500.500 Muir-Torre Syndrome
-	C04.700.305 Dysplastic Nevus Syndrome
-	C04.700.330 Exostoses, Multiple Hereditary
-	C04.700.435 Hamartoma Syndrome, Multiple
-	C04.700.517 Hereditary Breast and Ovarian Cancer Syndrome
-	C04.700.600 Li-Fraumeni Syndrome
-	C04.700.630 Multiple Endocrine Neoplasia
-	C04.700.630.500 Multiple Endocrine Neoplasia Type 1
-	C04.700.630.505 Multiple Endocrine Neoplasia Type 2a
-	C04.700.630.510 Multiple Endocrine Neoplasia Type 2b
-	C04.700.632 Tuberous Sclerosis
-	C04.700.635 Wilms Tumor
-	C04.700.635.220 Denys-Drash Syndrome
-	C04.700.635.950 WAGR Syndrome
-	C04.700.645 Neurofibromatoses
-	C04.700.645.650 Neurofibromatosis 1
-	C04.700.645.655 Neurofibromatosis 2
-	C04.700.705 Peutz-Jeghers Syndrome
-	C04.730 Paraneoplastic Syndromes
-	C04.730.713 Paraneoplastic Endocrine Syndromes
-	C04.730.713.317 ACTH Syndrome, Ectopic
-	C04.730.713.988 Zollinger-Ellison Syndrome
-	C04.730.856 Paraneoplastic Syndromes, Nervous System
-	C04.730.856.112 Anti-N-Methyl-D-Aspartate Receptor Encephalitis
-	C04.730.856.225 Lambert-Eaton Myasthenic Syndrome
-	C04.730.856.437 Limbic Encephalitis

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C04.730.856.543 Myelitis, Transverse
-	C04.730.856.596 Opsoclonus-Myoclonus Syndrome
-	C04.730.856.650 Paraneoplastic Cerebellar Degeneration
-	C04.730.856.700 Paraneoplastic Polyneuropathy
-	C04.730.900 Paraneoplastic Syndromes, Ocular
-	C04.834 Precancerous Conditions
-	C04.834.020 Aberrant Crypt Foci
-	C04.834.288 Erythroplasia
-	C04.834.450 Keratosis, Actinic
-	C04.834.512 Leukoplakia
-	C04.834.512.513 Leukoplakia, Oral
-	C04.834.512.513.500 Leukoplakia, Hairy
-	C04.834.567 Lymphomatoid Granulomatosis
-	C04.834.770 Preleukemia
-	C04.834.818 Uterine Cervical Dysplasia
-	C04.834.818.249 Atypical Squamous Cells of the Cervix
-	C04.834.818.500 Squamous Intraepithelial Lesions of the Cervix
-	C04.834.867 Xeroderma Pigmentosum
-	C04.850 Pregnancy Complications, Neoplastic
-	C04.850.908 Trophoblastic Neoplasms
-	C04.850.908.208 Choriocarcinoma
-	C04.850.908.208.438 Choriocarcinoma, Non-gestational
-	C04.850.908.208.875 Trophoblastic Tumor, Placental Site
-	C04.850.908.416 Gestational Trophoblastic Disease
-	C04.850.908.416.750 Hydatidiform Mole
-	C04.850.908.416.750.500 Hydatidiform Mole, Invasive
-	C05 Musculoskeletal Diseases
-	C05.116 Bone Diseases
-	C05.116.070 Bone Cysts
-	C05.116.070.265 Bone Cysts, Aneurysmal
-	C05.116.099 Bone Diseases, Developmental
-	C05.116.099.052 Acro-Osteolysis
-	C05.116.099.052.400 Hajdu-Cheney Syndrome
-	C05.116.099.105 Basal Cell Nevus Syndrome
-	C05.116.099.343 Dwarfism
-	C05.116.099.343.110 Achondroplasia

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C05.116.099.343.110.500                      Thanatophoric Dysplasia
-	C05.116.099.343.250                      Cockayne Syndrome
-	C05.116.099.343.347                      Congenital Hypothyroidism
-	C05.116.099.343.445                      Dwarfism, Pituitary
-	C05.116.099.343.679                      Laron Syndrome
-	C05.116.099.343.796                      Mulibrey Nanism
-	C05.116.099.343.957                      Weill-Marchesani Syndrome
-	C05.116.099.370                      Dysostoses
-	C05.116.099.370.231                      Craniofacial Dysostosis
-	C05.116.099.370.231.427                      Hallermann's Syndrome
-	C05.116.099.370.231.480                      Hypertelorism
-	C05.116.099.370.231.576                      Mandibulofacial Dysostosis
-	C05.116.099.370.231.576.410                      Goldenhar Syndrome
-	C05.116.099.370.380                      Focal Dermal Hypoplasia
-	C05.116.099.370.535                      Klippel-Feil Syndrome
-	C05.116.099.370.652                      Orofaciodigital Syndromes
-	C05.116.099.370.797                      Rubinstein-Taybi Syndrome
-	C05.116.099.370.894                      Synostosis
-	C05.116.099.370.894.115                      Antley-Bixler Syndrome Phenotype
-	C05.116.099.370.894.232                      Craniosynostoses
-	C05.116.099.370.894.232.015                      Acrocephalosyndactylia
-	C05.116.099.370.894.819                      Syndactyly
-	C05.116.099.370.894.819.100                      Acrocephalosyndactylia
New Tree	<a href="#">C05.116.099.370.894.819.428</a> <a href="#">Fraser Syndrome</a>
-	C05.116.099.370.894.819.756                      Poland Syndrome
New Heading	<b>C05.116.099.370.894.909</b> <b>Tarsal Coalition</b>
-	C05.116.099.386                      Funnel Chest
-	C05.116.099.492                      Gigantism
-	C05.116.099.655                      Leg Length Inequality
-	C05.116.099.674                      Marfan Syndrome
-	C05.116.099.708                      Osteochondrodysplasias
-	C05.116.099.708.017                      Achondroplasia
-	C05.116.099.708.017.500                      Thanatophoric Dysplasia
-	C05.116.099.708.025                      Acquired Hyperostosis Syndrome

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C05.116.099.708.180 Camurati-Engelmann Syndrome
-	C05.116.099.708.195 Chondrodysplasia Punctata
-	C05.116.099.708.195.200 Chondrodysplasia Punctata, Rhizomelic
-	C05.116.099.708.207 Cleidocranial Dysplasia
-	C05.116.099.708.327 Ellis-Van Creveld Syndrome
-	C05.116.099.708.338 Enchondromatosis
-	C05.116.099.708.375 Fibrous Dysplasia of Bone
-	C05.116.099.708.375.199 Cherubism
-	C05.116.099.708.375.372 Fibrous Dysplasia, Monostotic
-	C05.116.099.708.375.381 Fibrous Dysplasia, Polyostotic
-	C05.116.099.708.479 Hyperostosis, Cortical, Congenital
-	C05.116.099.708.486 Hyperostosis Frontalis Interna
-	C05.116.099.708.534 Kashin-Beck Disease
-	C05.116.099.708.582 Langer-Giedion Syndrome
-	C05.116.099.708.670 Osteochondroma
-	C05.116.099.708.670.615 Osteochondromatosis
-	C05.116.099.708.670.615.325 Exostoses, Multiple Hereditary
-	C05.116.099.708.685 Osteogenesis Imperfecta
-	C05.116.099.708.702 Osteosclerosis
-	C05.116.099.708.702.593 Melorheostosis
-	C05.116.099.708.702.678 Osteopetrosis
-	C05.116.099.708.702.685 Osteopoikilosis
-	C05.116.099.708.779 Pycnodysostosis
-	C05.116.099.708.857 Short Rib-Polydactyly Syndrome
-	C05.116.099.708.928 Slipped Capital Femoral Epiphyses
-	C05.116.099.736 Osteolysis, Essential
-	C05.116.099.739 Pectus Carinatum
-	C05.116.099.742 Platybasia
-	C05.116.099.750 Proteus Syndrome
-	C05.116.132 Bone Diseases, Endocrine
-	C05.116.132.082 Acromegaly
-	C05.116.132.256 Congenital Hypothyroidism
-	C05.116.132.358 Dwarfism, Pituitary
-	C05.116.132.479 Gigantism
-	C05.116.132.684 Osteitis Fibrosa Cystica
-	C05.116.165 Bone Diseases, Infectious

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C05.116.165.495 Osteomyelitis
-	C05.116.165.495.249 Mastoiditis
-	C05.116.165.495.500 Petrositis
-	C05.116.165.495.750 Pott Puffy Tumor
-	C05.116.165.595 Periostitis
-	C05.116.165.762 Spondylitis
-	C05.116.165.762.301 Discitis
-	C05.116.165.886 Tuberculosis, Osteoarticular
-	C05.116.165.886.722 Tuberculosis, Spinal
-	C05.116.198 Bone Diseases, Metabolic
-	C05.116.198.247 Bone Demineralization, Pathologic
-	C05.116.198.247.400 Decalcification, Pathologic
-	C05.116.198.371 Mucopolidoses
-	C05.116.198.579 Osteoporosis
-	C05.116.198.579.304 Female Athlete Triad Syndrome
-	C05.116.198.579.610 Osteoporosis, Postmenopausal
-	C05.116.198.709 Pseudohypoparathyroidism
-	C05.116.198.709.628 Pseudopseudohypoparathyroidism
-	C05.116.198.816 Rickets
-	C05.116.198.816.640 Osteomalacia
-	C05.116.198.816.750 Chronic Kidney Disease-Mineral and Bone Disorder
-	C05.116.198.816.750 Renal Osteodystrophy
-	C05.116.198.816.875 Rickets, Hypophosphatemic
-	C05.116.198.816.875.500 Familial Hypophosphatemic Rickets
-	C05.116.214 Bone Malalignment
-	C05.116.214.500 Bone Anteversion
-	C05.116.214.500.500 Coxa Vara
-	C05.116.214.750 Bone Retroversion
-	C05.116.214.750.500 Coxa Valga
-	C05.116.231 Bone Neoplasms
-	C05.116.231.030 Adamantinoma
-	C05.116.231.343 Femoral Neoplasms
-	C05.116.231.754 Skull Neoplasms
-	C05.116.231.754.450 Jaw Neoplasms
-	C05.116.231.754.450.583 Mandibular Neoplasms
-	C05.116.231.754.450.601 Maxillary Neoplasms

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C05.116.231.754.450.692 Palatal Neoplasms
-	C05.116.231.754.600 Nose Neoplasms
-	C05.116.231.754.659 Orbital Neoplasms
-	C05.116.231.754.829 Skull Base Neoplasms
-	C05.116.231.828 Spinal Neoplasms
-	C05.116.264 Bone Resorption
-	C05.116.264.143 Ainhum
-	C05.116.264.150 Alveolar Bone Loss
-	C05.116.264.579 Osteolysis
-	C05.116.264.579.052 Acro-Osteolysis
-	C05.116.264.579.052.400 Hajdu-Cheney Syndrome
-	C05.116.264.579.704 Osteolysis, Essential
New Heading	<b>C05.116.296 Coxa Magna</b>
-	C05.116.327 Coxa Valga
New Tree	<b>C05.116.359 Dislocations</b>
New Tree	<b>C05.116.359 Joint Dislocations</b>
New Heading	<b>C05.116.359.192 Diastasis, Bone</b>
New Tree	<b>C05.116.359.192.500 Pubic Symphysis Diastasis</b>
New Heading	<b>C05.116.359.288 Diastasis, Muscle</b>
New Heading	<b>C05.116.359.336 Fracture Dislocation</b>
New Tree	<b>C05.116.359.336.500 Colles' Fracture</b>
New Tree	<b>C05.116.359.336.750 Monteggia's Fracture</b>
New Heading	<b>C05.116.359.336.875 Salter-Harris Fractures</b>
New Tree	<b>C05.116.359.384 Hip Dislocation</b>
New Tree	<b>C05.116.359.500 Knee Dislocation</b>
New Tree	<b>C05.116.359.625 Patellar Dislocation</b>
New Tree	<b>C05.116.359.750 Shoulder Dislocation</b>

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C05.116.391 Eosinophilic Granuloma
-	C05.116.425 Epiphyses, Slipped
-	C05.116.425.500 Slipped Capital Femoral Epiphyses
-	C05.116.482 Genu Valgum
-	C05.116.511 Genu Varum
-	C05.116.540 Hyperostosis
-	C05.116.540.310 Exostoses
-	C05.116.540.310.500 Exostoses, Multiple Hereditary
-	C05.116.540.310.600 Heel Spur
-	C05.116.540.310.800 Osteophyte
-	C05.116.540.400 Hyperostosis, Cortical, Congenital
-	C05.116.540.410 Hyperostosis, Diffuse Idiopathic Skeletal
-	C05.116.540.420 Hyperostosis Frontalis Interna
-	C05.116.540.600 Hyperostosis, Sternocostoclavicular
-	C05.116.680 Osteitis
-	C05.116.692 Osteitis Deformans
-	C05.116.725 Osteoarthropathy, Primary Hypertrophic
-	C05.116.758 Osteoarthropathy, Secondary Hypertrophic
-	C05.116.791 Osteochondritis
-	C05.116.791.668 Osteochondritis Dissecans
-	C05.116.821 Osteochondrosis
-	C05.116.821.500 Spinal Osteochondrosis
-	C05.116.821.500.500 Scheuermann Disease
-	C05.116.852 Osteonecrosis
-	C05.116.852.087 Bisphosphonate-Associated Osteonecrosis of the Jaw
-	C05.116.852.175 Femur Head Necrosis
-	C05.116.852.175.570 Legg-Calve-Perthes Disease
-	C05.116.900 Spinal Diseases
-	C05.116.900.153 Intervertebral Disc Degeneration
-	C05.116.900.307 Intervertebral Disc Displacement
-	C05.116.900.480 Ossification of Posterior Longitudinal Ligament
-	C05.116.900.540 Platybasia
-	C05.116.900.596 Posterior Cervical Sympathetic Syndrome
-	C05.116.900.800 Spinal Curvatures
-	C05.116.900.800.500 Kyphosis
-	C05.116.900.800.500.500 Scheuermann Disease



## MeSH Tree Changes for 2017

Type	Tree - heading
-	C05.116.900.800.750 Lordosis
-	C05.116.900.800.875 Scoliosis
-	C05.116.900.801 Spinal Neoplasms
-	C05.116.900.808 Spinal Osteochondrosis
-	C05.116.900.808.500 Scheuermann Disease
-	C05.116.900.815 Spinal Osteophytosis
-	C05.116.900.815.651 Hyperostosis, Diffuse Idiopathic Skeletal
-	C05.116.900.825 Spinal Stenosis
-	C05.116.900.853 Spondylitis
-	C05.116.900.853.500 Discitis
-	C05.116.900.853.625 Spondylarthritis
-	C05.116.900.853.625.399 Osteoarthritis, Spine
-	C05.116.900.853.625.800 Spondylarthropathies
-	C05.116.900.853.625.800.424 Arthritis, Psoriatic
-	C05.116.900.853.625.800.637 Arthritis, Reactive
-	C05.116.900.853.625.800.850 Spondylitis, Ankylosing
-	C05.116.900.853.850 Tuberculosis, Spinal
-	C05.116.900.938 Spondylosis
-	C05.116.900.938.500 Spondylolysis
-	C05.116.900.938.500.500 Spondylolisthesis
-	C05.182 Cartilage Diseases
-	C05.182.100 Chondromalacia Patellae
-	C05.182.310 Laryngomalacia
-	C05.182.520 Osteochondritis
-	C05.182.525 Pectus Carinatum
-	C05.182.531 Polychondritis, Relapsing
-	C05.182.790 Tietze's Syndrome
-	C05.182.895 Tracheobronchomalacia
-	C05.182.895.249 Bronchomalacia
-	C05.182.895.500 Tracheomalacia
-	C05.321 Fasciitis
-	C05.321.550 Fasciitis, Necrotizing
-	C05.321.600 Fasciitis, Plantar
-	C05.330 Foot Deformities
Old Tree	<b>C05.330.448 Flatfoot</b>
-	C05.330.488 Foot Deformities, Acquired

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Heading	<b>C05.330.488.050</b>	<b>Bunion</b>
New Tree	<a href="#">C05.330.488.050.500</a>	<a href="#">Bunion, Tailor's</a>
Old Tree	<del>C05.330.488.099</del>	<del>Bunion, Tailor's</del>
Old Tree	<del>C05.330.488.200</del>	<del>Equinus Deformity</del>
-	C05.330.488.300	Hallux Limitus
-	C05.330.488.310	Hallux Rigidus
New Heading	<b>C05.330.488.655</b>	<b>Talipes</b>
New Tree	<a href="#">C05.330.488.655.063</a>	<a href="#">Clubfoot</a>
New Tree	<a href="#">C05.330.488.655.125</a>	<a href="#">Equinus Deformity</a>
New Tree	<a href="#">C05.330.488.655.250</a>	<a href="#">Flatfoot</a>
New Heading	<b>C05.330.488.655.500</b>	<b>Talipes Cavus</b>
-	C05.330.495	Foot Deformities, Congenital
Old Tree	<del>C05.330.495.150</del>	<del>Clubfoot</del>
New Heading	<b>C05.330.495.681</b>	<b>Talipes</b>
New Tree	<a href="#">C05.330.495.681.063</a>	<a href="#">Clubfoot</a>
New Tree	<a href="#">C05.330.495.681.125</a>	<a href="#">Equinus Deformity</a>
New Tree	<a href="#">C05.330.495.681.250</a>	<a href="#">Flatfoot</a>
New Heading	<b>C05.330.495.681.500</b>	<b>Talipes Cavus</b>
New Heading	<b>C05.330.495.787</b>	<b>Tarsal Coalition</b>
-	C05.330.610	Hallux Valgus
-	C05.330.612	Hallux Varus
-	C05.330.615	Hammer Toe Syndrome
New Heading	<b>C05.330.663</b>	<b>Metatarsal Valgus</b>
New Heading	<b>C05.330.711</b>	<b>Metatarsus Varus</b>
-	C05.360	Foot Diseases
-	C05.360.350	Fasciitis, Plantar

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Heading	<b>C05.360.375</b>	<b>Fibromatosis, Plantar</b>
-	C05.360.400	Heel Spur
-	C05.360.500	Metatarsalgia
New Heading	<b>C05.360.500.500</b>	<b>Morton Neuroma</b>
-	C05.360.750	Posterior Tibial Tendon Dysfunction
-	C05.390	Hand Deformities
-	C05.390.110	Hand Deformities, Acquired
-	C05.390.408	Hand Deformities, Congenital
-	C05.500	Jaw Diseases
-	C05.500.086	Bisphosphonate-Associated Osteonecrosis of the Jaw
-	C05.500.174	Cherubism
-	C05.500.368	Granuloma, Giant Cell
-	C05.500.460	Jaw Abnormalities
-	C05.500.460.185	Cleft Palate
-	C05.500.460.457	Micrognathism
-	C05.500.460.606	Pierre Robin Syndrome
-	C05.500.460.655	Prognathism
-	C05.500.460.827	Retrognathia
-	C05.500.470	Jaw Cysts
-	C05.500.470.660	Nonodontogenic Cysts
-	C05.500.470.690	Odontogenic Cysts
-	C05.500.470.690.150	Basal Cell Nevus Syndrome
-	C05.500.470.690.310	Dentigerous Cyst
-	C05.500.470.690.605	Odontogenic Cyst, Calcifying
-	C05.500.470.690.790	Periodontal Cyst
-	C05.500.470.690.790.820	Radicular Cyst
-	C05.500.480	Jaw, Edentulous
-	C05.500.480.450	Jaw, Edentulous, Partially
-	C05.500.499	Jaw Neoplasms
-	C05.500.499.583	Mandibular Neoplasms
-	C05.500.499.601	Maxillary Neoplasms
-	C05.500.499.692	Palatal Neoplasms
-	C05.500.607	Mandibular Diseases
-	C05.500.607.221	Craniomandibular Disorders

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C05.500.607.221.897 Temporomandibular Joint Disorders
-	C05.500.607.221.897.897 Temporomandibular Joint Dysfunction Syndrome
-	C05.500.607.442 Mandibular Neoplasms
-	C05.500.607.655 Prognathism
-	C05.500.693 Maxillary Diseases
-	C05.500.693.528 Maxillary Neoplasms
-	C05.550 Joint Diseases
-	C05.550.069 Ankylosis
-	C05.550.069.680 Spondylitis, Ankylosing
-	C05.550.091 Arthralgia
-	C05.550.091.700 Shoulder Pain
-	C05.550.114 Arthritis
-	C05.550.114.015 Arthritis, Experimental
-	C05.550.114.099 Arthritis, Infectious
-	C05.550.114.099.500 Arthritis, Reactive
-	C05.550.114.122 Arthritis, Juvenile
-	C05.550.114.145 Arthritis, Psoriatic
-	C05.550.114.154 Arthritis, Rheumatoid
-	C05.550.114.154.219 Caplan Syndrome
-	C05.550.114.154.389 Felty Syndrome
-	C05.550.114.154.683 Rheumatoid Nodule
-	C05.550.114.154.728 Rheumatoid Vasculitis
-	C05.550.114.154.774 Sjogren's Syndrome
-	C05.550.114.154.870 Still's Disease, Adult-Onset
-	C05.550.114.264 Chondrocalcinosis
New Heading	<b>C05.550.114.264.500 Rotator Cuff Tear Arthropathy</b>
-	C05.550.114.423 Gout
-	C05.550.114.423.410 Arthritis, Gouty
-	C05.550.114.606 Osteoarthritis
-	C05.550.114.606.400 Osteoarthritis, Hip
-	C05.550.114.606.500 Osteoarthritis, Knee
-	C05.550.114.606.750 Osteoarthritis, Spine
-	C05.550.114.678 Periarthritis
-	C05.550.114.843 Rheumatic Fever
-	C05.550.114.843.566 Rheumatic Nodule

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	C05.550.114.843.823	Wissler's Syndrome
-	C05.550.114.854	Sacroiliitis
-	C05.550.114.865	Spondylarthritis
-	C05.550.114.865.399	Osteoarthritis, Spine
-	C05.550.114.865.800	Spondylarthropathies
-	C05.550.114.865.800.424	Arthritis, Psoriatic
-	C05.550.114.865.800.637	Arthritis, Reactive
-	C05.550.114.865.800.850	Spondylitis, Ankylosing
-	C05.550.150	Arthrogyposis
-	C05.550.186	Arthropathy, Neurogenic
-	C05.550.251	Bursitis
-	C05.550.251.595	Periarthritis
-	C05.550.287	Chondromatosis, Synovial
-	C05.550.323	Contracture
-	C05.550.323.468	Hip Contracture
-	C05.550.323.734	Ischemic Contracture
New Heading	<b>C05.550.354</b>	<b>Crystal Arthropathies</b>
New Tree	<a href="#">C05.550.354.125</a>	<a href="#">Chondrocalcinosis</a>
New Heading	<b>C05.550.354.250</b>	<b>Rotator Cuff Tear Arthropathy</b>
New Tree	<a href="#">C05.550.354.500</a>	<a href="#">Gout</a>
New Tree	<a href="#">C05.550.354.500.500</a>	<a href="#">Arthritis, Gouty</a>
-	C05.550.384	Femoracetabular Impingement
-	C05.550.445	Hallux Limitus
-	C05.550.450	Hallux Rigidus
-	C05.550.459	Hemarthrosis
-	C05.550.509	Hydrarthrosis
-	C05.550.515	Joint Deformities, Acquired
-	C05.550.521	Joint Instability
-	C05.550.535	Joint Loose Bodies
-	C05.550.610	Metatarsalgia
New Heading	<b>C05.550.610.500</b>	<b>Morton Neuroma</b>
-	C05.550.629	Nail-Patella Syndrome

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	C05.550.648	Osteoarthropathy, Primary Hypertrophic
-	C05.550.684	Osteoarthropathy, Secondary Hypertrophic
-	C05.550.700	Patellofemoral Pain Syndrome
-	C05.550.840	Shoulder Impingement Syndrome
-	C05.550.870	Synovitis
New Heading	<b>C05.550.870.445</b>	<b>Giant Cell Tumor of Tendon Sheath</b>
New Tree	<a href="#">C05.550.870.445.500</a>	<a href="#">Synovitis, Pigmented Villonodular</a>
Old Tree	<del>C05.550.870.890</del>	<del>Synovitis, Pigmented Villonodular</del>
-	C05.550.905	Temporomandibular Joint Disorders
-	C05.550.905.905	Temporomandibular Joint Dysfunction Syndrome
-	C05.651	Muscular Diseases
-	C05.651.102	Arthrogryposis
-	C05.651.180	Compartment Syndromes
-	C05.651.180.063	Anterior Compartment Syndrome
-	C05.651.180.297	Intra-Abdominal Hypertension
-	C05.651.180.531	Ischemic Contracture
-	C05.651.197	Contracture
-	C05.651.197.270	Dupuytren Contracture
New Heading	<b>C05.651.197.369</b>	<b>Fibromatosis, Plantar</b>
-	C05.651.197.468	Hip Contracture
-	C05.651.197.734	Ischemic Contracture
-	C05.651.243	Craniomandibular Disorders
-	C05.651.243.897	Temporomandibular Joint Disorders
-	C05.651.243.897.897	Temporomandibular Joint Dysfunction Syndrome
-	C05.651.290	Eosinophilia-Myalgia Syndrome
-	C05.651.310	Fatigue Syndrome, Chronic
-	C05.651.324	Fibromyalgia
-	C05.651.392	Isaacs Syndrome
-	C05.651.426	Medial Tibial Stress Syndrome
-	C05.651.460	Mitochondrial Myopathies
-	C05.651.460.620	Mitochondrial Encephalomyopathies
-	C05.651.460.620.520	MELAS Syndrome
-	C05.651.460.620.530	MERRF Syndrome
-	C05.651.460.700	Ophthalmoplegia, Chronic Progressive External

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C05.651.460.700.500                      Kearns-Sayre Syndrome
-	C05.651.475                                    Muscle Cramp
-	C05.651.494                                    Muscle Neoplasms
-	C05.651.504                                    Muscle Rigidity
-	C05.651.512                                    Muscle Spasticity
-	C05.651.515                                    Muscle Weakness
-	C05.651.534                                    Muscular Disorders, Atrophic
-	C05.651.534.500                              Muscular Dystrophies
-	C05.651.534.500.074                        Distal Myopathies
-	C05.651.534.500.149                        Glycogen Storage Disease Type VII
-	C05.651.534.500.280                        Muscular Dystrophies, Limb-Girdle
-	C05.651.534.500.280.500                    Sarcoglycanopathies
-	C05.651.534.500.300                        Muscular Dystrophy, Duchenne
-	C05.651.534.500.350                        Muscular Dystrophy, Emery-Dreifuss
-	C05.651.534.500.400                        Muscular Dystrophy, Facioscapulohumeral
-	C05.651.534.500.450                        Muscular Dystrophy, Oculopharyngeal
-	C05.651.534.500.500                        Myotonic Dystrophy
-	C05.651.534.750                              Postpoliomyelitis Syndrome
-	C05.651.538                                    Musculoskeletal Pain
-	C05.651.542                                    Myalgia
-	C05.651.550                                    Myofascial Pain Syndromes
-	C05.651.550.905                              Temporomandibular Joint Dysfunction Syndrome
-	C05.651.575                                    Myopathies, Structural, Congenital
-	C05.651.575.290                              Myopathies, Nemaline
-	C05.651.575.300                              Myopathy, Central Core
-	C05.651.594                                    Myositis
-	C05.651.594.600                              Myositis, Inclusion Body
-	C05.651.594.638                              Myositis Ossificans
-	C05.651.594.728                              Orbital Myositis
-	C05.651.594.819                              Polymyositis
-	C05.651.594.819.500                        Dermatomyositis
-	C05.651.594.909                              Pyomyositis
-	C05.651.662                                    Myotonic Disorders
-	C05.651.662.500                              Myotonia Congenita
-	C05.651.662.750                              Myotonic Dystrophy
-	C05.651.701                                    Paralyses, Familial Periodic

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	C05.651.701.450	Hypokalemic Periodic Paralysis
-	C05.651.701.600	Paralysis, Hyperkalemic Periodic
-	C05.651.742	Polymyalgia Rheumatica
-	C05.651.807	Rhabdomyolysis
-	C05.651.807.628	Myoglobinuria
-	C05.651.869	Tendinopathy
New Heading	<b>C05.651.869.435</b>	<b>Elbow Tendinopathy</b>
New Tree	<a href="#">C05.651.869.435.500</a>	<a href="#">Tennis Elbow</a>
New Heading	<b>C05.651.869.653</b>	<b>Enthesopathy</b>
New Heading	<b>C05.651.869.762</b>	<b>Giant Cell Tumor of Tendon Sheath</b>
New Tree	<a href="#">C05.651.869.762.500</a>	<a href="#">Synovitis, Pigmented Villonodular</a>
New Tree	<a href="#">C05.651.869.816</a>	<a href="#">Tendon Entrapment</a>
New Tree	<a href="#">C05.651.869.816.200</a>	<a href="#">De Quervain Disease</a>
New Tree	<a href="#">C05.651.869.816.800</a>	<a href="#">Trigger Finger Disorder</a>
-	C05.651.869.870	Tenosynovitis
Old Tree	<del>C05.651.869.870.800</del>	<del>Tendon Entrapment</del>
Old Tree	<del>C05.651.869.870.800.200</del>	<del>De Quervain Disease</del>
Old Tree	<del>C05.651.869.870.800.800</del>	<del>Trigger Finger Disorder</del>
-	C05.660	Musculoskeletal Abnormalities
-	C05.660.077	Arthrogryposis
-	C05.660.142	Campomelic Dysplasia
-	C05.660.207	Craniofacial Abnormalities
-	C05.660.207.103	22q11 Deletion Syndrome
-	C05.660.207.103.500	DiGeorge Syndrome
-	C05.660.207.207	Cleidocranial Dysplasia
-	C05.660.207.219	Costello Syndrome
-	C05.660.207.231	Craniofacial Dysostosis
-	C05.660.207.231.427	Hallermann's Syndrome
-	C05.660.207.231.480	Hypertelorism
-	C05.660.207.231.576	Mandibulofacial Dysostosis
-	C05.660.207.231.576.410	Goldenhar Syndrome



## MeSH Tree Changes for 2017

Type	Tree - heading
-	C05.660.207.240 Craniosynostoses
-	C05.660.207.240.100 Acrocephalosyndactylia
-	C05.660.207.325 Donohue Syndrome
-	C05.660.207.410 Holoprosencephaly
-	C05.660.207.525 LEOPARD Syndrome
-	C05.660.207.532 Loeys-Dietz Syndrome
-	C05.660.207.536 Megalencephaly
-	C05.660.207.536.500 Hemimegalencephaly
-	C05.660.207.540 Maxillofacial Abnormalities
-	C05.660.207.540.229 Dentofacial Deformities
-	C05.660.207.540.460 Jaw Abnormalities
-	C05.660.207.540.460.185 Cleft Palate
-	C05.660.207.540.460.457 Micrognathism
-	C05.660.207.540.460.606 Pierre Robin Syndrome
-	C05.660.207.540.460.655 Prognathism
-	C05.660.207.540.460.827 Retrognathia
-	C05.660.207.620 Microcephaly
-	C05.660.207.620.500 Porencephaly
-	C05.660.207.690 Noonan Syndrome
-	C05.660.207.700 Orofaciodigital Syndromes
-	C05.660.207.707 Plagiocephaly
-	C05.660.207.707.624 Plagiocephaly, Nonsynostotic
-	C05.660.207.720 Platybasia
-	C05.660.207.850 Rubinstein-Taybi Syndrome
-	C05.660.207.925 Silver-Russell Syndrome
-	C05.660.386 Funnel Chest
-	C05.660.417 Gastroschisis
-	C05.660.449 Hip Dislocation, Congenital
-	C05.660.551 Klippel-Feil Syndrome
-	C05.660.585 Limb Deformities, Congenital
-	C05.660.585.174 Arachnodactyly
-	C05.660.585.262 Brachydactyly
-	C05.660.585.350 Ectromelia
-	C05.660.585.512 Lower Extremity Deformities, Congenital
-	C05.660.585.512.380 Foot Deformities, Congenital
Old Tree	<b>C05.660.585.512.380.500</b> <b>Clubfoot</b>

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Heading	<b>C05.660.585.512.380.813</b>	<b>Talipes</b>
New Tree	<a href="#">C05.660.585.512.380.813.063</a>	<a href="#">Clubfoot</a>
New Tree	<a href="#">C05.660.585.512.380.813.125</a>	<a href="#">Equinus Deformity</a>
New Tree	<a href="#">C05.660.585.512.380.813.250</a>	<a href="#">Flatfoot</a>
New Heading	<b>C05.660.585.512.380.813.500</b>	<b>Talipes Cavus</b>
New Heading	<b>C05.660.585.512.380.875</b>	<b>Tarsal Coalition</b>
-	C05.660.585.600	Polydactyly
-	C05.660.585.600.374	Pallister-Hall Syndrome
-	C05.660.585.600.750	Short Rib-Polydactyly Syndrome
-	C05.660.585.620	Proteus Syndrome
-	C05.660.585.800	Syndactyly
-	C05.660.585.800.100	Acrocephalosyndactylia
New Tree	<a href="#">C05.660.585.800.428</a>	<a href="#">Fraser Syndrome</a>
-	C05.660.585.800.756	Poland Syndrome
-	C05.660.585.984	Thanatophoric Dysplasia
-	C05.660.585.988	Upper Extremity Deformities, Congenital
-	C05.660.585.988.425	Hand Deformities, Congenital
-	C05.660.745	Pectus Carinatum
-	C05.660.906	Synostosis
-	C05.660.906.181	Antley-Bixler Syndrome Phenotype
-	C05.660.906.364	Craniosynostoses
-	C05.660.906.364.100	Acrocephalosyndactylia
-	C05.660.906.819	Syndactyly
-	C05.660.906.819.100	Acrocephalosyndactylia
New Tree	<a href="#">C05.660.906.819.428</a>	<a href="#">Fraser Syndrome</a>
-	C05.660.906.819.756	Poland Syndrome
New Heading	<b>C05.660.906.909</b>	<b>Tarsal Coalition</b>
-	C05.799	Rheumatic Diseases
-	C05.799.056	Arthritis, Juvenile
-	C05.799.114	Arthritis, Rheumatoid

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	C05.799.114.219	Caplan Syndrome
-	C05.799.114.389	Felty Syndrome
-	C05.799.114.683	Rheumatoid Nodule
-	C05.799.114.728	Rheumatoid Vasculitis
-	C05.799.114.774	Sjogren's Syndrome
-	C05.799.114.870	Still's Disease, Adult-Onset
-	C05.799.321	Fibromyalgia
-	C05.799.414	Gout
-	C05.799.414.410	Arthritis, Gouty
-	C05.799.440	Hyperostosis, Sternocostoclavicular
-	C05.799.613	Osteoarthritis
-	C05.799.613.400	Osteoarthritis, Hip
-	C05.799.613.500	Osteoarthritis, Knee
-	C05.799.613.750	Osteoarthritis, Spine
-	C05.799.720	Polymyalgia Rheumatica
-	C05.799.825	Rheumatic Fever
-	C05.799.825.566	Rheumatic Nodule
-	C05.799.825.823	Wissler's Syndrome
Old Tree	C05.906	Tennis Elbow
-	C06	Digestive System Diseases
-	C06.130	Biliary Tract Diseases
-	C06.130.120	Bile Duct Diseases
-	C06.130.120.120	Bile Duct Neoplasms
-	C06.130.120.120.280	Common Bile Duct Neoplasms
-	C06.130.120.123	Biliary Atresia
-	C06.130.120.127	Choledochal Cyst
-	C06.130.120.127.500	Caroli Disease
-	C06.130.120.135	Cholestasis
-	C06.130.120.135.150	Cholestasis, Extrahepatic
-	C06.130.120.135.250	Cholestasis, Intrahepatic
-	C06.130.120.135.250.125	Alagille Syndrome
-	C06.130.120.135.250.250	Liver Cirrhosis, Biliary
-	C06.130.120.135.812	Mirizzi Syndrome
-	C06.130.120.200	Cholangitis
-	C06.130.120.200.110	Cholangitis, Sclerosing
-	C06.130.120.250	Common Bile Duct Diseases

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C06.130.120.250.098 Biliary Dyskinesia
-	C06.130.120.250.098.800 Sphincter of Oddi Dysfunction
-	C06.130.120.250.174 Choledocholithiasis
-	C06.130.120.250.280 Common Bile Duct Neoplasms
-	C06.130.140 Bile Reflux
-	C06.130.320 Biliary Tract Neoplasms
-	C06.130.320.120 Bile Duct Neoplasms
-	C06.130.320.120.280 Common Bile Duct Neoplasms
-	C06.130.320.401 Gallbladder Neoplasms
-	C06.130.409 Cholelithiasis
-	C06.130.409.178 Cholecystolithiasis
-	C06.130.409.267 Choledocholithiasis
-	C06.130.409.633 Gallstones
-	C06.130.564 Gallbladder Diseases
-	C06.130.564.263 Cholecystitis
-	C06.130.564.263.249 Acalculous Cholecystitis
-	C06.130.564.263.500 Cholecystitis, Acute
-	C06.130.564.263.500.500 Emphysematous Cholecystitis
-	C06.130.564.332 Cholecystolithiasis
-	C06.130.564.332.500 Gallstones
-	C06.130.564.401 Gallbladder Neoplasms
-	C06.130.825 Postcholecystectomy Syndrome
-	C06.198 Digestive System Abnormalities
New Heading	<b>C06.198.025 Anorectal Malformations</b>
-	C06.198.050 Anus, Imperforate
-	C06.198.102 Barrett Esophagus
-	C06.198.125 Biliary Atresia
-	C06.198.184 Choledochal Cyst
-	C06.198.184.500 Caroli Disease
-	C06.198.257 Diaphragmatic Eventration
-	C06.198.330 Esophageal Atresia
-	C06.198.439 Hirschsprung Disease
-	C06.198.719 Intestinal Atresia
-	C06.198.859 Meckel Diverticulum
-	C06.267 Digestive System Fistula

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C06.267.150 Biliary Fistula
-	C06.267.250 Esophageal Fistula
-	C06.267.250.725 Tracheoesophageal Fistula
-	C06.267.375 Gastric Fistula
-	C06.267.550 Intestinal Fistula
-	C06.267.550.600 Rectal Fistula
-	C06.267.550.600.650 Rectovaginal Fistula
-	C06.267.775 Pancreatic Fistula
-	C06.301 Digestive System Neoplasms
-	C06.301.120 Biliary Tract Neoplasms
-	C06.301.120.250 Bile Duct Neoplasms
-	C06.301.120.250.250 Common Bile Duct Neoplasms
-	C06.301.120.401 Gallbladder Neoplasms
-	C06.301.371 Gastrointestinal Neoplasms
-	C06.301.371.205 Esophageal Neoplasms
-	C06.301.371.308 Gastrointestinal Stromal Tumors
-	C06.301.371.411 Intestinal Neoplasms
-	C06.301.371.411.184 Cecal Neoplasms
-	C06.301.371.411.184.290 Appendiceal Neoplasms
-	C06.301.371.411.307 Colorectal Neoplasms
-	C06.301.371.411.307.090 Adenomatous Polyposis Coli
-	C06.301.371.411.307.090.500 Gardner Syndrome
-	C06.301.371.411.307.180 Colonic Neoplasms
-	C06.301.371.411.307.180.800 Sigmoid Neoplasms
-	C06.301.371.411.307.190 Colorectal Neoplasms, Hereditary Nonpolyposis
-	C06.301.371.411.307.790 Rectal Neoplasms
-	C06.301.371.411.307.790.040 Anus Neoplasms
-	C06.301.371.411.307.790.040.040 Anal Gland Neoplasms
-	C06.301.371.411.445 Duodenal Neoplasms
-	C06.301.371.411.501 Ileal Neoplasms
-	C06.301.371.411.512 Immunoproliferative Small Intestinal Disease
-	C06.301.371.411.523 Jejunal Neoplasms
-	C06.301.371.767 Stomach Neoplasms
-	C06.301.371.883 Zollinger-Ellison Syndrome
-	C06.301.623 Liver Neoplasms
-	C06.301.623.040 Adenoma, Liver Cell

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C06.301.623.160                      Carcinoma, Hepatocellular
-	C06.301.623.460                      Liver Neoplasms, Experimental
-	C06.301.761                            Pancreatic Neoplasms
-	C06.301.761.249                      Adenoma, Islet Cell
-	C06.301.761.249.500                Insulinoma
-	C06.301.761.500                      Carcinoma, Islet Cell
-	C06.301.761.500.124                Gastrinoma
-	C06.301.761.500.249                Glucagonoma
-	C06.301.761.500.500                Somatostatinoma
-	C06.301.761.500.750                Vipoma
-	C06.301.761.750                      Carcinoma, Pancreatic Ductal
-	C06.301.780                            Peritoneal Neoplasms
-	C06.405                                 Gastrointestinal Diseases
-	C06.405.117                            Esophageal Diseases
-	C06.405.117.102                      Barrett Esophagus
-	C06.405.117.119                      Deglutition Disorders
-	C06.405.117.119.500                Esophageal Motility Disorders
-	C06.405.117.119.500.204            CREST Syndrome
-	C06.405.117.119.500.432            Esophageal Achalasia
-	C06.405.117.119.500.450            Esophageal Spasm, Diffuse
-	C06.405.117.119.500.484            Gastroesophageal Reflux
-	C06.405.117.119.500.484.500       Laryngopharyngeal Reflux
-	C06.405.117.119.500.484.500.500    Respiratory Aspiration of Gastric Contents
-	C06.405.117.119.500.742            Plummer-Vinson Syndrome
-	C06.405.117.136                      Diverticulosis, Esophageal
-	C06.405.117.240                      Esophageal and Gastric Varices
-	C06.405.117.260                      Esophageal Atresia
-	C06.405.117.316                      Esophageal Cyst
-	C06.405.117.367                      Esophageal Fistula
-	C06.405.117.367.725                Tracheoesophageal Fistula
-	C06.405.117.430                      Esophageal Neoplasms
-	C06.405.117.468                      Esophageal Perforation
-	C06.405.117.468.524                Mallory-Weiss Syndrome
-	C06.405.117.544                      Esophageal Stenosis
-	C06.405.117.620                      Esophagitis
-	C06.405.117.620.209                Eosinophilic Esophagitis

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C06.405.117.620.420 Esophagitis, Peptic
-	C06.405.205 Gastroenteritis
-	C06.405.205.099 Appendicitis
-	C06.405.205.200 Cholera Morbus
-	C06.405.205.265 Colitis
-	C06.405.205.265.115 Colitis, Ischemic
-	C06.405.205.265.173 Colitis, Microscopic
-	C06.405.205.265.173.500 Colitis, Collagenous
-	C06.405.205.265.173.750 Colitis, Lymphocytic
-	C06.405.205.265.231 Colitis, Ulcerative
-	C06.405.205.265.767 Proctocolitis
-	C06.405.205.298 Diverticulitis
-	C06.405.205.331 Dysentery
-	C06.405.205.331.312 Dysentery, Amebic
-	C06.405.205.331.479 Dysentery, Bacillary
-	C06.405.205.462 Enteritis
-	C06.405.205.462.249 Duodenitis
-	C06.405.205.462.624 Ileitis
-	C06.405.205.462.624.500 Pouchitis
-	C06.405.205.596 Enterocolitis
-	C06.405.205.596.700 Enterocolitis, Necrotizing
-	C06.405.205.596.750 Enterocolitis, Neutropenic
-	C06.405.205.596.800 Enterocolitis, Pseudomembranous
-	C06.405.205.663 Esophagitis
-	C06.405.205.663.209 Eosinophilic Esophagitis
-	C06.405.205.663.420 Esophagitis, Peptic
-	C06.405.205.697 Gastritis
-	C06.405.205.697.394 Gastritis, Atrophic
-	C06.405.205.697.410 Gastritis, Hypertrophic
-	C06.405.205.731 Inflammatory Bowel Diseases
-	C06.405.205.731.249 Colitis, Ulcerative
-	C06.405.205.731.500 Crohn Disease
-	C06.405.205.798 Mucositis
-	C06.405.205.865 Proctitis
-	C06.405.205.865.790 Proctocolitis
-	C06.405.205.932 Typhlitis

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C06.405.227                      Gastrointestinal Hemorrhage
-	C06.405.227.400                  Hematemesis
-	C06.405.227.600                  Melena
-	C06.405.227.700                  Peptic Ulcer Hemorrhage
-	C06.405.249                      Gastrointestinal Neoplasms
-	C06.405.249.205                  Esophageal Neoplasms
-	C06.405.249.308                  Gastrointestinal Stromal Tumors
-	C06.405.249.411                  Intestinal Neoplasms
-	C06.405.249.411.184              Cecal Neoplasms
-	C06.405.249.411.184.290        Appendiceal Neoplasms
-	C06.405.249.411.307              Colorectal Neoplasms
-	C06.405.249.411.307.090        Adenomatous Polyposis Coli
-	C06.405.249.411.307.090.500    Gardner Syndrome
-	C06.405.249.411.307.180        Colonic Neoplasms
-	C06.405.249.411.307.180.800    Sigmoid Neoplasms
-	C06.405.249.411.307.190        Colorectal Neoplasms, Hereditary Nonpolyposis
-	C06.405.249.411.307.790        Rectal Neoplasms
-	C06.405.249.411.307.790.040    Anus Neoplasms
-	C06.405.249.411.307.790.040.040    Anal Gland Neoplasms
-	C06.405.249.411.445              Duodenal Neoplasms
-	C06.405.249.411.501              Ileal Neoplasms
-	C06.405.249.411.512              Immunoproliferative Small Intestinal Disease
-	C06.405.249.411.523              Jejunal Neoplasms
-	C06.405.249.767                  Stomach Neoplasms
-	C06.405.249.883                  Zollinger-Ellison Syndrome
-	C06.405.469                      Intestinal Diseases
-	C06.405.469.110                  Cecal Diseases
-	C06.405.469.110.207              Appendicitis
-	C06.405.469.110.417              Cecal Neoplasms
-	C06.405.469.110.417.290        Appendiceal Neoplasms
-	C06.405.469.110.708              Typhlitis
-	C06.405.469.158                  Colonic Diseases
-	C06.405.469.158.093              Chilaiditi Syndrome
-	C06.405.469.158.188              Colitis
-	C06.405.469.158.188.115        Colitis, Ischemic
-	C06.405.469.158.188.173        Colitis, Microscopic



## MeSH Tree Changes for 2017

Type	Tree - heading
-	C06.405.469.158.188.173.500                      Colitis, Collagenous
-	C06.405.469.158.188.173.750                      Colitis, Lymphocytic
-	C06.405.469.158.188.231                          Colitis, Ulcerative
-	C06.405.469.158.188.767                        Proctocolitis
-	C06.405.469.158.272                            Colonic Diseases, Functional
-	C06.405.469.158.272.217                        Colonic Pseudo-Obstruction
-	C06.405.469.158.272.608                        Irritable Bowel Syndrome
-	C06.405.469.158.272.804                        Neurogenic Bowel
-	C06.405.469.158.356                            Colorectal Neoplasms
-	C06.405.469.158.356.090                        Adenomatous Polyposis Coli
-	C06.405.469.158.356.090.500                   Gardner Syndrome
-	C06.405.469.158.356.180                        Colonic Neoplasms
-	C06.405.469.158.356.180.800                   Sigmoid Neoplasms
-	C06.405.469.158.356.190                        Colorectal Neoplasms, Hereditary Nonpolyposis
-	C06.405.469.158.587                            Diverticulosis, Colonic
-	C06.405.469.158.587.500                        Diverticulitis, Colonic
-	C06.405.469.158.701                            Megacolon
-	C06.405.469.158.701.439                        Hirschsprung Disease
-	C06.405.469.158.701.591                        Megacolon, Toxic
-	C06.405.469.158.850                            Sigmoid Diseases
-	C06.405.469.158.850.790                        Proctocolitis
-	C06.405.469.158.850.850                        Sigmoid Neoplasms
-	C06.405.469.275                                Duodenal Diseases
-	C06.405.469.275.270                            Duodenal Neoplasms
-	C06.405.469.275.395                            Duodenal Obstruction
-	C06.405.469.275.395.890                        Superior Mesenteric Artery Syndrome
-	C06.405.469.275.600                            Duodenitis
-	C06.405.469.275.700                            Duodenogastric Reflux
-	C06.405.469.275.800                            Peptic Ulcer
-	C06.405.469.275.800.348                        Duodenal Ulcer
-	C06.405.469.275.800.523                        Esophagitis, Peptic
-	C06.405.469.275.800.698                        Peptic Ulcer Perforation
-	C06.405.469.275.800.849                        Stomach Ulcer
-	C06.405.469.275.800.924                        Zollinger-Ellison Syndrome
-	C06.405.469.300                                Dysentery
-	C06.405.469.300.312                            Dysentery, Amebic

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C06.405.469.300.479                      Dysentery, Bacillary
-	C06.405.469.326                            Enteritis
-	C06.405.469.326.750                      Duodenitis
-	C06.405.469.326.875                      Ileitis
-	C06.405.469.326.875.500                Pouchitis
-	C06.405.469.363                            Enterocolitis
-	C06.405.469.363.700                      Enterocolitis, Necrotizing
-	C06.405.469.363.750                      Enterocolitis, Neutropenic
-	C06.405.469.363.800                      Enterocolitis, Pseudomembranous
-	C06.405.469.400                            HIV Enteropathy
-	C06.405.469.420                            Ileal Diseases
-	C06.405.469.420.501                      Ileal Neoplasms
-	C06.405.469.420.520                      Ileitis
-	C06.405.469.420.520.500                Pouchitis
-	C06.405.469.432                            Inflammatory Bowel Diseases
-	C06.405.469.432.249                      Colitis, Ulcerative
-	C06.405.469.432.500                      Crohn Disease
-	C06.405.469.445                            Intestinal Atresia
-	C06.405.469.452                            Intestinal Diseases, Parasitic
-	C06.405.469.452.060                      Anisakiasis
-	C06.405.469.452.146                      Balantidiasis
-	C06.405.469.452.250                      Blastocystis Infections
-	C06.405.469.452.269                      Cryptosporidiosis
-	C06.405.469.452.275                      Dientamoebiasis
-	C06.405.469.452.396                      Dysentery, Amebic
-	C06.405.469.452.481                      Giardiasis
-	C06.405.469.471                            Intestinal Fistula
-	C06.405.469.471.600                      Rectal Fistula
-	C06.405.469.471.600.650                Rectovaginal Fistula
-	C06.405.469.491                            Intestinal Neoplasms
-	C06.405.469.491.184                      Cecal Neoplasms
-	C06.405.469.491.184.290                Appendiceal Neoplasms
-	C06.405.469.491.307                      Colorectal Neoplasms
-	C06.405.469.491.307.090                Adenomatous Polyposis Coli
-	C06.405.469.491.307.090.500        Gardner Syndrome
-	C06.405.469.491.307.180                Colonic Neoplasms

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C06.405.469.491.307.180.800 Sigmoid Neoplasms
-	C06.405.469.491.307.190 Colorectal Neoplasms, Hereditary Nonpolyposis
-	C06.405.469.491.307.790 Rectal Neoplasms
-	C06.405.469.491.307.790.040 Anus Neoplasms
-	C06.405.469.491.307.790.040.040 Anal Gland Neoplasms
-	C06.405.469.491.445 Duodenal Neoplasms
-	C06.405.469.491.501 Ileal Neoplasms
-	C06.405.469.491.505 Immunoproliferative Small Intestinal Disease
-	C06.405.469.491.523 Jejunal Neoplasms
-	C06.405.469.531 Intestinal Obstruction
-	C06.405.469.531.099 Afferent Loop Syndrome
-	C06.405.469.531.311 Duodenal Obstruction
-	C06.405.469.531.424 Fecal Impaction
-	C06.405.469.531.492 Ileus
-	C06.405.469.531.492.500 Intestinal Pseudo-Obstruction
-	C06.405.469.531.492.500.217 Colonic Pseudo-Obstruction
-	C06.405.469.531.568 Intestinal Volvulus
-	C06.405.469.531.577 Intussusception
-	C06.405.469.557 Intestinal Perforation
-	C06.405.469.578 Intestinal Polyposis
-	C06.405.469.578.249 Adenomatous Polyposis Coli
-	C06.405.469.578.249.393 Gardner Syndrome
-	C06.405.469.578.750 Peutz-Jeghers Syndrome
-	C06.405.469.600 Jejunal Diseases
-	C06.405.469.600.523 Jejunal Neoplasms
-	C06.405.469.637 Malabsorption Syndromes
-	C06.405.469.637.145 Blind Loop Syndrome
-	C06.405.469.637.250 Celiac Disease
-	C06.405.469.637.378 Collagenous Sprue
-	C06.405.469.637.506 Lactose Intolerance
-	C06.405.469.637.832 Short Bowel Syndrome
-	C06.405.469.637.850 Sprue, Tropical
-	C06.405.469.637.887 Steatorrhea
-	C06.405.469.637.925 Whipple Disease
-	C06.405.469.656 Mesenteric Ischemia
-	C06.405.469.675 Mesenteric Vascular Occlusion

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C06.405.469.778 Pneumatosis Cystoides Intestinalis
-	C06.405.469.818 Protein-Losing Enteropathies
-	C06.405.469.860 Rectal Diseases
-	C06.405.469.860.101 Anus Diseases
-	C06.405.469.860.101.163 Anus Neoplasms
-	C06.405.469.860.101.163.083 Anal Gland Neoplasms
-	C06.405.469.860.101.430 Fissure in Ano
-	C06.405.469.860.101.752 Pruritus Ani
-	C06.405.469.860.180 Colorectal Neoplasms
-	C06.405.469.860.180.500 Rectal Neoplasms
-	C06.405.469.860.180.500.040 Anus Neoplasms
-	C06.405.469.860.180.500.040.040 Anal Gland Neoplasms
-	C06.405.469.860.300 Fecal Incontinence
-	C06.405.469.860.401 Hemorrhoids
-	C06.405.469.860.622 Proctitis
-	C06.405.469.860.622.790 Proctocolitis
-	C06.405.469.860.752 Rectal Fistula
-	C06.405.469.860.752.650 Rectovaginal Fistula
-	C06.405.469.860.800 Rectal Prolapse
-	C06.405.469.860.810 Rectocele
-	C06.405.469.965 Zollinger-Ellison Syndrome
-	C06.405.748 Stomach Diseases
-	C06.405.748.045 Achlorhydria
-	C06.405.748.142 Diverticulosis, Stomach
-	C06.405.748.240 Duodenogastric Reflux
-	C06.405.748.240.140 Bile Reflux
-	C06.405.748.280 Gastric Antral Vascular Ectasia
-	C06.405.748.300 Gastric Dilatation
-	C06.405.748.340 Gastric Outlet Obstruction
-	C06.405.748.340.690 Pyloric Stenosis
-	C06.405.748.340.690.500 Pyloric Stenosis, Hypertrophic
-	C06.405.748.398 Gastritis
-	C06.405.748.398.394 Gastritis, Atrophic
-	C06.405.748.398.410 Gastritis, Hypertrophic
-	C06.405.748.543 Gastroparesis
-	C06.405.748.586 Peptic Ulcer

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C06.405.748.586.349 Duodenal Ulcer
-	C06.405.748.586.524 Esophagitis, Peptic
-	C06.405.748.586.698 Peptic Ulcer Perforation
-	C06.405.748.586.849 Stomach Ulcer
-	C06.405.748.586.924 Zollinger-Ellison Syndrome
-	C06.405.748.630 Postgastrectomy Syndromes
-	C06.405.748.630.310 Dumping Syndrome
-	C06.405.748.789 Stomach Neoplasms
-	C06.405.748.824 Stomach Rupture
-	C06.405.748.895 Stomach Volvulus
-	C06.405.748.947 Zollinger-Ellison Syndrome
-	C06.405.874 Tuberculosis, Gastrointestinal
-	C06.405.937 Visceral Prolapse
-	C06.552 Liver Diseases
-	C06.552.074 alpha 1-Antitrypsin Deficiency
-	C06.552.150 Cholestasis, Intrahepatic
-	C06.552.150.125 Alagille Syndrome
-	C06.552.150.250 Liver Cirrhosis, Biliary
-	C06.552.195 Chemical and Drug Induced Liver Injury
-	C06.552.195 Drug-Induced Liver Injury
-	C06.552.195.200 Drug-Induced Liver Injury, Chronic
-	C06.552.241 Fatty Liver
-	C06.552.241.390 Fatty Liver, Alcoholic
-	C06.552.241.519 Non-alcoholic Fatty Liver Disease
-	C06.552.241.649 Reye Syndrome
-	C06.552.270 Focal Nodular Hyperplasia
-	C06.552.308 Hepatic Insufficiency
-	C06.552.308.500 Liver Failure
-	C06.552.308.500.177 End Stage Liver Disease
-	C06.552.308.500.356 Hepatic Encephalopathy
-	C06.552.308.500.750 Liver Failure, Acute
-	C06.552.308.500.750.249 Acute-On-Chronic Liver Failure
-	C06.552.308.500.750.500 Massive Hepatic Necrosis
-	C06.552.347 Budd-Chiari Syndrome
-	C06.552.360 Hepatic Veno-Occlusive Disease
-	C06.552.380 Hepatitis

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C06.552.380.290 Hepatitis, Alcoholic
-	C06.552.380.315 Hepatitis, Animal
-	C06.552.380.315.430 Hepatitis, Viral, Animal
-	C06.552.380.315.430.440 Hepatitis, Infectious Canine
-	C06.552.380.315.430.812 Rift Valley Fever
-	C06.552.380.350 Hepatitis, Chronic
-	C06.552.380.350.050 Hepatitis, Autoimmune
-	C06.552.380.350.100 Hepatitis B, Chronic
-	C06.552.380.350.120 Hepatitis C, Chronic
-	C06.552.380.350.220 Hepatitis D, Chronic
-	C06.552.380.705 Hepatitis, Viral, Human
-	C06.552.380.705.422 Hepatitis A
-	C06.552.380.705.437 Hepatitis B
-	C06.552.380.705.437.100 Hepatitis B, Chronic
-	C06.552.380.705.440 Hepatitis C
-	C06.552.380.705.440.120 Hepatitis C, Chronic
-	C06.552.380.705.450 Hepatitis D
-	C06.552.380.705.450.100 Hepatitis D, Chronic
-	C06.552.380.705.470 Hepatitis E
-	C06.552.413 Hepatolenticular Degeneration
-	C06.552.416 Hepatomegaly
-	C06.552.455 Hepatopulmonary Syndrome
-	C06.552.465 Hepatorenal Syndrome
-	C06.552.494 Hypertension, Portal
-	C06.552.494.414 Esophageal and Gastric Varices
-	C06.552.597 Liver Abscess
-	C06.552.597.517 Liver Abscess, Amebic
-	C06.552.597.758 Liver Abscess, Pyogenic
-	C06.552.630 Liver Cirrhosis
-	C06.552.630.380 Liver Cirrhosis, Alcoholic
-	C06.552.630.400 Liver Cirrhosis, Biliary
-	C06.552.630.467 Liver Cirrhosis, Experimental
-	C06.552.645 Liver Diseases, Alcoholic
-	C06.552.645.390 Fatty Liver, Alcoholic
-	C06.552.645.490 Hepatitis, Alcoholic
-	C06.552.645.590 Liver Cirrhosis, Alcoholic

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C06.552.664 Liver Diseases, Parasitic
-	C06.552.664.272 Echinococcosis, Hepatic
-	C06.552.664.424 Fascioliasis
-	C06.552.664.642 Liver Abscess, Amebic
-	C06.552.697 Liver Neoplasms
-	C06.552.697.040 Adenoma, Liver Cell
-	C06.552.697.160 Carcinoma, Hepatocellular
-	C06.552.697.580 Liver Neoplasms, Experimental
-	C06.552.802 Peliosis Hepatis
-	C06.552.830 Porphyrrias, Hepatic
-	C06.552.830.074 Coproporphyrria, Hereditary
-	C06.552.830.150 Porphyrria, Acute Intermittent
-	C06.552.830.250 Porphyrria Cutanea Tarda
-	C06.552.830.437 Porphyrria, Hepatoerythropoietic
-	C06.552.830.625 Porphyrria, Variegate
-	C06.552.830.812 Protoporphyrria, Erythropoietic
-	C06.552.933 Tuberculosis, Hepatic
-	C06.552.970 Zellweger Syndrome
-	C06.689 Pancreatic Diseases
-	C06.689.150 Congenital Hyperinsulinism
-	C06.689.150.500 Nesidioblastosis
-	C06.689.202 Cystic Fibrosis
-	C06.689.276 Exocrine Pancreatic Insufficiency
-	C06.689.500 Pancreatic Cyst
-	C06.689.500.692 Pancreatic Pseudocyst
-	C06.689.583 Pancreatic Fistula
-	C06.689.667 Pancreatic Neoplasms
-	C06.689.667.249 Adenoma, Islet Cell
-	C06.689.667.249.500 Insulinoma
-	C06.689.667.500 Carcinoma, Islet Cell
-	C06.689.667.500.124 Gastrinoma
-	C06.689.667.500.249 Glucagonoma
-	C06.689.667.500.500 Somatostatinoma
-	C06.689.667.500.750 Vipoma
-	C06.689.667.625 Carcinoma, Pancreatic Ductal
-	C06.689.750 Pancreatitis

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	C06.689.750.650	Pancreatitis, Acute Necrotizing
-	C06.689.750.660	Pancreatitis, Alcoholic
-	C06.689.750.830	Pancreatitis, Chronic
-	C06.689.750.860	Pancreatitis, Graft
-	C06.844	Peritoneal Diseases
-	C06.844.200	Chylous Ascites
-	C06.844.400	Hemoperitoneum
-	C06.844.460	Mesenteric Ischemia
-	C06.844.520	Mesenteric Lymphadenitis
-	C06.844.550	Mesenteric Vascular Occlusion
-	C06.844.600	Panniculitis, Peritoneal
-	C06.844.610	Peritoneal Fibrosis
-	C06.844.620	Peritoneal Neoplasms
-	C06.844.620.500	Mesenteric Cyst
-	C06.844.640	Peritonitis
-	C06.844.640.249	Peritonitis, Tuberculous
-	C06.844.640.500	Subphrenic Abscess
-	C06.844.670	Pneumoperitoneum
-	C07	Stomatognathic Diseases
New Heading	<b>C07.160</b>	<b>Ankyloglossia</b>
-	C07.320	Jaw Diseases
-	C07.320.086	Bisphosphonate-Associated Osteonecrosis of the Jaw
-	C07.320.173	Cherubism
-	C07.320.391	Granuloma, Giant Cell
-	C07.320.440	Jaw Abnormalities
-	C07.320.440.185	Cleft Palate
-	C07.320.440.457	Micrognathism
-	C07.320.440.606	Pierre Robin Syndrome
-	C07.320.440.655	Prognathism
-	C07.320.440.827	Retrognathia
-	C07.320.450	Jaw Cysts
-	C07.320.450.640	Nonodontogenic Cysts
-	C07.320.450.670	Odontogenic Cysts
-	C07.320.450.670.130	Basal Cell Nevus Syndrome
-	C07.320.450.670.275	Dentigerous Cyst



## MeSH Tree Changes for 2017

Type	Tree - heading
-	C07.320.450.670.495 Odontogenic Cyst, Calcifying
-	C07.320.450.670.513 Periodontal Cyst
-	C07.320.450.670.513.811 Radicular Cyst
-	C07.320.480 Jaw, Edentulous
-	C07.320.480.450 Jaw, Edentulous, Partially
-	C07.320.515 Jaw Neoplasms
-	C07.320.515.583 Mandibular Neoplasms
-	C07.320.515.601 Maxillary Neoplasms
-	C07.320.515.692 Palatal Neoplasms
-	C07.320.610 Mandibular Diseases
-	C07.320.610.291 Craniomandibular Disorders
-	C07.320.610.291.897 Temporomandibular Joint Disorders
-	C07.320.610.291.897.897 Temporomandibular Joint Dysfunction Syndrome
-	C07.320.610.583 Mandibular Neoplasms
-	C07.320.610.655 Prognathism
-	C07.320.610.827 Retrognathia
-	C07.320.660 Maxillary Diseases
-	C07.320.660.601 Maxillary Neoplasms
-	C07.320.830 Periapical Diseases
-	C07.320.830.700 Periapical Periodontitis
-	C07.320.830.700.700 Periapical Abscess
-	C07.320.830.700.740 Periapical Granuloma
-	C07.320.830.820 Radicular Cyst
-	C07.465 Mouth Diseases
-	C07.465.075 Behcet Syndrome
-	C07.465.094 Bell Palsy
-	C07.465.114 Burning Mouth Syndrome
-	C07.465.130 Candidiasis, Oral
-	C07.465.227 Dry Socket
-	C07.465.284 Facial Hemiatrophy
-	C07.465.299 Facial Nerve Diseases
-	C07.465.299.250 Bell Palsy
-	C07.465.299.375 Facial Hemiatrophy
-	C07.465.299.500 Facial Nerve Injuries
-	C07.465.299.625 Facial Neuralgia
-	C07.465.299.625.500 Trigeminal Nerve Diseases

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C07.465.299.625.500.349 Trigeminal Nerve Injuries
-	C07.465.299.625.500.349.500 Lingual Nerve Injuries
-	C07.465.299.625.500.700 Trigeminal Neuralgia
-	C07.465.299.750 Herpes Zoster Oticus
-	C07.465.299.800 Melkersson-Rosenthal Syndrome
-	C07.465.299.825 Mobius Syndrome
-	C07.465.327 Facial Paralysis
-	C07.465.342 Focal Epithelial Hyperplasia
-	C07.465.353 Granulomatosis, Orofacial
-	C07.465.364 Hemifacial Spasm
-	C07.465.385 Leukoedema, Oral
-	C07.465.397 Lichen Planus, Oral
-	C07.465.409 Lip Diseases
-	C07.465.409.215 Cheilitis
-	C07.465.409.225 Cleft Lip
-	C07.465.409.466 Herpes Labialis
-	C07.465.409.640 Lip Neoplasms
-	C07.465.433 Ludwig's Angina
-	C07.465.466 Melkersson-Rosenthal Syndrome
-	C07.465.525 Mouth Abnormalities
-	C07.465.525.164 Cleft Lip
-	C07.465.525.185 Cleft Palate
-	C07.465.525.304 Fibromatosis, Gingival
-	C07.465.525.480 Macrostomia
-	C07.465.525.520 Microstomia
-	C07.465.525.955 Velopharyngeal Insufficiency
-	C07.465.550 Mouth, Edentulous
-	C07.465.550.425 Jaw, Edentulous
-	C07.465.550.425.450 Jaw, Edentulous, Partially
-	C07.465.565 Mouth Neoplasms
-	C07.465.565.402 Gingival Neoplasms
-	C07.465.565.545 Leukoplakia, Oral
-	C07.465.565.545.500 Leukoplakia, Hairy
-	C07.465.565.550 Lip Neoplasms
-	C07.465.565.692 Palatal Neoplasms
-	C07.465.565.824 Salivary Gland Neoplasms

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C07.465.565.824.695 Parotid Neoplasms
-	C07.465.565.824.882 Sublingual Gland Neoplasms
-	C07.465.565.824.885 Submandibular Gland Neoplasms
-	C07.465.565.925 Tongue Neoplasms
-	C07.465.584 Mucositis
-	C07.465.604 Noma
-	C07.465.614 Oral Fistula
-	C07.465.614.187 Dental Fistula
-	C07.465.614.421 Oroantral Fistula
-	C07.465.614.655 Salivary Gland Fistula
-	C07.465.625 Oral Hemorrhage
-	C07.465.625.446 Gingival Hemorrhage
-	C07.465.634 Oral Manifestations
-	C07.465.654 Oral Submucous Fibrosis
-	C07.465.672 Oral Ulcer
-	C07.465.714 Periodontal Diseases
-	C07.465.714.204 Furcation Defects
-	C07.465.714.258 Gingival Diseases
-	C07.465.714.258.250 Gingival Hemorrhage
-	C07.465.714.258.409 Gingival Neoplasms
-	C07.465.714.258.428 Gingival Overgrowth
-	C07.465.714.258.428.200 Fibromatosis, Gingival
-	C07.465.714.258.428.250 Gingival Hyperplasia
-	C07.465.714.258.428.260 Gingival Hypertrophy
-	C07.465.714.258.447 Gingival Recession
-	C07.465.714.258.480 Gingivitis
-	C07.465.714.258.480.360 Gingival Pocket
-	C07.465.714.258.480.446 Gingivitis, Necrotizing Ulcerative
-	C07.465.714.258.557 Granuloma, Giant Cell
-	C07.465.714.258.771 Pericoronitis
-	C07.465.714.282 Peri-Implantitis
-	C07.465.714.306 Periapical Diseases
-	C07.465.714.306.700 Periapical Periodontitis
-	C07.465.714.306.700.700 Periapical Abscess
-	C07.465.714.306.700.740 Periapical Granuloma
-	C07.465.714.306.820 Radicular Cyst

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C07.465.714.354                      Periodontal Atrophy
-	C07.465.714.354.500                      Alveolar Bone Loss
-	C07.465.714.354.625                      Gingival Recession
-	C07.465.714.354.750                      Periodontal Attachment Loss
-	C07.465.714.470                      Periodontal Cyst
-	C07.465.714.533                      Periodontitis
-	C07.465.714.533.161                      Aggressive Periodontitis
-	C07.465.714.533.324                      Chronic Periodontitis
-	C07.465.714.533.487                      Periapical Periodontitis
-	C07.465.714.533.487.700                      Periapical Abscess
-	C07.465.714.533.487.740                      Periapical Granuloma
-	C07.465.714.533.650                      Periodontal Abscess
-	C07.465.714.533.750                      Periodontal Pocket
-	C07.465.714.804                      Tooth Loss
-	C07.465.714.836                      Tooth Migration
-	C07.465.714.836.535                      Mesial Movement of Teeth
-	C07.465.714.898                      Tooth Mobility
-	C07.465.780                      Ranula
-	C07.465.815                      Salivary Gland Diseases
-	C07.465.815.355                      Mikulicz' Disease
-	C07.465.815.470                      Parotid Diseases
-	C07.465.815.470.770                      Parotid Neoplasms
-	C07.465.815.470.800                      Parotitis
-	C07.465.815.470.800.630                      Mumps
-	C07.465.815.497                      Salivary Calculi
-	C07.465.815.497.325                      Salivary Duct Calculi
-	C07.465.815.497.500                      Salivary Gland Calculi
-	C07.465.815.655                      Salivary Gland Fistula
-	C07.465.815.718                      Salivary Gland Neoplasms
-	C07.465.815.718.589                      Parotid Neoplasms
-	C07.465.815.718.870                      Sublingual Gland Neoplasms
-	C07.465.815.718.885                      Submandibular Gland Neoplasms
-	C07.465.815.793                      Sialadenitis
-	C07.465.815.793.500                      Parotitis
-	C07.465.815.802                      Sialometaplasia, Necrotizing
-	C07.465.815.815                      Sialorrhea

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C07.465.815.882 Submandibular Gland Diseases
-	C07.465.815.882.500 Submandibular Gland Neoplasms
-	C07.465.815.929 Xerostomia
-	C07.465.815.929.669 Sjogren's Syndrome
-	C07.465.864 Stomatitis
-	C07.465.864.500 Stevens-Johnson Syndrome
-	C07.465.864.750 Stomatitis, Aphthous
-	C07.465.864.875 Stomatitis, Denture
-	C07.465.864.937 Stomatitis, Herpetic
-	C07.465.864.968 Vesicular Stomatitis
-	C07.465.910 Tongue Diseases
-	C07.465.910.280 Glossalgia
-	C07.465.910.363 Glossitis
-	C07.465.910.363.447 Glossitis, Benign Migratory
-	C07.465.910.411 Glossoptosis
-	C07.465.910.460 Macroglossia
-	C07.465.910.708 Tongue, Fissured
-	C07.465.910.791 Tongue, Hairy
-	C07.465.910.874 Tongue Neoplasms
-	C07.465.943 Tuberculosis, Oral
-	C07.550 Pharyngeal Diseases
-	C07.550.174 Lemierre Syndrome
-	C07.550.350 Nasopharyngeal Diseases
-	C07.550.350.650 Nasopharyngeal Neoplasms
-	C07.550.350.700 Nasopharyngitis
-	C07.550.745 Pharyngeal Neoplasms
-	C07.550.745.436 Hypopharyngeal Neoplasms
-	C07.550.745.650 Nasopharyngeal Neoplasms
-	C07.550.745.671 Oropharyngeal Neoplasms
-	C07.550.745.671.800 Tonsillar Neoplasms
-	C07.550.781 Pharyngitis
-	C07.550.781.500 Nasopharyngitis
-	C07.550.781.625 Retropharyngeal Abscess
-	C07.550.781.750 Tonsillitis
-	C07.550.781.750.500 Peritonsillar Abscess
-	C07.550.966 Velopharyngeal Insufficiency

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C07.650 Stomatognathic System Abnormalities
-	C07.650.500 Maxillofacial Abnormalities
-	C07.650.500.229 Dentofacial Deformities
-	C07.650.500.460 Jaw Abnormalities
-	C07.650.500.460.185 Cleft Palate
-	C07.650.500.460.457 Micrognathism
-	C07.650.500.460.606 Pierre Robin Syndrome
-	C07.650.500.460.655 Prognathism
-	C07.650.500.460.827 Retrognathia
-	C07.650.525 Mouth Abnormalities
-	C07.650.525.164 Cleft Lip
-	C07.650.525.185 Cleft Palate
-	C07.650.525.304 Fibromatosis, Gingival
-	C07.650.525.480 Macrostomia
-	C07.650.525.520 Microstomia
-	C07.650.525.955 Velopharyngeal Insufficiency
-	C07.650.800 Tooth Abnormalities
-	C07.650.800.100 Anodontia
-	C07.650.800.250 Dens in Dente
-	C07.650.800.255 Dental Enamel Hypoplasia
-	C07.650.800.255.500 Amelogenesis Imperfecta
-	C07.650.800.260 Dentin Dysplasia
-	C07.650.800.270 Dentinogenesis Imperfecta
-	C07.650.800.320 Diastema
-	C07.650.800.370 Fused Teeth
-	C07.650.800.600 Odontodysplasia
-	C07.650.800.850 Tooth, Supernumerary
-	C07.678 Temporomandibular Joint Disorders
-	C07.678.949 Temporomandibular Joint Dysfunction Syndrome
-	C07.793 Tooth Diseases
-	C07.793.099 Bruxism
-	C07.793.099.500 Sleep Bruxism
-	C07.793.208 Dental Deposits
-	C07.793.208.250 Dental Calculus
-	C07.793.208.377 Dental Plaque
-	C07.793.208.688 Smear Layer

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C07.793.221 Dental Leakage
-	C07.793.237 Dental Pulp Diseases
-	C07.793.237.252 Dental Pulp Calcification
-	C07.793.237.283 Dental Pulp Exposure
-	C07.793.237.315 Dental Pulp Necrosis
-	C07.793.237.378 Dentin, Secondary
-	C07.793.237.820 Pulpitis
-	C07.793.237.910 Tooth, Nonvital
-	C07.793.266 Dentin Sensitivity
-	C07.793.330 Fluorosis, Dental
-	C07.793.348 Focal Infection, Dental
-	C07.793.422 Hypercementosis
-	C07.793.494 Malocclusion
-	C07.793.494.293 Dental Occlusion, Traumatic
-	C07.793.494.610 Malocclusion, Angle Class I
-	C07.793.494.630 Malocclusion, Angle Class II
-	C07.793.494.630.500 Overbite
-	C07.793.494.650 Malocclusion, Angle Class III
-	C07.793.494.825 Open Bite
-	C07.793.597 Mouth, Edentulous
-	C07.793.597.425 Jaw, Edentulous
-	C07.793.597.425.450 Jaw, Edentulous, Partially
-	C07.793.700 Tooth Abnormalities
-	C07.793.700.100 Anodontia
-	C07.793.700.250 Dens in Dente
-	C07.793.700.255 Dental Enamel Hypoplasia
-	C07.793.700.255.500 Amelogenesis Imperfecta
-	C07.793.700.260 Dentin Dysplasia
-	C07.793.700.270 Dentinogenesis Imperfecta
-	C07.793.700.320 Diastema
-	C07.793.700.370 Fused Teeth
-	C07.793.700.600 Odontodysplasia
-	C07.793.700.850 Tooth, Supernumerary
-	C07.793.710 Tooth Ankylosis
-	C07.793.720 Tooth Demineralization
-	C07.793.720.210 Dental Caries

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C07.793.720.210.220                      Dental Fissures
-	C07.793.720.210.650                      Root Caries
-	C07.793.735                                  Tooth Discoloration
-	C07.793.790                                  Tooth Eruption, Ectopic
-	C07.793.818                                  Tooth Wear
-	C07.793.818.124                            Tooth Abrasion
-	C07.793.818.249                            Tooth Attrition
-	C07.793.818.500                            Tooth Erosion
-	C07.793.846                                  Tooth, Impacted
-	C07.793.850                                  Tooth Injuries
-	C07.793.850.725                            Tooth Avulsion
-	C07.793.850.750                            Tooth Fractures
-	C07.793.850.750.300                      Cracked Tooth Syndrome
-	C07.793.870                                  Tooth Loss
-	C07.793.901                                  Tooth Resorption
-	C07.793.901.653                            Root Resorption
-	C07.793.915                                  Tooth, Unerupted
-	C07.793.929                                  Toothache
-	C08    Respiratory Tract Diseases
-	C08.127                                        Bronchial Diseases
-	C08.127.108                                  Asthma
-	C08.127.108.054                            Asthma, Aspirin-Induced
-	C08.127.108.110                            Asthma, Exercise-Induced
-	C08.127.108.495                            Asthma, Occupational
-	C08.127.108.880                            Status Asthmaticus
-	C08.127.196                                  Bronchial Fistula
-	C08.127.210                                  Bronchial Hyperreactivity
-	C08.127.265                                  Bronchial Neoplasms
-	C08.127.321                                  Bronchial Spasm
-	C08.127.384                                  Bronchiectasis
-	C08.127.384.500                            Kartagener Syndrome
-	C08.127.446                                  Bronchitis
-	C08.127.446.135                            Bronchiolitis
-	C08.127.446.135.140                      Bronchiolitis Obliterans
-	C08.127.446.135.140.200                Cryptogenic Organizing Pneumonia
-	C08.127.446.135.321                      Bronchiolitis, Viral



## MeSH Tree Changes for 2017

Type	Tree - heading
-	C08.127.446.567      Bronchitis, Chronic
-	C08.127.480      Bronchogenic Cyst
-	C08.127.509      Bronchopneumonia
-	C08.127.719      Tracheobronchomalacia
-	C08.127.719.500      Bronchomalacia
-	C08.127.930      Tracheobronchomegaly
-	C08.200      Ciliary Motility Disorders
-	C08.200.531      Kartagener Syndrome
-	C08.280      Granuloma, Respiratory Tract
-	C08.280.400      Granuloma, Laryngeal
-	C08.360      Laryngeal Diseases
-	C08.360.232      Granuloma, Laryngeal
-	C08.360.313      Laryngeal Edema
-	C08.360.369      Laryngeal Neoplasms
-	C08.360.424      Laryngeal Nerve Injuries
-	C08.360.424.500      Recurrent Laryngeal Nerve Injuries
-	C08.360.535      Laryngitis
-	C08.360.535.365      Croup
-	C08.360.549      Laryngocele
-	C08.360.563      Laryngomalacia
-	C08.360.577      Laryngopharyngeal Reflux
-	C08.360.591      Laryngostenosis
-	C08.360.840      Supraglottitis
-	C08.360.860      Tuberculosis, Laryngeal
-	C08.360.895      Vocal Cord Dysfunction
-	C08.360.895.500      Laryngismus
-	C08.360.931      Vocal Cord Paralysis
-	C08.360.940      Voice Disorders
-	C08.360.940.160      Aphonia
-	C08.360.940.325      Dysphonia
-	C08.360.940.490      Hoarseness
-	C08.381      Lung Diseases
-	C08.381.074      Acute Chest Syndrome
-	C08.381.112      alpha 1-Antitrypsin Deficiency
-	C08.381.150      Cystic Adenomatoid Malformation of Lung, Congenital
-	C08.381.187      Cystic Fibrosis

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C08.381.348 Hemoptysis
-	C08.381.385 Hepatopulmonary Syndrome
-	C08.381.423 Hypertension, Pulmonary
-	C08.381.423.694 Persistent Fetal Circulation Syndrome
-	C08.381.423.847 Familial Primary Pulmonary Hypertension
-	C08.381.450 Lung Abscess
-	C08.381.472 Lung Diseases, Fungal
-	C08.381.472.350 Blastomycosis
-	C08.381.472.700 Pneumonia, Pneumocystis
-	C08.381.472.850 Pulmonary Aspergillosis
-	C08.381.472.850.500 Aspergillosis, Allergic Bronchopulmonary
-	C08.381.472.850.750 Invasive Pulmonary Aspergillosis
-	C08.381.483 Lung Diseases, Interstitial
-	C08.381.483.125 Alveolitis, Extrinsic Allergic
-	C08.381.483.125.125 Bird Fancier's Lung
-	C08.381.483.125.365 Farmer's Lung
-	C08.381.483.125.682 Silo Filler's Disease
-	C08.381.483.125.841 Trichosporonosis
-	C08.381.483.156 Anti-Glomerular Basement Membrane Disease
-	C08.381.483.375 Histiocytosis, Langerhans-Cell
-	C08.381.483.375.500 Eosinophilic Granuloma
-	C08.381.483.487 Idiopathic Interstitial Pneumonias
-	C08.381.483.487.249 Cryptogenic Organizing Pneumonia
-	C08.381.483.487.500 Idiopathic Pulmonary Fibrosis
-	C08.381.483.581 Pneumoconiosis
-	C08.381.483.581.062 Anthracosis
-	C08.381.483.581.062.500 Anthracosilicosis
-	C08.381.483.581.125 Asbestosis
-	C08.381.483.581.225 Berylliosis
-	C08.381.483.581.275 Byssinosis
-	C08.381.483.581.300 Caplan Syndrome
-	C08.381.483.581.750 Siderosis
-	C08.381.483.581.760 Silicosis
-	C08.381.483.581.760.125 Anthracosilicosis
-	C08.381.483.581.760.750 Silicotuberculosis
-	C08.381.483.675 Radiation Pneumonitis

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C08.381.483.725 Sarcoidosis, Pulmonary
-	C08.381.483.950 Granulomatosis with Polyangiitis
-	C08.381.495 Lung Diseases, Obstructive
-	C08.381.495.108 Asthma
-	C08.381.495.146 Bronchitis
-	C08.381.495.146.135 Bronchiolitis
-	C08.381.495.146.135.140 Bronchiolitis Obliterans
-	C08.381.495.146.135.140.200 Cryptogenic Organizing Pneumonia
-	C08.381.495.146.135.321 Bronchiolitis, Viral
-	C08.381.495.146.567 Bronchitis, Chronic
-	C08.381.495.389 Pulmonary Disease, Chronic Obstructive
-	C08.381.495.389.500 Bronchitis, Chronic
-	C08.381.495.389.750 Pulmonary Emphysema
-	C08.381.517 Lung Diseases, Parasitic
-	C08.381.517.314 Echinococcosis, Pulmonary
-	C08.381.520 Lung Injury
-	C08.381.520.500 Acute Lung Injury
-	C08.381.520.687 Meconium Aspiration Syndrome
-	C08.381.520.702 Pneumoconiosis
-	C08.381.520.702.062 Anthracosis
-	C08.381.520.702.062.500 Anthracosilicosis
-	C08.381.520.702.125 Asbestosis
-	C08.381.520.702.225 Berylliosis
-	C08.381.520.702.275 Byssinosis
-	C08.381.520.702.300 Caplan Syndrome
-	C08.381.520.702.750 Siderosis
-	C08.381.520.702.760 Silicosis
-	C08.381.520.702.760.125 Anthracosilicosis
-	C08.381.520.702.760.750 Silicotuberculosis
-	C08.381.520.734 Radiation Pneumonitis
-	C08.381.520.750 Ventilator-Induced Lung Injury
-	C08.381.520.750.500 Bronchopulmonary Dysplasia
-	C08.381.520.750.750 Pneumonia, Ventilator-Associated
-	C08.381.540 Lung Neoplasms
-	C08.381.540.140 Carcinoma, Bronchogenic
-	C08.381.540.140.500 Carcinoma, Non-Small-Cell Lung

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C08.381.540.140.750 Small Cell Lung Carcinoma
-	C08.381.540.148 Multiple Pulmonary Nodules
-	C08.381.540.734 Pancoast Syndrome
-	C08.381.540.867 Pulmonary Sclerosing Hemangioma
-	C08.381.570 Lung, Hyperlucent
-	C08.381.600 Plasma Cell Granuloma, Pulmonary
-	C08.381.677 Pneumonia
-	C08.381.677.127 Bronchopneumonia
-	C08.381.677.473 Pleuropneumonia
-	C08.381.677.529 Pneumonia, Aspiration
-	C08.381.677.529.612 Pneumonia, Lipid
-	C08.381.677.540 Pneumonia, Bacterial
-	C08.381.677.540.249 Chlamydial Pneumonia
-	C08.381.677.540.500 Pneumonia, Mycoplasma
-	C08.381.677.540.530 Pneumonia of Calves, Enzootic
-	C08.381.677.540.540 Pneumonia of Swine, Mycoplasmal
-	C08.381.677.540.550 Pneumonia, Pneumococcal
-	C08.381.677.540.600 Pneumonia, Rickettsial
-	C08.381.677.540.620 Pneumonia, Staphylococcal
New Heading	<b>C08.381.677.608 Pneumonia, Necrotizing</b>
-	C08.381.677.675 Pneumonia, Pneumocystis
-	C08.381.677.800 Pneumonia, Ventilator-Associated
-	C08.381.677.807 Pneumonia, Viral
-	C08.381.719 Pulmonary Alveolar Proteinosis
-	C08.381.730 Pulmonary Atelectasis
-	C08.381.730.542 Middle Lobe Syndrome
-	C08.381.742 Pulmonary Edema
-	C08.381.746 Pulmonary Embolism
-	C08.381.746.500 Pulmonary Infarction
-	C08.381.750 Pulmonary Eosinophilia
-	C08.381.765 Pulmonary Fibrosis
-	C08.381.765.500 Idiopathic Pulmonary Fibrosis
-	C08.381.780 Pulmonary Veno-Occlusive Disease
-	C08.381.840 Respiratory Distress Syndrome, Adult
-	C08.381.842 Respiratory Distress Syndrome, Newborn

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C08.381.842.475 Hyaline Membrane Disease
-	C08.381.842.737 Transient Tachypnea of the Newborn
-	C08.381.844 Scimitar Syndrome
-	C08.381.884 Solitary Pulmonary Nodule
-	C08.381.922 Tuberculosis, Pulmonary
-	C08.381.922.669 Silicotuberculosis
-	C08.460 Nose Diseases
-	C08.460.171 Choanal Atresia
-	C08.460.261 Epistaxis
-	C08.460.393 Granuloma, Lethal Midline
-	C08.460.525 Nasal Obstruction
-	C08.460.572 Nasal Polyps
-	C08.460.595 Nasal Septal Perforation
-	C08.460.619 Nose Deformities, Acquired
-	C08.460.669 Nose Neoplasms
-	C08.460.669.693 Paranasal Sinus Neoplasms
-	C08.460.669.693.575 Maxillary Sinus Neoplasms
-	C08.460.692 Paranasal Sinus Diseases
-	C08.460.692.503 Paranasal Sinus Neoplasms
-	C08.460.692.503.503 Maxillary Sinus Neoplasms
-	C08.460.692.752 Sinusitis
-	C08.460.692.752.267 Ethmoid Sinusitis
-	C08.460.692.752.387 Frontal Sinusitis
-	C08.460.692.752.578 Maxillary Sinusitis
-	C08.460.692.752.827 Sphenoid Sinusitis
-	C08.460.799 Rhinitis
-	C08.460.799.315 Rhinitis, Allergic
-	C08.460.799.315.500 Rhinitis, Allergic, Perennial
-	C08.460.799.315.750 Rhinitis, Allergic, Seasonal
-	C08.460.799.649 Rhinitis, Atrophic
-	C08.460.799.910 Rhinitis, Vasomotor
-	C08.460.850 Rhinoscleroma
-	C08.528 Pleural Diseases
-	C08.528.142 Chylothorax
-	C08.528.240 Empyema, Pleural
-	C08.528.240.320 Empyema, Tuberculous

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	C08.528.338	Hemopneumothorax
-	C08.528.380	Hemothorax
-	C08.528.434	Hydropneumothorax
-	C08.528.476	Hydrothorax
-	C08.528.652	Pleural Effusion
-	C08.528.652.700	Pleural Effusion, Malignant
-	C08.528.694	Pleural Neoplasms
-	C08.528.694.700	Pleural Effusion, Malignant
-	C08.528.735	Pleurisy
-	C08.528.735.473	Pleuropneumonia
-	C08.528.778	Pneumothorax
-	C08.528.928	Tuberculosis, Pleural
-	C08.528.928.405	Empyema, Tuberculous
-	C08.618	Respiration Disorders
-	C08.618.009	Acute Chest Syndrome
-	C08.618.020	Altitude Sickness
-	C08.618.085	Apnea
-	C08.618.085.852	Sleep Apnea Syndromes
-	C08.618.085.852.800	Sleep Apnea, Central
-	C08.618.085.852.850	Sleep Apnea, Obstructive
-	C08.618.085.852.850.500	Obesity Hypoventilation Syndrome
-	C08.618.182	Cheyne-Stokes Respiration
-	C08.618.248	Cough
-	C08.618.326	Dyspnea
-	C08.618.326.396	Dyspnea, Paroxysmal
-	C08.618.490	Hoarseness
-	C08.618.490.500	Lipoid Proteinosis of Urbach and Wiethe
-	C08.618.501	Hyperventilation
-	C08.618.501.271	Alkalosis, Respiratory
-	C08.618.580	Meconium Aspiration Syndrome
-	C08.618.659	Mouth Breathing
-	C08.618.749	Respiratory Aspiration
-	C08.618.749.500	Respiratory Aspiration of Gastric Contents
-	C08.618.840	Respiratory Distress Syndrome, Adult
-	C08.618.842	Respiratory Distress Syndrome, Newborn
-	C08.618.842.475	Hyaline Membrane Disease

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C08.618.842.737      Transient Tachypnea of the Newborn
-	C08.618.846      Respiratory Insufficiency
-	C08.618.846.093      Acidosis, Respiratory
-	C08.618.846.185      Airway Obstruction
-	C08.618.846.185.525      Nasal Obstruction
-	C08.618.846.414      Granuloma, Laryngeal
-	C08.618.846.450      Hantavirus Pulmonary Syndrome
-	C08.618.846.565      Hypoventilation
-	C08.618.846.565.500      Obesity Hypoventilation Syndrome
-	C08.618.846.600      Infantile Apparent Life-Threatening Event
-	C08.618.846.688      Positive-Pressure Respiration, Intrinsic
-	C08.618.846.812      Respiratory Paralysis
-	C08.618.923      Sarcoglycanopathies
-	C08.618.961      Tachypnea
-	C08.618.961.500      Transient Tachypnea of the Newborn
-	C08.618.980      Vocal Cord Dysfunction
-	C08.618.980.500      Laryngismus
-	C08.674      Respiratory Hypersensitivity
-	C08.674.055      Alveolitis, Extrinsic Allergic
-	C08.674.055.125      Bird Fancier's Lung
-	C08.674.055.365      Farmer's Lung
-	C08.674.055.682      Trichosporonosis
-	C08.674.060      Aspergillosis, Allergic Bronchopulmonary
-	C08.674.095      Asthma
-	C08.674.095.054      Asthma, Aspirin-Induced
-	C08.674.095.110      Asthma, Exercise-Induced
-	C08.674.095.495      Asthma, Occupational
-	C08.674.095.880      Status Asthmaticus
-	C08.674.453      Rhinitis, Allergic
-	C08.674.453.500      Rhinitis, Allergic, Perennial
-	C08.674.453.750      Rhinitis, Allergic, Seasonal
-	C08.695      Respiratory System Abnormalities
-	C08.695.195      Bronchogenic Cyst
-	C08.695.214      Bronchopulmonary Sequestration
-	C08.695.271      Choanal Atresia
-	C08.695.290      Cystic Adenomatoid Malformation of Lung, Congenital

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C08.695.501 Kartagener Syndrome
-	C08.695.520 Laryngocele
-	C08.695.815 Scimitar Syndrome
-	C08.695.830 Tracheobronchomegaly
-	C08.702 Respiratory Tract Fistula
-	C08.702.196 Bronchial Fistula
-	C08.702.750 Tracheoesophageal Fistula
-	C08.730 Respiratory Tract Infections
-	C08.730.085 Bovine Respiratory Disease Complex
-	C08.730.085.600 Pasteurellosis, Pneumonic
-	C08.730.085.660 Pneumonia, Atypical Interstitial, of Cattle
-	C08.730.085.675 Pneumonia of Calves, Enzootic
-	C08.730.099 Bronchitis
-	C08.730.099.135 Bronchiolitis
-	C08.730.099.135.321 Bronchiolitis, Viral
-	C08.730.099.567 Bronchitis, Chronic
-	C08.730.162 Common Cold
-	C08.730.265 Empyema, Pleural
-	C08.730.265.320 Empyema, Tuberculous
-	C08.730.310 Influenza, Human
-	C08.730.368 Laryngitis
-	C08.730.382 Legionellosis
-	C08.730.382.380 Legionnaires' Disease
-	C08.730.407 Lung Abscess
-	C08.730.435 Lung Diseases, Fungal
-	C08.730.435.090 Aspergillosis, Allergic Bronchopulmonary
-	C08.730.435.395 Blastomycosis
-	C08.730.435.700 Pneumonia, Pneumocystis
-	C08.730.450 Lung Diseases, Parasitic
-	C08.730.450.314 Echinococcosis, Pulmonary
-	C08.730.561 Pharyngitis
-	C08.730.561.500 Nasopharyngitis
-	C08.730.561.625 Retropharyngeal Abscess
-	C08.730.561.750 Tonsillitis
-	C08.730.561.750.500 Peritonsillar Abscess
-	C08.730.582 Pleurisy



## MeSH Tree Changes for 2017

Type	Tree - heading
-	C08.730.582.473 Pleuropneumonia
-	C08.730.610 Pneumonia
-	C08.730.610.127 Bronchopneumonia
-	C08.730.610.473 Pleuropneumonia
-	C08.730.610.529 Pneumonia, Aspiration
-	C08.730.610.529.612 Pneumonia, Lipid
-	C08.730.610.540 Pneumonia, Bacterial
-	C08.730.610.540.249 Chlamydial Pneumonia
-	C08.730.610.540.500 Pneumonia, Mycoplasma
-	C08.730.610.540.530 Pneumonia of Calves, Enzootic
-	C08.730.610.540.540 Pneumonia of Swine, Mycoplasmal
-	C08.730.610.540.550 Pneumonia, Pneumococcal
-	C08.730.610.540.600 Pneumonia, Rickettsial
-	C08.730.610.540.620 Pneumonia, Staphylococcal
New Heading	<b>C08.730.610.608 Pneumonia, Necrotizing</b>
-	C08.730.610.675 Pneumonia, Pneumocystis
-	C08.730.610.750 Pneumonia, Ventilator-Associated
-	C08.730.610.763 Pneumonia, Viral
-	C08.730.674 Rhinitis
-	C08.730.702 Rhinoscleroma
-	C08.730.730 Severe Acute Respiratory Syndrome
-	C08.730.749 Sinusitis
-	C08.730.749.267 Ethmoid Sinusitis
-	C08.730.749.387 Frontal Sinusitis
-	C08.730.749.578 Maxillary Sinusitis
-	C08.730.749.827 Sphenoid Sinusitis
-	C08.730.798 Supraglottitis
-	C08.730.798.200 Epiglottitis
-	C08.730.848 Tracheitis
-	C08.730.860 Tuberculosis, Laryngeal
-	C08.730.912 Tuberculosis, Pleural
-	C08.730.912.405 Empyema, Tuberculous
-	C08.730.939 Tuberculosis, Pulmonary
-	C08.730.939.669 Silicotuberculosis
-	C08.730.969 Whooping Cough

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C08.785                      Respiratory Tract Neoplasms
-	C08.785.481                  Laryngeal Neoplasms
-	C08.785.520                  Lung Neoplasms
-	C08.785.520.100              Bronchial Neoplasms
-	C08.785.520.100.220          Carcinoma, Bronchogenic
-	C08.785.520.100.220.500      Carcinoma, Non-Small-Cell Lung
-	C08.785.520.100.220.750      Small Cell Lung Carcinoma
-	C08.785.520.148              Multiple Pulmonary Nodules
-	C08.785.520.734              Pancoast Syndrome
-	C08.785.520.867              Pulmonary Sclerosing Hemangioma
-	C08.785.600                  Nose Neoplasms
-	C08.785.600.693              Paranasal Sinus Neoplasms
-	C08.785.600.693.575          Maxillary Sinus Neoplasms
-	C08.785.640                  Pleural Neoplasms
-	C08.785.640.700              Pleural Effusion, Malignant
-	C08.785.760                  Tracheal Neoplasms
-	C08.846                      Thoracic Diseases
-	C08.846.187                  Mediastinal Diseases
-	C08.846.187.145              Mediastinal Cyst
-	C08.846.187.290              Mediastinal Emphysema
-	C08.846.187.580              Mediastinal Neoplasms
-	C08.846.187.790              Mediastinitis
-	C08.907                      Tracheal Diseases
-	C08.907.563                  Tracheal Neoplasms
-	C08.907.663                  Tracheal Stenosis
-	C08.907.763                  Tracheitis
-	C08.907.796                  Tracheobronchomalacia
-	C08.907.796.500              Tracheomalacia
-	C08.907.830                  Tracheobronchomegaly
-	C08.907.863                  Tracheoesophageal Fistula
-	C09                          Otorhinolaryngologic Diseases
-	C09.150                      Ciliary Motility Disorders
-	C09.150.531                  Kartagener Syndrome
-	C09.218                      Ear Diseases
-	C09.218.200                  Cholesteatoma, Middle Ear
-	C09.218.235                  Congenital Microtia

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	C09.218.271	Ear Deformities, Acquired
-	C09.218.334	Ear Neoplasms
-	C09.218.350	Earache
-	C09.218.458	Hearing Disorders
-	C09.218.458.341	Hearing Loss
-	C09.218.458.341.186	Deafness
-	C09.218.458.341.186.500	Deaf-Blind Disorders
-	C09.218.458.341.186.500.500	Usher Syndromes
-	C09.218.458.341.186.500.750	Wolfram Syndrome
-	C09.218.458.341.374	Hearing Loss, Bilateral
-	C09.218.458.341.562	Hearing Loss, Conductive
-	C09.218.458.341.750	Hearing Loss, Functional
-	C09.218.458.341.812	Hearing Loss, High-Frequency
-	C09.218.458.341.849	Hearing Loss, Mixed Conductive-Sensorineural
-	C09.218.458.341.887	Hearing Loss, Sensorineural
-	C09.218.458.341.887.432	Hearing Loss, Central
-	C09.218.458.341.887.460	Hearing Loss, Noise-Induced
-	C09.218.458.341.887.772	Presbycusis
-	C09.218.458.341.887.886	Usher Syndromes
-	C09.218.458.341.900	Hearing Loss, Sudden
-	C09.218.458.341.950	Hearing Loss, Unilateral
-	C09.218.458.505	Hyperacusis
-	C09.218.458.670	Tinnitus
-	C09.218.513	Herpes Zoster Oticus
-	C09.218.568	Labyrinth Diseases
-	C09.218.568.120	Cochlear Diseases
-	C09.218.568.217	Endolymphatic Hydrops
-	C09.218.568.217.500	Meniere Disease
-	C09.218.568.558	Labyrinthitis
-	C09.218.568.900	Vestibular Diseases
New Heading	<b>C09.218.568.900.442</b>	<b>Bilateral Vestibulopathy</b>
-	C09.218.568.900.883	Vertigo
-	C09.218.568.900.883.500	Benign Paroxysmal Positional Vertigo
-	C09.218.636	Myringosclerosis
-	C09.218.705	Otitis

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C09.218.705.371 Labyrinthitis
-	C09.218.705.496 Otitis Externa
-	C09.218.705.663 Otitis Media
-	C09.218.705.663.652 Mastoiditis
-	C09.218.705.663.680 Otitis Media, Suppurative
-	C09.218.705.663.683 Otitis Media with Effusion
-	C09.218.705.663.841 Petrositis
-	C09.218.736 Otomycosis
-	C09.218.768 Otosclerosis
-	C09.218.807 Retrocochlear Diseases
-	C09.218.807.186 Auditory Diseases, Central
-	C09.218.807.186.094 Auditory Perceptual Disorders
-	C09.218.807.186.432 Hearing Loss, Central
-	C09.218.807.800 Vestibulocochlear Nerve Diseases
-	C09.218.807.800.675 Neuroma, Acoustic
-	C09.218.807.800.675.500 Neurofibromatosis 2
-	C09.218.807.800.837 Vestibular Neuritis
-	C09.218.807.800.918 Vestibulocochlear Nerve Injuries
-	C09.218.855 Susac Syndrome
-	C09.218.903 Tympanic Membrane Perforation
-	C09.400 Laryngeal Diseases
-	C09.400.232 Granuloma, Laryngeal
-	C09.400.313 Laryngeal Edema
-	C09.400.369 Laryngeal Neoplasms
-	C09.400.424 Laryngeal Nerve Injuries
-	C09.400.424.500 Recurrent Laryngeal Nerve Injuries
-	C09.400.535 Laryngitis
-	C09.400.535.365 Croup
-	C09.400.549 Laryngocele
-	C09.400.563 Laryngomalacia
-	C09.400.591 Laryngostenosis
-	C09.400.840 Supraglottitis
-	C09.400.860 Tuberculosis, Laryngeal
-	C09.400.895 Vocal Cord Dysfunction
-	C09.400.895.500 Laryngismus
-	C09.400.931 Vocal Cord Paralysis

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C09.400.940 Voice Disorders
-	C09.400.940.160 Aphonia
-	C09.400.940.325 Dysphonia
-	C09.400.940.490 Hoarseness
-	C09.603 Nose Diseases
-	C09.603.171 Choanal Atresia
-	C09.603.261 Epistaxis
-	C09.603.393 Granuloma, Lethal Midline
-	C09.603.525 Nasal Obstruction
-	C09.603.557 Nasal Polyps
-	C09.603.588 Nasal Septal Perforation
-	C09.603.619 Nose Deformities, Acquired
-	C09.603.669 Nose Neoplasms
-	C09.603.669.693 Paranasal Sinus Neoplasms
-	C09.603.669.693.575 Maxillary Sinus Neoplasms
-	C09.603.692 Paranasal Sinus Diseases
-	C09.603.692.503 Paranasal Sinus Neoplasms
-	C09.603.692.503.503 Maxillary Sinus Neoplasms
-	C09.603.692.752 Sinusitis
-	C09.603.692.752.267 Ethmoid Sinusitis
-	C09.603.692.752.387 Frontal Sinusitis
-	C09.603.692.752.578 Maxillary Sinusitis
-	C09.603.692.752.827 Sphenoid Sinusitis
-	C09.603.799 Rhinitis
-	C09.603.799.315 Rhinitis, Allergic
-	C09.603.799.315.500 Rhinitis, Allergic, Perennial
-	C09.603.799.315.750 Rhinitis, Allergic, Seasonal
-	C09.603.799.649 Rhinitis, Atrophic
-	C09.603.799.910 Rhinitis, Vasomotor
-	C09.603.850 Rhinoscleroma
-	C09.647 Otorhinolaryngologic Neoplasms
-	C09.647.312 Ear Neoplasms
-	C09.647.481 Laryngeal Neoplasms
-	C09.647.675 Neuroma, Acoustic
-	C09.647.675.500 Neurofibromatosis 2
-	C09.647.685 Nose Neoplasms

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C09.647.685.693                      Paranasal Sinus Neoplasms
-	C09.647.685.693.575                      Maxillary Sinus Neoplasms
-	C09.647.710                              Pharyngeal Neoplasms
-	C09.647.710.485                        Hypopharyngeal Neoplasms
-	C09.647.710.650                        Nasopharyngeal Neoplasms
-	C09.647.710.685                        Oropharyngeal Neoplasms
-	C09.647.710.685.800                      Tonsillar Neoplasms
-	C09.775                                  Pharyngeal Diseases
-	C09.775.174                              Deglutition Disorders
-	C09.775.262                              Lemierre Syndrome
-	C09.775.350                              Nasopharyngeal Diseases
-	C09.775.350.650                        Nasopharyngeal Neoplasms
-	C09.775.350.700                        Nasopharyngitis
-	C09.775.549                              Pharyngeal Neoplasms
-	C09.775.549.485                        Hypopharyngeal Neoplasms
-	C09.775.549.650                        Nasopharyngeal Neoplasms
-	C09.775.549.685                        Oropharyngeal Neoplasms
-	C09.775.549.685.800                      Tonsillar Neoplasms
-	C09.775.649                              Pharyngitis
-	C09.775.649.500                        Nasopharyngitis
-	C09.775.649.625                        Retropharyngeal Abscess
-	C09.775.649.750                        Tonsillitis
-	C09.775.649.750.500                      Peritonsillar Abscess
-	C09.775.955                              Velopharyngeal Insufficiency
-	C10                                        Nervous System Diseases
-	C10.114                                  Autoimmune Diseases of the Nervous System
-	C10.114.375                              Demyelinating Autoimmune Diseases, CNS
-	C10.114.375.112                        Diffuse Cerebral Sclerosis of Schilder
-	C10.114.375.225                        Encephalomyelitis, Acute Disseminated
-	C10.114.375.225.500                      Leukoencephalitis, Acute Hemorrhagic
-	C10.114.375.362                        Leukoencephalitis, Acute Hemorrhagic
-	C10.114.375.500                        Multiple Sclerosis
-	C10.114.375.500.200                      Multiple Sclerosis, Chronic Progressive
-	C10.114.375.500.600                      Multiple Sclerosis, Relapsing-Remitting
-	C10.114.375.600                        Myelitis, Transverse
-	C10.114.375.600.500                      Neuromyelitis Optica

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C10.114.375.800 Neuromyelitis Optica
-	C10.114.468 Lambert-Eaton Myasthenic Syndrome
-	C10.114.656 Myasthenia Gravis
-	C10.114.656.300 Myasthenia Gravis, Autoimmune, Experimental
-	C10.114.656.650 Myasthenia Gravis, Neonatal
-	C10.114.703 Nervous System Autoimmune Disease, Experimental
-	C10.114.703.300 Encephalomyelitis, Autoimmune, Experimental
-	C10.114.703.350 Myasthenia Gravis, Autoimmune, Experimental
-	C10.114.703.700 Neuritis, Autoimmune, Experimental
-	C10.114.750 Polyradiculoneuropathy
-	C10.114.750.100 Guillain-Barre Syndrome
-	C10.114.750.100.500 Miller Fisher Syndrome
-	C10.114.750.175 Polyradiculoneuropathy, Chronic Inflammatory Demyelinating
-	C10.114.812 Stiff-Person Syndrome
-	C10.114.843 Uveomeningoencephalitic Syndrome
-	C10.114.875 Vasculitis, Central Nervous System
-	C10.114.875.175 AIDS Arteritis, Central Nervous System
-	C10.114.875.700 Giant Cell Arteritis
-	C10.114.875.850 Lupus Vasculitis, Central Nervous System
-	C10.177 Autonomic Nervous System Diseases
-	C10.177.045 Adie Syndrome
-	C10.177.090 Autonomic Dysreflexia
-	C10.177.195 Complex Regional Pain Syndromes
-	C10.177.195.200 Causalgia
-	C10.177.195.800 Reflex Sympathetic Dystrophy
-	C10.177.350 Horner Syndrome
-	C10.177.575 Primary Dysautonomias
-	C10.177.575.300 Dysautonomia, Familial
-	C10.177.575.550 Multiple System Atrophy
-	C10.177.575.550.375 Olivopontocerebellar Atrophies
-	C10.177.575.550.750 Shy-Drager Syndrome
-	C10.177.575.550.875 Striatonigral Degeneration
-	C10.177.575.600 Orthostatic Intolerance
-	C10.177.575.600.450 Hypotension, Orthostatic
-	C10.177.575.600.537 Post-Exercise Hypotension
-	C10.177.575.600.625 Postural Orthostatic Tachycardia Syndrome

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C10.177.575.600.800                      Syncope, Vasovagal
-	C10.177.575.650                              Pure Autonomic Failure
-	C10.177.825                                    Sweating, Gustatory
-	C10.228                                        Central Nervous System Diseases
-	C10.228.140                                  Brain Diseases
New Heading	<b>C10.228.140.021                              Acute Febrile Encephalopathy</b>
-	C10.228.140.042                              Akinetic Mutism
-	C10.228.140.055                              Amblyopia
-	C10.228.140.060                              Amnesia, Transient Global
-	C10.228.140.068                              Auditory Diseases, Central
-	C10.228.140.068.094                        Auditory Perceptual Disorders
-	C10.228.140.068.432                        Hearing Loss, Central
-	C10.228.140.079                              Basal Ganglia Diseases
-	C10.228.140.079.127                        Basal Ganglia Cerebrovascular Disease
-	C10.228.140.079.127.500                   Basal Ganglia Hemorrhage
-	C10.228.140.079.127.500.500            Putaminal Hemorrhage
-	C10.228.140.079.294                        Chorea Gravidarum
-	C10.228.140.079.357                        Dystonia Musculorum Deformans
-	C10.228.140.079.493                        Hepatolenticular Degeneration
-	C10.228.140.079.545                        Huntington Disease
-	C10.228.140.079.590                        Meige Syndrome
-	C10.228.140.079.612                        Multiple System Atrophy
-	C10.228.140.079.612.600                   Olivopontocerebellar Atrophies
-	C10.228.140.079.612.700                   Shy-Drager Syndrome
-	C10.228.140.079.612.800                   Striatonigral Degeneration
-	C10.228.140.079.737                        Neuroleptic Malignant Syndrome
-	C10.228.140.079.800                        Pantothenate Kinase-Associated Neurodegeneration
-	C10.228.140.079.862                        Parkinsonian Disorders
-	C10.228.140.079.862.400                   Lewy Body Disease
-	C10.228.140.079.862.500                   Parkinson Disease
-	C10.228.140.079.862.800                   Parkinson Disease, Secondary
-	C10.228.140.079.862.800.300            MPTP Poisoning
-	C10.228.140.079.862.800.600            Parkinson Disease, Postencephalitic
-	C10.228.140.079.882                        Supranuclear Palsy, Progressive
-	C10.228.140.079.898                        Tourette Syndrome



## MeSH Tree Changes for 2017

Type	Tree - heading
-	C10.228.140.116                      Brain Abscess
-	C10.228.140.116.800                      Toxoplasmosis, Cerebral
-	C10.228.140.140                      Brain Damage, Chronic
-	C10.228.140.140.127                      Brain Injury, Chronic
-	C10.228.140.140.254                      Cerebral Palsy
-	C10.228.140.140.627                      Persistent Vegetative State
-	C10.228.140.151                      Brain Death
-	C10.228.140.163                      Brain Diseases, Metabolic
-	C10.228.140.163.100                      Brain Diseases, Metabolic, Inborn
-	C10.228.140.163.100.084                      Adrenoleukodystrophy
-	C10.228.140.163.100.168                      Cerebral Amyloid Angiopathy, Familial
-	C10.228.140.163.100.320                      Galactosemias
-	C10.228.140.163.100.355                      Hartnup Disease
-	C10.228.140.163.100.360                      Hepatolenticular Degeneration
-	C10.228.140.163.100.362                      Hereditary Central Nervous System Demyelinating Diseases
-	C10.228.140.163.100.362.250                      Adrenoleukodystrophy
-	C10.228.140.163.100.362.312                      Alexander Disease
-	C10.228.140.163.100.362.375                      Canavan Disease
-	C10.228.140.163.100.362.500                      Leukodystrophy, Globoid Cell
-	C10.228.140.163.100.362.550                      Leukodystrophy, Metachromatic
-	C10.228.140.163.100.362.775                      Pelizaeus-Merzbacher Disease
-	C10.228.140.163.100.365                      Homocystinuria
-	C10.228.140.163.100.375                      Hyperglycinemia, Nonketotic
-	C10.228.140.163.100.380                      Hyperlysinemias
-	C10.228.140.163.100.412                      Leigh Disease
-	C10.228.140.163.100.425                      Lesch-Nyhan Syndrome
-	C10.228.140.163.100.435                      Lysosomal Storage Diseases, Nervous System
-	C10.228.140.163.100.435.295                      Fucosidosis
-	C10.228.140.163.100.435.340                      Glycogen Storage Disease Type II
-	C10.228.140.163.100.435.590                      Mucopolidoses
-	C10.228.140.163.100.435.810                      Sialic Acid Storage Disease
-	C10.228.140.163.100.435.825                      Sphingolipidoses
-	C10.228.140.163.100.435.825.200                      Fabry Disease
-	C10.228.140.163.100.435.825.250                      Farber Lipogranulomatosis
-	C10.228.140.163.100.435.825.300                      Gangliosidoses

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C10.228.140.163.100.435.825.300.300 Gangliosidoses, GM2
-	C10.228.140.163.100.435.825.300.300.249 Sandhoff Disease
-	C10.228.140.163.100.435.825.300.300.500 Tay-Sachs Disease
-	C10.228.140.163.100.435.825.300.300.750 Tay-Sachs Disease, AB Variant
-	C10.228.140.163.100.435.825.300.400 Gangliosidosis, GM1
-	C10.228.140.163.100.435.825.400 Gaucher Disease
-	C10.228.140.163.100.435.825.590 Leukodystrophy, Globoid Cell
-	C10.228.140.163.100.435.825.700 Niemann-Pick Diseases
-	C10.228.140.163.100.435.825.700.500 Niemann-Pick Disease, Type A
-	C10.228.140.163.100.435.825.700.750 Niemann-Pick Disease, Type B
-	C10.228.140.163.100.435.825.700.875 Niemann-Pick Disease, Type C
-	C10.228.140.163.100.435.825.775 Sea-Blue Histiocyte Syndrome
-	C10.228.140.163.100.435.825.850 Sulfatidosis
-	C10.228.140.163.100.435.825.850.500 Leukodystrophy, Metachromatic
-	C10.228.140.163.100.435.825.850.750 Multiple Sulfatase Deficiency Disease
-	C10.228.140.163.100.520 Maple Syrup Urine Disease
-	C10.228.140.163.100.535 MELAS Syndrome
-	C10.228.140.163.100.540 Menkes Kinky Hair Syndrome
-	C10.228.140.163.100.545 MERRF Syndrome
-	C10.228.140.163.100.593 Mevalonate Kinase Deficiency
-	C10.228.140.163.100.640 Oculocerebrorenal Syndrome
-	C10.228.140.163.100.687 Phenylketonurias
-	C10.228.140.163.100.687.500 Phenylketonuria, Maternal
-	C10.228.140.163.100.725 Pyruvate Carboxylase Deficiency Disease
-	C10.228.140.163.100.750 Pyruvate Dehydrogenase Complex Deficiency Disease
-	C10.228.140.163.100.813 Refsum Disease
-	C10.228.140.163.100.844 Refsum Disease, Infantile
-	C10.228.140.163.100.875 Tyrosinemias
-	C10.228.140.163.100.937 Urea Cycle Disorders, Inborn
-	C10.228.140.163.100.937.124 Argininosuccinic Aciduria
-	C10.228.140.163.100.937.249 Carbamoyl-Phosphate Synthase I Deficiency Disease
-	C10.228.140.163.100.937.374 Citrullinemia
-	C10.228.140.163.100.937.500 Hyperargininemia
-	C10.228.140.163.100.937.750 Ornithine Carbamoyltransferase Deficiency Disease

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	C10.228.140.163.100.968	Zellweger Syndrome
-	C10.228.140.163.360	Hepatic Encephalopathy
-	C10.228.140.163.480	Kernicterus
-	C10.228.140.163.510	Marchiafava-Bignami Disease
-	C10.228.140.163.540	Mitochondrial Encephalomyopathies
-	C10.228.140.163.560	Myelinolysis, Central Pontine
-	C10.228.140.163.780	Reye Syndrome
-	C10.228.140.163.960	Wernicke Encephalopathy
-	C10.228.140.187	Brain Edema
-	C10.228.140.199	Brain Injuries
Old Tree	<b>C10.228.140.199.250</b>	<b>Brain Concussion</b>
Old Tree	<b>C10.228.140.199.250.250</b>	<b>Contrecoup Injury</b>
Old Tree	<b>C10.228.140.199.250.500</b>	<b>Post-Concussion Syndrome</b>
-	C10.228.140.199.275	Brain Hemorrhage, Traumatic
-	C10.228.140.199.275.200	Brain Stem Hemorrhage, Traumatic
-	C10.228.140.199.275.300	Cerebral Hemorrhage, Traumatic
New Heading	<b>C10.228.140.199.388</b>	<b>Brain Injuries, Diffuse</b>
New Tree	<b>C10.228.140.199.388.500</b>	<b>Diffuse Axonal Injury</b>
New Heading	<b>C10.228.140.199.444</b>	<b>Brain Injuries, Traumatic</b>
New Tree	<b>C10.228.140.199.444.250</b>	<b>Brain Concussion</b>
New Heading	<b>C10.228.140.199.444.375</b>	<b>Brain Contusion</b>
New Heading	<b>C10.228.140.199.444.500</b>	<b>Chronic Traumatic Encephalopathy</b>
-	C10.228.140.199.500	Brain Injury, Chronic
New Heading	<b>C10.228.140.199.500.500</b>	<b>Chronic Traumatic Encephalopathy</b>
Old Tree	<b>C10.228.140.199.600</b>	<b>Diffuse Axonal Injury</b>
-	C10.228.140.199.650	Epilepsy, Post-Traumatic
-	C10.228.140.199.700	Pneumocephalus
-	C10.228.140.211	Brain Neoplasms
-	C10.228.140.211.280	Cerebral Ventricle Neoplasms
-	C10.228.140.211.280.300	Choroid Plexus Neoplasms
-	C10.228.140.211.280.300.500	Papilloma, Choroid Plexus

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C10.228.140.211.500                      Infratentorial Neoplasms
-	C10.228.140.211.500.100                      Brain Stem Neoplasms
-	C10.228.140.211.500.200                      Cerebellar Neoplasms
-	C10.228.140.211.692                      Neurocytoma
-	C10.228.140.211.788                      Pinealoma
-	C10.228.140.211.885                      Supratentorial Neoplasms
-	C10.228.140.211.885.500                      Hypothalamic Neoplasms
-	C10.228.140.211.885.500.299                      Pallister-Hall Syndrome
-	C10.228.140.211.885.500.600                      Pituitary Neoplasms
-	C10.228.140.252                      Cerebellar Diseases
-	C10.228.140.252.190                      Cerebellar Ataxia
-	C10.228.140.252.190.530                      Spinocerebellar Ataxias
-	C10.228.140.252.190.530.060                      Ataxia Telangiectasia
-	C10.228.140.252.190.530.530                      Machado-Joseph Disease
-	C10.228.140.252.200                      Cerebellar Neoplasms
-	C10.228.140.252.300                      Dandy-Walker Syndrome
-	C10.228.140.252.500                      Miller Fisher Syndrome
-	C10.228.140.252.620                      Paraneoplastic Cerebellar Degeneration
-	C10.228.140.252.700                      Spinocerebellar Degenerations
-	C10.228.140.252.700.150                      Friedreich Ataxia
-	C10.228.140.252.700.250                      Myoclonic Cerebellar Dyssynergia
-	C10.228.140.252.700.650                      Olivopontocerebellar Atrophies
-	C10.228.140.252.700.700                      Spinocerebellar Ataxias
-	C10.228.140.252.700.700.500                      Machado-Joseph Disease
-	C10.228.140.300                      Cerebrovascular Disorders
-	C10.228.140.300.100                      Basal Ganglia Cerebrovascular Disease
-	C10.228.140.300.100.200                      Basal Ganglia Hemorrhage
-	C10.228.140.300.100.200.500                      Putaminal Hemorrhage
-	C10.228.140.300.150                      Brain Ischemia
-	C10.228.140.300.150.477                      Brain Infarction
-	C10.228.140.300.150.477.100                      Brain Stem Infarctions
-	C10.228.140.300.150.477.100.500                      Lateral Medullary Syndrome
-	C10.228.140.300.150.477.200                      Cerebral Infarction
-	C10.228.140.300.150.477.200.199                      Dementia, Multi-Infarct
-	C10.228.140.300.150.477.200.400                      Infarction, Anterior Cerebral Artery
-	C10.228.140.300.150.477.200.450                      Infarction, Middle Cerebral Artery

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C10.228.140.300.150.477.200.475 Infarction, Posterior Cerebral Artery
-	C10.228.140.300.150.716 Hypoxia-Ischemia, Brain
-	C10.228.140.300.150.836 Ischemic Attack, Transient
-	C10.228.140.300.150.956 Vertebrobasilar Insufficiency
-	C10.228.140.300.150.956.700 Subclavian Steal Syndrome
-	C10.228.140.300.200 Carotid Artery Diseases
-	C10.228.140.300.200.345 Carotid Artery Injuries
-	C10.228.140.300.200.345.300 Carotid Artery, Internal, Dissection
-	C10.228.140.300.200.355 Carotid Artery Thrombosis
-	C10.228.140.300.200.360 Carotid Stenosis
-	C10.228.140.300.200.480 Carotid-Cavernous Sinus Fistula
-	C10.228.140.300.200.600 Moyamoya Disease
-	C10.228.140.300.275 Cerebral Small Vessel Diseases
-	C10.228.140.300.275.249 CADASIL
-	C10.228.140.300.275.311 Cerebral Amyloid Angiopathy, Familial
-	C10.228.140.300.275.374 Fabry Disease
-	C10.228.140.300.275.500 MELAS Syndrome
-	C10.228.140.300.275.600 Microscopic Polyangiitis
-	C10.228.140.300.275.800 Stroke, Lacunar
-	C10.228.140.300.350 Cerebrovascular Trauma
-	C10.228.140.300.350.500 Carotid Artery Injuries
-	C10.228.140.300.350.500.300 Carotid Artery, Internal, Dissection
-	C10.228.140.300.350.500.350 Carotid-Cavernous Sinus Fistula
-	C10.228.140.300.350.875 Vertebral Artery Dissection
-	C10.228.140.300.400 Dementia, Vascular
-	C10.228.140.300.400.203 CADASIL
-	C10.228.140.300.400.408 Dementia, Multi-Infarct
-	C10.228.140.300.510 Intracranial Arterial Diseases
-	C10.228.140.300.510.200 Cerebral Arterial Diseases
-	C10.228.140.300.510.200.175 CADASIL
-	C10.228.140.300.510.200.200 Cerebral Amyloid Angiopathy
-	C10.228.140.300.510.200.200.160 Cerebral Amyloid Angiopathy, Familial
-	C10.228.140.300.510.200.325 Infarction, Anterior Cerebral Artery
-	C10.228.140.300.510.200.387 Infarction, Middle Cerebral Artery
-	C10.228.140.300.510.200.418 Infarction, Posterior Cerebral Artery
-	C10.228.140.300.510.200.737 Moyamoya Disease

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C10.228.140.300.510.600                      Intracranial Aneurysm
-	C10.228.140.300.510.800                      Intracranial Arteriosclerosis
-	C10.228.140.300.510.800.500                      Dementia, Vascular
-	C10.228.140.300.520                      Intracranial Arteriovenous Malformations
-	C10.228.140.300.520.500                      Vein of Galen Malformations
-	C10.228.140.300.525                      Intracranial Embolism and Thrombosis
-	C10.228.140.300.525.400                      Intracranial Embolism
-	C10.228.140.300.525.425                      Intracranial Thrombosis
-	C10.228.140.300.525.425.500                      Sinus Thrombosis, Intracranial
-	C10.228.140.300.525.425.500.375                      Cavernous Sinus Thrombosis
-	C10.228.140.300.525.425.500.562                      Lateral Sinus Thrombosis
-	C10.228.140.300.525.425.500.750                      Sagittal Sinus Thrombosis
-	C10.228.140.300.535                      Intracranial Hemorrhages
-	C10.228.140.300.535.200                      Cerebral Hemorrhage
-	C10.228.140.300.535.200.150                      Basal Ganglia Hemorrhage
-	C10.228.140.300.535.200.150.500                      Putaminal Hemorrhage
-	C10.228.140.300.535.200.200                      Cerebral Hemorrhage, Traumatic
-	C10.228.140.300.535.325                      Intracranial Hemorrhage, Hypertensive
-	C10.228.140.300.535.450                      Intracranial Hemorrhage, Traumatic
-	C10.228.140.300.535.450.200                      Brain Hemorrhage, Traumatic
-	C10.228.140.300.535.450.200.500                      Brain Stem Hemorrhage, Traumatic
-	C10.228.140.300.535.450.200.750                      Cerebral Hemorrhage, Traumatic
-	C10.228.140.300.535.450.300                      Hematoma, Epidural, Cranial
-	C10.228.140.300.535.450.400                      Hematoma, Subdural
-	C10.228.140.300.535.450.400.050                      Hematoma, Subdural, Acute
-	C10.228.140.300.535.450.400.120                      Hematoma, Subdural, Chronic
-	C10.228.140.300.535.450.400.400                      Hematoma, Subdural, Intracranial
-	C10.228.140.300.535.450.650                      Subarachnoid Hemorrhage, Traumatic
-	C10.228.140.300.535.625                      Pituitary Apoplexy
-	C10.228.140.300.535.800                      Subarachnoid Hemorrhage
-	C10.228.140.300.535.800.700                      Subarachnoid Hemorrhage, Traumatic
-	C10.228.140.300.700                      Leukomalacia, Periventricular
-	C10.228.140.300.750                      Sneddon Syndrome
-	C10.228.140.300.775                      Stroke
-	C10.228.140.300.775.200                      Brain Infarction
-	C10.228.140.300.775.200.100                      Brain Stem Infarctions

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C10.228.140.300.775.200.100.500 Lateral Medullary Syndrome
-	C10.228.140.300.775.200.200 Cerebral Infarction
-	C10.228.140.300.775.200.200.199 Dementia, Multi-Infarct
-	C10.228.140.300.775.200.200.400 Infarction, Anterior Cerebral Artery
-	C10.228.140.300.775.200.200.450 Infarction, Middle Cerebral Artery
-	C10.228.140.300.775.200.200.475 Infarction, Posterior Cerebral Artery
-	C10.228.140.300.775.600 Stroke, Lacunar
-	C10.228.140.300.787 Susac Syndrome
-	C10.228.140.300.800 Vascular Headaches
-	C10.228.140.300.850 Vasculitis, Central Nervous System
-	C10.228.140.300.850.125 AIDS Arteritis, Central Nervous System
-	C10.228.140.300.850.500 Giant Cell Arteritis
-	C10.228.140.300.850.750 Lupus Vasculitis, Central Nervous System
-	C10.228.140.300.900 Vasospasm, Intracranial
-	C10.228.140.380 Dementia
-	C10.228.140.380.070 AIDS Dementia Complex
-	C10.228.140.380.100 Alzheimer Disease
-	C10.228.140.380.132 Aphasia, Primary Progressive
-	C10.228.140.380.132.600 Primary Progressive Nonfluent Aphasia
-	C10.228.140.380.165 Creutzfeldt-Jakob Syndrome
-	C10.228.140.380.230 Dementia, Vascular
-	C10.228.140.380.230.124 CADASIL
-	C10.228.140.380.230.250 Dementia, Multi-Infarct
-	C10.228.140.380.254 Diffuse Neurofibrillary Tangles with Calcification
-	C10.228.140.380.266 Frontotemporal Lobar Degeneration
-	C10.228.140.380.266.299 Frontotemporal Dementia
-	C10.228.140.380.266.299.500 Pick Disease of the Brain
-	C10.228.140.380.266.600 Primary Progressive Nonfluent Aphasia
-	C10.228.140.380.278 Huntington Disease
-	C10.228.140.380.326 Kluver-Bucy Syndrome
-	C10.228.140.380.422 Lewy Body Disease
-	C10.228.140.400 Diffuse Cerebral Sclerosis of Schilder
-	C10.228.140.430 Encephalitis
-	C10.228.140.430.124 Anti-N-Methyl-D-Aspartate Receptor Encephalitis
-	C10.228.140.430.249 Cerebral Ventriculitis
-	C10.228.140.430.520 Infectious Encephalitis

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C10.228.140.430.520.750                      Encephalitis, Viral
-	C10.228.140.430.520.750.300                  Encephalitis, Arbovirus
-	C10.228.140.430.520.750.300.200              Encephalitis, California
-	C10.228.140.430.520.750.300.400              Encephalitis, Japanese
-	C10.228.140.430.520.750.300.550              Encephalitis, St. Louis
-	C10.228.140.430.520.750.300.775              Encephalitis, Tick-Borne
-	C10.228.140.430.520.750.300.887              West Nile Fever
-	C10.228.140.430.520.750.350                  Encephalitis, Herpes Simplex
-	C10.228.140.430.520.750.400                  Encephalitis, Varicella Zoster
-	C10.228.140.430.520.750.450                  Encephalomyelitis, Equine
-	C10.228.140.430.520.750.450.200              Encephalomyelitis, Eastern Equine
-	C10.228.140.430.520.750.450.600              Encephalomyelitis, Venezuelan Equine
-	C10.228.140.430.520.750.450.800              Encephalomyelitis, Western Equine
-	C10.228.140.430.520.750.500                  Leukoencephalopathy, Progressive Multifocal
-	C10.228.140.430.520.750.600                  Subacute Sclerosing Panencephalitis
-	C10.228.140.430.525                              Limbic Encephalitis
-	C10.228.140.430.550                              Meningoencephalitis
-	C10.228.140.430.550.500                          Lupus Vasculitis, Central Nervous System
-	C10.228.140.461                                  Encephalomalacia
-	C10.228.140.461.550                              Leukomalacia, Periventricular
-	C10.228.140.490                                  Epilepsy
-	C10.228.140.490.125                              Drug Resistant Epilepsy
-	C10.228.140.490.250                              Epilepsies, Myoclonic
-	C10.228.140.490.250.650                          Myoclonic Epilepsies, Progressive
-	C10.228.140.490.250.650.500                  Lafora Disease
-	C10.228.140.490.250.650.700                  MERRF Syndrome
-	C10.228.140.490.250.650.900                  Unverricht-Lundborg Syndrome
-	C10.228.140.490.250.670                          Myoclonic Epilepsy, Juvenile
-	C10.228.140.490.360                              Epilepsies, Partial
-	C10.228.140.490.360.260                          Epilepsy, Complex Partial
-	C10.228.140.490.360.270                          Epilepsy, Frontal Lobe
-	C10.228.140.490.360.272                          Epilepsy, Partial, Motor
-	C10.228.140.490.360.275                          Epilepsy, Partial, Sensory
-	C10.228.140.490.360.280                          Epilepsy, Rolandic
-	C10.228.140.490.360.290                          Epilepsy, Temporal Lobe
-	C10.228.140.490.370                              Epilepsy, Benign Neonatal



## MeSH Tree Changes for 2017

Type	Tree - heading
-	C10.228.140.490.375 Epilepsy, Generalized
-	C10.228.140.490.375.260 Epilepsy, Absence
-	C10.228.140.490.375.290 Epilepsy, Tonic-Clonic
-	C10.228.140.490.375.525 Nodding Syndrome
-	C10.228.140.490.375.760 Spasms, Infantile
-	C10.228.140.490.380 Epilepsy, Post-Traumatic
-	C10.228.140.490.450 Epilepsy, Reflex
-	C10.228.140.490.535 Landau-Kleffner Syndrome
-	C10.228.140.490.583 Lennox Gastaut Syndrome
-	C10.228.140.490.631 Seizures
-	C10.228.140.490.650 Seizures, Febrile
-	C10.228.140.490.690 Status Epilepticus
-	C10.228.140.490.690.260 Epilepsia Partialis Continua
-	C10.228.140.546 Headache Disorders
-	C10.228.140.546.399 Headache Disorders, Primary
-	C10.228.140.546.399.750 Migraine Disorders
-	C10.228.140.546.399.750.124 Alice in Wonderland Syndrome
-	C10.228.140.546.399.750.250 Migraine with Aura
-	C10.228.140.546.399.750.450 Migraine without Aura
-	C10.228.140.546.399.750.725 Ophthalmoplegic Migraine
-	C10.228.140.546.399.875 Tension-Type Headache
-	C10.228.140.546.399.937 Trigeminal Autonomic Cephalalgias
-	C10.228.140.546.399.937.500 Cluster Headache
-	C10.228.140.546.399.937.750 Paroxysmal Hemicrania
-	C10.228.140.546.399.937.875 SUNCT Syndrome
-	C10.228.140.546.699 Headache Disorders, Secondary
-	C10.228.140.546.699.124 Post-Dural Puncture Headache
-	C10.228.140.546.699.249 Post-Traumatic Headache
-	C10.228.140.546.699.500 Vascular Headaches
-	C10.228.140.602 Hydrocephalus
-	C10.228.140.602.500 Dandy-Walker Syndrome
-	C10.228.140.602.750 Hydrocephalus, Normal Pressure
-	C10.228.140.617 Hypothalamic Diseases
-	C10.228.140.617.200 Bardet-Biedl Syndrome
-	C10.228.140.617.477 Hypothalamic Neoplasms
-	C10.228.140.617.477.299 Pallister-Hall Syndrome

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	C10.228.140.617.477.600	Pituitary Neoplasms
-	C10.228.140.617.500	Laurence-Moon Syndrome
-	C10.228.140.617.738	Pituitary Diseases
Old Tree	C10.228.140.617.738.100	Autoimmune Hypophysitis
-	C10.228.140.617.738.200	Empty Sella Syndrome
-	C10.228.140.617.738.250	Hyperpituitarism
-	C10.228.140.617.738.250.100	Acromegaly
-	C10.228.140.617.738.250.450	Hyperprolactinemia
-	C10.228.140.617.738.250.725	Pituitary ACTH Hypersecretion
New Heading	C10.228.140.617.738.275	Hypophysitis
New Tree	C10.228.140.617.738.275.500	Autoimmune Hypophysitis
-	C10.228.140.617.738.300	Hypopituitarism
-	C10.228.140.617.738.300.300	Dwarfism, Pituitary
-	C10.228.140.617.738.320	Inappropriate ADH Syndrome
-	C10.228.140.617.738.350	Pituitary Apoplexy
-	C10.228.140.617.738.675	Pituitary Neoplasms
-	C10.228.140.617.738.675.149	ACTH-Secreting Pituitary Adenoma
-	C10.228.140.617.738.675.149.500	Nelson Syndrome
-	C10.228.140.617.738.675.299	Growth Hormone-Secreting Pituitary Adenoma
-	C10.228.140.617.738.675.800	Prolactinoma
-	C10.228.140.624	Hypoxia, Brain
-	C10.228.140.624.500	Hypoxia-Ischemia, Brain
-	C10.228.140.631	Intracranial Hypertension
-	C10.228.140.631.500	Hypertensive Encephalopathy
-	C10.228.140.631.500.500	Posterior Leukoencephalopathy Syndrome
-	C10.228.140.631.750	Pseudotumor Cerebri
-	C10.228.140.638	Intracranial Hypotension
-	C10.228.140.695	Leukoencephalopathies
-	C10.228.140.695.500	Dementia, Vascular
-	C10.228.140.695.562	Demyelinating Autoimmune Diseases, CNS
-	C10.228.140.695.562.112	Diffuse Cerebral Sclerosis of Schilder
-	C10.228.140.695.562.225	Encephalomyelitis, Acute Disseminated
-	C10.228.140.695.562.225.500	Leukoencephalitis, Acute Hemorrhagic
-	C10.228.140.695.562.250	Encephalomyelitis, Autoimmune, Experimental

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C10.228.140.695.562.375                      Leukoencephalitis, Acute Hemorrhagic
-	C10.228.140.695.625                              Hereditary Central Nervous System Demyelinating Diseases
-	C10.228.140.695.625.250                      Adrenoleukodystrophy
-	C10.228.140.695.625.312                      Alexander Disease
-	C10.228.140.695.625.375                      Canavan Disease
-	C10.228.140.695.625.500                      Leukodystrophy, Globoid Cell
-	C10.228.140.695.625.550                      Leukodystrophy, Metachromatic
-	C10.228.140.695.625.775                      Pelizaeus-Merzbacher Disease
-	C10.228.140.695.750                              Leukoencephalopathy, Progressive Multifocal
-	C10.228.140.695.875                              Posterior Leukoencephalopathy Syndrome
-	C10.228.140.744                                      Neuroaxonal Dystrophies
-	C10.228.140.744.320                              Pantothenate Kinase-Associated Neurodegeneration
-	C10.228.140.807                                      Sepsis-Associated Encephalopathy
-	C10.228.140.870                                      Subdural Effusion
-	C10.228.140.915                                      Thalamic Diseases
-	C10.228.228    Central Nervous System Infections
-	C10.228.228.090                                      Brain Abscess
-	C10.228.228.090.800                              Toxoplasmosis, Cerebral
-	C10.228.228.180                                      Central Nervous System Bacterial Infections
-	C10.228.228.180.437                              Lyme Neuroborreliosis
-	C10.228.228.180.500                              Meningitis, Bacterial
-	C10.228.228.180.500.350                      Meningitis, Escherichia coli
-	C10.228.228.180.500.425                      Meningitis, Haemophilus
-	C10.228.228.180.500.500                      Meningitis, Listeria
-	C10.228.228.180.500.750                      Meningitis, Meningococcal
-	C10.228.228.180.500.750.500              Waterhouse-Friderichsen Syndrome
-	C10.228.228.180.500.875                      Meningitis, Pneumococcal
-	C10.228.228.180.500.937                      Tuberculosis, Meningeal
-	C10.228.228.180.600                              Neurosyphilis
-	C10.228.228.180.600.800                      Tabes Dorsalis
-	C10.228.228.180.850                              Tuberculosis, Central Nervous System
-	C10.228.228.180.850.400                      Tuberculoma, Intracranial
-	C10.228.228.180.850.800                      Tuberculosis, Meningeal
-	C10.228.228.198                                      Central Nervous System Fungal Infections
-	C10.228.228.198.500                              Meningitis, Fungal

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C10.228.228.198.500.500                      Meningitis, Cryptococcal
-	C10.228.228.198.750                              Neuroaspergillosis
-	C10.228.228.205                                  Central Nervous System Parasitic Infections
-	C10.228.228.205.250                              Central Nervous System Helminthiasis
-	C10.228.228.205.250.550                              Neurocysticercosis
-	C10.228.228.205.250.600                              Neuroschistosomiasis
-	C10.228.228.205.300                                  Central Nervous System Protozoal Infections
-	C10.228.228.205.300.500                              Malaria, Cerebral
-	C10.228.228.205.300.800                              Toxoplasmosis, Cerebral
-	C10.228.228.205.300.900                              Toxoplasmosis, Congenital
-	C10.228.228.245                                      Central Nervous System Viral Diseases
-	C10.228.228.245.169                                  Cerebral Ventriculitis
-	C10.228.228.245.340                                  Encephalitis, Viral
-	C10.228.228.245.340.300                              Encephalitis, Arbovirus
-	C10.228.228.245.340.300.200                              Encephalitis, California
-	C10.228.228.245.340.300.400                              Encephalitis, Japanese
-	C10.228.228.245.340.300.550                              Encephalitis, St. Louis
-	C10.228.228.245.340.300.775                              Encephalitis, Tick-Borne
-	C10.228.228.245.340.300.887                              West Nile Fever
-	C10.228.228.245.340.350                              Encephalitis, Herpes Simplex
-	C10.228.228.245.340.400                              Encephalitis, Varicella Zoster
-	C10.228.228.245.340.450                              Encephalomyelitis, Equine
-	C10.228.228.245.340.450.200                              Encephalomyelitis, Eastern Equine
-	C10.228.228.245.340.450.600                              Encephalomyelitis, Venezuelan Equine
-	C10.228.228.245.340.450.800                              Encephalomyelitis, Western Equine
-	C10.228.228.245.340.500                              Leukoencephalopathy, Progressive Multifocal
-	C10.228.228.245.340.600                              Subacute Sclerosing Panencephalitis
-	C10.228.228.245.500                                  Meningitis, Viral
-	C10.228.228.245.500.500                              Lymphocytic Choriomeningitis
-	C10.228.228.245.550                                  Meningoencephalitis
-	C10.228.228.245.550.500                              Lupus Vasculitis, Central Nervous System
-	C10.228.228.245.700                                  Limbic Encephalitis
-	C10.228.228.245.710                                  Pseudorabies
-	C10.228.228.291                                      Encephalomyelitis
-	C10.228.228.291.323                                  Encephalomyelitis, Equine
-	C10.228.228.291.323.162                              Encephalomyelitis, Eastern Equine

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C10.228.228.291.323.325                      Encephalomyelitis, Venezuelan Equine
-	C10.228.228.291.323.662                      Encephalomyelitis, Western Equine
-	C10.228.228.300                                  Epidural Abscess
-	C10.228.228.399                                  Infectious Encephalitis
-	C10.228.228.399.750                              Encephalitis, Viral
-	C10.228.228.399.750.300                          Encephalitis, Arbovirus
-	C10.228.228.399.750.300.200                      Encephalitis, California
-	C10.228.228.399.750.300.400                      Encephalitis, Japanese
-	C10.228.228.399.750.300.550                      Encephalitis, St. Louis
-	C10.228.228.399.750.300.775                      Encephalitis, Tick-Borne
-	C10.228.228.399.750.300.887                      West Nile Fever
-	C10.228.228.399.750.350                          Encephalitis, Herpes Simplex
-	C10.228.228.399.750.400                          Encephalitis, Varicella Zoster
-	C10.228.228.399.750.450                          Encephalomyelitis, Equine
-	C10.228.228.399.750.450.200                      Encephalomyelitis, Eastern Equine
-	C10.228.228.399.750.450.600                      Encephalomyelitis, Venezuelan Equine
-	C10.228.228.399.750.450.800                      Encephalomyelitis, Western Equine
-	C10.228.228.399.750.500                          Leukoencephalopathy, Progressive Multifocal
-	C10.228.228.399.750.600                          Subacute Sclerosing Panencephalitis
-	C10.228.228.570                                  Meningoencephalitis
-	C10.228.228.570.450                                  Lupus Vasculitis, Central Nervous System
-	C10.228.228.618                                  Myelitis
-	C10.228.228.618.250                                  Myelitis, Transverse
-	C10.228.228.618.500                                  Paraparesis, Tropical Spastic
-	C10.228.228.618.750                                  Poliomyelitis
-	C10.228.228.618.750.500                              Poliomyelitis, Bulbar
-	C10.228.228.618.750.750                              Postpoliomyelitis Syndrome
-	C10.228.228.709                                  Perimeningeal Infections
-	C10.228.228.709.300                                  Empyema, Subdural
-	C10.228.228.709.350                                  Epidural Abscess
-	C10.228.228.709.675                                  Subdural Effusion
-	C10.228.228.800                                  Prion Diseases
-	C10.228.228.800.230                                  Creutzfeldt-Jakob Syndrome
-	C10.228.228.800.260                                  Encephalopathy, Bovine Spongiform
-	C10.228.228.800.350                                  Gerstmann-Straussler-Scheinker Disease
-	C10.228.228.800.392                                  Insomnia, Fatal Familial

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C10.228.228.800.435 Kuru
-	C10.228.228.800.717 Scrapie
-	C10.228.228.800.858 Wasting Disease, Chronic
-	C10.228.440 Encephalomyelitis
-	C10.228.440.406 Encephalomyelitis, Equine
-	C10.228.440.406.200 Encephalomyelitis, Eastern Equine
-	C10.228.440.406.225 Encephalomyelitis, Venezuelan Equine
-	C10.228.440.406.250 Encephalomyelitis, Western Equine
-	C10.228.440.600 Fatigue Syndrome, Chronic
-	C10.228.566 High Pressure Neurological Syndrome
New Heading	<b>C10.228.590 Hyperekplexia</b>
-	C10.228.614 Meningitis
-	C10.228.614.097 Arachnoiditis
-	C10.228.614.220 Meningitis, Aseptic
-	C10.228.614.280 Meningitis, Bacterial
-	C10.228.614.280.350 Meningitis, Escherichia coli
-	C10.228.614.280.393 Meningitis, Haemophilus
-	C10.228.614.280.449 Meningitis, Listeria
-	C10.228.614.280.505 Meningitis, Meningococcal
-	C10.228.614.280.505.904 Waterhouse-Friderichsen Syndrome
-	C10.228.614.280.560 Meningitis, Pneumococcal
-	C10.228.614.280.915 Tuberculosis, Meningeal
-	C10.228.614.300 Meningitis, Fungal
-	C10.228.614.300.500 Meningitis, Cryptococcal
-	C10.228.614.400 Meningitis, Viral
-	C10.228.614.400.500 Lymphocytic Choriomeningitis
-	C10.228.614.500 Meningoencephalitis
-	C10.228.614.500.500 Lupus Vasculitis, Central Nervous System
-	C10.228.662 Movement Disorders
-	C10.228.662.037 Akathisia, Drug-Induced
-	C10.228.662.075 Angelman Syndrome
-	C10.228.662.262 Dyskinesias
-	C10.228.662.262.249 Chorea
-	C10.228.662.262.249.500 Chorea Gravidarum
-	C10.228.662.262.249.750 Huntington Disease

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C10.228.662.262.249.937                      Neuroacanthocytosis
-	C10.228.662.262.500                              Dyskinesia, Drug-Induced
New Heading	<b>C10.228.662.262.500.500</b> <b>Tardive Dyskinesia</b>
-	C10.228.662.300                                  Dystonic Disorders
-	C10.228.662.300.200                              Dystonia Musculorum Deformans
-	C10.228.662.300.500                              Meige Syndrome
-	C10.228.662.300.750                              Torticollis
-	C10.228.662.350                                  Essential Tremor
-	C10.228.662.400                                  Hepatolenticular Degeneration
-	C10.228.662.550                                  Multiple System Atrophy
-	C10.228.662.550.600                              Olivopontocerebellar Atrophies
-	C10.228.662.550.700                              Shy-Drager Syndrome
-	C10.228.662.550.800                              Striatonigral Degeneration
-	C10.228.662.575                                  Pantothenate Kinase-Associated Neurodegeneration
-	C10.228.662.600                                  Parkinsonian Disorders
-	C10.228.662.600.200                              Lewy Body Disease
-	C10.228.662.600.400                              Parkinson Disease
-	C10.228.662.600.700                              Parkinson Disease, Secondary
-	C10.228.662.600.700.250                              MPTP Poisoning
-	C10.228.662.600.700.500                              Parkinson Disease, Postencephalitic
-	C10.228.662.700                                  Supranuclear Palsy, Progressive
-	C10.228.662.825                                  Tic Disorders
-	C10.228.662.825.800                              Tourette Syndrome
-	C10.228.758                                      Ocular Motility Disorders
-	C10.228.758.500                                  Opsoclonus-Myoclonus Syndrome
-	C10.228.806                                      Pneumocephalus
-	C10.228.854                                      Spinal Cord Diseases
-	C10.228.854.139                                  Amyotrophic Lateral Sclerosis
-	C10.228.854.303                                  Epidural Abscess
-	C10.228.854.468                                  Muscular Atrophy, Spinal
-	C10.228.854.468.399                              Bulbo-Spinal Atrophy, X-Linked
-	C10.228.854.468.800                              Spinal Muscular Atrophies of Childhood
-	C10.228.854.525                                  Myelitis
-	C10.228.854.525.553                              Myelitis, Transverse
-	C10.228.854.525.700                              Paraparesis, Tropical Spastic

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C10.228.854.525.850                      Poliomyelitis
-	C10.228.854.525.850.500                      Poliomyelitis, Bulbar
-	C10.228.854.525.850.750                      Postpoliomyelitis Syndrome
-	C10.228.854.583                      Pneumorrhachis
-	C10.228.854.761                      Spinal Cord Compression
-	C10.228.854.765                      Spinal Cord Neoplasms
-	C10.228.854.765.342                      Epidural Neoplasms
-	C10.228.854.770                      Spinal Cord Injuries
-	C10.228.854.770.500                      Central Cord Syndrome
-	C10.228.854.785                      Spinal Cord Vascular Diseases
-	C10.228.854.785.650                      Spinal Cord Ischemia
-	C10.228.854.785.650.100                      Anterior Spinal Artery Syndrome
-	C10.228.854.787                      Spinocerebellar Degenerations
-	C10.228.854.787.200                      Friedreich Ataxia
-	C10.228.854.787.500                      Myoclonic Cerebellar Dyssynergia
-	C10.228.854.787.750                      Olivopontocerebellar Atrophies
-	C10.228.854.787.875                      Spinocerebellar Ataxias
-	C10.228.854.787.875.500                      Machado-Joseph Disease
-	C10.228.854.790                      Stiff-Person Syndrome
-	C10.228.854.811                      Subacute Combined Degeneration
-	C10.228.854.833                      Syringomyelia
-	C10.228.854.889                      Tabes Dorsalis
-	C10.281                      Chronobiology Disorders
-	C10.281.440                      Jet Lag Syndrome
-	C10.281.800                      Sleep Disorders, Circadian Rhythm
-	C10.281.900                      Smith-Magenis Syndrome
-	C10.292                      Cranial Nerve Diseases
-	C10.292.150                      Abducens Nerve Diseases
-	C10.292.150.100                      Abducens Nerve Injury
-	C10.292.175                      Accessory Nerve Diseases
-	C10.292.175.500                      Accessory Nerve Injuries
-	C10.292.225                      Cranial Nerve Neoplasms
-	C10.292.225.750                      Neuroma, Acoustic
-	C10.292.225.750.500                      Neurofibromatosis 2
-	C10.292.225.800                      Optic Nerve Neoplasms
-	C10.292.225.800.500                      Optic Nerve Glioma



## MeSH Tree Changes for 2017

Type	Tree - heading
-	C10.292.262                      Cranial Nerve Injuries
-	C10.292.262.200                  Abducens Nerve Injury
-	C10.292.262.350                  Accessory Nerve Injuries
-	C10.292.262.500                  Facial Nerve Injuries
-	C10.292.262.531                  Glossopharyngeal Nerve Injuries
-	C10.292.262.562                  Hypoglossal Nerve Injuries
-	C10.292.262.656                  Oculomotor Nerve Injuries
-	C10.292.262.687                  Olfactory Nerve Injuries
-	C10.292.262.750                  Optic Nerve Injuries
-	C10.292.262.875                  Trigeminal Nerve Injuries
-	C10.292.262.875.500              Lingual Nerve Injuries
-	C10.292.262.906                  Trochlear Nerve Injuries
-	C10.292.262.937                  Vagus Nerve Injuries
-	C10.292.262.937.750              Laryngeal Nerve Injuries
-	C10.292.262.937.750.500        Recurrent Laryngeal Nerve Injuries
-	C10.292.262.968                  Vestibulocochlear Nerve Injuries
-	C10.292.319                      Facial Nerve Diseases
-	C10.292.319.250                  Bell Palsy
-	C10.292.319.375                  Facial Hemiatrophy
-	C10.292.319.500                  Facial Nerve Injuries
-	C10.292.319.625                  Facial Neuralgia
-	C10.292.319.625.700              Trigeminal Nerve Diseases
-	C10.292.319.625.700.349        Trigeminal Nerve Injuries
-	C10.292.319.625.700.349.500    Lingual Nerve Injuries
-	C10.292.319.625.700.700        Trigeminal Neuralgia
-	C10.292.319.750                  Herpes Zoster Oticus
-	C10.292.319.800                  Melkersson-Rosenthal Syndrome
-	C10.292.319.825                  Mobius Syndrome
-	C10.292.450                      Glossopharyngeal Nerve Diseases
-	C10.292.450.500                  Glossopharyngeal Nerve Injuries
-	C10.292.525                      Hypoglossal Nerve Diseases
-	C10.292.525.500                  Hypoglossal Nerve Injuries
-	C10.292.562                      Ocular Motility Disorders
-	C10.292.562.250                  Duane Retraction Syndrome
-	C10.292.562.350                  Miller Fisher Syndrome
-	C10.292.562.675                  Nystagmus, Pathologic

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C10.292.562.675.300 Nystagmus, Congenital
-	C10.292.562.700 Oculomotor Nerve Diseases
-	C10.292.562.700.250 Adie Syndrome
-	C10.292.562.700.500 Oculomotor Nerve Injuries
-	C10.292.562.750 Ophthalmoplegia
-	C10.292.562.750.250 Ophthalmoplegia, Chronic Progressive External
-	C10.292.562.750.250.500 Kearns-Sayre Syndrome
-	C10.292.562.750.375 Ophthalmoplegic Migraine
-	C10.292.562.750.500 Supranuclear Palsy, Progressive
-	C10.292.562.831 Opsoclonus-Myoclonus Syndrome
-	C10.292.562.887 Strabismus
-	C10.292.562.887.300 Esotropia
-	C10.292.562.887.650 Exotropia
-	C10.292.562.900 Tolosa-Hunt Syndrome
-	C10.292.650 Olfactory Nerve Diseases
-	C10.292.650.200 Esthesioneuroblastoma, Olfactory
-	C10.292.650.600 Olfactory Nerve Injuries
-	C10.292.675 Ophthalmoplegic Migraine
-	C10.292.700 Optic Nerve Diseases
-	C10.292.700.225 Optic Atrophy
-	C10.292.700.225.500 Optic Atrophies, Hereditary
-	C10.292.700.225.500.100 Optic Atrophy, Autosomal Dominant
-	C10.292.700.225.500.400 Optic Atrophy, Hereditary, Leber
-	C10.292.700.225.500.980 Wolfram Syndrome
-	C10.292.700.450 Optic Disk Drusen
-	C10.292.700.475 Optic Nerve Injuries
-	C10.292.700.500 Optic Nerve Neoplasms
-	C10.292.700.500.500 Optic Nerve Glioma
-	C10.292.700.550 Optic Neuritis
-	C10.292.700.550.500 Neuromyelitis Optica
-	C10.292.700.600 Optic Neuropathy, Ischemic
-	C10.292.700.900 Papilledema
-	C10.292.850 Trochlear Nerve Diseases
-	C10.292.887 Vagus Nerve Diseases
-	C10.292.887.399 Vagus Nerve Injuries
-	C10.292.887.800 Vocal Cord Paralysis

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C10.292.910 Vestibulocochlear Nerve Diseases
-	C10.292.910.299 Cogan Syndrome
-	C10.292.910.600 Neuroma, Acoustic
-	C10.292.910.600.500 Neurofibromatosis 2
-	C10.292.910.850 Vestibular Neuronitis
-	C10.292.910.925 Vestibulocochlear Nerve Injuries
-	C10.314 Demyelinating Diseases
-	C10.314.350 Demyelinating Autoimmune Diseases, CNS
-	C10.314.350.112 Diffuse Cerebral Sclerosis of Schilder
-	C10.314.350.225 Encephalomyelitis, Acute Disseminated
-	C10.314.350.225.500 Leukoencephalitis, Acute Hemorrhagic
-	C10.314.350.250 Encephalomyelitis, Autoimmune, Experimental
-	C10.314.350.375 Leukoencephalitis, Acute Hemorrhagic
-	C10.314.350.500 Multiple Sclerosis
-	C10.314.350.500.200 Multiple Sclerosis, Chronic Progressive
-	C10.314.350.500.600 Multiple Sclerosis, Relapsing-Remitting
-	C10.314.350.600 Myelitis, Transverse
-	C10.314.350.600.500 Neuromyelitis Optica
-	C10.314.350.800 Neuromyelitis Optica
-	C10.314.400 Hereditary Central Nervous System Demyelinating Diseases
-	C10.314.400.250 Adrenoleukodystrophy
-	C10.314.400.312 Alexander Disease
-	C10.314.400.375 Canavan Disease
-	C10.314.400.500 Leukodystrophy, Globoid Cell
-	C10.314.400.550 Leukodystrophy, Metachromatic
-	C10.314.400.775 Pelizaeus-Merzbacher Disease
-	C10.314.450 Leukoencephalopathy, Progressive Multifocal
-	C10.314.475 Marchiafava-Bignami Disease
-	C10.314.500 Myelinolysis, Central Pontine
-	C10.314.687 Ophthalmoplegic Migraine
-	C10.314.750 Polyradiculoneuropathy
-	C10.314.750.450 Guillain-Barre Syndrome
-	C10.314.750.450.500 Miller Fisher Syndrome
-	C10.314.750.700 Polyradiculoneuropathy, Chronic Inflammatory Demyelinating
-	C10.314.875 Subacute Combined Degeneration
-	C10.500 Nervous System Malformations

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C10.500.034                      Agenesis of Corpus Callosum
-	C10.500.034.500                  Acrocallosal Syndrome
-	C10.500.034.687                  Aicardi Syndrome
-	C10.500.034.875                  Holoprosencephaly
-	C10.500.142                      Central Nervous System Cysts
-	C10.500.142.100                  Arachnoid Cysts
-	C10.500.142.200                  Colloid Cysts
-	C10.500.190                      Central Nervous System Vascular Malformations
-	C10.500.190.100                  Central Nervous System Venous Angioma
-	C10.500.190.200                  Hemangioma, Cavernous, Central Nervous System
-	C10.500.190.500                  Intracranial Arteriovenous Malformations
-	C10.500.190.500.500              Vein of Galen Malformations
-	C10.500.190.800                  Sinus Pericranii
-	C10.500.205                      Dandy-Walker Syndrome
-	C10.500.300                      Hereditary Sensory and Motor Neuropathy
-	C10.500.300.099                  Alstrom Syndrome
-	C10.500.300.200                  Charcot-Marie-Tooth Disease
-	C10.500.300.490                  Giant Axonal Neuropathy
-	C10.500.300.780                  Refsum Disease
-	C10.500.300.820                  Spastic Paraplegia, Hereditary
-	C10.500.310                      Hereditary Sensory and Autonomic Neuropathies
-	C10.500.310.309                  Dysautonomia, Familial
-	C10.500.450                      Hydranencephaly
-	C10.500.507                      Malformations of Cortical Development
-	C10.500.507.400                  Malformations of Cortical Development, Group I
-	C10.500.507.400.249              Megalencephaly
-	C10.500.507.400.249.500        Hemimegalencephaly
-	C10.500.507.400.500              Microcephaly
-	C10.500.507.400.750              Tuberous Sclerosis
-	C10.500.507.450                  Malformations of Cortical Development, Group II
-	C10.500.507.450.230              Classical Lissencephalies and Subcortical Band Heterotopias
-	C10.500.507.450.249              Cobblestone Lissencephaly
-	C10.500.507.450.499              Lissencephaly
-	C10.500.507.450.499.230        Classical Lissencephalies and Subcortical Band Heterotopias

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C10.500.507.450.499.249 Cobblestone Lissencephaly
-	C10.500.507.450.499.249.500 Walker-Warburg Syndrome
-	C10.500.507.450.750 Periventricular Nodular Heterotopia
-	C10.500.507.500 Malformations of Cortical Development, Group III
-	C10.500.507.500.500 Polymicrogyria
-	C10.500.507.500.625 Porencephaly
-	C10.500.507.500.750 Schizencephaly
-	C10.500.680 Neural Tube Defects
-	C10.500.680.196 Anencephaly
-	C10.500.680.291 Arnold-Chiari Malformation
-	C10.500.680.488 Encephalocele
-	C10.500.680.598 Meningocele
-	C10.500.680.610 Meningomyelocele
-	C10.500.680.705 Pentalogy of Cantrell
-	C10.500.680.800 Spinal Dysraphism
-	C10.500.680.800.730 Spina Bifida Cystica
-	C10.500.680.800.750 Spina Bifida Occulta
-	C10.500.840 Septo-Optic Dysplasia
-	C10.551 Nervous System Neoplasms
-	C10.551.240 Central Nervous System Neoplasms
-	C10.551.240.250 Brain Neoplasms
-	C10.551.240.250.200 Cerebral Ventricle Neoplasms
-	C10.551.240.250.200.200 Choroid Plexus Neoplasms
-	C10.551.240.250.200.200.500 Papilloma, Choroid Plexus
-	C10.551.240.250.400 Infratentorial Neoplasms
-	C10.551.240.250.400.200 Brain Stem Neoplasms
-	C10.551.240.250.400.300 Cerebellar Neoplasms
-	C10.551.240.250.550 Neurocytoma
-	C10.551.240.250.625 Pinealoma
-	C10.551.240.250.700 Supratentorial Neoplasms
-	C10.551.240.250.700.500 Hypothalamic Neoplasms
-	C10.551.240.250.700.500.249 Pallister-Hall Syndrome
-	C10.551.240.250.700.500.500 Pituitary Neoplasms
-	C10.551.240.375 Central Nervous System Cysts
-	C10.551.240.375.100 Arachnoid Cysts
-	C10.551.240.375.200 Colloid Cysts

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C10.551.240.500 Meningeal Neoplasms
-	C10.551.240.500.150 Meningeal Carcinomatosis
-	C10.551.240.500.500 Meningioma
-	C10.551.240.750 Spinal Cord Neoplasms
-	C10.551.240.750.200 Epidural Neoplasms
-	C10.551.360 Cranial Nerve Neoplasms
-	C10.551.360.500 Optic Nerve Neoplasms
-	C10.551.360.500.500 Optic Nerve Glioma
-	C10.551.775 Peripheral Nervous System Neoplasms
-	C10.551.775.250 Cranial Nerve Neoplasms
-	C10.551.775.250.500 Optic Nerve Neoplasms
-	C10.551.775.250.500.500 Optic Nerve Glioma
-	C10.551.775.500 Nerve Sheath Neoplasms
-	C10.551.775.500.750 Neurofibroma
-	C10.551.775.500.750.500 Neurofibroma, Plexiform
-	C10.551.775.500.750.750 Neurofibrosarcoma
-	C10.562 Neurocutaneous Syndromes
-	C10.562.100 Ataxia Telangiectasia
-	C10.562.600 Neurofibromatoses
-	C10.562.600.500 Neurofibromatosis 1
-	C10.562.600.750 Neurofibromatosis 2
-	C10.562.700 Nevus, Sebaceous of Jadassohn
-	C10.562.800 Sturge-Weber Syndrome
-	C10.562.850 Tuberous Sclerosis
-	C10.562.925 von Hippel-Lindau Disease
-	C10.574 Neurodegenerative Diseases
New Heading	<b>C10.574.250 Chronic Traumatic Encephalopathy</b>
-	C10.574.500 Heredodegenerative Disorders, Nervous System
-	C10.574.500.024 Alexander Disease
-	C10.574.500.050 Amyloid Neuropathies, Familial
-	C10.574.500.175 Bulbo-Spinal Atrophy, X-Linked
-	C10.574.500.300 Canavan Disease
-	C10.574.500.362 Cockayne Syndrome
-	C10.574.500.393 Dystonia Musculorum Deformans
-	C10.574.500.425 Gerstmann-Straussler-Scheinker Disease

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C10.574.500.487 Hepatolenticular Degeneration
-	C10.574.500.494 Hereditary Central Nervous System Demyelinating Diseases
-	C10.574.500.495 Hereditary Sensory and Motor Neuropathy
-	C10.574.500.495.099 Alstrom Syndrome
-	C10.574.500.495.200 Charcot-Marie-Tooth Disease
-	C10.574.500.495.490 Giant Axonal Neuropathy
-	C10.574.500.495.780 Refsum Disease
-	C10.574.500.495.820 Spastic Paraplegia, Hereditary
-	C10.574.500.496 Hereditary Sensory and Autonomic Neuropathies
-	C10.574.500.496.250 Dysautonomia, Familial
-	C10.574.500.497 Huntington Disease
-	C10.574.500.529 Lafora Disease
-	C10.574.500.545 Myotonia Congenita
-	C10.574.500.547 Myotonic Dystrophy
-	C10.574.500.549 Neurofibromatoses
-	C10.574.500.549.400 Neurofibromatosis 1
-	C10.574.500.549.700 Neurofibromatosis 2
-	C10.574.500.550 Neuronal Ceroid-Lipofuscinoses
-	C10.574.500.662 Optic Atrophies, Hereditary
-	C10.574.500.662.100 Optic Atrophy, Autosomal Dominant
-	C10.574.500.662.400 Optic Atrophy, Hereditary, Leber
-	C10.574.500.662.980 Wolfram Syndrome
-	C10.574.500.700 Pantothenate Kinase-Associated Neurodegeneration
-	C10.574.500.812 Spinal Muscular Atrophies of Childhood
-	C10.574.500.825 Spinocerebellar Degenerations
-	C10.574.500.825.200 Friedreich Ataxia
-	C10.574.500.825.250 Myoclonic Cerebellar Dyssynergia
-	C10.574.500.825.650 Olivopontocerebellar Atrophies
-	C10.574.500.825.700 Spinocerebellar Ataxias
-	C10.574.500.825.700.500 Machado-Joseph Disease
-	C10.574.500.850 Tourette Syndrome
-	C10.574.500.865 Tuberous Sclerosis
-	C10.574.500.875 Unverricht-Lundborg Syndrome
-	C10.574.531 Lewy Body Disease
-	C10.574.562 Motor Neuron Disease
-	C10.574.562.250 Amyotrophic Lateral Sclerosis

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C10.574.562.300      Bulbar Palsy, Progressive
-	C10.574.562.500      Muscular Atrophy, Spinal
-	C10.574.562.500.374      Bulbo-Spinal Atrophy, X-Linked
-	C10.574.562.500.750      Spinal Muscular Atrophies of Childhood
-	C10.574.625      Multiple System Atrophy
-	C10.574.625.600      Olivopontocerebellar Atrophies
-	C10.574.625.700      Shy-Drager Syndrome
-	C10.574.625.800      Striatonigral Degeneration
-	C10.574.781      Paraneoplastic Syndromes, Nervous System
-	C10.574.781.249      Anti-N-Methyl-D-Aspartate Receptor Encephalitis
-	C10.574.781.500      Lambert-Eaton Myasthenic Syndrome
-	C10.574.781.550      Limbic Encephalitis
-	C10.574.781.625      Myelitis, Transverse
-	C10.574.781.662      Opsoclonus-Myoclonus Syndrome
-	C10.574.781.700      Paraneoplastic Cerebellar Degeneration
-	C10.574.781.850      Paraneoplastic Polyneuropathy
-	C10.574.812      Parkinson Disease
-	C10.574.827      Postpoliomyelitis Syndrome
-	C10.574.843      Prion Diseases
-	C10.574.843.300      Encephalopathy, Bovine Spongiform
-	C10.574.843.400      Gerstmann-Straussler-Scheinker Disease
-	C10.574.843.512      Insomnia, Fatal Familial
-	C10.574.843.625      Kuru
-	C10.574.843.850      Scrapie
-	C10.574.843.925      Wasting Disease, Chronic
-	C10.574.910      Subacute Combined Degeneration
-	C10.574.945      Tauopathies
-	C10.574.945.249      Alzheimer Disease
-	C10.574.945.374      Diffuse Neurofibrillary Tangles with Calcification
-	C10.574.945.500      Supranuclear Palsy, Progressive
-	C10.574.950      TDP-43 Proteinopathies
-	C10.574.950.050      Amyotrophic Lateral Sclerosis
-	C10.574.950.300      Frontotemporal Lobar Degeneration
-	C10.574.950.300.299      Frontotemporal Dementia
-	C10.574.950.300.600      Primary Progressive Nonfluent Aphasia
-	C10.597      Neurologic Manifestations



## MeSH Tree Changes for 2017

Type	Tree - heading	
New Heading	<b>C10.597.057</b>	<b>Bilateral Vestibulopathy</b>
-	C10.597.114	Cerebrospinal Fluid Leak
-	C10.597.114.500	Cerebrospinal Fluid Otorrhea
-	C10.597.114.750	Cerebrospinal Fluid Rhinorrhea
-	C10.597.305	Decerebrate State
-	C10.597.350	Dyskinesias
-	C10.597.350.090	Ataxia
-	C10.597.350.090.500	Cerebellar Ataxia
-	C10.597.350.090.500.530	Spinocerebellar Ataxias
-	C10.597.350.090.500.530.060	Ataxia Telangiectasia
-	C10.597.350.090.500.530.530	Machado-Joseph Disease
-	C10.597.350.090.750	Gait Ataxia
-	C10.597.350.110	Athetosis
-	C10.597.350.200	Catalepsy
-	C10.597.350.250	Chorea
New Tree	<b>C10.597.350.275</b>	<b>Dyskinesia, Drug-Induced</b>
New Heading	<b>C10.597.350.275.500</b>	<b>Tardive Dyskinesia</b>
-	C10.597.350.300	Dystonia
-	C10.597.350.300.800	Torticollis
-	C10.597.350.350	Hyperkinesia
-	C10.597.350.400	Hypokinesia
-	C10.597.350.500	Myoclonus
-	C10.597.350.500.500	Opsoclonus-Myoclonus Syndrome
-	C10.597.350.600	Psychomotor Agitation
-	C10.597.350.675	Synkinesis
-	C10.597.350.700	Tics
-	C10.597.350.850	Tremor
-	C10.597.404	Gait Disorders, Neurologic
-	C10.597.404.400	Gait Apraxia
-	C10.597.404.450	Gait Ataxia
-	C10.597.544	Meningism
-	C10.597.606	Neurobehavioral Manifestations
-	C10.597.606.057	Anhedonia
-	C10.597.606.115	Catatonia

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C10.597.606.150 Communication Disorders
-	C10.597.606.150.500 Language Disorders
-	C10.597.606.150.500.050 Agraphia
-	C10.597.606.150.500.090 Anomia
-	C10.597.606.150.500.300 Dyslexia
-	C10.597.606.150.500.300.200 Dyslexia, Acquired
-	C10.597.606.150.500.300.200.100 Alexia, Pure
-	C10.597.606.150.500.550 Language Development Disorders
-	C10.597.606.150.500.800 Speech Disorders
-	C10.597.606.150.500.800.100 Aphasia
-	C10.597.606.150.500.800.100.100 Aphasia, Broca
-	C10.597.606.150.500.800.100.111 Aphasia, Conduction
-	C10.597.606.150.500.800.100.155 Aphasia, Primary Progressive
-	C10.597.606.150.500.800.100.155.600 Primary Progressive Nonfluent Aphasia
-	C10.597.606.150.500.800.100.166 Aphasia, Wernicke
-	C10.597.606.150.500.800.150 Articulation Disorders
-	C10.597.606.150.500.800.150.200 Dysarthria
-	C10.597.606.150.500.800.300 Echolalia
-	C10.597.606.150.500.800.500 Mutism
-	C10.597.606.150.500.800.750 Stuttering
-	C10.597.606.150.550 Learning Disorders
-	C10.597.606.150.550.099 Dyscalculia
-	C10.597.606.150.550.200 Dyslexia
-	C10.597.606.150.550.200.500 Dyslexia, Acquired
-	C10.597.606.337 Confusion
-	C10.597.606.337.500 Delirium
New Heading	<b>C10.597.606.337.500.500</b> <b>Emergence Delirium</b>
-	C10.597.606.358 Consciousness Disorders
-	C10.597.606.358.800 Unconsciousness
-	C10.597.606.358.800.200 Coma
-	C10.597.606.358.800.200.100 Brain Death
-	C10.597.606.358.800.200.200 Coma, Post-Head Injury
-	C10.597.606.358.800.200.600 Insulin Coma
-	C10.597.606.358.800.400 Persistent Vegetative State
-	C10.597.606.358.800.500 Stupor

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C10.597.606.358.800.600                      Syncope
-	C10.597.606.358.800.600.500                      Syncope, Vasovagal
-	C10.597.606.441                      Lethargy
-	C10.597.606.525                      Memory Disorders
-	C10.597.606.525.100                      Amnesia
-	C10.597.606.525.100.075                      Amnesia, Anterograde
-	C10.597.606.525.100.150                      Amnesia, Retrograde
-	C10.597.606.525.100.800                      Amnesia, Transient Global
-	C10.597.606.525.400                      Korsakoff Syndrome
-	C10.597.606.643                      Intellectual Disability
-	C10.597.606.643.180                      Cri-du-Chat Syndrome
-	C10.597.606.643.210                      De Lange Syndrome
-	C10.597.606.643.220                      Down Syndrome
-	C10.597.606.643.455                      Mental Retardation, X-Linked
-	C10.597.606.643.455.124                      Adrenoleukodystrophy
-	C10.597.606.643.455.249                      Coffin-Lowry Syndrome
-	C10.597.606.643.455.500                      Fragile X Syndrome
-	C10.597.606.643.455.562                      Glycogen Storage Disease Type IIb
-	C10.597.606.643.455.625                      Lesch-Nyhan Syndrome
-	C10.597.606.643.455.687                      Menkes Kinky Hair Syndrome
-	C10.597.606.643.455.750                      Mucopolysaccharidosis II
-	C10.597.606.643.455.875                      Pyruvate Dehydrogenase Complex Deficiency Disease
-	C10.597.606.643.455.937                      Rett Syndrome
-	C10.597.606.643.690                      Prader-Willi Syndrome
-	C10.597.606.643.700                      Rubinstein-Taybi Syndrome
-	C10.597.606.643.969                      WAGR Syndrome
-	C10.597.606.643.970                      Williams Syndrome
-	C10.597.606.762                      Perceptual Disorders
-	C10.597.606.762.100                      Agnosia
-	C10.597.606.762.100.300                      Gerstmann Syndrome
-	C10.597.606.762.100.650                      Prosopagnosia
-	C10.597.606.762.150                      Alice in Wonderland Syndrome
-	C10.597.606.762.175                      Allesthesia
-	C10.597.606.762.200                      Auditory Perceptual Disorders
-	C10.597.606.762.300                      Hallucinations

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C10.597.606.762.400                      Illusions
-	C10.597.606.762.700                      Phantom Limb
-	C10.597.606.881                            Psychomotor Disorders
-	C10.597.606.881.350                      Apraxias
-	C10.597.606.881.350.099                   Alien Hand Syndrome
-	C10.597.606.881.350.200                   Apraxia, Ideomotor
-	C10.597.606.881.350.600                   Gait Apraxia
-	C10.597.606.881.700                      Psychomotor Agitation
-	C10.597.609                                Neurogenic Inflammation
-	C10.597.613                                Neuromuscular Manifestations
-	C10.597.613.250                            Fasciculation
-	C10.597.613.500                            Muscle Cramp
-	C10.597.613.550                            Muscle Hypertonia
-	C10.597.613.550.500                      Muscle Rigidity
-	C10.597.613.550.550                      Muscle Spasticity
-	C10.597.613.575                            Muscle Hypotonia
-	C10.597.613.593                            Muscle Weakness
-	C10.597.613.612                            Muscular Atrophy
-	C10.597.613.612.500                      Sarcopenia
-	C10.597.613.650                            Myokymia
-	C10.597.613.700                            Myotonia
-	C10.597.613.750                            Spasm
-	C10.597.613.750.400                      Hemifacial Spasm
-	C10.597.613.750.700                      Trismus
-	C10.597.613.875                            Tetany
-	C10.597.617                                Pain
-	C10.597.617.044                            Abdominal Pain
-	C10.597.617.044.200                      Abdomen, Acute
-	C10.597.617.088                            Acute Pain
-	C10.597.617.133                            Arthralgia
-	C10.597.617.133.700                      Shoulder Pain
-	C10.597.617.140                            Back Pain
-	C10.597.617.140.199                      Failed Back Surgery Syndrome
-	C10.597.617.140.400                      Low Back Pain
-	C10.597.617.178                            Breakthrough Pain
New	<b>C10.597.617.185                            Cancer Pain</b>

## MeSH Tree Changes for 2017

Type	Tree - heading	
Heading		
-	C10.597.617.192	Chest Pain
-	C10.597.617.192.500	Angina Pectoris
-	C10.597.617.192.500.150	Angina, Unstable
-	C10.597.617.192.500.150.150	Angina Pectoris, Variant
-	C10.597.617.192.500.575	Angina, Stable
-	C10.597.617.196	Chronic Pain
-	C10.597.617.199	Earache
-	C10.597.617.202	Eye Pain
-	C10.597.617.203	Facial Pain
-	C10.597.617.203.500	Toothache
-	C10.597.617.204	Flank Pain
-	C10.597.617.217	Glossalgia
Old Tree	<del>C10.597.617.229</del>	<del>Mastodynia</del>
Old Tree	<del>C10.597.617.231</del>	<del>Musculoskeletal Pain</del>
Old Tree	<del>C10.597.617.231.249</del>	<del>Myalgia</del>
Old Tree	<del>C10.597.617.231.500</del>	<del>Pelvic Girdle Pain</del>
-	C10.597.617.470	Headache
-	C10.597.617.470.800	Slit Ventricle Syndrome
-	C10.597.617.515	Labor Pain
New Tree	<b>C10.597.617.550</b>	<b>Mastodynia</b>
-	C10.597.617.560	Metatarsalgia
New Heading	<b>C10.597.617.560.500</b>	<b>Morton Neuroma</b>
New Tree	<b>C10.597.617.570</b>	<b>Musculoskeletal Pain</b>
New Tree	<b>C10.597.617.570.249</b>	<b>Myalgia</b>
New Tree	<b>C10.597.617.570.500</b>	<b>Pelvic Girdle Pain</b>
-	C10.597.617.576	Neck Pain
-	C10.597.617.682	Neuralgia
New Heading	<b>C10.597.617.682.275</b>	<b>Morton Neuroma</b>
-	C10.597.617.682.550	Neuralgia, Postherpetic
-	C10.597.617.682.675	Piriformis Muscle Syndrome
-	C10.597.617.682.737	Pudendal Neuralgia

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C10.597.617.682.800                      Sciatica
-	C10.597.617.735                              Nociceptive Pain
-	C10.597.617.735.500                        Visceral Pain
-	C10.597.617.788                              Pain, Intractable
-	C10.597.617.841                              Pain, Postoperative
-	C10.597.617.841.500                        Phantom Limb
-	C10.597.617.894                              Pain, Referred
-	C10.597.617.947                              Renal Colic
-	C10.597.622                                    Paralysis
-	C10.597.622.214                              Facial Paralysis
-	C10.597.622.295                              Hemiplegia
-	C10.597.622.447                              Ophthalmoplegia
-	C10.597.622.447.511                        Ophthalmoplegia, Chronic Progressive External
-	C10.597.622.447.511.500                    Kearns-Sayre Syndrome
-	C10.597.622.447.600                        Ophthalmoplegic Migraine
-	C10.597.622.447.690                        Supranuclear Palsy, Progressive
-	C10.597.622.669                              Paraplegia
-	C10.597.622.669.300                        Brown-Sequard Syndrome
-	C10.597.622.714                              Pseudobulbar Palsy
-	C10.597.622.760                              Quadriplegia
-	C10.597.622.812                              Respiratory Paralysis
-	C10.597.622.943                              Vocal Cord Paralysis
-	C10.597.636                                    Paresis
-	C10.597.636.500                              Paraparesis
-	C10.597.636.500.500                        Paraparesis, Spastic
-	C10.597.690                                    Pupil Disorders
-	C10.597.690.150                              Anisocoria
-	C10.597.690.362                              Miosis
-	C10.597.690.362.500                        Horner Syndrome
-	C10.597.690.575                              Tonic Pupil
-	C10.597.704                                    Reflex, Abnormal
-	C10.597.742                                    Seizures
-	C10.597.742.143                              Alcohol Withdrawal Seizures
-	C10.597.751                                    Sensation Disorders
-	C10.597.751.237                              Dizziness
-	C10.597.751.418                              Hearing Disorders

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C10.597.751.418.341                      Hearing Loss
-	C10.597.751.418.341.186                      Deafness
-	C10.597.751.418.341.186.500                      Deaf-Blind Disorders
-	C10.597.751.418.341.186.500.500                      Usher Syndromes
-	C10.597.751.418.341.186.500.750                      Wolfram Syndrome
-	C10.597.751.418.341.374                      Hearing Loss, Bilateral
-	C10.597.751.418.341.562                      Hearing Loss, Conductive
-	C10.597.751.418.341.750                      Hearing Loss, Functional
-	C10.597.751.418.341.812                      Hearing Loss, High-Frequency
-	C10.597.751.418.341.849                      Hearing Loss, Mixed Conductive-Sensorineural
-	C10.597.751.418.341.887                      Hearing Loss, Sensorineural
-	C10.597.751.418.341.887.432                      Hearing Loss, Central
-	C10.597.751.418.341.887.460                      Hearing Loss, Noise-Induced
-	C10.597.751.418.341.887.772                      Presbycusis
-	C10.597.751.418.341.887.886                      Usher Syndromes
-	C10.597.751.418.341.900                      Hearing Loss, Sudden
-	C10.597.751.418.341.950                      Hearing Loss, Unilateral
-	C10.597.751.418.505                      Hyperacusis
-	C10.597.751.418.670                      Tinnitus
-	C10.597.751.600                      Olfaction Disorders
-	C10.597.751.791                      Somatosensory Disorders
-	C10.597.751.791.400                      Hyperalgesia
-	C10.597.751.791.450                      Hyperesthesia
-	C10.597.751.791.500                      Hypesthesia
-	C10.597.751.791.875                      Paresthesia
-	C10.597.751.861                      Taste Disorders
-	C10.597.751.861.184                      Ageusia
-	C10.597.751.861.479                      Dysgeusia
-	C10.597.751.941                      Vision Disorders
-	C10.597.751.941.036                      Alice in Wonderland Syndrome
-	C10.597.751.941.073                      Amblyopia
-	C10.597.751.941.162                      Blindness
-	C10.597.751.941.162.125                      Amaurosis Fugax
-	C10.597.751.941.162.250                      Blindness, Cortical
-	C10.597.751.941.162.625                      Deaf-Blind Disorders
-	C10.597.751.941.162.625.500                      Usher Syndromes

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C10.597.751.941.162.625.750 Wolfram Syndrome
-	C10.597.751.941.256 Color Vision Defects
-	C10.597.751.941.339 Diplopia
-	C10.597.751.941.512 Hemianopsia
-	C10.597.751.941.661 Photophobia
-	C10.597.751.941.811 Scotoma
-	C10.597.751.941.905 Vision, Low
-	C10.597.825 Susac Syndrome
-	C10.597.900 Urinary Bladder, Neurogenic
-	C10.597.951 Vertigo
-	C10.597.951.500 Benign Paroxysmal Positional Vertigo
-	C10.597.975 Voice Disorders
-	C10.597.975.100 Aphonia
-	C10.597.975.325 Dysphonia
-	C10.597.975.550 Hoarseness
-	C10.668 Neuromuscular Diseases
-	C10.668.364 Fatigue Syndrome, Chronic
-	C10.668.467 Motor Neuron Disease
-	C10.668.467.250 Amyotrophic Lateral Sclerosis
-	C10.668.467.300 Bulbar Palsy, Progressive
-	C10.668.467.500 Muscular Atrophy, Spinal
-	C10.668.467.500.186 Bulbo-Spinal Atrophy, X-Linked
-	C10.668.467.500.750 Spinal Muscular Atrophies of Childhood
-	C10.668.491 Muscular Diseases
-	C10.668.491.087 Medial Tibial Stress Syndrome
-	C10.668.491.175 Muscular Disorders, Atrophic
-	C10.668.491.175.500 Muscular Dystrophies
-	C10.668.491.175.500.074 Distal Myopathies
-	C10.668.491.175.500.112 Glycogen Storage Disease Type VII
-	C10.668.491.175.500.149 Muscular Dystrophies, Limb-Girdle
-	C10.668.491.175.500.149.500 Sarcoglycanopathies
-	C10.668.491.175.500.300 Muscular Dystrophy, Duchenne
-	C10.668.491.175.500.350 Muscular Dystrophy, Emery-Dreifuss
-	C10.668.491.175.500.400 Muscular Dystrophy, Facioscapulohumeral
-	C10.668.491.175.500.450 Muscular Dystrophy, Oculopharyngeal
-	C10.668.491.175.500.500 Myotonic Dystrophy



## MeSH Tree Changes for 2017

Type	Tree - heading
-	C10.668.491.175.750 Postpoliomyelitis Syndrome
-	C10.668.491.387 Eosinophilia-Myalgia Syndrome
-	C10.668.491.425 Fibromyalgia
-	C10.668.491.500 Mitochondrial Myopathies
-	C10.668.491.500.500 Mitochondrial Encephalomyopathies
-	C10.668.491.500.500.500 MELAS Syndrome
-	C10.668.491.500.500.550 MERRF Syndrome
-	C10.668.491.500.700 Ophthalmoplegia, Chronic Progressive External
-	C10.668.491.500.700.500 Kearns-Sayre Syndrome
-	C10.668.491.525 Myalgia
-	C10.668.491.550 Myopathies, Structural, Congenital
-	C10.668.491.550.290 Myopathies, Nemaline
-	C10.668.491.550.300 Myopathy, Central Core
-	C10.668.491.562 Myositis
-	C10.668.491.562.500 Myositis, Inclusion Body
-	C10.668.491.562.537 Orbital Myositis
-	C10.668.491.562.575 Polymyositis
-	C10.668.491.562.575.500 Dermatomyositis
-	C10.668.491.562.787 Pyomyositis
-	C10.668.491.606 Myotonic Disorders
-	C10.668.491.606.500 Myotonia Congenita
-	C10.668.491.606.750 Myotonic Dystrophy
-	C10.668.491.650 Paralyses, Familial Periodic
-	C10.668.491.650.450 Hypokalemic Periodic Paralysis
-	C10.668.491.650.600 Paralysis, Hyperkalemic Periodic
-	C10.668.758 Neuromuscular Junction Diseases
-	C10.668.758.200 Botulism
-	C10.668.758.450 Lambert-Eaton Myasthenic Syndrome
-	C10.668.758.725 Myasthenia Gravis
-	C10.668.758.725.300 Myasthenia Gravis, Autoimmune, Experimental
-	C10.668.758.725.650 Myasthenia Gravis, Neonatal
-	C10.668.758.800 Myasthenic Syndromes, Congenital
-	C10.668.829 Peripheral Nervous System Diseases
-	C10.668.829.025 Acrodynia
-	C10.668.829.050 Amyloid Neuropathies
-	C10.668.829.050.050 Amyloid Neuropathies, Familial

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C10.668.829.100      Brachial Plexus Neuropathies
-	C10.668.829.100.500      Brachial Plexus Neuritis
-	C10.668.829.250      Complex Regional Pain Syndromes
-	C10.668.829.250.200      Causalgia
-	C10.668.829.250.800      Reflex Sympathetic Dystrophy
-	C10.668.829.300      Diabetic Neuropathies
-	C10.668.829.325      Giant Axonal Neuropathy
-	C10.668.829.350      Guillain-Barre Syndrome
-	C10.668.829.350.500      Miller Fisher Syndrome
-	C10.668.829.380      Hand-Arm Vibration Syndrome
-	C10.668.829.425      Isaacs Syndrome
-	C10.668.829.500      Mononeuropathies
-	C10.668.829.500.200      Femoral Neuropathy
-	C10.668.829.500.500      Median Neuropathy
-	C10.668.829.500.500.200      Carpal Tunnel Syndrome
-	C10.668.829.500.600      Peroneal Neuropathies
-	C10.668.829.500.650      Radial Neuropathy
-	C10.668.829.500.675      Sciatic Neuropathy
-	C10.668.829.500.675.399      Piriformis Muscle Syndrome
-	C10.668.829.500.675.800      Sciatica
-	C10.668.829.500.700      Tibial Neuropathy
-	C10.668.829.500.700.800      Tarsal Tunnel Syndrome
-	C10.668.829.500.850      Ulnar Neuropathies
-	C10.668.829.500.850.200      Cubital Tunnel Syndrome
-	C10.668.829.500.850.600      Ulnar Nerve Compression Syndromes
-	C10.668.829.550      Nerve Compression Syndromes
-	C10.668.829.550.200      Carpal Tunnel Syndrome
-	C10.668.829.550.500      Piriformis Muscle Syndrome
-	C10.668.829.550.650      Pudendal Neuralgia
-	C10.668.829.550.800      Tarsal Tunnel Syndrome
-	C10.668.829.550.850      Thoracic Outlet Syndrome
-	C10.668.829.550.850.200      Cervical Rib Syndrome
-	C10.668.829.550.925      Ulnar Nerve Compression Syndromes
-	C10.668.829.550.925.200      Cubital Tunnel Syndrome
-	C10.668.829.600      Neuralgia
-	C10.668.829.600.200      Causalgia

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Heading	<b>C10.668.829.600.375</b>	<b>Morton Neuroma</b>
-	C10.668.829.600.550	Neuralgia, Postherpetic
-	C10.668.829.600.675	Piriformis Muscle Syndrome
-	C10.668.829.600.737	Pudendal Neuralgia
-	C10.668.829.600.800	Sciatica
-	C10.668.829.650	Neuritis
-	C10.668.829.650.250	Brachial Plexus Neuritis
-	C10.668.829.650.500	Neuritis, Autoimmune, Experimental
-	C10.668.829.675	Neurofibromatosis 1
-	C10.668.829.700	Pain Insensitivity, Congenital
-	C10.668.829.712	Peripheral Nerve Injuries
-	C10.668.829.725	Peripheral Nervous System Neoplasms
-	C10.668.829.725.500	Nerve Sheath Neoplasms
-	C10.668.829.725.500.600	Neurofibroma
-	C10.668.829.725.500.600.500	Neurofibroma, Plexiform
-	C10.668.829.725.500.600.600	Neurofibrosarcoma
-	C10.668.829.800	Polyneuropathies
-	C10.668.829.800.050	Alcoholic Neuropathy
-	C10.668.829.800.175	Hereditary Sensory and Autonomic Neuropathies
-	C10.668.829.800.175.250	Dysautonomia, Familial
-	C10.668.829.800.300	Hereditary Sensory and Motor Neuropathy
-	C10.668.829.800.300.099	Alstrom Syndrome
-	C10.668.829.800.300.200	Charcot-Marie-Tooth Disease
-	C10.668.829.800.300.490	Giant Axonal Neuropathy
-	C10.668.829.800.300.780	Refsum Disease
-	C10.668.829.800.300.820	Spastic Paraplegia, Hereditary
-	C10.668.829.800.662	Paraneoplastic Polyneuropathy
-	C10.668.829.800.700	POEMS Syndrome
-	C10.668.829.800.750	Polyradiculoneuropathy
-	C10.668.829.800.750.300	Guillain-Barre Syndrome
-	C10.668.829.800.750.300.500	Miller Fisher Syndrome
-	C10.668.829.800.750.600 Demyelinating	Polyradiculoneuropathy, Chronic Inflammatory
-	C10.668.829.800.750.700	Polyradiculopathy
-	C10.668.829.800.875	Tangier Disease

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	C10.668.829.820	Radiculopathy
New Heading	<b>C10.668.829.860</b>	<b>Small Fiber Neuropathy</b>
-	C10.668.829.900	Tarlov Cysts
-	C10.668.864	Poliomyelitis
-	C10.668.864.250	Poliomyelitis, Bulbar
-	C10.668.864.500	Postpoliomyelitis Syndrome
-	C10.668.900	Stiff-Person Syndrome
-	C10.720	Neurotoxicity Syndromes
-	C10.720.075	Akathisia, Drug-Induced
-	C10.720.112	Alcohol-Induced Disorders, Nervous System
-	C10.720.112.100	Alcohol Amnestic Disorder
-	C10.720.112.100.500	Korsakoff Syndrome
-	C10.720.112.200	Alcohol Withdrawal Delirium
-	C10.720.112.300	Alcohol Withdrawal Seizures
-	C10.720.112.400	Alcoholic Neuropathy
-	C10.720.150	Botulism
-	C10.720.312	Dyskinesia, Drug-Induced
-	C10.720.475	Heavy Metal Poisoning, Nervous System
-	C10.720.475.150	Arsenic Poisoning
-	C10.720.475.400	Lead Poisoning, Nervous System
-	C10.720.475.400.350	Lead Poisoning, Nervous System, Adult
-	C10.720.475.400.700	Lead Poisoning, Nervous System, Childhood
-	C10.720.475.500	Manganese Poisoning
-	C10.720.475.600	Mercury Poisoning, Nervous System
-	C10.720.475.600.150	Acrodynia
-	C10.720.606	MPTP Poisoning
-	C10.720.737	Neuroleptic Malignant Syndrome
-	C10.803	Restless Legs Syndrome
-	C10.886	Sleep Wake Disorders
-	C10.886.425	Dyssomnias
-	C10.886.425.175	Sleep Deprivation
-	C10.886.425.200	Sleep Disorders, Circadian Rhythm
-	C10.886.425.200.500	Jet Lag Syndrome
-	C10.886.425.800	Sleep Disorders, Intrinsic
-	C10.886.425.800.200	Disorders of Excessive Somnolence

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	C10.886.425.800.200.400	Hypersomnolence, Idiopathic
-	C10.886.425.800.200.500	Kleine-Levin Syndrome
-	C10.886.425.800.200.750	Narcolepsy
-	C10.886.425.800.200.750.500	Cataplexy
-	C10.886.425.800.600	Nocturnal Myoclonus Syndrome
-	C10.886.425.800.700	Restless Legs Syndrome
-	C10.886.425.800.750	Sleep Apnea Syndromes
-	C10.886.425.800.750.800	Sleep Apnea, Central
-	C10.886.425.800.750.850	Sleep Apnea, Obstructive
-	C10.886.425.800.750.850.500	Obesity Hypoventilation Syndrome
-	C10.886.425.800.800	Sleep Initiation and Maintenance Disorders
-	C10.886.425.800.800.400	Insomnia, Fatal Familial
-	C10.886.659	Parasomnias
-	C10.886.659.618	Nocturnal Myoclonus Syndrome
-	C10.886.659.627	Nocturnal Paroxysmal Dystonia
-	C10.886.659.633	REM Sleep Parasomnias
-	C10.886.659.633.700	REM Sleep Behavior Disorder
-	C10.886.659.633.800	Sleep Paralysis
-	C10.886.659.634	Restless Legs Syndrome
-	C10.886.659.635	Sleep Arousal Disorders
-	C10.886.659.635.600	Night Terrors
-	C10.886.659.635.700	Somnambulism
-	C10.886.659.637	Sleep Bruxism
-	C10.886.659.700	Sleep-Wake Transition Disorders
-	C10.900	Trauma, Nervous System
-	C10.900.250	Cerebrovascular Trauma
-	C10.900.250.300	Carotid Artery Injuries
-	C10.900.250.300.300	Carotid Artery, Internal, Dissection
-	C10.900.250.300.400	Carotid-Cavernous Sinus Fistula
-	C10.900.250.650	Vertebral Artery Dissection
-	C10.900.300	Craniocerebral Trauma
-	C10.900.300.087	Brain Injuries
Old Tree	<b>C10.900.300.087.125</b>	<b>Brain Concussion</b>
Old Tree	<b>C10.900.300.087.125.249</b>	<b>Contrecoup Injury</b>
Old Tree	<b>C10.900.300.087.125.500</b>	<b>Post-Concussion Syndrome</b>
-	C10.900.300.087.187	Brain Hemorrhage, Traumatic

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	C10.900.300.087.187.200	Brain Stem Hemorrhage, Traumatic
-	C10.900.300.087.187.300	Cerebral Hemorrhage, Traumatic
New Heading	<b>C10.900.300.087.219</b>	<b>Brain Injuries, Diffuse</b>
New Tree	<a href="#">C10.900.300.087.219.500</a>	<a href="#">Diffuse Axonal Injury</a>
New Heading	<b>C10.900.300.087.235</b>	<b>Brain Injuries, Traumatic</b>
New Tree	<a href="#">C10.900.300.087.235.250</a>	<a href="#">Brain Concussion</a>
New Heading	<b>C10.900.300.087.235.375</b>	<b>Brain Contusion</b>
New Heading	<b>C10.900.300.087.235.500</b>	<b>Chronic Traumatic Encephalopathy</b>
-	C10.900.300.087.250	Brain Injury, Chronic
New Heading	<b>C10.900.300.087.250.500</b>	<b>Chronic Traumatic Encephalopathy</b>
Old Tree	<del>C10.900.300.087.500</del>	<del>Diffuse Axonal Injury</del>
-	C10.900.300.087.600	Epilepsy, Post-Traumatic
-	C10.900.300.087.700	Pneumocephalus
-	C10.900.300.087.850	Shaken Baby Syndrome
-	C10.900.300.109	Cerebrospinal Fluid Leak
-	C10.900.300.109.500	Cerebrospinal Fluid Otorrhea
-	C10.900.300.109.750	Cerebrospinal Fluid Rhinorrhea
-	C10.900.300.175	Coma, Post-Head Injury
-	C10.900.300.218	Cranial Nerve Injuries
-	C10.900.300.218.150	Abducens Nerve Injury
-	C10.900.300.218.225	Accessory Nerve Injuries
-	C10.900.300.218.300	Facial Nerve Injuries
-	C10.900.300.218.331	Glossopharyngeal Nerve Injuries
-	C10.900.300.218.362	Hypoglossal Nerve Injuries
-	C10.900.300.218.456	Oculomotor Nerve Injuries
-	C10.900.300.218.487	Olfactory Nerve Injuries
-	C10.900.300.218.550	Optic Nerve Injuries
-	C10.900.300.218.775	Trigeminal Nerve Injuries
-	C10.900.300.218.775.500	Lingual Nerve Injuries
-	C10.900.300.218.831	Trochlear Nerve Injuries
-	C10.900.300.218.887	Vagus Nerve Injuries

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C10.900.300.218.887.750 Laryngeal Nerve Injuries
-	C10.900.300.218.887.750.500 Recurrent Laryngeal Nerve Injuries
-	C10.900.300.218.943 Vestibulocochlear Nerve Injuries
-	C10.900.300.284 Facial Injuries
-	C10.900.300.284.250 Eye Injuries
-	C10.900.300.284.250.124 Corneal Injuries
-	C10.900.300.284.250.124.500 Corneal Perforation
-	C10.900.300.284.250.250 Eye Burns
-	C10.900.300.284.250.260 Eye Foreign Bodies
-	C10.900.300.284.250.270 Eye Injuries, Penetrating
-	C10.900.300.284.500 Maxillofacial Injuries
-	C10.900.300.284.500.400 Jaw Fractures
-	C10.900.300.284.500.400.255 Mandibular Fractures
-	C10.900.300.284.500.400.510 Maxillary Fractures
-	C10.900.300.284.500.500 Mandibular Injuries
-	C10.900.300.284.500.550 Orbital Fractures
-	C10.900.300.284.500.950 Zygomatic Fractures
-	C10.900.300.350 Head Injuries, Closed
-	C10.900.300.350.300 Brain Concussion
-	C10.900.300.350.300.250 Contrecoup Injury
-	C10.900.300.350.300.500 Post-Concussion Syndrome
-	C10.900.300.675 Head Injuries, Penetrating
-	C10.900.300.837 Intracranial Hemorrhage, Traumatic
-	C10.900.300.837.150 Brain Hemorrhage, Traumatic
-	C10.900.300.837.150.300 Brain Stem Hemorrhage, Traumatic
-	C10.900.300.837.150.650 Cerebral Hemorrhage, Traumatic
-	C10.900.300.837.300 Hematoma, Epidural, Cranial
-	C10.900.300.837.600 Hematoma, Subdural
-	C10.900.300.837.600.050 Hematoma, Subdural, Acute
-	C10.900.300.837.600.120 Hematoma, Subdural, Chronic
-	C10.900.300.837.600.400 Hematoma, Subdural, Intracranial
-	C10.900.300.837.800 Subarachnoid Hemorrhage, Traumatic
-	C10.900.300.918 Skull Fractures
-	C10.900.300.918.150 Skull Fracture, Basilar
-	C10.900.300.918.300 Skull Fracture, Depressed
-	C10.900.575 Peripheral Nerve Injuries

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C10.900.850 Spinal Cord Injuries
-	C10.900.850.250 Autonomic Dysreflexia
-	C10.900.850.625 Central Cord Syndrome
-	C11 Eye Diseases
-	C11.093 Asthenopia
-	C11.180 Cogan Syndrome
-	C11.187 Conjunctival Diseases
-	C11.187.169 Conjunctival Neoplasms
-	C11.187.183 Conjunctivitis
-	C11.187.183.200 Conjunctivitis, Allergic
-	C11.187.183.220 Conjunctivitis, Bacterial
-	C11.187.183.220.250 Conjunctivitis, Inclusion
-	C11.187.183.220.538 Ophthalmia Neonatorum
-	C11.187.183.220.889 Trachoma
-	C11.187.183.240 Conjunctivitis, Viral
-	C11.187.183.240.216 Conjunctivitis, Acute Hemorrhagic
-	C11.187.183.394 Keratoconjunctivitis
-	C11.187.183.394.520 Keratoconjunctivitis, Infectious
-	C11.187.183.394.550 Keratoconjunctivitis Sicca
-	C11.187.482 Pemphigoid, Benign Mucous Membrane
-	C11.187.631 Pinguecula
-	C11.187.781 Pterygium
-	C11.187.810 Xerophthalmia
-	C11.204 Corneal Diseases
-	C11.204.236 Corneal Dystrophies, Hereditary
-	C11.204.236.218 Corneal Dystrophy, Juvenile Epithelial of Meesmann
-	C11.204.236.438 Fuchs' Endothelial Dystrophy
-	C11.204.267 Corneal Edema
-	C11.204.278 Corneal Endothelial Cell Loss
-	C11.204.284 Corneal Injuries
-	C11.204.284.500 Corneal Perforation
-	C11.204.290 Corneal Neovascularization
-	C11.204.299 Corneal Opacity
-	C11.204.299.070 Arcus Senilis
-	C11.204.431 Corneal Wavefront Aberration
-	C11.204.497 Iridocorneal Endothelial Syndrome



## MeSH Tree Changes for 2017

Type	Tree - heading
-	C11.204.564 Keratitis
-	C11.204.564.112 Acanthamoeba Keratitis
-	C11.204.564.225 Corneal Ulcer
-	C11.204.564.425 Keratitis, Herpetic
-	C11.204.564.425.450 Keratitis, Dendritic
-	C11.204.564.585 Keratoconjunctivitis
-	C11.204.564.585.500 Keratoconjunctivitis, Infectious
-	C11.204.564.585.630 Keratoconjunctivitis Sicca
-	C11.204.627 Keratoconus
-	C11.204.813 Trachoma
-	C11.250 Eye Abnormalities
-	C11.250.060 Aniridia
-	C11.250.060.950 WAGR Syndrome
-	C11.250.080 Anophthalmos
-	C11.250.090 Blepharophimosis
-	C11.250.110 Coloboma
-	C11.250.300 Ectopia Lentis
-	C11.250.390 Fraser Syndrome
-	C11.250.480 Hydrophthalmos
-	C11.250.566 Microphthalmos
-	C11.250.616 Persistent Hyperplastic Primary Vitreous
-	C11.250.666 Retinal Dysplasia
-	C11.270 Eye Diseases, Hereditary
-	C11.270.019 Aicardi Syndrome
-	C11.270.040 Albinism
-	C11.270.040.090 Albinism, Ocular
-	C11.270.040.545 Albinism, Oculocutaneous
-	C11.270.040.545.400 Hermanski-Pudlak Syndrome
-	C11.270.060 Aniridia
-	C11.270.060.950 WAGR Syndrome
-	C11.270.142 Choroideremia
New Heading	<b>C11.270.152 Cone-Rod Dystrophies</b>
-	C11.270.162 Corneal Dystrophies, Hereditary
-	C11.270.162.218 Corneal Dystrophy, Juvenile Epithelial of Meesmann
-	C11.270.162.438 Fuchs' Endothelial Dystrophy

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C11.270.235 Duane Retraction Syndrome
-	C11.270.468 Gyrate Atrophy
-	C11.270.516 Leber Congenital Amaurosis
-	C11.270.564 Optic Atrophies, Hereditary
-	C11.270.564.100 Optic Atrophy, Autosomal Dominant
-	C11.270.564.400 Optic Atrophy, Hereditary, Leber
-	C11.270.564.980 Wolfram Syndrome
-	C11.270.660 Retinal Dysplasia
-	C11.270.684 Retinitis Pigmentosa
-	C11.270.684.249 Alstrom Syndrome
-	C11.270.842 Graves Ophthalmopathy
-	C11.270.881 Walker-Warburg Syndrome
-	C11.270.921 Weill-Marchesani Syndrome
-	C11.290 Eye Hemorrhage
-	C11.290.195 Choroid Hemorrhage
-	C11.290.484 Hyphema
-	C11.290.807 Retinal Hemorrhage
-	C11.290.960 Vitreous Hemorrhage
-	C11.294 Eye Infections
-	C11.294.177 Corneal Ulcer
-	C11.294.265 Endophthalmitis
-	C11.294.265.500 Uveitis, Suppurative
-	C11.294.354 Eye Infections, Bacterial
-	C11.294.354.220 Conjunctivitis, Bacterial
-	C11.294.354.220.250 Conjunctivitis, Inclusion
-	C11.294.354.220.625 Ophthalmia Neonatorum
-	C11.294.354.220.800 Trachoma
-	C11.294.354.400 Hordeolum
-	C11.294.354.450 Keratoconjunctivitis, Infectious
-	C11.294.354.800 Tuberculosis, Ocular
-	C11.294.354.900 Uveitis, Suppurative
-	C11.294.354.900.675 Panophthalmitis
-	C11.294.450 Eye Infections, Fungal
-	C11.294.450.900 Uveitis, Suppurative
-	C11.294.450.900.675 Panophthalmitis
-	C11.294.725 Eye Infections, Parasitic

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C11.294.725.125 Acanthamoeba Keratitis
-	C11.294.725.562 Onchocerciasis, Ocular
-	C11.294.725.781 Toxoplasmosis, Ocular
-	C11.294.800 Eye Infections, Viral
-	C11.294.800.250 Conjunctivitis, Viral
-	C11.294.800.250.250 Conjunctivitis, Acute Hemorrhagic
-	C11.294.800.270 Cytomegalovirus Retinitis
-	C11.294.800.450 Herpes Zoster Ophthalmicus
-	C11.294.800.475 Keratitis, Herpetic
-	C11.294.800.475.450 Keratitis, Dendritic
-	C11.297 Eye Injuries
-	C11.297.249 Anterior Capsular Rupture, Ocular
-	C11.297.374 Corneal Injuries
-	C11.297.374.500 Corneal Perforation
-	C11.297.500 Posterior Capsular Rupture, Ocular
-	C11.300 Eye Manifestations
-	C11.300.500 Eye Pain
-	C11.319 Eye Neoplasms
-	C11.319.217 Conjunctival Neoplasms
-	C11.319.421 Eyelid Neoplasms
-	C11.319.457 Orbital Neoplasms
-	C11.319.466 Paraneoplastic Syndromes, Ocular
-	C11.319.475 Retinal Neoplasms
-	C11.319.475.760 Retinoblastoma
-	C11.319.494 Uveal Neoplasms
-	C11.319.494.198 Choroid Neoplasms
-	C11.319.494.400 Iris Neoplasms
-	C11.338 Eyelid Diseases
-	C11.338.133 Blepharitis
-	C11.338.190 Blepharophimosis
-	C11.338.204 Blepharoptosis
-	C11.338.250 Blepharospasm
-	C11.338.300 Chalazion
-	C11.338.362 Ectropion
-	C11.338.443 Entropion
-	C11.338.526 Eyelid Neoplasms

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C11.338.648                      Hordeolum
-	C11.338.912                      Trichiasis
-	C11.496                              Lacrimal Apparatus Diseases
-	C11.496.221                      Dacryocystitis
-	C11.496.221.500                  Canaliculitis
-	C11.496.260                      Dry Eye Syndromes
-	C11.496.260.394                  Keratoconjunctivitis Sicca
-	C11.496.260.719                  Sjogren's Syndrome
-	C11.496.260.892                  Xerophthalmia
-	C11.496.456                      Lacrimal Duct Obstruction
-	C11.510                              Lens Diseases
-	C11.510.103                      Aphakia
-	C11.510.103.110                  Aphakia, Postcataract
-	C11.510.174                      Artificial Lens Implant Migration
-	C11.510.245                      Cataract
-	C11.510.245.500                  Capsule Opacification
-	C11.510.598                      Lens Subluxation
-	C11.510.598.373                  Ectopia Lentis
-	C11.525                              Ocular Hypertension
-	C11.525.381                      Glaucoma
-	C11.525.381.056                  Glaucoma, Angle-Closure
-	C11.525.381.348                  Glaucoma, Neovascular
-	C11.525.381.407                  Glaucoma, Open-Angle
-	C11.525.381.407.480              Hydrophthalmos
-	C11.540                              Ocular Hypotension
-	C11.590                              Ocular Motility Disorders
-	C11.590.224                      Duane Retraction Syndrome
-	C11.590.312                      Miller Fisher Syndrome
-	C11.590.400                      Nystagmus, Pathologic
-	C11.590.400.300                  Nystagmus, Congenital
-	C11.590.436                      Oculomotor Nerve Diseases
-	C11.590.436.200                  Adie Syndrome
-	C11.590.436.600                  Oculomotor Nerve Injuries
-	C11.590.472                      Ophthalmoplegia
-	C11.590.472.250                  Ophthalmoplegia, Chronic Progressive External
-	C11.590.472.250.500              Kearns-Sayre Syndrome

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C11.590.472.375 Ophthalmoplegic Migraine
-	C11.590.472.500 Supranuclear Palsy, Progressive
-	C11.590.725 Opsoclonus-Myoclonus Syndrome
-	C11.590.810 Strabismus
-	C11.590.810.400 Esotropia
-	C11.590.810.440 Exotropia
-	C11.590.905 Tolosa-Hunt Syndrome
-	C11.640 Optic Nerve Diseases
-	C11.640.225 Low Tension Glaucoma
-	C11.640.451 Optic Atrophy
-	C11.640.451.451 Optic Atrophies, Hereditary
-	C11.640.451.451.100 Optic Atrophy, Autosomal Dominant
-	C11.640.451.451.400 Optic Atrophy, Hereditary, Leber
-	C11.640.451.451.980 Wolfram Syndrome
-	C11.640.513 Optic Disk Drusen
-	C11.640.530 Optic Nerve Injuries
-	C11.640.544 Optic Nerve Neoplasms
-	C11.640.544.500 Optic Nerve Glioma
-	C11.640.576 Optic Neuritis
-	C11.640.576.695 Neuromyelitis Optica
-	C11.640.643 Optic Neuropathy, Ischemic
-	C11.640.710 Papilledema
-	C11.675 Orbital Diseases
-	C11.675.319 Enophthalmos
-	C11.675.349 Exophthalmos
-	C11.675.349.500 Graves Disease
-	C11.675.349.500.500 Graves Ophthalmopathy
-	C11.675.387 Orbital Cellulitis
-	C11.675.426 Orbital Myositis
-	C11.675.659 Orbital Neoplasms
-	C11.675.679 Orbital Pseudotumor
-	C11.675.689 Pott Puffy Tumor
-	C11.675.700 Retrobulbar Hemorrhage
-	C11.710 Pupil Disorders
-	C11.710.090 Anisocoria
-	C11.710.528 Miosis

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C11.710.528.500                      Horner Syndrome
-	C11.710.570                              Mydriasis
-	C11.710.800                              Tonic Pupil
-	C11.710.800.180                      Adie Syndrome
-	C11.744                                      Refractive Errors
-	C11.744.116                              Aniseikonia
-	C11.744.126                              Anisometropia
-	C11.744.212                              Astigmatism
-	C11.744.345                              Corneal Wavefront Aberration
-	C11.744.479                              Hyperopia
-	C11.744.636                              Myopia
-	C11.744.636.500                      Myopia, Degenerative
-	C11.744.786                              Presbyopia
-	C11.768                                      Retinal Diseases
-	C11.768.094                              Angioid Streaks
-	C11.768.175                              Central Serous Chorioretinopathy
-	C11.768.257                              Diabetic Retinopathy
-	C11.768.328                              Epiretinal Membrane
-	C11.768.346                              Hypertensive Retinopathy
-	C11.768.364                              Leber Congenital Amaurosis
-	C11.768.400                              Retinal Artery Occlusion
-	C11.768.400.500                      Susac Syndrome
-	C11.768.585                              Retinal Degeneration
-	C11.768.585.439                      Macular Degeneration
-	C11.768.585.439.122                      Geographic Atrophy
-	C11.768.585.439.245                      Macular Edema
-	C11.768.585.439.433                      Vitelliform Macular Dystrophy
-	C11.768.585.439.622                      Wet Macular Degeneration
-	C11.768.585.585                      Retinal Drusen
-	C11.768.585.658                      Retinal Dystrophies
New Heading	<b>C11.768.585.658.250                      Cone-Rod Dystrophies</b>
-	C11.768.585.658.500                      Retinitis Pigmentosa
-	C11.768.585.658.500.627                      Kearns-Sayre Syndrome
-	C11.768.585.658.500.813                      Usher Syndromes
-	C11.768.585.865                      Retinoschisis

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C11.768.648 Retinal Detachment
-	C11.768.660 Retinal Dysplasia
-	C11.768.710 Retinal Hemorrhage
-	C11.768.717 Retinal Neoplasms
-	C11.768.717.760 Retinoblastoma
-	C11.768.725 Retinal Neovascularization
-	C11.768.740 Retinal Perforations
-	C11.768.748 Retinal Telangiectasis
-	C11.768.757 Retinal Vasculitis
-	C11.768.760 Retinal Vein Occlusion
-	C11.768.773 Retinitis
-	C11.768.773.348 Chorioretinitis
-	C11.768.773.360 Cytomegalovirus Retinitis
-	C11.768.773.674 Retinal Necrosis Syndrome, Acute
-	C11.768.836 Retinopathy of Prematurity
-	C11.768.890 Vitreoretinopathy, Proliferative
-	C11.790 Scleral Diseases
-	C11.790.500 Scleritis
-	C11.941 Uveal Diseases
-	C11.941.160 Choroid Diseases
-	C11.941.160.177 Choroid Hemorrhage
-	C11.941.160.238 Choroid Neoplasms
-	C11.941.160.244 Choroidal Neovascularization
-	C11.941.160.300 Choroideremia
-	C11.941.160.478 Choroiditis
-	C11.941.160.478.400 Chorioretinitis
-	C11.941.160.478.700 Pars Planitis
-	C11.941.160.578 Gyrate Atrophy
-	C11.941.375 Iris Diseases
-	C11.941.375.060 Aniridia
-	C11.941.375.060.950 WAGR Syndrome
-	C11.941.375.285 Exfoliation Syndrome
-	C11.941.375.322 Iridocorneal Endothelial Syndrome
-	C11.941.375.360 Iridocyclitis
-	C11.941.375.375 Iris Neoplasms
-	C11.941.375.385 Iritis

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C11.941.855 Uveal Neoplasms
-	C11.941.855.198 Choroid Neoplasms
-	C11.941.855.400 Iris Neoplasms
-	C11.941.879 Uveitis
-	C11.941.879.780 Panuveitis
-	C11.941.879.780.500 Ophthalmia, Sympathetic
-	C11.941.879.780.880 Uveitis, Anterior
-	C11.941.879.780.880.200 Behcet Syndrome
-	C11.941.879.780.880.400 Iridocyclitis
-	C11.941.879.780.880.448 Iritis
-	C11.941.879.780.900 Uveitis, Posterior
-	C11.941.879.780.900.300 Choroiditis
-	C11.941.879.780.900.300.318 Chorioretinitis
-	C11.941.879.780.900.300.659 Pars Planitis
-	C11.941.879.900 Uveitis, Intermediate
-	C11.941.879.900.500 Pars Planitis
-	C11.941.879.960 Uveitis, Suppurative
-	C11.941.879.960.580 Panophthalmitis
-	C11.941.879.980 Uveomeningoencephalitic Syndrome
-	C11.966 Vision Disorders
-	C11.966.073 Amblyopia
-	C11.966.075 Blindness
-	C11.966.075.125 Amaurosis Fugax
-	C11.966.075.250 Blindness, Cortical
-	C11.966.075.375 Deaf-Blind Disorders
-	C11.966.075.375.500 Usher Syndromes
-	C11.966.075.375.750 Wolfram Syndrome
-	C11.966.075.500 Hemianopsia
-	C11.966.256 Color Vision Defects
-	C11.966.339 Diplopia
-	C11.966.671 Night Blindness
-	C11.966.741 Photophobia
-	C11.966.811 Scotoma
-	C11.966.858 Susac Syndrome
-	C11.966.905 Vision, Low
-	C11.980 Vitreous Detachment



## MeSH Tree Changes for 2017

Type	Tree - heading
-	C12 Male Urogenital Diseases
-	C12.294 Genital Diseases, Male
-	C12.294.199 Epididymitis
-	C12.294.229 Fournier Gangrene
-	C12.294.260 Genital Neoplasms, Male
-	C12.294.260.500 Penile Neoplasms
-	C12.294.260.750 Prostatic Neoplasms
-	C12.294.260.750.500 Prostatic Neoplasms, Castration-Resistant
-	C12.294.260.937 Testicular Neoplasms
-	C12.294.260.937.500 Sertoli-Leydig Cell Tumor
-	C12.294.260.937.500.249 Leydig Cell Tumor
-	C12.294.260.937.500.500 Sertoli Cell Tumor
-	C12.294.287 Hematocele
-	C12.294.293 Hemospermia
-	C12.294.329 Herpes Genitalis
-	C12.294.365 Infertility
-	C12.294.365.700 Infertility, Male
-	C12.294.365.700.126 Aspermia
-	C12.294.365.700.253 Asthenozoospermia
-	C12.294.365.700.380 Azoospermia
-	C12.294.365.700.508 Oligospermia
-	C12.294.365.700.754 Sertoli Cell-Only Syndrome
New Heading	<b>C12.294.365.700.877 Teratozoospermia</b>
-	C12.294.494 Penile Diseases
-	C12.294.494.136 Balanitis
-	C12.294.494.136.500 Balanitis Xerotica Obliterans
-	C12.294.494.400 Hypospadias
-	C12.294.494.508 Penile Induration
-	C12.294.494.591 Penile Neoplasms
-	C12.294.494.684 Phimosis
-	C12.294.494.684.587 Paraphimosis
-	C12.294.494.786 Priapism
-	C12.294.565 Prostatic Diseases
-	C12.294.565.500 Prostatic Hyperplasia
-	C12.294.565.625 Prostatic Neoplasms

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C12.294.565.625.500 Prostatic Neoplasms, Castration-Resistant
-	C12.294.565.750 Prostatitis
-	C12.294.604 Reproductive Tract Infections
-	C12.294.644 Sexual Dysfunction, Physiological
-	C12.294.644.242 Dyspareunia
-	C12.294.644.486 Erectile Dysfunction
-	C12.294.644.486.500 Impotence, Vasculogenic
-	C12.294.644.743 Premature Ejaculation
-	C12.294.668 Sexually Transmitted Diseases
-	C12.294.668.281 Sexually Transmitted Diseases, Bacterial
-	C12.294.668.281.201 Chancroid
-	C12.294.668.281.301 Chlamydia Infections
-	C12.294.668.281.301.490 Lymphogranuloma Venereum
-	C12.294.668.281.401 Gonorrhea
-	C12.294.668.281.451 Granuloma Inguinale
-	C12.294.668.281.859 Syphilis
-	C12.294.693 Spermatic Cord Torsion
-	C12.294.731 Spermatocele
-	C12.294.829 Testicular Diseases
-	C12.294.829.258 Cryptorchidism
-	C12.294.829.493 Orchitis
-	C12.294.882 Testicular Hydrocele
-	C12.294.909 Tuberculosis, Male Genital
-	C12.294.936 Varicocele
-	C12.483 Pelvic Floor Disorders
-	C12.672 Tuberculosis, Urogenital
-	C12.672.721 Tuberculosis, Male Genital
-	C12.672.847 Tuberculosis, Renal
-	C12.706 Urogenital Abnormalities
-	C12.706.132 Bladder Exstrophy
-	C12.706.258 Cryptorchidism
-	C12.706.316 Disorders of Sex Development
-	C12.706.316.064 46, XX Disorders of Sex Development
-	C12.706.316.064.124 46, XX Testicular Disorders of Sex Development
-	C12.706.316.064.249 Gonadal Dysgenesis, 46,XX
-	C12.706.316.064.500 Hyperandrogenism

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C12.706.316.096 46, XY Disorders of Sex Development
-	C12.706.316.096.500 Androgen-Insensitivity Syndrome
-	C12.706.316.096.562 Denys-Drash Syndrome
-	C12.706.316.096.624 Frasier Syndrome
-	C12.706.316.096.687 Gonadal Dysgenesis, 46,XY
-	C12.706.316.096.687.500 Gonadoblastoma
-	C12.706.316.096.750 Kallmann Syndrome
-	C12.706.316.096.875 WAGR Syndrome
-	C12.706.316.129 Adrenogenital Syndrome
-	C12.706.316.129.500 Adrenal Hyperplasia, Congenital
-	C12.706.316.129.750 Hyperandrogenism
-	C12.706.316.309 Gonadal Dysgenesis
-	C12.706.316.309.193 Gonadal Dysgenesis, 46,XX
-	C12.706.316.309.388 Gonadal Dysgenesis, 46,XY
-	C12.706.316.309.388.500 Gonadoblastoma
-	C12.706.316.309.391 Gonadal Dysgenesis, Mixed
-	C12.706.316.309.631 Sexual Infantilism
-	C12.706.316.309.872 Turner Syndrome
-	C12.706.316.343 Ovotesticular Disorders of Sex Development
-	C12.706.316.795 Sex Chromosome Disorders of Sex Development
-	C12.706.316.795.124 Freemartinism
-	C12.706.316.795.249 Gonadal Dysgenesis, Mixed
-	C12.706.316.795.500 Klinefelter Syndrome
-	C12.706.316.795.750 Turner Syndrome
-	C12.706.374 Epispadias
New Tree	<a href="#">C12.706.410</a> <a href="#">Fraser Syndrome</a>
-	C12.706.445 Fused Kidney
-	C12.706.516 Hypospadias
-	C12.706.629 Multicystic Dysplastic Kidney
-	C12.706.742 Nephritis, Hereditary
-	C12.706.811 Retrocaval Ureter
-	C12.706.881 Urinary Fistula
-	C12.706.881.312 Urinary Bladder Fistula
-	C12.758 Urogenital Neoplasms
-	C12.758.409 Genital Neoplasms, Male

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C12.758.409.500 Penile Neoplasms
-	C12.758.409.750 Prostatic Neoplasms
-	C12.758.409.750.500 Prostatic Neoplasms, Castration-Resistant
-	C12.758.409.937 Testicular Neoplasms
-	C12.758.409.937.500 Sertoli-Leydig Cell Tumor
-	C12.758.409.937.500.249 Leydig Cell Tumor
-	C12.758.409.937.500.500 Sertoli Cell Tumor
-	C12.758.820 Urologic Neoplasms
-	C12.758.820.750 Kidney Neoplasms
-	C12.758.820.750.160 Carcinoma, Renal Cell
-	C12.758.820.750.372 Nephroma, Mesoblastic
-	C12.758.820.750.585 Wilms Tumor
-	C12.758.820.750.585.220 Denys-Drash Syndrome
-	C12.758.820.750.585.950 WAGR Syndrome
-	C12.758.820.875 Ureteral Neoplasms
-	C12.758.820.937 Urethral Neoplasms
-	C12.758.820.968 Urinary Bladder Neoplasms
-	C12.777 Urologic Diseases
-	C12.777.419 Kidney Diseases
-	C12.777.419.050 AIDS-Associated Nephropathy
-	C12.777.419.078 Anuria
-	C12.777.419.135 Diabetes Insipidus
-	C12.777.419.135.500 Diabetes Insipidus, Nephrogenic
-	C12.777.419.135.750 Diabetes Insipidus, Neurogenic
-	C12.777.419.135.875 Wolfram Syndrome
-	C12.777.419.192 Diabetic Nephropathies
-	C12.777.419.291 Hepatorenal Syndrome
-	C12.777.419.307 Hydronephrosis
-	C12.777.419.307.500 Pyonephrosis
-	C12.777.419.313 Hyperoxaluria
-	C12.777.419.313.500 Hyperoxaluria, Primary
-	C12.777.419.331 Hypertension, Renal
-	C12.777.419.331.490 Hypertension, Renovascular
-	C12.777.419.393 Kidney Cortex Necrosis
-	C12.777.419.403 Kidney Diseases, Cystic
-	C12.777.419.403.500 Medullary Sponge Kidney

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C12.777.419.403.750 Multicystic Dysplastic Kidney
-	C12.777.419.403.875 Polycystic Kidney Diseases
-	C12.777.419.403.875.500 Polycystic Kidney, Autosomal Dominant
-	C12.777.419.403.875.510 Polycystic Kidney, Autosomal Recessive
-	C12.777.419.473 Kidney Neoplasms
-	C12.777.419.473.160 Carcinoma, Renal Cell
-	C12.777.419.473.372 Nephroma, Mesoblastic
-	C12.777.419.473.585 Wilms Tumor
-	C12.777.419.473.585.220 Denys-Drash Syndrome
-	C12.777.419.473.585.950 WAGR Syndrome
-	C12.777.419.493 Kidney Papillary Necrosis
-	C12.777.419.570 Nephritis
-	C12.777.419.570.363 Glomerulonephritis
-	C12.777.419.570.363.304 Anti-Glomerular Basement Membrane Disease
-	C12.777.419.570.363.608 Glomerulonephritis, IGA
-	C12.777.419.570.363.615 Glomerulonephritis, Membranoproliferative
-	C12.777.419.570.363.625 Glomerulonephritis, Membranous
-	C12.777.419.570.363.660 Glomerulosclerosis, Focal Segmental
-	C12.777.419.570.363.680 Lupus Nephritis
-	C12.777.419.570.620 Nephritis, Hereditary
-	C12.777.419.570.643 Nephritis, Interstitial
-	C12.777.419.570.643.150 Balkan Nephropathy
-	C12.777.419.570.643.790 Pyelonephritis
-	C12.777.419.570.643.790.810 Pyelonephritis, Xanthogranulomatous
-	C12.777.419.570.821 Pyelitis
-	C12.777.419.570.821.358 Pyelocystitis
-	C12.777.419.570.821.717 Pyelonephritis
-	C12.777.419.570.821.717.810 Pyelonephritis, Xanthogranulomatous
-	C12.777.419.590 Nephrocalcinosis
-	C12.777.419.600 Nephrolithiasis
-	C12.777.419.600.500 Kidney Calculi
New Heading	<b>C12.777.419.600.500.500 Staghorn Calculi</b>
-	C12.777.419.610 Nephrosclerosis
-	C12.777.419.630 Nephrosis
-	C12.777.419.630.477 Nephrosis, Lipoid

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C12.777.419.630.643 Nephrotic Syndrome
-	C12.777.419.685 Perinephritis
-	C12.777.419.775 Renal Artery Obstruction
-	C12.777.419.780 Renal Insufficiency
-	C12.777.419.780.050 Acute Kidney Injury
-	C12.777.419.780.050.500 Kidney Tubular Necrosis, Acute
-	C12.777.419.780.400 Cardio-Renal Syndrome
-	C12.777.419.780.750 Renal Insufficiency, Chronic
-	C12.777.419.780.750.500 Kidney Failure, Chronic
-	C12.777.419.780.750.500.500 Frasier Syndrome
-	C12.777.419.787 Renal Nutcracker Syndrome
-	C12.777.419.795 Chronic Kidney Disease-Mineral and Bone Disorder
-	C12.777.419.795 Renal Osteodystrophy
-	C12.777.419.815 Renal Tubular Transport, Inborn Errors
-	C12.777.419.815.093 Acidosis, Renal Tubular
-	C12.777.419.815.279 Bartter Syndrome
-	C12.777.419.815.364 Dent Disease
-	C12.777.419.815.450 Fanconi Syndrome
-	C12.777.419.815.491 Gitelman Syndrome
-	C12.777.419.815.532 Glycosuria, Renal
-	C12.777.419.815.647 Hypophosphatemia, Familial
-	C12.777.419.815.647.500 Familial Hypophosphatemic Rickets
-	C12.777.419.815.683 Liddle Syndrome
-	C12.777.419.815.720 Oculocerebrorenal Syndrome
-	C12.777.419.815.770 Pseudohypoaldosteronism
-	C12.777.419.815.885 Renal Aminoacidurias
-	C12.777.419.815.885.250 Cystinuria
-	C12.777.419.815.885.457 Hartnup Disease
-	C12.777.419.912 Tuberculosis, Renal
-	C12.777.419.936 Uremia
-	C12.777.419.936.231 Azotemia
-	C12.777.419.936.463 Hemolytic-Uremic Syndrome
-	C12.777.419.936.463.500 Atypical Hemolytic Uremic Syndrome
-	C12.777.419.978 Zellweger Syndrome
-	C12.777.725 Ureteral Diseases
-	C12.777.725.676 Ureteral Neoplasms

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C12.777.725.776 Ureteral Obstruction
-	C12.777.725.876 Ureterocele
-	C12.777.725.938 Ureterolithiasis
-	C12.777.725.938.500 Ureteral Calculi
-	C12.777.767 Urethral Diseases
-	C12.777.767.374 Epispadias
-	C12.777.767.601 Urethral Neoplasms
-	C12.777.767.700 Urethral Obstruction
-	C12.777.767.700.700 Urethral Stricture
-	C12.777.767.700.962 Urinary Bladder Neck Obstruction
-	C12.777.767.851 Urethritis
-	C12.777.829 Urinary Bladder Diseases
-	C12.777.829.132 Bladder Exstrophy
-	C12.777.829.495 Cystitis
-	C12.777.829.495.500 Cystitis, Interstitial
-	C12.777.829.495.750 Pyelocystitis
-	C12.777.829.707 Cystocele
-	C12.777.829.720 Urinary Bladder Calculi
-	C12.777.829.733 Urinary Bladder Fistula
-	C12.777.829.760 Urinary Bladder Neck Obstruction
-	C12.777.829.813 Urinary Bladder Neoplasms
-	C12.777.829.839 Urinary Bladder, Neurogenic
-	C12.777.829.866 Urinary Bladder, Overactive
-	C12.777.829.920 Vesico-Ureteral Reflux
-	C12.777.892 Urinary Tract Infections
-	C12.777.892.219 Bacteriuria
-	C12.777.892.719 Pyuria
-	C12.777.892.775 Schistosomiasis haematobia
-	C12.777.934 Urination Disorders
-	C12.777.934.141 Anuria
-	C12.777.934.284 Enuresis
-	C12.777.934.284.249 Diurnal Enuresis
-	C12.777.934.284.500 Nocturnal Enuresis
-	C12.777.934.363 Glycosuria
-	C12.777.934.363.450 Glycosuria, Renal
-	C12.777.934.442 Hematuria

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	C12.777.934.600	Oliguria
-	C12.777.934.616	Polyuria
-	C12.777.934.734	Proteinuria
-	C12.777.934.734.269	Albuminuria
-	C12.777.934.734.634	Hemoglobinuria
-	C12.777.934.852	Urinary Incontinence
-	C12.777.934.852.249	Urinary Incontinence, Stress
-	C12.777.934.852.500	Urinary Incontinence, Urge
-	C12.777.934.880	Urinary Retention
-	C12.777.967	Urolithiasis
-	C12.777.967.249	Nephrolithiasis
-	C12.777.967.249.500	Kidney Calculi
New Heading	<b>C12.777.967.249.500.500</b>	<b>Staghorn Calculi</b>
-	C12.777.967.374	Ureterolithiasis
-	C12.777.967.374.500	Ureteral Calculi
-	C12.777.967.500	Urinary Calculi
-	C12.777.967.500.503	Kidney Calculi
New Heading	<b>C12.777.967.500.503.500</b>	<b>Staghorn Calculi</b>
-	C12.777.967.500.851	Ureteral Calculi
-	C12.777.967.500.925	Urinary Bladder Calculi
-	C13	Female Urogenital Diseases and Pregnancy Complications
-	C13.351	Female Urogenital Diseases
-	C13.351.500	Genital Diseases, Female
-	C13.351.500.056	Adnexal Diseases
-	C13.351.500.056.390	Fallopian Tube Diseases
-	C13.351.500.056.390.390	Fallopian Tube Neoplasms
-	C13.351.500.056.390.890	Salpingitis
-	C13.351.500.056.630	Ovarian Diseases
-	C13.351.500.056.630.050	Anovulation
-	C13.351.500.056.630.250	Menopause, Premature
-	C13.351.500.056.630.450	Oophoritis
-	C13.351.500.056.630.580	Ovarian Cysts
-	C13.351.500.056.630.580.765	Polycystic Ovary Syndrome
-	C13.351.500.056.630.642	Ovarian Hyperstimulation Syndrome



## MeSH Tree Changes for 2017

Type	Tree - heading
-	C13.351.500.056.630.705 Ovarian Neoplasms
-	C13.351.500.056.630.705.265 Brenner Tumor
-	C13.351.500.056.630.705.331 Carcinoma, Endometrioid
-	C13.351.500.056.630.705.398 Granulosa Cell Tumor
-	C13.351.500.056.630.705.431 Hereditary Breast and Ovarian Cancer Syndrome
-	C13.351.500.056.630.705.464 Luteoma
-	C13.351.500.056.630.705.531 Meigs Syndrome
-	C13.351.500.056.630.705.648 Sertoli-Leydig Cell Tumor
-	C13.351.500.056.630.705.765 Thecoma
-	C13.351.500.056.630.750 Primary Ovarian Insufficiency
-	C13.351.500.056.750 Pelvic Inflammatory Disease
-	C13.351.500.056.750.249 Endometritis
-	C13.351.500.056.750.500 Oophoritis
-	C13.351.500.056.750.750 Parametritis
-	C13.351.500.056.750.875 Salpingitis
-	C13.351.500.163 Endometriosis
-	C13.351.500.320 Gynatresia
-	C13.351.500.342 Herpes Genitalis
-	C13.351.500.365 Infertility
-	C13.351.500.365.700 Infertility, Female
-	C13.351.500.630 Reproductive Tract Infections
-	C13.351.500.665 Sexual Dysfunction, Physiological
-	C13.351.500.665.313 Dyspareunia
-	C13.351.500.665.656 Vaginismus
-	C13.351.500.711 Sexually Transmitted Diseases
-	C13.351.500.711.281 Sexually Transmitted Diseases, Bacterial
-	C13.351.500.711.281.201 Chancroid
-	C13.351.500.711.281.301 Chlamydia Infections
-	C13.351.500.711.281.301.490 Lymphogranuloma Venereum
-	C13.351.500.711.281.401 Gonorrhoea
-	C13.351.500.711.281.451 Granuloma Inguinale
-	C13.351.500.711.281.859 Syphilis
-	C13.351.500.758 Tuberculosis, Female Genital
-	C13.351.500.852 Uterine Diseases
-	C13.351.500.852.113 Adenomyosis

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C13.351.500.852.228      Endometrial Hyperplasia
-	C13.351.500.852.299      Endometritis
-	C13.351.500.852.495      Hematometra
-	C13.351.500.852.544      Pyometra
-	C13.351.500.852.593      Uterine Cervical Diseases
-	C13.351.500.852.593.074      Uterine Cervical Dysplasia
-	C13.351.500.852.593.074.249      Atypical Squamous Cells of the Cervix
-	C13.351.500.852.593.074.500      Squamous Intraepithelial Lesions of the Cervix
-	C13.351.500.852.593.112      Uterine Cervical Erosion
-	C13.351.500.852.593.120      Uterine Cervical Incompetence
-	C13.351.500.852.593.131      Uterine Cervical Neoplasms
-	C13.351.500.852.593.150      Uterine Cervicitis
-	C13.351.500.852.691      Uterine Hemorrhage
-	C13.351.500.852.691.449      Menorrhagia
-	C13.351.500.852.691.622      Metrorrhagia
-	C13.351.500.852.726      Uterine Inversion
-	C13.351.500.852.762      Uterine Neoplasms
-	C13.351.500.852.762.200      Endometrial Neoplasms
-	C13.351.500.852.762.200.500      Endometrial Stromal Tumors
-	C13.351.500.852.762.200.500.500      Sarcoma, Endometrial Stromal
-	C13.351.500.852.762.850      Uterine Cervical Neoplasms
-	C13.351.500.852.833      Uterine Prolapse
-	C13.351.500.852.904      Uterine Rupture
-	C13.351.500.852.904.500      Uterine Perforation
-	C13.351.500.852.952      Uterine Retroversion
-	C13.351.500.894      Vaginal Diseases
-	C13.351.500.894.300      Hematocolpos
-	C13.351.500.894.500      Hydrocolpos
-	C13.351.500.894.700      Vaginal Discharge
-	C13.351.500.894.700.500      Leukorrhoea
-	C13.351.500.894.767      Vaginal Fistula
-	C13.351.500.894.767.249      Rectovaginal Fistula
-	C13.351.500.894.767.500      Vesicovaginal Fistula
-	C13.351.500.894.834      Vaginal Neoplasms
-	C13.351.500.894.870      Vaginismus
-	C13.351.500.894.906      Vaginitis

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C13.351.500.894.906.316 Atrophic Vaginitis
-	C13.351.500.894.906.633 Trichomonas Vaginitis
-	C13.351.500.894.906.800 Vaginosis, Bacterial
-	C13.351.500.894.906.820 Vulvovaginitis
-	C13.351.500.894.906.820.500 Candidiasis, Vulvovaginal
-	C13.351.500.944 Vulvar Diseases
-	C13.351.500.944.626 Pruritus Vulvae
-	C13.351.500.944.815 Vulvar Lichen Sclerosus
-	C13.351.500.944.819 Vulvar Neoplasms
-	C13.351.500.944.902 Vulvitis
-	C13.351.500.944.902.368 Vulvar Vestibulitis
-	C13.351.500.944.902.737 Vulvovaginitis
-	C13.351.500.944.902.737.500 Candidiasis, Vulvovaginal
-	C13.351.500.944.951 Vulvodynia
-	C13.351.625 Pelvic Floor Disorders
-	C13.351.750 Tuberculosis, Urogenital
-	C13.351.750.940 Tuberculosis, Female Genital
-	C13.351.750.970 Tuberculosis, Renal
-	C13.351.875 Urogenital Abnormalities
-	C13.351.875.132 Bladder Exstrophy
-	C13.351.875.253 Disorders of Sex Development
-	C13.351.875.253.064 46, XX Disorders of Sex Development
-	C13.351.875.253.064.124 46, XX Testicular Disorders of Sex Development
-	C13.351.875.253.064.249 Gonadal Dysgenesis, 46,XX
-	C13.351.875.253.064.500 Hyperandrogenism
-	C13.351.875.253.096 46, XY Disorders of Sex Development
-	C13.351.875.253.096.500 Androgen-Insensitivity Syndrome
-	C13.351.875.253.096.562 Denys-Drash Syndrome
-	C13.351.875.253.096.624 Frasier Syndrome
-	C13.351.875.253.096.687 Gonadal Dysgenesis, 46,XY
-	C13.351.875.253.096.687.500 Gonadoblastoma
-	C13.351.875.253.096.750 Kallmann Syndrome
-	C13.351.875.253.096.875 WAGR Syndrome
-	C13.351.875.253.129 Adrenogenital Syndrome
-	C13.351.875.253.129.500 Adrenal Hyperplasia, Congenital
-	C13.351.875.253.129.750 Hyperandrogenism

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C13.351.875.253.309                      Gonadal Dysgenesis
-	C13.351.875.253.309.193                      Gonadal Dysgenesis, 46,XX
-	C13.351.875.253.309.388                      Gonadal Dysgenesis, 46,XY
-	C13.351.875.253.309.388.500                      Gonadoblastoma
-	C13.351.875.253.309.391                      Gonadal Dysgenesis, Mixed
-	C13.351.875.253.309.631                      Sexual Infantilism
-	C13.351.875.253.309.872                      Turner Syndrome
-	C13.351.875.253.343                      Ovotesticular Disorders of Sex Development
-	C13.351.875.253.795                      Sex Chromosome Disorders of Sex Development
-	C13.351.875.253.795.124                      Freemartinism
-	C13.351.875.253.795.249                      Gonadal Dysgenesis, Mixed
-	C13.351.875.253.795.500                      Klinefelter Syndrome
-	C13.351.875.253.795.750                      Turner Syndrome
-	C13.351.875.374                      Epispadias
New Tree	<a href="#">C13.351.875.397</a> <a href="#">Fraser Syndrome</a>
-	C13.351.875.420                      Fused Kidney
-	C13.351.875.466                      Hypospadias
-	C13.351.875.558                      Multicystic Dysplastic Kidney
-	C13.351.875.742                      Nephritis, Hereditary
-	C13.351.875.811                      Retrocaval Ureter
-	C13.351.875.881                      Urinary Fistula
-	C13.351.875.881.312                      Urinary Bladder Fistula
-	C13.351.875.881.312.733                      Vesicovaginal Fistula
-	C13.351.937                      Urogenital Neoplasms
-	C13.351.937.418                      Genital Neoplasms, Female
-	C13.351.937.418.365                      Fallopian Tube Neoplasms
-	C13.351.937.418.685                      Ovarian Neoplasms
-	C13.351.937.418.685.265                      Brenner Tumor
-	C13.351.937.418.685.331                      Carcinoma, Endometrioid
-	C13.351.937.418.685.398                      Granulosa Cell Tumor
-	C13.351.937.418.685.431                      Hereditary Breast and Ovarian Cancer Syndrome
-	C13.351.937.418.685.464                      Luteoma
-	C13.351.937.418.685.531                      Meigs Syndrome
-	C13.351.937.418.685.648                      Sertoli-Leydig Cell Tumor
-	C13.351.937.418.685.765                      Thecoma

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C13.351.937.418.875 Uterine Neoplasms
-	C13.351.937.418.875.200 Endometrial Neoplasms
-	C13.351.937.418.875.200.124 Carcinoma, Endometrioid
-	C13.351.937.418.875.200.374 Endometrial Stromal Tumors
-	C13.351.937.418.875.200.374.500 Sarcoma, Endometrial Stromal
-	C13.351.937.418.875.850 Uterine Cervical Neoplasms
-	C13.351.937.418.937 Vaginal Neoplasms
-	C13.351.937.418.968 Vulvar Neoplasms
-	C13.351.937.820 Urologic Neoplasms
-	C13.351.937.820.535 Kidney Neoplasms
-	C13.351.937.820.535.160 Carcinoma, Renal Cell
-	C13.351.937.820.535.585 Wilms Tumor
-	C13.351.937.820.535.585.220 Denys-Drash Syndrome
-	C13.351.937.820.535.585.950 WAGR Syndrome
-	C13.351.937.820.535.790 Nephroma, Mesoblastic
-	C13.351.937.820.875 Ureteral Neoplasms
-	C13.351.937.820.890 Urethral Neoplasms
-	C13.351.937.820.945 Urinary Bladder Neoplasms
-	C13.351.968 Urologic Diseases
-	C13.351.968.419 Kidney Diseases
-	C13.351.968.419.050 AIDS-Associated Nephropathy
-	C13.351.968.419.078 Anuria
-	C13.351.968.419.135 Diabetes Insipidus
-	C13.351.968.419.135.500 Diabetes Insipidus, Nephrogenic
-	C13.351.968.419.135.750 Diabetes Insipidus, Neurogenic
-	C13.351.968.419.135.875 Wolfram Syndrome
-	C13.351.968.419.192 Diabetic Nephropathies
-	C13.351.968.419.291 Hepatorenal Syndrome
-	C13.351.968.419.307 Hydronephrosis
-	C13.351.968.419.307.500 Pyonephrosis
-	C13.351.968.419.313 Hyperoxaluria
-	C13.351.968.419.313.500 Hyperoxaluria, Primary
-	C13.351.968.419.331 Hypertension, Renal
-	C13.351.968.419.331.490 Hypertension, Renovascular
-	C13.351.968.419.393 Kidney Cortex Necrosis
-	C13.351.968.419.403 Kidney Diseases, Cystic

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	C13.351.968.419.403.500	Medullary Sponge Kidney
-	C13.351.968.419.403.750	Multicystic Dysplastic Kidney
-	C13.351.968.419.403.875	Polycystic Kidney Diseases
-	C13.351.968.419.403.875.500	Polycystic Kidney, Autosomal Dominant
-	C13.351.968.419.403.875.510	Polycystic Kidney, Autosomal Recessive
-	C13.351.968.419.473	Kidney Neoplasms
-	C13.351.968.419.473.160	Carcinoma, Renal Cell
-	C13.351.968.419.473.372	Nephroma, Mesoblastic
-	C13.351.968.419.473.585	Wilms Tumor
-	C13.351.968.419.473.585.220	Denys-Drash Syndrome
-	C13.351.968.419.473.585.950	WAGR Syndrome
-	C13.351.968.419.493	Kidney Papillary Necrosis
-	C13.351.968.419.570	Nephritis
-	C13.351.968.419.570.363	Glomerulonephritis
-	C13.351.968.419.570.363.304	Anti-Glomerular Basement Membrane Disease
-	C13.351.968.419.570.363.608	Glomerulonephritis, IGA
-	C13.351.968.419.570.363.615	Glomerulonephritis, Membranoproliferative
-	C13.351.968.419.570.363.625	Glomerulonephritis, Membranous
-	C13.351.968.419.570.363.640	Glomerulosclerosis, Focal Segmental
-	C13.351.968.419.570.363.680	Lupus Nephritis
-	C13.351.968.419.570.620	Nephritis, Hereditary
-	C13.351.968.419.570.643	Nephritis, Interstitial
-	C13.351.968.419.570.643.150	Balkan Nephropathy
-	C13.351.968.419.570.643.790	Pyelonephritis
-	C13.351.968.419.570.643.790.810	Pyelonephritis, Xanthogranulomatous
-	C13.351.968.419.570.821	Pyelitis
-	C13.351.968.419.570.821.358	Pyelocystitis
-	C13.351.968.419.570.821.717	Pyelonephritis
-	C13.351.968.419.570.821.717.810	Pyelonephritis, Xanthogranulomatous
-	C13.351.968.419.590	Nephrocalcinosis
-	C13.351.968.419.600	Nephrolithiasis
-	C13.351.968.419.600.500	Kidney Calculi
New Heading	<b>C13.351.968.419.600.500.500</b>	<b>Staghorn Calculi</b>
-	C13.351.968.419.610	Nephrosclerosis
-	C13.351.968.419.630	Nephrosis

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C13.351.968.419.630.477 Nephrosis, Lipoid
-	C13.351.968.419.630.643 Nephrotic Syndrome
-	C13.351.968.419.685 Perinephritis
-	C13.351.968.419.775 Renal Artery Obstruction
-	C13.351.968.419.780 Renal Insufficiency
-	C13.351.968.419.780.050 Acute Kidney Injury
-	C13.351.968.419.780.050.500 Kidney Tubular Necrosis, Acute
-	C13.351.968.419.780.400 Cardio-Renal Syndrome
-	C13.351.968.419.780.750 Renal Insufficiency, Chronic
-	C13.351.968.419.780.750.500 Kidney Failure, Chronic
-	C13.351.968.419.780.750.500.500 Frasier Syndrome
-	C13.351.968.419.787 Renal Nutcracker Syndrome
-	C13.351.968.419.795 Chronic Kidney Disease-Mineral and Bone Disorder
-	C13.351.968.419.795 Renal Osteodystrophy
-	C13.351.968.419.815 Renal Tubular Transport, Inborn Errors
-	C13.351.968.419.815.093 Acidosis, Renal Tubular
-	C13.351.968.419.815.279 Bartter Syndrome
-	C13.351.968.419.815.364 Dent Disease
-	C13.351.968.419.815.450 Fanconi Syndrome
-	C13.351.968.419.815.491 Gitelman Syndrome
-	C13.351.968.419.815.532 Glycosuria, Renal
-	C13.351.968.419.815.647 Hypophosphatemia, Familial
-	C13.351.968.419.815.647.500 Familial Hypophosphatemic Rickets
-	C13.351.968.419.815.683 Liddle Syndrome
-	C13.351.968.419.815.720 Oculocerebrorenal Syndrome
-	C13.351.968.419.815.770 Pseudohypoaldosteronism
-	C13.351.968.419.815.885 Renal Aminoacidurias
-	C13.351.968.419.815.885.250 Cystinuria
-	C13.351.968.419.815.885.625 Hartnup Disease
-	C13.351.968.419.912 Tuberculosis, Renal
-	C13.351.968.419.936 Uremia
-	C13.351.968.419.936.231 Azotemia
-	C13.351.968.419.936.463 Hemolytic-Uremic Syndrome
-	C13.351.968.419.936.463.500 Atypical Hemolytic Uremic Syndrome
-	C13.351.968.419.978 Zellweger Syndrome
-	C13.351.968.725 Ureteral Diseases

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C13.351.968.725.676 Ureteral Neoplasms
-	C13.351.968.725.776 Ureteral Obstruction
-	C13.351.968.725.876 Ureterocele
-	C13.351.968.725.938 Ureterolithiasis
-	C13.351.968.725.938.500 Ureteral Calculi
-	C13.351.968.767 Urethral Diseases
-	C13.351.968.767.374 Epispadias
-	C13.351.968.767.601 Urethral Neoplasms
-	C13.351.968.767.700 Urethral Obstruction
-	C13.351.968.767.700.700 Urethral Stricture
-	C13.351.968.767.700.850 Urinary Bladder Neck Obstruction
-	C13.351.968.767.851 Urethritis
-	C13.351.968.829 Urinary Bladder Diseases
-	C13.351.968.829.132 Bladder Exstrophy
-	C13.351.968.829.495 Cystitis
-	C13.351.968.829.495.500 Cystitis, Interstitial
-	C13.351.968.829.495.750 Pyelocystitis
-	C13.351.968.829.508 Cystocele
-	C13.351.968.829.521 Urinary Bladder Calculi
-	C13.351.968.829.548 Urinary Bladder Fistula
-	C13.351.968.829.548.733 Vesicovaginal Fistula
-	C13.351.968.829.601 Urinary Bladder Neck Obstruction
-	C13.351.968.829.707 Urinary Bladder Neoplasms
-	C13.351.968.829.760 Urinary Bladder, Neurogenic
-	C13.351.968.829.813 Urinary Bladder, Overactive
-	C13.351.968.829.920 Vesico-Ureteral Reflux
-	C13.351.968.892 Urinary Tract Infections
-	C13.351.968.892.219 Bacteriuria
-	C13.351.968.892.719 Pyuria
-	C13.351.968.892.775 Schistosomiasis haematobia
-	C13.351.968.934 Urination Disorders
-	C13.351.968.934.070 Anuria
-	C13.351.968.934.252 Enuresis
-	C13.351.968.934.252.249 Diurnal Enuresis
-	C13.351.968.934.252.500 Nocturnal Enuresis
-	C13.351.968.934.363 Glycosuria



## MeSH Tree Changes for 2017

Type	Tree - heading	
-	C13.351.968.934.363.450	Glycosuria, Renal
-	C13.351.968.934.442	Hematuria
-	C13.351.968.934.600	Oliguria
-	C13.351.968.934.616	Polyuria
-	C13.351.968.934.734	Proteinuria
-	C13.351.968.934.734.269	Albuminuria
-	C13.351.968.934.734.634	Hemoglobinuria
-	C13.351.968.934.814	Urinary Incontinence
-	C13.351.968.934.814.500	Urinary Incontinence, Stress
-	C13.351.968.934.814.750	Urinary Incontinence, Urge
-	C13.351.968.934.880	Urinary Retention
-	C13.351.968.967	Urolithiasis
-	C13.351.968.967.249	Nephrolithiasis
-	C13.351.968.967.249.500	Kidney Calculi
New Heading	<b>C13.351.968.967.249.500.500</b>	<b>Staghorn Calculi</b>
-	C13.351.968.967.374	Ureterolithiasis
-	C13.351.968.967.374.500	Ureteral Calculi
-	C13.351.968.967.500	Urinary Calculi
-	C13.351.968.967.500.503	Kidney Calculi
New Heading	<b>C13.351.968.967.500.503.500</b>	<b>Staghorn Calculi</b>
-	C13.351.968.967.500.851	Ureteral Calculi
-	C13.351.968.967.500.925	Urinary Bladder Calculi
-	C13.703	Pregnancy Complications
-	C13.703.039	Abortion, Spontaneous
-	C13.703.039.089	Abortion, Habitual
-	C13.703.039.089.339	Uterine Cervical Incompetence
-	C13.703.039.093	Abortion, Incomplete
-	C13.703.039.173	Abortion, Missed
-	C13.703.039.256	Abortion, Septic
-	C13.703.039.339	Abortion, Threatened
-	C13.703.039.422	Abortion, Veterinary
-	C13.703.039.711	Embryo Loss
-	C13.703.141	Chorea Gravidarum
-	C13.703.170	Diabetes, Gestational

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C13.703.170.500                      Fetal Macrosomia
-	C13.703.223                              Fetal Death
-	C13.703.223.300                      Fetal Resorption
-	C13.703.223.650                      Stillbirth
-	C13.703.277                              Fetal Diseases
-	C13.703.277.030                      Chorioamnionitis
-	C13.703.277.050                      Echogenic Bowel
-	C13.703.277.060                      Erythroblastosis, Fetal
-	C13.703.277.060.480                      Hydrops Fetalis
-	C13.703.277.220                      Fetal Alcohol Spectrum Disorders
-	C13.703.277.370                      Fetal Growth Retardation
-	C13.703.277.390                      Fetal Hypoxia
-	C13.703.277.570                      Fetal Macrosomia
-	C13.703.277.677                      Fetal Nutrition Disorders
-	C13.703.277.785                      Meconium Aspiration Syndrome
-	C13.703.277.838                      Pyelectasis
-	C13.703.395                              Hypertension, Pregnancy-Induced
-	C13.703.395.124                      Eclampsia
-	C13.703.395.186                      HELLP Syndrome
-	C13.703.395.249                      Pre-Eclampsia
-	C13.703.401                              Maternal Death
-	C13.703.407                              Morning Sickness
-	C13.703.407.500                      Hyperemesis Gravidarum
-	C13.703.413                              Nuchal Cord
-	C13.703.420                              Obstetric Labor Complications
-	C13.703.420.078                      Abruptio Placentae
-	C13.703.420.183                      Breech Presentation
-	C13.703.420.235                      Cephalopelvic Disproportion
-	C13.703.420.288                      Dystocia
-	C13.703.420.288.728                      Uterine Inertia
-	C13.703.420.339                      Fetal Membranes, Premature Rupture
-	C13.703.420.339.260                      Chorioamnionitis
-	C13.703.420.491                      Obstetric Labor, Premature
-	C13.703.420.491.500                      Premature Birth
-	C13.703.420.643                      Placenta Accreta
-	C13.703.420.714                      Placenta Previa

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C13.703.420.725 Postpartum Hemorrhage
-	C13.703.420.814 Uterine Inversion
-	C13.703.420.904 Uterine Rupture
-	C13.703.420.952 Vasa Previa
-	C13.703.560 Oligohydramnios
-	C13.703.565 Pelvic Floor Disorders
-	C13.703.570 Pemphigoid Gestationis
-	C13.703.572 Perinatal Death
-	C13.703.575 Phenylketonuria, Maternal
-	C13.703.590 Placenta Diseases
-	C13.703.590.132 Abruptio Placentae
-	C13.703.590.268 Chorioamnionitis
-	C13.703.590.609 Placenta Accreta
-	C13.703.590.734 Placenta Previa
-	C13.703.590.767 Placenta, Retained
-	C13.703.590.800 Placental Insufficiency
-	C13.703.610 Polyhydramnios
-	C13.703.634 Pregnancy Complications, Cardiovascular
-	C13.703.634.404 Embolism, Amniotic Fluid
-	C13.703.667 Pregnancy Complications, Hematologic
-	C13.703.700 Pregnancy Complications, Infectious
-	C13.703.700.173 Abortion, Septic
-	C13.703.700.680 Pregnancy Complications, Parasitic
-	C13.703.700.715 Puerperal Infection
-	C13.703.720 Pregnancy Complications, Neoplastic
-	C13.703.720.949 Trophoblastic Neoplasms
-	C13.703.720.949.208 Choriocarcinoma
-	C13.703.720.949.208.438 Choriocarcinoma, Non-gestational
-	C13.703.720.949.208.875 Trophoblastic Tumor, Placental Site
-	C13.703.720.949.416 Gestational Trophoblastic Disease
-	C13.703.720.949.416.875 Hydatidiform Mole
-	C13.703.720.949.416.875.500 Hydatidiform Mole, Invasive
-	C13.703.726 Pregnancy in Diabetics
-	C13.703.726.570 Fetal Macrosomia
-	C13.703.733 Pregnancy, Ectopic
-	C13.703.733.536 Pregnancy, Abdominal

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C13.703.733.619 Pregnancy, Angular
-	C13.703.733.640 Pregnancy, Cornual
-	C13.703.733.650 Pregnancy, Heterotopic
-	C13.703.733.661 Pregnancy, Ovarian
-	C13.703.733.703 Pregnancy, Tubal
-	C13.703.733.703.500 Pregnancy, Interstitial
-	C13.703.805 Pregnancy, Prolonged
-	C13.703.824 Prenatal Injuries
-	C13.703.824.500 Prenatal Exposure Delayed Effects
-	C13.703.844 Puerperal Disorders
-	C13.703.844.253 Depression, Postpartum
-	C13.703.844.506 Lactation Disorders
-	C13.703.844.506.389 Galactorrhea
-	C13.703.844.506.389.500 Chiari-Frommel Syndrome
-	C13.703.844.603 Mastitis
-	C13.703.844.603.400 Granulomatous Mastitis
-	C13.703.844.700 Postpartum Hemorrhage
-	C13.703.844.714 Postpartum Thyroiditis
-	C13.703.844.728 Pubic Symphysis Diastasis
-	C13.703.844.757 Puerperal Infection
-	C14 Cardiovascular Diseases
-	C14.240 Cardiovascular Abnormalities
-	C14.240.400 Heart Defects, Congenital
-	C14.240.400.021 22q11 Deletion Syndrome
-	C14.240.400.021.500 DiGeorge Syndrome
-	C14.240.400.044 Alagille Syndrome
-	C14.240.400.090 Aortic Coarctation
-	C14.240.400.145 Arrhythmogenic Right Ventricular Dysplasia
-	C14.240.400.172 Barth Syndrome
-	C14.240.400.200 Cor Triatriatum
-	C14.240.400.210 Coronary Vessel Anomalies
-	C14.240.400.210.249 Bland White Garland Syndrome
-	C14.240.400.210.500 Myocardial Bridging
-	C14.240.400.220 Crisscross Heart
-	C14.240.400.280 Dextrocardia
-	C14.240.400.280.500 Kartagener Syndrome

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C14.240.400.340 Ductus Arteriosus, Patent
-	C14.240.400.395 Ebstein Anomaly
-	C14.240.400.422 Ectopia Cordis
-	C14.240.400.450 Eisenmenger Complex
-	C14.240.400.560 Heart Septal Defects
-	C14.240.400.560.098 Aortopulmonary Septal Defect
-	C14.240.400.560.098.500 Truncus Arteriosus, Persistent
-	C14.240.400.560.350 Endocardial Cushion Defects
-	C14.240.400.560.375 Heart Septal Defects, Atrial
-	C14.240.400.560.375.258 Foramen Ovale, Patent
-	C14.240.400.560.375.518 Lutembacher Syndrome
-	C14.240.400.560.540 Heart Septal Defects, Ventricular
-	C14.240.400.592 Heterotaxy Syndrome
-	C14.240.400.625 Hypoplastic Left Heart Syndrome
-	C14.240.400.660 Isolated Noncompaction of the Ventricular Myocardium
-	C14.240.400.695 LEOPARD Syndrome
-	C14.240.400.701 Levocardia
-	C14.240.400.725 Marfan Syndrome
-	C14.240.400.787 Noonan Syndrome
-	C14.240.400.849 Tetralogy of Fallot
-	C14.240.400.915 Transposition of Great Vessels
-	C14.240.400.915.300 Double Outlet Right Ventricle
-	C14.240.400.920 Tricuspid Atresia
-	C14.240.400.960 Trilogy of Fallot
-	C14.240.400.980 Turner Syndrome
-	C14.240.850 Vascular Malformations
-	C14.240.850.750 Arteriovenous Malformations
-	C14.240.850.750.147 Arteriovenous Fistula
-	C14.240.850.750.147.500 Carotid-Cavernous Sinus Fistula
-	C14.240.850.750.295 Intracranial Arteriovenous Malformations
-	C14.240.850.750.295.500 Vein of Galen Malformations
-	C14.240.850.875 Central Nervous System Vascular Malformations
-	C14.240.850.875.124 Central Nervous System Venous Angioma
-	C14.240.850.875.249 Hemangioma, Cavernous, Central Nervous System
-	C14.240.850.875.500 Intracranial Arteriovenous Malformations
-	C14.240.850.875.500.500 Vein of Galen Malformations

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C14.240.850.875.750 Sinus Pericranii
-	C14.240.850.906 May-Thurner Syndrome
-	C14.240.850.937 Pulmonary Atresia
-	C14.240.850.968 Scimitar Syndrome
-	C14.240.850.976 Single Umbilical Artery
-	C14.240.850.984 Vascular Fistula
-	C14.240.850.984.500 Arterio-Arterial Fistula
-	C14.240.850.984.500.500 Bland White Garland Syndrome
-	C14.240.850.984.750 Arteriovenous Fistula
-	C14.240.850.984.750.500 Carotid-Cavernous Sinus Fistula
-	C14.260 Cardiovascular Infections
-	C14.260.249 Endocarditis, Bacterial
-	C14.260.249.407 Endocarditis, Subacute Bacterial
-	C14.260.500 Syphilis, Cardiovascular
-	C14.260.750 Tuberculosis, Cardiovascular
-	C14.260.750.595 Pericarditis, Tuberculous
-	C14.280 Heart Diseases
-	C14.280.067 Arrhythmias, Cardiac
-	C14.280.067.093 Arrhythmia, Sinus
-	C14.280.067.093.249 Sick Sinus Syndrome
-	C14.280.067.093.500 Sinus Arrest, Cardiac
-	C14.280.067.198 Atrial Fibrillation
-	C14.280.067.248 Atrial Flutter
-	C14.280.067.319 Bradycardia
-	C14.280.067.322 Brugada Syndrome
-	C14.280.067.325 Cardiac Complexes, Premature
-	C14.280.067.325.250 Atrial Premature Complexes
-	C14.280.067.325.500 Ventricular Premature Complexes
-	C14.280.067.441 Commotio Cordis
-	C14.280.067.558 Heart Block
-	C14.280.067.558.137 Adams-Stokes Syndrome
-	C14.280.067.558.230 Atrioventricular Block
-	C14.280.067.558.323 Bundle-Branch Block
-	C14.280.067.558.536 Sick Sinus Syndrome
-	C14.280.067.558.750 Sinoatrial Block
-	C14.280.067.565 Long QT Syndrome

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C14.280.067.565.070 Andersen Syndrome
-	C14.280.067.565.440 Jervell-Lange Nielsen Syndrome
-	C14.280.067.565.720 Romano-Ward Syndrome
-	C14.280.067.672 Parasystole
-	C14.280.067.780 Pre-Excitation Syndromes
-	C14.280.067.780.560 Lown-Ganong-Levine Syndrome
-	C14.280.067.780.770 Pre-Excitation, Mahaim-Type
-	C14.280.067.780.977 Wolff-Parkinson-White Syndrome
-	C14.280.067.845 Tachycardia
-	C14.280.067.845.695 Tachycardia, Paroxysmal
-	C14.280.067.845.787 Tachycardia, Reciprocating
-	C14.280.067.845.787.249 Tachycardia, Atrioventricular Nodal Reentry
-	C14.280.067.845.787.500 Tachycardia, Sinoatrial Nodal Reentry
-	C14.280.067.845.880 Tachycardia, Supraventricular
-	C14.280.067.845.880.315 Tachycardia, Ectopic Atrial
-	C14.280.067.845.880.320 Tachycardia, Ectopic Junctional
-	C14.280.067.845.880.845 Tachycardia, Sinus
-	C14.280.067.845.940 Tachycardia, Ventricular
-	C14.280.067.845.940.349 Accelerated Idioventricular Rhythm
-	C14.280.067.845.940.700 Torsades de Pointes
-	C14.280.067.922 Ventricular Fibrillation
-	C14.280.067.961 Ventricular Flutter
-	C14.280.104 Carcinoid Heart Disease
-	C14.280.142 Cardiac Output, High
-	C14.280.148 Cardiac Output, Low
-	C14.280.155 Cardiac Tamponade
-	C14.280.195 Cardiomegaly
-	C14.280.195.160 Cardiomyopathy, Dilated
-	C14.280.195.400 Hypertrophy, Left Ventricular
-	C14.280.195.410 Hypertrophy, Right Ventricular
-	C14.280.238 Cardiomyopathies
-	C14.280.238.028 Arrhythmogenic Right Ventricular Dysplasia
-	C14.280.238.057 Cardiomyopathy, Alcoholic
-	C14.280.238.070 Cardiomyopathy, Dilated
-	C14.280.238.100 Cardiomyopathy, Hypertrophic
-	C14.280.238.100.500 Cardiomyopathy, Hypertrophic, Familial

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C14.280.238.160 Cardiomyopathy, Restrictive
-	C14.280.238.190 Chagas Cardiomyopathy
-	C14.280.238.235 Diabetic Cardiomyopathies
-	C14.280.238.281 Endocardial Fibroelastosis
-	C14.280.238.281.500 Isolated Noncompaction of the Ventricular Myocardium
-	C14.280.238.406 Endomyocardial Fibrosis
-	C14.280.238.458 Glycogen Storage Disease Type IIb
-	C14.280.238.510 Kearns-Sayre Syndrome
-	C14.280.238.615 Myocardial Reperfusion Injury
-	C14.280.238.625 Myocarditis
-	C14.280.238.812 Sarcoglycanopathies
-	C14.280.282 Endocarditis
-	C14.280.282.407 Endocarditis, Bacterial
-	C14.280.282.407.407 Endocarditis, Subacute Bacterial
-	C14.280.282.703 Endocarditis, Non-Infective
-	C14.280.358 Heart Aneurysm
-	C14.280.383 Heart Arrest
-	C14.280.383.220 Death, Sudden, Cardiac
-	C14.280.383.220.500 Karoshi Death
-	C14.280.383.610 Out-of-Hospital Cardiac Arrest
-	C14.280.400 Heart Defects, Congenital
-	C14.280.400.044 22q11 Deletion Syndrome
-	C14.280.400.044.500 DiGeorge Syndrome
-	C14.280.400.090 Aortic Coarctation
-	C14.280.400.145 Arrhythmogenic Right Ventricular Dysplasia
-	C14.280.400.172 Barth Syndrome
-	C14.280.400.200 Cor Triatriatum
-	C14.280.400.210 Coronary Vessel Anomalies
-	C14.280.400.210.249 Bland White Garland Syndrome
-	C14.280.400.210.500 Myocardial Bridging
-	C14.280.400.220 Crisscross Heart
-	C14.280.400.280 Dextrocardia
-	C14.280.400.280.500 Kartagener Syndrome
-	C14.280.400.340 Ductus Arteriosus, Patent
-	C14.280.400.395 Ebstein Anomaly
-	C14.280.400.450 Eisenmenger Complex



## MeSH Tree Changes for 2017

Type	Tree - heading
-	C14.280.400.560 Heart Septal Defects
-	C14.280.400.560.098 Aortopulmonary Septal Defect
-	C14.280.400.560.098.500 Truncus Arteriosus, Persistent
-	C14.280.400.560.350 Endocardial Cushion Defects
-	C14.280.400.560.375 Heart Septal Defects, Atrial
-	C14.280.400.560.375.258 Foramen Ovale, Patent
-	C14.280.400.560.375.518 Lutembacher Syndrome
-	C14.280.400.560.540 Heart Septal Defects, Ventricular
-	C14.280.400.592 Heterotaxy Syndrome
-	C14.280.400.625 Hypoplastic Left Heart Syndrome
-	C14.280.400.660 Isolated Noncompaction of the Ventricular Myocardium
-	C14.280.400.695 LEOPARD Syndrome
-	C14.280.400.701 Levocardia
-	C14.280.400.725 Marfan Syndrome
-	C14.280.400.787 Noonan Syndrome
-	C14.280.400.849 Tetralogy of Fallot
-	C14.280.400.915 Transposition of Great Vessels
-	C14.280.400.915.300 Double Outlet Right Ventricle
-	C14.280.400.920 Tricuspid Atresia
-	C14.280.400.960 Trilogy of Fallot
-	C14.280.400.980 Turner Syndrome
-	C14.280.434 Heart Failure
-	C14.280.434.156 Cardio-Renal Syndrome
-	C14.280.434.313 Dyspnea, Paroxysmal
-	C14.280.434.482 Edema, Cardiac
-	C14.280.434.611 Heart Failure, Diastolic
-	C14.280.434.676 Heart Failure, Systolic
-	C14.280.459 Heart Neoplasms
-	C14.280.459.500 Carney Complex
-	C14.280.470 Heart Rupture
-	C14.280.470.475 Heart Rupture, Post-Infarction
-	C14.280.470.475.900 Ventricular Septal Rupture
-	C14.280.484 Heart Valve Diseases
-	C14.280.484.095 Aortic Valve Insufficiency
-	C14.280.484.150 Aortic Valve Stenosis
-	C14.280.484.150.070 Aortic Stenosis, Subvalvular

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C14.280.484.150.070.160 Cardiomyopathy, Hypertrophic
-	C14.280.484.150.070.160.500 Cardiomyopathy, Hypertrophic, Familial
-	C14.280.484.150.070.210 Discrete Subaortic Stenosis
-	C14.280.484.150.535 Aortic Stenosis, Supravalvular
-	C14.280.484.150.535.960 Williams Syndrome
-	C14.280.484.400 Heart Valve Prolapse
-	C14.280.484.400.100 Aortic Valve Prolapse
-	C14.280.484.400.500 Mitral Valve Prolapse
-	C14.280.484.400.875 Tricuspid Valve Prolapse
-	C14.280.484.461 Mitral Valve Insufficiency
-	C14.280.484.517 Mitral Valve Stenosis
-	C14.280.484.640 Pulmonary Atresia
-	C14.280.484.660 Pulmonary Valve Insufficiency
-	C14.280.484.716 Pulmonary Valve Stenosis
-	C14.280.484.716.525 LEOPARD Syndrome
-	C14.280.484.716.762 Pulmonary Subvalvular Stenosis
-	C14.280.484.845 Tricuspid Atresia
-	C14.280.484.856 Tricuspid Valve Insufficiency
-	C14.280.484.911 Tricuspid Valve Stenosis
-	C14.280.647 Myocardial Ischemia
-	C14.280.647.124 Acute Coronary Syndrome
-	C14.280.647.187 Angina Pectoris
-	C14.280.647.187.150 Angina, Unstable
-	C14.280.647.187.150.150 Angina Pectoris, Variant
-	C14.280.647.187.362 Angina, Stable
-	C14.280.647.187.575 Microvascular Angina
-	C14.280.647.250 Coronary Disease
-	C14.280.647.250.250 Coronary Aneurysm
-	C14.280.647.250.260 Coronary Artery Disease
-	C14.280.647.250.272 Coronary Occlusion
-	C14.280.647.250.285 Coronary Stenosis
-	C14.280.647.250.285.200 Coronary Restenosis
-	C14.280.647.250.290 Coronary Thrombosis
-	C14.280.647.250.295 Coronary Vasospasm
-	C14.280.647.250.647 Coronary-Subclavian Steal Syndrome
-	C14.280.647.500 Myocardial Infarction

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C14.280.647.500.093 Anterior Wall Myocardial Infarction
-	C14.280.647.500.187 Inferior Wall Myocardial Infarction
New Heading	<b>C14.280.647.500.469 Non-ST Elevated Myocardial Infarction</b>
-	C14.280.647.500.750 Shock, Cardiogenic
New Heading	<b>C14.280.647.500.875 ST Elevation Myocardial Infarction</b>
-	C14.280.647.625 Myocardial Reperfusion Injury
-	C14.280.671 Myocardial Stunning
-	C14.280.695 Pericardial Effusion
-	C14.280.720 Pericarditis
-	C14.280.720.595 Pericarditis, Constrictive
-	C14.280.720.801 Pericarditis, Tuberculous
-	C14.280.763 Pneumopericardium
-	C14.280.793 Postpericardiotomy Syndrome
-	C14.280.832 Pulmonary Heart Disease
-	C14.280.874 Rheumatic Heart Disease
-	C14.280.945 Ventricular Dysfunction
-	C14.280.945.900 Ventricular Dysfunction, Left
-	C14.280.945.900.500 Takotsubo Cardiomyopathy
-	C14.280.945.910 Ventricular Dysfunction, Right
-	C14.280.955 Ventricular Outflow Obstruction
-	C14.280.955.249 Aortic Valve Stenosis
-	C14.280.955.249.070 Aortic Stenosis, Subvalvular
-	C14.280.955.249.070.210 Discrete Subaortic Stenosis
-	C14.280.955.249.535 Aortic Stenosis, Supravalvular
-	C14.280.955.750 Pulmonary Valve Stenosis
-	C14.280.955.750.762 Pulmonary Subvalvular Stenosis
-	C14.583 Pregnancy Complications, Cardiovascular
-	C14.583.404 Embolism, Amniotic Fluid
-	C14.907 Vascular Diseases
-	C14.907.055 Aneurysm
-	C14.907.055.050 Aneurysm, Dissecting
-	C14.907.055.050.150 Carotid Artery, Internal, Dissection
-	C14.907.055.050.362 Loeys-Dietz Syndrome
-	C14.907.055.050.575 Vertebral Artery Dissection

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C14.907.055.090 Aneurysm, False
-	C14.907.055.131 Aneurysm, Infected
-	C14.907.055.185 Aneurysm, Ruptured
-	C14.907.055.185.125 Aortic Rupture
-	C14.907.055.239 Aortic Aneurysm
-	C14.907.055.239.075 Aortic Aneurysm, Abdominal
-	C14.907.055.239.125 Aortic Aneurysm, Thoracic
-	C14.907.055.239.175 Aortic Rupture
-	C14.907.055.239.587 Loeys-Dietz Syndrome
-	C14.907.055.395 Coronary Aneurysm
-	C14.907.055.501 Endoleak
-	C14.907.055.608 Heart Aneurysm
-	C14.907.055.625 Iliac Aneurysm
-	C14.907.055.635 Intracranial Aneurysm
New Heading	<b>C14.907.055.817 Microaneurysm</b>
-	C14.907.075 Angiodysplasia
-	C14.907.075.280 Gastric Antral Vascular Ectasia
-	C14.907.077 Angiomatosis
-	C14.907.077.060 Angiomatosis, Bacillary
-	C14.907.077.410 Klippel-Trenaunay-Weber Syndrome
-	C14.907.077.850 Sturge-Weber Syndrome
-	C14.907.077.925 von Hippel-Lindau Disease
-	C14.907.079 Angioedema
-	C14.907.079.500 Angioedemas, Hereditary
-	C14.907.079.500.500 Hereditary Angioedema Type III
-	C14.907.079.500.750 Hereditary Angioedema Types I and II
-	C14.907.109 Aortic Diseases
-	C14.907.109.139 Aortic Aneurysm
-	C14.907.109.139.075 Aortic Aneurysm, Abdominal
-	C14.907.109.139.125 Aortic Aneurysm, Thoracic
-	C14.907.109.139.175 Aortic Rupture
-	C14.907.109.139.587 Loeys-Dietz Syndrome
-	C14.907.109.239 Aortic Arch Syndromes
-	C14.907.109.239.650 Takayasu Arteritis
-	C14.907.109.320 Aortitis

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C14.907.109.661                      Leriche Syndrome
-	C14.907.137                              Arterial Occlusive Diseases
-	C14.907.137.126                        Arteriosclerosis
-	C14.907.137.126.056                    Arteriolosclerosis
-	C14.907.137.126.114                    Arteriosclerosis Obliterans
-	C14.907.137.126.307                    Atherosclerosis
-	C14.907.137.126.307.500                Peripheral Arterial Disease
-	C14.907.137.126.339                    Coronary Artery Disease
-	C14.907.137.126.372                    Intracranial Arteriosclerosis
-	C14.907.137.126.372.500                Dementia, Vascular
-	C14.907.137.126.669                    Intermittent Claudication
-	C14.907.137.230                        Carotid Stenosis
-	C14.907.137.372                        Fibromuscular Dysplasia
-	C14.907.137.427                        Leriche Syndrome
-	C14.907.137.520                        Malignant Atrophic Papulosis
-	C14.907.137.534                        Mesenteric Vascular Occlusion
-	C14.907.137.615                        Moyamoya Disease
-	C14.907.137.727                        Renal Artery Obstruction
-	C14.907.137.780                        Retinal Artery Occlusion
-	C14.907.137.780.500                    Susac Syndrome
New Heading	<b>C14.907.137.825                        Stenosis, Pulmonary Artery</b>
-	C14.907.137.870                        Thromboangiitis Obliterans
-	C14.907.150                              Arteriovenous Malformations
-	C14.907.150.125                        Arteriovenous Fistula
-	C14.907.150.125.500                    Carotid-Cavernous Sinus Fistula
-	C14.907.150.295                        Intracranial Arteriovenous Malformations
-	C14.907.150.295.500                    Vein of Galen Malformations
-	C14.907.218                              Capillary Leak Syndrome
-	C14.907.253                              Cerebrovascular Disorders
-	C14.907.253.061                        Basal Ganglia Cerebrovascular Disease
-	C14.907.253.061.200                    Basal Ganglia Hemorrhage
-	C14.907.253.061.200.500                Putaminal Hemorrhage
-	C14.907.253.092                        Brain Ischemia
-	C14.907.253.092.477                    Brain Infarction
-	C14.907.253.092.477.100                Brain Stem Infarctions

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C14.907.253.092.477.100.500 Lateral Medullary Syndrome
-	C14.907.253.092.477.200 Cerebral Infarction
-	C14.907.253.092.477.200.199 Dementia, Multi-Infarct
-	C14.907.253.092.477.200.400 Infarction, Anterior Cerebral Artery
-	C14.907.253.092.477.200.450 Infarction, Middle Cerebral Artery
-	C14.907.253.092.477.200.475 Infarction, Posterior Cerebral Artery
-	C14.907.253.092.716 Hypoxia-Ischemia, Brain
-	C14.907.253.092.836 Ischemic Attack, Transient
-	C14.907.253.092.956 Vertebrobasilar Insufficiency
-	C14.907.253.092.956.700 Subclavian Steal Syndrome
-	C14.907.253.123 Carotid Artery Diseases
-	C14.907.253.123.345 Carotid Artery Injuries
-	C14.907.253.123.345.300 Carotid Artery, Internal, Dissection
-	C14.907.253.123.355 Carotid Artery Thrombosis
-	C14.907.253.123.357 Carotid-Cavernous Sinus Fistula
-	C14.907.253.123.360 Carotid Stenosis
-	C14.907.253.123.620 Moyamoya Disease
-	C14.907.253.329 Cerebral Small Vessel Diseases
-	C14.907.253.329.249 CADASIL
-	C14.907.253.329.311 Cerebral Amyloid Angiopathy, Familial
-	C14.907.253.329.374 Fabry Disease
-	C14.907.253.329.500 MELAS Syndrome
-	C14.907.253.329.600 Microscopic Polyangiitis
-	C14.907.253.329.800 Stroke, Lacunar
-	C14.907.253.535 Cerebrovascular Trauma
-	C14.907.253.535.500 Carotid Artery Injuries
-	C14.907.253.535.500.300 Carotid Artery, Internal, Dissection
-	C14.907.253.535.500.350 Carotid-Cavernous Sinus Fistula
-	C14.907.253.535.600 Subarachnoid Hemorrhage, Traumatic
-	C14.907.253.535.800 Vertebral Artery Dissection
-	C14.907.253.560 Intracranial Arterial Diseases
-	C14.907.253.560.200 Cerebral Arterial Diseases
-	C14.907.253.560.200.175 CADASIL
-	C14.907.253.560.200.200 Cerebral Amyloid Angiopathy
-	C14.907.253.560.200.200.160 Cerebral Amyloid Angiopathy, Familial
-	C14.907.253.560.200.325 Infarction, Anterior Cerebral Artery

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C14.907.253.560.200.387      Infarction, Middle Cerebral Artery
-	C14.907.253.560.200.418      Infarction, Posterior Cerebral Artery
-	C14.907.253.560.200.737      Moyamoya Disease
-	C14.907.253.560.300      Intracranial Aneurysm
-	C14.907.253.560.350      Intracranial Arteriosclerosis
-	C14.907.253.560.350.500      Dementia, Vascular
-	C14.907.253.560.400      Intracranial Arteriovenous Malformations
-	C14.907.253.560.400.500      Vein of Galen Malformations
-	C14.907.253.566      Intracranial Embolism and Thrombosis
-	C14.907.253.566.206      Carotid Artery Thrombosis
-	C14.907.253.566.300      Intracranial Embolism
-	C14.907.253.566.350      Intracranial Thrombosis
-	C14.907.253.566.350.500      Sinus Thrombosis, Intracranial
-	C14.907.253.566.350.500.375      Cavernous Sinus Thrombosis
-	C14.907.253.566.350.500.562      Lateral Sinus Thrombosis
-	C14.907.253.566.350.500.750      Sagittal Sinus Thrombosis
-	C14.907.253.573      Intracranial Hemorrhages
-	C14.907.253.573.200      Cerebral Hemorrhage
-	C14.907.253.573.200.150      Basal Ganglia Hemorrhage
-	C14.907.253.573.200.150.500      Putaminal Hemorrhage
-	C14.907.253.573.200.200      Cerebral Hemorrhage, Traumatic
-	C14.907.253.573.350      Intracranial Hemorrhage, Hypertensive
-	C14.907.253.573.400      Intracranial Hemorrhage, Traumatic
-	C14.907.253.573.400.150      Brain Hemorrhage, Traumatic
-	C14.907.253.573.400.150.200      Brain Stem Hemorrhage, Traumatic
-	C14.907.253.573.400.150.300      Cerebral Hemorrhage, Traumatic
-	C14.907.253.573.400.400      Hematoma, Epidural, Cranial
-	C14.907.253.573.400.450      Hematoma, Subdural
-	C14.907.253.573.400.450.050      Hematoma, Subdural, Acute
-	C14.907.253.573.400.450.120      Hematoma, Subdural, Chronic
-	C14.907.253.573.400.450.400      Hematoma, Subdural, Intracranial
-	C14.907.253.573.400.700      Subarachnoid Hemorrhage, Traumatic
-	C14.907.253.573.600      Pituitary Apoplexy
-	C14.907.253.573.800      Subarachnoid Hemorrhage
-	C14.907.253.573.800.700      Subarachnoid Hemorrhage, Traumatic
-	C14.907.253.612      Leukomalacia, Periventricular

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C14.907.253.774 Sneddon Syndrome
-	C14.907.253.855 Stroke
-	C14.907.253.855.200 Brain Infarction
-	C14.907.253.855.200.100 Brain Stem Infarctions
-	C14.907.253.855.200.100.500 Lateral Medullary Syndrome
-	C14.907.253.855.200.200 Cerebral Infarction
-	C14.907.253.855.200.200.199 Dementia, Multi-Infarct
-	C14.907.253.855.200.200.400 Infarction, Anterior Cerebral Artery
-	C14.907.253.855.200.200.450 Infarction, Middle Cerebral Artery
-	C14.907.253.855.200.200.475 Infarction, Posterior Cerebral Artery
-	C14.907.253.855.600 Stroke, Lacunar
-	C14.907.253.937 Vascular Headaches
-	C14.907.253.946 Vasculitis, Central Nervous System
-	C14.907.253.946.175 AIDS Arteritis, Central Nervous System
-	C14.907.253.946.700 Giant Cell Arteritis
-	C14.907.253.946.850 Lupus Vasculitis, Central Nervous System
-	C14.907.253.951 Vasospasm, Intracranial
-	C14.907.286 Colitis, Ischemic
-	C14.907.303 Compartment Syndromes
-	C14.907.303.063 Anterior Compartment Syndrome
-	C14.907.303.297 Intra-Abdominal Hypertension
-	C14.907.303.531 Ischemic Contracture
-	C14.907.320 Diabetic Angiopathies
-	C14.907.320.191 Diabetic Foot
-	C14.907.320.382 Diabetic Retinopathy
-	C14.907.355 Embolism and Thrombosis
-	C14.907.355.350 Embolism
-	C14.907.355.350.254 Embolism, Air
-	C14.907.355.350.354 Embolism, Amniotic Fluid
-	C14.907.355.350.454 Embolism, Fat
-	C14.907.355.350.454.500 Embolism, Cholesterol
-	C14.907.355.350.454.500.200 Blue Toe Syndrome
-	C14.907.355.350.700 Pulmonary Embolism
-	C14.907.355.350.700.500 Pulmonary Infarction
-	C14.907.355.590 Thromboembolism
-	C14.907.355.590.213 Intracranial Embolism and Thrombosis



## MeSH Tree Changes for 2017

Type	Tree - heading
-	C14.907.355.590.213.206 Carotid Artery Thrombosis
-	C14.907.355.590.213.300 Intracranial Embolism
-	C14.907.355.590.213.350 Intracranial Thrombosis
-	C14.907.355.590.213.350.500 Sinus Thrombosis, Intracranial
-	C14.907.355.590.213.350.500.375 Cavernous Sinus Thrombosis
-	C14.907.355.590.213.350.500.562 Lateral Sinus Thrombosis
-	C14.907.355.590.213.350.500.750 Sagittal Sinus Thrombosis
-	C14.907.355.590.400 Embolism, Paradoxical
-	C14.907.355.590.700 Venous Thromboembolism
-	C14.907.355.830 Thrombosis
-	C14.907.355.830.220 Coronary Thrombosis
-	C14.907.355.830.925 Venous Thrombosis
-	C14.907.355.830.925.275 Budd-Chiari Syndrome
-	C14.907.355.830.925.462 Postthrombotic Syndrome
-	C14.907.355.830.925.650 Retinal Vein Occlusion
-	C14.907.355.830.925.770 Thrombophlebitis
-	C14.907.355.830.925.770.500 Lemierre Syndrome
-	C14.907.355.830.925.885 Upper Extremity Deep Vein Thrombosis
-	C14.907.440 Hand-Arm Vibration Syndrome
-	C14.907.449 Hemorrhoids
-	C14.907.454 Hemostatic Disorders
-	C14.907.454.140 Cryoglobulinemia
-	C14.907.454.240 Ehlers-Danlos Syndrome
-	C14.907.454.385 Hemangioma, Cavernous
-	C14.907.454.385.500 Hemangioma, Cavernous, Central Nervous System
-	C14.907.454.460 Multiple Myeloma
-	C14.907.454.530 Pseudoxanthoma Elasticum
-	C14.907.454.550 Purpura, Hyperglobulinemic
-	C14.907.454.800 Scurvy
-	C14.907.454.810 Shwartzman Phenomenon
-	C14.907.454.900 Telangiectasia, Hereditary Hemorrhagic
-	C14.907.454.960 Waldenstrom Macroglobulinemia
-	C14.907.460 Hepatic Veno-Occlusive Disease
-	C14.907.474 Hyperemia
-	C14.907.489 Hypertension
-	C14.907.489.330 Hypertension, Malignant

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	C14.907.489.480	Hypertension, Pregnancy-Induced
-	C14.907.489.631	Hypertension, Renal
-	C14.907.489.631.485	Hypertension, Renovascular
-	C14.907.489.815	Hypertensive Retinopathy
-	C14.907.489.861	Masked Hypertension
-	C14.907.489.907	White Coat Hypertension
-	C14.907.514	Hypotension
-	C14.907.514.482	Hypotension, Orthostatic
-	C14.907.514.611	Post-Exercise Hypotension
-	C14.907.514.741	Shy-Drager Syndrome
-	C14.907.549	Mesenteric Ischemia
-	C14.907.585	Myocardial Ischemia
-	C14.907.585.124	Acute Coronary Syndrome
-	C14.907.585.187	Angina Pectoris
-	C14.907.585.187.150	Angina, Unstable
-	C14.907.585.187.150.500	Angina Pectoris, Variant
-	C14.907.585.187.362	Angina, Stable
-	C14.907.585.187.575	Microvascular Angina
-	C14.907.585.250	Coronary Disease
-	C14.907.585.250.250	Coronary Aneurysm
-	C14.907.585.250.260	Coronary Artery Disease
-	C14.907.585.250.272	Coronary Occlusion
-	C14.907.585.250.285	Coronary Stenosis
-	C14.907.585.250.285.200	Coronary Restenosis
-	C14.907.585.250.290	Coronary Thrombosis
-	C14.907.585.250.295	Coronary Vasospasm
-	C14.907.585.250.647	Coronary-Subclavian Steal Syndrome
-	C14.907.585.500	Myocardial Infarction
-	C14.907.585.500.093	Anterior Wall Myocardial Infarction
-	C14.907.585.500.187	Inferior Wall Myocardial Infarction
-	C14.907.585.500.562	No-Reflow Phenomenon
New Heading	<b>C14.907.585.500.656</b>	<b>Non-ST Elevated Myocardial Infarction</b>
-	C14.907.585.500.750	Shock, Cardiogenic
New Heading	<b>C14.907.585.500.875</b>	<b>ST Elevation Myocardial Infarction</b>

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C14.907.585.625 Myocardial Reperfusion Injury
-	C14.907.601 Optic Neuropathy, Ischemic
-	C14.907.617 Peripheral Vascular Diseases
-	C14.907.617.249 Blue Toe Syndrome
-	C14.907.617.500 Erythromelalgia
-	C14.907.617.625 Livedo Reticularis
-	C14.907.617.648 May-Thurner Syndrome
-	C14.907.617.671 Peripheral Arterial Disease
-	C14.907.617.718 Phlebitis
-	C14.907.617.718.760 Postphlebitic Syndrome
-	C14.907.617.718.788 Thrombophlebitis
-	C14.907.617.718.788.500 Lemierre Syndrome
-	C14.907.617.812 Raynaud Disease
-	C14.907.617.812.500 CREST Syndrome
-	C14.907.653 Prehypertension
-	C14.907.690 Pulmonary Veno-Occlusive Disease
-	C14.907.725 Reperfusion Injury
-	C14.907.725.600 Myocardial Reperfusion Injury
-	C14.907.725.675 Primary Graft Dysfunction
-	C14.907.760 Retinal Vein Occlusion
-	C14.907.780 Scimitar Syndrome
-	C14.907.790 Spinal Cord Vascular Diseases
-	C14.907.790.550 Spinal Cord Ischemia
-	C14.907.790.550.100 Anterior Spinal Artery Syndrome
-	C14.907.795 Splenic Infarction
New Heading	<b>C14.907.798 Stenosis, Pulmonary Vein</b>
-	C14.907.800 Superior Vena Cava Syndrome
-	C14.907.823 Telangiectasis
-	C14.907.823.213 Ataxia Telangiectasia
-	C14.907.823.225 CREST Syndrome
-	C14.907.823.502 Retinal Telangiectasis
-	C14.907.823.780 Telangiectasia, Hereditary Hemorrhagic
-	C14.907.863 Thoracic Outlet Syndrome
-	C14.907.863.200 Cervical Rib Syndrome
-	C14.907.903 Varicocele

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C14.907.927 Varicose Veins
-	C14.907.927.730 Varicose Ulcer
-	C14.907.933 Vascular Fistula
-	C14.907.933.110 Arterio-Arterial Fistula
-	C14.907.933.110.500 Bland White Garland Syndrome
-	C14.907.933.555 Arteriovenous Fistula
-	C14.907.933.555.500 Carotid-Cavernous Sinus Fistula
-	C14.907.936 Vascular Neoplasms
-	C14.907.937 Vascular System Injuries
-	C14.907.940 Vasculitis
-	C14.907.940.080 Aortitis
-	C14.907.940.090 Arteritis
-	C14.907.940.090.170 AIDS Arteritis, Central Nervous System
-	C14.907.940.090.340 Endarteritis
-	C14.907.940.090.530 Giant Cell Arteritis
-	C14.907.940.090.720 Polyarteritis Nodosa
-	C14.907.940.090.800 Takayasu Arteritis
-	C14.907.940.100 Behcet Syndrome
-	C14.907.940.320 Cogan Syndrome
-	C14.907.940.530 Malignant Atrophic Papulosis
-	C14.907.940.560 Mucocutaneous Lymph Node Syndrome
-	C14.907.940.740 Phlebitis
-	C14.907.940.740.910 Thrombophlebitis
-	C14.907.940.740.910.500 Lemierre Syndrome
-	C14.907.940.777 Purpura, Schoenlein-Henoch
-	C14.907.940.815 Retinal Vasculitis
-	C14.907.940.890 Shwartzman Phenomenon
-	C14.907.940.897 Systemic Vasculitis
-	C14.907.940.897.249 Anti-Neutrophil Cytoplasmic Antibody-Associated Vasculitis
-	C14.907.940.897.249.249 Churg-Strauss Syndrome
-	C14.907.940.897.249.500 Microscopic Polyangiitis
-	C14.907.940.897.249.750 Granulomatosis with Polyangiitis
-	C14.907.940.897.500 Polyarteritis Nodosa
-	C14.907.940.897.750 Rheumatoid Vasculitis
-	C14.907.940.905 Thromboangiitis Obliterans

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C14.907.940.907 Vasculitis, Central Nervous System
-	C14.907.940.907.175 AIDS Arteritis, Central Nervous System
-	C14.907.940.907.700 Giant Cell Arteritis
-	C14.907.940.907.850 Lupus Vasculitis, Central Nervous System
-	C14.907.940.910 Vasculitis, Leukocytoclastic, Cutaneous
-	C14.907.946 Vasoplegia
-	C14.907.952 Venous Insufficiency
-	C14.907.952.760 Postphlebitic Syndrome
-	C14.907.952.880 Postthrombotic Syndrome
-	C15 Hemic and Lymphatic Diseases
-	C15.378 Hematologic Diseases
-	C15.378.071 Anemia
-	C15.378.071.085 Anemia, Aplastic
-	C15.378.071.085.080 Anemia, Hypoplastic, Congenital
-	C15.378.071.085.080.090 Anemia, Diamond-Blackfan
-	C15.378.071.085.080.280 Fanconi Anemia
-	C15.378.071.141 Anemia, Hemolytic
-	C15.378.071.141.125 Anemia, Hemolytic, Autoimmune
-	C15.378.071.141.150 Anemia, Hemolytic, Congenital
-	C15.378.071.141.150.095 Anemia, Dyserythropoietic, Congenital
-	C15.378.071.141.150.100 Anemia, Hemolytic, Congenital Nonspherocytic
-	C15.378.071.141.150.150 Anemia, Sickle Cell
-	C15.378.071.141.150.150.219 Acute Chest Syndrome
-	C15.378.071.141.150.150.440 Hemoglobin SC Disease
-	C15.378.071.141.150.150.670 Sickle Cell Trait
-	C15.378.071.141.150.365 Elliptocytosis, Hereditary
-	C15.378.071.141.150.480 Glucosephosphate Dehydrogenase Deficiency
-	C15.378.071.141.150.480.370 Favism
-	C15.378.071.141.150.490 Hemoglobin C Disease
-	C15.378.071.141.150.785 Spherocytosis, Hereditary
-	C15.378.071.141.150.875 Thalassemia
-	C15.378.071.141.150.875.100 alpha-Thalassemia
-	C15.378.071.141.150.875.150 beta-Thalassemia
-	C15.378.071.141.150.875.575 delta-Thalassemia
-	C15.378.071.141.370 Favism
-	C15.378.071.141.560 Hemoglobinuria, Paroxysmal

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C15.378.071.141.610 Hemolytic-Uremic Syndrome
-	C15.378.071.141.610.500 Atypical Hemolytic Uremic Syndrome
-	C15.378.071.196 Anemia, Hypochromic
-	C15.378.071.196.300 Anemia, Iron-Deficiency
-	C15.378.071.252 Anemia, Macrocytic
-	C15.378.071.252.196 Anemia, Megaloblastic
-	C15.378.071.252.196.500 Anemia, Pernicious
-	C15.378.071.307 Anemia, Myelophthisic
-	C15.378.071.363 Anemia, Neonatal
-	C15.378.071.363.344 Fetofetal Transfusion
-	C15.378.071.363.511 Fetomaternal Transfusion
-	C15.378.071.400 Anemia, Refractory
-	C15.378.071.400.080 Anemia, Refractory, with Excess of Blasts
-	C15.378.071.419 Anemia, Sideroblastic
-	C15.378.071.750 Red-Cell Aplasia, Pure
-	C15.378.071.750.500 Anemia, Diamond-Blackfan
-	C15.378.100 Blood Coagulation Disorders
-	C15.378.100.100 Blood Coagulation Disorders, Inherited
-	C15.378.100.100.037 Activated Protein C Resistance
-	C15.378.100.100.056 Afibrinogenemia
-	C15.378.100.100.075 Antithrombin III Deficiency
-	C15.378.100.100.080 Bernard-Soulier Syndrome
-	C15.378.100.100.300 Factor V Deficiency
-	C15.378.100.100.310 Factor VII Deficiency
-	C15.378.100.100.320 Factor X Deficiency
-	C15.378.100.100.325 Factor XI Deficiency
-	C15.378.100.100.330 Factor XII Deficiency
-	C15.378.100.100.335 Factor XIII Deficiency
-	C15.378.100.100.500 Hemophilia A
-	C15.378.100.100.510 Hemophilia B
-	C15.378.100.100.515 Hermanski-Pudlak Syndrome
-	C15.378.100.100.550 Hypoprothrombinemias
-	C15.378.100.100.690 Protein C Deficiency
-	C15.378.100.100.820 Thrombasthenia
-	C15.378.100.100.900 von Willebrand Diseases
-	C15.378.100.100.900.100 von Willebrand Disease, Type 1

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C15.378.100.100.900.200                      von Willebrand Disease, Type 2
-	C15.378.100.100.900.300                      von Willebrand Disease, Type 3
-	C15.378.100.100.970                            Wiskott-Aldrich Syndrome
-	C15.378.100.141                                Coagulation Protein Disorders
-	C15.378.100.141.036                          Activated Protein C Resistance
-	C15.378.100.141.072                          Afibrinogenemia
-	C15.378.100.141.300                          Factor V Deficiency
-	C15.378.100.141.310                          Factor VII Deficiency
-	C15.378.100.141.320                          Factor X Deficiency
-	C15.378.100.141.325                          Factor XI Deficiency
-	C15.378.100.141.330                          Factor XII Deficiency
-	C15.378.100.141.335                          Factor XIII Deficiency
-	C15.378.100.141.500                          Hemophilia A
-	C15.378.100.141.510                          Hemophilia B
-	C15.378.100.141.550                          Hypoprothrombinemias
-	C15.378.100.141.900                          von Willebrand Diseases
-	C15.378.100.141.900.100                      von Willebrand Disease, Type 1
-	C15.378.100.141.900.200                      von Willebrand Disease, Type 2
-	C15.378.100.141.900.300                      von Willebrand Disease, Type 3
-	C15.378.100.220                                Disseminated Intravascular Coagulation
-	C15.378.100.452                                Ecchymosis
-	C15.378.100.685                                Platelet Storage Pool Deficiency
-	C15.378.100.685.400                          Hermanski-Pudlak Syndrome
-	C15.378.100.800                                Protein S Deficiency
-	C15.378.100.802                                Purpura
-	C15.378.100.802.230                          Purpura Fulminans
-	C15.378.100.802.250                          Purpura, Hyperglobulinemic
-	C15.378.100.802.375                          Purpura, Schoenlein-Henoch
-	C15.378.100.802.687                          Purpura, Thrombocytopenic
-	C15.378.100.802.687.600                      Purpura, Thrombocytopenic, Idiopathic
-	C15.378.100.802.687.680                      Purpura, Thrombotic Thrombocytopenic
-	C15.378.100.802.843                          Waterhouse-Friderichsen Syndrome
-	C15.378.100.832                                Thrombocythemia, Essential
-	C15.378.100.920                                Vitamin K Deficiency
-	C15.378.100.920.360                          Vitamin K Deficiency Bleeding
-	C15.378.140                                      Blood Platelet Disorders

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C15.378.140.120 Bernard-Soulier Syndrome
-	C15.378.140.427 Gray Platelet Syndrome
-	C15.378.140.735 Platelet Storage Pool Deficiency
-	C15.378.140.735.400 Hermanski-Pudlak Syndrome
-	C15.378.140.810 Thrombasthenia
-	C15.378.140.855 Thrombocytopenia
-	C15.378.140.855.440 Jacobsen Distal 11q Deletion Syndrome
-	C15.378.140.855.645 Kasabach-Merritt Syndrome
-	C15.378.140.855.850 Thrombocytopenia, Neonatal Alloimmune
-	C15.378.140.855.925 Thrombotic Microangiopathies
-	C15.378.140.855.925.500 Hemolytic-Uremic Syndrome
-	C15.378.140.855.925.500.500 Atypical Hemolytic Uremic Syndrome
-	C15.378.140.855.925.750 Purpura, Thrombocytopenic
-	C15.378.140.855.925.750.600 Purpura, Thrombocytopenic, Idiopathic
-	C15.378.140.855.925.750.680 Purpura, Thrombotic Thrombocytopenic
-	C15.378.140.860 Thrombocytosis
-	C15.378.140.860.800 Thrombocythemia, Essential
-	C15.378.140.900 von Willebrand Diseases
-	C15.378.147 Blood Protein Disorders
-	C15.378.147.142 Agammaglobulinemia
-	C15.378.147.150 Antithrombin III Deficiency
-	C15.378.147.333 Dysgammaglobulinemia
-	C15.378.147.333.249 Hyper-IgM Immunodeficiency Syndrome
-	C15.378.147.333.249.500 Hyper-IgM Immunodeficiency Syndrome, Type 1
-	C15.378.147.333.500 IgA Deficiency
-	C15.378.147.333.750 IgG Deficiency
-	C15.378.147.542 Hypergammaglobulinemia
-	C15.378.147.542.319 Mevalonate Kinase Deficiency
-	C15.378.147.542.640 Monoclonal Gammopathy of Undetermined Significance
-	C15.378.147.607 Hypoproteinemia
-	C15.378.147.607.500 Hypoalbuminemia
-	C15.378.147.780 Paraproteinemias
-	C15.378.147.780.243 Cryoglobulinemia
-	C15.378.147.780.490 Heavy Chain Disease
-	C15.378.147.780.490.512 Immunoproliferative Small Intestinal Disease
-	C15.378.147.780.650 Multiple Myeloma



## MeSH Tree Changes for 2017

Type	Tree - heading
-	C15.378.147.780.750 POEMS Syndrome
-	C15.378.147.780.925 Waldenstrom Macroglobulinemia
-	C15.378.147.880 Protein C Deficiency
-	C15.378.147.890 Protein S Deficiency
-	C15.378.190 Bone Marrow Diseases
-	C15.378.190.196 Anemia, Aplastic
-	C15.378.190.196.080 Anemia, Hypoplastic, Congenital
-	C15.378.190.196.080.090 Anemia, Diamond-Blackfan
-	C15.378.190.196.080.280 Fanconi Anemia
-	C15.378.190.250 Bone Marrow Neoplasms
New Tree	<a href="#">C15.378.190.250.500</a> Polycythemia Vera
-	C15.378.190.615 Myelodysplastic-Myeloproliferative Diseases
-	C15.378.190.615.500 Negative Leukemia, Myeloid, Chronic, Atypical, BCR-ABL
-	C15.378.190.615.510 Leukemia, Myelomonocytic, Chronic
-	C15.378.190.615.520 Leukemia, Myelomonocytic, Juvenile
-	C15.378.190.625 Myelodysplastic Syndromes
-	C15.378.190.625.062 Anemia, Refractory
-	C15.378.190.625.062.080 Anemia, Refractory, with Excess of Blasts
-	C15.378.190.625.070 Anemia, Sideroblastic
-	C15.378.190.625.460 Hemoglobinuria, Paroxysmal
-	C15.378.190.636 Myeloproliferative Disorders
-	C15.378.190.636.085 Anemia, Myelophthitic
-	C15.378.190.636.276 Leukemia, Erythroblastic, Acute
-	C15.378.190.636.370 Leukemia, Myelogenous, Chronic, BCR-ABL Positive
-	C15.378.190.636.370.100 Blast Crisis
-	C15.378.190.636.370.300 Leukemia, Myeloid, Accelerated Phase
-	C15.378.190.636.370.400 Leukemia, Myeloid, Chronic-Phase
-	C15.378.190.636.380 Leukemia, Neutrophilic, Chronic
-	C15.378.190.636.484 Leukemoid Reaction
-	C15.378.190.636.753 Polycythemia Vera
-	C15.378.190.636.765 Primary Myelofibrosis
-	C15.378.190.636.860 Thrombocytosis
-	C15.378.190.636.860.800 Thrombocythemia, Essential
-	C15.378.295 Erythroblastosis, Fetal

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C15.378.295.480 Hydrops Fetalis
-	C15.378.295.502 Kernicterus
-	C15.378.400 Hematologic Neoplasms
-	C15.378.400.200 Bone Marrow Neoplasms
New Tree	<a href="#">C15.378.400.200.500</a> Polycythemia Vera
-	C15.378.420 Hemoglobinopathies
-	C15.378.420.155 Anemia, Sickle Cell
-	C15.378.420.155.219 Acute Chest Syndrome
-	C15.378.420.155.440 Hemoglobin SC Disease
-	C15.378.420.155.668 Sickle Cell Trait
-	C15.378.420.463 Hemoglobin C Disease
-	C15.378.420.826 Thalassemia
-	C15.378.420.826.100 alpha-Thalassemia
-	C15.378.420.826.100.350 Hydrops Fetalis
-	C15.378.420.826.150 beta-Thalassemia
-	C15.378.420.826.200 delta-Thalassemia
-	C15.378.463 Hemorrhagic Disorders
-	C15.378.463.067 Afibrinogenemia
-	C15.378.463.080 Bernard-Soulier Syndrome
-	C15.378.463.250 Disseminated Intravascular Coagulation
-	C15.378.463.300 Factor V Deficiency
-	C15.378.463.310 Factor VII Deficiency
-	C15.378.463.320 Factor X Deficiency
-	C15.378.463.325 Factor XI Deficiency
-	C15.378.463.330 Factor XII Deficiency
-	C15.378.463.335 Factor XIII Deficiency
-	C15.378.463.500 Hemophilia A
-	C15.378.463.510 Hemophilia B
-	C15.378.463.515 Hemostatic Disorders
-	C15.378.463.515.140 Cryoglobulinemia
-	C15.378.463.515.240 Ehlers-Danlos Syndrome
-	C15.378.463.515.385 Hemangioma, Cavernous
-	C15.378.463.515.385.500 Hemangioma, Cavernous, Central Nervous System
-	C15.378.463.515.460 Multiple Myeloma
-	C15.378.463.515.530 Pseudoxanthoma Elasticum

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C15.378.463.515.550 Purpura, Hyperglobulinemic
-	C15.378.463.515.580 Purpura, Schoenlein-Henoch
-	C15.378.463.515.800 Scurvy
-	C15.378.463.515.810 Shwartzman Phenomenon
-	C15.378.463.515.900 Telangiectasia, Hereditary Hemorrhagic
-	C15.378.463.515.960 Waldenstrom Macroglobulinemia
-	C15.378.463.550 Hypoprothrombinemias
-	C15.378.463.735 Platelet Storage Pool Deficiency
-	C15.378.463.735.400 Hermanski-Pudlak Syndrome
-	C15.378.463.740 Purpura, Thrombocytopenic, Idiopathic
-	C15.378.463.810 Thrombasthenia
-	C15.378.463.825 Thrombocythemia, Essential
-	C15.378.463.841 Vitamin K Deficiency
-	C15.378.463.841.464 Vitamin K Deficiency Bleeding
-	C15.378.463.920 von Willebrand Diseases
-	C15.378.463.920.100 von Willebrand Disease, Type 1
-	C15.378.463.920.200 von Willebrand Disease, Type 2
-	C15.378.463.920.300 von Willebrand Disease, Type 3
-	C15.378.463.950 Waterhouse-Friderichsen Syndrome
-	C15.378.463.960 Wiskott-Aldrich Syndrome
-	C15.378.553 Leukocyte Disorders
-	C15.378.553.231 Eosinophilia
-	C15.378.553.231.085 Angiolymphoid Hyperplasia with Eosinophilia
-	C15.378.553.231.335 Eosinophilia-Myalgia Syndrome
-	C15.378.553.231.341 Eosinophilic Esophagitis
-	C15.378.553.231.348 Eosinophilic Granuloma
-	C15.378.553.231.549 Hypereosinophilic Syndrome
-	C15.378.553.231.549.750 Pulmonary Eosinophilia
-	C15.378.553.381 Infectious Mononucleosis
-	C15.378.553.475 Leukocytosis
-	C15.378.553.475.428 Leukemoid Reaction
-	C15.378.553.475.604 Lymphocytosis
-	C15.378.553.546 Leukopenia
-	C15.378.553.546.184 Agranulocytosis
-	C15.378.553.546.184.564 Neutropenia
-	C15.378.553.546.184.564.750 Febrile Neutropenia

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C15.378.553.546.184.564.750.500 Neutropenia Chemotherapy-Induced Febrile
-	C15.378.553.546.605 Lymphopenia
-	C15.378.553.546.605.800 T-Lymphocytopenia, Idiopathic CD4-Positive
-	C15.378.553.546.605.900 Wiskott-Aldrich Syndrome
-	C15.378.553.560 Leukostasis
-	C15.378.553.696 Pelger-Huet Anomaly
-	C15.378.553.774 Phagocyte Bactericidal Dysfunction
-	C15.378.553.774.257 Chediak-Higashi Syndrome
-	C15.378.553.774.535 Granulomatous Disease, Chronic
-	C15.378.553.774.600 Job Syndrome
-	C15.378.619 Methemoglobinemia
-	C15.378.700 Pancytopenia
-	C15.378.738 Polycythemia
-	C15.378.785 Pregnancy Complications, Hematologic
-	C15.378.800 Preleukemia
-	C15.378.896 Sulfhemoglobinemia
-	C15.378.925 Thrombophilia
-	C15.378.925.050 Activated Protein C Resistance
-	C15.378.925.075 Antithrombin III Deficiency
-	C15.378.925.220 Disseminated Intravascular Coagulation
-	C15.378.925.795 Protein C Deficiency
-	C15.378.925.800 Protein S Deficiency
-	C15.378.925.850 Purpura, Thrombotic Thrombocytopenic
-	C15.378.962 Transfusion Reaction
-	C15.604 Lymphatic Diseases
-	C15.604.250 Histiocytosis
-	C15.604.250.390 Histiocytic Disorders, Malignant
-	C15.604.250.390.190 Dendritic Cell Sarcoma, Follicular
-	C15.604.250.390.199 Dendritic Cell Sarcoma, Interdigitating
-	C15.604.250.390.380 Histiocytic Sarcoma
-	C15.604.250.390.500 Langerhans Cell Sarcoma
-	C15.604.250.400 Histiocytosis, Langerhans-Cell
-	C15.604.250.400.360 Eosinophilic Granuloma
-	C15.604.250.410 Histiocytosis, Non-Langerhans-Cell
-	C15.604.250.410.224 Erdheim-Chester Disease

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	C15.604.250.410.450	Histiocytosis, Sinus
-	C15.604.250.410.575	Lymphohistiocytosis, Hemophagocytic
-	C15.604.250.410.625	Niemann-Pick Diseases
-	C15.604.250.410.625.500	Niemann-Pick Disease, Type A
-	C15.604.250.410.625.750	Niemann-Pick Disease, Type B
-	C15.604.250.410.625.875	Niemann-Pick Disease, Type C
-	C15.604.250.410.800	Sea-Blue Histiocyte Syndrome
-	C15.604.250.410.900	Xanthogranuloma, Juvenile
-	C15.604.315	Lymphadenitis
-	C15.604.315.249	Cat-Scratch Disease
-	C15.604.315.300	Histiocytic Necrotizing Lymphadenitis
-	C15.604.315.618	Mesenteric Lymphadenitis
New Heading	<b>C15.604.338</b>	<b>Lymphadenopathy</b>
New Tree	<a href="#">C15.604.338.500</a>	<a href="#">Immunoblastic Lymphadenopathy</a>
-	C15.604.360	Lymphangiectasis
-	C15.604.360.500	Lymphangiectasis, Intestinal
-	C15.604.406	Lymphangitis
-	C15.604.451	Lymphatic Abnormalities
-	C15.604.451.249	22q11 Deletion Syndrome
-	C15.604.451.249.500	DiGeorge Syndrome
-	C15.604.451.500	Lymphangiectasis, Intestinal
-	C15.604.496	Lymphedema
New Heading	<b>C15.604.496.160</b>	<b>Breast Cancer Lymphedema</b>
-	C15.604.496.320	Elephantiasis
-	C15.604.496.490	Elephantiasis, Filarial
-	C15.604.496.660	Non-Filarial Lymphedema
-	C15.604.510	Lymphocele
-	C15.604.515	Lymphoproliferative Disorders
-	C15.604.515.032	Agammaglobulinemia
-	C15.604.515.138	Autoimmune Lymphoproliferative Syndrome
-	C15.604.515.245	Giant Lymph Node Hyperplasia
-	C15.604.515.292	Granuloma
-	C15.604.515.292.007	Angiolymphoid Hyperplasia with Eosinophilia
-	C15.604.515.292.015	Churg-Strauss Syndrome

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C15.604.515.292.507 Necrobiotic Xanthogranuloma
-	C15.604.515.435 Heavy Chain Disease
-	C15.604.515.435.512 Immunoproliferative Small Intestinal Disease
-	C15.604.515.509 Immunoblastic Lymphadenopathy
-	C15.604.515.516 Infectious Mononucleosis
-	C15.604.515.553 Leukemia, Hairy Cell
-	C15.604.515.560 Leukemia, Lymphoid
-	C15.604.515.560.080 Leukemia, B-Cell
-	C15.604.515.560.080.125 Leukemia, Lymphocytic, Chronic, B-Cell
-	C15.604.515.560.080.562 Leukemia, Prolymphocytic, B-Cell
-	C15.604.515.560.100 Leukemia, Biphenotypic, Acute
-	C15.604.515.560.550 Leukemia, Prolymphocytic
-	C15.604.515.560.550.745 Leukemia, Prolymphocytic, B-Cell
-	C15.604.515.560.550.750 Leukemia, Prolymphocytic, T-Cell
-	C15.604.515.560.575 Leukemia, T-Cell
-	C15.604.515.560.575.049 Leukemia, Large Granular Lymphocytic
-	C15.604.515.560.575.100 Leukemia-Lymphoma, Adult T-Cell
-	C15.604.515.560.575.125 Leukemia, Prolymphocytic, T-Cell
-	C15.604.515.560.600 Precursor Cell Lymphoblastic Leukemia-Lymphoma
-	C15.604.515.560.600.600 Precursor B-Cell Lymphoblastic Leukemia-Lymphoma
-	C15.604.515.560.600.620 Precursor T-Cell Lymphoblastic Leukemia-Lymphoma
-	C15.604.515.562 Lymphangiomyoma
-	C15.604.515.562.465 Lymphangioliomyomatosis
-	C15.604.515.569 Lymphoma
-	C15.604.515.569.150 Composite Lymphoma
-	C15.604.515.569.355 Hodgkin Disease
-	C15.604.515.569.417 Intraocular Lymphoma
-	C15.604.515.569.480 Lymphoma, Non-Hodgkin
-	C15.604.515.569.480.150 Lymphoma, B-Cell
-	C15.604.515.569.480.150.165 Burkitt Lymphoma
-	C15.604.515.569.480.150.450 Lymphoma, AIDS-Related
-	C15.604.515.569.480.150.570 Lymphoma, B-Cell, Marginal Zone
-	C15.604.515.569.480.150.585 Lymphoma, Large B-Cell, Diffuse
-	C15.604.515.569.480.150.585.500 Plasmablastic Lymphoma

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C15.604.515.569.480.150.592                      Lymphoma, Primary Effusion
-	C15.604.515.569.480.150.600                      Lymphomatoid Granulomatosis
-	C15.604.515.569.480.350                          Lymphoma, Follicular
-	C15.604.515.569.480.493                          Lymphoma, Large-Cell, Immunoblastic
-	C15.604.515.569.480.525                          Lymphoma, Mantle-Cell
-	C15.604.515.569.480.750                          Lymphoma, T-Cell
-	C15.604.515.569.480.750.399                      Enteropathy-Associated T-Cell Lymphoma
-	C15.604.515.569.480.750.600                      Lymphoma, Large-Cell, Anaplastic
-	C15.604.515.569.480.750.800                      Lymphoma, T-Cell, Cutaneous
-	C15.604.515.569.480.750.800.507 Large Cell    Lymphoma, Primary Cutaneous Anaplastic
-	C15.604.515.569.480.750.800.528                      Lymphomatoid Papulosis
-	C15.604.515.569.480.750.800.550                      Mycosis Fungoides
-	C15.604.515.569.480.750.800.550.600                      Pagetoid Reticulosis
-	C15.604.515.569.480.750.800.775                      Sezary Syndrome
-	C15.604.515.569.480.750.825                          Lymphoma, T-Cell, Peripheral
-	C15.604.515.700                                      Marek Disease
-	C15.604.515.827                                      Sarcoidosis
-	C15.604.515.827.725                                      Sarcoidosis, Pulmonary
-	C15.604.515.827.865                                      Uveoparotid Fever
-	C15.604.515.841                                      Sezary Syndrome
-	C15.604.515.880                                      Tumor Lysis Syndrome
-	C15.604.515.925                                      Waldenstrom Macroglobulinemia
-	C15.604.560    Mucocutaneous Lymph Node Syndrome
-	C15.604.613    Pseudolymphoma
-	C15.604.744    Splenic Diseases
-	C15.604.744.146    Heterotaxy Syndrome
-	C15.604.744.293    Hypersplenism
-	C15.604.744.617    Splenic Infarction
-	C15.604.744.680    Splenic Neoplasms
-	C15.604.744.742    Splenic Rupture
-	C15.604.744.742.500    Splenosis
-	C15.604.744.909    Tuberculosis, Splenic
-	C15.604.744.954    Wandering Spleen
-	C15.604.816    Thymus Hyperplasia
-	C15.604.861    Thymus Neoplasms

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	C15.604.861.800	Thymoma
-	C16	Congenital, Hereditary, and Neonatal Diseases and Abnormalities
-	C16.131	Congenital Abnormalities
-	C16.131.042	Abnormalities, Drug-Induced
-	C16.131.077	Abnormalities, Multiple
-	C16.131.077.019	22q11 Deletion Syndrome
-	C16.131.077.019.500	DiGeorge Syndrome
-	C16.131.077.065	Alagille Syndrome
Old Tree	<b>C16.131.077.080</b>	<b>Alstrom Syndrome</b>
-	C16.131.077.095	Angelman Syndrome
Old Tree	<b>C16.131.077.112</b>	<b>Bardet-Biedl Syndrome</b>
-	C16.131.077.121	Barth Syndrome
-	C16.131.077.130	Basal Cell Nevus Syndrome
-	C16.131.077.133	Beckwith-Wiedemann Syndrome
-	C16.131.077.137	Bloom Syndrome
-	C16.131.077.208	Branchio-Oto-Renal Syndrome
-	C16.131.077.229	Carney Complex
-	C16.131.077.239	CHARGE Syndrome
New Heading	<b>C16.131.077.245</b>	<b>Ciliopathies</b>
New Tree	<b>C16.131.077.245.063</b>	<b>Alstrom Syndrome</b>
New Tree	<b>C16.131.077.245.125</b>	<b>Bardet-Biedl Syndrome</b>
New Tree	<b>C16.131.077.245.250</b>	<b>Caroli Disease</b>
New Tree	<b>C16.131.077.245.500</b>	<b>Ciliary Motility Disorders</b>
New Tree	<b>C16.131.077.245.500.531</b>	<b>Kartagener Syndrome</b>
-	C16.131.077.250	Cockayne Syndrome
-	C16.131.077.256	Costello Syndrome
-	C16.131.077.262	Cri-du-Chat Syndrome
-	C16.131.077.272	De Lange Syndrome
-	C16.131.077.299	Deaf-Blind Disorders
-	C16.131.077.299.500	Usher Syndromes
-	C16.131.077.299.750	Wolfram Syndrome
-	C16.131.077.313	Donohue Syndrome



## MeSH Tree Changes for 2017

Type	Tree - heading	
-	C16.131.077.327	Down Syndrome
-	C16.131.077.350	Ectodermal Dysplasia
-	C16.131.077.350.198	Ectodermal Dysplasia 1, Anhidrotic
-	C16.131.077.350.298	Ectodermal Dysplasia 3, Anhidrotic
-	C16.131.077.350.348 Recessive	Ectodermal Dysplasia, Hypohidrotic, Autosomal
-	C16.131.077.350.398	Ellis-Van Creveld Syndrome
-	C16.131.077.350.424	Focal Dermal Hypoplasia
-	C16.131.077.350.712	Neurocutaneous Syndromes
-	C16.131.077.371	Fraser Syndrome
-	C16.131.077.393	Gardner Syndrome
-	C16.131.077.401	Heterotaxy Syndrome
-	C16.131.077.410	Holoprosencephaly
-	C16.131.077.445	Incontinentia Pigmenti
-	C16.131.077.477	Isolated Noncompaction of the Ventricular Myocardium
-	C16.131.077.509	Laurence-Moon Syndrome
-	C16.131.077.525	LEOPARD Syndrome
-	C16.131.077.537	Loeys-Dietz Syndrome
-	C16.131.077.550	Marfan Syndrome
-	C16.131.077.578	Mobius Syndrome
-	C16.131.077.592	Monilethrix
-	C16.131.077.606	Nail-Patella Syndrome
-	C16.131.077.619	Netherton Syndrome
-	C16.131.077.633	Nevus, Sebaceous of Jadassohn
Old Tree	<del>C16.131.077.661</del>	<del>Oculocerebrorenal Syndrome</del>
New Tree	C16.131.077.662	Oculocerebrorenal Syndrome
New Tree	C16.131.077.676	Orofaciodigital Syndromes
Old Tree	<del>C16.131.077.677</del>	<del>Orofaciodigital Syndromes</del>
-	C16.131.077.690	Pallister-Hall Syndrome
-	C16.131.077.696	Pentalogy of Cantrell
-	C16.131.077.703	POEMS Syndrome
New Tree	C16.131.077.717	Polycystic Kidney Diseases
New Tree	C16.131.077.717.500	Polycystic Kidney, Autosomal Dominant

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	<a href="#">C16.131.077.717.510</a> <a href="#">Polycystic Kidney, Autosomal Recessive</a>
-	C16.131.077.730 Prader-Willi Syndrome
-	C16.131.077.735 Prolidase Deficiency
-	C16.131.077.740 Proteus Syndrome
-	C16.131.077.745 Prune Belly Syndrome
-	C16.131.077.790 Rubella Syndrome, Congenital
-	C16.131.077.804 Rubinstein-Taybi Syndrome
-	C16.131.077.850 Short Rib-Polydactyly Syndrome
-	C16.131.077.855 Silver-Russell Syndrome
-	C16.131.077.860 Smith-Lemli-Opitz Syndrome
-	C16.131.077.879 Smith-Magenis Syndrome
-	C16.131.077.889 Sotos Syndrome
-	C16.131.077.899 Trichothiodystrophy Syndromes
-	C16.131.077.938 Waardenburg Syndrome
-	C16.131.077.941 Weill-Marchesani Syndrome
-	C16.131.077.944 Wolf-Hirschhorn Syndrome
-	C16.131.077.970 Zellweger Syndrome
-	C16.131.080 Abnormalities, Radiation-Induced
-	C16.131.085 Abnormalities, Severe Teratoid
-	C16.131.085.197 Anencephaly
-	C16.131.085.806 Twins, Conjoined
-	C16.131.162 Aicardi Syndrome
-	C16.131.240 Cardiovascular Abnormalities
-	C16.131.240.400 Heart Defects, Congenital
-	C16.131.240.400.021 22q11 Deletion Syndrome
-	C16.131.240.400.021.500 DiGeorge Syndrome
-	C16.131.240.400.044 Alagille Syndrome
-	C16.131.240.400.090 Aortic Coarctation
-	C16.131.240.400.145 Arrhythmogenic Right Ventricular Dysplasia
-	C16.131.240.400.172 Barth Syndrome
-	C16.131.240.400.200 Cor Triatriatum
-	C16.131.240.400.210 Coronary Vessel Anomalies
-	C16.131.240.400.210.249 Bland White Garland Syndrome
-	C16.131.240.400.210.500 Myocardial Bridging
-	C16.131.240.400.220 Crisscross Heart

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C16.131.240.400.280 Dextrocardia
-	C16.131.240.400.280.500 Kartagener Syndrome
-	C16.131.240.400.340 Ductus Arteriosus, Patent
-	C16.131.240.400.395 Ebstein Anomaly
-	C16.131.240.400.422 Ectopia Cordis
-	C16.131.240.400.450 Eisenmenger Complex
-	C16.131.240.400.560 Heart Septal Defects
-	C16.131.240.400.560.098 Aortopulmonary Septal Defect
-	C16.131.240.400.560.098.500 Truncus Arteriosus, Persistent
-	C16.131.240.400.560.350 Endocardial Cushion Defects
-	C16.131.240.400.560.375 Heart Septal Defects, Atrial
-	C16.131.240.400.560.375.258 Foramen Ovale, Patent
-	C16.131.240.400.560.375.518 Lutembacher Syndrome
-	C16.131.240.400.560.540 Heart Septal Defects, Ventricular
-	C16.131.240.400.592 Heterotaxy Syndrome
-	C16.131.240.400.625 Hypoplastic Left Heart Syndrome
-	C16.131.240.400.655 Isolated Noncompaction of the Ventricular Myocardium
-	C16.131.240.400.685 LEOPARD Syndrome
-	C16.131.240.400.701 Levocardia
-	C16.131.240.400.715 Long QT Syndrome
-	C16.131.240.400.715.070 Andersen Syndrome
-	C16.131.240.400.715.440 Jervell-Lange Nielsen Syndrome
-	C16.131.240.400.715.720 Romano-Ward Syndrome
-	C16.131.240.400.720 Marfan Syndrome
-	C16.131.240.400.784 Noonan Syndrome
-	C16.131.240.400.849 Tetralogy of Fallot
-	C16.131.240.400.915 Transposition of Great Vessels
-	C16.131.240.400.915.300 Double Outlet Right Ventricle
-	C16.131.240.400.920 Tricuspid Atresia
-	C16.131.240.400.960 Trilogy of Fallot
-	C16.131.240.400.970 Turner Syndrome
-	C16.131.240.400.980 Wolff-Parkinson-White Syndrome
-	C16.131.240.850 Vascular Malformations
-	C16.131.240.850.500 Arterio-Arterial Fistula
-	C16.131.240.850.500.500 Bland White Garland Syndrome
-	C16.131.240.850.750 Arteriovenous Malformations

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C16.131.240.850.750.125 Arteriovenous Fistula
-	C16.131.240.850.750.125.500 Carotid-Cavernous Sinus Fistula
-	C16.131.240.850.750.295 Intracranial Arteriovenous Malformations
-	C16.131.240.850.750.295.500 Vein of Galen Malformations
-	C16.131.240.850.875 Central Nervous System Vascular Malformations
-	C16.131.240.850.875.500 Intracranial Arteriovenous Malformations
-	C16.131.240.850.875.500.500 Vein of Galen Malformations
-	C16.131.240.850.890 May-Thurner Syndrome
-	C16.131.240.850.906 Pulmonary Atresia
-	C16.131.240.850.937 Scimitar Syndrome
-	C16.131.240.850.952 Single Umbilical Artery
-	C16.131.240.850.968 Telangiectasia, Hereditary Hemorrhagic
-	C16.131.260 Chromosome Disorders
-	C16.131.260.019 22q11 Deletion Syndrome
-	C16.131.260.019.500 DiGeorge Syndrome
-	C16.131.260.040 Angelman Syndrome
-	C16.131.260.080 Beckwith-Wiedemann Syndrome
-	C16.131.260.090 Branchio-Oto-Renal Syndrome
-	C16.131.260.190 Cri-du-Chat Syndrome
-	C16.131.260.210 De Lange Syndrome
-	C16.131.260.260 Down Syndrome
-	C16.131.260.380 Holoprosencephaly
-	C16.131.260.440 Jacobsen Distal 11q Deletion Syndrome
-	C16.131.260.700 Prader-Willi Syndrome
-	C16.131.260.790 Rubinstein-Taybi Syndrome
-	C16.131.260.830 Sex Chromosome Disorders
-	C16.131.260.830.300 Fragile X Syndrome
-	C16.131.260.830.670 Orofaciodigital Syndromes
-	C16.131.260.830.835 Sex Chromosome Disorders of Sex Development
-	C16.131.260.830.835.249 Gonadal Dysgenesis, Mixed
-	C16.131.260.830.835.500 Klinefelter Syndrome
-	C16.131.260.830.835.750 Turner Syndrome
-	C16.131.260.870 Silver-Russell Syndrome
-	C16.131.260.887 Smith-Magenis Syndrome
-	C16.131.260.905 Sotos Syndrome
-	C16.131.260.940 WAGR Syndrome

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	C16.131.260.970	Williams Syndrome
-	C16.131.260.985	Wolf-Hirschhorn Syndrome
-	C16.131.287	Congenital Microtia
-	C16.131.314	Digestive System Abnormalities
New Heading	<b>C16.131.314.047</b>	<b>Anorectal Malformations</b>
-	C16.131.314.094	Anus, Imperforate
-	C16.131.314.125	Biliary Atresia
-	C16.131.314.184	Choledochal Cyst
-	C16.131.314.184.500	Caroli Disease
-	C16.131.314.244	Diaphragmatic Eventration
-	C16.131.314.330	Esophageal Atresia
-	C16.131.314.439	Hirschsprung Disease
-	C16.131.314.466	Intestinal Atresia
-	C16.131.314.556	Meckel Diverticulum
-	C16.131.384	Eye Abnormalities
-	C16.131.384.079	Aniridia
-	C16.131.384.079.950	WAGR Syndrome
-	C16.131.384.159	Anophthalmos
-	C16.131.384.190	Blepharophimosis
-	C16.131.384.282	Coloboma
-	C16.131.384.405	Ectopia Lentis
-	C16.131.384.442	Fraser Syndrome
-	C16.131.384.480	Hydrophthalmos
-	C16.131.384.666	Microphthalmos
-	C16.131.384.725	Persistent Hyperplastic Primary Vitreous
-	C16.131.384.784	Retinal Dysplasia
-	C16.131.433	Hernias, Diaphragmatic, Congenital
-	C16.131.482	Lymphatic Abnormalities
-	C16.131.482.249	22q11 Deletion Syndrome
-	C16.131.482.249.500	DiGeorge Syndrome
-	C16.131.482.500	Lymphangiectasis, Intestinal
-	C16.131.621	Musculoskeletal Abnormalities
-	C16.131.621.077	Arthrogryposis
-	C16.131.621.142	Campomelic Dysplasia
-	C16.131.621.174	Cervical Rib Syndrome

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C16.131.621.207                      Craniofacial Abnormalities
-	C16.131.621.207.103                      22q11 Deletion Syndrome
-	C16.131.621.207.103.500                      DiGeorge Syndrome
-	C16.131.621.207.207                      Cleidocranial Dysplasia
-	C16.131.621.207.231                      Craniofacial Dysostosis
-	C16.131.621.207.231.427                      Hallermann's Syndrome
-	C16.131.621.207.231.480                      Hypertelorism
-	C16.131.621.207.231.576                      Mandibulofacial Dysostosis
-	C16.131.621.207.231.576.410                      Goldenhar Syndrome
-	C16.131.621.207.240                      Craniosynostoses
-	C16.131.621.207.240.100                      Acrocephalosyndactylia
-	C16.131.621.207.410                      Holoprosencephaly
-	C16.131.621.207.525                      LEOPARD Syndrome
-	C16.131.621.207.532                      Megalencephaly
-	C16.131.621.207.532.500                      Hemimegalencephaly
-	C16.131.621.207.540                      Maxillofacial Abnormalities
-	C16.131.621.207.540.170                      Cherubism
-	C16.131.621.207.540.315                      Dentofacial Deformities
-	C16.131.621.207.540.460                      Jaw Abnormalities
-	C16.131.621.207.540.460.185                      Cleft Palate
-	C16.131.621.207.540.460.457                      Micrognathism
-	C16.131.621.207.540.460.606                      Pierre Robin Syndrome
-	C16.131.621.207.540.460.655                      Prognathism
-	C16.131.621.207.540.460.827                      Retrognathia
-	C16.131.621.207.620                      Microcephaly
-	C16.131.621.207.620.500                      Porencephaly
-	C16.131.621.207.690                      Noonan Syndrome
-	C16.131.621.207.700                      Orofaciodigital Syndromes
-	C16.131.621.207.707                      Plagiocephaly
-	C16.131.621.207.707.624                      Plagiocephaly, Nonsynostotic
-	C16.131.621.207.720                      Platybasia
-	C16.131.621.207.850                      Rubinstein-Taybi Syndrome
-	C16.131.621.386                      Funnel Chest
-	C16.131.621.417                      Gastroschisis
-	C16.131.621.445                      Hajdu-Cheney Syndrome
-	C16.131.621.449                      Hip Dislocation, Congenital

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	C16.131.621.551	Klippel-Feil Syndrome
-	C16.131.621.568	Laryngomalacia
-	C16.131.621.585	Limb Deformities, Congenital
-	C16.131.621.585.174	Arachnodactyly
-	C16.131.621.585.262	Brachydactyly
-	C16.131.621.585.350	Ectromelia
Old Tree	C16.131.621.585.380	Foot Deformities, Congenital
Old Tree	C16.131.621.585.380.500	Clubfoot
Old Tree	C16.131.621.585.425	Hand Deformities, Congenital
-	C16.131.621.585.512	Lower Extremity Deformities, Congenital
New Tree	C16.131.621.585.512.500	Foot Deformities, Congenital
New Heading	<b>C16.131.621.585.512.500.681</b>	<b>Talipes</b>
New Tree	C16.131.621.585.512.500.681.063	Clubfoot
New Tree	C16.131.621.585.512.500.681.125	Equinus Deformity
New Tree	C16.131.621.585.512.500.681.250	Flatfoot
New Heading	<b>C16.131.621.585.512.500.681.500</b>	<b>Talipes Cavus</b>
New Heading	<b>C16.131.621.585.512.500.787</b>	<b>Tarsal Coalition</b>
-	C16.131.621.585.600	Polydactyly
-	C16.131.621.585.600.374	Pallister-Hall Syndrome
-	C16.131.621.585.600.750	Short Rib-Polydactyly Syndrome
-	C16.131.621.585.620	Proteus Syndrome
-	C16.131.621.585.800	Syndactyly
-	C16.131.621.585.800.100	Acrocephalosyndactyly
New Tree	C16.131.621.585.800.428	Fraser Syndrome
-	C16.131.621.585.800.756	Poland Syndrome
-	C16.131.621.585.984	Thanatophoric Dysplasia
-	C16.131.621.585.988	Upper Extremity Deformities, Congenital
New Tree	C16.131.621.585.988.500	Hand Deformities, Congenital
-	C16.131.621.745	Pectus Carinatum
-	C16.131.621.906	Synostosis

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	C16.131.621.906.181	Antley-Bixler Syndrome Phenotype
-	C16.131.621.906.364	Craniosynostoses
-	C16.131.621.906.364.100	Acrocephalosyndactylia
-	C16.131.621.906.819	Syndactyly
-	C16.131.621.906.819.100	Acrocephalosyndactylia
New Tree	<a href="#">C16.131.621.906.819.428</a>	<a href="#">Fraser Syndrome</a>
-	C16.131.621.906.819.756	Poland Syndrome
New Heading	<b>C16.131.621.906.909</b>	<b>Tarsal Coalition</b>
-	C16.131.621.953	Tracheobronchomalacia
-	C16.131.621.953.249	Bronchomalacia
-	C16.131.621.953.500	Tracheomalacia
-	C16.131.666	Nervous System Malformations
-	C16.131.666.034	Agenesis of Corpus Callosum
-	C16.131.666.034.500	Acrocallosal Syndrome
-	C16.131.666.034.687	Aicardi Syndrome
-	C16.131.666.034.875	Holoprosencephaly
-	C16.131.666.142	Central Nervous System Cysts
-	C16.131.666.142.100	Arachnoid Cysts
-	C16.131.666.142.200	Colloid Cysts
-	C16.131.666.190	Central Nervous System Vascular Malformations
-	C16.131.666.190.100	Central Nervous System Venous Angioma
-	C16.131.666.190.200	Hemangioma, Cavernous, Central Nervous System
-	C16.131.666.190.500	Intracranial Arteriovenous Malformations
-	C16.131.666.190.500.500	Vein of Galen Malformations
-	C16.131.666.190.800	Sinus Pericranii
-	C16.131.666.205	Dandy-Walker Syndrome
-	C16.131.666.300	Hereditary Sensory and Motor Neuropathy
-	C16.131.666.300.099	Alstrom Syndrome
-	C16.131.666.300.200	Charcot-Marie-Tooth Disease
-	C16.131.666.300.490	Giant Axonal Neuropathy
-	C16.131.666.300.780	Refsum Disease
-	C16.131.666.300.820	Spastic Paraplegia, Hereditary
-	C16.131.666.310	Hereditary Sensory and Autonomic Neuropathies
-	C16.131.666.310.309	Dysautonomia, Familial



## MeSH Tree Changes for 2017

Type	Tree - heading
-	C16.131.666.450 Hydranencephaly
-	C16.131.666.507 Malformations of Cortical Development
-	C16.131.666.507.400 Malformations of Cortical Development, Group I
-	C16.131.666.507.400.249 Megalencephaly
-	C16.131.666.507.400.249.500 Hemimegalencephaly
-	C16.131.666.507.400.500 Microcephaly
-	C16.131.666.507.400.750 Tuberous Sclerosis
-	C16.131.666.507.450 Malformations of Cortical Development, Group II
-	C16.131.666.507.450.230 Classical Lissencephalies and Subcortical Band Heterotopias
-	C16.131.666.507.450.249 Cobblestone Lissencephaly
-	C16.131.666.507.450.499 Lissencephaly
-	C16.131.666.507.450.499.230 Classical Lissencephalies and Subcortical Band Heterotopias
-	C16.131.666.507.450.499.249 Cobblestone Lissencephaly
-	C16.131.666.507.450.499.249.500 Walker-Warburg Syndrome
-	C16.131.666.507.450.750 Periventricular Nodular Heterotopia
-	C16.131.666.507.500 Malformations of Cortical Development, Group III
-	C16.131.666.507.500.500 Polymicrogyria
-	C16.131.666.507.500.625 Porencephaly
-	C16.131.666.507.500.750 Schizencephaly
-	C16.131.666.680 Neural Tube Defects
-	C16.131.666.680.196 Anencephaly
-	C16.131.666.680.291 Arnold-Chiari Malformation
-	C16.131.666.680.488 Encephalocele
-	C16.131.666.680.598 Meningocele
-	C16.131.666.680.610 Meningomyelocele
-	C16.131.666.680.705 Pentalogy of Cantrell
-	C16.131.666.680.800 Spinal Dysraphism
-	C16.131.666.680.800.730 Spina Bifida Cystica
-	C16.131.666.680.800.750 Spina Bifida Occulta
-	C16.131.666.845 Septo-Optic Dysplasia
-	C16.131.740 Respiratory System Abnormalities
-	C16.131.740.195 Bronchogenic Cyst
-	C16.131.740.214 Bronchopulmonary Sequestration
-	C16.131.740.271 Choanal Atresia

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C16.131.740.290 Cystic Adenomatoid Malformation of Lung, Congenital
-	C16.131.740.501 Kartagener Syndrome
-	C16.131.740.650 Laryngocele
-	C16.131.740.658 Laryngostenosis
-	C16.131.740.815 Scimitar Syndrome
-	C16.131.740.830 Tracheobronchomegaly
-	C16.131.810 Situs Inversus
-	C16.131.810.250 Dextrocardia
-	C16.131.810.250.500 Kartagener Syndrome
-	C16.131.810.700 Levocardia
-	C16.131.831 Skin Abnormalities
-	C16.131.831.066 Acrodermatitis
-	C16.131.831.108 Carney Complex
-	C16.131.831.150 Dyskeratosis Congenita
-	C16.131.831.350 Ectodermal Dysplasia
-	C16.131.831.350.198 Ectodermal Dysplasia 1, Anhidrotic
-	C16.131.831.350.298 Ectodermal Dysplasia 3, Anhidrotic
-	C16.131.831.350.348 Ectodermal Dysplasia, Hypohidrotic, Autosomal Recessive
-	C16.131.831.350.398 Ellis-Van Creveld Syndrome
-	C16.131.831.350.424 Focal Dermal Hypoplasia
-	C16.131.831.350.712 Neurocutaneous Syndromes
-	C16.131.831.350.856 Pachyonychia Congenita
-	C16.131.831.350.856.500 Steatocystoma Multiplex
-	C16.131.831.428 Ehlers-Danlos Syndrome
-	C16.131.831.493 Epidermolysis Bullosa
-	C16.131.831.493.080 Epidermolysis Bullosa Acquisita
-	C16.131.831.493.160 Epidermolysis Bullosa Dystrophica
-	C16.131.831.493.170 Epidermolysis Bullosa, Junctional
-	C16.131.831.493.180 Epidermolysis Bullosa Simplex
-	C16.131.831.512 Ichthyosis
-	C16.131.831.512.400 Ichthyosiform Erythroderma, Congenital
-	C16.131.831.512.400.375 Hyperkeratosis, Epidermolytic
-	C16.131.831.512.400.410 Ichthyosis, Lamellar
-	C16.131.831.512.400.705 Netherton Syndrome
-	C16.131.831.512.408 Ichthyosis Bullosa of Siemens

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C16.131.831.512.410 Ichthyosis Vulgaris
-	C16.131.831.512.420 Ichthyosis, X-Linked
-	C16.131.831.512.723 Sjogren-Larsson Syndrome
-	C16.131.831.580 Incontinentia Pigmenti
-	C16.131.831.675 Port-Wine Stain
-	C16.131.831.720 Prolidase Deficiency
-	C16.131.831.766 Pseudoxanthoma Elasticum
-	C16.131.831.775 Rothmund-Thomson Syndrome
-	C16.131.831.812 Sclerema Neonatorum
-	C16.131.831.874 Trichothiodystrophy Syndromes
-	C16.131.831.936 Xeroderma Pigmentosum
-	C16.131.850 Stomatognathic System Abnormalities
-	C16.131.850.500 Maxillofacial Abnormalities
-	C16.131.850.500.229 Dentofacial Deformities
-	C16.131.850.500.460 Jaw Abnormalities
-	C16.131.850.500.460.185 Cleft Palate
-	C16.131.850.500.460.457 Micrognathism
-	C16.131.850.500.460.606 Pierre Robin Syndrome
-	C16.131.850.500.460.655 Prognathism
-	C16.131.850.500.460.827 Retrognathia
-	C16.131.850.525 Mouth Abnormalities
-	C16.131.850.525.164 Cleft Lip
-	C16.131.850.525.185 Cleft Palate
-	C16.131.850.525.304 Fibromatosis, Gingival
-	C16.131.850.525.480 Macrostomia
-	C16.131.850.525.520 Microstomia
-	C16.131.850.525.955 Velopharyngeal Insufficiency
-	C16.131.850.800 Tooth Abnormalities
-	C16.131.850.800.100 Anodontia
-	C16.131.850.800.250 Dens in Dente
-	C16.131.850.800.255 Dental Enamel Hypoplasia
-	C16.131.850.800.255.500 Amelogenesis Imperfecta
-	C16.131.850.800.260 Dentin Dysplasia
-	C16.131.850.800.270 Dentinogenesis Imperfecta
-	C16.131.850.800.320 Diastema
-	C16.131.850.800.370 Fused Teeth

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C16.131.850.800.600 Odontodysplasia
-	C16.131.850.800.850 Tooth, Supernumerary
-	C16.131.894 Thyroid Dysgenesis
-	C16.131.894.500 Lingual Thyroid
-	C16.131.894.500.500 Lingual Goiter
-	C16.131.939 Urogenital Abnormalities
-	C16.131.939.132 Bladder Exstrophy
-	C16.131.939.258 Cryptorchidism
-	C16.131.939.316 Disorders of Sex Development
-	C16.131.939.316.064 46, XX Disorders of Sex Development
-	C16.131.939.316.064.124 46, XX Testicular Disorders of Sex Development
-	C16.131.939.316.064.249 Gonadal Dysgenesis, 46,XX
-	C16.131.939.316.064.500 Hyperandrogenism
-	C16.131.939.316.096 46, XY Disorders of Sex Development
-	C16.131.939.316.096.500 Androgen-Insensitivity Syndrome
-	C16.131.939.316.096.562 Denys-Drash Syndrome
-	C16.131.939.316.096.624 Frasier Syndrome
-	C16.131.939.316.096.687 Gonadal Dysgenesis, 46,XY
-	C16.131.939.316.096.687.500 Gonadoblastoma
-	C16.131.939.316.096.750 Kallmann Syndrome
-	C16.131.939.316.096.875 WAGR Syndrome
-	C16.131.939.316.129 Adrenogenital Syndrome
-	C16.131.939.316.129.500 Adrenal Hyperplasia, Congenital
-	C16.131.939.316.129.750 Hyperandrogenism
-	C16.131.939.316.309 Gonadal Dysgenesis
-	C16.131.939.316.309.193 Gonadal Dysgenesis, 46,XX
-	C16.131.939.316.309.388 Gonadal Dysgenesis, 46,XY
-	C16.131.939.316.309.388.500 Gonadoblastoma
-	C16.131.939.316.309.391 Gonadal Dysgenesis, Mixed
-	C16.131.939.316.309.631 Sexual Infantilism
-	C16.131.939.316.309.872 Turner Syndrome
-	C16.131.939.316.343 Ovotesticular Disorders of Sex Development
-	C16.131.939.316.795 Sex Chromosome Disorders of Sex Development
-	C16.131.939.316.795.124 Freemartinism
-	C16.131.939.316.795.249 Gonadal Dysgenesis, Mixed
-	C16.131.939.316.795.500 Klinefelter Syndrome

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C16.131.939.316.795.750 Turner Syndrome
-	C16.131.939.374 Epispadias
New Tree	<a href="#">C16.131.939.410</a> <a href="#">Fraser Syndrome</a>
-	C16.131.939.445 Fused Kidney
-	C16.131.939.516 Hypospadias
-	C16.131.939.629 Multicystic Dysplastic Kidney
-	C16.131.939.742 Nephritis, Hereditary
-	C16.131.939.831 Pyelectasis
-	C16.131.939.915 Retrocaval Ureter
-	C16.300 Fetal Diseases
-	C16.300.030 Chorioamnionitis
-	C16.300.050 Echogenic Bowel
-	C16.300.060 Erythroblastosis, Fetal
-	C16.300.060.480 Hydrops Fetalis
-	C16.300.070 Fetal Alcohol Spectrum Disorders
-	C16.300.390 Fetal Growth Retardation
-	C16.300.420 Fetal Hypoxia
-	C16.300.570 Fetal Macrosomia
-	C16.300.580 Meconium Aspiration Syndrome
-	C16.300.790 Nuchal Cord
-	C16.300.895 Pyelectasis
-	C16.320 Genetic Diseases, Inborn
-	C16.320.033 Adrenal Hyperplasia, Congenital
-	C16.320.051 Alagille Syndrome
-	C16.320.060 alpha 1-Antitrypsin Deficiency
-	C16.320.070 Anemia, Hemolytic, Congenital
-	C16.320.070.095 Anemia, Dyserythropoietic, Congenital
-	C16.320.070.100 Anemia, Hemolytic, Congenital Nonspherocytic
-	C16.320.070.150 Anemia, Sickle Cell
-	C16.320.070.150.219 Acute Chest Syndrome
-	C16.320.070.150.440 Hemoglobin SC Disease
-	C16.320.070.150.670 Sickle Cell Trait
-	C16.320.070.365 Elliptocytosis, Hereditary
-	C16.320.070.480 Glucosephosphate Dehydrogenase Deficiency
-	C16.320.070.480.370 Favism

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C16.320.070.490 Hemoglobin C Disease
-	C16.320.070.785 Spherocytosis, Hereditary
-	C16.320.070.875 Thalassemia
-	C16.320.070.875.100 alpha-Thalassemia
-	C16.320.070.875.150 beta-Thalassemia
-	C16.320.070.875.575 delta-Thalassemia
-	C16.320.077 Anemia, Hypoplastic, Congenital
-	C16.320.077.090 Anemia, Diamond-Blackfan
-	C16.320.077.280 Fanconi Anemia
-	C16.320.078 Angioedemas, Hereditary
-	C16.320.080 Ataxia Telangiectasia
-	C16.320.089 Autoimmune Lymphoproliferative Syndrome
-	C16.320.099 Blood Coagulation Disorders, Inherited
-	C16.320.099.037 Activated Protein C Resistance
-	C16.320.099.056 Afibrinogenemia
-	C16.320.099.075 Antithrombin III Deficiency
-	C16.320.099.080 Bernard-Soulier Syndrome
-	C16.320.099.300 Factor V Deficiency
-	C16.320.099.310 Factor VII Deficiency
-	C16.320.099.320 Factor X Deficiency
-	C16.320.099.325 Factor XI Deficiency
-	C16.320.099.330 Factor XII Deficiency
-	C16.320.099.335 Factor XIII Deficiency
-	C16.320.099.417 Gray Platelet Syndrome
-	C16.320.099.500 Hemophilia A
-	C16.320.099.510 Hemophilia B
-	C16.320.099.515 Hermanski-Pudlak Syndrome
-	C16.320.099.550 Hypoprothrombinemias
-	C16.320.099.690 Protein C Deficiency
-	C16.320.099.820 Thrombasthenia
-	C16.320.099.920 von Willebrand Diseases
-	C16.320.099.920.100 von Willebrand Disease, Type 1
-	C16.320.099.920.200 von Willebrand Disease, Type 2
-	C16.320.099.920.300 von Willebrand Disease, Type 3
-	C16.320.099.970 Wiskott-Aldrich Syndrome
-	C16.320.100 Brugada Syndrome

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	C16.320.129	CADASIL
-	C16.320.144	Camurati-Engelmann Syndrome
-	C16.320.160	Cardiomyopathy, Hypertrophic, Familial
-	C16.320.165	CHARGE Syndrome
-	C16.320.170	Cherubism
-	C16.320.180	Chromosome Disorders
-	C16.320.180.019	22q11 Deletion Syndrome
-	C16.320.180.019.500	DiGeorge Syndrome
-	C16.320.180.040	Angelman Syndrome
-	C16.320.180.080	Beckwith-Wiedemann Syndrome
-	C16.320.180.090	Branchio-Oto-Renal Syndrome
-	C16.320.180.190	Cri-du-Chat Syndrome
-	C16.320.180.210	De Lange Syndrome
-	C16.320.180.260	Down Syndrome
-	C16.320.180.380	Holoprosencephaly
-	C16.320.180.440	Jacobsen Distal 11q Deletion Syndrome
-	C16.320.180.700	Prader-Willi Syndrome
-	C16.320.180.790	Rubinstein-Taybi Syndrome
-	C16.320.180.830	Sex Chromosome Disorders
-	C16.320.180.830.300	Fragile X Syndrome
-	C16.320.180.830.670	Orofaciodigital Syndromes
-	C16.320.180.830.835	Sex Chromosome Disorders of Sex Development
-	C16.320.180.830.835.249	Gonadal Dysgenesis, Mixed
-	C16.320.180.830.835.500	Klinefelter Syndrome
-	C16.320.180.830.835.750	Turner Syndrome
-	C16.320.180.870	Silver-Russell Syndrome
-	C16.320.180.887	Smith-Magenis Syndrome
-	C16.320.180.905	Sotos Syndrome
-	C16.320.180.940	WAGR Syndrome
-	C16.320.180.970	Williams Syndrome
-	C16.320.180.985	Wolf-Hirschhorn Syndrome
New Heading	<b>C16.320.184</b>	<b>Ciliopathies</b>
New Tree	<a href="#">C16.320.184.063</a>	<a href="#">Alstrom Syndrome</a>
New Tree	<a href="#">C16.320.184.125</a>	<a href="#">Bardet-Biedl Syndrome</a>

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	<a href="#">C16.320.184.250</a>	<a href="#">Caroli Disease</a>
New Tree	<a href="#">C16.320.184.500</a>	<a href="#">Ciliary Motility Disorders</a>
New Tree	<a href="#">C16.320.184.500.531</a>	<a href="#">Kartagener Syndrome</a>
-	C16.320.185	Costello Syndrome
-	C16.320.190	Cystic Fibrosis
-	C16.320.215	Donohue Syndrome
-	C16.320.240	Dwarfism
-	C16.320.240.500	Achondroplasia
-	C16.320.240.500.500	Thanatophoric Dysplasia
-	C16.320.240.562	Cockayne Syndrome
-	C16.320.240.625	Congenital Hypothyroidism
-	C16.320.240.750	Laron Syndrome
-	C16.320.240.875	Mulibrey Nanism
-	C16.320.240.937	Silver-Russell Syndrome
-	C16.320.290	Eye Diseases, Hereditary
-	C16.320.290.019	Aicardi Syndrome
-	C16.320.290.040	Albinism
-	C16.320.290.040.090	Albinism, Ocular
-	C16.320.290.040.100	Albinism, Oculocutaneous
-	C16.320.290.040.100.400	Hermanski-Pudlak Syndrome
-	C16.320.290.040.600	Piebaldism
-	C16.320.290.078	Aniridia
-	C16.320.290.078.950	WAGR Syndrome
-	C16.320.290.142	Choroideremia
New Heading	<b><a href="#">C16.320.290.152</a></b>	<b><a href="#">Cone-Rod Dystrophies</a></b>
-	C16.320.290.162	Corneal Dystrophies, Hereditary
-	C16.320.290.162.204	Corneal Dystrophy, Juvenile Epithelial of Meesmann
-	C16.320.290.162.410	Fuchs' Endothelial Dystrophy
-	C16.320.290.235	Duane Retraction Syndrome
-	C16.320.290.468	Gyrate Atrophy
-	C16.320.290.564	Optic Atrophies, Hereditary
-	C16.320.290.564.100	Optic Atrophy, Autosomal Dominant
-	C16.320.290.564.400	Optic Atrophy, Hereditary, Leber



## MeSH Tree Changes for 2017

Type	Tree - heading
-	C16.320.290.564.980 Wolfram Syndrome
-	C16.320.290.660 Retinal Dysplasia
-	C16.320.290.684 Retinitis Pigmentosa
-	C16.320.290.684.249 Alstrom Syndrome
-	C16.320.290.684.500 Usher Syndromes
-	C16.320.290.763 Vitelliform Macular Dystrophy
-	C16.320.290.842 Weill-Marchesani Syndrome
New Heading	<b>C16.320.298</b> <b>Familial Multiple Lipomatosis</b>
-	C16.320.306 Frasier Syndrome
-	C16.320.322 Genetic Diseases, X-Linked
-	C16.320.322.030 Aicardi Syndrome
-	C16.320.322.061 Androgen-Insensitivity Syndrome
-	C16.320.322.068 Barth Syndrome
-	C16.320.322.076 Bulbo-Spinal Atrophy, X-Linked
-	C16.320.322.092 Choroideremia
-	C16.320.322.100 Dent Disease
-	C16.320.322.108 Dyskeratosis Congenita
-	C16.320.322.116 Ectodermal Dysplasia 1, Anhidrotic
-	C16.320.322.124 Fabry Disease
-	C16.320.322.186 Focal Dermal Hypoplasia
-	C16.320.322.201 Glycogen Storage Disease Type IIb
-	C16.320.322.217 Glycogen Storage Disease Type VIII
-	C16.320.322.233 Granulomatous Disease, Chronic
-	C16.320.322.235 Hemophilia B
-	C16.320.322.237 Hyper-IgM Immunodeficiency Syndrome, Type 1
-	C16.320.322.241 Ichthyosis, X-Linked
-	C16.320.322.370 Isolated Noncompaction of the Ventricular Myocardium
-	C16.320.322.500 Mental Retardation, X-Linked
-	C16.320.322.500.124 Adrenoleukodystrophy
-	C16.320.322.500.186 Classical Lissencephalies and Subcortical Band Heterotopias
-	C16.320.322.500.249 Coffin-Lowry Syndrome
-	C16.320.322.500.500 Fragile X Syndrome
-	C16.320.322.500.625 Lesch-Nyhan Syndrome
-	C16.320.322.500.687 Menkes Kinky Hair Syndrome

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C16.320.322.500.750 Mucopolysaccharidosis II
-	C16.320.322.500.875 Pyruvate Dehydrogenase Complex Deficiency Disease
-	C16.320.322.500.937 Rett Syndrome
-	C16.320.322.562 Muscular Dystrophy, Duchenne
-	C16.320.322.625 Muscular Dystrophy, Emery-Dreifuss
-	C16.320.322.750 Oculocerebrorenal Syndrome
-	C16.320.322.828 Ornithine Carbamoyltransferase Deficiency Disease
-	C16.320.322.906 Pelizaeus-Merzbacher Disease
-	C16.320.322.937 Wiskott-Aldrich Syndrome
-	C16.320.322.968 X-Linked Combined Immunodeficiency Diseases
-	C16.320.338 Genetic Diseases, Y-Linked
-	C16.320.355 Hajdu-Cheney Syndrome
-	C16.320.365 Hemoglobinopathies
-	C16.320.365.155 Anemia, Sickle Cell
-	C16.320.365.155.219 Acute Chest Syndrome
-	C16.320.365.155.440 Hemoglobin SC Disease
-	C16.320.365.155.668 Sickle Cell Trait
-	C16.320.365.463 Hemoglobin C Disease
-	C16.320.365.826 Thalassemia
-	C16.320.365.826.100 alpha-Thalassemia
-	C16.320.365.826.100.350 Hydrops Fetalis
-	C16.320.365.826.150 beta-Thalassemia
-	C16.320.365.826.575 delta-Thalassemia
-	C16.320.382 Hereditary Autoinflammatory Diseases
-	C16.320.382.250 Behcet Syndrome
-	C16.320.382.500 Cryopyrin-Associated Periodic Syndromes
-	C16.320.382.625 Familial Mediterranean Fever
-	C16.320.382.750 Mevalonate Kinase Deficiency
-	C16.320.400 Heredodegenerative Disorders, Nervous System
-	C16.320.400.024 Alexander Disease
-	C16.320.400.050 Amyloid Neuropathies, Familial
-	C16.320.400.150 Canavan Disease
-	C16.320.400.200 Cockayne Syndrome
-	C16.320.400.330 Dystonia Musculorum Deformans
-	C16.320.400.350 Gerstmann-Straussler-Scheinker Disease
-	C16.320.400.361 Hepatolenticular Degeneration

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C16.320.400.367 Hereditary Central Nervous System Demyelinating Diseases
-	C16.320.400.375 Hereditary Sensory and Motor Neuropathy
-	C16.320.400.375.099 Alstrom Syndrome
-	C16.320.400.375.200 Charcot-Marie-Tooth Disease
-	C16.320.400.375.490 Giant Axonal Neuropathy
-	C16.320.400.375.780 Refsum Disease
-	C16.320.400.375.820 Spastic Paraplegia, Hereditary
-	C16.320.400.415 Hereditary Sensory and Autonomic Neuropathies
-	C16.320.400.415.309 Dysautonomia, Familial
-	C16.320.400.430 Huntington Disease
-	C16.320.400.480 Lafora Disease
-	C16.320.400.525 Mental Retardation, X-Linked
-	C16.320.400.525.124 Adrenoleukodystrophy
-	C16.320.400.525.249 Coffin-Lowry Syndrome
-	C16.320.400.525.500 Fragile X Syndrome
-	C16.320.400.525.625 Lesch-Nyhan Syndrome
-	C16.320.400.525.687 Menkes Kinky Hair Syndrome
-	C16.320.400.525.750 Mucopolysaccharidosis II
-	C16.320.400.525.875 Pyruvate Dehydrogenase Complex Deficiency Disease
-	C16.320.400.525.937 Rett Syndrome
-	C16.320.400.540 Myotonia Congenita
-	C16.320.400.542 Myotonic Dystrophy
-	C16.320.400.550 Neuroacanthocytosis
-	C16.320.400.560 Neurofibromatoses
-	C16.320.400.560.400 Neurofibromatosis 1
-	C16.320.400.560.700 Neurofibromatosis 2
-	C16.320.400.600 Neuronal Ceroid-Lipofuscinoses
-	C16.320.400.630 Optic Atrophies, Hereditary
-	C16.320.400.630.100 Optic Atrophy, Autosomal Dominant
-	C16.320.400.630.400 Optic Atrophy, Hereditary, Leber
-	C16.320.400.630.980 Wolfram Syndrome
-	C16.320.400.650 Pantothenate Kinase-Associated Neurodegeneration
-	C16.320.400.765 Spinal Muscular Atrophies of Childhood
-	C16.320.400.780 Spinocerebellar Degenerations
-	C16.320.400.780.200 Friedreich Ataxia
-	C16.320.400.780.500 Myoclonic Cerebellar Dyssynergia

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C16.320.400.780.750 Olivopontocerebellar Atrophies
-	C16.320.400.780.875 Spinocerebellar Ataxias
-	C16.320.400.780.875.500 Machado-Joseph Disease
-	C16.320.400.820 Tourette Syndrome
-	C16.320.400.880 Tuberous Sclerosis
-	C16.320.400.940 Unverricht-Lundborg Syndrome
-	C16.320.413 Hyper-IgM Immunodeficiency Syndrome
-	C16.320.427 Hyperthyroxinemia, Familial Dysalbuminemic
-	C16.320.467 Kallmann Syndrome
-	C16.320.480 Kartagener Syndrome
-	C16.320.495 Lennox Gastaut Syndrome
-	C16.320.510 Loeys-Dietz Syndrome
-	C16.320.540 Marfan Syndrome
-	C16.320.565 Metabolism, Inborn Errors
-	C16.320.565.100 Amino Acid Metabolism, Inborn Errors
-	C16.320.565.100.102 Albinism
-	C16.320.565.100.102.090 Albinism, Ocular
-	C16.320.565.100.102.100 Albinism, Oculocutaneous
-	C16.320.565.100.102.100.400 Hermanski-Pudlak Syndrome
-	C16.320.565.100.102.600 Piebaldism
-	C16.320.565.100.187 Alkaptonuria
-	C16.320.565.100.477 Hyperglycinemia, Nonketotic
-	C16.320.565.100.480 Hyperhomocysteinemia
-	C16.320.565.100.480.500 Homocystinuria
-	C16.320.565.100.544 Hyperlysinemias
-	C16.320.565.100.608 Maple Syrup Urine Disease
-	C16.320.565.100.614 Multiple Acyl Coenzyme A Dehydrogenase Deficiency
-	C16.320.565.100.620 Multiple Carboxylase Deficiency
-	C16.320.565.100.620.100 Biotinidase Deficiency
-	C16.320.565.100.620.380 Holocarboxylase Synthetase Deficiency
-	C16.320.565.100.766 Phenylketonurias
-	C16.320.565.100.766.500 Phenylketonuria, Maternal
-	C16.320.565.100.794 Prolidase Deficiency
-	C16.320.565.100.823 Propionic Acidemia
-	C16.320.565.100.880 Tyrosinemias
-	C16.320.565.100.940 Urea Cycle Disorders, Inborn

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C16.320.565.100.940.124 Argininosuccinic Aciduria
-	C16.320.565.100.940.249 Carbamoyl-Phosphate Synthase I Deficiency Disease
-	C16.320.565.100.940.374 Citrullinemia
-	C16.320.565.100.940.500 Hyperargininemia
-	C16.320.565.100.940.750 Ornithine Carbamoyltransferase Deficiency Disease
-	C16.320.565.151 Amino Acid Transport Disorders, Inborn
-	C16.320.565.151.355 Hartnup Disease
-	C16.320.565.151.600 Oculocerebrorenal Syndrome
-	C16.320.565.176 Amyloidosis, Familial
-	C16.320.565.176.050 Amyloid Neuropathies, Familial
-	C16.320.565.176.160 Cerebral Amyloid Angiopathy, Familial
-	C16.320.565.189 Brain Diseases, Metabolic, Inborn
-	C16.320.565.189.084 Adrenoleukodystrophy
-	C16.320.565.189.168 Cerebral Amyloid Angiopathy, Familial
-	C16.320.565.189.320 Galactosemias
-	C16.320.565.189.355 Hartnup Disease
-	C16.320.565.189.360 Hepatolenticular Degeneration
-	C16.320.565.189.362 Hereditary Central Nervous System Demyelinating Diseases
-	C16.320.565.189.362.250 Adrenoleukodystrophy
-	C16.320.565.189.362.312 Alexander Disease
-	C16.320.565.189.362.375 Canavan Disease
-	C16.320.565.189.362.500 Leukodystrophy, Globoid Cell
-	C16.320.565.189.362.550 Leukodystrophy, Metachromatic
-	C16.320.565.189.362.775 Pelizaeus-Merzbacher Disease
-	C16.320.565.189.365 Homocystinuria
-	C16.320.565.189.375 Hyperglycinemia, Nonketotic
-	C16.320.565.189.380 Hyperlysinemias
-	C16.320.565.189.412 Leigh Disease
-	C16.320.565.189.425 Lesch-Nyhan Syndrome
-	C16.320.565.189.435 Lysosomal Storage Diseases, Nervous System
-	C16.320.565.189.435.295 Fucosidosis
-	C16.320.565.189.435.340 Glycogen Storage Disease Type II
-	C16.320.565.189.435.590 Mucopolidoses
-	C16.320.565.189.435.810 Sialic Acid Storage Disease

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C16.320.565.189.435.825 Sphingolipidoses
-	C16.320.565.189.435.825.200 Fabry Disease
-	C16.320.565.189.435.825.250 Farber Lipogranulomatosis
-	C16.320.565.189.435.825.300 Gangliosidoses
-	C16.320.565.189.435.825.300.300 Gangliosidoses, GM2
-	C16.320.565.189.435.825.300.300.249 Sandhoff Disease
-	C16.320.565.189.435.825.300.300.500 Tay-Sachs Disease
-	C16.320.565.189.435.825.300.300.750 Tay-Sachs Disease, AB Variant
-	C16.320.565.189.435.825.300.400 Gangliosidosis, GM1
-	C16.320.565.189.435.825.400 Gaucher Disease
-	C16.320.565.189.435.825.590 Leukodystrophy, Globoid Cell
-	C16.320.565.189.435.825.700 Niemann-Pick Diseases
-	C16.320.565.189.435.825.700.500 Niemann-Pick Disease, Type A
-	C16.320.565.189.435.825.700.750 Niemann-Pick Disease, Type B
-	C16.320.565.189.435.825.700.875 Niemann-Pick Disease, Type C
-	C16.320.565.189.435.825.775 Sea-Blue Histiocyte Syndrome
-	C16.320.565.189.435.825.850 Sulfatidosis
-	C16.320.565.189.435.825.850.500 Leukodystrophy, Metachromatic
-	C16.320.565.189.435.825.850.750 Multiple Sulfatase Deficiency Disease
-	C16.320.565.189.520 Maple Syrup Urine Disease
-	C16.320.565.189.535 MELAS Syndrome
-	C16.320.565.189.540 Menkes Kinky Hair Syndrome
-	C16.320.565.189.545 MERRF Syndrome
-	C16.320.565.189.593 Mevalonate Kinase Deficiency
-	C16.320.565.189.640 Oculocerebrorenal Syndrome
-	C16.320.565.189.687 Phenylketonurias
-	C16.320.565.189.687.500 Phenylketonuria, Maternal
-	C16.320.565.189.725 Pyruvate Carboxylase Deficiency Disease
-	C16.320.565.189.750 Pyruvate Dehydrogenase Complex Deficiency Disease
-	C16.320.565.189.813 Refsum Disease
-	C16.320.565.189.844 Refsum Disease, Infantile
-	C16.320.565.189.875 Tyrosinemias
-	C16.320.565.189.937 Urea Cycle Disorders, Inborn
-	C16.320.565.189.937.124 Argininosuccinic Aciduria
-	C16.320.565.189.937.249 Carbamoyl-Phosphate Synthase I Deficiency Disease

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C16.320.565.189.937.374 Citrullinemia
-	C16.320.565.189.937.500 Hyperargininemia
-	C16.320.565.189.937.750 Ornithine Carbamoyltransferase Deficiency Disease
-	C16.320.565.189.968 Zellweger Syndrome
-	C16.320.565.202 Carbohydrate Metabolism, Inborn Errors
-	C16.320.565.202.125 Congenital Disorders of Glycosylation
-	C16.320.565.202.251 Fructose Metabolism, Inborn Errors
-	C16.320.565.202.251.221 Fructose-1,6-Diphosphatase Deficiency
-	C16.320.565.202.251.271 Fructose Intolerance
-	C16.320.565.202.303 Fucosidosis
-	C16.320.565.202.355 Galactosemias
-	C16.320.565.202.402 Glucosephosphate Dehydrogenase Deficiency
-	C16.320.565.202.449 Glycogen Storage Disease
-	C16.320.565.202.449.448 Glycogen Storage Disease Type I
-	C16.320.565.202.449.500 Glycogen Storage Disease Type II
-	C16.320.565.202.449.510 Glycogen Storage Disease Type IIb
-	C16.320.565.202.449.520 Glycogen Storage Disease Type III
-	C16.320.565.202.449.540 Glycogen Storage Disease Type IV
-	C16.320.565.202.449.560 Glycogen Storage Disease Type V
-	C16.320.565.202.449.580 Glycogen Storage Disease Type VI
-	C16.320.565.202.449.600 Glycogen Storage Disease Type VII
-	C16.320.565.202.449.620 Glycogen Storage Disease Type VIII
-	C16.320.565.202.460 Hyperoxaluria, Primary
-	C16.320.565.202.589 Lactose Intolerance
-	C16.320.565.202.607 Mannosidase Deficiency Diseases
-	C16.320.565.202.607.500 alpha-Mannosidosis
-	C16.320.565.202.607.750 beta-Mannosidosis
-	C16.320.565.202.670 Mucopolipidoses
-	C16.320.565.202.715 Mucopolysaccharidoses
-	C16.320.565.202.715.640 Mucopolysaccharidosis I
-	C16.320.565.202.715.645 Mucopolysaccharidosis II
-	C16.320.565.202.715.650 Mucopolysaccharidosis III
-	C16.320.565.202.715.655 Mucopolysaccharidosis IV
-	C16.320.565.202.715.670 Mucopolysaccharidosis VI
-	C16.320.565.202.715.675 Mucopolysaccharidosis VII
-	C16.320.565.202.720 Multiple Carboxylase Deficiency

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C16.320.565.202.720.100 Biotinidase Deficiency
-	C16.320.565.202.720.380 Holocarboxylase Synthetase Deficiency
-	C16.320.565.202.810 Pyruvate Metabolism, Inborn Errors
-	C16.320.565.202.810.444 Leigh Disease
-	C16.320.565.202.810.666 Pyruvate Carboxylase Deficiency Disease
-	C16.320.565.202.810.766 Pyruvate Dehydrogenase Complex Deficiency Disease
-	C16.320.565.240 Cytochrome-c Oxidase Deficiency
-	C16.320.565.300 Hyperbilirubinemia, Hereditary
-	C16.320.565.300.281 Crigler-Najjar Syndrome
-	C16.320.565.300.528 Gilbert Disease
-	C16.320.565.300.764 Jaundice, Chronic Idiopathic
-	C16.320.565.398 Lipid Metabolism, Inborn Errors
-	C16.320.565.398.224 Barth Syndrome
-	C16.320.565.398.450 Hyperlipidemia, Familial Combined
-	C16.320.565.398.465 Hyperlipoproteinemia Type I
-	C16.320.565.398.481 Hyperlipoproteinemia Type II
-	C16.320.565.398.483 Hyperlipoproteinemia Type III
-	C16.320.565.398.487 Hyperlipoproteinemia Type IV
-	C16.320.565.398.493 Hyperlipoproteinemia Type V
-	C16.320.565.398.500 Hypolipoproteinemias
-	C16.320.565.398.500.330 Hypoalphalipoproteinemias
-	C16.320.565.398.500.330.500 Lecithin Acyltransferase Deficiency
-	C16.320.565.398.500.330.500 Lecithin Cholesterol Acyltransferase Deficiency
-	C16.320.565.398.500.330.750 Tangier Disease
-	C16.320.565.398.500.440 Hypobetalipoproteinemias
-	C16.320.565.398.500.440.500 Abetalipoproteinemia
-	C16.320.565.398.641 Lipidoses
-	C16.320.565.398.641.201 Cholesterol Ester Storage Disease
-	C16.320.565.398.641.201.500 Wolman Disease
-	C16.320.565.398.641.509 Neuronal Ceroid-Lipofuscinoses
-	C16.320.565.398.641.723 Sjogren-Larsson Syndrome
-	C16.320.565.398.641.803 Sphingolipidoses
-	C16.320.565.398.641.803.300 Fabry Disease
-	C16.320.565.398.641.803.325 Farber Lipogranulomatosis
-	C16.320.565.398.641.803.350 Gangliosidoses



## MeSH Tree Changes for 2017

Type	Tree - heading
-	C16.320.565.398.641.803.350.300 Gangliosidoses, GM2
-	C16.320.565.398.641.803.350.300.700 Sandhoff Disease
-	C16.320.565.398.641.803.350.300.850 Tay-Sachs Disease
-	C16.320.565.398.641.803.350.300.925 Tay-Sachs Disease, AB Variant
-	C16.320.565.398.641.803.350.360 Gangliosidosis, GM1
-	C16.320.565.398.641.803.441 Gaucher Disease
-	C16.320.565.398.641.803.585 Leukodystrophy, Globoid Cell
-	C16.320.565.398.641.803.730 Niemann-Pick Diseases
-	C16.320.565.398.641.803.730.500 Niemann-Pick Disease, Type A
-	C16.320.565.398.641.803.730.750 Niemann-Pick Disease, Type B
-	C16.320.565.398.641.803.730.875 Niemann-Pick Disease, Type C
-	C16.320.565.398.641.803.850 Sea-Blue Histiocyte Syndrome
-	C16.320.565.398.641.803.925 Sulfatidosis
-	C16.320.565.398.641.803.925.500 Leukodystrophy, Metachromatic
-	C16.320.565.398.641.803.925.750 Multiple Sulfatase Deficiency Disease
-	C16.320.565.398.745 Lipodystrophy, Congenital Generalized
-	C16.320.565.398.850 Smith-Lemli-Opitz Syndrome
-	C16.320.565.398.925 Xanthomatosis, Cerebrotendinous
-	C16.320.565.595 Lysosomal Storage Diseases
-	C16.320.565.595.100 Aspartylglucosaminuria
-	C16.320.565.595.201 Cholesterol Ester Storage Disease
-	C16.320.565.595.201.500 Wolman Disease
-	C16.320.565.595.377 Cystinosis
-	C16.320.565.595.554 Lysosomal Storage Diseases, Nervous System
-	C16.320.565.595.554.295 Fucosidosis
-	C16.320.565.595.554.340 Glycogen Storage Disease Type II
-	C16.320.565.595.554.590 Mucopolidoses
-	C16.320.565.595.554.810 Sialic Acid Storage Disease
-	C16.320.565.595.554.825 Sphingolipidoses
-	C16.320.565.595.554.825.200 Fabry Disease
-	C16.320.565.595.554.825.250 Farber Lipogranulomatosis
-	C16.320.565.595.554.825.300 Gangliosidoses
-	C16.320.565.595.554.825.300.300 Gangliosidoses, GM2
-	C16.320.565.595.554.825.300.300.800 Sandhoff Disease
-	C16.320.565.595.554.825.300.300.840 Tay-Sachs Disease
-	C16.320.565.595.554.825.300.300.920 Tay-Sachs Disease, AB Variant

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C16.320.565.595.554.825.300.400                      Gangliosidosis, GM1
-	C16.320.565.595.554.825.400                                      Gaucher Disease
-	C16.320.565.595.554.825.590                                      Leukodystrophy, Globoid Cell
-	C16.320.565.595.554.825.700                                      Niemann-Pick Diseases
-	C16.320.565.595.554.825.700.500                                      Niemann-Pick Disease, Type A
-	C16.320.565.595.554.825.700.750                                      Niemann-Pick Disease, Type B
-	C16.320.565.595.554.825.700.875                                      Niemann-Pick Disease, Type C
-	C16.320.565.595.554.825.775                                      Sea-Blue Histiocyte Syndrome
-	C16.320.565.595.554.825.850                                      Sulfatidosis
-	C16.320.565.595.554.825.850.500                                      Leukodystrophy, Metachromatic
-	C16.320.565.595.554.825.850.750                                      Multiple Sulfatase Deficiency Disease
-	C16.320.565.595.577                                      Mannosidase Deficiency Diseases
-	C16.320.565.595.577.500                                      alpha-Mannosidosis
-	C16.320.565.595.577.750                                      beta-Mannosidosis
-	C16.320.565.595.600                                      Mucopolysaccharidoses
-	C16.320.565.595.600.640                                      Mucopolysaccharidosis I
-	C16.320.565.595.600.645                                      Mucopolysaccharidosis II
-	C16.320.565.595.600.650                                      Mucopolysaccharidosis III
-	C16.320.565.595.600.655                                      Mucopolysaccharidosis IV
-	C16.320.565.595.600.670                                      Mucopolysaccharidosis VI
-	C16.320.565.595.600.675                                      Mucopolysaccharidosis VII
-	C16.320.565.595.800                                      Pycnodysostosis
-	C16.320.565.618                                      Metal Metabolism, Inborn Errors
-	C16.320.565.618.337                                      Hemochromatosis
-	C16.320.565.618.403                                      Hepatolenticular Degeneration
-	C16.320.565.618.482                                      Hypophosphatasia
-	C16.320.565.618.544                                      Hypophosphatemia, Familial
-	C16.320.565.618.544.500                                      Familial Hypophosphatemic Rickets
-	C16.320.565.618.590                                      Menkes Kinky Hair Syndrome
-	C16.320.565.618.711                                      Paralysis, Familial Periodic
-	C16.320.565.618.711.550                                      Hypokalemic Periodic Paralysis
-	C16.320.565.618.711.600                                      Paralysis, Hyperkalemic Periodic
-	C16.320.565.618.815                                      Pseudohypoparathyroidism
-	C16.320.565.618.815.815                                      Pseudopseudohypoparathyroidism
-	C16.320.565.663                                      Peroxisomal Disorders
-	C16.320.565.663.050                                      Acatlasia

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C16.320.565.663.100 Adrenoleukodystrophy
-	C16.320.565.663.265 Chondrodysplasia Punctata, Rhizomelic
-	C16.320.565.663.430 Mevalonate Kinase Deficiency
-	C16.320.565.663.760 Refsum Disease
-	C16.320.565.663.865 Refsum Disease, Infantile
-	C16.320.565.663.970 Zellweger Syndrome
-	C16.320.565.753 Progeria
-	C16.320.565.798 Purine-Pyrimidine Metabolism, Inborn Errors
-	C16.320.565.798.183 Dihydropyrimidine Dehydrogenase Deficiency
-	C16.320.565.798.368 Gout
-	C16.320.565.798.368.410 Arthritis, Gouty
-	C16.320.565.798.594 Lesch-Nyhan Syndrome
-	C16.320.565.861 Renal Tubular Transport, Inborn Errors
-	C16.320.565.861.093 Acidosis, Renal Tubular
-	C16.320.565.861.271 Dent Disease
-	C16.320.565.861.450 Fanconi Syndrome
-	C16.320.565.861.491 Gitelman Syndrome
-	C16.320.565.861.532 Glycosuria, Renal
-	C16.320.565.861.647 Hypophosphatemia, Familial
-	C16.320.565.861.647.500 Familial Hypophosphatemic Rickets
-	C16.320.565.861.698 Liddle Syndrome
-	C16.320.565.861.750 Oculocerebrorenal Syndrome
-	C16.320.565.861.770 Pseudohypoaldosteronism
-	C16.320.565.861.885 Renal Aminoacidurias
-	C16.320.565.861.885.250 Cystinuria
-	C16.320.565.861.885.457 Hartnup Disease
-	C16.320.565.925 Steroid Metabolism, Inborn Errors
-	C16.320.565.925.249 Adrenal Hyperplasia, Congenital
-	C16.320.565.925.324 Antley-Bixler Syndrome Phenotype
-	C16.320.565.925.400 Ichthyosis, X-Linked
-	C16.320.565.925.500 Mineralocorticoid Excess Syndrome, Apparent
-	C16.320.565.925.875 Smith-Lemli-Opitz Syndrome
-	C16.320.577 Muscular Dystrophies
-	C16.320.577.074 Distal Myopathies
-	C16.320.577.149 Glycogen Storage Disease Type VII
-	C16.320.577.280 Muscular Dystrophies, Limb-Girdle

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C16.320.577.280.500 Sarcoglycanopathies
-	C16.320.577.300 Muscular Dystrophy, Duchenne
-	C16.320.577.350 Muscular Dystrophy, Emery-Dreifuss
-	C16.320.577.400 Muscular Dystrophy, Facioscapulohumeral
-	C16.320.577.450 Muscular Dystrophy, Oculopharyngeal
-	C16.320.577.500 Myotonic Dystrophy
-	C16.320.577.750 Walker-Warburg Syndrome
-	C16.320.590 Myasthenic Syndromes, Congenital
-	C16.320.600 Nail-Patella Syndrome
-	C16.320.700 Neoplastic Syndromes, Hereditary
-	C16.320.700.100 Adenomatous Polyposis Coli
-	C16.320.700.100.393 Gardner Syndrome
-	C16.320.700.175 Basal Cell Nevus Syndrome
-	C16.320.700.212 Birt-Hogg-Dube Syndrome
-	C16.320.700.250 Colorectal Neoplasms, Hereditary Nonpolyposis
-	C16.320.700.250.500 Lynch Syndrome II
-	C16.320.700.250.500.500 Muir-Torre Syndrome
-	C16.320.700.305 Dysplastic Nevus Syndrome
-	C16.320.700.330 Exostoses, Multiple Hereditary
-	C16.320.700.435 Hamartoma Syndrome, Multiple
-	C16.320.700.517 Hereditary Breast and Ovarian Cancer Syndrome
-	C16.320.700.600 Li-Fraumeni Syndrome
-	C16.320.700.630 Multiple Endocrine Neoplasia
-	C16.320.700.630.500 Multiple Endocrine Neoplasia Type 1
-	C16.320.700.630.505 Multiple Endocrine Neoplasia Type 2a
-	C16.320.700.630.510 Multiple Endocrine Neoplasia Type 2b
-	C16.320.700.636 Tuberous Sclerosis
-	C16.320.700.642 Wilms Tumor
-	C16.320.700.642.220 Denys-Drash Syndrome
-	C16.320.700.642.950 WAGR Syndrome
-	C16.320.700.645 Neurofibromatoses
-	C16.320.700.645.650 Neurofibromatosis 1
-	C16.320.700.645.655 Neurofibromatosis 2
-	C16.320.700.705 Peutz-Jeghers Syndrome
New Tree	<a href="#">C16.320.709 Oculocerebrorenal Syndrome</a>

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	<a href="#">C16.320.714</a>	<a href="#">Orofaciodigital Syndromes</a>
-	C16.320.718	Osteoarthropathy, Primary Hypertrophic
-	C16.320.728	Osteochondrodysplasias
-	C16.320.737	Osteogenesis Imperfecta
-	C16.320.775	Pain Insensitivity, Congenital
-	C16.320.784	Pelger-Huet Anomaly
Old Tree	<a href="#">C16.320.793</a>	<a href="#">Polycystic Kidney, Autosomal Recessive</a>
New Tree	<a href="#">C16.320.798</a>	<a href="#">Polycystic Kidney Diseases</a>
New Tree	<a href="#">C16.320.798.500</a>	<a href="#">Polycystic Kidney, Autosomal Dominant</a>
New Tree	<a href="#">C16.320.798.510</a>	<a href="#">Polycystic Kidney, Autosomal Recessive</a>
-	C16.320.812	Pycnodysostosis
-	C16.320.850	Skin Diseases, Genetic
-	C16.320.850.080	Albinism
-	C16.320.850.080.090	Albinism, Ocular
-	C16.320.850.080.100	Albinism, Oculocutaneous
-	C16.320.850.080.100.400	Hermanski-Pudlak Syndrome
-	C16.320.850.080.600	Piebaldism
-	C16.320.850.180	Cutis Laxa
-	C16.320.850.190	Darier Disease
-	C16.320.850.210	Dermatitis, Atopic
-	C16.320.850.235	Dyskeratosis Congenita
-	C16.320.850.250	Ectodermal Dysplasia
-	C16.320.850.250.198	Ectodermal Dysplasia 1, Anhidrotic
-	C16.320.850.250.298	Ectodermal Dysplasia 3, Anhidrotic
-	C16.320.850.250.348 Recessive	Ectodermal Dysplasia, Hypohidrotic, Autosomal
-	C16.320.850.250.398	Ellis-Van Creveld Syndrome
-	C16.320.850.250.424	Focal Dermal Hypoplasia
-	C16.320.850.250.712	Neurocutaneous Syndromes
-	C16.320.850.250.856	Pachyonychia Congenita
-	C16.320.850.250.856.500	Steatocystoma Multiplex
-	C16.320.850.260	Ehlers-Danlos Syndrome
-	C16.320.850.275	Epidermolysis Bullosa

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C16.320.850.275.160      Epidermolysis Bullosa Dystrophica
-	C16.320.850.275.170      Epidermolysis Bullosa, Junctional
-	C16.320.850.275.180      Epidermolysis Bullosa Simplex
-	C16.320.850.337          Erythrokeratoderma Variabilis
-	C16.320.850.368          Hyalinosis, Systemic
-	C16.320.850.400          Ichthyosiform Erythroderma, Congenital
-	C16.320.850.400.375      Hyperkeratosis, Epidermolytic
-	C16.320.850.400.410      Ichthyosis, Lamellar
-	C16.320.850.402          Ichthyosis Bullosa of Siemens
-	C16.320.850.405          Ichthyosis Vulgaris
-	C16.320.850.408          Ichthyosis, X-Linked
-	C16.320.850.420          Incontinentia Pigmenti
-	C16.320.850.475          Keratoderma, Palmoplantar
-	C16.320.850.475.440      Keratoderma, Palmoplantar, Diffuse
-	C16.320.850.475.440.500   Keratoderma, Palmoplantar, Epidermolytic
-	C16.320.850.475.600      Papillon-Lefevre Disease
-	C16.320.850.542          Leukokeratosis, Hereditary Mucosal
-	C16.320.850.595          Lipoid Proteinosis of Urbach and Wiethe
-	C16.320.850.647          Monilethrix
-	C16.320.850.673          Netherton Syndrome
-	C16.320.850.700          Pemphigus, Benign Familial
-	C16.320.850.730          Prokeratosis
-	C16.320.850.738          Porphyria, Erythropoietic
-	C16.320.850.742          Porphyrias, Hepatic
-	C16.320.850.742.074      Coproporphyrinuria, Hereditary
-	C16.320.850.742.150      Porphyria, Acute Intermittent
-	C16.320.850.742.250      Porphyria Cutanea Tarda
-	C16.320.850.742.437      Porphyria, Hepatoerythropoietic
-	C16.320.850.742.625      Porphyria, Variegata
-	C16.320.850.742.812      Protoporphyrinuria, Erythropoietic
-	C16.320.850.746          Prolidase Deficiency
-	C16.320.850.750          Pseudoxanthoma Elasticum
-	C16.320.850.765          Rothmund-Thomson Syndrome
-	C16.320.850.820          Sjogren-Larsson Syndrome
-	C16.320.850.895          Trichothiodystrophy Syndromes
-	C16.320.850.970          Xeroderma Pigmentosum

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C16.320.925                      Werner Syndrome
-	C16.320.962                      Yellow Nail Syndrome
-	C16.614                              Infant, Newborn, Diseases
-	C16.614.042                      Amniotic Band Syndrome
-	C16.614.053                      Anemia, Neonatal
-	C16.614.053.344                      Fetofetal Transfusion
-	C16.614.053.511                      Fetomaternal Transfusion
-	C16.614.092                      Asphyxia Neonatorum
-	C16.614.131                      Birth Injuries
-	C16.614.131.587                      Paralysis, Obstetric
-	C16.614.166                      Colic
-	C16.614.200                      Congenital Hyperinsulinism
-	C16.614.200.500                      Nesidioblastosis
-	C16.614.213                      Cystic Fibrosis
-	C16.614.258                      Epilepsy, Benign Neonatal
-	C16.614.304                      Erythroblastosis, Fetal
-	C16.614.304.502                      Kernicterus
-	C16.614.378                      Hernia, Umbilical
-	C16.614.438                      Hydrophthalmos
-	C16.614.451                      Hyperbilirubinemia, Neonatal
-	C16.614.451.500                      Jaundice, Neonatal
-	C16.614.451.500.250                      Jaundice, Chronic Idiopathic
-	C16.614.465                      Hyperostosis, Cortical, Congenital
-	C16.614.492                      Ichthyosis
-	C16.614.492.400                      Ichthyosiform Erythroderma, Congenital
-	C16.614.492.400.375                      Hyperkeratosis, Epidermolytic
-	C16.614.492.400.410                      Ichthyosis, Lamellar
-	C16.614.492.400.705                      Netherton Syndrome
-	C16.614.492.410                      Ichthyosis Bullosa of Siemens
-	C16.614.492.420                      Ichthyosis, X-Linked
-	C16.614.492.723                      Sjogren-Larsson Syndrome
-	C16.614.521                      Infant, Premature, Diseases
-	C16.614.521.125                      Bronchopulmonary Dysplasia
-	C16.614.521.450                      Leukomalacia, Periventricular
-	C16.614.521.563                      Respiratory Distress Syndrome, Newborn
-	C16.614.521.563.475                      Hyaline Membrane Disease

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C16.614.521.563.737 Transient Tachypnea of the Newborn
-	C16.614.521.731 Retinopathy of Prematurity
-	C16.614.580 Meconium Aspiration Syndrome
-	C16.614.595 Mobius Syndrome
-	C16.614.610 Neonatal Abstinence Syndrome
New Heading	<b>C16.614.627 Neonatal Sepsis</b>
-	C16.614.643 Nystagmus, Congenital
-	C16.614.677 Ophthalmia Neonatorum
-	C16.614.694 Persistent Fetal Circulation Syndrome
-	C16.614.760 Rothmund-Thomson Syndrome
-	C16.614.810 Sclerema Neonatorum
-	C16.614.815 Severe Combined Immunodeficiency
-	C16.614.815.500 X-Linked Combined Immunodeficiency Diseases
-	C16.614.868 Syphilis, Congenital
-	C16.614.890 Thanatophoric Dysplasia
-	C16.614.899 Thrombocytopenia, Neonatal Alloimmune
-	C16.614.909 Toxoplasmosis, Congenital
-	C16.614.940 Vitamin K Deficiency Bleeding
-	C16.614.947 Wolman Disease
-	C17 Skin and Connective Tissue Diseases
-	C17.300 Connective Tissue Diseases
-	C17.300.116 Anetoderma
-	C17.300.182 Cartilage Diseases
-	C17.300.182.100 Chondromalacia Patellae
-	C17.300.182.310 Laryngomalacia
-	C17.300.182.520 Osteochondritis
-	C17.300.182.525 Pectus Carinatum
-	C17.300.182.531 Polychondritis, Relapsing
-	C17.300.182.790 Tietze's Syndrome
-	C17.300.182.895 Tracheobronchomalacia
-	C17.300.182.895.249 Bronchomalacia
-	C17.300.182.895.500 Tracheomalacia
-	C17.300.185 Cellulitis
-	C17.300.185.500 Orbital Cellulitis
-	C17.300.200 Collagen Diseases



## MeSH Tree Changes for 2017

Type	Tree - heading	
-	C17.300.200.310	Ehlers-Danlos Syndrome
-	C17.300.200.367	Epidermolysis Bullosa Dystrophica
-	C17.300.200.425	Keloid
-	C17.300.200.425.125	Acne Keloid
-	C17.300.200.495	Necrobiotic Disorders
-	C17.300.200.495.380	Granuloma Annulare
-	C17.300.200.495.545	Necrobiosis Lipoidica
-	C17.300.200.495.772	Necrobiotic Xanthogranuloma
-	C17.300.200.517	Nephritis, Hereditary
-	C17.300.200.540	Osteogenesis Imperfecta
-	C17.300.230	Cutis Laxa
-	C17.300.250	Dermatomyositis
-	C17.300.270	Dupuytren Contracture
New Heading	<b>C17.300.349</b>	<b>Fibromatosis, Plantar</b>
-	C17.300.428	Homocystinuria
-	C17.300.451	Lipedema
-	C17.300.475	Lupus Erythematosus, Cutaneous
-	C17.300.475.479	Lupus Erythematosus, Discoid
-	C17.300.475.479.400	Panniculitis, Lupus Erythematosus
-	C17.300.480	Lupus Erythematosus, Systemic
-	C17.300.480.680	Lupus Nephritis
-	C17.300.480.750	Lupus Vasculitis, Central Nervous System
-	C17.300.500	Marfan Syndrome
-	C17.300.540	Mixed Connective Tissue Disease
-	C17.300.550	Mucinoses
-	C17.300.550.274	Ganglion Cysts
-	C17.300.550.550	Mucinosis, Follicular
-	C17.300.550.575	Mucopolysaccharidoses
-	C17.300.550.575.640	Mucopolysaccharidosis I
-	C17.300.550.575.645	Mucopolysaccharidosis II
-	C17.300.550.575.650	Mucopolysaccharidosis III
-	C17.300.550.575.655	Mucopolysaccharidosis IV
-	C17.300.550.575.670	Mucopolysaccharidosis VI
-	C17.300.550.575.675	Mucopolysaccharidosis VII
-	C17.300.550.590	Myxedema

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C17.300.550.750 Scleredema Adultorum
-	C17.300.550.875 Scleromyxedema
-	C17.300.680 Neoplasms, Connective Tissue
-	C17.300.680.540 Myofibroma
-	C17.300.690 Noonan Syndrome
-	C17.300.705 Osteopoikilosis
-	C17.300.710 Panniculitis
-	C17.300.710.329 Erythema Induratum
-	C17.300.710.400 Panniculitis, Lupus Erythematosus
-	C17.300.710.500 Panniculitis, Nodular Nonsuppurative
-	C17.300.710.600 Panniculitis, Peritoneal
-	C17.300.715 Penile Induration
-	C17.300.766 Pseudoxanthoma Elasticum
-	C17.300.775 Rheumatic Diseases
-	C17.300.775.049 Arthritis, Juvenile
-	C17.300.775.099 Arthritis, Rheumatoid
-	C17.300.775.099.219 Caplan Syndrome
-	C17.300.775.099.389 Felty Syndrome
-	C17.300.775.099.683 Rheumatoid Nodule
-	C17.300.775.099.728 Rheumatoid Vasculitis
-	C17.300.775.099.774 Sjogren's Syndrome
-	C17.300.775.099.870 Still's Disease, Adult-Onset
-	C17.300.775.440 Hyperostosis, Sternocostoclavicular
-	C17.300.775.720 Polymyalgia Rheumatica
-	C17.300.787 Scleroderma, Localized
-	C17.300.799 Scleroderma, Systemic
-	C17.300.799.602 Scleroderma, Diffuse
-	C17.300.799.801 Scleroderma, Limited
-	C17.300.799.801.500 CREST Syndrome
-	C17.300.899 Weill-Marchesani Syndrome
-	C17.800 Skin Diseases
-	C17.800.030 Acneiform Eruptions
-	C17.800.030.030 Acne Keloid
-	C17.800.030.150 Acne Vulgaris
-	C17.800.030.150.500 Acne Conglobata
-	C17.800.030.575 Chloracne

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	C17.800.060	Angiolymphoid Hyperplasia with Eosinophilia
-	C17.800.090	Breast Diseases
-	C17.800.090.249	Breast Cyst
-	C17.800.090.500	Breast Neoplasms
New Heading	<b>C17.800.090.500.130</b>	<b>Breast Carcinoma In Situ</b>
-	C17.800.090.500.260	Breast Neoplasms, Male
-	C17.800.090.500.390	Carcinoma, Ductal, Breast
New Tree	<b>C17.800.090.500.437</b>	<b>Carcinoma, Lobular</b>
-	C17.800.090.500.483	Hereditary Breast and Ovarian Cancer Syndrome
-	C17.800.090.500.576	Inflammatory Breast Neoplasms
-	C17.800.090.500.682	Unilateral Breast Neoplasms
-	C17.800.090.500.788	Triple Negative Breast Neoplasms
-	C17.800.090.750	Fibrocystic Breast Disease
-	C17.800.090.875	Gynecomastia
-	C17.800.090.937	Lactation Disorders
-	C17.800.090.937.439	Galactorrhea
-	C17.800.090.937.439.500	Chiari-Frommel Syndrome
-	C17.800.090.968	Mastitis
-	C17.800.090.968.400	Granulomatous Mastitis
-	C17.800.135	Cutaneous Fistula
-	C17.800.174	Dermatitis
-	C17.800.174.100	Acrodermatitis
-	C17.800.174.146	Dandruff
-	C17.800.174.193	Dermatitis, Atopic
-	C17.800.174.255	Dermatitis, Contact
-	C17.800.174.255.100	Dermatitis, Allergic Contact
-	C17.800.174.255.100.600	Dermatitis, Photoallergic
-	C17.800.174.255.100.700	Dermatitis, Toxicodendron
-	C17.800.174.255.400	Dermatitis, Irritant
-	C17.800.174.255.400.225	Dermatitis, Phototoxic
-	C17.800.174.255.400.250	Diaper Rash
-	C17.800.174.255.700	Dermatitis, Occupational
-	C17.800.174.318	Dermatitis, Exfoliative
-	C17.800.174.360	Dermatitis Herpetiformis

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	C17.800.174.550	Dermatitis, Perioral
-	C17.800.174.580	Dermatitis, Seborrheic
-	C17.800.174.600	Drug Eruptions
-	C17.800.174.600.174	Acute Generalized Exanthematous Pustulosis
-	C17.800.174.600.262	Drug Hypersensitivity Syndrome
-	C17.800.174.600.375	Erythema Nodosum
-	C17.800.174.600.587	Hand-Foot Syndrome
-	C17.800.174.600.693	Nicolau Syndrome
-	C17.800.174.600.800	Serum Sickness
-	C17.800.174.600.900	Stevens-Johnson Syndrome
-	C17.800.174.620	Eczema
-	C17.800.174.620.300	Eczema, Dyshidrotic
-	C17.800.174.640	Intertrigo
-	C17.800.174.660	Neurodermatitis
-	C17.800.174.826	Radiodermatitis
-	C17.800.185	Dermatomyositis
-	C17.800.229	Erythema
-	C17.800.229.200	Erythema Chronicum Migrans
-	C17.800.229.329	Erythema Induratum
-	C17.800.229.335	Erythema Infectiosum
-	C17.800.229.400	Erythema Multiforme
-	C17.800.229.400.683	Stevens-Johnson Syndrome
-	C17.800.229.413	Erythema Nodosum
-	C17.800.229.606	Erythrokeratoderma Variabilis
-	C17.800.229.800	Sweet Syndrome
-	C17.800.257	Exanthema
-	C17.800.257.335	Exanthema Subitum
-	C17.800.271	Facial Dermatoses
-	C17.800.271.250	Dermatitis, Perioral
-	C17.800.321	Foot Diseases
New Heading	<b>C17.800.321.063</b>	<b>Fibromatosis, Plantar</b>
-	C17.800.321.125	Foot Dermatoses
-	C17.800.321.125.500	Tinea Pedis
-	C17.800.321.250	Foot Ulcer
-	C17.800.321.500	Immersion Foot

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C17.800.329 Hair Diseases
-	C17.800.329.500 Folliculitis
-	C17.800.329.500.261 Acne Keloid
-	C17.800.329.750 Hirsutism
-	C17.800.329.875 Hypertrichosis
-	C17.800.329.937 Hypotrichosis
-	C17.800.329.937.122 Alopecia
-	C17.800.329.937.122.147 Alopecia Areata
-	C17.800.329.937.122.348 Loose Anagen Hair Syndrome
-	C17.800.329.937.122.550 Mucinosis, Follicular
-	C17.800.329.968 Menkes Kinky Hair Syndrome
-	C17.800.329.984 Monilethrix
-	C17.800.329.992 Piedra
-	C17.800.338 Hand Dermatoses
-	C17.800.417 Keratoacanthoma
-	C17.800.428 Keratosis
-	C17.800.428.200 Callosities
-	C17.800.428.260 Cholesteatoma
-	C17.800.428.260.300 Cholesteatoma, Middle Ear
-	C17.800.428.275 Darier Disease
-	C17.800.428.304 Erythrokeratoderma Variabilis
-	C17.800.428.333 Ichthyosis
-	C17.800.428.333.250 Ichthyosiform Erythroderma, Congenital
-	C17.800.428.333.250.375 Hyperkeratosis, Epidermolytic
-	C17.800.428.333.250.410 Ichthyosis, Lamellar
-	C17.800.428.333.250.705 Netherton Syndrome
-	C17.800.428.333.330 Ichthyosis Bullosa of Siemens
-	C17.800.428.333.410 Ichthyosis Vulgaris
-	C17.800.428.333.420 Ichthyosis, X-Linked
-	C17.800.428.333.723 Sjogren-Larsson Syndrome
-	C17.800.428.435 Keratoderma, Palmoplantar
-	C17.800.428.435.440 Keratoderma, Palmoplantar, Diffuse
-	C17.800.428.435.440.500 Keratoderma, Palmoplantar, Epidermolytic
-	C17.800.428.435.600 Papillon-Lefevre Disease
-	C17.800.428.570 Keratosis, Actinic
-	C17.800.428.580 Keratosis, Seborrheic

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C17.800.428.736 Parakeratosis
-	C17.800.428.750 Porokeratosis
-	C17.800.446 Leg Dermatoses
-	C17.800.463 Lipomatosis
-	C17.800.463.249 Adiposis Dolorosa
New Heading	<b>C17.800.463.375 Familial Multiple Lipomatosis</b>
-	C17.800.463.500 Lipomatosis, Multiple Symmetrical
-	C17.800.480 Lupus Erythematosus, Cutaneous
-	C17.800.480.479 Lupus Erythematosus, Discoid
-	C17.800.480.479.400 Panniculitis, Lupus Erythematosus
-	C17.800.508 Mastocytosis
-	C17.800.508.236 Mastocytoma
-	C17.800.508.236.500 Mastocytoma, Skin
-	C17.800.508.473 Mastocytosis, Cutaneous
-	C17.800.508.473.500 Mastocytoma, Skin
-	C17.800.508.473.850 Urticaria Pigmentosa
-	C17.800.518 Morgellons Disease
-	C17.800.529 Nail Diseases
-	C17.800.529.400 Nail-Patella Syndrome
-	C17.800.529.406 Nails, Ingrown
-	C17.800.529.478 Onycholysis
-	C17.800.529.550 Onychomycosis
-	C17.800.529.594 Pachyonychia Congenita
-	C17.800.529.594.500 Steatocystoma Multiplex
-	C17.800.529.639 Paronychia
-	C17.800.529.819 Yellow Nail Syndrome
-	C17.800.550 Necrobiotic Disorders
-	C17.800.550.380 Granuloma Annulare
-	C17.800.550.545 Necrobiosis Lipoidica
-	C17.800.550.772 Necrobiotic Xanthogranuloma
-	C17.800.551 Necrolytic Migratory Erythema
-	C17.800.553 Nephrogenic Fibrosing Dermopathy
-	C17.800.566 Panniculitis
-	C17.800.566.329 Erythema Induratum
-	C17.800.566.400 Panniculitis, Lupus Erythematosus

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C17.800.566.500 Panniculitis, Nodular Nonsuppurative
-	C17.800.600 Photosensitivity Disorders
-	C17.800.600.325 Dermatitis, Photoallergic
-	C17.800.600.335 Dermatitis, Phototoxic
-	C17.800.600.425 Hydroa Vacciniforme
-	C17.800.600.725 Sunburn
-	C17.800.600.925 Xeroderma Pigmentosum
-	C17.800.621 Pigmentation Disorders
-	C17.800.621.166 Argyria
-	C17.800.621.250 Cafe-au-Lait Spots
-	C17.800.621.430 Hyperpigmentation
-	C17.800.621.430.530 Melanosis
-	C17.800.621.430.530.100 Acanthosis Nigricans
-	C17.800.621.430.530.550 Lentigo
-	C17.800.621.430.530.550.525 LEOPARD Syndrome
-	C17.800.621.430.530.550.625 Peutz-Jeghers Syndrome
-	C17.800.621.440 Hypopigmentation
-	C17.800.621.440.102 Albinism
-	C17.800.621.440.102.090 Albinism, Ocular
-	C17.800.621.440.102.100 Albinism, Oculocutaneous
-	C17.800.621.440.102.100.400 Hermanski-Pudlak Syndrome
-	C17.800.621.440.102.600 Piebaldism
-	C17.800.621.440.895 Vitiligo
-	C17.800.621.497 Incontinentia Pigmenti
-	C17.800.621.893 Urticaria Pigmentosa
-	C17.800.621.936 Xeroderma Pigmentosum
-	C17.800.621.968 Yellow Nail Syndrome
-	C17.800.674 Prurigo
-	C17.800.685 Pruritus
-	C17.800.685.544 Pruritus Ani
-	C17.800.685.710 Pruritus Vulvae
-	C17.800.695 Pyoderma
-	C17.800.695.210 Ecthyma
-	C17.800.695.675 Pyoderma Gangrenosum
-	C17.800.716 Rosacea
-	C17.800.716.500 Rhinophyma

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C17.800.738 Scalp Dermatoses
-	C17.800.738.353 Dandruff
-	C17.800.738.708 Tinea Capitis
-	C17.800.738.708.708 Tinea Favosa
-	C17.800.751 Scleredema Adultorum
-	C17.800.767 Scleroderma, Localized
-	C17.800.784 Scleroderma, Systemic
-	C17.800.784.602 Scleroderma, Diffuse
-	C17.800.784.801 Scleroderma, Limited
-	C17.800.784.801.500 CREST Syndrome
-	C17.800.794 Sebaceous Gland Diseases
-	C17.800.794.111 Acne Vulgaris
-	C17.800.794.111.500 Acne Conglobata
-	C17.800.794.230 Dermatitis, Seborrheic
-	C17.800.794.550 Mucinosis, Follicular
-	C17.800.794.650 Rhinophyma
-	C17.800.794.712 Sebaceous Gland Neoplasms
-	C17.800.794.712.500 Muir-Torre Syndrome
-	C17.800.804 Skin Abnormalities
-	C17.800.804.066 Acrodermatitis
-	C17.800.804.108 Anetoderma
-	C17.800.804.150 Dyskeratosis Congenita
-	C17.800.804.350 Ectodermal Dysplasia
-	C17.800.804.350.198 Ectodermal Dysplasia 1, Anhidrotic
-	C17.800.804.350.298 Ectodermal Dysplasia 3, Anhidrotic
-	C17.800.804.350.348 Ectodermal Dysplasia, Hypohidrotic, Autosomal Recessive
-	C17.800.804.350.398 Ellis-Van Creveld Syndrome
-	C17.800.804.350.424 Focal Dermal Hypoplasia
-	C17.800.804.350.712 Neurocutaneous Syndromes
-	C17.800.804.350.856 Pachyonychia Congenita
-	C17.800.804.350.856.500 Steatocystoma Multiplex
-	C17.800.804.428 Ehlers-Danlos Syndrome
-	C17.800.804.493 Epidermolysis Bullosa
-	C17.800.804.493.080 Epidermolysis Bullosa Acquisita
-	C17.800.804.493.160 Epidermolysis Bullosa Dystrophica



## MeSH Tree Changes for 2017

Type	Tree - heading
-	C17.800.804.493.170 Epidermolysis Bullosa, Junctional
-	C17.800.804.493.180 Epidermolysis Bullosa Simplex
-	C17.800.804.512 Ichthyosis
-	C17.800.804.512.400 Ichthyosiform Erythroderma, Congenital
-	C17.800.804.512.400.375 Hyperkeratosis, Epidermolytic
-	C17.800.804.512.400.410 Ichthyosis, Lamellar
-	C17.800.804.512.400.705 Netherton Syndrome
-	C17.800.804.512.405 Ichthyosis Bullosa of Siemens
-	C17.800.804.512.410 Ichthyosis Vulgaris
-	C17.800.804.512.420 Ichthyosis, X-Linked
-	C17.800.804.512.723 Sjogren-Larsson Syndrome
-	C17.800.804.580 Incontinentia Pigmenti
-	C17.800.804.675 Port-Wine Stain
-	C17.800.804.766 Pseudoxanthoma Elasticum
-	C17.800.804.775 Rothmund-Thomson Syndrome
-	C17.800.804.812 Sclerema Neonatorum
-	C17.800.804.874 Trichothiodystrophy Syndromes
-	C17.800.804.936 Xeroderma Pigmentosum
-	C17.800.815 Skin Diseases, Eczematous
-	C17.800.815.193 Dermatitis, Atopic
-	C17.800.815.255 Dermatitis, Contact
-	C17.800.815.255.100 Dermatitis, Allergic Contact
-	C17.800.815.255.100.600 Dermatitis, Photoallergic
-	C17.800.815.255.100.700 Dermatitis, Toxicodendron
-	C17.800.815.255.400 Dermatitis, Irritant
-	C17.800.815.255.400.225 Dermatitis, Phototoxic
-	C17.800.815.255.400.250 Diaper Rash
-	C17.800.815.255.700 Dermatitis, Occupational
-	C17.800.815.318 Dermatitis, Exfoliative
-	C17.800.815.580 Dermatitis, Seborrheic
-	C17.800.815.620 Eczema
-	C17.800.815.620.300 Eczema, Dyshidrotic
-	C17.800.815.650 Intertrigo
-	C17.800.815.660 Neurodermatitis
-	C17.800.827 Skin Diseases, Genetic
-	C17.800.827.080 Albinism

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C17.800.827.080.090 Albinism, Ocular
-	C17.800.827.080.100 Albinism, Oculocutaneous
-	C17.800.827.080.100.400 Hermanski-Pudlak Syndrome
-	C17.800.827.080.600 Piebaldism
-	C17.800.827.180 Cutis Laxa
-	C17.800.827.190 Darier Disease
-	C17.800.827.210 Dermatitis, Atopic
-	C17.800.827.235 Dyskeratosis Congenita
-	C17.800.827.250 Ectodermal Dysplasia
-	C17.800.827.250.198 Ectodermal Dysplasia 1, Anhidrotic
-	C17.800.827.250.298 Ectodermal Dysplasia 3, Anhidrotic
-	C17.800.827.250.348 Ectodermal Dysplasia, Hypohidrotic, Autosomal Recessive
-	C17.800.827.250.398 Ellis-Van Creveld Syndrome
-	C17.800.827.250.424 Focal Dermal Hypoplasia
-	C17.800.827.250.712 Neurocutaneous Syndromes
-	C17.800.827.250.856 Pachyonychia Congenita
-	C17.800.827.250.856.500 Steatocystoma Multiplex
-	C17.800.827.260 Ehlers-Danlos Syndrome
-	C17.800.827.275 Epidermolysis Bullosa
-	C17.800.827.275.080 Epidermolysis Bullosa Acquisita
-	C17.800.827.275.160 Epidermolysis Bullosa Dystrophica
-	C17.800.827.275.170 Epidermolysis Bullosa, Junctional
-	C17.800.827.275.180 Epidermolysis Bullosa Simplex
-	C17.800.827.337 Erythrokeratoderma Variabilis
-	C17.800.827.368 Hereditary Autoinflammatory Diseases
-	C17.800.827.368.250 Behcet Syndrome
-	C17.800.827.368.500 Cryopyrin-Associated Periodic Syndromes
-	C17.800.827.384 Hyalinosis, Systemic
-	C17.800.827.400 Ichthyosiform Erythroderma, Congenital
-	C17.800.827.400.375 Hyperkeratosis, Epidermolytic
-	C17.800.827.400.410 Ichthyosis, Lamellar
-	C17.800.827.403 Ichthyosis Bullosa of Siemens
-	C17.800.827.405 Ichthyosis Vulgaris
-	C17.800.827.408 Ichthyosis, X-Linked
-	C17.800.827.420 Incontinentia Pigmenti

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C17.800.827.475 Keratoderma, Palmoplantar
-	C17.800.827.475.440 Keratoderma, Palmoplantar, Diffuse
-	C17.800.827.475.440.500 Keratoderma, Palmoplantar, Epidermolytic
-	C17.800.827.475.600 Papillon-Lefevre Disease
-	C17.800.827.595 Leukokeratosis, Hereditary Mucosal
-	C17.800.827.602 Monilethrix
-	C17.800.827.610 Muir-Torre Syndrome
-	C17.800.827.655 Netherton Syndrome
-	C17.800.827.700 Pemphigus, Benign Familial
-	C17.800.827.730 Porokeratosis
-	C17.800.827.738 Porphyria, Erythropoietic
-	C17.800.827.742 Porphyrias, Hepatic
-	C17.800.827.742.074 Coproporphyrinuria, Hereditary
-	C17.800.827.742.150 Porphyria, Acute Intermittent
-	C17.800.827.742.250 Porphyria Cutanea Tarda
-	C17.800.827.742.437 Porphyria, Hepatoerythropoietic
-	C17.800.827.742.625 Porphyria, Variegate
-	C17.800.827.742.812 Protoporphyrinuria, Erythropoietic
-	C17.800.827.750 Pseudoxanthoma Elasticum
-	C17.800.827.775 Rothmund-Thomson Syndrome
-	C17.800.827.820 Sjogren-Larsson Syndrome
-	C17.800.827.895 Trichothiodystrophy Syndromes
-	C17.800.827.970 Xeroderma Pigmentosum
-	C17.800.838 Skin Diseases, Infectious
-	C17.800.838.208 Dermatomycoses
-	C17.800.838.208.055 Blastomycosis
-	C17.800.838.208.165 Candidiasis, Chronic Mucocutaneous
-	C17.800.838.208.170 Candidiasis, Cutaneous
-	C17.800.838.208.241 Chromoblastomycosis
-	C17.800.838.208.416 Hyalohyphomycosis
-	C17.800.838.208.416.125 Alternariosis
-	C17.800.838.208.416.249 Aspergillosis
-	C17.800.838.208.416.249.074 Aspergillosis, Allergic Bronchopulmonary
-	C17.800.838.208.416.249.537 Neuroaspergillosis
-	C17.800.838.208.416.375 Cerebral Phaeohyphomycosis
-	C17.800.838.208.416.500 Fusariosis

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C17.800.838.208.475 Lobomycosis
-	C17.800.838.208.557 Mycetoma
-	C17.800.838.208.675 Sporotrichosis
-	C17.800.838.208.883 Tinea
-	C17.800.838.208.883.458 Onychomycosis
-	C17.800.838.208.883.558 Tinea Capitis
-	C17.800.838.208.883.558.708 Tinea Favosa
-	C17.800.838.208.883.658 Tinea Pedis
-	C17.800.838.208.941 Tinea Versicolor
-	C17.800.838.486 Paronychia
-	C17.800.838.765 Skin Diseases, Bacterial
-	C17.800.838.765.110 Actinomycosis, Cervicofacial
-	C17.800.838.765.150 Angiomatosis, Bacillary
-	C17.800.838.765.180 Digital Dermatitis
-	C17.800.838.765.210 Ecthyma
-	C17.800.838.765.260 Erysipelas
-	C17.800.838.765.310 Erythema Chronicum Migrans
-	C17.800.838.765.320 Erythrasma
-	C17.800.838.765.360 Granuloma Inguinale
-	C17.800.838.765.420 Hidradenitis Suppurativa
-	C17.800.838.765.557 Mycetoma
-	C17.800.838.765.630 Pinta
-	C17.800.838.765.705 Rhinoscleroma
-	C17.800.838.765.770 Staphylococcal Skin Infections
-	C17.800.838.765.770.270 Furunculosis
-	C17.800.838.765.770.270.200 Carbuncle
-	C17.800.838.765.770.360 Impetigo
-	C17.800.838.765.770.770 Staphylococcal Scalded Skin Syndrome
-	C17.800.838.765.790 Syphilis, Cutaneous
-	C17.800.838.765.820 Tuberculosis, Cutaneous
-	C17.800.838.765.820.320 Erythema Induratum
-	C17.800.838.765.820.470 Lupus Vulgaris
-	C17.800.838.765.910 Yaws
-	C17.800.838.775 Skin Diseases, Parasitic
-	C17.800.838.775.424 Larva Migrans
-	C17.800.838.775.560 Leishmaniasis

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C17.800.838.775.560.400                      Leishmaniasis, Cutaneous
-	C17.800.838.775.560.400.350                      Leishmaniasis, Diffuse Cutaneous
-	C17.800.838.775.560.400.385                      Leishmaniasis, Mucocutaneous
-	C17.800.838.775.580                      Lice Infestations
-	C17.800.838.775.690                      Onchocerciasis
-	C17.800.838.775.800                      Scabies
-	C17.800.838.790                      Skin Diseases, Viral
-	C17.800.838.790.260                      Erythema Infectiosum
-	C17.800.838.790.290                      Exanthema Subitum
-	C17.800.838.790.320                      Herpes Simplex
-	C17.800.838.790.320.320                      Herpes Labialis
-	C17.800.838.790.320.410                      Kaposi Varicelliform Eruption
-	C17.800.838.790.550                      Molluscum Contagiosum
-	C17.800.838.790.810                      Warts
-	C17.800.838.790.810.110                      Condylomata Acuminata
-	C17.800.838.790.810.110.500                      Buschke-Lowenstein Tumor
-	C17.800.838.790.810.260                      Epidermodysplasia Verruciformis
-	C17.800.849                      Skin Diseases, Metabolic
-	C17.800.849.391                      Lipodystrophy
-	C17.800.849.391.400                      HIV-Associated Lipodystrophy Syndrome
-	C17.800.849.391.550                      Lipodystrophy, Congenital Generalized
-	C17.800.849.391.700                      Lipodystrophy, Familial Partial
-	C17.800.849.495                      Necrobiosis Lipoidica
-	C17.800.859                      Skin Diseases, Papulosquamous
-	C17.800.859.350                      Dermatitis, Seborrheic
-	C17.800.859.475                      Lichenoid Eruptions
-	C17.800.859.475.545                      Lichen Nitidus
-	C17.800.859.475.560                      Lichen Planus
-	C17.800.859.475.560.397                      Lichen Planus, Oral
-	C17.800.859.475.605                      Lichen Sclerosus et Atrophicus
-	C17.800.859.475.650                      Pityriasis Lichenoides
-	C17.800.859.575                      Parapsoriasis
-	C17.800.859.575.650                      Pityriasis Lichenoides
-	C17.800.859.600                      Pityriasis
-	C17.800.859.600.650                      Pityriasis Lichenoides
-	C17.800.859.600.675                      Pityriasis Rosea

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C17.800.859.600.685 Pityriasis Rubra Pilaris
-	C17.800.859.675 Psoriasis
-	C17.800.859.675.175 Arthritis, Psoriatic
-	C17.800.862 Skin Diseases, Vascular
-	C17.800.862.060 Angiomatosis, Bacillary
-	C17.800.862.150 Behcet Syndrome
-	C17.800.862.252 Giant Cell Arteritis
-	C17.800.862.355 Livedo Reticularis
-	C17.800.862.457 Malignant Atrophic Papulosis
-	C17.800.862.560 Mucocutaneous Lymph Node Syndrome
-	C17.800.862.625 Polyarteritis Nodosa
-	C17.800.862.675 Pyoderma Gangrenosum
-	C17.800.862.775 Sneddon Syndrome
-	C17.800.862.875 Takayasu Arteritis
-	C17.800.862.945 Urticaria
-	C17.800.862.945.066 Angioedema
-	C17.800.862.945.066.500 Angioedemas, Hereditary
-	C17.800.862.945.066.500.500 Hereditary Angioedema Type III
-	C17.800.862.945.066.500.750 Hereditary Angioedema Types I and II
-	C17.800.865 Skin Diseases, Vesiculobullous
-	C17.800.865.070 Acantholysis
-	C17.800.865.187 Blister
-	C17.800.865.360 Dermatitis Herpetiformis
-	C17.800.865.385 Eczema, Dyshidrotic
-	C17.800.865.410 Epidermolysis Bullosa
-	C17.800.865.410.080 Epidermolysis Bullosa Acquisita
-	C17.800.865.410.160 Epidermolysis Bullosa Dystrophica
-	C17.800.865.410.170 Epidermolysis Bullosa, Junctional
-	C17.800.865.410.180 Epidermolysis Bullosa Simplex
-	C17.800.865.475 Erythema Multiforme
-	C17.800.865.475.683 Stevens-Johnson Syndrome
-	C17.800.865.575 Hydroa Vacciniforme
-	C17.800.865.622 Linear IgA Bullous Dermatitis
-	C17.800.865.670 Pemphigoid, Benign Mucous Membrane
-	C17.800.865.690 Pemphigoid, Bullous
-	C17.800.865.700 Pemphigoid Gestationis

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C17.800.865.716 Pemphigus
-	C17.800.865.858 Pemphigus, Benign Familial
-	C17.800.882 Skin Neoplasms
-	C17.800.882.712 Sebaceous Gland Neoplasms
-	C17.800.882.712.500 Muir-Torre Syndrome
-	C17.800.882.743 Sweat Gland Neoplasms
-	C17.800.893 Skin Ulcer
-	C17.800.893.295 Buruli Ulcer
-	C17.800.893.592 Leg Ulcer
-	C17.800.893.592.450 Foot Ulcer
-	C17.800.893.592.450.200 Diabetic Foot
-	C17.800.893.592.730 Varicose Ulcer
-	C17.800.893.665 Pressure Ulcer
-	C17.800.893.675 Pyoderma Gangrenosum
-	C17.800.946 Sweat Gland Diseases
-	C17.800.946.315 Hidradenitis
-	C17.800.946.315.320 Hidradenitis Suppurativa
-	C17.800.946.350 Hyperhidrosis
-	C17.800.946.350.843 Sweating, Gustatory
-	C17.800.946.370 Hypohidrosis
-	C17.800.946.492 Miliaria
-	C17.800.946.492.285 Fox-Fordyce Disease
-	C17.800.946.743 Sweat Gland Neoplasms
-	C17.800.973 Xanthogranuloma, Juvenile
-	C18 Nutritional and Metabolic Diseases
-	C18.452 Metabolic Diseases
-	C18.452.076 Acid-Base Imbalance
-	C18.452.076.087 Achlorhydria
-	C18.452.076.176 Acidosis
-	C18.452.076.176.180 Acidosis, Lactic
-	C18.452.076.176.210 Acidosis, Renal Tubular
-	C18.452.076.176.310 Acidosis, Respiratory
-	C18.452.076.176.652 Ketosis
-	C18.452.076.176.652.500 Diabetic Ketoacidosis
-	C18.452.076.354 Alkalosis
-	C18.452.076.354.271 Alkalosis, Respiratory

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	C18.452.104 Bone Diseases, Metabolic
New Tree	C18.452.104.247 Bone Demineralization, Pathologic
New Tree	C18.452.104.247.400 Decalcification, Pathologic
New Tree	C18.452.104.579 Osteoporosis
New Tree	C18.452.104.579.304 Female Athlete Triad Syndrome
New Tree	C18.452.104.579.610 Osteoporosis, Postmenopausal
New Tree	C18.452.104.709 Pseudohypoparathyroidism
New Tree	C18.452.104.709.628 Pseudopseudohypoparathyroidism
New Tree	C18.452.104.816 Rickets
New Tree	C18.452.104.816.640 Osteomalacia
New Tree	C18.452.104.816.750 Chronic Kidney Disease-Mineral and Bone Disorder
New Tree	C18.452.104.816.750 Renal Osteodystrophy
New Tree	C18.452.104.816.875 Rickets, Hypophosphatemic
New Tree	C18.452.104.816.875.500 Familial Hypophosphatemic Rickets
-	C18.452.132 Brain Diseases, Metabolic
-	C18.452.132.100 Brain Diseases, Metabolic, Inborn
-	C18.452.132.100.084 Adrenoleukodystrophy
-	C18.452.132.100.168 Cerebral Amyloid Angiopathy, Familial
-	C18.452.132.100.320 Galactosemias
-	C18.452.132.100.355 Hartnup Disease
-	C18.452.132.100.360 Hepatolenticular Degeneration
-	C18.452.132.100.362 Hereditary Central Nervous System Demyelinating Diseases
-	C18.452.132.100.362.250 Adrenoleukodystrophy
-	C18.452.132.100.362.312 Alexander Disease
-	C18.452.132.100.362.375 Canavan Disease
-	C18.452.132.100.362.500 Leukodystrophy, Globoid Cell



## MeSH Tree Changes for 2017

Type	Tree - heading
-	C18.452.132.100.362.550                      Leukodystrophy, Metachromatic
-	C18.452.132.100.362.775                      Pelizaeus-Merzbacher Disease
-	C18.452.132.100.365                          Homocystinuria
-	C18.452.132.100.375                          Hyperglycinemia, Nonketotic
-	C18.452.132.100.380                          Hyperlysinemias
-	C18.452.132.100.412                          Leigh Disease
-	C18.452.132.100.425                          Lesch-Nyhan Syndrome
-	C18.452.132.100.435                          Lysosomal Storage Diseases, Nervous System
-	C18.452.132.100.435.295                      Fucosidosis
-	C18.452.132.100.435.340                      Glycogen Storage Disease Type II
-	C18.452.132.100.435.590                      Mucopolidoses
-	C18.452.132.100.435.810                      Sialic Acid Storage Disease
-	C18.452.132.100.435.825                      Sphingolipidoses
-	C18.452.132.100.435.825.200                  Fabry Disease
-	C18.452.132.100.435.825.250                  Farber Lipogranulomatosis
-	C18.452.132.100.435.825.300                  Gangliosidoses
-	C18.452.132.100.435.825.300.300              Gangliosidoses, GM2
-	C18.452.132.100.435.825.300.300.249              Sandhoff Disease
-	C18.452.132.100.435.825.300.300.500              Tay-Sachs Disease
-	C18.452.132.100.435.825.300.300.750              Tay-Sachs Disease, AB Variant
-	C18.452.132.100.435.825.300.400                  Gangliosidosis, GM1
-	C18.452.132.100.435.825.400                  Gaucher Disease
-	C18.452.132.100.435.825.590                  Leukodystrophy, Globoid Cell
-	C18.452.132.100.435.825.700                  Niemann-Pick Diseases
-	C18.452.132.100.435.825.700.500                  Niemann-Pick Disease, Type A
-	C18.452.132.100.435.825.700.750                  Niemann-Pick Disease, Type B
-	C18.452.132.100.435.825.700.875                  Niemann-Pick Disease, Type C
-	C18.452.132.100.435.825.775                  Sea-Blue Histiocyte Syndrome
-	C18.452.132.100.435.825.850                  Sulfatidosis
-	C18.452.132.100.435.825.850.500                  Leukodystrophy, Metachromatic
-	C18.452.132.100.435.825.850.750                  Multiple Sulfatase Deficiency Disease
-	C18.452.132.100.520                          Maple Syrup Urine Disease
-	C18.452.132.100.535                          MELAS Syndrome
-	C18.452.132.100.540                          Menkes Kinky Hair Syndrome
-	C18.452.132.100.545                          MERRF Syndrome
-	C18.452.132.100.593                          Mevalonate Kinase Deficiency

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C18.452.132.100.640 Oculocerebrorenal Syndrome
-	C18.452.132.100.687 Phenylketonurias
-	C18.452.132.100.687.500 Phenylketonuria, Maternal
-	C18.452.132.100.725 Pyruvate Carboxylase Deficiency Disease
-	C18.452.132.100.750 Pyruvate Dehydrogenase Complex Deficiency Disease
-	C18.452.132.100.813 Refsum Disease
-	C18.452.132.100.844 Refsum Disease, Infantile
-	C18.452.132.100.875 Tyrosinemias
-	C18.452.132.100.937 Urea Cycle Disorders, Inborn
-	C18.452.132.100.937.124 Argininosuccinic Aciduria
-	C18.452.132.100.937.249 Carbamoyl-Phosphate Synthase I Deficiency Disease
-	C18.452.132.100.937.374 Citrullinemia
-	C18.452.132.100.937.437 Hyperargininemia
-	C18.452.132.100.937.500 Ornithine Carbamoyltransferase Deficiency Disease
-	C18.452.132.100.968 Zellweger Syndrome
-	C18.452.132.360 Hepatic Encephalopathy
-	C18.452.132.480 Kernicterus
-	C18.452.132.540 Mitochondrial Encephalomyopathies
-	C18.452.132.560 Myelinolysis, Central Pontine
-	C18.452.132.780 Reye Syndrome
-	C18.452.132.960 Wernicke Encephalopathy
-	C18.452.174 Calcium Metabolism Disorders
-	C18.452.174.130 Calcinosis
-	C18.452.174.130.186 Calciphylaxis
-	C18.452.174.130.204 CREST Syndrome
-	C18.452.174.130.560 Nephrocalcinosis
-	C18.452.174.130.780 Vascular Calcification
-	C18.452.174.130.780.500 Monckeberg Medial Calcific Sclerosis
-	C18.452.174.289 Decalcification, Pathologic
-	C18.452.174.451 Hypercalcemia
-	C18.452.174.509 Hypocalcemia
-	C18.452.174.509.700 Tetany
-	C18.452.174.766 Pseudohypoparathyroidism
-	C18.452.174.766.815 Pseudopseudohypoparathyroidism
-	C18.452.174.845 Rickets

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C18.452.174.845.640 Osteomalacia
-	C18.452.174.845.750 Chronic Kidney Disease-Mineral and Bone Disorder
-	C18.452.174.845.750 Renal Osteodystrophy
-	C18.452.174.845.875 Rickets, Hypophosphatemic
-	C18.452.174.845.875.500 Familial Hypophosphatemic Rickets
-	C18.452.284 DNA Repair-Deficiency Disorders
-	C18.452.284.060 Ataxia Telangiectasia
-	C18.452.284.100 Bloom Syndrome
-	C18.452.284.250 Cockayne Syndrome
-	C18.452.284.255 Colorectal Neoplasms, Hereditary Nonpolyposis
-	C18.452.284.280 Fanconi Anemia
-	C18.452.284.520 Li-Fraumeni Syndrome
-	C18.452.284.600 Nijmegen Breakage Syndrome
-	C18.452.284.760 Rothmund-Thomson Syndrome
-	C18.452.284.800 Severe Combined Immunodeficiency
-	C18.452.284.960 Werner Syndrome
-	C18.452.284.975 Xeroderma Pigmentosum
-	C18.452.394 Glucose Metabolism Disorders
-	C18.452.394.750 Diabetes Mellitus
-	C18.452.394.750.074 Diabetes Mellitus, Experimental
-	C18.452.394.750.124 Diabetes Mellitus, Type 1
-	C18.452.394.750.124.960 Wolfram Syndrome
-	C18.452.394.750.149 Diabetes Mellitus, Type 2
-	C18.452.394.750.149.500 Diabetes Mellitus, Lipoatrophic
-	C18.452.394.750.448 Diabetes, Gestational
-	C18.452.394.750.535 Diabetic Ketoacidosis
-	C18.452.394.750.654 Donohue Syndrome
New Heading	<b>C18.452.394.750.714 Latent Autoimmune Diabetes in Adults</b>
-	C18.452.394.750.774 Prediabetic State
-	C18.452.394.937 Glycosuria
-	C18.452.394.937.450 Glycosuria, Renal
-	C18.452.394.952 Hyperglycemia
-	C18.452.394.952.500 Glucose Intolerance
-	C18.452.394.968 Hyperinsulinism
-	C18.452.394.968.250 Congenital Hyperinsulinism

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C18.452.394.968.250.500                      Nesidioblastosis
-	C18.452.394.968.500                              Insulin Resistance
-	C18.452.394.968.500.570                      Metabolic Syndrome X
-	C18.452.394.984                                  Hypoglycemia
-	C18.452.394.984.200                              Congenital Hyperinsulinism
-	C18.452.394.984.200.500                      Nesidioblastosis
-	C18.452.394.984.492                              Insulin Coma
-	C18.452.479    Hyperlactatemia
-	C18.452.565    Iron Metabolism Disorders
-	C18.452.565.100                                    Anemia, Iron-Deficiency
-	C18.452.565.500                                    Iron Overload
-	C18.452.565.500.480                              Hemochromatosis
-	C18.452.565.500.500                              Hemosiderosis
-	C18.452.584    Lipid Metabolism Disorders
-	C18.452.584.500                                    Dyslipidemias
-	C18.452.584.500.500                              Hyperlipidemias
-	C18.452.584.500.500.396                        Hypercholesterolemia
-	C18.452.584.500.500.438                        Hyperlipidemia, Familial Combined
-	C18.452.584.500.500.644                        Hyperlipoproteinemias
-	C18.452.584.500.500.644.237                    Hyperlipoproteinemia Type I
-	C18.452.584.500.500.644.475                    Hyperlipoproteinemia Type II
-	C18.452.584.500.500.644.485                    Hyperlipoproteinemia Type III
-	C18.452.584.500.500.644.490                    Hyperlipoproteinemia Type IV
-	C18.452.584.500.500.644.495                    Hyperlipoproteinemia Type V
-	C18.452.584.500.500.851                        Hypertriglyceridemia
-	C18.452.584.500.500.851.500                    Hyperlipoproteinemia Type IV
-	C18.452.584.500.500.851.750                    Hyperlipoproteinemia Type V
-	C18.452.584.500.500.851.875                    Hypertriglyceridemic Waist
-	C18.452.584.500.875                              Hypolipoproteinemias
-	C18.452.584.500.875.330                        Hypoalphalipoproteinemias
-	C18.452.584.500.875.330.500                    Lecithin Acyltransferase Deficiency
-	C18.452.584.500.875.330.500                    Lecithin Cholesterol Acyltransferase Deficiency
-	C18.452.584.500.875.330.750                    Tangier Disease
-	C18.452.584.500.875.440                        Hypobetalipoproteinemias
-	C18.452.584.500.875.440.500                    Abetalipoproteinemia
-	C18.452.584.500.875.440.750                    Hypobetalipoproteinemia, Familial,

## MeSH Tree Changes for 2017

Type	Tree - heading
	Apolipoprotein B
-	C18.452.584.500.937                      Smith-Lemli-Opitz Syndrome
-	C18.452.584.562                              Lipid Metabolism, Inborn Errors
-	C18.452.584.625                              Lipodystrophy
-	C18.452.584.625.400                        HIV-Associated Lipodystrophy Syndrome
-	C18.452.584.625.550                        Lipodystrophy, Congenital Generalized
-	C18.452.584.625.700                        Lipodystrophy, Familial Partial
-	C18.452.584.687                              Lipidoses
-	C18.452.584.687.201                        Cholesterol Ester Storage Disease
-	C18.452.584.687.201.500                    Wolman Disease
-	C18.452.584.687.509                        Neuronal Ceroid-Lipofuscinoses
-	C18.452.584.687.723                        Sjogren-Larsson Syndrome
-	C18.452.584.687.803                        Sphingolipidoses
-	C18.452.584.687.803.300                    Fabry Disease
-	C18.452.584.687.803.325                    Farber Lipogranulomatosis
-	C18.452.584.687.803.350                    Gangliosidoses
-	C18.452.584.687.803.350.300              Gangliosidoses, GM2
-	C18.452.584.687.803.350.300.700        Sandhoff Disease
-	C18.452.584.687.803.350.300.850        Tay-Sachs Disease
-	C18.452.584.687.803.350.300.925        Tay-Sachs Disease, AB Variant
-	C18.452.584.687.803.350.360              Gangliosidosis, GM1
-	C18.452.584.687.803.441                    Gaucher Disease
-	C18.452.584.687.803.585                    Leukodystrophy, Globoid Cell
-	C18.452.584.687.803.730                    Niemann-Pick Diseases
-	C18.452.584.687.803.730.500              Niemann-Pick Disease, Type A
-	C18.452.584.687.803.730.750              Niemann-Pick Disease, Type B
-	C18.452.584.687.803.730.875              Niemann-Pick Disease, Type C
-	C18.452.584.687.803.850                    Sea-Blue Histiocyte Syndrome
-	C18.452.584.687.803.925                    Sulfatidosis
-	C18.452.584.687.803.925.500              Leukodystrophy, Metachromatic
-	C18.452.584.687.803.925.750              Multiple Sulfatase Deficiency Disease
-	C18.452.584.718                              Lipomatosis
-	C18.452.584.718.500                        Adiposis Dolorosa
New Heading	<b>C18.452.584.718.625                        Familial Multiple Lipomatosis</b>
-	C18.452.584.718.750                        Lipomatosis, Multiple Symmetrical

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C18.452.584.750 Xanthomatosis
-	C18.452.584.750.487 Necrobiotic Xanthogranuloma
-	C18.452.584.750.975 Xanthomatosis, Cerebrotendinous
-	C18.452.603 Malabsorption Syndromes
-	C18.452.603.145 Blind Loop Syndrome
-	C18.452.603.250 Celiac Disease
-	C18.452.603.314 Collagenous Sprue
-	C18.452.603.378 Hyperhomocysteinemia
-	C18.452.603.506 Lactose Intolerance
-	C18.452.603.850 Sprue, Tropical
-	C18.452.603.887 Steatorrhea
-	C18.452.603.925 Whipple Disease
-	C18.452.625 Metabolic Syndrome X
-	C18.452.648 Metabolism, Inborn Errors
-	C18.452.648.100 Amino Acid Metabolism, Inborn Errors
-	C18.452.648.100.102 Albinism
-	C18.452.648.100.102.090 Albinism, Ocular
-	C18.452.648.100.102.100 Albinism, Oculocutaneous
-	C18.452.648.100.102.100.400 Hermanski-Pudlak Syndrome
-	C18.452.648.100.102.600 Piebaldism
-	C18.452.648.100.187 Alkaptonuria
-	C18.452.648.100.477 Hyperglycinemia, Nonketotic
-	C18.452.648.100.480 Hyperhomocysteinemia
-	C18.452.648.100.480.500 Homocystinuria
-	C18.452.648.100.544 Hyperlysinemias
-	C18.452.648.100.608 Maple Syrup Urine Disease
-	C18.452.648.100.614 Multiple Acyl Coenzyme A Dehydrogenase Deficiency
-	C18.452.648.100.620 Multiple Carboxylase Deficiency
-	C18.452.648.100.620.100 Biotinidase Deficiency
-	C18.452.648.100.620.380 Holocarboxylase Synthetase Deficiency
-	C18.452.648.100.766 Phenylketonurias
-	C18.452.648.100.766.500 Phenylketonuria, Maternal
-	C18.452.648.100.823 Propionic Acidemia
-	C18.452.648.100.880 Tyrosinemias
-	C18.452.648.100.940 Urea Cycle Disorders, Inborn
-	C18.452.648.100.940.124 Argininosuccinic Aciduria

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C18.452.648.100.940.249 Disease Carbamoyl-Phosphate Synthase I Deficiency
-	C18.452.648.100.940.374 Citrullinemia
-	C18.452.648.100.940.437 Hyperargininemia
-	C18.452.648.100.940.500 Ornithine Carbamoyltransferase Deficiency Disease
-	C18.452.648.151 Amino Acid Transport Disorders, Inborn
-	C18.452.648.151.355 Hartnup Disease
-	C18.452.648.151.600 Oculocerebrorenal Syndrome
-	C18.452.648.176 Amyloidosis, Familial
-	C18.452.648.176.050 Amyloid Neuropathies, Familial
-	C18.452.648.176.160 Cerebral Amyloid Angiopathy, Familial
-	C18.452.648.189 Brain Diseases, Metabolic, Inborn
-	C18.452.648.189.084 Adrenoleukodystrophy
-	C18.452.648.189.168 Cerebral Amyloid Angiopathy, Familial
-	C18.452.648.189.320 Galactosemias
-	C18.452.648.189.355 Hartnup Disease
-	C18.452.648.189.360 Hepatolenticular Degeneration
-	C18.452.648.189.362 Diseases Hereditary Central Nervous System Demyelinating
-	C18.452.648.189.362.250 Adrenoleukodystrophy
-	C18.452.648.189.362.312 Alexander Disease
-	C18.452.648.189.362.375 Canavan Disease
-	C18.452.648.189.362.500 Leukodystrophy, Globoid Cell
-	C18.452.648.189.362.550 Leukodystrophy, Metachromatic
-	C18.452.648.189.362.775 Pelizaeus-Merzbacher Disease
-	C18.452.648.189.365 Homocystinuria
-	C18.452.648.189.375 Hyperglycinemia, Nonketotic
-	C18.452.648.189.380 Hyperlysinemias
-	C18.452.648.189.412 Leigh Disease
-	C18.452.648.189.425 Lesch-Nyhan Syndrome
-	C18.452.648.189.435 Lysosomal Storage Diseases, Nervous System
-	C18.452.648.189.435.295 Fucosidosis
-	C18.452.648.189.435.340 Glycogen Storage Disease Type II
-	C18.452.648.189.435.590 Mucopolidoses
-	C18.452.648.189.435.810 Sialic Acid Storage Disease
-	C18.452.648.189.435.825 Sphingolipidoses

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C18.452.648.189.435.825.200 Fabry Disease
-	C18.452.648.189.435.825.250 Farber Lipogranulomatosis
-	C18.452.648.189.435.825.300 Gangliosidoses
-	C18.452.648.189.435.825.300.300 Gangliosidoses, GM2
-	C18.452.648.189.435.825.300.300.249 Sandhoff Disease
-	C18.452.648.189.435.825.300.300.500 Tay-Sachs Disease
-	C18.452.648.189.435.825.300.300.750 Tay-Sachs Disease, AB Variant
-	C18.452.648.189.435.825.300.400 Gangliosidosis, GM1
-	C18.452.648.189.435.825.400 Gaucher Disease
-	C18.452.648.189.435.825.590 Leukodystrophy, Globoid Cell
-	C18.452.648.189.435.825.700 Niemann-Pick Diseases
-	C18.452.648.189.435.825.700.500 Niemann-Pick Disease, Type A
-	C18.452.648.189.435.825.700.750 Niemann-Pick Disease, Type B
-	C18.452.648.189.435.825.700.875 Niemann-Pick Disease, Type C
-	C18.452.648.189.435.825.775 Sea-Blue Histiocyte Syndrome
-	C18.452.648.189.435.825.850 Sulfatidosis
-	C18.452.648.189.435.825.850.500 Leukodystrophy, Metachromatic
-	C18.452.648.189.435.825.850.750 Multiple Sulfatase Deficiency Disease
-	C18.452.648.189.520 Maple Syrup Urine Disease
-	C18.452.648.189.535 MELAS Syndrome
-	C18.452.648.189.540 Menkes Kinky Hair Syndrome
-	C18.452.648.189.545 MERRF Syndrome
-	C18.452.648.189.593 Mevalonate Kinase Deficiency
-	C18.452.648.189.640 Oculocerebrorenal Syndrome
-	C18.452.648.189.687 Phenylketonurias
-	C18.452.648.189.687.500 Phenylketonuria, Maternal
-	C18.452.648.189.725 Pyruvate Carboxylase Deficiency Disease
-	C18.452.648.189.750 Pyruvate Dehydrogenase Complex Deficiency Disease
-	C18.452.648.189.813 Refsum Disease
-	C18.452.648.189.844 Refsum Disease, Infantile
-	C18.452.648.189.875 Tyrosinemias
-	C18.452.648.189.937 Urea Cycle Disorders, Inborn
-	C18.452.648.189.937.124 Argininosuccinic Aciduria
-	C18.452.648.189.937.249 Carbamoyl-Phosphate Synthase I Deficiency Disease
-	C18.452.648.189.937.374 Citrullinemia



## MeSH Tree Changes for 2017

Type	Tree - heading
-	C18.452.648.189.937.437                      Hyperargininemia
-	C18.452.648.189.937.500                      Ornithine Carbamoyltransferase Deficiency Disease
-	C18.452.648.189.968                              Zellweger Syndrome
-	C18.452.648.202                                  Carbohydrate Metabolism, Inborn Errors
-	C18.452.648.202.125                              Congenital Disorders of Glycosylation
-	C18.452.648.202.251                              Fructose Metabolism, Inborn Errors
-	C18.452.648.202.251.221                              Fructose-1,6-Diphosphatase Deficiency
-	C18.452.648.202.251.271                              Fructose Intolerance
-	C18.452.648.202.303                              Fucosidosis
-	C18.452.648.202.355                              Galactosemias
-	C18.452.648.202.402                              Glucosephosphate Dehydrogenase Deficiency
-	C18.452.648.202.449                              Glycogen Storage Disease
-	C18.452.648.202.449.448                              Glycogen Storage Disease Type I
-	C18.452.648.202.449.500                              Glycogen Storage Disease Type II
-	C18.452.648.202.449.510                              Glycogen Storage Disease Type IIb
-	C18.452.648.202.449.520                              Glycogen Storage Disease Type III
-	C18.452.648.202.449.540                              Glycogen Storage Disease Type IV
-	C18.452.648.202.449.560                              Glycogen Storage Disease Type V
-	C18.452.648.202.449.580                              Glycogen Storage Disease Type VI
-	C18.452.648.202.449.600                              Glycogen Storage Disease Type VII
-	C18.452.648.202.449.620                              Glycogen Storage Disease Type VIII
-	C18.452.648.202.460                              Hyperoxaluria, Primary
-	C18.452.648.202.589                              Lactose Intolerance
-	C18.452.648.202.607                              Mannosidase Deficiency Diseases
-	C18.452.648.202.607.500                              alpha-Mannosidosis
-	C18.452.648.202.607.750                              beta-Mannosidosis
-	C18.452.648.202.670                              Mucopolidoses
-	C18.452.648.202.715                              Mucopolysaccharidoses
-	C18.452.648.202.715.640                              Mucopolysaccharidosis I
-	C18.452.648.202.715.645                              Mucopolysaccharidosis II
-	C18.452.648.202.715.650                              Mucopolysaccharidosis III
-	C18.452.648.202.715.655                              Mucopolysaccharidosis IV
-	C18.452.648.202.715.670                              Mucopolysaccharidosis VI
-	C18.452.648.202.715.675                              Mucopolysaccharidosis VII
-	C18.452.648.202.720                              Multiple Carboxylase Deficiency
-	C18.452.648.202.720.100                              Biotinidase Deficiency

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C18.452.648.202.720.380                      Holocarboxylase Synthetase Deficiency
-	C18.452.648.202.810                              Pyruvate Metabolism, Inborn Errors
-	C18.452.648.202.810.444                      Leigh Disease
-	C18.452.648.202.810.666                      Pyruvate Carboxylase Deficiency Disease
-	C18.452.648.202.810.766 Disease                              Pyruvate Dehydrogenase Complex Deficiency
-	C18.452.648.300                                  Hyperbilirubinemia, Hereditary
-	C18.452.648.300.281                              Crigler-Najjar Syndrome
-	C18.452.648.300.528                              Gilbert Disease
-	C18.452.648.300.764                              Jaundice, Chronic Idiopathic
-	C18.452.648.398                                  Lipid Metabolism, Inborn Errors
-	C18.452.648.398.224                              Barth Syndrome
-	C18.452.648.398.450                              Hyperlipidemia, Familial Combined
-	C18.452.648.398.465                              Hyperlipoproteinemia Type I
-	C18.452.648.398.481                              Hyperlipoproteinemia Type II
-	C18.452.648.398.483                              Hyperlipoproteinemia Type III
-	C18.452.648.398.487                              Hyperlipoproteinemia Type IV
-	C18.452.648.398.493                              Hyperlipoproteinemia Type V
-	C18.452.648.398.500                              Hypolipoproteinemias
-	C18.452.648.398.500.330                      Hypoalphalipoproteinemias
-	C18.452.648.398.500.330.500                  Lecithin Acyltransferase Deficiency
-	C18.452.648.398.500.330.500                  Lecithin Cholesterol Acyltransferase Deficiency
-	C18.452.648.398.500.330.750                  Tangier Disease
-	C18.452.648.398.500.440                      Hypobetalipoproteinemias
-	C18.452.648.398.500.440.500                  Abetalipoproteinemia
-	C18.452.648.398.641                              Lipidoses
-	C18.452.648.398.641.201                      Cholesterol Ester Storage Disease
-	C18.452.648.398.641.201.500                  Wolman Disease
-	C18.452.648.398.641.509                      Neuronal Ceroid-Lipofuscinoses
-	C18.452.648.398.641.723                      Sjogren-Larsson Syndrome
-	C18.452.648.398.641.803                      Sphingolipidoses
-	C18.452.648.398.641.803.300                  Fabry Disease
-	C18.452.648.398.641.803.325                  Farber Lipogranulomatosis
-	C18.452.648.398.641.803.350                  Gangliosidoses
-	C18.452.648.398.641.803.350.300              Gangliosidoses, GM2
-	C18.452.648.398.641.803.350.300.700      Sandhoff Disease

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C18.452.648.398.641.803.350.300.850 Tay-Sachs Disease
-	C18.452.648.398.641.803.350.300.925 Tay-Sachs Disease, AB Variant
-	C18.452.648.398.641.803.350.360 Gangliosidosis, GM1
-	C18.452.648.398.641.803.441 Gaucher Disease
-	C18.452.648.398.641.803.585 Leukodystrophy, Globoid Cell
-	C18.452.648.398.641.803.730 Niemann-Pick Diseases
-	C18.452.648.398.641.803.730.500 Niemann-Pick Disease, Type A
-	C18.452.648.398.641.803.730.750 Niemann-Pick Disease, Type B
-	C18.452.648.398.641.803.730.875 Niemann-Pick Disease, Type C
-	C18.452.648.398.641.803.850 Sea-Blue Histiocyte Syndrome
-	C18.452.648.398.641.803.925 Sulfatidosis
-	C18.452.648.398.641.803.925.500 Leukodystrophy, Metachromatic
-	C18.452.648.398.641.803.925.750 Multiple Sulfatase Deficiency Disease
-	C18.452.648.398.745 Lipodystrophy, Congenital Generalized
-	C18.452.648.398.850 Smith-Lemli-Opitz Syndrome
-	C18.452.648.398.925 Xanthomatosis, Cerebrotendinous
-	C18.452.648.595 Lysosomal Storage Diseases
-	C18.452.648.595.100 Aspartylglucosaminuria
-	C18.452.648.595.201 Cholesterol Ester Storage Disease
-	C18.452.648.595.201.500 Wolman Disease
-	C18.452.648.595.377 Cystinosis
-	C18.452.648.595.554 Lysosomal Storage Diseases, Nervous System
-	C18.452.648.595.554.295 Fucosidosis
-	C18.452.648.595.554.340 Glycogen Storage Disease Type II
-	C18.452.648.595.554.590 Mucopolidoses
-	C18.452.648.595.554.810 Sialic Acid Storage Disease
-	C18.452.648.595.554.825 Sphingolipidoses
-	C18.452.648.595.554.825.200 Fabry Disease
-	C18.452.648.595.554.825.250 Farber Lipogranulomatosis
-	C18.452.648.595.554.825.300 Gangliosidoses
-	C18.452.648.595.554.825.300.300 Gangliosidoses, GM2
-	C18.452.648.595.554.825.300.300.800 Sandhoff Disease
-	C18.452.648.595.554.825.300.300.840 Tay-Sachs Disease
-	C18.452.648.595.554.825.300.300.920 Tay-Sachs Disease, AB Variant
-	C18.452.648.595.554.825.300.400 Gangliosidosis, GM1
-	C18.452.648.595.554.825.400 Gaucher Disease

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C18.452.648.595.554.825.590      Leukodystrophy, Globoid Cell
-	C18.452.648.595.554.825.700      Niemann-Pick Diseases
-	C18.452.648.595.554.825.700.500      Niemann-Pick Disease, Type A
-	C18.452.648.595.554.825.700.750      Niemann-Pick Disease, Type B
-	C18.452.648.595.554.825.700.875      Niemann-Pick Disease, Type C
-	C18.452.648.595.554.825.775      Sea-Blue Histiocyte Syndrome
-	C18.452.648.595.554.825.850      Sulfatidosis
-	C18.452.648.595.554.825.850.500      Leukodystrophy, Metachromatic
-	C18.452.648.595.554.825.850.750      Multiple Sulfatase Deficiency Disease
-	C18.452.648.595.577      Mannosidase Deficiency Diseases
-	C18.452.648.595.577.500      alpha-Mannosidosis
-	C18.452.648.595.577.750      beta-Mannosidosis
-	C18.452.648.595.600      Mucopolysaccharidoses
-	C18.452.648.595.600.640      Mucopolysaccharidosis I
-	C18.452.648.595.600.645      Mucopolysaccharidosis II
-	C18.452.648.595.600.650      Mucopolysaccharidosis III
-	C18.452.648.595.600.655      Mucopolysaccharidosis IV
-	C18.452.648.595.600.670      Mucopolysaccharidosis VI
-	C18.452.648.595.600.675      Mucopolysaccharidosis VII
-	C18.452.648.595.800      Pycnodysostosis
-	C18.452.648.618      Metal Metabolism, Inborn Errors
-	C18.452.648.618.337      Hemochromatosis
-	C18.452.648.618.403      Hepatolenticular Degeneration
-	C18.452.648.618.482      Hypophosphatasia
-	C18.452.648.618.544      Hypophosphatemia, Familial
-	C18.452.648.618.544.500      Familial Hypophosphatemic Rickets
-	C18.452.648.618.590      Menkes Kinky Hair Syndrome
-	C18.452.648.618.711      Paralyzes, Familial Periodic
-	C18.452.648.618.711.550      Hypokalemic Periodic Paralysis
-	C18.452.648.618.711.600      Paralysis, Hyperkalemic Periodic
-	C18.452.648.618.815      Pseudohypoparathyroidism
-	C18.452.648.618.815.815      Pseudopseudohypoparathyroidism
-	C18.452.648.663      Peroxisomal Disorders
-	C18.452.648.663.050      Acatlasia
-	C18.452.648.663.100      Adrenoleukodystrophy
-	C18.452.648.663.265      Chondrodysplasia Punctata, Rhizomelic

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C18.452.648.663.430 Mevalonate Kinase Deficiency
-	C18.452.648.663.760 Refsum Disease
-	C18.452.648.663.865 Refsum Disease, Infantile
-	C18.452.648.663.970 Zellweger Syndrome
-	C18.452.648.753 Progeria
-	C18.452.648.798 Purine-Pyrimidine Metabolism, Inborn Errors
-	C18.452.648.798.183 Dihydropyrimidine Dehydrogenase Deficiency
-	C18.452.648.798.368 Gout
-	C18.452.648.798.368.410 Arthritis, Gouty
-	C18.452.648.798.594 Lesch-Nyhan Syndrome
-	C18.452.648.861 Renal Tubular Transport, Inborn Errors
-	C18.452.648.861.093 Acidosis, Renal Tubular
-	C18.452.648.861.271 Dent Disease
-	C18.452.648.861.450 Fanconi Syndrome
-	C18.452.648.861.491 Gitelman Syndrome
-	C18.452.648.861.532 Glycosuria, Renal
-	C18.452.648.861.647 Hypophosphatemia, Familial
-	C18.452.648.861.647.500 Familial Hypophosphatemic Rickets
-	C18.452.648.861.698 Liddle Syndrome
-	C18.452.648.861.750 Oculocerebrorenal Syndrome
-	C18.452.648.861.770 Pseudohypoaldosteronism
-	C18.452.648.861.885 Renal Aminoacidurias
-	C18.452.648.861.885.250 Cystinuria
-	C18.452.648.925 Steroid Metabolism, Inborn Errors
-	C18.452.648.925.249 Adrenal Hyperplasia, Congenital
-	C18.452.648.925.324 Antley-Bixler Syndrome Phenotype
-	C18.452.648.925.400 Ichthyosis, X-Linked
-	C18.452.648.925.500 Mineralocorticoid Excess Syndrome, Apparent
-	C18.452.648.925.875 Smith-Lemli-Opitz Syndrome
-	C18.452.660 Mitochondrial Diseases
-	C18.452.660.097 Carbamoyl-Phosphate Synthase I Deficiency Disease
-	C18.452.660.195 Cytochrome-c Oxidase Deficiency
-	C18.452.660.300 Friedreich Ataxia
-	C18.452.660.410 Kearns-Sayre Syndrome
-	C18.452.660.520 Leigh Disease
-	C18.452.660.560 Mitochondrial Myopathies

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C18.452.660.560.620 Mitochondrial Encephalomyopathies
-	C18.452.660.560.620.520 MELAS Syndrome
-	C18.452.660.560.620.530 MERRF Syndrome
-	C18.452.660.560.700 Ophthalmoplegia, Chronic Progressive External
-	C18.452.660.560.700.500 Kearns-Sayre Syndrome
-	C18.452.660.612 Multiple Acyl Coenzyme A Dehydrogenase Deficiency
-	C18.452.660.665 Optic Atrophy, Autosomal Dominant
-	C18.452.660.670 Optic Atrophy, Hereditary, Leber
-	C18.452.660.705 Pyruvate Carboxylase Deficiency Disease
-	C18.452.660.710 Pyruvate Dehydrogenase Complex Deficiency Disease
-	C18.452.750 Phosphorus Metabolism Disorders
-	C18.452.750.199 Hyperphosphatemia
-	C18.452.750.400 Hypophosphatemia
-	C18.452.750.400.500 Hypophosphatemia, Familial
-	C18.452.750.400.500.500 Familial Hypophosphatemic Rickets
-	C18.452.750.400.750 Rickets, Hypophosphatemic
-	C18.452.750.400.750.500 Familial Hypophosphatemic Rickets
-	C18.452.811 Porphyrrias
-	C18.452.811.250 Porphyria, Erythropoietic
-	C18.452.811.400 Porphyrrias, Hepatic
-	C18.452.811.400.074 Coproporphyrria, Hereditary
-	C18.452.811.400.150 Porphyria, Acute Intermittent
-	C18.452.811.400.250 Porphyria Cutanea Tarda
-	C18.452.811.400.437 Porphyria, Hepatoerythropoietic
-	C18.452.811.400.625 Porphyria, Variegate
-	C18.452.811.400.812 Protoporphyrria, Erythropoietic
-	C18.452.845 Proteostasis Deficiencies
-	C18.452.845.500 Amyloidosis
-	C18.452.845.500.050 Amyloid Neuropathies
-	C18.452.845.500.050.050 Amyloid Neuropathies, Familial
-	C18.452.845.500.075 Amyloidosis, Familial
-	C18.452.845.500.075.050 Amyloid Neuropathies, Familial
-	C18.452.845.500.075.160 Cerebral Amyloid Angiopathy, Familial
-	C18.452.845.500.100 Cerebral Amyloid Angiopathy
-	C18.452.845.500.100.160 Cerebral Amyloid Angiopathy, Familial
-	C18.452.845.800 TDP-43 Proteinopathies

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C18.452.845.800.050 Amyotrophic Lateral Sclerosis
-	C18.452.845.800.300 Frontotemporal Lobar Degeneration
-	C18.452.845.800.300.299 Frontotemporal Dementia
-	C18.452.845.800.300.600 Primary Progressive Nonfluent Aphasia
-	C18.452.880 Skin Diseases, Metabolic
-	C18.452.880.391 Lipodystrophy
-	C18.452.880.391.400 HIV-Associated Lipodystrophy Syndrome
-	C18.452.880.391.550 Lipodystrophy, Congenital Generalized
-	C18.452.880.391.700 Lipodystrophy, Familial Partial
-	C18.452.880.495 Necrobiosis Lipoidica
-	C18.452.915 Wasting Syndrome
-	C18.452.915.520 HIV Wasting Syndrome
-	C18.452.950 Water-Electrolyte Imbalance
-	C18.452.950.179 Dehydration
-	C18.452.950.340 Hypercalcemia
-	C18.452.950.396 Hyperkalemia
-	C18.452.950.452 Hyponatremia
-	C18.452.950.509 Hypocalcemia
-	C18.452.950.565 Hypokalemia
-	C18.452.950.620 Hyponatremia
-	C18.452.950.626 Inappropriate ADH Syndrome
-	C18.452.950.932 Water Intoxication
-	C18.654 Nutrition Disorders
-	C18.654.180 Child Nutrition Disorders
-	C18.654.301 Hypervitaminosis A
-	C18.654.422 Infant Nutrition Disorders
-	C18.654.422.360 Vitamin K Deficiency Bleeding
-	C18.654.521 Malnutrition
-	C18.654.521.500 Deficiency Diseases
-	C18.654.521.500.133 Avitaminosis
-	C18.654.521.500.133.115 Ascorbic Acid Deficiency
-	C18.654.521.500.133.115.661 Scurvy
-	C18.654.521.500.133.628 Vitamin A Deficiency
-	C18.654.521.500.133.699 Vitamin B Deficiency
-	C18.654.521.500.133.699.160 Choline Deficiency
-	C18.654.521.500.133.699.308 Folic Acid Deficiency

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C18.654.521.500.133.699.418      Hyperhomocysteinemia
-	C18.654.521.500.133.699.529      Pellagra
-	C18.654.521.500.133.699.713      Riboflavin Deficiency
-	C18.654.521.500.133.699.827      Thiamine Deficiency
-	C18.654.521.500.133.699.827.223      Beriberi
-	C18.654.521.500.133.699.827.822      Wernicke Encephalopathy
-	C18.654.521.500.133.699.901      Vitamin B 6 Deficiency
-	C18.654.521.500.133.699.923      Vitamin B 12 Deficiency
-	C18.654.521.500.133.699.923.280      Anemia, Pernicious
-	C18.654.521.500.133.699.923.640      Subacute Combined Degeneration
-	C18.654.521.500.133.770      Vitamin D Deficiency
-	C18.654.521.500.133.770.734      Rickets
-	C18.654.521.500.133.770.734.640      Osteomalacia
-	C18.654.521.500.133.770.734.750      Chronic Kidney Disease-Mineral and Bone Disorder
-	C18.654.521.500.133.770.734.750      Renal Osteodystrophy
-	C18.654.521.500.133.770.734.875      Rickets, Hypophosphatemic
-	C18.654.521.500.133.770.734.875.500      Familial Hypophosphatemic Rickets
-	C18.654.521.500.133.841      Vitamin E Deficiency
-	C18.654.521.500.133.841.682      Steatitis
-	C18.654.521.500.133.912      Vitamin K Deficiency
-	C18.654.521.500.133.912.360      Vitamin K Deficiency Bleeding
-	C18.654.521.500.439      Magnesium Deficiency
-	C18.654.521.500.617      Potassium Deficiency
-	C18.654.521.500.708      Protein Deficiency
-	C18.654.521.500.708.626      Protein-Energy Malnutrition
-	C18.654.521.500.857      Swayback
-	C18.654.521.625      Fetal Nutrition Disorders
-	C18.654.521.687      Refeeding Syndrome
-	C18.654.521.719      Severe Acute Malnutrition
-	C18.654.521.719.500      Kwashiorkor
-	C18.654.521.750      Starvation
-	C18.654.726      Overnutrition
-	C18.654.726.500      Obesity
-	C18.654.726.500.695      Obesity Hypoventilation Syndrome
-	C18.654.726.500.697      Obesity, Abdominal



## MeSH Tree Changes for 2017

Type	Tree - heading
-	C18.654.726.500.698                      Obesity, Metabolically Benign
-	C18.654.726.500.700                      Obesity, Morbid
-	C18.654.726.500.720                      Pediatric Obesity
-	C18.654.726.500.740                      Prader-Willi Syndrome
-	C18.654.940                                      Wasting Syndrome
-	C18.654.940.520                              HIV Wasting Syndrome
-	C19    Endocrine System Diseases
-	C19.053    Adrenal Gland Diseases
-	C19.053.098                                    Adrenal Cortex Diseases
-	C19.053.098.265                              Adrenal Cortex Neoplasms
-	C19.053.098.265.500                        Adrenocortical Adenoma
-	C19.053.098.265.750                        Adrenocortical Carcinoma
-	C19.053.347                                    Adrenal Gland Neoplasms
-	C19.053.347.500                              Adrenal Cortex Neoplasms
-	C19.053.347.500.500                        Adrenocortical Adenoma
-	C19.053.347.500.750                        Adrenocortical Carcinoma
-	C19.053.440                                    Adrenal Hyperplasia, Congenital
-	C19.053.500                                    Adrenal Insufficiency
-	C19.053.500.263                              Addison Disease
-	C19.053.500.270                              Adrenoleukodystrophy
-	C19.053.500.480                              Hypoaldosteronism
-	C19.053.500.740                              Waterhouse-Friderichsen Syndrome
-	C19.053.800                                    Adrenocortical Hyperfunction
-	C19.053.800.367                              Cushing Syndrome
-	C19.053.800.604                              Hyperaldosteronism
-	C19.053.800.604.249                        Bartter Syndrome
-	C19.149    Bone Diseases, Endocrine
-	C19.246    Diabetes Mellitus
-	C19.246.099                                    Diabetes Complications
-	C19.246.099.500                              Diabetic Angiopathies
-	C19.246.099.500.191                        Diabetic Foot
-	C19.246.099.500.382                        Diabetic Retinopathy
-	C19.246.099.625                              Diabetic Cardiomyopathies
-	C19.246.099.750                              Diabetic Coma
-	C19.246.099.750.490                        Hyperglycemic Hyperosmolar Nonketotic Coma
-	C19.246.099.812                              Diabetic Ketoacidosis

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C19.246.099.875 Diabetic Nephropathies
-	C19.246.099.937 Diabetic Neuropathies
-	C19.246.099.937.250 Diabetic Foot
-	C19.246.099.968 Fetal Macrosomia
-	C19.246.200 Diabetes, Gestational
-	C19.246.240 Diabetes Mellitus, Experimental
-	C19.246.267 Diabetes Mellitus, Type 1
-	C19.246.267.960 Wolfram Syndrome
-	C19.246.300 Diabetes Mellitus, Type 2
-	C19.246.300.500 Diabetes Mellitus, Lipoatrophic
-	C19.246.537 Donohue Syndrome
New Heading	<b>C19.246.656 Latent Autoimmune Diabetes in Adults</b>
-	C19.246.774 Prediabetic State
-	C19.297 Dwarfism
-	C19.297.155 Congenital Hypothyroidism
-	C19.297.312 Dwarfism, Pituitary
-	C19.297.656 Laron Syndrome
-	C19.344 Endocrine Gland Neoplasms
-	C19.344.078 Adrenal Gland Neoplasms
-	C19.344.078.265 Adrenal Cortex Neoplasms
-	C19.344.078.265.500 Adrenocortical Adenoma
-	C19.344.078.265.750 Adrenocortical Carcinoma
-	C19.344.400 Multiple Endocrine Neoplasia
-	C19.344.400.500 Multiple Endocrine Neoplasia Type 1
-	C19.344.400.505 Multiple Endocrine Neoplasia Type 2a
-	C19.344.400.510 Multiple Endocrine Neoplasia Type 2b
-	C19.344.410 Ovarian Neoplasms
-	C19.344.410.398 Granulosa Cell Tumor
-	C19.344.410.431 Hereditary Breast and Ovarian Cancer Syndrome
-	C19.344.410.464 Luteoma
-	C19.344.410.531 Meigs Syndrome
-	C19.344.410.648 Sertoli-Leydig Cell Tumor
-	C19.344.410.765 Thecoma
-	C19.344.421 Pancreatic Neoplasms
-	C19.344.421.249 Adenoma, Islet Cell

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C19.344.421.249.500 Insulinoma
-	C19.344.421.500 Carcinoma, Islet Cell
-	C19.344.421.500.124 Gastrinoma
-	C19.344.421.500.249 Glucagonoma
-	C19.344.421.500.500 Somatostatinoma
-	C19.344.421.500.750 Vipoma
-	C19.344.421.750 Carcinoma, Pancreatic Ductal
-	C19.344.473 Paraneoplastic Endocrine Syndromes
-	C19.344.525 Parathyroid Neoplasms
-	C19.344.609 Pituitary Neoplasms
-	C19.344.609.145 ACTH-Secreting Pituitary Adenoma
-	C19.344.609.145.500 Nelson Syndrome
-	C19.344.609.292 Growth Hormone-Secreting Pituitary Adenoma
-	C19.344.609.792 Prolactinoma
-	C19.344.762 Testicular Neoplasms
-	C19.344.762.500 Sertoli-Leydig Cell Tumor
-	C19.344.762.500.249 Leydig Cell Tumor
-	C19.344.762.500.500 Sertoli Cell Tumor
-	C19.344.894 Thyroid Neoplasms
-	C19.344.894.800 Thyroid Nodule
New Tree	<a href="#">C19.368</a> <a href="#">Female Athlete Triad Syndrome</a>
-	C19.391 Gonadal Disorders
-	C19.391.119 Disorders of Sex Development
-	C19.391.119.064 46, XX Disorders of Sex Development
-	C19.391.119.064.124 46, XX Testicular Disorders of Sex Development
-	C19.391.119.064.249 Gonadal Dysgenesis, 46,XX
-	C19.391.119.064.500 Hyperandrogenism
-	C19.391.119.096 46, XY Disorders of Sex Development
-	C19.391.119.096.500 Androgen-Insensitivity Syndrome
-	C19.391.119.096.562 Denys-Drash Syndrome
-	C19.391.119.096.624 Frasier Syndrome
-	C19.391.119.096.687 Gonadal Dysgenesis, 46,XY
-	C19.391.119.096.687.500 Gonadoblastoma
-	C19.391.119.096.750 Kallmann Syndrome
-	C19.391.119.096.875 WAGR Syndrome

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C19.391.119.129 Adrenogenital Syndrome
-	C19.391.119.129.500 Adrenal Hyperplasia, Congenital
-	C19.391.119.129.750 Hyperandrogenism
-	C19.391.119.309 Gonadal Dysgenesis
-	C19.391.119.309.193 Gonadal Dysgenesis, 46,XX
-	C19.391.119.309.388 Gonadal Dysgenesis, 46,XY
-	C19.391.119.309.388.500 Gonadoblastoma
-	C19.391.119.309.391 Gonadal Dysgenesis, Mixed
-	C19.391.119.309.631 Sexual Infantilism
-	C19.391.119.309.872 Turner Syndrome
-	C19.391.119.343 Ovotesticular Disorders of Sex Development
-	C19.391.119.795 Sex Chromosome Disorders of Sex Development
-	C19.391.119.795.124 Freemartinism
-	C19.391.119.795.249 Gonadal Dysgenesis, Mixed
-	C19.391.119.795.500 Klinefelter Syndrome
-	C19.391.119.795.750 Turner Syndrome
Old Tree	<b>C19.391.240 Female Athlete Triad Syndrome</b>
-	C19.391.482 Hypogonadism
-	C19.391.482.293 Eunuchism
-	C19.391.482.600 Kallmann Syndrome
-	C19.391.482.629 Klinefelter Syndrome
-	C19.391.482.814 Sexual Infantilism
-	C19.391.630 Ovarian Diseases
-	C19.391.630.050 Anovulation
-	C19.391.630.450 Oophoritis
-	C19.391.630.580 Ovarian Cysts
-	C19.391.630.580.765 Polycystic Ovary Syndrome
-	C19.391.630.642 Ovarian Hyperstimulation Syndrome
-	C19.391.630.705 Ovarian Neoplasms
-	C19.391.630.705.265 Brenner Tumor
-	C19.391.630.705.331 Carcinoma, Endometrioid
-	C19.391.630.705.398 Granulosa Cell Tumor
-	C19.391.630.705.431 Hereditary Breast and Ovarian Cancer Syndrome
-	C19.391.630.705.464 Luteoma
-	C19.391.630.705.531 Meigs Syndrome
-	C19.391.630.705.648 Sertoli-Leydig Cell Tumor

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	C19.391.630.705.765	Thecoma
-	C19.391.630.750	Primary Ovarian Insufficiency
-	C19.391.690	Puberty, Delayed
-	C19.391.693	Puberty, Precocious
-	C19.391.829	Testicular Diseases
-	C19.391.829.258	Cryptorchidism
-	C19.391.829.493	Orchitis
-	C19.391.829.782	Testicular Neoplasms
-	C19.391.829.782.500	Sertoli-Leydig Cell Tumor
-	C19.391.829.782.500.249	Leydig Cell Tumor
-	C19.391.829.782.500.500	Sertoli Cell Tumor
-	C19.642	Parathyroid Diseases
-	C19.642.355	Hyperparathyroidism
-	C19.642.355.239	Hyperparathyroidism, Primary
-	C19.642.355.480	Hyperparathyroidism, Secondary
-	C19.642.355.480.500	Chronic Kidney Disease-Mineral and Bone Disorder
-	C19.642.355.480.500	Renal Osteodystrophy
-	C19.642.482	Hypoparathyroidism
-	C19.642.482.500	22q11 Deletion Syndrome
-	C19.642.482.500.500	DiGeorge Syndrome
-	C19.642.713	Parathyroid Neoplasms
-	C19.700	Pituitary Diseases
Old Tree	<b>C19.700.080</b>	<b>Autoimmune Hypophysitis</b>
-	C19.700.159	Diabetes Insipidus
-	C19.700.159.750	Diabetes Insipidus, Neurogenic
-	C19.700.159.875	Wolfram Syndrome
-	C19.700.320	Empty Sella Syndrome
-	C19.700.355	Hyperpituitarism
-	C19.700.355.179	Acromegaly
-	C19.700.355.528	Gigantism
-	C19.700.355.600	Hyperprolactinemia
-	C19.700.355.800	Pituitary ACTH Hypersecretion
New Heading	<b>C19.700.419</b>	<b>Hypophysitis</b>
New Tree	<b>C19.700.419.500</b>	<b>Autoimmune Hypophysitis</b>

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C19.700.482 Hypopituitarism
-	C19.700.482.311 Dwarfism, Pituitary
-	C19.700.490 Inappropriate ADH Syndrome
-	C19.700.725 Pituitary Apoplexy
-	C19.700.734 Pituitary Neoplasms
-	C19.700.734.145 ACTH-Secreting Pituitary Adenoma
-	C19.700.734.145.500 Nelson Syndrome
-	C19.700.734.292 Growth Hormone-Secreting Pituitary Adenoma
-	C19.700.734.792 Prolactinoma
-	C19.787 Polyendocrinopathies, Autoimmune
-	C19.874 Thyroid Diseases
-	C19.874.255 Euthyroid Sick Syndromes
-	C19.874.283 Goiter
-	C19.874.283.300 Goiter, Endemic
-	C19.874.283.501 Goiter, Nodular
-	C19.874.283.601 Goiter, Substernal
-	C19.874.283.605 Graves Disease
-	C19.874.283.605.500 Graves Ophthalmopathy
-	C19.874.283.802 Lingual Goiter
-	C19.874.397 Hyperthyroidism
-	C19.874.397.370 Graves Disease
-	C19.874.397.370.500 Graves Ophthalmopathy
-	C19.874.397.685 Thyrotoxicosis
-	C19.874.397.685.905 Thyroid Crisis
-	C19.874.410 Hyperthyroxinemia
-	C19.874.410.249 Hyperthyroxinemia, Familial Dysalbuminemic
-	C19.874.410.500 Thyroid Hormone Resistance Syndrome
-	C19.874.482 Hypothyroidism
-	C19.874.482.281 Congenital Hypothyroidism
-	C19.874.482.638 Myxedema
-	C19.874.689 Thyroid Dysgenesis
-	C19.874.689.500 Lingual Thyroid
-	C19.874.689.500.500 Lingual Goiter
-	C19.874.788 Thyroid Neoplasms
-	C19.874.788.800 Thyroid Nodule
-	C19.874.871 Thyroiditis

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C19.874.871.102                      Thyroiditis, Autoimmune
-	C19.874.871.102.500                      Hashimoto Disease
-	C19.874.871.102.750                      Postpartum Thyroiditis
-	C19.874.871.900                      Thyroiditis, Subacute
-	C19.874.871.910                      Thyroiditis, Suppurative
-	C19.927                      Tuberculosis, Endocrine
-	C20                      Immune System Diseases
-	C20.111                      Autoimmune Diseases
-	C20.111.163                      Addison Disease
-	C20.111.175                      Anemia, Hemolytic, Autoimmune
-	C20.111.190                      Anti-Glomerular Basement Membrane Disease
-	C20.111.193                      Anti-Neutrophil Cytoplasmic Antibody-Associated Vasculitis
-	C20.111.193.500                      Churg-Strauss Syndrome
-	C20.111.193.750                      Microscopic Polyangiitis
-	C20.111.193.875                      Granulomatosis with Polyangiitis
-	C20.111.197                      Antiphospholipid Syndrome
-	C20.111.198                      Arthritis, Juvenile
-	C20.111.199                      Arthritis, Rheumatoid
-	C20.111.199.389                      Felty Syndrome
-	C20.111.199.581                      Rheumatoid Vasculitis
-	C20.111.199.774                      Sjogren's Syndrome
-	C20.111.199.870                      Still's Disease, Adult-Onset
-	C20.111.258                      Autoimmune Diseases of the Nervous System
-	C20.111.258.124                      Anti-N-Methyl-D-Aspartate Receptor Encephalitis
-	C20.111.258.250                      Demyelinating Autoimmune Diseases, CNS
-	C20.111.258.250.175                      Diffuse Cerebral Sclerosis of Schilder
-	C20.111.258.250.350                      Encephalomyelitis, Acute Disseminated
-	C20.111.258.250.350.500                      Leukoencephalitis, Acute Hemorrhagic
-	C20.111.258.250.425                      Leukoencephalitis, Acute Hemorrhagic
-	C20.111.258.250.500                      Multiple Sclerosis
-	C20.111.258.250.500.200                      Multiple Sclerosis, Chronic Progressive
-	C20.111.258.250.500.600                      Multiple Sclerosis, Relapsing-Remitting
-	C20.111.258.250.550                      Myelitis, Transverse
-	C20.111.258.250.550.500                      Neuromyelitis Optica
-	C20.111.258.250.775                      Neuromyelitis Optica
-	C20.111.258.350                      Lambert-Eaton Myasthenic Syndrome

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C20.111.258.500 Myasthenia Gravis
-	C20.111.258.500.300 Myasthenia Gravis, Autoimmune, Experimental
-	C20.111.258.500.650 Myasthenia Gravis, Neonatal
-	C20.111.258.625 Nervous System Autoimmune Disease, Experimental
-	C20.111.258.625.300 Encephalomyelitis, Autoimmune, Experimental
-	C20.111.258.625.350 Myasthenia Gravis, Autoimmune, Experimental
-	C20.111.258.625.700 Neuritis, Autoimmune, Experimental
-	C20.111.258.750 Polyradiculoneuropathy
-	C20.111.258.750.400 Guillain-Barre Syndrome
-	C20.111.258.750.400.500 Miller Fisher Syndrome
-	C20.111.258.750.800 Polyradiculoneuropathy, Chronic Inflammatory Demyelinating
-	C20.111.258.850 Stiff-Person Syndrome
-	C20.111.258.925 Uveomeningoencephalitic Syndrome
-	C20.111.258.962 Vasculitis, Central Nervous System
-	C20.111.258.962.800 Giant Cell Arteritis
-	C20.111.258.962.900 Lupus Vasculitis, Central Nervous System
-	C20.111.273 Autoimmune Hypophysitis
-	C20.111.288 Autoimmune Lymphoproliferative Syndrome
-	C20.111.318 Dermatitis Herpetiformis
-	C20.111.327 Diabetes Mellitus, Type 1
-	C20.111.525 Glomerulonephritis, IGA
-	C20.111.535 Glomerulonephritis, Membranous
-	C20.111.555 Graves Disease
-	C20.111.555.500 Graves Ophthalmopathy
-	C20.111.567 Hepatitis, Autoimmune
New Heading	<b>C20.111.576 Latent Autoimmune Diabetes in Adults</b>
-	C20.111.585 Linear IgA Bullous Dermatitis
-	C20.111.590 Lupus Erythematosus, Systemic
-	C20.111.590.560 Lupus Nephritis
-	C20.111.590.750 Lupus Vasculitis, Central Nervous System
-	C20.111.709 Ophthalmia, Sympathetic
-	C20.111.730 Pemphigoid, Bullous
-	C20.111.736 Pemphigus
-	C20.111.750 Polyendocrinopathies, Autoimmune



## MeSH Tree Changes for 2017

Type	Tree - heading
-	C20.111.759                      Purpura, Thrombocytopenic, Idiopathic
-	C20.111.809                      Thyroiditis, Autoimmune
-	C20.306                              Erythroblastosis, Fetal
-	C20.306.480                      Hydrops Fetalis
-	C20.306.502                      Kernicterus
-	C20.425                              Glomerulonephritis, Membranoproliferative
-	C20.452                              Graft vs Host Disease
-	C20.543                              Hypersensitivity
-	C20.543.206                      Drug Hypersensitivity
-	C20.543.206.189                      Asthma, Aspirin-Induced
-	C20.543.206.380                      Drug Eruptions
-	C20.543.206.380.174                      Acute Generalized Exanthematous Pustulosis
-	C20.543.206.380.262                      Drug Hypersensitivity Syndrome
-	C20.543.206.380.375                      Erythema Nodosum
-	C20.543.206.380.587                      Nicolau Syndrome
-	C20.543.206.380.800                      Serum Sickness
-	C20.543.206.380.900                      Stevens-Johnson Syndrome
-	C20.543.312                      Environmental Illness
-	C20.543.312.500                      Multiple Chemical Sensitivity
-	C20.543.312.750                      Sick Building Syndrome
-	C20.543.418                      Hypersensitivity, Delayed
-	C20.543.418.150                      Dermatitis, Allergic Contact
-	C20.543.418.150.600                      Dermatitis, Photoallergic
-	C20.543.418.150.700                      Dermatitis, Toxicodendron
-	C20.543.480                      Hypersensitivity, Immediate
-	C20.543.480.099                      Anaphylaxis
-	C20.543.480.149                      Asthma, Aspirin-Induced
-	C20.543.480.200                      Conjunctivitis, Allergic
-	C20.543.480.343                      Dermatitis, Atopic
-	C20.543.480.356                      Eosinophilic Esophagitis
-	C20.543.480.370                      Food Hypersensitivity
-	C20.543.480.370.150                      Egg Hypersensitivity
-	C20.543.480.370.500                      Milk Hypersensitivity
-	C20.543.480.370.550                      Nut Hypersensitivity
-	C20.543.480.370.675                      Peanut Hypersensitivity
-	C20.543.480.370.763                      Shellfish Hypersensitivity

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C20.543.480.370.850                      Wheat Hypersensitivity
-	C20.543.480.680                            Respiratory Hypersensitivity
-	C20.543.480.680.075                      Alveolitis, Extrinsic Allergic
-	C20.543.480.680.075.125                      Bird Fancier's Lung
-	C20.543.480.680.075.365                      Farmer's Lung
-	C20.543.480.680.075.682                      Trichosporonosis
-	C20.543.480.680.085                      Aspergillosis, Allergic Bronchopulmonary
-	C20.543.480.680.095                      Asthma
-	C20.543.480.680.095.110                      Asthma, Exercise-Induced
-	C20.543.480.680.095.495                      Asthma, Occupational
-	C20.543.480.680.095.880                      Status Asthmaticus
-	C20.543.480.680.443                      Rhinitis, Allergic
-	C20.543.480.680.443.500                      Rhinitis, Allergic, Perennial
-	C20.543.480.680.443.750                      Rhinitis, Allergic, Seasonal
-	C20.543.480.904                            Urticaria
-	C20.543.480.904.066                      Angioedema
-	C20.543.480.904.066.500                      Angioedemas, Hereditary
-	C20.543.480.904.066.500.500                      Hereditary Angioedema Type III
-	C20.543.480.904.066.500.750                      Hereditary Angioedema Types I and II
-	C20.543.520                                  Immune Complex Diseases
-	C20.543.520.100                              Arthus Reaction
-	C20.543.520.600                              Purpura, Schoenlein-Henoch
-	C20.543.520.770                              Serum Sickness
-	C20.543.520.910                              Vasculitis, Leukocytoclastic, Cutaneous
-	C20.543.600                                  Latex Hypersensitivity
-	C20.543.928                                  Wissler's Syndrome
-	C20.608    Immune Reconstitution Inflammatory Syndrome
-	C20.673    Immunologic Deficiency Syndromes
-	C20.673.088                                  Agammaglobulinemia
-	C20.673.290                                  Ataxia Telangiectasia
-	C20.673.330                                  Common Variable Immunodeficiency
-	C20.673.430                                  Dysgammaglobulinemia
-	C20.673.430.249                              Hyper-IgM Immunodeficiency Syndrome
-	C20.673.430.249.500                              Hyper-IgM Immunodeficiency Syndrome, Type 1
-	C20.673.430.500                              IgA Deficiency
-	C20.673.430.750                              IgG Deficiency

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C20.673.480 HIV Infections
-	C20.673.480.040 Acquired Immunodeficiency Syndrome
New Heading	<b>C20.673.480.044 Acute Retroviral Syndrome</b>
-	C20.673.480.048 AIDS Arteritis, Central Nervous System
-	C20.673.480.050 AIDS-Associated Nephropathy
-	C20.673.480.070 AIDS Dementia Complex
-	C20.673.480.080 AIDS-Related Complex
-	C20.673.480.100 AIDS-Related Opportunistic Infections
-	C20.673.480.400 HIV-Associated Lipodystrophy Syndrome
-	C20.673.480.480 HIV Enteropathy
-	C20.673.480.500 HIV Seropositivity
-	C20.673.480.520 HIV Wasting Syndrome
-	C20.673.483 Deltaretrovirus Infections
-	C20.673.483.260 Enzootic Bovine Leukosis
-	C20.673.483.470 HTLV-I Infections
-	C20.673.483.480 HTLV-II Infections
-	C20.673.600 Leukocyte-Adhesion Deficiency Syndrome
-	C20.673.627 Lymphopenia
-	C20.673.627.800 T-Lymphocytopenia, Idiopathic CD4-Positive
-	C20.673.627.900 Wiskott-Aldrich Syndrome
-	C20.673.774 Phagocyte Bactericidal Dysfunction
-	C20.673.774.257 Chediak-Higashi Syndrome
-	C20.673.774.257.270 Aleutian Mink Disease
-	C20.673.774.535 Granulomatous Disease, Chronic
-	C20.673.774.600 Job Syndrome
-	C20.673.815 Severe Combined Immunodeficiency
-	C20.673.815.500 X-Linked Combined Immunodeficiency Diseases
-	C20.683 Immunoproliferative Disorders
-	C20.683.460 Hypergammaglobulinemia
-	C20.683.460.319 Mevalonate Kinase Deficiency
-	C20.683.460.640 Monoclonal Gammopathy of Undetermined Significance
-	C20.683.515 Lymphoproliferative Disorders
-	C20.683.515.124 Autoimmune Lymphoproliferative Syndrome
-	C20.683.515.250 Giant Lymph Node Hyperplasia
-	C20.683.515.501 Immunoblastic Lymphadenopathy

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C20.683.515.512 Immunoproliferative Small Intestinal Disease
-	C20.683.515.515 Infectious Mononucleosis
-	C20.683.515.517 Leukemia, Hairy Cell
-	C20.683.515.528 Leukemia, Lymphoid
-	C20.683.515.528.080 Leukemia, B-Cell
-	C20.683.515.528.080.125 Leukemia, Lymphocytic, Chronic, B-Cell
-	C20.683.515.528.080.562 Leukemia, Prolymphocytic, B-Cell
-	C20.683.515.528.100 Leukemia, Biphenotypic, Acute
-	C20.683.515.528.565 Leukemia, Prolymphocytic
-	C20.683.515.528.565.745 Leukemia, Prolymphocytic, B-Cell
-	C20.683.515.528.565.750 Leukemia, Prolymphocytic, T-Cell
-	C20.683.515.528.582 Leukemia, T-Cell
-	C20.683.515.528.582.049 Leukemia, Large Granular Lymphocytic
-	C20.683.515.528.582.100 Leukemia-Lymphoma, Adult T-Cell
-	C20.683.515.528.582.125 Leukemia, Prolymphocytic, T-Cell
-	C20.683.515.528.600 Precursor Cell Lymphoblastic Leukemia-Lymphoma
-	C20.683.515.528.600.600 Precursor B-Cell Lymphoblastic Leukemia-Lymphoma
-	C20.683.515.528.600.620 Precursor T-Cell Lymphoblastic Leukemia-Lymphoma
-	C20.683.515.710 Lymphangiomyoma
-	C20.683.515.710.465 Lymphangioliomyomatosis
-	C20.683.515.761 Lymphoma
-	C20.683.515.761.150 Composite Lymphoma
-	C20.683.515.761.355 Hodgkin Disease
-	C20.683.515.761.417 Intraocular Lymphoma
-	C20.683.515.761.480 Lymphoma, Non-Hodgkin
-	C20.683.515.761.480.150 Lymphoma, B-Cell
-	C20.683.515.761.480.150.165 Burkitt Lymphoma
-	C20.683.515.761.480.150.450 Lymphoma, AIDS-Related
-	C20.683.515.761.480.150.570 Lymphoma, B-Cell, Marginal Zone
-	C20.683.515.761.480.150.585 Lymphoma, Large B-Cell, Diffuse
-	C20.683.515.761.480.150.585.500 Plasmablastic Lymphoma
-	C20.683.515.761.480.150.592 Lymphoma, Primary Effusion
-	C20.683.515.761.480.150.600 Lymphomatoid Granulomatosis
-	C20.683.515.761.480.350 Lymphoma, Follicular

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C20.683.515.761.480.487 Lymphoma, Large-Cell, Immunoblastic
-	C20.683.515.761.480.525 Lymphoma, Mantle-Cell
-	C20.683.515.761.480.750 Lymphoma, T-Cell
-	C20.683.515.761.480.750.199 Enteropathy-Associated T-Cell Lymphoma
-	C20.683.515.761.480.750.399 Lymphoma, Large-Cell, Anaplastic
-	C20.683.515.761.480.750.800 Lymphoma, T-Cell, Cutaneous
-	C20.683.515.761.480.750.800.507 Lymphoma, Primary Cutaneous Anaplastic Large Cell
-	C20.683.515.761.480.750.800.528 Lymphomatoid Papulosis
-	C20.683.515.761.480.750.800.550 Mycosis Fungoides
-	C20.683.515.761.480.750.800.550.600 Pagetoid Reticulosis
-	C20.683.515.761.480.750.800.775 Sezary Syndrome
-	C20.683.515.761.480.750.825 Lymphoma, T-Cell, Peripheral
-	C20.683.515.800 Macrophage Activation Syndrome
-	C20.683.515.840 Marek Disease
-	C20.683.515.845 Multiple Myeloma
-	C20.683.515.845.500 Leukemia, Plasma Cell
-	C20.683.515.880 Plasmacytoma
-	C20.683.515.920 Sezary Syndrome
-	C20.683.515.950 Tumor Lysis Syndrome
-	C20.683.780 Paraproteinemias
-	C20.683.780.250 Cryoglobulinemia
-	C20.683.780.490 Heavy Chain Disease
-	C20.683.780.490.512 Immunoproliferative Small Intestinal Disease
-	C20.683.780.640 Monoclonal Gammopathy of Undetermined Significance
-	C20.683.780.640.700 Schnitzler Syndrome
-	C20.683.780.650 Multiple Myeloma
-	C20.683.780.750 POEMS Syndrome
-	C20.683.780.925 Waldenstrom Macroglobulinemia
-	C20.841 Purpura, Thrombocytopenic
-	C20.841.600 Purpura, Thrombocytopenic, Idiopathic
-	C20.920 Transfusion Reaction
-	C21 Disorders of Environmental Origin
-	C21.223 Environmental Illness
-	C21.223.500 Multiple Chemical Sensitivity
-	C21.223.750 Sick Building Syndrome

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C21.676 Preconception Injuries
-	C22 Animal Diseases
-	C22.021 Abortion, Veterinary
-	C22.021.322 Brucellosis, Bovine
-	C22.039 Actinobacillosis
-	C22.062 Aleutian Mink Disease
-	C22.073 Anal Gland Neoplasms
-	C22.085 Anaplasmosis
-	C22.131 Bird Diseases
-	C22.131.094 Avian Leukosis
-	C22.131.321 Fowlpox
-	C22.131.450 Influenza in Birds
-	C22.131.498 Malaria, Avian
-	C22.131.546 Marek Disease
-	C22.131.630 Newcastle Disease
-	C22.131.728 Poultry Diseases
-	C22.131.728.250 Enteritis, Transmissible, of Turkeys
-	C22.131.728.650 Poult Enteritis Mortality Syndrome
-	C22.131.780 Reticuloendotheliosis, Avian
-	C22.131.800 Sarcoma, Avian
-	C22.131.921 Tuberculosis, Avian
-	C22.152 Borna Disease
-	C22.180 Cat Diseases
-	C22.180.350 Feline Acquired Immunodeficiency Syndrome
-	C22.180.440 Feline Infectious Peritonitis
-	C22.180.460 Feline Panleukopenia
-	C22.180.500 Leukemia, Feline
-	C22.196 Cattle Diseases
-	C22.196.090 Bovine Respiratory Disease Complex
-	C22.196.090.600 Pasteurellosis, Pneumonic
-	C22.196.090.660 Pneumonia, Atypical Interstitial, of Cattle
-	C22.196.090.675 Pneumonia of Calves, Enzootic
-	C22.196.106 Bovine Virus Diarrhea-Mucosal Disease
-	C22.196.148 Brucellosis, Bovine
-	C22.196.250 Encephalopathy, Bovine Spongiform
-	C22.196.260 Enzootic Bovine Leukosis

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	C22.196.274	Ephemeral Fever
-	C22.196.339	Freemartinism
-	C22.196.400	Hemorrhagic Syndrome, Bovine
-	C22.196.429	Infectious Bovine Rhinotracheitis
-	C22.196.497	Lumpy Skin Disease
-	C22.196.540	Malignant Catarrh
-	C22.196.581	Mastitis, Bovine
-	C22.196.831	Theileriosis
-	C22.196.888	Trypanosomiasis, Bovine
-	C22.196.927	Tuberculosis, Bovine
-	C22.196.963	White Heifer Disease
-	C22.214	Digital Dermatitis
-	C22.232	Disease Models, Animal
-	C22.268	Dog Diseases
-	C22.268.265	Distemper
-	C22.268.465	Hepatitis, Infectious Canine
-	C22.268.485	Hip Dysplasia, Canine
-	C22.313	Enterotoxemia
-	C22.331	Erysipelothrix Infections
-	C22.331.693	Swine Erysipelas
-	C22.362	Fish Diseases
-	C22.362.224	Furunculosis
-	C22.362.450	Hemorrhagic Septicemia, Viral
-	C22.380	Foot-and-Mouth Disease
-	C22.394	Foot Rot
-	C22.405	Goat Diseases
-	C22.434	Heartwater Disease
-	C22.467	Hepatitis, Animal
-	C22.467.435	Hepatitis, Viral, Animal
-	C22.467.435.442	Hepatitis, Infectious Canine
-	C22.467.435.812	Rift Valley Fever
-	C22.488	Horse Diseases
-	C22.488.088	African Horse Sickness
-	C22.488.304	Equine Infectious Anemia
-	C22.488.409	Glanders
-	C22.488.861	Strongyle Infections, Equine

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	C22.500	Keratoconjunctivitis, Infectious
-	C22.510	Lameness, Animal
-	C22.520	Mammary Neoplasms, Animal
-	C22.557	Mink Viral Enteritis
-	C22.595	Muscular Dystrophy, Animal
-	C22.595.740	White Muscle Disease
-	C22.627	Myxomatosis, Infectious
-	C22.674	Parasitic Diseases, Animal
-	C22.674.377	Helminthiasis, Animal
-	C22.674.377.315	Dictyocaulus Infections
-	C22.674.377.320	Dirofilariasis
-	C22.674.377.444	Fascioloidiasis
-	C22.674.377.656	Moniezia
-	C22.674.377.820	Setariasis
-	C22.674.377.840	Strongyle Infections, Equine
-	C22.674.377.868	Toxocariasis
-	C22.674.710	Protozoan Infections, Animal
-	C22.674.710.122	Babesiosis
-	C22.674.710.235	Cryptosporidiosis
-	C22.674.710.367	Dourine
-	C22.674.710.735	Theileriasis
-	C22.674.710.817	Toxoplasmosis, Animal
-	C22.674.710.896	Trypanosomiasis, Bovine
-	C22.688	Paratuberculosis
-	C22.695	Parturient Paresis
-	C22.706	Peste-des-Petits-Ruminants
-	C22.717	Pleuropneumonia, Contagious
-	C22.735	Primate Diseases
-	C22.735.050	Ape Diseases
-	C22.735.500	Monkey Diseases
-	C22.735.500.500	Marburg Virus Disease
-	C22.735.500.850	Simian Acquired Immunodeficiency Syndrome
-	C22.735.750	Monkeypox
-	C22.742	Pseudorabies
-	C22.761	Pythiosis
-	C22.780	Rinderpest



## MeSH Tree Changes for 2017

Type	Tree - heading
-	C22.795 Rodent Diseases
-	C22.795.239 Ectromelia, Infectious
-	C22.795.600 Monkeypox
-	C22.795.650 Murine Acquired Immunodeficiency Syndrome
-	C22.812 Salmonella Infections, Animal
-	C22.836 Sheep Diseases
-	C22.836.120 Bluetongue
-	C22.836.160 Border Disease
-	C22.836.259 Ecthyma, Contagious
-	C22.836.435 Louping Ill
-	C22.836.583 Nairobi Sheep Disease
-	C22.836.660 Pneumonia, Progressive Interstitial, of Sheep
-	C22.836.715 Pulmonary Adenomatosis, Ovine
-	C22.836.799 Scrapie
-	C22.836.886 Swayback
-	C22.836.900 Visna
-	C22.880 Steatitis
-	C22.905 Swine Diseases
-	C22.905.072 African Swine Fever
-	C22.905.170 Classical Swine Fever
-	C22.905.260 Edema Disease of Swine
-	C22.905.323 Encephalomyelitis, Enzootic Porcine
-	C22.905.382 Epidermitis, Exudative, of Swine
-	C22.905.469 Gastroenteritis, Transmissible, of Swine
-	C22.905.626 Pneumonia of Swine, Mycoplasmal
-	C22.905.690 Porcine Postweaning Multisystemic Wasting Syndrome
-	C22.905.700 Porcine Reproductive and Respiratory Syndrome
-	C22.905.832 Swine Erysipelas
-	C22.905.850 Swine Vesicular Disease
-	C22.905.927 Vesicular Exanthema of Swine
-	C22.950 Venereal Tumors, Veterinary
-	C22.952 Vesicular Stomatitis
-	C22.955 Wasting Disease, Chronic
-	C22.969 Zoonoses
-	C23 Pathological Conditions, Signs and Symptoms
-	C23.149 Morphological and Microscopic Findings

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	C23.149.249	Adenocarcinoma in Situ
-	C23.149.500	Atypical Squamous Cells of the Cervix
New Heading	<b>C23.149.625</b>	<b>Margins of Excision</b>
-	C23.149.750	Squamous Intraepithelial Lesions of the Cervix
-	C23.300	Pathological Conditions, Anatomical
-	C23.300.008	Agenesis of Corpus Callosum
-	C23.300.017	Airway Remodeling
-	C23.300.035	Alopecia
-	C23.300.035.500	Loose Anagen Hair Syndrome
-	C23.300.052	Atrial Remodeling
-	C23.300.070	Atrophy
-	C23.300.070.500	Muscular Atrophy
-	C23.300.070.500.500	Sarcopenia
-	C23.300.122	Blister
-	C23.300.175	Calculi
-	C23.300.175.350	Dental Calculus
-	C23.300.175.525	Gallstones
-	C23.300.175.700	Salivary Calculi
-	C23.300.175.700.325	Salivary Duct Calculi
-	C23.300.175.700.500	Salivary Gland Calculi
-	C23.300.175.850	Urinary Calculi
-	C23.300.175.850.550	Kidney Calculi
New Heading	<b>C23.300.175.850.550.500</b>	<b>Staghorn Calculi</b>
-	C23.300.175.850.750	Ureteral Calculi
-	C23.300.175.850.875	Urinary Bladder Calculi
-	C23.300.190	Accessory Atrioventricular Bundle
-	C23.300.250	Choristoma
-	C23.300.287	Constriction, Pathologic
-	C23.300.306	Cysts
-	C23.300.306.500	Parovarian Cyst
-	C23.300.306.750	Spermatocoele
-	C23.300.325	Dilatation, Pathologic
-	C23.300.415	Diverticulum
-	C23.300.415.500	Diverticulum, Colon

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C23.300.415.625                      Diverticulum, Esophageal
-	C23.300.415.625.900                      Zenker Diverticulum
-	C23.300.415.687                      Diverticulum, Stomach
-	C23.300.415.750                      Meckel Diverticulum
-	C23.300.505                      Facial Asymmetry
-	C23.300.575                      Fistula
-	C23.300.575.150                      Cutaneous Fistula
-	C23.300.575.185                      Digestive System Fistula
-	C23.300.575.185.150                      Biliary Fistula
-	C23.300.575.185.250                      Esophageal Fistula
-	C23.300.575.185.250.725                      Tracheoesophageal Fistula
-	C23.300.575.185.375                      Gastric Fistula
-	C23.300.575.185.550                      Intestinal Fistula
-	C23.300.575.185.550.600                      Rectal Fistula
-	C23.300.575.185.550.600.650                      Rectovaginal Fistula
-	C23.300.575.185.775                      Pancreatic Fistula
-	C23.300.575.500                      Oral Fistula
-	C23.300.575.500.275                      Dental Fistula
-	C23.300.575.500.550                      Oroantral Fistula
-	C23.300.575.500.775                      Salivary Gland Fistula
-	C23.300.575.687                      Respiratory Tract Fistula
-	C23.300.575.687.225                      Bronchial Fistula
-	C23.300.575.825                      Urinary Fistula
-	C23.300.575.825.250                      Urinary Bladder Fistula
-	C23.300.575.825.250.775                      Vesicovaginal Fistula
-	C23.300.575.925                      Vaginal Fistula
-	C23.300.575.925.558                      Rectovaginal Fistula
-	C23.300.575.925.816                      Vesicovaginal Fistula
-	C23.300.575.950                      Vascular Fistula
-	C23.300.575.950.150                      Arterio-Arterial Fistula
-	C23.300.575.950.150.500                      Bland White Garland Syndrome
-	C23.300.575.950.250                      Arteriovenous Fistula
-	C23.300.575.950.250.500                      Carotid-Cavernous Sinus Fistula
-	C23.300.707                      Hernia
-	C23.300.707.186                      Encephalocele
-	C23.300.707.374                      Hernia, Abdominal

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C23.300.707.374.500                      Gastroschisis
-	C23.300.707.374.750                      Hernia, Femoral
-	C23.300.707.374.875                      Hernia, Inguinal
-	C23.300.707.374.937                      Hernia, Ventral
-	C23.300.707.374.937.500                      Hernia, Umbilical
-	C23.300.707.500                            Hernia, Diaphragmatic
-	C23.300.707.500.116                      Hernias, Diaphragmatic, Congenital
-	C23.300.707.500.233                      Hernia, Diaphragmatic, Traumatic
-	C23.300.707.500.467                      Hernia, Hiatal
-	C23.300.707.937                           Hernia, Obturator
-	C23.300.707.945                           Incisional Hernia
-	C23.300.707.952                           Intervertebral Disc Displacement
-	C23.300.707.968                           Meningocele
-	C23.300.707.984                           Rectocele
-	C23.300.775                                Hypertrophy
-	C23.300.775.250                            Cardiomegaly
-	C23.300.775.250.400                      Hypertrophy, Left Ventricular
-	C23.300.775.250.401                      Hypertrophy, Right Ventricular
-	C23.300.775.525                           Hepatomegaly
-	C23.300.775.750                            Splenomegaly
-	C23.300.808                                Leg Length Inequality
-	C23.300.816                                Leukoplakia
-	C23.300.816.513                            Leukoplakia, Oral
-	C23.300.816.513.500                      Leukoplakia, Hairy
-	C23.300.820                                Nails, Malformed
-	C23.300.820.500                            Yellow Nail Syndrome
-	C23.300.821                                Plaque, Amyloid
-	C23.300.823                                Plaque, Atherosclerotic
-	C23.300.825                                Polyps
-	C23.300.825.411                            Intestinal Polyps
-	C23.300.825.411.235                      Colonic Polyps
-	C23.300.825.557                            Nasal Polyps
-	C23.300.842                                Prolapse
-	C23.300.842.624                            Pelvic Organ Prolapse
-	C23.300.842.624.249                      Cystocele
-	C23.300.842.624.500                      Rectal Prolapse

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C23.300.842.624.750 Uterine Prolapse
-	C23.300.842.624.875 Visceral Prolapse
-	C23.300.909 Rupture, Spontaneous
New Heading	<b>C23.300.940 Spontaneous Perforation</b>
New Heading	<b>C23.300.955 Tertiary Lymphoid Structures</b>
-	C23.300.970 Torsion Abnormality
-	C23.300.970.249 Bone Anteversion
-	C23.300.970.249.500 Coxa Vara
-	C23.300.970.374 Bone Retroversion
-	C23.300.970.374.500 Coxa Valga
-	C23.300.970.500 Intestinal Volvulus
-	C23.300.970.750 Uterine Retroversion
-	C23.300.977 Vascular Remodeling
-	C23.300.985 Ventricular Remodeling
-	C23.550 Pathologic Processes
-	C23.550.035 Acantholysis
-	C23.550.073 Arrhythmias, Cardiac
-	C23.550.073.093 Arrhythmia, Sinus
-	C23.550.073.093.249 Sick Sinus Syndrome
-	C23.550.073.093.500 Sinus Arrest, Cardiac
-	C23.550.073.198 Atrial Fibrillation
-	C23.550.073.248 Atrial Flutter
-	C23.550.073.300 Bradycardia
-	C23.550.073.325 Cardiac Complexes, Premature
-	C23.550.073.325.050 Atrial Premature Complexes
-	C23.550.073.325.800 Ventricular Premature Complexes
-	C23.550.073.425 Heart Block
-	C23.550.073.425.025 Adams-Stokes Syndrome
-	C23.550.073.425.062 Atrioventricular Block
-	C23.550.073.425.100 Bundle-Branch Block
-	C23.550.073.425.440 Sick Sinus Syndrome
-	C23.550.073.425.780 Sinoatrial Block
-	C23.550.073.547 Long QT Syndrome
-	C23.550.073.547.070 Andersen Syndrome

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C23.550.073.547.720 Romano-Ward Syndrome
-	C23.550.073.670 Parasystole
-	C23.550.073.845 Tachycardia
-	C23.550.073.845.695 Tachycardia, Paroxysmal
-	C23.550.073.845.787 Tachycardia, Reciprocating
-	C23.550.073.845.787.500 Tachycardia, Atrioventricular Nodal Reentry
-	C23.550.073.845.787.750 Tachycardia, Sinoatrial Nodal Reentry
-	C23.550.073.845.880 Tachycardia, Supraventricular
-	C23.550.073.845.880.315 Tachycardia, Ectopic Atrial
-	C23.550.073.845.880.320 Tachycardia, Ectopic Junctional
-	C23.550.073.845.880.845 Tachycardia, Sinus
-	C23.550.073.845.940 Tachycardia, Ventricular
-	C23.550.073.845.940.349 Accelerated Idioventricular Rhythm
-	C23.550.073.845.940.700 Torsades de Pointes
-	C23.550.073.922 Ventricular Fibrillation
-	C23.550.073.961 Ventricular Flutter
-	C23.550.081 Ascites
-	C23.550.113 Atrial Remodeling
-	C23.550.145 Azotemia
-	C23.550.161 Cardiotoxicity
-	C23.550.177 Channelopathies
-	C23.550.210 Chromosome Aberrations
-	C23.550.210.024 Abnormal Karyotype
-	C23.550.210.024.500 XYY Karyotype
-	C23.550.210.050 Aneuploidy
-	C23.550.210.050.500 Monosomy
-	C23.550.210.050.500.500 Chromosome Deletion
-	C23.550.210.050.625 Tetrasomy
-	C23.550.210.050.750 Trisomy
-	C23.550.210.110 Chromosomal Instability
-	C23.550.210.110.180 Chromosome Fragility
-	C23.550.210.170 Chromosome Breakage
-	C23.550.210.182 Chromosome Duplication
-	C23.550.210.182.249 Tetrasomy
-	C23.550.210.182.500 Trisomy
-	C23.550.210.190 Chromosome Inversion

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Heading	<b>C23.550.210.310</b>	<b>Chromothripsis</b>
-	C23.550.210.430	Isochromosomes
-	C23.550.210.570	Micronuclei, Chromosome-Defective
-	C23.550.210.645	Nondisjunction, Genetic
-	C23.550.210.645.890	Uniparental Disomy
-	C23.550.210.702	Polyploidy
-	C23.550.210.702.249	Tetraploidy
-	C23.550.210.702.500	Triploidy
-	C23.550.210.760	Ring Chromosomes
-	C23.550.210.815	Sex Chromosome Aberrations
-	C23.550.210.815.970	XYY Karyotype
-	C23.550.210.870	Translocation, Genetic
-	C23.550.210.870.680	Philadelphia Chromosome
-	C23.550.260	Death
-	C23.550.260.095	Asphyxia
New Heading	<b>C23.550.260.127</b>	<b>Body Remains</b>
-	C23.550.260.159	Brain Death
-	C23.550.260.224	Cadaver
-	C23.550.260.224.200	Corpse Dismemberment
-	C23.550.260.224.617	Postmortem Changes
-	C23.550.260.224.617.236	Autolysis
-	C23.550.260.224.617.839	Rigor Mortis
-	C23.550.260.322	Death, Sudden
-	C23.550.260.322.250	Death, Sudden, Cardiac
-	C23.550.260.322.250.500	Karoshi Death
-	C23.550.260.322.625	Sudden Infant Death
-	C23.550.260.393	Drowning
-	C23.550.260.440	Embryo Loss
-	C23.550.260.585	Fetal Death
-	C23.550.260.585.260	Fetal Resorption
-	C23.550.260.585.630	Stillbirth
-	C23.550.260.657	Infant Death
-	C23.550.260.657.500	Sudden Infant Death
-	C23.550.260.730	Parental Death

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C23.550.260.730.500 Maternal Death
-	C23.550.260.865 Perinatal Death
-	C23.550.274 Dehydration
-	C23.550.277 Delayed Graft Function
-	C23.550.288 Disease
-	C23.550.288.500 Syndrome
-	C23.550.291 Disease Attributes
-	C23.550.291.125 Acute Disease
-	C23.550.291.187 Asymptomatic Diseases
-	C23.550.291.187.500 Asymptomatic Infections
-	C23.550.291.250 Catastrophic Illness
-	C23.550.291.500 Chronic Disease
New Heading	<b>C23.550.291.500.500 Multiple Chronic Conditions</b>
-	C23.550.291.562 Convalescence
-	C23.550.291.625 Critical Illness
-	C23.550.291.656 Disease Progression
-	C23.550.291.656.700 Remission, Spontaneous
-	C23.550.291.671 Disease Resistance
-	C23.550.291.687 Disease Susceptibility
-	C23.550.291.687.500 Genetic Predisposition to Disease
-	C23.550.291.687.500.500 Anticipation, Genetic
-	C23.550.291.750 Diseases in Twins
-	C23.550.291.781 Emergencies
-	C23.550.291.812 Facies
-	C23.550.291.875 Iatrogenic Disease
-	C23.550.291.875.500 Cross Infection
-	C23.550.291.883 Late Onset Disorders
-	C23.550.291.890 Neglected Diseases
-	C23.550.291.906 Rare Diseases
-	C23.550.291.937 Recurrence
-	C23.550.291.937.500 Symptom Flare Up
-	C23.550.308 Dysbiosis
-	C23.550.325 Emphysema
-	C23.550.325.250 Mediastinal Emphysema
-	C23.550.325.500 Subcutaneous Emphysema



## MeSH Tree Changes for 2017

Type	Tree - heading
-	C23.550.325.500.500 alpha 1-Antitrypsin Deficiency
-	C23.550.340 Extravasation of Diagnostic and Therapeutic Materials
-	C23.550.347 Femoracetabular Impingement
-	C23.550.355 Fibrosis
-	C23.550.355.274 Cicatrix
-	C23.550.355.274.505 Cicatrix, Hypertrophic
-	C23.550.355.274.510 Keloid
-	C23.550.355.274.840 Tissue Adhesions
-	C23.550.355.550 Nephrogenic Fibrosing Dermopathy
-	C23.550.355.625 Peritoneal Fibrosis
-	C23.550.355.700 Retroperitoneal Fibrosis
-	C23.550.362 Genomic Instability
-	C23.550.362.180 Chromosomal Instability
-	C23.550.362.180.180 Chromosome Fragility
-	C23.550.362.590 Microsatellite Instability
-	C23.550.369 Gliosis
-	C23.550.382 Granuloma
-	C23.550.382.250 Eosinophilic Granuloma
-	C23.550.382.375 Granuloma Annulare
-	C23.550.382.437 Granuloma, Foreign-Body
-	C23.550.382.468 Granuloma, Giant Cell
-	C23.550.382.875 Granuloma, Plasma Cell
-	C23.550.382.937 Granuloma, Pyogenic
-	C23.550.382.968 Granuloma, Respiratory Tract
-	C23.550.382.968.500 Granuloma, Laryngeal
-	C23.550.382.984 Necrobiotic Xanthogranuloma
-	C23.550.384 Granulomatosis, Orofacial
-	C23.550.393 Growth Disorders
-	C23.550.393.450 Fetal Growth Retardation
-	C23.550.403 Hemolysis
-	C23.550.414 Hemorrhage
-	C23.550.414.300 Blood Loss, Surgical
-	C23.550.414.625 Ecchymosis
-	C23.550.414.712 Epistaxis
-	C23.550.414.734 Exsanguination
-	C23.550.414.756 Eye Hemorrhage

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C23.550.414.756.300 Choroid Hemorrhage
-	C23.550.414.756.550 Hyphema
-	C23.550.414.756.775 Retinal Hemorrhage
-	C23.550.414.756.887 Vitreous Hemorrhage
-	C23.550.414.788 Gastrointestinal Hemorrhage
-	C23.550.414.788.400 Hematemesis
-	C23.550.414.788.600 Melena
-	C23.550.414.788.700 Peptic Ulcer Hemorrhage
-	C23.550.414.794 Hemarthrosis
-	C23.550.414.817 Hematocele
-	C23.550.414.838 Hematoma
-	C23.550.414.838.349 Hematoma, Epidural, Cranial
-	C23.550.414.838.355 Hematoma, Epidural, Spinal
-	C23.550.414.838.700 Hematoma, Subdural
-	C23.550.414.838.700.100 Hematoma, Subdural, Acute
-	C23.550.414.838.700.200 Hematoma, Subdural, Chronic
-	C23.550.414.838.700.400 Hematoma, Subdural, Intracranial
-	C23.550.414.838.700.700 Hematoma, Subdural, Spinal
-	C23.550.414.849 Hematuria
-	C23.550.414.864 Hemobilia
-	C23.550.414.888 Hemoperitoneum
-	C23.550.414.896 Hemoptysis
-	C23.550.414.904 Hemothorax
-	C23.550.414.904.500 Hemopneumothorax
-	C23.550.414.913 Intracranial Hemorrhages
-	C23.550.414.913.100 Cerebral Hemorrhage
-	C23.550.414.913.100.200 Basal Ganglia Hemorrhage
-	C23.550.414.913.100.200.500 Putaminal Hemorrhage
-	C23.550.414.913.400 Hematoma, Epidural, Cranial
-	C23.550.414.913.700 Hematoma, Subdural
-	C23.550.414.913.700.100 Hematoma, Subdural, Acute
-	C23.550.414.913.700.200 Hematoma, Subdural, Chronic
-	C23.550.414.913.700.400 Hematoma, Subdural, Intracranial
-	C23.550.414.913.850 Subarachnoid Hemorrhage
-	C23.550.414.922 Oral Hemorrhage
-	C23.550.414.922.500 Gingival Hemorrhage

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C23.550.414.941 Postoperative Hemorrhage
-	C23.550.414.941.500 Endoleak
-	C23.550.414.950 Purpura
-	C23.550.414.950.230 Purpura Fulminans
-	C23.550.414.950.250 Purpura, Hyperglobulinemic
-	C23.550.414.950.375 Purpura, Schoenlein-Henoch
-	C23.550.414.950.687 Purpura, Thrombocytopenic
-	C23.550.414.950.687.600 Purpura, Thrombocytopenic, Idiopathic
-	C23.550.414.950.687.680 Purpura, Thrombotic Thrombocytopenic
-	C23.550.414.950.843 Waterhouse-Friderichsen Syndrome
-	C23.550.414.960 Retrobulbar Hemorrhage
-	C23.550.414.980 Shock, Hemorrhagic
-	C23.550.414.993 Uterine Hemorrhage
-	C23.550.414.993.700 Metrorrhagia
-	C23.550.414.993.850 Postpartum Hemorrhage
-	C23.550.421 Hyperammonemia
-	C23.550.425 Hyperamylasemia
-	C23.550.429 Hyperbilirubinemia
-	C23.550.429.249 Hyperbilirubinemia, Neonatal
-	C23.550.429.249.500 Jaundice, Neonatal
-	C23.550.429.500 Jaundice
-	C23.550.429.500.755 Jaundice, Obstructive
-	C23.550.429.750 Kernicterus
-	C23.550.444 Hyperplasia
-	C23.550.449 Hyperuricemia
-	C23.550.455 Hypovolemia
-	C23.550.470 Inflammation
-	C23.550.470.099 Acute-Phase Reaction
-	C23.550.470.251 Foreign-Body Reaction
-	C23.550.470.251.500 Implant Capsular Contracture
-	C23.550.470.448 Neurogenic Inflammation
-	C23.550.470.640 Seroma
-	C23.550.470.646 Serositis
-	C23.550.470.756 Suppuration
-	C23.550.470.756.100 Abscess
-	C23.550.470.756.200 Cellulitis

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C23.550.470.756.305                      Empyema
-	C23.550.470.756.305.500                      Empyema, Subdural
-	C23.550.470.790                      Systemic Inflammatory Response Syndrome
-	C23.550.470.790.500                      Sepsis
-	C23.550.470.790.500.100                      Bacteremia
-	C23.550.470.790.500.100.275                      Endotoxemia
-	C23.550.470.790.500.100.375                      Hemorrhagic Septicemia
-	C23.550.470.790.500.360                      Fungemia
-	C23.550.470.790.500.360.150                      Candidemia
New Heading	<b>C23.550.470.790.500.470                      Neonatal Sepsis</b>
-	C23.550.470.790.500.580                      Parasitemia
-	C23.550.470.790.500.800                      Shock, Septic
-	C23.550.470.790.500.900                      Viremia
-	C23.550.470.790.500.900.400                      Hemorrhagic Septicemia, Viral
-	C23.550.505                      Intraoperative Complications
-	C23.550.505.300                      Blood Loss, Surgical
-	C23.550.505.400                      Intraoperative Awareness
-	C23.550.505.700                      Malignant Hyperthermia
-	C23.550.513                      Ischemia
-	C23.550.513.355                      Infarction
-	C23.550.513.677                      No-Reflow Phenomenon
-	C23.550.522                      Leukoaraiosis
-	C23.550.526                      Leukocytosis
-	C23.550.537                      Lithiasis
-	C23.550.543                      Long Term Adverse Effects
-	C23.550.548                      Malacoplakia
-	C23.550.568                      Menstruation Disturbances
-	C23.550.568.500                      Amenorrhea
-	C23.550.568.750                      Dysmenorrhea
-	C23.550.568.875                      Menorrhagia
-	C23.550.568.937                      Oligomenorrhea
-	C23.550.568.968                      Premenstrual Syndrome
-	C23.550.568.968.500                      Premenstrual Dysphoric Disorder
-	C23.550.589                      Metaplasia
-	C23.550.589.500                      Neovascularization, Pathologic

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C23.550.589.500.145 Choroidal Neovascularization
-	C23.550.589.500.725 Retinal Neovascularization
-	C23.550.695 Muscle Weakness
-	C23.550.717 Necrosis
-	C23.550.717.182 Dental Pulp Necrosis
-	C23.550.717.273 DNA Degradation, Necrotic
-	C23.550.717.365 Fat Necrosis
-	C23.550.717.427 Gangrene
-	C23.550.717.489 Infarction
-	C23.550.717.732 Osteonecrosis
-	C23.550.717.732.183 Bisphosphonate-Associated Osteonecrosis of the Jaw
-	C23.550.717.732.368 Femur Head Necrosis
-	C23.550.722 Neointima
-	C23.550.727 Neoplastic Processes
-	C23.550.727.045 Anaplasia
-	C23.550.727.098 Carcinogenesis
-	C23.550.727.098.500 Cell Transformation, Neoplastic
-	C23.550.727.098.500.110 Blast Crisis
-	C23.550.727.098.500.160 Cell Transformation, Viral
-	C23.550.727.098.750 Cocarcinogenesis
-	C23.550.727.645 Neoplasm Invasiveness
-	C23.550.727.645.500 Leukemic Infiltration
-	C23.550.727.650 Neoplasm Metastasis
-	C23.550.727.650.560 Lymphatic Metastasis
-	C23.550.727.650.695 Neoplasm Micrometastasis
-	C23.550.727.650.830 Neoplasm Seeding
-	C23.550.727.650.895 Neoplasms, Unknown Primary
-	C23.550.727.650.900 Neoplastic Cells, Circulating
-	C23.550.727.655 Neoplasm Recurrence, Local
-	C23.550.727.670 Neoplasm Regression, Spontaneous
-	C23.550.727.700 Neoplasm, Residual
-	C23.550.737 Nerve Degeneration
-	C23.550.737.500 Retrograde Degeneration
-	C23.550.737.625 Subacute Combined Degeneration
-	C23.550.737.750 Wallerian Degeneration
-	C23.550.744 Ochronosis

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	C23.550.751	Ossification, Heterotopic
-	C23.550.751.500	Ossification of Posterior Longitudinal Ligament
New Tree	<b>C23.550.755</b>	<b>Pigmentation Disorders</b>
-	C23.550.759	Polydipsia
-	C23.550.759.500	Polydipsia, Psychogenic
-	C23.550.767	Postoperative Complications
-	C23.550.767.050	Afferent Loop Syndrome
-	C23.550.767.071	Anastomotic Leak
New Heading	<b>C23.550.767.082</b>	<b>Breast Cancer Lymphedema</b>
-	C23.550.767.093	Corneal Endothelial Cell Loss
-	C23.550.767.115	Coronary-Subclavian Steal Syndrome
-	C23.550.767.137	Delayed Emergence from Anesthesia
New Heading	<b>C23.550.767.181</b>	<b>Emergence Delirium</b>
-	C23.550.767.225	Failed Back Surgery Syndrome
-	C23.550.767.400	Graft Occlusion, Vascular
-	C23.550.767.500	Incisional Hernia
-	C23.550.767.600	Malignant Hyperthermia
-	C23.550.767.700	Pain, Postoperative
-	C23.550.767.700.500	Phantom Limb
-	C23.550.767.775	Postcholecystectomy Syndrome
-	C23.550.767.812	Postgastrectomy Syndromes
-	C23.550.767.812.500	Dumping Syndrome
-	C23.550.767.850	Postoperative Hemorrhage
-	C23.550.767.850.500	Endoleak
-	C23.550.767.859	Postoperative Nausea and Vomiting
-	C23.550.767.863	Postpericardiotomy Syndrome
-	C23.550.767.865	Prosthesis Failure
-	C23.550.767.865.500	Implant Capsular Contracture
-	C23.550.767.868	Prosthesis-Related Infections
-	C23.550.767.877	Reperfusion Injury
-	C23.550.767.877.500	Myocardial Reperfusion Injury
-	C23.550.767.877.750	Primary Graft Dysfunction
-	C23.550.767.879	Shock, Surgical
-	C23.550.767.882	Short Bowel Syndrome

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C23.550.767.884 Slit Ventricle Syndrome
-	C23.550.767.887 Surgical Wound Dehiscence
-	C23.550.767.925 Surgical Wound Infection
-	C23.550.767.962 Vasoplegia
-	C23.550.770 Protein Aggregation, Pathological
-	C23.550.773 Respiratory Aspiration
-	C23.550.773.500 Respiratory Aspiration of Gastric Contents
-	C23.550.794 Retropneumoperitoneum
-	C23.550.823 Sclerosis
-	C23.550.835 Shock
-	C23.550.835.525 Multiple Organ Failure
-	C23.550.835.550 Shock, Cardiogenic
-	C23.550.835.650 Shock, Hemorrhagic
-	C23.550.835.775 Shock, Surgical
-	C23.550.835.888 Shock, Traumatic
-	C23.550.835.900 Systemic Inflammatory Response Syndrome
-	C23.550.835.900.712 Shock, Septic
-	C23.550.863 Teratogenesis
-	C23.550.891 Ulcer
-	C23.550.918 Vascular Remodeling
-	C23.550.945 Yang Deficiency
-	C23.550.972 Yin Deficiency
-	C23.888 Signs and Symptoms
-	C23.888.069 Aging, Premature
-	C23.888.089 Asthenia
-	C23.888.119 Body Temperature Changes
-	C23.888.119.344 Fever
-	C23.888.119.344.345 Fever of Unknown Origin
-	C23.888.119.344.672 Sweating Sickness
-	C23.888.119.565 Hypothermia
-	C23.888.144 Body Weight
-	C23.888.144.186 Birth Weight
-	C23.888.144.186.500 Fetal Macrosomia
-	C23.888.144.243 Body Weight Changes
-	C23.888.144.243.926 Weight Gain
-	C23.888.144.243.963 Weight Loss

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C23.888.144.243.963.500 Emaciation
-	C23.888.144.243.963.500.500 Cachexia
-	C23.888.144.300 Fetal Weight
-	C23.888.144.699 Overweight
-	C23.888.144.699.500 Obesity
-	C23.888.144.699.500.250 Obesity, Metabolically Benign
-	C23.888.144.699.500.500 Obesity, Morbid
-	C23.888.144.699.500.750 Pediatric Obesity
-	C23.888.144.828 Thinness
-	C23.888.176 Cardiac Output, High
-	C23.888.192 Cardiac Output, Low
-	C23.888.208 Chills
-	C23.888.248 Cyanosis
-	C23.888.248.500 Infantile Apparent Life-Threatening Event
-	C23.888.277 Edema
-	C23.888.277.197 Edema, Cardiac
-	C23.888.277.395 Hydrops Fetalis
-	C23.888.307 Eye Manifestations
-	C23.888.307.500 Eye Hemorrhage
-	C23.888.307.625 Eye Pain
-	C23.888.307.750 Susac Syndrome
-	C23.888.338 Failure to Thrive
-	C23.888.369 Fatigue
-	C23.888.369.500 Mental Fatigue
New Heading	<b>C23.888.369.500.250 Alert Fatigue, Health Personnel</b>
-	C23.888.369.500.500 Compassion Fatigue
-	C23.888.378 Feminization
-	C23.888.380 Fetal Distress
-	C23.888.388 Flushing
-	C23.888.447 Heart Murmurs
-	C23.888.447.500 Systolic Murmurs
-	C23.888.475 Hot Flashes
-	C23.888.512 Hypergammaglobulinemia
-	C23.888.516 Hyperlactatemia
-	C23.888.521 Hypertriglyceridemic Waist



## MeSH Tree Changes for 2017

Type	Tree - heading	
-	C23.888.531	Intermittent Claudication
New Heading	<b>C23.888.541</b>	<b>Medically Unexplained Symptoms</b>
-	C23.888.550	Mobility Limitation
-	C23.888.571	Motion Sickness
-	C23.888.571.800	Space Motion Sickness
-	C23.888.582	Myocardial Stunning
-	C23.888.592	Neurologic Manifestations
New Heading	<b>C23.888.592.057</b>	<b>Bilateral Vestibulopathy</b>
-	C23.888.592.114	Cerebrospinal Fluid Leak
-	C23.888.592.114.249	Cerebrospinal Fluid Otorrhea
-	C23.888.592.114.624	Cerebrospinal Fluid Rhinorrhea
-	C23.888.592.298	Decerebrate State
-	C23.888.592.350	Dyskinesias
-	C23.888.592.350.090	Ataxia
-	C23.888.592.350.090.200	Cerebellar Ataxia
-	C23.888.592.350.090.600	Gait Ataxia
-	C23.888.592.350.110	Athetosis
-	C23.888.592.350.200	Catalepsy
-	C23.888.592.350.250	Chorea
New Tree	<a href="#">C23.888.592.350.275</a>	<a href="#">Dyskinesia, Drug-Induced</a>
New Heading	<b>C23.888.592.350.275.500</b>	<b>Tardive Dyskinesia</b>
-	C23.888.592.350.300	Dystonia
-	C23.888.592.350.300.800	Torticollis
-	C23.888.592.350.350	Hyperkinesia
-	C23.888.592.350.400	Hypokinesia
-	C23.888.592.350.500	Myoclonus
-	C23.888.592.350.600	Psychomotor Agitation
-	C23.888.592.350.600.500	Akathisia, Drug-Induced
-	C23.888.592.350.675	Synkinesia
-	C23.888.592.350.700	Tics
-	C23.888.592.350.850	Tremor
-	C23.888.592.413	Gait Disorders, Neurologic
-	C23.888.592.413.400	Gait Apraxia

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C23.888.592.413.450 Gait Ataxia
-	C23.888.592.544 Meningism
-	C23.888.592.604 Neurobehavioral Manifestations
-	C23.888.592.604.039 Anhedonia
-	C23.888.592.604.115 Catatonia
-	C23.888.592.604.150 Communication Disorders
-	C23.888.592.604.150.500 Language Disorders
-	C23.888.592.604.150.500.050 Agraphia
-	C23.888.592.604.150.500.090 Anomia
-	C23.888.592.604.150.500.300 Dyslexia
-	C23.888.592.604.150.500.300.200 Dyslexia, Acquired
-	C23.888.592.604.150.500.300.200.100 Alexia, Pure
-	C23.888.592.604.150.500.550 Language Development Disorders
-	C23.888.592.604.150.500.800 Speech Disorders
-	C23.888.592.604.150.500.800.100 Aphasia
-	C23.888.592.604.150.500.800.100.100 Aphasia, Broca
-	C23.888.592.604.150.500.800.100.111 Aphasia, Conduction
-	C23.888.592.604.150.500.800.100.155 Aphasia, Primary Progressive
-	C23.888.592.604.150.500.800.100.155.600 Aphasia Primary Progressive Nonfluent
-	C23.888.592.604.150.500.800.100.166 Aphasia, Wernicke
-	C23.888.592.604.150.500.800.150 Articulation Disorders
-	C23.888.592.604.150.500.800.150.200 Dysarthria
-	C23.888.592.604.150.500.800.300 Echolalia
-	C23.888.592.604.150.500.800.500 Mutism
-	C23.888.592.604.150.500.800.750 Stuttering
-	C23.888.592.604.150.550 Learning Disorders
-	C23.888.592.604.150.550.099 Dyscalculia
-	C23.888.592.604.150.550.200 Dyslexia
-	C23.888.592.604.150.550.200.500 Dyslexia, Acquired
-	C23.888.592.604.339 Confusion
-	C23.888.592.604.339.500 Delirium
New Heading	<b>C23.888.592.604.339.500.500 Emergence Delirium</b>
-	C23.888.592.604.359 Consciousness Disorders
-	C23.888.592.604.359.800 Unconsciousness

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C23.888.592.604.359.800.200 Coma
-	C23.888.592.604.359.800.400 Persistent Vegetative State
-	C23.888.592.604.359.800.500 Stupor
-	C23.888.592.604.359.800.600 Syncope
-	C23.888.592.604.359.800.600.500 Syncope, Vasovagal
-	C23.888.592.604.444 Lethargy
-	C23.888.592.604.529 Memory Disorders
-	C23.888.592.604.529.100 Amnesia
-	C23.888.592.604.529.100.075 Amnesia, Anterograde
-	C23.888.592.604.529.100.150 Amnesia, Retrograde
-	C23.888.592.604.529.100.800 Amnesia, Transient Global
-	C23.888.592.604.529.400 Korsakoff Syndrome
-	C23.888.592.604.646 Intellectual Disability
-	C23.888.592.604.764 Perceptual Disorders
-	C23.888.592.604.764.100 Agnosia
-	C23.888.592.604.764.100.300 Gerstmann Syndrome
-	C23.888.592.604.764.100.650 Prosopagnosia
-	C23.888.592.604.764.150 Alice in Wonderland Syndrome
-	C23.888.592.604.764.175 Allesthesia
-	C23.888.592.604.764.200 Auditory Perceptual Disorders
-	C23.888.592.604.764.300 Hallucinations
-	C23.888.592.604.764.400 Illusions
-	C23.888.592.604.764.700 Phantom Limb
-	C23.888.592.604.882 Psychomotor Disorders
-	C23.888.592.604.882.350 Apraxias
-	C23.888.592.604.882.350.099 Alien Hand Syndrome
-	C23.888.592.604.882.350.200 Apraxia, Ideomotor
-	C23.888.592.604.882.350.600 Gait Apraxia
-	C23.888.592.604.882.700 Psychomotor Agitation
-	C23.888.592.608 Neuromuscular Manifestations
-	C23.888.592.608.250 Fasciculation
-	C23.888.592.608.500 Muscle Cramp
-	C23.888.592.608.550 Muscle Hypertonia
-	C23.888.592.608.550.500 Muscle Rigidity
-	C23.888.592.608.550.550 Muscle Spasticity
-	C23.888.592.608.575 Muscle Hypotonia

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C23.888.592.608.593 Muscle Weakness
-	C23.888.592.608.612 Muscular Atrophy
-	C23.888.592.608.612.500 Sarcopenia
-	C23.888.592.608.650 Myokymia
-	C23.888.592.608.700 Myotonia
-	C23.888.592.608.750 Spasm
-	C23.888.592.608.750.400 Hemifacial Spasm
-	C23.888.592.608.750.700 Trismus
-	C23.888.592.608.875 Tetany
-	C23.888.592.610 Orthostatic Intolerance
-	C23.888.592.612 Pain
-	C23.888.592.612.054 Abdominal Pain
-	C23.888.592.612.054.200 Abdomen, Acute
-	C23.888.592.612.081 Acute Pain
-	C23.888.592.612.094 Arthralgia
-	C23.888.592.612.094.700 Shoulder Pain
-	C23.888.592.612.107 Back Pain
-	C23.888.592.612.107.200 Failed Back Surgery Syndrome
-	C23.888.592.612.107.400 Low Back Pain
-	C23.888.592.612.191 Breakthrough Pain
New Heading	<b>C23.888.592.612.212 Cancer Pain</b>
-	C23.888.592.612.233 Chest Pain
-	C23.888.592.612.233.500 Angina Pectoris
-	C23.888.592.612.233.500.150 Angina, Unstable
-	C23.888.592.612.233.500.150.150 Angina Pectoris, Variant
-	C23.888.592.612.233.500.575 Angina, Stable
-	C23.888.592.612.274 Chronic Pain
-	C23.888.592.612.302 Earache
-	C23.888.592.612.316 Eye Pain
-	C23.888.592.612.330 Facial Pain
-	C23.888.592.612.330.500 Toothache
-	C23.888.592.612.386 Flank Pain
-	C23.888.592.612.414 Glossalgia
-	C23.888.592.612.441 Headache
-	C23.888.592.612.441.500 Slit Ventricle Syndrome

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	C23.888.592.612.451	Labor Pain
-	C23.888.592.612.530	Mastodynia
-	C23.888.592.612.540	Metatarsalgia
New Heading	<b>C23.888.592.612.540.500</b>	<b>Morton Neuroma</b>
-	C23.888.592.612.547	Musculoskeletal Pain
-	C23.888.592.612.547.249	Myalgia
-	C23.888.592.612.547.500	Pelvic Girdle Pain
-	C23.888.592.612.553	Neck Pain
-	C23.888.592.612.664	Neuralgia
New Heading	<b>C23.888.592.612.664.275</b>	<b>Morton Neuroma</b>
-	C23.888.592.612.664.550	Neuralgia, Postherpetic
-	C23.888.592.612.664.675	Piriformis Muscle Syndrome
-	C23.888.592.612.664.737	Pudendal Neuralgia
-	C23.888.592.612.664.800	Sciatica
-	C23.888.592.612.720	Nociceptive Pain
-	C23.888.592.612.720.500	Visceral Pain
-	C23.888.592.612.776	Pain, Intractable
-	C23.888.592.612.832	Pain, Postoperative
-	C23.888.592.612.832.500	Phantom Limb
-	C23.888.592.612.888	Pain, Referred
-	C23.888.592.612.944	Pelvic Pain
-	C23.888.592.612.944.500	Dysmenorrhea
-	C23.888.592.612.944.625	Pelvic Girdle Pain
-	C23.888.592.612.944.750	Piriformis Muscle Syndrome
-	C23.888.592.612.972	Renal Colic
-	C23.888.592.636	Paralysis
-	C23.888.592.636.214	Facial Paralysis
-	C23.888.592.636.263	Gastroparesis
-	C23.888.592.636.312	Hemiplegia
-	C23.888.592.636.447	Ophthalmoplegia
-	C23.888.592.636.447.511	Ophthalmoplegia, Chronic Progressive External
-	C23.888.592.636.447.511.500	Kearns-Sayre Syndrome
-	C23.888.592.636.447.600	Ophthalmoplegic Migraine
-	C23.888.592.636.447.690	Supranuclear Palsy, Progressive

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C23.888.592.636.637 Paraplegia
-	C23.888.592.636.637.300 Brown-Sequard Syndrome
-	C23.888.592.636.711 Pseudobulbar Palsy
-	C23.888.592.636.786 Quadriplegia
-	C23.888.592.636.812 Respiratory Paralysis
-	C23.888.592.636.943 Vocal Cord Paralysis
-	C23.888.592.643 Paresis
-	C23.888.592.643.500 Paraparesis
-	C23.888.592.643.500.500 Paraparesis, Spastic
-	C23.888.592.700 Psychophysiologic Disorders
-	C23.888.592.708 Pupil Disorders
-	C23.888.592.708.150 Anisocoria
-	C23.888.592.708.362 Miosis
-	C23.888.592.708.362.500 Horner Syndrome
-	C23.888.592.708.575 Tonic Pupil
-	C23.888.592.717 Reflex, Abnormal
-	C23.888.592.742 Seizures
-	C23.888.592.763 Sensation Disorders
-	C23.888.592.763.237 Dizziness
-	C23.888.592.763.393 Hearing Disorders
-	C23.888.592.763.393.341 Hearing Loss
-	C23.888.592.763.393.341.186 Deafness
-	C23.888.592.763.393.341.186.500 Deaf-Blind Disorders
-	C23.888.592.763.393.341.374 Hearing Loss, Bilateral
-	C23.888.592.763.393.341.562 Hearing Loss, Conductive
-	C23.888.592.763.393.341.750 Hearing Loss, Functional
-	C23.888.592.763.393.341.812 Hearing Loss, High-Frequency
-	C23.888.592.763.393.341.849 Hearing Loss, Mixed Conductive-Sensorineural
-	C23.888.592.763.393.341.887 Hearing Loss, Sensorineural
-	C23.888.592.763.393.341.887.432 Hearing Loss, Central
-	C23.888.592.763.393.341.887.460 Hearing Loss, Noise-Induced
-	C23.888.592.763.393.341.887.772 Presbycusis
-	C23.888.592.763.393.341.887.886 Usher Syndromes
-	C23.888.592.763.393.341.900 Hearing Loss, Sudden
-	C23.888.592.763.393.341.950 Hearing Loss, Unilateral
-	C23.888.592.763.393.505 Hyperacusis

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C23.888.592.763.393.670 Tinnitus
-	C23.888.592.763.550 Olfaction Disorders
-	C23.888.592.763.770 Somatosensory Disorders
-	C23.888.592.763.770.400 Hyperalgesia
-	C23.888.592.763.770.450 Hyperesthesia
-	C23.888.592.763.770.500 Hypesthesia
-	C23.888.592.763.770.875 Paresthesia
-	C23.888.592.763.861 Taste Disorders
-	C23.888.592.763.861.184 Ageusia
-	C23.888.592.763.861.479 Dysgeusia
-	C23.888.592.763.941 Vision Disorders
-	C23.888.592.763.941.036 Alice in Wonderland Syndrome
-	C23.888.592.763.941.073 Amblyopia
-	C23.888.592.763.941.162 Blindness
-	C23.888.592.763.941.162.125 Amaurosis Fugax
-	C23.888.592.763.941.162.250 Blindness, Cortical
-	C23.888.592.763.941.162.625 Deaf-Blind Disorders
-	C23.888.592.763.941.256 Color Vision Defects
-	C23.888.592.763.941.339 Diplopia
-	C23.888.592.763.941.512 Hemianopsia
-	C23.888.592.763.941.661 Photophobia
-	C23.888.592.763.941.811 Scotoma
-	C23.888.592.763.941.848 Vision, Low
-	C23.888.592.796 Sleep Wake Disorders
-	C23.888.592.796.772 Sleep Deprivation
-	C23.888.592.848 Susac Syndrome
-	C23.888.592.900 Urinary Bladder, Neurogenic
-	C23.888.592.958 Vertigo
-	C23.888.592.958.500 Benign Paroxysmal Positional Vertigo
-	C23.888.592.979 Voice Disorders
-	C23.888.592.979.100 Aphonia
-	C23.888.592.979.325 Dysphonia
-	C23.888.592.979.550 Hoarseness
-	C23.888.619 Oral Manifestations
-	C23.888.619.500 Oral Hemorrhage
-	C23.888.663 Polydipsia

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C23.888.663.500 Polydipsia, Psychogenic
-	C23.888.672 Prodromal Symptoms
-	C23.888.681 Pseudophakia
-	C23.888.721 Renal Colic
-	C23.888.760 Reticulocytosis
-	C23.888.821 Signs and Symptoms, Digestive
-	C23.888.821.030 Abdominal Pain
-	C23.888.821.030.249 Abdomen, Acute
-	C23.888.821.061 Aerophagy
-	C23.888.821.108 Anorexia
-	C23.888.821.150 Constipation
-	C23.888.821.214 Diarrhea
-	C23.888.821.214.500 Diarrhea, Infantile
-	C23.888.821.236 Dyspepsia
-	C23.888.821.266 Encopresis
-	C23.888.821.297 Eructation
-	C23.888.821.360 Flatulence
-	C23.888.821.414 Gagging
-	C23.888.821.475 Halitosis
-	C23.888.821.525 Heartburn
-	C23.888.821.578 Hiccup
-	C23.888.821.645 Hyperphagia
-	C23.888.821.645.500 Bulimia
-	C23.888.821.712 Nausea
-	C23.888.821.712.700 Postoperative Nausea and Vomiting
New Tree	<a href="#">C23.888.821.825</a> <a href="#">Pica</a>
-	C23.888.821.937 Vomiting
-	C23.888.821.937.019 Hematemesis
-	C23.888.821.937.049 Morning Sickness
-	C23.888.821.937.049.500 Hyperemesis Gravidarum
-	C23.888.821.937.059 Postoperative Nausea and Vomiting
-	C23.888.821.937.080 Vomiting, Anticipatory
-	C23.888.852 Signs and Symptoms, Respiratory
-	C23.888.852.079 Anoxia
-	C23.888.852.079 Hypoxia



## MeSH Tree Changes for 2017

Type	Tree - heading	
-	C23.888.852.079.594	Fetal Hypoxia
-	C23.888.852.130	Apnea
-	C23.888.852.227	Cheyne-Stokes Respiration
-	C23.888.852.293	Cough
-	C23.888.852.371	Dyspnea
-	C23.888.852.371.396	Dyspnea, Paroxysmal
-	C23.888.852.430	Hemoptysis
-	C23.888.852.490	Hoarseness
-	C23.888.852.544	Hypercapnia
-	C23.888.852.567	Hyperoxia
-	C23.888.852.591	Hyperventilation
-	C23.888.852.614	Hypocapnia
-	C23.888.852.638	Hypoventilation
-	C23.888.852.700	Infantile Apparent Life-Threatening Event
-	C23.888.852.761	Mouth Breathing
-	C23.888.852.779	Respiratory Sounds
-	C23.888.852.779.850	Snoring
-	C23.888.852.889	Sneezing
-	C23.888.852.944	Tachypnea
-	C23.888.852.944.500	Transient Tachypnea of the Newborn
-	C23.888.885	Skin Manifestations
-	C23.888.885.250	Cafe-au-Lait Spots
New Heading	<b>C23.888.885.281</b>	<b>Cellulite</b>
-	C23.888.885.312	Ecchymosis
-	C23.888.885.375	Jaundice
-	C23.888.885.375.500	Jaundice, Obstructive
-	C23.888.885.437	Livedo Reticularis
-	C23.888.885.468	Necrolytic Migratory Erythema
-	C23.888.885.500	Pallor
-	C23.888.885.625	Pruritus
-	C23.888.885.625.500	Tinea Pedis
-	C23.888.885.687	Purpura
-	C23.888.885.687.230	Purpura Fulminans
-	C23.888.885.687.250	Purpura, Hyperglobulinemic
-	C23.888.885.687.375	Purpura, Schoenlein-Henoch

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C23.888.885.687.687                      Purpura, Thrombocytopenic
-	C23.888.885.687.687.600                      Purpura, Thrombocytopenic, Idiopathic
-	C23.888.885.687.687.680                      Purpura, Thrombotic Thrombocytopenic
-	C23.888.885.687.843                      Waterhouse-Friderichsen Syndrome
-	C23.888.885.800                      Striae Distensae
-	C23.888.942                      Urological Manifestations
-	C23.888.942.337                      Hypercalciuria
-	C23.888.942.343                      Lower Urinary Tract Symptoms
-	C23.888.942.343.274                      Dysuria
-	C23.888.942.343.550                      Nocturia
-	C23.888.942.343.600                      Prostatism
-	C23.888.942.343.780                      Urinary Bladder, Overactive
-	C23.888.942.343.800                      Urinary Incontinence
-	C23.888.942.343.800.500                      Urinary Incontinence, Stress
-	C23.888.942.343.800.750                      Urinary Incontinence, Urge
-	C23.888.942.400                      Oliguria
-	C23.888.942.600                      Polyuria
-	C23.888.942.750                      Proteinuria
-	C23.888.942.750.269                      Albuminuria
-	C23.888.942.750.634                      Hemoglobinuria
-	C23.888.942.900                      Urinoma
-	C23.888.971                      Virilism
-	C23.888.971.468                      Hirsutism
-	C24                      Occupational Diseases
-	C24.080                      Agricultural Workers' Diseases
-	C24.080.365                      Farmer's Lung
-	C24.080.752                      Silo Filler's Disease
-	C24.102                      Asthma, Occupational
-	C24.125                      Bird Fancier's Lung
-	C24.198                      Burnout, Professional
-	C24.270                      Dermatitis, Occupational
-	C24.400                      Hand-Arm Vibration Syndrome
-	C24.410                      High Pressure Neurological Syndrome
-	C24.426                      Inert Gas Narcosis
-	C24.506                      Laboratory Infection
-	C24.653                      Persian Gulf Syndrome

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	C24.800	Pneumoconiosis
-	C24.800.127	Asbestosis
-	C24.800.225	Berylliosis
-	C24.800.323	Byssinosis
-	C24.800.340	Caplan Syndrome
-	C24.800.773	Siderosis
-	C24.800.834	Silicosis
-	C24.800.834.201	Anthracosilicosis
-	C24.800.834.752	Silicotuberculosis
-	C24.900	Sleep Disorders, Circadian Rhythm
-	C25	Chemically-Induced Disorders
-	C25.100	Drug-Related Side Effects and Adverse Reactions
-	C25.100.249	Akathisia, Drug-Induced
-	C25.100.311	Anticholinergic Syndrome
-	C25.100.389	Cardiotoxicity
-	C25.100.468	Drug Hypersensitivity
-	C25.100.468.189	Asthma, Aspirin-Induced
-	C25.100.468.380	Drug Eruptions
-	C25.100.468.380.174	Acute Generalized Exanthematous Pustulosis
-	C25.100.468.380.262	Drug Hypersensitivity Syndrome
-	C25.100.468.380.375	Erythema Nodosum
-	C25.100.468.380.587	Hand-Foot Syndrome
-	C25.100.468.380.693	Nicolau Syndrome
-	C25.100.468.380.800	Serum Sickness
-	C25.100.468.380.900	Stevens-Johnson Syndrome
-	C25.100.562	Chemical and Drug Induced Liver Injury
-	C25.100.562	Drug-Induced Liver Injury
-	C25.100.562.200	Drug-Induced Liver Injury, Chronic
-	C25.100.750	Dyskinesia, Drug-Induced
-	C25.100.812	Metabolic Side Effects of Drugs and Substances
New Heading	<b>C25.100.844</b>	<b>Propofol Infusion Syndrome</b>
-	C25.100.875	Serotonin Syndrome
-	C25.723	Poisoning
-	C25.723.068	Argyria
-	C25.723.097	Arsenic Poisoning

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C25.723.127 Bites and Stings
-	C25.723.127.071 Insect Bites and Stings
-	C25.723.127.142 Scorpion Stings
-	C25.723.127.442 Snake Bites
-	C25.723.127.723 Spider Bites
-	C25.723.127.789 Tick Bites
-	C25.723.127.789.500 Tick Toxicoses
-	C25.723.127.789.500.707 Tick Paralysis
-	C25.723.165 Cadmium Poisoning
-	C25.723.177 Carbon Tetrachloride Poisoning
-	C25.723.260 Chemical and Drug Induced Liver Injury
-	C25.723.260 Drug-Induced Liver Injury
-	C25.723.260.200 Drug-Induced Liver Injury, Chronic
-	C25.723.380 Fluoride Poisoning
-	C25.723.415 Foodborne Diseases
-	C25.723.415.151 Botulism
-	C25.723.415.246 Ciguatera Poisoning
-	C25.723.415.341 Favism
-	C25.723.415.551 Mushroom Poisoning
-	C25.723.415.738 Salmonella Food Poisoning
-	C25.723.415.792 Shellfish Poisoning
-	C25.723.415.846 Staphylococcal Food Poisoning
-	C25.723.455 Gas Poisoning
-	C25.723.455.245 Carbon Monoxide Poisoning
-	C25.723.455.571 Inert Gas Narcosis
-	C25.723.589 Lead Poisoning
-	C25.723.589.500 Lead Poisoning, Nervous System
-	C25.723.589.500.400 Lead Poisoning, Nervous System, Adult
-	C25.723.589.500.700 Lead Poisoning, Nervous System, Childhood
-	C25.723.618 Manganese Poisoning
-	C25.723.647 Mercury Poisoning
-	C25.723.647.500 Mercury Poisoning, Nervous System
-	C25.723.647.500.100 Acrodynia
-	C25.723.680 Mycotoxicosis
-	C25.723.680.262 Ergotism
-	C25.723.680.551 Mushroom Poisoning

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C25.723.705                      Neurotoxicity Syndromes
-	C25.723.705.100                  Akathisia, Drug-Induced
-	C25.723.705.150                  Alcohol-Induced Disorders, Nervous System
-	C25.723.705.150.100              Alcohol Amnestic Disorder
-	C25.723.705.150.100.500          Korsakoff Syndrome
-	C25.723.705.150.200              Alcohol Withdrawal Delirium
-	C25.723.705.150.300              Alcohol Withdrawal Seizures
-	C25.723.705.150.400              Alcoholic Neuropathy
-	C25.723.705.200                  Dyskinesia, Drug-Induced
-	C25.723.705.400                  MPTP Poisoning
-	C25.723.705.600                  Neuroleptic Malignant Syndrome
-	C25.723.717                      Organophosphate Poisoning
-	C25.723.756                      Plant Poisoning
-	C25.723.756.375                      Favism
-	C25.723.756.558                      Lathyrism
-	C25.723.756.600                      Milk Sickness
-	C25.723.809                      Psychoses, Substance-Induced
-	C25.723.809.750                      Psychoses, Alcoholic
-	C25.723.932                      Water Intoxication
-	C25.775                          Substance-Related Disorders
-	C25.775.100                      Alcohol-Related Disorders
-	C25.775.100.087                      Alcohol-Induced Disorders
-	C25.775.100.087.193                  Alcohol-Induced Disorders, Nervous System
-	C25.775.100.087.193.100              Alcohol Amnestic Disorder
-	C25.775.100.087.193.100.500          Korsakoff Syndrome
-	C25.775.100.087.193.200              Alcohol Withdrawal Delirium
-	C25.775.100.087.193.300              Alcohol Withdrawal Seizures
-	C25.775.100.087.193.400              Alcoholic Neuropathy
-	C25.775.100.087.250                  Cardiomyopathy, Alcoholic
-	C25.775.100.087.323                  Fetal Alcohol Spectrum Disorders
-	C25.775.100.087.645                  Liver Diseases, Alcoholic
-	C25.775.100.087.645.390              Fatty Liver, Alcoholic
-	C25.775.100.087.645.490              Hepatitis, Alcoholic
-	C25.775.100.087.645.550              Liver Cirrhosis, Alcoholic
-	C25.775.100.087.730                  Pancreatitis, Alcoholic
-	C25.775.100.087.750                  Psychoses, Alcoholic

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	C25.775.100.175	Alcoholic Intoxication
-	C25.775.100.250	Alcoholism
-	C25.775.100.437	Binge Drinking
-	C25.775.100.625	Wernicke Encephalopathy
-	C25.775.225	Amphetamine-Related Disorders
-	C25.775.300	Cocaine-Related Disorders
-	C25.775.383	Drug Overdose
-	C25.775.467	Inhalant Abuse
-	C25.775.635	Marijuana Abuse
-	C25.775.650	Neonatal Abstinence Syndrome
-	C25.775.675	Opioid-Related Disorders
-	C25.775.675.400	Heroin Dependence
-	C25.775.675.600	Morphine Dependence
-	C25.775.700	Phencyclidine Abuse
-	C25.775.746	Psychoses, Substance-Induced
-	C25.775.793	Substance Abuse, Intravenous
-	C25.775.835	Substance Withdrawal Syndrome
-	C25.775.835.250	Alcohol Withdrawal Delirium
-	C25.775.835.500	Alcohol Withdrawal Seizures
-	C25.775.912	Tobacco Use Disorder
-	C26	Wounds and Injuries
-	C26.017	Abdominal Injuries
-	C26.017.258	Hernia, Diaphragmatic, Traumatic
-	C26.017.680	Splenic Rupture
-	C26.017.680.500	Splenosis
-	C26.017.809	Stomach Rupture
-	C26.062	Amputation, Traumatic
-	C26.088	Arm Injuries
-	C26.088.268	Forearm Injuries
-	C26.088.268.556	Radius Fractures
-	C26.088.268.807	Ulna Fractures
-	C26.088.390	Humeral Fractures
Old Tree	<b>C26.088.666</b>	<b>Shoulder Dislocation</b>
Old Tree	<b>C26.088.749</b>	<b>Shoulder Fractures</b>
-	C26.088.890	Tennis Elbow
-	C26.088.906	Wrist Injuries

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	C26.103	Asphyxia
-	C26.115	Athletic Injuries
-	C26.117	Back Injuries
-	C26.117.500	Spinal Injuries
-	C26.117.500.500	Spinal Fractures
-	C26.120	Barotrauma
-	C26.120.126	Blast Injuries
-	C26.120.248	Decompression Sickness
-	C26.130	Battered Child Syndrome
-	C26.141	Birth Injuries
-	C26.141.587	Paralysis, Obstetric
-	C26.176	Bites and Stings
-	C26.176.071	Bites, Human
-	C26.176.143	Insect Bites and Stings
-	C26.176.443	Scorpion Stings
-	C26.176.724	Snake Bites
-	C26.176.790	Spider Bites
-	C26.176.793	Tick Bites
-	C26.200	Burns
-	C26.200.156	Burns, Chemical
-	C26.200.239	Burns, Electric
-	C26.200.322	Burns, Inhalation
-	C26.200.322.800	Smoke Inhalation Injury
-	C26.200.503	Eye Burns
-	C26.200.855	Sunburn
-	C26.212	Cold Injury
-	C26.212.500	Frostbite
-	C26.212.500.217	Chilblains
-	C26.224	Contrecoup Injury
New Heading	<b>C26.257</b>	<b>Crush Injuries</b>
New Tree	<b>C26.257.500</b>	<b>Crush Syndrome</b>
-	C26.289	Dislocations
-	C26.289	Joint Dislocations
New Heading	<b>C26.289.192</b>	<b>Diastasis, Bone</b>

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	<a href="#">C26.289.192.500</a>	<a href="#">Pubic Symphysis Diastasis</a>
New Heading	<b>C26.289.288</b>	<b>Diastasis, Muscle</b>
New Heading	<b>C26.289.336</b>	<b>Fracture Dislocation</b>
New Tree	<a href="#">C26.289.336.500</a>	<a href="#">Colles' Fracture</a>
New Tree	<a href="#">C26.289.336.750</a>	<a href="#">Monteggia's Fracture</a>
New Heading	<b>C26.289.336.875</b>	<b>Salter-Harris Fractures</b>
-	C26.289.384	Hip Dislocation
-	C26.289.500	Knee Dislocation
-	C26.289.625	Patellar Dislocation
Old Tree	<b>C26.289.650</b>	<b>Pubic Symphysis Diastasis</b>
-	C26.289.750	Shoulder Dislocation
-	C26.304	Drowning
-	C26.304.500	Near Drowning
-	C26.324	Electric Injuries
-	C26.324.323	Burns, Electric
-	C26.324.436	Conducted Energy Weapon Injuries
-	C26.324.550	Lightning Injuries
-	C26.348	Esophageal Perforation
-	C26.371	Extravasation of Diagnostic and Therapeutic Materials
-	C26.392	Foreign Bodies
-	C26.392.183	Bezoars
-	C26.392.421	Eye Foreign Bodies
-	C26.392.500	Foreign-Body Migration
-	C26.392.500.249	Artificial Lens Implant Migration
-	C26.392.500.500	Intrauterine Device Migration
-	C26.392.560	Foreign-Body Reaction
-	C26.392.560.325	Granuloma, Foreign-Body
-	C26.404	Fractures, Bone
-	C26.404.014	Ankle Fractures
New Heading	<b>C26.404.026</b>	<b>Fracture Dislocation</b>
New Heading	<b>C26.404.026.500</b>	<b>Salter-Harris Fractures</b>



## MeSH Tree Changes for 2017

Type	Tree - heading	
New Heading	<b>C26.404.038</b>	<b>Fractures, Avulsion</b>
-	C26.404.061	Femoral Fractures
-	C26.404.061.425	Hip Fractures
-	C26.404.061.425.500	Femoral Neck Fractures
-	C26.404.124	Fractures, Closed
-	C26.404.186	Fractures, Comminuted
-	C26.404.195	Fractures, Compression
-	C26.404.249	Fractures, Malunited
-	C26.404.280	Fractures, Multiple
-	C26.404.311	Fractures, Open
-	C26.404.374	Fractures, Spontaneous
-	C26.404.437	Fractures, Stress
-	C26.404.468	Fractures, Ununited
-	C26.404.468.627	Pseudarthrosis
-	C26.404.500	Humeral Fractures
-	C26.404.505	Intra-Articular Fractures
-	C26.404.545	Osteoporotic Fractures
-	C26.404.550	Periprosthetic Fractures
-	C26.404.562	Radius Fractures
-	C26.404.562.356	Colles' Fracture
-	C26.404.593	Rib Fractures
-	C26.404.625	Shoulder Fractures
New Heading	<b>C26.404.625.500</b>	<b>Bankart Lesions</b>
-	C26.404.750	Skull Fractures
-	C26.404.750.467	Jaw Fractures
-	C26.404.750.467.441	Mandibular Fractures
-	C26.404.750.467.611	Maxillary Fractures
-	C26.404.750.684	Orbital Fractures
-	C26.404.750.821	Skull Fracture, Basilar
-	C26.404.750.890	Skull Fracture, Depressed
-	C26.404.750.959	Zygomatic Fractures
-	C26.404.812	Spinal Fractures
-	C26.404.875	Tibial Fractures
-	C26.404.937	Ulna Fractures

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C26.404.937.547 Monteggia's Fracture
-	C26.411 Fractures, Cartilage
-	C26.417 Frostbite
-	C26.417.217 Chilblains
-	C26.448 Hand Injuries
-	C26.448.429 Finger Injuries
-	C26.522 Heat Stress Disorders
-	C26.522.250 Heat Exhaustion
-	C26.522.500 Heat Stroke
-	C26.522.500.500 Sunstroke
-	C26.531 Hip Injuries
-	C26.531.500 Hip Dislocation
-	C26.531.750 Hip Fractures
-	C26.531.750.500 Femoral Neck Fractures
-	C26.540 Lacerations
-	C26.558 Leg Injuries
-	C26.558.100 Ankle Injuries
-	C26.558.276 Femoral Fractures
-	C26.558.276.425 Hip Fractures
-	C26.558.276.425.500 Femoral Neck Fractures
-	C26.558.300 Foot Injuries
New Tree	<a href="#">C26.558.300.500</a> <a href="#">Hallux Limitus</a>
-	C26.558.554 Knee Injuries
New Heading	<b>C26.558.554.213</b> <b>Anterior Cruciate Ligament Injuries</b>
-	C26.558.554.425 Iliotibial Band Syndrome
-	C26.558.554.500 Knee Dislocation
-	C26.558.554.750 Patellar Dislocation
-	C26.558.705 Medial Tibial Stress Syndrome
New Heading	<b>C26.558.781</b> <b>Tibial Meniscus Injuries</b>
-	C26.558.857 Tibial Fractures
New Heading	<b>C26.599</b> <b>Microtrauma, Physical</b>
-	C26.640 Multiple Trauma
-	C26.640.500 Fractures, Multiple

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	C26.670	Nasal Septal Perforation
-	C26.700	Neck Injuries
-	C26.700.500	Whiplash Injuries
-	C26.716	Occupational Injuries
-	C26.733	Radiation Injuries
-	C26.733.031	Abnormalities, Radiation-Induced
-	C26.733.188	Acute Radiation Syndrome
-	C26.733.266	Cardiotoxicity
-	C26.733.345	Leukemia, Radiation-Induced
-	C26.733.476	Neoplasms, Radiation-Induced
-	C26.733.579	Osteoradionecrosis
-	C26.733.720	Radiation Injuries, Experimental
-	C26.733.762	Radiation Pneumonitis
-	C26.733.804	Radiodermatitis
-	C26.748	Retropneumoperitoneum
-	C26.761	Rupture
-	C26.761.125	Aortic Rupture
New Heading	<b>C26.761.340</b>	<b>Rotator Cuff Injuries</b>
-	C26.761.555	Splenic Rupture
-	C26.761.555.500	Splenosis
-	C26.761.684	Stomach Rupture
-	C26.761.853	Uterine Rupture
-	C26.761.853.500	Uterine Perforation
-	C26.780	Self Mutilation
-	C26.797	Shock, Traumatic
-	C26.797.240	Crush Syndrome
New Heading	<b>C26.803</b>	<b>Shoulder Injuries</b>
New Heading	<b>C26.803.063</b>	<b>Rotator Cuff Injuries</b>
New Tree	<a href="#">C26.803.125</a>	<a href="#">Shoulder Dislocation</a>
New Tree	<a href="#">C26.803.250</a>	<a href="#">Shoulder Fractures</a>
New Heading	<b>C26.803.250.500</b>	<b>Bankart Lesions</b>
New	<a href="#">C26.803.500</a>	<a href="#">Shoulder Impingement Syndrome</a>

## MeSH Tree Changes for 2017

Type	Tree - heading	
Tree		
-	C26.808	Soft Tissue Injuries
New Heading	<b>C26.808.500</b>	<b>Degloving Injuries</b>
-	C26.819	Spinal Cord Injuries
-	C26.819.339	Central Cord Syndrome
-	C26.819.678	Spinal Cord Compression
-	C26.844	Sprains and Strains
-	C26.844.150	Cumulative Trauma Disorders
-	C26.844.150.206	Carpal Tunnel Syndrome
-	C26.844.150.425	Iliotibial Band Syndrome
-	C26.844.150.957	Ulnar Nerve Compression Syndromes
-	C26.844.150.957.200	Cubital Tunnel Syndrome
New Heading	<b>C26.859</b>	<b>Surgical Wound</b>
-	C26.874	Tendon Injuries
New Heading	<b>C26.874.400</b>	<b>Rotator Cuff Injuries</b>
-	C26.874.800	Tendinopathy
New Heading	<b>C26.874.800.500</b>	<b>Elbow Tendinopathy</b>
New Tree	<a href="#">C26.874.800.500.500</a>	<a href="#">Tennis Elbow</a>
New Heading	<b>C26.874.800.750</b>	<b>Enthesopathy</b>
-	C26.891	Thoracic Injuries
-	C26.891.315	Flail Chest
-	C26.891.375	Heart Injuries
New Heading	<b>C26.891.375.750</b>	<b>Myocardial Contusions</b>
New Tree	<a href="#">C26.891.375.750.500</a>	<a href="#">Commotio Cordis</a>
-	C26.891.554	Lung Injury
-	C26.891.733	Rib Fractures
-	C26.900	Tooth Injuries
-	C26.900.725	Tooth Avulsion
-	C26.900.750	Tooth Fractures
-	C26.900.750.300	Cracked Tooth Syndrome
-	C26.915	Trauma, Nervous System

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	C26.915.200	Cerebrovascular Trauma
-	C26.915.200.200	Carotid Artery Injuries
-	C26.915.200.200.500	Carotid Artery, Internal, Dissection
-	C26.915.200.200.550	Carotid-Cavernous Sinus Fistula
-	C26.915.200.600	Vertebral Artery Dissection
-	C26.915.300	Craniocerebral Trauma
-	C26.915.300.200	Brain Injuries
Old Tree	<del>C26.915.300.200.150</del>	<del>Brain Concussion</del>
Old Tree	<del>C26.915.300.200.150.249</del>	<del>Contrecoup Injury</del>
Old Tree	<del>C26.915.300.200.150.500</del>	<del>Post-Concussion Syndrome</del>
-	C26.915.300.200.175	Brain Hemorrhage, Traumatic
-	C26.915.300.200.175.200	Brain Stem Hemorrhage, Traumatic
-	C26.915.300.200.175.300	Cerebral Hemorrhage, Traumatic
New Heading	<b>C26.915.300.200.188</b>	<b>Brain Injuries, Diffuse</b>
New Tree	C26.915.300.200.188.500	Diffuse Axonal Injury
New Heading	<b>C26.915.300.200.194</b>	<b>Brain Injuries, Traumatic</b>
New Tree	C26.915.300.200.194.250	Brain Concussion
New Heading	<b>C26.915.300.200.194.375</b>	<b>Brain Contusion</b>
New Heading	<b>C26.915.300.200.194.500</b>	<b>Chronic Traumatic Encephalopathy</b>
-	C26.915.300.200.200	Brain Injury, Chronic
New Heading	<b>C26.915.300.200.200.500</b>	<b>Chronic Traumatic Encephalopathy</b>
Old Tree	<del>C26.915.300.200.300</del>	<del>Diffuse Axonal Injury</del>
-	C26.915.300.200.475	Epilepsy, Post-Traumatic
-	C26.915.300.200.650	Pneumocephalus
-	C26.915.300.200.825	Shaken Baby Syndrome
-	C26.915.300.225	Cerebrospinal Fluid Leak
-	C26.915.300.225.500	Cerebrospinal Fluid Otorrhea
-	C26.915.300.225.750	Cerebrospinal Fluid Rhinorrhea
-	C26.915.300.350	Coma, Post-Head Injury
-	C26.915.300.400	Cranial Nerve Injuries
-	C26.915.300.400.100	Abducens Nerve Injury

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C26.915.300.400.200      Accessory Nerve Injuries
-	C26.915.300.400.300      Facial Nerve Injuries
-	C26.915.300.400.343      Glossopharyngeal Nerve Injuries
-	C26.915.300.400.387      Hypoglossal Nerve Injuries
-	C26.915.300.400.518      Oculomotor Nerve Injuries
-	C26.915.300.400.562      Olfactory Nerve Injuries
-	C26.915.300.400.650      Optic Nerve Injuries
-	C26.915.300.400.825      Trigeminal Nerve Injuries
-	C26.915.300.400.825.500      Lingual Nerve Injuries
-	C26.915.300.400.868      Trochlear Nerve Injuries
-	C26.915.300.400.912      Vagus Nerve Injuries
-	C26.915.300.400.912.750      Laryngeal Nerve Injuries
-	C26.915.300.400.912.750.500      Recurrent Laryngeal Nerve Injuries
-	C26.915.300.400.956      Vestibulocochlear Nerve Injuries
-	C26.915.300.425      Facial Injuries
-	C26.915.300.425.250      Eye Injuries
-	C26.915.300.425.250.124      Corneal Injuries
-	C26.915.300.425.250.124.500      Corneal Perforation
-	C26.915.300.425.250.250      Eye Burns
-	C26.915.300.425.250.260      Eye Foreign Bodies
-	C26.915.300.425.250.270      Eye Injuries, Penetrating
-	C26.915.300.425.500      Maxillofacial Injuries
-	C26.915.300.425.500.400      Jaw Fractures
-	C26.915.300.425.500.400.255      Mandibular Fractures
-	C26.915.300.425.500.400.510      Maxillary Fractures
-	C26.915.300.425.500.500      Mandibular Injuries
-	C26.915.300.425.500.550      Orbital Fractures
-	C26.915.300.425.500.950      Zygomatic Fractures
-	C26.915.300.450      Head Injuries, Closed
-	C26.915.300.450.500      Brain Concussion
-	C26.915.300.450.500.250      Contrecoup Injury
-	C26.915.300.450.500.500      Post-Concussion Syndrome
-	C26.915.300.475      Head Injuries, Penetrating
-	C26.915.300.490      Intracranial Hemorrhage, Traumatic
-	C26.915.300.490.150      Brain Hemorrhage, Traumatic
-	C26.915.300.490.150.200      Brain Stem Hemorrhage, Traumatic

## MeSH Tree Changes for 2017

Type	Tree - heading
-	C26.915.300.490.150.300 Cerebral Hemorrhage, Traumatic
-	C26.915.300.490.400 Hematoma, Epidural, Cranial
-	C26.915.300.490.450 Hematoma, Subdural
-	C26.915.300.490.450.050 Hematoma, Subdural, Acute
-	C26.915.300.490.450.120 Hematoma, Subdural, Chronic
-	C26.915.300.490.450.400 Hematoma, Subdural, Intracranial
-	C26.915.300.490.700 Subarachnoid Hemorrhage, Traumatic
-	C26.915.300.745 Skull Fractures
-	C26.915.300.745.300 Skull Fracture, Basilar
-	C26.915.300.745.350 Skull Fracture, Depressed
-	C26.915.650 Peripheral Nerve Injuries
-	C26.930 Tympanic Membrane Perforation
-	C26.940 Vascular System Injuries
-	C26.946 War-Related Injuries
Old Tree	<b>C26.951</b> <b>Wound Infection</b>
-	C26.974 Wounds, Nonpenetrating
-	C26.974.250 Contusions
New Heading	<b>C26.974.250.500</b> <b>Brain Contusion</b>
New Heading	<b>C26.974.250.875</b> <b>Myocardial Contusions</b>
New Tree	<b>C26.974.250.875.500</b> <b>Commotio Cordis</b>
-	C26.974.382 Head Injuries, Closed
-	C26.974.382.200 Brain Concussion
-	C26.974.382.200.250 Contrecoup Injury
-	C26.974.382.200.500 Post-Concussion Syndrome
-	C26.986 Wounds, Penetrating
-	C26.986.111 Corpse Dismemberment
-	C26.986.224 Decapitation
-	C26.986.450 Eye Injuries, Penetrating
-	C26.986.500 Head Injuries, Penetrating
-	C26.986.900 Wounds, Gunshot
-	C26.986.950 Wounds, Stab
-	C26.986.950.500 Needlestick Injuries
-	D01 Inorganic Chemicals
-	D01.029 Acids

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D01.029.260 Acids, Noncarboxylic
-	D01.029.260.093 Boric Acids
-	D01.029.260.100 Borinic Acids
-	D01.029.260.110 Boronic Acids
-	D01.029.260.110.500 Bortezomib
-	D01.029.260.150 Carbonic Acid
-	D01.029.260.300 Hydrobromic Acid
-	D01.029.260.326 Hydrochloric Acid
-	D01.029.260.328 Hydrofluoric Acid
-	D01.029.260.330 Hydrogen Cyanide
-	D01.029.260.340 Hydrogen Sulfide
-	D01.029.260.365 Hypochlorous Acid
-	D01.029.260.550 Nitric Acid
-	D01.029.260.575 Nitrous Acid
-	D01.029.260.650 Perchlorates
-	D01.029.260.675 Periodic Acid
-	D01.029.260.700 Phosphorus Acids
-	D01.029.260.700.600 Phosphinic Acids
-	D01.029.260.700.675 Phosphoric Acids
-	D01.029.260.700.675.374 Phosphates
-	D01.029.260.700.675.374.025 Acidulated Phosphate Fluoride
-	D01.029.260.700.675.374.075 Calcium Phosphates
-	D01.029.260.700.675.374.075.025 Apatites
-	D01.029.260.700.675.374.075.025.300 Hydroxyapatites
-	D01.029.260.700.675.374.075.025.300.150 Durapatite
-	D01.029.260.700.675.374.075.150 Calcium Pyrophosphate
-	D01.029.260.700.675.374.425 Phosphoramides
-	D01.029.260.700.675.374.775 Polyphosphates
-	D01.029.260.700.675.374.775.150 Diphosphates
-	D01.029.260.700.675.374.775.150.150 Calcium Pyrophosphate
-	D01.029.260.700.675.374.775.150.900 Technetium Tc 99m Pyrophosphate
-	D01.029.260.700.675.374.775.950 Tin Polyphosphates
New Heading	<b>D01.029.260.700.675.374.887 Struvite</b>
-	D01.029.260.700.675.750 Phosphotungstic Acid
-	D01.029.260.700.700 Phosphorous Acids



## MeSH Tree Changes for 2017

Type	Tree - heading
-	D01.029.260.863 Silicic Acid
-	D01.029.260.877 Sulfur Acids
-	D01.029.260.877.700 Sulfinic Acids
-	D01.029.260.877.740 Sulfonic Acids
-	D01.029.260.877.740.900 Thiosulfonic Acids
-	D01.029.260.877.800 Sulfuric Acids
-	D01.029.260.877.800.900 Tetrathionic Acid
-	D01.029.630 Lewis Acids
-	D01.045 Alkalies
-	D01.045.125 Carbonates
-	D01.045.125.500 Lithium Carbonate
-	D01.045.250 Hydroxides
-	D01.045.250.050 Aluminum Hydroxide
-	D01.045.250.225 Ammonium Hydroxide
-	D01.045.250.313 Calcium Hydroxide
-	D01.045.250.357 Hydroxyl Radical
-	D01.045.250.400 Lye
-	D01.045.250.575 Magnesium Hydroxide
-	D01.045.250.750 Sodium Hydroxide
-	D01.045.250.875 Water
-	D01.045.250.875.200 Deuterium Oxide
-	D01.045.250.875.300 Drinking Water
-	D01.045.250.875.400 Ice
-	D01.045.250.875.600 Mineral Waters
-	D01.045.250.875.600.500 Carbonated Water
-	D01.045.250.875.800 Steam
-	D01.045.625 Lewis Bases
-	D01.056 Aluminum Compounds
-	D01.056.025 Alum Compounds
-	D01.056.037 Aluminum Hydroxide
-	D01.056.050 Aluminum Oxide
-	D01.056.050.075 Aluminum Silicates
-	D01.056.050.075.100 Bentonite
-	D01.056.050.075.350 Kaolin
-	D01.056.050.075.975 Zeolites
-	D01.075 Arsenicals

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D01.075.025                      Arsenates
-	D01.075.050                      Arsenites
-	D01.103                              Barium Compounds
-	D01.103.075                      Barium Sulfate
-	D01.132                              Boron Compounds
-	D01.132.105                      Boranes
-	D01.132.105.050                      Borohydrides
-	D01.132.250                      Boric Acids
-	D01.132.250.075                      Borates
-	D01.132.275                      Borinic Acids
-	D01.132.285                      Boronic Acids
-	D01.132.285.500                      Bortezomib
-	D01.139                              Bromine Compounds
-	D01.139.100                      Bromates
-	D01.139.300                      Hydrobromic Acid
-	D01.139.300.050                      Bromides
-	D01.139.300.050.100                      Cyanogen Bromide
-	D01.142                              Cadmium Compounds
-	D01.142.175                      Cadmium Chloride
-	D01.146                              Calcium Compounds
-	D01.146.275                      Calcium Carbonate
-	D01.146.275.500                      Nacre
-	D01.146.300                      Calcium Chloride
-	D01.146.307                      Calcium Citrate
-	D01.146.315                      Calcium, Dietary
-	D01.146.325                      Calcium Fluoride
-	D01.146.335                      Calcium Hydroxide
-	D01.146.360                      Calcium Phosphates
-	D01.146.360.050                      Apatites
-	D01.146.360.050.300                      Hydroxyapatites
-	D01.146.360.050.300.200                      Durapatite
-	D01.146.360.150                      Calcium Pyrophosphate
-	D01.146.375                      Calcium Sulfate
-	D01.200                              Carbon Compounds, Inorganic
-	D01.200.200                      Carbon Dioxide
-	D01.200.200.250                      Dry Ice

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D01.200.210	Carbon Disulfide
-	D01.200.250	Carbon Monoxide
-	D01.200.275	Carbonic Acid
-	D01.200.275.150	Carbonates
-	D01.200.275.150.100	Bicarbonates
-	D01.200.275.150.100.800	Sodium Bicarbonate
-	D01.200.275.150.150	Calcium Carbonate
-	D01.200.275.150.150.500	Nacre
-	D01.200.275.150.550	Lithium Carbonate
-	D01.210	Chlorine Compounds
New Tree	<a href="#">D01.210.150</a>	<a href="#">Chloramines</a>
-	D01.210.300	Chlorates
-	D01.210.375	Cisplatin
-	D01.210.450	Hydrochloric Acid
-	D01.210.450.150	Chlorides
-	D01.210.450.150.050	Ammonium Chloride
-	D01.210.450.150.125	Cadmium Chloride
-	D01.210.450.150.150	Calcium Chloride
-	D01.210.450.150.450	Lithium Chloride
-	D01.210.450.150.500	Magnesium Chloride
-	D01.210.450.150.525	Mercuric Chloride
-	D01.210.450.150.750	Potassium Chloride
-	D01.210.450.150.875	Sodium Chloride
-	D01.210.465	Hypochlorous Acid
-	D01.210.465.800	Sodium Hypochlorite
-	D01.210.675	Perchlorates
-	D01.210.837	Ruthenium Red
-	D01.220	Chromium Compounds
-	D01.220.150	Chromates
-	D01.220.150.650	Potassium Dichromate
-	D01.220.175	Chromium Alloys
-	D01.220.175.950	Vitallium
-	D01.234	Coordination Complexes
-	D01.248	Electrolytes
New	<a href="#">D01.248.249</a>	<a href="#">Polyelectrolytes</a>

## MeSH Tree Changes for 2017

Type	Tree - heading
Heading	
-	D01.248.497 Ions
-	D01.248.497.158 Anions
-	D01.248.497.158.050 Arsenates
-	D01.248.497.158.055 Arsenites
-	D01.248.497.158.076 Borates
-	D01.248.497.158.100 Bromates
-	D01.248.497.158.125 Bromides
-	D01.248.497.158.165 Carbonates
-	D01.248.497.158.165.100 Bicarbonates
-	D01.248.497.158.190 Chlorates
-	D01.248.497.158.215 Chlorides
-	D01.248.497.158.235 Chromates
-	D01.248.497.158.291 Cyanides
-	D01.248.497.158.291.350 Ferricyanides
-	D01.248.497.158.291.350.550 Nitroprusside
-	D01.248.497.158.291.370 Ferrocyanides
-	D01.248.497.158.380 Fluorides
-	D01.248.497.158.459 Hydroxides
-	D01.248.497.158.459.075 Aluminum Hydroxide
-	D01.248.497.158.459.113 Ammonium Hydroxide
-	D01.248.497.158.459.150 Calcium Hydroxide
-	D01.248.497.158.459.300 Hydroxyl Radical
-	D01.248.497.158.459.344 Lye
-	D01.248.497.158.459.388 Magnesium Hydroxide
-	D01.248.497.158.459.475 Sodium Hydroxide
-	D01.248.497.158.459.650 Water
-	D01.248.497.158.459.650.200 Deuterium Oxide
-	D01.248.497.158.459.650.300 Drinking Water
-	D01.248.497.158.459.650.400 Ice
-	D01.248.497.158.459.650.600 Mineral Waters
-	D01.248.497.158.459.650.600.500 Carbonated Water
-	D01.248.497.158.459.650.800 Steam
-	D01.248.497.158.480 Iodates
-	D01.248.497.158.490 Iodides
-	D01.248.497.158.606 Nitrates

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D01.248.497.158.635 Nitrites
-	D01.248.497.158.685 Oxides
-	D01.248.497.158.685.750 Peroxides
-	D01.248.497.158.685.750.212 Artemisinins
-	D01.248.497.158.685.750.424 Hydrogen Peroxide
-	D01.248.497.158.685.750.637 Lipid Peroxides
-	D01.248.497.158.685.750.744 Prostaglandin Endoperoxides
-	D01.248.497.158.685.750.744.600 Prostaglandins G
-	D01.248.497.158.685.750.744.650 Prostaglandins H
-	D01.248.497.158.685.750.744.650.500 Prostaglandin H2
-	D01.248.497.158.685.750.744.650.500.500 (epoxymethano)prosta-5,13-dienoic Acid 15-Hydroxy-11 alpha,9 alpha-
-	D01.248.497.158.685.750.850 Superoxides
-	D01.248.497.158.685.750.925 tert-Butylhydroperoxide
-	D01.248.497.158.685.750.962 Tetraoxanes
-	D01.248.497.158.730 Phosphates
-	D01.248.497.158.730.650 Polyphosphates
-	D01.248.497.158.730.650.200 Diphosphates
-	D01.248.497.158.750 Phosphites
-	D01.248.497.158.845 Sulfates
-	D01.248.497.158.845.703 Thiosulfates
-	D01.248.497.158.874 Sulfides
-	D01.248.497.158.874.390 Disulfides
-	D01.248.497.158.874.390.369 Cystine
-	D01.248.497.158.904 Sulfites
-	D01.248.497.158.904.150 Dithionite
-	D01.248.497.158.952 Vanadates
-	D01.248.497.300 Cations
-	D01.248.497.300.333 Cations, Divalent
-	D01.248.497.300.459 Cations, Monovalent
-	D01.248.497.300.459.700 Protons
-	D01.268 Elements
-	D01.268.150 Carbon
-	D01.268.150.150 Charcoal
-	D01.268.150.200 Diamond
-	D01.268.150.200.550 Nanodiamonds

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D01.268.150.250 Fullerenes
-	D01.268.150.250.500 Nanotubes, Carbon
-	D01.268.150.300 Graphite
-	D01.268.185 Chalcogens
-	D01.268.185.550 Oxygen
-	D01.268.185.600 Polonium
-	D01.268.185.850 Selenium
-	D01.268.185.900 Sulfur
-	D01.268.185.950 Tellurium
-	D01.268.271 Elements, Radioactive
-	D01.268.271.100 Actinoid Series Elements
-	D01.268.271.100.033 Actinium
-	D01.268.271.100.050 Americium
-	D01.268.271.100.100 Berkelium
-	D01.268.271.100.150 Californium
-	D01.268.271.100.175 Curium
-	D01.268.271.100.250 Einsteinium
-	D01.268.271.100.275 Fermium
-	D01.268.271.100.500 Lawrencium
-	D01.268.271.100.525 Mendelevium
-	D01.268.271.100.550 Neptunium
-	D01.268.271.100.575 Nobelium
-	D01.268.271.100.700 Plutonium
-	D01.268.271.100.725 Protactinium
-	D01.268.271.100.900 Thorium
-	D01.268.271.100.950 Uranium
-	D01.268.271.110 Astatine
-	D01.268.271.380 Francium
-	D01.268.271.680 Polonium
-	D01.268.271.720 Promethium
-	D01.268.271.770 Radium
-	D01.268.271.800 Radon
-	D01.268.271.800.800 Radon Daughters
-	D01.268.271.870 Technetium
-	D01.268.380 Halogens
-	D01.268.380.075 Astatine

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D01.268.380.112 Bromine
-	D01.268.380.150 Chlorine
-	D01.268.380.300 Fluorine
-	D01.268.380.400 Iodine
-	D01.268.406 Hydrogen
-	D01.268.406.500 Deuterium
-	D01.268.406.500.250 Deuterium Oxide
-	D01.268.406.750 Protons
-	D01.268.406.875 Tritium
-	D01.268.513 Metalloids
-	D01.268.513.124 Antimony
-	D01.268.513.249 Arsenic
-	D01.268.513.500 Boron
-	D01.268.513.750 Germanium
-	D01.268.513.875 Polonium
-	D01.268.513.937 Silicon
-	D01.268.513.968 Tellurium
-	D01.268.549 Metals, Alkali
-	D01.268.549.125 Cesium
-	D01.268.549.250 Francium
-	D01.268.549.450 Lithium
-	D01.268.549.550 Potassium
-	D01.268.549.650 Rubidium
-	D01.268.549.750 Sodium
-	D01.268.552 Metals, Alkaline Earth
-	D01.268.552.050 Barium
-	D01.268.552.075 Beryllium
-	D01.268.552.100 Calcium
-	D01.268.552.437 Magnesium
-	D01.268.552.775 Radium
-	D01.268.552.850 Strontium
-	D01.268.556 Metals, Heavy
-	D01.268.556.025 Actinium
-	D01.268.556.037 Americium
-	D01.268.556.050 Antimony
-	D01.268.556.062 Barium

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D01.268.556.075 Berkelium
-	D01.268.556.100 Bismuth
-	D01.268.556.137 Cadmium
-	D01.268.556.156 Californium
-	D01.268.556.165 Cesium
-	D01.268.556.175 Chromium
-	D01.268.556.185 Cobalt
-	D01.268.556.195 Copper
-	D01.268.556.200 Curium
-	D01.268.556.244 Einsteinium
-	D01.268.556.266 Fermium
-	D01.268.556.277 Francium
-	D01.268.556.289 Gallium
-	D01.268.556.305 Germanium
-	D01.268.556.322 Gold
-	D01.268.556.337 Hafnium
-	D01.268.556.381 Indium
-	D01.268.556.401 Iridium
-	D01.268.556.412 Iron
-	D01.268.556.423 Lawrencium
-	D01.268.556.435 Lead
-	D01.268.556.484 Manganese
-	D01.268.556.494 Mendeleevium
-	D01.268.556.504 Mercury
-	D01.268.556.555 Molybdenum
-	D01.268.556.600 Neptunium
-	D01.268.556.607 Nickel
-	D01.268.556.615 Niobium
-	D01.268.556.620 Nobelium
-	D01.268.556.660 Osmium
-	D01.268.556.680 Palladium
-	D01.268.556.690 Platinum
-	D01.268.556.700 Plutonium
-	D01.268.556.750 Protactinium
-	D01.268.556.775 Radium
-	D01.268.556.787 Rhenium



## MeSH Tree Changes for 2017

Type	Tree - heading
-	D01.268.556.793 Rhodium
-	D01.268.556.800 Rubidium
-	D01.268.556.805 Ruthenium
-	D01.268.556.812 Silver
-	D01.268.556.825 Strontium
-	D01.268.556.837 Tantalum
-	D01.268.556.843 Technetium
-	D01.268.556.847 Thallium
-	D01.268.556.850 Thorium
-	D01.268.556.875 Tin
-	D01.268.556.887 Tungsten
-	D01.268.556.900 Uranium
-	D01.268.556.920 Vanadium
-	D01.268.556.940 Zinc
-	D01.268.556.950 Zirconium
-	D01.268.557 Metals, Light
-	D01.268.557.050 Aluminum
-	D01.268.557.080 Beryllium
-	D01.268.557.290 Lithium
-	D01.268.557.500 Magnesium
-	D01.268.557.575 Potassium
-	D01.268.557.650 Sodium
-	D01.268.557.800 Titanium
-	D01.268.558 Metals, Rare Earth
-	D01.268.558.362 Lanthanoid Series Elements
-	D01.268.558.362.249 Cerium
-	D01.268.558.362.374 Dysprosium
-	D01.268.558.362.437 Erbium
-	D01.268.558.362.468 Europium
-	D01.268.558.362.484 Gadolinium
-	D01.268.558.362.492 Holmium
-	D01.268.558.362.500 Lanthanum
-	D01.268.558.362.562 Lutetium
-	D01.268.558.362.625 Neodymium
-	D01.268.558.362.750 Praseodymium
-	D01.268.558.362.875 Promethium

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D01.268.558.362.937                      Samarium
-	D01.268.558.362.968                      Terbium
-	D01.268.558.362.984                      Thulium
-	D01.268.558.362.992                      Ytterbium
-	D01.268.558.800                          Scandium
-	D01.268.558.975                          Yttrium
-	D01.268.604                                Nitrogen
-	D01.268.613                                Noble Gases
-	D01.268.613.050                          Argon
-	D01.268.613.350                          Helium
-	D01.268.613.425                          Krypton
-	D01.268.613.600                          Neon
-	D01.268.613.700                          Radon
-	D01.268.613.900                          Xenon
-	D01.268.666                                Phosphorus
-	D01.268.956                                Transition Elements
-	D01.268.956.061                          Cadmium
-	D01.268.956.124                          Chromium
-	D01.268.956.155                          Cobalt
-	D01.268.956.170                          Copper
-	D01.268.956.186                          Gold
-	D01.268.956.249                          Hafnium
-	D01.268.956.280                          Iridium
-	D01.268.956.287                          Iron
-	D01.268.956.295                          Lawrencium
-	D01.268.956.311                          Lutetium
-	D01.268.956.374                          Manganese
-	D01.268.956.437                          Mercury
-	D01.268.956.500                          Molybdenum
-	D01.268.956.625                          Nickel
-	D01.268.956.687                          Niobium
-	D01.268.956.702                          Osmium
-	D01.268.956.718                          Palladium
-	D01.268.956.734                          Platinum
-	D01.268.956.750                          Rhenium
-	D01.268.956.781                          Rhodium

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D01.268.956.812 Ruthenium
-	D01.268.956.827 Scandium
-	D01.268.956.843 Silver
-	D01.268.956.859 Tantalum
-	D01.268.956.875 Technetium
-	D01.268.956.878 Titanium
-	D01.268.956.882 Tungsten
-	D01.268.956.886 Vanadium
-	D01.268.956.890 Yttrium
-	D01.268.956.906 Zinc
-	D01.268.956.937 Zirconium
-	D01.303 Fluorine Compounds
-	D01.303.350 Hydrofluoric Acid
-	D01.303.350.300 Fluorides
-	D01.303.350.300.150 Calcium Fluoride
-	D01.303.350.300.512 Fluorides, Topical
-	D01.303.350.300.875 Sodium Fluoride
-	D01.303.350.300.875.050 Acidulated Phosphate Fluoride
-	D01.303.350.300.900 Sulfur Hexafluoride
-	D01.303.350.300.950 Tin Fluorides
-	D01.339 Free Radicals
-	D01.339.387 Nitric Oxide
-	D01.339.431 Reactive Oxygen Species
-	D01.339.431.249 Hydroxyl Radical
-	D01.339.431.311 Hypochlorous Acid
-	D01.339.431.374 Peroxides
-	D01.339.431.374.212 Artemisinins
-	D01.339.431.374.424 Hydrogen Peroxide
-	D01.339.431.374.637 Lipid Peroxides
-	D01.339.431.374.744 Prostaglandin Endoperoxides
-	D01.339.431.374.744.600 Prostaglandins G
-	D01.339.431.374.744.650 Prostaglandins H
-	D01.339.431.374.744.650.500 Prostaglandin H2
-	D01.339.431.374.744.650.500.500 15-Hydroxy-11 alpha,9 alpha-(epoxymethano)prosta-5,13-dienoic Acid
-	D01.339.431.374.850 Superoxides

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D01.339.431.374.925 tert-Butylhydroperoxide
-	D01.339.431.374.962 Tetraoxanes
-	D01.339.431.500 Singlet Oxygen
-	D01.362 Gases
-	D01.362.075 Ammonia
-	D01.362.150 Carbon Dioxide
-	D01.362.200 Carbon Monoxide
-	D01.362.225 Chlorine
-	D01.362.282 Gasotransmitters
-	D01.362.340 Hydrogen
-	D01.362.340.500 Deuterium
-	D01.362.340.500.250 Deuterium Oxide
-	D01.362.340.750 Protons
-	D01.362.340.875 Tritium
-	D01.362.350 Hydrogen Sulfide
-	D01.362.625 Nitrogen
-	D01.362.635 Nitrogen Oxides
-	D01.362.635.600 Nitrogen Dioxide
-	D01.362.635.625 Nitrous Oxide
-	D01.362.641 Noble Gases
-	D01.362.641.113 Argon
-	D01.362.641.352 Helium
-	D01.362.641.468 Krypton
-	D01.362.641.592 Neon
-	D01.362.641.745 Radon
-	D01.362.641.745.800 Radon Daughters
-	D01.362.641.918 Xenon
-	D01.362.670 Oxygen
-	D01.362.670.600 Ozone
-	D01.362.670.600.500 Stratospheric Ozone
-	D01.362.740 Plasma Gases
-	D01.362.810 Sulfur Dioxide
-	D01.362.820 Sulfur Hexafluoride
-	D01.379 Gold Compounds
-	D01.379.375 Gold Alloys
-	D01.379.400 Gold Colloid

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D01.379.400.350 Gold Colloid, Radioactive
-	D01.379.425 Gold Sodium Thiosulfate
-	D01.475 Iodine Compounds
-	D01.475.400 Iodates
-	D01.475.410 Iodides
-	D01.475.410.700 Potassium Iodide
-	D01.475.410.800 Sodium Iodide
-	D01.475.557 Iodophors
-	D01.475.557.500 Povidone-Iodine
-	D01.475.705 Periodic Acid
-	D01.490 Iron Compounds
-	D01.490.100 Ferric Compounds
-	D01.490.100.300 Ferricyanides
-	D01.490.100.300.550 Nitroprusside
-	D01.490.100.375 Ferrosferric Oxide
-	D01.490.100.375.500 Magnetite Nanoparticles
-	D01.490.200 Ferrous Compounds
-	D01.490.200.250 Ferrocyanides
-	D01.490.200.350 Ferrosferric Oxide
-	D01.490.200.350.500 Magnetite Nanoparticles
-	D01.490.500 Iron Carbonyl Compounds
-	D01.490.600 Iron, Dietary
-	D01.490.800 Steel
-	D01.490.800.900 Stainless Steel
-	D01.496 Isotopes
-	D01.496.098 Calcium Isotopes
-	D01.496.098.325 Calcium Radioisotopes
-	D01.496.123 Carbon Isotopes
-	D01.496.123.328 Carbon Radioisotopes
-	D01.496.156 Cerium Isotopes
-	D01.496.156.300 Cerium Radioisotopes
-	D01.496.180 Cesium Isotopes
-	D01.496.180.300 Cesium Radioisotopes
-	D01.496.212 Chromium Isotopes
-	D01.496.212.349 Chromium Radioisotopes
-	D01.496.239 Cobalt Isotopes

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D01.496.239.354 Cobalt Radioisotopes
-	D01.496.289 Deuterium
-	D01.496.360 Gallium Isotopes
-	D01.496.360.400 Gallium Radioisotopes
-	D01.496.381 Gold Isotopes
-	D01.496.381.550 Gold Radioisotopes
-	D01.496.381.550.400 Gold Colloid, Radioactive
-	D01.496.448 Iodine Isotopes
-	D01.496.448.496 Iodine Radioisotopes
-	D01.496.448.496.665 Serum Albumin, Radio-Iodinated
-	D01.496.473 Iron Isotopes
-	D01.496.473.498 Iron Radioisotopes
-	D01.496.532 Mercury Isotopes
-	D01.496.532.500 Mercury Radioisotopes
-	D01.496.586 Nitrogen Isotopes
-	D01.496.586.520 Nitrogen Radioisotopes
-	D01.496.625 Oxygen Isotopes
-	D01.496.625.600 Oxygen Radioisotopes
-	D01.496.669 Phosphorus Isotopes
-	D01.496.669.604 Phosphorus Radioisotopes
-	D01.496.705 Potassium Isotopes
-	D01.496.705.700 Potassium Radioisotopes
-	D01.496.749 Radioisotopes
-	D01.496.749.075 Barium Radioisotopes
-	D01.496.749.090 Bromine Radioisotopes
-	D01.496.749.108 Cadmium Radioisotopes
-	D01.496.749.113 Calcium Radioisotopes
-	D01.496.749.154 Carbon Radioisotopes
-	D01.496.749.185 Cerium Radioisotopes
-	D01.496.749.190 Cesium Radioisotopes
-	D01.496.749.213 Chromium Radioisotopes
-	D01.496.749.256 Cobalt Radioisotopes
-	D01.496.749.270 Copper Radioisotopes
-	D01.496.749.340 Fluorine Radioisotopes
-	D01.496.749.360 Gallium Radioisotopes
-	D01.496.749.410 Gold Radioisotopes

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D01.496.749.410.400 Gold Colloid, Radioactive
-	D01.496.749.460 Indium Radioisotopes
-	D01.496.749.474 Iodine Radioisotopes
-	D01.496.749.474.665 Serum Albumin, Radio-Iodinated
-	D01.496.749.500 Iridium Radioisotopes
-	D01.496.749.514 Iron Radioisotopes
-	D01.496.749.540 Krypton Radioisotopes
-	D01.496.749.560 Lead Radioisotopes
-	D01.496.749.590 Mercury Radioisotopes
-	D01.496.749.615 Nitrogen Radioisotopes
-	D01.496.749.635 Oxygen Radioisotopes
-	D01.496.749.658 Phosphorus Radioisotopes
-	D01.496.749.690 Potassium Radioisotopes
-	D01.496.749.731 Radioactive Tracers
-	D01.496.749.740 Rubidium Radioisotopes
-	D01.496.749.745 Ruthenium Radioisotopes
-	D01.496.749.780 Selenium Radioisotopes
-	D01.496.749.795 Sodium Radioisotopes
-	D01.496.749.815 Strontium Radioisotopes
-	D01.496.749.858 Sulfur Radioisotopes
-	D01.496.749.900 Thallium Radioisotopes
-	D01.496.749.910 Tin Radioisotopes
-	D01.496.749.925 Tritium
-	D01.496.749.945 Xenon Radioisotopes
-	D01.496.749.960 Yttrium Radioisotopes
-	D01.496.749.980 Zinc Radioisotopes
-	D01.496.807 Sodium Isotopes
-	D01.496.807.800 Sodium Radioisotopes
-	D01.496.840 Strontium Isotopes
-	D01.496.840.685 Strontium Radioisotopes
-	D01.496.868 Sulfur Isotopes
-	D01.496.868.690 Sulfur Radioisotopes
-	D01.496.920 Xenon Isotopes
-	D01.496.920.700 Xenon Radioisotopes
-	D01.496.943 Yttrium Isotopes
-	D01.496.943.800 Yttrium Radioisotopes

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D01.496.966 Zinc Isotopes
-	D01.496.966.800 Zinc Radioisotopes
-	D01.510 Lithium Compounds
-	D01.510.475 Lithium Carbonate
-	D01.510.500 Lithium Chloride
-	D01.524 Magnesium Compounds
-	D01.524.475 Magnesium Chloride
-	D01.524.485 Magnesium Hydroxide
-	D01.524.490 Magnesium Oxide
-	D01.524.500 Magnesium Silicates
-	D01.524.500.040 Asbestos, Amosite
-	D01.524.500.050 Asbestos, Serpentine
-	D01.524.500.850 Talc
-	D01.524.550 Magnesium Sulfate
New Heading	<b>D01.524.775 Struvite</b>
-	D01.530 Manganese Compounds
-	D01.530.700 Potassium Permanganate
-	D01.538 Mercury Compounds
-	D01.538.500 Mercuric Chloride
-	D01.552 Metals
-	D01.552.020 Actinoid Series Elements
-	D01.552.020.042 Actinium
-	D01.552.020.086 Americium
-	D01.552.020.150 Berkelium
-	D01.552.020.200 Californium
-	D01.552.020.272 Curium
-	D01.552.020.337 Einsteinium
-	D01.552.020.394 Fermium
-	D01.552.020.507 Lawrencium
-	D01.552.020.561 Mendelevium
-	D01.552.020.621 Neptunium
-	D01.552.020.659 Nobelium
-	D01.552.020.734 Plutonium
-	D01.552.020.778 Protactinium
-	D01.552.020.889 Thorium



## MeSH Tree Changes for 2017

Type	Tree - heading
-	D01.552.020.940 Uranium
-	D01.552.033 Alloys
-	D01.552.033.182 Chromium Alloys
-	D01.552.033.182.900 Vitallium
-	D01.552.033.533 Gold Alloys
-	D01.552.033.690 Metal Ceramic Alloys
-	D01.552.033.690.250 Cermet Cements
-	D01.552.033.847 Steel
-	D01.552.033.847.681 Stainless Steel
-	D01.552.528 Metals, Alkali
-	D01.552.528.160 Cesium
-	D01.552.528.328 Francium
-	D01.552.528.480 Lithium
-	D01.552.528.652 Potassium
-	D01.552.528.759 Rubidium
-	D01.552.528.850 Sodium
-	D01.552.539 Metals, Alkaline Earth
-	D01.552.539.124 Barium
-	D01.552.539.288 Calcium
-	D01.552.539.745 Radium
-	D01.552.539.861 Strontium
-	D01.552.544 Metals, Heavy
-	D01.552.544.025 Actinium
-	D01.552.544.037 Americium
-	D01.552.544.050 Antimony
-	D01.552.544.062 Barium
-	D01.552.544.075 Berkelium
-	D01.552.544.100 Bismuth
-	D01.552.544.137 Cadmium
-	D01.552.544.156 Californium
-	D01.552.544.165 Cesium
-	D01.552.544.175 Chromium
-	D01.552.544.185 Cobalt
-	D01.552.544.195 Copper
-	D01.552.544.200 Curium
-	D01.552.544.244 Einsteinium

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D01.552.544.266	Fermium
-	D01.552.544.277	Francium
-	D01.552.544.289	Gallium
-	D01.552.544.305	Germanium
-	D01.552.544.322	Gold
-	D01.552.544.337	Hafnium
-	D01.552.544.381	Indium
-	D01.552.544.401	Iridium
-	D01.552.544.412	Iron
-	D01.552.544.423	Lawrencium
-	D01.552.544.435	Lead
-	D01.552.544.484	Manganese
-	D01.552.544.494	Mendelevium
-	D01.552.544.504	Mercury
-	D01.552.544.555	Molybdenum
-	D01.552.544.600	Neptunium
-	D01.552.544.607	Nickel
-	D01.552.544.615	Niobium
-	D01.552.544.620	Nobelium
-	D01.552.544.660	Osmium
-	D01.552.544.680	Palladium
-	D01.552.544.690	Platinum
-	D01.552.544.700	Plutonium
-	D01.552.544.750	Protactinium
-	D01.552.544.775	Radium
-	D01.552.544.787	Rhenium
-	D01.552.544.793	Rhodium
-	D01.552.544.800	Rubidium
-	D01.552.544.805	Ruthenium
-	D01.552.544.812	Silver
-	D01.552.544.825	Strontium
-	D01.552.544.837	Tantalum
-	D01.552.544.843	Technetium
-	D01.552.544.847	Thallium
-	D01.552.544.850	Thorium
-	D01.552.544.875	Tin

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D01.552.544.887 Tungsten
-	D01.552.544.900 Uranium
-	D01.552.544.920 Vanadium
-	D01.552.544.940 Zinc
-	D01.552.544.950 Zirconium
-	D01.552.547 Metals, Light
-	D01.552.547.050 Aluminum
-	D01.552.547.080 Beryllium
-	D01.552.547.290 Lithium
-	D01.552.547.500 Magnesium
-	D01.552.547.650 Potassium
-	D01.552.547.725 Sodium
-	D01.552.547.800 Titanium
-	D01.552.550 Metals, Rare Earth
-	D01.552.550.399 Lanthanoid Series Elements
-	D01.552.550.399.249 Cerium
-	D01.552.550.399.374 Dysprosium
-	D01.552.550.399.437 Erbium
-	D01.552.550.399.468 Europium
-	D01.552.550.399.484 Gadolinium
-	D01.552.550.399.492 Holmium
-	D01.552.550.399.500 Lanthanum
-	D01.552.550.399.562 Lutetium
-	D01.552.550.399.625 Neodymium
-	D01.552.550.399.750 Praseodymium
-	D01.552.550.399.875 Promethium
-	D01.552.550.399.937 Samarium
-	D01.552.550.399.968 Terbium
-	D01.552.550.399.984 Thulium
-	D01.552.550.399.992 Ytterbium
-	D01.552.550.800 Scandium
-	D01.552.550.975 Yttrium
-	D01.578 Minerals
-	D01.578.122 Apatites
-	D01.578.122.477 Hydroxyapatites
-	D01.578.122.477.300 Durapatite

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D01.578.200 Calcium Carbonate
-	D01.578.200.500 Nacre
-	D01.578.215 Calcium Sulfate
-	D01.578.285 Ferrosferric Oxide
-	D01.578.285.500 Magnetite Nanoparticles
-	D01.578.300 Graphite
-	D01.578.500 Mineral Fibers
-	D01.578.700 Selenium
-	D01.578.725 Silicates
-	D01.578.725.025 Aluminum Silicates
-	D01.578.725.025.100 Bentonite
-	D01.578.725.025.350 Kaolin
-	D01.578.725.025.975 Zeolites
-	D01.578.725.050 Asbestos
-	D01.578.725.050.050 Asbestos, Amphibole
-	D01.578.725.050.050.050 Asbestos, Amosite
-	D01.578.725.050.050.100 Asbestos, Crocidolite
-	D01.578.725.050.075 Asbestos, Serpentine
-	D01.578.725.500 Magnesium Silicates
-	D01.578.725.500.040 Asbestos, Amosite
-	D01.578.725.500.050 Asbestos, Serpentine
-	D01.578.725.500.800 Talc
-	D01.578.750 Silicon Dioxide
-	D01.578.750.300 Diatomaceous Earth
-	D01.578.750.600 Quartz
-	D01.625 Nitrogen Compounds
-	D01.625.050 Ammonia
-	D01.625.062 Ammonium Compounds
-	D01.625.062.249 Ammonium Chloride
-	D01.625.062.311 Ammonium Hydroxide
-	D01.625.062.374 Ammonium Sulfate
-	D01.625.062.437 Hydroxylamine
-	D01.625.062.500 Quaternary Ammonium Compounds
-	D01.625.100 Azides
-	D01.625.100.750 Sodium Azide
-	D01.625.125 Cisplatin

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D01.625.175 Cyanogen Bromide
-	D01.625.400 Hydrogen Cyanide
-	D01.625.400.100 Cyanides
-	D01.625.400.100.150 Cyanamide
-	D01.625.400.100.325 Ferricyanides
-	D01.625.400.100.325.550 Nitroprusside
-	D01.625.400.100.350 Ferrocyanides
-	D01.625.400.100.750 Potassium Cyanide
-	D01.625.400.100.875 Sodium Cyanide
-	D01.625.525 Nitric Acid
-	D01.625.525.550 Nitrates
-	D01.625.525.550.850 Silver Nitrate
-	D01.625.525.550.950 Uranyl Nitrate
-	D01.625.550 Nitrogen Oxides
-	D01.625.550.500 Nitric Oxide
-	D01.625.550.525 Nitrogen Dioxide
-	D01.625.550.550 Nitrous Oxide
-	D01.625.600 Nitrous Acid
-	D01.625.600.600 Nitrites
-	D01.625.600.600.800 Sodium Nitrite
-	D01.625.600.800 Peroxynitrous Acid
-	D01.625.700 Reactive Nitrogen Species
-	D01.625.700.500 Nitric Oxide
-	D01.625.700.750 Peroxynitrous Acid
-	D01.625.800 Ruthenium Red
-	D01.640 Osmium Compounds
-	D01.640.500 Osmium Tetroxide
-	D01.650 Oxygen Compounds
-	D01.650.550 Oxides
-	D01.650.550.050 Aluminum Oxide
-	D01.650.550.050.075 Aluminum Silicates
-	D01.650.550.050.075.100 Bentonite
-	D01.650.550.050.075.350 Kaolin
-	D01.650.550.050.075.975 Zeolites
-	D01.650.550.200 Carbon Dioxide
-	D01.650.550.200.250 Dry Ice

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D01.650.550.250 Carbon Monoxide
-	D01.650.550.400 Hypochlorous Acid
-	D01.650.550.400.800 Sodium Hypochlorite
-	D01.650.550.575 Magnesium Oxide
-	D01.650.550.587 Nitrogen Oxides
-	D01.650.550.587.600 Nitric Oxide
-	D01.650.550.587.625 Nitrogen Dioxide
-	D01.650.550.587.650 Nitrous Oxide
-	D01.650.550.600 Osmium Tetroxide
-	D01.650.550.750 Peroxides
-	D01.650.550.750.200 Artemisinins
-	D01.650.550.750.400 Hydrogen Peroxide
-	D01.650.550.750.600 Lipid Peroxides
-	D01.650.550.750.700 Prostaglandin Endoperoxides
-	D01.650.550.750.700.600 Prostaglandins G
-	D01.650.550.750.700.650 Prostaglandins H
-	D01.650.550.750.700.650.500 Prostaglandin H2
-	D01.650.550.750.700.650.500.500 (epoxymethano)prosta-5,13-dienoic Acid 15-Hydroxy-11 alpha,9 alpha-
-	D01.650.550.750.800 Superoxides
-	D01.650.550.750.900 tert-Butylhydroperoxide
-	D01.650.550.750.950 Tetraoxanes
-	D01.650.550.825 Silicon Dioxide
-	D01.650.550.825.400 Diatomaceous Earth
-	D01.650.550.825.600 Quartz
-	D01.650.550.850 Sulfur Oxides
-	D01.650.550.850.925 Sulfur Dioxide
-	D01.650.550.875 Thorium Dioxide
-	D01.650.550.925 Water
-	D01.650.550.925.199 Drinking Water
-	D01.650.550.925.400 Ice
-	D01.650.550.925.600 Mineral Waters
-	D01.650.550.925.600.500 Carbonated Water
-	D01.650.550.925.800 Steam
-	D01.650.550.975 Zinc Oxide
-	D01.650.775 Reactive Oxygen Species

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D01.695 Phosphorus Compounds
-	D01.695.525 Phosphines
-	D01.695.550 Phosphoranes
-	D01.695.625 Phosphorus Acids
-	D01.695.625.600 Phosphinic Acids
-	D01.695.625.675 Phosphoric Acids
-	D01.695.625.675.650 Phosphates
-	D01.695.625.675.650.025 Acidulated Phosphate Fluoride
-	D01.695.625.675.650.075 Calcium Phosphates
-	D01.695.625.675.650.075.025 Apatites
-	D01.695.625.675.650.075.025.300 Hydroxyapatites
-	D01.695.625.675.650.075.025.300.150 Durapatite
-	D01.695.625.675.650.075.150 Calcium Pyrophosphate
-	D01.695.625.675.650.425 Phosphoramides
-	D01.695.625.675.650.775 Polyphosphates
-	D01.695.625.675.650.775.150 Diphosphates
-	D01.695.625.675.650.775.150.150 Calcium Pyrophosphate
-	D01.695.625.675.650.775.150.900 Technetium Tc 99m Pyrophosphate
-	D01.695.625.675.650.775.950 Tin Polyphosphates
New Heading	<b>D01.695.625.675.650.887 Struvite</b>
-	D01.695.625.675.750 Phosphotungstic Acid
-	D01.695.625.700 Phosphorous Acids
-	D01.695.625.700.675 Phosphites
-	D01.695.635 Phosphorus, Dietary
-	D01.710 Platinum Compounds
-	D01.710.100 Cisplatin
-	D01.745 Potassium Compounds
-	D01.745.312 Potassium Acetate
-	D01.745.625 Potassium Chloride
-	D01.745.635 Potassium Cyanide
-	D01.745.650 Potassium Dichromate
-	D01.745.660 Potassium, Dietary
-	D01.745.680 Potassium Iodide
-	D01.745.750 Potassium Permanganate
-	D01.765 Ruthenium Compounds

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D01.765.765 Ruthenium Red
-	D01.786 Salts
-	D01.810 Selenium Compounds
-	D01.810.449 Selenic Acid
-	D01.810.900 Selenious Acid
-	D01.810.900.500 Sodium Selenite
-	D01.810.950 Selenium Oxides
-	D01.837 Silicon Compounds
-	D01.837.700 Silanes
-	D01.837.725 Silicon Dioxide
-	D01.837.725.400 Diatomaceous Earth
-	D01.837.725.600 Quartz
-	D01.837.725.700 Silicic Acid
-	D01.837.725.700.379 Silica Gel
-	D01.837.725.700.760 Silicates
-	D01.837.725.700.760.050 Aluminum Silicates
-	D01.837.725.700.760.050.100 Bentonite
-	D01.837.725.700.760.050.400 Kaolin
-	D01.837.725.700.760.050.950 Zeolites
-	D01.837.725.700.760.070 Asbestos
-	D01.837.725.700.760.070.050 Asbestos, Amphibole
-	D01.837.725.700.760.070.050.060 Asbestos, Amosite
-	D01.837.725.700.760.070.050.090 Asbestos, Crocidolite
-	D01.837.725.700.760.070.110 Asbestos, Serpentine
-	D01.837.725.700.760.535 Magnesium Silicates
-	D01.837.725.700.760.535.040 Asbestos, Amosite
-	D01.837.725.700.760.535.400 Asbestos, Serpentine
-	D01.837.725.700.760.535.800 Talc
-	D01.837.800 Siloxanes
-	D01.847 Silver Compounds
-	D01.847.900 Silver Nitrate
-	D01.857 Sodium Compounds
-	D01.857.300 Gold Sodium Thiosulfate
-	D01.857.462 Sodium Azide
-	D01.857.625 Sodium Bicarbonate
-	D01.857.650 Sodium Chloride



## MeSH Tree Changes for 2017

Type	Tree - heading
-	D01.857.650.705 Sodium Chloride, Dietary
-	D01.857.660 Sodium Cyanide
-	D01.857.700 Sodium, Dietary
-	D01.857.700.705 Sodium Chloride, Dietary
-	D01.857.725 Sodium Fluoride
-	D01.857.725.050 Acidulated Phosphate Fluoride
-	D01.857.745 Sodium Hydroxide
-	D01.857.750 Sodium Hypochlorite
-	D01.857.760 Sodium Iodide
-	D01.857.775 Sodium Nitrite
-	D01.857.850 Sodium Selenite
-	D01.875 Sulfur Compounds
-	D01.875.350 Hydrogen Sulfide
-	D01.875.350.850 Sulfides
-	D01.875.350.850.074 Carbon Disulfide
-	D01.875.350.850.150 Disulfides
-	D01.875.350.850.150.369 Cystine
-	D01.875.750 Sulfites
-	D01.875.750.200 Dithionite
-	D01.875.800 Sulfur Acids
-	D01.875.800.700 Sulfinic Acids
-	D01.875.800.740 Sulfonic Acids
-	D01.875.800.740.900 Thiosulfonic Acids
-	D01.875.800.800 Sulfuric Acids
-	D01.875.800.800.850 Sulfates
-	D01.875.800.800.850.025 Alum Compounds
-	D01.875.800.800.850.050 Ammonium Sulfate
-	D01.875.800.800.850.075 Barium Sulfate
-	D01.875.800.800.850.125 Calcium Sulfate
-	D01.875.800.800.850.150 Copper Sulfate
-	D01.875.800.800.850.500 Magnesium Sulfate
-	D01.875.800.800.850.900 Thiosulfates
-	D01.875.800.800.850.900.250 Gold Sodium Thiosulfate
-	D01.875.800.800.850.950 Zinc Sulfate
-	D01.875.800.800.900 Tetrathionic Acid
-	D01.875.812 Sulfur Hexafluoride

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D01.875.825 Sulfur Oxides
-	D01.875.825.925 Sulfur Dioxide
-	D01.875.900 Technetium Tc 99m Sulfur Colloid
-	D01.925 Technetium Compounds
-	D01.925.800 Sodium Pertechnetate Tc 99m
-	D01.925.925 Technetium Tc 99m Pyrophosphate
-	D01.925.950 Technetium Tc 99m Sulfur Colloid
-	D01.930 Thorium Compounds
-	D01.930.930 Thorium Dioxide
-	D01.935 Tin Compounds
-	D01.935.925 Tin Fluorides
-	D01.935.950 Tin Polyphosphates
-	D01.940 Tungsten Compounds
-	D01.940.700 Phosphotungstic Acid
-	D01.950 Uranium Compounds
-	D01.950.950 Uranyl Nitrate
-	D01.960 Vanadium Compounds
-	D01.960.960 Vanadates
-	D01.975 Zinc Compounds
-	D01.975.975 Zinc Oxide
-	D01.975.987 Zinc Sulfate
-	D02 Organic Chemicals
-	D02.033 Alcohols
-	D02.033.100 Amino Alcohols
-	D02.033.100.291 Ethanolamines
-	D02.033.100.291.057 Albuterol
-	D02.033.100.291.057.125 Albuterol, Ipratropium Drug Combination
-	D02.033.100.291.057.500 Levalbuterol
-	D02.033.100.291.057.750 Salmeterol Xinafoate
-	D02.033.100.291.057.750.500 Fluticasone Propionate, Salmeterol Xinafoate Drug Combination
-	D02.033.100.291.211 Choline
-	D02.033.100.291.211.500 Platelet Activating Factor
-	D02.033.100.291.231 Clenbuterol
-	D02.033.100.291.274 Deanol
-	D02.033.100.291.310 Epinephrine

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.033.100.291.310.500 Racepinephrine
-	D02.033.100.291.375 Ethanolamine
-	D02.033.100.291.410 2-Hydroxyphenethylamine
-	D02.033.100.291.425 Formoterol Fumarate
-	D02.033.100.291.425.250 Budesonide, Formoterol Fumarate Drug Combination
-	D02.033.100.291.425.500 Mometasone Furoate, Formoterol Fumarate Drug Combination
-	D02.033.100.291.439 Isoproterenol
-	D02.033.100.291.460 Labetalol
-	D02.033.100.291.465 Metaproterenol
-	D02.033.100.291.465.300 Fenoterol
-	D02.033.100.291.480 Midodrine
-	D02.033.100.291.491 Nebivolol
-	D02.033.100.291.502 Norepinephrine
-	D02.033.100.291.502.651 Normetanephrine
-	D02.033.100.291.525 Octopamine
-	D02.033.100.291.617 Phenylephrine
-	D02.033.100.291.617.300 Etilefrine
-	D02.033.100.291.630 Procaterol
-	D02.033.100.291.805 Sotalol
-	D02.033.100.291.870 Synephrine
-	D02.033.100.291.905 Terbutaline
-	D02.033.100.350 Heptaminol
-	D02.033.100.400 Isoetharine
-	D02.033.100.624 Propanolamines
-	D02.033.100.624.302 Ephedrine
-	D02.033.100.624.380 Histidinol
-	D02.033.100.624.427 Isoxsuprine
-	D02.033.100.624.536 Methoxamine
-	D02.033.100.624.605 Nylidrin
-	D02.033.100.624.675 Oxyfedrine
-	D02.033.100.624.698 Phenoxypropanolamines
-	D02.033.100.624.698.025 Acebutolol
-	D02.033.100.624.698.055 Alprenolol
-	D02.033.100.624.698.055.200 Dihydroalprenolol

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.033.100.624.698.070                      Atenolol
-	D02.033.100.624.698.077                      Betaxolol
-	D02.033.100.624.698.085                      Bisoprolol
-	D02.033.100.624.698.115                      Bunolol
-	D02.033.100.624.698.115.500                      Levobunolol
-	D02.033.100.624.698.146                      Bupranolol
-	D02.033.100.624.698.207                      Carteolol
-	D02.033.100.624.698.268                      Celiprolol
-	D02.033.100.624.698.542                      Metipranolol
-	D02.033.100.624.698.573                      Metoprolol
-	D02.033.100.624.698.601                      Nadolol
-	D02.033.100.624.698.633                      Oxprenolol
-	D02.033.100.624.698.688                      Penbutolol
-	D02.033.100.624.698.699                      Pindolol
-	D02.033.100.624.698.699.420                      Iodocyanopindolol
-	D02.033.100.624.698.705                      Practolol
-	D02.033.100.624.698.708                      Prenalterol
-	D02.033.100.624.698.711                      Propranolol
-	D02.033.100.624.698.855                      Xamoterol
-	D02.033.100.624.706                      Phenylpropanolamine
-	D02.033.100.624.706.400                      p-Hydroxynorephedrine
-	D02.033.100.624.706.500                      Metaraminol
-	D02.033.100.624.706.750                      Tolterodine Tartrate
-	D02.033.100.624.853                      Pseudoephedrine
-	D02.033.100.624.870                      Ritodrine
-	D02.033.100.624.890                      Suloctidil
-	D02.033.100.624.915                      Timolol
-	D02.033.100.624.915.500                      Brimonidine Tartrate, Timolol Maleate Drug Combination
-	D02.033.100.700                      Sphingosine
-	D02.033.100.700.350                      Fingolimod Hydrochloride
-	D02.033.100.700.700                      Psychosine
-	D02.033.160                      Benzyl Alcohols
-	D02.033.160.100                      Benzyl Alcohol
-	D02.033.260                      Chlorohydrins
-	D02.033.260.100                      alpha-Chlorohydrin

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.033.260.220 Chlorobutanol
-	D02.033.260.335 Ethchlorvynol
-	D02.033.260.350 Ethylene Chlorohydrin
-	D02.033.375 Ethanol
-	D02.033.375.135 Blood Alcohol Content
-	D02.033.375.270 Ethamoxytriphetol
-	D02.033.375.291 Ethanolamines
-	D02.033.375.291.375 Ethanolamine
-	D02.033.375.310 Ethylene Chlorohydrin
-	D02.033.375.534 Mercaptoethanol
-	D02.033.375.650 Phenylethyl Alcohol
-	D02.033.375.880 Trifluoroethanol
-	D02.033.415 Fatty Alcohols
-	D02.033.415.054 Acetogenins
-	D02.033.415.110 Butanols
-	D02.033.415.110.175 1-Butanol
-	D02.033.415.110.220 Chlorobutanol
-	D02.033.415.110.855 tert-Butyl Alcohol
-	D02.033.415.220 Dodecanol
-	D02.033.415.220.720 Sodium Dodecyl Sulfate
-	D02.033.415.230 Dolichol
-	D02.033.415.230.250 Dolichol Phosphates
-	D02.033.415.400 Farnesol
-	D02.033.415.500 Heptanol
-	D02.033.415.510 Hexanols
-	D02.033.415.510.500 Cyclohexanols
-	D02.033.415.510.500.303 Desvenlafaxine Succinate
-	D02.033.415.510.500.605 Menthol
-	D02.033.415.510.500.802 Tramadol
-	D02.033.415.510.500.901 Venlafaxine Hydrochloride
-	D02.033.415.600 Octanols
-	D02.033.415.600.600 1-Octanol
-	D02.033.415.640 Pentanols
-	D02.033.415.900 Sodium Tetradecyl Sulfate
-	D02.033.455 Glycols
-	D02.033.455.125 Butylene Glycols

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.033.455.125.125 Busulfan
-	D02.033.455.250 Ethylene Glycols
-	D02.033.455.250.130 Chloral Hydrate
-	D02.033.455.250.130.150 Chloralose
-	D02.033.455.250.268 Ethylene Glycol
-	D02.033.455.250.610 Methoxyhydroxyphenylglycol
-	D02.033.455.250.700 Polyethylene Glycols
-	D02.033.455.250.700.150 Cetomacrogol
-	D02.033.455.250.700.485 Hydrogel
-	D02.033.455.250.700.620 Nonoxynol
-	D02.033.455.250.700.660 Octoxynol
-	D02.033.455.250.700.680 Poloxalene
-	D02.033.455.250.700.682 Poloxamer
-	D02.033.455.250.700.685 Polyhydroxyethyl Methacrylate
-	D02.033.455.250.700.690 Polysorbates
-	D02.033.455.706 Propylene Glycols
-	D02.033.455.706.100 alpha-Chlorohydrin
-	D02.033.455.706.300 Chloramphenicol
-	D02.033.455.706.300.850 Thiamphenicol
-	D02.033.455.706.345 Chlorphenesin
-	D02.033.455.706.431 Fingolimod Hydrochloride
-	D02.033.455.706.517 Mephenesin
-	D02.033.455.706.690 Pentaerythritol Tetranitrate
-	D02.033.455.706.725 Propylene Glycol
-	D02.033.455.706.900 Tromethamine
-	D02.033.455.843 Sphingosine
-	D02.033.623 Methanol
-	D02.033.750 Polyvinyl Alcohol
-	D02.033.755 Propanols
-	D02.033.755.600 1-Propanol
-	D02.033.755.615 2-Propanol
-	D02.033.755.624 Propanolamines
-	D02.033.755.624.302 Ephedrine
-	D02.033.755.624.380 Histidinol
-	D02.033.755.624.427 Isoxsuprine
-	D02.033.755.624.536 Methoxamine

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.033.755.624.605 Nylidrin
-	D02.033.755.624.675 Oxyfedrine
-	D02.033.755.624.698 Phenoxypropanolamines
-	D02.033.755.624.698.025 Acebutolol
-	D02.033.755.624.698.055 Alprenolol
-	D02.033.755.624.698.055.200 Dihydroalprenolol
-	D02.033.755.624.698.070 Atenolol
-	D02.033.755.624.698.077 Betaxolol
-	D02.033.755.624.698.085 Bisoprolol
-	D02.033.755.624.698.115 Bunolol
-	D02.033.755.624.698.115.500 Levobunolol
-	D02.033.755.624.698.146 Bupranolol
-	D02.033.755.624.698.207 Carteolol
-	D02.033.755.624.698.268 Celiprolol
-	D02.033.755.624.698.542 Metipranolol
-	D02.033.755.624.698.573 Metoprolol
-	D02.033.755.624.698.601 Nadolol
-	D02.033.755.624.698.633 Oxprenolol
-	D02.033.755.624.698.688 Penbutolol
-	D02.033.755.624.698.699 Pindolol
-	D02.033.755.624.698.699.420 Iodocyanopindolol
-	D02.033.755.624.698.705 Practolol
-	D02.033.755.624.698.708 Prenalterol
-	D02.033.755.624.698.711 Propranolol
-	D02.033.755.624.698.855 Xamoterol
-	D02.033.755.624.706 Phenylpropanolamine
-	D02.033.755.624.706.400 p-Hydroxynorephedrine
-	D02.033.755.624.706.500 Metaraminol
-	D02.033.755.624.706.750 Tolterodine Tartrate
-	D02.033.755.624.853 Pseudoephedrine
-	D02.033.755.624.870 Ritodrine
-	D02.033.755.624.890 Suloctidil
-	D02.033.755.624.915 Timolol
-	D02.033.755.624.915.500 Combination Brimonidine Tartrate, Timolol Maleate Drug
-	D02.033.800 Sugar Alcohols

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.033.800.175 Dithioerythritol
-	D02.033.800.196 Dithiothreitol
-	D02.033.800.329 Erythritol
-	D02.033.800.329.225 Erythrityl Tetranitrate
-	D02.033.800.400 Galactitol
-	D02.033.800.400.350 Dianhydrogalactitol
-	D02.033.800.400.500 Mitolactol
-	D02.033.800.421 Glycerol
-	D02.033.800.421.400 Glyceryl Ethers
-	D02.033.800.421.400.750 Phospholipid Ethers
-	D02.033.800.421.560 Nitroglycerin
-	D02.033.800.519 Inositol
-	D02.033.800.519.400 Inositol Phosphates
-	D02.033.800.519.400.350 Inositol 1,4,5-Trisphosphate
-	D02.033.800.519.400.700 Phytic Acid
-	D02.033.800.609 Mannitol
-	D02.033.800.609.450 Mannitol Phosphates
-	D02.033.800.609.500 Mitobronitol
-	D02.033.800.780 Ribitol
-	D02.033.800.813 Sorbitol
-	D02.033.800.813.480 Isosorbide
-	D02.033.800.813.480.500 Isosorbide Dinitrate
-	D02.033.800.813.550 Meglumine
-	D02.033.800.813.550.500 Diatrizoate Meglumine
-	D02.033.800.813.550.600 Iothalamate Meglumine
-	D02.033.800.936 Xylitol
-	D02.047 Aldehydes
-	D02.047.064 Acetaldehyde
-	D02.047.064.586 Paraldehyde
-	D02.047.122 Acrolein
-	D02.047.222 Benzaldehydes
-	D02.047.407 Formaldehyde
-	D02.047.407.518 Formocresols
-	D02.047.466 Furaldehyde
-	D02.047.532 Glutaral
-	D02.047.587 Glyceraldehyde



## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.047.587.450                      Glyceraldehyde 3-Phosphate
-	D02.047.644                              Glyoxal
-	D02.047.644.620                      Phenylglyoxal
-	D02.047.644.650                      Pyruvaldehyde
-	D02.047.700                              Malondialdehyde
-	D02.047.700.700                      Thiobarbituric Acid Reactive Substances
-	D02.047.750                              o-Phthalaldehyde
-	D02.047.850                              Retinaldehyde
-	D02.065                                      Amides
-	D02.065.064                              Acetamides
-	D02.065.064.150                      2-Acetylaminofluorene
-	D02.065.064.150.100                      Acetoxyacetylaminofluorene
-	D02.065.064.150.400                      Hydroxyacetylaminofluorene
-	D02.065.064.189                      Allylisopropylacetamide
-	D02.065.064.294                      Benzeneacetamides
-	D02.065.064.294.088                      Bufexamac
-	D02.065.064.400                      Iodoacetamide
-	D02.065.064.463                      Linezolid
-	D02.065.064.525                      Oseltamivir
-	D02.065.064.650                      Piracetam
-	D02.065.064.786                      Thioacetamide
-	D02.065.122                              Acrylamides
-	D02.065.122.015                      Acrylamide
-	D02.065.141                              Bimatoprost
-	D02.065.160                              Polyunsaturated Alkamides
-	D02.065.160.500                      Capsaicin
-	D02.065.199                              Anilides
-	D02.065.199.092                      Acetanilides
-	D02.065.199.092.040                      Acetaminophen
-	D02.065.199.092.250                      Diamfenetide
-	D02.065.199.092.325                      Etidocaine
-	D02.065.199.092.450                      Inosine Pranobex
-	D02.065.199.092.500                      Lidocaine
-	D02.065.199.092.700                      Phenacetin
-	D02.065.199.092.800                      Practolol
-	D02.065.199.092.850                      Ranolazine

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.065.199.092.900 Trimecaine
-	D02.065.199.200 Benzoylarginine Nitroanilide
-	D02.065.199.239 Bupivacaine
-	D02.065.199.326 Carbanilides
-	D02.065.199.326.400 Imidocarb
-	D02.065.199.326.550 Nicarbazin
-	D02.065.199.350 Carboxin
-	D02.065.199.385 Encainide
-	D02.065.199.420 Flutamide
-	D02.065.199.750 Prilocaine
-	D02.065.199.789 Propanil
-	D02.065.199.860 Salicylanilides
-	D02.065.199.860.470 Niclosamide
-	D02.065.199.860.500 Oxyclozanide
-	D02.065.199.860.600 Rafoxanide
-	D02.065.199.920 Tocainide
-	D02.065.277 Benzamides
-	D02.065.277.067 Bezafibrate
-	D02.065.277.135 Cisapride
-	D02.065.277.194 DEET
-	D02.065.277.225 Dinitolmide
-	D02.065.277.431 Hippurates
-	D02.065.277.431.192 Aminohippuric Acids
-	D02.065.277.431.192.100 p-Aminohippuric Acid
-	D02.065.277.431.579 Iodohippuric Acid
-	D02.065.277.456 Imatinib Mesylate
-	D02.065.277.480 Indoramin
-	D02.065.277.573 Metoclopramide
-	D02.065.277.600 Moclobemide
-	D02.065.277.650 Procainamide
-	D02.065.277.727 Procarbazine
-	D02.065.277.761 Raclopride
-	D02.065.277.796 Remoxipride
-	D02.065.277.866 Sulpiride
-	D02.065.300 Benzoylarginine-2-Naphthylamide
-	D02.065.313 Ceramides

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.065.313.250 Cerebrosides
-	D02.065.313.250.450 Galactosylceramides
-	D02.065.313.250.490 Glucosylceramides
-	D02.065.313.425 Globosides
-	D02.065.313.612 Lactosylceramides
-	D02.065.313.906 Trihexosylceramides
-	D02.065.327 Cerulenin
-	D02.065.355 Dibucaine
-	D02.065.463 Formamides
-	D02.065.463.387 Dimethylformamide
-	D02.065.589 Lactams
-	D02.065.589.099 beta-Lactams
-	D02.065.589.099.124 Carbapenems
-	D02.065.589.099.124.300 Thienamycins
-	D02.065.589.099.124.300.500 Imipenem
-	D02.065.589.099.249 Cephalosporins
-	D02.065.589.099.249.150 Cefamandole
-	D02.065.589.099.249.150.160 Cefoperazone
-	D02.065.589.099.249.160 Cefazolin
-	D02.065.589.099.249.177 Cefonicid
-	D02.065.589.099.249.185 Cefsulodin
-	D02.065.589.099.249.190 Cephacetrile
-	D02.065.589.099.249.190.190 Cefotaxime
-	D02.065.589.099.249.190.190.115 Cefixime
-	D02.065.589.099.249.190.190.125 Cefmenoxime
-	D02.065.589.099.249.190.190.135 Cefotiam
-	D02.065.589.099.249.190.190.145 Ceftizoxime
-	D02.065.589.099.249.190.190.155 Ceftriaxone
-	D02.065.589.099.249.190.190.165 Cefuroxime
-	D02.065.589.099.249.190.210 Cephalothin
-	D02.065.589.099.249.190.230 Cephapirin
-	D02.065.589.099.249.200 Cephalixin
-	D02.065.589.099.249.200.155 Cefaclor
-	D02.065.589.099.249.200.165 Cefadroxil
-	D02.065.589.099.249.200.165.125 Cefatrizine
-	D02.065.589.099.249.200.180 Cephaloglycin

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.065.589.099.249.200.185 Cephadrine
-	D02.065.589.099.249.210 Cephaloridine
-	D02.065.589.099.249.210.150 Ceftazidime
-	D02.065.589.099.249.250 Cephamycins
-	D02.065.589.099.249.250.177 Cefmetazole
-	D02.065.589.099.249.250.199 Cefotetan
-	D02.065.589.099.249.250.222 Cefoxitin
-	D02.065.589.099.374 Clavulanic Acids
-	D02.065.589.099.374.160 Clavulanic Acid
-	D02.065.589.099.374.160.060 Amoxicillin-Potassium Clavulanate Combination
-	D02.065.589.099.500 Monobactams
-	D02.065.589.099.500.044 Aztreonam
-	D02.065.589.099.625 Moxalactam
-	D02.065.589.099.750 Penicillins
-	D02.065.589.099.750.124 Amdinocillin
-	D02.065.589.099.750.124.036 Amdinocillin Pivoxil
-	D02.065.589.099.750.249 Cyclacillin
-	D02.065.589.099.750.500 Methicillin
-	D02.065.589.099.750.562 Nafcillin
-	D02.065.589.099.750.625 Oxacillin
-	D02.065.589.099.750.625.150 Cloxacillin
-	D02.065.589.099.750.625.150.205 Dicloxacillin
-	D02.065.589.099.750.625.150.250 Floxacillin
-	D02.065.589.099.750.687 Penicillanic Acid
-	D02.065.589.099.750.750 Penicillin G
-	D02.065.589.099.750.750.050 Ampicillin
-	D02.065.589.099.750.750.050.050 Amoxicillin
-	D02.065.589.099.750.750.050.050.060 Amoxicillin-Potassium Clavulanate Combination
-	D02.065.589.099.750.750.050.075 Azlocillin
-	D02.065.589.099.750.750.050.500 Mezlocillin
-	D02.065.589.099.750.750.050.650 Piperacillin
-	D02.065.589.099.750.750.050.700 Pivampicillin
-	D02.065.589.099.750.750.050.900 Talampicillin
-	D02.065.589.099.750.750.170 Carbenicillin
-	D02.065.589.099.750.750.170.200 Carfecillin

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.065.589.099.750.750.685 Penicillin G Benzathine
-	D02.065.589.099.750.750.695 Penicillin G Procaine
-	D02.065.589.099.750.750.875 Sulbenicillin
-	D02.065.589.099.750.781 Penicillin V
-	D02.065.589.099.750.812 Sulbactam
-	D02.065.589.099.750.875 Ticarcillin
-	D02.065.589.200 Caprolactam
-	D02.065.589.327 Lactams, Macrocyclic
-	D02.065.793 Salicylamides
-	D02.065.793.324 Labetalol
-	D02.065.793.650 Salicylanilides
-	D02.065.793.650.470 Niclosamide
-	D02.065.793.650.500 Oxyclozanide
-	D02.065.793.650.600 Rafoxanide
-	D02.065.884 Sulfonamides
-	D02.065.884.120 Benzamide
-	D02.065.884.150 Bumetanide
-	D02.065.884.247 Celecoxib
Old Tree	D02.065.884.344 Chloramines
-	D02.065.884.365 Chlorthalidone
-	D02.065.884.390 Clopamide
-	D02.065.884.438 Darunavir
-	D02.065.884.485 Dichlorphenamide
-	D02.065.884.500 Ethoxzolamide
-	D02.065.884.540 Indapamide
-	D02.065.884.570 Mafenide
-	D02.065.884.575 Mefruside
-	D02.065.884.600 Metolazone
-	D02.065.884.625 Probenecid
-	D02.065.884.650 Rosuvastatin Calcium
-	D02.065.884.675 Sildenafil Citrate
-	D02.065.884.725 Sulfanilamides
-	D02.065.884.725.300 Furosemide
-	D02.065.884.725.650 Sulfacetamide
-	D02.065.884.725.700 Sulfachlorpyridazine
-	D02.065.884.725.755 Sulfadiazine

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.065.884.725.755.800 Silver Sulfadiazine
-	D02.065.884.725.760 Sulfadimethoxine
-	D02.065.884.725.765 Sulfadoxine
-	D02.065.884.725.810 Sulfaguanidine
-	D02.065.884.725.850 Sulfalene
-	D02.065.884.725.853 Sulfamerazine
-	D02.065.884.725.857 Sulfameter
-	D02.065.884.725.862 Sulfamethazine
-	D02.065.884.725.867 Sulfamethoxazole
-	D02.065.884.725.867.500 Trimethoprim, Sulfamethoxazole Drug Combination
-	D02.065.884.725.872 Sulfamethoxyipyridazine
-	D02.065.884.725.877 Sulfamonomethoxine
-	D02.065.884.725.882 Sulfamoxole
-	D02.065.884.725.887 Sulfaphenazole
-	D02.065.884.725.900 Sulfapyridine
-	D02.065.884.725.925 Sulfaquinoxaline
-	D02.065.884.725.935 Sulfathiazoles
-	D02.065.884.725.935.813 Sulfamethizole
-	D02.065.884.725.940 Sulfisomidine
-	D02.065.884.725.945 Sulfisoxazole
-	D02.065.884.730 Sulfasalazine
-	D02.065.884.750 Sumatriptan
-	D02.065.884.975 Xipamide
-	D02.065.900 Thioamides
-	D02.065.900.890 Thioacetamide
-	D02.078 Amidines
-	D02.078.100 Benzamidines
-	D02.078.100.260 Diminazene
-	D02.078.100.700 Pentamidine
-	D02.078.200 Chlorphenamidine
-	D02.078.370 Guanidines
-	D02.078.370.060 Agmatine
-	D02.078.370.120 Bethanidine
-	D02.078.370.141 Biguanides
-	D02.078.370.141.075 Buformin
-	D02.078.370.141.100 Chlorhexidine

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.078.370.141.450 Metformin
-	D02.078.370.141.450.500 Sitagliptin Phosphate, Metformin Hydrochloride Drug Combination
-	D02.078.370.141.700 Phenformin
-	D02.078.370.141.710 Proguanil
-	D02.078.370.200 Cimetidine
-	D02.078.370.280 Creatine
-	D02.078.370.380 Gabexate
-	D02.078.370.435 Guanabenz
-	D02.078.370.460 Guanethidine
-	D02.078.370.465 Guanfacine
-	D02.078.370.472 Guanidine
-	D02.078.370.500 Impromidine
-	D02.078.370.510 3-Iodobenzylguanidine
-	D02.078.370.570 Methylguanidine
-	D02.078.370.600 Mitoguazone
-	D02.078.370.649 Nitrosoguanidines
-	D02.078.370.649.400 Methylnitronitrosoguanidine
-	D02.078.370.729 Pinacidil
-	D02.078.370.810 Robenidine
-	D02.078.370.853 Sulfaguanidine
-	D02.078.370.926 Zanamivir
-	D02.078.766 Stilbamidines
-	D02.092 Amines
-	D02.092.050 Allylamine
-	D02.092.050.500 Colesevelam Hydrochloride
-	D02.092.063 Amino Alcohols
-	D02.092.063.291 Ethanolamines
-	D02.092.063.291.057 Albuterol
-	D02.092.063.291.057.125 Albuterol, Ipratropium Drug Combination
-	D02.092.063.291.057.500 Levalbuterol
-	D02.092.063.291.057.750 Salmeterol Xinafoate
-	D02.092.063.291.057.750.500 Fluticasone Propionate, Salmeterol Xinafoate Drug Combination
-	D02.092.063.291.211 Choline
-	D02.092.063.291.211.500 Platelet Activating Factor

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.092.063.291.231                      Clenbuterol
-	D02.092.063.291.274                      Deanol
-	D02.092.063.291.310                      Epinephrine
-	D02.092.063.291.310.500                      Racepinephrine
-	D02.092.063.291.375                      Ethanolamine
-	D02.092.063.291.410                      2-Hydroxyphenethylamine
-	D02.092.063.291.425                      Formoterol Fumarate
-	D02.092.063.291.425.250 Combination                      Budesonide, Formoterol Fumarate Drug
-	D02.092.063.291.425.500 Combination                      Mometasone Furoate, Formoterol Fumarate Drug
-	D02.092.063.291.439                      Isoproterenol
-	D02.092.063.291.460                      Labetalol
-	D02.092.063.291.465                      Metaproterenol
-	D02.092.063.291.465.300                      Fenoterol
-	D02.092.063.291.480                      Midodrine
-	D02.092.063.291.503                      Nebivolol
-	D02.092.063.291.525                      Octopamine
-	D02.092.063.291.617                      Phenylephrine
-	D02.092.063.291.617.300                      Etilefrine
-	D02.092.063.291.647                      Procaterol
-	D02.092.063.291.805                      Sotalol
-	D02.092.063.291.870                      Synephrine
-	D02.092.063.291.905                      Terbutaline
-	D02.092.063.350                      Heptaminol
-	D02.092.063.400                      Isoetharine
-	D02.092.063.480                      Norepinephrine
-	D02.092.063.480.651                      Normetanephine
-	D02.092.063.624                      Propanolamines
-	D02.092.063.624.302                      Ephedrine
-	D02.092.063.624.380                      Histidinol
-	D02.092.063.624.536                      Methoxamine
-	D02.092.063.624.698                      Phenoxypropanolamines
-	D02.092.063.624.698.025                      Acebutolol
-	D02.092.063.624.698.055                      Alprenolol
-	D02.092.063.624.698.055.200                      Dihydroalprenolol



## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.092.063.624.698.070                      Atenolol
-	D02.092.063.624.698.077                      Betaxolol
-	D02.092.063.624.698.085                      Bisoprolol
-	D02.092.063.624.698.115                      Bunolol
-	D02.092.063.624.698.115.500                      Levobunolol
-	D02.092.063.624.698.146                      Bupranolol
-	D02.092.063.624.698.207                      Carteolol
-	D02.092.063.624.698.268                      Celiprolol
-	D02.092.063.624.698.542                      Metipranolol
-	D02.092.063.624.698.573                      Metoprolol
-	D02.092.063.624.698.601                      Nadolol
-	D02.092.063.624.698.633                      Oxprenolol
-	D02.092.063.624.698.688                      Penbutolol
-	D02.092.063.624.698.699                      Pindolol
-	D02.092.063.624.698.699.420                      Iodocyanopindolol
-	D02.092.063.624.698.705                      Practolol
-	D02.092.063.624.698.708                      Prenalterol
-	D02.092.063.624.698.711                      Propranolol
-	D02.092.063.624.698.855                      Xamoterol
-	D02.092.063.624.706                      Phenylpropanolamine
-	D02.092.063.624.706.400                      p-Hydroxynorephedrine
-	D02.092.063.624.706.500                      Metaraminol
-	D02.092.063.624.706.550                      Nyldrin
-	D02.092.063.624.706.600                      Oxyfedrine
-	D02.092.063.624.706.800                      Tolterodine Tartrate
-	D02.092.063.624.788                      Pseudoephedrine
-	D02.092.063.624.870                      Ritodrine
-	D02.092.063.624.890                      Suloctidil
-	D02.092.063.624.915                      Timolol
-	D02.092.063.624.915.500                      Brimonidine Tartrate, Timolol Maleate Drug Combination
-	D02.092.063.700                      Sphingosine
-	D02.092.063.700.350                      Fingolimod Hydrochloride
-	D02.092.063.700.700                      Psychosine
-	D02.092.080                      Aminopyridines
-	D02.092.080.060                      4-Aminopyridine

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.092.080.085 Amrinone
-	D02.092.080.085.543 Milrinone
-	D02.092.146 Aniline Compounds
-	D02.092.146.100 Aminophenols
-	D02.092.146.100.700 Phenetidine
New Tree	<a href="#">D02.092.146.113</a> Anilides
New Tree	<a href="#">D02.092.146.113.092</a> Acetanilides
New Tree	<a href="#">D02.092.146.113.092.040</a> Acetaminophen
New Tree	<a href="#">D02.092.146.113.092.250</a> Diamfenetide
New Tree	<a href="#">D02.092.146.113.092.325</a> Etidocaine
New Tree	<a href="#">D02.092.146.113.092.450</a> Inosine Pranobex
New Tree	<a href="#">D02.092.146.113.092.500</a> Lidocaine
New Tree	<a href="#">D02.092.146.113.092.700</a> Phenacetin
New Tree	<a href="#">D02.092.146.113.092.800</a> Practolol
New Tree	<a href="#">D02.092.146.113.092.850</a> Ranolazine
New Tree	<a href="#">D02.092.146.113.092.900</a> Trimecaine
New Tree	<a href="#">D02.092.146.113.200</a> Benzoylarginine Nitroanilide
New Tree	<a href="#">D02.092.146.113.239</a> Bupivacaine
New Tree	<a href="#">D02.092.146.113.326</a> Carbanilides
New Tree	<a href="#">D02.092.146.113.326.400</a> Imidocarb
New Tree	<a href="#">D02.092.146.113.326.550</a> Nicarbazin
New Tree	<a href="#">D02.092.146.113.350</a> Carboxin
New Tree	<a href="#">D02.092.146.113.385</a> Encainide
New Tree	<a href="#">D02.092.146.113.420</a> Flutamide

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	<a href="#">D02.092.146.113.750</a>	<a href="#">Prilocaine</a>
New Tree	<a href="#">D02.092.146.113.789</a>	<a href="#">Propanil</a>
New Tree	<a href="#">D02.092.146.113.860</a>	<a href="#">Salicylanilides</a>
New Tree	<a href="#">D02.092.146.113.860.470</a>	<a href="#">Niclosamide</a>
New Tree	<a href="#">D02.092.146.113.860.500</a>	<a href="#">Oxyclozanide</a>
New Tree	<a href="#">D02.092.146.113.860.600</a>	<a href="#">Rafoxanide</a>
New Tree	<a href="#">D02.092.146.113.920</a>	<a href="#">Tocainide</a>
-	D02.092.146.125	Anilino Naphthalenesulfonates
-	D02.092.146.215	Benzophenoneidum
-	D02.092.146.271	Bromhexine
-	D02.092.146.271.100	Ambroxol
-	D02.092.146.290 dimethyl-N-2-propenyl-), Dibromide	Benzenaminium, 4,4'-(3-oxo-1,5-pentanediy)bis(N,N-
-	D02.092.146.307	Fenamates
-	D02.092.146.316	p-Aminoazobenzene
-	D02.092.146.325	p-Dimethylaminoazobenzene
-	D02.092.146.325.500	Methyldimethylaminoazobenzene
-	D02.092.146.350	Diphenylamine
-	D02.092.146.400	Gentian Violet
-	D02.092.146.475	Methyl Green
-	D02.092.146.500	Methylenebis(chloroaniline)
-	D02.092.146.651	Phenylenediamines
-	D02.092.146.651.900	Tetramethylphenylenediamine
-	D02.092.146.755	Rosaniline Dyes
-	D02.092.146.807	Sulfanilamides
-	D02.092.146.807.300	Furosemide
-	D02.092.146.807.650	Sulfacetamide
-	D02.092.146.807.700	Sulfachlorpyridazine
-	D02.092.146.807.755	Sulfadiazine
-	D02.092.146.807.755.800	Silver Sulfadiazine
-	D02.092.146.807.760	Sulfadimethoxine
-	D02.092.146.807.765	Sulfadoxine

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.092.146.807.810 Sulfaguanidine
-	D02.092.146.807.850 Sulfalene
-	D02.092.146.807.853 Sulfamerazine
-	D02.092.146.807.857 Sulfameter
-	D02.092.146.807.862 Sulfamethazine
-	D02.092.146.807.867 Sulfamethoxazole
-	D02.092.146.807.867.500 Trimethoprim, Sulfamethoxazole Drug Combination
-	D02.092.146.807.872 Sulfamethoxypyridazine
-	D02.092.146.807.877 Sulfamonomethoxine
-	D02.092.146.807.882 Sulfamoxole
-	D02.092.146.807.887 Sulfaphenazole
-	D02.092.146.807.900 Sulfapyridine
-	D02.092.146.807.925 Sulfaquinoxaline
-	D02.092.146.807.935 Sulfathiazoles
-	D02.092.146.807.935.813 Sulfamethizole
-	D02.092.146.807.940 Sulfisomidine
-	D02.092.146.807.945 Sulfisoxazole
-	D02.092.146.859 Toluidines
-	D02.092.146.859.717 Trifluralin
-	D02.092.200 Benzylamines
-	D02.092.200.650 Pargyline
-	D02.092.211 Biogenic Amines
-	D02.092.211.111 Acetylcholine
-	D02.092.211.215 Biogenic Monoamines
-	D02.092.211.215.311 Catecholamines
-	D02.092.211.215.311.342 Dopamine
-	D02.092.211.215.311.461 Epinephrine
-	D02.092.211.215.311.461.400 Metanephrine
-	D02.092.211.215.311.461.700 Racepinephrine
-	D02.092.211.215.311.465 Metaproterenol
-	D02.092.211.215.311.465.300 Fenoterol
-	D02.092.211.215.311.560 Norepinephrine
-	D02.092.211.215.311.560.220 Droxidopa
-	D02.092.211.215.311.560.651 Normetanephrine
-	D02.092.211.215.501 Histamine
-	D02.092.211.215.501.621 Methylhistamines

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.092.211.215.581 2-Hydroxyphenethylamine
-	D02.092.211.215.801 Tryptamines
-	D02.092.211.215.801.451 N,N-Dimethyltryptamine
-	D02.092.211.215.801.451.601 Methoxydimethyltryptamines
-	D02.092.211.215.801.852 Serotonin
-	D02.092.211.215.801.852.611 5-Methoxytryptamine
-	D02.092.211.215.811 Tyramine
-	D02.092.211.215.811.651 Octopamine
-	D02.092.211.215.811.875 Synephrine
-	D02.092.211.415 Biogenic Polyamines
-	D02.092.211.415.261 Cadaverine
-	D02.092.211.415.701 Putrescine
-	D02.092.211.415.701.801 Spermidine
-	D02.092.211.415.701.801.821 Spermine
-	D02.092.264 Butylamines
-	D02.092.311 Catecholamines
-	D02.092.311.200 Dihydroxyphenylalanine
-	D02.092.311.200.180 Cysteinyldopa
-	D02.092.311.200.480 Levodopa
-	D02.092.311.200.538 Methylidopa
-	D02.092.311.200.538.200 Carbidopa
-	D02.092.311.220 Dobutamine
-	D02.092.311.342 Dopamine
-	D02.092.311.342.300 Deoxyepinephrine
-	D02.092.311.342.478 Hydroxydopamines
-	D02.092.311.342.478.650 Oxidopamine
-	D02.092.311.461 Epinephrine
-	D02.092.311.461.200 Deoxyepinephrine
-	D02.092.311.461.400 Metanephrine
-	D02.092.311.461.484 Norepinephrine
-	D02.092.311.461.484.220 Droxidopa
-	D02.092.311.461.484.400 Nordefrin
-	D02.092.311.461.651 Normetanephrine
-	D02.092.311.461.825 Racepinephrine
-	D02.092.311.649 Isoproterenol
-	D02.092.311.660 Metaproterenol

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.092.311.660.300 Fenoterol
New Tree	<a href="#">D02.092.348</a> Chloramines
-	D02.092.384 Cyclohexylamines
-	D02.092.384.175 Bromhexine
-	D02.092.384.175.100 Ambroxol
-	D02.092.471 Ethylamines
-	D02.092.471.200 Cystamine
-	D02.092.471.245 Dibenzylchloroethamine
-	D02.092.471.302 Diethylamines
-	D02.092.471.302.500 Propanidid
-	D02.092.471.308 Diethylpropion
-	D02.092.471.320 Diphenhydramine
-	D02.092.471.320.250 Dimenhydrinate
-	D02.092.471.440 Histamine
-	D02.092.471.440.500 Methylhistamines
-	D02.092.471.562 Mercaptoethylamines
-	D02.092.471.562.369 Cysteamine
-	D02.092.471.600 Orphenadrine
-	D02.092.471.683 Phenethylamines
-	D02.092.471.683.061 Albuterol
-	D02.092.471.683.061.125 Albuterol, Ipratropium Drug Combination
-	D02.092.471.683.061.500 Levalbuterol
-	D02.092.471.683.061.750 Salmeterol Xinafoate
-	D02.092.471.683.061.750.500 Fluticasone Propionate, Salmeterol Xinafoate Drug Combination
-	D02.092.471.683.152 Amphetamines
-	D02.092.471.683.152.110 Amphetamine
-	D02.092.471.683.152.110.200 Dextroamphetamine
-	D02.092.471.683.152.110.200.500 Lisdexamfetamine Dimesylate
-	D02.092.471.683.152.200 p-Chloroamphetamine
-	D02.092.471.683.152.391 2,5-Dimethoxy-4-Methylamphetamine
-	D02.092.471.683.152.500 p-Hydroxyamphetamine
-	D02.092.471.683.152.535 lofetamine
-	D02.092.471.683.152.619 Methamphetamine
-	D02.092.471.683.152.619.307 Benzphetamine

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.092.471.683.152.660 3,4-Methylenedioxyamphetamine
-	D02.092.471.683.152.670 N-Methyl-3,4-methylenedioxyamphetamine
-	D02.092.471.683.152.766 Phentermine
-	D02.092.471.683.152.766.262 Chlorphentermine
-	D02.092.471.683.152.766.617 Mephentermine
-	D02.092.471.683.221 Butoxamine
-	D02.092.471.683.338 Dexfenfluramine
-	D02.092.471.683.386 Dimethoxyphenylethylamine
-	D02.092.471.683.410 Dobutamine
-	D02.092.471.683.430 Ephedrine
-	D02.092.471.683.450 Fendiline
-	D02.092.471.683.467 Fenfluramine
-	D02.092.471.683.467.500 Norfenfluramine
-	D02.092.471.683.530 Hexoprenaline
-	D02.092.471.683.545 2-Hydroxyphenethylamine
-	D02.092.471.683.560 Isoxsuprine
-	D02.092.471.683.635 Mescaline
-	D02.092.471.683.661 Methoxamine
-	D02.092.471.683.670 p-Methoxy-N-methylphenethylamine
-	D02.092.471.683.715 Nylidrin
-	D02.092.471.683.725 Octopamine
-	D02.092.471.683.775 Oxyfedrine
-	D02.092.471.683.825 Prenylamine
-	D02.092.471.683.847 Pseudoephedrine
-	D02.092.471.683.870 Ritodrine
-	D02.092.471.683.915 Selegiline
-	D02.092.471.683.934 Suloctidil
-	D02.092.471.683.943 Tiapamil Hydrochloride
-	D02.092.471.683.948 Venlafaxine Hydrochloride
-	D02.092.471.683.953 Verapamil
-	D02.092.471.683.953.395 Gallopamil
-	D02.092.471.739 Phenoxybenzamine
-	D02.092.471.869 Tiapride Hydrochloride
-	D02.092.570 Hydroxylamines
-	D02.092.570.050 Aminoxyacetic Acid
-	D02.092.570.394 Hydroxamic Acids

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.092.570.394.150                      Bufexamac
-	D02.092.570.394.265                      Deferoxamine
-	D02.092.570.394.380                      Ferrichrome
-	D02.092.570.450                          4-Hydroxyaminoquinoline-1-oxide
-	D02.092.570.665                          Oximes
-	D02.092.570.665.250                      Fluvoxamine
-	D02.092.570.665.500                      Obidoxime Chloride
-	D02.092.570.665.535                      Pralidoxime Compounds
-	D02.092.570.665.810                      Technetium Tc 99m Exametazime
-	D02.092.570.665.850                      Trimedoxime
-	D02.092.650                                  Mannich Bases
-	D02.092.668                                  Methylamines
-	D02.092.668.387                              Dimethylamines
-	D02.092.668.387.500                          Moxisylyte
-	D02.092.668.387.750                          Tramadol
-	D02.092.705                                  1-Naphthylamine
-	D02.092.705.800                              Sertraline
-	D02.092.710                                  2-Naphthylamine
-	D02.092.782                                  Polyamines
-	D02.092.782.200                              Colestipol
-	D02.092.782.258                              Diamines
-	D02.092.782.258.174                          Cadaverine
-	D02.092.782.258.368                          Ethylenediamines
-	D02.092.782.258.368.050                      Aminophylline
-	D02.092.782.258.368.250                      Edetic Acid
-	D02.092.782.258.368.257                      Egtazic Acid
-	D02.092.782.258.368.265                      Ethambutol
-	D02.092.782.258.368.500                      Methapyrilene
-	D02.092.782.258.368.880                      Trientine
-	D02.092.782.258.368.900                      Tripelennamine
-	D02.092.782.258.651                          Phenylenediamines
-	D02.092.782.258.651.900                      Tetramethylphenylenediamine
-	D02.092.782.258.784                          Putrescine
-	D02.092.782.420                              Hexadimethrine Bromide
-	D02.092.782.500                              Methenamine
-	D02.092.782.590                              Pentetic Acid



## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.092.782.590.401 Gadolinium DTPA
-	D02.092.782.590.883 Technetium Tc 99m Pentetate
-	D02.092.782.634 Sevelamer
-	D02.092.782.677 Spermidine
-	D02.092.782.802 Spermine
-	D02.092.831 Propylamines
-	D02.092.831.085 Atomoxetine Hydrochloride
-	D02.092.831.170 Citalopram
-	D02.092.831.180 Clorgyline
-	D02.092.831.280 Fluoxetine
-	D02.092.831.500 Mexiletine
-	D02.092.831.690 Promethazine
-	D02.092.831.845 Tranylcypromine
-	D02.092.877 Quaternary Ammonium Compounds
-	D02.092.877.096 Benzylammonium Compounds
-	D02.092.877.096.019 Ambenonium Chloride
-	D02.092.877.096.040 Benzalkonium Compounds
-	D02.092.877.096.082 Benzethonium
-	D02.092.877.096.166 Bephenium Compounds
-	D02.092.877.096.333 Breylium Compounds
-	D02.092.877.096.333.150 Breylium Tosylate
-	D02.092.877.233 Betalains
-	D02.092.877.233.500 Betacyanins
-	D02.092.877.233.555 Betaxanthins
-	D02.092.877.250 Bis-Trimethylammonium Compounds
-	D02.092.877.250.372 Decamethonium Compounds
-	D02.092.877.250.592 Hexamethonium Compounds
-	D02.092.877.250.592.400 Hexamethonium
-	D02.092.877.310 Butylscopolammonium Bromide
-	D02.092.877.370 Emepronium
-	D02.092.877.400 Gallamine Triethiodide
-	D02.092.877.425 Glycopyrrolate
-	D02.092.877.435 Hemicholinium 3
-	D02.092.877.475 Lissamine Green Dyes
-	D02.092.877.648 Oxyphenonium
-	D02.092.877.674 Phenylammonium Compounds

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.092.877.674.033 butynyl)trimethylammonium Chloride (4-(m-Chlorophenylcarbamoxy)-2-
-	D02.092.877.674.333 Edrophonium
-	D02.092.877.674.666 Neostigmine
-	D02.092.877.700 Propantheline
-	D02.092.877.787 Tetraethylammonium Compounds
-	D02.092.877.787.500 Tetraethylammonium
-	D02.092.877.844 Toxiferine
-	D02.092.877.844.050 Alcuronium
-	D02.092.877.883 Trimethyl Ammonium Compounds
-	D02.092.877.883.077 Betaine
-	D02.092.877.883.088 Bethanechol Compounds
-	D02.092.877.883.088.100 Bethanechol
-	D02.092.877.883.099 Carnitine
-	D02.092.877.883.099.090 Acetylcarnitine
-	D02.092.877.883.099.700 Palmitoylcarnitine
-	D02.092.877.883.111 Cetrimonium Compounds
-	D02.092.877.883.222 Chlorisondamine
-	D02.092.877.883.277 Chlormequat
-	D02.092.877.883.333 Choline
-	D02.092.877.883.333.100 Benzoylcholine
-	D02.092.877.883.333.115 Carbachol
-	D02.092.877.883.333.130 Cytidine Diphosphate Choline
-	D02.092.877.883.333.700 Phosphorylcholine
-	D02.092.877.883.333.710 Platelet Activating Factor
-	D02.092.877.883.333.720 Propylbenzilylcholine Mustard
-	D02.092.877.883.333.780 Succinylcholine
-	D02.092.877.883.333.800 Thiocholine
-	D02.092.877.883.333.800.030 Acetylthiocholine
-	D02.092.877.883.333.800.200 Butyrylthiocholine
-	D02.092.877.883.555 Methacholine Compounds
-	D02.092.877.883.555.500 Methacholine Chloride
-	D02.092.877.883.777 Muscarine
-	D02.092.877.922 Tubocurarine
-	D02.113 Anhydrides
-	D02.113.075 Acetic Anhydrides

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.113.250 Citraconic Anhydrides
-	D02.113.450 Maleic Anhydrides
-	D02.113.600 Phthalic Anhydrides
-	D02.113.650 Polyanhydrides
-	D02.113.700 Succinic Anhydrides
-	D02.145 Aza Compounds
-	D02.145.074 Azabicyclo Compounds
-	D02.145.074.444 Granisetron
-	D02.145.074.722 Tropanes
-	D02.145.074.722.229 Atropine Derivatives
-	D02.145.074.722.229.199 Atropine
-	D02.145.074.722.229.199.500 Hyoscyamine
-	D02.145.074.722.229.400 Ipratropium
-	D02.145.074.722.229.400.500 Albuterol, Ipratropium Drug Combination
-	D02.145.074.722.270 Benztropine
-	D02.145.074.722.388 Cocaine
-	D02.145.074.722.388.250 Crack Cocaine
-	D02.145.074.722.744 Nortropanes
-	D02.145.074.722.822 Scopolamine Derivatives
-	D02.145.074.722.822.200 Butylscopolammonium Bromide
-	D02.145.074.722.822.550 N-Methylscopolamine
-	D02.145.074.722.822.775 Scopolamine Hydrobromide
-	D02.145.074.722.822.887 Tiotropium Bromide
-	D02.145.150 Azacitidine
-	D02.145.250 Azaguanine
-	D02.145.451 Azauridine
-	D02.159 Azides
-	D02.172 Azo Compounds
-	D02.172.025 Amarant Dye
-	D02.172.029 Amido Black
-	D02.172.030 o-Aminoazotoluene
-	D02.172.032 p-Aminoazobenzene
-	D02.172.040 Arsenazo III
-	D02.172.045 p-Azobenzene arsonate
-	D02.172.080 Azoxymethane
-	D02.172.080.200 Cycasin

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.172.080.600 Methylazoxymethanol Acetate
-	D02.172.300 Diamide
-	D02.172.383 Diazonium Compounds
-	D02.172.383.383 Diazomethane
-	D02.172.508 Diazooxonorleucine
-	D02.172.530 Dithizone
-	D02.172.560 Evans Blue
-	D02.172.600 Formazans
-	D02.172.950 Tartrazine
-	D02.172.975 Trypan Blue
-	D02.203 Boron Compounds
-	D02.203.087 Boranes
-	D02.203.087.050 Borohydrides
-	D02.203.087.050.750 Tetraphenylborate
-	D02.203.130 Boric Acids
-	D02.203.130.075 Borates
-	D02.203.187 Borinic Acids
-	D02.203.200 Boronic Acids
-	D02.203.200.500 Bortezomib
-	D02.241 Carboxylic Acids
-	D02.241.081 Acids, Acyclic
-	D02.241.081.018 Acetates
-	D02.241.081.018.110 Acetamides
-	D02.241.081.018.110.080 2-Acetylaminofluorene
-	D02.241.081.018.110.080.070 Acetoxyacetylaminofluorene
-	D02.241.081.018.110.080.400 Hydroxyacetylaminofluorene
-	D02.241.081.018.110.189 Allylisopropylacetamide
-	D02.241.081.018.110.400 Iodoacetamide
-	D02.241.081.018.110.525 Linezolid
-	D02.241.081.018.110.650 Piracetam
-	D02.241.081.018.110.786 Thioacetamide
-	D02.241.081.018.165 Acetic Acid
-	D02.241.081.018.165.249 Potassium Acetate
-	D02.241.081.018.165.500 Sodium Acetate
-	D02.241.081.018.165.750 Zinc Acetate
-	D02.241.081.018.193 Acetic Anhydrides

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.241.081.018.207 Aminoxyacetic Acid
-	D02.241.081.018.214 Chloroacetates
-	D02.241.081.018.214.500 Dichloroacetic Acid
-	D02.241.081.018.214.750 Trichloroacetic Acid
-	D02.241.081.018.253 Edetic Acid
-	D02.241.081.018.269 Egtazic Acid
-	D02.241.081.018.285 Fluoroacetates
-	D02.241.081.018.285.500 Trifluoroacetic Acid
-	D02.241.081.018.386 Glycolates
-	D02.241.081.018.386.682 Phenoxyacetates
-	D02.241.081.018.386.682.224 2,4-Dichlorophenoxyacetic Acid
-	D02.241.081.018.386.682.300 Ethacrynic Acid
-	D02.241.081.018.386.682.500 2-Methyl-4-chlorophenoxyacetic Acid
-	D02.241.081.018.386.682.800 2,4,5-Trichlorophenoxyacetic Acid
-	D02.241.081.018.386.682.850 Halofenate
-	D02.241.081.018.386.682.875 Meclofenoxate
-	D02.241.081.018.386.682.900 Ticrynafen
-	D02.241.081.018.487 Iodoacetates
-	D02.241.081.018.487.249 Iodoacetamide
-	D02.241.081.018.487.500 Iodoacetic Acid
-	D02.241.081.018.588 Nitrilotriacetic Acid
-	D02.241.081.018.639 Pentetic Acid
-	D02.241.081.018.639.400 Gadolinium DTPA
-	D02.241.081.018.639.883 Technetium Tc 99m Pentetate
-	D02.241.081.018.664 Peracetic Acid
-	D02.241.081.018.677 Phosphonoacetic Acid
-	D02.241.081.018.677.250 Foscarnet
-	D02.241.081.018.900 Thioglycolates
-	D02.241.081.069 Acrylates
-	D02.241.081.069.094 Acrylamides
-	D02.241.081.069.094.015 Acrylamide
-	D02.241.081.069.366 Cyanoacrylates
-	D02.241.081.069.366.200 Bucrylate
-	D02.241.081.069.366.350 Enbucrilate
-	D02.241.081.069.600 Methacrylates
-	D02.241.081.069.600.150 Bisphenol A-Glycidyl Methacrylate

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.241.081.069.800 Polymethacrylic Acids
-	D02.241.081.069.800.550 Methylmethacrylates
-	D02.241.081.069.800.550.450 Methylmethacrylate
-	D02.241.081.069.800.550.500 Polymethyl Methacrylate
-	D02.241.081.069.800.700 Polyhydroxyethyl Methacrylate
-	D02.241.081.069.920 Urocanic Acid
-	D02.241.081.114 Butyrates
-	D02.241.081.114.500 Aminobutyrate
-	D02.241.081.114.500.350 gamma-Aminobutyric Acid
-	D02.241.081.114.500.350.100 Baclofen
-	D02.241.081.114.500.350.500 Pregabalin
-	D02.241.081.114.500.350.900 Vigabatrin
-	D02.241.081.114.625 Bendamustine Hydrochloride
-	D02.241.081.114.750 Butyric Acid
-	D02.241.081.114.875 Crotonates
-	D02.241.081.114.937 Hydroxybutyrates
-	D02.241.081.114.937.349 3-Hydroxybutyric Acid
-	D02.241.081.114.937.700 Sodium Oxybate
-	D02.241.081.114.968 Isobutyrate
-	D02.241.081.114.968.249 Aminoisobutyric Acids
-	D02.241.081.114.968.500 Fibrates
-	D02.241.081.114.968.500.249 Bezafibrate
-	D02.241.081.114.968.500.374 Clofenapate
-	D02.241.081.114.968.500.500 Clofibrate
-	D02.241.081.114.968.500.500.195 Clofibrate
-	D02.241.081.114.968.500.625 Fenofibrate
-	D02.241.081.114.968.500.750 Gemfibrozil
-	D02.241.081.193 Caproates
-	D02.241.081.193.150 Aminocaproates
-	D02.241.081.193.150.075 Aminocaproic Acid
-	D02.241.081.193.678 Mycophenolic Acid
-	D02.241.081.193.839 Penicillic Acid
-	D02.241.081.222 Caprylates
-	D02.241.081.251 Carbamates
-	D02.241.081.251.022 Albendazole
-	D02.241.081.251.046 Aldicarb

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D02.241.081.251.125	Benomyl
New Tree	<a href="#">D02.241.081.251.133</a>	<a href="#">Bethanechol Compounds</a>
New Tree	<a href="#">D02.241.081.251.133.100</a>	<a href="#">Bethanechol</a>
-	D02.241.081.251.140	Carbadox
-	D02.241.081.251.145	Carbamyl Phosphate
-	D02.241.081.251.150	Carbaryl
-	D02.241.081.251.165	Carisoprodol
-	D02.241.081.251.203	Cobicistat
-	D02.241.081.251.203.500 Disoproxil Fumarate Drug Combination	Elvitegravir, Cobicistat, Emtricitabine, Tenofovir
-	D02.241.081.251.222	Darunavir
-	D02.241.081.251.240	Diethylcarbamazine
-	D02.241.081.251.320	Fenbendazole
-	D02.241.081.251.415	Mebendazole
-	D02.241.081.251.510	Meprobamate
-	D02.241.081.251.525	Methomyl
-	D02.241.081.251.583	Phenylcarbamates
-	D02.241.081.251.583.166	Carbofuran
-	D02.241.081.251.583.177	Chlorpropham
-	D02.241.081.251.583.555	Methiocarb
-	D02.241.081.251.583.577	Methocarbamol
-	D02.241.081.251.583.682	Physostigmine
-	D02.241.081.251.583.788	Propoxur
-	D02.241.081.251.583.799	Pyridinolcarbamate
-	D02.241.081.251.583.899	Rivastigmine
-	D02.241.081.251.869	Thiocarbamates
-	D02.241.081.251.869.210	Dimethyldithiocarbamate
-	D02.241.081.251.869.210.800	Thiram
-	D02.241.081.251.869.210.925	Ziram
-	D02.241.081.251.869.220	Ditiocarb
-	D02.241.081.251.869.220.250	Disulfiram
-	D02.241.081.251.869.265	Ethylenebis(dithiocarbamates)
-	D02.241.081.251.869.265.441	Maneb
-	D02.241.081.251.869.265.908	Zineb
-	D02.241.081.251.869.791	Tolnaftate

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.241.081.251.869.820                      Triallate
-	D02.241.081.251.890                            Thiophanate
-	D02.241.081.251.944                            Urethane
-	D02.241.081.251.944.500                      Nitrosomethylurethane
-	D02.241.081.251.944.750                      Polyurethanes
-	D02.241.081.337                                Dicarboxylic Acids
-	D02.241.081.337.052                            Adipates
-	D02.241.081.337.052.075                      2-Aminoadipic Acid
-	D02.241.081.337.052.250                      Dimethyl Adipimidate
-	D02.241.081.337.075                            Bongkreic Acid
-	D02.241.081.337.302                            Fumarates
-	D02.241.081.337.302.500                      Dimethyl Fumarate
-	D02.241.081.337.351                            Glutarates
-	D02.241.081.337.351.300                      Formiminoglutamic Acid
-	D02.241.081.337.351.502                      Ketoglutaric Acids
-	D02.241.081.337.351.550                      Meglutol
-	D02.241.081.337.463                            Malates
-	D02.241.081.337.463.703                      Thiomalates
-	D02.241.081.337.463.703.451                      Gold Sodium Thiomalate
-	D02.241.081.337.502                            Maleates
-	D02.241.081.337.502.524                      Maleimides
-	D02.241.081.337.502.524.418                      Ethylmaleimide
-	D02.241.081.337.540                            Malonates
-	D02.241.081.337.540.500                      Methylmalonic Acid
-	D02.241.081.337.540.700                      Phenylethylmalonamide
-	D02.241.081.337.593                            Oxalates
-	D02.241.081.337.593.750                      Oxalic Acid
-	D02.241.081.337.593.750.500                      Calcium Oxalate
-	D02.241.081.337.593.812                      Oxaloacetates
-	D02.241.081.337.593.812.750                      Oxaloacetic Acid
-	D02.241.081.337.699                            Pimelic Acids
-	D02.241.081.337.699.250                      Diaminopimelic Acid
-	D02.241.081.337.759                            Succinates
-	D02.241.081.337.759.124                      Argininosuccinic Acid
-	D02.241.081.337.759.249                      Dioctyl Sulfosuccinic Acid
-	D02.241.081.337.759.500                      Succimer



## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.241.081.337.759.500.725                                      Technetium Tc 99m Dimercaptosuccinic Acid
-	D02.241.081.337.759.625                                      Succinic Acid
-	D02.241.081.337.759.750                                      Succinic Anhydrides
-	D02.241.081.337.759.875                                      Succinylcholine
-	D02.241.081.337.864                                      Tartrates
-	D02.241.081.337.902                                      Tartronates
-	D02.241.081.420                                      Formates
-	D02.241.081.420.500                                      Formamides
-	D02.241.081.420.500.387                                      Dimethylformamide
-	D02.241.081.420.750                                      Formic Acid Esters
-	D02.241.081.420.750.250                                      Diethyl Pyrocarbonate
-	D02.241.081.583                                      Imino Acids
-	D02.241.081.583.100                                      Azetidinecarboxylic Acid
-	D02.241.081.583.400                                      Technetium Tc 99m Diethyl-iminodiacetic Acid
-	D02.241.081.583.450                                      Technetium Tc 99m Disofenin
-	D02.241.081.583.900                                      Technetium Tc 99m Lidofenin
-	D02.241.081.751                                      Propionates
-	D02.241.081.751.080                                      3-Mercaptopropionic Acid
-	D02.241.081.751.120                                      Dextropropoxyphene
-	D02.241.081.751.161                                      Flurbiprofen
-	D02.241.081.751.324                                      Levopropoxyphene
-	D02.241.081.751.512                                      Nafenopin
-	D02.241.081.844                                      Sugar Acids
-	D02.241.081.844.107                                      Ascorbic Acid
-	D02.241.081.844.107.260                                      Dehydroascorbic Acid
-	D02.241.081.844.200                                      2,3-Diketogulonic Acid
-	D02.241.081.844.300                                      Glucaric Acid
-	D02.241.081.844.322                                      Gluconates
-	D02.241.081.844.322.060                                      Antimony Sodium Gluconate
-	D02.241.081.844.322.200                                      Calcium Gluconate
-	D02.241.081.844.387                                      Glyceric Acids
-	D02.241.081.844.387.388                                      Diphosphoglyceric Acids
-	D02.241.081.844.387.388.175                                      2,3-Diphosphoglycerate
-	D02.241.081.844.520                                      Muramic Acids
-	D02.241.081.844.562                                      Neuraminic Acids
-	D02.241.081.844.562.668                                      Sialic Acids

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.241.081.844.562.668.050 N-Acetylneuraminic Acid
-	D02.241.081.844.562.668.250 Cytidine Monophosphate N-Acetylneuraminic Acid
-	D02.241.081.844.562.668.775 Zanamivir
-	D02.241.081.844.759 Tartrates
-	D02.241.081.844.821 Tartronates
-	D02.241.081.844.915 Uronic Acids
-	D02.241.081.844.915.162 Glucuronates
-	D02.241.081.844.915.162.249 Glucuronic Acid
-	D02.241.081.844.915.162.500 Glucuronides
-	D02.241.081.844.915.400 Hexuronic Acids
-	D02.241.081.844.915.400.500 Iduronic Acid
-	D02.241.081.901 Tricarboxylic Acids
-	D02.241.081.901.177 Aconitic Acid
-	D02.241.081.901.400 1-Carboxyglutamic Acid
-	D02.241.081.901.434 Citrates
-	D02.241.081.901.434.249 Citric Acid
-	D02.241.081.901.434.249.500 Calcium Citrate
-	D02.241.081.901.434.249.750 Potassium Citrate
-	D02.241.081.901.434.498 Isocitrates
-	D02.241.081.944 Valerates
-	D02.241.081.944.509 Pentanoic Acids
-	D02.241.081.944.509.350 Gemfibrozil
-	D02.241.081.944.509.900 Valproic Acid
-	D02.241.081.944.550 Proadifen
-	D02.241.152 Acids, Aldehydic
-	D02.241.152.367 Glyoxylates
-	D02.241.152.367.600 Oxamic Acid
-	D02.241.152.811 Uronic Acids
-	D02.241.152.811.162 Glucuronates
-	D02.241.152.811.162.500 Glucuronic Acid
-	D02.241.152.811.162.750 Glucuronides
-	D02.241.152.811.400 Hexuronic Acids
-	D02.241.152.811.400.500 Iduronic Acid
-	D02.241.223 Acids, Carbocyclic
-	D02.241.223.100 Benzoates

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.241.223.100.050                      Aminobenzoates
-	D02.241.223.100.050.300                      meta-Aminobenzoates
-	D02.241.223.100.050.300.100                      Acetrizic Acid
-	D02.241.223.100.050.300.200                      Bumetanide
-	D02.241.223.100.050.300.500                      Mesalamine
-	D02.241.223.100.050.400                      ortho-Aminobenzoates
-	D02.241.223.100.050.400.200                      Fenamates
-	D02.241.223.100.050.400.200.249                      Flufenamic Acid
-	D02.241.223.100.050.400.200.500                      Meclofenamic Acid
-	D02.241.223.100.050.400.200.555                      Mefenamic Acid
-	D02.241.223.100.050.400.200.777                      Niflumic Acid
-	D02.241.223.100.050.400.300                      Glafenine
-	D02.241.223.100.050.400.400                      3-Hydroxyanthranilic Acid
-	D02.241.223.100.050.500                      para-Aminobenzoates
-	D02.241.223.100.050.500.625                      Acecainide
-	D02.241.223.100.050.500.640                      4-Aminobenzoic Acid
-	D02.241.223.100.050.500.642                      Aminosalicylic Acid
-	D02.241.223.100.050.500.645                      Ethopabate
-	D02.241.223.100.050.500.647                      Metoclopramide
-	D02.241.223.100.050.500.650                      p-Aminohippuric Acid
-	D02.241.223.100.050.500.687                      Benzocaine
-	D02.241.223.100.050.500.700                      Cisapride
-	D02.241.223.100.050.500.875                      Procainamide
-	D02.241.223.100.050.500.906                      Procaine
-	D02.241.223.100.050.500.906.666                      Penicillin G Procaine
-	D02.241.223.100.050.500.937                      Propoxycaine
-	D02.241.223.100.050.500.968                      Tetracaine
-	D02.241.223.100.100                      Benzamides
-	D02.241.223.100.100.033                      Acecainide
-	D02.241.223.100.100.100                      Aminohippuric Acids
-	D02.241.223.100.100.100.100                      p-Aminohippuric Acid
-	D02.241.223.100.100.120                      Bezafibrate
-	D02.241.223.100.100.135                      Cisapride
-	D02.241.223.100.100.230                      DEET
-	D02.241.223.100.100.240                      Dinitolmide
-	D02.241.223.100.100.400                      Hippurates

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.241.223.100.100.400.500 Iodohippuric Acid
-	D02.241.223.100.100.435 Imatinib Mesylate
-	D02.241.223.100.100.470 Indoramin
-	D02.241.223.100.100.510 Metoclopramide
-	D02.241.223.100.100.600 Moclobemide
-	D02.241.223.100.100.650 Procainamide
-	D02.241.223.100.100.655 Procarbazine
-	D02.241.223.100.100.761 Raclopride
-	D02.241.223.100.100.796 Remoxipride
-	D02.241.223.100.100.866 Sulpiride
-	D02.241.223.100.100.933 Tiapride Hydrochloride
-	D02.241.223.100.120 Benzoic Acid
-	D02.241.223.100.120.500 Sodium Benzoate
-	D02.241.223.100.133 Benzoyl Peroxide
-	D02.241.223.100.133.500 Adapalene, Benzoyl Peroxide Drug Combination
-	D02.241.223.100.141 Benzoylcholine
-	D02.241.223.100.150 Bromobenzoates
-	D02.241.223.100.150.500 Remoxipride
-	D02.241.223.100.200 Chlorobenzoates
-	D02.241.223.100.200.249 Bezafibrate
-	D02.241.223.100.200.311 Chloromercuribenzoates
-	D02.241.223.100.200.311.275 p-Chloromercuribenzoic Acid
-	D02.241.223.100.200.374 Cisapride
-	D02.241.223.100.200.500 Dicamba
-	D02.241.223.100.200.750 Metoclopramide
-	D02.241.223.100.200.875 Moclobemide
-	D02.241.223.100.200.937 Raclopride
-	D02.241.223.100.300 Hydroxybenzoates
-	D02.241.223.100.300.100 Depsides
-	D02.241.223.100.300.200 Gallic Acid
-	D02.241.223.100.300.200.299 Hydrolyzable Tannins
-	D02.241.223.100.300.200.600 Propyl Gallate
-	D02.241.223.100.300.300 3-Hydroxyanthranilic Acid
-	D02.241.223.100.300.350 Hydroxybenzoate Ethers
-	D02.241.223.100.300.350.500 Hexobendine
-	D02.241.223.100.300.350.625 Metoclopramide

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.241.223.100.300.350.687 Propoxycaine
-	D02.241.223.100.300.350.750 Trimebutine
-	D02.241.223.100.300.350.875 Vanillic Acid
-	D02.241.223.100.300.400 Hydroxymercuribenzoates
-	D02.241.223.100.300.450 Pactamycin
-	D02.241.223.100.300.460 Parabens
-	D02.241.223.100.300.595 Salicylates
-	D02.241.223.100.300.595.100 Aminosalicic Acids
-	D02.241.223.100.300.595.100.060 Aminosalicic Acid
-	D02.241.223.100.300.595.100.540 Mesalamine
-	D02.241.223.100.300.595.202 Dicamba
-	D02.241.223.100.300.595.405 Gentisates
-	D02.241.223.100.300.595.608 Salicylic Acid
-	D02.241.223.100.300.595.608.500 Sodium Salicylate
-	D02.241.223.100.400 Iodobenzoates
-	D02.241.223.100.400.880 Triiodobenzoic Acids
-	D02.241.223.100.400.880.020 Acetrizic Acid
-	D02.241.223.100.400.880.270 Diatrizoate
-	D02.241.223.100.400.880.270.500 Diatrizoate Meglumine
-	D02.241.223.100.400.880.375 Iodamide
-	D02.241.223.100.400.880.380 Iodipamide
-	D02.241.223.100.400.880.390 Ioglycamic Acid
-	D02.241.223.100.400.880.400 Iohexol
-	D02.241.223.100.400.880.410 Iopamidol
-	D02.241.223.100.400.880.430 Iothalamic Acid
-	D02.241.223.100.400.880.430.500 Iothalamate Meglumine
-	D02.241.223.100.400.880.450 Ioxaglic Acid
-	D02.241.223.100.400.880.520 Metrizamide
-	D02.241.223.100.400.880.530 Metrizoate
-	D02.241.223.100.500 Mercuribenzoates
-	D02.241.223.100.500.261 Chloromercuribenzoates
-	D02.241.223.100.500.261.275 p-Chloromercuribenzoic Acid
-	D02.241.223.100.500.450 Hydroxymercuribenzoates
-	D02.241.223.100.600 Nitrobenzoates
-	D02.241.223.100.600.300 Dithionitrobenzoic Acid
-	D02.241.223.200 Cinnamates

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.241.223.200.054 Caffeic Acids
-	D02.241.223.200.054.500 Eugenol
-	D02.241.223.200.185 Chlorogenic Acid
-	D02.241.223.200.200 Cinanserin
-	D02.241.223.200.210 Coumaric Acids
-	D02.241.223.200.380 Puromycin
-	D02.241.223.200.380.650 Puromycin Aminonucleoside
-	D02.241.223.268 Cyclohexanecarboxylic Acids
-	D02.241.223.268.034 Abscisic Acid
-	D02.241.223.268.070 Aurintricarboxylic Acid
-	D02.241.223.268.220 Chlorogenic Acid
-	D02.241.223.268.250 Chorismic Acid
-	D02.241.223.268.300 Dicyclomine
-	D02.241.223.268.637 Quinic Acid
-	D02.241.223.268.792 Shikimic Acid
-	D02.241.223.268.825 Tilidine
-	D02.241.223.268.860 Tranexamic Acid
-	D02.241.223.475 Mandelic Acids
-	D02.241.223.475.180 Cyclandelate
-	D02.241.223.475.730 Vanilmandelic Acid
-	D02.241.223.601 Phenylacetates
-	D02.241.223.601.200 Cyclopentolate
-	D02.241.223.601.210 Diclofenac
-	D02.241.223.601.220 3,4-Dihydroxyphenylacetic Acid
-	D02.241.223.601.238 Diphenylacetic Acids
-	D02.241.223.601.238.306 Benzilates
-	D02.241.223.601.238.306.140 Benactyzine
-	D02.241.223.601.238.306.740 Quinuclidinyl Benzilate
-	D02.241.223.601.329 Guanfacine
-	D02.241.223.601.375 Halofenate
-	D02.241.223.601.421 Homogentisic Acid
-	D02.241.223.601.521 Homovanillic Acid
-	D02.241.223.601.600 Methylphenidate
-	D02.241.223.601.600.500 Dexmethylphenidate Hydrochloride
-	D02.241.223.601.774 Propanidid
-	D02.241.223.651 Phenylbutyrates

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.241.223.701 Phenylpropionates
-	D02.241.223.701.350 Fenoprofen
-	D02.241.223.701.430 Ibuprofen
-	D02.241.223.701.450 Indoprofen
-	D02.241.223.701.480 Ketoprofen
-	D02.241.223.701.700 Suprofen
-	D02.241.223.751 Phenylpyruvic Acids
-	D02.241.223.805 Phthalic Acids
-	D02.241.223.805.225 Dibutyl Phthalate
-	D02.241.223.805.250 Diethylhexyl Phthalate
-	D02.241.223.805.780 o-Phthalaldehyde
-	D02.241.223.805.800 Phthalic Anhydrides
-	D02.241.223.805.810 Phthalimides
-	D02.241.223.805.810.800 Thalidomide
-	D02.241.400 Esters
-	D02.241.444 Humic Substances
-	D02.241.511 Hydroxy Acids
-	D02.241.511.085 Benzilates
-	D02.241.511.085.140 Benactyzine
-	D02.241.511.085.740 Quinuclidinyl Benzilate
-	D02.241.511.316 Glycolates
-	D02.241.511.316.455 Halofenate
-	D02.241.511.316.495 Meclofenoxate
-	D02.241.511.316.682 Phenoxyacetates
-	D02.241.511.316.682.149 2,4-Dichlorophenoxyacetic Acid
-	D02.241.511.316.682.200 Ethacrynic Acid
-	D02.241.511.316.682.224 2-Methyl-4-chlorophenoxyacetic Acid
-	D02.241.511.316.682.800 2,4,5-Trichlorophenoxyacetic Acid
-	D02.241.511.316.682.900 Ticrynafen
-	D02.241.511.372 Hydroxamic Acids
-	D02.241.511.372.150 Bufexamac
-	D02.241.511.372.265 Deferoxamine
-	D02.241.511.372.380 Ferrichrome
-	D02.241.511.390 Hydroxybenzoates
-	D02.241.511.390.100 Depsides
-	D02.241.511.390.200 Gallic Acid

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.241.511.390.200.299 Hydrolyzable Tannins
-	D02.241.511.390.200.600 Propyl Gallate
-	D02.241.511.390.300 3-Hydroxyanthranilic Acid
-	D02.241.511.390.350 Hydroxybenzoate Ethers
-	D02.241.511.390.350.500 Hexobendine
-	D02.241.511.390.350.625 Metoclopramide
-	D02.241.511.390.350.687 Propoxycaine
-	D02.241.511.390.350.750 Trimebutine
-	D02.241.511.390.350.875 Vanillic Acid
-	D02.241.511.390.400 Hydroxymercuribenzoates
-	D02.241.511.390.450 Pactamycin
-	D02.241.511.390.460 Parabens
-	D02.241.511.390.595 Salicylates
-	D02.241.511.390.595.100 Aminosalicilic Acids
-	D02.241.511.390.595.100.060 Aminosalicilic Acid
-	D02.241.511.390.595.100.540 Mesalamine
-	D02.241.511.390.595.202 Dicamba
-	D02.241.511.390.595.405 Gentisates
-	D02.241.511.390.595.608 Salicylic Acid
-	D02.241.511.390.595.608.500 Sodium Salicylate
-	D02.241.511.429 Hydroxybutyrates
-	D02.241.511.429.349 3-Hydroxybutyric Acid
-	D02.241.511.429.700 Sodium Oxybate
-	D02.241.511.459 Lactates
-	D02.241.511.459.450 Lactic Acid
-	D02.241.511.459.800 Sodium Lactate
-	D02.241.511.505 Malates
-	D02.241.511.536 Mandelic Acids
-	D02.241.511.536.180 Cyclandelate
-	D02.241.511.536.730 Vanilmandelic Acid
-	D02.241.511.579 Mevalonic Acid
-	D02.241.511.621 Mycolic Acids
-	D02.241.511.701 Phosphoenolpyruvate
-	D02.241.511.764 Quinic Acid
-	D02.241.511.852 Shikimic Acid
-	D02.241.511.902 Sugar Acids



## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.241.511.902.107                      Ascorbic Acid
-	D02.241.511.902.107.260                      Dehydroascorbic Acid
-	D02.241.511.902.200                      2,3-Diketogulonic Acid
-	D02.241.511.902.300                      Glucaric Acid
-	D02.241.511.902.322                      Gluconates
-	D02.241.511.902.322.060                      Antimony Sodium Gluconate
-	D02.241.511.902.322.200                      Calcium Gluconate
-	D02.241.511.902.387                      Glyceric Acids
-	D02.241.511.902.387.388                      Diphosphoglyceric Acids
-	D02.241.511.902.387.388.175                      2,3-Diphosphoglycerate
-	D02.241.511.902.522                      Muramic Acids
-	D02.241.511.902.562                      Neuraminic Acids
-	D02.241.511.902.562.668                      Sialic Acids
-	D02.241.511.902.562.668.050                      N-Acetylneuraminic Acid
-	D02.241.511.902.562.668.250                      Cytidine Monophosphate N-Acetylneuraminic Acid
-	D02.241.511.902.562.668.775                      Zanamivir
-	D02.241.511.902.759                      Tartrates
-	D02.241.511.902.821                      Tartronates
-	D02.241.511.902.915                      Uronic Acids
-	D02.241.511.902.915.162                      Glucuronates
-	D02.241.511.902.915.162.500                      Glucuronic Acid
-	D02.241.511.902.915.162.750                      Glucuronides
-	D02.241.511.902.915.400                      Hexuronic Acids
-	D02.241.511.902.915.400.500                      Iduronic Acid
-	D02.241.755                      Keto Acids
-	D02.241.755.080                      Acetoacetates
-	D02.241.755.360                      Hippurates
-	D02.241.755.360.192                      Aminohippuric Acids
-	D02.241.755.360.192.100                      p-Aminohippuric Acid
-	D02.241.755.360.579                      Iodohippuric Acid
-	D02.241.755.465                      Ketoglutaric Acids
-	D02.241.755.547                      Levulinic Acids
-	D02.241.755.547.276                      Aminolevulinic Acid
-	D02.241.755.648                      Oxaloacetates
-	D02.241.755.648.750                      Oxaloacetic Acid

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.241.755.812 Pyruvates
-	D02.241.755.812.601 Phenylpyruvic Acids
-	D02.241.755.812.800 Pyruvic Acid
-	D02.241.803 Thioctic Acid
-	D02.251 Catenanes
-	D02.251.500 DNA, Catenated
-	D02.262 Cyanates
-	D02.262.775 Thiocyanates
New Heading	<b>D02.309 Dioxins and Dioxin-like Compounds</b>
New Heading	<b>D02.309.250 Dibenzofurans, Polychlorinated</b>
New Tree	<a href="#">D02.309.500 Dioxins</a>
New Heading	<b>D02.309.500.450 Polychlorinated Dibenzodioxins</b>
New Tree	<a href="#">D02.309.750 Polychlorinated Biphenyls</a>
New Tree	<a href="#">D02.309.750.500 Aroclors</a>
New Tree	<a href="#">D02.309.750.500.077 Chlorodiphenyl (54% Chlorine)</a>
-	D02.355 Ethers
-	D02.355.071 Acetals
-	D02.355.291 Ethers, Cyclic
-	D02.355.291.205 Ciguatoxins
-	D02.355.291.308 Crown Ethers
-	D02.355.291.411 Epoxy Compounds
-	D02.355.291.411.400 Epichlorohydrin
-	D02.355.291.411.408 Ethoglucid
-	D02.355.291.411.417 Ethylene Oxide
-	D02.355.291.411.658 Mupirocin
-	D02.355.291.411.900 Trichloroepoxypropane
-	D02.355.291.705 Okadaic Acid
-	D02.355.291.852 Oxepins
-	D02.355.291.852.500 Doxepin
-	D02.355.291.866 Oxocins
-	D02.355.354 Ethoglucid

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.355.417 Ethyl Ethers
-	D02.355.417.332 Ether
-	D02.355.417.515 Flurothyl
-	D02.355.417.600 Methoxyflurane
-	D02.355.417.650 Phenetidine
-	D02.355.460 Glyceryl Ethers
-	D02.355.460.750 Phospholipid Ethers
-	D02.355.601 Methyl Ethers
-	D02.355.601.200 Anisoles
-	D02.355.601.200.100 Anethole Trithione
-	D02.355.601.200.324 Butylated Hydroxyanisole
-	D02.355.601.250 Bis(Chloromethyl) Ether
-	D02.355.601.400 Enflurane
-	D02.355.601.536 Guaiacol
-	D02.355.601.536.400 Guaifenesin
-	D02.355.601.536.400.500 Methocarbamol
-	D02.355.601.570 Isoflurane
-	D02.355.601.620 Methoxyflurane
-	D02.355.726 Phenyl Ethers
-	D02.355.726.158 Anisoles
-	D02.355.726.158.100 Anethole Trithione
-	D02.355.726.158.324 Butylated Hydroxyanisole
-	D02.355.726.305 Fibric Acids
-	D02.355.726.305.249 Bezafibrate
-	D02.355.726.305.374 Clofenapate
-	D02.355.726.305.500 Clofibric Acid
-	D02.355.726.305.500.195 Clofibrate
-	D02.355.726.305.625 Fenofibrate
-	D02.355.726.305.750 Gemfibrozil
-	D02.355.726.453 Guaiacol
-	D02.355.726.453.400 Guaifenesin
-	D02.355.726.453.400.500 Methocarbamol
-	D02.355.726.601 Halogenated Diphenyl Ethers
-	D02.355.726.675 Hydroxybenzoate Ethers
-	D02.355.726.750 Phenetidine
-	D02.355.726.825 Tiapamil Hydrochloride

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.355.726.900 Triclosan
-	D02.389 Free Radicals
-	D02.389.338 Peroxides
-	D02.389.338.055 Artemisinins
-	D02.389.338.253 Hydrogen Peroxide
-	D02.389.338.450 Lipid Peroxides
-	D02.389.338.638 Prostaglandin Endoperoxides
-	D02.389.338.638.600 Prostaglandins G
-	D02.389.338.638.650 Prostaglandins H
-	D02.389.338.638.650.500 Prostaglandin H2
-	D02.389.338.638.650.500.500 15-Hydroxy-11 alpha,9 alpha-(epoxymethano)prosta-5,13-dienoic Acid
-	D02.389.338.732 Superoxides
-	D02.389.338.825 tert-Butylhydroperoxide
-	D02.389.338.912 Tetraoxanes
-	D02.389.678 Spin Labels
-	D02.389.678.900 Triacetoneamine-N-Oxyl
-	D02.442 Hydrazines
-	D02.442.175 Benserazide
-	D02.442.200 Carbidopa
-	D02.442.288 Hydrazones
-	D02.442.288.200 Carbonyl Cyanide m-Chlorophenyl Hydrazone
-	D02.442.288.220 Carbonyl Cyanide p-Trifluoromethoxyphenylhydrazone
-	D02.442.372 Iproniazid
-	D02.442.436 Isoniazid
-	D02.442.600 Methylhydrazines
-	D02.442.600.400 Dimethylhydrazines
-	D02.442.600.400.200 1,2-Dimethylhydrazine
-	D02.442.600.601 Monomethylhydrazine
-	D02.442.700 Phenelzine
-	D02.442.726 Phenylhydrazines
-	D02.442.726.200 Dithizone
-	D02.442.925 Todralazine
-	D02.455 Hydrocarbons
-	D02.455.326 Hydrocarbons, Acyclic
-	D02.455.326.146 Alkanes

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.455.326.146.049 Acetogenins
-	D02.455.326.146.100 Alkanesulfonic Acids
-	D02.455.326.146.100.050 Alkanesulfonates
-	D02.455.326.146.100.050.490 Mesna
-	D02.455.326.146.100.050.500 Mesylates
-	D02.455.326.146.100.050.500.100 Busulfan
-	D02.455.326.146.100.050.500.300 Ethyl Methanesulfonate
-	D02.455.326.146.100.050.500.500 Methyl Methanesulfonate
-	D02.455.326.146.100.250 HEPES
-	D02.455.326.146.100.300 Isethionic Acid
-	D02.455.326.146.100.850 Taurine
-	D02.455.326.146.100.850.875 Taurocholic Acid
-	D02.455.326.146.100.850.875.900 Taurodeoxycholic Acid
-	D02.455.326.146.100.850.875.900.875 Taurochenodeoxycholic Acid
-	D02.455.326.146.100.850.875.925 Taurolithocholic Acid
-	D02.455.326.146.216 Butanes
New Tree	<a href="#">D02.455.326.146.216.500</a> Butylamines
-	D02.455.326.146.379 Ethane
-	D02.455.326.146.379.350 Ethylene Dichlorides
-	D02.455.326.146.432 Fumonisin
-	D02.455.326.146.485 Heptanes
-	D02.455.326.146.485.222 Diarylheptanoids
-	D02.455.326.146.485.222.222 Curcumin
-	D02.455.326.146.500 Hexanes
-	D02.455.326.146.571 Methane
-	D02.455.326.146.571.850 Tetranitromethane
-	D02.455.326.146.672 Nitroparaffins
-	D02.455.326.146.672.800 Tetranitromethane
-	D02.455.326.146.720 Octanes
-	D02.455.326.146.770 Pentanes
-	D02.455.326.146.800 Propane
-	D02.455.326.271 Alkenes
-	D02.455.326.271.060 Polyunsaturated Alkamides
-	D02.455.326.271.060.222 Capsaicin
-	D02.455.326.271.122 Allyl Compounds

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D02.455.326.271.122.100	Allylamine
-	D02.455.326.271.122.100.500	Colesevelam Hydrochloride
-	D02.455.326.271.122.260	Allylglycine
-	D02.455.326.271.122.272	Allylisopropylacetamide
-	D02.455.326.271.244	Enediynes
-	D02.455.326.271.244.500	Zinostatin
-	D02.455.326.271.367	Ethylenes
-	D02.455.326.271.367.300	Dichloroethylenes
-	D02.455.326.271.665	Polyenes
-	D02.455.326.271.665.146	Alkadienes
-	D02.455.326.271.665.146.240	Butadienes
-	D02.455.326.271.665.146.280	Chloroprene
-	D02.455.326.271.665.155	Aurodox
-	D02.455.326.271.665.202	Carotenoids
-	D02.455.326.271.665.202.061	Abscisic Acid
-	D02.455.326.271.665.202.123	beta Carotene
-	D02.455.326.271.665.202.309	Norisoprenoids
-	D02.455.326.271.665.202.495	Retinoids
-	D02.455.326.271.665.202.495.050	Acitretin
-	D02.455.326.271.665.202.495.250	Etretinate
-	D02.455.326.271.665.202.495.270	Fenretinide
-	D02.455.326.271.665.202.495.325	Isotretinoin
-	D02.455.326.271.665.202.495.690	Retinaldehyde
-	D02.455.326.271.665.202.495.818	Vitamin A
-	D02.455.326.271.665.202.495.818.500	Tretinoin
-	D02.455.326.271.665.202.868	Xanthophylls
-	D02.455.326.271.665.202.868.249	Canthaxanthin
-	D02.455.326.271.665.202.868.374	Cryptoxanthins
New Heading	<b>D02.455.326.271.665.202.868.374.500</b>	<b>Beta-Cryptoxanthin</b>
-	D02.455.326.271.665.202.868.500	Lutein
-	D02.455.326.271.665.202.868.750	Zeaxanthins
-	D02.455.326.271.665.202.912	zeta Carotene
-	D02.455.326.271.665.250	Diphenylhexatriene
-	D02.455.326.271.665.325	Filipin
-	D02.455.326.271.665.400	Mepartricin

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.455.326.271.665.480 Neoprene
-	D02.455.326.271.665.550 Polyethylenes
-	D02.455.326.271.665.550.500 Polyethylene
-	D02.455.326.271.665.550.600 Polyethyleneimine
-	D02.455.326.271.665.590 Polypropylenes
-	D02.455.326.271.665.616 Polyvinyls
-	D02.455.326.271.665.806 Squalene
-	D02.455.326.271.884 Vinyl Compounds
-	D02.455.326.271.884.533 Polyvinyls
-	D02.455.326.271.884.533.532 Polyvinyl Alcohol
-	D02.455.326.271.884.533.565 Polyvinyl Chloride
-	D02.455.326.271.884.533.660 Polyvinylpyridine N-Oxide
-	D02.455.326.271.884.533.699 Povidone
-	D02.455.326.271.884.533.710 Povidone-Iodine
-	D02.455.326.271.884.750 Vinyl Chloride
-	D02.455.326.397 Alkynes
-	D02.455.326.397.259 Acetylene
-	D02.455.326.397.300 Carbocyanines
-	D02.455.326.397.325 (4-(m-Chlorophenylcarbamoyloxy)-2-butynyl)trimethylammonium Chloride
-	D02.455.326.397.662 Polyacetylenes
-	D02.455.326.397.662.500 Diynes
-	D02.455.326.397.662.500.500 Ene-diyne
-	D02.455.326.397.662.500.500.500 Zinostatin
-	D02.455.326.397.675 Polyunsaturated Alkarnides
-	D02.455.426 Hydrocarbons, Cyclic
-	D02.455.426.100 Bridged Compounds
-	D02.455.426.100 Bridged-Ring Compounds
-	D02.455.426.100.050 Adamantane
-	D02.455.426.100.050.035 Amantadine
-	D02.455.426.100.050.035.500 Memantine
-	D02.455.426.100.050.392 Methenamine
-	D02.455.426.100.050.750 Rimantadine
-	D02.455.426.100.080 Bicyclo Compounds
-	D02.455.426.100.080 Bridged Bicyclo Compounds
Old Tree	<b>D02.455.426.100.080.085 Bicyclo Compounds, Heterocyclic</b>

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	<b>D02.455.426.100.080.085</b> <span style="float: right;"><b>Bridged Bicyclo Compounds, Heterocyclic</b></span>
-	D02.455.426.100.080.100 <span style="float: right;">Biperiden</span>
-	D02.455.426.392 <span style="float: right;">Hydrocarbons, Alicyclic</span>
-	D02.455.426.392.368 <span style="float: right;">Cycloparaffins</span>
-	D02.455.426.392.368.075 <span style="float: right;">Adamantane</span>
-	D02.455.426.392.368.075.750 <span style="float: right;">Rimantadine</span>
-	D02.455.426.392.368.138 <span style="float: right;">Cyclitols</span>
-	D02.455.426.392.368.201 <span style="float: right;">Cyclobutanes</span>
-	D02.455.426.392.368.242 <span style="float: right;">Cyclodecanes</span>
-	D02.455.426.392.368.242.888 <span style="float: right;">Taxoids</span>
-	D02.455.426.392.368.242.888.777 <span style="float: right;">Paclitaxel</span>
-	D02.455.426.392.368.242.888.777.500 <span style="float: right;">Albumin-Bound Paclitaxel</span>
-	D02.455.426.392.368.284 <span style="float: right;">Cycloheptanes</span>
-	D02.455.426.392.368.284.049 <span style="float: right;">Azulenes</span>
-	D02.455.426.392.368.284.100 <span style="float: right;">Bencyclane</span>
-	D02.455.426.392.368.284.500 <span style="float: right;">Sesquiterpenes, Guaiane</span>
-	D02.455.426.392.368.284.500.888 <span style="float: right;">Thapsigargin</span>
-	D02.455.426.392.368.284.900 <span style="float: right;">Tropolone</span>
-	D02.455.426.392.368.367 <span style="float: right;">Cyclohexanes</span>
-	D02.455.426.392.368.367.190 <span style="float: right;">Cuprizone</span>
-	D02.455.426.392.368.367.204 <span style="float: right;">Cyclamates</span>
-	D02.455.426.392.368.367.218 <span style="float: right;">Cyclohexanecarboxylic Acids</span>
-	D02.455.426.392.368.367.318 <span style="float: right;">Cyclohexanols</span>
-	D02.455.426.392.368.367.318.500 <span style="float: right;">Desvenlafaxine Succinate</span>
-	D02.455.426.392.368.367.318.750 <span style="float: right;">Venlafaxine Hydrochloride</span>
-	D02.455.426.392.368.367.340 <span style="float: right;">Cyclohexanones</span>
-	D02.455.426.392.368.367.379 <span style="float: right;">Cyclohexenes</span>
-	D02.455.426.392.368.367.379.249 <span style="float: right;">Carotenoids</span>
-	D02.455.426.392.368.367.379.249.024 <span style="float: right;">Abscisic Acid</span>
-	D02.455.426.392.368.367.379.249.050 <span style="float: right;">beta Carotene</span>
-	D02.455.426.392.368.367.379.249.375 <span style="float: right;">Norisoprenoids</span>
-	D02.455.426.392.368.367.379.249.700 <span style="float: right;">Retinoids</span>
-	D02.455.426.392.368.367.379.249.700.270 <span style="float: right;">Fenretinide</span>
-	D02.455.426.392.368.367.379.249.700.325 <span style="float: right;">Isotretinoin</span>
-	D02.455.426.392.368.367.379.249.700.690 <span style="float: right;">Retinaldehyde</span>
-	D02.455.426.392.368.367.379.249.700.860 <span style="float: right;">Vitamin A</span>



## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.455.426.392.368.367.379.249.700.860.500 <span style="float: right;">Tretinoin</span>
-	D02.455.426.392.368.367.379.249.887 <span style="float: right;">Xanthophylls</span>
-	D02.455.426.392.368.367.379.249.887.249 <span style="float: right;">Canthaxanthin</span>
-	D02.455.426.392.368.367.379.249.887.374 <span style="float: right;">Cryptoxanthins</span>
New Heading	<b>D02.455.426.392.368.367.379.249.887.374.500</b> <span style="float: right;"><b>Beta-Cryptoxanthin</b></span>
-	D02.455.426.392.368.367.379.249.887.500 <span style="float: right;">Lutein</span>
-	D02.455.426.392.368.367.379.249.887.750 <span style="float: right;">Zeaxanthins</span>
-	D02.455.426.392.368.367.379.249.925 <span style="float: right;">zeta Carotene</span>
-	D02.455.426.392.368.367.379.374 <span style="float: right;">Chorismic Acid</span>
-	D02.455.426.392.368.367.379.500 <span style="float: right;">Oseltamivir</span>
-	D02.455.426.392.368.367.379.750 <span style="float: right;">Shikimic Acid</span>
-	D02.455.426.392.368.367.418 <span style="float: right;">Cyclohexylamines</span>
-	D02.455.426.392.368.367.440 <span style="float: right;">Dicyclomine</span>
-	D02.455.426.392.368.367.652 <span style="float: right;">Ketamine</span>
-	D02.455.426.392.368.367.800 <span style="float: right;">Picrotoxin</span>
-	D02.455.426.392.368.367.850 <span style="float: right;">Tiletamine</span>
-	D02.455.426.392.368.367.900 <span style="float: right;">trans-1,4-Bis(2-</span> chlorobenzaminomethyl)cyclohexane Dihydrochloride
-	D02.455.426.392.368.408 <span style="float: right;">Cyclooctanes</span>
-	D02.455.426.392.368.408.333 <span style="float: right;">Iprindole</span>
-	D02.455.426.392.368.450 <span style="float: right;">Cyclopentanes</span>
-	D02.455.426.392.368.450.350 <span style="float: right;">Cycloleucine</span>
-	D02.455.426.392.368.533 <span style="float: right;">Cyclopropanes</span>
-	D02.455.426.392.368.533.200 <span style="float: right;">Cilastatin</span>
-	D02.455.426.392.368.533.450 <span style="float: right;">Hypoglycins</span>
-	D02.455.426.559 <span style="float: right;">Hydrocarbons, Aromatic</span>
-	D02.455.426.559.389 <span style="float: right;">Benzene Derivatives</span>
-	D02.455.426.559.389.023 <span style="float: right;">Benzene</span>
-	D02.455.426.559.389.048 <span style="float: right;">Benzeneacetamides</span>
-	D02.455.426.559.389.048.088 <span style="float: right;">Bufexamac</span>
-	D02.455.426.559.389.097 <span style="float: right;">Benzenesulfonates</span>
-	D02.455.426.559.389.097.120 <span style="float: right;">Calcium Dobesilate</span>
-	D02.455.426.559.389.097.150 <span style="float: right;">4-Chloromercuribenzenesulfonate</span>
-	D02.455.426.559.389.097.175 <span style="float: right;">1,2-Dihydroxybenzene-3,5-Disulfonic Acid</span> Disodium Salt
-	D02.455.426.559.389.097.250 <span style="float: right;">Ethamsylate</span>

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.455.426.559.389.097.280 Ferrozine
-	D02.455.426.559.389.097.675 Polyanetholesulfonate
-	D02.455.426.559.389.115 Benzhydryl Compounds
Old Tree	<b>D02.455.426.559.389.115.124 Benzophenones</b>
Old Tree	<b>D02.455.426.559.389.115.124.500 Chlorthalidone</b>
Old Tree	<b>D02.455.426.559.389.115.124.750 Fenofibrate</b>
-	D02.455.426.559.389.115.250 Diphenhydramine
-	D02.455.426.559.389.115.250.250 Dimenhydrinate
-	D02.455.426.559.389.115.475 Meclizine
-	D02.455.426.559.389.115.500 Methylenebis(chloroaniline)
-	D02.455.426.559.389.115.600 Orphenadrine
-	D02.455.426.559.389.115.800 Terfenadine
-	D02.455.426.559.389.115.900 Tolterodine Tartrate
-	D02.455.426.559.389.127 Benzoates
-	D02.455.426.559.389.127.020 Aminobenzoates
-	D02.455.426.559.389.127.020.452 meta-Aminobenzoates
-	D02.455.426.559.389.127.020.452.249 Acetrizic Acid
-	D02.455.426.559.389.127.020.452.500 Bumetanide
-	D02.455.426.559.389.127.020.452.750 Mesalamine
-	D02.455.426.559.389.127.020.906 ortho-Aminobenzoates
-	D02.455.426.559.389.127.020.906.249 3-Hydroxyanthranilic Acid
-	D02.455.426.559.389.127.020.906.750 Fenamates
-	D02.455.426.559.389.127.020.906.750.249 Flufenamic Acid
-	D02.455.426.559.389.127.020.906.750.500 Meclofenamic Acid
-	D02.455.426.559.389.127.020.906.750.555 Mefenamic Acid
-	D02.455.426.559.389.127.020.906.750.777 Niflumic Acid
-	D02.455.426.559.389.127.020.906.875 Glafenine
-	D02.455.426.559.389.127.020.937 para-Aminobenzoates
-	D02.455.426.559.389.127.020.937.625 Acecainide
-	D02.455.426.559.389.127.020.937.640 4-Aminobenzoic Acid
-	D02.455.426.559.389.127.020.937.642 Aminosalicylic Acid
-	D02.455.426.559.389.127.020.937.645 Ethopabate
-	D02.455.426.559.389.127.020.937.647 Metoclopramide
-	D02.455.426.559.389.127.020.937.650 p-Aminohippuric Acid
-	D02.455.426.559.389.127.020.937.687 Benzocaine
-	D02.455.426.559.389.127.020.937.700 Cisapride

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.455.426.559.389.127.020.937.875 Procainamide
-	D02.455.426.559.389.127.020.937.906 Procaine
-	D02.455.426.559.389.127.020.937.906.666 Penicillin G Procaine
-	D02.455.426.559.389.127.020.937.937 Propoxycaine
-	D02.455.426.559.389.127.020.937.968 Tetracaine
-	D02.455.426.559.389.127.085 Benzamides
-	D02.455.426.559.389.127.085.033 Acecainide
-	D02.455.426.559.389.127.085.067 Aminohippuric Acids
-	D02.455.426.559.389.127.085.067.100 p-Aminohippuric Acid
-	D02.455.426.559.389.127.085.101 Bezafibrate
-	D02.455.426.559.389.127.085.135 Cisapride
-	D02.455.426.559.389.127.085.230 DEET
-	D02.455.426.559.389.127.085.240 Dinitolmide
-	D02.455.426.559.389.127.085.460 Hippurates
-	D02.455.426.559.389.127.085.460.500 Iodohippuric Acid
-	D02.455.426.559.389.127.085.465 Imatinib Mesylate
-	D02.455.426.559.389.127.085.470 Indoramin
-	D02.455.426.559.389.127.085.510 Metoclopramide
-	D02.455.426.559.389.127.085.600 Moclobemide
-	D02.455.426.559.389.127.085.650 Procainamide
-	D02.455.426.559.389.127.085.655 Procarbazine
-	D02.455.426.559.389.127.085.761 Raclopride
-	D02.455.426.559.389.127.085.796 Remoxipride
-	D02.455.426.559.389.127.085.866 Sulpiride
-	D02.455.426.559.389.127.085.933 Tiapride Hydrochloride
-	D02.455.426.559.389.127.117 Benzoic Acid
-	D02.455.426.559.389.127.117.500 Sodium Benzoate
-	D02.455.426.559.389.127.133 Benzoyl Peroxide
-	D02.455.426.559.389.127.133.500 Combination Adapalene, Benzoyl Peroxide Drug
-	D02.455.426.559.389.127.141 Benzoylcholine
-	D02.455.426.559.389.127.150 Bromobenzoates
-	D02.455.426.559.389.127.150.500 Remoxipride
-	D02.455.426.559.389.127.250 Chlorobenzoates
-	D02.455.426.559.389.127.250.249 Bezafibrate
-	D02.455.426.559.389.127.250.311 Chloromercuribenzoates

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.455.426.559.389.127.250.311.275 p-Chloromercuribenzoic Acid
-	D02.455.426.559.389.127.250.374 Cisapride
-	D02.455.426.559.389.127.250.500 Dicamba
-	D02.455.426.559.389.127.250.750 Metoclopramide
-	D02.455.426.559.389.127.250.875 Moclobemide
-	D02.455.426.559.389.127.250.937 Racioipride
-	D02.455.426.559.389.127.281 Hydroxybenzoates
-	D02.455.426.559.389.127.281.100 Depsides
-	D02.455.426.559.389.127.281.200 Gallic Acid
-	D02.455.426.559.389.127.281.200.299 Hydrolyzable Tannins
-	D02.455.426.559.389.127.281.200.600 Propyl Gallate
-	D02.455.426.559.389.127.281.350 Hydroxybenzoate Ethers
-	D02.455.426.559.389.127.281.350.500 Hexobendine
-	D02.455.426.559.389.127.281.350.625 Metoclopramide
-	D02.455.426.559.389.127.281.350.687 Propoxycaine
-	D02.455.426.559.389.127.281.350.750 Trimebutine
-	D02.455.426.559.389.127.281.350.875 Vanillic Acid
-	D02.455.426.559.389.127.281.400 Hydroxymercuribenzoates
-	D02.455.426.559.389.127.281.450 Pactamycin
-	D02.455.426.559.389.127.281.460 Parabens
-	D02.455.426.559.389.127.281.595 Salicylates
-	D02.455.426.559.389.127.281.595.100 Aminosalicylic Acids
-	D02.455.426.559.389.127.281.595.100.060 Aminosalicylic Acid
-	D02.455.426.559.389.127.281.595.100.540 Mesalamine
-	D02.455.426.559.389.127.281.595.202 Dicamba
-	D02.455.426.559.389.127.281.595.405 Gentsitates
-	D02.455.426.559.389.127.281.595.608 Salicylic Acid
-	D02.455.426.559.389.127.281.595.608.500 Sodium Salicylate
-	D02.455.426.559.389.127.375 Iodobenzoates
-	D02.455.426.559.389.127.375.880 Triiodobenzoic Acids
-	D02.455.426.559.389.127.375.880.020 Acetrizic Acid
-	D02.455.426.559.389.127.375.880.270 Diatrizoate
-	D02.455.426.559.389.127.375.880.270.500 Diatrizoate Meglumine
-	D02.455.426.559.389.127.375.880.375 Iodamide
-	D02.455.426.559.389.127.375.880.380 Iodipamide
-	D02.455.426.559.389.127.375.880.390 Ioglycamic Acid

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D02.455.426.559.389.127.375.880.400	Iohexol
-	D02.455.426.559.389.127.375.880.410	Iopamidol
-	D02.455.426.559.389.127.375.880.420	Iothalamate Meglumine
-	D02.455.426.559.389.127.375.880.430	Iothalamic Acid
-	D02.455.426.559.389.127.375.880.450	Ioxaglic Acid
-	D02.455.426.559.389.127.375.880.520	Metrizamide
-	D02.455.426.559.389.127.375.880.530	Metrizoate
-	D02.455.426.559.389.127.500	Mercuribenzoates
-	D02.455.426.559.389.127.500.261	Chloromercuribenzoates
-	D02.455.426.559.389.127.500.261.275	p-Chloromercuribenzoic Acid
-	D02.455.426.559.389.127.500.450	Hydroxymercuribenzoates
-	D02.455.426.559.389.127.650	Nitrobenzoates
-	D02.455.426.559.389.127.650.300	Dithionitrobenzoic Acid
New Tree	<a href="#">D02.455.426.559.389.134</a>	<a href="#">Benzophenones</a>
New Tree	<a href="#">D02.455.426.559.389.134.500</a>	<a href="#">Chlorthalidone</a>
New Tree	<a href="#">D02.455.426.559.389.134.750</a>	<a href="#">Fenofibrate</a>
-	D02.455.426.559.389.140	Benzyl Compounds
-	D02.455.426.559.389.140.200	Benzyl Alcohols
-	D02.455.426.559.389.140.200.100	Benzyl Alcohol
-	D02.455.426.559.389.140.210	Benzylamines
-	D02.455.426.559.389.140.210.650	Pargyline
-	D02.455.426.559.389.140.308	Bibenzyls
-	D02.455.426.559.389.140.400	2-Hydroxy-5-nitrobenzyl Bromide
-	D02.455.426.559.389.140.450	Lignans
-	D02.455.426.559.389.140.450.388	Flavonolignans
-	D02.455.426.559.389.140.450.582	Masoprocol
-	D02.455.426.559.389.140.450.777	Podophyllotoxin
-	D02.455.426.559.389.150	Benzylidene Compounds
-	D02.455.426.559.389.150.700	Stilbenes
-	D02.455.426.559.389.150.700.050 disulfonic Acid	4-Acetamido-4'-isothiocyanatostilbene-2,2'-
-	D02.455.426.559.389.150.700.075	Bibenzyls
-	D02.455.426.559.389.150.700.100	Chlorotrianisene
-	D02.455.426.559.389.150.700.125	Clomiphene

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.455.426.559.389.150.700.125.500 Enclomiphene
-	D02.455.426.559.389.150.700.125.750 Zuclomiphene
-	D02.455.426.559.389.150.700.175 Diethylstilbestrol
-	D02.455.426.559.389.150.700.175.450 Hexestrol
-	D02.455.426.559.389.150.700.200 Acid 4,4'-Diisothiocyanostilbene-2,2'-Disulfonic
-	D02.455.426.559.389.150.700.550 Stilbamidines
-	D02.455.426.559.389.150.700.900 Tamoxifen
-	D02.455.426.559.389.150.700.900.775 Raloxifene Hydrochloride
-	D02.455.426.559.389.150.700.900.900 Toremifene
-	D02.455.426.559.389.150.750 Styrenes
-	D02.455.426.559.389.150.750.800 Styrene
-	D02.455.426.559.389.150.750.800.830 Polystyrenes
-	D02.455.426.559.389.150.795 Tyrphostins
-	D02.455.426.559.389.185 Biphenyl Compounds
-	D02.455.426.559.389.185.060 Aminobiphenyl Compounds
-	D02.455.426.559.389.185.100 Benzidines
-	D02.455.426.559.389.185.200 3,3'-Diaminobenzidine
-	D02.455.426.559.389.185.210 Clofenapate
-	D02.455.426.559.389.185.220 Dianisidine
-	D02.455.426.559.389.185.226 3,3'-Dichlorobenzidine
-	D02.455.426.559.389.185.350 Flurbiprofen
-	D02.455.426.559.389.185.475 Losartan
-	D02.455.426.559.389.185.600 Niclofolan
-	D02.455.426.559.389.185.680 Polybrominated Biphenyls
-	D02.455.426.559.389.185.698 Polychlorinated Biphenyls
-	D02.455.426.559.389.220 Bromobenzenes
-	D02.455.426.559.389.261 Chlorobenzenes
-	D02.455.426.559.389.261.190 Chlorophenols
-	D02.455.426.559.389.261.190.350 Hexachlorophene
-	D02.455.426.559.389.261.220 Dicofol
-	D02.455.426.559.389.261.250 Dinitrochlorobenzene
-	D02.455.426.559.389.261.400 Hexachlorobenzene
-	D02.455.426.559.389.454 Iodobenzenes
-	D02.455.426.559.389.454.300 3-Iodobenzylguanidine
-	D02.455.426.559.389.454.385 Iopanoic Acid

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.455.426.559.389.454.400 Iophendylate
-	D02.455.426.559.389.454.450 Ipodate
-	D02.455.426.559.389.454.950 Tyropanoate
-	D02.455.426.559.389.565 Nitrobenzenes
-	D02.455.426.559.389.565.175 Chloramphenicol
-	D02.455.426.559.389.565.175.850 Thiamphenicol
-	D02.455.426.559.389.565.225 Dinitrobenzenes
-	D02.455.426.559.389.565.250 Dinitrochlorobenzene
-	D02.455.426.559.389.565.280 Dinitrofluorobenzene
-	D02.455.426.559.389.565.880 Trinitrobenzenes
-	D02.455.426.559.389.565.880.880 Trinitrobenzenesulfonic Acid
-	D02.455.426.559.389.657 Phenols
-	D02.455.426.559.389.657.050 Aminophenols
-	D02.455.426.559.389.657.100 Bisphenol A-Glycidyl Methacrylate
-	D02.455.426.559.389.657.120 Bithionol
-	D02.455.426.559.389.657.150 Bromphenol Blue
-	D02.455.426.559.389.657.166 Catechols
-	D02.455.426.559.389.657.166.099 Capsaicin
-	D02.455.426.559.389.657.166.200 Curcumin
-	D02.455.426.559.389.657.166.210 1,2-Dihydroxybenzene-3,5-Disulfonic Acid Disodium Salt
-	D02.455.426.559.389.657.166.350 Estrogens, Catechol
-	D02.455.426.559.389.657.166.350.350 Hydroxyestrones
-	D02.455.426.559.389.657.166.453 Guaiacol
-	D02.455.426.559.389.657.166.453.400 Guaifenesin
-	D02.455.426.559.389.657.166.500 3-Methoxy-4-hydroxyphenylethanol
-	D02.455.426.559.389.657.166.550 Masoprocol
-	D02.455.426.559.389.657.190 Chlorophenols
-	D02.455.426.559.389.657.190.270 Dichlorophen
-	D02.455.426.559.389.657.190.350 Hexachlorophene
-	D02.455.426.559.389.657.190.633 Pentachlorophenol
-	D02.455.426.559.389.657.239 Cresols
-	D02.455.426.559.389.657.239.132 Bisacodyl
-	D02.455.426.559.389.657.239.180 Brocresine
-	D02.455.426.559.389.657.239.190 Bromcresol Green
-	D02.455.426.559.389.657.239.195 Bromcresol Purple

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.455.426.559.389.657.239.216 Butylated Hydroxytoluene
-	D02.455.426.559.389.657.239.327 Creosote
-	D02.455.426.559.389.657.239.402 Cyclofenil
-	D02.455.426.559.389.657.239.493 Dinitrocresols
-	D02.455.426.559.389.657.239.611 Formocresols
-	D02.455.426.559.389.657.239.756 Tolterodine Tartrate
-	D02.455.426.559.389.657.239.900 Tritolyl Phosphates
-	D02.455.426.559.389.657.265 Desvenlafaxine Succinate
-	D02.455.426.559.389.657.290 Dienestrol
-	D02.455.426.559.389.657.377 Humic Substances
-	D02.455.426.559.389.657.393 Hydroquinones
-	D02.455.426.559.389.657.410 Hydroxybenzoates
-	D02.455.426.559.389.657.410.100 Depsides
-	D02.455.426.559.389.657.410.200 Gallic Acid
-	D02.455.426.559.389.657.410.200.299 Hydrolyzable Tannins
-	D02.455.426.559.389.657.410.200.600 Propyl Gallate
-	D02.455.426.559.389.657.410.400 Hydroxymercuribenzoates
-	D02.455.426.559.389.657.410.450 Pactamycin
-	D02.455.426.559.389.657.410.460 Parabens
-	D02.455.426.559.389.657.410.595 Salicylates
-	D02.455.426.559.389.657.410.595.100 Aminosalicylic Acids
-	D02.455.426.559.389.657.410.595.100.060 Aminosalicylic Acid
-	D02.455.426.559.389.657.410.595.100.540 Mesalamine
-	D02.455.426.559.389.657.410.595.151 Anacardic Acids
-	D02.455.426.559.389.657.410.595.176 Aspirin
-	D02.455.426.559.389.657.410.595.176.500 Aspirin, Dipyridamole Drug Combination
-	D02.455.426.559.389.657.410.595.202 Dicamba
-	D02.455.426.559.389.657.410.595.303 Diflunisal
-	D02.455.426.559.389.657.410.595.405 Gentisates
-	D02.455.426.559.389.657.410.595.608 Salicylic Acid
-	D02.455.426.559.389.657.410.595.608.500 Sodium Salicylate
-	D02.455.426.559.389.657.446 Indophenol
-	D02.455.426.559.389.657.446.250 2,6-Dichloroindophenol
-	D02.455.426.559.389.657.566 Nitrophenols
-	D02.455.426.559.389.657.566.304 Dinitrophenols



## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.455.426.559.389.657.566.304.500 2,4-Dinitrophenol
-	D02.455.426.559.389.657.566.400 2-Hydroxy-5-nitrobenzyl Bromide
-	D02.455.426.559.389.657.566.600 Niclofolan
-	D02.455.426.559.389.657.566.610 Nitrohydroxyiodophenylacetate
-	D02.455.426.559.389.657.566.620 Nitroxinil
-	D02.455.426.559.389.657.566.690 Picrates
-	D02.455.426.559.389.657.595 Phenol
-	D02.455.426.559.389.657.625 Phenolphthaleins
-	D02.455.426.559.389.657.625.555 Phenolphthalein
-	D02.455.426.559.389.657.625.560 Phenolsulfonphthalein
-	D02.455.426.559.389.657.625.617 Sulfobromophthalein
-	D02.455.426.559.389.657.625.808 Thymolphthalein
-	D02.455.426.559.389.657.654 Phenyl Ethers
-	D02.455.426.559.389.657.654.158 Anisoles
-	D02.455.426.559.389.657.654.158.100 Anethole Trithione
-	D02.455.426.559.389.657.654.158.324 Butylated Hydroxyanisole
-	D02.455.426.559.389.657.654.305 Fibric Acids
-	D02.455.426.559.389.657.654.305.249 Bezafibrate
-	D02.455.426.559.389.657.654.305.374 Clofenapate
-	D02.455.426.559.389.657.654.305.500 Clofibrilic Acid
-	D02.455.426.559.389.657.654.305.500.195 Clofibrate
-	D02.455.426.559.389.657.654.305.625 Fenofibrate
-	D02.455.426.559.389.657.654.305.750 Gemfibrozil
-	D02.455.426.559.389.657.654.453 Guaiacol
-	D02.455.426.559.389.657.654.453.400 Guaifenesin
-	D02.455.426.559.389.657.654.453.400.500 Methocarbamol
-	D02.455.426.559.389.657.654.601 Halogenated Diphenyl Ethers
-	D02.455.426.559.389.657.654.638 Hydroxybenzoate Ethers
-	D02.455.426.559.389.657.654.638.500 Hexobendine
-	D02.455.426.559.389.657.654.638.625 Metoclopramide
-	D02.455.426.559.389.657.654.638.750 Trimebutine
-	D02.455.426.559.389.657.654.638.875 Vanillic Acid
-	D02.455.426.559.389.657.654.675 Mexiletine
-	D02.455.426.559.389.657.654.750 Phenetidine
-	D02.455.426.559.389.657.654.900 Triclosan
-	D02.455.426.559.389.657.684 Phloroglucinol

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.455.426.559.389.657.684.602 Phloretin
-	D02.455.426.559.389.657.684.602.615 Polyphloretin Phosphate
-	D02.455.426.559.389.657.715 Polyphenols
-	D02.455.426.559.389.657.746 Probucol
-	D02.455.426.559.389.657.773 Propofol
-	D02.455.426.559.389.657.800 Pyrogallol
-	D02.455.426.559.389.657.852 Resorcinols
-	D02.455.426.559.389.657.852.467 Hexylresorcinol
-	D02.455.426.559.389.657.852.900 Zearalenone
-	D02.455.426.559.389.657.852.900.975 Zeranol
-	D02.455.426.559.389.703 Phenylurea Compounds
-	D02.455.426.559.389.703.202 Carbanilides
-	D02.455.426.559.389.703.202.400 Imidocarb
-	D02.455.426.559.389.703.202.550 Nicarbazin
-	D02.455.426.559.389.703.241 Celiprolol
-	D02.455.426.559.389.703.280 Diflubenzuron
-	D02.455.426.559.389.703.397 Diuron
-	D02.455.426.559.389.703.515 Linuron
-	D02.455.426.559.389.750 Sulfanilic Acids
-	D02.455.426.559.389.805 Terphenyl Compounds
-	D02.455.426.559.389.805.700 Polychloroterphenyl Compounds
-	D02.455.426.559.389.832 Toluene
-	D02.455.426.559.389.832.140 Bromcresol Green
-	D02.455.426.559.389.832.150 Bromcresol Purple
-	D02.455.426.559.389.832.180 Bromphenol Blue
-	D02.455.426.559.389.832.200 Bromthymol Blue
-	D02.455.426.559.389.832.500 Toluene 2,4-Diisocyanate
-	D02.455.426.559.389.832.559 Toluidines
-	D02.455.426.559.389.832.661 Tosyl Compounds
-	D02.455.426.559.389.832.670 Tosylarginine Methyl Ester
-	D02.455.426.559.389.832.690 Tosyllysine Chloromethyl Ketone
-	D02.455.426.559.389.832.710 Tosylphenylalanyl Chloromethyl Ketone
-	D02.455.426.559.389.832.767 Trifluralin
-	D02.455.426.559.389.832.868 Trinitrotoluene
-	D02.455.426.559.389.875 Triparanol
-	D02.455.426.559.389.884 Trityl Compounds

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.455.426.559.389.948 Xylenes
-	D02.455.426.559.694 Diarylheptanoids
-	D02.455.426.559.694.222 Curcumin
-	D02.455.426.559.847 Polycyclic Aromatic Hydrocarbons
-	D02.455.426.559.847 Polycyclic Hydrocarbons, Aromatic
-	D02.455.426.559.847.117 Anthracenes
-	D02.455.426.559.847.117.050 Anthralin
-	D02.455.426.559.847.117.159 Anthraquinones
-	D02.455.426.559.847.117.159.200 Carmine
-	D02.455.426.559.847.117.159.205 Cascara
-	D02.455.426.559.847.117.159.205.400 Emodin
-	D02.455.426.559.847.117.159.500 Mitoxantrone
-	D02.455.426.559.847.117.159.750 Senna Extract
-	D02.455.426.559.847.117.600 Maprotiline
-	D02.455.426.559.847.133 Azulenes
-	D02.455.426.559.847.149 Benz(a)Anthracenes
-	D02.455.426.559.847.149.301 9,10-Dimethyl-1,2-benzanthracene
-	D02.455.426.559.847.149.500 Methylcholanthrene
-	D02.455.426.559.847.149.700 Perylene
-	D02.455.426.559.847.181 Benzocycloheptenes
-	D02.455.426.559.847.181.384 Dibenzocycloheptenes
-	D02.455.426.559.847.181.384.100 Amitriptyline
-	D02.455.426.559.847.181.384.230 Butaclamol
-	D02.455.426.559.847.181.384.340 Cyproheptadine
-	D02.455.426.559.847.181.384.340.500 Loratadine
-	D02.455.426.559.847.181.384.380 Dizocilpine Maleate
-	D02.455.426.559.847.181.384.535 Nortriptyline
-	D02.455.426.559.847.181.384.650 Protriptyline
-	D02.455.426.559.847.389 Fluorenes
-	D02.455.426.559.847.389.050 2-Acetylaminofluorene
-	D02.455.426.559.847.389.050.060 Acetoxyacetylaminofluorene
-	D02.455.426.559.847.389.050.400 Hydroxyacetylaminofluorene
-	D02.455.426.559.847.389.850 Tilorone
-	D02.455.426.559.847.486 Indenes
-	D02.455.426.559.847.486.250 Dimethindene
-	D02.455.426.559.847.486.487 Indans

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.455.426.559.847.486.487.060 Aprindine
-	D02.455.426.559.847.486.487.500 Ninhydrin
-	D02.455.426.559.847.486.487.750 Phenindione
-	D02.455.426.559.847.486.875 Sulindac
-	D02.455.426.559.847.562 Naphthacenes
-	D02.455.426.559.847.562.050 Anthracyclines
-	D02.455.426.559.847.562.050.050 Aclarubicin
-	D02.455.426.559.847.562.050.200 Daunorubicin
-	D02.455.426.559.847.562.050.200.150 Carubicin
-	D02.455.426.559.847.562.050.200.175 Doxorubicin
-	D02.455.426.559.847.562.050.200.175.200 Epirubicin
-	D02.455.426.559.847.562.050.200.300 Idarubicin
-	D02.455.426.559.847.562.050.200.650 Nogalamycin
-	D02.455.426.559.847.562.050.200.650.500 Menogaril
-	D02.455.426.559.847.562.050.650 Plicamycin
-	D02.455.426.559.847.562.900 Tetracyclines
-	D02.455.426.559.847.562.900.146 Chlortetracycline
-	D02.455.426.559.847.562.900.185 Demeclocycline
-	D02.455.426.559.847.562.900.200 Doxycycline
-	D02.455.426.559.847.562.900.450 Lymecycline
-	D02.455.426.559.847.562.900.525 Methacycline
-	D02.455.426.559.847.562.900.550 Minocycline
-	D02.455.426.559.847.562.900.600 Oxytetracycline
-	D02.455.426.559.847.562.900.720 Rolitetracycline
-	D02.455.426.559.847.562.900.875 Tetracycline
-	D02.455.426.559.847.638 Naphthalenes
-	D02.455.426.559.847.638.059 Acenaphthenes
-	D02.455.426.559.847.638.090 Adapalene
-	D02.455.426.559.847.638.090.500 Adapalene, Benzoyl Peroxide Drug Combination
-	D02.455.426.559.847.638.120 Bunaftine
-	D02.455.426.559.847.638.162 Carbaryl
-	D02.455.426.559.847.638.183 Cinacalcet Hydrochloride
-	D02.455.426.559.847.638.204 Dansyl Compounds
-	D02.455.426.559.847.638.400 Lovastatin
-	D02.455.426.559.847.638.400.900 Simvastatin

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.455.426.559.847.638.400.900.500 Ezetimibe, Simvastatin Drug Combination
-	D02.455.426.559.847.638.472 Naphthaleneacetic Acids
-	D02.455.426.559.847.638.472.500 Naproxen
-	D02.455.426.559.847.638.555 Naphthalenesulfonates
-	D02.455.426.559.847.638.555.200 Amaranth Dye
-	D02.455.426.559.847.638.555.220 Amido Black
-	D02.455.426.559.847.638.555.282 Anilino Naphthalenesulfonates
-	D02.455.426.559.847.638.555.300 Congo Red
-	D02.455.426.559.847.638.555.400 Evans Blue
-	D02.455.426.559.847.638.555.750 Suramin
-	D02.455.426.559.847.638.555.875 Trypan Blue
-	D02.455.426.559.847.638.638 Naphthols
-	D02.455.426.559.847.638.721 Naphthoquinones
-	D02.455.426.559.847.638.721.374 Vitamin K
-	D02.455.426.559.847.638.721.374.689 Vitamin K 1
-	D02.455.426.559.847.638.721.374.844 Vitamin K 2
-	D02.455.426.559.847.638.721.374.922 Vitamin K 3
-	D02.455.426.559.847.638.845 1-Naphthylamine
-	D02.455.426.559.847.638.845.800 Sertraline
-	D02.455.426.559.847.638.850 2-Naphthylamine
-	D02.455.426.559.847.638.870 1-Naphthylisothiocyanate
-	D02.455.426.559.847.638.900 Naphthylvinylpyridine
-	D02.455.426.559.847.638.930 Pravastatin
-	D02.455.426.559.847.638.945 Propranolol
-	D02.455.426.559.847.638.960 Tetrahydronaphthalenes
-	D02.455.426.559.847.638.960.400 8-Hydroxy-2-(di-n-propylamino)tetralin
-	D02.455.426.559.847.638.960.446 Bunolol
-	D02.455.426.559.847.638.960.446.500 Levobunolol
-	D02.455.426.559.847.638.960.585 Mibefradil
-	D02.455.426.559.847.638.960.675 Podophyllotoxin
-	D02.455.426.559.847.638.960.675.250 Etoposide
-	D02.455.426.559.847.638.960.837 Tetralones
-	D02.455.426.559.847.638.975 Tolnaftate
-	D02.455.426.559.847.680 Phenalenenes
-	D02.455.426.559.847.680.500 Perylene

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.455.426.559.847.723 Phenanthrenes
-	D02.455.426.559.847.723.089 Aristolochic Acids
-	D02.455.426.559.847.723.180 Chrysenes
-	D02.455.426.559.847.723.590 Diterpenes, Abietane
-	D02.455.426.559.847.799 Pyrenes
-	D02.455.426.559.847.799.306 Benzopyrenes
-	D02.455.426.559.847.799.306.300 Benzo(a)pyrene
-	D02.455.426.559.847.799.306.400 Dihydroxydihydrobenzopyrenes
-	D02.455.426.559.847.799.306.400.350 dihydroxybenzo(a)pyrene 9,10-oxide 7,8-Dihydro-7,8-
-	D02.455.426.779 Spiro Compounds
-	D02.455.426.779.120 Buspirone
-	D02.455.426.779.345 Fluorescamine
-	D02.455.426.779.347 Fluoresceins
-	D02.455.426.779.347.300 Eosine I Bluish
-	D02.455.426.779.347.325 Eosine Yellowish-(YS)
-	D02.455.426.779.347.350 Erythrosine
-	D02.455.426.779.347.390 Fluorescein
-	D02.455.426.779.347.400 Fluorescein-5-isothiocyanate
-	D02.455.426.779.347.700 Rose Bengal
-	D02.455.426.779.350 Fluspirilene
-	D02.455.426.779.511 Leucogenenol
-	D02.455.426.779.800 Spiperone
-	D02.455.526 Hydrocarbons, Halogenated
-	D02.455.526.340 Halothane
-	D02.455.526.368 Hydrocarbons, Brominated
-	D02.455.526.368.150 Bromobenzenes
-	D02.455.526.368.175 Bromochlorofluorocarbons
-	D02.455.526.368.200 Bromotrichloromethane
-	D02.455.526.368.225 Ethylene Dibromide
-	D02.455.526.368.700 Polybrominated Biphenyls
-	D02.455.526.439 Hydrocarbons, Chlorinated
-	D02.455.526.439.042 Aldrin
-	D02.455.526.439.150 Carbon Tetrachloride
-	D02.455.526.439.180 Chlordan
-	D02.455.526.439.190 Chlordecone

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.455.526.439.196 Chloroacetates
-	D02.455.526.439.196.500 Dichloroacetic Acid
-	D02.455.526.439.196.750 Trichloroacetic Acid
-	D02.455.526.439.202 Chlorobenzenes
-	D02.455.526.439.202.190 Dicofol
-	D02.455.526.439.202.200 Dinitrochlorobenzene
-	D02.455.526.439.202.400 Hexachlorobenzene
-	D02.455.526.439.220 Chlorofluorocarbons
-	D02.455.526.439.220.149 Bromochlorofluorocarbons
-	D02.455.526.439.220.224 Chlorofluorocarbons, Ethane
-	D02.455.526.439.220.300 Chlorofluorocarbons, Methane
-	D02.455.526.439.224 Chloroform
-	D02.455.526.439.224.200 Bromotrichloromethane
-	D02.455.526.439.255 DDT
-	D02.455.526.439.292 Dichlorodiphenyl Dichloroethylene
-	D02.455.526.439.294 Dichlorodiphenyldichloroethane
-	D02.455.526.439.350 Dichloroethylenes
-	D02.455.526.439.371 Dieldrin
-	D02.455.526.439.416 Endrin
-	D02.455.526.439.447 Ethyl Chloride
-	D02.455.526.439.458 Ethylene Dichlorides
-	D02.455.526.439.516 Heptachlor
-	D02.455.526.439.516.350 Heptachlor Epoxide
-	D02.455.526.439.600 Lindane
-	D02.455.526.439.610 Methoxychlor
-	D02.455.526.439.632 Methyl Chloride
-	D02.455.526.439.642 Methylene Chloride
-	D02.455.526.439.659 Mirex
-	D02.455.526.439.681 Mitotane
-	D02.455.526.439.750 Picryl Chloride
-	D02.455.526.439.773 Polychlorinated Biphenyls
-	D02.455.526.439.773.292 Aroclors
-	D02.455.526.439.773.292.077 Chlorodiphenyl (54% Chlorine)
-	D02.455.526.439.785 Polychloroterphenyl Compounds
-	D02.455.526.439.785.292 Aroclors
-	D02.455.526.439.785.292.077 Chlorodiphenyl (54% Chlorine)

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.455.526.439.880 Tetrachloroethylene
-	D02.455.526.439.913 Toxaphene
-	D02.455.526.439.920 Trichloroepoxypropane
-	D02.455.526.439.927 Trichloroethanes
-	D02.455.526.439.939 Trichloroethylene
-	D02.455.526.439.975 Vinyl Chloride
-	D02.455.526.510 Hydrocarbons, Fluorinated
-	D02.455.526.510.140 Chlorofluorocarbons
-	D02.455.526.510.140.149 Bromochlorofluorocarbons
-	D02.455.526.510.140.224 Chlorofluorocarbons, Ethane
-	D02.455.526.510.140.300 Chlorofluorocarbons, Methane
-	D02.455.526.510.286 Fluoroacetates
-	D02.455.526.510.286.500 Trifluoroacetic Acid
-	D02.455.526.510.432 Fluorobenzenes
-	D02.455.526.510.432.500 Rosuvastatin Calcium
-	D02.455.526.510.435 Fluorocarbons
-	D02.455.526.581 Hydrocarbons, Iodinated
-	D02.455.526.581.247 Iodoacetates
-	D02.455.526.581.247.249 Iodoacetamide
-	D02.455.526.581.247.500 Iodoacetic Acid
-	D02.455.526.581.496 Iodobenzenes
-	D02.455.526.581.496.300 3-Iodobenzylguanidine
-	D02.455.526.581.496.385 Iopanoic Acid
-	D02.455.526.581.496.400 Iophendylate
-	D02.455.526.581.496.450 Ipodate
-	D02.455.526.581.496.950 Tyropanoate
-	D02.455.526.728 Mustard Compounds
-	D02.455.526.728.468 Mustard Gas
-	D02.455.526.728.650 Nitrogen Mustard Compounds
-	D02.455.526.728.650.050 Aniline Mustard
-	D02.455.526.728.650.103 Bendamustine Hydrochloride
-	D02.455.526.728.650.156 Chlorambucil
-	D02.455.526.728.650.156.700 Prednimustine
-	D02.455.526.728.650.275 Estramustine
-	D02.455.526.728.650.463 Mannomustine
-	D02.455.526.728.650.529 Mechlorethamine



## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.455.526.728.650.594                      Melphalan
-	D02.455.526.728.650.594.690                      Peptichemio
-	D02.455.526.728.650.730                      Phosphoramide Mustards
-	D02.455.526.728.650.730.243                      Cyclophosphamide
-	D02.455.526.728.650.730.243.250                      Ifosfamide
-	D02.455.526.728.650.740                      Propylbenzilylcholine Mustard
-	D02.455.526.728.650.760                      Quinacrine Mustard
-	D02.455.526.728.650.913                      Uracil Mustard
-	D02.455.526.913                      Trihalomethanes
-	D02.455.526.913.810                      Chloroform
-	D02.455.612                      Paraffin
-	D02.455.699                      Petrolatum
-	D02.455.699.500                      Mineral Oil
-	D02.455.849                      Terpenes
-	D02.455.849.090                      Cannabinoids
-	D02.455.849.090.100                      Cannabidiol
-	D02.455.849.090.110                      Cannabinol
-	D02.455.849.090.810                      Dronabinol
-	D02.455.849.131                      Carotenoids
-	D02.455.849.131.061                      Abscisic Acid
-	D02.455.849.131.123                      beta Carotene
-	D02.455.849.131.309                      Norisoprenoids
-	D02.455.849.131.495                      Retinoids
-	D02.455.849.131.495.050                      Acitretin
-	D02.455.849.131.495.250                      Etretinate
-	D02.455.849.131.495.270                      Fenretinide
-	D02.455.849.131.495.325                      Isotretinoin
-	D02.455.849.131.495.690                      Retinaldehyde
-	D02.455.849.131.495.818                      Vitamin A
-	D02.455.849.131.495.818.800                      Tretinoin
-	D02.455.849.131.868                      Xanthophylls
-	D02.455.849.131.868.249                      Canthaxanthin
-	D02.455.849.131.868.374                      Cryptoxanthins
New Heading	<b>D02.455.849.131.868.374.500                      Beta-Cryptoxanthin</b>
-	D02.455.849.131.868.500                      Lutein

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.455.849.131.868.750 Zeaxanthins
-	D02.455.849.131.912 zeta Carotene
-	D02.455.849.291 Diterpenes
-	D02.455.849.291.037 Aconitine
-	D02.455.849.291.075 Aphidicolin
-	D02.455.849.291.162 Atractyloside
-	D02.455.849.291.206 Diterpenes, Abietane
-	D02.455.849.291.228 Diterpenes, Clerodane
-	D02.455.849.291.239 Diterpenes, Kaurane
-	D02.455.849.291.239.500 Gibberellins
-	D02.455.849.291.300 Colforsin
-	D02.455.849.291.400 Ginkgolides
-	D02.455.849.291.400.500 Bilobalides
-	D02.455.849.291.500 Phorbols
-	D02.455.849.291.500.510 Phorbol Esters
-	D02.455.849.291.500.510.700 Phorbol 12,13-Dibutyrate
-	D02.455.849.291.500.510.850 Tetradecanoylphorbol Acetate
-	D02.455.849.291.515 Phytanic Acid
-	D02.455.849.291.523 Phytol
-	D02.455.849.291.523.500 Vitamin K
-	D02.455.849.291.523.500.689 Vitamin K 1
-	D02.455.849.291.523.500.844 Vitamin K 2
-	D02.455.849.291.523.500.922 Vitamin K 3
-	D02.455.849.291.686 Ryanodine
-	D02.455.849.291.850 Taxoids
-	D02.455.849.291.850.777 Paclitaxel
-	D02.455.849.291.850.777.500 Albumin-Bound Paclitaxel
-	D02.455.849.365 Dolichol
-	D02.455.849.365.250 Dolichol Phosphates
-	D02.455.849.440 Gefarnate
-	D02.455.849.486 Hemiterpenes
-	D02.455.849.575 Monoterpenes
-	D02.455.849.575.500 Iridoids
-	D02.455.849.575.500.500 Iridoid Glycosides
-	D02.455.849.575.500.500.500 Iridoid Glucosides
-	D02.455.849.575.750 Menthol

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.455.849.575.781 Norbornanes
-	D02.455.849.575.781.234 Bornanes
-	D02.455.849.575.781.234.326 Camphor
-	D02.455.849.575.781.500 Mecamylamine
-	D02.455.849.575.781.795 Toxaphene
-	D02.455.849.575.812 Pyrethrins
-	D02.455.849.575.812.272 Allethrin
-	D02.455.849.575.812.636 Permethrin
-	D02.455.849.575.875 Thymol
-	D02.455.849.575.875.200 Bromthymol Blue
-	D02.455.849.690 Polyisoprenyl Phosphates
-	D02.455.849.690.250 Dolichol Phosphates
-	D02.455.849.690.700 Polyisoprenyl Phosphate Sugars
-	D02.455.849.690.700.700 Polyisoprenyl Phosphate Monosaccharides
-	D02.455.849.690.700.700.250 Dolichol Monophosphate Mannose
-	D02.455.849.690.700.710 Polyisoprenyl Phosphate Oligosaccharides
-	D02.455.849.765 Sesquiterpenes
-	D02.455.849.765.033 Abscisic Acid
-	D02.455.849.765.211 Artemisinins
-	D02.455.849.765.424 Farnesol
-	D02.455.849.765.444 Gossypol
-	D02.455.849.765.750 Santonin
-	D02.455.849.765.775 Sesquiterpenes, Eudesmane
-	D02.455.849.765.787 Sesquiterpenes, Germacrane
-	D02.455.849.765.793 Sesquiterpenes, Guaiane
-	D02.455.849.765.793.888 Thapsigargin
-	D02.455.849.765.850 Trichothecenes
-	D02.455.849.765.850.870 T-2 Toxin
-	D02.455.849.765.850.900 Trichodermin
-	D02.455.849.842 Sesterterpenes
-	D02.455.849.919 Triterpenes
-	D02.455.849.919.138 Cucurbitacins
-	D02.455.849.919.277 Ginsenosides
-	D02.455.849.919.383 Lanosterol
-	D02.455.849.919.490 Limonins
-	D02.455.849.919.530 Pentacyclic Triterpenes

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.455.849.919.530.333 Escin
-	D02.455.849.919.530.444 Glycyrrhetic Acid
-	D02.455.849.919.530.444.250 Carbenoxolone
-	D02.455.849.919.530.466 Glycyrrhizic Acid
-	D02.455.849.919.530.733 Oleanolic Acid
-	D02.455.849.919.570 Quassins
-	D02.455.849.919.570.500 Glaucarubin
-	D02.455.849.919.650 Sapogenins
-	D02.455.849.919.650.600 Oleanolic Acid
-	D02.455.849.919.681 Squalene
-	D02.478 Imides
-	D02.478.400 Imidoesters
-	D02.478.400.175 Dimethyl Adipimidate
-	D02.478.400.200 Dimethyl Suberimidate
-	D02.478.440 Maleimides
-	D02.478.440.418 Ethylmaleimide
-	D02.478.480 Naphthalimides
-	D02.478.520 Pantothenic Acid
-	D02.478.600 Phthalimides
-	D02.478.600.500 Chlorthalidone
-	D02.478.770 Succinimides
-	D02.478.770.150 Bromosuccinimide
-	D02.478.770.333 Ethosuximide
-	D02.491 Imines
-	D02.491.203 Carbodiimides
-	D02.491.203.340 CME-Carbodiimide
-	D02.491.203.362 Cyanamide
-	D02.491.203.385 Dicyclohexylcarbodiimide
-	D02.491.203.425 Ethyldimethylaminopropyl Carbodiimide
-	D02.491.485 Imino Acids
-	D02.491.485.100 Azetidinecarboxylic Acid
-	D02.491.485.400 Technetium Tc 99m Diethyl-iminodiacetic Acid
-	D02.491.485.450 Technetium Tc 99m Disofenin
-	D02.491.485.900 Technetium Tc 99m Lidofenin
-	D02.491.567 Imino Sugars
-	D02.491.567.249 Imino Furanoses

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.491.567.500 Imino Pyranoses
-	D02.491.567.500.033 1-Deoxynojirimycin
-	D02.491.650 Polyethyleneimine
-	D02.491.784 Schiff Bases
-	D02.500 Isocyanates
-	D02.500.375 Isothiocyanates
-	D02.500.375.050 4-Acetamido-4'-isothiocyanatostilbene-2,2'-disulfonic Acid
-	D02.500.375.125 4,4'-Diisothiocyanostilbene-2,2'-Disulfonic Acid
-	D02.500.375.250 Fluorescein-5-isothiocyanate
-	D02.500.375.625 1-Naphthylisothiocyanate
-	D02.500.800 Toluene 2,4-Diisocyanate
-	D02.522 Ketones
-	D02.522.064 Acetone
-	D02.522.120 Acetophenones
-	D02.522.120.100 Benzoin
-	D02.522.120.575 omega-Chloroacetophenone
-	D02.522.223 Benzophenones
-	D02.522.223.500 Chlorthalidone
-	D02.522.223.750 Fenofibrate
-	D02.522.296 Butanones
-	D02.522.296.100 Acetoin
-	D02.522.296.400 Diacetyl
-	D02.522.296.900 Thenoyltrifluoroacetone
-	D02.522.352 Butyrophenones
-	D02.522.352.110 Azaperone
-	D02.522.352.155 Benperidol
-	D02.522.352.343 Droperidol
-	D02.522.352.506 Haloperidol
-	D02.522.352.800 Spiperone
-	D02.522.352.867 Trifluoperidol
-	D02.522.376 Camphor
-	D02.522.400 Cyclohexanones
-	D02.522.400.850 Tiletamine
-	D02.522.520 Hexanones
-	D02.522.520.500 Methyl n-Butyl Ketone
-	D02.522.585 Ketone Bodies

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.522.585.087 3-Hydroxybutyric Acid
-	D02.522.585.175 Acetoacetates
-	D02.522.650 Mannich Bases
-	D02.522.675 Methadone
-	D02.522.675.500 Methadyl Acetate
-	D02.522.720 Pentanones
-	D02.522.741 Phosgene
-	D02.522.818 Propiophenones
-	D02.522.818.110 Bupropion
-	D02.522.818.222 Chalcones
-	D02.522.818.222.500 Chalcone
-	D02.522.818.561 Hydroxypropiophenone
-	D02.522.818.561.602 Phloretin
-	D02.522.818.561.602.615 Polyphloretin Phosphate
-	D02.522.818.600 Kynuramine
-	D02.522.818.800 Oxymfedrine
-	D02.522.818.830 Propafenone
-	D02.522.818.950 Tolperisone
-	D02.540 Lactones
-	D02.540.150 4-Butyrolactone
-	D02.540.205 Acetogenins
-	D02.540.232 Acyl-Butyrolactones
-	D02.540.260 Dehydroascorbic Acid
-	D02.540.505 Macrolides
-	D02.540.505.450 Lucensomycin
-	D02.540.505.495 Maytansine
-	D02.540.505.497 Mepartricin
-	D02.540.505.500 Miocamycin
-	D02.540.505.550 Natamycin
-	D02.540.505.575 Nystatin
-	D02.540.505.600 Oleandomycin
-	D02.540.505.600.800 Troleandomycin
-	D02.540.505.620 Oligomycins
-	D02.540.505.620.750 Rutamycin
-	D02.540.505.760 Sirolimus
-	D02.540.505.760.500 Everolimus

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.540.505.810 Tacrolimus
-	D02.540.505.905 Tylosin
-	D02.540.552 Picrotoxin
-	D02.540.576 Polyketides
-	D02.540.576.500 Macrolides
-	D02.540.576.500.500 Amphotericin B
-	D02.540.576.500.750 Antimycin A
-	D02.540.576.500.875 Brefeldin A
-	D02.540.576.500.937 Bryostatins
-	D02.540.576.500.968 Candicidin
-	D02.540.576.500.984 Epothilones
-	D02.540.576.500.992 Erythromycin
-	D02.540.576.500.992.050 Azithromycin
-	D02.540.576.500.992.100 Clarithromycin
-	D02.540.576.500.992.250 Erythromycin Estolate
-	D02.540.576.500.992.260 Erythromycin Ethylsuccinate
-	D02.540.576.500.992.445 Ketolides
-	D02.540.576.500.992.630 Roxithromycin
-	D02.540.576.500.996 Filipin
-	D02.540.576.500.997 Ivermectin
-	D02.540.576.500.998 Josamycin
-	D02.540.576.500.999 Leucomycins
-	D02.540.576.500.999.450 Kitasamycin
-	D02.540.576.500.999.725 Spiramycin
-	D02.540.600 Propiolactone
-	D02.540.679 Spironolactone
-	D02.540.800 Venturicidins
-	D02.540.950 Zearalenone
-	D02.540.950.975 Zeranol
-	D02.561 Lewis Acids
-	D02.572 Lewis Bases
-	D02.583 Nitrates
-	D02.583.225 Erythrityl Tetranitrate
-	D02.583.575 Nicorandil
-	D02.626 Nitriles
-	D02.626.080 Acetonitriles

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.626.080.085 Aminoacetonitrile
-	D02.626.095 Acrylonitrile
-	D02.626.151 Aminopropionitrile
-	D02.626.175 Amygdalin
-	D02.626.260 Carbonyl Cyanide m-Chlorophenyl Hydrazone
-	D02.626.270 Carbonyl Cyanide p-Trifluoromethoxyphenylhydrazone
-	D02.626.290 Cyanoacrylates
-	D02.626.290.200 Bucrylate
-	D02.626.290.350 Enbucrilate
-	D02.626.310 o-Chlorobenzylidenemalonitrile
-	D02.626.320 Citalopram
-	D02.626.400 Fadrozole
-	D02.626.745 Pregnenolone Carbonitrile
-	D02.626.809 Rilpivirine
-	D02.626.809.500 Emtricitabine, Rilpivirine, Tenofovir Drug Combination
-	D02.626.872 Technetium Tc 99m Sestamibi
-	D02.626.886 Tyrphostins
-	D02.633 Nitrites
-	D02.633.025 Amyl Nitrite
-	D02.640 Nitro Compounds
-	D02.640.139 Aristolochic Acids
-	D02.640.280 Dinitrocresols
-	D02.640.529 Nitrobenzenes
-	D02.640.529.175 Chloramphenicol
-	D02.640.529.175.850 Thiamphenicol
-	D02.640.529.240 Dinitrobenzenes
-	D02.640.529.240.250 Dinitrochlorobenzene
-	D02.640.529.240.280 Dinitrofluorobenzene
-	D02.640.529.880 Trinitrobenzenes
-	D02.640.529.880.607 Picryl Chloride
-	D02.640.529.880.900 Trinitrobenzenesulfonic Acid
-	D02.640.600 Nitrofurans
-	D02.640.600.200 5-Amino-3-((5-nitro-2-furyl)vinyl)-1,2,4-oxadiazole
-	D02.640.600.290 FANFT
-	D02.640.600.308 Furagin
-	D02.640.600.313 Furazolidone



## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.640.600.330 Furfurylamide
-	D02.640.600.410 Nifuratel
-	D02.640.600.480 Nifurtimox
-	D02.640.600.566 Nitrofurantoin
-	D02.640.600.691 Nitrofurazone
-	D02.640.600.750 Nitrovin
-	D02.640.636 Nitroglycerin
-	D02.640.672 Nitroimidazoles
-	D02.640.672.200 Dimetridazole
-	D02.640.672.250 Etanidazole
-	D02.640.672.400 Ipronidazole
-	D02.640.672.500 Metronidazole
-	D02.640.672.530 Misonidazole
-	D02.640.672.580 Nimorazole
-	D02.640.672.700 Ornidazole
-	D02.640.672.730 Ronidazole
-	D02.640.672.900 Tinidazole
-	D02.640.720 Nitroparaffins
-	D02.640.720.800 Tetranitromethane
-	D02.640.743 Nitrophenols
-	D02.640.743.304 Dinitrophenols
-	D02.640.743.304.250 2,4-Dinitrophenol
-	D02.640.743.400 2-Hydroxy-5-nitrobenzyl Bromide
-	D02.640.743.600 Niclofolan
-	D02.640.743.610 Nitrohydroxyiodophenylacetate
-	D02.640.743.620 Nitroxinil
-	D02.640.743.690 Picrates
-	D02.640.820 Nitroquinolines
-	D02.640.820.600 4-Nitroquinoline-1-oxide
-	D02.654 Nitroso Compounds
-	D02.654.442 Nitrosamines
-	D02.654.442.100 Butylhydroxybutylnitrosamine
-	D02.654.442.200 Diethylnitrosamine
-	D02.654.442.225 Dimethylnitrosamine
-	D02.654.442.550 N-Nitrosopyrrolidine
-	D02.654.567 Nitrosoguanidines

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.654.567.400 Methylnitronitrosoguanidine
-	D02.654.585 Nitrosomethylurethane
-	D02.654.692 Nitrosoarea Compounds
-	D02.654.692.247 Carmustine
-	D02.654.692.300 Ethylnitrosoarea
-	D02.654.692.440 Lomustine
-	D02.654.692.440.700 Semustine
-	D02.654.692.480 Methylnitrosoarea
-	D02.654.692.550 Nimustine
-	D02.654.692.768 Streptozocin
-	D02.654.846 Reactive Nitrogen Species
-	D02.654.846.500 S-Nitrosothiols
-	D02.654.846.500.124 S-Nitroso-N-Acetylpenicillamine
-	D02.654.846.500.249 S-Nitrosoglutathione
-	D02.675 Onium Compounds
-	D02.675.276 Quaternary Ammonium Compounds
-	D02.675.276.046 Ambenonium Chloride
-	D02.675.276.080 Benzalkonium Compounds
-	D02.675.276.090 Benzethonium
-	D02.675.276.102 Bephenium Compounds
-	D02.675.276.125 Betaine
-	D02.675.276.136 Betalains
-	D02.675.276.136.500 Betacyanins
-	D02.675.276.136.555 Betaxanthins
-	D02.675.276.148 Bethanechol Compounds
-	D02.675.276.148.100 Bethanechol
Old Tree	<b>D02.675.276.175 Bretylium Compounds</b>
Old Tree	<b>D02.675.276.175.150 Bretylium Tosylate</b>
-	D02.675.276.190 Cetrimonium Compounds
-	D02.675.276.200 Chlorisondamine
-	D02.675.276.207 Chlormequat
-	D02.675.276.210 (4-(m-Chlorophenylcarbamoyloxy)-2-butynyl)trimethylammonium Chloride
-	D02.675.276.232 Choline
-	D02.675.276.232.100 Benzoylcholine
-	D02.675.276.232.115 Carbachol

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.675.276.232.130 Cytidine Diphosphate Choline
-	D02.675.276.232.700 Phosphorylcholine
-	D02.675.276.232.710 Platelet Activating Factor
-	D02.675.276.232.720 Propylbenzylcholine Mustard
-	D02.675.276.232.780 Succinylcholine
-	D02.675.276.232.800 Thiocholine
-	D02.675.276.232.800.030 Acetylthiocholine
-	D02.675.276.232.800.200 Butyrylthiocholine
-	D02.675.276.352 Edrophonium
-	D02.675.276.370 Emepronium
-	D02.675.276.400 Gallamine Triethiodide
-	D02.675.276.425 Glycopyrrolate
-	D02.675.276.435 Hemicholinium 3
-	D02.675.276.475 Lissamine Green Dyes
-	D02.675.276.534 Methacholine Compounds
-	D02.675.276.534.500 Methacholine Chloride
-	D02.675.276.558 Bis-Trimethylammonium Compounds
-	D02.675.276.558.372 Decamethonium Compounds
-	D02.675.276.558.592 Hexamethonium Compounds
-	D02.675.276.558.592.500 Hexamethonium
-	D02.675.276.580 Muscarine
-	D02.675.276.602 Neostigmine
-	D02.675.276.648 Oxyphenonium
-	D02.675.276.700 Propantheline
-	D02.675.276.787 Tetraethylammonium Compounds
-	D02.675.276.787.500 Tetraethylammonium
-	D02.675.276.844 Toxiferine
-	D02.675.276.844.050 Alcuronium
-	D02.675.276.922 Tubocurarine
-	D02.675.800 Sulfonium Compounds
-	D02.691 Organometallic Compounds
-	D02.691.075 Antimony Potassium Tartrate
-	D02.691.080 Antimony Sodium Gluconate
-	D02.691.088 Arsenicals
-	D02.691.088.030 Arsanilic Acid
-	D02.691.088.040 Arsenamide

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.691.088.042 Arsenates
-	D02.691.088.044 Arsenazo III
-	D02.691.088.047 Arsenites
-	D02.691.088.050 Arsphenamine
-	D02.691.088.060 p-Azobenzene arsonate
-	D02.691.088.100 Cacodylic Acid
-	D02.691.088.500 Melarsoprol
-	D02.691.088.700 Roxarsone
-	D02.691.214 Coordination Complexes
-	D02.691.400 Gadolinium DTPA
-	D02.691.550 Iron Compounds
-	D02.691.550.100 Ferric Compounds
-	D02.691.550.100.500 Iron-Dextran Complex
-	D02.691.550.200 Ferrous Compounds
-	D02.691.550.500 Iron Carbonyl Compounds
-	D02.691.550.600 Iron, Dietary
-	D02.691.600 Maneb
-	D02.691.675 Organogold Compounds
-	D02.691.675.249 Aurothioglucose
-	D02.691.675.249.150 Auranofin
-	D02.691.675.500 Gold Sodium Thiomalate
-	D02.691.750 Organomercury Compounds
-	D02.691.750.100 Alkylmercury Compounds
-	D02.691.750.100.229 Chlormerodrin
-	D02.691.750.100.347 Ethylmercury Compounds
-	D02.691.750.100.347.350 Cialit
-	D02.691.750.100.347.417 Ethylmercuric Chloride
-	D02.691.750.100.347.850 Thimerosal
-	D02.691.750.100.710 Mersalyl
-	D02.691.750.100.738 Methylmercury Compounds
-	D02.691.750.575 Merbromin
-	D02.691.750.740 Phenylmercury Compounds
-	D02.691.750.740.220 4-Chloromercuribenzenesulfonate
-	D02.691.750.740.225 Chloromercurinitrophenols
-	D02.691.750.740.644 Mercuribenzoates
-	D02.691.750.740.644.261 Chloromercuribenzoates

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.691.750.740.644.261.275 p-Chloromercuribenzoic Acid
-	D02.691.750.740.644.450 Hydroxymercuribenzoates
-	D02.691.750.740.760 Phenylmercuric Acetate
-	D02.691.800 Organoplatinum Compounds
-	D02.691.800.338 Carboplatin
-	D02.691.825 Organotechnetium Compounds
-	D02.691.825.375 Technetium Tc 99m Aggregated Albumin
-	D02.691.825.445 Technetium Tc 99m Diethyl-iminodiacetic Acid
-	D02.691.825.468 Technetium Tc 99m Dimercaptosuccinic Acid
-	D02.691.825.475 Technetium Tc 99m Disofenin
-	D02.691.825.562 Technetium Tc 99m Exametazime
-	D02.691.825.710 Technetium Tc 99m Lidofenin
-	D02.691.825.750 Technetium Tc 99m Medronate
-	D02.691.825.775 Technetium Tc 99m Mertiatide
-	D02.691.825.875 Technetium Tc 99m Pentetate
-	D02.691.825.937 Technetium Tc 99m Sestamibi
-	D02.691.850 Organotin Compounds
-	D02.691.850.900 Trialkyltin Compounds
-	D02.691.850.900.910 Triethyltin Compounds
-	D02.691.850.900.950 Trimethyltin Compounds
-	D02.691.925 Sucralfate
-	D02.691.950 Tetraethyl Lead
-	D02.691.975 Zineb
-	D02.705 Organophosphorus Compounds
-	D02.705.400 Organophosphates
-	D02.705.400.100 Carbamyl Phosphate
-	D02.705.400.120 Chlorfenvinphos
-	D02.705.400.130 Dichlorvos
-	D02.705.400.140 Diphosphoglyceric Acids
-	D02.705.400.140.175 2,3-Diphosphoglycerate
-	D02.705.400.501 Mevinphos
-	D02.705.400.550 Monocrotophos
-	D02.705.400.600 Naled
-	D02.705.400.625 Organothiophosphates
-	D02.705.400.625.050 Amifostine
-	D02.705.400.625.060 Azinphosmethyl

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.705.400.625.100 Chlorpyrifos
-	D02.705.400.625.120 Coumaphos
-	D02.705.400.625.130 Cystaphos
-	D02.705.400.625.200 Diazinon
-	D02.705.400.625.220 Dimethoate
-	D02.705.400.625.230 Disulfoton
-	D02.705.400.625.240 Echothiophate Iodide
-	D02.705.400.625.300 Fenitrothion
-	D02.705.400.625.320 Fenthion
-	D02.705.400.625.500 Malathion
-	D02.705.400.625.600 Parathion
-	D02.705.400.625.600.500 Methyl Parathion
-	D02.705.400.625.650 Phorate
-	D02.705.400.625.670 Phosmet
-	D02.705.400.625.800 Temefos
-	D02.705.400.650 Paraoxon
-	D02.705.400.700 Phosphamidon
-	D02.705.400.725 Polyisoprenyl Phosphates
-	D02.705.400.725.200 Dolichol Phosphates
-	D02.705.400.750 Tetrachlorvinphos
-	D02.705.400.875 Tritolyl Phosphates
-	D02.705.429 Organophosphonates
-	D02.705.429.249 Aminoethylphosphonic Acid
-	D02.705.429.374 Armin
-	D02.705.429.500 Diphosphonates
-	D02.705.429.500.100 Alendronate
-	D02.705.429.500.200 Clodronic Acid
-	D02.705.429.500.830 Etidronic Acid
-	D02.705.429.500.830.500 Risedronate Sodium
-	D02.705.429.500.885 Technetium Tc 99m Medronate
-	D02.705.429.625 Fosfomycin
-	D02.705.429.750 Organofluorophosphonates
-	D02.705.429.750.249 Isofluorophate
-	D02.705.429.750.500 Sarin
-	D02.705.429.750.750 Soman
-	D02.705.429.812 Organothiophosphonates

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.705.429.812.249 Fonofos
-	D02.705.429.812.500 Leptophos
-	D02.705.429.812.750 Phenylphosphonothioic Acid, 2-Ethyl 2-(4-Nitrophenyl) Ester
-	D02.705.429.875 Phosphonoacetic Acid
-	D02.705.429.875.500 Foscarnet
-	D02.705.429.906 Tenofovir
-	D02.705.429.906.125 Efavirenz, Emtricitabine, Tenofovir Disoproxil Fumarate Drug Combination
-	D02.705.429.906.250 Elvitegravir, Cobicistat, Emtricitabine, Tenofovir Disoproxil Fumarate Drug Combination
-	D02.705.429.906.500 Emtricitabine, Rilpivirine, Tenofovir Drug Combination
-	D02.705.429.906.750 Emtricitabine, Tenofovir Disoproxil Fumarate Drug Combination
-	D02.705.429.937 Trichlorfon
-	D02.705.539 Organothiophosphorus Compounds
-	D02.705.539.345 Organothiophosphates
-	D02.705.539.345.050 Amifostine
-	D02.705.539.345.060 Azinphosmethyl
-	D02.705.539.345.100 Chlorpyrifos
-	D02.705.539.345.120 Coumaphos
-	D02.705.539.345.130 Cystaphos
-	D02.705.539.345.200 Diazinon
-	D02.705.539.345.220 Dimethoate
-	D02.705.539.345.230 Disulfoton
-	D02.705.539.345.240 Echothiophate Iodide
-	D02.705.539.345.300 Fenitrothion
-	D02.705.539.345.320 Fenthion
-	D02.705.539.345.500 Malathion
-	D02.705.539.345.600 Parathion
-	D02.705.539.345.600.500 Methyl Parathion
-	D02.705.539.345.650 Phorate
-	D02.705.539.345.670 Phosmet
-	D02.705.539.345.800 Temefos
-	D02.705.539.692 Organothiophosphonates
-	D02.705.539.692.249 Fonofos
-	D02.705.539.692.500 Leptophos

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.705.539.692.750 Phenylphosphonothioic Acid, 2-Ethyl 2-(4-Nitrophenyl) Ester
-	D02.705.621 Phosphines
-	D02.705.629 Phosphinic Acids
-	D02.705.629.500 Fosinopril
-	D02.705.672 Phosphoramides
-	D02.705.672.249 Hempa
-	D02.705.672.500 Phosphoramide Mustards
-	D02.705.672.500.243 Cyclophosphamide
-	D02.705.672.500.243.250 Ifosfamide
-	D02.705.672.750 Tetraisopropylpyrophosphamide
-	D02.705.672.875 Thiotepa
-	D02.705.675 Phosphoranes
-	D02.712 Organoselenium Compounds
-	D02.712.600 Selenocysteine
-	D02.712.700 Selenomethionine
-	D02.715 Organosilicon Compounds
-	D02.715.312 Organically Modified Ceramics
-	D02.715.625 Silanes
-	D02.715.650 Siloxanes
-	D02.715.650.700 Silicones
-	D02.715.650.700.150 Dimethylpolysiloxanes
-	D02.715.650.700.150.750 Simethicone
-	D02.715.650.700.755 Silicone Gels
-	D02.715.650.700.800 Silicone Oils
-	D02.715.650.850 Silorane Resins
-	D02.715.715 Trimethylsilyl Compounds
-	D02.806 Quinones
-	D02.806.100 Anthraquinones
-	D02.806.100.200 Carmine
-	D02.806.100.205 Cascara
-	D02.806.100.205.400 Emodin
-	D02.806.100.500 Mitoxantrone
-	D02.806.100.750 Senna Extract
-	D02.806.250 Benzoquinones
-	D02.806.250.180 Carbazilquinone



## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.806.250.210 Chloranil
-	D02.806.250.280 Dibromothymoquinone
-	D02.806.250.700 Plastoquinone
-	D02.806.250.900 Ubiquinone
-	D02.806.400 Indolequinones
-	D02.806.400.249 Mitomycins
-	D02.806.400.249.350 Mitomycin
-	D02.806.400.249.700 Porfiromycin
-	D02.806.400.500 Pyrroloiminoquinones
-	D02.806.550 Naphthoquinones
-	D02.806.550.312 Atovaquone
-	D02.806.550.625 Vitamin K 1
-	D02.806.550.750 Vitamin K 2
-	D02.806.550.875 Vitamin K 3
-	D02.825 Rotaxanes
-	D02.845 Semicarbazides
-	D02.845.250 Diphenylcarbazine
-	D02.845.746 Semicarbazones
-	D02.845.746.703 Thiosemicarbazones
-	D02.845.746.703.450 Methisazone
-	D02.845.746.703.702 Thioacetazone
-	D02.886 Sulfur Compounds
-	D02.886.030 Amino Acids, Sulfur
-	D02.886.030.175 Cystathionine
-	D02.886.030.215 Cysteic Acid
-	D02.886.030.230 Cysteine
-	D02.886.030.230.259 Acetylcysteine
-	D02.886.030.230.310 Carbocysteine
-	D02.886.030.230.330 Cysteinyldopa
-	D02.886.030.230.369 Cystine
-	D02.886.030.230.700 Selenocysteine
-	D02.886.030.388 Ethionine
-	D02.886.030.498 Homocysteine
-	D02.886.030.498.050 S-Adenosylhomocysteine
-	D02.886.030.554 Homocystine
-	D02.886.030.676 Methionine

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.886.030.676.089 Racemethionine
-	D02.886.030.676.180 S-Adenosylmethionine
-	D02.886.030.676.450 N-Formylmethionine
-	D02.886.030.676.450.440 N-Formylmethionine Leucyl-Phenylalanine
-	D02.886.030.676.620 Methionine Sulfoximine
-	D02.886.030.676.620.125 Buthionine Sulfoximine
-	D02.886.030.676.900 Selenomethionine
-	D02.886.030.676.950 Vitamin U
-	D02.886.030.786 Penicillamine
-	D02.886.030.786.500 S-Nitroso-N-Acetylpenicillamine
-	D02.886.030.893 Thiorphan
-	D02.886.030.896 Tiopronin
-	D02.886.092 Benzothiepins
-	D02.886.092.408 Endosulfan
-	D02.886.108 beta-Lactams
-	D02.886.108.500 Monobactams
-	D02.886.108.500.044 Aztreonam
-	D02.886.108.750 Penicillins
-	D02.886.108.750.124 Amdinocillin
-	D02.886.108.750.124.036 Amdinocillin Pivoxil
-	D02.886.108.750.249 Cyclacillin
-	D02.886.108.750.500 Methicillin
-	D02.886.108.750.562 Nafcillin
-	D02.886.108.750.625 Oxacillin
-	D02.886.108.750.625.150 Cloxacillin
-	D02.886.108.750.625.150.205 Dicloxacillin
-	D02.886.108.750.625.150.250 Floxacillin
-	D02.886.108.750.687 Penicillanic Acid
-	D02.886.108.750.750 Penicillin G
-	D02.886.108.750.750.050 Ampicillin
-	D02.886.108.750.750.050.050 Amoxicillin
-	D02.886.108.750.750.050.050.060 Amoxicillin-Potassium Clavulanate Combination
-	D02.886.108.750.750.050.075 Azlocillin
-	D02.886.108.750.750.050.500 Mezlocillin
-	D02.886.108.750.750.050.650 Piperacillin

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.886.108.750.750.050.700 Pivampicillin
-	D02.886.108.750.750.050.900 Talampicillin
-	D02.886.108.750.750.170 Carbenicillin
-	D02.886.108.750.750.170.200 Carfecillin
-	D02.886.108.750.750.685 Penicillin G Benzathine
-	D02.886.108.750.750.695 Penicillin G Procaine
-	D02.886.108.750.750.875 Sulbenicillin
-	D02.886.108.750.781 Penicillin V
-	D02.886.108.750.812 Sulbactam
-	D02.886.108.750.875 Ticarcillin
-	D02.886.124 Cyclic S-Oxides
-	D02.886.250 Isothiocyanates
-	D02.886.250.050 4-Acetamido-4'-isothiocyanatostilbene-2,2'-disulfonic Acid
-	D02.886.250.125 4,4'-Diisothiocyanatostilbene-2,2'-Disulfonic Acid
-	D02.886.250.250 Fluorescein-5-isothiocyanate
-	D02.886.250.625 1-Naphthylisothiocyanate
-	D02.886.300 Organothiophosphorus Compounds
-	D02.886.300.692 Organothiophosphates
-	D02.886.300.692.050 Amifostine
-	D02.886.300.692.060 Azinphosmethyl
-	D02.886.300.692.100 Chlorpyrifos
-	D02.886.300.692.120 Coumaphos
-	D02.886.300.692.130 Cystaphos
-	D02.886.300.692.200 Diazinon
-	D02.886.300.692.220 Dimethoate
-	D02.886.300.692.230 Disulfoton
-	D02.886.300.692.240 Echothiophate Iodide
-	D02.886.300.692.300 Fenitrothion
-	D02.886.300.692.320 Fenthion
-	D02.886.300.692.500 Malathion
-	D02.886.300.692.600 Parathion
-	D02.886.300.692.600.500 Methyl Parathion
-	D02.886.300.692.650 Phorate
-	D02.886.300.692.670 Phosmet
-	D02.886.300.692.800 Temefos
-	D02.886.300.846 Organothiophosphonates

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.886.300.846.249 Fonofos
-	D02.886.300.846.500 Leptophos
-	D02.886.300.846.750 Phenylphosphonothioic Acid, 2-Ethyl 2-(4-Nitrophenyl) Ester
-	D02.886.369 Phenothiazines
-	D02.886.369.029 Acepromazine
-	D02.886.369.080 Azure Stains
-	D02.886.369.198 Chlorpromazine
-	D02.886.369.326 Fluphenazine
-	D02.886.369.470 Mesoridazine
-	D02.886.369.513 Methotrimeprazine
-	D02.886.369.517 Methylene Blue
-	D02.886.369.533 Moricizine
-	D02.886.369.550 Nonachlazine
-	D02.886.369.575 Perazine
-	D02.886.369.593 Perphenazine
-	D02.886.369.639 Prochlorperazine
-	D02.886.369.661 Promazine
-	D02.886.369.670 Promethazine
-	D02.886.369.780 Thiethylperazine
-	D02.886.369.843 Thioridazine
-	D02.886.369.869 Tolonium Chloride
-	D02.886.369.898 Trifluoperazine
-	D02.886.369.918 Triflupromazine
-	D02.886.369.939 Trimeprazine
-	D02.886.489 Sulfhydryl Compounds
-	D02.886.489.155 Cysteine
-	D02.886.489.180 Dimercaprol
-	D02.886.489.180.900 Unithiol
-	D02.886.489.225 Ergothioneine
-	D02.886.489.409 Mercaptoethanol
-	D02.886.489.472 Mercaptoethylamines
-	D02.886.489.472.369 Cysteamine
-	D02.886.489.520 3-Mercaptopropionic Acid
-	D02.886.489.534 6-Mercaptopurine
-	D02.886.489.590 Mesna

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.886.489.600 Methimazole
-	D02.886.489.650 S-Nitrosothiols
-	D02.886.489.650.500 S-Nitroso-N-Acetylpenicillamine
-	D02.886.489.650.750 S-Nitrosoglutathione
-	D02.886.489.700 Pantetheine
-	D02.886.489.750 Succimer
-	D02.886.489.750.725 Technetium Tc 99m Dimercaptosuccinic Acid
-	D02.886.489.789 Thiocholine
-	D02.886.489.789.030 Acetylthiocholine
-	D02.886.489.789.200 Butyrylthiocholine
-	D02.886.489.828 Thioglycolates
-	D02.886.489.891 Thiomalates
-	D02.886.489.891.451 Gold Sodium Thiomalate
-	D02.886.520 Sulfides
-	D02.886.520.150 Disulfides
-	D02.886.520.150.087 Cystine
-	D02.886.520.150.175 Disulfiram
-	D02.886.520.575 Moxalactam
-	D02.886.590 Sulfones
-	D02.886.590.192 Dansyl Compounds
-	D02.886.590.263 Dapsone
-	D02.886.590.263.050 Acedapsone
-	D02.886.590.625 Phenylmethylsulfonyl Fluoride
-	D02.886.590.700 Sulfonamides
-	D02.886.590.700.120 Benzamide
-	D02.886.590.700.135 Benzothiadiazines
-	D02.886.590.700.135.138 Bendroflumethiazide
-	D02.886.590.700.135.261 Chlorothiazide
-	D02.886.590.700.135.261.476 Hydrochlorothiazide
-	D02.886.590.700.135.261.476.716 Trichlormethiazide
-	D02.886.590.700.135.285 Cyclopenthiiazide
-	D02.886.590.700.135.300 Diazoxide
-	D02.886.590.700.135.475 Hydroflumethiazide
-	D02.886.590.700.135.620 Methyclothiazide
-	D02.886.590.700.135.781 Polythiazide
-	D02.886.590.700.150 Bumetanide

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D02.886.590.700.247	Celecoxib
Old Tree	D02.886.590.700.344	Chloramines
-	D02.886.590.700.365	Chlorthalidone
-	D02.886.590.700.390	Clopamide
-	D02.886.590.700.400	Darunavir
-	D02.886.590.700.410	Dichlorphenamide
-	D02.886.590.700.420	Ethoxzolamide
-	D02.886.590.700.540	Indapamide
-	D02.886.590.700.545	1-(5-Isoquinolinesulfonyl)-2-Methylpiperazine
-	D02.886.590.700.570	Mafenide
-	D02.886.590.700.575	Mefruside
-	D02.886.590.700.600	Metolazone
-	D02.886.590.700.625	Probenecid
-	D02.886.590.700.650	Rosuvastatin Calcium
-	D02.886.590.700.675	Sildenafil Citrate
-	D02.886.590.700.700	Simeprevir
-	D02.886.590.700.725	Sulfanilamides
-	D02.886.590.700.725.300	Furosemide
-	D02.886.590.700.725.650	Sulfacetamide
-	D02.886.590.700.725.700	Sulfachlorpyridazine
-	D02.886.590.700.725.755	Sulfadiazine
-	D02.886.590.700.725.755.800	Silver Sulfadiazine
-	D02.886.590.700.725.760	Sulfadimethoxine
-	D02.886.590.700.725.765	Sulfadoxine
-	D02.886.590.700.725.810	Sulfaguanidine
-	D02.886.590.700.725.850	Sulfalene
-	D02.886.590.700.725.853	Sulfamerazine
-	D02.886.590.700.725.857	Sulfameter
-	D02.886.590.700.725.862	Sulfamethazine
-	D02.886.590.700.725.867	Sulfamethoxazole
-	D02.886.590.700.725.867.500 Combination	Trimethoprim, Sulfamethoxazole Drug Combination
-	D02.886.590.700.725.872	Sulfamethoxypyridazine
-	D02.886.590.700.725.877	Sulfamonomethoxine
-	D02.886.590.700.725.882	Sulfamoxole
-	D02.886.590.700.725.887	Sulfaphenazole

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.886.590.700.725.900 Sulfapyridine
-	D02.886.590.700.725.925 Sulfaquinoxaline
-	D02.886.590.700.725.935 Sulfathiazoles
-	D02.886.590.700.725.935.813 Sulfamethizole
-	D02.886.590.700.725.940 Sulfisomidine
-	D02.886.590.700.725.945 Sulfisoxazole
-	D02.886.590.700.730 Sulfasalazine
-	D02.886.590.700.750 Sumatriptan
-	D02.886.590.700.975 Xipamide
-	D02.886.590.795 Sulfonyleurea Compounds
-	D02.886.590.795.071 Acetohexamide
-	D02.886.590.795.204 Carbutamide
-	D02.886.590.795.283 Chlorpropamide
-	D02.886.590.795.475 Gliclazide
-	D02.886.590.795.500 Glipizide
-	D02.886.590.795.575 Glyburide
-	D02.886.590.795.834 Tolazamide
-	D02.886.590.795.896 Tolbutamide
-	D02.886.590.887 Tosyl Compounds
-	D02.886.590.887.140 Bromcresol Green
-	D02.886.590.887.150 Bromcresol Purple
-	D02.886.590.887.180 Bromphenol Blue
-	D02.886.590.887.200 Bromthymol Blue
-	D02.886.590.887.570 Tosylarginine Methyl Ester
-	D02.886.590.887.600 Tosyllysine Chloromethyl Ketone
-	D02.886.590.887.660 Tosylphenylalanyl Chloromethyl Ketone
-	D02.886.620 Sulfonium Compounds
-	D02.886.640 Sulfoxides
-	D02.886.640.074 2-Pyridinylmethylsulfinylbenzimidazoles
-	D02.886.640.074.249 Lansoprazole
-	D02.886.640.074.249.500 Dexlansoprazole
-	D02.886.640.074.500 Omeprazole
-	D02.886.640.074.500.500 Esomeprazole
-	D02.886.640.074.750 Rabeprazole
-	D02.886.640.150 Dimethyl Sulfoxide
-	D02.886.645 Sulfur Acids

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.886.645.575 Sulfenic Acids
-	D02.886.645.585 Sulfinic Acids
-	D02.886.645.600 Sulfonic Acids
-	D02.886.645.600.055 Alkanesulfonic Acids
-	D02.886.645.600.055.050 Alkanesulfonates
-	D02.886.645.600.055.050.500 Mesna
-	D02.886.645.600.055.050.510 Mesylates
-	D02.886.645.600.055.050.510.100 Busulfan
-	D02.886.645.600.055.050.510.300 Ethyl Methanesulfonate
-	D02.886.645.600.055.050.510.500 Methyl Methanesulfonate
-	D02.886.645.600.055.050.632 Sodium Dodecyl Sulfate
-	D02.886.645.600.055.050.755 Sodium Tetradecyl Sulfate
-	D02.886.645.600.055.250 HEPES
-	D02.886.645.600.055.300 Isethionic Acid
-	D02.886.645.600.055.850 Taurine
-	D02.886.645.600.055.850.800 Taurocholic Acid
-	D02.886.645.600.055.850.800.900 Taurodeoxycholic Acid
-	D02.886.645.600.055.850.800.900.875 Taurochenodeoxycholic Acid
-	D02.886.645.600.055.850.800.925 Tauroolithocholic Acid
-	D02.886.645.600.080 Arylsulfonic Acids
-	D02.886.645.600.080.050 Arylsulfonates
-	D02.886.645.600.080.050.100 Benzenesulfonates
-	D02.886.645.600.080.050.100.075 Calcium Dobesilate
-	D02.886.645.600.080.050.100.100 4-Chloromercuribenzenesulfonate
-	D02.886.645.600.080.050.100.150 1,2-Dihydroxybenzene-3,5-Disulfonic Acid Disodium Salt
-	D02.886.645.600.080.050.100.300 Ethamsylate
-	D02.886.645.600.080.050.100.325 Ferrozine
-	D02.886.645.600.080.050.100.600 Polyanetholesulfonate
-	D02.886.645.600.080.050.550 Lissamine Green Dyes
-	D02.886.645.600.080.050.650 Naphthalenesulfonates
-	D02.886.645.600.080.050.650.012 Amaranth Dye
-	D02.886.645.600.080.050.650.025 Amido Black
-	D02.886.645.600.080.050.650.050 Anilino Naphthalenesulfonates
-	D02.886.645.600.080.050.650.250 Congo Red
-	D02.886.645.600.080.050.650.300 Evans Blue



## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.886.645.600.080.050.650.750 Suramin
-	D02.886.645.600.080.050.650.875 Trypan Blue
-	D02.886.645.600.080.850 Sulfanilic Acids
-	D02.886.645.600.080.900 Trinitrobenzenesulfonic Acid
-	D02.886.645.600.925 Thiosulfonic Acids
-	D02.886.645.655 Sulfuric Acids
-	D02.886.645.655.424 Pentosan Sulfuric Polyester
-	D02.886.645.655.850 Sulfuric Acid Esters
-	D02.886.655 Thiazides
-	D02.886.655.500 Benzothiadiazines
-	D02.886.655.500.138 Bendroflumethiazide
-	D02.886.655.500.261 Chlorothiazide
-	D02.886.655.500.261.476 Hydrochlorothiazide
-	D02.886.655.500.261.476.716 Trichlormethiazide
-	D02.886.655.500.285 Cyclopenthiazide
-	D02.886.655.500.300 Diazoxide
-	D02.886.655.500.475 Hydroflumethiazide
-	D02.886.655.500.620 Methyclothiazide
-	D02.886.655.500.781 Polythiazide
-	D02.886.665 Thiazines
-	D02.886.665.074 Cephalosporins
-	D02.886.665.074.150 Cefamandole
-	D02.886.665.074.150.160 Cefoperazone
-	D02.886.665.074.160 Cefazolin
-	D02.886.665.074.177 Cefonicid
-	D02.886.665.074.185 Cefsulodin
-	D02.886.665.074.190 Cephacetrile
-	D02.886.665.074.190.190 Cefotaxime
-	D02.886.665.074.190.190.115 Cefixime
-	D02.886.665.074.190.190.125 Cefmenoxime
-	D02.886.665.074.190.190.135 Cefotiam
-	D02.886.665.074.190.190.145 Ceftizoxime
-	D02.886.665.074.190.190.155 Ceftriaxone
-	D02.886.665.074.190.190.165 Cefuroxime
-	D02.886.665.074.190.210 Cephalothin
-	D02.886.665.074.190.230 Cephapirin

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.886.665.074.200 Cephalexin
-	D02.886.665.074.200.155 Cefaclor
-	D02.886.665.074.200.165 Cefadroxil
-	D02.886.665.074.200.165.125 Cefatrizine
-	D02.886.665.074.200.180 Cephaloglycin
-	D02.886.665.074.200.185 Cephradine
-	D02.886.665.074.210 Cephaloridine
-	D02.886.665.074.210.150 Ceftazidime
-	D02.886.665.074.250 Cephamycins
-	D02.886.665.074.250.177 Cefmetazole
-	D02.886.665.074.250.199 Cefotetan
-	D02.886.665.074.250.222 Cefoxitin
-	D02.886.665.150 Chlormezanone
-	D02.886.665.400 Nifurtimox
-	D02.886.665.500 Piroxicam
-	D02.886.665.785 Thiadiazines
-	D02.886.665.985 Xylazine
-	D02.886.675 Thiazoles
-	D02.886.675.180 Chlormethiazole
-	D02.886.675.182 Cobicistat
-	D02.886.675.182.500 Elvitegravir, Cobicistat, Emtricitabine, Tenofovir Disoproxil Fumarate Drug Combination
-	D02.886.675.184 Dasatinib
-	D02.886.675.188 Dithiazanine
-	D02.886.675.197 Ethoxzolamide
-	D02.886.675.215 Famotidine
-	D02.886.675.250 FANFT
-	D02.886.675.274 Febuxostat
-	D02.886.675.298 Firefly Luciferin
-	D02.886.675.346 Levamisole
-	D02.886.675.414 Lurasidone Hydrochloride
-	D02.886.675.482 Niridazole
-	D02.886.675.500 Nizatidine
-	D02.886.675.540 Oxythiamine
-	D02.886.675.649 Rhodanine
-	D02.886.675.651 Riluzole

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.886.675.653 Ritonavir
-	D02.886.675.687 Saccharin
-	D02.886.675.725 Sulfathiazoles
-	D02.886.675.782 Tetramisole
-	D02.886.675.840 Thiabendazole
-	D02.886.675.867 Thiadiazoles
-	D02.886.675.867.060 Acetazolamide
-	D02.886.675.867.120 Benzolamide
-	D02.886.675.867.537 Methazolamide
-	D02.886.675.867.768 Timolol
-	D02.886.675.867.768.500 Combination Brimonidine Tartrate, Timolol Maleate Drug
-	D02.886.675.900 Thiamine
-	D02.886.675.900.300 Fursultiamin
-	D02.886.675.900.600 Thiamine Monophosphate
-	D02.886.675.900.702 Thiamine Pyrophosphate
-	D02.886.675.900.900 Thiamine Triphosphate
-	D02.886.675.933 Thiazolidinediones
-	D02.886.675.966 Thiazolidines
-	D02.886.680 Thiepins
-	D02.886.680.350 Dibenzothiepins
-	D02.886.680.350.250 Dothiepin
-	D02.886.680.350.520 Methiothepin
-	D02.886.680.702 Thiazepines
-	D02.886.680.702.500 Dibenzothiazepines
-	D02.886.680.702.500.500 Quetiapine Fumarate
-	D02.886.685 Thioamides
-	D02.886.685.800 Thioacetamide
-	D02.886.706 Thiocarbamates
-	D02.886.706.175 Dimethyldithiocarbamate
-	D02.886.706.175.900 Thiram
-	D02.886.706.175.950 Ziram
-	D02.886.706.200 Ditiocarb
-	D02.886.706.200.200 Disulfiram
-	D02.886.706.250 Ethylenebis(dithiocarbamates)
-	D02.886.706.250.500 Maneb

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.886.706.250.950 Zineb
-	D02.886.706.537 Methomyl
-	D02.886.706.825 Tolnaftate
-	D02.886.706.850 Triallate
-	D02.886.728 Thiocyanates
-	D02.886.740 Thioglycosides
-	D02.886.740.149 Dithioerythritol
-	D02.886.740.224 Dithiothreitol
-	D02.886.740.299 Sucralfate
-	D02.886.740.600 Thiogalactosides
-	D02.886.740.600.500 Isopropyl Thiogalactoside
-	D02.886.740.703 Thioglucosides
-	D02.886.740.703.350 Glucosinolates
-	D02.886.753 Thiones
-	D02.886.753.077 Anethole Trithione
-	D02.886.759 Thionucleosides
-	D02.886.759.111 Azathioprine
-	D02.886.759.800 Thioinosine
-	D02.886.759.800.500 Methylthioinosine
-	D02.886.759.829 Thiouridine
-	D02.886.765 Thionucleotides
-	D02.886.765.380 Guanosine 5'-O-(3-Thiotriphosphate)
-	D02.886.778 Thiophenes
-	D02.886.778.075 Canagliflozin
-	D02.886.778.150 Carticaine
-	D02.886.778.260 Duloxetine Hydrochloride
-	D02.886.778.315 Prasugrel Hydrochloride
-	D02.886.778.370 Ketotifen
-	D02.886.778.440 Morantel
-	D02.886.778.580 Pizotyline
-	D02.886.778.634 Pyrantel
-	D02.886.778.634.600 Pyrantel Pamoate
-	D02.886.778.634.670 Pyrantel Tartrate
-	D02.886.778.727 Rivaroxaban
-	D02.886.778.820 Thenoyltrifluoroacetone
-	D02.886.778.823 Thienopyridines

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.886.778.823.500 Ticlopidine
-	D02.886.778.827 Thioctic Acid
-	D02.886.778.840 Ticrynafen
-	D02.886.778.850 Tiletamine
-	D02.886.803 Thiosemicarbazones
-	D02.886.803.450 Methisazone
-	D02.886.803.702 Thioacetazone
-	D02.886.904 Thiourea
-	D02.886.904.100 beta-Aminoethyl Isothiourea
-	D02.886.904.200 Burimamide
-	D02.886.904.282 Dimaprit
-	D02.886.904.365 Guanylthiourea
-	D02.886.904.390 Isothiuronium
-	D02.886.904.425 Methallibure
-	D02.886.904.455 Metiamide
-	D02.886.904.500 Noxythiolin
-	D02.886.904.601 Phenylthiourea
-	D02.886.904.601.600 Phenylthiazolylthiourea
-	D02.886.904.601.900 Thiophanate
-	D02.886.952 Thioxanthenes
-	D02.886.952.250 Chlorprothixene
-	D02.886.952.305 Clopenthixol
-	D02.886.952.360 Flupenthixol
-	D02.886.952.450 Hycanthon
-	D02.886.952.500 Lucanthon
-	D02.886.952.787 Thiothixene
-	D02.925 Triazenes
-	D02.925.200 Dacarbazine
-	D02.925.240 Diminazene
-	D02.948 Urea
-	D02.948.058 Allantoin
-	D02.948.117 Biureas
-	D02.948.130 Biuret
-	D02.948.160 Bromisovalum
-	D02.948.395 Hydroxyurea
-	D02.948.511 Methylurea Compounds

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.948.594 Nitrosourea Compounds
-	D02.948.594.247 Carmustine
-	D02.948.594.310 Ethylnitrosourea
-	D02.948.594.440 Lomustine
-	D02.948.594.440.700 Semustine
-	D02.948.594.490 Methylnitrosourea
-	D02.948.594.550 Nimustine
-	D02.948.594.768 Streptozocin
-	D02.948.681 Phenylurea Compounds
-	D02.948.681.202 Carbanilides
-	D02.948.681.202.400 Imidocarb
-	D02.948.681.202.550 Nicarbazin
-	D02.948.681.241 Celiprolol
-	D02.948.681.280 Diflubenzuron
-	D02.948.681.397 Diuron
-	D02.948.681.515 Linuron
-	D02.948.828 Sulfonylurea Compounds
-	D02.948.828.071 Acetohexamide
-	D02.948.828.204 Carbutamide
-	D02.948.828.283 Chlorpropamide
-	D02.948.828.475 Gliclazide
-	D02.948.828.575 Glyburide
-	D02.948.828.834 Tolazamide
-	D02.948.828.896 Tolbutamide
-	D02.948.898 Thiourea
-	D02.948.898.100 beta-Aminoethyl Isothiourea
-	D02.948.898.200 Burimamide
-	D02.948.898.282 Dimaprit
-	D02.948.898.323 Ethylenethiourea
-	D02.948.898.365 Guanylthiourea
-	D02.948.898.390 Isothiuronium
-	D02.948.898.425 Methallibure
-	D02.948.898.455 Metiamide
-	D02.948.898.500 Noxythiolin
-	D02.948.898.601 Phenylthiourea
-	D02.948.898.601.600 Phenylthiazolylthiourea

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D02.948.898.601.900 Thiophanate
-	D02.974 Volatile Organic Compounds
-	D03 Heterocyclic Compounds
-	D03.066 Acids, Heterocyclic
-	D03.066.160 Cinoxacin
-	D03.066.288 Indoleacetic Acids
-	D03.066.288.200 Etodolac
-	D03.066.288.478 Hydroxyindoleacetic Acid
-	D03.066.349 Isonicotinic Acids
-	D03.066.349.220 Ethionamide
-	D03.066.349.390 Iproniazid
-	D03.066.349.410 Isoniazid
-	D03.066.349.490 Nialamide
-	D03.066.349.570 Prothionamide
-	D03.066.349.600 Pyridoxic Acid
-	D03.066.399 Isonipectic Acids
-	D03.066.399.300 Diphenoxylate
-	D03.066.399.450 Meperidine
-	D03.066.399.450.700 Promedol
-	D03.066.399.600 Phenoperidine
-	D03.066.399.650 Pirinitramide
-	D03.066.515 Nicotinic Acids
-	D03.066.515.292 Arecoline
-	D03.066.515.375 Clonixin
-	D03.066.515.400 Etazolate
-	D03.066.515.475 Niacin
-	D03.066.515.530 Niacinamide
-	D03.066.515.530.248 6-Aminonicotinamide
-	D03.066.515.530.498 Nicorandil
-	D03.066.515.530.500 Nikethamide
-	D03.066.515.540 Niceritrol
-	D03.066.515.550 Niflumic Acid
-	D03.066.515.635 Pipemidic Acid
-	D03.066.515.650 Piromidic Acid
-	D03.066.515.950 Xanthinol Niacinate
-	D03.066.566 Nipecotic Acids

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D03.066.627 Orotic Acid
-	D03.066.707 Picolinic Acids
-	D03.066.707.200 Fusaric Acid
-	D03.066.707.523 Picloram
-	D03.066.707.768 Streptonigrin
-	D03.066.758 Pipecolic Acids
-	D03.066.942 Xanthurenates
-	D03.066.942.505 Kynurenic Acid
-	D03.132 Alkaloids
-	D03.132.014 1-Deoxynojirimycin
-	D03.132.030 Aconitine
-	D03.132.032 Acridones
-	D03.132.032.077 Acronine
-	D03.132.052 Amaryllidaceae Alkaloids
-	D03.132.052.500 Galantamine
-	D03.132.066 Anabasine
-	D03.132.080 Arecoline
-	D03.132.089 Benzophenanthridines
-	D03.132.098 Benzyloquinolines
-	D03.132.098.038 Aporphines
-	D03.132.098.038.290 Apomorphine
-	D03.132.098.057 Berberine Alkaloids
-	D03.132.098.057.100 Berberine
-	D03.132.098.077 Bicuculline
-	D03.132.098.666 Papaverine
-	D03.132.098.666.850 Tetrahydropapaveroline
-	D03.132.098.833 Toxiferine
-	D03.132.098.833.050 Alcuronium
-	D03.132.098.916 Tubocurarine
-	D03.132.124 Betalains
-	D03.132.124.500 Betacyanins
-	D03.132.124.750 Betaxanthins
-	D03.132.151 Camptothecin
-	D03.132.151.850 Topotecan
-	D03.132.206 Cinchona Alkaloids
-	D03.132.206.636 Quinidine



## MeSH Tree Changes for 2017

Type	Tree - heading
-	D03.132.206.719 Quinine
-	D03.132.225 Colchicine
-	D03.132.225.374 Demecolcine
-	D03.132.225.500 Lumicolchicines
-	D03.132.285 Dihydro-beta-Erythroidine
-	D03.132.316 Emetine
-	D03.132.327 Ergot Alkaloids
-	D03.132.327.287 Ergolines
-	D03.132.327.287.262 Ergonovine
-	D03.132.327.287.262.455 Methylergonovine
-	D03.132.327.287.417 Lisuride
-	D03.132.327.287.572 Lysergic Acid
-	D03.132.327.287.572.522 Lysergic Acid Diethylamide
-	D03.132.327.287.630 Metergoline
-	D03.132.327.287.689 Methysergide
-	D03.132.327.287.730 Nicergoline
-	D03.132.327.287.800 Pergolide
-	D03.132.327.412 Ergotamines
-	D03.132.327.412.100 Bromocriptine
-	D03.132.327.412.150 Dihydroergocornine
-	D03.132.327.412.175 Dihydroergocristine
-	D03.132.327.412.200 Dihydroergocryptine
-	D03.132.327.412.225 Dihydroergotamine
-	D03.132.327.412.250 Dihydroergotoxine
-	D03.132.327.412.250.500 Ergoloid Mesylates
-	D03.132.327.412.400 Ergotamine
-	D03.132.420 Harringtonines
-	D03.132.436 Indole Alkaloids
-	D03.132.436.444 Harmaline
-	D03.132.436.477 Harmine
-	D03.132.436.545 Physostigmine
-	D03.132.436.613 Psilocybin
-	D03.132.436.613 Psilocybine
-	D03.132.436.681 Secologanin Tryptamine Alkaloids
-	D03.132.436.681.077 Ajmaline
-	D03.132.436.681.077.324 Lorajmine

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D03.132.436.681.077.650                      Prajmaline
-	D03.132.436.681.333                              Ellipticines
-	D03.132.436.681.444                              Ibogaine
-	D03.132.436.681.722                              Strychnine
-	D03.132.436.681.827                              Vinca Alkaloids
-	D03.132.436.681.827.650                          Vinblastine
-	D03.132.436.681.827.750                          Vincamine
-	D03.132.436.681.827.817                          Vincristine
-	D03.132.436.681.827.830                          Vindesine
-	D03.132.436.681.933                              Yohimbine
-	D03.132.436.681.933.500                          Reserpine
-	D03.132.436.750                                    Staurosporine
-	D03.132.478                                        Lobeline
-	D03.132.524                                        Mescaline
-	D03.132.545                                        Muscarine
-	D03.132.577                                        Opiate Alkaloids
-	D03.132.577.249                                  Morphinans
-	D03.132.577.249.106                              Benzomorphans
-	D03.132.577.249.106.700                          Pentazocine
-	D03.132.577.249.106.715                          Phenazocine
-	D03.132.577.249.150                              Buprenorphine
-	D03.132.577.249.150.500                          Buprenorphine, Naloxone Drug Combination
-	D03.132.577.249.165                              Butorphanol
-	D03.132.577.249.200                              Dextromethorphan
-	D03.132.577.249.235                              Dextrophan
-	D03.132.577.249.250                              Diprenorphine
-	D03.132.577.249.270                              Etorphine
-	D03.132.577.249.413                              Levallorphan
-	D03.132.577.249.463                              Levorphanol
-	D03.132.577.249.562                              Morphine Derivatives
-	D03.132.577.249.562.149                          Codeine
-	D03.132.577.249.562.149.287                      Hydrocodone
-	D03.132.577.249.562.149.575                      Oxycodone
-	D03.132.577.249.562.350                          Dihydromorphine
-	D03.132.577.249.562.430                          Ethylmorphine
-	D03.132.577.249.562.445                          Heroin

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D03.132.577.249.562.450	Hydromorphone
-	D03.132.577.249.562.571	Morphine
-	D03.132.577.249.562.692	Oxymorphone
-	D03.132.577.249.562.856	Thebaine
-	D03.132.577.249.606	Nalbuphine
-	D03.132.577.249.656	Nalorphine
-	D03.132.577.249.706	Naloxone
-	D03.132.577.249.706.275	Buprenorphine, Naloxone Drug Combination
-	D03.132.577.249.706.550	Naltrexone
-	D03.132.577.500	Noscapine
-	D03.132.577.750	Papaverine
Old Tree	D03.132.577.750.850	Tetrahydropapaveroline
-	D03.132.672	Pilocarpine
-	D03.132.716	Pyrrolizidine Alkaloids
-	D03.132.716.500	Monocrotaline
-	D03.132.740	Ryanodine
-	D03.132.750	Salsoline Alkaloids
-	D03.132.760	Solanaceous Alkaloids
-	D03.132.760.180	Belladonna Alkaloids
-	D03.132.760.180.572	Atropine Derivatives
-	D03.132.760.180.572.199	Atropine
-	D03.132.760.180.572.199.500	Hyoscyamine
-	D03.132.760.180.572.400	Ipratropium
-	D03.132.760.180.572.400.500	Albuterol, Ipratropium Drug Combination
-	D03.132.760.180.848	Scopolamine Hydrobromide
-	D03.132.760.200	Capsaicin
-	D03.132.760.570	Nicotine
-	D03.132.760.769	Solanine
-	D03.132.760.864	Tomatine
-	D03.132.786	Sparteine
-	D03.132.830	Swainsonine
-	D03.132.889	Tropanes
-	D03.132.889.180	Belladonna Alkaloids
-	D03.132.889.180.648	Atropine Derivatives
-	D03.132.889.180.648.199	Atropine
-	D03.132.889.180.648.199.500	Hyoscyamine

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D03.132.889.180.648.400 Ipratropium
-	D03.132.889.180.648.400.500 Albuterol, Ipratropium Drug Combination
-	D03.132.889.244 Benztropine
-	D03.132.889.354 Cocaine
-	D03.132.889.354.250 Crack Cocaine
-	D03.132.889.601 Scopolamine Derivatives
-	D03.132.889.601.200 Butylscopolammonium Bromide
-	D03.132.889.601.550 N-Methylscopolamine
-	D03.132.889.601.775 Scopolamine Hydrobromide
-	D03.132.889.601.887 Tiotropium Bromide
-	D03.132.920 Veratrum Alkaloids
-	D03.132.920.256 Cevanes
-	D03.132.920.256.310 Germine Acetates
-	D03.132.920.256.543 Protoveratrine
-	D03.132.920.256.679 Veratridine
-	D03.132.920.256.815 Veratrine
-	D03.132.960 Xanthines
-	D03.132.960.175 Caffeine
-	D03.132.960.651 Theobromine
-	D03.132.960.751 Theophylline
-	D03.132.960.877 Uric Acid
-	D03.132.960.938 Xanthine
-	D03.383 Heterocyclic Compounds, 1-Ring
-	D03.383.066 Azepines
-	D03.383.066.150 Caprolactam
-	D03.383.066.200 Dilazep
-	D03.383.066.400 Meptazinol
-	D03.383.066.498 Oxazepines
-	D03.383.066.600 Pentylenetetrazole
-	D03.383.066.785 Thiazepines
Old Tree	<b>D03.383.066.985 Zolazepam</b>
-	D03.383.082 Azetines
-	D03.383.082.301 Azetidines
-	D03.383.082.301.100 Azetidinecarboxylic Acid
-	D03.383.082.301.550 Ezetimibe
-	D03.383.082.301.550.500 Ezetimibe, Simvastatin Drug Combination

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D03.383.097 Azirines
-	D03.383.097.217 Aziridines
-	D03.383.097.217.300 Carbazilquinone
-	D03.383.097.217.900 Triaziquone
-	D03.383.097.217.924 Triethylenemelamine
-	D03.383.097.217.935 Triethylenephosphoramidate
-	D03.383.097.217.935.960 Thiotepa
-	D03.383.097.500 Mitomycins
-	D03.383.097.500.350 Mitomycin
-	D03.383.097.500.700 Porfiromycin
-	D03.383.113 Azocines
-	D03.383.113.367 Cyclazocine
-	D03.383.113.367.350 Ethylketocyclazocine
-	D03.383.113.767 Oxazocines
-	D03.383.113.767.500 Nefopam
-	D03.383.129 Azoles
-	D03.383.129.308 Imidazoles
-	D03.383.129.308.030 Aminoimidazole Carboxamide
-	D03.383.129.308.040 Antazoline
-	D03.383.129.308.080 Biotin
-	D03.383.129.308.090 Bis(4-Methyl-1-Homopiperazinylthiocarbonyl)disulfide
-	D03.383.129.308.100 4-(3-Butoxy-4-methoxybenzyl)-2-imidazolidinone
-	D03.383.129.308.108 Carbimazole
-	D03.383.129.308.130 Cimetidine
-	D03.383.129.308.175 Clotrimazole
-	D03.383.129.308.207 Creatinine
-	D03.383.129.308.240 Dacarbazine
-	D03.383.129.308.245 Dexmedetomidine
-	D03.383.129.308.250 Econazole
-	D03.383.129.308.253 Enoximone
-	D03.383.129.308.260 Etimizol
-	D03.383.129.308.265 Etomidate
-	D03.383.129.308.270 Fadzozole
-	D03.383.129.308.290 Fluspirilene
-	D03.383.129.308.373 Histamine
-	D03.383.129.308.373.500 Methylhistamines

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D03.383.129.308.385 Histidinol
-	D03.383.129.308.428 Idazoxan
-	D03.383.129.308.432 Imidazolidines
-	D03.383.129.308.432.500 Ethylenethiourea
-	D03.383.129.308.432.555 Hydantoins
-	D03.383.129.308.432.555.105 Allantoin
-	D03.383.129.308.432.555.287 Dantrolene
-	D03.383.129.308.432.555.620 Mephenytoin
-	D03.383.129.308.432.555.730 Phenytoin
-	D03.383.129.308.432.555.868 Thiohydantoins
-	D03.383.129.308.432.555.868.650 Phenylthiohydantoin
-	D03.383.129.308.436 Imidazolines
-	D03.383.129.308.436.500 Clonidine
-	D03.383.129.308.436.750 Tolazoline
-	D03.383.129.308.445 Impromidine
-	D03.383.129.308.480 Levamisole
-	D03.383.129.308.507 Losartan
-	D03.383.129.308.521 Medetomidine
-	D03.383.129.308.535 Methimazole
-	D03.383.129.308.550 Miconazole
-	D03.383.129.308.585 Naphazoline
-	D03.383.129.308.626 Niridazole
-	D03.383.129.308.658 Nitroimidazoles
-	D03.383.129.308.658.200 Dimetridazole
-	D03.383.129.308.658.250 Etanidazole
-	D03.383.129.308.658.400 Ipronidazole
-	D03.383.129.308.658.500 Metronidazole
-	D03.383.129.308.658.530 Misonidazole
-	D03.383.129.308.658.580 Nimorazole
-	D03.383.129.308.658.700 Ornidazole
-	D03.383.129.308.658.730 Ronidazole
-	D03.383.129.308.658.900 Tinidazole
-	D03.383.129.308.674 Olmesartan Medoxomil
-	D03.383.129.308.674.500 Amlodipine Besylate, Olmesartan Medoxomil Drug Combination
-	D03.383.129.308.690 Ondansetron

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D03.383.129.308.704 Oxymetazoline
-	D03.383.129.308.754 Phentolamine
-	D03.383.129.308.884 Tetramisole
-	D03.383.129.308.935 Trimethaphan
-	D03.383.129.308.960 Urocanic Acid
-	D03.383.129.308.980 Vardenafil Dihydrochloride
-	D03.383.129.385 Isoxazoles
-	D03.383.129.385.025 Acid alpha-Amino-3-hydroxy-5-methyl-4-isoxazolepropionic
-	D03.383.129.385.162 Cycloserine
-	D03.383.129.385.231 Ibotenic Acid
-	D03.383.129.385.300 Isocarboxazid
-	D03.383.129.385.650 Paliperidone Palmitate
-	D03.383.129.462 Oxazoles
-	D03.383.129.462.050 Aminorex
-	D03.383.129.462.175 Dimethadione
-	D03.383.129.462.285 Fura-2
-	D03.383.129.462.470 Muscimol
-	D03.383.129.462.580 Oxadiazoles
-	D03.383.129.462.580.200 5-Amino-3-((5-nitro-2-furyl)vinyl)-1,2,4-oxadiazole
-	D03.383.129.462.580.400 4-Chloro-7-nitrobenzofurazan
-	D03.383.129.462.580.600 Quisqualic Acid
-	D03.383.129.462.580.693 Sydnones
-	D03.383.129.462.580.693.450 Molsidomine
-	D03.383.129.462.600 Oxazolidinones
-	D03.383.129.462.600.162 Cycloserine
-	D03.383.129.462.600.500 Furazolidone
-	D03.383.129.462.600.550 Linezolid
-	D03.383.129.462.600.600 Nifuratel
-	D03.383.129.462.620 Oxazolone
-	D03.383.129.462.673 Pemoline
-	D03.383.129.462.883 Trimethadione
-	D03.383.129.539 Pyrazoles
-	D03.383.129.539.100 Betazole
-	D03.383.129.539.120 amine 4,5-Dihydro-1-(3-(trifluoromethyl)phenyl)-1H-pyrazol-3-

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D03.383.129.539.160 Celecoxib
-	D03.383.129.539.200 Epirizole
-	D03.383.129.539.487 Indazoles
-	D03.383.129.539.487.130 Benzydamine
-	D03.383.129.539.487.350 Granisetron
-	D03.383.129.539.550 Muzolimine
-	D03.383.129.539.650 Oxypurinol
-	D03.383.129.539.850 Pyrazolones
-	D03.383.129.539.850.077 Aminopyrine
-	D03.383.129.539.850.077.025 Ampyrone
-	D03.383.129.539.850.077.150 Dipyrone
-	D03.383.129.539.850.088 Antipyrine
-	D03.383.129.539.850.777 Phenylbutazone
-	D03.383.129.539.850.777.670 Feprazone
-	D03.383.129.539.850.777.700 Oxyphenbutazone
-	D03.383.129.539.850.777.800 Sulfinpyrazone
-	D03.383.129.539.925 Sulfaphenazole
-	D03.383.129.578 Pyrroles
-	D03.383.129.578.075 Atorvastatin Calcium
-	D03.383.129.578.150 Cromakalim
-	D03.383.129.578.399 Maleimides
-	D03.383.129.578.399.418 Ethylmaleimide
-	D03.383.129.578.617 Porphobilinogen
-	D03.383.129.578.748 Prodigiosin
-	D03.383.129.578.770 Pyrrolnitrin
-	D03.383.129.578.805 Ryanodine
-	D03.383.129.578.840 Tetrapyrroles
-	D03.383.129.578.840.249 Bile Pigments
-	D03.383.129.578.840.249.184 Bilirubin
-	D03.383.129.578.840.249.184.200 Biliverdine
-	D03.383.129.578.840.249.727 Urobilin
-	D03.383.129.578.840.249.852 Urobilinogen
-	D03.383.129.578.840.374 Chlorophyll
-	D03.383.129.578.840.374.100 Bacteriochlorophylls
-	D03.383.129.578.840.374.180 Chlorophyllides
-	D03.383.129.578.840.374.700 Pheophytins



## MeSH Tree Changes for 2017

Type	Tree - heading
-	D03.383.129.578.840.374.725 Protochlorophyllide
-	D03.383.129.578.840.437 Corrinoids
-	D03.383.129.578.840.437.777 Vitamin B 12
-	D03.383.129.578.840.437.777.270 Cobamides
-	D03.383.129.578.840.437.777.560 Hydroxocobalamin
-	D03.383.129.578.840.468 Phycobilins
-	D03.383.129.578.840.500 Porphyrins
-	D03.383.129.578.840.500.250 Coproporphyrins
-	D03.383.129.578.840.500.280 Deuteroporphyrins
-	D03.383.129.578.840.500.340 Etioporphyrins
-	D03.383.129.578.840.500.462 Hematoporphyrins
-	D03.383.129.578.840.500.462.400 Hematoporphyrin Derivative
-	D03.383.129.578.840.500.462.400.200 Dihematoporphyrin Ether
-	D03.383.129.578.840.500.620 Mesoporphyrins
-	D03.383.129.578.840.500.640 Metalloporphyrins
-	D03.383.129.578.840.500.640.220 Chlorophyll
-	D03.383.129.578.840.500.640.220.100 Bacteriochlorophylls
-	D03.383.129.578.840.500.640.220.180 Chlorophyllides
-	D03.383.129.578.840.500.640.220.453 Pheophytins
-	D03.383.129.578.840.500.640.220.725 Protochlorophyllide
-	D03.383.129.578.840.500.640.587 Heme
-	D03.383.129.578.840.500.640.587.462 Hemin
-	D03.383.129.578.840.500.700 Porphyrinogens
-	D03.383.129.578.840.500.700.250 Coproporphyrinogens
-	D03.383.129.578.840.500.700.900 Uroporphyrinogens
-	D03.383.129.578.840.500.725 Protoporphyrins
-	D03.383.129.578.840.500.880 Uroporphyrins
-	D03.383.129.578.910 Tolmetin
-	D03.383.129.617 Tetrazoles
-	D03.383.129.617.235 Cefotetan
-	D03.383.129.617.467 Losartan
-	D03.383.129.617.584 Olmesartan Medoxomil
-	D03.383.129.617.584.500 Amlodipine Besylate, Olmesartan Medoxomil Drug Combination
-	D03.383.129.617.700 Tetrazolium Salts
-	D03.383.129.617.700.500 Nitroblue Tetrazolium

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D03.383.129.617.850 Valsartan
-	D03.383.129.617.850.500 Amlodipine, Valsartan Drug Combination
-	D03.383.129.708 Thiazoles
-	D03.383.129.708.089 Benzothiazoles
-	D03.383.129.708.089.222 Dithiazanine
-	D03.383.129.708.089.416 Ethoxzolamide
-	D03.383.129.708.089.611 Riluzole
-	D03.383.129.708.089.708 Saccharin
-	D03.383.129.708.180 Chlormethiazole
-	D03.383.129.708.189 Cobicistat
-	D03.383.129.708.189.500 Elvitegravir, Cobicistat, Emtricitabine, Tenofovir Disoproxil Fumarate Drug Combination
-	D03.383.129.708.198 Dasatinib
-	D03.383.129.708.215 Famotidine
-	D03.383.129.708.250 FANFT
-	D03.383.129.708.274 Febuxostat
-	D03.383.129.708.298 Firefly Luciferin
-	D03.383.129.708.346 Levamisole
-	D03.383.129.708.414 Lurasidone Hydrochloride
-	D03.383.129.708.482 Niridazole
-	D03.383.129.708.500 Nizatidine
-	D03.383.129.708.540 Oxythiamine
-	D03.383.129.708.649 Rhodanine
-	D03.383.129.708.653 Ritonavir
-	D03.383.129.708.725 Sulfathiazoles
-	D03.383.129.708.782 Tetramisole
-	D03.383.129.708.824 Thiabendazole
-	D03.383.129.708.867 Thiadiazoles
-	D03.383.129.708.867.060 Acetazolamide
-	D03.383.129.708.867.120 Benzolamide
-	D03.383.129.708.867.537 Methazolamide
-	D03.383.129.708.867.768 Timolol
-	D03.383.129.708.867.768.500 Brimonidine Tartrate, Timolol Maleate Drug Combination
-	D03.383.129.708.900 Thiamine
-	D03.383.129.708.900.300 Fursultiamin

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D03.383.129.708.900.600 Thiamine Monophosphate
-	D03.383.129.708.900.702 Thiamine Pyrophosphate
-	D03.383.129.708.900.900 Thiamine Triphosphate
-	D03.383.129.708.933 Thiazolidinediones
-	D03.383.129.799 Triazoles
-	D03.383.129.799.100 Amitrole
-	D03.383.129.799.450 Fluconazole
-	D03.383.129.799.500 Guanazole
-	D03.383.129.799.550 Itraconazole
-	D03.383.129.799.725 Sitagliptin Phosphate
-	D03.383.129.799.725.500 Sitagliptin Phosphate, Metformin Hydrochloride Drug Combination
-	D03.383.129.799.900 Trapidil
-	D03.383.129.799.950 Voriconazole
New Tree	<a href="#">D03.383.188</a> <a href="#">Dioxanes</a>
New Tree	<a href="#">D03.383.188.425</a> <a href="#">Idazoxan</a>
New Tree	<a href="#">D03.383.188.568</a> <a href="#">Piperoxan</a>
New Tree	<a href="#">D03.383.188.712</a> <a href="#">Spectinomycin</a>
-	D03.383.231 Dioxins
Old Tree	<del><a href="#">D03.383.231.388</a></del> <del><a href="#">Dioxanes</a></del>
Old Tree	<del><a href="#">D03.383.231.388.425</a></del> <del><a href="#">Idazoxan</a></del>
Old Tree	<del><a href="#">D03.383.231.388.568</a></del> <del><a href="#">Piperoxan</a></del>
Old Tree	<del><a href="#">D03.383.231.388.712</a></del> <del><a href="#">Spectinomycin</a></del>
Old Tree	<del><a href="#">D03.383.231.900</a></del> <del><a href="#">Tetrachlorodibenzodioxin</a></del>
-	D03.383.246 Dioxoles
-	D03.383.246.118 Benzodioxoles
-	D03.383.246.118.600 Piperonyl Butoxide
-	D03.383.246.118.750 Safrole
-	D03.383.246.238 Dioxolanes
-	D03.383.312 Furans
-	D03.383.312.150 4-Butyrolactone
-	D03.383.312.200 Acetogenins
-	D03.383.312.250 Citraconic Anhydrides

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D03.383.312.303 Darunavir
-	D03.383.312.355 Furaldehyde
-	D03.383.312.490 Lasalocid
-	D03.383.312.520 Maleic Anhydrides
-	D03.383.312.600 Monensin
-	D03.383.312.620 Nafronyl
-	D03.383.312.634 Nigericin
-	D03.383.312.649 Nitrofurans
-	D03.383.312.649.200 5-Amino-3-((5-nitro-2-furyl)vinyl)-1,2,4-oxadiazole
-	D03.383.312.649.290 FANFT
-	D03.383.312.649.308 Furagin
-	D03.383.312.649.313 Furazolidone
-	D03.383.312.649.330 Furylfuramide
-	D03.383.312.649.410 Nifuratel
-	D03.383.312.649.480 Nifurtimox
-	D03.383.312.649.566 Nitrofurantoin
-	D03.383.312.649.691 Nitrofurazone
-	D03.383.312.649.750 Nitrovin
-	D03.383.312.750 Ranitidine
-	D03.383.411 Lactams
-	D03.383.411.200 Caprolactam
-	D03.383.411.350 Monobactams
-	D03.383.411.350.044 Aztreonam
-	D03.383.517 Oxathiins
-	D03.383.517.200 Carboxin
-	D03.383.533 Oxazines
-	D03.383.533.249 Benzoxazines
-	D03.383.533.375 Efavirenz, Emtricitabine, Tenofovir Disoproxil Fumarate Drug Combination
-	D03.383.533.500 Ifosfamide
-	D03.383.533.640 Morpholines
-	D03.383.533.640.250 Dextromoramide
-	D03.383.533.640.362 Molsidomine
-	D03.383.533.640.475 Morcizine
-	D03.383.533.640.562 Morpholinos
-	D03.383.533.640.650 Phenmetrazine

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D03.383.533.640.713 Rivaroxaban
-	D03.383.533.640.775 Timolol
-	D03.383.533.640.900 Viloxazine
-	D03.383.533.640.950 Xamoterol
-	D03.383.564 Oxepins
-	D03.383.585 Oxocins
-	D03.383.606 Piperazines
-	D03.383.606.130 Almitrine
-	D03.383.606.170 Aripiprazole
-	D03.383.606.210 Buspirone
-	D03.383.606.290 Cinnarizine
-	D03.383.606.320 Cyclizine
-	D03.383.606.350 Delavirdine
-	D03.383.606.380 Diethylcarbamazine
-	D03.383.606.385 Diketopiperazines
-	D03.383.606.385.500 Razoxane
-	D03.383.606.385.500.500 Dexrazoxane
-	D03.383.606.390 Dimethylphenylpiperazinium Iodide
-	D03.383.606.398 Eszopiclone
-	D03.383.606.405 Imatinib Mesylate
-	D03.383.606.420 Prasugrel Hydrochloride
-	D03.383.606.450 Flunarizine
-	D03.383.606.460 Flupenthixol
-	D03.383.606.500 HEPES
-	D03.383.606.515 Hydroxyzine
-	D03.383.606.515.200 Cetirizine
-	D03.383.606.527 1-(5-Isoquinolinesulfonyl)-2-Methylpiperazine
-	D03.383.606.530 Itraconazole
-	D03.383.606.560 Ketoconazole
-	D03.383.606.596 Lidoflazine
-	D03.383.606.650 Meclizine
-	D03.383.606.760 Pipecuronium
-	D03.383.606.765 Pipemidic Acid
-	D03.383.606.773 Pipobroman
-	D03.383.606.790 Piribedil
-	D03.383.606.810 Prospidium

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D03.383.606.827 Quipazine
-	D03.383.606.841 Ranolazine
-	D03.383.606.854 Sildenafil Citrate
-	D03.383.606.880 Thiethylperazine
-	D03.383.606.900 Trazodone
-	D03.383.606.920 Trimetazidine
-	D03.383.606.960 Vardenafil Dihydrochloride
-	D03.383.606.980 Vilazodone Hydrochloride
-	D03.383.621 Piperidines
-	D03.383.621.040 Alphaprodine
-	D03.383.621.050 Anabasine
-	D03.383.621.080 Betalains
-	D03.383.621.080.500 Betacyanins
-	D03.383.621.080.750 Betaxanthins
-	D03.383.621.110 Biperiden
-	D03.383.621.135 Cisapride
-	D03.383.621.147 Clopamide
-	D03.383.621.160 Cyproheptadine
-	D03.383.621.160.500 Loratadine
-	D03.383.621.210 Domperidone
-	D03.383.621.265 Fentanyl
-	D03.383.621.265.100 Alfentanil
-	D03.383.621.265.900 Sufentanil
-	D03.383.621.270 Flecainide
-	D03.383.621.280 Fluspirilene
-	D03.383.621.300 Imino Pyranoses
-	D03.383.621.300.033 1-Deoxynojirimycin
-	D03.383.621.320 Indoramin
-	D03.383.621.349 Isonipectic Acids
-	D03.383.621.349.300 Diphenoxylate
-	D03.383.621.349.450 Meperidine
-	D03.383.621.349.450.700 Promedol
-	D03.383.621.349.683 Phenoperidine
-	D03.383.621.349.700 Pirinitramide
-	D03.383.621.365 Ketanserin
-	D03.383.621.370 Ketotifen

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D03.383.621.418 Lobeline
-	D03.383.621.440 Loperamide
-	D03.383.621.450 Mepivacaine
-	D03.383.621.460 Methylphenidate
-	D03.383.621.460.500 Dexmethylphenidate Hydrochloride
-	D03.383.621.493 Minoxidil
-	D03.383.621.566 Nipecotic Acids
-	D03.383.621.600 Paroxetine
-	D03.383.621.610 Pempidine
-	D03.383.621.620 Penfluridol
-	D03.383.621.644 Perhexiline
-	D03.383.621.699 Phencyclidine
-	D03.383.621.758 Pipecolic Acids
-	D03.383.621.758.323 Bupivacaine
-	D03.383.621.808 Piperidones
-	D03.383.621.808.240 Cycloheximide
-	D03.383.621.808.250 Dexetimide
-	D03.383.621.808.425 Glutethimide
-	D03.383.621.808.425.100 Aminoglutethimide
-	D03.383.621.808.612 Naphthalimides
-	D03.383.621.808.800 Thalidomide
-	D03.383.621.808.930 Triacetoneamine-N-Oxyl
-	D03.383.621.815 Piperoxan
-	D03.383.621.840 Ritanserine
-	D03.383.621.855 Terfenadine
-	D03.383.621.900 Trihexyphenidyl
-	D03.383.663 Pyrans
-	D03.383.663.075 Aurovertins
-	D03.383.663.283 Benzopyrans
-	D03.383.663.283.119 Aflatoxins
-	D03.383.663.283.119.075 Aflatoxin B1
-	D03.383.663.283.119.100 Aflatoxin M1
-	D03.383.663.283.240 Chromans
-	D03.383.663.283.240.190 Catechin
-	D03.383.663.283.240.225 Centchroman
-	D03.383.663.283.266 Chromones

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D03.383.663.283.266.300 Cromolyn Sodium
-	D03.383.663.283.266.450 Flavonoids
-	D03.383.663.283.266.450.087 Anthocyanins
-	D03.383.663.283.266.450.175 Benzoflavones
-	D03.383.663.283.266.450.175.100 beta-Naphthoflavone
-	D03.383.663.283.266.450.190 Biflavonoids
-	D03.383.663.283.266.450.206 Catechin
-	D03.383.663.283.266.450.221 Chalcones
-	D03.383.663.283.266.450.221.500 Chalcone
-	D03.383.663.283.266.450.252 Flavanones
-	D03.383.663.283.266.450.252.500 Hesperidin
-	D03.383.663.283.266.450.260 Flavones
-	D03.383.663.283.266.450.260.110 Apigenin
-	D03.383.663.283.266.450.260.222 Diosmin
-	D03.383.663.283.266.450.260.444 Flavoxate
-	D03.383.663.283.266.450.260.555 Luteolin
-	D03.383.663.283.266.450.268 Flavonolignans
-	D03.383.663.283.266.450.268.777 Silymarin
-	D03.383.663.283.266.450.284 Flavonols
-	D03.383.663.283.266.450.284.388 Kaempferols
-	D03.383.663.283.266.450.284.777 Quercetin
-	D03.383.663.283.266.450.284.888 Rutin
-	D03.383.663.283.266.450.284.888.500 Hydroxyethylrutoside
-	D03.383.663.283.266.450.400 Isoflavones
-	D03.383.663.283.266.450.400.187 Coumestrol
-	D03.383.663.283.266.450.400.375 Genistein
-	D03.383.663.283.266.450.400.687 Pterocarpans
-	D03.383.663.283.266.450.400.843 Rotenone
-	D03.383.663.283.266.450.700 Proanthocyanidins
-	D03.383.663.283.320 Citrinin
-	D03.383.663.283.446 Coumarins
-	D03.383.663.283.446.139 Aminocoumarins
-	D03.383.663.283.446.139.500 Novobiocin
-	D03.383.663.283.446.280 Chromonar
-	D03.383.663.283.446.350 Coumestrol
-	D03.383.663.283.446.400 Esculin



## MeSH Tree Changes for 2017

Type	Tree - heading
-	D03.383.663.283.446.520 4-Hydroxycoumarins
-	D03.383.663.283.446.520.079 Acenocoumarol
-	D03.383.663.283.446.520.203 Dicumarol
-	D03.383.663.283.446.520.451 Ethyl Biscoumacetate
-	D03.383.663.283.446.520.750 Phenprocoumon
-	D03.383.663.283.446.520.914 Warfarin
-	D03.383.663.283.446.598 Isocoumarins
-	D03.383.663.283.446.598.500 Ochratoxins
-	D03.383.663.283.446.794 Furocoumarins
-	D03.383.663.283.446.794.200 Ficusin
-	D03.383.663.283.446.794.300 Khellin
-	D03.383.663.283.446.794.500 Methoxsalen
-	D03.383.663.283.446.794.875 Trioxsalen
-	D03.383.663.283.446.853 Pyranocoumarins
-	D03.383.663.283.446.912 Umbelliferones
-	D03.383.663.283.446.912.326 Coumaphos
-	D03.383.663.283.446.912.531 Hymecromone
-	D03.383.663.283.446.912.850 Scopoletin
-	D03.383.663.283.455 Cromakalim
-	D03.383.663.283.500 Ellagic Acid
-	D03.383.663.283.600 Hematoxylin
-	D03.383.663.283.755 Nebivolol
-	D03.383.663.283.909 Vitamin E
-	D03.383.663.283.909.750 Tocopherols
-	D03.383.663.283.909.750.249 alpha-Tocopherol
-	D03.383.663.283.909.750.374 beta-Tocopherol
-	D03.383.663.283.909.750.500 gamma-Tocopherol
-	D03.383.663.283.909.875 Tocotrienols
-	D03.383.663.387 Glaucarubin
-	D03.383.663.491 Iridoids
-	D03.383.663.491.500 Iridoid Glycosides
-	D03.383.663.491.500.500 Iridoid Glucosides
-	D03.383.663.555 Mupirocin
-	D03.383.663.620 Nigericin
-	D03.383.663.663 Patulin
-	D03.383.663.705 Pyran Copolymer

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D03.383.663.712 Pyranocoumarins
-	D03.383.663.718 Pyrones
-	D03.383.663.859 Zanamivir
-	D03.383.679 Pyrazines
-	D03.383.679.149 Amiloride
-	D03.383.679.450 Bortezomib
-	D03.383.679.600 Eszopiclone
-	D03.383.679.750 Pyrazinamide
-	D03.383.679.875 Sitagliptin Phosphate
-	D03.383.679.875.500 Combination Sitagliptin Phosphate, Metformin Hydrochloride Drug
-	D03.383.710 Pyridazines
-	D03.383.710.100 Cilazapril
-	D03.383.710.350 Luminol
-	D03.383.710.400 Maleic Hydrazide
-	D03.383.710.605 Phthalazines
-	D03.383.710.605.500 Hydralazine
-	D03.383.710.605.500.250 Dihydralazine
-	D03.383.710.605.925 Todralazine
-	D03.383.725 Pyridines
-	D03.383.725.024 2-Pyridinylmethylsulfanylbenzimidazoles
-	D03.383.725.024.249 Lansoprazole
-	D03.383.725.024.249.500 Dexlansoprazole
-	D03.383.725.024.500 Omeprazole
-	D03.383.725.024.500.500 Esomeprazole
-	D03.383.725.024.750 Rabeprazole
-	D03.383.725.050 Aminopyridines
-	D03.383.725.050.060 4-Aminopyridine
-	D03.383.725.050.085 Amrinone
-	D03.383.725.050.085.543 Milrinone
-	D03.383.725.050.440 Methapyrilene
-	D03.383.725.050.610 Phenazopyridine
-	D03.383.725.050.805 Ppyrilamine
-	D03.383.725.050.920 Tripelennamine
-	D03.383.725.068 Atazanavir Sulfate
-	D03.383.725.086 Betahistine

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D03.383.725.150 Carbolines
-	D03.383.725.150.500 Tadalafil
-	D03.383.725.180 Clopidol
-	D03.383.725.192 Dabigatran
-	D03.383.725.203 Dihydropyridines
-	D03.383.725.203.065 Amlodipine
-	D03.383.725.203.065.500 Amlodipine Besylate, Olmesartan Medoxomil Drug Combination
-	D03.383.725.203.065.750 Amlodipine, Valsartan Drug Combination
-	D03.383.725.203.200 Dicarbethoxydihydrocollidine
-	D03.383.725.203.250 Felodipine
-	D03.383.725.203.310 Isradipine
-	D03.383.725.203.510 Nicardipine
-	D03.383.725.203.540 Nifedipine
-	D03.383.725.203.570 Nimodipine
-	D03.383.725.203.580 Nisoldipine
-	D03.383.725.203.590 Nitrendipine
-	D03.383.725.203.600 3-Pyridinecarboxylic acid, 1,4-dihydro-2,6-dimethyl-5- nitro-4-(2-(trifluoromethyl)phenyl)-, Methyl ester
-	D03.383.725.210 Dimethindene
-	D03.383.725.220 2,2'-Dipyridyl
-	D03.383.725.227 Disopyramide
-	D03.383.725.259 Doxylamine
-	D03.383.725.322 Eszopiclone
-	D03.383.725.385 Indinavir
-	D03.383.725.394 Isonicotinic Acids
-	D03.383.725.394.220 Ethionamide
-	D03.383.725.394.414 Iproniazid
-	D03.383.725.394.582 Isoniazid
-	D03.383.725.394.628 Nialamide
-	D03.383.725.394.675 Prothionamide
-	D03.383.725.394.700 Pyridoxic Acid
-	D03.383.725.450 1-Methyl-4-phenyl-1,2,3,6-tetrahydropyridine
-	D03.383.725.463 Metyrapone
-	D03.383.725.495 Naphthylvinylpyridine
-	D03.383.725.506 Nevirapine

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D03.383.725.518 Nicotine
-	D03.383.725.547 Nicotinic Acids
-	D03.383.725.547.292 Arecoline
-	D03.383.725.547.375 Clonixin
-	D03.383.725.547.400 Etazolate
-	D03.383.725.547.475 Niacin
-	D03.383.725.547.530 Niacinamide
-	D03.383.725.547.530.248 6-Aminonicotinamide
-	D03.383.725.547.530.498 Nicorandil
-	D03.383.725.547.530.500 Nikethamide
-	D03.383.725.547.540 Niceritrol
-	D03.383.725.547.550 Niflumic Acid
-	D03.383.725.547.570 Nimodipine
-	D03.383.725.547.635 Pipemidic Acid
-	D03.383.725.547.650 Piromidic Acid
-	D03.383.725.547.900 3-Pyridinecarboxylic acid, 1,4-dihydro-2,6-dimethyl-5-nitro-4-(2-(trifluoromethyl)phenyl)-, Methyl ester
-	D03.383.725.547.950 Xanthinol Niacinate
-	D03.383.725.565 Nicotiny Alcohol
-	D03.383.725.620 Pheniramine
-	D03.383.725.620.129 Brompheniramine
-	D03.383.725.620.129.960 Zimeldine
-	D03.383.725.620.150 Chlorpheniramine
-	D03.383.725.676 Picolines
-	D03.383.725.676.152 Amprolium
-	D03.383.725.676.925 Vitamin B 6
-	D03.383.725.676.925.500 Pyridoxal
-	D03.383.725.676.925.500.500 Pyridoxal Phosphate
-	D03.383.725.676.925.750 Pyridoxamine
-	D03.383.725.676.925.875 Pyridoxine
-	D03.383.725.705 Picolinic Acids
-	D03.383.725.705.200 Fusaric Acid
-	D03.383.725.705.523 Picloram
-	D03.383.725.705.645 Pyridoxic Acid
-	D03.383.725.705.768 Streptonigrin
-	D03.383.725.740 Polyvinylpyridine N-Oxide

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D03.383.725.762                      Pyridinium Compounds
-	D03.383.725.762.232                      Cetylpyridinium
-	D03.383.725.762.300                      Desmosine
-	D03.383.725.762.352                      Diquat
-	D03.383.725.762.500                      Isodesmosine
-	D03.383.725.762.550                      1-Methyl-4-phenylpyridinium
-	D03.383.725.762.600                      Obidoxime Chloride
-	D03.383.725.762.621                      Paraquat
-	D03.383.725.762.740                      Pyridostigmine Bromide
-	D03.383.725.762.760                      Pyrithiamine
-	D03.383.725.762.900                      Trimedoxime
-	D03.383.725.762.925                      Viologens
-	D03.383.725.762.925.100                      Benzyl Viologen
-	D03.383.725.791                      Pyridones
-	D03.383.725.791.100                      Bemegride
-	D03.383.725.791.496                      Iodopyridones
-	D03.383.725.791.496.371                      Iodopyracet
-	D03.383.725.791.496.750                      Propylidone
-	D03.383.725.791.550                      Mimosine
-	D03.383.725.791.900                      Trazodone
-	D03.383.725.812                      Pyrithioxin
-	D03.383.725.822                      Quinolinic Acids
-	D03.383.725.822.700                      Quinolinic Acid
-	D03.383.725.849                      Thienopyridines
-	D03.383.725.849.500                      Ticlopidine
-	D03.383.725.877                      Tolperisone
-	D03.383.725.932                      Triprolidine
-	D03.383.725.942                      Tropicamide
-	D03.383.742                      Pyrimidines
-	D03.383.742.120                      Buspirone
-	D03.383.742.148                      Dasatinib
-	D03.383.742.175                      Dipyridamole
-	D03.383.742.175.500                      Aspirin, Dipyridamole Drug Combination
-	D03.383.742.200                      Epirizole
-	D03.383.742.279                      Hexetidine
-	D03.383.742.349                      Imatinib Mesylate

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D03.383.742.418 Minoxidil
-	D03.383.742.430 Mopidamol
-	D03.383.742.440 Morantel
-	D03.383.742.470 Nicarbazin
-	D03.383.742.550 Oxypurinol
-	D03.383.742.592 Paliperidone Palmitate
-	D03.383.742.634 Pyrantel
-	D03.383.742.634.600 Pyrantel Pamoate
-	D03.383.742.634.670 Pyrantel Tartrate
-	D03.383.742.675 Pyrimethamine
-	D03.383.742.680 Pyrimidine Nucleosides
-	D03.383.742.680.245 Cytidine
-	D03.383.742.680.245.217 Azacitidine
-	D03.383.742.680.245.453 Cytarabine
-	D03.383.742.680.245.453.050 Ancitabine
-	D03.383.742.680.245.500 Deoxycytidine
-	D03.383.742.680.245.500.250 Bromodeoxycytidine
-	D03.383.742.680.245.500.425 Capecitabine
-	D03.383.742.680.245.500.600 Emtricitabine
-	D03.383.742.680.245.500.600.125 Efavirenz, Emtricitabine, Tenofovir Disoproxil Fumarate Drug Combination
-	D03.383.742.680.245.500.600.250 Elvitegravir, Cobicistat, Emtricitabine, Tenofovir Disoproxil Fumarate Drug Combination
-	D03.383.742.680.245.500.600.500 Emtricitabine, Rilpivirine, Tenofovir Drug Combination
-	D03.383.742.680.245.500.600.750 Emtricitabine, Tenofovir Disoproxil Fumarate Drug Combination
-	D03.383.742.680.245.500.950 Zalcitabine
-	D03.383.742.680.245.500.950.500 Lamivudine
-	D03.383.742.680.350 Formycins
-	D03.383.742.680.350.200 Coformycin
-	D03.383.742.680.350.200.700 Pentostatin
-	D03.383.742.680.705 Thymidine
-	D03.383.742.680.705.875 Stavudine
-	D03.383.742.680.705.900 Trifluridine
-	D03.383.742.680.705.950 Zidovudine
-	D03.383.742.680.725 Tunicamycin

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D03.383.742.680.852 Uridine
-	D03.383.742.680.852.150 Arabinofuranosyluracil
-	D03.383.742.680.852.176 Azauridine
-	D03.383.742.680.852.250 3-Deazauridine
-	D03.383.742.680.852.300 Deoxyuridine
-	D03.383.742.680.852.300.150 Bromodeoxyuridine
-	D03.383.742.680.852.300.350 Floxuridine
-	D03.383.742.680.852.300.400 Idoxuridine
-	D03.383.742.680.852.628 Pseudouridine
-	D03.383.742.680.852.800 Tetrahydrouridine
-	D03.383.742.680.852.829 Thiouridine
-	D03.383.742.686 Pyrimidine Nucleotides
-	D03.383.742.686.050 Apurinic Acid
-	D03.383.742.686.246 Cytosine Nucleotides
-	D03.383.742.686.246.050 Arabinofuranosylcytosine Triphosphate
-	D03.383.742.686.246.115 Cyclic CMP
-	D03.383.742.686.246.150 Cytidine Diphosphate
-	D03.383.742.686.246.150.180 Cytidine Diphosphate Choline
-	D03.383.742.686.246.150.210 Cytidine Diphosphate Diglycerides
-	D03.383.742.686.246.370 Cytidine Monophosphate
-	D03.383.742.686.246.370.250 Cytidine Monophosphate N-Acetylneuraminic Acid
-	D03.383.742.686.246.400 Cytidine Triphosphate
-	D03.383.742.686.246.425 Deoxycytosine Nucleotides
-	D03.383.742.686.246.425.300 Deoxycytidine Monophosphate
-	D03.383.742.686.600 Pyrimidine Dimers
-	D03.383.742.686.706 Thymine Nucleotides
-	D03.383.742.686.706.788 Thymidine Monophosphate
-	D03.383.742.686.850 Uracil Nucleotides
-	D03.383.742.686.850.210 Deoxyuracil Nucleotides
-	D03.383.742.686.850.210.200 Fluorodeoxyuridylate
-	D03.383.742.686.850.600 Uridine Diphosphate
-	D03.383.742.686.850.600.677 Uridine Diphosphate Sugars
-	D03.383.742.686.850.600.677.100 Uridine Diphosphate N-Acetylgalactosamine
-	D03.383.742.686.850.600.677.120 Uridine Diphosphate N-Acetylglucosamine

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D03.383.742.686.850.600.677.150 Uridine Diphosphate N-Acetylmuramic Acid
-	D03.383.742.686.850.600.677.300 Uridine Diphosphate Galactose
-	D03.383.742.686.850.600.677.350 Uridine Diphosphate Glucose
-	D03.383.742.686.850.600.677.375 Uridine Diphosphate Glucuronic Acid
-	D03.383.742.686.850.600.677.800 Uridine Diphosphate Xylose
-	D03.383.742.686.850.877 Uridine Monophosphate
-	D03.383.742.686.850.877.500 Sofosbuvir
-	D03.383.742.686.850.950 Uridine Triphosphate
-	D03.383.742.698 Pyrimidinones
-	D03.383.742.698.122 Alloxan
-	D03.383.742.698.253 Barbiturates
-	D03.383.742.698.253.077 Amobarbital
-	D03.383.742.698.253.150 Barbital
-	D03.383.742.698.253.317 Hexobarbital
-	D03.383.742.698.253.434 Mephobarbital
-	D03.383.742.698.253.488 Methohexital
-	D03.383.742.698.253.500 Murexide
-	D03.383.742.698.253.593 Pentobarbital
-	D03.383.742.698.253.650 Phenobarbital
-	D03.383.742.698.253.650.700 Primidone
-	D03.383.742.698.253.762 Secobarbital
-	D03.383.742.698.253.800 Thiobarbiturates
-	D03.383.742.698.253.800.750 Thiamylal
-	D03.383.742.698.253.800.810 Thiopental
-	D03.383.742.698.421 Cytosine
-	D03.383.742.698.421.431 Flucytosine
-	D03.383.742.698.421.500 5-Methylcytosine
-	D03.383.742.698.553 Lopinavir
-	D03.383.742.698.685 Risperidone
-	D03.383.742.698.700 Ritanserin
-	D03.383.742.698.715 Sparsomycin
-	D03.383.742.698.875 Uracil
-	D03.383.742.698.875.237 Bromouracil
-	D03.383.742.698.875.404 Fluorouracil
-	D03.383.742.698.875.404.425 Capecitabine
-	D03.383.742.698.875.404.850 Tegafur



## MeSH Tree Changes for 2017

Type	Tree - heading
-	D03.383.742.698.875.480 Hydroxyphenylazouracil
-	D03.383.742.698.875.660 Orotic Acid
-	D03.383.742.698.875.700 Pentoxyl
-	D03.383.742.698.875.842 Thiouracil
-	D03.383.742.698.875.842.455 Methylthiouracil
-	D03.383.742.698.875.842.708 Propylthiouracil
-	D03.383.742.698.875.899 Thymine
-	D03.383.742.698.875.949 Uracil Mustard
-	D03.383.742.715 Pyrithiamine
-	D03.383.742.755 Rilpivirine
-	D03.383.742.755.500 Emtricitabine, Rilpivirine, Tenofovir Drug Combination
-	D03.383.742.775 Rosuvastatin Calcium
-	D03.383.742.795 Thiamine
-	D03.383.742.795.300 Fursultiamin
-	D03.383.742.795.600 Thiamine Monophosphate
-	D03.383.742.795.702 Thiamine Pyrophosphate
-	D03.383.742.795.900 Thiamine Triphosphate
-	D03.383.742.875 Trapidil
-	D03.383.742.906 Trimethoprim
-	D03.383.742.906.500 Trimethoprim, Sulfamethoxazole Drug Combination
-	D03.383.773 Pyrrolidines
-	D03.383.773.050 Anisomycin
-	D03.383.773.107 Bepidil
-	D03.383.773.165 Clemastine
-	D03.383.773.170 3,4-Dichloro-N-methyl-N-(2-(1-pyrrolidinyl)-cyclohexyl)-benzeneacetamide, (trans)-Isomer
-	D03.383.773.342 Glycopyrrolate
-	D03.383.773.371 Imino Furanoses
-	D03.383.773.400 Kainic Acid
-	D03.383.773.532 Lincosamides
-	D03.383.773.532.500 Lincomycin
-	D03.383.773.532.500.125 Clindamycin
-	D03.383.773.589 Nafoxidine
-	D03.383.773.610 Nitromifene
-	D03.383.773.630 N-Nitrosopyrrolidine
-	D03.383.773.700 Pentolinium Tartrate

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D03.383.773.728 Procyclidine
-	D03.383.773.812 Pyrrolidinones
-	D03.383.773.812.180 Cotinine
-	D03.383.773.812.226 Doxapram
-	D03.383.773.812.498 Oxotremorine
-	D03.383.773.812.555 Piracetam
-	D03.383.773.812.615 Povidone
-	D03.383.773.812.615.630 Povidone-Iodine
-	D03.383.773.812.718 Pyrrolidonecarboxylic Acid
-	D03.383.773.812.752 Raltegravir Potassium
-	D03.383.773.812.785 Rolipram
-	D03.383.773.812.852 Succinimides
-	D03.383.773.812.852.150 Bromosuccinimide
-	D03.383.773.812.852.333 Ethosuximide
-	D03.383.773.812.900 Tenuazonic Acid
-	D03.383.773.906 Tremorine
-	D03.383.855 Thiazines
-	D03.383.855.150 Chlormezanone
-	D03.383.855.400 Nifurtimox
-	D03.383.855.500 Piroxicam
-	D03.383.855.785 Thiadiazines
-	D03.383.855.985 Xylazine
-	D03.383.886 Thiepins
-	D03.383.886.702 Thiazepines
-	D03.383.903 Thiophenes
-	D03.383.903.075 Canagliflozin
-	D03.383.903.150 Carticaine
-	D03.383.903.260 Duloxetine Hydrochloride
-	D03.383.903.315 Prasugrel Hydrochloride
-	D03.383.903.370 Ketotifen
-	D03.383.903.440 Morantel
-	D03.383.903.580 Pizotyline
-	D03.383.903.634 Pyrantel
-	D03.383.903.634.600 Pyrantel Pamoate
-	D03.383.903.634.670 Pyrantel Tartrate
-	D03.383.903.727 Rivaroxaban

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D03.383.903.820      Thenoyltrifluoroacetone
-	D03.383.903.830      Thienopyridines
-	D03.383.903.830.500      Ticlopidine
-	D03.383.903.840      Ticrynafen
-	D03.383.903.850      Tiletamine
-	D03.383.931      Triazines
-	D03.383.931.095      Almitrine
-	D03.383.931.135      Altretamine
-	D03.383.931.190      Apazone
-	D03.383.931.247      Atrazine
-	D03.383.931.320      Ferrozine
-	D03.383.931.640      Oxonic Acid
-	D03.383.931.674      Prometryne
-	D03.383.931.819      Simazine
-	D03.383.931.915      Triethylenemelamine
Old Tree	<b>D03.438</b> <b>Heterocyclic Compounds, 2-Ring</b>
Old Tree	<b>D03.438.079</b> <b>Benzazepines</b>
Old Tree	<b>D03.438.079.080</b> <b>Benzodiazepines</b>
Old Tree	<b>D03.438.079.080.030</b> <b>Alprazolam</b>
Old Tree	<b>D03.438.079.080.070</b> <b>Benzodiazepinones</b>
Old Tree	<b>D03.438.079.080.070.050</b> <b>Anthramycin</b>
Old Tree	<b>D03.438.079.080.070.110</b> <b>Bromazepam</b>
Old Tree	<b>D03.438.079.080.070.150</b> <b>Clonazepam</b>
Old Tree	<b>D03.438.079.080.070.200</b> <b>Devazepide</b>
Old Tree	<b>D03.438.079.080.070.216</b> <b>Diazepam</b>
Old Tree	<b>D03.438.079.080.070.216.500</b> <b>Nordazepam</b>
Old Tree	<b>D03.438.079.080.070.305</b> <b>Flumazenil</b>
Old Tree	<b>D03.438.079.080.070.320</b> <b>Flunitrazepam</b>
Old Tree	<b>D03.438.079.080.070.348</b> <b>Flurazepam</b>
Old Tree	<b>D03.438.079.080.070.450</b> <b>Lorazepam</b>
Old Tree	<b>D03.438.079.080.070.565</b> <b>Nitrazepam</b>
Old Tree	<b>D03.438.079.080.070.663</b> <b>Oxazepam</b>
Old Tree	<b>D03.438.079.080.070.750</b> <b>Pirenzepine</b>
Old Tree	<b>D03.438.079.080.070.784</b> <b>Prazepam</b>
Old Tree	<b>D03.438.079.080.070.880</b> <b>Temazepam</b>
Old Tree	<b>D03.438.079.080.150</b> <b>Chlordiazepoxide</b>

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	D03.438.079.080.180 Clorazepate Dipotassium
Old Tree	D03.438.079.080.250 Estazolam
Old Tree	D03.438.079.080.550 Medazepam
Old Tree	D03.438.079.080.575 Midazolam
Old Tree	D03.438.079.080.900 Triazolam
Old Tree	D03.438.079.150 Diltiazem
Old Tree	D03.438.079.300 Fenoldopam
Old Tree	D03.438.079.525 Galantamine
Old Tree	D03.438.079.533 Harringtonines
Old Tree	D03.438.079.800 benzazepine 2,3,4,5-Tetrahydro-7,8-dihydroxy-1-phenyl-1H-3-
Old Tree	D03.438.079.900 Varenicline
Old Tree	D03.438.103 Benzimidazoles
Old Tree	D03.438.103.034 2-Pyridinylmethylsulfinylbenzimidazoles
Old Tree	D03.438.103.034.249 Lansoprazole
Old Tree	D03.438.103.034.249.500 Dexlansoprazole
Old Tree	D03.438.103.034.500 Omeprazole
Old Tree	D03.438.103.034.500.500 Esomeprazole
Old Tree	D03.438.103.034.750 Rabeprazole
Old Tree	D03.438.103.070 Albendazole
Old Tree	D03.438.103.105 Astemizole
Old Tree	D03.438.103.123 Bendamustine Hydrochloride
Old Tree	D03.438.103.140 Benomyl
Old Tree	D03.438.103.145 Bisbenzimidazole
Old Tree	D03.438.103.190 Cambendazole
Old Tree	D03.438.103.280 Dabigatran
Old Tree	D03.438.103.370 Domperidone
Old Tree	D03.438.103.393 Droperidol
Old Tree	D03.438.103.450 Fenbendazole
Old Tree	D03.438.103.600 Mebendazole
Old Tree	D03.438.103.618 Mibefradil
Old Tree	D03.438.103.637 Nocodazole
Old Tree	D03.438.103.732 Pimozide
Old Tree	D03.438.103.850 Thiabendazole
Old Tree	D03.438.115 Benzodioxoles
Old Tree	D03.438.115.600 Piperonyl Butoxide

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	D03.438.115.750 Safrole
Old Tree	D03.438.127 Benzofurans
Old Tree	D03.438.127.075 Amiodarone
Old Tree	D03.438.127.110 Benzbromarone
Old Tree	D03.438.127.125 Cantharidin
Old Tree	D03.438.127.187 Citalopram
Old Tree	D03.438.127.218 Fluorescamine
Old Tree	D03.438.127.250 Fura-2
Old Tree	D03.438.127.275 Griseofulvin
Old Tree	D03.438.127.637 Pterocarpans
Old Tree	D03.438.127.818 Vilazodone Hydrochloride
Old Tree	D03.438.150 Benzopyrans
Old Tree	D03.438.150.119 Aflatoxins
Old Tree	D03.438.150.119.075 Aflatoxin B1
Old Tree	D03.438.150.119.100 Aflatoxin M1
Old Tree	D03.438.150.240 Chromans
Old Tree	D03.438.150.240.190 Catechin
Old Tree	D03.438.150.240.225 Centchroman
Old Tree	D03.438.150.266 Chromones
Old Tree	D03.438.150.266.300 Cromolyn Sodium
Old Tree	D03.438.150.266.450 Flavonoids
Old Tree	D03.438.150.266.450.087 Anthocyanins
Old Tree	D03.438.150.266.450.175 Benzoflavones
Old Tree	D03.438.150.266.450.175.100 beta-Naphthoflavone
Old Tree	D03.438.150.266.450.190 Biflavonoids
Old Tree	D03.438.150.266.450.206 Catechin
Old Tree	D03.438.150.266.450.221 Chalcones
Old Tree	D03.438.150.266.450.221.500 Chalcone
Old Tree	D03.438.150.266.450.252 Flavanones
Old Tree	D03.438.150.266.450.252.500 Hesperidin
Old Tree	D03.438.150.266.450.260 Flavones
Old Tree	D03.438.150.266.450.260.110 Apigenin
Old Tree	D03.438.150.266.450.260.222 Diosmin
Old Tree	D03.438.150.266.450.260.444 Flavoxate
Old Tree	D03.438.150.266.450.260.555 Luteolin
Old Tree	D03.438.150.266.450.260.777 Polyphenols

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	D03.438.150.266.450.268                      Flavonolignans
Old Tree	D03.438.150.266.450.268.777                      Silymarin
Old Tree	D03.438.150.266.450.284                      Flavonols
Old Tree	D03.438.150.266.450.284.388                      Kaempferols
Old Tree	D03.438.150.266.450.284.777                      Quercetin
Old Tree	D03.438.150.266.450.284.888                      Rutin
Old Tree	D03.438.150.266.450.284.888.500                      Hydroxyethylrutoside
Old Tree	D03.438.150.266.450.400                      Isoflavones
Old Tree	D03.438.150.266.450.400.187                      Coumestrol
Old Tree	D03.438.150.266.450.400.281                      Equol
Old Tree	D03.438.150.266.450.400.375                      Genistein
Old Tree	D03.438.150.266.450.400.687                      Pterocarpans
Old Tree	D03.438.150.266.450.400.843                      Rotenone
Old Tree	D03.438.150.266.450.700                      Proanthocyanidins
Old Tree	D03.438.150.320                      Citrinin
Old Tree	D03.438.150.446                      Coumarins
Old Tree	D03.438.150.446.139                      Aminocoumarins
Old Tree	D03.438.150.446.139.500                      Novobiocin
Old Tree	D03.438.150.446.280                      Chromonar
Old Tree	D03.438.150.446.350                      Coumestrol
Old Tree	D03.438.150.446.400                      Esculin
Old Tree	D03.438.150.446.520                      4-Hydroxycoumarins
Old Tree	D03.438.150.446.520.079                      Acenocoumarol
Old Tree	D03.438.150.446.520.203                      Dicumarol
Old Tree	D03.438.150.446.520.451                      Ethyl Biscoumacetate
Old Tree	D03.438.150.446.520.750                      Phenprocoumon
Old Tree	D03.438.150.446.520.914                      Warfarin
Old Tree	D03.438.150.446.598                      Isocoumarins
Old Tree	D03.438.150.446.598.500                      Ochratoxins
Old Tree	D03.438.150.446.794                      Furocoumarins
Old Tree	D03.438.150.446.794.200                      Ficusin
Old Tree	D03.438.150.446.794.300                      Khellin
Old Tree	D03.438.150.446.794.500                      Methoxsalen
Old Tree	D03.438.150.446.794.875                      Trioxsalen
Old Tree	D03.438.150.446.853                      Pyranocoumarins
Old Tree	D03.438.150.446.912                      Umbelliferones

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	D03.438.150.446.912.326                      Coumaphos
Old Tree	D03.438.150.446.912.531                      Hymecromone
Old Tree	D03.438.150.446.912.850                      Scopoletin
Old Tree	D03.438.150.455                                  Cromakalim
Old Tree	D03.438.150.500                                  Ellagic Acid
Old Tree	D03.438.150.600                                  Hematoxylin
Old Tree	D03.438.150.755                                  Nebivolol
Old Tree	D03.438.150.909                                  Vitamin E
Old Tree	D03.438.150.909.750                              Tocopherols
Old Tree	D03.438.150.909.750.249                              alpha-Tocopherol
Old Tree	D03.438.150.909.750.374                              beta-Tocopherol
Old Tree	D03.438.150.909.750.500                              gamma-Tocopherol
Old Tree	D03.438.150.909.875                                  Tocotrienols
Old Tree	D03.438.174    Benzothiadiazines
Old Tree	D03.438.174.138    Bendroflumethiazide
Old Tree	D03.438.174.261    Chlorothiazide
Old Tree	D03.438.174.261.476    Hydrochlorothiazide
Old Tree	D03.438.174.261.476.716    Trichlormethiazide
Old Tree	D03.438.174.285    Cyclopenthiazide
Old Tree	D03.438.174.300    Diazoxide
Old Tree	D03.438.174.475    Hydroflumethiazide
Old Tree	D03.438.174.620    Methyclothiazide
Old Tree	D03.438.174.781    Polythiazide
Old Tree	D03.438.185    Benzothiazoles
Old Tree	D03.438.185.222    Dithiazanine
Old Tree	D03.438.185.416    Ethoxzolamide
Old Tree	D03.438.185.611    Riluzole
Old Tree	D03.438.185.708    Saccharin
Old Tree	D03.438.197    Benzothiepins
Old Tree	D03.438.197.408    Endosulfan
Old Tree	D03.438.209    Benzoxazines
Old Tree	D03.438.221    Benzoxazoles
Old Tree	D03.438.221.173    Calcimycin
Old Tree	D03.438.221.346    Chlorzoxazone
Old Tree	D03.438.221.370    Cialit
Old Tree	D03.438.221.950    Zoxazolamine

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	D03.438.245 Benzoxepins
Old Tree	D03.438.260 Bicyclo Compounds, Heterocyclic
Old Tree	D03.438.260 Bridged Bicyclo Compounds, Heterocyclic
Old Tree	D03.438.260.825 Penicillins
Old Tree	D03.438.260.825.124 Amdinocillin
Old Tree	D03.438.260.825.124.036 Amdinocillin Pivoxil
Old Tree	D03.438.260.825.249 Cyclacillin
Old Tree	D03.438.260.825.500 Methicillin
Old Tree	D03.438.260.825.562 Nafcillin
Old Tree	D03.438.260.825.625 Oxacillin
Old Tree	D03.438.260.825.625.150 Cloxacillin
Old Tree	D03.438.260.825.625.150.205 Dicloxacillin
Old Tree	D03.438.260.825.625.150.250 Floxacillin
Old Tree	D03.438.260.825.687 Penicillanic Acid
Old Tree	D03.438.260.825.750 Penicillin G
Old Tree	D03.438.260.825.750.050 Ampicillin
Old Tree	D03.438.260.825.750.050.050 Amoxicillin
Old Tree	D03.438.260.825.750.050.050.060 Amoxicillin-Potassium Clavulanate Combination
Old Tree	D03.438.260.825.750.050.075 Azlocillin
Old Tree	D03.438.260.825.750.050.500 Mezlocillin
Old Tree	D03.438.260.825.750.050.650 Piperacillin
Old Tree	D03.438.260.825.750.050.700 Pivampicillin
Old Tree	D03.438.260.825.750.050.900 Talampicillin
Old Tree	D03.438.260.825.750.170 Carbenicillin
Old Tree	D03.438.260.825.750.170.200 Carfecillin
Old Tree	D03.438.260.825.750.685 Penicillin G Benzathine
Old Tree	D03.438.260.825.750.695 Penicillin G Procaine
Old Tree	D03.438.260.825.750.875 Sulbenicillin
Old Tree	D03.438.260.825.781 Penicillin V
Old Tree	D03.438.260.825.812 Sulbactam
Old Tree	D03.438.260.825.875 Ticarcillin
Old Tree	D03.438.449 Indazoles
Old Tree	D03.438.449.130 Benzydamine
Old Tree	D03.438.449.350 Granisetron
Old Tree	D03.438.473 Indoles



## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	D03.438.473.025 Adrenochrome
Old Tree	D03.438.473.050 Alcian Blue
Old Tree	D03.438.473.144 Carbazoles
Old Tree	D03.438.473.144.249 Ellipticines
Old Tree	D03.438.473.144.500 Ondansetron
Old Tree	D03.438.473.144.750 Staurosporine
Old Tree	D03.438.473.155 Carbolines
Old Tree	D03.438.473.155.500 Tadalafil
Old Tree	D03.438.473.231 Cytochalasins
Old Tree	D03.438.473.231.370 Cytochalasin B
Old Tree	D03.438.473.231.450 Cytochalasin D
Old Tree	D03.438.473.250 Delavirdine
Old Tree	D03.438.473.360 Gliotoxin
Old Tree	D03.438.473.385 Hydroxytryptophol
Old Tree	D03.438.473.390 Ibogaine
Old Tree	D03.438.473.391 Indapamide
Old Tree	D03.438.473.393 Indican
Old Tree	D03.438.473.396 Indigo Carmine
Old Tree	D03.438.473.400 Indocyanine Green
Old Tree	D03.438.473.402 Indole Alkaloids
Old Tree	D03.438.473.402.444 Harmaline
Old Tree	D03.438.473.402.477 Harmine
Old Tree	D03.438.473.402.511 Lyngbya Toxins
Old Tree	D03.438.473.402.545 Physostigmine
Old Tree	D03.438.473.402.613 Psilocybin
Old Tree	D03.438.473.402.613 Psilocybine
Old Tree	D03.438.473.402.681 Secologanin Tryptamine Alkaloids
Old Tree	D03.438.473.402.681.077 Ajmaline
Old Tree	D03.438.473.402.681.077.324 Lorajmine
Old Tree	D03.438.473.402.681.077.650 Prajmaline
Old Tree	D03.438.473.402.681.333 Ellipticines
Old Tree	D03.438.473.402.681.444 Ibogaine
Old Tree	D03.438.473.402.681.722 Strychnine
Old Tree	D03.438.473.402.681.827 Vinca Alkaloids
Old Tree	D03.438.473.402.681.827.650 Vinblastine
Old Tree	D03.438.473.402.681.827.750 Vincamine

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	D03.438.473.402.681.827.817 Vincristine
Old Tree	D03.438.473.402.681.827.830 Vindesine
Old Tree	D03.438.473.402.681.933 Yohimbine
Old Tree	D03.438.473.402.681.933.500 Reserpine
Old Tree	D03.438.473.402.750 Staurosporine
Old Tree	D03.438.473.404 Indoleacetic Acids
Old Tree	D03.438.473.404.200 Etodolac
Old Tree	D03.438.473.404.478 Hydroxyindoleacetic Acid
Old Tree	D03.438.473.412 Indolequinones
Old Tree	D03.438.473.412.249 Mitomycins
Old Tree	D03.438.473.412.249.350 Mitomycin
Old Tree	D03.438.473.412.249.700 Porfiromycin
Old Tree	D03.438.473.412.500 Pyrroloiminoquinones
Old Tree	D03.438.473.420 Indomethacin
Old Tree	D03.438.473.420.485 Ketorolac
Old Tree	D03.438.473.420.742 Ketorolac Tromethamine
Old Tree	D03.438.473.432 Indoramin
Old Tree	D03.438.473.469 Iprindole
Old Tree	D03.438.473.525 Isatin
Old Tree	D03.438.473.615 Methisazone
Old Tree	D03.438.473.629 Molindone
Old Tree	D03.438.473.725 Oxyphenisatin Acetate
Old Tree	D03.438.473.766 Perindopril
Old Tree	D03.438.473.808 Skatole
Old Tree	D03.438.473.840 Sporidesmins
Old Tree	D03.438.473.914 Tryptamines
Old Tree	D03.438.473.914.201 Dihydroxytryptamines
Old Tree	D03.438.473.914.201.259 5,6-Dihydroxytryptamine
Old Tree	D03.438.473.914.201.263 5,7-Dihydroxytryptamine
Old Tree	D03.438.473.914.237 N,N-Dimethyltryptamine
Old Tree	D03.438.473.914.237.150 Bufotenin
Old Tree	D03.438.473.914.237.150.500 Methoxydimethyltryptamines
Old Tree	D03.438.473.914.481 Melatonin
Old Tree	D03.438.473.914.700 Psilocybin
Old Tree	D03.438.473.914.700 Psilocybine
Old Tree	D03.438.473.914.814 Serotonin

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	D03.438.473.914.814.150 <span style="float: right;">Bufotenin</span>
Old Tree	D03.438.473.914.814.150.500 <span style="float: right;">Methoxydimethyltryptamines</span>
Old Tree	D03.438.473.914.814.400 <span style="float: right;">5-Methoxytryptamine</span>
Old Tree	D03.438.473.914.907 <span style="float: right;">Sumatriptan</span>
Old Tree	D03.438.473.957 <span style="float: right;">Vilazodone Hydrochloride</span>
Old Tree	D03.438.496 <span style="float: right;">Indolizines</span>
Old Tree	D03.438.496.500 <span style="float: right;">Indolizidines</span>
Old Tree	D03.438.513 <span style="float: right;">Isoindoles</span>
Old Tree	D03.438.513.124 <span style="float: right;">Captan</span>
Old Tree	D03.438.513.249 <span style="float: right;">Chlorisondamine</span>
Old Tree	D03.438.513.374 <span style="float: right;">Indoprofen</span>
Old Tree	D03.438.513.437 <span style="float: right;">Lurasidone Hydrochloride</span>
Old Tree	D03.438.513.500 <span style="float: right;">Mazindol</span>
Old Tree	D03.438.513.750 <span style="float: right;">Phthalimides</span>
Old Tree	D03.438.513.750.500 <span style="float: right;">Chlorthalidone</span>
Old Tree	D03.438.513.750.625 <span style="float: right;">Phosmet</span>
Old Tree	D03.438.513.750.750 <span style="float: right;">Thalidomide</span>
Old Tree	D03.438.531 <span style="float: right;">Isoquinolines</span>
Old Tree	D03.438.531.085 <span style="float: right;">Benzylisoquinolines</span>
Old Tree	D03.438.531.085.030 <span style="float: right;">Aporphines</span>
Old Tree	D03.438.531.085.030.290 <span style="float: right;">Apomorphine</span>
Old Tree	D03.438.531.085.061 <span style="float: right;">Atracurium</span>
Old Tree	D03.438.531.085.077 <span style="float: right;">Bicuculline</span>
Old Tree	D03.438.531.085.666 <span style="float: right;">Papaverine</span>
Old Tree	D03.438.531.085.666.850 <span style="float: right;">Tetrahydropapaveroline</span>
Old Tree	D03.438.531.085.777 <span style="float: right;">Toxiferine</span>
Old Tree	D03.438.531.085.777.050 <span style="float: right;">Alcuronium</span>
Old Tree	D03.438.531.085.888 <span style="float: right;">Tretoquinol</span>
Old Tree	D03.438.531.085.944 <span style="float: right;">Tubocurarine</span>
Old Tree	D03.438.531.205 <span style="float: right;">Debrisoquin</span>
Old Tree	D03.438.531.321 <span style="float: right;">Emetine</span>
Old Tree	D03.438.531.400 <span style="float: right;">1-(5-Isoquinolinesulfonyl)-2-Methylpiperazine</span>
Old Tree	D03.438.531.460 <span style="float: right;">Naphthalimides</span>
Old Tree	D03.438.531.520 <span style="float: right;">Nelfinavir</span>
Old Tree	D03.438.531.535 <span style="float: right;">Nomifensine</span>
Old Tree	D03.438.531.567 <span style="float: right;">Noscapine</span>

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	D03.438.531.690 Praziquantel
Old Tree	D03.438.531.770 Saquinavir
Old Tree	D03.438.531.820 Tetrahydroisoquinolines
Old Tree	D03.438.531.820.500 Salsoline Alkaloids
Old Tree	D03.438.531.820.594 Solifenacin Succinate
Old Tree	D03.438.531.820.687 Toxiferine
Old Tree	D03.438.531.820.687.050 Alcuronium
Old Tree	D03.438.531.820.875 Tubocurarine
Old Tree	D03.438.612 Naphthyridines
Old Tree	D03.438.612.500 Nalidixic Acid
Old Tree	D03.438.733 Pteridines
Old Tree	D03.438.733.315 Flavins
Old Tree	D03.438.733.315.650 Riboflavin
Old Tree	D03.438.733.315.650.249 Flavin-Adenine Dinucleotide
Old Tree	D03.438.733.315.650.500 Flavin Mononucleotide
Old Tree	D03.438.733.631 Pterins
Old Tree	D03.438.733.631.192 Aminopterin
Old Tree	D03.438.733.631.192.500 Methotrexate
Old Tree	D03.438.733.631.202 Biopterin
Old Tree	D03.438.733.631.202.500 Neopterin
Old Tree	D03.438.733.631.400 Folic Acid
Old Tree	D03.438.733.631.400.600 Pteroylpolyglutamic Acids
Old Tree	D03.438.733.631.400.800 Tetrahydrofolates
Old Tree	D03.438.733.631.400.800.350 Formyltetrahydrofolates
Old Tree	D03.438.733.631.400.800.350.450 Leucovorin
Old Tree	D03.438.733.631.400.800.350.450.500 Levoleucovorin
Old Tree	D03.438.733.631.825 Xanthopterin
Old Tree	D03.438.733.900 Triamterene
Old Tree	D03.438.759 Purines
Old Tree	D03.438.759.138 Adenine
Old Tree	D03.438.759.138.050 2-Aminopurine
Old Tree	D03.438.759.138.525 Cytokinins
Old Tree	D03.438.759.138.525.350 Isopentenyladenosine
Old Tree	D03.438.759.138.525.400 Kinetin
Old Tree	D03.438.759.138.525.700 Zeatin
Old Tree	D03.438.759.138.881 Tenofovir

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	D03.438.759.138.881.125 Fumarate Drug Combination Efavirenz, Emtricitabine, Tenofovir Disoproxil
Old Tree	D03.438.759.138.881.250 Disoproxil Fumarate Drug Combination Elvitegravir, Cobicistat, Emtricitabine, Tenofovir
Old Tree	D03.438.759.138.881.500 Combination Emtricitabine, Rilpivirine, Tenofovir Drug
Old Tree	D03.438.759.138.881.750 Combination Emtricitabine, Tenofovir Disoproxil Fumarate Drug
Old Tree	D03.438.759.160 Allopurinol
Old Tree	D03.438.759.534 6-Mercaptopurine
Old Tree	D03.438.759.534.090 Azathioprine
Old Tree	D03.438.759.562 Linagliptin
Old Tree	D03.438.759.590 Purine Nucleosides
Old Tree	D03.438.759.590.138 Adenosine
Old Tree	D03.438.759.590.138.025 Adenosine-5'-(N-ethylcarboxamide)
Old Tree	D03.438.759.590.138.240 S-Adenosylhomocysteine
Old Tree	D03.438.759.590.138.264 S-Adenosylmethionine
Old Tree	D03.438.759.590.138.300 2-Chloroadenosine
Old Tree	D03.438.759.590.138.300.200 Cladribine
Old Tree	D03.438.759.590.138.325 Deoxyadenosines
Old Tree	D03.438.759.590.138.325.075 Cladribine
Old Tree	D03.438.759.590.138.325.105 Dideoxyadenosine
Old Tree	D03.438.759.590.138.325.800 Puromycin Aminonucleoside
Old Tree	D03.438.759.590.138.500 Isopentenyladenosine
Old Tree	D03.438.759.590.138.630 Phenylisopropyladenosine
Old Tree	D03.438.759.590.138.711 Puromycin
Old Tree	D03.438.759.590.138.711.650 Puromycin Aminonucleoside
Old Tree	D03.438.759.590.138.900 Vidarabine
Old Tree	D03.438.759.590.454 Guanosine
Old Tree	D03.438.759.590.454.240 Deoxyguanosine
Old Tree	D03.438.759.590.454.500 Nucleoside Q
Old Tree	D03.438.759.590.616 Inosine
Old Tree	D03.438.759.590.616.130 Didanosine
Old Tree	D03.438.759.590.616.450 Inosine Pranobex
Old Tree	D03.438.759.590.616.900 Thioinosine
Old Tree	D03.438.759.590.616.900.500 Methylthioinosine
Old Tree	D03.438.759.590.910 Tubercidin

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	D03.438.759.646 Purine Nucleotides
Old Tree	D03.438.759.646.138 Adenine Nucleotides
Old Tree	D03.438.759.646.138.124 Adenosine Diphosphate
Old Tree	D03.438.759.646.138.124.070 Adenosine Diphosphate Sugars
Old Tree	D03.438.759.646.138.124.070.075 Adenosine Diphosphate Glucose
Old Tree	D03.438.759.646.138.124.070.125 Adenosine Diphosphate Ribose
Old Tree	D03.438.759.646.138.124.070.125.040 O-Acetyl-ADP-Ribose
Old Tree	D03.438.759.646.138.124.070.125.195 Cyclic ADP-Ribose
Old Tree	D03.438.759.646.138.180 Adenosine Monophosphate
Old Tree	D03.438.759.646.138.180.080 Adenosine Phosphosulfate
Old Tree	D03.438.759.646.138.236 Adenosine Triphosphate
Old Tree	D03.438.759.646.138.236.050 Adenylyl Imidodiphosphate
Old Tree	D03.438.759.646.138.236.250 Ethenoadenosine Triphosphate
Old Tree	D03.438.759.646.138.382 Coenzyme A
Old Tree	D03.438.759.646.138.382.300 Acyl Coenzyme A
Old Tree	D03.438.759.646.138.382.300.020 Acetyl Coenzyme A
Old Tree	D03.438.759.646.138.382.300.500 Malonyl Coenzyme A
Old Tree	D03.438.759.646.138.382.300.700 Palmitoyl Coenzyme A
Old Tree	D03.438.759.646.138.395 Cyclic AMP
Old Tree	D03.438.759.646.138.395.225 8-Bromo Cyclic Adenosine Monophosphate
Old Tree	D03.438.759.646.138.395.250 Bucladesine
Old Tree	D03.438.759.646.138.410 Deoxyadenine Nucleotides
Old Tree	D03.438.759.646.138.506 Flavin-Adenine Dinucleotide
Old Tree	D03.438.759.646.138.694 NAD
Old Tree	D03.438.759.646.138.749 NADP
Old Tree	D03.438.759.646.138.850 Phosphoadenosine Phosphosulfate
Old Tree	D03.438.759.646.138.925 Vidarabine Phosphate
Old Tree	D03.438.759.646.454 Guanine Nucleotides
Old Tree	D03.438.759.646.454.160 Cyclic GMP
Old Tree	D03.438.759.646.454.160.325 Dibutyryl Cyclic GMP
Old Tree	D03.438.759.646.454.200 Deoxyguanine Nucleotides
Old Tree	D03.438.759.646.454.340 Guanosine Diphosphate
Old Tree	D03.438.759.646.454.340.350 Guanosine Diphosphate Sugars
Old Tree	D03.438.759.646.454.340.350.400 Guanosine Diphosphate Fucose
Old Tree	D03.438.759.646.454.340.350.500 Guanosine Diphosphate Mannose
Old Tree	D03.438.759.646.454.440 Guanosine Pentaphosphate

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	D03.438.759.646.454.480 Guanosine Tetraphosphate
Old Tree	D03.438.759.646.454.504 Guanosine Triphosphate
Old Tree	D03.438.759.646.454.504.400 Guanylyl Imidodiphosphate
Old Tree	D03.438.759.646.454.525 Guanosine Monophosphate
Old Tree	D03.438.759.646.454.700 RNA Caps
Old Tree	D03.438.759.646.454.700.710 RNA Cap Analogs
Old Tree	D03.438.759.646.616 Inosine Nucleotides
Old Tree	D03.438.759.646.616.300 Cyclic IMP
Old Tree	D03.438.759.646.616.400 Inosine Diphosphate
Old Tree	D03.438.759.646.616.500 Inosine Monophosphate
Old Tree	D03.438.759.646.616.800 Inosine Triphosphate
Old Tree	D03.438.759.758 Purinones
Old Tree	D03.438.759.758.399 Hypoxanthines
Old Tree	D03.438.759.758.399.454 Guanine
Old Tree	D03.438.759.758.399.454.250 Acyclovir
Old Tree	D03.438.759.758.399.454.250.300 Ganciclovir
Old Tree	D03.438.759.758.399.454.300 Azaguanine
Old Tree	D03.438.759.758.399.454.650 Pemetrexed
Old Tree	D03.438.759.758.399.475 Hypoxanthine
Old Tree	D03.438.759.758.824 Xanthines
Old Tree	D03.438.759.758.824.175 Caffeine
Old Tree	D03.438.759.758.824.651 Theobromine
Old Tree	D03.438.759.758.824.651.700 Pentoxifylline
Old Tree	D03.438.759.758.824.751 Theophylline
Old Tree	D03.438.759.758.824.751.075 Aminophylline
Old Tree	D03.438.759.758.824.751.162 Dimenhydrinate
Old Tree	D03.438.759.758.824.751.250 Dyphylline
Old Tree	D03.438.759.758.824.751.500 1-Methyl-3-isobutylxanthine
Old Tree	D03.438.759.758.824.751.950 Xanthinol Niacinate
Old Tree	D03.438.759.758.824.877 Uric Acid
Old Tree	D03.438.759.758.824.938 Xanthine
Old Tree	D03.438.759.794 Saxitoxin
Old Tree	D03.438.759.824 Sildenafil Citrate
Old Tree	D03.438.759.854 Thioguanine
Old Tree	D03.438.772 Pyrrolizidine Alkaloids
Old Tree	D03.438.772.500 Monocrotaline

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	D03.438.786 Quinazolines
Old Tree	D03.438.786.375 Erlotinib Hydrochloride
Old Tree	D03.438.786.563 Linagliptin
Old Tree	D03.438.786.750 Prazosin
Old Tree	D03.438.786.750.200 Doxazosin
Old Tree	D03.438.786.830 Quinazolinones
Old Tree	D03.438.786.830.333 Ketanserin
Old Tree	D03.438.786.830.666 Methaqualone
Old Tree	D03.438.786.830.688 Metolazone
Old Tree	D03.438.786.910 Tetrodotoxin
Old Tree	D03.438.786.925 Trimetrexate
Old Tree	D03.438.810 Quinolines
Old Tree	D03.438.810.050 Aminoquinolines
Old Tree	D03.438.810.050.060 Amodiaquine
Old Tree	D03.438.810.050.180 Chloroquine
Old Tree	D03.438.810.050.180.350 Hydroxychloroquine
Old Tree	D03.438.810.050.440 4-Hydroxyaminoquinoline-1-oxide
Old Tree	D03.438.810.050.650 Primaquine
Old Tree	D03.438.810.087 Diarylquinolines
Old Tree	D03.438.810.125 Dibucaine
Old Tree	D03.438.810.200 Ethoxyquin
Old Tree	D03.438.810.350 Hydroxyquinolines
Old Tree	D03.438.810.350.250 Decoquinate
Old Tree	D03.438.810.350.400 Kynurenic Acid
Old Tree	D03.438.810.350.600 Oxamniquine
Old Tree	D03.438.810.350.625 Oxyquinoline
Old Tree	D03.438.810.350.625.250 Chloroquinolinols
Old Tree	D03.438.810.350.625.250.260 Chlorquinaldol
Old Tree	D03.438.810.350.625.400 Clioquinol
Old Tree	D03.438.810.350.625.420 Iodoquinol
Old Tree	D03.438.810.350.700 Procaterol
Old Tree	D03.438.810.410 Mefloquine
Old Tree	D03.438.810.470 Nitroquinolines
Old Tree	D03.438.810.470.450 4-Nitroquinoline-1-oxide
Old Tree	D03.438.810.470.600 Oxamniquine
Old Tree	D03.438.810.553 Pyrroloiminoquinones



## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	D03.438.810.637 Quinaldines
Old Tree	D03.438.810.699 Quinidine
Old Tree	D03.438.810.762 Quinine
Old Tree	D03.438.810.824 Quinolinium Compounds
Old Tree	D03.438.810.824.200 Dequalinium
Old Tree	D03.438.810.824.700 Pyrvinium Compounds
Old Tree	D03.438.810.835 Quinolones
Old Tree	D03.438.810.835.055 4-Quinolones
Old Tree	D03.438.810.835.055.500 Nalidixic Acid
Old Tree	D03.438.810.835.055.550 Nedocromil
Old Tree	D03.438.810.835.055.580 Oxolinic Acid
Old Tree	D03.438.810.835.122 Aripiprazole
Old Tree	D03.438.810.835.188 Carteolol
Old Tree	D03.438.810.835.322 Fluoroquinolones
Old Tree	D03.438.810.835.322.186 Ciprofloxacin
Old Tree	D03.438.810.835.322.186.400 Fleroxacin
Old Tree	D03.438.810.835.322.280 Enoxacin
Old Tree	D03.438.810.835.322.374 Norfloxacin
Old Tree	D03.438.810.835.322.500 Ofloxacin
Old Tree	D03.438.810.835.322.500.500 Levofloxacin
Old Tree	D03.438.810.835.322.750 Pefloxacin
Old Tree	D03.438.810.835.661 PQQ Cofactor
Old Tree	D03.438.810.842 Quinpirole
Old Tree	D03.438.810.850 Quipazine
Old Tree	D03.438.810.900 Saquinavir
Old Tree	D03.438.834 Quinolizines
Old Tree	D03.438.834.700 2H-Benzo(a)quinolizin-2-ol, 2-Ethyl-1,3,4,6,7,11b-hexahydro-3-isobutyl-9,10-dimethoxy-
Old Tree	D03.438.834.737 Quinolizidines
Old Tree	D03.438.834.737.500 Sparteine
Old Tree	D03.438.834.850 Tetrabenazine
Old Tree	D03.438.857 Quinoxalines
Old Tree	D03.438.857.070 Brimonidine Tartrate
Old Tree	D03.438.857.070.500 Brimonidine Tartrate, Timolol Maleate Drug Combination
Old Tree	D03.438.857.140 Carbadox
Old Tree	D03.438.857.160 6-Cyano-7-nitroquinoxaline-2,3-dione

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	D03.438.857.233 Echinomycin
Old Tree	D03.438.857.885 Tyrphostins
Old Tree	D03.438.857.942 Varenicline
Old Tree	D03.438.928 Thienopyridines
Old Tree	D03.438.928.500 Ticlopidine
Old Tree	D03.494 Heterocyclic Compounds, 3-Ring
Old Tree	D03.494.046 Acridines
Old Tree	D03.494.046.109 Acridones
Old Tree	D03.494.046.109.077 Acronine
Old Tree	D03.494.046.250 Aminoacridines
Old Tree	D03.494.046.250.150 Acridine Orange
Old Tree	D03.494.046.250.177 Acriflavine
Old Tree	D03.494.046.250.200 Aminacrine
Old Tree	D03.494.046.250.225 Amsacrine
Old Tree	D03.494.046.250.450 Ethacridine
Old Tree	D03.494.046.250.650 Nitracrine
Old Tree	D03.494.046.250.720 Proflavine
Old Tree	D03.494.046.250.760 Quinacrine
Old Tree	D03.494.046.250.760.750 Quinacrine Mustard
Old Tree	D03.494.046.250.900 Tacrine
Old Tree	D03.494.060 Anthramycin
Old Tree	D03.494.148 Carbazoles
Old Tree	D03.494.148.249 Ellipticines
Old Tree	D03.494.148.500 Ondansetron
Old Tree	D03.494.148.750 Staurosporine
Old Tree	D03.494.154 Carbolines
Old Tree	D03.494.154.333 Harmaline
Old Tree	D03.494.154.344 Harmine
Old Tree	D03.494.154.672 Tadalafil
Old Tree	D03.494.160 Cinoxacin
Old Tree	D03.494.200 Dactinomycin
Old Tree	D03.494.240 Dibenzazepines
Old Tree	D03.494.240.127 Carbamazepine
Old Tree	D03.494.240.194 Clomipramine
Old Tree	D03.494.240.220 Clozapine
Old Tree	D03.494.240.281 Desipramine

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	D03.494.240.485 Imipramine
Old Tree	D03.494.240.520 Lofepamine
Old Tree	D03.494.240.550 Mianserin
Old Tree	D03.494.240.625 Opipramol
Old Tree	D03.494.240.918 Trimipramine
Old Tree	D03.494.276 Dibenzothiazepines
Old Tree	D03.494.276.500 Quetiapine Fumarate
Old Tree	D03.494.311 Dibenzothiepins
Old Tree	D03.494.311.250 Dothiepin
Old Tree	D03.494.311.520 Methiothepin
Old Tree	D03.494.347 Dibenzoxazepines
Old Tree	D03.494.347.200 2-acetylhydrazide Dibenz(b,f)(1,4)oxazepine-10(11H)-carboxylic acid, 8-chloro-,
Old Tree	D03.494.347.500 Loxapine
Old Tree	D03.494.347.500.040 Amoxapine
Old Tree	D03.494.382 Dibenzoxepins
Old Tree	D03.494.382.393 Doxepin
Old Tree	D03.494.382.696 Olopatadine Hydrochloride
Old Tree	D03.494.507 Flavins
Old Tree	D03.494.507.650 Riboflavin
Old Tree	D03.494.507.650.249 Flavin-Adenine Dinucleotide
Old Tree	D03.494.507.650.500 Flavin Mononucleotide
Old Tree	D03.494.633 Phenanthridines
Old Tree	D03.494.633.207 Benzophenanthridines
Old Tree	D03.494.633.416 Ethidium
Old Tree	D03.494.633.700 Propidium
Old Tree	D03.494.669 Phenanthrolines
Old Tree	D03.494.704 Phenazines
Old Tree	D03.494.704.353 Clofazimine
Old Tree	D03.494.704.550 Methylphenazonium Methosulfate
Old Tree	D03.494.704.600 Neutral Red
Old Tree	D03.494.704.700 Pyocyanine
Old Tree	D03.494.741 Phenothiazines
Old Tree	D03.494.741.029 Acepromazine
Old Tree	D03.494.741.080 Azure Stains
Old Tree	D03.494.741.198 Chlorpromazine

## MeSH Tree Changes for 2017

Type	Tree - heading	
Old Tree	D03.494.741.326	Fluphenazine
Old Tree	D03.494.741.470	Mesoridazine
Old Tree	D03.494.741.513	Methotrimeprazine
Old Tree	D03.494.741.517	Methylene Blue
Old Tree	D03.494.741.533	Moricizine
Old Tree	D03.494.741.550	Nonachlazine
Old Tree	D03.494.741.575	Perazine
Old Tree	D03.494.741.593	Perphenazine
Old Tree	D03.494.741.639	Prochlorperazine
Old Tree	D03.494.741.661	Promazine
Old Tree	D03.494.741.670	Promethazine
Old Tree	D03.494.741.780	Thiethylperazine
Old Tree	D03.494.741.843	Thioridazine
Old Tree	D03.494.741.869	Tolonium Chloride
Old Tree	D03.494.741.898	Trifluoperazine
Old Tree	D03.494.741.918	Triflupromazine
Old Tree	D03.494.741.939	Trimeprazine
Old Tree	D03.494.770	Furocoumarins
Old Tree	D03.494.770.200	Ficusin
Old Tree	D03.494.770.300	Khellin
Old Tree	D03.494.770.500	Methoxsalen
Old Tree	D03.494.770.875	Trioxsalen
Old Tree	D03.494.802	Quinpirole
Old Tree	D03.494.819	Simeprevir
Old Tree	D03.494.835	Spectinomycin
Old Tree	D03.494.953	Xanthenes
Old Tree	D03.494.953.275	Fluoresceins
Old Tree	D03.494.953.275.300	Eosine I Bluish
Old Tree	D03.494.953.275.325	Eosine Yellowish-(YS)
Old Tree	D03.494.953.275.350	Erythrosine
Old Tree	D03.494.953.275.390	Fluorescein
Old Tree	D03.494.953.275.400	Fluorescein-5-isothiocyanate
Old Tree	D03.494.953.275.700	Rose Bengal
Old Tree	D03.494.953.558	Propantheline
Old Tree	D03.494.953.570	Pyronine
Old Tree	D03.494.953.600	Rhodamines

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	D03.494.953.600.500 Rhodamine 123
Old Tree	D03.494.953.704 Thioxanthenes
Old Tree	D03.494.953.704.250 Chlorprothixene
Old Tree	D03.494.953.704.269 Clopenthixol
Old Tree	D03.494.953.704.360 Flupenthixol
Old Tree	D03.494.953.704.450 Hycanthone
Old Tree	D03.494.953.704.500 Lucanthone
Old Tree	D03.494.953.704.787 Thiothixene
Old Tree	D03.494.953.852 Xanthenes
Old Tree	D03.494.953.852.388 Lucanthone
Old Tree	D03.494.953.852.777 Sterigmatocystin
Old Tree	D03.549 Heterocyclic Compounds with 4 or More Rings
Old Tree	D03.549 Heterocyclic Compounds, 4 or More Rings
Old Tree	D03.549.095 Aporphines
Old Tree	D03.549.095.290 Apomorphine
Old Tree	D03.549.131 Benzophenanthridines
Old Tree	D03.549.168 Berberine Alkaloids
Old Tree	D03.549.168.100 Berberine
Old Tree	D03.549.256 Cevanes
Old Tree	D03.549.256.310 Germine Acetates
Old Tree	D03.549.256.543 Protoveratrine
Old Tree	D03.549.256.679 Veratridine
Old Tree	D03.549.256.815 Veratrine
Old Tree	D03.549.380 Dihydro-beta-Erythroidine
Old Tree	D03.549.439 Ergolines
Old Tree	D03.549.439.131 Bromocriptine
Old Tree	D03.549.439.262 Ergonovine
Old Tree	D03.549.439.262.538 Methylergonovine
Old Tree	D03.549.439.560 Lisuride
Old Tree	D03.549.439.572 Lysergic Acid
Old Tree	D03.549.439.572.522 Lysergic Acid Diethylamide
Old Tree	D03.549.439.630 Metergoline
Old Tree	D03.549.439.689 Methysergide
Old Tree	D03.549.439.730 Nicergoline
Old Tree	D03.549.439.800 Pergolide
Old Tree	D03.549.562 Ergotamines

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	D03.549.562.100 Bromocriptine
Old Tree	D03.549.562.150 Dihydroergocornine
Old Tree	D03.549.562.200 Dihydroergocristine
Old Tree	D03.549.562.250 Dihydroergocryptine
Old Tree	D03.549.562.300 Dihydroergotamine
Old Tree	D03.549.562.400 Dihydroergotoxine
Old Tree	D03.549.562.400.500 Ergoloid Mesylates
Old Tree	D03.549.562.500 Ergotamine
Old Tree	D03.549.624 Harringtonines
Old Tree	D03.549.686 Morphinans
Old Tree	D03.549.686.150 Buprenorphine
Old Tree	D03.549.686.150.500 Buprenorphine, Naloxone Drug Combination
Old Tree	D03.549.686.165 Butorphanol
Old Tree	D03.549.686.212 Dextromethorphan
Old Tree	D03.549.686.260 Dextrorphan
Old Tree	D03.549.686.275 Diprenorphine
Old Tree	D03.549.686.320 Etorphine
Old Tree	D03.549.686.429 Levallorphan
Old Tree	D03.549.686.485 Levorphanol
Old Tree	D03.549.686.607 Morphine Derivatives
Old Tree	D03.549.686.607.204 Codeine
Old Tree	D03.549.686.607.204.540 Hydrocodone
Old Tree	D03.549.686.607.204.650 Oxycodone
Old Tree	D03.549.686.607.420 Dihydromorphine
Old Tree	D03.549.686.607.460 Ethylmorphine
Old Tree	D03.549.686.607.490 Heroin
Old Tree	D03.549.686.607.500 Hydromorphone
Old Tree	D03.549.686.607.587 Morphine
Old Tree	D03.549.686.607.675 Oxymorphone
Old Tree	D03.549.686.607.851 Thebaine
Old Tree	D03.549.686.639 Nalbuphine
Old Tree	D03.549.686.694 Nalorphine
Old Tree	D03.549.686.750 Naloxone
Old Tree	D03.549.686.750.275 Buprenorphine, Naloxone Drug Combination
Old Tree	D03.549.686.750.550 Naltrexone
Old Tree	D03.549.748 Pterocarpanes

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	D03.549.811 Rifamycins
Old Tree	D03.549.811.650 Rifabutin
Old Tree	D03.549.811.650.700 Streptovaricin
Old Tree	D03.549.811.700 Rifampin
Old Tree	D03.549.825 Rotenone
Old Tree	D03.549.909 Tetrapyrroles
Old Tree	D03.549.909.249 Bile Pigments
Old Tree	D03.549.909.249.184 Bilirubin
Old Tree	D03.549.909.249.184.200 Biliverdine
Old Tree	D03.549.909.249.727 Urobilin
Old Tree	D03.549.909.249.852 Urobilinogen
Old Tree	D03.549.909.374 Chlorophyll
Old Tree	D03.549.909.374.100 Bacteriochlorophylls
Old Tree	D03.549.909.374.180 Chlorophyllides
Old Tree	D03.549.909.374.700 Pheophytins
Old Tree	D03.549.909.374.725 Protochlorophyllide
Old Tree	D03.549.909.437 Corrinoids
Old Tree	D03.549.909.437.777 Vitamin B 12
Old Tree	D03.549.909.437.777.270 Cobamides
Old Tree	D03.549.909.437.777.560 Hydroxocobalamin
Old Tree	D03.549.909.468 Phycobilins
Old Tree	D03.549.909.500 Porphyrins
Old Tree	D03.549.909.500.250 Coproporphyrins
Old Tree	D03.549.909.500.280 Deuteroporphyrins
Old Tree	D03.549.909.500.340 Etioporphyrins
Old Tree	D03.549.909.500.462 Hematoporphyrins
Old Tree	D03.549.909.500.462.400 Hematoporphyrin Derivative
Old Tree	D03.549.909.500.462.400.200 Dihematoporphyrin Ether
Old Tree	D03.549.909.500.620 Mesoporphyrins
Old Tree	D03.549.909.500.640 Metalloporphyrins
Old Tree	D03.549.909.500.640.220 Chlorophyll
Old Tree	D03.549.909.500.640.220.100 Bacteriochlorophylls
Old Tree	D03.549.909.500.640.220.180 Chlorophyllides
Old Tree	D03.549.909.500.640.220.453 Pheophytins
Old Tree	D03.549.909.500.640.220.725 Protochlorophyllide
Old Tree	D03.549.909.500.640.587 Heme

## MeSH Tree Changes for 2017

Type	Tree - heading	
Old Tree	D03.549.909.500.640.587.462	Hemin
Old Tree	D03.549.909.500.700	Porphyrinogens
Old Tree	D03.549.909.500.700.250	Coproporphyrinogens
Old Tree	D03.549.909.500.700.900	Uroporphyrinogens
Old Tree	D03.549.909.500.725	Protoporphyrins
Old Tree	D03.549.909.500.880	Uroporphyrins
-	D03.605	Heterocyclic Compounds, Bridged-Ring
-	D03.605.084	Bicyclo Compounds, Heterocyclic
-	D03.605.084	Bridged Bicyclo Compounds, Heterocyclic
New Tree	D03.605.084.500	Azabicyclo Compounds
New Tree	D03.605.084.500.332	Biperiden
New Tree	D03.605.084.500.444	Granisetron
New Tree	D03.605.084.500.722	Tropanes
New Tree	D03.605.084.500.722.229	Atropine Derivatives
New Tree	D03.605.084.500.722.229.199	Atropine
New Tree	D03.605.084.500.722.229.199.500	Hyoscyamine
New Tree	D03.605.084.500.722.229.400	Ipratropium
New Tree	D03.605.084.500.722.229.400.500	Albuterol, Ipratropium Drug Combination
New Tree	D03.605.084.500.722.270	Benztropine
New Tree	D03.605.084.500.722.388	Cocaine
New Tree	D03.605.084.500.722.388.250	Crack Cocaine
New Tree	D03.605.084.500.722.744	Nortropanes
New Tree	D03.605.084.500.722.822	Scopolamine Derivatives
New Tree	D03.605.084.500.722.822.200	Butylscopolammonium Bromide
New Tree	D03.605.084.500.722.822.550	N-Methylscopolamine
New	D03.605.084.500.722.822.775	Scopolamine Hydrobromide



## MeSH Tree Changes for 2017

Type	Tree - heading	
Tree		
New Tree	<a href="#">D03.605.084.500.722.822.887</a>	<a href="#">Tiotropium Bromide</a>
Old Tree	<a href="#">D03.605.084.737</a>	<a href="#">Penicillins</a>
Old Tree	<a href="#">D03.605.084.737.124</a>	<a href="#">Amdinocillin</a>
Old Tree	<a href="#">D03.605.084.737.124.036</a>	<a href="#">Amdinocillin Pivoxil</a>
Old Tree	<a href="#">D03.605.084.737.249</a>	<a href="#">Cyclacillin</a>
Old Tree	<a href="#">D03.605.084.737.500</a>	<a href="#">Methicillin</a>
Old Tree	<a href="#">D03.605.084.737.562</a>	<a href="#">Nafcillin</a>
Old Tree	<a href="#">D03.605.084.737.625</a>	<a href="#">Oxacillin</a>
Old Tree	<a href="#">D03.605.084.737.625.150</a>	<a href="#">Cloxacillin</a>
Old Tree	<a href="#">D03.605.084.737.625.150.205</a>	<a href="#">Dicloxacillin</a>
Old Tree	<a href="#">D03.605.084.737.625.150.250</a>	<a href="#">Floxacillin</a>
Old Tree	<a href="#">D03.605.084.737.687</a>	<a href="#">Penicillanic Acid</a>
Old Tree	<a href="#">D03.605.084.737.750</a>	<a href="#">Penicillin G</a>
Old Tree	<a href="#">D03.605.084.737.750.050</a>	<a href="#">Ampicillin</a>
Old Tree	<a href="#">D03.605.084.737.750.050.050</a>	<a href="#">Amoxicillin</a>
Old Tree	<a href="#">D03.605.084.737.750.050.050.060</a> <a href="#">Combination</a>	<a href="#">Amoxicillin-Potassium Clavulanate</a>
Old Tree	<a href="#">D03.605.084.737.750.050.075</a>	<a href="#">Azlocillin</a>
Old Tree	<a href="#">D03.605.084.737.750.050.500</a>	<a href="#">Mezlocillin</a>
Old Tree	<a href="#">D03.605.084.737.750.050.650</a>	<a href="#">Piperacillin</a>
Old Tree	<a href="#">D03.605.084.737.750.050.700</a>	<a href="#">Pivampicillin</a>
Old Tree	<a href="#">D03.605.084.737.750.050.900</a>	<a href="#">Talampicillin</a>
Old Tree	<a href="#">D03.605.084.737.750.170</a>	<a href="#">Carbenicillin</a>
Old Tree	<a href="#">D03.605.084.737.750.170.200</a>	<a href="#">Carfecillin</a>
Old Tree	<a href="#">D03.605.084.737.750.685</a>	<a href="#">Penicillin G Benzathine</a>
Old Tree	<a href="#">D03.605.084.737.750.695</a>	<a href="#">Penicillin G Procaine</a>
Old Tree	<a href="#">D03.605.084.737.750.875</a>	<a href="#">Sulbenicillin</a>
Old Tree	<a href="#">D03.605.084.737.781</a>	<a href="#">Penicillin V</a>
Old Tree	<a href="#">D03.605.084.737.812</a>	<a href="#">Sulbactam</a>
Old Tree	<a href="#">D03.605.084.737.875</a>	<a href="#">Ticarcillin</a>
Old Tree	<a href="#">D03.605.084.750</a>	<a href="#">Ramipril</a>
-	<a href="#">D03.605.168</a>	<a href="#">Cyclazocine</a>
-	<a href="#">D03.605.168.350</a>	<a href="#">Ethylketocyclazocine</a>
-	<a href="#">D03.605.497</a>	<a href="#">Morphinans</a>

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D03.605.497.150 Buprenorphine
-	D03.605.497.150.500 Buprenorphine, Naloxone Drug Combination
-	D03.605.497.165 Butorphanol
-	D03.605.497.212 Dextromethorphan
-	D03.605.497.260 Dextrophan
-	D03.605.497.280 Diprenorphine
-	D03.605.497.320 Etorphine
-	D03.605.497.429 Levallorphan
-	D03.605.497.485 Levorphanol
-	D03.605.497.607 Morphine Derivatives
-	D03.605.497.607.204 Codeine
-	D03.605.497.607.204.540 Hydrocodone
-	D03.605.497.607.204.650 Oxycodone
-	D03.605.497.607.420 Dihydromorphine
-	D03.605.497.607.460 Ethylmorphine
-	D03.605.497.607.490 Heroin
-	D03.605.497.607.500 Hydromorphone
-	D03.605.497.607.587 Morphine
-	D03.605.497.607.675 Oxymorphone
-	D03.605.497.607.851 Thebaine
-	D03.605.497.639 Nalbuphine
-	D03.605.497.694 Nalorphine
-	D03.605.497.750 Naloxone
-	D03.605.497.750.275 Buprenorphine, Naloxone Drug Combination
-	D03.605.497.750.550 Naltrexone
-	D03.605.687 Quinuclidines
-	D03.605.687.637 Quinidine
-	D03.605.687.762 Quinine
-	D03.605.687.800 Quinuclidinyl Benzilate
-	D03.605.687.900 Solifenacin Succinate
-	D03.605.869 Tropanes
-	D03.605.869.229 Atropine Derivatives
-	D03.605.869.229.199 Atropine
-	D03.605.869.229.199.500 Hyoscyamine
-	D03.605.869.229.400 Ipratropium
-	D03.605.869.229.400.500 Albuterol, Ipratropium Drug Combination

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D03.605.869.270	Benztropine
-	D03.605.869.388	Cocaine
-	D03.605.869.388.250	Crack Cocaine
-	D03.605.869.744	Nortropanes
-	D03.605.869.822	Scopolamine Derivatives
-	D03.605.869.822.200	Butylscopolammonium Bromide
-	D03.605.869.822.550	N-Methylscopolamine
-	D03.605.869.822.775	Scopolamine Hydrobromide
-	D03.605.869.822.887	Tiotropium Bromide
New Heading	<b>D03.633</b>	<b>Heterocyclic Compounds, Fused-Ring</b>
New Tree	D03.633.100	Heterocyclic Compounds, 2-Ring
New Tree	D03.633.100.079	Benzazepines
New Tree	D03.633.100.079.080	Benzodiazepines
New Tree	D03.633.100.079.080.030	Alprazolam
New Tree	D03.633.100.079.080.070	Benzodiazepinones
New Tree	D03.633.100.079.080.070.050	Anthramycin
New Tree	D03.633.100.079.080.070.110	Bromazepam
New Tree	D03.633.100.079.080.070.150	Clonazepam
New Tree	D03.633.100.079.080.070.200	Devazepide
New Tree	D03.633.100.079.080.070.216	Diazepam
New Tree	D03.633.100.079.080.070.216.500	Nordazepam
New Tree	D03.633.100.079.080.070.305	Flumazenil
New Tree	D03.633.100.079.080.070.320	Flunitrazepam
New Tree	D03.633.100.079.080.070.348	Flurazepam
New Tree	D03.633.100.079.080.070.450	Lorazepam
New	D03.633.100.079.080.070.565	Nitrazepam

## MeSH Tree Changes for 2017

Type	Tree - heading	
Tree		
New Tree	<a href="#">D03.633.100.079.080.070.663</a>	Oxazepam
New Tree	<a href="#">D03.633.100.079.080.070.750</a>	Pirenzepine
New Tree	<a href="#">D03.633.100.079.080.070.784</a>	Prazepam
New Tree	<a href="#">D03.633.100.079.080.070.880</a>	Temazepam
New Tree	<a href="#">D03.633.100.079.080.150</a>	Chlordiazepoxide
New Tree	<a href="#">D03.633.100.079.080.180</a>	Clorazepate Dipotassium
New Tree	<a href="#">D03.633.100.079.080.250</a>	Estazolam
New Tree	<a href="#">D03.633.100.079.080.550</a>	Medazepam
New Tree	<a href="#">D03.633.100.079.080.575</a>	Midazolam
New Tree	<a href="#">D03.633.100.079.080.900</a>	Triazolam
New Tree	<a href="#">D03.633.100.079.150</a>	Diltiazem
New Tree	<a href="#">D03.633.100.079.300</a>	Fenoldopam
New Tree	<a href="#">D03.633.100.079.525</a>	Galantamine
New Tree	<a href="#">D03.633.100.079.533</a>	Harringtonines
New Tree	<a href="#">D03.633.100.079.800</a> benzazepine	2,3,4,5-Tetrahydro-7,8-dihydroxy-1-phenyl-1H-3-
New Tree	<a href="#">D03.633.100.079.900</a>	Varenicline
New Tree	<a href="#">D03.633.100.103</a>	Benzimidazoles
New Tree	<a href="#">D03.633.100.103.034</a>	2-Pyridinylmethylsulfinylbenzimidazoles
New Tree	<a href="#">D03.633.100.103.034.249</a>	Lansoprazole
New Tree	<a href="#">D03.633.100.103.034.249.500</a>	Dexlansoprazole
New Tree	<a href="#">D03.633.100.103.034.500</a>	Omeprazole

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	D03.633.100.103.034.500.500 <span style="float: right;">Esomeprazole</span>
New Tree	D03.633.100.103.034.750 <span style="float: right;">Rabeprazole</span>
New Tree	D03.633.100.103.070 <span style="float: right;">Albendazole</span>
New Tree	D03.633.100.103.105 <span style="float: right;">Astemizole</span>
New Tree	D03.633.100.103.123 <span style="float: right;">Bendamustine Hydrochloride</span>
New Tree	D03.633.100.103.140 <span style="float: right;">Benomyl</span>
New Tree	D03.633.100.103.145 <span style="float: right;">Bisbenzimidazole</span>
New Tree	D03.633.100.103.190 <span style="float: right;">Cambendazole</span>
New Tree	D03.633.100.103.280 <span style="float: right;">Dabigatran</span>
New Tree	D03.633.100.103.370 <span style="float: right;">Domperidone</span>
New Tree	D03.633.100.103.393 <span style="float: right;">Droperidol</span>
New Tree	D03.633.100.103.450 <span style="float: right;">Fenbendazole</span>
New Tree	D03.633.100.103.600 <span style="float: right;">Mebendazole</span>
New Tree	D03.633.100.103.618 <span style="float: right;">Mibefradil</span>
New Tree	D03.633.100.103.637 <span style="float: right;">Nocodazole</span>
New Tree	D03.633.100.103.732 <span style="float: right;">Pimozide</span>
New Tree	D03.633.100.103.850 <span style="float: right;">Thiabendazole</span>
New Tree	D03.633.100.115 <span style="float: right;">Benzodioxoles</span>
New Tree	D03.633.100.115.600 <span style="float: right;">Piperonyl Butoxide</span>
New Tree	D03.633.100.115.750 <span style="float: right;">Safrole</span>
New Tree	D03.633.100.127 <span style="float: right;">Benzofurans</span>
New Tree	D03.633.100.127.075 <span style="float: right;">Amiodarone</span>

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	D03.633.100.127.110 Benzbromarone
New Tree	D03.633.100.127.125 Cantharidin
New Tree	D03.633.100.127.187 Citalopram
New Tree	D03.633.100.127.218 Fluorescamine
New Tree	D03.633.100.127.250 Fura-2
New Tree	D03.633.100.127.275 Griseofulvin
New Tree	D03.633.100.127.637 Pterocarpans
New Tree	D03.633.100.127.818 Vilazodone Hydrochloride
New Tree	D03.633.100.150 Benzopyrans
New Tree	D03.633.100.150.119 Aflatoxins
New Tree	D03.633.100.150.119.075 Aflatoxin B1
New Tree	D03.633.100.150.119.100 Aflatoxin M1
New Tree	D03.633.100.150.240 Chromans
New Tree	D03.633.100.150.240.190 Catechin
New Tree	D03.633.100.150.240.225 Centchroman
New Tree	D03.633.100.150.266 Chromones
New Tree	D03.633.100.150.266.300 Cromolyn Sodium
New Tree	D03.633.100.150.266.450 Flavonoids
New Tree	D03.633.100.150.266.450.087 Anthocyanins
New Tree	D03.633.100.150.266.450.175 Benzoflavones
New Tree	D03.633.100.150.266.450.175.100 beta-Naphthoflavone
New Tree	D03.633.100.150.266.450.190 Biflavonoids

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	D03.633.100.150.266.450.206 Catechin
New Tree	D03.633.100.150.266.450.221 Chalcones
New Tree	D03.633.100.150.266.450.221.500 Chalcone
New Tree	D03.633.100.150.266.450.252 Flavanones
New Tree	D03.633.100.150.266.450.252.500 Hesperidin
New Tree	D03.633.100.150.266.450.260 Flavones
New Tree	D03.633.100.150.266.450.260.110 Apigenin
New Tree	D03.633.100.150.266.450.260.222 Diosmin
New Tree	D03.633.100.150.266.450.260.444 Flavoxate
New Tree	D03.633.100.150.266.450.260.555 Luteolin
New Tree	D03.633.100.150.266.450.260.777 Polyphenols
New Tree	D03.633.100.150.266.450.268 Flavonolignans
New Tree	D03.633.100.150.266.450.268.777 Silymarin
New Tree	D03.633.100.150.266.450.284 Flavonols
New Tree	D03.633.100.150.266.450.284.388 Kaempferols
New Tree	D03.633.100.150.266.450.284.777 Quercetin
New Tree	D03.633.100.150.266.450.284.888 Rutin
New Tree	D03.633.100.150.266.450.284.888.500 Hydroxyethylrutoside
New Tree	D03.633.100.150.266.450.400 Isoflavones
New Tree	D03.633.100.150.266.450.400.187 Coumestrol
New Tree	D03.633.100.150.266.450.400.281 Equol
New Tree	D03.633.100.150.266.450.400.375 Genistein

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	D03.633.100.150.266.450.400.687 Pterocarpan
New Tree	D03.633.100.150.266.450.400.843 Rotenone
New Tree	D03.633.100.150.266.450.700 Proanthocyanidins
New Tree	D03.633.100.150.320 Citrinin
New Tree	D03.633.100.150.446 Coumarins
New Tree	D03.633.100.150.446.139 Aminocoumarins
New Tree	D03.633.100.150.446.139.500 Novobiocin
New Tree	D03.633.100.150.446.280 Chromonar
New Tree	D03.633.100.150.446.350 Coumestrol
New Tree	D03.633.100.150.446.400 Esculin
New Tree	D03.633.100.150.446.520 4-Hydroxycoumarins
New Tree	D03.633.100.150.446.520.079 Acenocoumarol
New Tree	D03.633.100.150.446.520.203 Dicumarol
New Tree	D03.633.100.150.446.520.451 Ethyl Biscoumacetate
New Tree	D03.633.100.150.446.520.750 Phenprocoumon
New Tree	D03.633.100.150.446.520.914 Warfarin
New Tree	D03.633.100.150.446.598 Isocoumarins
New Tree	D03.633.100.150.446.598.500 Ochratoxins
New Tree	D03.633.100.150.446.794 Furocoumarins
New Tree	D03.633.100.150.446.794.200 Ficusin
New Tree	D03.633.100.150.446.794.300 Khellin
New Tree	D03.633.100.150.446.794.500 Methoxsalen



## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	D03.633.100.150.446.794.875                      Trioxsalen
New Tree	D03.633.100.150.446.853                              Pyranocoumarins
New Tree	D03.633.100.150.446.912                              Umbelliferones
New Tree	D03.633.100.150.446.912.326                      Coumaphos
New Tree	D03.633.100.150.446.912.531                      Hymecromone
New Tree	D03.633.100.150.446.912.850                      Scopoletin
New Tree	D03.633.100.150.455                                  Cromakalim
New Tree	D03.633.100.150.500                                  Ellagic Acid
New Tree	D03.633.100.150.600                                  Hematoxylin
New Tree	D03.633.100.150.755                                  Nebivolol
New Tree	D03.633.100.150.909                                  Vitamin E
New Tree	D03.633.100.150.909.750                              Tocopherols
New Tree	D03.633.100.150.909.750.249                      alpha-Tocopherol
New Tree	D03.633.100.150.909.750.374                      beta-Tocopherol
New Tree	D03.633.100.150.909.750.500                      gamma-Tocopherol
New Tree	D03.633.100.150.909.875                              Tocotrienols
New Tree	D03.633.100.174    Benzothiadiazines
New Tree	D03.633.100.174.138                                  Bendroflumethiazide
New Tree	D03.633.100.174.261                                  Chlorothiazide
New Tree	D03.633.100.174.261.476                              Hydrochlorothiazide
New Tree	D03.633.100.174.261.476.716                      Trichlormethiazide
New Tree	D03.633.100.174.285                                  Cyclopenthiazide

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	D03.633.100.174.300 Diazoxide
New Tree	D03.633.100.174.475 Hydroflumethiazide
New Tree	D03.633.100.174.620 Methyclothiazide
New Tree	D03.633.100.174.781 Polythiazide
New Tree	D03.633.100.185 Benzothiazoles
New Tree	D03.633.100.185.222 Dithiazanine
New Tree	D03.633.100.185.416 Ethoxzolamide
New Tree	D03.633.100.185.611 Riluzole
New Tree	D03.633.100.185.708 Saccharin
New Tree	D03.633.100.197 Benzothiepins
New Tree	D03.633.100.197.408 Endosulfan
New Tree	D03.633.100.209 Benzoxazines
New Tree	D03.633.100.221 Benzoxazoles
New Tree	D03.633.100.221.173 Calcimycin
New Tree	D03.633.100.221.346 Chlorzoxazone
New Tree	D03.633.100.221.370 Cialit
New Tree	D03.633.100.221.950 Zoxazolamine
New Tree	D03.633.100.245 Benzoxepins
New Tree	D03.633.100.300 beta-Lactams
New Tree	D03.633.100.300.124 Carbapenems
New Tree	D03.633.100.300.124.300 Thienamycins
New Tree	D03.633.100.300.124.300.500 Imipenem

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	D03.633.100.300.249 Cephalsporins
New Tree	D03.633.100.300.249.150 Cefamandole
New Tree	D03.633.100.300.249.150.160 Cefoperazone
New Tree	D03.633.100.300.249.160 Cefazolin
New Tree	D03.633.100.300.249.177 Cefonicid
New Tree	D03.633.100.300.249.185 Cefsulodin
New Tree	D03.633.100.300.249.190 Cephacetrile
New Tree	D03.633.100.300.249.190.190 Cefotaxime
New Tree	D03.633.100.300.249.190.190.115 Cefixime
New Tree	D03.633.100.300.249.190.190.125 Cefmenoxime
New Tree	D03.633.100.300.249.190.190.135 Cefotiam
New Tree	D03.633.100.300.249.190.190.145 Ceftizoxime
New Tree	D03.633.100.300.249.190.190.155 Ceftriaxone
New Tree	D03.633.100.300.249.190.190.165 Cefuroxime
New Tree	D03.633.100.300.249.190.210 Cephalothin
New Tree	D03.633.100.300.249.190.230 Cephapirin
New Tree	D03.633.100.300.249.200 Cephalixin
New Tree	D03.633.100.300.249.200.155 Cefaclor
New Tree	D03.633.100.300.249.200.165 Cefadroxil
New Tree	D03.633.100.300.249.200.165.125 Cefatrizine
New Tree	D03.633.100.300.249.200.180 Cephaloglycin
New Tree	D03.633.100.300.249.200.185 Cephradine

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	D03.633.100.300.249.210 Cephaloridine
New Tree	D03.633.100.300.249.210.150 Ceftazidime
New Tree	D03.633.100.300.249.250 Cephamycins
New Tree	D03.633.100.300.249.250.177 Cefmetazole
New Tree	D03.633.100.300.249.250.199 Cefotetan
New Tree	D03.633.100.300.249.250.222 Cefoxitin
New Tree	D03.633.100.300.374 Clavulanic Acids
New Tree	D03.633.100.300.374.160 Clavulanic Acid
New Tree	D03.633.100.300.374.160.060 Amoxicillin-Potassium Clavulanate Combination
New Tree	D03.633.100.300.500 Monobactams
New Tree	D03.633.100.300.500.044 Aztreonam
New Tree	D03.633.100.300.625 Moxalactam
New Tree	D03.633.100.300.750 Penicillins
New Tree	D03.633.100.300.750.124 Amdinocillin
New Tree	D03.633.100.300.750.124.036 Amdinocillin Pivoxil
New Tree	D03.633.100.300.750.249 Cyclacillin
New Tree	D03.633.100.300.750.500 Methicillin
New Tree	D03.633.100.300.750.562 Nafcillin
New Tree	D03.633.100.300.750.625 Oxacillin
New Tree	D03.633.100.300.750.625.150 Cloxacillin
New Tree	D03.633.100.300.750.625.150.205 Dicloxacillin
New Tree	D03.633.100.300.750.625.150.250 Floxacillin

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	D03.633.100.300.750.687 Penicillanic Acid
New Tree	D03.633.100.300.750.750 Penicillin G
New Tree	D03.633.100.300.750.750.050 Ampicillin
New Tree	D03.633.100.300.750.750.050.050 Amoxicillin
New Tree	D03.633.100.300.750.750.050.050.500 Amoxicillin-Potassium Clavulanate Combination
New Tree	D03.633.100.300.750.750.050.075 Azlocillin
New Tree	D03.633.100.300.750.750.050.500 Mezlocillin
New Tree	D03.633.100.300.750.750.050.650 Piperacillin
New Tree	D03.633.100.300.750.750.050.700 Pivampicillin
New Tree	D03.633.100.300.750.750.050.900 Talampicillin
New Tree	D03.633.100.300.750.750.170 Carbenicillin
New Tree	D03.633.100.300.750.750.170.200 Carfecillin
New Tree	D03.633.100.300.750.750.685 Penicillin G Benzathine
New Tree	D03.633.100.300.750.750.695 Penicillin G Procaine
New Tree	D03.633.100.300.750.750.875 Sulbenicillin
New Tree	D03.633.100.300.750.781 Penicillin V
New Tree	D03.633.100.300.750.812 Sulbactam
New Tree	D03.633.100.300.750.875 Ticarcillin
New Tree	D03.633.100.355 Idazoxan
New Tree	D03.633.100.449 Indazoles
New Tree	D03.633.100.449.130 Benzydamine
New Tree	D03.633.100.449.350 Granisetron

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	D03.633.100.473 Indoles
New Tree	D03.633.100.473.025 Adrenochrome
New Tree	D03.633.100.473.050 Alcian Blue
New Tree	D03.633.100.473.144 Carbazoles
New Tree	D03.633.100.473.144.249 Ellipticines
New Tree	D03.633.100.473.144.500 Ondansetron
New Tree	D03.633.100.473.144.750 Staurosporine
New Tree	D03.633.100.473.155 Carbolines
New Tree	D03.633.100.473.155.500 Tadalafil
New Tree	D03.633.100.473.231 Cytochalasins
New Tree	D03.633.100.473.231.370 Cytochalasin B
New Tree	D03.633.100.473.231.450 Cytochalasin D
New Tree	D03.633.100.473.250 Delavirdine
New Tree	D03.633.100.473.360 Gliotoxin
New Tree	D03.633.100.473.385 Hydroxytryptophol
New Tree	D03.633.100.473.390 Ibogaine
New Tree	D03.633.100.473.391 Indapamide
New Tree	D03.633.100.473.393 Indican
New Tree	D03.633.100.473.396 Indigo Carmine
New Tree	D03.633.100.473.400 Indocyanine Green
New Tree	D03.633.100.473.402 Indole Alkaloids
New Tree	D03.633.100.473.402.444 Harmaline

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	D03.633.100.473.402.477 Harmine
New Tree	D03.633.100.473.402.511 Lyngbya Toxins
New Tree	D03.633.100.473.402.545 Physostigmine
New Tree	D03.633.100.473.402.613 Psilocybin
New Tree	D03.633.100.473.402.613 Psilocybine
New Tree	D03.633.100.473.402.681 Secologanin Tryptamine Alkaloids
New Tree	D03.633.100.473.402.681.077 Ajmaline
New Tree	D03.633.100.473.402.681.077.324 Lorajmine
New Tree	D03.633.100.473.402.681.077.650 Prajmaline
New Tree	D03.633.100.473.402.681.333 Ellipticines
New Tree	D03.633.100.473.402.681.444 Ibogaine
New Tree	D03.633.100.473.402.681.722 Strychnine
New Tree	D03.633.100.473.402.681.827 Vinca Alkaloids
New Tree	D03.633.100.473.402.681.827.650 Vinblastine
New Tree	D03.633.100.473.402.681.827.750 Vincamine
New Tree	D03.633.100.473.402.681.827.817 Vincristine
New Tree	D03.633.100.473.402.681.827.830 Vindesine
New Tree	D03.633.100.473.402.681.933 Yohimbine
New Tree	D03.633.100.473.402.681.933.500 Reserpine
New Tree	D03.633.100.473.402.750 Staurosporine
New Tree	D03.633.100.473.404 Indoleacetic Acids
New Tree	D03.633.100.473.404.200 Etodolac

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	D03.633.100.473.404.478                      Hydroxyindoleacetic Acid
New Tree	D03.633.100.473.412                              Indolequinones
New Tree	D03.633.100.473.412.249                      Mitomycins
New Tree	D03.633.100.473.412.249.350                      Mitomycin
New Tree	D03.633.100.473.412.249.700                      Porfiromycin
New Tree	D03.633.100.473.412.500                      Pyrroloiminoquinones
New Tree	D03.633.100.473.420                              Indomethacin
New Tree	D03.633.100.473.420.485                      Ketorolac
New Tree	D03.633.100.473.420.742                      Ketorolac Tromethamine
New Tree	D03.633.100.473.432                              Indoramin
New Tree	D03.633.100.473.469                              Iprindole
New Tree	D03.633.100.473.525                              Isatin
New Tree	D03.633.100.473.615                              Methisazone
New Tree	D03.633.100.473.629                              Molindone
New Tree	D03.633.100.473.725                              Oxyphenisatin Acetate
New Tree	D03.633.100.473.766                              Perindopril
New Tree	D03.633.100.473.808                              Skatole
New Tree	D03.633.100.473.840                              Sporidesmins
New Tree	D03.633.100.473.914                              Tryptamines
New Tree	D03.633.100.473.914.201                      Dihydroxytryptamines
New Tree	D03.633.100.473.914.201.259                      5,6-Dihydroxytryptamine
New Tree	D03.633.100.473.914.201.263                      5,7-Dihydroxytryptamine



## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	D03.633.100.473.914.237 N,N-Dimethyltryptamine
New Tree	D03.633.100.473.914.237.150 Bufotenin
New Tree	D03.633.100.473.914.237.150.500 Methoxydimethyltryptamines
New Tree	D03.633.100.473.914.481 Melatonin
New Tree	D03.633.100.473.914.700 Psilocybin
New Tree	D03.633.100.473.914.700 Psilocybine
New Tree	D03.633.100.473.914.814 Serotonin
New Tree	D03.633.100.473.914.814.150 Bufotenin
New Tree	D03.633.100.473.914.814.150.500 Methoxydimethyltryptamines
New Tree	D03.633.100.473.914.814.400 5-Methoxytryptamine
New Tree	D03.633.100.473.914.907 Sumatriptan
New Tree	D03.633.100.473.957 Vilazodone Hydrochloride
New Tree	D03.633.100.496 Indolizines
New Tree	D03.633.100.496.500 Indolizidines
New Tree	D03.633.100.513 Isoindoles
New Tree	D03.633.100.513.124 Captan
New Tree	D03.633.100.513.249 Chlorisondamine
New Tree	D03.633.100.513.374 Indoprofen
New Tree	D03.633.100.513.437 Lurasidone Hydrochloride
New Tree	D03.633.100.513.500 Mazindol
New Tree	D03.633.100.513.750 Phthalimides
New Tree	D03.633.100.513.750.500 Chlorthalidone

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	D03.633.100.513.750.625 Phosmet
New Tree	D03.633.100.513.750.750 Thalidomide
New Tree	D03.633.100.531 Isoquinolines
New Tree	D03.633.100.531.085 Benzylisoquinolines
New Tree	D03.633.100.531.085.030 Aporphines
New Tree	D03.633.100.531.085.030.290 Apomorphine
New Tree	D03.633.100.531.085.061 Atracurium
New Tree	D03.633.100.531.085.077 Bicuculline
New Tree	D03.633.100.531.085.666 Papaverine
New Tree	D03.633.100.531.085.666.850 Tetrahydropapaveroline
New Tree	D03.633.100.531.085.777 Toxiferine
New Tree	D03.633.100.531.085.777.050 Alcuronium
New Tree	D03.633.100.531.085.888 Tretoquinol
New Tree	D03.633.100.531.085.944 Tubocurarine
New Tree	D03.633.100.531.205 Debrisoquin
New Tree	D03.633.100.531.321 Emetine
New Tree	D03.633.100.531.400 1-(5-Isoquinolinesulfonyl)-2-Methylpiperazine
New Tree	D03.633.100.531.460 Naphthalimides
New Tree	D03.633.100.531.520 Nelfinavir
New Tree	D03.633.100.531.535 Nomifensine
New Tree	D03.633.100.531.567 Noscaphine
New Tree	D03.633.100.531.690 Praziquantel

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	D03.633.100.531.770 Saquinavir
New Tree	D03.633.100.531.820 Tetrahydroisoquinolines
New Tree	D03.633.100.531.820.500 Salsoline Alkaloids
New Tree	D03.633.100.531.820.594 Solifenacin Succinate
New Tree	D03.633.100.531.820.687 Toxiferine
New Tree	D03.633.100.531.820.687.050 Alcuronium
New Tree	D03.633.100.531.820.875 Tubocurarine
New Tree	D03.633.100.612 Naphthyridines
New Tree	D03.633.100.612.500 Nalidixic Acid
New Tree	D03.633.100.673 Piperoxan
New Tree	D03.633.100.733 Pteridines
New Tree	D03.633.100.733.315 Flavins
New Tree	D03.633.100.733.315.650 Riboflavin
New Tree	D03.633.100.733.315.650.249 Flavin-Adenine Dinucleotide
New Tree	D03.633.100.733.315.650.500 Flavin Mononucleotide
New Tree	D03.633.100.733.631 Pterins
New Tree	D03.633.100.733.631.192 Aminopterin
New Tree	D03.633.100.733.631.192.500 Methotrexate
New Tree	D03.633.100.733.631.202 Biopterin
New Tree	D03.633.100.733.631.202.500 Neopterin
New Tree	D03.633.100.733.631.400 Folic Acid
New Tree	D03.633.100.733.631.400.600 Pteroylpolyglutamic Acids

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	D03.633.100.733.631.400.800 <span style="float: right;">Tetrahydrofolates</span>
New Tree	D03.633.100.733.631.400.800.350 <span style="float: right;">Formyltetrahydrofolates</span>
New Tree	D03.633.100.733.631.400.800.350.450 <span style="float: right;">Leucovorin</span>
New Tree	D03.633.100.733.631.400.800.350.450.500 <span style="float: right;">Levoleucovorin</span>
New Tree	D03.633.100.733.631.825 <span style="float: right;">Xanthopterin</span>
New Tree	D03.633.100.733.900 <span style="float: right;">Triamterene</span>
New Tree	D03.633.100.759 <span style="float: right;">Purines</span>
New Tree	D03.633.100.759.138 <span style="float: right;">Adenine</span>
New Tree	D03.633.100.759.138.050 <span style="float: right;">2-Aminopurine</span>
New Tree	D03.633.100.759.138.525 <span style="float: right;">Cytokinins</span>
New Tree	D03.633.100.759.138.525.350 <span style="float: right;">Isopentenyladenosine</span>
New Tree	D03.633.100.759.138.525.400 <span style="float: right;">Kinetin</span>
New Tree	D03.633.100.759.138.525.700 <span style="float: right;">Zeatin</span>
New Tree	D03.633.100.759.138.881 <span style="float: right;">Tenofovir</span>
New Tree	D03.633.100.759.138.881.125 <span style="float: right;">Efavirenz, Emtricitabine, Tenofovir Disoproxil Fumarate Drug Combination</span>
New Tree	D03.633.100.759.138.881.250 <span style="float: right;">Elvitegravir, Cobicistat, Emtricitabine, Tenofovir Disoproxil Fumarate Drug Combination</span>
New Tree	D03.633.100.759.138.881.500 <span style="float: right;">Emtricitabine, Rilpivirine, Tenofovir Drug Combination</span>
New Tree	D03.633.100.759.138.881.750 <span style="float: right;">Emtricitabine, Tenofovir Disoproxil Fumarate Drug Combination</span>
New Tree	D03.633.100.759.160 <span style="float: right;">Allopurinol</span>
New Tree	D03.633.100.759.534 <span style="float: right;">6-Mercaptopurine</span>
New Tree	D03.633.100.759.534.090 <span style="float: right;">Azathioprine</span>
New Tree	D03.633.100.759.562 <span style="float: right;">Linagliptin</span>

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	D03.633.100.759.590 <span style="float: right;">Purine Nucleosides</span>
New Tree	D03.633.100.759.590.138 <span style="float: right;">Adenosine</span>
New Tree	D03.633.100.759.590.138.025 <span style="float: right;">Adenosine-5'-(N-ethylcarboxamide)</span>
New Tree	D03.633.100.759.590.138.240 <span style="float: right;">S-Adenosylhomocysteine</span>
New Tree	D03.633.100.759.590.138.264 <span style="float: right;">S-Adenosylmethionine</span>
New Tree	D03.633.100.759.590.138.300 <span style="float: right;">2-Chloroadenosine</span>
New Tree	D03.633.100.759.590.138.300.200 <span style="float: right;">Cladribine</span>
New Tree	D03.633.100.759.590.138.325 <span style="float: right;">Deoxyadenosines</span>
New Tree	D03.633.100.759.590.138.325.075 <span style="float: right;">Cladribine</span>
New Tree	D03.633.100.759.590.138.325.105 <span style="float: right;">Dideoxyadenosine</span>
New Tree	D03.633.100.759.590.138.325.800 <span style="float: right;">Puromycin Aminonucleoside</span>
New Tree	D03.633.100.759.590.138.500 <span style="float: right;">Isopentenyladenosine</span>
New Tree	D03.633.100.759.590.138.630 <span style="float: right;">Phenylisopropyladenosine</span>
New Tree	D03.633.100.759.590.138.711 <span style="float: right;">Puromycin</span>
New Tree	D03.633.100.759.590.138.711.650 <span style="float: right;">Puromycin Aminonucleoside</span>
New Tree	D03.633.100.759.590.138.900 <span style="float: right;">Vidarabine</span>
New Tree	D03.633.100.759.590.454 <span style="float: right;">Guanosine</span>
New Tree	D03.633.100.759.590.454.240 <span style="float: right;">Deoxyguanosine</span>
New Tree	D03.633.100.759.590.454.500 <span style="float: right;">Nucleoside Q</span>
New Tree	D03.633.100.759.590.616 <span style="float: right;">Inosine</span>
New Tree	D03.633.100.759.590.616.130 <span style="float: right;">Didanosine</span>
New Tree	D03.633.100.759.590.616.450 <span style="float: right;">Inosine Pranobex</span>

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	D03.633.100.759.590.616.900 Thioinosine
New Tree	D03.633.100.759.590.616.900.500 Methylthioinosine
New Tree	D03.633.100.759.590.910 Tubercidin
New Tree	D03.633.100.759.646 Purine Nucleotides
New Tree	D03.633.100.759.646.138 Adenine Nucleotides
New Tree	D03.633.100.759.646.138.124 Adenosine Diphosphate
New Tree	D03.633.100.759.646.138.124.070 Adenosine Diphosphate Sugars
New Tree	D03.633.100.759.646.138.124.070.075 Adenosine Diphosphate Glucose
New Tree	D03.633.100.759.646.138.124.070.125 Adenosine Diphosphate Ribose
New Tree	D03.633.100.759.646.138.124.070.125.040 O-Acetyl-ADP-Ribose
New Tree	D03.633.100.759.646.138.124.070.125.195 Cyclic ADP-Ribose
New Tree	D03.633.100.759.646.138.180 Adenosine Monophosphate
New Tree	D03.633.100.759.646.138.180.080 Adenosine Phosphosulfate
New Tree	D03.633.100.759.646.138.236 Adenosine Triphosphate
New Tree	D03.633.100.759.646.138.236.050 Adenylyl Imidodiphosphate
New Tree	D03.633.100.759.646.138.236.250 Ethenoadenosine Triphosphate
New Tree	D03.633.100.759.646.138.382 Coenzyme A
New Tree	D03.633.100.759.646.138.382.300 Acyl Coenzyme A
New Tree	D03.633.100.759.646.138.382.300.020 Acetyl Coenzyme A
New Tree	D03.633.100.759.646.138.382.300.500 Malonyl Coenzyme A
New Tree	D03.633.100.759.646.138.382.300.700 Palmitoyl Coenzyme A
New Tree	D03.633.100.759.646.138.395 Cyclic AMP

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	D03.633.100.759.646.138.395.225 <span style="float: right;">8-Bromo Cyclic Adenosine Monophosphate</span>
New Tree	D03.633.100.759.646.138.395.250 <span style="float: right;">Bucladesine</span>
New Tree	D03.633.100.759.646.138.410 <span style="float: right;">Deoxyadenine Nucleotides</span>
New Tree	D03.633.100.759.646.138.506 <span style="float: right;">Flavin-Adenine Dinucleotide</span>
New Tree	D03.633.100.759.646.138.694 <span style="float: right;">NAD</span>
New Tree	D03.633.100.759.646.138.749 <span style="float: right;">NADP</span>
New Tree	D03.633.100.759.646.138.850 <span style="float: right;">Phosphoadenosine Phosphosulfate</span>
New Tree	D03.633.100.759.646.138.925 <span style="float: right;">Vidarabine Phosphate</span>
New Tree	D03.633.100.759.646.454 <span style="float: right;">Guanine Nucleotides</span>
New Tree	D03.633.100.759.646.454.160 <span style="float: right;">Cyclic GMP</span>
New Tree	D03.633.100.759.646.454.160.325 <span style="float: right;">Dibutryl Cyclic GMP</span>
New Tree	D03.633.100.759.646.454.200 <span style="float: right;">Deoxyguanine Nucleotides</span>
New Tree	D03.633.100.759.646.454.340 <span style="float: right;">Guanosine Diphosphate</span>
New Tree	D03.633.100.759.646.454.340.350 <span style="float: right;">Guanosine Diphosphate Sugars</span>
New Tree	D03.633.100.759.646.454.340.350.400 <span style="float: right;">Guanosine Diphosphate Fucose</span>
New Tree	D03.633.100.759.646.454.340.350.500 <span style="float: right;">Guanosine Diphosphate Mannose</span>
New Tree	D03.633.100.759.646.454.440 <span style="float: right;">Guanosine Pentaphosphate</span>
New Tree	D03.633.100.759.646.454.480 <span style="float: right;">Guanosine Tetraphosphate</span>
New Tree	D03.633.100.759.646.454.504 <span style="float: right;">Guanosine Triphosphate</span>
New Tree	D03.633.100.759.646.454.504.400 <span style="float: right;">Guanylyl Imidodiphosphate</span>
New Tree	D03.633.100.759.646.454.525 <span style="float: right;">Guanosine Monophosphate</span>
New Tree	D03.633.100.759.646.454.700 <span style="float: right;">RNA Caps</span>

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	D03.633.100.759.646.454.700.710 RNA Cap Analogs
New Tree	D03.633.100.759.646.616 Inosine Nucleotides
New Tree	D03.633.100.759.646.616.300 Cyclic IMP
New Tree	D03.633.100.759.646.616.400 Inosine Diphosphate
New Tree	D03.633.100.759.646.616.500 Inosine Monophosphate
New Tree	D03.633.100.759.646.616.800 Inosine Triphosphate
New Tree	D03.633.100.759.758 Purinones
New Tree	D03.633.100.759.758.399 Hypoxanthines
New Tree	D03.633.100.759.758.399.454 Guanine
New Tree	D03.633.100.759.758.399.454.250 Acyclovir
New Tree	D03.633.100.759.758.399.454.250.300 Ganciclovir
New Tree	D03.633.100.759.758.399.454.300 Azaguanine
New Tree	D03.633.100.759.758.399.454.650 Pemetrexed
New Tree	D03.633.100.759.758.399.475 Hypoxanthine
New Tree	D03.633.100.759.758.824 Xanthines
New Tree	D03.633.100.759.758.824.175 Caffeine
New Tree	D03.633.100.759.758.824.651 Theobromine
New Tree	D03.633.100.759.758.824.651.700 Pentoxifylline
New Tree	D03.633.100.759.758.824.751 Theophylline
New Tree	D03.633.100.759.758.824.751.075 Aminophylline
New Tree	D03.633.100.759.758.824.751.162 Dimenhydrinate
New Tree	D03.633.100.759.758.824.751.250 Dyphylline



## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	D03.633.100.759.758.824.751.500 1-Methyl-3-isobutylxanthine
New Tree	D03.633.100.759.758.824.751.950 Xanthinol Niacinate
New Tree	D03.633.100.759.758.824.877 Uric Acid
New Tree	D03.633.100.759.758.824.938 Xanthine
New Tree	D03.633.100.759.794 Saxitoxin
New Tree	D03.633.100.759.824 Sildenafil Citrate
New Tree	D03.633.100.759.854 Thioguanine
New Tree	D03.633.100.772 Pyrrolizidine Alkaloids
New Tree	D03.633.100.772.500 Monocrotaline
New Tree	D03.633.100.786 Quinazolines
New Tree	D03.633.100.786.375 Erlotinib Hydrochloride
New Tree	D03.633.100.786.563 Linagliptin
New Tree	D03.633.100.786.750 Prazosin
New Tree	D03.633.100.786.750.200 Doxazosin
New Tree	D03.633.100.786.830 Quinazolinones
New Tree	D03.633.100.786.830.333 Ketanserin
New Tree	D03.633.100.786.830.666 Methaqualone
New Tree	D03.633.100.786.830.688 Metolazone
New Tree	D03.633.100.786.910 Tetrodotoxin
New Tree	D03.633.100.786.925 Trimetrexate
New Tree	D03.633.100.810 Quinolines
New Tree	D03.633.100.810.050 Aminoquinolines

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	D03.633.100.810.050.060                      Amodiaquine
New Tree	D03.633.100.810.050.180                      Chloroquine
New Tree	D03.633.100.810.050.180.350                      Hydroxychloroquine
New Tree	D03.633.100.810.050.440                      4-Hydroxyaminoquinoline-1-oxide
New Tree	D03.633.100.810.050.650                      Primaquine
New Tree	D03.633.100.810.087                      Diarylquinolines
New Tree	D03.633.100.810.125                      Dibucaine
New Tree	D03.633.100.810.200                      Ethoxyquin
New Tree	D03.633.100.810.350                      Hydroxyquinolines
New Tree	D03.633.100.810.350.250                      Decoquinatate
New Tree	D03.633.100.810.350.400                      Kynurenic Acid
New Tree	D03.633.100.810.350.600                      Oxamniquine
New Tree	D03.633.100.810.350.625                      Oxyquinoline
New Tree	D03.633.100.810.350.625.250                      Chloroquinolinols
New Tree	D03.633.100.810.350.625.250.260                      Chlorquinaldol
New Tree	D03.633.100.810.350.625.400                      Clioquinol
New Tree	D03.633.100.810.350.625.420                      Iodoquinol
New Tree	D03.633.100.810.350.700                      Procaterol
New Tree	D03.633.100.810.410                      Mefloquine
New Tree	D03.633.100.810.470                      Nitroquinolines
New Tree	D03.633.100.810.470.450                      4-Nitroquinoline-1-oxide
New Tree	D03.633.100.810.470.600                      Oxamniquine

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	D03.633.100.810.553	Pyrroloiminoquinones
New Tree	D03.633.100.810.637	Quinaldines
New Tree	D03.633.100.810.699	Quinidine
New Tree	D03.633.100.810.762	Quinine
New Tree	D03.633.100.810.824	Quinolinium Compounds
New Tree	D03.633.100.810.824.200	Dequalinium
New Tree	D03.633.100.810.824.700	Pyrvinium Compounds
New Tree	D03.633.100.810.835	Quinolones
New Tree	D03.633.100.810.835.055	4-Quinolones
New Tree	D03.633.100.810.835.055.500	Nalidixic Acid
New Tree	D03.633.100.810.835.055.550	Nedocromil
New Tree	D03.633.100.810.835.055.580	Oxolinic Acid
New Tree	D03.633.100.810.835.122	Aripiprazole
New Tree	D03.633.100.810.835.188	Carteolol
New Tree	D03.633.100.810.835.322	Fluoroquinolones
New Tree	D03.633.100.810.835.322.186	Ciprofloxacin
New Tree	D03.633.100.810.835.322.186.400	Fleroxacin
New Tree	D03.633.100.810.835.322.280	Enoxacin
New Tree	D03.633.100.810.835.322.374	Norfloxacin
New Tree	D03.633.100.810.835.322.500	Ofloxacin
New Tree	D03.633.100.810.835.322.500.500	Levofloxacin
New Tree	D03.633.100.810.835.322.750	Pefloxacin

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	D03.633.100.810.835.661 PQQ Cofactor
New Tree	D03.633.100.810.842 Quinpirole
New Tree	D03.633.100.810.850 Quipazine
New Tree	D03.633.100.810.900 Saquinavir
New Tree	D03.633.100.834 Quinolizines
New Tree	D03.633.100.834.700 2H-Benzo(a)quinolizin-2-ol, 2-Ethyl-1,3,4,6,7,11b-hexahydro-3-isobutyl-9,10-dimethoxy-
New Tree	D03.633.100.834.737 Quinolizidines
New Tree	D03.633.100.834.737.500 Sparteine
New Tree	D03.633.100.834.850 Tetrabenazine
New Tree	D03.633.100.857 Quinoxalines
New Tree	D03.633.100.857.070 Brimonidine Tartrate
New Tree	D03.633.100.857.070.500 Brimonidine Tartrate, Timolol Maleate Drug Combination
New Tree	D03.633.100.857.140 Carbadox
New Tree	D03.633.100.857.160 6-Cyano-7-nitroquinoxaline-2,3-dione
New Tree	D03.633.100.857.233 Echinomycin
New Tree	D03.633.100.857.885 Tyrphostins
New Tree	D03.633.100.857.942 Varenicline
New Tree	D03.633.100.893 Ramipril
New Tree	D03.633.100.928 Thienopyridines
New Tree	D03.633.100.928.500 Ticlopidine
New Tree	D03.633.100.964 Zolazepam
New Tree	D03.633.300 Heterocyclic Compounds, 3-Ring

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	D03.633.300.046 <span style="float: right;">Acridines</span>
New Tree	D03.633.300.046.109 <span style="float: right;">Acridones</span>
New Tree	D03.633.300.046.109.077 <span style="float: right;">Acronine</span>
New Tree	D03.633.300.046.250 <span style="float: right;">Aminoacridines</span>
New Tree	D03.633.300.046.250.150 <span style="float: right;">Acridine Orange</span>
New Tree	D03.633.300.046.250.177 <span style="float: right;">Acriflavine</span>
New Tree	D03.633.300.046.250.200 <span style="float: right;">Aminacrine</span>
New Tree	D03.633.300.046.250.225 <span style="float: right;">Amsacrine</span>
New Tree	D03.633.300.046.250.450 <span style="float: right;">Ethacridine</span>
New Tree	D03.633.300.046.250.650 <span style="float: right;">Nitracrine</span>
New Tree	D03.633.300.046.250.720 <span style="float: right;">Proflavine</span>
New Tree	D03.633.300.046.250.760 <span style="float: right;">Quinacrine</span>
New Tree	D03.633.300.046.250.760.750 <span style="float: right;">Quinacrine Mustard</span>
New Tree	D03.633.300.046.250.900 <span style="float: right;">Tacrine</span>
New Tree	D03.633.300.060 <span style="float: right;">Anthramycin</span>
New Tree	D03.633.300.148 <span style="float: right;">Carbazoles</span>
New Tree	D03.633.300.148.249 <span style="float: right;">Ellipticines</span>
New Tree	D03.633.300.148.500 <span style="float: right;">Ondansetron</span>
New Tree	D03.633.300.148.750 <span style="float: right;">Staurosporine</span>
New Tree	D03.633.300.154 <span style="float: right;">Carbolines</span>
New Tree	D03.633.300.154.333 <span style="float: right;">Harmaline</span>
New Tree	D03.633.300.154.344 <span style="float: right;">Harmine</span>

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	D03.633.300.154.672 Tadalafil
New Tree	D03.633.300.160 Cinoxacin
New Tree	D03.633.300.200 Dactinomycin
New Tree	D03.633.300.240 Dibenzazepines
New Tree	D03.633.300.240.127 Carbamazepine
New Tree	D03.633.300.240.194 Clomipramine
New Tree	D03.633.300.240.220 Clozapine
New Tree	D03.633.300.240.281 Desipramine
New Tree	D03.633.300.240.485 Imipramine
New Tree	D03.633.300.240.520 Lofepramine
New Tree	D03.633.300.240.550 Mianserin
New Tree	D03.633.300.240.625 Opipramol
New Tree	D03.633.300.240.918 Trimipramine
New Heading	<b>D03.633.300.258 Dibenzofurans</b>
New Heading	<b>D03.633.300.258.500 Dibenzofurans, Polychlorinated</b>
New Tree	D03.633.300.276 Dibenzothiazepines
New Tree	D03.633.300.276.500 Quetiapine Fumarate
New Tree	D03.633.300.311 Dibenzothiepins
New Tree	D03.633.300.311.250 Dothiepin
New Tree	D03.633.300.311.520 Methiothepin
New Tree	D03.633.300.347 Dibenzoxazepines
New Tree	D03.633.300.347.200 chloro-, 2-acetylhydrazide Dibenz(b,f)(1,4)oxazepine-10(11H)-carboxylic acid, 8-

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	D03.633.300.347.500 Loxapine
New Tree	D03.633.300.347.500.040 Amoxapine
New Tree	D03.633.300.382 Dibenzoxepins
New Tree	D03.633.300.382.393 Doxepin
New Tree	D03.633.300.382.696 Olopatadine Hydrochloride
New Tree	D03.633.300.507 Flavins
New Tree	D03.633.300.507.650 Riboflavin
New Tree	D03.633.300.507.650.249 Flavin-Adenine Dinucleotide
New Tree	D03.633.300.507.650.500 Flavin Mononucleotide
New Tree	D03.633.300.633 Phenanthridines
New Tree	D03.633.300.633.207 Benzophenanthridines
New Tree	D03.633.300.633.416 Ethidium
New Tree	D03.633.300.633.700 Propidium
New Tree	D03.633.300.669 Phenanthrolines
New Tree	D03.633.300.704 Phenazines
New Tree	D03.633.300.704.353 Clofazimine
New Tree	D03.633.300.704.550 Methylphenazonium Methosulfate
New Tree	D03.633.300.704.600 Neutral Red
New Tree	D03.633.300.704.700 Pyocyanine
New Tree	D03.633.300.741 Phenothiazines
New Tree	D03.633.300.741.029 Acepromazine
New Tree	D03.633.300.741.080 Azure Stains

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	<a href="#">D03.633.300.741.198</a>	Chlorpromazine
New Tree	<a href="#">D03.633.300.741.326</a>	Fluphenazine
New Tree	<a href="#">D03.633.300.741.470</a>	Mesoridazine
New Tree	<a href="#">D03.633.300.741.513</a>	Methotrimeprazine
New Tree	<a href="#">D03.633.300.741.517</a>	Methylene Blue
New Tree	<a href="#">D03.633.300.741.533</a>	Moricizine
New Tree	<a href="#">D03.633.300.741.550</a>	Nonachlazine
New Tree	<a href="#">D03.633.300.741.575</a>	Perazine
New Tree	<a href="#">D03.633.300.741.593</a>	Perphenazine
New Tree	<a href="#">D03.633.300.741.639</a>	Prochlorperazine
New Tree	<a href="#">D03.633.300.741.661</a>	Promazine
New Tree	<a href="#">D03.633.300.741.670</a>	Promethazine
New Tree	<a href="#">D03.633.300.741.780</a>	Thiethylperazine
New Tree	<a href="#">D03.633.300.741.843</a>	Thioridazine
New Tree	<a href="#">D03.633.300.741.869</a>	Tolonium Chloride
New Tree	<a href="#">D03.633.300.741.898</a>	Trifluoperazine
New Tree	<a href="#">D03.633.300.741.918</a>	Triflupromazine
New Tree	<a href="#">D03.633.300.741.939</a>	Trimeprazine
New Tree	<a href="#">D03.633.300.770</a>	Furocoumarins
New Tree	<a href="#">D03.633.300.770.200</a>	Ficusin
New Tree	<a href="#">D03.633.300.770.300</a>	Khellin
New Tree	<a href="#">D03.633.300.770.500</a>	Methoxsalen



## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	D03.633.300.770.875 Trioxsalen
New Heading	<b>D03.633.300.786</b> <b>Polychlorinated Dibenzodioxins</b>
New Tree	D03.633.300.802 Quinpirole
New Tree	D03.633.300.819 Simeprevir
New Tree	D03.633.300.835 Spectinomycin
New Tree	D03.633.300.953 Xanthenes
New Tree	D03.633.300.953.275 Fluoresceins
New Tree	D03.633.300.953.275.300 Eosine I Bluish
New Tree	D03.633.300.953.275.325 Eosine Yellowish-(YS)
New Tree	D03.633.300.953.275.350 Erythrosine
New Tree	D03.633.300.953.275.390 Fluorescein
New Tree	D03.633.300.953.275.400 Fluorescein-5-isothiocyanate
New Tree	D03.633.300.953.275.700 Rose Bengal
New Tree	D03.633.300.953.558 Propantheline
New Tree	D03.633.300.953.570 Pyronine
New Tree	D03.633.300.953.600 Rhodamines
New Tree	D03.633.300.953.600.500 Rhodamine 123
New Tree	D03.633.300.953.704 Thioxanthenes
New Tree	D03.633.300.953.704.250 Chlorprothixene
New Tree	D03.633.300.953.704.269 Clopenthixol
New Tree	D03.633.300.953.704.360 Flupenthixol
New Tree	D03.633.300.953.704.450 Hycanthon

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	D03.633.300.953.704.500 <span style="float: right;">Lucanthone</span>
New Tree	D03.633.300.953.704.787 <span style="float: right;">Thiothixene</span>
New Tree	D03.633.300.953.852 <span style="float: right;">Xanthones</span>
New Tree	D03.633.300.953.852.388 <span style="float: right;">Lucanthone</span>
New Tree	D03.633.300.953.852.777 <span style="float: right;">Sterigmatocystin</span>
New Tree	D03.633.400 <span style="float: right;">Heterocyclic Compounds with 4 or More Rings</span>
New Tree	D03.633.400 <span style="float: right;">Heterocyclic Compounds, 4 or More Rings</span>
New Tree	D03.633.400.095 <span style="float: right;">Aporphines</span>
New Tree	D03.633.400.095.290 <span style="float: right;">Apomorphine</span>
New Tree	D03.633.400.131 <span style="float: right;">Benzophenanthridines</span>
New Tree	D03.633.400.168 <span style="float: right;">Berberine Alkaloids</span>
New Tree	D03.633.400.168.100 <span style="float: right;">Berberine</span>
New Tree	D03.633.400.256 <span style="float: right;">Cevanes</span>
New Tree	D03.633.400.256.310 <span style="float: right;">Germine Acetates</span>
New Tree	D03.633.400.256.543 <span style="float: right;">Protoveratrines</span>
New Tree	D03.633.400.256.679 <span style="float: right;">Veratridine</span>
New Tree	D03.633.400.256.815 <span style="float: right;">Veratrine</span>
New Tree	D03.633.400.380 <span style="float: right;">Dihydro-beta-Erythroidine</span>
New Tree	D03.633.400.439 <span style="float: right;">Ergolines</span>
New Tree	D03.633.400.439.131 <span style="float: right;">Bromocriptine</span>
New Tree	D03.633.400.439.262 <span style="float: right;">Ergonovine</span>
New Tree	D03.633.400.439.262.538 <span style="float: right;">Methylergonovine</span>

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	D03.633.400.439.560 Lisuride
New Tree	D03.633.400.439.572 Lysergic Acid
New Tree	D03.633.400.439.572.522 Lysergic Acid Diethylamide
New Tree	D03.633.400.439.630 Metergoline
New Tree	D03.633.400.439.689 Methysergide
New Tree	D03.633.400.439.730 Nicergoline
New Tree	D03.633.400.439.800 Pergolide
New Tree	D03.633.400.562 Ergotamines
New Tree	D03.633.400.562.100 Bromocriptine
New Tree	D03.633.400.562.150 Dihydroergocornine
New Tree	D03.633.400.562.200 Dihydroergocristine
New Tree	D03.633.400.562.250 Dihydroergocryptine
New Tree	D03.633.400.562.300 Dihydroergotamine
New Tree	D03.633.400.562.400 Dihydroergotoxine
New Tree	D03.633.400.562.400.500 Ergoloid Mesylates
New Tree	D03.633.400.562.500 Ergotamine
New Tree	D03.633.400.624 Harringtonines
New Tree	D03.633.400.686 Morphinans
New Tree	D03.633.400.686.150 Buprenorphine
New Tree	D03.633.400.686.150.500 Buprenorphine, Naloxone Drug Combination
New Tree	D03.633.400.686.165 Butorphanol
New Tree	D03.633.400.686.212 Dextromethorphan

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	D03.633.400.686.260                      Dextrophan
New Tree	D03.633.400.686.275                      Diprenorphine
New Tree	D03.633.400.686.320                      Etorphine
New Tree	D03.633.400.686.429                      Levallorphan
New Tree	D03.633.400.686.485                      Levorphanol
New Tree	D03.633.400.686.607                      Morphine Derivatives
New Tree	D03.633.400.686.607.204                      Codeine
New Tree	D03.633.400.686.607.204.540                      Hydrocodone
New Tree	D03.633.400.686.607.204.650                      Oxycodone
New Tree	D03.633.400.686.607.420                      Dihydromorphine
New Tree	D03.633.400.686.607.460                      Ethylmorphine
New Tree	D03.633.400.686.607.490                      Heroin
New Tree	D03.633.400.686.607.500                      Hydromorphone
New Tree	D03.633.400.686.607.587                      Morphine
New Tree	D03.633.400.686.607.675                      Oxymorphone
New Tree	D03.633.400.686.607.851                      Thebaine
New Tree	D03.633.400.686.639                      Nalbuphine
New Tree	D03.633.400.686.694                      Nalorphine
New Tree	D03.633.400.686.750                      Naloxone
New Tree	D03.633.400.686.750.275                      Buprenorphine, Naloxone Drug Combination
New Tree	D03.633.400.686.750.550                      Naltrexone
New Tree	D03.633.400.748                      Pterocarpans

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	D03.633.400.811 Rifamycins
New Tree	D03.633.400.811.650 Rifabutin
New Tree	D03.633.400.811.650.700 Streptovaricin
New Tree	D03.633.400.811.700 Rifampin
New Tree	D03.633.400.825 Rotenone
New Tree	D03.633.400.909 Tetrapyrroles
New Tree	D03.633.400.909.249 Bile Pigments
New Tree	D03.633.400.909.249.184 Bilirubin
New Tree	D03.633.400.909.249.184.200 Biliverdine
New Tree	D03.633.400.909.249.727 Urobilin
New Tree	D03.633.400.909.249.852 Urobilinogen
New Tree	D03.633.400.909.374 Chlorophyll
New Tree	D03.633.400.909.374.100 Bacteriochlorophylls
New Tree	D03.633.400.909.374.180 Chlorophyllides
New Tree	D03.633.400.909.374.700 Pheophytins
New Tree	D03.633.400.909.374.725 Protochlorophyllide
New Tree	D03.633.400.909.437 Corrinoids
New Tree	D03.633.400.909.437.777 Vitamin B 12
New Tree	D03.633.400.909.437.777.270 Cobamides
New Tree	D03.633.400.909.437.777.560 Hydroxocobalamin
New Tree	D03.633.400.909.468 Phycobilins
New Tree	D03.633.400.909.500 Porphyrins

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	D03.633.400.909.500.250	Coproporphyrins
New Tree	D03.633.400.909.500.280	Deuteroporphyrins
New Tree	D03.633.400.909.500.340	Etioporphyrins
New Tree	D03.633.400.909.500.462	Hematoporphyrins
New Tree	D03.633.400.909.500.462.400	Hematoporphyrin Derivative
New Tree	D03.633.400.909.500.462.400.200	Dihematoporphyrin Ether
New Tree	D03.633.400.909.500.620	Mesoporphyrins
New Tree	D03.633.400.909.500.640	Metalloporphyrins
New Tree	D03.633.400.909.500.640.220	Chlorophyll
New Tree	D03.633.400.909.500.640.220.100	Bacteriochlorophylls
New Tree	D03.633.400.909.500.640.220.180	Chlorophyllides
New Tree	D03.633.400.909.500.640.220.453	Pheophytins
New Tree	D03.633.400.909.500.640.220.725	Protochlorophyllide
New Tree	D03.633.400.909.500.640.587	Heme
New Tree	D03.633.400.909.500.640.587.462	Hemin
New Tree	D03.633.400.909.500.700	Porphyrinogens
New Tree	D03.633.400.909.500.700.250	Coproporphyrinogens
New Tree	D03.633.400.909.500.700.900	Uroporphyrinogens
New Tree	D03.633.400.909.500.725	Protoporphyrins
New Tree	D03.633.400.909.500.880	Uroporphyrins
-	D03.661	Heterocyclic Oxides
-	D03.661.243	Cyclic N-Oxides
-	D03.661.243.320	4-Hydroxyaminoquinoline-1-oxide

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D03.661.243.500	4-Nitroquinoline-1-oxide
-	D03.661.243.930	Triacetoneamine-N-Oxyl
-	D03.661.368	Cyclic P-Oxides
-	D03.661.493	Cyclic S-Oxides
Old Tree	<b>D03.787</b>	<b>Phytochemicals</b>
Old Tree	<b>D03.915</b>	<b>Spiro Compounds</b>
Old Tree	<b>D03.915.500</b>	<b>Prospidium</b>
-	D04	Polycyclic Compounds
-	D04.075	Bridged Compounds
-	D04.075	Bridged-Ring Compounds
-	D04.075.080	Bicyclo Compounds
-	D04.075.080	Bridged Bicyclo Compounds
Old Tree	<b>D04.075.080.875</b>	<b>Bicyclo Compounds, Heterocyclic</b>
Old Tree	<b>D04.075.080.875</b>	<b>Bridged Bicyclo Compounds, Heterocyclic</b>
Old Tree	<b>D04.075.080.875.099</b>	<b>Azabicyclo Compounds</b>
Old Tree	<b>D04.075.080.875.099.221</b>	<b>beta-Lactams</b>
Old Tree	<b>D04.075.080.875.099.221.124</b>	<b>Carbapenems</b>
Old Tree	<b>D04.075.080.875.099.221.124.300</b>	<b>Thienamycins</b>
Old Tree	<b>D04.075.080.875.099.221.124.300.500</b>	<b>Imipenem</b>
Old Tree	<b>D04.075.080.875.099.221.249</b>	<b>Cephalosporins</b>
Old Tree	<b>D04.075.080.875.099.221.249.150</b>	<b>Cefamandole</b>
Old Tree	<b>D04.075.080.875.099.221.249.150.160</b>	<b>Cefoperazone</b>
Old Tree	<b>D04.075.080.875.099.221.249.160</b>	<b>Cefazolin</b>
Old Tree	<b>D04.075.080.875.099.221.249.177</b>	<b>Cefonicid</b>
Old Tree	<b>D04.075.080.875.099.221.249.185</b>	<b>Cefsulodin</b>
Old Tree	<b>D04.075.080.875.099.221.249.190</b>	<b>Cephacetrile</b>
Old Tree	<b>D04.075.080.875.099.221.249.190.190</b>	<b>Cefotaxime</b>
Old Tree	<b>D04.075.080.875.099.221.249.190.190.115</b>	<b>Cefixime</b>
Old Tree	<b>D04.075.080.875.099.221.249.190.190.125</b>	<b>Cefmenoxime</b>
Old Tree	<b>D04.075.080.875.099.221.249.190.190.135</b>	<b>Cefotiam</b>
Old Tree	<b>D04.075.080.875.099.221.249.190.190.145</b>	<b>Ceftizoxime</b>
Old Tree	<b>D04.075.080.875.099.221.249.190.190.155</b>	<b>Ceftriaxone</b>
Old Tree	<b>D04.075.080.875.099.221.249.190.190.165</b>	<b>Cefuroxime</b>
Old Tree	<b>D04.075.080.875.099.221.249.190.210</b>	<b>Cephalothin</b>
Old Tree	<b>D04.075.080.875.099.221.249.190.230</b>	<b>Cephapirin</b>
Old Tree	<b>D04.075.080.875.099.221.249.200</b>	<b>Cephalexin</b>

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	D04.075.080.875.099.221.249.200.155 Cefaclor
Old Tree	D04.075.080.875.099.221.249.200.165 Cefadroxil
Old Tree	D04.075.080.875.099.221.249.200.165.125 Cefatrizine
Old Tree	D04.075.080.875.099.221.249.200.180 Cephaloglycin
Old Tree	D04.075.080.875.099.221.249.200.185 Cephradine
Old Tree	D04.075.080.875.099.221.249.210 Cephaloridine
Old Tree	D04.075.080.875.099.221.249.210.150 Ceftazidime
Old Tree	D04.075.080.875.099.221.249.250 Cephamycins
Old Tree	D04.075.080.875.099.221.249.250.177 Cefmetazole
Old Tree	D04.075.080.875.099.221.249.250.199 Cefotetan
Old Tree	D04.075.080.875.099.221.249.250.222 Cefoxitin
Old Tree	D04.075.080.875.099.221.374 Clavulanic Acids
Old Tree	D04.075.080.875.099.221.374.160 Clavulanic Acid
Old Tree	D04.075.080.875.099.221.374.160.060 Amoxicillin-Potassium Clavulanate Combination
Old Tree	D04.075.080.875.099.221.500 Monobactams
Old Tree	D04.075.080.875.099.221.500.044 Aztreonam
Old Tree	D04.075.080.875.099.221.625 Moxalactam
Old Tree	D04.075.080.875.099.221.750 Penicillins
Old Tree	D04.075.080.875.099.221.750.124 Amdinocillin
Old Tree	D04.075.080.875.099.221.750.124.036 Amdinocillin Pivoxil
Old Tree	D04.075.080.875.099.221.750.249 Cyclacillin
Old Tree	D04.075.080.875.099.221.750.500 Methicillin
Old Tree	D04.075.080.875.099.221.750.562 Nafcillin
Old Tree	D04.075.080.875.099.221.750.625 Oxacillin
Old Tree	D04.075.080.875.099.221.750.625.150 Cloxacillin
Old Tree	D04.075.080.875.099.221.750.625.150.205 Dicloxacillin
Old Tree	D04.075.080.875.099.221.750.625.150.250 Floxacillin
Old Tree	D04.075.080.875.099.221.750.687 Penicillanic Acid
Old Tree	D04.075.080.875.099.221.750.750 Penicillin G
Old Tree	D04.075.080.875.099.221.750.750.050 Ampicillin
Old Tree	D04.075.080.875.099.221.750.750.050.050 Amoxicillin
Old Tree	D04.075.080.875.099.221.750.750.050.050.500 Amoxicillin-Potassium Clavulanate Combination
Old Tree	D04.075.080.875.099.221.750.750.050.075 Azlocillin
Old Tree	D04.075.080.875.099.221.750.750.050.500 Mezlocillin



## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	D04.075.080.875.099.221.750.750.050.650 Piperacillin
Old Tree	D04.075.080.875.099.221.750.750.050.700 Pivampicillin
Old Tree	D04.075.080.875.099.221.750.750.050.900 Talampicillin
Old Tree	D04.075.080.875.099.221.750.750.170 Carbenicillin
Old Tree	D04.075.080.875.099.221.750.750.170.200 Carfecillin
Old Tree	D04.075.080.875.099.221.750.750.685 Penicillin G Benzathine
Old Tree	D04.075.080.875.099.221.750.750.695 Penicillin G Procaine
Old Tree	D04.075.080.875.099.221.750.750.875 Sulbenicillin
Old Tree	D04.075.080.875.099.221.750.781 Penicillin V
Old Tree	D04.075.080.875.099.221.750.812 Sulbactam
Old Tree	D04.075.080.875.099.221.750.875 Ticarcillin
Old Tree	D04.075.080.875.099.332 Biperiden
Old Tree	D04.075.080.875.099.444 Granisetron
Old Tree	D04.075.080.875.099.722 Tropanes
Old Tree	D04.075.080.875.099.722.229 Atropine Derivatives
Old Tree	D04.075.080.875.099.722.229.199 Atropine
Old Tree	D04.075.080.875.099.722.229.199.500 Hyoscyamine
Old Tree	D04.075.080.875.099.722.229.400 Ipratropium
Old Tree	D04.075.080.875.099.722.229.400.500 Albuterol, Ipratropium Drug Combination
Old Tree	D04.075.080.875.099.722.270 Benztropine
Old Tree	D04.075.080.875.099.722.388 Cocaine
Old Tree	D04.075.080.875.099.722.388.250 Crack Cocaine
Old Tree	D04.075.080.875.099.722.744 Nortropanes
Old Tree	D04.075.080.875.099.722.822 Scopolamine Derivatives
Old Tree	D04.075.080.875.099.722.822.200 Butylscopolammonium Bromide
Old Tree	D04.075.080.875.099.722.822.550 N-Methylscopolamine
Old Tree	D04.075.080.875.099.722.822.775 Scopolamine Hydrobromide
Old Tree	D04.075.080.875.099.722.822.887 Tiotropium Bromide
New Heading	<b>D04.210 Fused-Ring Compounds</b>
New Tree	D04.210.500 Steroids
New Tree	D04.210.500.054 Androstanes
New Tree	D04.210.500.054.040 Androstanols

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	D04.210.500.054.040.080      Androstane-3,17-diol
New Tree	D04.210.500.054.040.129      Androsterone
New Tree	D04.210.500.054.040.248      Dihydrotestosterone
New Tree	D04.210.500.054.040.248.450      Mesterolone
New Tree	D04.210.500.054.040.368      Etiocholanolone
New Tree	D04.210.500.054.040.581      Oxandrolone
New Tree	D04.210.500.054.040.632      Oxymetholone
New Tree	D04.210.500.054.040.685      Pancuronium
New Tree	D04.210.500.054.040.881      Stanozolol
New Tree	D04.210.500.054.040.920      Vecuronium Bromide
New Tree	D04.210.500.054.079      Androstenes
New Tree	D04.210.500.054.079.065      Abiraterone Acetate
New Tree	D04.210.500.054.079.129      Androstadienes
New Tree	D04.210.500.054.079.129.114      Fluticasone
New Tree	D04.210.500.054.079.129.114.500      Fluticasone Propionate, Salmeterol Xinafoate Drug Combination
New Tree	D04.210.500.054.079.129.284      Loteprednol Etabonate
New Tree	D04.210.500.054.079.129.453      Methandrostenolone
New Tree	D04.210.500.054.079.129.782      Testolactone
New Tree	D04.210.500.054.079.229      Androstatrienes
New Tree	D04.210.500.054.079.329      Androstenedione
New Tree	D04.210.500.054.079.429      Androstenols
New Tree	D04.210.500.054.079.429.154      Androstenediols

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	D04.210.500.054.079.429.154.050	Androstenediol
New Tree	D04.210.500.054.079.429.154.349	Fluoxymesterone
New Tree	D04.210.500.054.079.429.154.600	Methandriol
New Tree	D04.210.500.054.079.429.320	Cyanoketone
New Tree	D04.210.500.054.079.429.625	Dehydroepiandrosterone
New Tree	D04.210.500.054.079.429.625.300	Dehydroepiandrosterone Sulfate
New Tree	D04.210.500.054.079.429.824	Testosterone
New Tree	D04.210.500.054.079.429.824.275	Epitestosterone
New Tree	D04.210.500.054.079.429.824.375	Hydroxytestosterones
New Tree	D04.210.500.054.079.429.824.537	Methenolone
New Tree	D04.210.500.054.079.429.824.664	Methyltestosterone
New Tree	D04.210.500.054.079.429.824.832	Testosterone Propionate
New Tree	D04.210.500.054.079.500	Finasteride
New Tree	D04.210.500.105	Bile Acids and Salts
New Tree	D04.210.500.105.225	Cholic Acids
New Tree	D04.210.500.105.225.130	Cholic Acid
New Tree	D04.210.500.105.225.130.330	Cholates
New Tree	D04.210.500.105.225.130.330.850	Sodium Cholate
New Tree	D04.210.500.105.225.261	Dehydrocholic Acid
New Tree	D04.210.500.105.225.272	Deoxycholic Acid
New Tree	D04.210.500.105.225.272.150	Chenodeoxycholic Acid
New Tree	D04.210.500.105.225.272.150.350	Glycochenodeoxycholic Acid

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	D04.210.500.105.225.272.150.850 Taurochenodeoxycholic Acid
New Tree	D04.210.500.105.225.272.411 Glycodeoxycholic Acid
New Tree	D04.210.500.105.225.272.411.360 Glycochenodeoxycholic Acid
New Tree	D04.210.500.105.225.272.925 Taurodeoxycholic Acid
New Tree	D04.210.500.105.225.272.925.875 Taurochenodeoxycholic Acid
New Tree	D04.210.500.105.225.272.962 Ursodeoxycholic Acid
New Tree	D04.210.500.105.225.400 Glycocholic Acid
New Tree	D04.210.500.105.225.400.380 Glycodeoxycholic Acid
New Tree	D04.210.500.105.225.400.380.360 Glycochenodeoxycholic Acid
New Tree	D04.210.500.105.225.480 Lithocholic Acid
New Tree	D04.210.500.105.225.480.880 Taurolithocholic Acid
New Tree	D04.210.500.105.225.900 Taurocholic Acid
New Tree	D04.210.500.105.225.900.900 Taurodeoxycholic Acid
New Tree	D04.210.500.105.225.900.900.870 Taurochenodeoxycholic Acid
New Tree	D04.210.500.105.225.900.920 Taurolithocholic Acid
New Tree	D04.210.500.155 Cardanolides
New Tree	D04.210.500.155.580 Cardiac Glycosides
New Tree	D04.210.500.155.580.064 Bufanolides
New Tree	D04.210.500.155.580.064.580 Proscillaridin
New Tree	D04.210.500.155.580.130 Cardenolides
New Tree	D04.210.500.155.580.130.500 Digitalis Glycosides
New Tree	D04.210.500.155.580.130.500.236 Digitonin

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	D04.210.500.155.580.130.500.336 Digitoxin
New Tree	D04.210.500.155.580.130.500.336.259 Acetyldigitoxins
New Tree	D04.210.500.155.580.130.500.436 Digoxin
New Tree	D04.210.500.155.580.130.500.436.050 Acetyldigoxins
New Tree	D04.210.500.155.580.130.500.436.500 Medigoxin
New Tree	D04.210.500.155.580.130.500.657 Lanatosides
New Tree	D04.210.500.155.580.130.500.657.200 Deslanoside
New Tree	D04.210.500.155.580.130.625 Digitoxin
New Tree	D04.210.500.155.580.130.625.259 Acetyldigitoxins
New Tree	D04.210.500.155.580.130.625.350 Digitoxigenin
New Tree	D04.210.500.155.580.130.688 Digoxin
New Tree	D04.210.500.155.580.130.688.050 Acetyldigoxins
New Tree	D04.210.500.155.580.130.688.350 Digoxigenin
New Tree	D04.210.500.155.580.130.688.500 Medigoxin
New Tree	D04.210.500.155.580.130.750 Strophanthins
New Tree	D04.210.500.155.580.130.750.250 Cymarine
New Tree	D04.210.500.155.580.130.750.600 Ouabain
New Tree	D04.210.500.155.580.130.750.800 Strophanthidin
New Tree	D04.210.500.221 Cholanes
New Tree	D04.210.500.221.263 Cholenes
New Tree	D04.210.500.221.430 Cholic Acids
New Tree	D04.210.500.221.430.130 Cholic Acid

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	D04.210.500.221.430.130.330 Cholates
New Tree	D04.210.500.221.430.130.330.850 Sodium Cholate
New Tree	D04.210.500.221.430.265 Dehydrocholic Acid
New Tree	D04.210.500.221.430.342 Deoxycholic Acid
New Tree	D04.210.500.221.430.342.300 Chenodeoxycholic Acid
New Tree	D04.210.500.221.430.342.300.400 Glycochenodeoxycholic Acid
New Tree	D04.210.500.221.430.342.300.900 Taurochenodeoxycholic Acid
New Tree	D04.210.500.221.430.342.400 Glycodeoxycholic Acid
New Tree	D04.210.500.221.430.342.400.450 Glycochenodeoxycholic Acid
New Tree	D04.210.500.221.430.342.900 Taurodeoxycholic Acid
New Tree	D04.210.500.221.430.342.900.910 Taurochenodeoxycholic Acid
New Tree	D04.210.500.221.430.342.925 Ursodeoxycholic Acid
New Tree	D04.210.500.221.430.484 Glycocholic Acid
New Tree	D04.210.500.221.430.484.430 Glycodeoxycholic Acid
New Tree	D04.210.500.221.430.484.430.420 Glycochenodeoxycholic Acid
New Tree	D04.210.500.221.430.622 Lithocholic Acid
New Tree	D04.210.500.221.430.622.900 Taurolithocholic Acid
New Tree	D04.210.500.221.430.873 Taurocholic Acid
New Tree	D04.210.500.221.430.873.920 Taurodeoxycholic Acid
New Tree	D04.210.500.221.430.873.920.900 Taurochenodeoxycholic Acid
New Tree	D04.210.500.221.430.873.940 Taurolithocholic Acid
New Tree	D04.210.500.247 Cholestanes

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	D04.210.500.247.100 Cholestanols
New Tree	D04.210.500.247.100.250 Cholestanol
New Tree	D04.210.500.247.125 Cholestanones
New Tree	D04.210.500.247.222 Cholestenes
New Tree	D04.210.500.247.222.159 Cholecalciferol
New Tree	D04.210.500.247.222.159.478 Hydroxycholecalciferols
New Tree	D04.210.500.247.222.159.478.250 Calcifediol
New Tree	D04.210.500.247.222.159.478.387 Dihydroxycholecalciferols
New Tree	D04.210.500.247.222.159.478.387.300 Calcitriol
New Tree	D04.210.500.247.222.159.478.387.400 24,25-Dihydroxyvitamin D 3
New Tree	D04.210.500.247.222.222 Cholestadienes
New Tree	D04.210.500.247.222.222.347 Cholestadienols
New Tree	D04.210.500.247.222.222.347.200 Dehydrocholesterols
New Tree	D04.210.500.247.222.222.347.231 Desmosterol
New Tree	D04.210.500.247.222.222.347.389 Fusidic Acid
New Tree	D04.210.500.247.222.222.347.557 Lanosterol
New Tree	D04.210.500.247.222.222.347.833 Stigmasterol
New Tree	D04.210.500.247.222.265 Cholestenones
New Tree	D04.210.500.247.222.265.165 Ecdysteroids
New Tree	D04.210.500.247.222.265.165.500 Ecdysone
New Tree	D04.210.500.247.222.265.165.750 Ecdysterone
New Tree	D04.210.500.247.222.265.450 Ketocholesterols

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	D04.210.500.247.222.284 Cholesterol
New Tree	D04.210.500.247.222.284.070 Azacosterol
New Tree	D04.210.500.247.222.284.200 Cholesterol Esters
New Tree	D04.210.500.247.222.284.510 19-Iodocholesterol
New Heading	<b>D04.210.500.247.222.284.800 Oxysterols</b>
New Tree	D04.210.500.247.222.284.800.500 Hydroxycholesterols
New Tree	D04.210.500.247.222.284.800.750 Ketocholesterols
New Tree	D04.210.500.247.222.474 Ergocalciferols
New Tree	D04.210.500.247.222.474.250 Dihydrotachysterol
New Tree	D04.210.500.247.222.474.400 25-Hydroxyvitamin D 2
New Tree	D04.210.500.247.222.537 Ergosterol
New Tree	D04.210.500.247.222.537.888 Withanolides
New Tree	D04.210.500.247.222.857 Sitosterols
New Tree	D04.210.500.247.515 Spirostans
New Tree	D04.210.500.247.515.500 Diosgenin
New Tree	D04.210.500.247.515.500.500 Solanine
New Tree	D04.210.500.247.808 Sterols
New Tree	D04.210.500.247.808.050 Adosterol
New Tree	D04.210.500.247.808.146 Cholecalciferol
New Tree	D04.210.500.247.808.146.478 Hydroxycholecalciferols
New Tree	D04.210.500.247.808.146.478.250 Calcifediol
New Tree	D04.210.500.247.808.146.478.387 Dihydroxycholecalciferols



## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	D04.210.500.247.808.146.478.387.300 <span style="float: right;">Calcitriol</span>
New Tree	D04.210.500.247.808.146.478.387.400 <span style="float: right;">24,25-Dihydroxyvitamin D 3</span>
New Tree	D04.210.500.247.808.197 <span style="float: right;">Cholesterol</span>
New Tree	D04.210.500.247.808.197.070 <span style="float: right;">Azacosterol</span>
New Tree	D04.210.500.247.808.197.135 <span style="float: right;">Cholestanol</span>
New Tree	D04.210.500.247.808.197.200 <span style="float: right;">Cholesterol Esters</span>
New Tree	D04.210.500.247.808.197.225 <span style="float: right;">Cholesterol, Dietary</span>
New Tree	D04.210.500.247.808.197.238 <span style="float: right;">Cholesterol, HDL</span>
New Tree	D04.210.500.247.808.197.244 <span style="float: right;">Cholesterol, LDL</span>
New Tree	D04.210.500.247.808.197.247 <span style="float: right;">Cholesterol, VLDL</span>
New Tree	D04.210.500.247.808.197.250 <span style="float: right;">Dehydrocholesterols</span>
New Tree	D04.210.500.247.808.197.250.281 <span style="float: right;">Desmosterol</span>
New Tree	D04.210.500.247.808.197.580 <span style="float: right;">19-Iodocholesterol</span>
New Heading	<b>D04.210.500.247.808.197.800</b> <span style="float: right;"><b>Oxysterols</b></span>
New Tree	D04.210.500.247.808.197.800.500 <span style="float: right;">Hydroxycholesterols</span>
New Tree	D04.210.500.247.808.197.800.750 <span style="float: right;">Ketocholesterols</span>
New Tree	D04.210.500.247.808.412 <span style="float: right;">Ergocalciferols</span>
New Tree	D04.210.500.247.808.412.250 <span style="float: right;">Dihydrotachysterol</span>
New Tree	D04.210.500.247.808.412.400 <span style="float: right;">25-Hydroxyvitamin D 2</span>
New Tree	D04.210.500.247.808.489 <span style="float: right;">Fusidic Acid</span>
New Tree	D04.210.500.247.808.607 <span style="float: right;">Lanosterol</span>
New Tree	D04.210.500.247.808.756 <span style="float: right;">Phytosterols</span>

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	D04.210.500.247.808.756.071 <span style="float: right;">Brassinosteroids</span>
New Tree	D04.210.500.247.808.756.143 <span style="float: right;">Ecdysteroids</span>
New Tree	D04.210.500.247.808.756.669 <span style="float: right;">Sitosterols</span>
New Tree	D04.210.500.247.808.756.808 <span style="float: right;">Stigmasterol</span>
New Tree	D04.210.500.247.808.756.904 <span style="float: right;">Withanolides</span>
New Tree	D04.210.500.247.808.874 <span style="float: right;">Solanine</span>
New Tree	D04.210.500.294 <span style="float: right;">Cyclosteroids</span>
New Tree	D04.210.500.365 <span style="float: right;">Estranes</span>
New Tree	D04.210.500.365.415 <span style="float: right;">Estrenes</span>
New Tree	D04.210.500.365.415.050 <span style="float: right;">Allylestrenol</span>
New Tree	D04.210.500.365.415.200 <span style="float: right;">Epimestrol</span>
New Tree	D04.210.500.365.415.215 <span style="float: right;">Equilenin</span>
New Tree	D04.210.500.365.415.220 <span style="float: right;">Equilin</span>
New Tree	D04.210.500.365.415.248 <span style="float: right;">Estradiol</span>
New Tree	D04.210.500.365.415.248.320 <span style="float: right;">Estramustine</span>
New Tree	D04.210.500.365.415.331 <span style="float: right;">Estriol</span>
New Tree	D04.210.500.365.415.331.320 <span style="float: right;">Estetrol</span>
New Tree	D04.210.500.365.415.414 <span style="float: right;">Estrone</span>
New Tree	D04.210.500.365.415.414.400 <span style="float: right;">Hydroxyestrones</span>
New Tree	D04.210.500.365.415.550 <span style="float: right;">Metribolone</span>
New Tree	D04.210.500.365.415.580 <span style="float: right;">Mifepristone</span>
New Tree	D04.210.500.365.415.638 <span style="float: right;">Nandrolone</span>

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	D04.210.500.365.415.930 <span style="float: right;">Trenbolone Acetate</span>
New Tree	D04.210.500.451 <span style="float: right;">Gonanes</span>
New Tree	D04.210.500.496 <span style="float: right;">Homosteroids</span>
New Tree	D04.210.500.496.699 <span style="float: right;">Testolactone</span>
New Tree	D04.210.500.528 <span style="float: right;">Hydroxysteroids</span>
New Tree	D04.210.500.578 <span style="float: right;">Ketosteroids</span>
New Tree	D04.210.500.578.502 <span style="float: right;">17-Ketosteroids</span>
New Tree	D04.210.500.578.502.112 <span style="float: right;">Androstenedione</span>
New Tree	D04.210.500.578.502.195 <span style="float: right;">Androsterone</span>
New Tree	D04.210.500.578.502.400 <span style="float: right;">Dehydroepiandrosterone</span>
New Tree	D04.210.500.578.502.400.300 <span style="float: right;">Dehydroepiandrosterone Sulfate</span>
New Tree	D04.210.500.578.502.470 <span style="float: right;">Equilenin</span>
New Tree	D04.210.500.578.502.475 <span style="float: right;">Equilin</span>
New Tree	D04.210.500.578.502.497 <span style="float: right;">Estrone</span>
New Tree	D04.210.500.578.502.497.400 <span style="float: right;">Hydroxyestrones</span>
New Tree	D04.210.500.578.502.583 <span style="float: right;">Etiocholanolone</span>
New Tree	D04.210.500.668 <span style="float: right;">Norsteroids</span>
New Tree	D04.210.500.668.100 <span style="float: right;">Norandrostanes</span>
New Tree	D04.210.500.668.651 <span style="float: right;">Norpregnanes</span>
New Tree	D04.210.500.668.651.443 <span style="float: right;">Norpregnadienes</span>
New Tree	D04.210.500.668.651.443.680 <span style="float: right;">Promegestone</span>
New Tree	D04.210.500.668.651.568 <span style="float: right;">Norpregnatrienes</span>

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	D04.210.500.668.651.568.291 Ethinyl Estradiol
New Tree	D04.210.500.668.651.568.291.250 Ethinyl Estradiol-Norgestrel Combination
New Tree	D04.210.500.668.651.568.291.500 Mestranol
New Tree	D04.210.500.668.651.568.291.750 Quinestrol
New Tree	D04.210.500.668.651.568.620 Norgestrienone
New Tree	D04.210.500.668.651.568.620.400 Gestrinone
New Tree	D04.210.500.668.651.693 Norpregnenes
New Tree	D04.210.500.668.651.693.175 Desogestrel
New Tree	D04.210.500.668.651.693.223 Ethylestrenol
New Tree	D04.210.500.668.651.693.279 Ethynodiol Diacetate
New Tree	D04.210.500.668.651.693.362 Gestonorone Caproate
New Tree	D04.210.500.668.651.693.494 Lynestrenol
New Tree	D04.210.500.668.651.693.595 Norethandrolone
New Tree	D04.210.500.668.651.693.651 Norethindrone
New Tree	D04.210.500.668.651.693.706 Norethynodrel
New Tree	D04.210.500.668.651.693.762 Norgestrel
New Tree	D04.210.500.668.651.693.762.225 Ethinyl Estradiol-Norgestrel Combination
New Tree	D04.210.500.668.651.693.762.450 Levonorgestrel
New Tree	D04.210.500.668.651.693.820 Norprogesterones
New Tree	D04.210.500.745 Pregnanes
New Tree	D04.210.500.745.432 Pregnadienes
New Tree	D04.210.500.745.432.100 Canrenoic Acid

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	D04.210.500.745.432.120 <span style="float: right;">Canrenone</span>
New Tree	D04.210.500.745.432.144 <span style="float: right;">Chlormadinone Acetate</span>
New Tree	D04.210.500.745.432.219 <span style="float: right;">Cyproterone</span>
New Tree	D04.210.500.745.432.219.150 <span style="float: right;">Cyproterone Acetate</span>
New Tree	D04.210.500.745.432.235 <span style="float: right;">Danazol</span>
New Tree	D04.210.500.745.432.296 <span style="float: right;">Dydrogesterone</span>
New Tree	D04.210.500.745.432.370 <span style="float: right;">Fluocinolone Acetonide</span>
New Tree	D04.210.500.745.432.370.325 <span style="float: right;">Fluocinonide</span>
New Tree	D04.210.500.745.432.481 <span style="float: right;">Medrogestone</span>
New Tree	D04.210.500.745.432.531 <span style="float: right;">Megestrol</span>
New Tree	D04.210.500.745.432.531.500 <span style="float: right;">Megestrol Acetate</span>
New Tree	D04.210.500.745.432.581 <span style="float: right;">Melengestrol Acetate</span>
New Tree	D04.210.500.745.432.719 <span style="float: right;">Pregnadienediols</span>
New Tree	D04.210.500.745.432.719.260 <span style="float: right;">Desoximetasone</span>
New Tree	D04.210.500.745.432.719.320 <span style="float: right;">Fluocortolone</span>
New Tree	D04.210.500.745.432.719.320.250 <span style="float: right;">Diflucortolone</span>
New Tree	D04.210.500.745.432.719.349 <span style="float: right;">Fluorometholone</span>
New Tree	D04.210.500.745.432.719.526 <span style="float: right;">Mometasone Furoate</span>
New Tree	D04.210.500.745.432.719.526.500 <span style="float: right;">Mometasone Furoate, Formoterol</span> Fumarate Drug Combination
New Tree	D04.210.500.745.432.719.702 <span style="float: right;">Prednisone</span>
New Tree	D04.210.500.745.432.769 <span style="float: right;">Pregnadienetriols</span>
New Tree	D04.210.500.745.432.769.125 <span style="float: right;">Beclomethasone</span>

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	D04.210.500.745.432.769.199 <span style="float: right;">Betamethasone</span>
New Tree	D04.210.500.745.432.769.199.150 <span style="float: right;">Betamethasone Valerate</span>
New Tree	D04.210.500.745.432.769.325 <span style="float: right;">Desonide</span>
New Tree	D04.210.500.745.432.769.344 <span style="float: right;">Dexamethasone</span>
New Tree	D04.210.500.745.432.769.344.300 <span style="float: right;">Dexamethasone Isonicotinate</span>
New Tree	D04.210.500.745.432.769.465 <span style="float: right;">Flumethasone</span>
New Tree	D04.210.500.745.432.769.692 <span style="float: right;">Paramethasone</span>
New Tree	D04.210.500.745.432.769.795 <span style="float: right;">Prednisolone</span>
New Tree	D04.210.500.745.432.769.795.307 <span style="float: right;">Fluprednisolone</span>
New Tree	D04.210.500.745.432.769.795.539 <span style="float: right;">Methylprednisolone</span>
New Tree	D04.210.500.745.432.769.795.539.500 <span style="float: right;">Methylprednisolone Hemisuccinate</span>
New Tree	D04.210.500.745.432.769.795.700 <span style="float: right;">Prednimustine</span>
New Tree	D04.210.500.745.432.915 <span style="float: right;">Triamcinolone</span>
New Tree	D04.210.500.745.432.915.715 <span style="float: right;">Triamcinolone Acetonide</span>
New Tree	D04.210.500.745.495 <span style="float: right;">Pregnanediol</span>
New Tree	D04.210.500.745.558 <span style="float: right;">Pregnenediones</span>
New Tree	D04.210.500.745.558.050 <span style="float: right;">Alfaxalone Alfadolone Mixture</span>
New Tree	D04.210.500.745.558.075 <span style="float: right;">5-alpha-Dihydroprogesterone</span>
New Tree	D04.210.500.745.558.783 <span style="float: right;">Tetrahydrocortisone</span>
New Tree	D04.210.500.745.620 <span style="float: right;">Pregnanetriol</span>
New Tree	D04.210.500.745.640 <span style="float: right;">Pregnanolone</span>
New Tree	D04.210.500.745.683 <span style="float: right;">Pregnatrienes</span>

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	D04.210.500.745.745 <span style="float: right;">Pregnenes</span>
New Tree	D04.210.500.745.745.183 <span style="float: right;">Cortisone</span>
New Tree	D04.210.500.745.745.279 <span style="float: right;">Dimethisterone</span>
New Tree	D04.210.500.745.745.379 <span style="float: right;">Ethisterone</span>
New Tree	D04.210.500.745.745.654 <span style="float: right;">Pregnenediones</span>
New Tree	D04.210.500.745.745.654.062 <span style="float: right;">Aldosterone</span>
New Tree	D04.210.500.745.745.654.105 <span style="float: right;">Budesonide</span>
New Tree	D04.210.500.745.745.654.105.500 <span style="float: right;">Budesonide, Formoterol Fumarate Drug</span> Combination
New Tree	D04.210.500.745.745.654.237 <span style="float: right;">Corticosterone</span>
New Tree	D04.210.500.745.745.654.237.400 <span style="float: right;">18-Hydroxycorticosterone</span>
New Tree	D04.210.500.745.745.654.252 <span style="float: right;">Cortodoxone</span>
New Tree	D04.210.500.745.745.654.339 <span style="float: right;">Desoxycorticosterone</span>
New Tree	D04.210.500.745.745.654.339.400 <span style="float: right;">18-Hydroxydesoxycorticosterone</span>
New Tree	D04.210.500.745.745.654.339.700 <span style="float: right;">Desoxycorticosterone Acetate</span>
New Tree	D04.210.500.745.745.654.473 <span style="float: right;">Flurandrenolone</span>
New Tree	D04.210.500.745.745.654.485 <span style="float: right;">Flurogestone Acetate</span>
New Tree	D04.210.500.745.745.654.527 <span style="float: right;">Gestonorone Caproate</span>
New Tree	D04.210.500.745.745.654.570 <span style="float: right;">Halcinonide</span>
New Tree	D04.210.500.745.745.654.600 <span style="float: right;">Hydrocortisone</span>
New Tree	D04.210.500.745.745.654.600.500 <span style="float: right;">Fludrocortisone</span>
New Tree	D04.210.500.745.745.654.829 <span style="float: right;">Progesterone</span>
New Tree	D04.210.500.745.745.654.829.025 <span style="float: right;">Algestone</span>

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	D04.210.500.745.745.654.829.025.025 <span style="float: right;">Algestone Acetophenide</span>
New Tree	D04.210.500.745.745.654.829.100 <span style="float: right;">20-alpha-Dihydroprogesterone</span>
New Tree	D04.210.500.745.745.654.829.395 <span style="float: right;">Hydroxyprogesterones</span>
New Tree	D04.210.500.745.745.654.829.395.400 <span style="float: right;">17-alpha-Hydroxyprogesterone</span>
New Tree	D04.210.500.745.745.654.829.395.700 <span style="float: right;">Medroxyprogesterone</span>
New Tree	D04.210.500.745.745.654.829.395.700.500 <span style="float: right;">Medroxyprogesterone Acetate</span>
New Tree	D04.210.500.745.745.725 <span style="float: right;">Pregnenolone</span>
New Tree	D04.210.500.745.745.725.395 <span style="float: right;">17-alpha-Hydroxypregnenolone</span>
New Tree	D04.210.500.745.745.725.703 <span style="float: right;">Pregnenolone Carbonitrile</span>
New Tree	D04.210.500.745.745.855 <span style="float: right;">Spironolactone</span>
New Tree	D04.210.500.745.887 <span style="float: right;">Tetrahydrocortisol</span>
New Tree	D04.210.500.795 <span style="float: right;">Sapogenins</span>
New Tree	D04.210.500.812 <span style="float: right;">Secosteroids</span>
New Tree	D04.210.500.812.768 <span style="float: right;">Vitamin D</span>
New Tree	D04.210.500.812.768.196 <span style="float: right;">Cholecalciferol</span>
New Tree	D04.210.500.812.768.196.478 <span style="float: right;">Hydroxycholecalciferols</span>
New Tree	D04.210.500.812.768.196.478.250 <span style="float: right;">Calcifediol</span>
New Tree	D04.210.500.812.768.196.478.387 <span style="float: right;">Dihydroxycholecalciferols</span>
New Tree	D04.210.500.812.768.196.478.387.300 <span style="float: right;">Calcitriol</span>
New Tree	D04.210.500.812.768.196.478.387.400 <span style="float: right;">24,25-Dihydroxyvitamin D 3</span>
New Tree	D04.210.500.812.768.462 <span style="float: right;">Ergocalciferols</span>
New Tree	D04.210.500.812.768.462.250 <span style="float: right;">Dihydrotachysterol</span>



## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	D04.210.500.812.768.462.400	25-Hydroxyvitamin D 2
New Tree	D04.210.500.875	Steroids, Brominated
New Tree	D04.210.500.883	Steroids, Chlorinated
New Tree	D04.210.500.883.154	Beclomethasone
New Tree	D04.210.500.883.294	Chlormadinone Acetate
New Tree	D04.210.500.883.419	Cyproterone
New Tree	D04.210.500.883.419.150	Cyproterone Acetate
New Tree	D04.210.500.908	Steroids, Fluorinated
New Tree	D04.210.500.908.093	Betamethasone
New Tree	D04.210.500.908.093.150	Betamethasone Valerate
New Tree	D04.210.500.908.093.250	Clobetasol
New Tree	D04.210.500.908.238	Dexamethasone
New Tree	D04.210.500.908.238.250	Desoximetasone
New Tree	D04.210.500.908.238.300	Dexamethasone Isonicotinate
New Tree	D04.210.500.908.359	Flumethasone
New Tree	D04.210.500.908.394	Fluocinolone Acetonide
New Tree	D04.210.500.908.394.300	Fluocinonide
New Tree	D04.210.500.908.405	Fluocortolone
New Tree	D04.210.500.908.405.250	Difluocortolone
New Tree	D04.210.500.908.431	Fluorometholone
New Tree	D04.210.500.908.466	Fluoxymesterone
New Tree	D04.210.500.908.502	Fluprednisolone

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	D04.210.500.908.537	Flurandrenolone
New Tree	D04.210.500.908.550	Flurogestone Acetate
New Tree	D04.210.500.908.727	Paramethasone
New Tree	D04.210.500.908.891	Triamcinolone
New Tree	D04.210.500.908.891.927	Triamcinolone Acetonide
New Tree	D04.210.500.925	Steroids, Heterocyclic
New Tree	D04.210.500.925.100	Azasteroids
New Tree	D04.210.500.925.100.125	Dutasteride
New Tree	D04.210.500.925.100.250	Finasteride
-	D04.345	Macrocyclic Compounds
-	D04.345.025	Calixarenes
-	D04.345.051	Crown Compounds
-	D04.345.051.500	Crown Ethers
-	D04.345.103	Cyclodextrins
-	D04.345.103.222	alpha-Cyclodextrins
-	D04.345.103.333	beta-Cyclodextrins
-	D04.345.103.444	gamma-Cyclodextrins
-	D04.345.133	Cycloparaffins
-	D04.345.241	Ethers, Cyclic
-	D04.345.241.308	Crown Ethers
-	D04.345.295	Lactams, Macrocyclic
-	D04.345.295.500	Maytansine
-	D04.345.295.750	Rifamycins
-	D04.345.295.750.650	Rifabutin
-	D04.345.295.750.650.700	Streptovaricin
-	D04.345.295.750.700	Rifampin
-	D04.345.566	Peptides, Cyclic
-	D04.345.566.040	Alamethicin
-	D04.345.566.050	Amanitins
-	D04.345.566.050.111	Alpha-Amanitin

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D04.345.566.075 Bacitracin
-	D04.345.566.142 Capreomycin
-	D04.345.566.235 Cyclosporins
-	D04.345.566.235.300 Cyclosporine
-	D04.345.566.252 Dactinomycin
-	D04.345.566.270 Daptomycin
-	D04.345.566.297 Depsipeptides
-	D04.345.566.297.500 Valinomycin
-	D04.345.566.325 Echinomycin
-	D04.345.566.352 Enterobactin
-	D04.345.566.380 Ferrichrome
-	D04.345.566.447 Microcystins
-	D04.345.566.515 Mycobacillin
-	D04.345.566.582 Nisin
-	D04.345.566.650 Octreotide
-	D04.345.566.735 Phalloidine
-	D04.345.566.780 Polymyxins
-	D04.345.566.780.110 Colistin
-	D04.345.566.780.750 Polymyxin B
-	D04.345.566.802 Streptogramins
-	D04.345.566.802.249 Mikamycin
-	D04.345.566.802.374 Pristinamycin
-	D04.345.566.802.500 Streptogramin Group A
-	D04.345.566.802.500.500 Streptogramin A
-	D04.345.566.802.750 Streptogramin Group B
-	D04.345.566.802.750.500 Streptogramin B
-	D04.345.566.802.812 Vernamycin B
-	D04.345.566.802.875 Virginiamycin
-	D04.345.566.802.875.500 Streptogramin A
-	D04.345.566.825 Thiostrepton
-	D04.345.566.850 Tyrothricin
-	D04.345.566.850.300 Gramicidin
-	D04.345.566.850.920 Tyrocidine
-	D04.345.566.875 Viomycin
-	D04.345.566.875.250 Enviomycin
-	D04.345.674 Polyketides

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D04.345.674.500 Macrolides
-	D04.345.783 Tetrapyrroles
-	D04.345.783.249 Bile Pigments
-	D04.345.783.249.184 Bilirubin
-	D04.345.783.249.184.200 Biliverdine
-	D04.345.783.249.727 Urobilin
-	D04.345.783.249.852 Urobilinogen
-	D04.345.783.374 Chlorophyll
-	D04.345.783.374.100 Bacteriochlorophylls
-	D04.345.783.374.180 Chlorophyllides
-	D04.345.783.374.700 Pheophytins
-	D04.345.783.374.725 Protochlorophyllide
-	D04.345.783.437 Corrinoids
-	D04.345.783.437.777 Vitamin B 12
-	D04.345.783.437.777.270 Cobamides
-	D04.345.783.437.777.560 Hydroxocobalamin
-	D04.345.783.468 Phycobilins
-	D04.345.783.500 Porphyrins
-	D04.345.783.500.250 Coproporphyrins
-	D04.345.783.500.280 Deuteroporphyrins
-	D04.345.783.500.340 Etioporphyrins
-	D04.345.783.500.462 Hematoporphyrins
-	D04.345.783.500.462.400 Hematoporphyrin Derivative
-	D04.345.783.500.462.400.200 Dihematoporphyrin Ether
-	D04.345.783.500.620 Mesoporphyrins
-	D04.345.783.500.640 Metalloporphyrins
-	D04.345.783.500.640.220 Chlorophyll
-	D04.345.783.500.640.220.100 Bacteriochlorophylls
-	D04.345.783.500.640.220.180 Chlorophyllides
-	D04.345.783.500.640.220.453 Pheophytins
-	D04.345.783.500.640.220.725 Protochlorophyllide
-	D04.345.783.500.640.587 Heme
-	D04.345.783.500.640.587.462 Hemin
-	D04.345.783.500.700 Porphyrinogens
-	D04.345.783.500.700.250 Coproporphyrinogens
-	D04.345.783.500.700.900 Uroporphyrinogens

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D04.345.783.500.725                      Protoporphyrins
-	D04.345.783.500.880                      Uroporphyrins
-	D04.345.891                                  Trichothecenes
-	D04.345.891.870                            T-2 Toxin
-	D04.345.891.900                            Trichodermin
-	D04.615                                        Polycyclic Aromatic Hydrocarbons
-	D04.615                                        Polycyclic Hydrocarbons, Aromatic
-	D04.615.117                                  Anthracenes
-	D04.615.117.050                            Anthralin
-	D04.615.117.159                            Anthraquinones
-	D04.615.117.159.200                      Carmine
-	D04.615.117.159.205                      Cascara
-	D04.615.117.159.205.400                Emodin
-	D04.615.117.159.500                      Mitoxantrone
-	D04.615.117.600                            Maprotiline
-	D04.615.133                                  Azulenes
-	D04.615.149                                  Benz(a)Anthracenes
-	D04.615.149.301                            9,10-Dimethyl-1,2-benzanthracene
-	D04.615.149.500                            Methylcholanthrene
-	D04.615.149.700                            Perylene
-	D04.615.181                                  Benzocycloheptenes
-	D04.615.181.384                            Dibenzocycloheptenes
-	D04.615.181.384.100                      Amitriptyline
-	D04.615.181.384.230                      Butaclamol
-	D04.615.181.384.340                      Cyproheptadine
-	D04.615.181.384.340.500                Loratadine
-	D04.615.181.384.380                      Dizocilpine Maleate
-	D04.615.181.384.535                      Nortriptyline
-	D04.615.181.384.650                      Protriptyline
-	D04.615.389                                  Fluorenes
-	D04.615.389.050                            2-Acetylaminofluorene
-	D04.615.389.050.060                      Acetoxyacetylaminofluorene
-	D04.615.389.050.400                      Hydroxyacetylaminofluorene
-	D04.615.389.850                            Tilorone
-	D04.615.486                                  Indenes
-	D04.615.486.250                            Dimethindene

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D04.615.486.487 Indans
-	D04.615.486.487.060 Aprindine
-	D04.615.486.487.500 Ninhydrin
-	D04.615.486.487.750 Phenindione
-	D04.615.486.875 Sulindac
-	D04.615.562 Naphthacenes
-	D04.615.562.050 Anthracyclines
-	D04.615.562.050.050 Aclarubicin
-	D04.615.562.050.200 Daunorubicin
-	D04.615.562.050.200.150 Carubicin
-	D04.615.562.050.200.175 Doxorubicin
-	D04.615.562.050.200.175.200 Epirubicin
-	D04.615.562.050.200.300 Idarubicin
-	D04.615.562.050.200.650 Nogalamycin
-	D04.615.562.050.200.650.500 Menogaril
-	D04.615.562.050.650 Plicamycin
-	D04.615.562.900 Tetracyclines
-	D04.615.562.900.146 Chlortetracycline
-	D04.615.562.900.185 Demeclocycline
-	D04.615.562.900.200 Doxycycline
-	D04.615.562.900.450 Lymecycline
-	D04.615.562.900.525 Methacycline
-	D04.615.562.900.550 Minocycline
-	D04.615.562.900.600 Oxytetracycline
-	D04.615.562.900.720 Rolitetracycline
-	D04.615.562.900.875 Tetracycline
-	D04.615.638 Naphthalenes
-	D04.615.638.059 Acenaphthenes
-	D04.615.638.090 Adapalene
-	D04.615.638.090.500 Adapalene, Benzoyl Peroxide Drug Combination
-	D04.615.638.120 Bunaftine
-	D04.615.638.162 Carbaryl
-	D04.615.638.183 Cinacalcet Hydrochloride
-	D04.615.638.204 Dansyl Compounds
-	D04.615.638.400 Lovastatin
-	D04.615.638.400.900 Simvastatin

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D04.615.638.400.900.500 Ezetimibe, Simvastatin Drug Combination
-	D04.615.638.472 Naphthaleneacetic Acids
-	D04.615.638.472.450 Naproxen
-	D04.615.638.555 Naphthalenesulfonates
-	D04.615.638.555.200 Amaranth Dye
-	D04.615.638.555.220 Amido Black
-	D04.615.638.555.282 Anilino Naphthalenesulfonates
-	D04.615.638.555.300 Congo Red
-	D04.615.638.555.400 Evans Blue
-	D04.615.638.555.750 Suramin
-	D04.615.638.555.875 Trypan Blue
-	D04.615.638.596 Naphthalimides
-	D04.615.638.638 Naphthols
-	D04.615.638.721 Naphthoquinones
-	D04.615.638.721.186 Atovaquone
-	D04.615.638.721.374 Vitamin K
-	D04.615.638.721.374.689 Vitamin K 1
-	D04.615.638.721.374.844 Vitamin K 2
-	D04.615.638.721.374.922 Vitamin K 3
-	D04.615.638.845 1-Naphthylamine
-	D04.615.638.845.800 Sertraline
-	D04.615.638.850 2-Naphthylamine
-	D04.615.638.870 1-Naphthylisothiocyanate
-	D04.615.638.900 Naphthylvinylpyridine
-	D04.615.638.930 Pravastatin
-	D04.615.638.945 Propranolol
-	D04.615.638.960 Tetrahydronaphthalenes
-	D04.615.638.960.400 8-Hydroxy-2-(di-n-propylamino)tetralin
-	D04.615.638.960.446 Bunolol
-	D04.615.638.960.446.500 Levobunolol
-	D04.615.638.960.585 Mibefradil
-	D04.615.638.960.675 Podophyllotoxin
-	D04.615.638.960.675.250 Etoposide
-	D04.615.638.960.837 Tetralones
-	D04.615.638.975 Tolnaftate
-	D04.615.680 Phenalenes

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D04.615.680.500 Perylene
-	D04.615.723 Phenanthrenes
-	D04.615.723.089 Aristolochic Acids
-	D04.615.723.180 Chrysenes
-	D04.615.723.590 Diterpenes, Abietane
-	D04.615.723.795 Morphinans
-	D04.615.723.795.106 Benzomorphans
-	D04.615.723.795.106.700 Pentazocine
-	D04.615.723.795.106.715 Phenazocine
-	D04.615.723.795.150 Buprenorphine
-	D04.615.723.795.150.500 Buprenorphine, Naloxone Drug Combination
-	D04.615.723.795.165 Butorphanol
-	D04.615.723.795.200 Dextromethorphan
-	D04.615.723.795.235 Dextrophan
-	D04.615.723.795.250 Diprenorphine
-	D04.615.723.795.270 Etorphine
-	D04.615.723.795.413 Levallorphan
-	D04.615.723.795.463 Levorphanol
-	D04.615.723.795.576 Morphine Derivatives
-	D04.615.723.795.576.149 Codeine
-	D04.615.723.795.576.149.287 Hydrocodone
-	D04.615.723.795.576.149.575 Oxycodone
-	D04.615.723.795.576.350 Dihydromorphine
-	D04.615.723.795.576.430 Ethylmorphine
-	D04.615.723.795.576.445 Heroin
-	D04.615.723.795.576.450 Hydromorphone
-	D04.615.723.795.576.571 Morphine
-	D04.615.723.795.576.692 Oxymorphone
-	D04.615.723.795.576.856 Thebaine
-	D04.615.723.795.606 Nalbuphine
-	D04.615.723.795.656 Nalorphine
-	D04.615.723.795.706 Naloxone
-	D04.615.723.795.706.275 Buprenorphine, Naloxone Drug Combination
-	D04.615.723.795.706.550 Naltrexone
-	D04.615.799 Pyrenes
-	D04.615.799.306 Benzopyrenes



## MeSH Tree Changes for 2017

Type	Tree - heading
-	D04.615.799.306.300 Benzo(a)pyrene
-	D04.615.799.306.400 Dihydroxydihydrobenzopyrenes
-	D04.615.799.306.400.350 7,8-Dihydro-7,8-dihydroxybenzo(a)pyrene 9,10-oxide
-	D04.711 Spiro Compounds
-	D04.711.120 Buspirone
-	D04.711.345 Fluorescamine
-	D04.711.347 Fluoresceins
-	D04.711.347.300 Eosine I Bluish
-	D04.711.347.325 Eosine Yellowish-(YS)
-	D04.711.347.350 Erythrosine
-	D04.711.347.390 Fluorescein
-	D04.711.347.400 Fluorescein-5-isothiocyanate
-	D04.711.347.700 Rose Bengal
-	D04.711.350 Fluspirilene
-	D04.711.511 Leucogenenol
-	D04.711.656 Prospidium
-	D04.711.800 Spiperone
Old Tree	<b>D04.808 Steroids</b>
Old Tree	<b>D04.808.054 Androstanes</b>
Old Tree	<b>D04.808.054.040 Androstanols</b>
Old Tree	<b>D04.808.054.040.080 Androstane-3,17-diol</b>
Old Tree	<b>D04.808.054.040.129 Androsterone</b>
Old Tree	<b>D04.808.054.040.248 Dihydrotestosterone</b>
Old Tree	<b>D04.808.054.040.248.450 Mesterolone</b>
Old Tree	<b>D04.808.054.040.368 Etiocholanolone</b>
Old Tree	<b>D04.808.054.040.581 Oxandrolone</b>
Old Tree	<b>D04.808.054.040.632 Oxymetholone</b>
Old Tree	<b>D04.808.054.040.685 Pancuronium</b>
Old Tree	<b>D04.808.054.040.881 Stanozolol</b>
Old Tree	<b>D04.808.054.040.920 Vecuronium Bromide</b>
Old Tree	<b>D04.808.054.079 Androstenes</b>
Old Tree	<b>D04.808.054.079.065 Abiraterone Acetate</b>
Old Tree	<b>D04.808.054.079.129 Androstadienes</b>
Old Tree	<b>D04.808.054.079.129.114 Fluticasone</b>
Old Tree	<b>D04.808.054.079.129.114.500 Fluticasone Propionate, Salmeterol Xinafoate</b>

## MeSH Tree Changes for 2017

Type	Tree - heading
	<b>Drug Combination</b>
Old Tree	D04.808.054.079.129.284 Loteprednol Etabonate
Old Tree	D04.808.054.079.129.453 Methandrostenolone
Old Tree	D04.808.054.079.129.782 Testolactone
Old Tree	D04.808.054.079.229 Androstatrienes
Old Tree	D04.808.054.079.329 Androstenedione
Old Tree	D04.808.054.079.429 Androstenols
Old Tree	D04.808.054.079.429.154 Androstenediols
Old Tree	D04.808.054.079.429.154.050 Androstenediol
Old Tree	D04.808.054.079.429.154.349 Fluoxymesterone
Old Tree	D04.808.054.079.429.154.600 Methandriol
Old Tree	D04.808.054.079.429.320 Cyanoketone
Old Tree	D04.808.054.079.429.625 Dehydroepiandrosterone
Old Tree	D04.808.054.079.429.625.300 Dehydroepiandrosterone Sulfate
Old Tree	D04.808.054.079.429.824 Testosterone
Old Tree	D04.808.054.079.429.824.275 Epitestosterone
Old Tree	D04.808.054.079.429.824.375 Hydroxytestosterones
Old Tree	D04.808.054.079.429.824.537 Methenolone
Old Tree	D04.808.054.079.429.824.664 Methyltestosterone
Old Tree	D04.808.054.079.429.824.832 Testosterone Propionate
Old Tree	D04.808.054.079.500 Finasteride
Old Tree	D04.808.105 Bile Acids and Salts
Old Tree	D04.808.105.225 Cholic Acids
Old Tree	D04.808.105.225.130 Cholic Acid
Old Tree	D04.808.105.225.130.330 Cholates
Old Tree	D04.808.105.225.130.330.850 Sodium Cholate
Old Tree	D04.808.105.225.261 Dehydrocholic Acid
Old Tree	D04.808.105.225.272 Deoxycholic Acid
Old Tree	D04.808.105.225.272.150 Chenodeoxycholic Acid
Old Tree	D04.808.105.225.272.150.350 Glycochenodeoxycholic Acid
Old Tree	D04.808.105.225.272.150.850 Taurochenodeoxycholic Acid
Old Tree	D04.808.105.225.272.411 Glycodeoxycholic Acid
Old Tree	D04.808.105.225.272.411.360 Glycochenodeoxycholic Acid
Old Tree	D04.808.105.225.272.925 Taurodeoxycholic Acid
Old Tree	D04.808.105.225.272.925.875 Taurochenodeoxycholic Acid
Old Tree	D04.808.105.225.272.962 Ursodeoxycholic Acid

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	D04.808.105.225.400 Glycocholic Acid
Old Tree	D04.808.105.225.400.380 Glycodeoxycholic Acid
Old Tree	D04.808.105.225.400.380.360 Glycochenodeoxycholic Acid
Old Tree	D04.808.105.225.480 Lithocholic Acid
Old Tree	D04.808.105.225.480.880 Taurolithocholic Acid
Old Tree	D04.808.105.225.900 Taurocholic Acid
Old Tree	D04.808.105.225.900.900 Taurodeoxycholic Acid
Old Tree	D04.808.105.225.900.900.870 Taurochenodeoxycholic Acid
Old Tree	D04.808.105.225.900.920 Taurolithocholic Acid
Old Tree	D04.808.155 Cardanolides
Old Tree	D04.808.155.580 Cardiac Glycosides
Old Tree	D04.808.155.580.064 Bufanolides
Old Tree	D04.808.155.580.064.580 Proscillaridin
Old Tree	D04.808.155.580.130 Cardenolides
Old Tree	D04.808.155.580.130.500 Digitalis Glycosides
Old Tree	D04.808.155.580.130.500.236 Digitonin
Old Tree	D04.808.155.580.130.500.336 Digitoxin
Old Tree	D04.808.155.580.130.500.336.259 Acetyldigitoxins
Old Tree	D04.808.155.580.130.500.436 Digoxin
Old Tree	D04.808.155.580.130.500.436.050 Acetyldigoxins
Old Tree	D04.808.155.580.130.500.436.500 Medigoxin
Old Tree	D04.808.155.580.130.500.657 Lanatosides
Old Tree	D04.808.155.580.130.500.657.200 Deslanoside
Old Tree	D04.808.155.580.130.625 Digitoxin
Old Tree	D04.808.155.580.130.625.259 Acetyldigitoxins
Old Tree	D04.808.155.580.130.625.350 Digitoxigenin
Old Tree	D04.808.155.580.130.688 Digoxin
Old Tree	D04.808.155.580.130.688.050 Acetyldigoxins
Old Tree	D04.808.155.580.130.688.350 Digoxigenin
Old Tree	D04.808.155.580.130.688.500 Medigoxin
Old Tree	D04.808.155.580.130.750 Strophanthins
Old Tree	D04.808.155.580.130.750.250 Cymarine
Old Tree	D04.808.155.580.130.750.600 Ouabain
Old Tree	D04.808.155.580.130.750.800 Strophanthidin
Old Tree	D04.808.221 Cholanes
Old Tree	D04.808.221.263 Cholenes

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	D04.808.221.430 Cholic Acids
Old Tree	D04.808.221.430.130 Cholic Acid
Old Tree	D04.808.221.430.130.330 Cholates
Old Tree	D04.808.221.430.130.330.850 Sodium Cholate
Old Tree	D04.808.221.430.265 Dehydrocholic Acid
Old Tree	D04.808.221.430.342 Deoxycholic Acid
Old Tree	D04.808.221.430.342.300 Chenodeoxycholic Acid
Old Tree	D04.808.221.430.342.300.400 Glycochenodeoxycholic Acid
Old Tree	D04.808.221.430.342.300.900 Taurochenodeoxycholic Acid
Old Tree	D04.808.221.430.342.400 Glycodeoxycholic Acid
Old Tree	D04.808.221.430.342.400.450 Glycochenodeoxycholic Acid
Old Tree	D04.808.221.430.342.900 Taurodeoxycholic Acid
Old Tree	D04.808.221.430.342.900.910 Taurochenodeoxycholic Acid
Old Tree	D04.808.221.430.342.925 Ursodeoxycholic Acid
Old Tree	D04.808.221.430.484 Glycocholic Acid
Old Tree	D04.808.221.430.484.430 Glycodeoxycholic Acid
Old Tree	D04.808.221.430.484.430.420 Glycochenodeoxycholic Acid
Old Tree	D04.808.221.430.622 Lithocholic Acid
Old Tree	D04.808.221.430.622.900 Taurolithocholic Acid
Old Tree	D04.808.221.430.873 Taurocholic Acid
Old Tree	D04.808.221.430.873.920 Taurodeoxycholic Acid
Old Tree	D04.808.221.430.873.920.900 Taurochenodeoxycholic Acid
Old Tree	D04.808.221.430.873.940 Taurolithocholic Acid
Old Tree	D04.808.247 Cholestanes
Old Tree	D04.808.247.100 Cholestanols
Old Tree	D04.808.247.100.250 Cholestanol
Old Tree	D04.808.247.125 Cholestanones
Old Tree	D04.808.247.222 Cholestenes
Old Tree	D04.808.247.222.159 Cholecalciferol
Old Tree	D04.808.247.222.159.478 Hydroxycholecalciferols
Old Tree	D04.808.247.222.159.478.250 Calcifediol
Old Tree	D04.808.247.222.159.478.387 Dihydroxycholecalciferols
Old Tree	D04.808.247.222.159.478.387.300 Calcitriol
Old Tree	D04.808.247.222.159.478.387.400 24,25-Dihydroxyvitamin D 3
Old Tree	D04.808.247.222.222 Cholestadienes
Old Tree	D04.808.247.222.222.347 Cholestadienols

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	D04.808.247.222.222.347.200 Dehydrocholesterols
Old Tree	D04.808.247.222.222.347.231 Desmosterol
Old Tree	D04.808.247.222.222.347.389 Fusidic Acid
Old Tree	D04.808.247.222.222.347.557 Lanosterol
Old Tree	D04.808.247.222.222.347.833 Stigmasterol
Old Tree	D04.808.247.222.265 Cholestenones
Old Tree	D04.808.247.222.265.165 Ecdysteroids
Old Tree	D04.808.247.222.265.165.500 Ecdysone
Old Tree	D04.808.247.222.265.165.750 Ecdysterone
Old Tree	D04.808.247.222.265.450 Ketocholesterols
Old Tree	D04.808.247.222.284 Cholesterol
Old Tree	D04.808.247.222.284.070 Azacosterol
Old Tree	D04.808.247.222.284.200 Cholesterol Esters
Old Tree	D04.808.247.222.284.478 Hydroxycholesterols
Old Tree	D04.808.247.222.284.510 19-Iodocholesterol
Old Tree	D04.808.247.222.284.600 Ketocholesterols
Old Tree	D04.808.247.222.474 Ergocalciferols
Old Tree	D04.808.247.222.474.250 Dihydrotachysterol
Old Tree	D04.808.247.222.474.400 25-Hydroxyvitamin D 2
Old Tree	D04.808.247.222.537 Ergosterol
Old Tree	D04.808.247.222.537.888 Withanolides
Old Tree	D04.808.247.222.857 Sitosterols
Old Tree	D04.808.247.515 Spirostans
Old Tree	D04.808.247.515.500 Diosgenin
Old Tree	D04.808.247.515.500.500 Solanine
Old Tree	D04.808.247.808 Sterols
Old Tree	D04.808.247.808.050 Adosterol
Old Tree	D04.808.247.808.146 Cholecalciferol
Old Tree	D04.808.247.808.146.478 Hydroxycholecalciferols
Old Tree	D04.808.247.808.146.478.250 Calcifediol
Old Tree	D04.808.247.808.146.478.387 Dihydroxycholecalciferols
Old Tree	D04.808.247.808.146.478.387.300 Calcitriol
Old Tree	D04.808.247.808.146.478.387.400 24,25-Dihydroxyvitamin D 3
Old Tree	D04.808.247.808.197 Cholesterol
Old Tree	D04.808.247.808.197.070 Azacosterol
Old Tree	D04.808.247.808.197.135 Cholestanol

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	D04.808.247.808.197.200 Cholesterol Esters
Old Tree	D04.808.247.808.197.225 Cholesterol, Dietary
Old Tree	D04.808.247.808.197.238 Cholesterol, HDL
Old Tree	D04.808.247.808.197.244 Cholesterol, LDL
Old Tree	D04.808.247.808.197.247 Cholesterol, VLDL
Old Tree	D04.808.247.808.197.250 Dehydrocholesterols
Old Tree	D04.808.247.808.197.250.281 Desmosterol
Old Tree	D04.808.247.808.197.561 Hydroxycholesterols
Old Tree	D04.808.247.808.197.580 19-Iodocholesterol
Old Tree	D04.808.247.808.197.600 Ketocholesterols
Old Tree	D04.808.247.808.412 Ergocalciferols
Old Tree	D04.808.247.808.412.250 Dihydrotachysterol
Old Tree	D04.808.247.808.412.400 25-Hydroxyvitamin D 2
Old Tree	D04.808.247.808.489 Fusidic Acid
Old Tree	D04.808.247.808.607 Lanosterol
Old Tree	D04.808.247.808.756 Phytosterols
Old Tree	D04.808.247.808.756.071 Brassinosteroids
Old Tree	D04.808.247.808.756.143 Ecdysteroids
Old Tree	D04.808.247.808.756.287 Ergosterol
Old Tree	D04.808.247.808.756.287.888 Withanolides
Old Tree	D04.808.247.808.756.669 Sitosterols
Old Tree	D04.808.247.808.756.808 Stigmasterol
Old Tree	D04.808.247.808.874 Solanine
Old Tree	D04.808.294 Cyclosteroids
Old Tree	D04.808.365 Estranes
Old Tree	D04.808.365.415 Estrenes
Old Tree	D04.808.365.415.050 Allylestrenol
Old Tree	D04.808.365.415.200 Epimestrol
Old Tree	D04.808.365.415.215 Equilenin
Old Tree	D04.808.365.415.220 Equilin
Old Tree	D04.808.365.415.248 Estradiol
Old Tree	D04.808.365.415.248.320 Estramustine
Old Tree	D04.808.365.415.331 Estriol
Old Tree	D04.808.365.415.331.320 Estetrol
Old Tree	D04.808.365.415.414 Estrone
Old Tree	D04.808.365.415.414.400 Hydroxyestrones

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	D04.808.365.415.550                      Metribolone
Old Tree	D04.808.365.415.580                      Mifepristone
Old Tree	D04.808.365.415.638                      Nandrolone
Old Tree	D04.808.365.415.930                      Trenbolone Acetate
Old Tree	D04.808.451                                  Gonanes
Old Tree	D04.808.496                                  Homosteroids
Old Tree	D04.808.496.699                              Testolactone
Old Tree	D04.808.528                                  Hydroxysteroids
Old Tree	D04.808.578                                  Ketosteroids
Old Tree	D04.808.578.502                              17-Ketosteroids
Old Tree	D04.808.578.502.112                          Androstenedione
Old Tree	D04.808.578.502.195                          Androsterone
Old Tree	D04.808.578.502.400                          Dehydroepiandrosterone
Old Tree	D04.808.578.502.400.300                      Dehydroepiandrosterone Sulfate
Old Tree	D04.808.578.502.470                          Equilenin
Old Tree	D04.808.578.502.475                          Equilin
Old Tree	D04.808.578.502.497                          Estrone
Old Tree	D04.808.578.502.497.400                      Hydroxyestrones
Old Tree	D04.808.578.502.583                          Etiocholanolone
Old Tree	D04.808.668                                  Norsteroids
Old Tree	D04.808.668.100                              Norandrostanes
Old Tree	D04.808.668.651                              Norpregnanes
Old Tree	D04.808.668.651.443                          Norpregnadienes
Old Tree	D04.808.668.651.443.680                      Promegestone
Old Tree	D04.808.668.651.568                          Norpregnatrienes
Old Tree	D04.808.668.651.568.291                      Ethinyl Estradiol
Old Tree	D04.808.668.651.568.291.250                  Ethinyl Estradiol-Norgestrel Combination
Old Tree	D04.808.668.651.568.291.500                  Mestranol
Old Tree	D04.808.668.651.568.291.750                  Quinestrol
Old Tree	D04.808.668.651.568.620                      Norgestrienone
Old Tree	D04.808.668.651.568.620.400                  Gestrinone
Old Tree	D04.808.668.651.693                          Norpregnenes
Old Tree	D04.808.668.651.693.175                      Desogestrel
Old Tree	D04.808.668.651.693.223                      Ethylestrenol
Old Tree	D04.808.668.651.693.279                      Ethynodiol Diacetate
Old Tree	D04.808.668.651.693.362                      Gestonorone Caproate

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	D04.808.668.651.693.494      Lynestrenol
Old Tree	D04.808.668.651.693.595      Norethandrolone
Old Tree	D04.808.668.651.693.651      Norethindrone
Old Tree	D04.808.668.651.693.706      Norethynodrel
Old Tree	D04.808.668.651.693.762      Norgestrel
Old Tree	D04.808.668.651.693.762.225      Ethinyl Estradiol-Norgestrel Combination
Old Tree	D04.808.668.651.693.762.450      Levonorgestrel
Old Tree	D04.808.668.651.693.820      Norprogesterones
Old Tree	D04.808.745      Pregnanes
Old Tree	D04.808.745.432      Pregnadienes
Old Tree	D04.808.745.432.100      Canrenoic Acid
Old Tree	D04.808.745.432.120      Canrenone
Old Tree	D04.808.745.432.144      Chlormadinone Acetate
Old Tree	D04.808.745.432.219      Cyproterone
Old Tree	D04.808.745.432.219.150      Cyproterone Acetate
Old Tree	D04.808.745.432.235      Danazol
Old Tree	D04.808.745.432.296      Dydrogesterone
Old Tree	D04.808.745.432.370      Fluocinolone Acetonide
Old Tree	D04.808.745.432.370.325      Fluocinonide
Old Tree	D04.808.745.432.481      Medrogestone
Old Tree	D04.808.745.432.531      Megestrol
Old Tree	D04.808.745.432.531.500      Megestrol Acetate
Old Tree	D04.808.745.432.581      Melengestrol Acetate
Old Tree	D04.808.745.432.719      Pregnadienediols
Old Tree	D04.808.745.432.719.260      Desoximetasone
Old Tree	D04.808.745.432.719.320      Fluocortolone
Old Tree	D04.808.745.432.719.320.250      Diflucortolone
Old Tree	D04.808.745.432.719.349      Fluorometholone
Old Tree	D04.808.745.432.719.526      Mometasone Furoate
Old Tree	D04.808.745.432.719.526.500      Mometasone Furoate, Formoterol Fumarate Drug Combination
Old Tree	D04.808.745.432.719.702      Prednisone
Old Tree	D04.808.745.432.769      Pregnadienetriols
Old Tree	D04.808.745.432.769.125      Beclomethasone
Old Tree	D04.808.745.432.769.199      Betamethasone
Old Tree	D04.808.745.432.769.199.150      Betamethasone Valerate



## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	D04.808.745.432.769.325      Desonide
Old Tree	D04.808.745.432.769.344      Dexamethasone
Old Tree	D04.808.745.432.769.344.300      Dexamethasone Isonicotinate
Old Tree	D04.808.745.432.769.465      Flumethasone
Old Tree	D04.808.745.432.769.692      Paramethasone
Old Tree	D04.808.745.432.769.795      Prednisolone
Old Tree	D04.808.745.432.769.795.307      Fluprednisolone
Old Tree	D04.808.745.432.769.795.539      Methylprednisolone
Old Tree	D04.808.745.432.769.795.539.500      Methylprednisolone Hemisuccinate
Old Tree	D04.808.745.432.769.795.700      Prednimustine
Old Tree	D04.808.745.432.915      Triamcinolone
Old Tree	D04.808.745.432.915.715      Triamcinolone Acetonide
Old Tree	D04.808.745.495      Pregnanediol
Old Tree	D04.808.745.558      Pregnanediones
Old Tree	D04.808.745.558.050      Alfaxalone Alfadolone Mixture
Old Tree	D04.808.745.558.075      5-alpha-Dihydroprogesterone
Old Tree	D04.808.745.558.783      Tetrahydrocortisone
Old Tree	D04.808.745.620      Pregnanetriol
Old Tree	D04.808.745.640      Pregnanolone
Old Tree	D04.808.745.683      Pregnatrienes
Old Tree	D04.808.745.745      Pregnenes
Old Tree	D04.808.745.745.183      Cortisone
Old Tree	D04.808.745.745.279      Dimethisterone
Old Tree	D04.808.745.745.379      Ethisterone
Old Tree	D04.808.745.745.654      Pregnenediones
Old Tree	D04.808.745.745.654.062      Aldosterone
Old Tree	D04.808.745.745.654.105      Budesonide
Old Tree	D04.808.745.745.654.105.500 Combination      Budesonide, Formoterol Fumarate Drug
Old Tree	D04.808.745.745.654.237      Corticosterone
Old Tree	D04.808.745.745.654.237.400      18-Hydroxycorticosterone
Old Tree	D04.808.745.745.654.252      Cortodoxone
Old Tree	D04.808.745.745.654.339      Desoxycorticosterone
Old Tree	D04.808.745.745.654.339.400      18-Hydroxydesoxycorticosterone
Old Tree	D04.808.745.745.654.339.700      Desoxycorticosterone Acetate
Old Tree	D04.808.745.745.654.473      Flurandrenolone

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	D04.808.745.745.654.485 Flurogestone Acetate
Old Tree	D04.808.745.745.654.527 Gestonorone Caproate
Old Tree	D04.808.745.745.654.570 Halcinonide
Old Tree	D04.808.745.745.654.600 Hydrocortisone
Old Tree	D04.808.745.745.654.600.500 Fludrocortisone
Old Tree	D04.808.745.745.654.829 Progesterone
Old Tree	D04.808.745.745.654.829.025 Algestone
Old Tree	D04.808.745.745.654.829.025.025 Algestone Acetophenide
Old Tree	D04.808.745.745.654.829.100 20-alpha-Dihydroprogesterone
Old Tree	D04.808.745.745.654.829.395 Hydroxyprogesterones
Old Tree	D04.808.745.745.654.829.395.400 17-alpha-Hydroxyprogesterone
Old Tree	D04.808.745.745.654.829.395.700 Medroxyprogesterone
Old Tree	D04.808.745.745.654.829.395.700.500 Medroxyprogesterone Acetate
Old Tree	D04.808.745.745.725 Pregnenolone
Old Tree	D04.808.745.745.725.395 17-alpha-Hydroxypregnenolone
Old Tree	D04.808.745.745.725.703 Pregnenolone Carbonitrile
Old Tree	D04.808.745.745.855 Spironolactone
Old Tree	D04.808.745.887 Tetrahydrocortisol
Old Tree	D04.808.795 Sapogenins
Old Tree	D04.808.812 Secosteroids
Old Tree	D04.808.812.768 Vitamin D
Old Tree	D04.808.812.768.196 Cholecalciferol
Old Tree	D04.808.812.768.196.478 Hydroxycholecalciferols
Old Tree	D04.808.812.768.196.478.250 Calcifediol
Old Tree	D04.808.812.768.196.478.387 Dihydroxycholecalciferols
Old Tree	D04.808.812.768.196.478.387.300 Calcitriol
Old Tree	D04.808.812.768.196.478.387.400 24,25-Dihydroxyvitamin D 3
Old Tree	D04.808.812.768.462 Ergocalciferols
Old Tree	D04.808.812.768.462.250 Dihydrotachysterol
Old Tree	D04.808.812.768.462.400 25-Hydroxyvitamin D 2
Old Tree	D04.808.875 Steroids, Brominated
Old Tree	D04.808.883 Steroids, Chlorinated
Old Tree	D04.808.883.154 Beclomethasone
Old Tree	D04.808.883.294 Chlormadinone Acetate
Old Tree	D04.808.883.419 Cyproterone
Old Tree	D04.808.883.419.150 Cyproterone Acetate

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	D04.808.908 Steroids, Fluorinated
Old Tree	D04.808.908.093 Betamethasone
Old Tree	D04.808.908.093.150 Betamethasone Valerate
Old Tree	D04.808.908.093.250 Clobetasol
Old Tree	D04.808.908.238 Dexamethasone
Old Tree	D04.808.908.238.250 Desoximetasone
Old Tree	D04.808.908.238.300 Dexamethasone Isonicotinate
Old Tree	D04.808.908.359 Flumethasone
Old Tree	D04.808.908.394 Fluocinolone Acetonide
Old Tree	D04.808.908.394.300 Fluocinonide
Old Tree	D04.808.908.405 Fluocortolone
Old Tree	D04.808.908.405.250 Difluocortolone
Old Tree	D04.808.908.431 Fluorometholone
Old Tree	D04.808.908.466 Fluoxymesterone
Old Tree	D04.808.908.502 Fluprednisolone
Old Tree	D04.808.908.537 Flurandrenolone
Old Tree	D04.808.908.550 Flurogestone Acetate
Old Tree	D04.808.908.727 Paramethasone
Old Tree	D04.808.908.891 Triamcinolone
Old Tree	D04.808.908.891.927 Triamcinolone Acetonide
Old Tree	D04.808.925 Steroids, Heterocyclic
Old Tree	D04.808.925.100 Azasteroids
Old Tree	D04.808.925.100.125 Dutasteride
Old Tree	D04.808.925.100.250 Finasteride
-	D05 Macromolecular Substances
-	D05.374 Micelles
-	D05.500 Multiprotein Complexes
-	D05.500.049 Amyloid
-	D05.500.099 Apoptosomes
-	D05.500.117 Axin Signaling Complex
-	D05.500.117.249 Adenomatous Polyposis Coli Protein
-	D05.500.117.500 Axin Protein
-	D05.500.117.750 Casein Kinase I
-	D05.500.117.750.100 Casein Kinase Ialpha
-	D05.500.117.750.200 Casein Kinase Idelta
-	D05.500.117.750.300 Casein Kinase Iepsilon

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D05.500.117.875	Glycogen Synthase Kinase 3
New Heading	<b>D05.500.117.875.500</b>	<b>Glycogen Synthase Kinase 3 beta</b>
-	D05.500.135	Calcifying Nanoparticles
New Heading	<b>D05.500.142</b>	<b>Dynactin Complex</b>
-	D05.500.149	Dystrophin-Associated Protein Complex
-	D05.500.199	Endosomal Sorting Complexes Required for Transport
-	D05.500.224	Inflammasomes
-	D05.500.299	Lactose Synthase
-	D05.500.374	Mediator Complex
-	D05.500.374.249	Cyclin C
-	D05.500.374.500	Cyclin-Dependent Kinase 8
-	D05.500.374.750	Mediator Complex Subunit 1
-	D05.500.500	Molecular Motor Proteins
-	D05.500.562	Multienzyme Complexes
-	D05.500.562.249	Electron Transport Complex I
-	D05.500.562.374	Electron Transport Complex IV
-	D05.500.562.405	Exosome Multienzyme Ribonuclease Complex
-	D05.500.562.444	Fatty Acid Synthase, Type II
-	D05.500.562.452	Glycine Decarboxylase Complex
-	D05.500.562.452.100	Aminomethyltransferase
-	D05.500.562.452.150	Dihydrolipoamide Dehydrogenase
-	D05.500.562.452.175	Glycine Decarboxylase Complex H-Protein
-	D05.500.562.452.200	Glycine Dehydrogenase (Decarboxylating)
-	D05.500.562.468	Ketoglutarate Dehydrogenase Complex
-	D05.500.562.468.500	Dihydrolipoamide Dehydrogenase
-	D05.500.562.484	Phosphoenolpyruvate Sugar Phosphotransferase System
-	D05.500.562.488	Photosynthetic Reaction Center Complex Proteins
-	D05.500.562.488.374	Cytochrome b6f Complex
-	D05.500.562.488.374.500	Cytochromes b6
-	D05.500.562.488.374.750	Cytochromes f
-	D05.500.562.488.374.875	Plastoquinol-Plastocyanin Reductase
-	D05.500.562.488.490	Light-Harvesting Protein Complexes
-	D05.500.562.488.490.249	Chlorophyll Binding Proteins
-	D05.500.562.488.490.500	Phycobilisomes

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D05.500.562.488.490.500.500 Phycobiliproteins
-	D05.500.562.488.490.500.500.755 Phycocyanin
-	D05.500.562.488.490.500.500.777 Phycoerythrin
-	D05.500.562.488.500 Photosystem I Protein Complex
-	D05.500.562.488.750 Photosystem II Protein Complex
-	D05.500.562.500 Proteasome Endopeptidase Complex
-	D05.500.562.625 Pyruvate Dehydrogenase Complex
-	D05.500.562.625.500 Dihydrolipoamide Dehydrogenase
-	D05.500.562.625.750 Dihydrolipoyllysine-Residue Acetyltransferase
-	D05.500.562.625.875 Pyruvate Dehydrogenase (Lipoamide)
-	D05.500.562.750 Succinate Cytochrome c Oxidoreductase
-	D05.500.562.750.249 Electron Transport Complex II
-	D05.500.562.750.249.500 Succinate Dehydrogenase
-	D05.500.562.750.500 Electron Transport Complex III
-	D05.500.562.875 Sucrase-Isomaltase Complex
-	D05.500.781 Polycomb-Group Proteins
-	D05.500.781.100 Polycomb Repressive Complex 1
-	D05.500.781.750 Polycomb Repressive Complex 2
New Heading	<b>D05.500.781.750.250 Enhancer of Zeste Homolog 2 Protein</b>
-	D05.500.781.750.500 Retinoblastoma-Binding Protein 4
-	D05.500.781.750.750 Retinoblastoma-Binding Protein 7
-	D05.500.890 Protein Translocation Systems
-	D05.500.890.500 Bacterial Secretion Systems
-	D05.500.890.500.500 Type I Secretion Systems
-	D05.500.890.500.750 Type II Secretion Systems
-	D05.500.890.500.875 Type III Secretion Systems
-	D05.500.890.500.937 Type IV Secretion Systems
-	D05.500.890.500.968 Type V Secretion Systems
-	D05.500.890.500.984 Type VI Secretion Systems
-	D05.500.890.500.992 Type VII Secretion Systems
New Heading	<b>D05.500.890.625 SEC Translocation Channels</b>
-	D05.500.890.750 Twin-Arginine-Translocation System
New Tree	<a href="#">D05.500.945</a> <a href="#">Troponin</a>
New	<a href="#">D05.500.945.900</a> <a href="#">Troponin C</a>

## MeSH Tree Changes for 2017

Type	Tree - heading	
Tree		
New Tree	<a href="#">D05.500.945.925</a>	Troponin I
New Tree	<a href="#">D05.500.945.962</a>	Troponin T
-	D05.750	Polymers
-	D05.750.078	Biopolymers
-	D05.750.078.139	Chitin
-	D05.750.078.139.500	Chitosan
-	D05.750.078.280	Collagen
-	D05.750.078.280.300	Fibrillar Collagens
-	D05.750.078.280.300.100	Collagen Type I
-	D05.750.078.280.300.200	Collagen Type II
-	D05.750.078.280.300.300	Collagen Type III
-	D05.750.078.280.300.400	Collagen Type V
-	D05.750.078.280.300.500	Collagen Type XI
-	D05.750.078.421	Elastin
-	D05.750.078.562	Glucans
-	D05.750.078.562.180	Cellulose
-	D05.750.078.562.180.200	Cellulose, Oxidized
-	D05.750.078.562.180.357	Hypromellose Derivatives
-	D05.750.078.562.180.515	Lignin
-	D05.750.078.562.272	Dextrans
-	D05.750.078.562.388	Glycogen
-	D05.750.078.562.388.518	Liver Glycogen
-	D05.750.078.562.855	Starch
-	D05.750.078.562.855.750	Inulin
-	D05.750.078.593	Intermediate Filament Proteins
-	D05.750.078.593.200	Desmin
-	D05.750.078.593.400	Glial Fibrillary Acidic Protein
-	D05.750.078.593.450	Keratins
-	D05.750.078.593.450.074	beta-Keratins
-	D05.750.078.593.450.149	Keratins, Hair-Specific
-	D05.750.078.593.450.300	Keratins, Type I
-	D05.750.078.593.450.300.100	Keratin-10
-	D05.750.078.593.450.300.200	Keratin-12

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D05.750.078.593.450.300.300 Keratin-13
-	D05.750.078.593.450.300.400 Keratin-14
-	D05.750.078.593.450.300.600 Keratin-16
-	D05.750.078.593.450.300.700 Keratin-17
-	D05.750.078.593.450.300.800 Keratin-18
-	D05.750.078.593.450.300.900 Keratin-19
-	D05.750.078.593.450.300.950 Keratin-20
-	D05.750.078.593.450.600 Keratins, Type II
-	D05.750.078.593.450.600.100 Keratin-1
-	D05.750.078.593.450.600.200 Keratin-2
-	D05.750.078.593.450.600.300 Keratin-3
-	D05.750.078.593.450.600.400 Keratin-4
-	D05.750.078.593.450.600.500 Keratin-5
-	D05.750.078.593.450.600.600 Keratin-6
-	D05.750.078.593.450.600.700 Keratin-7
-	D05.750.078.593.450.600.800 Keratin-8
-	D05.750.078.593.450.600.900 Keratin-9
-	D05.750.078.593.540 Nestin
-	D05.750.078.593.630 Neurofilament Proteins
-	D05.750.078.593.765 Peripherins
-	D05.750.078.593.900 Vimentin
-	D05.750.078.625 Latex
-	D05.750.078.687 Lignin
-	D05.750.078.730 Microfilament Proteins
-	D05.750.078.730.032 Actin Capping Proteins
-	D05.750.078.730.032.500 CapZ Actin Capping Protein
-	D05.750.078.730.032.750 Tropomodulin
-	D05.750.078.730.212 Actin Depolymerizing Factors
-	D05.750.078.730.212.500 Cofilin 1
-	D05.750.078.730.212.750 Cofilin 2
-	D05.750.078.730.212.875 Destrin
-	D05.750.078.730.246 Actin-Related Protein 2-3 Complex
-	D05.750.078.730.246.500 Actin-Related Protein 2
-	D05.750.078.730.246.750 Actin-Related Protein 3
-	D05.750.078.730.248 Actinin
-	D05.750.078.730.250 Actins

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D05.750.078.730.281 Cortactin
-	D05.750.078.730.315 Filamins
-	D05.750.078.730.350 Gelsolin
-	D05.750.078.730.475 Myosins
-	D05.750.078.730.475.100 Myosin Heavy Chains
-	D05.750.078.730.475.200 Myosin Light Chains
-	D05.750.078.730.475.300 Myosin Subfragments
-	D05.750.078.730.475.470 Myosin Type I
-	D05.750.078.730.475.475 Myosin Type II
-	D05.750.078.730.475.475.124 Cardiac Myosins
-	D05.750.078.730.475.475.124.249 Atrial Myosins
-	D05.750.078.730.475.475.124.500 Ventricular Myosins
-	D05.750.078.730.475.475.249 Nonmuscle Myosin Type IIA
-	D05.750.078.730.475.475.500 Nonmuscle Myosin Type IIB
-	D05.750.078.730.475.475.750 Skeletal Muscle Myosins
-	D05.750.078.730.475.475.875 Smooth Muscle Myosins
-	D05.750.078.730.475.612 Myosin Type III
-	D05.750.078.730.475.681 Myosin Type IV
-	D05.750.078.730.475.750 Myosin Type V
-	D05.750.078.730.637 Profilins
New Heading	<b>D05.750.078.730.719 Tensins</b>
-	D05.750.078.730.800 Tropomyosin
-	D05.750.078.730.825 Troponin
-	D05.750.078.730.825.900 Troponin C
-	D05.750.078.730.825.925 Troponin I
-	D05.750.078.730.825.962 Troponin T
-	D05.750.078.730.912 Wiskott-Aldrich Syndrome Protein Family
-	D05.750.078.730.912.500 Wiskott-Aldrich Syndrome Protein
-	D05.750.078.730.912.550 Wiskott-Aldrich Syndrome Protein, Neuronal
-	D05.750.078.734 Microtubule Proteins
-	D05.750.078.734.800 Tubulin
-	D05.750.078.738 Pectins
-	D05.750.078.739 Plant Gums
-	D05.750.078.739.249 Chewing Gum
-	D05.750.078.739.500 Gum Arabic



## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D05.750.078.739.625	Karaya Gum
-	D05.750.078.739.750	Tragacanth
-	D05.750.078.764	Plant Mucilage
-	D05.750.078.789	Polyhydroxyalkanoates
-	D05.750.078.840	Resins, Plant
-	D05.750.078.840.109	Amber
-	D05.750.078.840.219	Balsams
-	D05.750.078.840.490	Frankincense
-	D05.750.078.840.762	Propolis
-	D05.750.078.850	Reticulin
-	D05.750.078.875	Silk
-	D05.750.078.875.500	Fibroins
-	D05.750.078.875.750	Sericins
-	D05.750.078.937	Tannins
-	D05.750.078.937.214	Hydrolyzable Tannins
-	D05.750.078.937.429	Proanthocyanidins
-	D05.750.200	Colestipol
New Heading	<b>D05.750.230</b>	<b>Polyelectrolytes</b>
-	D05.750.259	Cyanoacrylates
-	D05.750.259.341	Enbucrilate
-	D05.750.259.341.110	Bucrylate
-	D05.750.327	Dendrimers
-	D05.750.395	Fluorocarbon Polymers
-	D05.750.395.616	Polytetrafluoroethylene
-	D05.750.395.616.755	Proplast
-	D05.750.470	Hexadimethrine Bromide
-	D05.750.593	Organically Modified Ceramics
-	D05.750.716	Plastics
-	D05.750.716.195	Biodegradable Plastics
-	D05.750.716.392	Nylons
-	D05.750.716.507	Polyethylenes
-	D05.750.716.507.500	Polyethylene
-	D05.750.716.507.600	Polyethyleneimine
-	D05.750.716.550	Polypropylenes
-	D05.750.716.579	Polystyrenes

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D05.750.716.579.159 Cholestyramine Resin
-	D05.750.716.650 Polyurethanes
-	D05.750.716.721 Polyvinyls
-	D05.750.716.721.616 Polyvinyl Alcohol
-	D05.750.716.721.812 Polyvinyl Chloride
-	D05.750.716.721.838 Povidone
-	D05.750.716.721.838.745 Povidone-Iodine
-	D05.750.716.822 Resins, Synthetic
-	D05.750.716.822.111 Acrylic Resins
-	D05.750.716.822.111.650 Polymethacrylic Acids
-	D05.750.716.822.111.650.605 Methylmethacrylates
-	D05.750.716.822.111.650.605.450 Methylmethacrylate
-	D05.750.716.822.111.650.605.500 Polymethyl Methacrylate
-	D05.750.716.822.111.650.750 Polyhydroxyethyl Methacrylate
-	D05.750.716.822.300 Bone Cements
-	D05.750.716.822.308 Composite Resins
-	D05.750.716.822.308.200 Bisphenol A-Glycidyl Methacrylate
-	D05.750.716.822.308.300 Compomers
-	D05.750.716.822.461 Epoxy Resins
-	D05.750.716.822.461.500 Silorane Resins
-	D05.750.716.822.730 Resin Cements
-	D05.750.722 Polyanetholesulfonate
-	D05.750.725 Polyanhydrides
-	D05.750.728 Polyesters
-	D05.750.728.700 Polydioxanone
-	D05.750.728.764 Polyethylene Terephthalates
-	D05.750.728.772 Polyglactin 910
-	D05.750.728.780 Polyglycolic Acid
-	D05.750.728.890 Polyhydroxyalkanoates
-	D05.750.741 Polyethylene Glycols
-	D05.750.741.125 Certolizumab Pegol
-	D05.750.741.250 Cetomacrogol
-	D05.750.741.485 Hydrogel
-	D05.750.741.575 Nonoxynol
-	D05.750.741.610 Octoxynol
-	D05.750.741.650 Poloxalene

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D05.750.741.667 Poloxamer
-	D05.750.741.685 Polyhydroxyethyl Methacrylate
-	D05.750.741.700 Polysorbates
-	D05.750.780 Polygeline
-	D05.750.795 Polyphlorethin Phosphate
-	D05.750.830 Pyran Copolymer
-	D05.750.900 Siloxanes
-	D05.750.900.850 Silicones
-	D05.750.900.850.150 Dimethylpolysiloxanes
-	D05.750.900.850.150.750 Simethicone
-	D05.750.900.850.900 Silicone Elastomers
-	D05.750.900.850.905 Silicone Gels
-	D05.750.900.850.950 Silicone Oils
-	D05.750.900.925 Silorane Resins
-	D05.875 Protein Aggregates
-	D06 Hormones, Hormone Substitutes, and Hormone Antagonists
-	D06.347 Hormone Antagonists
-	D06.347.065 Androgen Antagonists
-	D06.347.065.249 Androgen Receptor Antagonists
-	D06.347.065.500 Nonsteroidal Anti-Androgens
-	D06.347.100 Antithyroid Agents
-	D06.347.230 Calcimimetic Agents
-	D06.347.295 Estrogen Antagonists
-	D06.347.295.500 Estrogen Receptor Antagonists
-	D06.347.360 Estrogen Receptor Modulators
-	D06.347.360.827 Selective Estrogen Receptor Modulators
-	D06.347.420 Insulin Antagonists
-	D06.347.565 Leukotriene Antagonists
-	D06.347.700 Mineralocorticoid Receptor Antagonists
-	D06.347.710 Prostaglandin Antagonists
-	D06.472 Hormones
-	D06.472.040 Adrenal Cortex Hormones
-	D06.472.040.502 17-Ketosteroids
-	D06.472.040.502.112 Androstenedione
-	D06.472.040.502.195 Androsterone
-	D06.472.040.502.400 Dehydroepiandrosterone

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D06.472.040.502.400.300 Dehydroepiandrosterone Sulfate
-	D06.472.040.502.497 Estrone
-	D06.472.040.502.583 Etiocholanolone
-	D06.472.040.543 Glucocorticoids
-	D06.472.040.585 Hydroxycorticosteroids
-	D06.472.040.585.353 11-Hydroxycorticosteroids
-	D06.472.040.585.353.118 Aldosterone
-	D06.472.040.585.353.237 Corticosterone
-	D06.472.040.585.353.476 Hydrocortisone
-	D06.472.040.585.353.500 18-Hydroxycorticosterone
-	D06.472.040.585.353.825 Tetrahydrocortisol
-	D06.472.040.585.478 17-Hydroxycorticosteroids
-	D06.472.040.585.478.195 Cortisone
-	D06.472.040.585.478.225 Cortodoxone
-	D06.472.040.585.478.392 Hydrocortisone
-	D06.472.040.585.478.782 Tetrahydrocortisol
-	D06.472.040.585.478.865 Tetrahydrocortisone
-	D06.472.040.585.611 Desoxycorticosterone
-	D06.472.040.585.611.400 18-Hydroxydesoxycorticosterone
-	D06.472.040.585.611.700 Desoxycorticosterone Acetate
-	D06.472.040.585.745 Pregnenolone
-	D06.472.040.585.745.500 17-alpha-Hydroxypregnenolone
-	D06.472.317 Gastrointestinal Hormones
-	D06.472.317.152 Cholecystokinin
-	D06.472.317.152.700 Sincalide
-	D06.472.317.350 Epidermal Growth Factor
-	D06.472.317.400 Gastric Inhibitory Polypeptide
-	D06.472.317.410 Gastrin-Releasing Peptide
-	D06.472.317.413 Gastrins
-	D06.472.317.413.593 Pentagastrin
-	D06.472.317.413.800 Tetragastrin
-	D06.472.317.469 Proglucagon
-	D06.472.317.469.249 Glicentin
-	D06.472.317.469.500 Glucagon-Like Peptides
-	D06.472.317.469.500.500 Glucagon-Like Peptide 1
-	D06.472.317.469.500.500.500 Liraglutide

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D06.472.317.469.500.750                      Glucagon-Like Peptide 2
-	D06.472.317.469.750                              Oxyntomodulin
-	D06.472.317.525                                  Motilin
-	D06.472.317.662                                  Peptide YY
-	D06.472.317.800                                  Secretin
-	D06.472.317.950                                  Vasoactive Intestinal Peptide
-	D06.472.334    Gonadal Hormones
-	D06.472.334.500                                  Activins
-	D06.472.334.500.500                              Inhibin-beta Subunits
-	D06.472.334.734                                  Corpus Luteum Hormones
-	D06.472.334.734.623                              Progesterone
-	D06.472.334.734.769                              Relaxin
-	D06.472.334.851                                  Gonadal Steroid Hormones
-	D06.472.334.851.437                              Estradiol Congeners
-	D06.472.334.851.437.249                          Equilenin
-	D06.472.334.851.437.374                          Equilin
-	D06.472.334.851.437.500                          Estradiol
-	D06.472.334.851.437.750                          Estriol
-	D06.472.334.851.437.750.320                      Estetrol
-	D06.472.334.851.437.968                          Estrogenic Steroids, Alkylated
-	D06.472.334.851.437.968.500                      Ethinyl Estradiol
-	D06.472.334.851.437.968.500.250                  Ethinyl Estradiol-Norgestrel Combination
-	D06.472.334.851.437.968.500.500                  Mestranol
-	D06.472.334.851.437.968.500.750                  Quinestrol
-	D06.472.334.851.437.984                          Estrogens, Catechol
-	D06.472.334.851.437.984.350                      Hydroxyestrones
-	D06.472.334.851.437.988                          Estrogens, Conjugated (USP)
-	D06.472.334.851.437.994                          Estrogens, Esterified (USP)
-	D06.472.334.851.437.996                          Estrone
-	D06.472.334.851.687                              Progesterone Congeners
-	D06.472.334.851.687.500                          Pregnenolone
-	D06.472.334.851.687.500.500                      17-alpha-Hydroxypregnenolone
-	D06.472.334.851.687.750                          Progesterone
-	D06.472.334.851.687.750.070                      5-alpha-Dihydroprogesterone
-	D06.472.334.851.687.750.074                      20-alpha-Dihydroprogesterone
-	D06.472.334.851.687.750.478                      Hydroxyprogesterones

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D06.472.334.851.687.750.478.400 17-alpha-Hydroxyprogesterone
-	D06.472.334.851.687.750.478.400.500 Pregnanetriol
-	D06.472.334.851.687.750.739 Pregnanediol
-	D06.472.334.851.968 Testosterone Congeners
-	D06.472.334.851.968.500 Androstane-3,17-diol
-	D06.472.334.851.968.750 Androstenediol
-	D06.472.334.851.968.875 Androstenedione
-	D06.472.334.851.968.937 Androsterone
-	D06.472.334.851.968.952 Dehydroepiandrosterone
-	D06.472.334.851.968.952.300 Dehydroepiandrosterone Sulfate
-	D06.472.334.851.968.964 Dihydrotestosterone
-	D06.472.334.851.968.968 Etiocholanolone
-	D06.472.334.851.968.976 Nandrolone
-	D06.472.334.851.968.984 Testosterone
-	D06.472.334.851.968.984.500 Epitestosterone
-	D06.472.334.851.968.984.750 Testosterone Propionate
-	D06.472.334.968 Inhibins
-	D06.472.334.968.500 Inhibin-beta Subunits
-	D06.472.334.984 Testicular Hormones
-	D06.472.334.984.500 Anti-Mullerian Hormone
-	D06.472.397 Hormones, Ectopic
-	D06.472.445 Invertebrate Hormones
-	D06.472.445.573 Insect Hormones
-	D06.472.445.573.271 Ecdysteroids
-	D06.472.445.573.271.500 Ecdysone
-	D06.472.445.573.271.750 Ecdysterone
-	D06.472.445.573.666 Juvenile Hormones
-	D06.472.445.573.666.170 Diflubenzuron
-	D06.472.445.573.666.500 Methoprene
-	D06.472.506 Melatonin
-	D06.472.699 Peptide Hormones
-	D06.472.699.009 Activins
-	D06.472.699.009.500 Inhibin-beta Subunits
-	D06.472.699.042 Adipokines
-	D06.472.699.042.249 Adiponectin
-	D06.472.699.042.500 Leptin

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D06.472.699.042.750 Resistin
-	D06.472.699.077 Adrenomedullin
-	D06.472.699.094 Angiotensins
-	D06.472.699.094.075 Angiotensin I
-	D06.472.699.094.078 Angiotensin II
-	D06.472.699.094.080 Angiotensin III
-	D06.472.699.100 Bombesin
-	D06.472.699.150 Calcitonin
-	D06.472.699.275 Gastric Inhibitory Polypeptide
-	D06.472.699.280 Gastrins
-	D06.472.699.301 Ghrelin
-	D06.472.699.322 Gonadotropins
-	D06.472.699.322.326 Chorionic Gonadotropin
-	D06.472.699.322.326.125 Chorionic Gonadotropin, beta Subunit, Human
-	D06.472.699.322.326.562 Glycoprotein Hormones, alpha Subunit
-	D06.472.699.322.451 Gonadotropins, Equine
-	D06.472.699.322.576 Gonadotropins, Pituitary
-	D06.472.699.322.576.288 Follicle Stimulating Hormone
-	D06.472.699.322.576.288.500 Follicle Stimulating Hormone, beta Subunit
-	D06.472.699.322.576.288.750 Glycoprotein Hormones, alpha Subunit
-	D06.472.699.322.576.463 Luteinizing Hormone
-	D06.472.699.322.576.463.249 Glycoprotein Hormones, alpha Subunit
-	D06.472.699.322.576.463.500 Luteinizing Hormone, beta Subunit
-	D06.472.699.322.576.583 Menotropins
-	D06.472.699.322.576.583.500 Urofollitropin
-	D06.472.699.322.576.773 Prolactin
-	D06.472.699.327 Hypothalamic Hormones
-	D06.472.699.327.700 Pituitary Hormone Release Inhibiting Hormones
-	D06.472.699.327.700.500 MSH Release-Inhibiting Hormone
-	D06.472.699.327.700.750 Prolactin Release-Inhibiting Factors
-	D06.472.699.327.700.875 Somatostatin
-	D06.472.699.327.700.875.500 Somatostatin-28
-	D06.472.699.327.740 Pituitary Hormone-Releasing Hormones
-	D06.472.699.327.740.140 Corticotropin-Releasing Hormone
-	D06.472.699.327.740.320 Gonadotropin-Releasing Hormone
-	D06.472.699.327.740.320.100 Buserelin

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D06.472.699.327.740.320.340 Goserelin
-	D06.472.699.327.740.320.400 Leuprolide
-	D06.472.699.327.740.320.580 Nafarelin
-	D06.472.699.327.740.320.790 Triptorelin Pamoate
-	D06.472.699.327.740.860 Growth Hormone-Releasing Hormone
-	D06.472.699.327.740.860.780 Sermorelin
-	D06.472.699.327.740.880 Thyrotropin-Releasing Hormone
-	D06.472.699.327.935 Pro-Opiomelanocortin
-	D06.472.699.327.935.119 alpha-Endorphin
-	D06.472.699.327.935.179 alpha-MSH
-	D06.472.699.327.935.239 beta-Endorphin
-	D06.472.699.327.935.480 beta-Lipotropin
-	D06.472.699.327.935.492 beta-MSH
-	D06.472.699.327.935.498 Corticotropin-Like Intermediate Lobe Peptide
-	D06.472.699.327.935.505 gamma-Endorphin
-	D06.472.699.327.935.518 gamma-Lipotropin
-	D06.472.699.327.935.524 gamma-MSH
-	D06.472.699.327.935.531 Melanocortins
-	D06.472.699.327.935.531.500 Adrenocorticotrophic Hormone
-	D06.472.699.327.935.531.500.200 Cosyntropin
-	D06.472.699.327.935.531.750 Melanocyte-Stimulating Hormones
-	D06.472.699.327.935.531.750.050 alpha-MSH
-	D06.472.699.327.935.531.750.075 beta-MSH
-	D06.472.699.327.935.531.750.115 gamma-MSH
-	D06.472.699.327.967 Prolactin-Releasing Hormone
-	D06.472.699.337 Inhibins
-	D06.472.699.337.500 Inhibin-beta Subunits
-	D06.472.699.500 Motilin
-	D06.472.699.584 Natriuretic Peptides
-	D06.472.699.584.500 Atrial Natriuretic Factor
-	D06.472.699.584.625 Natriuretic Peptide, Brain
-	D06.472.699.584.750 Natriuretic Peptide, C-Type
-	D06.472.699.587 Pancreatic Hormones
-	D06.472.699.587.200 Insulins
-	D06.472.699.587.200.200 Biphasic Insulins
-	D06.472.699.587.200.300 Insulin, Long-Acting



## MeSH Tree Changes for 2017

Type	Tree - heading
-	D06.472.699.587.200.300.050      Insulin Detemir
-	D06.472.699.587.200.300.100      Insulin Glargine
-	D06.472.699.587.200.300.200      Insulin, Isophane
-	D06.472.699.587.200.300.200.500      Isophane Insulin, Human
-	D06.472.699.587.200.300.300      Insulin, Lente
-	D06.472.699.587.200.300.800      Insulin, Ultralente
-	D06.472.699.587.200.400      Insulin, Short-Acting
-	D06.472.699.587.200.400.100      Insulin Aspart
-	D06.472.699.587.200.400.500      Insulin Lispro
-	D06.472.699.587.200.500      Proinsulin
-	D06.472.699.587.200.500.250      C-Peptide
-	D06.472.699.587.200.500.625      Insulin
-	D06.472.699.587.200.500.625.500      Insulin, Regular, Human
-	D06.472.699.587.200.500.625.500.500      Isophane Insulin, Human
-	D06.472.699.587.200.500.625.700      Insulin, Regular, Pork
-	D06.472.699.587.334      Islet Amyloid Polypeptide
-	D06.472.699.587.469      Proglucagon
-	D06.472.699.587.469.500      Glucagon
-	D06.472.699.587.700      Pancreatic Polypeptide
-	D06.472.699.587.780      Somatostatin
-	D06.472.699.587.780.500      Somatostatin-28
-	D06.472.699.590      Parathyroid Hormone
-	D06.472.699.590.850      Teriparatide
-	D06.472.699.591      Parathyroid Hormone-Related Protein
-	D06.472.699.592      Peptide PHI
-	D06.472.699.595      Peptide YY
-	D06.472.699.631      Pituitary Hormones
-	D06.472.699.631.525      Pituitary Hormones, Anterior
-	D06.472.699.631.525.343      Gonadotropins, Pituitary
-	D06.472.699.631.525.343.288      Follicle Stimulating Hormone
-	D06.472.699.631.525.343.288.500      Follicle Stimulating Hormone, beta Subunit
-	D06.472.699.631.525.343.288.625      Follicle Stimulating Hormone, Human
-	D06.472.699.631.525.343.288.750      Glycoprotein Hormones, alpha Subunit
-	D06.472.699.631.525.343.463      Luteinizing Hormone
-	D06.472.699.631.525.343.463.249      Glycoprotein Hormones, alpha Subunit
-	D06.472.699.631.525.343.463.500      Luteinizing Hormone, beta Subunit

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D06.472.699.631.525.343.583 Menotropins
-	D06.472.699.631.525.343.583.500 Urofollitropin
-	D06.472.699.631.525.425 Growth Hormone
-	D06.472.699.631.525.425.875 Human Growth Hormone
-	D06.472.699.631.525.525 Prolactin
-	D06.472.699.631.525.600 Pro-Opiomelanocortin
-	D06.472.699.631.525.600.119 alpha-Endorphin
-	D06.472.699.631.525.600.179 alpha-MSH
-	D06.472.699.631.525.600.239 beta-Endorphin
-	D06.472.699.631.525.600.480 beta-Lipotropin
-	D06.472.699.631.525.600.492 beta-MSH
-	D06.472.699.631.525.600.498 Corticotropin-Like Intermediate Lobe Peptide
-	D06.472.699.631.525.600.505 gamma-Endorphin
-	D06.472.699.631.525.600.518 gamma-Lipotropin
-	D06.472.699.631.525.600.524 gamma-MSH
-	D06.472.699.631.525.600.531 Melanocortins
-	D06.472.699.631.525.600.531.500 Adrenocorticotrophic Hormone
-	D06.472.699.631.525.600.531.500.200 Cosyntropin
-	D06.472.699.631.525.600.531.750 Melanocyte-Stimulating Hormones
-	D06.472.699.631.525.600.531.750.050 alpha-MSH
-	D06.472.699.631.525.600.531.750.075 beta-MSH
-	D06.472.699.631.525.600.531.750.115 gamma-MSH
-	D06.472.699.631.525.883 Thyrotropin
-	D06.472.699.631.525.883.249 Glycoprotein Hormones, alpha Subunit
-	D06.472.699.631.525.883.374 Thyrotropin Alfa
-	D06.472.699.631.525.883.500 Thyrotropin, beta Subunit
-	D06.472.699.631.692 Pituitary Hormones, Posterior
-	D06.472.699.631.692.433 Oxytocin
-	D06.472.699.631.692.781 Vasopressins
-	D06.472.699.631.692.781.100 Arginine Vasopressin
-	D06.472.699.631.692.781.100.250 Deamino Arginine Vasopressin
-	D06.472.699.631.692.781.400 Lypressin
-	D06.472.699.631.692.781.400.350 Felypressin
-	D06.472.699.631.692.781.700 Ornipressin
-	D06.472.699.631.692.881 Vasotocin
-	D06.472.699.649 Placental Hormones

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D06.472.699.649.367 Chorionic Gonadotropin
-	D06.472.699.649.367.125 Chorionic Gonadotropin, beta Subunit, Human
-	D06.472.699.649.367.562 Glycoprotein Hormones, alpha Subunit
-	D06.472.699.649.451 Gonadotropins, Equine
-	D06.472.699.649.692 Placental Lactogen
-	D06.472.699.715 Relaxin
-	D06.472.699.810 Secretin
-	D06.472.699.857 Urocortins
-	D06.472.699.905 Urotensins
-	D06.472.699.952 Vasoactive Intestinal Peptide
-	D06.472.910 Thymus Hormones
-	D06.472.910.750 Thymic Factor, Circulating
-	D06.472.910.800 Thymopoietins
-	D06.472.910.800.850 Thymopentin
-	D06.472.910.850 Thymosin
-	D06.472.931 Thyroid Hormones
-	D06.472.931.103 Dextrothyroxine
-	D06.472.931.208 Diiodotyrosine
-	D06.472.931.388 Monoiodotyrosine
-	D06.472.931.669 Thyroid (USP)
-	D06.472.931.740 Thyronines
-	D06.472.931.740.180 Diiodothyronines
-	D06.472.931.740.385 Triiodothyronine
-	D06.472.931.740.590 Triiodothyronine, Reverse
-	D06.472.931.812 Thyroxine
-	D08 Enzymes and Coenzymes
-	D08.211 Coenzymes
-	D08.211.090 Biopterin
-	D08.211.090.500 Neopterin
-	D08.211.096 Biotin
-	D08.211.175 Cobamides
-	D08.211.211 Coenzyme A
-	D08.211.211.300 Acyl Coenzyme A
-	D08.211.211.300.075 Acetyl Coenzyme A
-	D08.211.211.300.500 Malonyl Coenzyme A
-	D08.211.211.300.700 Palmitoyl Coenzyme A

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D08.211.474                      Flavins
-	D08.211.474.650                  Riboflavin
-	D08.211.474.650.249              Flavin-Adenine Dinucleotide
-	D08.211.474.650.500              Flavin Mononucleotide
-	D08.211.589                      NAD
-	D08.211.625                      NADP
-	D08.211.682                      PQQ Cofactor
-	D08.211.740                      Pyridoxal Phosphate
-	D08.211.790                      Sphingolipid Activator Proteins
-	D08.211.790.249                  G(M2) Activator Protein
-	D08.211.790.500                  Saposins
-	D08.211.840                      Tetrahydrofolates
-	D08.211.840.300                  Formyltetrahydrofolates
-	D08.211.840.300.500              Leucovorin
-	D08.211.878                      Thiamine Pyrophosphate
-	D08.211.906                      Thioctic Acid
-	D08.211.935                      Ubiquinone
-	D08.244                          Cytochromes
-	D08.244.175                      Cytochrome a Group
-	D08.244.175.249                  Cytochromes a
-	D08.244.175.500                  Cytochromes a1
-	D08.244.175.600                  Cytochromes a3
-	D08.244.187                      Cytochrome b Group
-	D08.244.187.249                  Cytochromes b
-	D08.244.187.500                  Cytochromes b5
-	D08.244.187.600                  Cytochromes b6
-	D08.244.286                      Cytochrome c Group
-	D08.244.286.100                  Cytochromes c
-	D08.244.286.150                  Cytochromes c'
-	D08.244.286.200                  Cytochromes c1
-	D08.244.286.300                  Cytochromes c2
-	D08.244.286.600                  Cytochromes c6
-	D08.244.300                      Cytochrome d Group
-	D08.244.453                      Cytochrome P-450 Enzyme System
New Tree	<a href="#">D08.244.453.005</a> <a href="#">Aryl Hydrocarbon Hydroxylases</a>

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	D08.244.453.005.050	Aniline Hydroxylase
New Tree	D08.244.453.005.110	Benzopyrene Hydroxylase
New Tree	D08.244.453.005.332	Cytochrome P-450 CYP1A1
New Tree	D08.244.453.005.443	Cytochrome P-450 CYP1A2
New Tree	D08.244.453.005.500	Cytochrome P-450 CYP1B1
New Tree	D08.244.453.005.550	Cytochrome P-450 CYP2B1
New Tree	D08.244.453.005.575	Cytochrome P-450 CYP2B6
New Tree	D08.244.453.005.590	Cytochrome P-450 CYP2C8
New Tree	D08.244.453.005.600	Cytochrome P-450 CYP2D6
New Tree	D08.244.453.012	Camphor 5-Monooxygenase
Old Tree	D08.244.453.040	Aryl Hydrocarbon Hydroxylases
Old Tree	D08.244.453.040.050	Aniline Hydroxylase
Old Tree	D08.244.453.040.110	Benzopyrene Hydroxylase
Old Tree	D08.244.453.040.332	Cytochrome P-450 CYP1A1
Old Tree	D08.244.453.040.443	Cytochrome P-450 CYP1A2
Old Tree	D08.244.453.040.500	Cytochrome P-450 CYP1B1
Old Tree	D08.244.453.040.550	Cytochrome P-450 CYP2B1
Old Tree	D08.244.453.040.575	Cytochrome P-450 CYP2B6
Old Tree	D08.244.453.040.590	Cytochrome P-450 CYP2C8
Old Tree	D08.244.453.040.600	Cytochrome P-450 CYP2D6
Old Tree	D08.244.453.085	Camphor 5-Monooxygenase
Old Tree	D08.244.453.100	Cytochrome P-450 CYP2A6
New Heading	<b>D08.244.453.100</b>	<b>Cytochrome P450 Family 1</b>
New Tree	D08.244.453.100.500	Cytochrome P-450 CYP1A1
New Tree	D08.244.453.100.750	Cytochrome P-450 CYP1A2
New Tree	D08.244.453.100.875	Cytochrome P-450 CYP1B1
Old Tree	D08.244.453.300	Cytochrome P-450 CYP2E1

## MeSH Tree Changes for 2017

Type	Tree - heading	
Old Tree	D08.244.453.400	Cytochrome P-450 CYP3A
Old Tree	D08.244.453.450	Cytochrome P-450 CYP4A
New Heading	<b>D08.244.453.484</b>	<b>Cytochrome P450 Family 11</b>
New Tree	D08.244.453.484.250	Cholesterol Side-Chain Cleavage Enzyme
New Tree	D08.244.453.484.500	Cytochrome P-450 CYP11B2
New Tree	D08.244.453.484.750	Steroid 11-beta-Hydroxylase
New Heading	<b>D08.244.453.485</b>	<b>Cytochrome P450 Family 12</b>
New Heading	<b>D08.244.453.487</b>	<b>Cytochrome P450 Family 17</b>
New Tree	D08.244.453.487.500	Steroid 17-alpha-Hydroxylase
New Heading	<b>D08.244.453.489</b>	<b>Cytochrome P450 Family 19</b>
New Tree	D08.244.453.489.500	Aromatase
New Heading	<b>D08.244.453.491</b>	<b>Cytochrome P450 Family 2</b>
New Tree	D08.244.453.491.250	Cytochrome P-450 CYP2A6
New Tree	D08.244.453.491.313	Cytochrome P-450 CYP2B1
New Tree	D08.244.453.491.344	Cytochrome P-450 CYP2B6
New Tree	D08.244.453.491.368	Cytochrome P-450 CYP2C8
New Tree	D08.244.453.491.372	Cytochrome P-450 CYP2D6
New Tree	D08.244.453.491.375	Cytochrome P-450 CYP2E1
New Tree	D08.244.453.491.500	Limonene Hydroxylases
New Tree	D08.244.453.491.500.500	Cytochrome P-450 CYP2C9
New Tree	D08.244.453.491.500.700	Cytochrome P-450 CYP2C19
New Heading	<b>D08.244.453.493</b>	<b>Cytochrome P450 Family 21</b>
New	D08.244.453.493.500	Steroid 21-Hydroxylase

## MeSH Tree Changes for 2017

Type	Tree - heading	
Tree		
New Heading	<b>D08.244.453.496</b>	<b>Cytochrome P450 Family 24</b>
New Tree	<a href="#">D08.244.453.496.500</a>	<a href="#">Vitamin D3 24-Hydroxylase</a>
New Heading	<b>D08.244.453.498</b>	<b>Cytochrome P450 Family 26</b>
New Heading	<b>D08.244.453.498.500</b>	<b>Retinoic Acid 4-Hydroxylase</b>
New Heading	<b>D08.244.453.499</b>	<b>Cytochrome P450 Family 27</b>
New Tree	<a href="#">D08.244.453.499.500</a>	<a href="#">25-Hydroxyvitamin D3 1-alpha-Hydroxylase</a>
New Tree	<a href="#">D08.244.453.499.750</a>	<a href="#">Cholestanetriol 26-Monooxygenase</a>
Old Tree	<a href="#">D08.244.453.500</a>	<a href="#">Limonene Hydroxylases</a>
Old Tree	<a href="#">D08.244.453.500.500</a>	<a href="#">Cytochrome P-450 CYP2C9</a>
Old Tree	<a href="#">D08.244.453.500.700</a>	<a href="#">Cytochrome P-450 CYP2C19</a>
New Heading	<b>D08.244.453.860</b>	<b>Cytochrome P450 Family 3</b>
New Tree	<a href="#">D08.244.453.860.500</a>	<a href="#">Cytochrome P-450 CYP3A</a>
New Heading	<b>D08.244.453.870</b>	<b>Cytochrome P450 Family 4</b>
New Tree	<a href="#">D08.244.453.870.500</a>	<a href="#">Cytochrome P-450 CYP4A</a>
New Heading	<b>D08.244.453.875</b>	<b>Cytochrome P450 Family 46</b>
New Heading	<b>D08.244.453.875.500</b>	<b>Cholesterol 24-Hydroxylase</b>
New Heading	<b>D08.244.453.878</b>	<b>Cytochrome P450 Family 51</b>
New Tree	<a href="#">D08.244.453.878.500</a>	<a href="#">Sterol 14-Demethylase</a>
New Heading	<b>D08.244.453.880</b>	<b>Cytochrome P450 Family 6</b>
New Heading	<b>D08.244.453.890</b>	<b>Cytochrome P450 Family 7</b>
New Tree	<a href="#">D08.244.453.890.500</a>	<a href="#">Cholesterol 7-alpha-Hydroxylase</a>
New Heading	<b>D08.244.453.900</b>	<b>Cytochrome P450 Family 8</b>
New	<a href="#">D08.244.453.900.500</a>	<a href="#">Steroid 12-alpha-Hydroxylase</a>

## MeSH Tree Changes for 2017

Type	Tree - heading	
Tree		
-	D08.244.453.915	Steroid Hydroxylases
New Heading	<b>D08.244.453.915.025</b>	<b>Cholesterol 24-Hydroxylase</b>
-	D08.244.453.915.050	Cytochrome P-450 CYP11B2
-	D08.244.453.915.099	Aromatase
-	D08.244.453.915.149	Cholestanetriol 26-Monooxygenase
-	D08.244.453.915.200	Cholesterol 7-alpha-Hydroxylase
-	D08.244.453.915.212	Cholesterol Side-Chain Cleavage Enzyme
-	D08.244.453.915.400	25-Hydroxyvitamin D3 1-alpha-Hydroxylase
-	D08.244.453.915.730	Steroid 12-alpha-Hydroxylase
-	D08.244.453.915.737	Steroid 16-alpha-Hydroxylase
-	D08.244.453.915.748	Steroid 17-alpha-Hydroxylase
-	D08.244.453.915.750	Steroid 11-beta-Hydroxylase
-	D08.244.453.915.760	Steroid 21-Hydroxylase
-	D08.244.453.915.880	Sterol 14-Demethylase
-	D08.244.453.957	Trans-Cinnamate 4-Monooxygenase
Old Tree	<b>D08.244.453.978</b>	<b>Vitamin D3 24-Hydroxylase</b>
-	D08.244.726	Cytochromes f
-	D08.622	Enzyme Precursors
-	D08.622.184	Chymotrypsinogen
-	D08.622.200	Complement Factor B
-	D08.622.509	Pepsinogens
-	D08.622.509.700	Pepsinogen A
-	D08.622.509.725	Pepsinogen C
-	D08.622.610	Plasminogen
-	D08.622.610.500	Angiostatins
-	D08.622.705	Protein C
-	D08.622.709	Prothrombin
-	D08.622.885	Trypsinogen
-	D08.811	Enzymes
New Heading	<b>D08.811.037</b>	<b>Deubiquitinating Enzymes</b>
New Tree	<a href="#">D08.811.037.250</a>	<a href="#">Ataxin-3</a>
New Tree	<a href="#">D08.811.037.500</a>	<a href="#">Ubiquitin Thiolesterase</a>



## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	<a href="#">D08.811.037.750</a>	<a href="#">Ubiquitin-Specific Proteases</a>
New Heading	<b>D08.811.037.750.500</b>	<b>Tumor Necrosis Factor alpha-Induced Protein 3</b>
-	D08.811.074	DNA Repair Enzymes
New Heading	<b>D08.811.074.062</b>	<b>AlkB Enzymes</b>
New Heading	<b>D08.811.074.062.500</b>	<b>AlkB Homolog 1, Histone H2a Dioxygenase</b>
New Heading	<b>D08.811.074.062.750</b> <b>Dioxygenase</b>	<b>AlkB Homolog 2, Alpha-Ketoglutarate-Dependent Dioxygenase</b>
New Heading	<b>D08.811.074.062.875</b> <b>Dioxygenase</b>	<b>AlkB Homolog 3, Alpha-Ketoglutarate-Dependent Dioxygenase</b>
New Heading	<b>D08.811.074.062.937</b>	<b>AlkB Homolog 5, RNA Demethylase</b>
New Heading	<b>D08.811.074.062.968</b>	<b>Alpha-Ketoglutarate-Dependent Dioxygenase FTO</b>
-	D08.811.074.124	Deoxyribodipyrimidine Photo-Lyase
-	D08.811.074.249	DNA Glycosylases
-	D08.811.074.249.500	DNA-Formamidopyrimidine Glycosylase
-	D08.811.074.249.750	Thymine DNA Glycosylase
-	D08.811.074.249.875	Uracil-DNA Glycosidase
-	D08.811.074.500	DNA Ligases
New Heading	<b>D08.811.074.500.500</b>	<b>DNA Ligase ATP</b>
-	D08.811.074.750	DNA-(Apurinic or Apyrimidinic Site) Lyase
New Heading	<b>D08.811.074.766</b>	<b>MutL Proteins</b>
New Heading	<b>D08.811.074.766.250</b>	<b>Mismatch Repair Endonuclease PMS2</b>
New Heading	<b>D08.811.074.766.500</b>	<b>MutL Protein Homolog 1</b>
-	D08.811.074.781	MutS DNA Mismatch-Binding Protein
-	D08.811.074.812	MutS Homolog 2 Protein
-	D08.811.074.875	Polynucleotide 5'-Hydroxyl-Kinase
-	D08.811.150	DNA Restriction-Modification Enzymes
-	D08.811.150.240	DNA Modification Methylases
-	D08.811.150.280	DNA Restriction Enzymes
-	D08.811.150.280.250	Deoxyribonucleases, Type I Site-Specific
-	D08.811.150.280.260	Deoxyribonucleases, Type II Site-Specific

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D08.811.150.280.260.240	Deoxyribonuclease BamHI
-	D08.811.150.280.260.300	Deoxyribonuclease EcoRI
-	D08.811.150.280.260.400	Deoxyribonuclease HindIII
-	D08.811.150.280.260.420	Deoxyribonuclease HpaII
-	D08.811.150.280.270	Deoxyribonucleases, Type III Site-Specific
-	D08.811.165	DNA, Catalytic
-	D08.811.180	Enzymes, Immobilized
-	D08.811.255	Holoenzymes
-	D08.811.255.500	Apoenzymes
-	D08.811.277	Hydrolases
-	D08.811.277.040	Acid Anhydride Hydrolases
-	D08.811.277.040.025	Adenosine Triphosphatases
-	D08.811.277.040.025.047	Arsenite Transporting ATPases
-	D08.811.277.040.025.095	Ca(2+) Mg(2+)-ATPase
-	D08.811.277.040.025.125	Calcium-Transporting ATPases
-	D08.811.277.040.025.125.249 ATPases	Plasma Membrane Calcium-Transporting
-	D08.811.277.040.025.125.500 ATPases	Sarcoplasmic Reticulum Calcium-Transporting
-	D08.811.277.040.025.142	Chaperonins
-	D08.811.277.040.025.142.500	Group I Chaperonins
-	D08.811.277.040.025.142.500.500	Chaperonin 60
-	D08.811.277.040.025.142.750	Group II Chaperonins
-	D08.811.277.040.025.142.750.500	Chaperonin Containing TCP-1
-	D08.811.277.040.025.142.750.750	Thermosomes
-	D08.811.277.040.025.159	DNA Helicases
-	D08.811.277.040.025.159.124	DnaB Helicases
New Heading	<b>D08.811.277.040.025.159.155</b>	<b>Ku Autoantigen</b>
-	D08.811.277.040.025.159.186	Minichromosome Maintenance Proteins
-	D08.811.277.040.025.159.186.200 Component 2	Minichromosome Maintenance Complex
-	D08.811.277.040.025.159.186.300 Component 3	Minichromosome Maintenance Complex
-	D08.811.277.040.025.159.186.400 Component 4	Minichromosome Maintenance Complex
-	D08.811.277.040.025.159.186.500 Component 5	Minichromosome Maintenance Complex

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D08.811.277.040.025.159.186.600 Component 6	Minichromosome Maintenance Complex
-	D08.811.277.040.025.159.186.700 Component 7	Minichromosome Maintenance Complex
-	D08.811.277.040.025.159.186.800 Component 8	Minichromosome Maintenance Complex
-	D08.811.277.040.025.159.186.900 Component 9	Minichromosome Maintenance Complex
-	D08.811.277.040.025.159.249	RecQ Helicases
New Heading	<b>D08.811.277.040.025.159.249.500</b>	<b>Werner Syndrome Helicase</b>
-	D08.811.277.040.025.159.500	Xeroderma Pigmentosum Group D Protein
-	D08.811.277.040.025.176 Complex	Mi-2 Nucleosome Remodeling and Deacetylase Complex
-	D08.811.277.040.025.193	Molecular Motor Proteins
-	D08.811.277.040.025.193.249	Dyneins
-	D08.811.277.040.025.193.249.500	Axonemal Dyneins
-	D08.811.277.040.025.193.249.750	Cytoplasmic Dyneins
-	D08.811.277.040.025.193.500	Kinesin
-	D08.811.277.040.025.193.750	Myosins
-	D08.811.277.040.025.193.750.249	Myosin Heavy Chains
-	D08.811.277.040.025.193.750.500	Myosin Type I
-	D08.811.277.040.025.193.750.750	Myosin Type II
-	D08.811.277.040.025.193.750.750.124	Cardiac Myosins
-	D08.811.277.040.025.193.750.750.124.249	Atrial Myosins
-	D08.811.277.040.025.193.750.750.124.500	Ventricular Myosins
-	D08.811.277.040.025.193.750.750.374	Nonmuscle Myosin Type IIA
-	D08.811.277.040.025.193.750.750.500	Nonmuscle Myosin Type IIB
-	D08.811.277.040.025.193.750.750.750	Skeletal Muscle Myosins
-	D08.811.277.040.025.193.750.750.875	Smooth Muscle Myosins
-	D08.811.277.040.025.193.750.812	Myosin Type III
-	D08.811.277.040.025.193.750.843	Myosin Type IV
-	D08.811.277.040.025.193.750.875	Myosin Type V
New Heading	<b>D08.811.277.040.025.215</b>	<b>MutL Proteins</b>
New Heading	<b>D08.811.277.040.025.215.250</b>	<b>Mismatch Repair Endonuclease PMS2</b>
New Heading	<b>D08.811.277.040.025.215.500</b>	<b>MutL Protein Homolog 1</b>

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D08.811.277.040.025.237 MutS DNA Mismatch-Binding Protein
-	D08.811.277.040.025.281 MutS Homolog 2 Protein
-	D08.811.277.040.025.303 N-Ethylmaleimide-Sensitive Proteins
-	D08.811.277.040.025.325 Proton-Translocating ATPases
-	D08.811.277.040.025.325.249 Bacterial Proton-Translocating ATPases
-	D08.811.277.040.025.325.500 Chloroplast Proton-Translocating ATPases
-	D08.811.277.040.025.325.625 H(+)-K(+)-Exchanging ATPase
-	D08.811.277.040.025.325.750 Mitochondrial Proton-Translocating ATPases
-	D08.811.277.040.025.325.875 Vacuolar Proton-Translocating ATPases
-	D08.811.277.040.025.810 Sodium-Potassium-Exchanging ATPase
-	D08.811.277.040.050 Apyrase
-	D08.811.277.040.330 GTP Phosphohydrolases
-	D08.811.277.040.330.200 Dynamins
-	D08.811.277.040.330.200.100 Dynamin I
-	D08.811.277.040.330.200.200 Dynamin II
-	D08.811.277.040.330.200.300 Dynamin III
-	D08.811.277.040.330.300 GTP-Binding Proteins
-	D08.811.277.040.330.300.100 GTP Phosphohydrolase-Linked Elongation Factors
-	D08.811.277.040.330.300.100.101 Peptide Elongation Factor 1
-	D08.811.277.040.330.300.100.102 Peptide Elongation Factor 2
-	D08.811.277.040.330.300.100.200 Peptide Elongation Factor G
-	D08.811.277.040.330.300.100.700 Peptide Elongation Factor Tu
-	D08.811.277.040.330.300.200 Heterotrimeric GTP-Binding Proteins
-	D08.811.277.040.330.300.200.100 GTP-Binding Protein alpha Subunits
-	D08.811.277.040.330.300.200.100.100 GTP-Binding Protein alpha Subunits, G12-G13
-	D08.811.277.040.330.300.200.100.200 GTP-Binding Protein alpha Subunits, Gi-Go
-	D08.811.277.040.330.300.200.100.200.500 GTP-Binding Protein alpha Subunit, Gi2
-	D08.811.277.040.330.300.200.100.300 GTP-Binding Protein alpha Subunits, Gq-G11
-	D08.811.277.040.330.300.200.100.400 GTP-Binding Protein alpha Subunits, Gs
-	D08.811.277.040.330.300.200.800 Transducin
-	D08.811.277.040.330.300.400 Monomeric GTP-Binding Proteins
-	D08.811.277.040.330.300.400.100 ADP-Ribosylation Factors

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D08.811.277.040.330.300.400.100.100 ADP-Ribosylation Factor 1
-	D08.811.277.040.330.300.400.400 rab GTP-Binding Proteins
-	D08.811.277.040.330.300.400.400.025 rab1 GTP-Binding Proteins
-	D08.811.277.040.330.300.400.400.050 rab2 GTP-Binding Protein
-	D08.811.277.040.330.300.400.400.100 rab3 GTP-Binding Proteins
-	D08.811.277.040.330.300.400.400.100.500 rab3A GTP-Binding Protein
-	D08.811.277.040.330.300.400.400.150 rab4 GTP-Binding Proteins
-	D08.811.277.040.330.300.400.400.200 rab5 GTP-Binding Proteins
-	D08.811.277.040.330.300.400.450 ral GTP-Binding Proteins
-	D08.811.277.040.330.300.400.462 ran GTP-Binding Protein
-	D08.811.277.040.330.300.400.475 rap GTP-Binding Proteins
-	D08.811.277.040.330.300.400.475.100 rap1 GTP-Binding Proteins
-	D08.811.277.040.330.300.400.500 ras Proteins
-	D08.811.277.040.330.300.400.500.300 Oncogene Protein p21(ras)
-	D08.811.277.040.330.300.400.500.600 Proto-Oncogene Proteins p21(ras)
-	D08.811.277.040.330.300.400.700 rho GTP-Binding Proteins
-	D08.811.277.040.330.300.400.700.050 cdc42 GTP-Binding Protein
-	D08.811.277.040.330.300.400.700.060 cdc42 GTP-Binding Protein, Saccharomyces cerevisiae
-	D08.811.277.040.330.300.400.700.100 rac GTP-Binding Proteins
-	D08.811.277.040.330.300.400.700.100.500 rac1 GTP-Binding Protein
-	D08.811.277.040.330.300.400.700.200 rhoA GTP-Binding Protein
-	D08.811.277.040.330.300.400.700.300 rhoB GTP-Binding Protein
-	D08.811.277.040.330.300.550 Myxovirus Resistance Proteins
-	D08.811.277.040.330.300.700 Septins
-	D08.811.277.040.465 Nucleoside-Triphosphatase
-	D08.811.277.040.600 Pyrophosphatases
-	D08.811.277.040.600.399 Inorganic Pyrophosphatase
-	D08.811.277.040.600.800 Thiamine Pyrophosphatase
-	D08.811.277.040.850 Thiamin-Triphosphatase
-	D08.811.277.063 Adenosylhomocysteinase
-	D08.811.277.087 Amidohydrolases
-	D08.811.277.087.030 N-Acetylmuramoyl-L-alanine Amidase
-	D08.811.277.087.060 Allophanate Hydrolase
-	D08.811.277.087.100 Arylformamidase
-	D08.811.277.087.116 Asparaginase

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D08.811.277.087.125 Aspartylglucosylaminase
-	D08.811.277.087.180 beta-Lactamases
-	D08.811.277.087.180.229 Cephalosporinase
-	D08.811.277.087.180.697 Penicillinase
-	D08.811.277.087.200 Biotinidase
-	D08.811.277.087.250 Ceramidases
-	D08.811.277.087.250.100 Acid Ceramidase
-	D08.811.277.087.250.300 Alkaline Ceramidase
-	D08.811.277.087.250.650 Neutral Ceramidase
-	D08.811.277.087.280 Dihydroorotase
-	D08.811.277.087.483 Glutaminase
-	D08.811.277.087.520 Histone Deacetylases
-	D08.811.277.087.520.200 Group III Histone Deacetylases
-	D08.811.277.087.520.200.650 Sirtuins
-	D08.811.277.087.520.200.650.100 Sirtuin 1
-	D08.811.277.087.520.200.650.200 Sirtuin 2
-	D08.811.277.087.520.200.650.600 Sirtuin 3
-	D08.811.277.087.520.500 Mi-2 Nucleosome Remodeling and Deacetylase Complex
-	D08.811.277.087.520.500.100 Histone Deacetylase 1
-	D08.811.277.087.520.500.200 Histone Deacetylase 2
-	D08.811.277.087.520.500.775 Retinoblastoma-Binding Protein 4
-	D08.811.277.087.520.500.887 Retinoblastoma-Binding Protein 7
-	D08.811.277.087.520.750 Sin3 Histone Deacetylase and Corepressor Complex
-	D08.811.277.087.520.750.100 Histone Deacetylase 1
-	D08.811.277.087.520.750.200 Histone Deacetylase 2
-	D08.811.277.087.520.750.775 Retinoblastoma-Binding Protein 4
-	D08.811.277.087.520.750.887 Retinoblastoma-Binding Protein 7
-	D08.811.277.087.610 Nicotinamidase
-	D08.811.277.087.690 Penicillin Amidase
-	D08.811.277.087.725 Peptide-N4-(N-acetyl-beta-glucosaminy) Asparagine Amidase
-	D08.811.277.087.760 Pyroglutamate Hydrolase
-	D08.811.277.087.902 Urease
-	D08.811.277.151 Aminohydrolases
-	D08.811.277.151.300 GTP Cyclohydrolase

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D08.811.277.151.350                      Guanine Deaminase
-	D08.811.277.151.418                      Methenyltetrahydrofolate Cyclohydrolase
-	D08.811.277.151.486                      Nucleoside Deaminases
-	D08.811.277.151.486.075                      Adenosine Deaminase
-	D08.811.277.151.486.250                      Cytidine Deaminase
New Heading	<b>D08.811.277.151.486.250.500                      APOBEC Deaminases</b>
New Heading	<b>D08.811.277.151.486.250.500.500                      APOBEC-1 Deaminase</b>
New Heading	<b>D08.811.277.151.486.250.500.750                      APOBEC-3G Deaminase</b>
-	D08.811.277.151.486.625                      Cytosine Deaminase
-	D08.811.277.151.653                      Nucleotide Deaminases
-	D08.811.277.151.653.060                      AMP Deaminase
-	D08.811.277.151.653.200                      DCMP Deaminase
-	D08.811.277.300                      Complement Activating Enzymes
-	D08.811.277.300.280                      Complement C1r
-	D08.811.277.300.290                      Complement C1s
-	D08.811.277.300.450                      Complement Factor D
-	D08.811.277.340                      Epoxide Hydrolases
-	D08.811.277.352                      Esterases
-	D08.811.277.352.100                      Carboxylic Ester Hydrolases
-	D08.811.277.352.100.050                      Acetylesterase
-	D08.811.277.352.100.100                      Carboxylesterase
-	D08.811.277.352.100.170                      Cholinesterases
-	D08.811.277.352.100.170.176                      Acetylcholinesterase
-	D08.811.277.352.100.170.250                      Butyrylcholinesterase
-	D08.811.277.352.100.170.710                      Pseudocholinesterase
-	D08.811.277.352.100.220                      Dehydroascorbatase
-	D08.811.277.352.100.400                      Lipase
-	D08.811.277.352.100.400.745                      Pancrelipase
-	D08.811.277.352.100.430                      Lipoprotein Lipase
-	D08.811.277.352.100.500                      Monoacylglycerol Lipases
-	D08.811.277.352.100.550                      Naphthol AS D Esterase
-	D08.811.277.352.100.680                      Phospholipases
-	D08.811.277.352.100.680.510                      Lysophospholipase
-	D08.811.277.352.100.680.750                      Phospholipases A

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D08.811.277.352.100.680.750.875      Phospholipases A1
-	D08.811.277.352.100.680.750.937      Phospholipases A2
-	D08.811.277.352.100.680.750.937.249 Esterase      1-Alkyl-2-acetylglycerophosphocholine
-	D08.811.277.352.100.680.750.937.300 Independent      Phospholipases A2, Calcium-
-	D08.811.277.352.100.680.750.937.300.249      Group VI Phospholipases A2
-	D08.811.277.352.100.680.750.937.300.500      Peroxiredoxin VI
-	D08.811.277.352.100.680.750.937.625      Phospholipases A2, Cytosolic
-	D08.811.277.352.100.680.750.937.625.249      Group IV Phospholipases A2
-	D08.811.277.352.100.680.750.937.625.624      Peroxiredoxin VI
-	D08.811.277.352.100.680.750.937.750      Phospholipases A2, Secretory
-	D08.811.277.352.100.680.750.937.750.100      Group I Phospholipases A2
-	D08.811.277.352.100.680.750.937.750.100.500      Group IA Phospholipases A2
-	D08.811.277.352.100.680.750.937.750.100.750      Group IB Phospholipases A2
-	D08.811.277.352.100.680.750.937.750.550      Group II Phospholipases A2
-	D08.811.277.352.100.680.750.937.750.662      Group III Phospholipases A2
-	D08.811.277.352.100.680.750.937.750.775      Group V Phospholipases A2
-	D08.811.277.352.100.680.750.937.750.887      Group X Phospholipases A2
-	D08.811.277.352.100.700      Sterol Esterase
-	D08.811.277.352.335      Deoxyribonucleases
-	D08.811.277.352.335.350      Endodeoxyribonucleases
-	D08.811.277.352.335.350.137      Deoxyribonuclease (Pyrimidine Dimer)
-	D08.811.277.352.335.350.250      Deoxyribonuclease I
-	D08.811.277.352.335.350.250.900      Streptodornase and Streptokinase
-	D08.811.277.352.335.350.275      Deoxyribonuclease IV (Phage T4-Induced)
-	D08.811.277.352.335.350.300      DNA Restriction Enzymes
-	D08.811.277.352.335.350.300.250      Deoxyribonucleases, Type I Site-Specific
-	D08.811.277.352.335.350.300.260      Deoxyribonucleases, Type II Site-Specific
-	D08.811.277.352.335.350.300.260.240      Deoxyribonuclease BamHI
-	D08.811.277.352.335.350.300.260.250      Deoxyribonuclease EcoRI
-	D08.811.277.352.335.350.300.260.260      Deoxyribonuclease HindIII
-	D08.811.277.352.335.350.300.260.300      Deoxyribonuclease HpaII
-	D08.811.277.352.335.350.300.270      Deoxyribonucleases, Type III Site-Specific
-	D08.811.277.352.335.350.400      Holliday Junction Resolvases
-	D08.811.277.352.335.350.500      Micrococcal Nuclease



## MeSH Tree Changes for 2017

Type	Tree - heading	
New Heading	<b>D08.811.277.352.335.350.600</b>	<b>Mismatch Repair Endonuclease PMS2</b>
-	D08.811.277.352.335.350.700 Endonucleases	Single-Strand Specific DNA and RNA
New Heading	<b>D08.811.277.352.335.350.850</b> <b>Nucleases</b>	<b>Transcription Activator-Like Effector</b>
-	D08.811.277.352.335.375	Exodeoxyribonucleases
-	D08.811.277.352.335.375.750	Exodeoxyribonuclease V
New Heading	<b>D08.811.277.352.335.375.875</b>	<b>Werner Syndrome Helicase</b>
-	D08.811.277.352.355	Endonucleases
-	D08.811.277.352.355.325	Endodeoxyribonucleases
-	D08.811.277.352.355.325.300	DNA Restriction Enzymes
-	D08.811.277.352.355.325.300.250	Deoxyribonucleases, Type I Site-Specific
-	D08.811.277.352.355.325.300.260	Deoxyribonucleases, Type II Site-Specific
-	D08.811.277.352.355.325.300.260.240	Deoxyribonuclease BamHI
-	D08.811.277.352.355.325.300.260.250	Deoxyribonuclease EcoRI
-	D08.811.277.352.355.325.300.260.260	Deoxyribonuclease HindIII
-	D08.811.277.352.355.325.300.260.300	Deoxyribonuclease HpaII
-	D08.811.277.352.355.325.300.270	Deoxyribonucleases, Type III Site-Specific
-	D08.811.277.352.355.325.350	Flap Endonucleases
-	D08.811.277.352.355.325.400	Holliday Junction Resolvases
-	D08.811.277.352.355.325.500	Micrococcal Nuclease
-	D08.811.277.352.355.325.700 Endonucleases	Single-Strand Specific DNA and RNA
-	D08.811.277.352.355.350	Endoribonucleases
-	D08.811.277.352.355.350.500	Micrococcal Nuclease
-	D08.811.277.352.355.350.700	Ribonuclease H
-	D08.811.277.352.355.350.700.500 Virus	Ribonuclease H, Human Immunodeficiency
-	D08.811.277.352.355.350.715	Ribonuclease, Pancreatic
-	D08.811.277.352.355.350.725	Ribonuclease T1
-	D08.811.277.352.355.350.810	RNA-Induced Silencing Complex
-	D08.811.277.352.355.350.810.500	Argonaute Proteins
-	D08.811.277.352.355.350.820 Endonucleases	Single-Strand Specific DNA and RNA
-	D08.811.277.352.365	Exonucleases
-	D08.811.277.352.365.290	Exodeoxyribonucleases

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Heading	<b>D08.811.277.352.365.290.500</b>	<b>Werner Syndrome Helicase</b>
-	D08.811.277.352.365.300	Exoribonucleases
-	D08.811.277.352.640	Phosphoric Diester Hydrolases
-	D08.811.277.352.640.050	Annexin A3
-	D08.811.277.352.640.150	3',5'-Cyclic-AMP Phosphodiesterases
-	D08.811.277.352.640.150.100	Cyclic Nucleotide Phosphodiesterases, Type 1
-	D08.811.277.352.640.150.200	Cyclic Nucleotide Phosphodiesterases, Type 2
-	D08.811.277.352.640.150.300	Cyclic Nucleotide Phosphodiesterases, Type 3
-	D08.811.277.352.640.150.400	Cyclic Nucleotide Phosphodiesterases, Type 4
-	D08.811.277.352.640.150.500	Cyclic Nucleotide Phosphodiesterases, Type 5
-	D08.811.277.352.640.150.600	Cyclic Nucleotide Phosphodiesterases, Type 6
-	D08.811.277.352.640.150.700	Cyclic Nucleotide Phosphodiesterases, Type 7
-	D08.811.277.352.640.155	3',5'-Cyclic-GMP Phosphodiesterases
-	D08.811.277.352.640.155.100	Cyclic Nucleotide Phosphodiesterases, Type 1
-	D08.811.277.352.640.155.200	Cyclic Nucleotide Phosphodiesterases, Type 2
-	D08.811.277.352.640.155.500	Cyclic Nucleotide Phosphodiesterases, Type 5
-	D08.811.277.352.640.155.750	Cyclic Nucleotide Phosphodiesterases, Type 6
-	D08.811.277.352.640.160	2',3'-Cyclic-Nucleotide Phosphodiesterases
-	D08.811.277.352.640.160.500	2',3'-Cyclic Nucleotide 3'-Phosphodiesterase
-	D08.811.277.352.640.295	Glycerophosphoinositol Inositolphosphodiesterase
-	D08.811.277.352.640.430	Phosphodiesterase I
-	D08.811.277.352.640.700	Phospholipases
-	D08.811.277.352.640.700.700	Type C Phospholipases
-	D08.811.277.352.640.700.700.249 Lyase	Glycosylphosphatidylinositol Diacylglycerol-Lyase
-	D08.811.277.352.640.700.700.500	Phosphatidylinositol Diacylglycerol-Lyase
-	D08.811.277.352.640.700.700.562	Phosphoinositide Phospholipase C
-	D08.811.277.352.640.700.700.562.500	Phospholipase C beta
-	D08.811.277.352.640.700.700.562.625	Phospholipase C delta
-	D08.811.277.352.640.700.700.562.750	Phospholipase C gamma
-	D08.811.277.352.640.700.710	Phospholipase D
-	D08.811.277.352.640.750	Sphingomyelin Phosphodiesterase
-	D08.811.277.352.650	Phosphoric Monoester Hydrolases
-	D08.811.277.352.650.025	Acid Phosphatase
New Heading	<b>D08.811.277.352.650.025.500</b>	<b>Tartrate-Resistant Acid Phosphatase</b>

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D08.811.277.352.650.035	Alkaline Phosphatase
-	D08.811.277.352.650.200	Fructose-Bisphosphatase
-	D08.811.277.352.650.225	Glucose-6-Phosphatase
-	D08.811.277.352.650.300	Histidinol-Phosphatase
-	D08.811.277.352.650.575	4-Nitrophenylphosphatase
-	D08.811.277.352.650.587	Mitogen-Activated Protein Kinase Phosphatases
-	D08.811.277.352.650.587.200	Dual Specificity Phosphatase 1
-	D08.811.277.352.650.587.250	Dual Specificity Phosphatase 2
-	D08.811.277.352.650.587.300	Dual Specificity Phosphatase 3
-	D08.811.277.352.650.587.600	Dual Specificity Phosphatase 6
-	D08.811.277.352.650.600	Nucleotidases
-	D08.811.277.352.650.600.600	5'-Nucleotidase
-	D08.811.277.352.650.620	Phosphatidate Phosphatase
-	D08.811.277.352.650.622	Phosphofructokinase-2
New Heading	<b>D08.811.277.352.650.624</b>	<b>Phosphoinositide Phosphatases</b>
New Heading	<b>D08.811.277.352.650.624.500</b>	<b>Inositol Polyphosphate 5-Phosphatases</b>
New Heading	<b>D08.811.277.352.650.624.750</b> <b>Phosphatases</b>	<b>Phosphatidylinositol-3,4,5-Trisphosphate 5-</b>
-	D08.811.277.352.650.625	Phosphoprotein Phosphatases
-	D08.811.277.352.650.625.150	Calcineurin
-	D08.811.277.352.650.625.225	Dual-Specificity Phosphatases
-	D08.811.277.352.650.625.225.100	cdc25 Phosphatases
-	D08.811.277.352.650.625.225.200	Dual Specificity Phosphatase 1
-	D08.811.277.352.650.625.225.250	Dual Specificity Phosphatase 2
-	D08.811.277.352.650.625.225.300	Dual Specificity Phosphatase 3
-	D08.811.277.352.650.625.225.600	Dual Specificity Phosphatase 6
-	D08.811.277.352.650.625.300	Glycogen-Synthase-D Phosphatase
-	D08.811.277.352.650.625.475	Myosin-Light-Chain Phosphatase
-	D08.811.277.352.650.625.650	Phosphorylase Phosphatase
-	D08.811.277.352.650.625.687	Protein Phosphatase 1
-	D08.811.277.352.650.625.706	Protein Phosphatase 2
New Heading	<b>D08.811.277.352.650.625.716</b>	<b>Protein Phosphatase 2C</b>
-	D08.811.277.352.650.625.725 Phosphatase	Pyruvate Dehydrogenase (Lipoamide)- Phosphatase

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Heading	<b>D08.811.277.352.650.625.862</b>	<b>Tartrate-Resistant Acid Phosphatase</b>
-	D08.811.277.352.650.700	6-Phytase
-	D08.811.277.352.650.775	Protein Tyrosine Phosphatases
-	D08.811.277.352.650.775.250	Dual-Specificity Phosphatases
-	D08.811.277.352.650.775.250.100	cdc25 Phosphatases
-	D08.811.277.352.650.775.250.200	Dual Specificity Phosphatase 1
-	D08.811.277.352.650.775.250.250	Dual Specificity Phosphatase 2
-	D08.811.277.352.650.775.250.300	Dual Specificity Phosphatase 3
-	D08.811.277.352.650.775.250.600	Dual Specificity Phosphatase 6
-	D08.811.277.352.650.775.300	Protein Tyrosine Phosphatases, Non-Receptor
-	D08.811.277.352.650.775.300.100 Receptor Type 1	Protein Tyrosine Phosphatase, Non-Receptor Type 1
-	D08.811.277.352.650.775.300.200 Receptor Type 2	Protein Tyrosine Phosphatase, Non-Receptor Type 2
-	D08.811.277.352.650.775.300.300 Receptor Type 3	Protein Tyrosine Phosphatase, Non-Receptor Type 3
-	D08.811.277.352.650.775.300.400 Receptor Type 4	Protein Tyrosine Phosphatase, Non-Receptor Type 4
-	D08.811.277.352.650.775.300.600 Receptor Type 6	Protein Tyrosine Phosphatase, Non-Receptor Type 6
-	D08.811.277.352.650.775.300.800 Receptor Type 11	Protein Tyrosine Phosphatase, Non-Receptor Type 11
-	D08.811.277.352.650.775.300.850 Receptor Type 12	Protein Tyrosine Phosphatase, Non-Receptor Type 12
-	D08.811.277.352.650.775.300.860 Receptor Type 13	Protein Tyrosine Phosphatase, Non-Receptor Type 13
-	D08.811.277.352.650.775.300.930 Receptor Type 22	Protein Tyrosine Phosphatase, Non-Receptor Type 22
-	D08.811.277.352.650.775.400	Receptor-Like Protein Tyrosine Phosphatases
-	D08.811.277.352.650.775.400.100 Phosphatases, Class 1	Receptor-Like Protein Tyrosine Phosphatases, Class 1
-	D08.811.277.352.650.775.400.100.500	Antigens, CD45
-	D08.811.277.352.650.775.400.200 Phosphatases, Class 2	Receptor-Like Protein Tyrosine Phosphatases, Class 2
-	D08.811.277.352.650.775.400.300 Phosphatases, Class 3	Receptor-Like Protein Tyrosine Phosphatases, Class 3
-	D08.811.277.352.650.775.400.400 Phosphatases, Class 4	Receptor-Like Protein Tyrosine Phosphatases, Class 4
-	D08.811.277.352.650.775.400.500 Phosphatases, Class 5	Receptor-Like Protein Tyrosine Phosphatases, Class 5

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D08.811.277.352.650.775.400.700 Phosphatases, Class 7
-	D08.811.277.352.650.775.400.800 Phosphatases, Class 8
-	D08.811.277.352.650.775.700 Phosphatases
-	D08.811.277.352.650.775.700.200 Receptor Type 6
-	D08.811.277.352.650.775.700.800 Receptor Type 11
-	D08.811.277.352.650.850
-	D08.811.277.352.660
-	D08.811.277.352.660.500
-	D08.811.277.352.700
-	D08.811.277.352.700.350
-	D08.811.277.352.700.350.262
-	D08.811.277.352.700.350.381
-	D08.811.277.352.700.350.500
-	D08.811.277.352.700.350.700
-	D08.811.277.352.700.350.700.500 Virus
-	D08.811.277.352.700.350.707
-	D08.811.277.352.700.350.711
-	D08.811.277.352.700.350.715
-	D08.811.277.352.700.350.725
-	D08.811.277.352.700.350.810
-	D08.811.277.352.700.350.810.500
-	D08.811.277.352.700.350.850 Endonucleases
-	D08.811.277.352.700.375
-	D08.811.277.352.700.687
-	D08.811.277.352.827
-	D08.811.277.352.827.070
-	D08.811.277.352.827.070.060
-	D08.811.277.352.827.070.250
-	D08.811.277.352.827.070.625
-	D08.811.277.352.827.180
-	D08.811.277.352.827.180.175

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D08.811.277.352.827.180.175.060 N-Acetylgalactosamine-4-Sulfatase
-	D08.811.277.352.827.180.175.275 Chondro-4-Sulfatase
-	D08.811.277.352.827.500 Iduronate Sulfatase
-	D08.811.277.352.897 Thiolester Hydrolases
-	D08.811.277.352.897.075 Acetyl-CoA Hydrolase
-	D08.811.277.352.897.387 Fatty Acid Synthases
-	D08.811.277.352.897.387.100 Fatty Acid Synthase, Type I
-	D08.811.277.352.897.387.200 Fatty Acid Synthase, Type II
-	D08.811.277.352.897.700 Palmitoyl-CoA Hydrolase
-	D08.811.277.352.897.850 Ubiquitin Thiolesterase
-	D08.811.277.450 Glycoside Hydrolases
-	D08.811.277.450.050 alpha-L-Fucosidase
-	D08.811.277.450.066 Amylases
-	D08.811.277.450.066.050 alpha-Amylases
-	D08.811.277.450.066.050.249 Pancreatic alpha-Amylases
-	D08.811.277.450.066.050.500 Salivary alpha-Amylases
-	D08.811.277.450.066.100 beta-Amylase
-	D08.811.277.450.114 beta-Fructofuranosidase
-	D08.811.277.450.207 Chitinase
-	D08.811.277.450.207 Chitinases
New Heading	<b>D08.811.277.450.207.500 Chitinase-3-Like Protein 1</b>
-	D08.811.277.450.283 Dextranase
-	D08.811.277.450.329 Disaccharidases
-	D08.811.277.450.329.738 Sucrase
-	D08.811.277.450.329.738.700 Sucrase-Isomaltase Complex
-	D08.811.277.450.329.865 Trehalase
-	D08.811.277.450.410 Galactosidases
-	D08.811.277.450.410.050 alpha-Galactosidase
-	D08.811.277.450.410.100 beta-Galactosidase
-	D08.811.277.450.410.100.500 Lactase
-	D08.811.277.450.410.100.500.500 Lactase-Phlorizin Hydrolase
-	D08.811.277.450.410.120 Galactosylceramidase
-	D08.811.277.450.410.150 Galactosylgalactosylglucosylceramidase
-	D08.811.277.450.420 Glucosidases
-	D08.811.277.450.420.050 alpha-Glucosidases

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D08.811.277.450.420.200 Cellulases
-	D08.811.277.450.420.200.100 beta-Glucosidase
-	D08.811.277.450.420.200.200 Cellulase
-	D08.811.277.450.420.200.400 Cellulose 1,4-beta-Cellobiosidase
-	D08.811.277.450.420.200.450 Endo-1,3(4)-beta-Glucanase
-	D08.811.277.450.420.200.500 Glucan 1,3-beta-Glucosidase
-	D08.811.277.450.420.200.550 Glucan 1,4-beta-Glucosidase
-	D08.811.277.450.420.200.600 Glucan Endo-1,3-beta-D-Glucosidase
-	D08.811.277.450.420.375 Glucan 1,4-alpha-Glucosidase
-	D08.811.277.450.420.412 Glucosylceramidase
-	D08.811.277.450.420.450 Glycogen Debranching Enzyme System
-	D08.811.277.450.420.475 Lactase-Phlorizin Hydrolase
-	D08.811.277.450.426 Glucuronidase
-	D08.811.277.450.430 N-Glycosyl Hydrolases
-	D08.811.277.450.430.099 DNA Glycosylases
-	D08.811.277.450.430.099.500 DNA-Formamidopyrimidine Glycosylase
-	D08.811.277.450.430.099.750 Thymine DNA Glycosylase
-	D08.811.277.450.430.400 NAD+ Nucleosidase
-	D08.811.277.450.430.400.060 ADP-ribosyl Cyclase
-	D08.811.277.450.430.400.060.500 Antigens, CD38
-	D08.811.277.450.430.700 Ribosome Inactivating Proteins
-	D08.811.277.450.430.700.500 Ribosome Inactivating Proteins, Type 1
-	D08.811.277.450.430.700.750 Ribosome Inactivating Proteins, Type 2
-	D08.811.277.450.430.700.750.111 Abrin
-	D08.811.277.450.430.700.750.666 Ricin
-	D08.811.277.450.430.700.750.750 Shiga Toxins
-	D08.811.277.450.430.700.750.750.100 Shiga Toxin
-	D08.811.277.450.430.700.750.750.120 Shiga Toxin 1
-	D08.811.277.450.430.700.750.750.124 Shiga Toxin 2
-	D08.811.277.450.483 Hexosaminidases
-	D08.811.277.450.483.044 alpha-N-Acetylgalactosaminidase
-	D08.811.277.450.483.112 beta-N-Acetyl-Galactosaminidase
-	D08.811.277.450.483.180 beta-N-Acetylhexosaminidases
-	D08.811.277.450.483.180.500 Acetylglucosaminidase
-	D08.811.277.450.483.180.750 Hexosaminidase A
-	D08.811.277.450.483.180.750.500 beta-Hexosaminidase alpha Chain

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D08.811.277.450.483.180.750.750	beta-Hexosaminidase beta Chain
-	D08.811.277.450.483.180.875	Hexosaminidase B
-	D08.811.277.450.483.180.875.500	beta-Hexosaminidase beta Chain
-	D08.811.277.450.483.765 Acetylglucosaminidase	Mannosyl-Glycoprotein Endo-beta-N-
-	D08.811.277.450.529	Hyaluronoglucosaminidase
-	D08.811.277.450.560	Iduronidase
-	D08.811.277.450.585	Isoamylase
-	D08.811.277.450.625	Mannosidases
-	D08.811.277.450.625.500	alpha-Mannosidase
-	D08.811.277.450.625.750	beta-Mannosidase
-	D08.811.277.450.642	Muramidase
-	D08.811.277.450.692	Neuraminidase
-	D08.811.277.450.770	Oligo-1,6-Glucosidase
-	D08.811.277.450.770.800	Sucrase-Isomaltase Complex
-	D08.811.277.450.800	Polygalacturonase
-	D08.811.277.450.950	Xylosidases
-	D08.811.277.450.950.249	Endo-1,4-beta Xylanases
-	D08.811.277.450.950.500	Xylan Endo-1,3-beta-Xylosidase
-	D08.811.277.610	Pancreatin
-	D08.811.277.656	Peptide Hydrolases
-	D08.811.277.656.074	Aspartic Acid Proteases
-	D08.811.277.656.074.500	Aspartic Acid Endopeptidases
-	D08.811.277.656.074.500.180	Cathepsin D
-	D08.811.277.656.074.500.185	Cathepsin E
-	D08.811.277.656.074.500.200	Chymosin
New Heading	<b>D08.811.277.656.074.500.270</b>	<b>Endothelin-Converting Enzymes</b>
-	D08.811.277.656.074.500.340	HIV Protease
-	D08.811.277.656.074.500.700	Pepsin A
-	D08.811.277.656.074.500.780	Renin
-	D08.811.277.656.149	ATP-Dependent Proteases
-	D08.811.277.656.149.099	ATP-Dependent Endopeptidases
-	D08.811.277.656.149.099.500	Endopeptidase Clp
-	D08.811.277.656.149.099.750	Protease La
-	D08.811.277.656.224	Cathepsins



## MeSH Tree Changes for 2017

Type	Tree - heading
-	D08.811.277.656.224.062                      Cathepsin A
-	D08.811.277.656.224.125                      Cathepsin B
-	D08.811.277.656.224.130                      Cathepsin C
-	D08.811.277.656.224.187                      Cathepsin D
-	D08.811.277.656.224.250                      Cathepsin E
-	D08.811.277.656.224.300                      Cathepsin F
-	D08.811.277.656.224.350                      Cathepsin G
-	D08.811.277.656.224.400                      Cathepsin H
-	D08.811.277.656.224.405                      Cathepsin K
-	D08.811.277.656.224.407                      Cathepsin L
-	D08.811.277.656.224.417                      Cathepsin W
-	D08.811.277.656.224.425                      Cathepsin Z
-	D08.811.277.656.262                          Cysteine Proteases
-	D08.811.277.656.262.186                      Cathepsin C
-	D08.811.277.656.262.249                      Cathepsin Z
-	D08.811.277.656.262.500                      Cysteine Endopeptidases
-	D08.811.277.656.262.500.096                      Bromelains
-	D08.811.277.656.262.500.120                      Calpain
-	D08.811.277.656.262.500.126                      Caspases
-	D08.811.277.656.262.500.126.350                      Caspases, Effector
-	D08.811.277.656.262.500.126.350.300                      Caspase 3
-	D08.811.277.656.262.500.126.350.600                      Caspase 6
-	D08.811.277.656.262.500.126.350.700                      Caspase 7
-	D08.811.277.656.262.500.126.350.900                      Caspase 14
-	D08.811.277.656.262.500.126.550                      Caspases, Initiator
-	D08.811.277.656.262.500.126.550.100                      Caspase 1
-	D08.811.277.656.262.500.126.550.200                      Caspase 2
-	D08.811.277.656.262.500.126.550.800                      Caspase 8
-	D08.811.277.656.262.500.126.550.900                      Caspase 9
-	D08.811.277.656.262.500.126.550.910                      Caspase 10
-	D08.811.277.656.262.500.126.550.920                      Caspase 12
-	D08.811.277.656.262.500.133                      Cathepsin B
-	D08.811.277.656.262.500.146                      Cathepsin F
-	D08.811.277.656.262.500.153                      Cathepsin H
-	D08.811.277.656.262.500.156                      Cathepsin K
-	D08.811.277.656.262.500.157                      Cathepsin L

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D08.811.277.656.262.500.158	Cathepsin W
-	D08.811.277.656.262.500.160	Chymopapain
-	D08.811.277.656.262.500.350	Ficain
-	D08.811.277.656.262.500.585	Papain
-	D08.811.277.656.262.500.792	Separase
-	D08.811.277.656.300	Endopeptidases
-	D08.811.277.656.300.032	Amyloid Precursor Protein Secretases
-	D08.811.277.656.300.048	Aspartic Acid Endopeptidases
-	D08.811.277.656.300.048.180	Cathepsin D
-	D08.811.277.656.300.048.185	Cathepsin E
-	D08.811.277.656.300.048.200	Chymosin
New Heading	<b>D08.811.277.656.300.048.270</b>	<b>Endothelin-Converting Enzymes</b>
-	D08.811.277.656.300.048.340	HIV Protease
-	D08.811.277.656.300.048.700	Pepsin A
-	D08.811.277.656.300.048.780	Renin
-	D08.811.277.656.300.065	ATP-Dependent Endopeptidases
-	D08.811.277.656.300.065.500	Endopeptidase Clp
-	D08.811.277.656.300.065.750	Protease La
-	D08.811.277.656.300.099	Brinolase
-	D08.811.277.656.300.174	Coagulase
-	D08.811.277.656.300.200	Cysteine Endopeptidases
-	D08.811.277.656.300.200.096	Bromelains
-	D08.811.277.656.300.200.120	Calpain
-	D08.811.277.656.300.200.126	Caspases
-	D08.811.277.656.300.200.126.350	Caspases, Effector
-	D08.811.277.656.300.200.126.350.300	Caspase 3
-	D08.811.277.656.300.200.126.350.600	Caspase 6
-	D08.811.277.656.300.200.126.350.700	Caspase 7
-	D08.811.277.656.300.200.126.350.900	Caspase 14
-	D08.811.277.656.300.200.126.550	Caspases, Initiator
-	D08.811.277.656.300.200.126.550.100	Caspase 1
-	D08.811.277.656.300.200.126.550.200	Caspase 2
-	D08.811.277.656.300.200.126.550.800	Caspase 8
-	D08.811.277.656.300.200.126.550.900	Caspase 9
-	D08.811.277.656.300.200.126.550.910	Caspase 10

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D08.811.277.656.300.200.126.550.920	Caspase 12
-	D08.811.277.656.300.200.133	Cathepsin B
-	D08.811.277.656.300.200.146	Cathepsin F
-	D08.811.277.656.300.200.153	Cathepsin H
-	D08.811.277.656.300.200.156	Cathepsin K
-	D08.811.277.656.300.200.157	Cathepsin L
-	D08.811.277.656.300.200.158	Cathepsin W
-	D08.811.277.656.300.200.160	Chymopapain
-	D08.811.277.656.300.200.350	Ficain
-	D08.811.277.656.300.200.585	Papain
-	D08.811.277.656.300.200.792	Separase
-	D08.811.277.656.300.480	Metalloendopeptidases
-	D08.811.277.656.300.480.153	Botulinum Toxins
-	D08.811.277.656.300.480.153.100	Botulinum Toxins, Type A
-	D08.811.277.656.300.480.205	Collagenases
-	D08.811.277.656.300.480.205.351	Matrix Metalloproteinase 1
-	D08.811.277.656.300.480.205.352	Matrix Metalloproteinase 2
-	D08.811.277.656.300.480.205.358	Matrix Metalloproteinase 8
-	D08.811.277.656.300.480.205.360	Matrix Metalloproteinase 9
-	D08.811.277.656.300.480.205.363	Matrix Metalloproteinase 13
-	D08.811.277.656.300.480.205.500	Microbial Collagenase
New Heading	<b>D08.811.277.656.300.480.229</b>	<b>Endothelin-Converting Enzymes</b>
-	D08.811.277.656.300.480.252	Gelatinases
-	D08.811.277.656.300.480.252.420	Matrix Metalloproteinase 2
-	D08.811.277.656.300.480.252.445	Matrix Metalloproteinase 9
-	D08.811.277.656.300.480.300	Insulysin
-	D08.811.277.656.300.480.452	Lysostaphin
-	D08.811.277.656.300.480.525	Matrix Metalloproteinases
-	D08.811.277.656.300.480.525.300 Associated	Matrix Metalloproteinases, Membrane-Associated
-	D08.811.277.656.300.480.525.300.500	Matrix Metalloproteinase 14
-	D08.811.277.656.300.480.525.300.550	Matrix Metalloproteinase 15
-	D08.811.277.656.300.480.525.300.600	Matrix Metalloproteinase 16
-	D08.811.277.656.300.480.525.300.650	Matrix Metalloproteinase 17
-	D08.811.277.656.300.480.525.700	Matrix Metalloproteinases, Secreted

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D08.811.277.656.300.480.525.700.049 Bone Morphogenetic Protein 1
-	D08.811.277.656.300.480.525.700.100 Matrix Metalloproteinase 1
-	D08.811.277.656.300.480.525.700.150 Matrix Metalloproteinase 2
-	D08.811.277.656.300.480.525.700.200 Matrix Metalloproteinase 3
-	D08.811.277.656.300.480.525.700.250 Matrix Metalloproteinase 7
-	D08.811.277.656.300.480.525.700.300 Matrix Metalloproteinase 8
-	D08.811.277.656.300.480.525.700.350 Matrix Metalloproteinase 9
-	D08.811.277.656.300.480.525.700.400 Matrix Metalloproteinase 10
-	D08.811.277.656.300.480.525.700.450 Matrix Metalloproteinase 11
-	D08.811.277.656.300.480.525.700.500 Matrix Metalloproteinase 12
-	D08.811.277.656.300.480.525.700.550 Matrix Metalloproteinase 13
-	D08.811.277.656.300.480.525.700.800 Matrix Metalloproteinase 20
-	D08.811.277.656.300.480.600 Neprilysin
-	D08.811.277.656.300.480.616 PHEX Phosphate Regulating Neutral Endopeptidase
-	D08.811.277.656.300.480.632 Pregnancy-Associated Plasma Protein-A
-	D08.811.277.656.300.480.664 Procollagen N-Endopeptidase
-	D08.811.277.656.300.480.680 Pronase
-	D08.811.277.656.300.480.827 Thermolysin
-	D08.811.277.656.300.760 Serine Endopeptidases
-	D08.811.277.656.300.760.030 Acrosin
-	D08.811.277.656.300.760.066 Cathepsin G
-	D08.811.277.656.300.760.103 Chymases
-	D08.811.277.656.300.760.176 Chymotrypsin
-	D08.811.277.656.300.760.198 Complement Factor B
-	D08.811.277.656.300.760.200 Complement Factor D
-	D08.811.277.656.300.760.210 Complement Factor I
-	D08.811.277.656.300.760.247 Endopeptidase K
-	D08.811.277.656.300.760.284 Enteropeptidase
-	D08.811.277.656.300.760.300 Factor VIIa
-	D08.811.277.656.300.760.310 Factor IXa
-	D08.811.277.656.300.760.315 Factor Xa
-	D08.811.277.656.300.760.320 Factor XIa
-	D08.811.277.656.300.760.324 Factor XIIa
-	D08.811.277.656.300.760.330 Fibrinolysin
-	D08.811.277.656.300.760.353 Furin

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D08.811.277.656.300.760.397	Granzymes
-	D08.811.277.656.300.760.442	Kallikreins
-	D08.811.277.656.300.760.442.700	Plasma Kallikrein
-	D08.811.277.656.300.760.442.725	Prekallikrein
-	D08.811.277.656.300.760.442.750	Prostate-Specific Antigen
-	D08.811.277.656.300.760.442.875	Tissue Kallikreins
-	D08.811.277.656.300.760.471	Lactoferrin
-	D08.811.277.656.300.760.501 Proteases	Mannose-Binding Protein-Associated Serine
-	D08.811.277.656.300.760.530	Myeloblastin
-	D08.811.277.656.300.760.560	Pancreatic Elastase
-	D08.811.277.656.300.760.560.500	Leukocyte Elastase
-	D08.811.277.656.300.760.635	Plasminogen Activators
-	D08.811.277.656.300.760.635.075	Anistreplase
-	D08.811.277.656.300.760.638	Pronase
-	D08.811.277.656.300.760.640	Proprotein Convertase 1
-	D08.811.277.656.300.760.646	Proprotein Convertase 2
-	D08.811.277.656.300.760.648	Proprotein Convertase 5
New Heading	<b>D08.811.277.656.300.760.718</b>	<b>Proprotein Convertase 9</b>
-	D08.811.277.656.300.760.787	Subtilisins
-	D08.811.277.656.300.760.787.805	Subtilisin
-	D08.811.277.656.300.760.855	Thrombin
-	D08.811.277.656.300.760.875	Tissue Plasminogen Activator
-	D08.811.277.656.300.760.895	Trypsin
-	D08.811.277.656.300.760.902	Tryptases
-	D08.811.277.656.300.760.910	Urokinase-Type Plasminogen Activator
-	D08.811.277.656.300.760.955	Venombin A
-	D08.811.277.656.300.760.955.060	Ancrod
-	D08.811.277.656.300.760.955.135	Batroxobin
-	D08.811.277.656.300.775	Streptokinase
-	D08.811.277.656.300.775.075	Anistreplase
-	D08.811.277.656.300.775.900	Streptodornase and Streptokinase
-	D08.811.277.656.300.887	Ubiquitin-Specific Proteases
Old Tree	<b>D08.811.277.656.300.887.500</b>	<b>Ataxin-3</b>
New Heading	<b>D08.811.277.656.300.887.750 Protein 3</b>	<b>Tumor Necrosis Factor alpha-Induced</b>

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D08.811.277.656.350 Exopeptidases
-	D08.811.277.656.350.100 Aminopeptidases
-	D08.811.277.656.350.100.160 Antigens, CD13
-	D08.811.277.656.350.100.235 Cystinyl Aminopeptidase
-	D08.811.277.656.350.100.373 Glutamyl Aminopeptidase
-	D08.811.277.656.350.100.511 Leucyl Aminopeptidase
-	D08.811.277.656.350.100.633 Methionyl Aminopeptidases
-	D08.811.277.656.350.100.755 Pyroglutamyl-Peptidase I
-	D08.811.277.656.350.245 Carboxypeptidases
-	D08.811.277.656.350.245.083 Carboxypeptidase B
-	D08.811.277.656.350.245.125 Carboxypeptidase B2
-	D08.811.277.656.350.245.167 Carboxypeptidase H
-	D08.811.277.656.350.245.250 Carboxypeptidases A
-	D08.811.277.656.350.245.260 Cathepsin A
-	D08.811.277.656.350.245.270 Cathepsin Z
-	D08.811.277.656.350.245.280 gamma-Glutamyl Hydrolase
-	D08.811.277.656.350.245.400 Glutamate Carboxypeptidase II
-	D08.811.277.656.350.245.450 Lysine Carboxypeptidase
-	D08.811.277.656.350.245.500 Muramoylpentapeptide Carboxypeptidase
-	D08.811.277.656.350.245.800 Serine-Type D-Ala-D-Ala Carboxypeptidase
-	D08.811.277.656.350.297 Dipeptidases
-	D08.811.277.656.350.350 Dipeptidyl-Peptidases and Tripeptidyl-Peptidases
-	D08.811.277.656.350.350.100 Cathepsin C
-	D08.811.277.656.350.350.126 Dipeptidyl Peptidase 4
-	D08.811.277.656.350.350.687 Peptidyl-Dipeptidase A
-	D08.811.277.656.350.555 Metalloexopeptidases
-	D08.811.277.656.350.555.100 Antigens, CD13
-	D08.811.277.656.350.555.200 Carboxypeptidase B
-	D08.811.277.656.350.555.225 Carboxypeptidase B2
-	D08.811.277.656.350.555.250 Carboxypeptidase H
-	D08.811.277.656.350.555.350 Carboxypeptidases A
-	D08.811.277.656.350.555.400 Cystinyl Aminopeptidase
-	D08.811.277.656.350.555.500 Glutamate Carboxypeptidase II
-	D08.811.277.656.350.555.600 Glutamyl Aminopeptidase
-	D08.811.277.656.350.555.700 Leucyl Aminopeptidase
-	D08.811.277.656.350.555.750 Lysine Carboxypeptidase

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D08.811.277.656.675	Metalloproteases
-	D08.811.277.656.675.374	Metalloendopeptidases
-	D08.811.277.656.675.374.102	ADAM Proteins
New Heading	<b>D08.811.277.656.675.374.102.125</b>	<b>ADAM12 Protein</b>
New Heading	<b>D08.811.277.656.675.374.102.250</b>	<b>ADAM10 Protein</b>
New Heading	<b>D08.811.277.656.675.374.102.375</b>	<b>ADAM17 Protein</b>
New Heading	<b>D08.811.277.656.675.374.102.500</b>	<b>ADAMTS Proteins</b>
New Heading	<b>D08.811.277.656.675.374.102.500.500</b>	<b>ADAMTS1 Protein</b>
New Heading	<b>D08.811.277.656.675.374.102.500.813</b>	<b>ADAMTS13 Protein</b>
New Heading	<b>D08.811.277.656.675.374.102.500.844</b>	<b>ADAMTS4 Protein</b>
New Heading	<b>D08.811.277.656.675.374.102.500.875</b>	<b>ADAMTS5 Protein</b>
New Heading	<b>D08.811.277.656.675.374.102.500.937</b>	<b>ADAMTS7 Protein</b>
New Heading	<b>D08.811.277.656.675.374.102.500.968</b>	<b>ADAMTS9 Protein</b>
New Heading	<b>D08.811.277.656.675.374.102.750</b>	<b>Fertilins</b>
-	D08.811.277.656.675.374.153	Botulinum Toxins
-	D08.811.277.656.675.374.153.100	Botulinum Toxins, Type A
-	D08.811.277.656.675.374.205	Collagenases
-	D08.811.277.656.675.374.205.351	Matrix Metalloproteinase 1
-	D08.811.277.656.675.374.205.352	Matrix Metalloproteinase 2
-	D08.811.277.656.675.374.205.358	Matrix Metalloproteinase 8
-	D08.811.277.656.675.374.205.360	Matrix Metalloproteinase 9
-	D08.811.277.656.675.374.205.363	Matrix Metalloproteinase 13
-	D08.811.277.656.675.374.205.500	Microbial Collagenase
New Heading	<b>D08.811.277.656.675.374.229</b>	<b>Endothelin-Converting Enzymes</b>
-	D08.811.277.656.675.374.252	Gelatinases
-	D08.811.277.656.675.374.252.420	Matrix Metalloproteinase 2
-	D08.811.277.656.675.374.252.445	Matrix Metalloproteinase 9
-	D08.811.277.656.675.374.300	Insulysin

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D08.811.277.656.675.374.452                      Lysostaphin
-	D08.811.277.656.675.374.525                      Matrix Metalloproteinases
-	D08.811.277.656.675.374.525.300                      Matrix Metalloproteinases, Membrane-Associated
-	D08.811.277.656.675.374.525.300.500                      Matrix Metalloproteinase 14
-	D08.811.277.656.675.374.525.300.625                      Matrix Metalloproteinase 15
-	D08.811.277.656.675.374.525.300.750                      Matrix Metalloproteinase 16
-	D08.811.277.656.675.374.525.300.875                      Matrix Metalloproteinase 17
-	D08.811.277.656.675.374.525.700                      Matrix Metalloproteinases, Secreted
-	D08.811.277.656.675.374.525.700.100                      Matrix Metalloproteinase 1
-	D08.811.277.656.675.374.525.700.150                      Matrix Metalloproteinase 2
-	D08.811.277.656.675.374.525.700.200                      Matrix Metalloproteinase 3
-	D08.811.277.656.675.374.525.700.250                      Matrix Metalloproteinase 7
-	D08.811.277.656.675.374.525.700.300                      Matrix Metalloproteinase 8
-	D08.811.277.656.675.374.525.700.350                      Matrix Metalloproteinase 9
-	D08.811.277.656.675.374.525.700.400                      Matrix Metalloproteinase 10
-	D08.811.277.656.675.374.525.700.450                      Matrix Metalloproteinase 11
-	D08.811.277.656.675.374.525.700.500                      Matrix Metalloproteinase 12
-	D08.811.277.656.675.374.525.700.550                      Matrix Metalloproteinase 13
-	D08.811.277.656.675.374.525.700.800                      Matrix Metalloproteinase 20
-	D08.811.277.656.675.374.525.700.900                      Tolloid-Like Metalloproteinases
-	D08.811.277.656.675.374.525.700.900.500                      Bone Morphogenetic Protein 1
-	D08.811.277.656.675.374.600                      Neprilysin
-	D08.811.277.656.675.374.616                      PHEX Phosphate Regulating Neutral Endopeptidase
-	D08.811.277.656.675.374.632                      Pregnancy-Associated Plasma Protein-A
-	D08.811.277.656.675.374.664                      Procollagen N-Endopeptidase
-	D08.811.277.656.675.374.680                      Pronase
-	D08.811.277.656.675.374.827                      Thermolysin
-	D08.811.277.656.675.555                      Metalloexopeptidases
-	D08.811.277.656.675.555.100                      Antigens, CD13
-	D08.811.277.656.675.555.200                      Carboxypeptidase B
-	D08.811.277.656.675.555.225                      Carboxypeptidase B2
-	D08.811.277.656.675.555.250                      Carboxypeptidase H
-	D08.811.277.656.675.555.350                      Carboxypeptidases A
-	D08.811.277.656.675.555.400                      Cystinyl Aminopeptidase



## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D08.811.277.656.675.555.500	Glutamate Carboxypeptidase II
-	D08.811.277.656.675.555.600	Glutamyl Aminopeptidase
-	D08.811.277.656.675.555.700	Leucyl Aminopeptidase
-	D08.811.277.656.675.555.750	Lysine Carboxypeptidase
-	D08.811.277.656.837	Proprotein Convertases
-	D08.811.277.656.837.061	Carboxypeptidase B
-	D08.811.277.656.837.092	Carboxypeptidase B2
-	D08.811.277.656.837.124	Carboxypeptidase H
-	D08.811.277.656.837.249	Furin
-	D08.811.277.656.837.500	Proprotein Convertase 1
-	D08.811.277.656.837.562	Proprotein Convertase 2
-	D08.811.277.656.837.625	Proprotein Convertase 5
New Heading	<b>D08.811.277.656.837.688</b>	<b>Proprotein Convertase 9</b>
-	D08.811.277.656.837.750	Renin
-	D08.811.277.656.918	Proteasome Endopeptidase Complex
-	D08.811.277.656.959	Serine Proteases
-	D08.811.277.656.959.200	Cathepsin A
-	D08.811.277.656.959.250	Dipeptidyl Peptidase 4
-	D08.811.277.656.959.350	Serine Endopeptidases
-	D08.811.277.656.959.350.030	Acrosin
-	D08.811.277.656.959.350.066	Cathepsin G
-	D08.811.277.656.959.350.103	Chymases
-	D08.811.277.656.959.350.176	Chymotrypsin
-	D08.811.277.656.959.350.198	Complement Factor B
-	D08.811.277.656.959.350.200	Complement Factor D
-	D08.811.277.656.959.350.210	Complement Factor I
-	D08.811.277.656.959.350.247	Endopeptidase K
-	D08.811.277.656.959.350.284	Enteropeptidase
-	D08.811.277.656.959.350.300	Factor VIIa
-	D08.811.277.656.959.350.310	Factor IXa
-	D08.811.277.656.959.350.315	Factor Xa
-	D08.811.277.656.959.350.320	Factor XIa
-	D08.811.277.656.959.350.324	Factor XIIa
-	D08.811.277.656.959.350.330	Fibrinolysin
-	D08.811.277.656.959.350.353	Furin

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D08.811.277.656.959.350.397	Granzymes
-	D08.811.277.656.959.350.442	Kallikreins
-	D08.811.277.656.959.350.442.700	Plasma Kallikrein
-	D08.811.277.656.959.350.442.725	Prekallikrein
-	D08.811.277.656.959.350.442.750	Prostate-Specific Antigen
-	D08.811.277.656.959.350.442.875	Tissue Kallikreins
-	D08.811.277.656.959.350.471	Lactoferrin
-	D08.811.277.656.959.350.501 Proteases	Mannose-Binding Protein-Associated Serine
-	D08.811.277.656.959.350.530	Myeloblastin
-	D08.811.277.656.959.350.560	Pancreatic Elastase
-	D08.811.277.656.959.350.560.500	Leukocyte Elastase
-	D08.811.277.656.959.350.635	Plasminogen Activators
-	D08.811.277.656.959.350.635.075	Anistreplase
-	D08.811.277.656.959.350.638	Pronase
-	D08.811.277.656.959.350.640	Proprotein Convertase 1
-	D08.811.277.656.959.350.646	Proprotein Convertase 2
-	D08.811.277.656.959.350.648	Proprotein Convertase 5
New Heading	<b>D08.811.277.656.959.350.718</b>	<b>Proprotein Convertase 9</b>
-	D08.811.277.656.959.350.787	Subtilisins
-	D08.811.277.656.959.350.787.805	Subtilisin
-	D08.811.277.656.959.350.855	Thrombin
-	D08.811.277.656.959.350.875	Tissue Plasminogen Activator
-	D08.811.277.656.959.350.895	Trypsin
-	D08.811.277.656.959.350.902	Tryptases
-	D08.811.277.656.959.350.910	Urokinase-Type Plasminogen Activator
-	D08.811.277.656.959.350.955	Venombin A
-	D08.811.277.656.959.350.955.060	Ancrod
-	D08.811.277.656.959.350.955.135	Batroxobin
-	D08.811.277.656.959.560	Serine-Type D-Ala-D-Ala Carboxypeptidase
New Heading	<b>D08.811.277.785</b>	<b>Protein Deglycase DJ-1</b>
-	D08.811.277.913	Ureohydrolases
-	D08.811.277.913.292	Arginase
-	D08.811.348	Isoenzymes
-	D08.811.399	Isomerases

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D08.811.399.325 cis-trans-Isomerases
-	D08.811.399.325.500 Peptidylprolyl Isomerase
-	D08.811.399.325.500.400 Immunophilins
-	D08.811.399.325.500.400.300 Cyclophilins
-	D08.811.399.325.500.400.300.500 Cyclophilin A
New Heading	<b>D08.811.399.325.500.400.300.750 Cyclophilin C</b>
-	D08.811.399.325.500.400.700 Tacrolimus Binding Proteins
-	D08.811.399.325.500.400.700.500 Tacrolimus Binding Protein 1A
New Heading	<b>D08.811.399.325.500.700 NIMA-Interacting Peptidylprolyl Isomerase</b>
New Heading	<b>D08.811.399.325.500.850 Peptidyl-Prolyl Cis-Trans Isomerase NIMA-Interacting 4</b>
-	D08.811.399.340 DNA Helicases
-	D08.811.399.340.124 DnaB Helicases
New Heading	<b>D08.811.399.340.155 Ku Autoantigen</b>
-	D08.811.399.340.186 Minichromosome Maintenance Proteins
-	D08.811.399.340.186.200.2 Minichromosome Maintenance Complex Component
-	D08.811.399.340.186.300.3 Minichromosome Maintenance Complex Component
-	D08.811.399.340.186.400.4 Minichromosome Maintenance Complex Component
-	D08.811.399.340.186.500.5 Minichromosome Maintenance Complex Component
-	D08.811.399.340.186.600.6 Minichromosome Maintenance Complex Component
-	D08.811.399.340.186.700.7 Minichromosome Maintenance Complex Component
-	D08.811.399.340.186.800.8 Minichromosome Maintenance Complex Component
-	D08.811.399.340.186.900.9 Minichromosome Maintenance Complex Component
-	D08.811.399.340.249 RecQ Helicases
New Heading	<b>D08.811.399.340.249.500 Werner Syndrome Helicase</b>
-	D08.811.399.340.500 Xeroderma Pigmentosum Group D Protein
-	D08.811.399.403 DNA Topoisomerases
-	D08.811.399.403.483 DNA Topoisomerases, Type I

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D08.811.399.403.741 DNA Topoisomerases, Type II
-	D08.811.399.403.741.149 DNA Gyrase
-	D08.811.399.403.741.224 DNA Topoisomerase IV
-	D08.811.399.430 Intramolecular Lyases
-	D08.811.399.430.500 Myo-Inositol-1-Phosphate Synthase
-	D08.811.399.475 Intramolecular Oxidoreductases
-	D08.811.399.475.200 Aldose-Ketose Isomerases
-	D08.811.399.475.200.350 Glucose-6-Phosphate Isomerase
-	D08.811.399.475.200.550 Mannose-6-Phosphate Isomerase
-	D08.811.399.475.200.775 Triose-Phosphate Isomerase
-	D08.811.399.475.400 Carbon-Carbon Double Bond Isomerases
-	D08.811.399.475.400.349 Dodecenoyl-CoA Isomerase
-	D08.811.399.475.400.349.500 Peroxisomal Bifunctional Enzyme
-	D08.811.399.475.400.700 Steroid Isomerases
New Heading	<b>D08.811.399.475.600 Prostaglandin-E Synthases</b>
-	D08.811.399.475.800 Sulfur-Sulfur Bond Isomerases
-	D08.811.399.475.800.550 Protein Disulfide-Isomerases
-	D08.811.399.475.900 Thromboxane-A Synthase
-	D08.811.399.520 Intramolecular Transferases
-	D08.811.399.520.100 2-Acetolactate Mutase
-	D08.811.399.520.250 Chorismate Mutase
-	D08.811.399.520.250.500 Prephenate Dehydratase
-	D08.811.399.520.250.750 Prephenate Dehydrogenase
-	D08.811.399.520.625 Methylmalonyl-CoA Mutase
-	D08.811.399.520.750 Phosphotransferases (Phosphomutases)
-	D08.811.399.520.750.250 Bisphosphoglycerate Mutase
-	D08.811.399.520.750.625 Phosphoglucomutase
-	D08.811.399.520.750.700 Phosphoglycerate Mutase
-	D08.811.399.894 Racemases and Epimerases
-	D08.811.399.894.200 Amino Acid Isomerases
-	D08.811.399.894.200.200 Alanine Racemase
-	D08.811.399.894.500 Carbohydrate Epimerases
-	D08.811.399.894.500.700 UDPglucose 4-Epimerase
-	D08.811.464 Ligases
-	D08.811.464.257 Carbon-Carbon Ligases

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D08.811.464.257.050 Acetyl-CoA Carboxylase
-	D08.811.464.257.275 Polyketide Synthases
-	D08.811.464.257.500 Pyruvate Carboxylase
-	D08.811.464.259 Carbon-Nitrogen Ligases
-	D08.811.464.259.100 Adenylosuccinate Synthase
-	D08.811.464.259.200 Amide Synthases
-	D08.811.464.259.200.200 Aspartate-Ammonia Ligase
-	D08.811.464.259.200.600 Glutamate-Ammonia Ligase
-	D08.811.464.259.300 Argininosuccinate Synthase
-	D08.811.464.259.350 Carbamoyl-Phosphate Synthase (Ammonia)
-	D08.811.464.259.400 Carbon-Nitrogen Ligases with Glutamine as Amide-N-Donor
-	D08.811.464.259.400.300 Carbamoyl-Phosphate Synthase (Glutamine-Hydrolyzing)
-	D08.811.464.259.550 Formate-Tetrahydrofolate Ligase
-	D08.811.464.259.850 Peptide Synthases
-	D08.811.464.259.850.400 Glutamate-Cysteine Ligase
-	D08.811.464.259.850.500 Glutathione Synthase
-	D08.811.464.263 Carbon-Oxygen Ligases
-	D08.811.464.263.200 Amino Acyl-tRNA Synthetases
-	D08.811.464.263.200.050 Alanine-tRNA Ligase
-	D08.811.464.263.200.100 Arginine-tRNA Ligase
-	D08.811.464.263.200.150 Aspartate-tRNA Ligase
-	D08.811.464.263.200.250 Glutamate-tRNA Ligase
-	D08.811.464.263.200.350 Glycine-tRNA Ligase
-	D08.811.464.263.200.400 Histidine-tRNA Ligase
-	D08.811.464.263.200.450 Isoleucine-tRNA Ligase
-	D08.811.464.263.200.500 Leucine-tRNA Ligase
-	D08.811.464.263.200.550 Lysine-tRNA Ligase
-	D08.811.464.263.200.600 Methionine-tRNA Ligase
-	D08.811.464.263.200.650 Phenylalanine-tRNA Ligase
-	D08.811.464.263.200.750 Serine-tRNA Ligase
-	D08.811.464.263.200.800 Threonine-tRNA Ligase
-	D08.811.464.263.200.850 Tryptophan-tRNA Ligase
-	D08.811.464.263.200.900 Tyrosine-tRNA Ligase
-	D08.811.464.263.200.950 Valine-tRNA Ligase

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D08.811.464.267 Carbon-Sulfur Ligases
-	D08.811.464.267.500 Coenzyme A Ligases
-	D08.811.464.267.500.200 Acetate-CoA Ligase
-	D08.811.464.267.500.600 Succinate-CoA Ligases
-	D08.811.464.754 Polynucleotide Ligases
-	D08.811.464.754.600 DNA Ligases
New Heading	<b>D08.811.464.754.600.500</b> <b>DNA Ligase ATP</b>
-	D08.811.464.754.720 RNA Ligase (ATP)
-	D08.811.464.938 Ubiquitin-Protein Ligase Complexes
-	D08.811.464.938.249 Ubiquitin-Activating Enzymes
-	D08.811.464.938.500 Ubiquitin-Conjugating Enzymes
-	D08.811.464.938.750 Ubiquitin-Protein Ligases
-	D08.811.464.938.750.092 Anaphase-Promoting Complex-Cyclosome
-	D08.811.464.938.750.092.500 Cyclosome Apc1 Subunit, Anaphase-Promoting Complex-
-	D08.811.464.938.750.092.625 Cyclosome Apc10 Subunit, Anaphase-Promoting Complex-
-	D08.811.464.938.750.092.687 Cyclosome Apc11 Subunit, Anaphase-Promoting Complex-
-	D08.811.464.938.750.092.750 Cyclosome Apc2 Subunit, Anaphase-Promoting Complex-
-	D08.811.464.938.750.092.875 Cyclosome Apc3 Subunit, Anaphase-Promoting Complex-
-	D08.811.464.938.750.092.937 Cyclosome Apc4 Subunit, Anaphase-Promoting Complex-
-	D08.811.464.938.750.092.968 Cyclosome Apc5 Subunit, Anaphase-Promoting Complex-
-	D08.811.464.938.750.092.984 Cyclosome Apc6 Subunit, Anaphase-Promoting Complex-
-	D08.811.464.938.750.092.992 Cyclosome Apc7 Subunit, Anaphase-Promoting Complex-
-	D08.811.464.938.750.092.996 Cyclosome Apc8 Subunit, Anaphase-Promoting Complex-
-	D08.811.464.938.750.092.998 Cdc20 Proteins
-	D08.811.464.938.750.092.999 Cdh1 Proteins
-	D08.811.464.938.750.186 Fanconi Anemia Complementation Group L Protein
-	D08.811.464.938.750.280 Polycomb Repressive Complex 1
-	D08.811.464.938.750.374 Proto-Oncogene Proteins c-cbl
-	D08.811.464.938.750.562 Proto-Oncogene Proteins c-mdm2

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D08.811.464.938.750.656 Receptors, Autocrine Motility Factor
-	D08.811.464.938.750.750 SKP Cullin F-Box Protein Ligases
-	D08.811.464.938.750.750.500 Cullin Proteins
New Heading	<b>D08.811.464.938.750.813 Tumor Necrosis Factor alpha-Induced Protein 3</b>
-	D08.811.464.938.750.875 Von Hippel-Lindau Tumor Suppressor Protein
-	D08.811.520 Lyases
-	D08.811.520.224 Carbon-Carbon Lyases
-	D08.811.520.224.062 Aldehyde-Lyases
-	D08.811.520.224.062.400 Fructose-Bisphosphate Aldolase
-	D08.811.520.224.125 Carboxy-Lyases
-	D08.811.520.224.125.050 Adenosylmethionine Decarboxylase
-	D08.811.520.224.125.100 Aromatic-L-Amino-Acid Decarboxylases
-	D08.811.520.224.125.100.500 Dopa Decarboxylase
-	D08.811.520.224.125.250 Glutamate Decarboxylase
-	D08.811.520.224.125.300 Histidine Decarboxylase
-	D08.811.520.224.125.350 Indole-3-Glycerol-Phosphate Synthase
-	D08.811.520.224.125.387 Methylmalonyl-CoA Decarboxylase
-	D08.811.520.224.125.425 Ornithine Decarboxylase
-	D08.811.520.224.125.450 Orotidine-5'-Phosphate Decarboxylase
-	D08.811.520.224.125.500 Phosphoenolpyruvate Carboxykinase (ATP)
-	D08.811.520.224.125.550 Phosphoenolpyruvate Carboxykinase (GTP)
-	D08.811.520.224.125.650 Phosphoenolpyruvate Carboxylase
-	D08.811.520.224.125.750 Pyruvate Decarboxylase
-	D08.811.520.224.125.800 Ribulose-Bisphosphate Carboxylase
-	D08.811.520.224.125.875 Tyrosine Decarboxylase
-	D08.811.520.224.125.900 Uroporphyrinogen Decarboxylase
-	D08.811.520.224.187 Deoxyribodipyrimidine Photo-Lyase
-	D08.811.520.224.600 Oxo-Acid-Lyases
-	D08.811.520.224.600.200 Anthranilate Synthase
-	D08.811.520.224.600.700 Isocitrate Lyase
-	D08.811.520.224.800 Tryptophanase
-	D08.811.520.224.900 Tyrosine Phenol-Lyase
-	D08.811.520.232 Carbon-Nitrogen Lyases
-	D08.811.520.232.300 Amidine-Lyases
-	D08.811.520.232.300.200 Adenylosuccinate Lyase

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D08.811.520.232.300.400 Argininosuccinate Lyase
-	D08.811.520.232.300.400.500 delta-Crystallins
-	D08.811.520.232.400 Ammonia-Lyases
-	D08.811.520.232.400.200 Aspartate Ammonia-Lyase
-	D08.811.520.232.400.350 Ethanolamine Ammonia-Lyase
-	D08.811.520.232.400.500 Histidine Ammonia-Lyase
-	D08.811.520.232.400.600 L-Serine Dehydratase
-	D08.811.520.232.400.700 Phenylalanine Ammonia-Lyase
-	D08.811.520.232.400.850 Threonine Dehydratase
-	D08.811.520.241 Carbon-Oxygen Lyases
-	D08.811.520.241.225 DNA-(Apurinic or Apyrimidinic Site) Lyase
-	D08.811.520.241.300 Hydro-Lyases
-	D08.811.520.241.300.050 Aconitate Hydratase
-	D08.811.520.241.300.050.500 Iron Regulatory Protein 1
-	D08.811.520.241.300.050.750 Iron Regulatory Protein 2
-	D08.811.520.241.300.150 Carbonic Anhydrases
-	D08.811.520.241.300.150.100 Carbonic Anhydrase I
-	D08.811.520.241.300.150.200 Carbonic Anhydrase II
-	D08.811.520.241.300.150.300 Carbonic Anhydrase III
-	D08.811.520.241.300.150.400 Carbonic Anhydrase IV
New Heading	<b>D08.811.520.241.300.150.450 Carbonic Anhydrase IX</b>
-	D08.811.520.241.300.150.500 Carbonic Anhydrase V
-	D08.811.520.241.300.200 Cystathionine beta-Synthase
-	D08.811.520.241.300.250 Enoyl-CoA Hydratase
-	D08.811.520.241.300.250.249 Mitochondrial Trifunctional Protein
-	D08.811.520.241.300.250.249.500 Mitochondrial Trifunctional Protein, alpha Subunit
-	D08.811.520.241.300.250.750 Peroxisomal Bifunctional Enzyme
-	D08.811.520.241.300.275 Enoyl-CoA Hydratase 2
-	D08.811.520.241.300.275.500 Peroxisomal Multifunctional Protein-2
-	D08.811.520.241.300.287 Fatty Acid Synthases
-	D08.811.520.241.300.287.099 Fatty Acid Synthase, Type I
-	D08.811.520.241.300.287.200 Fatty Acid Synthase, Type II
-	D08.811.520.241.300.300 Fumarate Hydratase
-	D08.811.520.241.300.500 Phosphopyruvate Hydratase



## MeSH Tree Changes for 2017

Type	Tree - heading
-	D08.811.520.241.300.500.500                      tau-Crystallins
-	D08.811.520.241.300.550                      Porphobilinogen Synthase
-	D08.811.520.241.300.600                      Prephenate Dehydratase
-	D08.811.520.241.300.650                      Propanediol Dehydratase
-	D08.811.520.241.300.850                      Tryptophan Synthase
-	D08.811.520.241.300.900                      Urocanate Hydratase
-	D08.811.520.241.300.950                      Uroporphyrinogen III Synthetase
-	D08.811.520.241.700                          Polysaccharide-Lyases
-	D08.811.520.241.700.350                      Chondroitinases and Chondroitin Lyases
-	D08.811.520.241.700.350.500                  Chondroitin Lyases
-	D08.811.520.241.700.350.500.500              Chondroitin ABC Lyase
-	D08.811.520.241.700.512                      Heparin Lyase
-	D08.811.520.241.700.675                      Hyaluronoglucosaminidase
-	D08.811.520.300                                  Carbon-Sulfur Lyases
-	D08.811.520.300.250                              Cystathionine gamma-Lyase
-	D08.811.520.300.500                              Lactoylglutathione Lyase
-	D08.811.520.500                                  Ferrochelataze
-	D08.811.520.650                                  Phosphorus-Oxygen Lyases
-	D08.811.520.650.200                              Adenylyl Cyclases
-	D08.811.520.650.200.040                          Adenylate Cyclase Toxin
-	D08.811.520.650.600                              Guanylate Cyclase
-	D08.811.520.650.600.500                          Receptors, Guanylate Cyclase-Coupled
-	D08.811.520.650.600.500.500                      Receptors, Atrial Natriuretic Factor
New Heading	<b>D08.811.520.650.600.750                          Soluble Guanylyl Cyclase</b>
-	D08.811.520.650.800                              Phosphatidylinositol Diacylglycerol-Lyase
-	D08.811.600    Multienzyme Complexes
-	D08.811.600.075                                      Anthranilate Phosphoribosyltransferase
-	D08.811.600.085                                      Anthranilate Synthase
-	D08.811.600.250                                      Electron Transport Chain Complex Proteins
-	D08.811.600.250.500                                  Electron-Transferring Flavoproteins
-	D08.811.600.250.500.500                              Electron Transport Complex I
-	D08.811.600.250.500.750                              Electron Transport Complex II
-	D08.811.600.250.500.750.500                          Succinate Dehydrogenase
-	D08.811.600.250.687                                  Electron Transport Complex IV
-	D08.811.600.250.875                                  Succinate Cytochrome c Oxidoreductase

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D08.811.600.250.875.249 Electron Transport Complex II
-	D08.811.600.250.875.249.500 Succinate Dehydrogenase
-	D08.811.600.250.875.500 Electron Transport Complex III
-	D08.811.600.283 Exosome Multienzyme Ribonuclease Complex
-	D08.811.600.354 Fatty Acid Synthase, Type II
-	D08.811.600.391 Glycine Decarboxylase Complex
-	D08.811.600.391.100 Aminomethyltransferase
-	D08.811.600.391.150 Dihydrolipoamide Dehydrogenase
-	D08.811.600.391.175 Glycine Decarboxylase Complex H-Protein
-	D08.811.600.391.200 Glycine Dehydrogenase (Decarboxylating)
-	D08.811.600.465 Ketoglutarate Dehydrogenase Complex
-	D08.811.600.465.500 Dihydrolipoamide Dehydrogenase
-	D08.811.600.620 Mi-2 Nucleosome Remodeling and Deacetylase Complex
-	D08.811.600.620.100 Histone Deacetylase 1
-	D08.811.600.620.200 Histone Deacetylase 2
-	D08.811.600.620.775 Retinoblastoma-Binding Protein 4
-	D08.811.600.620.887 Retinoblastoma-Binding Protein 7
-	D08.811.600.700 Phosphoenolpyruvate Sugar Phosphotransferase System
-	D08.811.600.710 Photosynthetic Reaction Center Complex Proteins
-	D08.811.600.710.374 Cytochrome b6f Complex
-	D08.811.600.710.374.500 Cytochromes b6
-	D08.811.600.710.374.750 Cytochromes f
-	D08.811.600.710.374.875 Plastoquinol-Plastocyanin Reductase
-	D08.811.600.710.490 Light-Harvesting Protein Complexes
-	D08.811.600.710.490.249 Chlorophyll Binding Proteins
-	D08.811.600.710.490.500 Phycobilisomes
-	D08.811.600.710.490.500.500 Phycobiliproteins
-	D08.811.600.710.490.500.500.755 Phycocyanin
-	D08.811.600.710.490.500.500.777 Phycoerythrin
-	D08.811.600.710.500 Photosystem I Protein Complex
-	D08.811.600.710.750 Photosystem II Protein Complex
-	D08.811.600.715 Polyketide Synthases
-	D08.811.600.720 Prostaglandin-Endoperoxide Synthases
-	D08.811.600.720.500 Cyclooxygenase 1
-	D08.811.600.720.750 Cyclooxygenase 2
-	D08.811.600.730 Proteasome Endopeptidase Complex

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D08.811.600.741 Pyruvate Dehydrogenase Complex
-	D08.811.600.741.525 Dihydrolipoamide Dehydrogenase
-	D08.811.600.741.625 Dihydrolipoyllysine-Residue Acetyltransferase
-	D08.811.600.741.725 Pyruvate Dehydrogenase (Lipoamide)
-	D08.811.600.795 Sin3 Histone Deacetylase and Corepressor Complex
-	D08.811.600.795.100 Histone Deacetylase 1
-	D08.811.600.795.200 Histone Deacetylase 2
-	D08.811.600.795.775 Retinoblastoma-Binding Protein 4
-	D08.811.600.795.887 Retinoblastoma-Binding Protein 7
-	D08.811.600.850 Sucrase-Isomaltase Complex
-	D08.811.600.896 Tryptophan Synthase
-	D08.811.641 Multifunctional Enzymes
-	D08.811.641.249 Acetyl-CoA Carboxylase
-	D08.811.641.260 N-Acetyllactosamine Synthase
-	D08.811.641.500 Aspartate Carbamoyltransferase
-	D08.811.641.750 Aspartokinase Homoserine Dehydrogenase
-	D08.811.641.752 Cyclin-Dependent Kinase 9
-	D08.811.641.755 Dual-Specificity Phosphatases
-	D08.811.641.755.100 cdc25 Phosphatases
-	D08.811.641.755.200 Dual Specificity Phosphatase 1
-	D08.811.641.755.250 Dual Specificity Phosphatase 2
-	D08.811.641.755.300 Dual Specificity Phosphatase 3
-	D08.811.641.755.600 Dual Specificity Phosphatase 6
-	D08.811.641.758 Fatty Acid Synthase, Type I
-	D08.811.641.761 Lactase-Phlorizin Hydrolase
-	D08.811.641.765 Mitochondrial Trifunctional Protein
-	D08.811.641.765.200 Mitochondrial Trifunctional Protein, alpha Subunit
-	D08.811.641.765.500 Mitochondrial Trifunctional Protein, beta Subunit
-	D08.811.641.773 Peroxiredoxin VI
-	D08.811.641.781 Peroxisomal Bifunctional Enzyme
-	D08.811.641.812 Peroxisomal Multifunctional Protein-2
-	D08.811.641.843 Phosphofructokinase-2
-	D08.811.682 Oxidoreductases
-	D08.811.682.047 Alcohol Oxidoreductases
-	D08.811.682.047.150 Carbohydrate Dehydrogenases
-	D08.811.682.047.150.225 Fructuronate Reductase

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D08.811.682.047.150.250 Galactose Dehydrogenases
-	D08.811.682.047.150.270 Glucose Dehydrogenases
-	D08.811.682.047.150.270.500 Glucose 1-Dehydrogenase
-	D08.811.682.047.150.300 Glucosephosphate Dehydrogenase
-	D08.811.682.047.150.600 Phosphogluconate Dehydrogenase
-	D08.811.682.047.150.650 Phosphoglycerate Dehydrogenase
-	D08.811.682.047.150.700 Sugar Alcohol Dehydrogenases
-	D08.811.682.047.150.700.075 Aldehyde Reductase
-	D08.811.682.047.150.700.237 D-Xylulose Reductase
-	D08.811.682.047.150.700.400 Glycerolphosphate Dehydrogenase
-	D08.811.682.047.150.700.400.500 (NAD+) Glycerol-3-Phosphate Dehydrogenase
-	D08.811.682.047.150.700.437 L-Gulonolactone Oxidase
-	D08.811.682.047.150.700.475 L-Iditol 2-Dehydrogenase
-	D08.811.682.047.150.700.649 Mannitol Dehydrogenases
-	D08.811.682.047.150.900 Uridine Diphosphate Glucose Dehydrogenase
-	D08.811.682.047.180 Choline Dehydrogenase
-	D08.811.682.047.210 Galactose Oxidase
-	D08.811.682.047.239 Glucose Oxidase
-	D08.811.682.047.436 Hydroxysteroid Dehydrogenases
-	D08.811.682.047.436.174 11-beta-Hydroxysteroid Dehydrogenases
-	D08.811.682.047.436.174.300 11-beta-Hydroxysteroid Dehydrogenase Type 1
-	D08.811.682.047.436.174.600 11-beta-Hydroxysteroid Dehydrogenase Type 2
-	D08.811.682.047.436.350 3-Hydroxysteroid Dehydrogenases
-	D08.811.682.047.436.350.100 3-alpha-Hydroxysteroid Dehydrogenase (B-Specific)
-	D08.811.682.047.436.350.150 Cholesterol Oxidase
-	D08.811.682.047.436.350.700 Progesterone Reductase
-	D08.811.682.047.436.375 17-Hydroxysteroid Dehydrogenases
-	D08.811.682.047.436.375.280 Estradiol Dehydrogenases
-	D08.811.682.047.436.375.640 Peroxisomal Multifunctional Protein-2
-	D08.811.682.047.551 Lactate Dehydrogenases
-	D08.811.682.047.551.249 epsilon-Crystallins
-	D08.811.682.047.551.400 L-Lactate Dehydrogenase
-	D08.811.682.047.551.500 L-Lactate Dehydrogenase (Cytochrome)
-	D08.811.682.047.820 NAD (+) and NADP (+) Dependent Alcohol Oxidoreductases

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D08.811.682.047.820.100 11-beta-Hydroxysteroid Dehydrogenases
-	D08.811.682.047.820.100.300 11-beta-Hydroxysteroid Dehydrogenase Type 1
-	D08.811.682.047.820.100.600 11-beta-Hydroxysteroid Dehydrogenase Type 2
-	D08.811.682.047.820.125 20-Hydroxysteroid Dehydrogenases
-	D08.811.682.047.820.125.074 20-alpha-Hydroxysteroid Dehydrogenase
-	D08.811.682.047.820.125.150 Cortisone Reductase
-	D08.811.682.047.820.150 3-Hydroxyacyl CoA Dehydrogenases
-	D08.811.682.047.820.150.207 3-Hydroxyacyl-CoA Dehydrogenase
-	D08.811.682.047.820.150.207.500 Peroxisomal Bifunctional Enzyme
-	D08.811.682.047.820.150.415 Hydroxymethylglutaryl CoA Reductases
-	D08.811.682.047.820.150.415.250 NAD-Dependent Hydroxymethylglutaryl-CoA Reductases,
-	D08.811.682.047.820.150.415.750 NADP-dependent Hydroxymethylglutaryl-CoA-Reductases,
-	D08.811.682.047.820.150.707 Long-Chain-3-Hydroxyacyl-CoA Dehydrogenase
-	D08.811.682.047.820.150.707.249 Mitochondrial Trifunctional Protein
-	D08.811.682.047.820.150.707.249.500 Mitochondrial Trifunctional Protein, alpha Subunit
-	D08.811.682.047.820.150.926 Peroxisomal Multifunctional Protein-2
-	D08.811.682.047.820.186 3-alpha-Hydroxysteroid Dehydrogenase (B-Specific)
-	D08.811.682.047.820.193 3-Isopropylmalate Dehydrogenase
-	D08.811.682.047.820.196 3-Oxoacyl-(Acyl-Carrier-Protein) Reductase
-	D08.811.682.047.820.196.500 Fatty Acid Synthases
-	D08.811.682.047.820.196.500.099 Fatty Acid Synthase, Type I
-	D08.811.682.047.820.196.500.200 Fatty Acid Synthase, Type II
-	D08.811.682.047.820.200 Acetoin Dehydrogenase
-	D08.811.682.047.820.250 Alcohol Dehydrogenase
-	D08.811.682.047.820.275 Aldehyde Reductase
-	D08.811.682.047.820.293 Glycerol-3-Phosphate Dehydrogenase (NAD+)
-	D08.811.682.047.820.300 Homoserine Dehydrogenase
-	D08.811.682.047.820.300.060 Aspartokinase Homoserine Dehydrogenase
-	D08.811.682.047.820.350 Hydroxybutyrate Dehydrogenase
-	D08.811.682.047.820.375 Hydroxyprostaglandin Dehydrogenases
-	D08.811.682.047.820.400 Hydroxypyruvate Reductase
-	D08.811.682.047.820.450 IMP Dehydrogenase
-	D08.811.682.047.820.475 Isocitrate Dehydrogenase

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D08.811.682.047.820.487                      Ketol-Acid Reductoisomerase
-	D08.811.682.047.820.493                      L-Lactate Dehydrogenase
-	D08.811.682.047.820.496                      Malate Dehydrogenase
-	D08.811.682.047.820.498                      Malate Dehydrogenase (NADP+)
-	D08.811.682.047.820.500                      Progesterone Reductase
-	D08.811.682.047.820.800                      D-Xylulose Reductase
-	D08.811.682.047.892                              Xanthine Dehydrogenase
-	D08.811.682.047.928                              Xanthine Oxidase
-	D08.811.682.113                                      Arsenate Reductases
-	D08.811.682.180                                      Ascorbate Oxidase
-	D08.811.682.226                                      Ceruloplasmin
-	D08.811.682.285                                      Electron Transport Complex IV
-	D08.811.682.400                                      Hydrogenase
-	D08.811.682.494                                      Laccase
-	D08.811.682.517                                      Luciferases
-	D08.811.682.517.500                              Luciferases, Bacterial
-	D08.811.682.517.750                              Luciferases, Firefly
-	D08.811.682.517.875                              Luciferases, Renilla
-	D08.811.682.550                                      5,10-Methylenetetrahydrofolate Reductase (FADH2)
-	D08.811.682.608                                      NADH, NADPH Oxidoreductases
-	D08.811.682.608.047                              Apoptosis Inducing Factor
-	D08.811.682.608.191                              Cytochrome Reductases
-	D08.811.682.608.191.237                              Cytochrome-B(5) Reductase
-	D08.811.682.608.191.500                              NADPH-Ferrihemoprotein Reductase
-	D08.811.682.608.504                              Electron Transport Complex I
-	D08.811.682.608.504.500                              NADH Dehydrogenase
-	D08.811.682.608.530                              NADH Tetrazolium Reductase
-	D08.811.682.608.540                              NADP Transhydrogenases
-	D08.811.682.608.540.500                              NADP Transhydrogenase, AB-Specific
-	D08.811.682.608.540.750                              NADP Transhydrogenase, B-Specific
-	D08.811.682.608.550                              NADPH Dehydrogenase
-	D08.811.682.608.575                              NADPH Oxidase
-	D08.811.682.608.800                              Quinone Reductases
-	D08.811.682.608.800.500                              NAD(P)H Dehydrogenase (Quinone)
-	D08.811.682.608.800.500.500                              zeta-Crystallins
-	D08.811.682.647                                      Nitrogenase

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D08.811.682.647.200 Dinitrogenase Reductase
-	D08.811.682.647.550 Molybdoferredoxin
-	D08.811.682.655 Nitroreductases
-	D08.811.682.655.249 GMP Reductase
-	D08.811.682.655.500 Nitrate Reductases
-	D08.811.682.655.500.124 Nitrate Reductase
-	D08.811.682.655.500.200 Nitrate Reductase (NADH)
-	D08.811.682.655.500.249 Nitrate Reductase (NAD(P)H)
-	D08.811.682.655.500.374 Nitrate Reductase (NADPH)
-	D08.811.682.655.750 Nitrite Reductases
-	D08.811.682.655.750.249 Ferredoxin-Nitrite Reductase
-	D08.811.682.655.750.500 Nitrite Reductase (NAD(P)H)
-	D08.811.682.657 Oxidoreductases Acting on Aldehyde or Oxo Group Donors
-	D08.811.682.657.163 Aldehyde Oxidoreductases
-	D08.811.682.657.163.249 Aldehyde Dehydrogenase
New Heading	<b>D08.811.682.657.163.249.375 Aldehyde Dehydrogenase, Mitochondrial</b>
-	D08.811.682.657.163.249.750 omega-Crystallins
-	D08.811.682.657.163.311 Aldehyde Oxidase
-	D08.811.682.657.163.342 Aminomuconate-Semialdehyde Dehydrogenase
-	D08.811.682.657.163.374 Aspartate-Semialdehyde Dehydrogenase
-	D08.811.682.657.163.468 Benzaldehyde Dehydrogenase (NADP+)
-	D08.811.682.657.163.515 Betaine-Aldehyde Dehydrogenase
-	D08.811.682.657.163.562 Glutamate-5-Semialdehyde Dehydrogenase
-	D08.811.682.657.163.750 Glycerinaldehyde-3-Phosphate Dehydrogenases
-	D08.811.682.657.163.750.250 Glycerinaldehyde 3-Phosphate Dehydrogenase (NADP+)
-	D08.811.682.657.163.750.300 Glycerinaldehyde-3-Phosphate Dehydrogenase (NADP+)(Phosphorylating)
-	D08.811.682.657.163.750.350 Glycerinaldehyde-3-Phosphate Dehydrogenase (Phosphorylating)
-	D08.811.682.657.163.781 Glycolaldehyde Dehydrogenase
-	D08.811.682.657.163.796 L-Amino adipate-Semialdehyde Dehydrogenase
-	D08.811.682.657.163.812 Malonate-Semialdehyde Dehydrogenase (Acetylating)
-	D08.811.682.657.163.827 Methylmalonate-Semialdehyde Dehydrogenase (Acylyating)
-	D08.811.682.657.163.835 Retinal Dehydrogenase

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D08.811.682.657.163.843 Succinate-Semialdehyde Dehydrogenase
-	D08.811.682.657.163.875 Succinate-Semialdehyde Dehydrogenase (NADP+)
-	D08.811.682.657.180 Formate Dehydrogenases
-	D08.811.682.657.350 Ketone Oxidoreductases
-	D08.811.682.657.350.750 Ketoglutarate Dehydrogenase Complex
-	D08.811.682.657.350.750.500 Dihydrolipoamide Dehydrogenase
-	D08.811.682.657.350.760 (Lipoamide) 3-Methyl-2-Oxobutanoate Dehydrogenase
-	D08.811.682.657.350.825 2-Oxoisovalerate Dehydrogenase (Acylation)
-	D08.811.682.657.350.875 Pyruvate Dehydrogenase (Lipoamide)
-	D08.811.682.657.350.937 Pyruvate Oxidase
-	D08.811.682.657.350.968 Pyruvate Synthase
-	D08.811.682.660 Oxidoreductases Acting on CH-CH Group Donors
-	D08.811.682.660.150 Acyl-CoA Dehydrogenases
-	D08.811.682.660.150.100 Acyl-CoA Dehydrogenase
-	D08.811.682.660.150.150 Acyl-CoA Dehydrogenase, Long-Chain
-	D08.811.682.660.150.200 Acyl-CoA Oxidase
-	D08.811.682.660.150.300 Butyryl-CoA Dehydrogenase
-	D08.811.682.660.200 Cholestenone 5 alpha-Reductase
-	D08.811.682.660.250 Coproporphyrinogen Oxidase
-	D08.811.682.660.275 Dihydrodipicolinate Reductase
-	D08.811.682.660.300 Dihydroorotate Oxidase
-	D08.811.682.660.325 Dihydrouracil Dehydrogenase (NAD+)
-	D08.811.682.660.350 Dihydrouracil Dehydrogenase (NADP)
-	D08.811.682.660.385 Electron Transport Complex II
-	D08.811.682.660.385.500 Succinate Dehydrogenase
-	D08.811.682.660.387 Enoyl-(Acyl-Carrier-Protein) Reductase (NADH)
-	D08.811.682.660.390 Enoyl-(Acyl-Carrier Protein) Reductase (NADPH, B-Specific)
-	D08.811.682.660.425 Glutaryl-CoA Dehydrogenase
-	D08.811.682.660.462 Isovaleryl-CoA Dehydrogenase
-	D08.811.682.660.465 3-Oxo-5-alpha-Steroid 4-Dehydrogenase
-	D08.811.682.660.490 15-Oxoprostaglandin 13-Reductase
-	D08.811.682.660.500 Prephenate Dehydrogenase
-	D08.811.682.660.600 Protoporphyrinogen Oxidase
-	D08.811.682.662 Oxidoreductases Acting on CH-NH Group Donors



## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D08.811.682.662.171	FMN Reductase
-	D08.811.682.662.217	Methylenetetrahydrofolate Dehydrogenase (NAD+)
-	D08.811.682.662.253	Methylenetetrahydrofolate Dehydrogenase (NADP)
-	D08.811.682.662.290	Methylenetetrahydrofolate Reductase (NADPH2)
-	D08.811.682.662.582	Oxidoreductases, N-Demethylating
New Heading	<b>D08.811.682.662.582.138</b>	<b>AlkB Homolog 4, Lysine Demethylase</b>
New Heading	<b>D08.811.682.662.582.207</b>	<b>AlkB Homolog 5, RNA Demethylase</b>
New Heading	<b>D08.811.682.662.582.242</b> <b>FTO</b>	<b>Alpha-Ketoglutarate-Dependent Dioxygenase</b>
-	D08.811.682.662.582.276	Aminopyrine N-Demethylase
-	D08.811.682.662.582.338	Cytochrome P-450 CYP2E1
-	D08.811.682.662.582.353	Cytochrome P-450 CYP3A
-	D08.811.682.662.582.361	Dihydropteridine Reductase
-	D08.811.682.662.582.369	Dimethylglycine Dehydrogenase
-	D08.811.682.662.582.400	Ethylmorphine-N-Demethylase
-	D08.811.682.662.582.475	Histone Demethylases
-	D08.811.682.662.582.475.500	Jumonji Domain-Containing Histone Demethylases
-	D08.811.682.662.582.475.500.500	Retinoblastoma-Binding Protein 2
-	D08.811.682.662.582.550	Sarcosine Dehydrogenase
-	D08.811.682.662.582.700	Sarcosine Oxidase
-	D08.811.682.662.640	Proline Oxidase
-	D08.811.682.662.680	Pyridoxaminephosphate Oxidase
-	D08.811.682.662.693	1-Pyrroline-5-Carboxylate Dehydrogenase
-	D08.811.682.662.695	Pyrroline Carboxylate Reductases
-	D08.811.682.662.750	Saccharopine Dehydrogenases
-	D08.811.682.662.825	Tetrahydrofolate Dehydrogenase
-	D08.811.682.664	Oxidoreductases Acting on CH-NH2 Group Donors
-	D08.811.682.664.249	Amine Oxidase (Copper-Containing)
-	D08.811.682.664.500	Amino Acid Oxidoreductases
-	D08.811.682.664.500.062	Alanine Dehydrogenase
-	D08.811.682.664.500.125	D-Amino-Acid Oxidase
-	D08.811.682.664.500.261	D-Aspartate Oxidase
-	D08.811.682.664.500.398	Glutamate Dehydrogenase
-	D08.811.682.664.500.410	Glutamate Dehydrogenase (NADP+)

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D08.811.682.664.500.470      Glutamate Synthase
-	D08.811.682.664.500.484      Glutamate Synthase (NADH)
-	D08.811.682.664.500.498      Glycine Decarboxylase Complex
-	D08.811.682.664.500.498.500      Glycine Dehydrogenase (Decarboxylating)
-	D08.811.682.664.500.526      Glycine Dehydrogenase
-	D08.811.682.664.500.677      L-Amino Acid Oxidase
-	D08.811.682.664.500.724      Leucine Dehydrogenase
-	D08.811.682.664.500.772      Nitric Oxide Synthase
-	D08.811.682.664.500.772.249      Nitric Oxide Synthase Type I
-	D08.811.682.664.500.772.500      Nitric Oxide Synthase Type II
-	D08.811.682.664.500.772.750      Nitric Oxide Synthase Type III
-	D08.811.682.664.500.810      Proline Oxidase
-	D08.811.682.664.500.848      Protein-Lysine 6-Oxidase
-	D08.811.682.664.500.924      Valine Dehydrogenase (NADP+)
-	D08.811.682.664.750      Monoamine Oxidase
-	D08.811.682.664.750.100      Benzylamine Oxidase
-	D08.811.682.667      Oxidoreductases Acting on Sulfur Group Donors
-	D08.811.682.667.061      Dihydrolipoamide Dehydrogenase
-	D08.811.682.667.076      Ferredoxin-NADP Reductase
-	D08.811.682.667.084      Glutaredoxins
-	D08.811.682.667.092      Glutathione Reductase
-	D08.811.682.667.124      Hydrogensulfite Reductase
-	D08.811.682.667.186      Protein Disulfide Reductase (Glutathione)
-	D08.811.682.667.217      Sulfite Dehydrogenase
-	D08.811.682.667.249      Sulfite Oxidase
-	D08.811.682.667.374      Sulfite Reductase (Ferredoxin)
-	D08.811.682.667.500      Sulfite Reductase (NADPH)
-	D08.811.682.667.750      Thioredoxin-Disulfide Reductase
-	D08.811.682.667.750.500      Thioredoxin Reductase 1
-	D08.811.682.667.750.750      Thioredoxin Reductase 2
-	D08.811.682.670      Oxidoreductases, O-Demethylating
-	D08.811.682.670.550      Nitroanisole O-Demethylase
-	D08.811.682.690      Oxygenases
-	D08.811.682.690.416      Dioxygenases
New Heading	<b>D08.811.682.690.416.139      AlkB Enzymes</b>

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Heading	<b>D08.811.682.690.416.139.500</b>	<b>AlkB Homolog 1, Histone H2a Dioxygenase</b>
New Heading	<b>D08.811.682.690.416.139.750</b> <b>Dependent Dioxygenase</b>	<b>AlkB Homolog 2, Alpha-Ketoglutarate-</b>
New Heading	<b>D08.811.682.690.416.139.875</b> <b>Dependent Dioxygenase</b>	<b>AlkB Homolog 3, Alpha-Ketoglutarate-</b>
New Heading	<b>D08.811.682.690.416.139.937</b>	<b>AlkB Homolog 4, Lysine Demethylase</b>
New Heading	<b>D08.811.682.690.416.139.968</b>	<b>AlkB Homolog 5, RNA Demethylase</b>
New Heading	<b>D08.811.682.690.416.139.984</b>	<b>AlkB Homolog 8, tRNA Methyltransferase</b>
New Heading	<b>D08.811.682.690.416.139.992</b> <b>Dioxygenase FTO</b>	<b>Alpha-Ketoglutarate-Dependent</b>
-	D08.811.682.690.416.277	Catechol 1,2-Dioxygenase
-	D08.811.682.690.416.305	Catechol 2,3-Dioxygenase
-	D08.811.682.690.416.319	Cysteine Dioxygenase
-	D08.811.682.690.416.326	Homogentisate 1,2-Dioxygenase
-	D08.811.682.690.416.328	3-Hydroxyanthranilate 3,4-Dioxygenase
-	D08.811.682.690.416.330	4-Hydroxyphenylpyruvate Dioxygenase
-	D08.811.682.690.416.333	Indoleamine-Pyrrole 2,3,-Dioxygenase
-	D08.811.682.690.416.388	Jumonji Domain-Containing Histone Demethylases
-	D08.811.682.690.416.388.200	Retinoblastoma-Binding Protein 2
-	D08.811.682.690.416.583	Lipoxygenases
-	D08.811.682.690.416.583.500	Arachidonate Lipoxygenases
-	D08.811.682.690.416.583.500.055	Arachidonate 5-Lipoxygenase
-	D08.811.682.690.416.583.500.060	Arachidonate 12-Lipoxygenase
-	D08.811.682.690.416.583.500.065	Arachidonate 15-Lipoxygenase
-	D08.811.682.690.416.583.625	Lipoxygenase
-	D08.811.682.690.416.617	Prolyl Hydroxylases
-	D08.811.682.690.416.617.500	Hypoxia-Inducible Factor-Proline Dioxygenases
-	D08.811.682.690.416.617.750	Procollagen-Proline Dioxygenase
-	D08.811.682.690.416.652	Protocatechuate-3,4-Dioxygenase
-	D08.811.682.690.416.722	Tryptophan Oxygenase
-	D08.811.682.690.562	Inositol Oxygenase
-	D08.811.682.690.708	Mixed Function Oxygenases
-	D08.811.682.690.708.062	Benzoate 4-Monooxygenase
-	D08.811.682.690.708.125	Catechol Oxidase

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D08.811.682.690.708.125.500	Monophenol Monooxygenase
-	D08.811.682.690.708.170	Cytochrome P-450 Enzyme System
New Tree	<a href="#">D08.811.682.690.708.170.010</a>	<a href="#">Aryl Hydrocarbon Hydroxylases</a>
New Tree	<a href="#">D08.811.682.690.708.170.010.024</a>	<a href="#">7-Alkoxy coumarin O-Dealkylase</a>
New Tree	<a href="#">D08.811.682.690.708.170.010.050</a>	<a href="#">Aniline Hydroxylase</a>
New Tree	<a href="#">D08.811.682.690.708.170.010.110</a>	<a href="#">Benzopyrene Hydroxylase</a>
New Tree	<a href="#">D08.811.682.690.708.170.010.277</a>	<a href="#">Cytochrome P-450 CYP1A1</a>
New Tree	<a href="#">D08.811.682.690.708.170.010.443</a>	<a href="#">Cytochrome P-450 CYP1A2</a>
New Tree	<a href="#">D08.811.682.690.708.170.010.500</a>	<a href="#">Cytochrome P-450 CYP1B1</a>
New Tree	<a href="#">D08.811.682.690.708.170.010.550</a>	<a href="#">Cytochrome P-450 CYP2B1</a>
New Tree	<a href="#">D08.811.682.690.708.170.010.575</a>	<a href="#">Cytochrome P-450 CYP2B6</a>
New Tree	<a href="#">D08.811.682.690.708.170.010.590</a>	<a href="#">Cytochrome P-450 CYP2C8</a>
New Tree	<a href="#">D08.811.682.690.708.170.010.600</a>	<a href="#">Cytochrome P-450 CYP2D6</a>
New Tree	<a href="#">D08.811.682.690.708.170.012</a>	<a href="#">Camphor 5-Monooxygenase</a>
New Heading	<b><a href="#">D08.811.682.690.708.170.020</a></b>	<b><a href="#">Cytochrome P450 Family 1</a></b>
New Tree	<a href="#">D08.811.682.690.708.170.020.500</a>	<a href="#">Cytochrome P-450 CYP1A1</a>
New Tree	<a href="#">D08.811.682.690.708.170.020.750</a>	<a href="#">Cytochrome P-450 CYP1A2</a>
New Tree	<a href="#">D08.811.682.690.708.170.020.875</a>	<a href="#">Cytochrome P-450 CYP1B1</a>
Old Tree	<del><a href="#">D08.811.682.690.708.170.040</a></del>	<del><a href="#">Aryl Hydrocarbon Hydroxylases</a></del>
Old Tree	<del><a href="#">D08.811.682.690.708.170.040.024</a></del>	<del><a href="#">7-Alkoxy coumarin O-Dealkylase</a></del>
Old Tree	<del><a href="#">D08.811.682.690.708.170.040.050</a></del>	<del><a href="#">Aniline Hydroxylase</a></del>
Old Tree	<del><a href="#">D08.811.682.690.708.170.040.110</a></del>	<del><a href="#">Benzopyrene Hydroxylase</a></del>
Old Tree	<del><a href="#">D08.811.682.690.708.170.040.332</a></del>	<del><a href="#">Cytochrome P-450 CYP1A1</a></del>
Old Tree	<del><a href="#">D08.811.682.690.708.170.040.443</a></del>	<del><a href="#">Cytochrome P-450 CYP1A2</a></del>
Old Tree	<del><a href="#">D08.811.682.690.708.170.040.500</a></del>	<del><a href="#">Cytochrome P-450 CYP1B1</a></del>
Old Tree	<del><a href="#">D08.811.682.690.708.170.040.550</a></del>	<del><a href="#">Cytochrome P-450 CYP2B1</a></del>

## MeSH Tree Changes for 2017

Type	Tree - heading	
Old Tree	D08.811.682.690.708.170.040.575	Cytochrome P-450 CYP2B6
Old Tree	D08.811.682.690.708.170.040.590	Cytochrome P-450 CYP2C8
Old Tree	D08.811.682.690.708.170.040.600	Cytochrome P-450 CYP2D6
Old Tree	D08.811.682.690.708.170.085	Camphor 5-Monooxygenase
Old Tree	D08.811.682.690.708.170.100	Cytochrome P-450 CYP2A6
Old Tree	D08.811.682.690.708.170.200	Cytochrome P-450 CYP2E1
Old Tree	D08.811.682.690.708.170.250	Cytochrome P-450 CYP3A
Old Tree	D08.811.682.690.708.170.300	Cytochrome P-450 CYP4A
New Heading	<b>D08.811.682.690.708.170.425</b>	<b>Cytochrome P450 Family 11</b>
New Tree	D08.811.682.690.708.170.425.250	Cholesterol Side-Chain Cleavage Enzyme
New Tree	D08.811.682.690.708.170.425.500	Cytochrome P-450 CYP11B2
New Tree	D08.811.682.690.708.170.425.750	Steroid 11-beta-Hydroxylase
New Heading	<b>D08.811.682.690.708.170.438</b>	<b>Cytochrome P450 Family 12</b>
New Heading	<b>D08.811.682.690.708.170.444</b>	<b>Cytochrome P450 Family 17</b>
New Tree	D08.811.682.690.708.170.444.500	Steroid 17-alpha-Hydroxylase
New Heading	<b>D08.811.682.690.708.170.447</b>	<b>Cytochrome P450 Family 19</b>
New Tree	D08.811.682.690.708.170.447.500	Aromatase
New Heading	<b>D08.811.682.690.708.170.450</b>	<b>Cytochrome P450 Family 2</b>
New Tree	D08.811.682.690.708.170.450.250	Cytochrome P-450 CYP2A6
New Tree	D08.811.682.690.708.170.450.313	Cytochrome P-450 CYP2B1
New Tree	D08.811.682.690.708.170.450.344	Cytochrome P-450 CYP2B6
New Tree	D08.811.682.690.708.170.450.360	Cytochrome P-450 CYP2C8
New Tree	D08.811.682.690.708.170.450.368	Cytochrome P-450 CYP2D6
New Tree	D08.811.682.690.708.170.450.375	Cytochrome P-450 CYP2E1
New Tree	D08.811.682.690.708.170.450.500	Limonene Hydroxylases

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	D08.811.682.690.708.170.450.500.500	Cytochrome P-450 CYP2C9
New Tree	D08.811.682.690.708.170.450.500.700	Cytochrome P-450 CYP2C19
New Heading	<b>D08.811.682.690.708.170.463</b>	<b>Cytochrome P450 Family 21</b>
New Tree	D08.811.682.690.708.170.463.500	Steroid 21-Hydroxylase
New Heading	<b>D08.811.682.690.708.170.469</b>	<b>Cytochrome P450 Family 24</b>
New Tree	D08.811.682.690.708.170.469.500	Vitamin D3 24-Hydroxylase
New Heading	<b>D08.811.682.690.708.170.485</b>	<b>Cytochrome P450 Family 26</b>
New Heading	<b>D08.811.682.690.708.170.485.500</b>	<b>Retinoic Acid 4-Hydroxylase</b>
New Heading	<b>D08.811.682.690.708.170.493</b>	<b>Cytochrome P450 Family 27</b>
New Tree	D08.811.682.690.708.170.493.500	25-Hydroxyvitamin D3 1-alpha-Hydroxylase
New Tree	D08.811.682.690.708.170.493.750	Cholestanetriol 26-Monooxygenase
New Heading	<b>D08.811.682.690.708.170.495</b>	<b>Cytochrome P450 Family 3</b>
New Tree	D08.811.682.690.708.170.495.500	Cytochrome P-450 CYP3A
New Heading	<b>D08.811.682.690.708.170.496</b>	<b>Cytochrome P450 Family 4</b>
New Tree	D08.811.682.690.708.170.496.500	Cytochrome P-450 CYP4A
New Heading	<b>D08.811.682.690.708.170.497</b>	<b>Cytochrome P450 Family 46</b>
New Heading	<b>D08.811.682.690.708.170.497.500</b>	<b>Cholesterol 24-Hydroxylase</b>
New Heading	<b>D08.811.682.690.708.170.499</b>	<b>Cytochrome P450 Family 51</b>
New Tree	D08.811.682.690.708.170.499.500	Sterol 14-Demethylase
Old Tree	D08.811.682.690.708.170.500	Limonene Hydroxylases
Old Tree	D08.811.682.690.708.170.500.500	Cytochrome P-450 CYP2C9
Old Tree	D08.811.682.690.708.170.500.700	Cytochrome P-450 CYP2C19
New Heading	<b>D08.811.682.690.708.170.880</b>	<b>Cytochrome P450 Family 6</b>

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Heading	<b>D08.811.682.690.708.170.890</b>	<b>Cytochrome P450 Family 7</b>
New Tree	D08.811.682.690.708.170.890.500	Cholesterol 7-alpha-Hydroxylase
New Heading	<b>D08.811.682.690.708.170.900</b>	<b>Cytochrome P450 Family 8</b>
New Tree	D08.811.682.690.708.170.900.500	Steroid 12-alpha-Hydroxylase
-	D08.811.682.690.708.170.915	Steroid Hydroxylases
New Heading	<b>D08.811.682.690.708.170.915.025</b>	<b>Cholesterol 24-Hydroxylase</b>
-	D08.811.682.690.708.170.915.050	Cytochrome P-450 CYP11B2
-	D08.811.682.690.708.170.915.099	Aromatase
-	D08.811.682.690.708.170.915.150	Cholestanetriol 26-Monooxygenase
-	D08.811.682.690.708.170.915.200	Cholesterol 7-alpha-Hydroxylase
-	D08.811.682.690.708.170.915.212	Cholesterol Side-Chain Cleavage Enzyme
-	D08.811.682.690.708.170.915.400	25-Hydroxyvitamin D3 1-alpha-Hydroxylase
-	D08.811.682.690.708.170.915.730	Steroid 12-alpha-Hydroxylase
-	D08.811.682.690.708.170.915.737	Steroid 16-alpha-Hydroxylase
-	D08.811.682.690.708.170.915.748	Steroid 17-alpha-Hydroxylase
-	D08.811.682.690.708.170.915.750	Steroid 11-beta-Hydroxylase
-	D08.811.682.690.708.170.915.760	Steroid 21-Hydroxylase
-	D08.811.682.690.708.170.915.880	Sterol 14-Demethylase
Old Tree	<b>D08.811.682.690.708.170.957</b>	<b>Vitamin D3 24-Hydroxylase</b>
-	D08.811.682.690.708.292	Dopamine beta-Hydroxylase
-	D08.811.682.690.708.392	Fatty Acid Desaturases
-	D08.811.682.690.708.392.312	beta-Carotene 15,15'-Monooxygenase
-	D08.811.682.690.708.392.468	Linoleoyl-CoA Desaturase
-	D08.811.682.690.708.392.625	Stearoyl-CoA Desaturase
-	D08.811.682.690.708.401	gamma-Butyrobetaine Dioxygenase
-	D08.811.682.690.708.410	Heme Oxygenase (Decyclizing)
-	D08.811.682.690.708.410.500	Heme Oxygenase-1
-	D08.811.682.690.708.425	4-Hydroxybenzoate-3-Monooxygenase
-	D08.811.682.690.708.557	Kynurenine 3-Monooxygenase
-	D08.811.682.690.708.601	Phenylalanine Hydroxylase
-	D08.811.682.690.708.660	Procollagen-Lysine, 2-Oxoglutarate 5-Dioxygenase
-	D08.811.682.690.708.694	Prolyl Hydroxylases

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D08.811.682.690.708.694.500 Hypoxia-Inducible Factor-Proline Dioxygenases
-	D08.811.682.690.708.694.750 Procollagen-Proline Dioxygenase
-	D08.811.682.690.708.715 Prostaglandin-Endoperoxide Synthases
-	D08.811.682.690.708.749 Squalene Monooxygenase
-	D08.811.682.690.708.826 Trans-Cinnamate 4-Monooxygenase
-	D08.811.682.690.708.870 Tryptophan Hydroxylase
-	D08.811.682.690.708.923 Tyrosine 3-Monooxygenase
-	D08.811.682.690.708.961 Vitamin K Epoxide Reductases
-	D08.811.682.730 Methionine Sulfoxide Reductases
-	D08.811.682.732 Peroxidases
-	D08.811.682.732.165 Ascorbate Peroxidases
-	D08.811.682.732.332 Catalase
-	D08.811.682.732.360 Chloride Peroxidase
-	D08.811.682.732.380 Cytochrome-c Peroxidase
-	D08.811.682.732.440 Eosinophil Peroxidase
-	D08.811.682.732.500 Glutathione Peroxidase
-	D08.811.682.732.512 Horseradish Peroxidase
-	D08.811.682.732.512.900 Wheat Germ Agglutinin-Horseradish Peroxidase Conjugate
-	D08.811.682.732.525 Iodide Peroxidase
-	D08.811.682.732.550 Lactoperoxidase
-	D08.811.682.732.700 Peroxidase
-	D08.811.682.732.850 Peroxiredoxins
-	D08.811.682.732.850.249 Peroxiredoxin III
-	D08.811.682.732.850.500 Peroxiredoxin VI
-	D08.811.682.771 Plastoquinol-Plastocyanin Reductase
-	D08.811.682.810 Ribonucleotide Reductases
-	D08.811.682.810.567 Ribonucleoside Diphosphate Reductase
-	D08.811.682.830 Succinate Cytochrome c Oxidoreductase
-	D08.811.682.830.249 Electron Transport Complex II
-	D08.811.682.830.249.500 Succinate Dehydrogenase
-	D08.811.682.830.500 Electron Transport Complex III
-	D08.811.682.881 Superoxide Dismutase
New Heading	<b>D08.811.682.881.500 Superoxide Dismutase-1</b>
-	D08.811.682.943 Urate Oxidase



## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D08.811.710	Penicillin-Binding Proteins
-	D08.811.739	Recombinases
-	D08.811.739.250	Holliday Junction Resolvases
-	D08.811.739.500	Integrases
-	D08.811.739.500.667	Transposases
-	D08.811.739.500.667.500	HIV Integrase
-	D08.811.739.650	Rec A Recombinases
-	D08.811.739.650.500	Rad51 Recombinase
-	D08.811.739.800	Transposon Resolvases
-	D08.811.739.900	VDJ Recombinases
-	D08.811.797	RNA, Catalytic
-	D08.811.797.500	Ribonuclease P
-	D08.811.797.750	RNA, Ribosomal, Self-Splicing
-	D08.811.913	Transferases
-	D08.811.913.050	Acyltransferases
-	D08.811.913.050.080	Acetyl-CoA C-Acyltransferase
-	D08.811.913.050.080.249	Mitochondrial Trifunctional Protein
-	D08.811.913.050.080.249.500	Mitochondrial Trifunctional Protein, beta Subunit
-	D08.811.913.050.134	Acetyltransferases
-	D08.811.913.050.134.025	Acetyl-CoA C-Acetyltransferase
-	D08.811.913.050.134.029	Acyl-Carrier Protein S-Acetyltransferase
-	D08.811.913.050.134.029.500	Fatty Acid Synthases
-	D08.811.913.050.134.029.500.100	Fatty Acid Synthase, Type I
-	D08.811.913.050.134.029.500.200	Fatty Acid Synthase, Type II
-	D08.811.913.050.134.105	Amino-Acid N-Acetyltransferase
-	D08.811.913.050.134.127	Arylalkylamine N-Acetyltransferase
-	D08.811.913.050.134.138	Arylamine N-Acetyltransferase
-	D08.811.913.050.134.150	Carnitine O-Acetyltransferase
-	D08.811.913.050.134.170	Chloramphenicol O-Acetyltransferase
-	D08.811.913.050.134.180	Choline O-Acetyltransferase
-	D08.811.913.050.134.310	Dihydrolipoyllysine-Residue Acetyltransferase
-	D08.811.913.050.134.375	Glucosamine 6-Phosphate N-Acetyltransferase
Old Tree	D08.811.913.050.134.407	Histone Acetyltransferases
Old Tree	D08.811.913.050.134.407.049	CLOCK Proteins
Old Tree	D08.811.913.050.134.407.074	N-Terminal Acetyltransferase D
Old Tree	D08.811.913.050.134.407.100	Nuclear Receptor Coactivator 1

## MeSH Tree Changes for 2017

Type	Tree - heading	
Old Tree	<a href="#">D08.811.913.050.134.407.150</a>	<b>Nuclear Receptor Coactivator 3</b>
New Heading	<b><a href="#">D08.811.913.050.134.415</a></b>	<b>Lysine Acetyltransferases</b>
New Tree	<a href="#">D08.811.913.050.134.415.500</a>	Histone Acetyltransferases
New Tree	<a href="#">D08.811.913.050.134.415.500.049</a>	CLOCK Proteins
New Tree	<a href="#">D08.811.913.050.134.415.500.074</a>	N-Terminal Acetyltransferase D
New Tree	<a href="#">D08.811.913.050.134.415.500.100</a>	Nuclear Receptor Coactivator 1
New Tree	<a href="#">D08.811.913.050.134.415.500.150</a>	Nuclear Receptor Coactivator 3
New Tree	<a href="#">D08.811.913.050.134.415.500.575</a>	p300-CBP Transcription Factors
New Tree	<a href="#">D08.811.913.050.134.415.500.575.249</a>	CREB-Binding Protein
New Tree	<a href="#">D08.811.913.050.134.415.500.575.600</a>	E1A-Associated p300 Protein
-	<a href="#">D08.811.913.050.134.423</a>	N-Terminal Acetyltransferases
-	<a href="#">D08.811.913.050.134.423.100</a>	N-Terminal Acetyltransferase A
-	<a href="#">D08.811.913.050.134.423.200</a>	N-Terminal Acetyltransferase B
-	<a href="#">D08.811.913.050.134.423.300</a>	N-Terminal Acetyltransferase C
-	<a href="#">D08.811.913.050.134.423.400</a>	N-Terminal Acetyltransferase D
-	<a href="#">D08.811.913.050.134.423.500</a>	N-Terminal Acetyltransferase E
-	<a href="#">D08.811.913.050.134.423.600</a>	N-Terminal Acetyltransferase F
Old Tree	<a href="#">D08.811.913.050.134.440</a>	<b>p300-CBP Transcription Factors</b>
Old Tree	<a href="#">D08.811.913.050.134.440.249</a>	<b>CREB-Binding Protein</b>
Old Tree	<a href="#">D08.811.913.050.134.440.600</a>	<b>E1A-Associated p300 Protein</b>
-	<a href="#">D08.811.913.050.134.700</a>	Phosphate Acetyltransferase
-	<a href="#">D08.811.913.050.134.850</a>	Serine O-Acetyltransferase
-	<a href="#">D08.811.913.050.170</a>	Acyl-Carrier Protein S-Malonyltransferase
-	<a href="#">D08.811.913.050.170.500</a>	Fatty Acid Synthases
-	<a href="#">D08.811.913.050.170.500.100</a>	Fatty Acid Synthase, Type I
-	<a href="#">D08.811.913.050.170.500.200</a>	Fatty Acid Synthase, Type II
-	<a href="#">D08.811.913.050.173</a>	1-Acylglycerol-3-Phosphate O-Acyltransferase
-	<a href="#">D08.811.913.050.175</a>	1-Acylglycerophosphocholine O-Acyltransferase
-	<a href="#">D08.811.913.050.200</a>	Aminoacyltransferases
-	<a href="#">D08.811.913.050.200.400</a>	gamma-Glutamylcyclotransferase

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D08.811.913.050.200.500                      gamma-Glutamyltransferase
-	D08.811.913.050.200.700                      Peptidyl Transferases
-	D08.811.913.050.200.800                      Transglutaminases
-	D08.811.913.050.200.800.300                      Factor XIIIa
-	D08.811.913.050.276                      5-Aminolevulinate Synthetase
-	D08.811.913.050.331                      ATP Citrate (pro-S)-Lyase
-	D08.811.913.050.350                      Carnitine Acyltransferases
-	D08.811.913.050.350.170                      Carnitine O-Acetyltransferase
-	D08.811.913.050.350.200                      Carnitine O-Palmitoyltransferase
-	D08.811.913.050.368                      Citrate (si)-Synthase
-	D08.811.913.050.387                      Diacylglycerol O-Acyltransferase
-	D08.811.913.050.425                      Glycerol-3-Phosphate O-Acyltransferase
-	D08.811.913.050.600                      Homoserine O-Succinyltransferase
-	D08.811.913.050.612                      Hydroxymethylglutaryl-CoA Synthase
-	D08.811.913.050.614                      2-Isopropylmalate Synthase
-	D08.811.913.050.618                      Malate Synthase
-	D08.811.913.050.622                      3-Oxoacyl-(Acyl-Carrier-Protein) Synthase
-	D08.811.913.050.625                      Phosphatidylcholine-Sterol O-Acyltransferase
-	D08.811.913.050.646                      Retinol O-Fatty-Acyltransferase
-	D08.811.913.050.668                      Serine C-Palmitoyltransferase
-	D08.811.913.050.712                      Sphingosine N-Acyltransferase
-	D08.811.913.050.799                      Sterol O-Acyltransferase
-	D08.811.913.200                      Aldehyde-Ketone Transferases
-	D08.811.913.200.324                      Acetolactate Synthase
-	D08.811.913.200.650                      Transaldolase
-	D08.811.913.200.825                      Transketolase
-	D08.811.913.225                      Alkyl and Aryl Transferases
-	D08.811.913.225.224                      Cysteine Synthase
-	D08.811.913.225.250                      3-Deoxy-7-Phosphoheptulonate Synthase
-	D08.811.913.225.300                      Dihydropteroate Synthase
-	D08.811.913.225.400                      Dimethylallyltranstransferase
-	D08.811.913.225.431                      Farnesyl-Diphosphate Farnesyltransferase
-	D08.811.913.225.437                      Farnesyltranstransferase
-	D08.811.913.225.443                      Geranylgeranyl-Diphosphate Geranylgeranyltransferase
-	D08.811.913.225.450                      Geranyltranstransferase
-	D08.811.913.225.500                      Glutathione Transferase

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D08.811.913.225.500.500                                  Glutathione S-Transferase pi
-	D08.811.913.225.575    Hydroxymethylbilane Synthase
-	D08.811.913.225.650    Methionine Adenosyltransferase
-	D08.811.913.225.735    3-Phosphoshikimate 1-Carboxyvinyltransferase
-	D08.811.913.225.750    Riboflavin Synthase
-	D08.811.913.225.825    Spermidine Synthase
-	D08.811.913.225.912    Spermine Synthase
-	D08.811.913.400    Glycosyltransferases
-	D08.811.913.400.100    N-Acetylhexosaminyltransferases
-	D08.811.913.400.100.200                                      N-Acetylgalactosaminyltransferases
-	D08.811.913.400.100.200.300                                Fucosyl Galactose alpha-N-Acetylgalactosaminyltransferase
-	D08.811.913.400.100.250                                      N-Acetylglucosaminyltransferases
-	D08.811.913.400.450    Hexosyltransferases
-	D08.811.913.400.450.300                                      Fucosyltransferases
-	D08.811.913.400.450.400                                      Galactosyltransferases
-	D08.811.913.400.450.400.450                                beta-N-Acetylglucosaminylglycopeptide
-	D08.811.913.400.450.400.450                                beta-1,4-Galactosyltransferase
-	D08.811.913.400.450.400.460                                N-Acylsphingosine Galactosyltransferase
-	D08.811.913.400.450.400.475                                Ganglioside Galactosyltransferase
-	D08.811.913.400.450.400.500                                Lactose Synthase
-	D08.811.913.400.450.400.500.100                            N-Acetyllactosamine Synthase
-	D08.811.913.400.450.460                                      Glucosyltransferases
-	D08.811.913.400.450.460.100                                1,4-alpha-Glucan Branching Enzyme
-	D08.811.913.400.450.460.200                                Chitin Synthase
-	D08.811.913.400.450.460.350                                Glycogen Debranching Enzyme System
-	D08.811.913.400.450.460.375                                Glycogen Synthase
-	D08.811.913.400.450.460.400                                Phosphorylases
-	D08.811.913.400.450.460.400.186                            Glycogen Phosphorylase
-	D08.811.913.400.450.460.400.186.061                      Glycogen Phosphorylase, Brain Form
-	D08.811.913.400.450.460.400.186.124                      Glycogen Phosphorylase, Liver Form
-	D08.811.913.400.450.460.400.186.312                      Glycogen Phosphorylase, Muscle Form
-	D08.811.913.400.450.460.400.280                            Phosphorylase a
-	D08.811.913.400.450.460.400.327                            Phosphorylase b
-	D08.811.913.400.450.460.400.374                            Starch Phosphorylase
-	D08.811.913.400.450.460.750                                Starch Synthase

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D08.811.913.400.450.480                      Glucuronosyltransferase
-	D08.811.913.400.450.560                      Mannosyltransferases
-	D08.811.913.400.450.780                      Peptidoglycan Glycosyltransferase
-	D08.811.913.400.725                            Pentosyltransferases
-	D08.811.913.400.725.100                      Adenine Phosphoribosyltransferase
-	D08.811.913.400.725.115                      ADP Ribose Transferases
-	D08.811.913.400.725.115.180                      Cholera Toxin
-	D08.811.913.400.725.115.220                      Diphtheria Toxin
-	D08.811.913.400.725.115.660                      NAD+ Nucleosidase
-	D08.811.913.400.725.115.660.060                      ADP-ribosyl Cyclase
-	D08.811.913.400.725.115.680                      Pertussis Toxin
-	D08.811.913.400.725.115.690                      Poly(ADP-ribose) Polymerases
New Heading	<b>D08.811.913.400.725.115.690.420                      Poly (ADP-Ribose) Polymerase-1</b>
-	D08.811.913.400.725.115.690.840                      Tankyrases
-	D08.811.913.400.725.115.961                      Sirtuins
-	D08.811.913.400.725.115.961.200                      Sirtuin 2
-	D08.811.913.400.725.115.961.600                      Sirtuin 3
-	D08.811.913.400.725.130                      Amidophosphoribosyltransferase
-	D08.811.913.400.725.160                      Anthranilate Phosphoribosyltransferase
-	D08.811.913.400.725.200                      ATP Phosphoribosyltransferase
-	D08.811.913.400.725.450                      Hypoxanthine Phosphoribosyltransferase
-	D08.811.913.400.725.575                      Nicotinamide Phosphoribosyltransferase
-	D08.811.913.400.725.700                      Orotate Phosphoribosyltransferase
-	D08.811.913.400.725.800                      Purine-Nucleoside Phosphorylase
-	D08.811.913.400.725.850                      Pyrimidine Phosphorylases
-	D08.811.913.400.725.850.500                      Thymidine Phosphorylase
-	D08.811.913.400.725.850.750                      Uridine Phosphorylase
-	D08.811.913.400.800                            Sialyltransferases
-	D08.811.913.477                                Nitrogenous Group Transferases
-	D08.811.913.477.700                            Transaminases
-	D08.811.913.477.700.100                      Alanine Transaminase
-	D08.811.913.477.700.120                      2-Aminoadipate Transaminase
-	D08.811.913.477.700.200                      4-Aminobutyrate Transaminase
-	D08.811.913.477.700.225                      Aspartate Aminotransferases
-	D08.811.913.477.700.225.249                      Aspartate Aminotransferase, Cytoplasmic

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D08.811.913.477.700.225.500 Aspartate Aminotransferase, Mitochondrial
-	D08.811.913.477.700.286 beta-Alanine-Pyruvate Transaminase
-	D08.811.913.477.700.347 D-Alanine Transaminase
-	D08.811.913.477.700.470 Glutamate Synthase
-	D08.811.913.477.700.500 (Isomerizing) Glutamine-Fructose-6-Phosphate Transaminase
-	D08.811.913.477.700.525 Glycine Transaminase
-	D08.811.913.477.700.535 Leucine Transaminase
-	D08.811.913.477.700.550 L-Lysine 6-Transaminase
-	D08.811.913.477.700.700 Ornithine-Oxo-Acid Transaminase
-	D08.811.913.477.700.800 Succinyldiaminopimelate Transaminase
-	D08.811.913.477.700.850 Tryptophan Transaminase
-	D08.811.913.477.700.900 Tyrosine Transaminase
-	D08.811.913.555 One-Carbon Group Transferases
-	D08.811.913.555.150 Amidinotransferases
-	D08.811.913.555.275 Carboxyl and Carbamoyl Transferases
-	D08.811.913.555.275.200 Aspartate Carbamoyltransferase
-	D08.811.913.555.275.600 Ornithine Carbamoyltransferase
-	D08.811.913.555.400 Hydroxymethyl and Formyl Transferases
-	D08.811.913.555.400.100 Aminomethyltransferase
-	D08.811.913.555.400.300 Glutamate Formimidoyltransferase
-	D08.811.913.555.400.500 Glycine Hydroxymethyltransferase
-	D08.811.913.555.400.625 Phosphoribosylaminoimidazolecarboxamide Formyltransferase
-	D08.811.913.555.400.750 Phosphoribosylglycinamide Formyltransferase
-	D08.811.913.555.500 Methyltransferases
-	D08.811.913.555.500.100 Acetylserotonin O-Methyltransferase
-	D08.811.913.555.500.175 Betaine-Homocysteine S-Methyltransferase
-	D08.811.913.555.500.250 Catechol O-Methyltransferase
-	D08.811.913.555.500.350 DNA Modification Methylases
-	D08.811.913.555.500.350.100 DNA-Cytosine Methylases
-	D08.811.913.555.500.350.100.500 DNA (Cytosine-5-)-Methyltransferase
-	D08.811.913.555.500.350.100.750 (Cytosine-N(4)-Specific) Site-Specific DNA-Methyltransferase
-	D08.811.913.555.500.350.700 Site-Specific DNA-Methyltransferase (Adenine-Specific)
-	D08.811.913.555.500.387 Glycine N-Methyltransferase

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D08.811.913.555.500.425	Guanidinoacetate N-Methyltransferase
-	D08.811.913.555.500.500	Histamine N-Methyltransferase
-	D08.811.913.555.500.625	Homocysteine S-Methyltransferase
-	D08.811.913.555.500.645	5-Methyltetrahydrofolate-Homocysteine S-Methyltransferase
-	D08.811.913.555.500.650	Nicotinamide N-Methyltransferase
-	D08.811.913.555.500.700	Phenylethanolamine N-Methyltransferase
-	D08.811.913.555.500.710	Phosphatidyl-N-Methylethanolamine N-Methyltransferase
-	D08.811.913.555.500.712	Phosphatidylethanolamine N-Methyltransferase
-	D08.811.913.555.500.800	Protein Methyltransferases
-	D08.811.913.555.500.800.400	Histone-Lysine N-Methyltransferase
-	D08.811.913.555.500.800.400.500	Polycomb Repressive Complex 2
New Heading	<b>D08.811.913.555.500.800.400.500.500</b>	<b>Enhancer of Zeste Homolog 2 Protein</b>
-	D08.811.913.555.500.800.650	O(6)-Methylguanine-DNA Methyltransferase
-	D08.811.913.555.500.800.750	Protein-Arginine N-Methyltransferases
-	D08.811.913.555.500.800.800	Protein O-Methyltransferase
-	D08.811.913.555.500.800.800.700	Protein D-Aspartate-L-Isoaspartate Methyltransferase
-	D08.811.913.555.500.862	Thymidylate Synthase
-	D08.811.913.555.500.925	tRNA Methyltransferases
New Heading	<b>D08.811.913.555.500.925.500</b>	<b>AlkB Homolog 8, tRNA Methyltransferase</b>
-	D08.811.913.696	Phosphotransferases
-	D08.811.913.696.175	Diphosphotransferases
-	D08.811.913.696.175.300	GTP Pyrophosphokinase
-	D08.811.913.696.175.650	Ribose-Phosphate Pyrophosphokinase
-	D08.811.913.696.175.825	Thiamin Pyrophosphokinase
-	D08.811.913.696.310	Myosin Type III
-	D08.811.913.696.445	Nucleotidyltransferases
-	D08.811.913.696.445.035	N-Acylneuraminate Cytidyltransferase
-	D08.811.913.696.445.184	Choline-Phosphate Cytidyltransferase
-	D08.811.913.696.445.308	DNA Nucleotidyltransferases
-	D08.811.913.696.445.308.300	DNA-Directed DNA Polymerase
-	D08.811.913.696.445.308.300.112	DNA Polymerase beta
-	D08.811.913.696.445.308.300.225	DNA Polymerase I

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D08.811.913.696.445.308.300.230	DNA Polymerase II
-	D08.811.913.696.445.308.300.235	DNA Polymerase III
-	D08.811.913.696.445.308.300.750	RNA-Directed DNA Polymerase
-	D08.811.913.696.445.308.300.750.187	HIV Reverse Transcriptase
-	D08.811.913.696.445.308.300.750.750	Telomerase
-	D08.811.913.696.445.308.300.875	Taq Polymerase
-	D08.811.913.696.445.308.325	DNA Nucleotidyltransferase
-	D08.811.913.696.445.400	Glucose-1-Phosphate Adenylyltransferase
-	D08.811.913.696.445.600	Nicotinamide-Nucleotide Adenylyltransferase
-	D08.811.913.696.445.625	2',5'-Oligoadenylate Synthetase
-	D08.811.913.696.445.650	Polynucleotide Adenylyltransferase
-	D08.811.913.696.445.692	Rec A Recombinases
-	D08.811.913.696.445.735	RNA Nucleotidyltransferases
-	D08.811.913.696.445.735.265	DNA, Catalytic
-	D08.811.913.696.445.735.270	DNA-Directed RNA Polymerases
-	D08.811.913.696.445.735.270.375	DNA Primase
-	D08.811.913.696.445.735.270.750	RNA Polymerase I
-	D08.811.913.696.445.735.270.762	RNA Polymerase II
-	D08.811.913.696.445.735.270.775	RNA Polymerase III
-	D08.811.913.696.445.735.270.887	RNA Polymerase Sigma 54
-	D08.811.913.696.445.735.532	Polyribonucleotide Nucleotidyltransferase
-	D08.811.913.696.445.735.630	Q beta Replicase
-	D08.811.913.696.445.735.720	RNA Helicases
-	D08.811.913.696.445.735.720.249	DEAD-box RNA Helicases
-	D08.811.913.696.445.735.720.249.500	DEAD Box Protein 20
New Heading	<b>D08.811.913.696.445.735.720.249.750</b>	<b>DEAD Box Protein 58</b>
New Heading	<b>D08.811.913.696.445.735.720.249.875</b>	<b>Interferon-Induced Helicase, IFIH1</b>
-	D08.811.913.696.445.735.720.500	Eukaryotic Initiation Factor-4A
-	D08.811.913.696.445.735.780	RNA Replicase
-	D08.811.913.696.445.735.917	RNA, Ribosomal, Self-Splicing
-	D08.811.913.696.445.800	Sulfate Adenylyltransferase
-	D08.811.913.696.445.825	Transposases
-	D08.811.913.696.445.825.500	HIV Integrase
-	D08.811.913.696.445.837	Transposon Resolvases



## MeSH Tree Changes for 2017

Type	Tree - heading
-	D08.811.913.696.445.850 Uridyltransferase
-	D08.811.913.696.445.875 UDPglucose-Hexose-1-Phosphate Uridyltransferase
-	D08.811.913.696.445.900 UTP-Glucose-1-Phosphate Uridyltransferase
-	D08.811.913.696.445.950 UTP-Hexose-1-Phosphate Uridyltransferase
-	D08.811.913.696.445.950 VDJ Recombinases
-	D08.811.913.696.620 Phosphotransferases (Alcohol Group Acceptor)
-	D08.811.913.696.620.010 Adenosine Kinase
-	D08.811.913.696.620.155 Choline Kinase
-	D08.811.913.696.620.175 Deoxycytidine Kinase
-	D08.811.913.696.620.200 Diacylglycerol Kinase
-	D08.811.913.696.620.225 Fructokinases
-	D08.811.913.696.620.225.850 Phosphofructokinases
-	D08.811.913.696.620.225.850.500 Phosphofructokinase-1
-	D08.811.913.696.620.225.850.500.249 Phosphofructokinase-1, Liver Type
-	D08.811.913.696.620.225.850.500.500 Phosphofructokinase-1, Muscle Type
-	D08.811.913.696.620.225.850.500.750 Phosphofructokinase-1, Type C
-	D08.811.913.696.620.225.850.750 Phosphofructokinase-2
-	D08.811.913.696.620.240 Galactokinase
-	D08.811.913.696.620.250 Glucokinase
-	D08.811.913.696.620.275 Glycerol Kinase
-	D08.811.913.696.620.300 Hexokinase
-	D08.811.913.696.620.475 Kanamycin Kinase
-	D08.811.913.696.620.500 Phosphatidylinositol 3-Kinases
-	D08.811.913.696.620.500.100 Phosphatidylinositol 3-Kinase
-	D08.811.913.696.620.500.100.100 Class I Phosphatidylinositol 3-Kinases
-	D08.811.913.696.620.500.100.100.100 Class Ia Phosphatidylinositol 3-Kinase
-	D08.811.913.696.620.500.100.100.200 Class Ib Phosphatidylinositol 3-Kinase
-	D08.811.913.696.620.500.100.200 Class II Phosphatidylinositol 3-Kinases
-	D08.811.913.696.620.500.100.300 Class III Phosphatidylinositol 3-Kinases
-	D08.811.913.696.620.500.100.300.500 Vacuolar Sorting Protein VPS15
-	D08.811.913.696.620.500.200 Phosphatidylinositol-4-Phosphate 3-Kinase
-	D08.811.913.696.620.500.200.100 Class I Phosphatidylinositol 3-Kinases
-	D08.811.913.696.620.500.200.100.100 Class Ia Phosphatidylinositol 3-Kinase
-	D08.811.913.696.620.500.200.100.200 Class Ib Phosphatidylinositol 3-Kinase
-	D08.811.913.696.620.500.200.200 Class II Phosphatidylinositol 3-Kinases
-	D08.811.913.696.620.550 1-Phosphatidylinositol 4-Kinase

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D08.811.913.696.620.650 System	Phosphoenolpyruvate Sugar Phosphotransferase
-	D08.811.913.696.620.680	Polynucleotide 5'-Hydroxyl-Kinase
-	D08.811.913.696.620.682	Protein Kinases
-	D08.811.913.696.620.682.324	Connectin
-	D08.811.913.696.620.682.650	Phosphorylase Kinase
New Heading	<b>D08.811.913.696.620.682.675</b>	<b>Histidine Kinase</b>
-	D08.811.913.696.620.682.700	Protein-Serine-Threonine Kinases
-	D08.811.913.696.620.682.700.062	Activin Receptors
-	D08.811.913.696.620.682.700.062.500	Activin Receptors, Type I
-	D08.811.913.696.620.682.700.062.750	Activin Receptors, Type II
-	D08.811.913.696.620.682.700.085	AMP-Activated Protein Kinases
-	D08.811.913.696.620.682.700.097	Ataxia Telangiectasia Mutated Proteins
-	D08.811.913.696.620.682.700.103	Aurora Kinases
-	D08.811.913.696.620.682.700.103.500	Aurora Kinase A
-	D08.811.913.696.620.682.700.103.750	Aurora Kinase B
-	D08.811.913.696.620.682.700.103.875	Aurora Kinase C
New Heading	<b>D08.811.913.696.620.682.700.108</b>	<b>Autophagy-Related Protein-1 Homolog</b>
-	D08.811.913.696.620.682.700.109	Bone Morphogenetic Protein Receptors
-	D08.811.913.696.620.682.700.109.500 Receptors, Type I	Bone Morphogenetic Protein
-	D08.811.913.696.620.682.700.109.750 Receptors, Type II	Bone Morphogenetic Protein
-	D08.811.913.696.620.682.700.125 Kinases	Calcium-Calmodulin-Dependent Protein
-	D08.811.913.696.620.682.700.125.049 Protein Kinase Kinase	Calcium-Calmodulin-Dependent
-	D08.811.913.696.620.682.700.125.100 Protein Kinase Type 1	Calcium-Calmodulin-Dependent
-	D08.811.913.696.620.682.700.125.200 Protein Kinase Type 2	Calcium-Calmodulin-Dependent
-	D08.811.913.696.620.682.700.125.350 Protein Kinase Type 4	Calcium-Calmodulin-Dependent
-	D08.811.913.696.620.682.700.125.387	Death-Associated Protein Kinases
-	D08.811.913.696.620.682.700.125.425	Elongation Factor 2 Kinase
-	D08.811.913.696.620.682.700.125.500	Myosin-Light-Chain Kinase
-	D08.811.913.696.620.682.700.140	Casein Kinases

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D08.811.913.696.620.682.700.140.300	Casein Kinase I
-	D08.811.913.696.620.682.700.140.300.100	Casein Kinase Ialpha
-	D08.811.913.696.620.682.700.140.300.200	Casein Kinase Idelta
-	D08.811.913.696.620.682.700.140.300.300	Casein Kinase Iepsilon
-	D08.811.913.696.620.682.700.140.600	Casein Kinase II
New Heading	<b>D08.811.913.696.620.682.700.143</b>	<b>Checkpoint Kinase 1</b>
-	D08.811.913.696.620.682.700.145	Checkpoint Kinase 2
-	D08.811.913.696.620.682.700.150 Kinases	Cyclic Nucleotide-Regulated Protein
-	D08.811.913.696.620.682.700.150.125 Kinases	Cyclic AMP-Dependent Protein
-	D08.811.913.696.620.682.700.150.125.750 Kinase Type I	Cyclic AMP-Dependent Protein
-	D08.811.913.696.620.682.700.150.125.750.500 Protein Kinase Catalytic Subunits	Cyclic AMP-Dependent
-	D08.811.913.696.620.682.700.150.125.750.625 Protein Kinase RIalpha Subunit	Cyclic AMP-Dependent
-	D08.811.913.696.620.682.700.150.125.750.750 Protein Kinase RIIbeta Subunit	Cyclic AMP-Dependent
-	D08.811.913.696.620.682.700.150.125.875 Kinase Type II	Cyclic AMP-Dependent Protein
-	D08.811.913.696.620.682.700.150.125.875.500 Protein Kinase Catalytic Subunits	Cyclic AMP-Dependent
-	D08.811.913.696.620.682.700.150.125.875.750 Protein Kinase RIIalpha Subunit	Cyclic AMP-Dependent
-	D08.811.913.696.620.682.700.150.125.875.875 Protein Kinase RIIbeta Subunit	Cyclic AMP-Dependent
-	D08.811.913.696.620.682.700.150.150 Kinases	Cyclic GMP-Dependent Protein
-	D08.811.913.696.620.682.700.150.150.500 Kinase Type I	Cyclic GMP-Dependent Protein
-	D08.811.913.696.620.682.700.150.150.750 Kinase Type II	Cyclic GMP-Dependent Protein
-	D08.811.913.696.620.682.700.150.575	Protamine Kinase
-	D08.811.913.696.620.682.700.250	DNA-Activated Protein Kinase
-	D08.811.913.696.620.682.700.300	eIF-2 Kinase
-	D08.811.913.696.620.682.700.364	G-Protein-Coupled Receptor Kinases
-	D08.811.913.696.620.682.700.364.049	beta-Adrenergic Receptor Kinases
-	D08.811.913.696.620.682.700.364.049.200 Kinase 2	G-Protein-Coupled Receptor

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D08.811.913.696.620.682.700.364.049.300 Kinase 3 <span style="float: right;">G-Protein-Coupled Receptor</span>
-	D08.811.913.696.620.682.700.364.100 <span style="float: right;">G-Protein-Coupled Receptor Kinase 1</span>
-	D08.811.913.696.620.682.700.364.550 <span style="float: right;">G-Protein-Coupled Receptor Kinase 4</span>
-	D08.811.913.696.620.682.700.364.775 <span style="float: right;">G-Protein-Coupled Receptor Kinase 5</span>
-	D08.811.913.696.620.682.700.429 <span style="float: right;">Glycogen Synthase Kinases</span>
-	D08.811.913.696.620.682.700.429.500 <span style="float: right;">Glycogen Synthase Kinase 3</span>
New Heading	<b>D08.811.913.696.620.682.700.429.500.500</b> <span style="float: right;"><b>Glycogen Synthase Kinase 3</b></span> <b>beta</b>
-	D08.811.913.696.620.682.700.494 <span style="float: right;">I-kappa B Kinase</span>
-	D08.811.913.696.620.682.700.526 <span style="float: right;">Interleukin-1 Receptor-Associated Kinases</span>
New Heading	<b>D08.811.913.696.620.682.700.534</b> <span style="float: right;"><b>Leucine-Rich Repeat Serine-Threonine</b></span> <b>Protein Kinase-2</b>
-	D08.811.913.696.620.682.700.542 <span style="float: right;">Lim Kinases</span>
-	D08.811.913.696.620.682.700.559 <span style="float: right;">MAP Kinase Kinase Kinases</span>
-	D08.811.913.696.620.682.700.559.100 <span style="float: right;">MAP Kinase Kinase Kinase 1</span>
-	D08.811.913.696.620.682.700.559.200 <span style="float: right;">MAP Kinase Kinase Kinase 2</span>
-	D08.811.913.696.620.682.700.559.300 <span style="float: right;">MAP Kinase Kinase Kinase 3</span>
-	D08.811.913.696.620.682.700.559.400 <span style="float: right;">MAP Kinase Kinase Kinase 4</span>
-	D08.811.913.696.620.682.700.559.500 <span style="float: right;">MAP Kinase Kinase Kinase 5</span>
-	D08.811.913.696.620.682.700.559.800 <span style="float: right;">Proto-Oncogene Proteins c-mos</span>
-	D08.811.913.696.620.682.700.559.842 <span style="float: right;">raf Kinases</span>
-	D08.811.913.696.620.682.700.559.842.249 <span style="float: right;">Oncogene Proteins v-raf</span>
-	D08.811.913.696.620.682.700.559.842.374 <span style="float: right;">Proto-Oncogene Proteins B-raf</span>
-	D08.811.913.696.620.682.700.559.842.500 <span style="float: right;">Proto-Oncogene Proteins c-raf</span>
-	D08.811.913.696.620.682.700.565 <span style="float: right;">Mitogen-Activated Protein Kinase Kinases</span>
-	D08.811.913.696.620.682.700.565.100 <span style="float: right;">MAP Kinase Kinase 1</span>
-	D08.811.913.696.620.682.700.565.200 <span style="float: right;">MAP Kinase Kinase 2</span>
-	D08.811.913.696.620.682.700.565.300 <span style="float: right;">MAP Kinase Kinase 3</span>
-	D08.811.913.696.620.682.700.565.400 <span style="float: right;">MAP Kinase Kinase 4</span>
-	D08.811.913.696.620.682.700.565.500 <span style="float: right;">MAP Kinase Kinase 5</span>
-	D08.811.913.696.620.682.700.565.600 <span style="float: right;">MAP Kinase Kinase 6</span>
-	D08.811.913.696.620.682.700.565.700 <span style="float: right;">MAP Kinase Kinase 7</span>
-	D08.811.913.696.620.682.700.567 <span style="float: right;">Mitogen-Activated Protein Kinases</span>
-	D08.811.913.696.620.682.700.567.249 <span style="float: right;">Extracellular Signal-Regulated MAP</span> Kinases
-	D08.811.913.696.620.682.700.567.249.500 <span style="float: right;">Mitogen-Activated Protein Kinase</span>

## MeSH Tree Changes for 2017

Type	Tree - heading	
	1	
-	D08.811.913.696.620.682.700.567.249.750 3	Mitogen-Activated Protein Kinase
-	D08.811.913.696.620.682.700.567.249.875 6	Mitogen-Activated Protein Kinase
-	D08.811.913.696.620.682.700.567.249.937 7	Mitogen-Activated Protein Kinase
-	D08.811.913.696.620.682.700.567.374	JNK Mitogen-Activated Protein Kinases
-	D08.811.913.696.620.682.700.567.374.500 8	Mitogen-Activated Protein Kinase
-	D08.811.913.696.620.682.700.567.374.750 9	Mitogen-Activated Protein Kinase
-	D08.811.913.696.620.682.700.567.374.800 10	Mitogen-Activated Protein Kinase
-	D08.811.913.696.620.682.700.567.843	p38 Mitogen-Activated Protein Kinases
-	D08.811.913.696.620.682.700.567.843.500 11	Mitogen-Activated Protein Kinase
-	D08.811.913.696.620.682.700.567.843.750 12	Mitogen-Activated Protein Kinase
-	D08.811.913.696.620.682.700.567.843.875 13	Mitogen-Activated Protein Kinase
-	D08.811.913.696.620.682.700.567.843.937 14	Mitogen-Activated Protein Kinase
-	D08.811.913.696.620.682.700.576	Myotonin-Protein Kinase
New Heading	<b>D08.811.913.696.620.682.700.581</b>	<b>NIMA-Related Kinases</b>
New Heading	<b>D08.811.913.696.620.682.700.581.500</b>	<b>NIMA-Related Kinase 1</b>
-	D08.811.913.696.620.682.700.586	Oncogene Protein v-akt
-	D08.811.913.696.620.682.700.596	p21-Activated Kinases
-	D08.811.913.696.620.682.700.606	Phytochrome A
-	D08.811.913.696.620.682.700.626	Positive Transcriptional Elongation Factor B
-	D08.811.913.696.620.682.700.646	Proline-Directed Protein Kinases
-	D08.811.913.696.620.682.700.646.500	Cyclin-Dependent Kinases
-	D08.811.913.696.620.682.700.646.500.500	CDC2-CDC28 Kinases
-	D08.811.913.696.620.682.700.646.500.500.250	CDC2 Protein Kinase
-	D08.811.913.696.620.682.700.646.500.500.375 cerevisiae	CDC28 Protein Kinase, S
-	D08.811.913.696.620.682.700.646.500.500.500	Cyclin-Dependent Kinase 5
-	D08.811.913.696.620.682.700.646.500.500.750	Cyclin-Dependent Kinase 9

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D08.811.913.696.620.682.700.646.500.750	Cyclin-Dependent Kinase 2
-	D08.811.913.696.620.682.700.646.500.812	Cyclin-Dependent Kinase 3
-	D08.811.913.696.620.682.700.646.500.875	Cyclin-Dependent Kinase 4
-	D08.811.913.696.620.682.700.646.500.937	Cyclin-Dependent Kinase 6
-	D08.811.913.696.620.682.700.646.500.968	Cyclin-Dependent Kinase 8
-	D08.811.913.696.620.682.700.646.500.984	Maturation-Promoting Factor
-	D08.811.913.696.620.682.700.646.500.984.500	CDC2 Protein Kinase
-	D08.811.913.696.620.682.700.646.625	Glycogen Synthase Kinase 3
New Heading	<b>D08.811.913.696.620.682.700.646.625.500 beta</b>	<b>Glycogen Synthase Kinase 3</b>
-	D08.811.913.696.620.682.700.725	Protein Kinase C
-	D08.811.913.696.620.682.700.725.049	Protein Kinase C beta
-	D08.811.913.696.620.682.700.725.100	Protein Kinase C-alpha
-	D08.811.913.696.620.682.700.725.400	Protein Kinase C-delta
-	D08.811.913.696.620.682.700.725.750	Protein Kinase C-epsilon
-	D08.811.913.696.620.682.700.745 Kinases	3-Phosphoinositide-Dependent Protein
-	D08.811.913.696.620.682.700.755	Proto-Oncogene Proteins c-akt
-	D08.811.913.696.620.682.700.759	Proto-Oncogene Proteins c-bcr
-	D08.811.913.696.620.682.700.776	Proto-Oncogene Proteins c-pim-1
-	D08.811.913.696.620.682.700.801 Threonine Kinases	Receptor-Interacting Protein Serine-
-	D08.811.913.696.620.682.700.801.500 Threonine Kinase 2	Receptor-Interacting Protein Serine-
-	D08.811.913.696.620.682.700.814	rho-Associated Kinases
-	D08.811.913.696.620.682.700.862	Ribosomal Protein S6 Kinases
-	D08.811.913.696.620.682.700.862.249	Ribosomal Protein S6 Kinases, 70-kDa
-	D08.811.913.696.620.682.700.862.500	Ribosomal Protein S6 Kinases, 90-kDa
-	D08.811.913.696.620.682.700.931	TOR Serine-Threonine Kinases
-	D08.811.913.696.620.682.700.965	Vacuolar Sorting Protein VPS15
-	D08.811.913.696.620.682.725	Protein-Tyrosine Kinases
-	D08.811.913.696.620.682.725.049	Focal Adhesion Protein-Tyrosine Kinases
-	D08.811.913.696.620.682.725.049.500	Focal Adhesion Kinase 1
-	D08.811.913.696.620.682.725.049.750	Focal Adhesion Kinase 2
-	D08.811.913.696.620.682.725.124	Janus Kinases
-	D08.811.913.696.620.682.725.124.100	Janus Kinase 1
-	D08.811.913.696.620.682.725.124.200	Janus Kinase 2

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D08.811.913.696.620.682.725.124.300	Janus Kinase 3
-	D08.811.913.696.620.682.725.124.650	TYK2 Kinase
-	D08.811.913.696.620.682.725.200	Mitogen-Activated Protein Kinase Kinases
-	D08.811.913.696.620.682.725.200.100	MAP Kinase Kinase 1
-	D08.811.913.696.620.682.725.200.200	MAP Kinase Kinase 2
-	D08.811.913.696.620.682.725.200.300	MAP Kinase Kinase 3
-	D08.811.913.696.620.682.725.200.400	MAP Kinase Kinase 4
-	D08.811.913.696.620.682.725.200.500	MAP Kinase Kinase 5
-	D08.811.913.696.620.682.725.200.600	MAP Kinase Kinase 6
-	D08.811.913.696.620.682.725.200.700	MAP Kinase Kinase 7
-	D08.811.913.696.620.682.725.300	Proto-Oncogene Proteins c-fes
-	D08.811.913.696.620.682.725.400	Receptor Protein-Tyrosine Kinases
New Heading	<b>D08.811.913.696.620.682.725.400.005</b>	<b>Discoidin Domain Receptors</b>
New Heading	<b>D08.811.913.696.620.682.725.400.005.500</b>	<b>Discoidin Domain Receptor 1</b>
New Heading	<b>D08.811.913.696.620.682.725.400.005.750</b>	<b>Discoidin Domain Receptor 2</b>
-	D08.811.913.696.620.682.725.400.009	ErbB Receptors
-	D08.811.913.696.620.682.725.400.009.300 Factor	Receptor, Epidermal Growth
-	D08.811.913.696.620.682.725.400.009.400	Receptor, ErbB-2
-	D08.811.913.696.620.682.725.400.009.500	Receptor, ErbB-3
-	D08.811.913.696.620.682.725.400.009.600	Receptor, ErbB-4
-	D08.811.913.696.620.682.725.400.020	fms-Like Tyrosine Kinase 3
-	D08.811.913.696.620.682.725.400.050	Proto-Oncogene Proteins c-kit
-	D08.811.913.696.620.682.725.400.075	Proto-Oncogene Proteins c-met
-	D08.811.913.696.620.682.725.400.087	Proto-Oncogene Proteins c-ret
-	D08.811.913.696.620.682.725.400.093 Receptors	Receptor Tyrosine Kinase-like Orphan
-	D08.811.913.696.620.682.725.400.177 Type 1	Receptor, Fibroblast Growth Factor,
-	D08.811.913.696.620.682.725.400.178 Type 2	Receptor, Fibroblast Growth Factor,
-	D08.811.913.696.620.682.725.400.179 Type 3	Receptor, Fibroblast Growth Factor,
-	D08.811.913.696.620.682.725.400.180 Type 4	Receptor, Fibroblast Growth Factor,
-	D08.811.913.696.620.682.725.400.185	Receptor, IGF Type 1

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D08.811.913.696.620.682.725.400.200 Receptor, Insulin
-	D08.811.913.696.620.682.725.400.500 Receptor, Macrophage Colony-Stimulating Factor
-	D08.811.913.696.620.682.725.400.660 Receptor, trkA
-	D08.811.913.696.620.682.725.400.700 Receptor, trkB
-	D08.811.913.696.620.682.725.400.800 Receptor, trkC
-	D08.811.913.696.620.682.725.400.850 Receptors, Eph Family
-	D08.811.913.696.620.682.725.400.850.050 Receptor, EphA1
-	D08.811.913.696.620.682.725.400.850.100 Receptor, EphA2
-	D08.811.913.696.620.682.725.400.850.150 Receptor, EphA3
-	D08.811.913.696.620.682.725.400.850.200 Receptor, EphA4
-	D08.811.913.696.620.682.725.400.850.250 Receptor, EphA5
-	D08.811.913.696.620.682.725.400.850.300 Receptor, EphA6
-	D08.811.913.696.620.682.725.400.850.400 Receptor, EphA7
-	D08.811.913.696.620.682.725.400.850.500 Receptor, EphA8
-	D08.811.913.696.620.682.725.400.850.600 Receptor, EphB1
-	D08.811.913.696.620.682.725.400.850.650 Receptor, EphB2
-	D08.811.913.696.620.682.725.400.850.700 Receptor, EphB3
-	D08.811.913.696.620.682.725.400.850.750 Receptor, EphB4
-	D08.811.913.696.620.682.725.400.850.800 Receptor, EphB5
-	D08.811.913.696.620.682.725.400.900 Receptors, Platelet-Derived Growth Factor
-	D08.811.913.696.620.682.725.400.900.500 Receptor, Platelet-Derived Growth Factor alpha
-	D08.811.913.696.620.682.725.400.900.750 Receptor, Platelet-Derived Growth Factor beta
-	D08.811.913.696.620.682.725.400.925 Receptors, TIE
-	D08.811.913.696.620.682.725.400.925.249 Receptor, TIE-1
-	D08.811.913.696.620.682.725.400.925.500 Receptor, TIE-2
-	D08.811.913.696.620.682.725.400.950 Receptors, Vascular Endothelial Growth Factor
-	D08.811.913.696.620.682.725.400.950.100 Vascular Endothelial Growth Factor Receptor-1
-	D08.811.913.696.620.682.725.400.950.200 Vascular Endothelial Growth Factor Receptor-2
-	D08.811.913.696.620.682.725.400.950.300 Vascular Endothelial Growth Factor Receptor-3
-	D08.811.913.696.620.682.725.500 Proto-Oncogene Proteins c-abl
-	D08.811.913.696.620.682.725.500.500 Fusion Proteins, bcr-abl



## MeSH Tree Changes for 2017

Type	Tree - heading	
New Heading	<b>D08.811.913.696.620.682.725.650</b>	<b>Syk Kinase</b>
-	D08.811.913.696.620.682.725.800	src-Family Kinases
-	D08.811.913.696.620.682.725.800.315 Kinase p56(lck)	Lymphocyte Specific Protein Tyrosine
-	D08.811.913.696.620.682.725.800.472	Oncogene Protein pp60(v-src)
-	D08.811.913.696.620.682.725.800.551	Proto-Oncogene Proteins c-fyn
-	D08.811.913.696.620.682.725.800.590	Proto-Oncogene Proteins c-hck
-	D08.811.913.696.620.682.725.800.610	Proto-Oncogene Proteins c-yes
-	D08.811.913.696.620.682.725.800.630	Proto-Oncogene Proteins pp60(c-src)
-	D08.811.913.696.620.682.725.900	ZAP-70 Protein-Tyrosine Kinase
-	D08.811.913.696.620.685	Pyridoxal Kinase
-	D08.811.913.696.620.695	Pyruvate Kinase
-	D08.811.913.696.620.750	Thymidine Kinase
-	D08.811.913.696.620.800	Uridine Kinase
-	D08.811.913.696.630	Phosphotransferases (Carboxyl Group Acceptor)
-	D08.811.913.696.630.025	Acetate Kinase
-	D08.811.913.696.630.050	Aspartate Kinase
-	D08.811.913.696.630.050.050	Aspartokinase Homoserine Dehydrogenase
-	D08.811.913.696.630.700	Phosphoglycerate Kinase
-	D08.811.913.696.640	Phosphotransferases (Nitrogenous Group Acceptor)
-	D08.811.913.696.640.025	Arginine Kinase
-	D08.811.913.696.640.150	Creatine Kinase
-	D08.811.913.696.640.150.500	Creatine Kinase, BB Form
-	D08.811.913.696.640.150.625	Creatine Kinase, MB Form
-	D08.811.913.696.640.150.750	Creatine Kinase, Mitochondrial Form
-	D08.811.913.696.640.150.875	Creatine Kinase, MM Form
-	D08.811.913.696.645	Phosphotransferases (Paired Acceptors)
-	D08.811.913.696.645.700	Pyruvate, Orthophosphate Dikinase
-	D08.811.913.696.650	Phosphotransferases (Phosphate Group Acceptor)
-	D08.811.913.696.650.025	Adenylate Kinase
-	D08.811.913.696.650.150	ATP Synthetase Complexes
-	D08.811.913.696.650.150.500	Proton-Translocating ATPases
-	D08.811.913.696.650.150.500.249	Bacterial Proton-Translocating ATPases
-	D08.811.913.696.650.150.500.500	Chloroplast Proton-Translocating ATPases
-	D08.811.913.696.650.150.500.750 ATPases	Mitochondrial Proton-Translocating

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D08.811.913.696.650.150.500.875 Vacuolar Proton-Translocating ATPases
-	D08.811.913.696.650.450 Guanylate Kinase
-	D08.811.913.696.650.450 Guanylate Kinases
-	D08.811.913.696.650.550 Nucleoside-Diphosphate Kinase
-	D08.811.913.696.650.550.200 NM23 Nucleoside Diphosphate Kinases
-	D08.811.913.696.650.550.200.500 Nucleoside Diphosphate Kinase D
-	D08.811.913.696.650.575 Nucleoside-Phosphate Kinase
-	D08.811.913.696.900 Transferases (Other Substituted Phosphate Groups)
-	D08.811.913.696.900.074 CDP-Diacylglycerol-Inositol 3-Phosphatidyltransferase
-	D08.811.913.696.900.150 CDPdiacylglycerol-Serine O-Phosphatidyltransferase
-	D08.811.913.696.900.200 Diacylglycerol Cholinephosphotransferase
-	D08.811.913.696.900.250 Ethanolaminephosphotransferase
-	D08.811.913.817 Sulfur Group Transferases
-	D08.811.913.817.200 Coenzyme A-Transferases
-	D08.811.913.817.400 Sulfotransferases
-	D08.811.913.817.400.300 Arylsulfotransferase
-	D08.811.913.817.500 Sulfurtransferases
-	D08.811.913.817.500.500 Thiosulfate Sulfurtransferase
-	D09 Carbohydrates
-	D09.067 Amino Sugars
-	D09.067.342 Hexosamines
-	D09.067.342.300 Fructosamine
-	D09.067.342.356 Galactosamine
-	D09.067.342.356.050 Acetylgalactosamine
-	D09.067.342.531 Glucosamine
-	D09.067.342.531.050 Acetylglucosamine
-	D09.067.342.600 Meglumine
-	D09.067.342.600.500 Diatrizoate Meglumine
-	D09.067.342.600.600 Iothalamate Meglumine
-	D09.067.550 Muramic Acids
-	D09.067.550.050 Acetylmuramyl-Alanyl-Isoglutamine
-	D09.067.687 Neuraminic Acids
-	D09.067.687.668 Sialic Acids
-	D09.067.687.668.030 N-Acetylneuraminic Acid

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D09.067.687.668.250 Cytidine Monophosphate N-Acetylneuraminic Acid
-	D09.067.687.668.775 Zanamivir
-	D09.254 Deoxy Sugars
-	D09.254.229 Deoxyglucose
-	D09.254.229.500 Fluorodeoxyglucose F18
-	D09.254.330 Deoxyribose
-	D09.254.488 Fucose
-	D09.254.799 Rhamnose
-	D09.301 Dietary Carbohydrates
-	D09.301.325 Dietary Sucrose
-	D09.301.662 High Fructose Corn Syrup
-	D09.400 Glycoconjugates
-	D09.400.410 Glycolipids
-	D09.400.410.209 Galactolipids
-	D09.400.410.420 Glycosphingolipids
-	D09.400.410.420.025 Acidic Glycosphingolipids
-	D09.400.410.420.025.475 Gangliosides
-	D09.400.410.420.025.475.390 G(M1) Ganglioside
-	D09.400.410.420.025.475.400 G(M2) Ganglioside
-	D09.400.410.420.025.475.510 G(M3) Ganglioside
-	D09.400.410.420.025.837 Sulfoglycosphingolipids
-	D09.400.410.420.525 Neutral Glycosphingolipids
-	D09.400.410.420.525.200 Ceramides
-	D09.400.410.420.525.200.250 Cerebrosides
-	D09.400.410.420.525.200.250.450 Galactosylceramides
-	D09.400.410.420.525.200.250.490 Glucosylceramides
-	D09.400.410.420.525.200.425 Globosides
-	D09.400.410.420.525.200.612 Lactosylceramides
-	D09.400.410.420.525.200.906 Trihexosylceramides
-	D09.400.410.420.525.870 Sphingomyelins
-	D09.400.410.420.806 Psychosine
-	D09.400.410.475 Glycosylphosphatidylinositols
-	D09.400.420 Glycopeptides
-	D09.400.420.110 Bleomycin
-	D09.400.420.110.690 Peplomycin
-	D09.400.420.110.710 Phleomycins

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D09.400.420.700	Peptidoglycan
-	D09.400.420.800	Ristocetin
-	D09.400.420.900	Teicoplanin
-	D09.400.420.925	Vancomycin
-	D09.400.430	Glycoproteins
New Heading	<b>D09.400.430.250</b>	<b>AC133 Antigen</b>
-	D09.400.430.500	ADAM Proteins
New Heading	<b>D09.400.430.500.125</b>	<b>ADAM12 Protein</b>
New Heading	<b>D09.400.430.500.250</b>	<b>ADAM10 Protein</b>
New Heading	<b>D09.400.430.500.375</b>	<b>ADAM17 Protein</b>
New Heading	<b>D09.400.430.500.500</b>	<b>ADAMTS Proteins</b>
New Heading	<b>D09.400.430.500.500.500</b>	<b>ADAMTS1 Protein</b>
New Heading	<b>D09.400.430.500.500.813</b>	<b>ADAMTS13 Protein</b>
New Heading	<b>D09.400.430.500.500.844</b>	<b>ADAMTS4 Protein</b>
New Heading	<b>D09.400.430.500.500.875</b>	<b>ADAMTS5 Protein</b>
New Heading	<b>D09.400.430.500.500.937</b>	<b>ADAMTS7 Protein</b>
New Heading	<b>D09.400.430.500.500.968</b>	<b>ADAMTS9 Protein</b>
New Heading	<b>D09.400.430.500.750</b>	<b>Fertilins</b>
-	D09.400.430.750	Cholesterol Ester Transfer Proteins
New Heading	<b>D09.400.430.875</b>	<b>Fibrillins</b>
New Heading	<b>D09.400.430.875.500</b>	<b>Fibrillin-1</b>
New Heading	<b>D09.400.430.875.750</b>	<b>Fibrillin-2</b>
-	D09.400.500	Lipopolysaccharides
-	D09.408	Glycosides
-	D09.408.051	Aminoglycosides
-	D09.408.051.059	Anthracyclines

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D09.408.051.059.050 Aclarubicin
-	D09.408.051.059.200 Daunorubicin
-	D09.408.051.059.200.150 Carubicin
-	D09.408.051.059.200.175 Doxorubicin
-	D09.408.051.059.200.175.200 Epirubicin
-	D09.408.051.059.200.300 Idarubicin
-	D09.408.051.059.200.650 Nogalamycin
-	D09.408.051.059.200.650.500 Menogaril
-	D09.408.051.059.650 Plicamycin
-	D09.408.051.120 Butirosin Sulfate
-	D09.408.051.374 Gentamicins
-	D09.408.051.374.785 Sisomicin
-	D09.408.051.374.785.525 Netilmicin
-	D09.408.051.420 Hygromycin B
-	D09.408.051.476 Kanamycin
-	D09.408.051.476.060 Amikacin
-	D09.408.051.476.200 Dibekacin
-	D09.408.051.476.600 Nebramycin
-	D09.408.051.476.600.800 Tobramycin
-	D09.408.051.545 Metrizamide
-	D09.408.051.623 Neomycin
-	D09.408.051.623.300 Framycetin
-	D09.408.051.623.635 Paromomycin
-	D09.408.051.623.800 Ribostamycin
-	D09.408.051.788 Puromycin
-	D09.408.051.788.650 Puromycin Aminonucleoside
-	D09.408.051.836 Spectinomycin
-	D09.408.051.885 Streptomycin
-	D09.408.051.885.387 Dihydrostreptomycin Sulfate
-	D09.408.051.892 Streptothricins
-	D09.408.051.900 Streptozocin
-	D09.408.084 Anthocyanins
-	D09.408.105 Atractyloside
-	D09.408.180 Cardiac Glycosides
-	D09.408.180.261 Digitalis Glycosides
-	D09.408.180.261.236 Digitonin

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D09.408.180.261.336 Digitoxin
-	D09.408.180.261.336.259 Acetyldigitoxins
-	D09.408.180.261.436 Digoxin
-	D09.408.180.261.436.050 Acetyldigoxins
-	D09.408.180.261.436.500 Medigoxin
-	D09.408.180.261.657 Lanatosides
-	D09.408.180.261.657.200 Deslanoside
-	D09.408.180.660 Proscillaridin
-	D09.408.180.810 Strophanthins
-	D09.408.180.810.250 Cymarine
-	D09.408.180.810.600 Ouabain
-	D09.408.210 Chromomycins
-	D09.408.210.209 Chromomycin A3
-	D09.408.320 Galactosides
-	D09.408.320.500 Methylgalactosides
-	D09.408.320.550 Nitrophenylgalactosides
-	D09.408.320.820 Thiogalactosides
-	D09.408.320.820.500 Isopropyl Thiogalactoside
-	D09.408.348 Glucosides
-	D09.408.348.050 Amygdalin
-	D09.408.348.075 Arbutin
-	D09.408.348.113 Canagliflozin
-	D09.408.348.150 Chloralose
-	D09.408.348.250 Esculin
-	D09.408.348.275 Etoposide
-	D09.408.348.387 Iridoid Glucosides
-	D09.408.348.500 Methylglucosides
-	D09.408.348.500.500 3-O-Methylglucose
-	D09.408.348.800 Teniposide
-	D09.408.348.820 Thioglucosides
-	D09.408.348.820.350 Glucosinolates
-	D09.408.375 Hemoglobin A, Glycosylated
-	D09.408.423 Iridoid Glycosides
-	D09.408.423.500 Iridoid Glucosides
-	D09.408.471 Lincosamides
-	D09.408.471.500 Lincomycin

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D09.408.471.500.125 Clindamycin
-	D09.408.490 Mannosides
-	D09.408.490.500 Methylmannosides
-	D09.408.514 Methylglycosides
-	D09.408.514.455 Methylgalactosides
-	D09.408.514.622 Methylglucosides
-	D09.408.514.622.500 3-O-Methylglucose
-	D09.408.514.700 Methylmannosides
-	D09.408.554 Novobiocin
-	D09.408.595 Nucleosides
-	D09.408.620 Nucleotides
-	D09.408.620.569 Nucleoside Diphosphate Sugars
-	D09.408.620.569.070 Adenosine Diphosphate Sugars
-	D09.408.620.569.070.075 Adenosine Diphosphate Glucose
-	D09.408.620.569.070.125 Adenosine Diphosphate Ribose
-	D09.408.620.569.070.125.040 O-Acetyl-ADP-Ribose
-	D09.408.620.569.070.125.195 Cyclic ADP-Ribose
-	D09.408.620.569.070.125.600 Poly Adenosine Diphosphate Ribose
-	D09.408.620.569.200 Cytidine Diphosphate Diglycerides
-	D09.408.620.569.400 Guanosine Diphosphate Sugars
-	D09.408.620.569.400.410 Guanosine Diphosphate Fucose
-	D09.408.620.569.400.500 Guanosine Diphosphate Mannose
-	D09.408.620.569.727 Uridine Diphosphate Sugars
-	D09.408.620.569.727.100 Uridine Diphosphate N-Acetylgalactosamine
-	D09.408.620.569.727.120 Uridine Diphosphate N-Acetylglucosamine
-	D09.408.620.569.727.150 Uridine Diphosphate N-Acetylmuramic Acid
-	D09.408.620.569.727.300 Uridine Diphosphate Galactose
-	D09.408.620.569.727.350 Uridine Diphosphate Glucose
-	D09.408.620.569.727.375 Uridine Diphosphate Glucuronic Acid
-	D09.408.620.569.727.800 Uridine Diphosphate Xylose
-	D09.408.661 Olivomycins
-	D09.408.702 Phlorhizin
-	D09.408.782 Saponins
-	D09.408.782.250 Escin
-	D09.408.782.300 Ginsenosides
-	D09.408.782.350 Holothurin

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D09.408.782.512 Quillaja Saponins
-	D09.408.782.674 Solanine
-	D09.408.872 Teichoic Acids
-	D09.408.903 Thioglycosides
-	D09.408.903.600 Thiogalactosides
-	D09.408.903.600.500 Isopropyl Thiogalactoside
-	D09.408.903.703 Thioglucosides
-	D09.408.903.703.350 Glucosinolates
-	D09.408.934 Tomatine
-	D09.546 Monosaccharides
-	D09.546.142 Carbasugars
-	D09.546.286 Heptoses
-	D09.546.286.500 Mannoheptulose
-	D09.546.359 Hexoses
-	D09.546.359.250 Fructose
-	D09.546.359.313 Fucose
-	D09.546.359.377 Galactose
-	D09.546.359.448 Glucose
-	D09.546.359.448.500 Blood Glucose
-	D09.546.359.588 Mannose
-	D09.546.359.774 Rhamnose
-	D09.546.359.862 Sorbose
-	D09.546.412 Imino Sugars
-	D09.546.412.249 Imino Furanoses
-	D09.546.412.500 Imino Pyranoses
-	D09.546.412.500.033 1-Deoxynojirimycin
-	D09.546.465 Ketoses
-	D09.546.465.200 Dihydroxyacetone
-	D09.546.465.354 Fructose
-	D09.546.465.500 Mannoheptulose
-	D09.546.465.758 Sorbose
-	D09.546.465.920 Xylulose
-	D09.546.627 Pentoses
-	D09.546.627.166 Arabinose
-	D09.546.627.651 Ribose
-	D09.546.627.867 Xylose



## MeSH Tree Changes for 2017

Type	Tree - heading
-	D09.546.627.885 Xylulose
-	D09.546.805 Tetroses
-	D09.546.849 Thiosugars
-	D09.546.894 Trioses
-	D09.546.894.200 Dihydroxyacetone
-	D09.546.894.449 Glyceraldehyde
-	D09.546.894.449.450 Glyceraldehyde 3-Phosphate
-	D09.698 Polysaccharides
-	D09.698.068 Alginates
-	D09.698.152 Carrageenan
-	D09.698.211 Chitin
-	D09.698.211.500 Chitosan
-	D09.698.330 Ficoll
-	D09.698.350 Fructans
-	D09.698.350.500 Inulin
-	D09.698.357 Fungal Polysaccharides
-	D09.698.360 Galactans
-	D09.698.360.041 Agar
-	D09.698.365 Glucans
-	D09.698.365.089 beta-Glucans
-	D09.698.365.089.500 Lentinan
-	D09.698.365.089.625 Sizofiran
-	D09.698.365.089.750 Zymosan
-	D09.698.365.180 Cellulose
-	D09.698.365.180.180 Cellobiose
-	D09.698.365.180.200 Cellulose, Oxidized
-	D09.698.365.180.247 DEAE-Cellulose
-	D09.698.365.180.455 Hypromellose Derivatives
-	D09.698.365.180.663 Methylcellulose
-	D09.698.365.180.663.329 Carboxymethylcellulose Sodium
-	D09.698.365.272 Dextrans
-	D09.698.365.272.250 DEAE-Dextran
-	D09.698.365.272.300 Dextran Sulfate
-	D09.698.365.272.400 Iron-Dextran Complex
-	D09.698.365.388 Glycogen
-	D09.698.365.388.518 Liver Glycogen

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D09.698.365.410 Isomaltose
-	D09.698.365.450 Maltose
-	D09.698.365.855 Starch
-	D09.698.365.855.194 Amylopectin
-	D09.698.365.855.361 Amylose
-	D09.698.365.855.400 Dextrins
-	D09.698.365.855.400.375 Cyclodextrins
-	D09.698.365.855.400.375.222 alpha-Cyclodextrins
-	D09.698.365.855.400.375.333 beta-Cyclodextrins
-	D09.698.365.855.400.375.444 gamma-Cyclodextrins
-	D09.698.365.855.500 Hydroxyethyl Starch Derivatives
-	D09.698.365.855.750 Inulin
-	D09.698.365.900 Trehalose
-	D09.698.373 Glycosaminoglycans
-	D09.698.373.200 Chondroitin
-	D09.698.373.200.300 Chondroitin Sulfates
-	D09.698.373.200.380 Dermatan Sulfate
-	D09.698.373.400 Heparin
-	D09.698.373.400.300 Heparin, Low-Molecular-Weight
-	D09.698.373.400.300.150 Dalteparin
-	D09.698.373.400.300.200 Enoxaparin
-	D09.698.373.400.300.600 Nadroparin
-	D09.698.373.400.320 Heparinoids
-	D09.698.373.425 Heparitin Sulfate
-	D09.698.373.425.500 Heparan Sulfate Proteoglycans
-	D09.698.373.475 Hyaluronic Acid
-	D09.698.373.550 Keratan Sulfate
-	D09.698.550 Mannans
-	D09.698.629 Oligosaccharides
-	D09.698.629.305 Disaccharides
-	D09.698.629.305.200 Cellobiose
-	D09.698.629.305.320 Isomaltose
-	D09.698.629.305.340 Lactose
-	D09.698.629.305.423 Lactulose
-	D09.698.629.305.523 Maltose
-	D09.698.629.305.540 Melibiose

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D09.698.629.305.770 Sucrose
-	D09.698.629.305.770.200 Dietary Sucrose
-	D09.698.629.305.770.850 Sucralfate
-	D09.698.629.305.880 Trehalose
-	D09.698.629.553 Oligosaccharides, Branched-Chain
-	D09.698.629.802 Trisaccharides
-	D09.698.629.802.100 Acarbose
-	D09.698.629.802.700 Raffinose
-	D09.698.670 Pectins
-	D09.698.682 Pentosan Sulfuric Polyester
-	D09.698.700 Plant Gums
-	D09.698.700.249 Chewing Gum
-	D09.698.700.500 Gum Arabic
-	D09.698.700.625 Karaya Gum
-	D09.698.700.750 Tragacanth
-	D09.698.709 Plant Mucilage
-	D09.698.718 Polysaccharides, Bacterial
-	D09.698.718.220 Bambermycins
-	D09.698.718.450 Lipopolysaccharides
-	D09.698.718.450.500 Lipid A
-	D09.698.718.450.600 O Antigens
-	D09.698.718.594 Peptidoglycan
-	D09.698.718.622 Prebiotics
-	D09.698.718.650 Prodigiozan
-	D09.698.718.825 Teichoic Acids
-	D09.698.735 Proteoglycans
-	D09.698.735.200 Chondroitin Sulfate Proteoglycans
-	D09.698.735.200.500 Aggrecans
New Tree	<a href="#">D09.698.735.200.625</a> <a href="#">Antigens, CD44</a>
-	D09.698.735.200.750 Versicans
-	D09.698.735.400 Heparan Sulfate Proteoglycans
New Heading	<b>D09.698.735.700</b> <b>Small Leucine-Rich Proteoglycans</b>
New Tree	<a href="#">D09.698.735.700.500</a> <a href="#">Biglycan</a>
New	<a href="#">D09.698.735.700.750</a> <a href="#">Decorin</a>

## MeSH Tree Changes for 2017

Type	Tree - heading	
Tree		
New Heading	<b>D09.698.735.700.813</b>	<b>Fibromodulin</b>
New Heading	<b>D09.698.735.700.875</b>	<b>Lumican</b>
-	D09.698.813	Sepharose
-	D09.698.925	Xylans
-	D09.811	Sugar Acids
-	D09.811.100	Ascorbic Acid
-	D09.811.100.260	Dehydroascorbic Acid
-	D09.811.200	2,3-Diketogulonic Acid
-	D09.811.295	Glucaric Acid
-	D09.811.308	Gluconates
-	D09.811.308.060	Antimony Sodium Gluconate
-	D09.811.308.200	Calcium Gluconate
-	D09.811.366	Glyceric Acids
-	D09.811.366.388	Diphosphoglyceric Acids
-	D09.811.366.388.175	2,3-Diphosphoglycerate
-	D09.811.522	Muramic Acids
-	D09.811.522.050	Acetylmuramyl-Alanyl-Isoglutamine
-	D09.811.589	Neuraminic Acids
-	D09.811.589.668	Sialic Acids
-	D09.811.589.668.030	N-Acetylneuraminic Acid
-	D09.811.589.668.250	Cytidine Monophosphate N-Acetylneuraminic Acid
-	D09.811.589.668.775	Zanamivir
-	D09.811.779	Tartrates
-	D09.811.835	Tartronates
-	D09.811.922	Uronic Acids
-	D09.811.922.162	Glucuronates
-	D09.811.922.162.500	Glucuronic Acid
-	D09.811.922.162.750	Glucuronides
-	D09.811.922.400	Hexuronic Acids
-	D09.811.922.400.500	Iduronic Acid
-	D09.853	Sugar Alcohols
-	D09.853.175	Dithioerythritol
-	D09.853.196	Dithiothreitol

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D09.853.329 Erythritol
-	D09.853.329.225 Erythrityl Tetranitrate
-	D09.853.400 Galactitol
-	D09.853.400.350 Dianhydrogalactitol
-	D09.853.400.500 Mitolactol
-	D09.853.421 Glycerol
-	D09.853.519 Inositol
-	D09.853.519.400 Inositol Phosphates
-	D09.853.519.400.350 Inositol 1,4,5-Trisphosphate
-	D09.853.519.400.700 Phytic Acid
-	D09.853.609 Mannitol
-	D09.853.609.450 Mannitol Phosphates
-	D09.853.609.500 Mitobronitol
-	D09.853.780 Ribitol
-	D09.853.813 Sorbitol
-	D09.853.813.480 Isosorbide
-	D09.853.813.480.500 Isosorbide Dinitrate
-	D09.853.813.550 Meglumine
-	D09.853.813.550.500 Diatrizoate Meglumine
-	D09.853.813.550.600 Iothalamate Meglumine
-	D09.853.936 Xylitol
-	D09.894 Sugar Phosphates
-	D09.894.150 Dihydroxyacetone Phosphate
-	D09.894.299 Glycerophosphates
-	D09.894.299.449 Glycerolphosphorylcholine
-	D09.894.417 Hexosephosphates
-	D09.894.417.313 Fructosephosphates
-	D09.894.417.313.300 Fructosediphosphates
-	D09.894.417.370 Galactosephosphates
-	D09.894.417.448 Glucosephosphates
-	D09.894.417.448.500 Glucose-6-Phosphate
-	D09.894.417.592 Hexosediphosphates
-	D09.894.417.592.300 Fructosediphosphates
-	D09.894.417.650 Mannosephosphates
-	D09.894.480 Inositol Phosphates
-	D09.894.480.350 Inositol 1,4,5-Trisphosphate

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D09.894.480.700 Phytic Acid
-	D09.894.500 Mannitol Phosphates
-	D09.894.643 Pentosephosphates
-	D09.894.643.650 Phosphoribosyl Pyrophosphate
-	D09.894.643.700 Ribosemonophosphates
-	D09.894.643.720 Ribulosephosphates
-	D09.894.680 Polyisoprenyl Phosphate Sugars
-	D09.894.680.700 Polyisoprenyl Phosphate Monosaccharides
-	D09.894.680.700.250 Dolichol Monophosphate Mannose
-	D09.894.680.710 Polyisoprenyl Phosphate Oligosaccharides
-	D09.894.847 Teichoic Acids
-	D10 Lipids
-	D10.150 Ceroid
-	D10.212 Fats
-	D10.212.302 Dietary Fats
-	D10.212.302.199 Butter
-	D10.212.302.199.500 Ghee
-	D10.212.302.347 Cholesterol, Dietary
-	D10.212.302.380 Dietary Fats, Unsaturated
-	D10.212.302.380.360 Cod Liver Oil
-	D10.212.302.380.370 Corn Oil
-	D10.212.302.380.380 Cottonseed Oil
-	D10.212.302.380.410 Fatty Acids, Omega-3
-	D10.212.302.380.410.100 alpha-Linolenic Acid
-	D10.212.302.380.410.210 Docosahexaenoic Acids
-	D10.212.302.380.410.385 Eicosapentaenoic Acid
-	D10.212.302.380.580 Olive Oil
-	D10.212.302.380.750 Safflower Oil
-	D10.212.302.380.775 Sesame Oil
-	D10.212.302.380.800 Soybean Oil
-	D10.212.302.651 Margarine
-	D10.212.507 Fats, Unsaturated
-	D10.212.507.300 Castor Oil
-	D10.212.507.300.500 Ricinoleic Acids
-	D10.212.507.325 Cod Liver Oil
-	D10.212.507.340 Corn Oil

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D10.212.507.350	Cottonseed Oil
-	D10.212.507.375	Croton Oil
-	D10.212.507.550	Linseed Oil
-	D10.212.507.650	Olive Oil
-	D10.212.507.750	Safflower Oil
-	D10.212.507.775	Sesame Oil
-	D10.212.507.800	Soybean Oil
-	D10.212.507.850	Triolein
-	D10.251	Fatty Acids
-	D10.251.122	Caprylates
-	D10.251.175	Decanoic Acids
-	D10.251.175.200	Decanoates
-	D10.251.220	Eicosanoic Acids
-	D10.251.265	Endocannabinoids
-	D10.251.310	Fatty Acids, Nonesterified
-	D10.251.355	Fatty Acids, Unsaturated
-	D10.251.355.255	Eicosanoids
-	D10.251.355.255.100	Arachidonic Acids
-	D10.251.355.255.100.100	Arachidonic Acid
-	D10.251.355.255.100.300	Hydroxyeicosatetraenoic Acids
-	D10.251.355.255.100.300.425	12-Hydroxy-5,8,10,14-eicosatetraenoic Acid
-	D10.251.355.255.100.375	Isoprostanes
-	D10.251.355.255.100.375.500	F2-Isoprostanes
-	D10.251.355.255.100.375.750	Neuroprostanes
-	D10.251.355.255.100.450	Leukotrienes
-	D10.251.355.255.100.450.405	Leukotriene A4
-	D10.251.355.255.100.450.411	Leukotriene B4
-	D10.251.355.255.100.450.855	SRS-A
-	D10.251.355.255.100.450.855.455	Leukotriene C4
-	D10.251.355.255.100.450.855.461	Leukotriene D4
-	D10.251.355.255.100.450.855.470	Leukotriene E4
-	D10.251.355.255.100.825	Thromboxanes
-	D10.251.355.255.100.825.800	Thromboxane A2
-	D10.251.355.255.100.825.810	Thromboxane B2
-	D10.251.355.255.200	Eicosapentaenoic Acid
-	D10.251.355.255.205	5,8,11,14-Eicosatetraenoic Acid

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D10.251.355.255.207 8,11,14-Eicosatrienoic Acid
-	D10.251.355.255.375 Lipoxins
-	D10.251.355.255.550 Prostaglandins
-	D10.251.355.255.550.025 Prostaglandin Endoperoxides
-	D10.251.355.255.550.025.600 Prostaglandins G
-	D10.251.355.255.550.025.650 Prostaglandins H
-	D10.251.355.255.550.025.650.500 Prostaglandin H2
-	D10.251.355.255.550.025.650.500.500 (epoxymethano)prosta-5,13-dienoic Acid 15-Hydroxy-11 alpha,9 alpha-
-	D10.251.355.255.550.100 Prostaglandins A
-	D10.251.355.255.550.150 Prostaglandins B
-	D10.251.355.255.550.200 Prostaglandins D
-	D10.251.355.255.550.200.200 Prostaglandin D2
-	D10.251.355.255.550.250 Prostaglandins E
-	D10.251.355.255.550.250.100 Alprostadil
-	D10.251.355.255.550.250.200 Dinoprostone
-	D10.251.355.255.550.400 Prostaglandins F
-	D10.251.355.255.550.400.200 Dinoprost
-	D10.251.355.255.550.400.350 6-Ketoprostaglandin F1 alpha
-	D10.251.355.255.550.550 Prostaglandins I
-	D10.251.355.255.550.550.500 Epoprostenol
-	D10.251.355.255.550.775 Prostaglandins, Synthetic
-	D10.251.355.255.550.775.125 Iloprost
-	D10.251.355.255.550.775.250 Prostaglandin Endoperoxides, Synthetic
-	D10.251.355.255.550.775.350 Prostaglandins A, Synthetic
-	D10.251.355.255.550.775.450 Prostaglandins E, Synthetic
-	D10.251.355.255.550.775.450.050 Arbabprostil
-	D10.251.355.255.550.775.450.300 16,16-Dimethylprostaglandin E2
-	D10.251.355.255.550.775.450.350 Enprostil
-	D10.251.355.255.550.775.450.500 Misoprostol
-	D10.251.355.255.550.775.450.750 Rioprostil
-	D10.251.355.255.550.775.500 Prostaglandins F, Synthetic
-	D10.251.355.255.550.775.500.150 Carboprost
-	D10.251.355.255.550.775.500.175 Cloprostenol
-	D10.251.355.255.550.775.500.175.250 Bimatoprost
-	D10.251.355.255.550.775.500.175.500 Travoprost



## MeSH Tree Changes for 2017

Type	Tree - heading
-	D10.251.355.310 Fatty Acids, Essential
-	D10.251.355.310.166 Arachidonic Acids
-	D10.251.355.310.166.100 Arachidonic Acid
-	D10.251.355.310.166.550 Hydroxyeicosatetraenoic Acids
-	D10.251.355.310.166.550.425 12-Hydroxy-5,8,10,14-eicosatetraenoic Acid
-	D10.251.355.310.166.775 Isoprostanes
-	D10.251.355.310.166.775.500 F2-Isoprostanes
-	D10.251.355.310.166.775.750 Neuroprostanes
-	D10.251.355.310.166.887 Leukotrienes
-	D10.251.355.310.166.887.405 Leukotriene A4
-	D10.251.355.310.166.887.411 Leukotriene B4
-	D10.251.355.310.166.887.855 SRS-A
-	D10.251.355.310.166.887.855.455 Leukotriene C4
-	D10.251.355.310.166.887.855.461 Leukotriene D4
-	D10.251.355.310.166.887.855.470 Leukotriene E4
-	D10.251.355.310.166.971 Thromboxanes
-	D10.251.355.310.166.971.800 Thromboxane A2
-	D10.251.355.310.166.971.810 Thromboxane B2
-	D10.251.355.310.515 Linoleic Acids
-	D10.251.355.310.515.500 Linoleic Acid
-	D10.251.355.310.640 Linolenic Acids
-	D10.251.355.310.640.400 alpha-Linolenic Acid
-	D10.251.355.310.640.425 gamma-Linolenic Acid
-	D10.251.355.325 Fatty Acids, Monounsaturated
-	D10.251.355.325.050 Alprostadil
-	D10.251.355.325.050.500 Lubiprostone
-	D10.251.355.325.190 Capsaicin
-	D10.251.355.325.200 Cilastatin
-	D10.251.355.325.300 Erucic Acids
-	D10.251.355.325.450 6-Ketoprostaglandin F1 alpha
-	D10.251.355.325.600 Oleic Acids
-	D10.251.355.325.600.525 Oleic Acid
-	D10.251.355.325.600.780 Ricinoleic Acids
-	D10.251.355.325.919 Undecylenic Acids
-	D10.251.355.337 Fatty Acids, Omega-3
-	D10.251.355.337.100 alpha-Linolenic Acid

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D10.251.355.337.250 Docosahexaenoic Acids
-	D10.251.355.337.250.500 Neuroprostanes
-	D10.251.355.337.290 Eicosapentaenoic Acid
-	D10.251.355.343 Fatty Acids, Omega-6
-	D10.251.355.343.249 gamma-Linolenic Acid
-	D10.251.355.343.500 Linoleic Acids
-	D10.251.355.343.500.500 Linoleic Acid
-	D10.251.355.343.500.750 Linoleic Acids, Conjugated
-	D10.251.355.350 Gefarnate
-	D10.251.355.391 Ionomycin
-	D10.251.355.411 Isoprostanes
-	D10.251.355.411.500 F2-Isoprostanes
-	D10.251.355.411.750 Neuroprostanes
-	D10.251.355.645 Oxylipins
-	D10.251.355.840 Sorbic Acid
-	D10.251.355.920 Trans Fatty Acids
-	D10.251.400 Fatty Acids, Volatile
-	D10.251.400.045 Acetates
-	D10.251.400.045.500 Acetic Acid
-	D10.251.400.143 Butyrates
-	D10.251.400.143.500 Butyric Acid
-	D10.251.400.143.687 Crotonates
-	D10.251.400.143.781 Hydroxybutyrates
-	D10.251.400.143.781.500 3-Hydroxybutyric Acid
-	D10.251.400.143.875 Isobutyrate
-	D10.251.400.326 Caproates
-	D10.251.400.706 Propionates
-	D10.251.400.895 Valerates
-	D10.251.400.895.593 Pentanoic Acids
-	D10.251.400.895.593.350 Gemfibrozil
-	D10.251.400.895.593.900 Valproic Acid
-	D10.251.450 Heptanoic Acids
-	D10.251.450.200 Atorvastatin Calcium
-	D10.251.450.400 Heptanoates
-	D10.251.500 Lauric Acids
-	D10.251.500.410 Laurates

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D10.251.536 Mupirocin
-	D10.251.572 Mycolic Acids
-	D10.251.618 Mycophenolic Acid
-	D10.251.640 Myristic Acids
-	D10.251.640.610 Myristates
-	D10.251.640.630 Myristic Acid
-	D10.251.694 Palmitic Acids
-	D10.251.694.600 Palmitates
-	D10.251.694.600.500 Palmitoyl Coenzyme A
-	D10.251.694.750 Palmitic Acid
-	D10.251.740 Prostanoic Acids
-	D10.251.860 Sodium Morrhuate
-	D10.251.882 Stearic Acids
-	D10.251.882.800 Stearates
-	D10.251.941 Thioctic Acid
-	D10.289 Fatty Alcohols
-	D10.289.054 Acetogenins
-	D10.289.110 Butanols
-	D10.289.110.175 1-Butanol
-	D10.289.110.220 Chlorobutanol
-	D10.289.110.855 tert-Butyl Alcohol
-	D10.289.220 Dodecanol
-	D10.289.220.720 Sodium Dodecyl Sulfate
-	D10.289.230 Dolichol
-	D10.289.230.250 Dolichol Phosphates
-	D10.289.400 Farnesol
-	D10.289.500 Heptanol
-	D10.289.510 Hexanols
-	D10.289.510.500 Cyclohexanols
-	D10.289.510.500.303 Desvenlafaxine Succinate
-	D10.289.510.500.605 Menthol
-	D10.289.510.500.802 Tramadol
-	D10.289.510.500.901 Venlafaxine Hydrochloride
-	D10.289.600 Octanols
-	D10.289.600.610 1-Octanol
-	D10.289.640 Pentanols

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D10.289.900 Sodium Tetradecyl Sulfate
-	D10.351 Glycerides
-	D10.351.303 Diglycerides
-	D10.351.676 Monoglycerides
-	D10.351.801 Triglycerides
-	D10.351.801.632 Triacetin
-	D10.351.801.801 Triolein
-	D10.390 Glycolipids
-	D10.390.240 Cord Factors
-	D10.390.355 Galactolipids
-	D10.390.470 Glycosphingolipids
-	D10.390.470.025 Acidic Glycosphingolipids
-	D10.390.470.025.475 Gangliosides
-	D10.390.470.025.475.390 G(M1) Ganglioside
-	D10.390.470.025.475.400 G(M2) Ganglioside
-	D10.390.470.025.475.510 G(M3) Ganglioside
-	D10.390.470.025.837 Sulfoglycosphingolipids
-	D10.390.470.675 Neutral Glycosphingolipids
-	D10.390.470.675.200 Ceramides
-	D10.390.470.675.200.250 Cerebrosides
-	D10.390.470.675.200.250.450 Galactosylceramides
-	D10.390.470.675.200.250.490 Glucosylceramides
-	D10.390.470.675.200.425 Globosides
-	D10.390.470.675.200.612 Lactosylceramides
-	D10.390.470.675.200.906 Trihexosylceramides
-	D10.390.470.675.870 Sphingomyelins
-	D10.390.470.806 Psychosine
-	D10.390.475 Glycosylphosphatidylinositols
-	D10.390.700 Polyisoprenyl Phosphate Sugars
-	D10.390.700.700 Polyisoprenyl Phosphate Monosaccharides
-	D10.390.700.700.250 Dolichol Monophosphate Mannose
-	D10.390.700.710 Polyisoprenyl Phosphate Oligosaccharides
-	D10.440 Lipid Peroxides
-	D10.460 Lipofuscin
-	D10.477 Lipopeptides
-	D10.477.500 Daptomycin

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D10.477.750	Polymyxins
-	D10.477.750.110	Colistin
-	D10.477.750.750	Polymyxin B
-	D10.494	Lipopolysaccharides
-	D10.494.500	Lipid A
-	D10.494.600	O Antigens
-	D10.532	Lipoproteins
-	D10.532.091	Apolipoproteins
-	D10.532.091.200	Apolipoproteins A
-	D10.532.091.200.100	Apolipoprotein A-I
-	D10.532.091.200.150	Apolipoprotein A-II
New Heading	<b>D10.532.091.200.575</b>	<b>Apolipoprotein A-V</b>
-	D10.532.091.300	Apolipoproteins B
-	D10.532.091.300.240	Apolipoprotein B-48
-	D10.532.091.300.249	Apolipoprotein B-100
-	D10.532.091.400	Apolipoproteins C
-	D10.532.091.400.500	Apolipoprotein C-I
-	D10.532.091.400.750	Apolipoprotein C-II
-	D10.532.091.400.875	Apolipoprotein C-III
-	D10.532.091.450	Apolipoproteins D
-	D10.532.091.500	Apolipoproteins E
-	D10.532.091.500.249	Apolipoprotein E2
-	D10.532.091.500.500	Apolipoprotein E3
-	D10.532.091.500.750	Apolipoprotein E4
New Heading	<b>D10.532.137</b>	<b>ATP Binding Cassette Transporter, Sub-Family G, Member 5</b>
New Heading	<b>D10.532.160</b>	<b>ATP Binding Cassette Transporter, Sub-Family G, Member 8</b>
-	D10.532.183	Chylomicrons
-	D10.532.183.500	Chylomicron Remnants
-	D10.532.350	Lipoprotein(a)
-	D10.532.350.500	Apoprotein(a)
-	D10.532.400	Lipoprotein-X
-	D10.532.432	Lipoproteins, HDL
-	D10.532.432.400	Cholesterol, HDL
-	D10.532.432.550	High-Density Lipoproteins, Pre-beta

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D10.532.432.700 Lipoproteins, HDL2
-	D10.532.432.850 Lipoproteins, HDL3
-	D10.532.473 Lipoproteins, IDL
-	D10.532.515 Lipoproteins, LDL
-	D10.532.515.500 Cholesterol, LDL
-	D10.532.599 Lipoproteins, VLDL
-	D10.532.599.700 Cholesterol, VLDL
-	D10.570 Membrane Lipids
-	D10.570.510 Lipid Bilayers
-	D10.570.755 Phospholipids
-	D10.570.755.375 Glycerophosphates
-	D10.570.755.375.760 Phosphatidic Acids
-	D10.570.755.375.760.400 Glycerophospholipids
-	D10.570.755.375.760.400.386 Glycerolphosphorylcholine
-	D10.570.755.375.760.400.800 Phosphatidylcholines
-	D10.570.755.375.760.400.800.200 Dimyristoylphosphatidylcholine
-	D10.570.755.375.760.400.800.224 1,2-Dipalmitoylphosphatidylcholine
-	D10.570.755.375.760.400.800.806 Lecithins
-	D10.570.755.375.760.400.800.806.500 Glycerolphosphorylcholine
-	D10.570.755.375.760.400.840 Phosphatidylethanolamines
-	D10.570.755.375.760.400.885 Phosphatidylglycerols
-	D10.570.755.375.760.400.885.185 Cardiolipins
-	D10.570.755.375.760.400.942 Phosphatidylinositols
-	D10.570.755.375.760.400.942.250 Glycosylphosphatidylinositols
-	D10.570.755.375.760.400.942.625 Phosphatidylinositol Phosphates
-	D10.570.755.375.760.400.942.625.900 Phosphatidylinositol 4,5-Diphosphate
-	D10.570.755.375.760.400.971 Phosphatidylserines
-	D10.570.755.375.760.400.985 Phospholipid Ethers
-	D10.570.755.375.760.400.985.820 Plasmalogens
-	D10.570.755.375.760.400.985.910 Platelet Activating Factor
-	D10.570.755.375.760.550 Lysophospholipids
-	D10.570.755.375.760.550.550 Lysophosphatidylcholines
-	D10.570.755.893 Sphingomyelins
-	D10.570.780 Proteolipids
-	D10.570.780.249 Myelin and Lymphocyte-Associated Proteolipid Proteins
-	D10.570.780.500 Myelin Proteolipid Protein

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D10.570.877 Sphingolipids
-	D10.570.877.360 Glycosphingolipids
-	D10.570.877.360.025 Acidic Glycosphingolipids
-	D10.570.877.360.025.475 Gangliosides
-	D10.570.877.360.025.475.390 G(M1) Ganglioside
-	D10.570.877.360.025.475.400 G(M2) Ganglioside
-	D10.570.877.360.025.475.510 G(M3) Ganglioside
-	D10.570.877.360.025.837 Sulfoglycosphingolipids
-	D10.570.877.360.612 Neutral Glycosphingolipids
-	D10.570.877.360.612.200 Ceramides
-	D10.570.877.360.612.200.250 Cerebrosides
-	D10.570.877.360.612.200.250.450 Galactosylceramides
-	D10.570.877.360.612.200.250.490 Glucosylceramides
-	D10.570.877.360.612.200.425 Globosides
-	D10.570.877.360.612.200.612 Lactosylceramides
-	D10.570.877.360.612.200.906 Trihexosylceramides
-	D10.570.877.360.612.870 Sphingomyelins
-	D10.570.877.360.806 Psychosine
-	D10.570.938 Sterols
-	D10.570.938.073 Adosterol
-	D10.570.938.146 Cholecalciferol
-	D10.570.938.146.478 Hydroxycholecalciferols
-	D10.570.938.146.478.250 Calcifediol
-	D10.570.938.146.478.387 Dihydroxycholecalciferols
-	D10.570.938.146.478.387.300 Calcitriol
-	D10.570.938.146.478.387.400 24,25-Dihydroxyvitamin D 3
-	D10.570.938.208 Cholesterol
-	D10.570.938.208.070 Azacosterol
-	D10.570.938.208.160 Cholestanol
-	D10.570.938.208.222 Cholesterol, Dietary
-	D10.570.938.208.250 Cholesterol Esters
-	D10.570.938.208.270 Cholesterol, HDL
-	D10.570.938.208.275 Cholesterol, LDL
-	D10.570.938.208.285 Cholesterol, VLDL
-	D10.570.938.208.320 Dehydrocholesterols
-	D10.570.938.208.320.297 Desmosterol

## MeSH Tree Changes for 2017

Type	Tree - heading	
Old Tree	<b>D10.570.938.208.603</b>	<b>Hydroxycholesterols</b>
-	D10.570.938.208.627	19-Iodocholesterol
Old Tree	<b>D10.570.938.208.650</b>	<b>Ketocholesterols</b>
New Heading	<b>D10.570.938.208.825</b>	<b>Oxysterols</b>
New Tree	<b>D10.570.938.208.825.500</b>	<b>Hydroxycholesterols</b>
New Tree	<b>D10.570.938.208.825.750</b>	<b>Ketocholesterols</b>
-	D10.570.938.439	Ergocalciferols
-	D10.570.938.439.500	25-Hydroxyvitamin D 2
-	D10.570.938.439.750	Dihydrotachysterol
-	D10.570.938.515	Fusidic Acid
-	D10.570.938.590	Lanosterol
-	D10.570.938.795	Phytosterols
-	D10.570.938.795.071	Brassinosteroids
-	D10.570.938.795.143	Ecdysteroids
Old Tree	<b>D10.570.938.795.287</b>	<b>Ergosterol</b>
Old Tree	<b>D10.570.938.795.287.888</b>	<b>Withanolides</b>
-	D10.570.938.795.669	Sitosterols
-	D10.570.938.795.808	Stigmasterol
New Tree	<b>D10.570.938.795.904</b>	<b>Withanolides</b>
-	D10.570.938.897	Solanine
-	D10.627	Oils
-	D10.627.430	Fish Oils
-	D10.627.430.354	Cod Liver Oil
-	D10.627.430.450	Fatty Acids, Omega-3
-	D10.627.430.450.375	Docosahexaenoic Acids
-	D10.627.430.450.390	Eicosapentaenoic Acid
-	D10.627.675	Oils, Volatile
-	D10.627.675.775	Tea Tree Oil
-	D10.627.675.800	Turpentine
-	D10.627.700	Plant Oils
-	D10.627.700.132	Castor Oil
-	D10.627.700.240	Corn Oil
-	D10.627.700.263	Cottonseed Oil



## MeSH Tree Changes for 2017

Type	Tree - heading
-	D10.627.700.315 Croton Oil
-	D10.627.700.550 Iodized Oil
-	D10.627.700.550.300 Ethiodized Oil
-	D10.627.700.615 Linseed Oil
-	D10.627.700.728 Olive Oil
-	D10.627.700.840 Safflower Oil
-	D10.627.700.865 Sesame Oil
-	D10.627.700.880 Soybean Oil
-	D10.627.700.940 Tea Tree Oil
-	D10.751 Polyhydroxyalkanoates
-	D10.945 Waxes
-	D10.945.507 Lanolin
-	D12 Amino Acids, Peptides, and Proteins
-	D12.125 Amino Acids
-	D12.125.042 Alanine
-	D12.125.042.070 beta-Alanine
-	D12.125.042.070.500 Pantothenic Acid
-	D12.125.042.475 Lysinoalanine
-	D12.125.042.500 Mimosine
-	D12.125.065 Amino Acid Chloromethyl Ketones
-	D12.125.065.850 Tosyllysine Chloromethyl Ketone
-	D12.125.065.930 Tosylphenylalanyl Chloromethyl Ketone
-	D12.125.067 Amino Acids, Acidic
-	D12.125.067.500 Aspartic Acid
-	D12.125.067.500.150 D-Aspartic Acid
-	D12.125.067.500.275 Isoaspartic Acid
-	D12.125.067.500.400 N-Methylaspartate
-	D12.125.067.500.700 Potassium Magnesium Aspartate
-	D12.125.067.625 Glutamates
-	D12.125.067.625.174 1-Carboxyglutamic Acid
-	D12.125.067.625.349 Glutamic Acid
-	D12.125.067.625.349.850 Sodium Glutamate
-	D12.125.067.625.525 Pemetrexed
-	D12.125.067.625.700 Polyglutamic Acid
-	D12.125.067.625.850 Pyrrolidonecarboxylic Acid
-	D12.125.068 Amino Acids, Basic

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.125.068.050 Arginine
-	D12.125.068.050.075 Argininosuccinic Acid
-	D12.125.068.050.095 Benzoylarginine-2-Naphthylamide
-	D12.125.068.050.100 Benzoylarginine Nitroanilide
-	D12.125.068.050.400 Homoarginine
-	D12.125.068.050.525 NG-Nitroarginine Methyl Ester
-	D12.125.068.050.587 Nitroarginine
-	D12.125.068.050.650 omega-N-Methylarginine
-	D12.125.068.050.900 Tosylarginine Methyl Ester
-	D12.125.068.060 Asparagine
-	D12.125.068.330 Glutamine
-	D12.125.068.330.700 Proglumide
-	D12.125.068.555 Lysine
-	D12.125.068.555.478 Hydroxylysine
-	D12.125.068.555.575 Lysinoalanine
-	D12.125.068.555.750 Polylysine
-	D12.125.068.665 Ornithine
-	D12.125.068.665.340 Eflornithine
-	D12.125.070 Amino Acids, Branched-Chain
-	D12.125.070.075 Aminoisobutyric Acids
-	D12.125.070.577 Isoleucine
-	D12.125.070.637 Leucine
-	D12.125.070.950 Valine
-	D12.125.070.950.100 2-Amino-5-phosphonovalerate
-	D12.125.070.950.550 Valsartan
-	D12.125.070.950.550.500 Amlodipine, Valsartan Drug Combination
-	D12.125.072 Amino Acids, Cyclic
-	D12.125.072.050 Amino Acids, Aromatic
-	D12.125.072.050.342 Dextrothyroxine
-	D12.125.072.050.685 Phenylalanine
-	D12.125.072.050.685.400 Dihydroxyphenylalanine
-	D12.125.072.050.685.400.180 Cysteinyldopa
-	D12.125.072.050.685.400.500 Levodopa
-	D12.125.072.050.685.400.600 Methylodopa
-	D12.125.072.050.685.440 Fenclonine
-	D12.125.072.050.685.445 N-Formylmethionine Leucyl-Phenylalanine

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.125.072.050.685.450                      p-Fluorophenylalanine
-	D12.125.072.050.685.500                      Melphalan
-	D12.125.072.050.767                            Thyroxine
-	D12.125.072.050.767.741                      Thyronines
-	D12.125.072.050.767.741.180                Diiodothyronines
-	D12.125.072.050.767.741.894                Triiodothyronine
-	D12.125.072.050.767.741.947                Triiodothyronine, Reverse
-	D12.125.072.050.850                            Tryptophan
-	D12.125.072.050.850.479                      5-Hydroxytryptophan
-	D12.125.072.050.875                            Tyrosine
-	D12.125.072.050.875.064                      Betalains
-	D12.125.072.050.875.064.500                Betacyanins
-	D12.125.072.050.875.130                      Dihydroxyphenylalanine
-	D12.125.072.050.875.130.180                Cysteinyldopa
-	D12.125.072.050.875.130.500                Levodopa
-	D12.125.072.050.875.130.600                Methyldopa
-	D12.125.072.050.875.262                      Diiodotyrosine
-	D12.125.072.050.875.379                      Melanins
-	D12.125.072.050.875.485                      Methyltyrosines
-	D12.125.072.050.875.485.050                alpha-Methyltyrosine
-	D12.125.072.050.875.496                      Monoiodotyrosine
-	D12.125.072.050.875.750                      Phosphotyrosine
-	D12.125.072.170                                Cycloleucine
-	D12.125.072.200                                Desmosine
-	D12.125.072.329                                Histidine
-	D12.125.072.329.269                            Ergothioneine
-	D12.125.072.329.539                            Methylhistidines
-	D12.125.072.401                                Imino Acids
-	D12.125.072.401.200                            Azetidinecarboxylic Acid
-	D12.125.072.401.623                            Proline
-	D12.125.072.401.623.270                      Captopril
-	D12.125.072.401.623.374                      Fosinopril
-	D12.125.072.401.623.478                      Hydroxyproline
-	D12.125.072.401.761                            Pyrrolidonecarboxylic Acid
-	D12.125.072.401.830                            Technetium Tc 99m Diethyl-iminodiacetic Acid
-	D12.125.072.401.840                            Technetium Tc 99m Disofenin

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.125.072.401.900                      Technetium Tc 99m Lidofenin
-	D12.125.072.415                              Isodesmosine
-	D12.125.095                                    Amino Acids, Diamino
-	D12.125.095.104                              Arginine
-	D12.125.095.104.075                        Argininosuccinic Acid
-	D12.125.095.104.095                        Benzoylarginine-2-Naphthylamide
-	D12.125.095.104.100                        Benzoylarginine Nitroanilide
-	D12.125.095.104.400                        Homoarginine
-	D12.125.095.104.525                        NG-Nitroarginine Methyl Ester
-	D12.125.095.104.587                        Nitroarginine
-	D12.125.095.104.650                        omega-N-Methylarginine
-	D12.125.095.104.900                        Tosylarginine Methyl Ester
-	D12.125.095.165                              Asparagine
-	D12.125.095.226                              Citrulline
-	D12.125.095.307                              Cystathionine
-	D12.125.095.369                              Cystine
-	D12.125.095.390                              Diaminopimelic Acid
-	D12.125.095.461                              Glutamine
-	D12.125.095.461.700                        Proglumide
-	D12.125.095.533                              Homocystine
-	D12.125.095.647                              Lysine
-	D12.125.095.647.478                        Hydroxylysine
-	D12.125.095.647.575                        Lysinoalanine
-	D12.125.095.647.750                        Polylysine
-	D12.125.095.765                              Ornithine
-	D12.125.095.765.340                        Eflornithine
-	D12.125.119                                    Amino Acids, Dicarboxylic
-	D12.125.119.075                              2-Aminoadipic Acid
-	D12.125.119.170                              Aspartic Acid
-	D12.125.119.170.150                        D-Aspartic Acid
-	D12.125.119.170.275                        Isoaspartic Acid
-	D12.125.119.170.400                        N-Methylaspartate
-	D12.125.119.170.700                        Potassium Magnesium Aspartate
-	D12.125.119.270                              Carbocysteine
-	D12.125.119.307                              Cystathionine
-	D12.125.119.369                              Cystine

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.125.119.409                      Glutamates
-	D12.125.119.409.174                      1-Carboxyglutamic Acid
-	D12.125.119.409.349                      Glutamic Acid
-	D12.125.119.409.349.575                      Sodium Glutamate
-	D12.125.119.409.525                      Pemetrexed
-	D12.125.119.409.700                      Polyglutamic Acid
-	D12.125.119.658                      Homocystine
-	D12.125.142                      Amino Acids, Essential
-	D12.125.142.087                      Arginine
-	D12.125.142.087.500                      omega-N-Methylarginine
-	D12.125.142.308                      Histidine
-	D12.125.142.383                      Isoleucine
-	D12.125.142.441                      Leucine
-	D12.125.142.497                      Lysine
-	D12.125.142.557                      Methionine
-	D12.125.142.557.500                      Racemethionine
-	D12.125.142.666                      Phenylalanine
-	D12.125.142.666.500                      N-Formylmethionine Leucyl-Phenylalanine
-	D12.125.142.815                      Threonine
-	D12.125.142.815.500                      Phosphothreonine
-	D12.125.142.875                      Tryptophan
-	D12.125.142.930                      Valine
-	D12.125.142.930.500                      Valsartan
-	D12.125.142.930.500.500                      Amlodipine, Valsartan Drug Combination
-	D12.125.154                      Amino Acids, Neutral
-	D12.125.154.049                      Asparagine
-	D12.125.154.299                      Cysteine
-	D12.125.154.424                      Glutamine
-	D12.125.154.549                      Methionine
-	D12.125.154.549.500                      Racemethionine
-	D12.125.154.800                      Serine
-	D12.125.154.800.500                      Azaserine
-	D12.125.154.800.750                      Cycloserine
-	D12.125.154.800.875                      Droxidopa
-	D12.125.154.800.937                      Enterobactin
-	D12.125.154.800.968                      Phosphoserine

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.125.154.900 Threonine
-	D12.125.154.900.500 Phosphothreonine
-	D12.125.166 Amino Acids, Sulfur
-	D12.125.166.175 Cystathionine
-	D12.125.166.215 Cysteic Acid
-	D12.125.166.230 Cysteine
-	D12.125.166.230.259 Acetylcysteine
-	D12.125.166.230.310 Carbocysteine
-	D12.125.166.230.330 Cysteinyl-dopa
-	D12.125.166.230.369 Cystine
-	D12.125.166.230.700 Selenocysteine
-	D12.125.166.388 Ethionine
-	D12.125.166.498 Homocysteine
-	D12.125.166.498.050 S-Adenosylhomocysteine
-	D12.125.166.554 Homocystine
-	D12.125.166.676 Methionine
-	D12.125.166.676.089 Racemethionine
-	D12.125.166.676.180 S-Adenosylmethionine
-	D12.125.166.676.450 N-Formylmethionine
-	D12.125.166.676.450.440 N-Formylmethionine Leucyl-Phenylalanine
-	D12.125.166.676.620 Methionine Sulfoximine
-	D12.125.166.676.620.125 Buthionine Sulfoximine
-	D12.125.166.676.900 Selenomethionine
-	D12.125.166.676.950 Vitamin U
-	D12.125.166.786 Penicillamine
-	D12.125.166.786.500 S-Nitroso-N-Acetylpenicillamine
-	D12.125.166.893 Thiorphan
-	D12.125.166.896 Tiopronin
-	D12.125.190 Aminobutyrate
-	D12.125.190.055 Aminoisobutyric Acids
-	D12.125.190.350 gamma-Aminobutyric Acid
-	D12.125.190.350.450 Pregabalin
-	D12.125.190.350.900 Vigabatrin
-	D12.125.213 Aminocaproate
-	D12.125.213.075 Aminocaproic Acid
-	D12.125.213.568 Norleucine

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.125.213.568.175 Diazooxonorleucine
-	D12.125.262 Aminolevulinic Acid
-	D12.125.311 Canavanine
-	D12.125.373 Creatine
-	D12.125.373.603 Phosphocreatine
-	D12.125.427 Excitatory Amino Acids
-	D12.125.427.040 Aspartic Acid
-	D12.125.427.300 Glutamic Acid
-	D12.125.481 Glycine
-	D12.125.481.100 Allylglycine
-	D12.125.481.700 N-substituted Glycines
-	D12.125.481.700.249 Glycocholic Acid
-	D12.125.481.700.249.420 Glycodeoxycholic Acid
-	D12.125.481.700.249.420.400 Glycochenodeoxycholic Acid
-	D12.125.481.700.374 Sarcosine
-	D12.125.481.700.750 Thiorphan
-	D12.125.481.700.760 Tiopronin
-	D12.125.526 Homoserine
-	D12.125.608 Kynurenine
-	D12.125.730 Oxamic Acid
-	D12.125.740 Phosphoamino Acids
-	D12.125.740.025 2-Amino-5-phosphonovalerate
-	D12.125.740.675 Phosphocreatine
-	D12.125.740.700 Phosphoserine
-	D12.125.740.725 Phosphothreonine
-	D12.125.740.740 Phosphotyrosine
-	D12.125.755 Quisqualic Acid
-	D12.125.780 RNA, Transfer, Amino Acyl
-	D12.644 Peptides
-	D12.644.024 Amyloid beta-Peptides
-	D12.644.050 Antimicrobial Cationic Peptides
-	D12.644.050.099 Cathelicidins
-	D12.644.050.149 Cecropins
-	D12.644.050.200 Defensins
-	D12.644.050.200.050 alpha-Defensins
-	D12.644.050.200.075 beta-Defensins

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.644.050.400 Dermcidins
-	D12.644.050.450 Histatins
-	D12.644.050.500 Magainins
-	D12.644.050.550 Melittin
-	D12.644.050.600 Polymyxins
-	D12.644.050.600.110 Colistin
-	D12.644.050.600.750 Polymyxin B
-	D12.644.050.800 Thionins
-	D12.644.082 Aptamers, Peptide
-	D12.644.098 Cell-Penetrating Peptides
-	D12.644.115 Charybdotoxin
-	D12.644.120 Cholecystokinin
-	D12.644.120.500 Sincalide
-	D12.644.128 Circadian Rhythm Signaling Peptides and Proteins
-	D12.644.128.049 ARNTL Transcription Factors
-	D12.644.128.100 CLOCK Proteins
-	D12.644.128.150 Cryptochromes
-	D12.644.128.575 Period Circadian Proteins
-	D12.644.132 Cystine-Knot Miniproteins
-	D12.644.136 Diazepam Binding Inhibitor
-	D12.644.138 Disintegrins
-	D12.644.140 Distamycins
-	D12.644.146 Edeine
-	D12.644.173 Erabutoxins
-	D12.644.200 Fibrinopeptide A
-	D12.644.205 Fibrinopeptide B
-	D12.644.219 Glatiramer Acetate
-	D12.644.233 Glycopeptides
-	D12.644.233.050 Acetylmuramyl-Alanyl-Isoglutamine
-	D12.644.233.110 Bleomycin
-	D12.644.233.110.690 Peplomycin
-	D12.644.233.110.710 Phleomycins
-	D12.644.233.594 Peptidoglycan
-	D12.644.233.697 Ristocetin
-	D12.644.233.800 Sialoglycoproteins
-	D12.644.233.800.174 Antigens, CD43



## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D12.644.233.800.350	Glycophorin
-	D12.644.233.900	Teicoplanin
-	D12.644.233.925	Vancomycin
-	D12.644.276	Intercellular Signaling Peptides and Proteins
-	D12.644.276.024	Adipokines
-	D12.644.276.024.249	Adiponectin
-	D12.644.276.024.500	Leptin
-	D12.644.276.024.750	Resistin
-	D12.644.276.049	Agouti Signaling Protein
-	D12.644.276.074	Agouti-Related Protein
Old Tree	D12.644.276.087	Alarmins
Old Tree	D12.644.276.087.044	alpha-Defensins
Old Tree	D12.644.276.087.054	beta-Defensins
Old Tree	D12.644.276.087.161	Calgranulin A
Old Tree	D12.644.276.087.171	Calgranulin B
Old Tree	D12.644.276.087.178	Cathelicidins
Old Tree	D12.644.276.087.185	Chaperonin 60
Old Tree	D12.644.276.087.371	HMGB1 Protein
Old Tree	D12.644.276.087.391	HSC70 Heat-Shock Proteins
Old Tree	D12.644.276.087.597	Interleukin-33
Old Tree	D12.644.276.087.802	S100 Calcium Binding Protein beta Subunit
Old Tree	D12.644.276.087.901	S100A12 Protein
-	D12.644.276.100	Angiogenic Proteins
-	D12.644.276.100.100	Angiopoietins
-	D12.644.276.100.100.100	Angiopoietin-1
-	D12.644.276.100.100.200	Angiopoietin-2
-	D12.644.276.100.450	Angiostatic Proteins
-	D12.644.276.100.450.500	Angiostatins
-	D12.644.276.100.450.750	Endostatins
-	D12.644.276.100.800	Vascular Endothelial Growth Factors
New Heading	<b>D12.644.276.100.800.100</b>	<b>Placenta Growth Factor</b>
-	D12.644.276.100.800.200	Vascular Endothelial Growth Factor A
-	D12.644.276.100.800.300	Vascular Endothelial Growth Factor B
-	D12.644.276.100.800.400	Vascular Endothelial Growth Factor C
-	D12.644.276.100.800.500	Vascular Endothelial Growth Factor D

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.644.276.100.800.600 Vascular Endothelial Growth Factor, Endocrine-Gland-Derived
-	D12.644.276.200 CCN Intercellular Signaling Proteins
-	D12.644.276.200.100 Connective Tissue Growth Factor
-	D12.644.276.200.200 Cysteine-Rich Protein 61
-	D12.644.276.200.500 Nephroblastoma Overexpressed Protein
-	D12.644.276.374 Cytokines
-	D12.644.276.374.200 Chemokines
-	D12.644.276.374.200.070 beta-Thromboglobulin
-	D12.644.276.374.200.100 Chemokines, C
-	D12.644.276.374.200.110 Chemokines, CC
-	D12.644.276.374.200.110.050 Chemokine CCL1
-	D12.644.276.374.200.110.150 Chemokine CCL3
-	D12.644.276.374.200.110.200 Chemokine CCL4
-	D12.644.276.374.200.110.250 Chemokine CCL5
-	D12.644.276.374.200.110.550 Chemokine CCL11
-	D12.644.276.374.200.110.850 Chemokine CCL17
-	D12.644.276.374.200.110.870 Chemokine CCL19
-	D12.644.276.374.200.110.880 Chemokine CCL20
-	D12.644.276.374.200.110.890 Chemokine CCL21
-	D12.644.276.374.200.110.900 Chemokine CCL22
-	D12.644.276.374.200.110.910 Chemokine CCL24
-	D12.644.276.374.200.110.930 Chemokine CCL27
-	D12.644.276.374.200.110.990 Monocyte Chemoattractant Proteins
-	D12.644.276.374.200.110.990.600 Chemokine CCL2
-	D12.644.276.374.200.110.990.800 Chemokine CCL7
-	D12.644.276.374.200.110.990.900 Chemokine CCL8
-	D12.644.276.374.200.120 Chemokines, CXC
-	D12.644.276.374.200.120.050 Chemokine CXCL1
-	D12.644.276.374.200.120.100 Chemokine CXCL2
-	D12.644.276.374.200.120.250 Chemokine CXCL5
-	D12.644.276.374.200.120.300 Chemokine CXCL6
-	D12.644.276.374.200.120.450 Chemokine CXCL9
-	D12.644.276.374.200.120.500 Chemokine CXCL10
-	D12.644.276.374.200.120.550 Chemokine CXCL11
-	D12.644.276.374.200.120.600 Chemokine CXCL12

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.644.276.374.200.120.650 Chemokine CXCL13
-	D12.644.276.374.200.120.800 Interleukin-8
-	D12.644.276.374.200.120.900 Platelet Factor 4
-	D12.644.276.374.200.130 Chemokines, CX3C
-	D12.644.276.374.200.130.500 Chemokine CX3CL1
-	D12.644.276.374.200.600 Macrophage Inflammatory Proteins
-	D12.644.276.374.200.600.150 Chemokine CCL3
-	D12.644.276.374.200.600.200 Chemokine CCL4
-	D12.644.276.374.200.600.870 Chemokine CCL19
-	D12.644.276.374.200.600.880 Chemokine CCL20
-	D12.644.276.374.200.600.940 Chemokine CXCL2
-	D12.644.276.374.305 Growth Differentiation Factor 15
-	D12.644.276.374.410 Hematopoietic Cell Growth Factors
-	D12.644.276.374.410.240 Colony-Stimulating Factors
-	D12.644.276.374.410.240.150 Erythropoietin
-	D12.644.276.374.410.240.150.500 Epoetin Alfa
-	D12.644.276.374.410.240.350 Granulocyte Colony-Stimulating Factor
-	D12.644.276.374.410.240.350.500 Filgrastim
-	D12.644.276.374.410.240.375 Granulocyte-Macrophage Colony-Stimulating Factor
-	D12.644.276.374.410.240.400 Interleukin-3
-	D12.644.276.374.410.240.500 Macrophage Colony-Stimulating Factor
-	D12.644.276.374.410.240.750 Thrombopoietin
-	D12.644.276.374.410.800 Stem Cell Factor
-	D12.644.276.374.420 Hepatocyte Growth Factor
-	D12.644.276.374.440 Interferons
-	D12.644.276.374.440.890 Interferon Type I
-	D12.644.276.374.440.890.250 Interferon-alpha
-	D12.644.276.374.440.890.275 Interferon-beta
-	D12.644.276.374.440.890.275.500 Interferon beta-1a
-	D12.644.276.374.440.890.275.750 Interferon beta-1b
-	D12.644.276.374.440.893 Interferon-gamma
-	D12.644.276.374.460 Interleukin 1 Receptor Antagonist Protein
-	D12.644.276.374.465 Interleukins
-	D12.644.276.374.465.010 Interleukin-1
-	D12.644.276.374.465.010.300 Interleukin-1alpha

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.644.276.374.465.010.600 Interleukin-1beta
-	D12.644.276.374.465.021 Interleukin-2
-	D12.644.276.374.465.032 Interleukin-3
-	D12.644.276.374.465.186 Interleukin-4
-	D12.644.276.374.465.202 Interleukin-5
-	D12.644.276.374.465.224 Interleukin-6
-	D12.644.276.374.465.246 Interleukin-7
-	D12.644.276.374.465.312 Interleukin-8
-	D12.644.276.374.465.377 Interleukin-9
-	D12.644.276.374.465.510 Interleukin-10
-	D12.644.276.374.465.511 Interleukin-11
-	D12.644.276.374.465.512 Interleukin-12
-	D12.644.276.374.465.512.249 Interleukin-12 Subunit p35
-	D12.644.276.374.465.512.500 Interleukin-12 Subunit p40
-	D12.644.276.374.465.513 Interleukin-13
-	D12.644.276.374.465.515 Interleukin-15
-	D12.644.276.374.465.516 Interleukin-16
-	D12.644.276.374.465.517 Interleukin-17
-	D12.644.276.374.465.518 Interleukin-18
-	D12.644.276.374.465.759 Interleukin-23
-	D12.644.276.374.465.759.249 Interleukin-12 Subunit p40
-	D12.644.276.374.465.759.500 Interleukin-23 Subunit p19
-	D12.644.276.374.465.800 Interleukin-27
-	D12.644.276.374.465.850 Interleukin-33
-	D12.644.276.374.470 Leukemia Inhibitory Factor
-	D12.644.276.374.480 Lymphokines
-	D12.644.276.374.480.372 Interleukin-2
-	D12.644.276.374.480.428 Leukocyte Migration-Inhibitory Factors
-	D12.644.276.374.480.438 Lymphotoxin-alpha
-	D12.644.276.374.480.615 Macrophage-Activating Factors
-	D12.644.276.374.480.615.350 Interferon-gamma
-	D12.644.276.374.480.625 Macrophage Migration-Inhibitory Factors
-	D12.644.276.374.480.700 Suppressor Factors, Immunologic
-	D12.644.276.374.480.750 Transfer Factor
-	D12.644.276.374.500 Monokines
-	D12.644.276.374.500.400 Interleukin-1

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.644.276.374.500.400.300 Interleukin-1alpha
-	D12.644.276.374.500.400.600 Interleukin-1beta
-	D12.644.276.374.500.800 Tumor Necrosis Factor-alpha
-	D12.644.276.374.562 Oncostatin M
-	D12.644.276.374.625 Osteopontin
-	D12.644.276.374.687 Transforming Growth Factor beta
-	D12.644.276.374.687.100 Transforming Growth Factor beta1
-	D12.644.276.374.687.200 Transforming Growth Factor beta2
-	D12.644.276.374.687.300 Transforming Growth Factor beta3
-	D12.644.276.374.750 Tumor Necrosis Factors
-	D12.644.276.374.750.030 Antigens, CD70
-	D12.644.276.374.750.061 B-Cell Activating Factor
-	D12.644.276.374.750.065 4-1BB Ligand
-	D12.644.276.374.750.092 CD30 Ligand
-	D12.644.276.374.750.124 CD40 Ligand
-	D12.644.276.374.750.186 Ectodysplasins
-	D12.644.276.374.750.249 Fas Ligand Protein
-	D12.644.276.374.750.500 Lymphotoxin-alpha
-	D12.644.276.374.750.505 Lymphotoxin alpha1, beta2 Heterotrimer
-	D12.644.276.374.750.515 Lymphotoxin-beta
-	D12.644.276.374.750.531 OX40 Ligand
-	D12.644.276.374.750.562 RANK Ligand
-	D12.644.276.374.750.625 TNF-Related Apoptosis-Inducing Ligand
-	D12.644.276.374.750.626 Tumor Necrosis Factor-alpha
-	D12.644.276.374.750.656 Tumor Necrosis Factor Ligand Superfamily Member 13
-	D12.644.276.374.750.690 Tumor Necrosis Factor Ligand Superfamily Member 14
-	D12.644.276.374.750.720 Tumor Necrosis Factor Ligand Superfamily Member 15
-	D12.644.276.382 EGF Family of Proteins
-	D12.644.276.382.249 Amphiregulin
-	D12.644.276.382.374 Betacellulin
-	D12.644.276.382.500 Epidermal Growth Factor
-	D12.644.276.382.531 Epigen
-	D12.644.276.382.562 Epiregulin
-	D12.644.276.382.625 Heparin-binding EGF-like Growth Factor

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.644.276.382.750 Transforming Growth Factor alpha
-	D12.644.276.390 Endothelial Growth Factors
-	D12.644.276.400 Endothelins
-	D12.644.276.400.225 Endothelin-1
-	D12.644.276.400.235 Endothelin-2
-	D12.644.276.400.245 Endothelin-3
-	D12.644.276.500 Ephrins
-	D12.644.276.500.100 Ephrin-A1
-	D12.644.276.500.200 Ephrin-A2
-	D12.644.276.500.300 Ephrin-A3
-	D12.644.276.500.400 Ephrin-A4
-	D12.644.276.500.500 Ephrin-A5
-	D12.644.276.500.600 Ephrin-B1
-	D12.644.276.500.700 Ephrin-B2
-	D12.644.276.500.800 Ephrin-B3
-	D12.644.276.624 Fibroblast Growth Factors
-	D12.644.276.624.110 Fibroblast Growth Factor 1
-	D12.644.276.624.120 Fibroblast Growth Factor 2
-	D12.644.276.624.130 Fibroblast Growth Factor 3
-	D12.644.276.624.140 Fibroblast Growth Factor 4
-	D12.644.276.624.150 Fibroblast Growth Factor 5
-	D12.644.276.624.160 Fibroblast Growth Factor 6
-	D12.644.276.624.170 Fibroblast Growth Factor 7
-	D12.644.276.624.180 Fibroblast Growth Factor 8
-	D12.644.276.624.190 Fibroblast Growth Factor 9
-	D12.644.276.624.200 Fibroblast Growth Factor 10
-	D12.644.276.671 Hedgehog Proteins
-	D12.644.276.812 Kinins
-	D12.644.276.812.169 Bradykinin
-	D12.644.276.812.169.400 Kallidin
-	D12.644.276.812.654 Kininogens
-	D12.644.276.812.654.350 Kininogen, High-Molecular-Weight
-	D12.644.276.812.654.400 Kininogen, Low-Molecular-Weight
-	D12.644.276.812.900 Tachykinins
-	D12.644.276.812.900.354 Eledoisin
-	D12.644.276.812.900.475 Kassinin

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D12.644.276.812.900.500	Neurokinin A
-	D12.644.276.812.900.550	Neurokinin B
-	D12.644.276.812.900.800	Physalaemin
-	D12.644.276.812.900.866	Substance P
-	D12.644.276.836	Kisspeptins
New Tree	<a href="#">D12.644.276.848</a>	<a href="#">Matrix Metalloproteinases, Secreted</a>
New Tree	<a href="#">D12.644.276.848.049</a>	<a href="#">Bone Morphogenetic Protein 1</a>
New Tree	<a href="#">D12.644.276.848.100</a>	<a href="#">Matrix Metalloproteinase 1</a>
New Tree	<a href="#">D12.644.276.848.150</a>	<a href="#">Matrix Metalloproteinase 2</a>
New Tree	<a href="#">D12.644.276.848.200</a>	<a href="#">Matrix Metalloproteinase 3</a>
New Tree	<a href="#">D12.644.276.848.250</a>	<a href="#">Matrix Metalloproteinase 7</a>
New Tree	<a href="#">D12.644.276.848.300</a>	<a href="#">Matrix Metalloproteinase 8</a>
New Tree	<a href="#">D12.644.276.848.350</a>	<a href="#">Matrix Metalloproteinase 9</a>
New Tree	<a href="#">D12.644.276.848.400</a>	<a href="#">Matrix Metalloproteinase 10</a>
New Tree	<a href="#">D12.644.276.848.450</a>	<a href="#">Matrix Metalloproteinase 11</a>
New Tree	<a href="#">D12.644.276.848.500</a>	<a href="#">Matrix Metalloproteinase 12</a>
New Tree	<a href="#">D12.644.276.848.550</a>	<a href="#">Matrix Metalloproteinase 13</a>
New Tree	<a href="#">D12.644.276.848.800</a>	<a href="#">Matrix Metalloproteinase 20</a>
-	D12.644.276.860	Nerve Growth Factors
-	D12.644.276.860.100	Brain-Derived Neurotrophic Factor
-	D12.644.276.860.212	Ciliary Neurotrophic Factor
-	D12.644.276.860.325	Glial Maturation Factor
-	D12.644.276.860.381	Glial Cell Line-Derived Neurotrophic Factors
-	D12.644.276.860.381.500	Glial Cell Line-Derived Neurotrophic Factor
-	D12.644.276.860.381.750	Neurturin
-	D12.644.276.860.437	Nerve Growth Factor
-	D12.644.276.860.550	Neuregulins

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D12.644.276.860.550.750	Neuregulin-1
-	D12.644.276.860.775	Neurotrophin 3
-	D12.644.276.860.887	Pituitary Adenylate Cyclase-Activating Polypeptide
-	D12.644.276.908	Parathyroid Hormone-Related Protein
-	D12.644.276.910	Platelet-Derived Growth Factor
-	D12.644.276.910.650	Proto-Oncogene Proteins c-sis
-	D12.644.276.923	Semaphorins
-	D12.644.276.923.374	Semaphorin-3A
New Heading	<b>D12.644.276.930</b>	<b>Serrate-Jagged Proteins</b>
New Heading	<b>D12.644.276.930.500</b>	<b>Jagged-1 Protein</b>
New Heading	<b>D12.644.276.930.750</b>	<b>Jagged-2 Protein</b>
-	D12.644.276.937	Somatomedins
-	D12.644.276.937.400	Insulin-Like Growth Factor I
-	D12.644.276.937.420	Insulin-Like Growth Factor II
-	D12.644.276.948	Tolloid-Like Metalloproteinases
-	D12.644.276.948.500	Bone Morphogenetic Protein 1
-	D12.644.276.954	TGF-beta Superfamily Proteins
-	D12.644.276.954.200	Bone Morphogenetic Proteins
-	D12.644.276.954.200.100	Bone Morphogenetic Protein 1
-	D12.644.276.954.200.200	Bone Morphogenetic Protein 2
-	D12.644.276.954.200.300	Bone Morphogenetic Protein 3
-	D12.644.276.954.200.400	Bone Morphogenetic Protein 4
-	D12.644.276.954.200.500	Bone Morphogenetic Protein 5
-	D12.644.276.954.200.600	Bone Morphogenetic Protein 6
-	D12.644.276.954.200.700	Bone Morphogenetic Protein 7
-	D12.644.276.954.200.900	Bone Morphogenetic Protein 15
-	D12.644.276.954.200.910	Growth Differentiation Factor 2
-	D12.644.276.954.200.950	Growth Differentiation Factor 10
-	D12.644.276.954.300	Growth Differentiation Factors
-	D12.644.276.954.300.049	Bone Morphogenetic Protein 15
-	D12.644.276.954.300.100	Growth Differentiation Factor 1
-	D12.644.276.954.300.200	Growth Differentiation Factor 2
-	D12.644.276.954.300.300	Growth Differentiation Factor 3
-	D12.644.276.954.300.500	Growth Differentiation Factor 5



## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.644.276.954.300.600 Growth Differentiation Factor 6
-	D12.644.276.954.300.800 Growth Differentiation Factor 9
-	D12.644.276.954.300.900 Growth Differentiation Factor 10
-	D12.644.276.954.300.915 Growth Differentiation Factor 15
-	D12.644.276.954.300.925 Myostatin
-	D12.644.276.954.550 Nodal Signaling Ligands
-	D12.644.276.954.550.100 Growth Differentiation Factor 1
-	D12.644.276.954.550.300 Growth Differentiation Factor 3
-	D12.644.276.954.550.475 Left-Right Determination Factors
-	D12.644.276.954.550.650 Nodal Protein
-	D12.644.276.954.775 Transforming Growth Factor beta
-	D12.644.276.954.775.100 Transforming Growth Factor beta1
-	D12.644.276.954.775.200 Transforming Growth Factor beta2
-	D12.644.276.954.775.300 Transforming Growth Factor beta3
-	D12.644.276.963 Transforming Growth Factors
-	D12.644.276.963.360 Transforming Growth Factor alpha
-	D12.644.360 Intracellular Signaling Peptides and Proteins
-	D12.644.360.008 3',5'-Cyclic-AMP Phosphodiesterases
-	D12.644.360.008.100 Cyclic Nucleotide Phosphodiesterases, Type 1
-	D12.644.360.008.200 Cyclic Nucleotide Phosphodiesterases, Type 2
-	D12.644.360.008.300 Cyclic Nucleotide Phosphodiesterases, Type 3
-	D12.644.360.008.400 Cyclic Nucleotide Phosphodiesterases, Type 4
-	D12.644.360.008.500 Cyclic Nucleotide Phosphodiesterases, Type 5
-	D12.644.360.008.600 Cyclic Nucleotide Phosphodiesterases, Type 6
-	D12.644.360.008.700 Cyclic Nucleotide Phosphodiesterases, Type 7
-	D12.644.360.009 3',5'-Cyclic-GMP Phosphodiesterases
-	D12.644.360.009.100 Cyclic Nucleotide Phosphodiesterases, Type 1
-	D12.644.360.009.200 Cyclic Nucleotide Phosphodiesterases, Type 2
-	D12.644.360.009.500 Cyclic Nucleotide Phosphodiesterases, Type 5
-	D12.644.360.009.750 Cyclic Nucleotide Phosphodiesterases, Type 6
-	D12.644.360.011 Activating Transcription Factor 6
-	D12.644.360.024 Adaptor Proteins, Signal Transducing
-	D12.644.360.024.050 14-3-3 Proteins
-	D12.644.360.024.065 A Kinase Anchor Proteins
New Tree	<a href="#">D12.644.360.024.098</a> <a href="#">Arrestins</a>

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	<a href="#">D12.644.360.024.098.050</a>	<a href="#">Arrestin</a>
New Heading	<b>D12.644.360.024.098.525</b>	<b>Beta-Arrestins</b>
New Heading	<b>D12.644.360.024.098.525.500</b>	<b>beta-Arrestin 1</b>
New Heading	<b>D12.644.360.024.098.525.750</b>	<b>Beta-Arrestin 2</b>
-	D12.644.360.024.131	CARD Signaling Adaptor Proteins
-	D12.644.360.024.131.124	Apoptotic Protease-Activating Factor 1
New Tree	<a href="#">D12.644.360.024.131.155</a>	<a href="#">Caspase 9</a>
-	D12.644.360.024.131.186	CRADD Signaling Adaptor Protein
-	D12.644.360.024.131.249	Nod1 Signaling Adaptor Protein
-	D12.644.360.024.131.500	Nod2 Signaling Adaptor Protein
-	D12.644.360.024.131.750 Kinase 2	Receptor-Interacting Protein Serine-Threonine
-	D12.644.360.024.264	Caveolin 1
-	D12.644.360.024.272	Caveolin 2
-	D12.644.360.024.280	Cortactin
New Tree	<a href="#">D12.644.360.024.282</a>	<a href="#">Crk-Associated Substrate Protein</a>
New Tree	<a href="#">D12.644.360.024.285</a>	<a href="#">Death Domain Receptor Signaling Adaptor Proteins</a>
New Tree	<a href="#">D12.644.360.024.285.024</a> Protein	<a href="#">CASP8 and FADD-Like Apoptosis Regulating</a>
New Tree	<a href="#">D12.644.360.024.285.037</a>	<a href="#">Caspase 8</a>
New Tree	<a href="#">D12.644.360.024.285.050</a>	<a href="#">CRADD Signaling Adaptor Protein</a>
New Tree	<a href="#">D12.644.360.024.285.100</a>	<a href="#">Edar-Associated Death Domain Protein</a>
New Tree	<a href="#">D12.644.360.024.285.200</a>	<a href="#">Fas-Associated Death Domain Protein</a>
New Tree	<a href="#">D12.644.360.024.285.400</a> Kinases	<a href="#">Receptor-Interacting Protein Serine-Threonine</a>
New Tree	<a href="#">D12.644.360.024.285.600</a>	<a href="#">TNF Receptor-Associated Death Domain Protein</a>
New Heading	<b>D12.644.360.024.288</b>	<b>Dishevelled Proteins</b>
New Tree	<a href="#">D12.644.360.024.290</a>	<a href="#">GRB2 Adaptor Protein</a>

## MeSH Tree Changes for 2017

Type	Tree - heading	
Old Tree	D12.644.360.024.295	Crk-Associated Substrate Protein
Old Tree	D12.644.360.024.296	Death Domain Receptor Signaling Adaptor Proteins
Old Tree	D12.644.360.024.296.024 Protein	CASP8 and FADD-Like Apoptosis Regulating
Old Tree	D12.644.360.024.296.050	CRADD Signaling Adaptor Protein
Old Tree	D12.644.360.024.296.100	Edar-Associated Death Domain Protein
Old Tree	D12.644.360.024.296.200	Fas-Associated Death Domain Protein
Old Tree	D12.644.360.024.296.400 Kinases	Receptor-Interacting Protein Serine-Threonine
Old Tree	D12.644.360.024.296.600	TNF Receptor-Associated Death Domain Protein
Old Tree	D12.644.360.024.297	GRB2 Adaptor Protein
Old Tree	D12.644.360.024.298	GRB7 Adaptor Protein
New Tree	D12.644.360.024.299	GRB7 Adaptor Protein
-	D12.644.360.024.300	GRB10 Adaptor Protein
-	D12.644.360.024.301	Insulin Receptor Substrate Proteins
-	D12.644.360.024.302	Interferon Regulatory Factors
-	D12.644.360.024.302.124	Interferon Regulatory Factor-1
-	D12.644.360.024.302.249	Interferon Regulatory Factor-2
-	D12.644.360.024.302.374	Interferon Regulatory Factor-3
-	D12.644.360.024.302.437	Interferon Regulatory Factor-7
-	D12.644.360.024.302.500 Subunit	Interferon-Stimulated Gene Factor 3, gamma
-	D12.644.360.024.303	Interferon-Stimulated Gene Factor 3
-	D12.644.360.024.303.500	Interferon-Stimulated Gene Factor 3, alpha Subunit
-	D12.644.360.024.303.500.500	STAT1 Transcription Factor
-	D12.644.360.024.303.500.750	STAT2 Transcription Factor
-	D12.644.360.024.303.750 Subunit	Interferon-Stimulated Gene Factor 3, gamma
Old Tree	D12.644.360.024.304	Mediator Complex
Old Tree	D12.644.360.024.304.249	Cyclin C
Old Tree	D12.644.360.024.304.500	Cyclin-Dependent Kinase 8
Old Tree	D12.644.360.024.304.750	Mediator Complex Subunit 1
Old Tree	D12.644.360.024.305	Myeloid Differentiation Factor 88
New Heading	<b>D12.644.360.024.306</b>	<b>Kelch-Like ECH-Associated Protein 1</b>
Old Tree	D12.644.360.024.307	Nod Signaling Adaptor Proteins
Old Tree	D12.644.360.024.307.249	Nod1 Signaling Adaptor Protein

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	D12.644.360.024.307.500 Nod2 Signaling Adaptor Protein
Old Tree	D12.644.360.024.308 Nuclear Receptor Coactivators
Old Tree	D12.644.360.024.308.049 Mediator Complex Subunit 1
Old Tree	D12.644.360.024.308.100 Nuclear Receptor Coactivator 1
Old Tree	D12.644.360.024.308.200 Nuclear Receptor Coactivator 2
Old Tree	D12.644.360.024.308.300 Nuclear Receptor Coactivator 3
New Tree	D12.644.360.024.309 Mediator Complex
Old Tree	D12.644.360.024.309 Paxillin
New Tree	D12.644.360.024.309.249 Cyclin C
New Tree	D12.644.360.024.309.500 Cyclin-Dependent Kinase 8
New Tree	D12.644.360.024.309.750 Mediator Complex Subunit 1
Old Tree	D12.644.360.024.310 PII Nitrogen Regulatory Proteins
New Tree	D12.644.360.024.311 Myeloid Differentiation Factor 88
Old Tree	D12.644.360.024.311 Protein Inhibitors of Activated STAT
New Tree	D12.644.360.024.313 Nod Signaling Adaptor Proteins
New Tree	D12.644.360.024.313.249 Nod1 Signaling Adaptor Protein
New Tree	D12.644.360.024.313.500 Nod2 Signaling Adaptor Protein
New Tree	D12.644.360.024.314 Nuclear Receptor Coactivators
New Tree	D12.644.360.024.314.049 Mediator Complex Subunit 1
New Tree	D12.644.360.024.314.100 Nuclear Receptor Coactivator 1
New Tree	D12.644.360.024.314.200 Nuclear Receptor Coactivator 2
New Tree	D12.644.360.024.314.300 Nuclear Receptor Coactivator 3
New Heading	<b>D12.644.360.024.314.650</b> <b>Peroxisome Proliferator-Activated Receptor</b> <b>Gamma Coactivator 1-alpha</b>
New Tree	D12.644.360.024.316 Paxillin
New Tree	D12.644.360.024.318 PII Nitrogen Regulatory Proteins
Old Tree	D12.644.360.024.318 Proto-Oncogene Proteins c-crk

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	<a href="#">D12.644.360.024.319</a>	<a href="#">Protein Inhibitors of Activated STAT</a>
New Tree	<a href="#">D12.644.360.024.321</a>	<a href="#">Proto-Oncogene Proteins c-crk</a>
-	D12.644.360.024.326	Proto-Oncogene Proteins c-vav
-	D12.644.360.024.328	Retinoblastoma Binding Proteins
-	D12.644.360.024.328.049	E2F1 Transcription Factor
-	D12.644.360.024.328.100	Retinoblastoma-Binding Protein 1
-	D12.644.360.024.328.200	Retinoblastoma-Binding Protein 2
-	D12.644.360.024.328.600	Retinoblastoma-Binding Protein 4
-	D12.644.360.024.328.700	Retinoblastoma-Binding Protein 7
New Heading	<b><a href="#">D12.644.360.024.329</a></b>	<b><a href="#">Sequestosome-1 Protein</a></b>
-	D12.644.360.024.330	Shc Signaling Adaptor Proteins
New Heading	<b><a href="#">D12.644.360.024.330.500</a></b> <b><a href="#">Transforming Protein 1</a></b>	<b><a href="#">Src Homology 2 Domain-Containing,</a></b>
New Heading	<b><a href="#">D12.644.360.024.330.750</a></b> <b><a href="#">Transforming Protein 2</a></b>	<b><a href="#">Src Homology 2 Domain-Containing,</a></b>
New Heading	<b><a href="#">D12.644.360.024.330.875</a></b> <b><a href="#">Transforming Protein 3</a></b>	<b><a href="#">Src Homology 2 Domain-Containing,</a></b>
New Heading	<b><a href="#">D12.644.360.024.332</a></b> <b><a href="#">Associated Protein</a></b>	<b><a href="#">Signaling Lymphocytic Activation Molecule</a></b>
-	D12.644.360.024.334	Smad Proteins
-	D12.644.360.024.334.200	Smad Proteins, Inhibitory
-	D12.644.360.024.334.200.600	Smad6 Protein
-	D12.644.360.024.334.200.700	Smad7 Protein
-	D12.644.360.024.334.500	Smad Proteins, Receptor-Regulated
-	D12.644.360.024.334.500.100	Smad1 Protein
-	D12.644.360.024.334.500.200	Smad2 Protein
-	D12.644.360.024.334.500.300	Smad3 Protein
-	D12.644.360.024.334.500.500	Smad5 Protein
-	D12.644.360.024.334.500.800	Smad8 Protein
-	D12.644.360.024.334.750	Smad4 Protein
-	D12.644.360.024.342	STAT Transcription Factors
-	D12.644.360.024.342.100	STAT1 Transcription Factor
-	D12.644.360.024.342.200	STAT2 Transcription Factor
-	D12.644.360.024.342.300	STAT3 Transcription Factor
-	D12.644.360.024.342.400	STAT4 Transcription Factor

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D12.644.360.024.342.500	STAT5 Transcription Factor
-	D12.644.360.024.342.600	STAT6 Transcription Factor
-	D12.644.360.024.374	Suppressor of Cytokine Signaling Proteins
New Heading	<b>D12.644.360.024.374.500</b>	<b>Suppressor of Cytokine Signaling 1 Protein</b>
New Heading	<b>D12.644.360.024.374.750</b>	<b>Suppressor of Cytokine Signaling 3 Protein</b>
-	D12.644.360.024.437	Syntenins
-	D12.644.360.024.500 and Proteins	Tumor Necrosis Factor Receptor-Associated Peptides
-	D12.644.360.024.500.024 Protein	CASP8 and FADD-Like Apoptosis Regulating
-	D12.644.360.024.500.050	CRADD Signaling Adaptor Protein
-	D12.644.360.024.500.061	Edar-Associated Death Domain Protein
-	D12.644.360.024.500.124	Fas-Associated Death Domain Protein
-	D12.644.360.024.500.186 Kinases	Receptor-Interacting Protein Serine-Threonine
-	D12.644.360.024.500.186.500 Kinase 2	Receptor-Interacting Protein Serine-Threonine
-	D12.644.360.024.500.249	TNF Receptor-Associated Death Domain Protein
-	D12.644.360.024.500.500	TNF Receptor-Associated Factor 1
-	D12.644.360.024.500.750	TNF Receptor-Associated Factor 2
-	D12.644.360.024.500.875	TNF Receptor-Associated Factor 3
-	D12.644.360.024.500.906	TNF Receptor-Associated Factor 4
-	D12.644.360.024.500.937	TNF Receptor-Associated Factor 5
-	D12.644.360.024.500.968	TNF Receptor-Associated Factor 6
-	D12.644.360.050	Adenylyl Cyclases
-	D12.644.360.062	AMP-Activated Protein Kinases
-	D12.644.360.075	Apoptosis Regulatory Proteins
-	D12.644.360.075.311	Apoptosis Inducing Factor
New Heading	<b>D12.644.360.075.323</b>	<b>Bcl-2-Like Protein 11</b>
New Heading	<b>D12.644.360.075.335</b>	<b>Beclin-1</b>
-	D12.644.360.075.358	CARD Signaling Adaptor Proteins
-	D12.644.360.075.358.124	Apoptotic Protease-Activating Factor 1
New Tree	<b>D12.644.360.075.358.155</b>	<b>Caspase 9</b>
-	D12.644.360.075.358.186	CRADD Signaling Adaptor Protein

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D12.644.360.075.358.249	Nod1 Signaling Adaptor Protein
-	D12.644.360.075.358.500	Nod2 Signaling Adaptor Protein
-	D12.644.360.075.358.750 Kinase 2	Receptor-Interacting Protein Serine-Threonine
-	D12.644.360.075.405	Caspases
-	D12.644.360.075.405.350	Caspases, Effector
-	D12.644.360.075.405.350.300	Caspase 3
-	D12.644.360.075.405.350.600	Caspase 6
-	D12.644.360.075.405.350.700	Caspase 7
-	D12.644.360.075.405.350.900	Caspase 14
-	D12.644.360.075.405.550	Caspases, Initiator
-	D12.644.360.075.405.550.100	Caspase 1
-	D12.644.360.075.405.550.200	Caspase 2
-	D12.644.360.075.405.550.800	Caspase 8
-	D12.644.360.075.405.550.900	Caspase 9
-	D12.644.360.075.405.550.910	Caspase 10
-	D12.644.360.075.405.550.920	Caspase 12
-	D12.644.360.075.421	Death Domain Receptor Signaling Adaptor Proteins
-	D12.644.360.075.421.024 Protein	CASP8 and FADD-Like Apoptosis Regulating
New Tree	<a href="#">D12.644.360.075.421.037</a>	<a href="#">Caspase 8</a>
-	D12.644.360.075.421.050	CRADD Signaling Adaptor Protein
-	D12.644.360.075.421.100	Edar-Associated Death Domain Protein
-	D12.644.360.075.421.200	Fas-Associated Death Domain Protein
-	D12.644.360.075.421.400 Kinases	Receptor-Interacting Protein Serine-Threonine
-	D12.644.360.075.421.600	TNF Receptor-Associated Death Domain Protein
-	D12.644.360.075.437	Inhibitor of Apoptosis Proteins
-	D12.644.360.075.437.500	Neuronal Apoptosis-Inhibitory Protein
-	D12.644.360.075.437.750	X-Linked Inhibitor of Apoptosis Protein
-	D12.644.360.075.718	Proto-Oncogene Proteins c-bcl-2
-	D12.644.360.075.718.100	bcl-Associated Death Protein
-	D12.644.360.075.718.400	bcl-2-Associated X Protein
-	D12.644.360.075.718.750	bcl-2 Homologous Antagonist-Killer Protein
-	D12.644.360.075.718.937	bcl-X Protein
-	D12.644.360.075.718.968	BH3 Interacting Domain Death Agonist Protein

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.644.360.075.718.984 Myeloid Cell Leukemia Sequence 1 Protein
New Heading	<b>D12.644.360.081 Autophagy-Related Protein-1 Homolog</b>
-	D12.644.360.087 Calcineurin
-	D12.644.360.100 Calcium-Calmodulin-Dependent Protein Kinases
-	D12.644.360.100.049 Calcium-Calmodulin-Dependent Protein Kinase Kinase
-	D12.644.360.100.100 Calcium-Calmodulin-Dependent Protein Kinase Type 1
-	D12.644.360.100.200 Calcium-Calmodulin-Dependent Protein Kinase Type 2
-	D12.644.360.100.350 Calcium-Calmodulin-Dependent Protein Kinase Type 4
-	D12.644.360.100.387 Death-Associated Protein Kinases
-	D12.644.360.100.425 Elongation Factor 2 Kinase
-	D12.644.360.100.500 Myosin-Light-Chain Kinase
-	D12.644.360.175 Cold Shock Proteins and Peptides
-	D12.644.360.200 Cyclic Nucleotide-Regulated Protein Kinases
-	D12.644.360.200.125 Cyclic AMP-Dependent Protein Kinases
-	D12.644.360.200.125.750 Cyclic AMP-Dependent Protein Kinase Type I
-	D12.644.360.200.125.750.500 Cyclic AMP-Dependent Protein Kinase Catalytic Subunits
-	D12.644.360.200.125.750.625 Cyclic AMP-Dependent Protein Kinase RIalpha Subunit
-	D12.644.360.200.125.750.750 Cyclic AMP-Dependent Protein Kinase RIIbeta Subunit
-	D12.644.360.200.125.875 Cyclic AMP-Dependent Protein Kinase Type II
-	D12.644.360.200.125.875.500 Cyclic AMP-Dependent Protein Kinase Catalytic Subunits
-	D12.644.360.200.125.875.750 Cyclic AMP-Dependent Protein Kinase RIIIalpha Subunit
-	D12.644.360.200.125.875.875 Cyclic AMP-Dependent Protein Kinase RIIbeta Subunit
-	D12.644.360.200.150 Cyclic GMP-Dependent Protein Kinases
-	D12.644.360.200.150.500 Cyclic GMP-Dependent Protein Kinase Type I
-	D12.644.360.200.150.750 Cyclic GMP-Dependent Protein Kinase Type II
-	D12.644.360.200.575 Protamine Kinase
-	D12.644.360.225 Cyclin-Dependent Kinase Inhibitor Proteins
-	D12.644.360.225.100 Cyclin-Dependent Kinase Inhibitor p15
-	D12.644.360.225.200 Cyclin-Dependent Kinase Inhibitor p16
-	D12.644.360.225.300 Cyclin-Dependent Kinase Inhibitor p18
-	D12.644.360.225.400 Cyclin-Dependent Kinase Inhibitor p19



## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.644.360.225.500 Cyclin-Dependent Kinase Inhibitor p21
-	D12.644.360.225.600 Cyclin-Dependent Kinase Inhibitor p27
-	D12.644.360.225.700 Cyclin-Dependent Kinase Inhibitor p57
-	D12.644.360.250 Cyclin-Dependent Kinases
-	D12.644.360.250.067 CDC2-CDC28 Kinases
-	D12.644.360.250.067.249 CDC2 Protein Kinase
-	D12.644.360.250.067.500 CDC28 Protein Kinase, S cerevisiae
-	D12.644.360.250.067.875 Cyclin-Dependent Kinase 5
-	D12.644.360.250.067.900 Cyclin-Dependent Kinase 9
-	D12.644.360.250.323 Cyclin-Dependent Kinase 2
-	D12.644.360.250.387 Cyclin-Dependent Kinase 3
-	D12.644.360.250.451 Cyclin-Dependent Kinase 4
-	D12.644.360.250.515 Cyclin-Dependent Kinase 6
-	D12.644.360.250.547 Cyclin-Dependent Kinase 8
-	D12.644.360.250.580 Maturation-Promoting Factor
-	D12.644.360.250.580.500 CDC2 Protein Kinase
-	D12.644.360.262 Cyclins
-	D12.644.360.262.100 Cyclin A
-	D12.644.360.262.100.100 Cyclin A1
-	D12.644.360.262.100.200 Cyclin A2
-	D12.644.360.262.120 Cyclin B
-	D12.644.360.262.120.100 Cyclin B1
-	D12.644.360.262.120.200 Cyclin B2
-	D12.644.360.262.135 Cyclin C
-	D12.644.360.262.150 Cyclin D
-	D12.644.360.262.150.100 Cyclin D1
-	D12.644.360.262.150.200 Cyclin D2
-	D12.644.360.262.150.300 Cyclin D3
-	D12.644.360.262.180 Cyclin E
-	D12.644.360.262.200 Cyclin G
-	D12.644.360.262.200.100 Cyclin G1
-	D12.644.360.262.200.200 Cyclin G2
-	D12.644.360.262.300 Cyclin H
-	D12.644.360.262.400 Cyclin I
-	D12.644.360.262.700 Cyclin T
-	D12.644.360.268 Dual-Specificity Phosphatases

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.644.360.268.100 cdc25 Phosphatases
-	D12.644.360.268.200 Dual Specificity Phosphatase 1
-	D12.644.360.268.250 Dual Specificity Phosphatase 2
-	D12.644.360.268.300 Dual Specificity Phosphatase 3
-	D12.644.360.268.600 Dual Specificity Phosphatase 6
-	D12.644.360.275 eIF-2 Kinase
-	D12.644.360.287 Focal Adhesion Protein-Tyrosine Kinases
-	D12.644.360.293 G-Protein-Coupled Receptor Kinases
-	D12.644.360.293.249 beta-Adrenergic Receptor Kinases
-	D12.644.360.293.249.200 G-Protein-Coupled Receptor Kinase 2
-	D12.644.360.293.249.300 G-Protein-Coupled Receptor Kinase 3
-	D12.644.360.293.500 G-Protein-Coupled Receptor Kinase 1
-	D12.644.360.293.750 G-Protein-Coupled Receptor Kinase 4
-	D12.644.360.293.875 G-Protein-Coupled Receptor Kinase 5
-	D12.644.360.300 Glycogen Synthase Kinases
-	D12.644.360.300.500 Glycogen Synthase Kinase 3
New Heading	<b>D12.644.360.300.500.500 Glycogen Synthase Kinase 3 beta</b>
-	D12.644.360.325 GTP-Binding Protein Regulators
-	D12.644.360.325.150 GTPase-Activating Proteins
-	D12.644.360.325.150.100 Chimerin Proteins
-	D12.644.360.325.150.100.200 Chimerin 1
-	D12.644.360.325.150.300 Eukaryotic Initiation Factor-5
-	D12.644.360.325.150.500 ras GTPase-Activating Proteins
-	D12.644.360.325.150.500.460 Neurofibromin 1
-	D12.644.360.325.150.500.500 p120 GTPase Activating Protein
-	D12.644.360.325.150.750 RGS Proteins
-	D12.644.360.325.225 Guanine Nucleotide Dissociation Inhibitors
-	D12.644.360.325.225.500 rho-Specific Guanine Nucleotide Dissociation Inhibitors
-	D12.644.360.325.225.500.100 alpha rho Guanine Nucleotide Dissociation Inhibitor
-	D12.644.360.325.225.500.200 beta rho Guanine Nucleotide Dissociation Inhibitor
-	D12.644.360.325.225.500.300 gamma rho Guanine Nucleotide Dissociation Inhibitor
-	D12.644.360.325.300 Guanine Nucleotide Exchange Factors

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D12.644.360.325.300.050	Eukaryotic Initiation Factor-2B
-	D12.644.360.325.300.099	Rho Guanine Nucleotide Exchange Factors
New Tree	<a href="#">D12.644.360.325.300.099.500</a>	<a href="#">Proto-Oncogene Proteins c-bcr</a>
New Tree	<a href="#">D12.644.360.325.300.099.750</a>	<a href="#">Proto-Oncogene Proteins c-vav</a>
-	D12.644.360.325.300.300	Guanine Nucleotide-Releasing Factor 2
Old Tree	<del>D12.644.360.325.300.450</del>	<del>Proto-Oncogene Proteins c-vav</del>
-	D12.644.360.325.300.600	ral Guanine Nucleotide Exchange Factor
-	D12.644.360.325.300.700	ras Guanine Nucleotide Exchange Factors
-	D12.644.360.325.300.700.500	ras-GRF1
-	D12.644.360.325.300.700.700	Son of Sevenless Proteins
-	D12.644.360.325.300.700.700.600	Son of Sevenless Protein, Drosophila
-	D12.644.360.325.300.700.700.630	SOS1 Protein
-	D12.644.360.350	Guanylate Cyclase
New Heading	<del>D12.644.360.350.500</del>	<del>Soluble Guanylyl Cyclase</del>
-	D12.644.360.375	Heterotrimeric GTP-Binding Proteins
-	D12.644.360.375.100	GTP-Binding Protein alpha Subunits
-	D12.644.360.375.100.100	GTP-Binding Protein alpha Subunits, G12-G13
-	D12.644.360.375.100.200	GTP-Binding Protein alpha Subunits, Gi-Go
-	D12.644.360.375.100.200.500	GTP-Binding Protein alpha Subunit, Gi2
-	D12.644.360.375.100.300	GTP-Binding Protein alpha Subunits, Gq-G11
-	D12.644.360.375.100.400	GTP-Binding Protein alpha Subunits, Gs
-	D12.644.360.375.520	GTP-Binding Protein beta Subunits
-	D12.644.360.375.730	GTP-Binding Protein gamma Subunits
-	D12.644.360.375.940	Transducin
-	D12.644.360.376	I-kappa B Kinase
-	D12.644.360.378	I-kappa B Proteins
New Heading	<del>D12.644.360.378.500</del>	<del>NF-KappaB Inhibitor alpha</del>
-	D12.644.360.379	Interleukin-1 Receptor-Associated Kinases
-	D12.644.360.381	Intracellular Calcium-Sensing Proteins
-	D12.644.360.381.249	Calmodulin
-	D12.644.360.381.311	Calnexin
-	D12.644.360.381.374	Calreticulin
-	D12.644.360.381.437	Gelsolin

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.644.360.381.500      Neuronal Calcium-Sensor Proteins
-	D12.644.360.381.500.124      Guanylate Cyclase-Activating Proteins
-	D12.644.360.381.500.249      Hippocalcin
-	D12.644.360.381.500.374      Kv Channel-Interacting Proteins
-	D12.644.360.381.500.500      Neurocalcin
-	D12.644.360.381.500.750      Recoverin
-	D12.644.360.390      Lim Kinases
-	D12.644.360.400      MAP Kinase Kinase Kinases
-	D12.644.360.400.100      MAP Kinase Kinase Kinase 1
-	D12.644.360.400.200      MAP Kinase Kinase Kinase 2
-	D12.644.360.400.300      MAP Kinase Kinase Kinase 3
-	D12.644.360.400.400      MAP Kinase Kinase Kinase 4
-	D12.644.360.400.500      MAP Kinase Kinase Kinase 5
-	D12.644.360.400.800      Proto-Oncogene Proteins c-mos
-	D12.644.360.400.842      raf Kinases
-	D12.644.360.400.842.249      Oncogene Proteins v-raf
-	D12.644.360.400.842.374      Proto-Oncogene Proteins B-raf
-	D12.644.360.400.842.500      Proto-Oncogene Proteins c-raf
New Heading	<b>D12.644.360.420      Methyl-Accepting Chemotaxis Proteins</b>
-	D12.644.360.440      Mitogen-Activated Protein Kinase Kinases
-	D12.644.360.440.100      MAP Kinase Kinase 1
-	D12.644.360.440.200      MAP Kinase Kinase 2
-	D12.644.360.440.300      MAP Kinase Kinase 3
-	D12.644.360.440.400      MAP Kinase Kinase 4
-	D12.644.360.440.500      MAP Kinase Kinase 5
-	D12.644.360.440.600      MAP Kinase Kinase 6
-	D12.644.360.440.700      MAP Kinase Kinase 7
-	D12.644.360.445      Mitogen-Activated Protein Kinase Phosphatases
-	D12.644.360.445.100      Dual Specificity Phosphatase 1
-	D12.644.360.445.250      Dual Specificity Phosphatase 2
-	D12.644.360.445.300      Dual Specificity Phosphatase 3
-	D12.644.360.445.600      Dual Specificity Phosphatase 6
-	D12.644.360.450      Mitogen-Activated Protein Kinases
-	D12.644.360.450.169      Extracellular Signal-Regulated MAP Kinases
-	D12.644.360.450.169.500      Mitogen-Activated Protein Kinase 1

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.644.360.450.169.750 Mitogen-Activated Protein Kinase 3
-	D12.644.360.450.169.875 Mitogen-Activated Protein Kinase 6
-	D12.644.360.450.169.937 Mitogen-Activated Protein Kinase 7
-	D12.644.360.450.340 JNK Mitogen-Activated Protein Kinases
-	D12.644.360.450.340.500 Mitogen-Activated Protein Kinase 8
-	D12.644.360.450.340.750 Mitogen-Activated Protein Kinase 9
-	D12.644.360.450.340.800 Mitogen-Activated Protein Kinase 10
-	D12.644.360.450.835 p38 Mitogen-Activated Protein Kinases
-	D12.644.360.450.835.200 Mitogen-Activated Protein Kinase 11
-	D12.644.360.450.835.400 Mitogen-Activated Protein Kinase 12
-	D12.644.360.450.835.600 Mitogen-Activated Protein Kinase 13
-	D12.644.360.450.835.800 Mitogen-Activated Protein Kinase 14
-	D12.644.360.525 Monomeric GTP-Binding Proteins
-	D12.644.360.525.100 ADP-Ribosylation Factors
-	D12.644.360.525.100.100 ADP-Ribosylation Factor 1
-	D12.644.360.525.400 rab GTP-Binding Proteins
-	D12.644.360.525.400.025 rab1 GTP-Binding Proteins
-	D12.644.360.525.400.050 rab2 GTP-Binding Protein
-	D12.644.360.525.400.100 rab3 GTP-Binding Proteins
-	D12.644.360.525.400.100.100 rab3A GTP-Binding Protein
-	D12.644.360.525.400.150 rab4 GTP-Binding Proteins
-	D12.644.360.525.400.200 rab5 GTP-Binding Proteins
-	D12.644.360.525.450 ral GTP-Binding Proteins
-	D12.644.360.525.462 ran GTP-Binding Protein
-	D12.644.360.525.475 rap GTP-Binding Proteins
-	D12.644.360.525.475.100 rap1 GTP-Binding Proteins
-	D12.644.360.525.500 ras Proteins
-	D12.644.360.525.500.300 Oncogene Protein p21(ras)
-	D12.644.360.525.500.600 Proto-Oncogene Proteins p21(ras)
-	D12.644.360.525.700 rho GTP-Binding Proteins
-	D12.644.360.525.700.050 cdc42 GTP-Binding Protein
-	D12.644.360.525.700.050.500 cdc42 GTP-Binding Protein, <i>Saccharomyces cerevisiae</i>
-	D12.644.360.525.700.100 rac GTP-Binding Proteins
-	D12.644.360.525.700.100.100 rac1 GTP-Binding Protein
-	D12.644.360.525.700.200 rhoA GTP-Binding Protein

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D12.644.360.525.700.300	rhoB GTP-Binding Protein
-	D12.644.360.534	Mucin-4
New Heading	<b>D12.644.360.539</b>	<b>NLR Proteins</b>
New Heading	<b>D12.644.360.539.250</b>	<b>NLR Family, Pyrin Domain-Containing 3 Protein</b>
New Tree	<a href="#">D12.644.360.539.500</a>	<a href="#">Nod Signaling Adaptor Proteins</a>
New Tree	<a href="#">D12.644.360.539.500.249</a>	<a href="#">Nod1 Signaling Adaptor Protein</a>
New Tree	<a href="#">D12.644.360.539.500.500</a>	<a href="#">Nod2 Signaling Adaptor Protein</a>
-	D12.644.360.543	Olfactory Marker Protein
-	D12.644.360.552	p21-Activated Kinases
-	D12.644.360.562	Phosphatidylethanolamine Binding Protein
-	D12.644.360.571	Phosphoinositide Phospholipase C
-	D12.644.360.571.500	Phospholipase C beta
-	D12.644.360.571.625	Phospholipase C delta
-	D12.644.360.571.750	Phospholipase C gamma
-	D12.644.360.583	Protein Phosphatase 2
-	D12.644.360.585	Protein Tyrosine Phosphatases, Non-Receptor
-	D12.644.360.585.100	Protein Tyrosine Phosphatase, Non-Receptor Type 1
-	D12.644.360.585.200	Protein Tyrosine Phosphatase, Non-Receptor Type 2
-	D12.644.360.585.300	Protein Tyrosine Phosphatase, Non-Receptor Type 3
-	D12.644.360.585.400	Protein Tyrosine Phosphatase, Non-Receptor Type 4
-	D12.644.360.585.600	Protein Tyrosine Phosphatase, Non-Receptor Type 6
-	D12.644.360.585.800	Protein Tyrosine Phosphatase, Non-Receptor Type 11
-	D12.644.360.585.850	Protein Tyrosine Phosphatase, Non-Receptor Type 12
-	D12.644.360.585.860	Protein Tyrosine Phosphatase, Non-Receptor Type 13
-	D12.644.360.585.930	Protein Tyrosine Phosphatase, Non-Receptor Type 22
-	D12.644.360.587	Receptor-Like Protein Tyrosine Phosphatases
-	D12.644.360.587.100	Receptor-Like Protein Tyrosine Phosphatases, Class 1
-	D12.644.360.587.100.500	Antigens, CD45
-	D12.644.360.587.200	Receptor-Like Protein Tyrosine Phosphatases, Class 2
-	D12.644.360.587.300	Receptor-Like Protein Tyrosine Phosphatases, Class 3
-	D12.644.360.587.400	Receptor-Like Protein Tyrosine Phosphatases, Class 4
-	D12.644.360.587.500	Receptor-Like Protein Tyrosine Phosphatases, Class 5

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D12.644.360.587.700	Receptor-Like Protein Tyrosine Phosphatases, Class 7
-	D12.644.360.587.800	Receptor-Like Protein Tyrosine Phosphatases, Class 8
-	D12.644.360.590	rho-Associated Kinases
-	D12.644.360.600	Ribosomal Protein S6 Kinases
-	D12.644.360.600.249	Ribosomal Protein S6 Kinases, 70-kDa
-	D12.644.360.600.500	Ribosomal Protein S6 Kinases, 90-kDa
-	D12.644.360.800	SH2 Domain-Containing Protein Tyrosine Phosphatases
-	D12.644.360.800.200	Protein Tyrosine Phosphatase, Non-Receptor Type 6
New Heading	<b>D12.644.360.900</b>	<b>Syk Kinase</b>
New Heading	<b>D12.644.360.950</b>	<b>Tumor Necrosis Factor alpha-Induced Protein 3</b>
-	D12.644.365	Lipopeptides
-	D12.644.365.500	Daptomycin
New Heading	<b>D12.644.368</b>	<b>Mating Factor</b>
-	D12.644.370	Myocardial Depressant Factor
-	D12.644.400	Neuropeptides
-	D12.644.400.070	Angiotensins
-	D12.644.400.070.075	Angiotensin I
-	D12.644.400.070.078	Angiotensin II
-	D12.644.400.070.080	Angiotensin III
-	D12.644.400.085	Bombesin
-	D12.644.400.090	Bradykinin
-	D12.644.400.095	Calcitonin
-	D12.644.400.097	Calcitonin Gene-Related Peptide
-	D12.644.400.100	Carnosine
-	D12.644.400.200	Delta Sleep-Inducing Peptide
-	D12.644.400.235	FMRFamide
-	D12.644.400.250	Galanin
-	D12.644.400.275	Galanin-Like Peptide
-	D12.644.400.300	Gastric Inhibitory Polypeptide
-	D12.644.400.315	Gastrin-Releasing Peptide
-	D12.644.400.320	Gastrins
-	D12.644.400.360	Orexins
-	D12.644.400.400	Hypothalamic Hormones
-	D12.644.400.400.700	Pituitary Hormone Release Inhibiting Hormones

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.644.400.400.700.500 MSH Release-Inhibiting Hormone
-	D12.644.400.400.700.750 Prolactin Release-Inhibiting Factors
-	D12.644.400.400.700.875 Somatostatin
-	D12.644.400.400.700.875.500 Somatostatin-28
-	D12.644.400.400.740 Pituitary Hormone-Releasing Hormones
-	D12.644.400.400.740.140 Corticotropin-Releasing Hormone
-	D12.644.400.400.740.320 Gonadotropin-Releasing Hormone
-	D12.644.400.400.740.320.100 Buserelin
-	D12.644.400.400.740.320.340 Goserelin
-	D12.644.400.400.740.320.400 Leuprolide
-	D12.644.400.400.740.320.580 Nafarelin
-	D12.644.400.400.740.320.790 Triptorelin Pamoate
-	D12.644.400.400.740.860 Growth Hormone-Releasing Hormone
-	D12.644.400.400.740.860.780 Sermorelin
-	D12.644.400.400.740.880 Thyrotropin-Releasing Hormone
-	D12.644.400.400.935 Pro-Opiomelanocortin
-	D12.644.400.400.935.119 alpha-Endorphin
-	D12.644.400.400.935.179 alpha-MSH
-	D12.644.400.400.935.239 beta-Endorphin
-	D12.644.400.400.935.480 beta-Lipotropin
-	D12.644.400.400.935.492 beta-MSH
-	D12.644.400.400.935.498 Corticotropin-Like Intermediate Lobe Peptide
-	D12.644.400.400.935.505 gamma-Endorphin
-	D12.644.400.400.935.518 gamma-Lipotropin
-	D12.644.400.400.935.524 gamma-MSH
-	D12.644.400.400.935.531 Melanocortins
-	D12.644.400.400.935.531.500 Adrenocorticotrophic Hormone
-	D12.644.400.400.935.531.500.200 Cosyntropin
-	D12.644.400.400.935.531.750 Melanocyte-Stimulating Hormones
-	D12.644.400.400.935.531.750.050 alpha-MSH
-	D12.644.400.400.935.531.750.075 beta-MSH
-	D12.644.400.400.935.531.750.115 gamma-MSH
-	D12.644.400.400.967 Prolactin-Releasing Hormone
-	D12.644.400.430 Kisspeptins
-	D12.644.400.460 Melanocyte-Stimulating Hormones
-	D12.644.400.460.050 alpha-MSH



## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.644.400.460.075                      beta-MSH
-	D12.644.400.460.115                      gamma-MSH
-	D12.644.400.470                            Motilin
-	D12.644.400.500                            Neuropeptide Y
-	D12.644.400.525                            Neurophysins
-	D12.644.400.550                            Neurotensin
-	D12.644.400.562                            omega-Agatoxin IVA
-	D12.644.400.575                            Opioid Peptides
-	D12.644.400.575.180                        Dynorphins
-	D12.644.400.575.241                        Endorphins
-	D12.644.400.575.241.039                    alpha-Endorphin
-	D12.644.400.575.241.080                    beta-Endorphin
-	D12.644.400.575.241.540                    gamma-Endorphin
-	D12.644.400.575.281                        Enkephalins
-	D12.644.400.575.281.075                    Enkephalin, Ala(2)-MePhe(4)-Gly(5)-
-	D12.644.400.575.281.231                    Enkephalin, Leucine
-	D12.644.400.575.281.231.500                Enkephalin, Leucine-2-Alanine
-	D12.644.400.575.281.381                    Enkephalin, Methionine
-	D12.644.400.575.281.381.500                D-Ala(2),MePhe(4),Met(0)-ol-enkephalin
-	D12.644.400.575.281.600                    Enkephalin, D-Penicillamine (2,5)-
-	D12.644.400.600                            Pancreatic Polypeptide
-	D12.644.400.610                            Peptide PHI
-	D12.644.400.625                            Pituitary Adenylate Cyclase-Activating Polypeptide
-	D12.644.400.705                            Secretin
-	D12.644.400.800                            Tachykinins
-	D12.644.400.800.354                        Eledoisin
-	D12.644.400.800.475                        Kassinin
-	D12.644.400.800.500                        Neurokinin A
-	D12.644.400.800.550                        Neurokinin B
-	D12.644.400.800.625                        Physalaemin
-	D12.644.400.800.750                        Substance P
-	D12.644.400.837                            Urocortins
-	D12.644.400.875                            Vasoactive Intestinal Peptide
-	D12.644.400.900                            Vasopressins
-	D12.644.400.900.100                        Arginine Vasopressin
-	D12.644.400.900.100.250                    Deamino Arginine Vasopressin

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.644.400.900.400 Lypressin
-	D12.644.400.900.400.350 Felypressin
-	D12.644.400.900.700 Ornipressin
-	D12.644.456 Oligopeptides
-	D12.644.456.050 Amanitins
-	D12.644.456.050.111 Alpha-Amanitin
-	D12.644.456.073 Angiotensins
-	D12.644.456.073.021 Angiotensin I
-	D12.644.456.073.041 Angiotensin II
-	D12.644.456.073.041.050 Angiotensin Amide
-	D12.644.456.073.041.800 Saralasin
-	D12.644.456.073.041.815 1-Sarcosine-8-Isoleucine Angiotensin II
-	D12.644.456.073.055 Angiotensin III
-	D12.644.456.073.070 Angiotensinogen
-	D12.644.456.120 Antipain
-	D12.644.456.157 Atazanavir Sulfate
-	D12.644.456.193 Bradykinin
-	D12.644.456.193.400 Kallidin
-	D12.644.456.241 Ceruletide
-	D12.644.456.270 Chalcones
-	D12.644.456.300 Delta Sleep-Inducing Peptide
-	D12.644.456.345 Dipeptides
-	D12.644.456.345.159 Anserine
-	D12.644.456.345.190 Aspartame
-	D12.644.456.345.331 Carnosine
-	D12.644.456.345.360 Enalapril
-	D12.644.456.345.360.300 Enalaprilat
-	D12.644.456.345.575 Glycylglycine
-	D12.644.456.345.600 Lisinopril
-	D12.644.456.400 N-Formylmethionine Leucyl-Phenylalanine
-	D12.644.456.448 Glutathione
-	D12.644.456.448.500 Glutathione Disulfide
-	D12.644.456.448.625 Phytochelatins
-	D12.644.456.448.750 S-Nitrosoglutathione
-	D12.644.456.460 Gonadotropin-Releasing Hormone
-	D12.644.456.460.150 Buserelin

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.644.456.460.315 Goserelin
-	D12.644.456.460.480 Leuprolide
-	D12.644.456.460.600 Nafarelin
-	D12.644.456.460.800 Triptorelin Pamoate
-	D12.644.456.580 Leupeptins
-	D12.644.456.650 Netropsin
-	D12.644.456.716 Pentagastrin
-	D12.644.456.724 Pepstatins
-	D12.644.456.726 Peptichemio
-	D12.644.456.729 Peptide T
-	D12.644.456.735 Phalloidine
-	D12.644.456.800 Tachykinins
-	D12.644.456.800.354 Eledoisin
-	D12.644.456.800.475 Kassinin
-	D12.644.456.800.500 Neurokinin A
-	D12.644.456.800.550 Neurokinin B
-	D12.644.456.800.745 Physalaemin
-	D12.644.456.800.866 Substance P
-	D12.644.456.805 Technetium Tc 99m Mertiatide
-	D12.644.456.810 Teprotide
-	D12.644.456.830 Tetragastrin
-	D12.644.456.835 Thymic Factor, Circulating
-	D12.644.456.837 Thyrotropin-Releasing Hormone
-	D12.644.456.840 Tuftsin
-	D12.644.456.925 Vasopressins
-	D12.644.456.925.100 Arginine Vasopressin
-	D12.644.456.925.100.250 Deamino Arginine Vasopressin
-	D12.644.456.925.480 Lypressin
-	D12.644.456.925.480.500 Felypressin
-	D12.644.456.925.700 Ornipressin
-	D12.644.504 Peptaibols
-	D12.644.504.111 Alamethicin
-	D12.644.541 Peptide Fragments
-	D12.644.541.500 Immunoglobulin Fragments
-	D12.644.541.500.650 Immunoglobulin Fab Fragments
-	D12.644.541.500.650.250 Certolizumab Pegol

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.644.541.500.650.500 Immunoglobulin Variable Region
-	D12.644.541.500.650.500.180 Complementarity Determining Regions
-	D12.644.541.500.650.500.590 Immunoglobulin Joining Region
-	D12.644.541.500.650.500.800 Single-Chain Antibodies
-	D12.644.541.500.650.500.900 Single-Domain Antibodies
-	D12.644.541.500.650.750 Tuftsin
-	D12.644.541.500.697 Immunoglobulin Fc Fragments
-	D12.644.541.500.697.249 CD4 Immunoadhesins
-	D12.644.541.500.745 Immunoglobulin Idiotypes
-	D12.644.548 Peptide Hormones
-	D12.644.548.009 Activins
-	D12.644.548.009.500 Inhibin-beta Subunits
-	D12.644.548.011 Adipokines
-	D12.644.548.011.249 Adiponectin
-	D12.644.548.011.500 Leptin
-	D12.644.548.011.750 Resistin
-	D12.644.548.017 Adrenomedullin
-	D12.644.548.058 Angiotensins
-	D12.644.548.058.075 Angiotensin I
-	D12.644.548.058.078 Angiotensin II
-	D12.644.548.058.080 Angiotensin III
-	D12.644.548.100 Bombesin
-	D12.644.548.150 Calcitonin
-	D12.644.548.275 Gastric Inhibitory Polypeptide
-	D12.644.548.280 Gastrins
-	D12.644.548.322 Ghrelin
-	D12.644.548.365 Hypothalamic Hormones
-	D12.644.548.365.700 Pituitary Hormone Release Inhibiting Hormones
-	D12.644.548.365.700.500 MSH Release-Inhibiting Hormone
-	D12.644.548.365.700.750 Prolactin Release-Inhibiting Factors
-	D12.644.548.365.700.875 Somatostatin
-	D12.644.548.365.700.875.500 Somatostatin-28
-	D12.644.548.365.740 Pituitary Hormone-Releasing Hormones
-	D12.644.548.365.740.140 Corticotropin-Releasing Hormone
-	D12.644.548.365.740.320 Gonadotropin-Releasing Hormone
-	D12.644.548.365.740.320.100 Buserelin

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.644.548.365.740.320.340 Goserelin
-	D12.644.548.365.740.320.400 Leuprolide
-	D12.644.548.365.740.320.580 Nafarelin
-	D12.644.548.365.740.320.790 Triptorelin Pamoate
-	D12.644.548.365.740.860 Growth Hormone-Releasing Hormone
-	D12.644.548.365.740.860.780 Sermorelin
-	D12.644.548.365.740.880 Thyrotropin-Releasing Hormone
-	D12.644.548.365.935 Pro-Opiomelanocortin
-	D12.644.548.365.935.119 alpha-Endorphin
-	D12.644.548.365.935.179 alpha-MSH
-	D12.644.548.365.935.239 beta-Endorphin
-	D12.644.548.365.935.480 beta-Lipotropin
-	D12.644.548.365.935.492 beta-MSH
-	D12.644.548.365.935.498 Corticotropin-Like Intermediate Lobe Peptide
-	D12.644.548.365.935.505 gamma-Endorphin
-	D12.644.548.365.935.518 gamma-Lipotropin
-	D12.644.548.365.935.524 gamma-MSH
-	D12.644.548.365.935.531 Melanocortins
-	D12.644.548.365.935.531.500 Adrenocorticotrophic Hormone
-	D12.644.548.365.935.531.500.200 Cosyntropin
-	D12.644.548.365.935.531.750 Melanocyte-Stimulating Hormones
-	D12.644.548.365.935.531.750.050 alpha-MSH
-	D12.644.548.365.935.531.750.075 beta-MSH
-	D12.644.548.365.935.531.750.115 gamma-MSH
-	D12.644.548.365.967 Prolactin-Releasing Hormone
-	D12.644.548.387 Inhibins
-	D12.644.548.387.500 Inhibin-beta Subunits
-	D12.644.548.500 Motilin
-	D12.644.548.585 Natriuretic Peptides
-	D12.644.548.585.500 Atrial Natriuretic Factor
-	D12.644.548.585.625 Natriuretic Peptide, Brain
-	D12.644.548.585.750 Natriuretic Peptide, C-Type
-	D12.644.548.586 Pancreatic Hormones
-	D12.644.548.586.200 Insulins
-	D12.644.548.586.200.200 Biphasic Insulins
-	D12.644.548.586.200.300 Insulin, Long-Acting

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.644.548.586.200.300.050      Insulin Detemir
-	D12.644.548.586.200.300.100      Insulin Glargine
-	D12.644.548.586.200.300.200      Insulin, Isophane
-	D12.644.548.586.200.300.200.500      Isophane Insulin, Human
-	D12.644.548.586.200.300.300      Insulin, Lente
-	D12.644.548.586.200.300.800      Insulin, Ultralente
-	D12.644.548.586.200.400      Insulin, Short-Acting
-	D12.644.548.586.200.400.100      Insulin Aspart
-	D12.644.548.586.200.400.500      Insulin Lispro
-	D12.644.548.586.200.500      Proinsulin
-	D12.644.548.586.200.500.250      C-Peptide
-	D12.644.548.586.200.500.625      Insulin
-	D12.644.548.586.200.500.625.500      Insulin, Regular, Human
-	D12.644.548.586.200.500.625.500.500      Isophane Insulin, Human
-	D12.644.548.586.200.500.625.700      Insulin, Regular, Pork
-	D12.644.548.586.234      Islet Amyloid Polypeptide
-	D12.644.548.586.469      Proglucagon
-	D12.644.548.586.469.500      Glucagon
-	D12.644.548.586.700      Pancreatic Polypeptide
-	D12.644.548.586.780      Somatostatin
-	D12.644.548.586.780.500      Somatostatin-28
-	D12.644.548.587      Parathyroid Hormone
-	D12.644.548.587.850      Teriparatide
-	D12.644.548.588      Parathyroid Hormone-Related Protein
-	D12.644.548.592      Peptide PHI
-	D12.644.548.595      Peptide YY
-	D12.644.548.691      Pituitary Hormones
-	D12.644.548.691.525      Pituitary Hormones, Anterior
-	D12.644.548.691.525.343      Gonadotropins, Pituitary
-	D12.644.548.691.525.343.288      Follicle Stimulating Hormone
-	D12.644.548.691.525.343.288.500      Follicle Stimulating Hormone, beta Subunit
-	D12.644.548.691.525.343.288.750      Glycoprotein Hormones, alpha Subunit
-	D12.644.548.691.525.343.463      Luteinizing Hormone
-	D12.644.548.691.525.343.463.249      Glycoprotein Hormones, alpha Subunit
-	D12.644.548.691.525.343.463.500      Luteinizing Hormone, beta Subunit
-	D12.644.548.691.525.343.583      Menotropins

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.644.548.691.525.343.583.500 Urofollitropin
-	D12.644.548.691.525.425 Growth Hormone
-	D12.644.548.691.525.425.875 Human Growth Hormone
-	D12.644.548.691.525.525 Prolactin
-	D12.644.548.691.525.690 Pro-Opiomelanocortin
-	D12.644.548.691.525.690.119 alpha-Endorphin
-	D12.644.548.691.525.690.179 alpha-MSH
-	D12.644.548.691.525.690.239 beta-Endorphin
-	D12.644.548.691.525.690.480 beta-Lipotropin
-	D12.644.548.691.525.690.492 beta-MSH
-	D12.644.548.691.525.690.498 Corticotropin-Like Intermediate Lobe Peptide
-	D12.644.548.691.525.690.505 gamma-Endorphin
-	D12.644.548.691.525.690.518 gamma-Lipotropin
-	D12.644.548.691.525.690.524 gamma-MSH
-	D12.644.548.691.525.690.531 Melanocortins
-	D12.644.548.691.525.690.531.500 Adrenocorticotropin Hormone
-	D12.644.548.691.525.690.531.500.200 Cosyntropin
-	D12.644.548.691.525.690.531.750 Melanocyte-Stimulating Hormones
-	D12.644.548.691.525.690.531.750.050 alpha-MSH
-	D12.644.548.691.525.690.531.750.075 beta-MSH
-	D12.644.548.691.525.690.531.750.115 gamma-MSH
-	D12.644.548.691.525.883 Thyrotropin
-	D12.644.548.691.525.883.249 Glycoprotein Hormones, alpha Subunit
-	D12.644.548.691.525.883.500 Thyrotropin, beta Subunit
-	D12.644.548.691.692 Pituitary Hormones, Posterior
-	D12.644.548.691.692.433 Oxytocin
-	D12.644.548.691.692.781 Vasopressins
-	D12.644.548.691.692.781.100 Arginine Vasopressin
-	D12.644.548.691.692.781.100.250 Deamino Arginine Vasopressin
-	D12.644.548.691.692.781.400 Lypressin
-	D12.644.548.691.692.781.400.350 Felypressin
-	D12.644.548.691.692.781.700 Ornipressin
-	D12.644.548.691.692.881 Vasotocin
-	D12.644.548.726 Placental Hormones
-	D12.644.548.726.367 Chorionic Gonadotropin
-	D12.644.548.726.367.125 Chorionic Gonadotropin, beta Subunit, Human

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.644.548.726.367.562 Glycoprotein Hormones, alpha Subunit
-	D12.644.548.726.451 Gonadotropins, Equine
-	D12.644.548.726.692 Placental Lactogen
-	D12.644.548.762 Relaxin
-	D12.644.548.810 Secretin
-	D12.644.548.869 Thymosin
-	D12.644.548.887 Urocortins
-	D12.644.548.905 Urotensins
-	D12.644.548.952 Vasoactive Intestinal Peptide
-	D12.644.555 Peptide Library
-	D12.644.641 Peptides, Cyclic
-	D12.644.641.040 Alamethicin
-	D12.644.641.050 Amanitins
-	D12.644.641.050.111 Alpha-Amanitin
-	D12.644.641.075 Bacitracin
-	D12.644.641.142 Capreomycin
-	D12.644.641.235 Cyclosporins
-	D12.644.641.235.300 Cyclosporine
-	D12.644.641.243 Cyclotides
-	D12.644.641.252 Dactinomycin
-	D12.644.641.270 Daptomycin
-	D12.644.641.297 Depsipeptides
-	D12.644.641.297.500 Valinomycin
-	D12.644.641.311 Echinocandins
-	D12.644.641.325 Echinomycin
-	D12.644.641.352 Enterobactin
-	D12.644.641.380 Ferrichrome
-	D12.644.641.447 Microcystins
-	D12.644.641.515 Mycobacillin
-	D12.644.641.548 Nanotubes, Peptide
-	D12.644.641.582 Nisin
-	D12.644.641.650 Octreotide
-	D12.644.641.735 Phalloidine
-	D12.644.641.780 Polymyxins
-	D12.644.641.780.110 Colistin
-	D12.644.641.780.750 Polymyxin B



## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.644.641.802 Streptogramins
-	D12.644.641.802.249 Mikamycin
-	D12.644.641.802.374 Pristinamycin
-	D12.644.641.802.500 Streptogramin Group A
-	D12.644.641.802.500.500 Streptogramin A
-	D12.644.641.802.750 Streptogramin Group B
-	D12.644.641.802.750.500 Streptogramin B
-	D12.644.641.802.812 Vernamycin B
-	D12.644.641.802.875 Virginiamycin
-	D12.644.641.802.875.500 Streptogramin A
-	D12.644.641.825 Thiostrepton
-	D12.644.641.850 Tyrothricin
-	D12.644.641.850.300 Gramicidin
-	D12.644.641.850.920 Tyrocidine
-	D12.644.641.875 Viomycin
-	D12.644.641.875.250 Enviomycin
-	D12.644.679 Peptoids
-	D12.644.679.500 N-substituted Glycines
-	D12.644.717 Phosphopeptides
-	D12.644.740 Polygeline
-	D12.644.748 Polyglutamic Acid
-	D12.644.760 Polylysine
-	D12.644.770 Protein Sorting Signals
-	D12.644.770.600 Nuclear Export Signals
-	D12.644.770.610 Nuclear Localization Signals
-	D12.644.822 Proteinase Inhibitory Proteins, Secretory
-	D12.644.822.437 Elafin
-	D12.644.822.468 Protease Nexins
-	D12.644.822.468.750 Serpin E2
-	D12.644.822.500 Secretory Leukocyte Peptidase Inhibitor
-	D12.644.848 Salivary Proteins and Peptides
-	D12.644.848.249 Glue Proteins, Drosophila
-	D12.644.848.500 Histatins
-	D12.644.848.750 Mucin-5B
-	D12.644.848.875 Salivary alpha-Amylases
-	D12.644.848.906 Salivary Cystatins

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.644.848.937 Salivary Proline-Rich Proteins
-	D12.644.861 Serpins
-	D12.644.861.020 Angiotensinogen
-	D12.644.861.030 alpha 1-Antichymotrypsin
-	D12.644.861.035 alpha 1-Antitrypsin
-	D12.644.861.050 alpha-2-Antiplasmin
-	D12.644.861.060 Antithrombin Proteins
-	D12.644.861.060.500 Antithrombin III
-	D12.644.861.060.750 Heparin Cofactor II
-	D12.644.861.060.875 Hirudins
-	D12.644.861.140 Complement C1 Inactivator Proteins
-	D12.644.861.140.500 Complement C1 Inhibitor Protein
-	D12.644.861.418 HSP47 Heat-Shock Proteins
-	D12.644.861.557 Ovalbumin
-	D12.644.861.695 Plasminogen Inactivators
-	D12.644.861.695.500 Plasminogen Activator Inhibitor 1
-	D12.644.861.695.520 Plasminogen Activator Inhibitor 2
-	D12.644.861.695.700 Protein C Inhibitor
-	D12.644.861.695.850 Serpin E2
-	D12.644.861.847 Thyroxine-Binding Globulin
-	D12.644.861.923 Transcortin
-	D12.644.875 Tissue Polypeptide Antigen
New Heading	<b>D12.644.937 Trefoil Factors</b>
New Heading	<b>D12.644.937.500 Trefoil Factor-1</b>
New Heading	<b>D12.644.937.750 Trefoil Factor-2</b>
New Heading	<b>D12.644.937.875 Trefoil Factor-3</b>
-	D12.776 Proteins
-	D12.776.034 Albumins
-	D12.776.034.073 Albumin-Bound Paclitaxel
-	D12.776.034.145 C-Reactive Protein
-	D12.776.034.180 Conalbumin
-	D12.776.034.398 Lactalbumin
-	D12.776.034.614 Ovalbumin

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.034.614.300 Avidin
-	D12.776.034.700 Parvalbumins
-	D12.776.034.756 Ricin
-	D12.776.034.841 Serum Albumin
-	D12.776.034.841.350 Methemalbumin
-	D12.776.034.841.450 Prealbumin
-	D12.776.034.841.540 Serum Albumin, Bovine
-	D12.776.034.841.665 Serum Albumin, Radio-Iodinated
-	D12.776.034.900 Technetium Tc 99m Aggregated Albumin
-	D12.776.037 Algal Proteins
-	D12.776.045 Amphibian Proteins
-	D12.776.045.500 Xenopus Proteins
-	D12.776.049 Amyloid
-	D12.776.049.407 Amyloidogenic Proteins
-	D12.776.049.407.249 Amyloid beta-Protein Precursor
-	D12.776.049.407.249.500 Amyloid beta-Peptides
-	D12.776.049.407.500 Islet Amyloid Polypeptide
-	D12.776.049.407.750 Serum Amyloid A Protein
-	D12.776.049.407.875 Serum Amyloid P-Component
-	D12.776.053 Antifreeze Proteins
-	D12.776.053.100 Antifreeze Proteins, Type I
-	D12.776.053.200 Antifreeze Proteins, Type II
-	D12.776.053.350 Antifreeze Proteins, Type III
-	D12.776.053.500 Antifreeze Proteins, Type IV
-	D12.776.070 Apoproteins
-	D12.776.070.290 Apoenzymes
-	D12.776.070.345 Apoferritins
-	D12.776.070.400 Apolipoproteins
-	D12.776.070.400.200 Apolipoproteins A
-	D12.776.070.400.200.100 Apolipoprotein A-I
-	D12.776.070.400.200.150 Apolipoprotein A-II
New Heading	<b>D12.776.070.400.200.575 Apolipoprotein A-V</b>
-	D12.776.070.400.300 Apolipoproteins B
-	D12.776.070.400.300.240 Apolipoprotein B-48
-	D12.776.070.400.300.249 Apolipoprotein B-100

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.070.400.400 Apolipoproteins C
-	D12.776.070.400.400.500 Apolipoprotein C-I
-	D12.776.070.400.400.750 Apolipoprotein C-II
-	D12.776.070.400.400.875 Apolipoprotein C-III
-	D12.776.070.400.450 Apolipoproteins D
-	D12.776.070.400.500 Apolipoproteins E
-	D12.776.070.400.500.249 Apolipoprotein E2
-	D12.776.070.400.500.500 Apolipoprotein E3
-	D12.776.070.400.500.750 Apolipoprotein E4
-	D12.776.083 Aprotinin
-	D12.776.090 Archaeal Proteins
-	D12.776.090.200 Bacteriorhodopsins
-	D12.776.090.300 Halorhodopsins
-	D12.776.090.650 Periplasmic Proteins
-	D12.776.090.825 Thermosomes
-	D12.776.091 Armadillo Domain Proteins
-	D12.776.091.249 beta Catenin
-	D12.776.091.500 gamma Catenin
-	D12.776.091.750 Plakophilins
-	D12.776.093 Arthropod Proteins
-	D12.776.093.249 Agatoxins
-	D12.776.093.249.500 omega-Agatoxin IVA
-	D12.776.093.500 Insect Proteins
-	D12.776.093.500.462 Drosophila Proteins
-	D12.776.093.500.462.500 Glue Proteins, Drosophila
-	D12.776.093.500.577 Luciferases, Firefly
-	D12.776.093.500.925 Vitellogenins
-	D12.776.093.750 Silk
-	D12.776.093.750.500 Fibroins
-	D12.776.093.750.750 Sericins
New Heading	<b>D12.776.094 Autophagy-Related Proteins</b>
New Heading	<b>D12.776.094.250 Autophagy-Related Protein 5</b>
New Heading	<b>D12.776.094.391 Autophagy-Related Protein 12</b>
New	<b>D12.776.094.399 Autophagy-Related Protein 7</b>

## MeSH Tree Changes for 2017

Type	Tree - heading	
Heading		
New Heading	<b>D12.776.094.407</b>	<b>Autophagy-Related Protein 8 Family</b>
New Heading	<b>D12.776.094.438</b>	<b>Autophagy-Related Protein-1 Homolog</b>
New Heading	<b>D12.776.094.500</b>	<b>Beclin-1</b>
New Heading	<b>D12.776.094.750</b>	<b>Sequestosome-1 Protein</b>
-	D12.776.095	Avian Proteins
-	D12.776.097	Bacterial Proteins
-	D12.776.097.049	AraC Transcription Factor
-	D12.776.097.100	Azurin
-	D12.776.097.120	Bacterial Outer Membrane Proteins
-	D12.776.097.120.050	Adhesins, Bacterial
-	D12.776.097.120.050.040	Adhesins, Escherichia coli
-	D12.776.097.120.300	Bacterial Transferrin Receptor Complex
-	D12.776.097.120.300.500	Transferrin-Binding Protein A
-	D12.776.097.120.300.750	Transferrin-Binding Protein B
-	D12.776.097.120.425	Fimbriae Proteins
-	D12.776.097.130	Bacterial Proton-Translocating ATPases
-	D12.776.097.151	Bacteriocins
-	D12.776.097.151.190	Cloacin
-	D12.776.097.151.615	Megacins
-	D12.776.097.151.700	Nisin
New Heading	<b>D12.776.097.151.743</b>	<b>Pediocins</b>
-	D12.776.097.151.785	Pyocins
-	D12.776.097.156	Botulinum Toxins
-	D12.776.097.156.100	Botulinum Toxins, Type A
-	D12.776.097.162	Cell Wall Skeleton
-	D12.776.097.181	Coagulase
-	D12.776.097.200	Colicins
-	D12.776.097.237	DNA Gyrase
-	D12.776.097.256	DNA Topoisomerase IV
-	D12.776.097.275	Escherichia coli Proteins
-	D12.776.097.275.500	Adhesins, Escherichia coli

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.097.275.877 Shiga Toxin 1
-	D12.776.097.275.879 Shiga Toxin 2
-	D12.776.097.293 Exfoliatins
-	D12.776.097.312 Factor For Inversion Stimulation Protein
-	D12.776.097.350 Ferredoxins
-	D12.776.097.350.450 Molybdoferredoxin
-	D12.776.097.350.650 Rubredoxins
-	D12.776.097.380 Flagellin
-	D12.776.097.400 Flavodoxin
-	D12.776.097.513 Integration Host Factors
-	D12.776.097.513.500 Host Factor 1 Protein
-	D12.776.097.521 Lac Repressors
-	D12.776.097.529 Luciferases, Bacterial
New Heading	<b>D12.776.097.533 Methyl-Accepting Chemotaxis Proteins</b>
-	D12.776.097.537 MutS DNA Mismatch-Binding Protein
-	D12.776.097.541 NADP Transhydrogenase, B-Specific
-	D12.776.097.545 Penicillin-Binding Proteins
-	D12.776.097.577 Periplasmic Proteins
-	D12.776.097.577.500 Periplasmic Binding Proteins
-	D12.776.097.577.500.500 Maltose-Binding Proteins
-	D12.776.097.698 RNA Polymerase Sigma 54
-	D12.776.097.820 Staphylococcal Protein A
-	D12.776.097.835 Streptavidin
-	D12.776.097.872 Tetanus Toxin
New Heading	<b>D12.776.097.890 Transcription Activator-Like Effectors</b>
-	D12.776.124 Blood Proteins
-	D12.776.124.050 Acute-Phase Proteins
-	D12.776.124.050.050 alpha 1-Antichymotrypsin
-	D12.776.124.050.070 alpha 1-Antitrypsin
-	D12.776.124.050.080 alpha-Macroglobulins
-	D12.776.124.050.120 C-Reactive Protein
-	D12.776.124.050.130 Ceruloplasmin
-	D12.776.124.050.140 Complement C3
-	D12.776.124.050.250 Fibrinogen

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.124.050.250.265 Fibrinogens, Abnormal
-	D12.776.124.050.300 Haptoglobins
-	D12.776.124.050.350 Hemopexin
New Heading	<b>D12.776.124.050.475 Lipocalin-2</b>
-	D12.776.124.050.600 Orosomuroid
-	D12.776.124.050.725 Serum Amyloid A Protein
-	D12.776.124.050.730 Serum Amyloid P-Component
-	D12.776.124.050.800 Transferrin
-	D12.776.124.050.850 Trypsin Inhibitor, Kazal Pancreatic
-	D12.776.124.078 Anion Exchange Protein 1, Erythrocyte
-	D12.776.124.080 Ankyrins
-	D12.776.124.117 beta 2-Glycoprotein I
-	D12.776.124.125 Blood Coagulation Factors
-	D12.776.124.125.050 beta-Thromboglobulin
-	D12.776.124.125.300 Factor V
-	D12.776.124.125.300.300 Factor Va
-	D12.776.124.125.325 Factor VII
-	D12.776.124.125.325.300 Factor VIIa
-	D12.776.124.125.350 Factor VIII
-	D12.776.124.125.350.300 Factor VIIIa
-	D12.776.124.125.375 Factor IX
-	D12.776.124.125.375.310 Factor IXa
-	D12.776.124.125.400 Factor X
-	D12.776.124.125.400.315 Factor Xa
-	D12.776.124.125.425 Factor XI
-	D12.776.124.125.425.320 Factor XIa
-	D12.776.124.125.450 Factor XII
-	D12.776.124.125.450.324 Factor XIIa
-	D12.776.124.125.475 Factor XIII
-	D12.776.124.125.475.500 Factor XIIIa
-	D12.776.124.125.500 Fibrinogen
-	D12.776.124.125.500.265 Fibrinogens, Abnormal
-	D12.776.124.125.515 Fibrinopeptide A
-	D12.776.124.125.530 Fibrinopeptide B
-	D12.776.124.125.597 Kallikreins

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.124.125.597.712                      Prekallikrein
-	D12.776.124.125.635                              Kininogens
-	D12.776.124.125.635.480                      Kininogen, High-Molecular-Weight
-	D12.776.124.125.635.490                      Kininogen, Low-Molecular-Weight
-	D12.776.124.125.640                              Plasminogen Activator Inhibitor 1
-	D12.776.124.125.645                              Plasminogen Activator Inhibitor 2
-	D12.776.124.125.662                              Plasminogen Activators
-	D12.776.124.125.662.537                      Streptokinase
-	D12.776.124.125.662.537.075                      Anistreplase
-	D12.776.124.125.662.537.900                      Streptodornase and Streptokinase
-	D12.776.124.125.662.768                      Tissue Plasminogen Activator
-	D12.776.124.125.662.884                      Urokinase-Type Plasminogen Activator
-	D12.776.124.125.691                              Platelet Factor 3
-	D12.776.124.125.720                              Platelet Factor 4
-	D12.776.124.125.800                              Prothrombin
-	D12.776.124.125.890                              Thrombin
-	D12.776.124.125.900                              Thromboplastin
-	D12.776.124.125.920                              von Willebrand Factor
-	D12.776.124.197                                  Cholesterol Ester Transfer Proteins
-	D12.776.124.270                                  Fibrin
-	D12.776.124.270.300                              Fibrin Fibrinogen Degradation Products
-	D12.776.124.270.302                              Fibrin Foam
-	D12.776.124.270.305                              Fibrin Tissue Adhesive
-	D12.776.124.270.310                              Fibrinopeptide A
-	D12.776.124.270.320                              Fibrinopeptide B
-	D12.776.124.300                                  Glycophorin
-	D12.776.124.337                                  Hemocyanin
-	D12.776.124.400                                  Hemoglobins
-	D12.776.124.400.141                              Carboxyhemoglobin
-	D12.776.124.400.220                              Erythrocruorins
-	D12.776.124.400.303                              Fetal Hemoglobin
-	D12.776.124.400.405                              Hemoglobin A
-	D12.776.124.400.405.440                      Hemoglobin A, Glycosylated
-	D12.776.124.400.405.450                      Hemoglobin A2
-	D12.776.124.400.434                              Hemoglobin Subunits
-	D12.776.124.400.434.320                      alpha-Globins



## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.124.400.434.320.500 zeta-Globins
-	D12.776.124.400.434.325 beta-Globins
-	D12.776.124.400.434.325.249 delta-Globins
-	D12.776.124.400.434.325.374 epsilon-Globins
-	D12.776.124.400.434.325.500 gamma-Globins
-	D12.776.124.400.463 Hemoglobins, Abnormal
-	D12.776.124.400.463.338 Hemoglobin C
-	D12.776.124.400.463.375 Hemoglobin E
-	D12.776.124.400.463.463 Hemoglobin H
-	D12.776.124.400.463.480 Hemoglobin J
-	D12.776.124.400.463.510 Hemoglobin M
-	D12.776.124.400.463.588 Hemoglobin, Sickle
-	D12.776.124.400.599 Methemoglobin
-	D12.776.124.400.707 Oxyhemoglobins
-	D12.776.124.400.877 Sulfhemoglobin
-	D12.776.124.486 Immunoproteins
-	D12.776.124.486.157 C-Reactive Protein
-	D12.776.124.486.274 Complement System Proteins
-	D12.776.124.486.274.024 Anaphylatoxins
-	D12.776.124.486.274.024.250 Complement C3a
-	D12.776.124.486.274.024.260 Complement C4a
-	D12.776.124.486.274.024.270 Complement C5a
-	D12.776.124.486.274.024.270.255 Complement C5a, des-Arginine
-	D12.776.124.486.274.045 Complement Activating Enzymes
-	D12.776.124.486.274.045.280 Complement C1r
-	D12.776.124.486.274.045.290 Complement C1s
-	D12.776.124.486.274.045.387 Complement C3-C5 Convertases
-	D12.776.124.486.274.045.387.500 Complement C3-C5 Convertases, Alternative Pathway
-	D12.776.124.486.274.045.387.500.374 Complement C3 Convertase, Alternative Pathway
-	D12.776.124.486.274.045.387.500.750 Complement C5 Convertase, Alternative Pathway
-	D12.776.124.486.274.045.387.750 Complement C3-C5 Convertases, Classical Pathway
-	D12.776.124.486.274.045.387.750.500 Complement C3 Convertase, Classical Pathway

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.124.486.274.045.387.750.750 Complement C5 Convertase, Classical Pathway
-	D12.776.124.486.274.045.387.875 Mannose-Binding Protein-Associated Serine Proteases
-	D12.776.124.486.274.045.450 Complement Factor D
-	D12.776.124.486.274.050 Complement C1
-	D12.776.124.486.274.050.270 Complement C1q
-	D12.776.124.486.274.050.280 Complement C1r
-	D12.776.124.486.274.050.290 Complement C1s
-	D12.776.124.486.274.150 Complement C2
-	D12.776.124.486.274.150.500 Complement C2a
-	D12.776.124.486.274.150.750 Complement C2b
-	D12.776.124.486.274.250 Complement C3
-	D12.776.124.486.274.250.250 Complement C3a
-	D12.776.124.486.274.250.260 Complement C3b
-	D12.776.124.486.274.250.260.500 Complement C3c
-	D12.776.124.486.274.250.260.750 Complement C3d
-	D12.776.124.486.274.350 Complement C4
-	D12.776.124.486.274.350.250 Complement C4a
-	D12.776.124.486.274.350.260 Complement C4b
-	D12.776.124.486.274.450 Complement C5
-	D12.776.124.486.274.450.250 Complement C5a
-	D12.776.124.486.274.450.250.255 Complement C5a, des-Arginine
-	D12.776.124.486.274.450.625 Complement C5b
-	D12.776.124.486.274.550 Complement C6
-	D12.776.124.486.274.650 Complement C7
-	D12.776.124.486.274.750 Complement C8
-	D12.776.124.486.274.850 Complement C9
-	D12.776.124.486.274.900 Complement Factor B
-	D12.776.124.486.274.920 Complement Inactivator Proteins
-	D12.776.124.486.274.920.124 Antigens, CD46
-	D12.776.124.486.274.920.250 Complement C1 Inactivator Proteins
-	D12.776.124.486.274.920.250.500 Complement C1 Inhibitor Protein
-	D12.776.124.486.274.920.287 Complement C3 Nephritic Factor
-	D12.776.124.486.274.920.325 Complement C3b Inactivator Proteins
-	D12.776.124.486.274.920.325.200 Complement Factor H

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.124.486.274.920.325.210 Complement Factor I
-	D12.776.124.486.274.920.662 Complement C4b-Binding Protein
-	D12.776.124.486.274.930 Complement Membrane Attack Complex
-	D12.776.124.486.274.965 Properdin
-	D12.776.124.486.379 Eosinophil Granule Proteins
-	D12.776.124.486.379.249 Eosinophil Cationic Protein
-	D12.776.124.486.379.275 Eosinophil-Derived Neurotoxin
-	D12.776.124.486.379.500 Eosinophil Major Basic Protein
-	D12.776.124.486.379.600 Eosinophil Peroxidase
-	D12.776.124.486.485 Immunoglobulins
-	D12.776.124.486.485.114 Antibodies
-	D12.776.124.486.485.114.071 Antibodies, Anti-Idiotypic
-	D12.776.124.486.485.114.071.500 Omalizumab
-	D12.776.124.486.485.114.089 Antibodies, Archaeal
-	D12.776.124.486.485.114.107 Antibodies, Bacterial
-	D12.776.124.486.485.114.107.288 Antistreptolysin
-	D12.776.124.486.485.114.125 Antibodies, Bispecific
-	D12.776.124.486.485.114.143 Antibodies, Blocking
-	D12.776.124.486.485.114.167 Antibodies, Catalytic
-	D12.776.124.486.485.114.179 Antibodies, Fungal
-	D12.776.124.486.485.114.185 Antibodies, Helminth
-	D12.776.124.486.485.114.191 Antibodies, Heterophile
-	D12.776.124.486.485.114.207 Antibodies, Immobilized
-	D12.776.124.486.485.114.224 Antibodies, Monoclonal
-	D12.776.124.486.485.114.224.060 Antibodies, Monoclonal, Humanized
-	D12.776.124.486.485.114.224.060.250 Adalimumab
-	D12.776.124.486.485.114.224.060.375 Bevacizumab
-	D12.776.124.486.485.114.224.060.500 Certolizumab Pegol
-	D12.776.124.486.485.114.224.060.750 Cetuximab
-	D12.776.124.486.485.114.224.060.782 Denosumab
-	D12.776.124.486.485.114.224.060.813 Natalizumab
-	D12.776.124.486.485.114.224.060.844 Omalizumab
-	D12.776.124.486.485.114.224.060.860 Palivizumab
-	D12.776.124.486.485.114.224.060.868 Ranibizumab
-	D12.776.124.486.485.114.224.060.875 Trastuzumab
-	D12.776.124.486.485.114.224.060.937 Ustekinumab

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.124.486.485.114.224.075      Antibodies, Monoclonal, Murine-Derived
-	D12.776.124.486.485.114.224.075.570      Muromonab-CD3
-	D12.776.124.486.485.114.224.075.785      Rituximab
-	D12.776.124.486.485.114.224.608      Infliximab
-	D12.776.124.486.485.114.224.785      Single-Chain Antibodies
-	D12.776.124.486.485.114.240      Antibodies, Neoplasm
-	D12.776.124.486.485.114.244      Antibodies, Neutralizing
-	D12.776.124.486.485.114.248      Antibodies, Phospho-Specific
-	D12.776.124.486.485.114.252      Antibodies, Protozoan
-	D12.776.124.486.485.114.254      Antibodies, Viral
-	D12.776.124.486.485.114.254.150      Deltaretrovirus Antibodies
-	D12.776.124.486.485.114.254.150.440      HIV Antibodies
-	D12.776.124.486.485.114.254.150.500      HTLV-I Antibodies
-	D12.776.124.486.485.114.254.150.510      HTLV-II Antibodies
-	D12.776.124.486.485.114.254.450      Hepatitis Antibodies
-	D12.776.124.486.485.114.254.450.251      Hepatitis A Antibodies
-	D12.776.124.486.485.114.254.450.504      Hepatitis B Antibodies
-	D12.776.124.486.485.114.254.450.510      Hepatitis C Antibodies
-	D12.776.124.486.485.114.257      Antigen-Antibody Complex
-	D12.776.124.486.485.114.323      Autoantibodies
-	D12.776.124.486.485.114.323.190      Antibodies, Antineutrophil Cytoplasmic
-	D12.776.124.486.485.114.323.204      Antibodies, Antinuclear
-	D12.776.124.486.485.114.323.210      Antibodies, Antiphospholipid
-	D12.776.124.486.485.114.323.210.100      Antibodies, Anticardiolipin
-	D12.776.124.486.485.114.323.210.600      Lupus Coagulation Inhibitor
-	D12.776.124.486.485.114.323.300      Complement C3 Nephritic Factor
-	D12.776.124.486.485.114.323.390      Immunoconglutinins
-	D12.776.124.486.485.114.323.480      Immunoglobulins, Thyroid-Stimulating
-	D12.776.124.486.485.114.323.480.500      Long-Acting Thyroid Stimulator
-	D12.776.124.486.485.114.323.732      Rheumatoid Factor
-	D12.776.124.486.485.114.573      Immune Sera
-	D12.776.124.486.485.114.573.203      Antilymphocyte Serum
-	D12.776.124.486.485.114.573.601      Antitoxins
-	D12.776.124.486.485.114.573.601.138      Antivenins
-	D12.776.124.486.485.114.573.601.268      Botulinum Antitoxin
-	D12.776.124.486.485.114.573.601.438      Diphtheria Antitoxin

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.124.486.485.114.573.601.849 Tetanus Antitoxin
-	D12.776.124.486.485.114.580 Immunoconjugates
-	D12.776.124.486.485.114.580.225 Abatacept
-	D12.776.124.486.485.114.580.450 Immunotoxins
-	D12.776.124.486.485.114.606 Immunoglobulin Allotypes
-	D12.776.124.486.485.114.606.586 Immunoglobulin Gm Allotypes
-	D12.776.124.486.485.114.606.587 Immunoglobulin Km Allotypes
-	D12.776.124.486.485.114.619 Immunoglobulin Isotypes
-	D12.776.124.486.485.114.619.026 Immunoglobulin A
-	D12.776.124.486.485.114.619.026.030 Immunoglobulin A, Secretory
-	D12.776.124.486.485.114.619.026.030.500 Secretory Component
-	D12.776.124.486.485.114.619.026.515 Immunoglobulin alpha-Chains
-	D12.776.124.486.485.114.619.251 Immunoglobulin D
-	D12.776.124.486.485.114.619.251.500 Immunoglobulin delta-Chains
-	D12.776.124.486.485.114.619.312 Immunoglobulin E
-	D12.776.124.486.485.114.619.312.500 Immunoglobulin epsilon-Chains
-	D12.776.124.486.485.114.619.393 Immunoglobulin G
-	D12.776.124.486.485.114.619.393.261 Etanercept
-	D12.776.124.486.485.114.619.393.522 Immunoglobulin gamma-Chains
-	D12.776.124.486.485.114.619.393.522.400 Immunoglobulin Gm Allotypes
-	D12.776.124.486.485.114.619.393.536 Immunoglobulins, Intravenous
-	D12.776.124.486.485.114.619.393.550 Long-Acting Thyroid Stimulator
-	D12.776.124.486.485.114.619.393.570 Muromonab-CD3
-	D12.776.124.486.485.114.619.393.700 Rho(D) Immune Globulin
-	D12.776.124.486.485.114.619.574 Immunoglobulin M
-	D12.776.124.486.485.114.619.574.500 Immunoglobulin mu-Chains
-	D12.776.124.486.485.114.632 Immunoglobulins, Intravenous
-	D12.776.124.486.485.114.656 Insulin Antibodies
-	D12.776.124.486.485.114.664 Isoantibodies
-	D12.776.124.486.485.114.715 Oligoclonal Bands
-	D12.776.124.486.485.114.767 Opsonin Proteins
-	D12.776.124.486.485.114.820 Plantibodies
-	D12.776.124.486.485.114.837 Precipitins
-	D12.776.124.486.485.114.890 Reagins
-	D12.776.124.486.485.397 gamma-Globulins
-	D12.776.124.486.485.397.500 Tuftsin

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.124.486.485.538 Immunoglobulin Constant Regions
-	D12.776.124.486.485.538.500 Immunoglobulin Fc Fragments
-	D12.776.124.486.485.538.500.249 CD4 Immunoadhesins
-	D12.776.124.486.485.680 Immunoglobulin Fragments
-	D12.776.124.486.485.680.650 Immunoglobulin Fab Fragments
-	D12.776.124.486.485.680.650.250 Certolizumab Pegol
-	D12.776.124.486.485.680.650.500 Immunoglobulin Variable Region
-	D12.776.124.486.485.680.650.500.180 Complementarity Determining Regions
-	D12.776.124.486.485.680.650.500.590 Immunoglobulin Joining Region
-	D12.776.124.486.485.680.650.500.800 Single-Chain Antibodies
-	D12.776.124.486.485.680.650.500.900 Single-Domain Antibodies
-	D12.776.124.486.485.680.650.750 Tuftsin
-	D12.776.124.486.485.680.697 Immunoglobulin Fc Fragments
-	D12.776.124.486.485.680.697.249 CD4 Immunoadhesins
-	D12.776.124.486.485.680.745 Immunoglobulin Idiotypes
-	D12.776.124.486.485.705 Immunoglobulin Subunits
-	D12.776.124.486.485.705.500 Immunoglobulin Heavy Chains
-	D12.776.124.486.485.705.500.350 Immunoglobulin alpha-Chains
-	D12.776.124.486.485.705.500.360 Immunoglobulin delta-Chains
-	D12.776.124.486.485.705.500.370 Immunoglobulin epsilon-Chains
-	D12.776.124.486.485.705.500.380 Immunoglobulin gamma-Chains
-	D12.776.124.486.485.705.500.380.500 Immunoglobulin Gm Allotypes
-	D12.776.124.486.485.705.500.500 Immunoglobulin mu-Chains
-	D12.776.124.486.485.705.625 Immunoglobulin J-Chains
-	D12.776.124.486.485.705.750 Immunoglobulin Light Chains
-	D12.776.124.486.485.705.750.530 Immunoglobulin kappa-Chains
-	D12.776.124.486.485.705.750.530.500 Immunoglobulin Km Allotypes
-	D12.776.124.486.485.705.750.550 Immunoglobulin lambda-Chains
-	D12.776.124.486.485.705.750.775 Immunoglobulin Light Chains, Surrogate
-	D12.776.124.486.485.705.875 Secretory Component
-	D12.776.124.486.485.797 Immunoglobulin Variable Region
-	D12.776.124.486.485.797.180 Complementarity Determining Regions
-	D12.776.124.486.485.797.590 Immunoglobulin Joining Region
-	D12.776.124.486.485.900 Paraproteins
-	D12.776.124.486.485.900.120 Bence Jones Protein
-	D12.776.124.486.485.900.225 Cryoglobulins

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.124.486.485.900.500 Myeloma Proteins
-	D12.776.124.486.485.900.700 Pyroglobulins
-	D12.776.124.486.485.950 Receptors, Antigen, B-Cell
-	D12.776.124.486.485.950.500 Antigens, CD79
-	D12.776.124.486.485.950.750 Pre-B Cell Receptors
-	D12.776.124.486.485.950.750.500 Immunoglobulin Light Chains, Surrogate
-	D12.776.124.486.657 Opsonin Proteins
-	D12.776.124.625 Platelet-Derived Growth Factor
-	D12.776.124.625.650 Proto-Oncogene Proteins c-sis
-	D12.776.124.650 Protein C
-	D12.776.124.670 Protein S
-	D12.776.124.698 Retinol-Binding Proteins, Plasma
-	D12.776.124.727 Serum Albumin
-	D12.776.124.727.500 Methemalbumin
-	D12.776.124.727.750 Prealbumin
-	D12.776.124.727.875 Serum Albumin, Bovine
-	D12.776.124.727.937 Serum Albumin, Radio-Iodinated
-	D12.776.124.790 Serum Globulins
-	D12.776.124.790.106 Alpha-Globulins
-	D12.776.124.790.106.050 alpha 1-Antichymotrypsin
-	D12.776.124.790.106.085 alpha 1-Antitrypsin
-	D12.776.124.790.106.090 alpha-2-Antiplasmin
-	D12.776.124.790.106.092 alpha-Fetoproteins
-	D12.776.124.790.106.100 alpha-Macroglobulins
-	D12.776.124.790.106.125 Antithrombin III
-	D12.776.124.790.106.214 Ceruloplasmin
-	D12.776.124.790.106.304 Fetuins
-	D12.776.124.790.106.304.500 alpha-2-HS-Glycoprotein
-	D12.776.124.790.106.304.750 Fetuin-B
-	D12.776.124.790.106.394 Haptoglobins
-	D12.776.124.790.106.450 Heparin Cofactor II
-	D12.776.124.790.106.640 Orosomuroid
-	D12.776.124.790.106.740 Progesterone-Binding Globulin
-	D12.776.124.790.106.745 Retinol-Binding Proteins, Plasma
-	D12.776.124.790.106.901 Transcortin
-	D12.776.124.790.223 Beta-Globulins

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.124.790.223.100      beta 2-Microglobulin
-	D12.776.124.790.223.160      beta-Thromboglobulin
-	D12.776.124.790.223.200      Complement Factor H
-	D12.776.124.790.223.338      Hemopexin
-	D12.776.124.790.223.580      Plasminogen
-	D12.776.124.790.223.580.500      Angiostatins
-	D12.776.124.790.223.624      Properdin
-	D12.776.124.790.223.800      Sex Hormone-Binding Globulin
-	D12.776.124.790.223.839      Transferrin
-	D12.776.124.790.651      Immunoglobulins
-	D12.776.124.790.651.114      Antibodies
-	D12.776.124.790.651.114.071      Antibodies, Anti-Idiotypic
-	D12.776.124.790.651.114.071.500      Omalizumab
-	D12.776.124.790.651.114.107      Antibodies, Archaeal
-	D12.776.124.790.651.114.125      Antibodies, Bacterial
-	D12.776.124.790.651.114.125.288      Antistreptolysin
-	D12.776.124.790.651.114.134      Antibodies, Bispecific
-	D12.776.124.790.651.114.143      Antibodies, Blocking
-	D12.776.124.790.651.114.167      Antibodies, Catalytic
-	D12.776.124.790.651.114.179      Antibodies, Fungal
-	D12.776.124.790.651.114.185      Antibodies, Helminth
-	D12.776.124.790.651.114.191      Antibodies, Heterophile
-	D12.776.124.790.651.114.207      Antibodies, Immobilized
-	D12.776.124.790.651.114.224      Antibodies, Monoclonal
-	D12.776.124.790.651.114.224.060      Antibodies, Monoclonal, Humanized
-	D12.776.124.790.651.114.224.060.250      Adalimumab
-	D12.776.124.790.651.114.224.060.500      Certolizumab Pegol
-	D12.776.124.790.651.114.224.060.750      Cetuximab
-	D12.776.124.790.651.114.224.060.782      Denosumab
-	D12.776.124.790.651.114.224.060.813      Natalizumab
-	D12.776.124.790.651.114.224.060.844      Omalizumab
-	D12.776.124.790.651.114.224.060.860      Palivizumab
-	D12.776.124.790.651.114.224.060.868      Ranibizumab
-	D12.776.124.790.651.114.224.060.875      Trastuzumab
-	D12.776.124.790.651.114.224.060.937      Ustekinumab
-	D12.776.124.790.651.114.224.075      Antibodies, Monoclonal, Murine-Derived



## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.124.790.651.114.224.075.570 Muromonab-CD3
-	D12.776.124.790.651.114.224.075.785 Rituximab
-	D12.776.124.790.651.114.224.537 Infliximab
-	D12.776.124.790.651.114.240 Antibodies, Neoplasm
-	D12.776.124.790.651.114.244 Antibodies, Neutralizing
-	D12.776.124.790.651.114.248 Antibodies, Phospho-Specific
-	D12.776.124.790.651.114.252 Antibodies, Protozoan
-	D12.776.124.790.651.114.254 Antibodies, Viral
-	D12.776.124.790.651.114.254.150 Deltaretrovirus Antibodies
-	D12.776.124.790.651.114.254.150.440 HIV Antibodies
-	D12.776.124.790.651.114.254.150.500 HTLV-I Antibodies
-	D12.776.124.790.651.114.254.150.510 HTLV-II Antibodies
-	D12.776.124.790.651.114.254.450 Hepatitis Antibodies
-	D12.776.124.790.651.114.254.450.251 Hepatitis A Antibodies
-	D12.776.124.790.651.114.254.450.504 Hepatitis B Antibodies
-	D12.776.124.790.651.114.254.450.510 Hepatitis C Antibodies
-	D12.776.124.790.651.114.257 Antigen-Antibody Complex
-	D12.776.124.790.651.114.323 Autoantibodies
-	D12.776.124.790.651.114.323.190 Antibodies, Antineutrophil Cytoplasmic
-	D12.776.124.790.651.114.323.204 Antibodies, Antinuclear
-	D12.776.124.790.651.114.323.210 Antibodies, Antiphospholipid
-	D12.776.124.790.651.114.323.210.100 Antibodies, Anticardiolipin
-	D12.776.124.790.651.114.323.210.600 Lupus Coagulation Inhibitor
-	D12.776.124.790.651.114.323.300 Complement C3 Nephritic Factor
-	D12.776.124.790.651.114.323.390 Immunoconglutinins
-	D12.776.124.790.651.114.323.480 Immunoglobulins, Thyroid-Stimulating
-	D12.776.124.790.651.114.323.480.500 Long-Acting Thyroid Stimulator
-	D12.776.124.790.651.114.323.732 Rheumatoid Factor
-	D12.776.124.790.651.114.573 Immune Sera
-	D12.776.124.790.651.114.573.203 Antilymphocyte Serum
-	D12.776.124.790.651.114.573.601 Antitoxins
-	D12.776.124.790.651.114.573.601.138 Antivenins
-	D12.776.124.790.651.114.573.601.268 Botulinum Antitoxin
-	D12.776.124.790.651.114.573.601.438 Diphtheria Antitoxin
-	D12.776.124.790.651.114.573.601.849 Tetanus Antitoxin
-	D12.776.124.790.651.114.580 Immunoconjugates

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.124.790.651.114.580.225 Abatacept
-	D12.776.124.790.651.114.580.450 Immunotoxins
-	D12.776.124.790.651.114.606 Immunoglobulin Allotypes
-	D12.776.124.790.651.114.606.586 Immunoglobulin Gm Allotypes
-	D12.776.124.790.651.114.606.587 Immunoglobulin Km Allotypes
-	D12.776.124.790.651.114.619 Immunoglobulin Isotypes
-	D12.776.124.790.651.114.619.026 Immunoglobulin A
-	D12.776.124.790.651.114.619.026.030 Immunoglobulin A, Secretory
-	D12.776.124.790.651.114.619.026.030.500 Secretory Component
-	D12.776.124.790.651.114.619.026.515 Immunoglobulin alpha-Chains
-	D12.776.124.790.651.114.619.251 Immunoglobulin D
-	D12.776.124.790.651.114.619.251.500 Immunoglobulin delta-Chains
-	D12.776.124.790.651.114.619.312 Immunoglobulin E
-	D12.776.124.790.651.114.619.312.500 Immunoglobulin epsilon-Chains
-	D12.776.124.790.651.114.619.393 Immunoglobulin G
-	D12.776.124.790.651.114.619.393.261 Etanercept
-	D12.776.124.790.651.114.619.393.522 Immunoglobulin gamma-Chains
-	D12.776.124.790.651.114.619.393.522.400 Immunoglobulin Gm Allotypes
-	D12.776.124.790.651.114.619.393.550 Long-Acting Thyroid Stimulator
-	D12.776.124.790.651.114.619.393.570 Muromonab-CD3
-	D12.776.124.790.651.114.619.393.700 Rho(D) Immune Globulin
-	D12.776.124.790.651.114.619.574 Immunoglobulin M
-	D12.776.124.790.651.114.619.574.500 Immunoglobulin mu-Chains
-	D12.776.124.790.651.114.632 Immunoglobulins, Intravenous
-	D12.776.124.790.651.114.656 Insulin Antibodies
-	D12.776.124.790.651.114.664 Isoantibodies
-	D12.776.124.790.651.114.715 Oligoclonal Bands
-	D12.776.124.790.651.114.767 Opsonin Proteins
-	D12.776.124.790.651.114.820 Plantibodies
-	D12.776.124.790.651.114.837 Precipitins
-	D12.776.124.790.651.114.890 Reagins
-	D12.776.124.790.651.397 gamma-Globulins
-	D12.776.124.790.651.397.500 Tuftsin
-	D12.776.124.790.651.538 Immunoglobulin Constant Regions
-	D12.776.124.790.651.538.500 Immunoglobulin Fc Fragments
-	D12.776.124.790.651.538.500.249 CD4 Immunoadhesins

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.124.790.651.680 Immunoglobulin Fragments
-	D12.776.124.790.651.680.650 Immunoglobulin Fab Fragments
-	D12.776.124.790.651.680.650.250 Certolizumab Pegol
-	D12.776.124.790.651.680.650.500 Immunoglobulin Variable Region
-	D12.776.124.790.651.680.650.500.180 Complementarity Determining Regions
-	D12.776.124.790.651.680.650.500.590 Immunoglobulin Joining Region
-	D12.776.124.790.651.680.650.500.800 Single-Chain Antibodies
-	D12.776.124.790.651.680.650.500.900 Single-Domain Antibodies
-	D12.776.124.790.651.680.650.750 Tuftsin
-	D12.776.124.790.651.680.660 Immunoglobulin Fc Fragments
-	D12.776.124.790.651.680.660.249 CD4 Immunoadhesins
-	D12.776.124.790.651.680.745 Immunoglobulin Idiotypes
-	D12.776.124.790.651.705 Immunoglobulin Subunits
-	D12.776.124.790.651.705.500 Immunoglobulin Heavy Chains
-	D12.776.124.790.651.705.500.350 Immunoglobulin alpha-Chains
-	D12.776.124.790.651.705.500.360 Immunoglobulin delta-Chains
-	D12.776.124.790.651.705.500.370 Immunoglobulin epsilon-Chains
-	D12.776.124.790.651.705.500.380 Immunoglobulin gamma-Chains
-	D12.776.124.790.651.705.500.380.500 Immunoglobulin Gm Allotypes
-	D12.776.124.790.651.705.500.500 Immunoglobulin mu-Chains
-	D12.776.124.790.651.705.625 Immunoglobulin J-Chains
-	D12.776.124.790.651.705.750 Immunoglobulin Light Chains
-	D12.776.124.790.651.705.750.530 Immunoglobulin kappa-Chains
-	D12.776.124.790.651.705.750.530.500 Immunoglobulin Km Allotypes
-	D12.776.124.790.651.705.750.550 Immunoglobulin lambda-Chains
-	D12.776.124.790.651.705.750.775 Immunoglobulin Light Chains, Surrogate
-	D12.776.124.790.651.705.875 Secretory Component
-	D12.776.124.790.651.797 Immunoglobulin Variable Region
-	D12.776.124.790.651.797.180 Complementarity Determining Regions
-	D12.776.124.790.651.797.590 Immunoglobulin Joining Region
-	D12.776.124.790.651.900 Paraproteins
-	D12.776.124.790.651.900.120 Bence Jones Protein
-	D12.776.124.790.651.900.225 Cryoglobulins
-	D12.776.124.790.651.900.500 Myeloma Proteins
-	D12.776.124.790.651.900.700 Pyroglobulins
-	D12.776.124.790.651.950 Receptors, Antigen, B-Cell

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D12.776.124.790.651.950.500	Antigens, CD79
-	D12.776.124.790.651.950.750	Pre-B Cell Receptors
-	D12.776.124.790.651.950.750.500	Immunoglobulin Light Chains, Surrogate
-	D12.776.124.790.720	Macroglobulins
-	D12.776.124.790.720.100	alpha-Macroglobulins
-	D12.776.124.790.720.100.500	Pregnancy-Associated alpha 2-Macroglobulins
-	D12.776.124.790.900	Transcobalamins
-	D12.776.124.862	Somatomedins
-	D12.776.124.862.400	Insulin-Like Growth Factor I
-	D12.776.124.862.425	Insulin-Like Growth Factor II
-	D12.776.124.862.575	Nonsuppressible Insulin-Like Activity
-	D12.776.124.875	Spectrin
-	D12.776.124.885	Thyroxine-Binding Proteins
-	D12.776.124.885.500	Thyroxine-Binding Globulin
-	D12.776.124.900	Tumor Necrosis Factor-alpha
-	D12.776.124.920	Vitronectin
-	D12.776.157	Carrier Proteins
-	D12.776.157.050	Acyl Carrier Protein
-	D12.776.157.057	Adaptor Proteins, Signal Transducing
-	D12.776.157.057.001	5-Lipoxygenase-Activating Proteins
-	D12.776.157.057.002	14-3-3 Proteins
-	D12.776.157.057.003	A Kinase Anchor Proteins
Old Tree	D12.776.157.057.004	CARD Signaling Adaptor Proteins
Old Tree	D12.776.157.057.004.124	Apoptotic Protease-Activating Factor 1
Old Tree	D12.776.157.057.004.186	CRADD Signaling Adaptor Protein
Old Tree	D12.776.157.057.004.249	Nod1 Signaling Adaptor Protein
Old Tree	D12.776.157.057.004.500	Nod2 Signaling Adaptor Protein
Old Tree	D12.776.157.057.004.750 Kinase 2	Receptor-Interacting Protein Serine-Threonine
New Tree	D12.776.157.057.005	Arrestins
New Tree	D12.776.157.057.005.050	Arrestin
New Heading	D12.776.157.057.005.525	Beta-Arrestins
New Heading	D12.776.157.057.005.525.500	beta-Arrestin 1

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Heading	<b>D12.776.157.057.005.525.750</b>	<b>Beta-Arrestin 2</b>
New Tree	D12.776.157.057.006	CARD Signaling Adaptor Proteins
New Tree	D12.776.157.057.006.124	Apoptotic Protease-Activating Factor 1
New Tree	D12.776.157.057.006.155	Caspase 9
New Tree	D12.776.157.057.006.186	CRADD Signaling Adaptor Protein
New Tree	D12.776.157.057.006.249	Nod1 Signaling Adaptor Protein
New Tree	D12.776.157.057.006.500	Nod2 Signaling Adaptor Protein
New Tree	D12.776.157.057.006.750 Kinase 2	Receptor-Interacting Protein Serine-Threonine
-	D12.776.157.057.010	Caveolin 1
-	D12.776.157.057.012	Caveolin 2
-	D12.776.157.057.014	Cortactin
-	D12.776.157.057.022	Crk-Associated Substrate Protein
-	D12.776.157.057.024	Death Domain Receptor Signaling Adaptor Proteins
-	D12.776.157.057.024.024 Protein	CASP8 and FADD-Like Apoptosis Regulating
New Tree	D12.776.157.057.024.037	Caspase 8
-	D12.776.157.057.024.050	CRADD Signaling Adaptor Protein
-	D12.776.157.057.024.100	Edar-Associated Death Domain Protein
-	D12.776.157.057.024.200	Fas-Associated Death Domain Protein
-	D12.776.157.057.024.400 Kinases	Receptor-Interacting Protein Serine-Threonine
-	D12.776.157.057.024.600	TNF Receptor-Associated Death Domain Protein
New Heading	<b>D12.776.157.057.025</b>	<b>Dishevelled Proteins</b>
-	D12.776.157.057.026	GRB2 Adaptor Protein
-	D12.776.157.057.028	GRB7 Adaptor Protein
-	D12.776.157.057.029	GRB10 Adaptor Protein
-	D12.776.157.057.039	Insulin Receptor Substrate Proteins
-	D12.776.157.057.050	Interferon Regulatory Factors
-	D12.776.157.057.050.124	Interferon Regulatory Factor-1
-	D12.776.157.057.050.249	Interferon Regulatory Factor-2

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D12.776.157.057.050.374	Interferon Regulatory Factor-3
-	D12.776.157.057.050.437	Interferon Regulatory Factor-7
-	D12.776.157.057.050.500 Subunit	Interferon-Stimulated Gene Factor 3, gamma
-	D12.776.157.057.061	Interferon-Stimulated Gene Factor 3
-	D12.776.157.057.061.500	Interferon-Stimulated Gene Factor 3, alpha Subunit
-	D12.776.157.057.061.500.500	STAT1 Transcription Factor
-	D12.776.157.057.061.500.750	STAT2 Transcription Factor
-	D12.776.157.057.061.750 Subunit	Interferon-Stimulated Gene Factor 3, gamma
Old Tree	D12.776.157.057.062	Mediator Complex
Old Tree	D12.776.157.057.062.249	Cyclin C
Old Tree	D12.776.157.057.062.500	Cyclin-Dependent Kinase 8
Old Tree	D12.776.157.057.062.750	Mediator Complex Subunit 1
Old Tree	D12.776.157.057.064	Myeloid Differentiation Factor 88
New Heading	<b>D12.776.157.057.067</b>	<b>Kelch-Like ECH-Associated Protein 1</b>
Old Tree	D12.776.157.057.068	Nod Signaling Adaptor Proteins
Old Tree	D12.776.157.057.068.249	Nod1 Signaling Adaptor Protein
Old Tree	D12.776.157.057.068.500	Nod2 Signaling Adaptor Protein
New Tree	D12.776.157.057.072	Mediator Complex
New Tree	D12.776.157.057.072.249	Cyclin C
New Tree	D12.776.157.057.072.500	Cyclin-Dependent Kinase 8
New Tree	D12.776.157.057.072.750	Mediator Complex Subunit 1
New Tree	D12.776.157.057.074	Myeloid Differentiation Factor 88
New Tree	D12.776.157.057.078	Nod Signaling Adaptor Proteins
New Tree	D12.776.157.057.078.249	Nod1 Signaling Adaptor Protein
New Tree	D12.776.157.057.078.500	Nod2 Signaling Adaptor Protein
-	D12.776.157.057.080	Nuclear Receptor Coactivators
-	D12.776.157.057.080.049	Mediator Complex Subunit 1
-	D12.776.157.057.080.100	Nuclear Receptor Coactivator 1
-	D12.776.157.057.080.200	Nuclear Receptor Coactivator 2

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.157.057.080.300 Nuclear Receptor Coactivator 3
New Heading	<b>D12.776.157.057.080.650</b> <b>Peroxisome Proliferator-Activated Receptor</b> <b>Gamma Coactivator 1-alpha</b>
-	D12.776.157.057.092 Paxillin
-	D12.776.157.057.110 PII Nitrogen Regulatory Proteins
-	D12.776.157.057.124 Protein Inhibitors of Activated STAT
-	D12.776.157.057.139 Proto-Oncogene Proteins c-crk
-	D12.776.157.057.155 Proto-Oncogene Proteins c-vav
-	D12.776.157.057.158 Retinoblastoma Binding Proteins
-	D12.776.157.057.158.049 E2F1 Transcription Factor
-	D12.776.157.057.158.100 Retinoblastoma-Binding Protein 1
-	D12.776.157.057.158.200 Retinoblastoma-Binding Protein 2
-	D12.776.157.057.158.600 Retinoblastoma-Binding Protein 4
-	D12.776.157.057.158.700 Retinoblastoma-Binding Protein 7
New Heading	<b>D12.776.157.057.160</b> <b>Sequestosome-1 Protein</b>
-	D12.776.157.057.162 Shc Signaling Adaptor Proteins
New Heading	<b>D12.776.157.057.162.500</b> <b>Src Homology 2 Domain-Containing,</b> <b>Transforming Protein 1</b>
New Heading	<b>D12.776.157.057.162.750</b> <b>Src Homology 2 Domain-Containing,</b> <b>Transforming Protein 2</b>
New Heading	<b>D12.776.157.057.162.875</b> <b>Src Homology 2 Domain-Containing,</b> <b>Transforming Protein 3</b>
New Heading	<b>D12.776.157.057.166</b> <b>Signaling Lymphocytic Activation Molecule</b> <b>Associated Protein</b>
-	D12.776.157.057.170 Smad Proteins
-	D12.776.157.057.170.249 Smad Proteins, Inhibitory
-	D12.776.157.057.170.249.600 Smad6 Protein
-	D12.776.157.057.170.249.700 Smad7 Protein
-	D12.776.157.057.170.500 Smad Proteins, Receptor-Regulated
-	D12.776.157.057.170.500.100 Smad1 Protein
-	D12.776.157.057.170.500.200 Smad2 Protein
-	D12.776.157.057.170.500.300 Smad3 Protein
-	D12.776.157.057.170.500.500 Smad5 Protein
-	D12.776.157.057.170.500.800 Smad8 Protein
-	D12.776.157.057.170.750 Smad4 Protein
-	D12.776.157.057.186 STAT Transcription Factors
-	D12.776.157.057.186.100 STAT1 Transcription Factor

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D12.776.157.057.186.200	STAT2 Transcription Factor
-	D12.776.157.057.186.300	STAT3 Transcription Factor
-	D12.776.157.057.186.400	STAT4 Transcription Factor
-	D12.776.157.057.186.500	STAT5 Transcription Factor
-	D12.776.157.057.186.600	STAT6 Transcription Factor
-	D12.776.157.057.249	Suppressor of Cytokine Signaling Proteins
New Heading	<b>D12.776.157.057.249.500</b>	<b>Suppressor of Cytokine Signaling 1 Protein</b>
New Heading	<b>D12.776.157.057.249.750</b>	<b>Suppressor of Cytokine Signaling 3 Protein</b>
-	D12.776.157.057.374	Syntenins
-	D12.776.157.057.500 and Proteins	Tumor Necrosis Factor Receptor-Associated Peptides
-	D12.776.157.057.500.050	CRADD Signaling Adaptor Protein
-	D12.776.157.057.500.061	Edar-Associated Death Domain Protein
-	D12.776.157.057.500.124	Fas-Associated Death Domain Protein
-	D12.776.157.057.500.186 Kinases	Receptor-Interacting Protein Serine-Threonine
-	D12.776.157.057.500.186.500 Kinase 2	Receptor-Interacting Protein Serine-Threonine
-	D12.776.157.057.500.249	TNF Receptor-Associated Death Domain Protein
-	D12.776.157.057.500.500	TNF Receptor-Associated Factor 1
-	D12.776.157.057.500.750	TNF Receptor-Associated Factor 2
-	D12.776.157.057.500.875	TNF Receptor-Associated Factor 3
-	D12.776.157.057.500.906	TNF Receptor-Associated Factor 4
-	D12.776.157.057.500.937	TNF Receptor-Associated Factor 5
-	D12.776.157.057.500.968	TNF Receptor-Associated Factor 6
-	D12.776.157.065	Androgen-Binding Protein
-	D12.776.157.125	Calcium-Binding Proteins
-	D12.776.157.125.050	Annexins
-	D12.776.157.125.050.050	Annexin A1
-	D12.776.157.125.050.060	Annexin A2
-	D12.776.157.125.050.070	Annexin A3
-	D12.776.157.125.050.080	Annexin A4
-	D12.776.157.125.050.100	Annexin A5
-	D12.776.157.125.050.110	Annexin A6
-	D12.776.157.125.050.120	Annexin A7



## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D12.776.157.125.090	Calbindins
-	D12.776.157.125.090.124	Calbindin 1
-	D12.776.157.125.090.249	Calbindin 2
-	D12.776.157.125.090.500	S100 Calcium Binding Protein G
-	D12.776.157.125.155	Calsequestrin
-	D12.776.157.125.283	Fetuin
-	D12.776.157.125.283.500	alpha-2-HS-Glycoprotein
-	D12.776.157.125.283.750	Fetuin-B
-	D12.776.157.125.412	Intracellular Calcium-Sensing Proteins
-	D12.776.157.125.412.249	Calmodulin
-	D12.776.157.125.412.311	Calnexin
-	D12.776.157.125.412.374	Calreticulin
-	D12.776.157.125.412.437	Gelsolin
-	D12.776.157.125.412.500	Neuronal Calcium-Sensor Proteins
-	D12.776.157.125.412.500.124	Guanylate Cyclase-Activating Proteins
-	D12.776.157.125.412.500.249	Hippocalcin
-	D12.776.157.125.412.500.374	Kv Channel-Interacting Proteins
-	D12.776.157.125.412.500.500	Neurocalcin
-	D12.776.157.125.412.500.750	Recoverin
-	D12.776.157.125.475	Myosin Light Chains
-	D12.776.157.125.700	Osteocalcin
-	D12.776.157.125.715	Osteonectin
-	D12.776.157.125.750	S100 Proteins
-	D12.776.157.125.750.500	Leukocyte L1 Antigen Complex
-	D12.776.157.125.750.500.100	Calgranulin A
-	D12.776.157.125.750.500.200	Calgranulin B
-	D12.776.157.125.750.625	S100 Calcium Binding Protein beta Subunit
-	D12.776.157.125.750.750	S100 Calcium Binding Protein G
New Heading	<b>D12.776.157.125.750.813</b>	<b>S100 Calcium-Binding Protein A4</b>
-	D12.776.157.125.750.875	S100A12 Protein
-	D12.776.157.125.787	Secretagogins
New Heading	<b>D12.776.157.125.797</b>	<b>Serrate-Jagged Proteins</b>
New Heading	<b>D12.776.157.125.797.500</b>	<b>Jagged-1 Protein</b>
New	<b>D12.776.157.125.797.750</b>	<b>Jagged-2 Protein</b>

## MeSH Tree Changes for 2017

Type	Tree - heading	
Heading		
New Heading	<b>D12.776.157.125.806</b>	<b>Stromal Interaction Molecules</b>
New Heading	<b>D12.776.157.125.806.500</b>	<b>Stromal Interaction Molecule 1</b>
New Heading	<b>D12.776.157.125.806.750</b>	<b>Stromal Interaction Molecule 2</b>
-	D12.776.157.125.825	Synaptotagmins
-	D12.776.157.125.825.249	Synaptotagmin I
-	D12.776.157.125.825.500	Synaptotagmin II
-	D12.776.157.125.900	Troponin C
-	D12.776.157.142	Calmodulin-Binding Proteins
-	D12.776.157.142.500	Neurogranin
-	D12.776.157.160	Ceruloplasmin
-	D12.776.157.165	Cholesterol Ester Transfer Proteins
-	D12.776.157.169	F-Box Proteins
-	D12.776.157.169.500	beta-Transducin Repeat-Containing Proteins
-	D12.776.157.170	Fatty Acid-Binding Proteins
New Heading	<b>D12.776.157.170.250</b>	<b>Fatty Acid-Binding Protein 7</b>
-	D12.776.157.170.500	Myelin P2 Protein
-	D12.776.157.247	Follistatin
-	D12.776.157.286	Follistatin-Related Proteins
-	D12.776.157.325	GTP-Binding Proteins
-	D12.776.157.325.150	GTP Phosphohydrolase-Linked Elongation Factors
-	D12.776.157.325.150.101	Peptide Elongation Factor 1
-	D12.776.157.325.150.102	Peptide Elongation Factor 2
-	D12.776.157.325.150.200	Peptide Elongation Factor G
-	D12.776.157.325.150.700	Peptide Elongation Factor Tu
-	D12.776.157.325.332	Heterotrimeric GTP-Binding Proteins
-	D12.776.157.325.332.100	GTP-Binding Protein alpha Subunits
-	D12.776.157.325.332.100.100	GTP-Binding Protein alpha Subunits, G12-G13
-	D12.776.157.325.332.100.200	GTP-Binding Protein alpha Subunits, Gi-Go
-	D12.776.157.325.332.100.200.500	GTP-Binding Protein alpha Subunit, Gi2
-	D12.776.157.325.332.100.300	GTP-Binding Protein alpha Subunits, Gq-G11
-	D12.776.157.325.332.100.400	GTP-Binding Protein alpha Subunits, Gs
-	D12.776.157.325.332.520	GTP-Binding Protein beta Subunits

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.157.325.332.730 GTP-Binding Protein gamma Subunits
-	D12.776.157.325.332.940 Transducin
-	D12.776.157.325.515 Monomeric GTP-Binding Proteins
-	D12.776.157.325.515.100 ADP-Ribosylation Factors
-	D12.776.157.325.515.100.100 ADP-Ribosylation Factor 1
-	D12.776.157.325.515.400 rab GTP-Binding Proteins
-	D12.776.157.325.515.400.025 rab1 GTP-Binding Proteins
-	D12.776.157.325.515.400.050 rab2 GTP-Binding Protein
-	D12.776.157.325.515.400.100 rab3 GTP-Binding Proteins
-	D12.776.157.325.515.400.100.100 rab3A GTP-Binding Protein
-	D12.776.157.325.515.400.150 rab4 GTP-Binding Proteins
-	D12.776.157.325.515.400.200 rab5 GTP-Binding Proteins
-	D12.776.157.325.515.450 ral GTP-Binding Proteins
-	D12.776.157.325.515.462 ran GTP-Binding Protein
-	D12.776.157.325.515.475 rap GTP-Binding Proteins
-	D12.776.157.325.515.475.100 rap1 GTP-Binding Proteins
-	D12.776.157.325.515.500 ras Proteins
-	D12.776.157.325.515.500.300 Oncogene Protein p21(ras)
-	D12.776.157.325.515.500.600 Proto-Oncogene Proteins p21(ras)
-	D12.776.157.325.515.700 rho GTP-Binding Proteins
-	D12.776.157.325.515.700.050 cdc42 GTP-Binding Protein
-	D12.776.157.325.515.700.050.500 cdc42 GTP-Binding Protein, Saccharomyces cerevisiae
-	D12.776.157.325.515.700.100 rac GTP-Binding Proteins
-	D12.776.157.325.515.700.100.100 rac1 GTP-Binding Protein
-	D12.776.157.325.515.700.200 rhoA GTP-Binding Protein
-	D12.776.157.325.515.700.300 rhoB GTP-Binding Protein
-	D12.776.157.325.636 Myxovirus Resistance Proteins
-	D12.776.157.325.757 Septins
-	D12.776.157.420 Insulin-Like Growth Factor Binding Proteins
-	D12.776.157.420.250 Insulin-Like Growth Factor Binding Protein 1
-	D12.776.157.420.260 Insulin-Like Growth Factor Binding Protein 2
-	D12.776.157.420.270 Insulin-Like Growth Factor Binding Protein 3
-	D12.776.157.420.280 Insulin-Like Growth Factor Binding Protein 4
-	D12.776.157.420.290 Insulin-Like Growth Factor Binding Protein 5
-	D12.776.157.420.300 Insulin-Like Growth Factor Binding Protein 6

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D12.776.157.427	Iron-Binding Proteins
-	D12.776.157.427.249	Ferritins
-	D12.776.157.427.249.290	Apo ferritins
-	D12.776.157.427.374	Nonheme Iron Proteins
-	D12.776.157.427.374.187	Hemerythrin
-	D12.776.157.427.374.375	Iron-Sulfur Proteins
-	D12.776.157.427.374.375.025	Adrenodoxin
-	D12.776.157.427.374.375.150	Ferredoxin-Nitrite Reductase
-	D12.776.157.427.374.375.275	Ferredoxins
-	D12.776.157.427.374.375.275.450	Molybdoferredoxin
-	D12.776.157.427.374.375.275.725	Rubredoxins
-	D12.776.157.427.374.375.863	Electron Transport Complex I
-	D12.776.157.427.374.375.863.500	NADH Dehydrogenase
-	D12.776.157.427.374.375.909	Electron Transport Complex II
-	D12.776.157.427.374.375.909.500	Succinate Dehydrogenase
-	D12.776.157.427.374.375.954	Electron Transport Complex III
-	D12.776.157.427.374.375.957	Ferrochelatase
-	D12.776.157.427.374.375.960	Iron Regulatory Protein 1
-	D12.776.157.427.374.375.961	Iron Regulatory Protein 2
-	D12.776.157.427.374.375.977	Nitrate Reductase (NAD(P)H)
-	D12.776.157.427.374.375.988	Nitrate Reductase (NADPH)
-	D12.776.157.427.750	Transferrins
-	D12.776.157.427.750.124	Conalbumin
-	D12.776.157.427.750.249	Lactoferrin
-	D12.776.157.427.750.500	Transferrin
Old Tree	D12.776.157.458	Latent TGF-beta Binding Proteins
New Heading	D12.776.157.464	Lipid Droplet Associated Proteins
New Heading	D12.776.157.464.500	Perilipins
New Heading	D12.776.157.464.500.500	Perilipin-1
New Heading	D12.776.157.464.500.750	Perilipin-2
New Heading	D12.776.157.464.500.875	Perilipin-3
New Heading	D12.776.157.464.500.906	Perilipin-4

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Heading	<b>D12.776.157.464.500.937</b>	<b>Perilipin-5</b>
-	D12.776.157.469	Lipocalins
New Heading	<b>D12.776.157.469.050</b>	<b>Glycodelin</b>
-	D12.776.157.469.100	Lipocalin 1
New Heading	<b>D12.776.157.469.325</b>	<b>Lipocalin-2</b>
-	D12.776.157.469.550	Retinol-Binding Proteins, Plasma
-	D12.776.157.478	Lymphocyte Antigen 96
-	D12.776.157.530	Membrane Transport Proteins
-	D12.776.157.530.100	ATP-Binding Cassette Transporters
-	D12.776.157.530.100.151	ATP Binding Cassette Transporter 1
New Heading	<b>D12.776.157.530.100.228 G</b>	<b>ATP Binding Cassette Transporter, Sub-Family G</b>
New Heading	<b>D12.776.157.530.100.228.250 Family G, Member 1</b>	<b>ATP Binding Cassette Transporter, Sub-Family G, Member 1</b>
New Heading	<b>D12.776.157.530.100.228.500 Family G, Member 2</b>	<b>ATP Binding Cassette Transporter, Sub-Family G, Member 2</b>
New Heading	<b>D12.776.157.530.100.228.750 Family G, Member 5</b>	<b>ATP Binding Cassette Transporter, Sub-Family G, Member 5</b>
New Heading	<b>D12.776.157.530.100.228.875 Family G, Member 8</b>	<b>ATP Binding Cassette Transporter, Sub-Family G, Member 8</b>
-	D12.776.157.530.100.304	Multidrug Resistance-Associated Proteins
New Tree	<a href="#">D12.776.157.530.100.304.500 Regulator</a>	<a href="#">Cystic Fibrosis Transmembrane Conductance Regulator</a>
-	D12.776.157.530.100.652	P-Glycoproteins
New Heading	<b>D12.776.157.530.100.652.250</b>	<b>Antigen Peptide Transporter-1</b>
New Heading	<b>D12.776.157.530.100.652.375</b>	<b>Antigen Peptide Transporter-2</b>
-	D12.776.157.530.100.652.500	P-Glycoprotein
-	D12.776.157.530.200	Amino Acid Transport Systems
-	D12.776.157.530.200.249	Amino Acid Transport Systems, Acidic
-	D12.776.157.530.200.249.500	Amino Acid Transport System X-AG
-	D12.776.157.530.200.249.500.500 Proteins	Glutamate Plasma Membrane Transport Proteins
-	D12.776.157.530.200.249.500.500.500	Excitatory Amino Acid Transporter 1
-	D12.776.157.530.200.249.500.500.750	Excitatory Amino Acid Transporter 2
-	D12.776.157.530.200.249.500.500.875	Excitatory Amino Acid Transporter 3

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D12.776.157.530.200.249.500.500.937	Excitatory Amino Acid Transporter 4
-	D12.776.157.530.200.249.500.500.968	Excitatory Amino Acid Transporter 5
-	D12.776.157.530.200.374	Amino Acid Transport Systems, Basic
-	D12.776.157.530.200.374.600	Amino Acid Transport System y+
-	D12.776.157.530.200.374.600.200	Cationic Amino Acid Transporter 1
-	D12.776.157.530.200.374.600.300	Cationic Amino Acid Transporter 2
-	D12.776.157.530.200.374.750	Amino Acid Transport System y+L
-	D12.776.157.530.200.374.750.500	Antigens, CD98
-	D12.776.157.530.200.374.750.500.250	Antigens, CD98 Heavy Chain
-	D12.776.157.530.200.374.750.500.625	Antigens, CD98 Light Chains
-	D12.776.157.530.200.500	Amino Acid Transport Systems, Neutral
-	D12.776.157.530.200.500.100	Amino Acid Transport System A
-	D12.776.157.530.200.500.200	Amino Acid Transport System ASC
-	D12.776.157.530.200.500.500	Amino Acid Transport System L
-	D12.776.157.530.200.500.500.500	Antigens, CD98
-	D12.776.157.530.200.500.500.500.250	Antigens, CD98 Heavy Chain
-	D12.776.157.530.200.500.500.500.300	Antigens, CD98 Light Chains
-	D12.776.157.530.200.500.500.500.300.500 Transporter 1	Large Neutral Amino Acid-
-	D12.776.157.530.300	Fatty Acid Transport Proteins
-	D12.776.157.530.300.500	Antigens, CD36
-	D12.776.157.530.400	Ion Channels
-	D12.776.157.530.400.150	Calcium Channels
-	D12.776.157.530.400.150.400	Calcium Channels, L-Type
-	D12.776.157.530.400.150.585	Calcium Channels, N-Type
-	D12.776.157.530.400.150.585.792	Calcium Channels, P-Type
-	D12.776.157.530.400.150.585.826	Calcium Channels, Q-Type
-	D12.776.157.530.400.150.585.867	Calcium Channels, R-Type
-	D12.776.157.530.400.150.720	Calcium Channels, T-Type
New Heading	<b>D12.776.157.530.400.150.740 Channels</b>	<b>Calcium Release Activated Calcium</b>
New Heading	<b>D12.776.157.530.400.150.740.500</b>	<b>ORAI1 Protein</b>
New Heading	<b>D12.776.157.530.400.150.740.750</b>	<b>ORAI2 Protein</b>
-	D12.776.157.530.400.150.760	Inositol 1,4,5-Trisphosphate Receptors
-	D12.776.157.530.400.150.800	Ryanodine Receptor Calcium Release Channel

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.157.530.400.150.900 TRPP Cation Channels
-	D12.776.157.530.400.175 Chloride Channels
-	D12.776.157.530.400.175.125 Cystic Fibrosis Transmembrane Conductance Regulator
-	D12.776.157.530.400.175.562 Receptors, GABA-A
-	D12.776.157.530.400.175.781 Receptors, Glycine
-	D12.776.157.530.400.337 Cyclic Nucleotide-Gated Cation Channels
-	D12.776.157.530.400.368 Hyperpolarization-Activated Cyclic Nucleotide-Gated Channels
-	D12.776.157.530.400.400 Ligand-Gated Ion Channels
-	D12.776.157.530.400.400.100 Cysteine Loop Ligand-Gated Ion Channel Receptors
-	D12.776.157.530.400.400.100.100 Receptors, GABA-A
-	D12.776.157.530.400.400.100.200 Receptors, Glycine
-	D12.776.157.530.400.400.100.500 Receptors, Nicotinic
-	D12.776.157.530.400.400.100.500.500 alpha7 Nicotinic Acetylcholine Receptor
-	D12.776.157.530.400.400.100.700 Receptors, Serotonin, 5-HT3
-	D12.776.157.530.400.400.500 Receptors, Ionotropic Glutamate
-	D12.776.157.530.400.400.500.100 Receptors, AMPA
-	D12.776.157.530.400.400.500.200 Receptors, Kainic Acid
-	D12.776.157.530.400.400.500.500 Receptors, N-Methyl-D-Aspartate
-	D12.776.157.530.400.400.750 Receptors, Purinergic P2X
-	D12.776.157.530.400.400.750.100 Receptors, Purinergic P2X1
-	D12.776.157.530.400.400.750.200 Receptors, Purinergic P2X2
-	D12.776.157.530.400.400.750.300 Receptors, Purinergic P2X3
-	D12.776.157.530.400.400.750.400 Receptors, Purinergic P2X4
-	D12.776.157.530.400.400.750.500 Receptors, Purinergic P2X5
-	D12.776.157.530.400.400.750.700 Receptors, Purinergic P2X7
-	D12.776.157.530.400.500 Porins
-	D12.776.157.530.400.500.040 Aquaporins
-	D12.776.157.530.400.500.040.249 Aquaglyceroporins
-	D12.776.157.530.400.500.040.249.500 Aquaporin 3
-	D12.776.157.530.400.500.040.249.750 Aquaporin 6
-	D12.776.157.530.400.500.040.374 Aquaporin 1
-	D12.776.157.530.400.500.040.437 Aquaporin 2
-	D12.776.157.530.400.500.040.468 Aquaporin 4

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D12.776.157.530.400.500.040.484	Aquaporin 5
-	D12.776.157.530.400.500.520	Voltage-Dependent Anion Channels
-	D12.776.157.530.400.500.520.500	Voltage-Dependent Anion Channel 1
-	D12.776.157.530.400.500.520.750	Voltage-Dependent Anion Channel 2
-	D12.776.157.530.400.600	Potassium Channels
-	D12.776.157.530.400.600.150	Potassium Channels, Calcium-Activated
-	D12.776.157.530.400.600.150.249 Activated Potassium Channels	Intermediate-Conductance Calcium-
-	D12.776.157.530.400.600.150.500 Potassium Channels	Large-Conductance Calcium-Activated
-	D12.776.157.530.400.600.150.500.500 Potassium Channel alpha Subunits	Large-Conductance Calcium-Activated
-	D12.776.157.530.400.600.150.500.750 Potassium Channel beta Subunits	Large-Conductance Calcium-Activated
-	D12.776.157.530.400.600.150.750 Potassium Channels	Small-Conductance Calcium-Activated
-	D12.776.157.530.400.600.450	Potassium Channels, Inwardly Rectifying
-	D12.776.157.530.400.600.450.500 Potassium Channels	G Protein-Coupled Inwardly-Rectifying
-	D12.776.157.530.400.600.450.550	KATP Channels
-	D12.776.157.530.400.600.450.550.500	Sulfonylurea Receptors
-	D12.776.157.530.400.600.850	Potassium Channels, Tandem Pore Domain
-	D12.776.157.530.400.600.900	Potassium Channels, Voltage-Gated
-	D12.776.157.530.400.600.900.124	Delayed Rectifier Potassium Channels
-	D12.776.157.530.400.600.900.124.249	KCNQ Potassium Channels
-	D12.776.157.530.400.600.900.124.249.500	KCNQ1 Potassium Channel
-	D12.776.157.530.400.600.900.124.249.750	KCNQ2 Potassium Channel
-	D12.776.157.530.400.600.900.124.249.875	KCNQ3 Potassium Channel
-	D12.776.157.530.400.600.900.124.374	Kv1.5 Potassium Channel
-	D12.776.157.530.400.600.900.124.500	Shab Potassium Channels
-	D12.776.157.530.400.600.900.249	Ether-A-Go-Go Potassium Channels
New Heading	<b>D12.776.157.530.400.600.900.249.500</b>	<b>ERG1 Potassium Channel</b>
-	D12.776.157.530.400.600.900.500	Shaker Superfamily of Potassium Channels
-	D12.776.157.530.400.600.900.500.124	Kv1.1 Potassium Channel
-	D12.776.157.530.400.600.900.500.186	Kv1.2 Potassium Channel
-	D12.776.157.530.400.600.900.500.217	Kv1.3 Potassium Channel
-	D12.776.157.530.400.600.900.500.233	Kv1.4 Potassium Channel



## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.157.530.400.600.900.500.241 Kv1.5 Potassium Channel
-	D12.776.157.530.400.600.900.500.245 Kv1.6 Potassium Channel
-	D12.776.157.530.400.600.900.500.249 Shab Potassium Channels
-	D12.776.157.530.400.600.900.500.625 Shal Potassium Channels
-	D12.776.157.530.400.600.900.500.750 Shaw Potassium Channels
-	D12.776.157.530.400.875 Sodium Channels
-	D12.776.157.530.400.875.050 Acid Sensing Ion Channels
-	D12.776.157.530.400.875.100 Degenerin Sodium Channels
-	D12.776.157.530.400.875.200 Epithelial Sodium Channels
-	D12.776.157.530.400.875.750 Voltage-Gated Sodium Channels
-	D12.776.157.530.400.875.750.100 NAV1.1 Voltage-Gated Sodium Channel
-	D12.776.157.530.400.875.750.200 NAV1.2 Voltage-Gated Sodium Channel
-	D12.776.157.530.400.875.750.300 NAV1.3 Voltage-Gated Sodium Channel
-	D12.776.157.530.400.875.750.400 NAV1.4 Voltage-Gated Sodium Channel
-	D12.776.157.530.400.875.750.500 NAV1.5 Voltage-Gated Sodium Channel
-	D12.776.157.530.400.875.750.600 NAV1.6 Voltage-Gated Sodium Channel
-	D12.776.157.530.400.875.750.700 NAV1.7 Voltage-Gated Sodium Channel
-	D12.776.157.530.400.875.750.800 NAV1.8 Voltage-Gated Sodium Channel
-	D12.776.157.530.400.875.750.900 NAV1.9 Voltage-Gated Sodium Channel
-	D12.776.157.530.400.875.750.960 Voltage-Gated Sodium Channel beta Subunits
-	D12.776.157.530.400.875.750.960.100 Voltage-Gated Sodium Channel beta-1 Subunit
-	D12.776.157.530.400.875.750.960.200 Voltage-Gated Sodium Channel beta-2 Subunit
-	D12.776.157.530.400.875.750.960.300 Voltage-Gated Sodium Channel beta-3 Subunit
-	D12.776.157.530.400.875.750.960.650 Voltage-Gated Sodium Channel beta-4 Subunit
-	D12.776.157.530.400.901 Transient Receptor Potential Channels
-	D12.776.157.530.400.901.500 TRPC Cation Channels
-	D12.776.157.530.400.901.555 TRPM Cation Channels
-	D12.776.157.530.400.901.777 TRPP Cation Channels
-	D12.776.157.530.400.901.888 TRPV Cation Channels
-	D12.776.157.530.450 Ion Pumps
-	D12.776.157.530.450.074 Anion Transport Proteins
-	D12.776.157.530.450.074.249 Halorhodopsins

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D12.776.157.530.450.074.500	Organic Anion Transporters
-	D12.776.157.530.450.074.500.199	Dicarboxylic Acid Transporters
-	D12.776.157.530.450.074.500.299	Folic Acid Transporters
-	D12.776.157.530.450.074.500.299.500	Folate Receptors, GPI-Anchored
-	D12.776.157.530.450.074.500.299.500.500	Folate Receptor 1
-	D12.776.157.530.450.074.500.299.500.750	Folate Receptor 2
-	D12.776.157.530.450.074.500.299.625	Proton-Coupled Folate Transporter
-	D12.776.157.530.450.074.500.299.750	Reduced Folate Carrier Protein
-	D12.776.157.530.450.074.500.400	Monocarboxylic Acid Transporters
-	D12.776.157.530.450.074.500.500 Dependent	Organic Anion Transporters, ATP-
-	D12.776.157.530.450.074.500.500.500 Proteins	Multidrug Resistance-Associated
New Tree	<a href="#">D12.776.157.530.450.074.500.500.500.500.500</a> <a href="#">Conductance Regulator</a>	<a href="#">Cystic Fibrosis Transmembrane</a>
Old Tree	<del><a href="#">D12.776.157.530.450.074.500.500.750</a></del>	<del><a href="#">P-Glycoprotein</a></del>
-	D12.776.157.530.450.074.500.500.875	P-Glycoproteins
New Heading	<b><a href="#">D12.776.157.530.450.074.500.500.875.250</a></b>	<b><a href="#">Antigen Peptide Transporter-1</a></b>
New Heading	<b><a href="#">D12.776.157.530.450.074.500.500.875.375</a></b>	<b><a href="#">Antigen Peptide Transporter-2</a></b>
New Tree	<a href="#">D12.776.157.530.450.074.500.500.875.500</a>	<a href="#">P-Glycoprotein</a>
-	D12.776.157.530.450.074.500.687 Dependent	Organic Anion Transporters, Sodium-
-	D12.776.157.530.450.074.500.687.500 Transporters	Sodium-Coupled Vitamin C
-	D12.776.157.530.450.074.500.781 Independent	Organic Anion Transporters, Sodium-
-	D12.776.157.530.450.074.500.781.500	Organic Anion Transport Polypeptide C
-	D12.776.157.530.450.074.500.781.500 Transporter Family Member 1b1	Solute Carrier Organic Anion
-	D12.776.157.530.450.074.500.781.750	Organic Anion Transport Protein 1
-	D12.776.157.530.450.074.750	Phosphate Transport Proteins
-	D12.776.157.530.450.074.750.500	Proton-Phosphate Symporters
-	D12.776.157.530.450.074.750.750	Sodium-Phosphate Cotransporter Proteins
-	D12.776.157.530.450.074.750.750.500 Proteins, Type I	Sodium-Phosphate Cotransporter
-	D12.776.157.530.450.074.750.750.750 Proteins, Type II	Sodium-Phosphate Cotransporter

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.157.530.450.074.750.750.750.124 Proteins, Type IIa <span style="float: right;">Sodium-Phosphate Cotransporter</span>
-	D12.776.157.530.450.074.750.750.750.249 Proteins, Type IIb <span style="float: right;">Sodium-Phosphate Cotransporter</span>
-	D12.776.157.530.450.074.750.750.750.500 Proteins, Type IIc <span style="float: right;">Sodium-Phosphate Cotransporter</span>
-	D12.776.157.530.450.074.750.750.875 Proteins, Type III <span style="float: right;">Sodium-Phosphate Cotransporter</span>
-	D12.776.157.530.450.162 <span style="float: right;">Antiporters</span>
-	D12.776.157.530.450.162.193 <span style="float: right;">Chloride-Bicarbonate Antiporters</span>
-	D12.776.157.530.450.162.193.500 <span style="float: right;">Anion Exchange Protein 1, Erythrocyte</span>
-	D12.776.157.530.450.162.276 <span style="float: right;">Potassium-Hydrogen Antiporters</span>
-	D12.776.157.530.450.162.359 <span style="float: right;">Reduced Folate Carrier Protein</span>
-	D12.776.157.530.450.162.442 <span style="float: right;">Sodium-Calcium Exchanger</span>
-	D12.776.157.530.450.162.775 <span style="float: right;">Sodium-Hydrogen Antiporter</span>
-	D12.776.157.530.450.162.780 <span style="float: right;">Sodium-Potassium-Exchanging ATPase</span>
-	D12.776.157.530.450.162.887 <span style="float: right;">Vesicular Neurotransmitter Transport Proteins</span>
-	D12.776.157.530.450.162.887.500 Proteins <span style="float: right;">Vesicular Biogenic Amine Transport</span>
-	D12.776.157.530.450.162.887.500.249 Proteins <span style="float: right;">Vesicular Acetylcholine Transport</span>
-	D12.776.157.530.450.162.887.500.500 Proteins <span style="float: right;">Vesicular Monoamine Transport</span>
-	D12.776.157.530.450.162.887.625 <span style="float: right;">Vesicular Glutamate Transport Proteins</span>
-	D12.776.157.530.450.162.887.625.500 1 <span style="float: right;">Vesicular Glutamate Transport Protein</span>
-	D12.776.157.530.450.162.887.625.750 2 <span style="float: right;">Vesicular Glutamate Transport Protein</span>
-	D12.776.157.530.450.162.887.750 Proteins <span style="float: right;">Vesicular Inhibitory Amino Acid Transport</span>
-	D12.776.157.530.450.250 <span style="float: right;">Cation Transport Proteins</span>
-	D12.776.157.530.450.250.249 <span style="float: right;">Arsenite Transporting ATPases</span>
-	D12.776.157.530.450.250.500 <span style="float: right;">Calcium-Transporting ATPases</span>
-	D12.776.157.530.450.250.500.249 ATPases <span style="float: right;">Plasma Membrane Calcium-Transporting</span>
-	D12.776.157.530.450.250.500.500 Transporting ATPases <span style="float: right;">Sarcoplasmic Reticulum Calcium-</span>
-	D12.776.157.530.450.250.812 <span style="float: right;">Organic Cation Transport Proteins</span>
-	D12.776.157.530.450.250.812.500 <span style="float: right;">Organic Cation Transporter 1</span>
-	D12.776.157.530.450.250.875 <span style="float: right;">Proton Pumps</span>

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.157.530.450.250.875.249 Bacteriorhodopsins
-	D12.776.157.530.450.250.875.300 Electron Transport Complex I
-	D12.776.157.530.450.250.875.303 Electron Transport Complex III
-	D12.776.157.530.450.250.875.304 Electron Transport Complex IV
-	D12.776.157.530.450.250.875.487 Inorganic Pyrophosphatase
-	D12.776.157.530.450.250.875.492 Photosystem I Protein Complex
-	D12.776.157.530.450.250.875.500 Proton-Translocating ATPases
-	D12.776.157.530.450.250.875.500.249 ATPases Bacterial Proton-Translocating
-	D12.776.157.530.450.250.875.500.500 ATPases Chloroplast Proton-Translocating
-	D12.776.157.530.450.250.875.500.625 H(+)-K(+)-Exchanging ATPase
-	D12.776.157.530.450.250.875.500.750 ATPases Mitochondrial Proton-Translocating
-	D12.776.157.530.450.250.875.500.875 ATPases Vacuolar Proton-Translocating
-	D12.776.157.530.450.250.880 Sodium-Potassium-Exchanging ATPase
-	D12.776.157.530.450.437 SLC4A Proteins
-	D12.776.157.530.450.437.249 Chloride-Bicarbonate Antiporters
-	D12.776.157.530.450.437.249.500 Anion Exchange Protein 1, Erythrocyte
-	D12.776.157.530.450.437.500 Sodium-Bicarbonate Symporters
-	D12.776.157.530.450.625 Symporters
-	D12.776.157.530.450.625.124 Proteins Dopamine Plasma Membrane Transport
-	D12.776.157.530.450.625.139 GABA Plasma Membrane Transport Proteins
-	D12.776.157.530.450.625.147 Proteins Glutamate Plasma Membrane Transport
-	D12.776.157.530.450.625.147.500 Excitatory Amino Acid Transporter 1
-	D12.776.157.530.450.625.147.750 Excitatory Amino Acid Transporter 2
-	D12.776.157.530.450.625.147.875 Excitatory Amino Acid Transporter 3
-	D12.776.157.530.450.625.147.937 Excitatory Amino Acid Transporter 4
-	D12.776.157.530.450.625.147.968 Excitatory Amino Acid Transporter 5
-	D12.776.157.530.450.625.155 Glycine Plasma Membrane Transport Proteins
-	D12.776.157.530.450.625.186 Proteins Norepinephrine Plasma Membrane Transport
-	D12.776.157.530.450.625.217 Proton-Coupled Folate Transporter
-	D12.776.157.530.450.625.249 Proton-Phosphate Symporters
-	D12.776.157.530.450.625.311 Proteins Serotonin Plasma Membrane Transport

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.157.530.450.625.374 Sodium Chloride Symporters
-	D12.776.157.530.450.625.374.500 Solute Carrier Family 12, Member 3
-	D12.776.157.530.450.625.405 Sodium-Coupled Vitamin C Transporters
-	D12.776.157.530.450.625.437 Sodium-Glucose Transport Proteins
-	D12.776.157.530.450.625.437.500 Sodium-Glucose Transporter 1
-	D12.776.157.530.450.625.437.750 Sodium-Glucose Transporter 2
-	D12.776.157.530.450.625.500 Sodium-Bicarbonate Symporters
-	D12.776.157.530.450.625.625 Sodium-Phosphate Cotransporter Proteins
-	D12.776.157.530.450.625.625.500 Sodium-Phosphate Cotransporter Proteins, Type I
-	D12.776.157.530.450.625.625.750 Sodium-Phosphate Cotransporter Proteins, Type II
-	D12.776.157.530.450.625.625.750.124 Sodium-Phosphate Cotransporter Proteins, Type IIa
-	D12.776.157.530.450.625.625.750.249 Sodium-Phosphate Cotransporter Proteins, Type IIb
-	D12.776.157.530.450.625.625.750.500 Sodium-Phosphate Cotransporter Proteins, Type IIc
-	D12.776.157.530.450.625.625.875 Sodium-Phosphate Cotransporter Proteins, Type III
-	D12.776.157.530.450.625.750 Sodium-Potassium-Chloride Symporters
-	D12.776.157.530.450.625.750.500 Solute Carrier Family 12, Member 1
-	D12.776.157.530.450.625.750.625 Solute Carrier Family 12, Member 2
-	D12.776.157.530.450.625.750.875 Solute Carrier Family 12, Member 4
-	D12.776.157.530.500 Monosaccharide Transport Proteins
-	D12.776.157.530.500.500 Glucose Transport Proteins, Facilitative
-	D12.776.157.530.500.500.500 Glucose Transporter Type 1
-	D12.776.157.530.500.500.750 Glucose Transporter Type 2
-	D12.776.157.530.500.500.875 Glucose Transporter Type 3
-	D12.776.157.530.500.500.937 Glucose Transporter Type 4
-	D12.776.157.530.500.500.968 Glucose Transporter Type 5
-	D12.776.157.530.500.750 Sodium-Glucose Transport Proteins
-	D12.776.157.530.500.750.500 Sodium-Glucose Transporter 1
-	D12.776.157.530.500.750.750 Sodium-Glucose Transporter 2
-	D12.776.157.530.562 Neurotransmitter Transport Proteins
-	D12.776.157.530.562.374 Plasma Membrane Neurotransmitter Transport Proteins
-	D12.776.157.530.562.374.500 Catecholamine Plasma Membrane Transport Proteins

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.157.530.562.374.500.500 Dopamine Plasma Membrane Transport Proteins
-	D12.776.157.530.562.374.500.750 Norepinephrine Plasma Membrane Transport Proteins
-	D12.776.157.530.562.374.750 GABA Plasma Membrane Transport Proteins
-	D12.776.157.530.562.374.781 Glutamate Plasma Membrane Transport Proteins
-	D12.776.157.530.562.374.781.500 Excitatory Amino Acid Transporter 1
-	D12.776.157.530.562.374.781.750 Excitatory Amino Acid Transporter 2
-	D12.776.157.530.562.374.781.781 Excitatory Amino Acid Transporter 3
-	D12.776.157.530.562.374.781.812 Excitatory Amino Acid Transporter 4
-	D12.776.157.530.562.374.781.875 Excitatory Amino Acid Transporter 5
-	D12.776.157.530.562.374.812 Glycine Plasma Membrane Transport Proteins
-	D12.776.157.530.562.374.875 Serotonin Plasma Membrane Transport Proteins
-	D12.776.157.530.562.750 Vesicular Neurotransmitter Transport Proteins
-	D12.776.157.530.562.750.500 Vesicular Biogenic Amine Transport Proteins
-	D12.776.157.530.562.750.500.249 Vesicular Acetylcholine Transport Proteins
-	D12.776.157.530.562.750.500.500 Vesicular Monoamine Transport Proteins
-	D12.776.157.530.562.750.625 Vesicular Glutamate Transport Proteins
-	D12.776.157.530.562.750.625.500 Vesicular Glutamate Transport Protein 1
-	D12.776.157.530.562.750.625.750 Vesicular Glutamate Transport Protein 2
-	D12.776.157.530.562.750.750 Vesicular Inhibitory Amino Acid Transport Proteins
-	D12.776.157.530.625 Nucleobase, Nucleoside, Nucleotide, and Nucleic Acid Transport Proteins
-	D12.776.157.530.625.500 Nucleobase Transport Proteins
-	D12.776.157.530.625.750 Nucleoside Transport Proteins
-	D12.776.157.530.625.750.500 Equilibrative Nucleoside Transport Proteins
-	D12.776.157.530.625.750.500.500 Equilibrative Nucleoside Transporter 1
-	D12.776.157.530.625.750.500.750 Equilibrative-Nucleoside Transporter 2
-	D12.776.157.530.625.875 Nucleotide Transport Proteins
-	D12.776.157.530.625.875.500 Mitochondrial ADP, ATP Translocases
-	D12.776.157.530.625.875.500.100 Adenine Nucleotide Translocator 1
-	D12.776.157.530.625.875.500.200 Adenine Nucleotide Translocator 2
-	D12.776.157.530.625.875.500.300 Adenine Nucleotide Translocator 3
-	D12.776.157.530.750 Nucleocytoplasmic Transport Proteins
-	D12.776.157.530.750.100 Aryl Hydrocarbon Receptor Nuclear Translocator

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D12.776.157.530.750.200	Cellular Apoptosis Susceptibility Protein
-	D12.776.157.530.750.500	Karyopherins
-	D12.776.157.530.750.500.100	alpha Karyopherins
-	D12.776.157.530.750.500.249	beta Karyopherins
-	D12.776.157.530.750.625	Nuclear Pore Complex Proteins
-	D12.776.157.530.750.687	Period Circadian Proteins
-	D12.776.157.530.750.750	ran GTP-Binding Protein
New Heading	<b>D12.776.157.530.875</b>	<b>SEC Translocation Channels</b>
New Heading	<b>D12.776.157.530.937</b>	<b>Solute Carrier Proteins</b>
New Tree	D12.776.157.530.937.250	Amino Acid Transport System X-AG
New Tree	D12.776.157.530.937.250.500	Glutamate Plasma Membrane Transport Proteins
New Tree	D12.776.157.530.937.250.500.500	Excitatory Amino Acid Transporter 1
New Tree	D12.776.157.530.937.250.500.750	Excitatory Amino Acid Transporter 2
New Tree	D12.776.157.530.937.250.500.875	Excitatory Amino Acid Transporter 3
New Tree	D12.776.157.530.937.250.500.937	Excitatory Amino Acid Transporter 4
New Tree	D12.776.157.530.937.250.500.968	Excitatory Amino Acid Transporter 5
New Tree	D12.776.157.530.937.313	Amino Acid Transport System y+L
New Tree	D12.776.157.530.937.313.500	Antigens, CD98
New Tree	D12.776.157.530.937.313.500.250	Antigens, CD98 Heavy Chain
New Tree	D12.776.157.530.937.313.500.625	Antigens, CD98 Light Chains
New Tree	D12.776.157.530.937.313.500.625.500 1	Large Neutral Amino Acid-Transporter 1
New Tree	D12.776.157.530.937.375	Amino Acid Transport System y+
New Tree	D12.776.157.530.937.375.200	Cationic Amino Acid Transporter 1
New Tree	D12.776.157.530.937.375.300	Cationic Amino Acid Transporter 2
New	D12.776.157.530.937.500	Dopamine Plasma Membrane Transport Proteins

## MeSH Tree Changes for 2017

Type	Tree - heading	
Tree		
New Tree	<a href="#">D12.776.157.530.937.532</a>	<a href="#">Equilibrative Nucleoside Transport Proteins</a>
New Tree	<a href="#">D12.776.157.530.937.532.500</a>	<a href="#">Equilibrative Nucleoside Transporter 1</a>
New Tree	<a href="#">D12.776.157.530.937.532.750</a>	<a href="#">Equilibrative-Nucleoside Transporter 2</a>
New Tree	<a href="#">D12.776.157.530.937.563</a>	<a href="#">Glucose Transport Proteins, Facilitative</a>
New Tree	<a href="#">D12.776.157.530.937.563.500</a>	<a href="#">Glucose Transporter Type 1</a>
New Tree	<a href="#">D12.776.157.530.937.563.750</a>	<a href="#">Glucose Transporter Type 2</a>
New Tree	<a href="#">D12.776.157.530.937.563.875</a>	<a href="#">Glucose Transporter Type 3</a>
New Tree	<a href="#">D12.776.157.530.937.563.937</a>	<a href="#">Glucose Transporter Type 4</a>
New Tree	<a href="#">D12.776.157.530.937.563.968</a>	<a href="#">Glucose Transporter Type 5</a>
New Tree	<a href="#">D12.776.157.530.937.594</a>	<a href="#">Mitochondrial ADP, ATP Translocases</a>
New Tree	<a href="#">D12.776.157.530.937.594.100</a>	<a href="#">Adenine Nucleotide Translocator 1</a>
New Tree	<a href="#">D12.776.157.530.937.594.200</a>	<a href="#">Adenine Nucleotide Translocator 2</a>
New Tree	<a href="#">D12.776.157.530.937.594.300</a>	<a href="#">Adenine Nucleotide Translocator 3</a>
New Heading	<b><a href="#">D12.776.157.530.937.598</a></b>	<b><a href="#">Mitochondrial Uncoupling Proteins</a></b>
New Heading	<b><a href="#">D12.776.157.530.937.598.500</a></b>	<b><a href="#">Uncoupling Protein 1</a></b>
New Heading	<b><a href="#">D12.776.157.530.937.598.750</a></b>	<b><a href="#">Uncoupling Protein 2</a></b>
New Heading	<b><a href="#">D12.776.157.530.937.598.875</a></b>	<b><a href="#">Uncoupling Protein 3</a></b>
New Tree	<a href="#">D12.776.157.530.937.600</a>	<a href="#">Norepinephrine Plasma Membrane Transport Proteins</a>
New Tree	<a href="#">D12.776.157.530.937.602</a>	<a href="#">Organic Anion Transport Polypeptide C</a>
New Tree	<a href="#">D12.776.157.530.937.602</a>	<a href="#">Solute Carrier Organic Anion Transporter Family Member 1b1</a>
New Tree	<a href="#">D12.776.157.530.937.610</a>	<a href="#">Organic Anion Transport Protein 1</a>



## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	D12.776.157.530.937.614	Organic Cation Transporter 1
New Tree	D12.776.157.530.937.618	Proton-Coupled Folate Transporter
New Tree	D12.776.157.530.937.622	Reduced Folate Carrier Protein
New Tree	D12.776.157.530.937.624	Serotonin Plasma Membrane Transport Proteins
New Tree	D12.776.157.530.937.625	SLC4A Proteins
New Tree	D12.776.157.530.937.625.249	Chloride-Bicarbonate Antiporters
New Tree	D12.776.157.530.937.625.249.500	Anion Exchange Protein 1, Erythrocyte
New Tree	D12.776.157.530.937.625.500	Sodium-Bicarbonate Symporters
New Tree	D12.776.157.530.937.688	Sodium-Coupled Vitamin C Transporters
New Tree	D12.776.157.530.937.696	Sodium-Glucose Transporter 1
New Tree	D12.776.157.530.937.700	Sodium-Glucose Transporter 2
New Tree	D12.776.157.530.937.704	Sodium-Phosphate Cotransporter Proteins, Type II
New Tree	D12.776.157.530.937.704.124 Type IIa	Sodium-Phosphate Cotransporter Proteins, Type IIa
New Tree	D12.776.157.530.937.704.249 Type IIb	Sodium-Phosphate Cotransporter Proteins, Type IIb
New Tree	D12.776.157.530.937.704.500 Type IIc	Sodium-Phosphate Cotransporter Proteins, Type IIc
New Tree	D12.776.157.530.937.719	Sodium-Phosphate Cotransporter Proteins, Type III
New Tree	D12.776.157.530.937.750	Sodium-Potassium-Chloride Symporters
New Tree	D12.776.157.530.937.750.500	Solute Carrier Family 12, Member 1
New Tree	D12.776.157.530.937.750.625	Solute Carrier Family 12, Member 2
New Tree	D12.776.157.530.937.750.750	Solute Carrier Family 12, Member 3
New Tree	D12.776.157.530.937.750.875	Solute Carrier Family 12, Member 4
-	D12.776.157.597	Neurophysins

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.157.622 Periplasmic Binding Proteins
-	D12.776.157.648 Phosphate-Binding Proteins
-	D12.776.157.648.500 Phosphate Transport Proteins
-	D12.776.157.648.500.500 Proton-Phosphate Symporters
-	D12.776.157.648.500.750 Sodium-Phosphate Cotransporter Proteins
-	D12.776.157.648.500.750.500 Sodium-Phosphate Cotransporter Proteins, Type I
-	D12.776.157.648.500.750.750 Sodium-Phosphate Cotransporter Proteins, Type II
-	D12.776.157.648.500.750.750.124 Sodium-Phosphate Cotransporter Proteins, Type IIa
-	D12.776.157.648.500.750.750.249 Sodium-Phosphate Cotransporter Proteins, Type IIb
-	D12.776.157.648.500.750.750.500 Sodium-Phosphate Cotransporter Proteins, Type IIc
-	D12.776.157.648.500.750.875 Sodium-Phosphate Cotransporter Proteins, Type III
-	D12.776.157.661 Phosphatidylethanolamine Binding Protein
-	D12.776.157.674 Phospholipid Transfer Proteins
-	D12.776.157.700 Retinol-Binding Proteins
-	D12.776.157.700.249 Retinol-Binding Proteins, Cellular
-	D12.776.157.700.500 Retinol-Binding Proteins, Plasma
-	D12.776.157.725 RNA-Binding Proteins
-	D12.776.157.725.030 Butyrate Response Factor 1
-	D12.776.157.725.061 Fragile X Mental Retardation Protein
-	D12.776.157.725.076 Gene Products, rev
-	D12.776.157.725.076.500 rev Gene Products, Human Immunodeficiency Virus
-	D12.776.157.725.124 mRNA Cleavage and Polyadenylation Factors
-	D12.776.157.725.124.240 Cleavage And Polyadenylation Specificity Factor
-	D12.776.157.725.124.249 Cleavage Stimulation Factor
-	D12.776.157.725.249 Host Factor 1 Protein
-	D12.776.157.725.374 Iron Regulatory Protein 1
-	D12.776.157.725.437 Iron Regulatory Protein 2
-	D12.776.157.725.444 Nuclear Factor 90 Proteins
-	D12.776.157.725.444.500 Nuclear Factor 45 Protein
-	D12.776.157.725.452 Poly(A)-Binding Proteins
-	D12.776.157.725.452.125 Ataxin-2
-	D12.776.157.725.452.249 Poly(A)-Binding Protein I

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.157.725.452.500 Poly(A)-Binding Protein II
Old Tree	<b>D12.776.157.725.468 Polypyrimidine Tract-Binding Protein</b>
-	D12.776.157.725.500 Ribonucleoproteins
-	D12.776.157.725.500.625 Ribonuclease P
-	D12.776.157.725.500.750 Ribonucleoproteins, Small Cytoplasmic
-	D12.776.157.725.500.750.800 Signal Recognition Particle
-	D12.776.157.725.500.875 Ribonucleoproteins, Small Nuclear
-	D12.776.157.725.500.875.590 Ribonucleoproteins, Small Nucleolar
-	D12.776.157.725.500.875.600 Ribonucleoprotein, U1 Small Nuclear
-	D12.776.157.725.500.875.605 Ribonucleoprotein, U2 Small Nuclear
-	D12.776.157.725.500.875.615 Ribonucleoprotein, U4-U6 Small Nuclear
-	D12.776.157.725.500.875.620 Ribonucleoprotein, U5 Small Nuclear
-	D12.776.157.725.500.875.625 Ribonucleoprotein, U7 Small Nuclear
-	D12.776.157.725.500.875.700 snRNP Core Proteins
-	D12.776.157.725.500.906 RNA-Induced Silencing Complex
-	D12.776.157.725.500.906.500 Argonaute Proteins
-	D12.776.157.725.500.921 Telomerase
-	D12.776.157.725.500.937 Vault Ribonucleoprotein Particles
-	D12.776.157.725.750 RNA Cap-Binding Proteins
-	D12.776.157.725.750.374 Eukaryotic Initiation Factor-4E
-	D12.776.157.725.750.750 Nuclear Cap-Binding Protein Complex
-	D12.776.157.725.813 RNA Recognition Motif (RRM) Proteins
-	D12.776.157.725.813 RNA Recognition Motif Proteins
-	D12.776.157.725.813.250 CELF Proteins
-	D12.776.157.725.813.250.500 CELF1 Protein
-	D12.776.157.725.813.500 ELAV Proteins
-	D12.776.157.725.813.500.250 ELAV-Like Protein 1
-	D12.776.157.725.813.500.500 ELAV-Like Protein 2
-	D12.776.157.725.813.500.750 ELAV-Like Protein 3
-	D12.776.157.725.813.500.875 ELAV-Like Protein 4
-	D12.776.157.725.813.750 Heterogeneous-Nuclear Ribonucleoproteins
-	D12.776.157.725.813.750.100 Heterogeneous-Nuclear Ribonucleoprotein Group A-B
-	D12.776.157.725.813.750.200 Heterogeneous-Nuclear Ribonucleoprotein Group C
-	D12.776.157.725.813.750.300 Heterogeneous-Nuclear Ribonucleoprotein D

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D12.776.157.725.813.750.400 Group F-H	Heterogeneous-Nuclear Ribonucleoprotein
-	D12.776.157.725.813.750.500	Heterogeneous-Nuclear Ribonucleoprotein K
-	D12.776.157.725.813.750.600	Heterogeneous-Nuclear Ribonucleoprotein L
-	D12.776.157.725.813.750.700 Group M	Heterogeneous-Nuclear Ribonucleoprotein
-	D12.776.157.725.813.750.800	Heterogeneous-Nuclear Ribonucleoprotein U
-	D12.776.157.725.813.750.900	RNA-Binding Protein EWS
-	D12.776.157.725.813.750.905	RNA-Binding Protein FUS
New Heading	<b>D12.776.157.725.813.875 Gamma Coactivator 1-alpha</b>	<b>Peroxisome Proliferator-Activated Receptor</b>
New Heading	<b>D12.776.157.725.829</b>	<b>RNA Splicing Factors</b>
New Tree	<a href="#">D12.776.157.725.829.250</a>	<a href="#">Polypyrimidine Tract-Binding Protein</a>
New Heading	<b>D12.776.157.725.829.282</b>	<b>PTB-Associated Splicing Factor</b>
New Tree	<a href="#">D12.776.157.725.829.313</a>	<a href="#">Ribonucleoprotein, U1 Small Nuclear</a>
New Tree	<a href="#">D12.776.157.725.829.344</a>	<a href="#">Ribonucleoprotein, U2 Small Nuclear</a>
New Tree	<a href="#">D12.776.157.725.829.375</a>	<a href="#">Ribonucleoprotein, U5 Small Nuclear</a>
New Tree	<a href="#">D12.776.157.725.829.500</a>	<a href="#">Serine-Arginine Splicing Factors</a>
New Heading	<b>D12.776.157.725.829.750</b>	<b>Splicing Factor U2AF</b>
Old Tree	<a href="#">D12.776.157.725.844</a>	<a href="#">Serine-Arginine Splicing Factors</a>
-	D12.776.157.725.875	SMN Complex Proteins
-	D12.776.157.725.875.249	DEAD Box Protein 20
-	D12.776.157.725.875.500	Survival of Motor Neuron 1 Protein
-	D12.776.157.725.875.750	Survival of Motor Neuron 2 Protein
-	D12.776.157.743	S-Phase Kinase-Associated Proteins
-	D12.776.157.762	Sex Hormone-Binding Globulin
-	D12.776.157.785	Thyroxine-Binding Proteins
-	D12.776.157.785.500	Thyroxine-Binding Globulin
-	D12.776.157.800	Transcobalamins
-	D12.776.157.818	Transcortin
-	D12.776.157.905	Transferrin-Binding Proteins
-	D12.776.157.905.500	Receptors, Transferrin

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.157.905.500.249                      Bacterial Transferrin Receptor Complex
-	D12.776.157.905.500.249.500                      Transferrin-Binding Protein A
-	D12.776.157.905.500.249.750                      Transferrin-Binding Protein B
-	D12.776.157.920                      Vitamin D-Binding Protein
-	D12.776.167                      Cell Cycle Proteins
-	D12.776.167.024                      Anaphase-Promoting Complex-Cyclosome
-	D12.776.167.024.500                      Apc1 Subunit, Anaphase-Promoting Complex-Cyclosome
-	D12.776.167.024.625                      Apc10 Subunit, Anaphase-Promoting Complex-Cyclosome
-	D12.776.167.024.687                      Apc11 Subunit, Anaphase-Promoting Complex-Cyclosome
-	D12.776.167.024.750                      Apc2 Subunit, Anaphase-Promoting Complex-Cyclosome
-	D12.776.167.024.875                      Apc3 Subunit, Anaphase-Promoting Complex-Cyclosome
-	D12.776.167.024.937                      Apc4 Subunit, Anaphase-Promoting Complex-Cyclosome
-	D12.776.167.024.968                      Apc5 Subunit, Anaphase-Promoting Complex-Cyclosome
-	D12.776.167.024.984                      Apc6 Subunit, Anaphase-Promoting Complex-Cyclosome
-	D12.776.167.024.992                      Apc7 Subunit, Anaphase-Promoting Complex-Cyclosome
-	D12.776.167.024.996                      Apc8 Subunit, Anaphase-Promoting Complex-Cyclosome
-	D12.776.167.024.998                      Cdc20 Proteins
-	D12.776.167.024.999                      Cdh1 Proteins
-	D12.776.167.049                      Aurora Kinases
-	D12.776.167.049.500                      Aurora Kinase A
-	D12.776.167.049.750                      Aurora Kinase B
-	D12.776.167.049.875                      Aurora Kinase C
-	D12.776.167.100                      cdc25 Phosphatases
-	D12.776.167.150                      Cellular Apoptosis Susceptibility Protein
-	D12.776.167.175                      Cullin Proteins
-	D12.776.167.187                      Cyclin-Dependent Kinase Inhibitor Proteins
-	D12.776.167.187.100                      Cyclin-Dependent Kinase Inhibitor p15
-	D12.776.167.187.200                      Cyclin-Dependent Kinase Inhibitor p16
-	D12.776.167.187.300                      Cyclin-Dependent Kinase Inhibitor p18

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.167.187.400 Cyclin-Dependent Kinase Inhibitor p19
-	D12.776.167.187.500 Cyclin-Dependent Kinase Inhibitor p21
-	D12.776.167.187.600 Cyclin-Dependent Kinase Inhibitor p27
-	D12.776.167.187.700 Cyclin-Dependent Kinase Inhibitor p57
-	D12.776.167.200 Cyclin-Dependent Kinases
-	D12.776.167.200.067 CDC2-CDC28 Kinases
-	D12.776.167.200.067.249 CDC2 Protein Kinase
-	D12.776.167.200.067.500 CDC28 Protein Kinase, S cerevisiae
-	D12.776.167.200.067.875 Cyclin-Dependent Kinase 5
-	D12.776.167.200.067.900 Cyclin-Dependent Kinase 9
-	D12.776.167.200.323 Cyclin-Dependent Kinase 2
-	D12.776.167.200.387 Cyclin-Dependent Kinase 3
-	D12.776.167.200.451 Cyclin-Dependent Kinase 4
-	D12.776.167.200.515 Cyclin-Dependent Kinase 6
-	D12.776.167.200.548 Cyclin-Dependent Kinase 8
-	D12.776.167.200.580 Maturation-Promoting Factor
-	D12.776.167.200.580.500 CDC2 Protein Kinase
-	D12.776.167.218 Cyclins
-	D12.776.167.218.100 Cyclin A
-	D12.776.167.218.100.100 Cyclin A1
-	D12.776.167.218.100.200 Cyclin A2
-	D12.776.167.218.120 Cyclin B
-	D12.776.167.218.120.100 Cyclin B1
-	D12.776.167.218.120.200 Cyclin B2
-	D12.776.167.218.135 Cyclin C
-	D12.776.167.218.150 Cyclin D
-	D12.776.167.218.150.100 Cyclin D1
-	D12.776.167.218.150.200 Cyclin D2
-	D12.776.167.218.150.300 Cyclin D3
-	D12.776.167.218.180 Cyclin E
-	D12.776.167.218.200 Cyclin G
-	D12.776.167.218.200.100 Cyclin G1
-	D12.776.167.218.200.200 Cyclin G2
-	D12.776.167.218.300 Cyclin H
-	D12.776.167.218.400 Cyclin I
-	D12.776.167.218.700 Cyclin T

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.167.265 Geminin
-	D12.776.167.313 Mad2 Proteins
-	D12.776.167.409 Minichromosome Maintenance Proteins
-	D12.776.167.409.100 Minichromosome Maintenance 1 Protein
-	D12.776.167.409.200 Minichromosome Maintenance Complex Component 2
-	D12.776.167.409.300 Minichromosome Maintenance Complex Component 3
-	D12.776.167.409.400 Minichromosome Maintenance Complex Component 4
-	D12.776.167.409.500 Minichromosome Maintenance Complex Component 5
-	D12.776.167.409.600 Minichromosome Maintenance Complex Component 6
-	D12.776.167.409.700 Minichromosome Maintenance Complex Component 7
-	D12.776.167.409.800 Minichromosome Maintenance Complex Component 8
-	D12.776.167.409.900 Minichromosome Maintenance Complex Component 9
New Heading	<b>D12.776.167.457 NIMA-Related Kinases</b>
New Heading	<b>D12.776.167.457.500 NIMA-Related Kinase 1</b>
-	D12.776.167.504 Securin
-	D12.776.167.552 Separase
-	D12.776.167.600 Tumor Suppressor Protein p14ARF
-	D12.776.178 Cerebrospinal Fluid Proteins
-	D12.776.189 Circadian Rhythm Signaling Peptides and Proteins
-	D12.776.189.100 ARNTL Transcription Factors
-	D12.776.189.200 CLOCK Proteins
-	D12.776.189.250 Cryptochromes
-	D12.776.189.300 Nuclear Receptor Subfamily 1, Group D, Member 1
-	D12.776.189.437 Nuclear Receptor Subfamily 1, Group F, Member 1
-	D12.776.189.531 Nuclear Receptor Subfamily 1, Group F, Member 2
-	D12.776.189.625 Period Circadian Proteins
-	D12.776.200 Colipases
-	D12.776.210 Contractile Proteins
-	D12.776.210.249 Filamins
-	D12.776.210.500 Muscle Proteins
-	D12.776.210.500.095 Actinin
-	D12.776.210.500.100 Actins
-	D12.776.210.500.154 Actomyosin
-	D12.776.210.500.220 Calsequestrin

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.210.500.227 CapZ Actin Capping Protein
-	D12.776.210.500.235 Caveolin 3
-	D12.776.210.500.242 Cofilin 2
-	D12.776.210.500.246 Connectin
-	D12.776.210.500.250 Dystrophin
-	D12.776.210.500.410 Dystrophin-Associated Proteins
-	D12.776.210.500.410.500 Dystroglycans
-	D12.776.210.500.410.750 Sarcoglycans
-	D12.776.210.500.570 Myogenic Regulatory Factors
-	D12.776.210.500.570.294 MEF2 Transcription Factors
-	D12.776.210.500.570.590 MyoD Protein
-	D12.776.210.500.570.595 Myogenic Regulatory Factor 5
-	D12.776.210.500.570.600 Myogenin
-	D12.776.210.500.588 Myoglobin
-	D12.776.210.500.600 Myosins
-	D12.776.210.500.600.100 Myosin Heavy Chains
-	D12.776.210.500.600.200 Myosin Light Chains
-	D12.776.210.500.600.300 Myosin Subfragments
-	D12.776.210.500.600.465 Myosin Type I
-	D12.776.210.500.600.470 Myosin Type II
-	D12.776.210.500.600.470.249 Cardiac Myosins
-	D12.776.210.500.600.470.249.249 Atrial Myosins
-	D12.776.210.500.600.470.249.500 Ventricular Myosins
-	D12.776.210.500.600.470.374 Nonmuscle Myosin Type IIA
-	D12.776.210.500.600.470.500 Nonmuscle Myosin Type IIB
-	D12.776.210.500.600.470.750 Skeletal Muscle Myosins
-	D12.776.210.500.600.470.875 Smooth Muscle Myosins
-	D12.776.210.500.675 NAV1.4 Voltage-Gated Sodium Channel
-	D12.776.210.500.750 Parvalbumins
-	D12.776.210.500.775 Profilins
-	D12.776.210.500.800 Ryanodine Receptor Calcium Release Channel
-	D12.776.210.500.847 Tropomodulin
-	D12.776.210.500.895 Tropomyosin
-	D12.776.210.500.910 Troponin
-	D12.776.210.500.910.900 Troponin C
-	D12.776.210.500.910.925 Troponin I



## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.210.500.910.962 Troponin T
-	D12.776.212 CRISPR-Associated Proteins
-	D12.776.215 Cystatins
-	D12.776.215.100 Cystatin A
-	D12.776.215.200 Cystatin B
-	D12.776.215.300 Cystatin C
-	D12.776.215.500 Cystatin M
-	D12.776.215.625 Fetuins
-	D12.776.215.625.500 alpha-2-HS-Glycoprotein
-	D12.776.215.625.750 Fetuin-B
-	D12.776.215.750 Salivary Cystatins
-	D12.776.217 Cystine-Knot Miniproteins
-	D12.776.220 Cytoskeletal Proteins
-	D12.776.220.040 Adenomatous Polyposis Coli Protein
-	D12.776.220.145 Catenins
-	D12.776.220.145.249 alpha Catenin
-	D12.776.220.145.500 beta Catenin
-	D12.776.220.145.750 gamma Catenin
-	D12.776.220.150 Cornified Envelope Proline-Rich Proteins
-	D12.776.220.250 Dystrophin
-	D12.776.220.362 Dystrophin-Associated Proteins
-	D12.776.220.362.249 Dystroglycans
-	D12.776.220.475 Intermediate Filament Proteins
-	D12.776.220.475.200 Desmin
-	D12.776.220.475.400 Glial Fibrillary Acidic Protein
-	D12.776.220.475.450 Keratins
-	D12.776.220.475.450.074 beta-Keratins
-	D12.776.220.475.450.149 Keratins, Hair-Specific
-	D12.776.220.475.450.300 Keratins, Type I
-	D12.776.220.475.450.300.100 Keratin-10
-	D12.776.220.475.450.300.200 Keratin-12
-	D12.776.220.475.450.300.300 Keratin-13
-	D12.776.220.475.450.300.400 Keratin-14
-	D12.776.220.475.450.300.500 Keratin-15
-	D12.776.220.475.450.300.600 Keratin-16
-	D12.776.220.475.450.300.700 Keratin-17

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.220.475.450.300.800 Keratin-18
-	D12.776.220.475.450.300.900 Keratin-19
-	D12.776.220.475.450.300.950 Keratin-20
-	D12.776.220.475.450.600 Keratins, Type II
-	D12.776.220.475.450.600.100 Keratin-1
-	D12.776.220.475.450.600.200 Keratin-2
-	D12.776.220.475.450.600.300 Keratin-3
-	D12.776.220.475.450.600.400 Keratin-4
-	D12.776.220.475.450.600.500 Keratin-5
-	D12.776.220.475.450.600.600 Keratin-6
-	D12.776.220.475.450.600.700 Keratin-7
-	D12.776.220.475.450.600.800 Keratin-8
-	D12.776.220.475.450.600.900 Keratin-9
-	D12.776.220.475.540 Nestin
-	D12.776.220.475.630 Neurofilament Proteins
-	D12.776.220.475.900 Vimentin
-	D12.776.220.525 Microfilament Proteins
-	D12.776.220.525.032 Actin Capping Proteins
-	D12.776.220.525.032.500 CapZ Actin Capping Protein
-	D12.776.220.525.032.750 Tropomodulin
-	D12.776.220.525.212 Actin Depolymerizing Factors
-	D12.776.220.525.212.500 Cofilin 1
-	D12.776.220.525.212.750 Cofilin 2
-	D12.776.220.525.212.875 Destrin
-	D12.776.220.525.246 Actin-Related Protein 2-3 Complex
-	D12.776.220.525.246.500 Actin-Related Protein 2
-	D12.776.220.525.246.750 Actin-Related Protein 3
-	D12.776.220.525.250 Actinin
-	D12.776.220.525.255 Actins
-	D12.776.220.525.281 Cortactin
-	D12.776.220.525.315 Filamins
-	D12.776.220.525.350 Gelsolin
-	D12.776.220.525.475 Myosins
-	D12.776.220.525.475.100 Myosin Heavy Chains
-	D12.776.220.525.475.200 Myosin Light Chains
-	D12.776.220.525.475.300 Myosin Subfragments

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D12.776.220.525.475.470	Myosin Type I
-	D12.776.220.525.475.475	Myosin Type II
-	D12.776.220.525.475.475.124	Cardiac Myosins
-	D12.776.220.525.475.475.124.249	Atrial Myosins
-	D12.776.220.525.475.475.124.500	Ventricular Myosins
-	D12.776.220.525.475.475.249	Nonmuscle Myosin Type IIA
-	D12.776.220.525.475.475.500	Nonmuscle Myosin Type IIB
-	D12.776.220.525.475.475.750	Skeletal Muscle Myosins
-	D12.776.220.525.475.475.875	Smooth Muscle Myosins
-	D12.776.220.525.475.612	Myosin Type III
-	D12.776.220.525.475.681	Myosin Type IV
-	D12.776.220.525.475.750	Myosin Type V
-	D12.776.220.525.637	Profilins
New Heading	<b>D12.776.220.525.719</b>	<b>Tensins</b>
-	D12.776.220.525.800	Tropomyosin
-	D12.776.220.525.825	Troponin
-	D12.776.220.525.825.900	Troponin C
-	D12.776.220.525.825.925	Troponin I
-	D12.776.220.525.825.962	Troponin T
-	D12.776.220.525.912	Wiskott-Aldrich Syndrome Protein Family
-	D12.776.220.525.912.500	Wiskott-Aldrich Syndrome Protein
-	D12.776.220.525.912.550	Wiskott-Aldrich Syndrome Protein, Neuronal
-	D12.776.220.600	Microtubule Proteins
-	D12.776.220.600.200	Dyneins
-	D12.776.220.600.450	Microtubule-Associated Proteins
New Heading	<b>D12.776.220.600.450.100</b>	<b>Autophagy-Related Protein 8 Family</b>
New Heading	<b>D12.776.220.600.450.150</b>	<b>Dynactin Complex</b>
-	D12.776.220.600.450.200	Dynamins
-	D12.776.220.600.450.200.100	Dynamin I
-	D12.776.220.600.450.200.200	Dynamin II
-	D12.776.220.600.450.200.300	Dynamin III
-	D12.776.220.600.450.450	Kinesin
-	D12.776.220.600.450.480	Stathmin
-	D12.776.220.600.450.510	tau Proteins

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.220.600.800 Tubulin
-	D12.776.220.790 Plakins
-	D12.776.220.790.500 Desmoplakins
New Heading	<b>D12.776.220.790.625 Dystonin</b>
-	D12.776.220.790.750 Plectin
-	D12.776.220.885 Plakophilins
New Heading	<b>D12.776.220.909 Pyrin</b>
-	D12.776.220.932 Septins
-	D12.776.220.980 Spectrin
-	D12.776.220.985 Talin
-	D12.776.220.987 Utrophin
-	D12.776.220.990 Vinculin
-	D12.776.220.995 Zyxin
-	D12.776.231 Dental Enamel Proteins
-	D12.776.231.500 Amelogenin
-	D12.776.256 Dietary Proteins
-	D12.776.256.317 Egg Proteins, Dietary
-	D12.776.256.317.180 Conalbumin
-	D12.776.256.317.663 Ovalbumin
-	D12.776.256.317.663.300 Avidin
-	D12.776.256.317.675 Ovomucin
-	D12.776.256.317.700 Phosvitin
-	D12.776.256.626 Milk Proteins
-	D12.776.256.626.207 Caseins
-	D12.776.256.626.816 Whey Proteins
-	D12.776.256.626.816.250 Lactalbumin
-	D12.776.256.626.816.500 Lactoglobulins
-	D12.776.256.626.816.500.507 Lactoferrin
-	D12.776.256.920 Vegetable Proteins
-	D12.776.260 DNA-Binding Proteins
-	D12.776.260.060 Adenovirus E2 Proteins
-	D12.776.260.103 Basic Helix-Loop-Helix Transcription Factors
-	D12.776.260.103.249 ARNTL Transcription Factors
-	D12.776.260.103.500 Basic Helix-Loop-Helix Leucine Zipper Transcription Factors

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D12.776.260.103.500.500	Microphthalmia-Associated Transcription Factor
New Heading	<b>D12.776.260.103.500.625</b>	<b>N-Myc Proto-Oncogene Protein</b>
-	D12.776.260.103.500.750	Sterol Regulatory Element Binding Proteins
-	D12.776.260.103.500.750.500	Sterol Regulatory Element Binding Protein 1
-	D12.776.260.103.500.750.750	Sterol Regulatory Element Binding Protein 2
-	D12.776.260.103.500.875	Transcription Factor 3
-	D12.776.260.103.562	CLOCK Proteins
-	D12.776.260.103.625	Hypoxia-Inducible Factor 1
-	D12.776.260.103.625.500	Aryl Hydrocarbon Receptor Nuclear Translocator
-	D12.776.260.103.625.750	Hypoxia-Inducible Factor 1, alpha Subunit
-	D12.776.260.103.750	Myogenic Regulatory Factors
-	D12.776.260.103.750.294	MEF2 Transcription Factors
-	D12.776.260.103.750.590	MyoD Protein
-	D12.776.260.103.750.595	Myogenic Regulatory Factor 5
-	D12.776.260.103.750.600	Myogenin
New Tree	<a href="#">D12.776.260.103.782</a>	<a href="#">Nuclear Receptor Coactivator 1</a>
New Tree	<a href="#">D12.776.260.103.813</a>	<a href="#">Proto-Oncogene Proteins c-myc</a>
New Heading	<b>D12.776.260.103.844</b>	<b>Transcription Factor HES-1</b>
Old Tree	<a href="#">D12.776.260.103.875</a>	<a href="#">Twist Transcription Factor</a>
Old Tree	<a href="#">D12.776.260.103.875</a>	<a href="#">Twist-Related Protein 1</a>
New Heading	<b>D12.776.260.103.906</b>	<b>Twist Transcription Factors</b>
New Tree	<a href="#">D12.776.260.103.906.250</a>	<a href="#">Twist Transcription Factor</a>
New Tree	<a href="#">D12.776.260.103.906.250</a>	<a href="#">Twist-Related Protein 1</a>
New Heading	<b>D12.776.260.103.906.500</b>	<b>Twist-Related Protein 2</b>
-	D12.776.260.103.937	Upstream Stimulatory Factors
-	D12.776.260.108	Basic-Leucine Zipper Transcription Factors
-	D12.776.260.108.061	Activating Transcription Factors
-	D12.776.260.108.061.500	Activating Transcription Factor 1
-	D12.776.260.108.061.750	Activating Transcription Factor 2
-	D12.776.260.108.061.875	Activating Transcription Factor 3
-	D12.776.260.108.061.937	Activating Transcription Factor 4

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.260.108.061.968      Activating Transcription Factor 6
-	D12.776.260.108.092      Basic Helix-Loop-Helix Leucine Zipper Transcription Factors
-	D12.776.260.108.092.500      Microphthalmia-Associated Transcription Factor
New Heading	<b>D12.776.260.108.092.625      N-Myc Proto-Oncogene Protein</b>
-	D12.776.260.108.092.750      Sterol Regulatory Element Binding Proteins
-	D12.776.260.108.092.750.500      Sterol Regulatory Element Binding Protein 1
-	D12.776.260.108.092.750.750      Sterol Regulatory Element Binding Protein 2
-	D12.776.260.108.092.875      Transcription Factor 3
-	D12.776.260.108.124      CCAAT-Enhancer-Binding Proteins
-	D12.776.260.108.124.249      CCAAT-Binding Factor
-	D12.776.260.108.124.500      CCAAT-Enhancer-Binding Protein-alpha
-	D12.776.260.108.124.750      CCAAT-Enhancer-Binding Protein-beta
-	D12.776.260.108.124.812      CCAAT-Enhancer-Binding Protein-delta
-	D12.776.260.108.124.875      Transcription Factor CHOP
-	D12.776.260.108.124.937      Y-Box-Binding Protein 1
-	D12.776.260.108.184      Cyclic AMP Response Element-Binding Protein
-	D12.776.260.108.186      Cyclic AMP Response Element-Binding Protein A
-	D12.776.260.108.217      Cyclic AMP Response Element Modulator
-	D12.776.260.108.311      Fos-Related Antigen-2
-	D12.776.260.108.374      Interferon Regulatory Factor-1
-	D12.776.260.108.500      Maf Transcription Factors
-	D12.776.260.108.500.061      Maf Transcription Factors, Large
-	D12.776.260.108.500.061.500      MafB Transcription Factor
-	D12.776.260.108.500.061.625      Oncogene Protein v-maf
-	D12.776.260.108.500.061.750      Proto-Oncogene Proteins c-maf
-	D12.776.260.108.500.500      Maf Transcription Factors, Small
-	D12.776.260.108.500.500.249      MafF Transcription Factor
-	D12.776.260.108.500.500.500      MafG Transcription Factor
-	D12.776.260.108.500.500.750      MafK Transcription Factor
-	D12.776.260.108.656      NF-E2 Transcription Factor
-	D12.776.260.108.656.750      Maf Transcription Factors, Small
-	D12.776.260.108.656.750.249      MafF Transcription Factor
-	D12.776.260.108.656.750.500      MafG Transcription Factor
-	D12.776.260.108.656.750.750      MafK Transcription Factor

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.260.108.656.770 NF-E2 Transcription Factor, p45 Subunit
-	D12.776.260.108.710 NF-E2-Related Factor 1
-	D12.776.260.108.737 NF-E2-Related Factor 2
-	D12.776.260.108.765 Proto-Oncogene Proteins c-fos
-	D12.776.260.108.820 Proto-Oncogene Proteins c-jun
-	D12.776.260.108.875 Transcription Factor AP-1
New Heading	<b>D12.776.260.108.937 X-Box Binding Protein 1</b>
-	D12.776.260.114 Butyrate Response Factor 1
-	D12.776.260.125 Centromere Protein B
-	D12.776.260.158 Early Growth Response Transcription Factors
-	D12.776.260.158.500 Early Growth Response Protein 1
-	D12.776.260.158.750 Early Growth Response Protein 2
-	D12.776.260.158.875 Early Growth Response Protein 3
-	D12.776.260.235 Erythroid-Specific DNA-Binding Factors
-	D12.776.260.235.500 GATA1 Transcription Factor
-	D12.776.260.235.625 GATA2 Transcription Factor
-	D12.776.260.235.687 GATA3 Transcription Factor
-	D12.776.260.235.750 NF-E2 Transcription Factor
-	D12.776.260.235.750.750 Maf Transcription Factors, Small
-	D12.776.260.235.750.750.249 MafF Transcription Factor
-	D12.776.260.235.750.750.500 MafG Transcription Factor
-	D12.776.260.235.750.750.750 MafK Transcription Factor
-	D12.776.260.235.750.770 NF-E2 Transcription Factor, p45 Subunit
-	D12.776.260.235.875 YY1 Transcription Factor
-	D12.776.260.246 Factor For Inversion Stimulation Protein
-	D12.776.260.254 G-Box Binding Factors
-	D12.776.260.257 GATA Transcription Factors
-	D12.776.260.257.100 GATA1 Transcription Factor
-	D12.776.260.257.200 GATA2 Transcription Factor
-	D12.776.260.257.300 GATA3 Transcription Factor
-	D12.776.260.257.400 GATA4 Transcription Factor
-	D12.776.260.257.500 GATA5 Transcription Factor
-	D12.776.260.257.600 GATA6 Transcription Factor
-	D12.776.260.262 Hepatocyte Nuclear Factors
-	D12.776.260.262.500 Hepatocyte Nuclear Factor 1

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.260.262.500.500 Hepatocyte Nuclear Factor 1-alpha
-	D12.776.260.262.500.750 Hepatocyte Nuclear Factor 1-beta
-	D12.776.260.262.750 Hepatocyte Nuclear Factor 3-alpha
-	D12.776.260.262.875 Hepatocyte Nuclear Factor 3-beta
-	D12.776.260.262.937 Hepatocyte Nuclear Factor 3-gamma
-	D12.776.260.262.968 Hepatocyte Nuclear Factor 4
-	D12.776.260.262.984 Hepatocyte Nuclear Factor 6
-	D12.776.260.268 Heterogeneous-Nuclear Ribonucleoprotein K
-	D12.776.260.312 HMGA Proteins
-	D12.776.260.312.500 HMGA1a Protein
-	D12.776.260.312.750 HMGA1b Protein
-	D12.776.260.312.875 HMGA1c Protein
-	D12.776.260.312.937 HMGA2 Protein
-	D12.776.260.356 HMGB Proteins
-	D12.776.260.356.300 HMGB1 Protein
-	D12.776.260.356.600 HMGB2 Protein
-	D12.776.260.356.800 HMGB3 Protein
-	D12.776.260.400 Homeodomain Proteins
-	D12.776.260.400.124 Antennapedia Homeodomain Protein
New Heading	<b>D12.776.260.400.140 CDX2 Transcription Factor</b>
-	D12.776.260.400.155 Fushi Tarazu Transcription Factors
-	D12.776.260.400.186 Goosecoid Protein
New Tree	<a href="#">D12.776.260.400.218 Hepatocyte Nuclear Factor 1</a>
New Tree	<a href="#">D12.776.260.400.218.500 Hepatocyte Nuclear Factor 1-alpha</a>
New Tree	<a href="#">D12.776.260.400.218.750 Hepatocyte Nuclear Factor 1-beta</a>
New Heading	<b>D12.776.260.400.234 Homeobox Protein Nkx-2.5</b>
-	D12.776.260.400.249 MADS Domain Proteins
-	D12.776.260.400.249.249 AGAMOUS Protein, Arabidopsis
-	D12.776.260.400.249.374 DEFICIENS Protein
-	D12.776.260.400.249.624 MEF2 Transcription Factors
-	D12.776.260.400.249.875 Serum Response Factor
-	D12.776.260.400.436 MSX1 Transcription Factor



## MeSH Tree Changes for 2017

Type	Tree - heading	
New Heading	<b>D12.776.260.400.530</b>	<b>Nanog Homeobox Protein</b>
-	D12.776.260.400.624	Onecut Transcription Factors
-	D12.776.260.400.624.500	Hepatocyte Nuclear Factor 6
-	D12.776.260.400.718	Otx Transcription Factors
-	D12.776.260.400.812	PAX7 Transcription Factor
New Heading	<b>D12.776.260.400.906</b>	<b>Zinc Finger E-box-Binding Homeobox 1</b>
-	D12.776.260.420	I-kappa B Proteins
New Heading	<b>D12.776.260.420.500</b>	<b>NF-KappaB Inhibitor alpha</b>
-	D12.776.260.457 Protein	Immunoglobulin J Recombination Signal Sequence-Binding
-	D12.776.260.495	Integration Host Factors
-	D12.776.260.504	Interferon Regulatory Factors
-	D12.776.260.504.124	Interferon Regulatory Factor-1
-	D12.776.260.504.249	Interferon Regulatory Factor-2
-	D12.776.260.504.374	Interferon Regulatory Factor-3
-	D12.776.260.504.437	Interferon Regulatory Factor-7
-	D12.776.260.504.500	Interferon-Stimulated Gene Factor 3, gamma Subunit
-	D12.776.260.513	Interferon-Stimulated Gene Factor 3
-	D12.776.260.513.249	Interferon-Stimulated Gene Factor 3, alpha Subunit
-	D12.776.260.513.249.500	STAT1 Transcription Factor
-	D12.776.260.513.249.750	STAT2 Transcription Factor
-	D12.776.260.513.500	Interferon-Stimulated Gene Factor 3, gamma Subunit
-	D12.776.260.522	Kruppel-Like Transcription Factors
-	D12.776.260.522.500	Ikaros Transcription Factor
-	D12.776.260.522.750	Sp Transcription Factors
-	D12.776.260.522.750.249	Sp1 Transcription Factor
-	D12.776.260.522.750.500	Sp2 Transcription Factor
-	D12.776.260.522.750.750	Sp3 Transcription Factor
-	D12.776.260.522.750.875	Sp4 Transcription Factor
New Heading	<b>D12.776.260.525</b>	<b>Ku Autoantigen</b>
-	D12.776.260.527	Leucine-Responsive Regulatory Protein
-	D12.776.260.529	LIM-Homeodomain Proteins
New Heading	<b>D12.776.260.531</b>	<b>Liver X Receptors</b>

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D12.776.260.532	Matrix Attachment Region Binding Proteins
-	D12.776.260.536	Methyl-CpG-Binding Protein 2
-	D12.776.260.538	Minichromosome Maintenance 1 Protein
New Heading	<b>D12.776.260.540</b>	<b>MutL Proteins</b>
New Heading	<b>D12.776.260.540.250</b>	<b>Mismatch Repair Endonuclease PMS2</b>
New Heading	<b>D12.776.260.540.500</b>	<b>MutL Protein Homolog 1</b>
-	D12.776.260.541	MutS DNA Mismatch-Binding Protein
-	D12.776.260.551	MutS Homolog 2 Protein
-	D12.776.260.560	Myeloid-Lymphoid Leukemia Protein
-	D12.776.260.600	NF-kappa B
-	D12.776.260.600.124	NF-kappa B p50 Subunit
-	D12.776.260.600.186	NF-kappa B p52 Subunit
-	D12.776.260.600.249	Transcription Factor RelA
-	D12.776.260.600.500	Transcription Factor RelB
-	D12.776.260.605	NFI Transcription Factors
-	D12.776.260.615	Nuclear Respiratory Factors
-	D12.776.260.615.249	GA-Binding Protein Transcription Factor
-	D12.776.260.615.500	Nuclear Respiratory Factor 1
-	D12.776.260.630	Oncogene Protein p55(v-myc)
-	D12.776.260.640	Origin Recognition Complex
-	D12.776.260.643	Orphan Nuclear Receptors
-	D12.776.260.643.039	COUP Transcription Factors
-	D12.776.260.643.039.249	COUP Transcription Factor I
-	D12.776.260.643.039.500	COUP Transcription Factor II
-	D12.776.260.643.080	Nuclear Receptor Subfamily 1, Group D, Member 1
-	D12.776.260.643.100	Nuclear Receptor Subfamily 1, Group F, Member 1
-	D12.776.260.643.110	Nuclear Receptor Subfamily 1, Group F, Member 2
-	D12.776.260.643.182	Nuclear Receptor Subfamily 1, Group F, Member 3
-	D12.776.260.643.255	Nuclear Receptor Subfamily 2, Group C, Member 1
-	D12.776.260.643.327	Nuclear Receptor Subfamily 2, Group C, Member 2
-	D12.776.260.643.400	Nuclear Receptor Subfamily 4, Group A, Member 1
-	D12.776.260.643.415	Nuclear Receptor Subfamily 4, Group A, Member 2
-	D12.776.260.643.430	Nuclear Receptor Subfamily 4, Group A, Member 3
-	D12.776.260.643.572	Nuclear Receptor Subfamily 6, Group A, Member 1

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D12.776.260.643.715	Receptors, Aryl Hydrocarbon
-	D12.776.260.645	Paired Box Transcription Factors
-	D12.776.260.645.500	B-Cell-Specific Activator Protein
-	D12.776.260.645.500	PAX5 Transcription Factor
-	D12.776.260.645.750	PAX2 Transcription Factor
New Heading	<b>D12.776.260.645.782</b>	<b>PAX3 Transcription Factor</b>
New Heading	<b>D12.776.260.645.813</b>	<b>PAX6 Transcription Factor</b>
-	D12.776.260.645.875	PAX7 Transcription Factor
New Heading	<b>D12.776.260.645.906</b>	<b>PAX8 Transcription Factor</b>
-	D12.776.260.645.937	PAX9 Transcription Factor
New Heading	<b>D12.776.260.650</b>	<b>Peptidyl-Prolyl Cis-Trans Isomerase NIMA-Interacting 4</b>
-	D12.776.260.655	POU Domain Factors
-	D12.776.260.655.500	Octamer Transcription Factors
-	D12.776.260.655.500.100	Octamer Transcription Factor-1
-	D12.776.260.655.500.200	Octamer Transcription Factor-2
-	D12.776.260.655.500.300	Octamer Transcription Factor-3
-	D12.776.260.655.500.600	Octamer Transcription Factor-6
-	D12.776.260.655.625	Transcription Factor Brn-3
-	D12.776.260.655.625.124	Transcription Factor Brn-3A
-	D12.776.260.655.625.249	Transcription Factor Brn-3B
-	D12.776.260.655.625.500	Transcription Factor Brn-3C
-	D12.776.260.655.750	Transcription Factor Pit-1
-	D12.776.260.660	Proto-Oncogene Proteins c-bcl-6
-	D12.776.260.665	Proto-Oncogene Proteins c-ets
-	D12.776.260.665.100	Proto-Oncogene Protein c-ets-1
-	D12.776.260.665.200	Proto-Oncogene Protein c-ets-2
-	D12.776.260.665.400	Proto-Oncogene Protein c-fli-1
-	D12.776.260.665.600	Ternary Complex Factors
-	D12.776.260.665.600.100	ets-Domain Protein Elk-1
-	D12.776.260.665.600.300	ets-Domain Protein Elk-4
-	D12.776.260.675	Proto-Oncogene Proteins c-myb
Old Tree	<b>D12.776.260.676</b>	<b>Proto-Oncogene Proteins c-myc</b>
-	D12.776.260.682	Proto-Oncogene Proteins c-rel

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D12.776.260.690	Proto-Oncogene Proteins c-sis
New Heading	<b>D12.776.260.693</b>	<b>PTB-Associated Splicing Factor</b>
-	D12.776.260.695	Rad51 Recombinase
-	D12.776.260.696	Rad52 DNA Repair and Recombination Protein
-	D12.776.260.700	Replication Protein A
-	D12.776.260.702	Replication Protein C
-	D12.776.260.703	Repressor Proteins
-	D12.776.260.703.099	AraC Transcription Factor
-	D12.776.260.703.200	COUP Transcription Factor II
-	D12.776.260.703.600	Lac Repressors
New Heading	<b>D12.776.260.703.800</b>	<b>Zinc Finger E-box-Binding Homeobox 1</b>
-	D12.776.260.704	Retinoblastoma Protein
-	D12.776.260.713	Smad Proteins
-	D12.776.260.713.500	Smad Proteins, Receptor-Regulated
-	D12.776.260.713.500.100	Smad1 Protein
-	D12.776.260.713.500.300	Smad3 Protein
-	D12.776.260.713.500.500	Smad5 Protein
-	D12.776.260.713.500.800	Smad8 Protein
-	D12.776.260.713.750	Smad4 Protein
New Heading	<b>D12.776.260.716</b>	<b>SMARCB1 Protein</b>
-	D12.776.260.719	SOX Transcription Factors
-	D12.776.260.719.049	Sex-Determining Region Y Protein
-	D12.776.260.719.100	SOXB1 Transcription Factors
-	D12.776.260.719.200	SOXB2 Transcription Factors
-	D12.776.260.719.300	SOXC Transcription Factors
-	D12.776.260.719.400	SOXD Transcription Factors
-	D12.776.260.719.500	SOXE Transcription Factors
-	D12.776.260.719.500.500	SOX9 Transcription Factor
-	D12.776.260.719.600	SOXF Transcription Factors
-	D12.776.260.725	T-Box Domain Proteins
-	D12.776.260.730	TCF Transcription Factors
-	D12.776.260.730.500	Lymphoid Enhancer-Binding Factor 1
-	D12.776.260.730.750	T Cell Transcription Factor 1
-	D12.776.260.730.812	Transcription Factor 7-Like 1 Protein

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D12.776.260.730.875	Transcription Factor 7-Like 2 Protein
-	D12.776.260.735	Telomere-Binding Proteins
-	D12.776.260.735.500	Heterogeneous-Nuclear Ribonucleoprotein Group A-B
-	D12.776.260.735.750	Telomeric Repeat Binding Protein 1
-	D12.776.260.735.875	Telomeric Repeat Binding Protein 2
-	D12.776.260.750	Toll-Like Receptor 9
-	D12.776.260.755	Trans-Activators
New Heading	<b>D12.776.260.755.100</b>	<b>Transcriptional Regulator ERG</b>
-	D12.776.260.755.199	Gene Products, tat
-	D12.776.260.755.199.500	tat Gene Products, Human Immunodeficiency Virus
-	D12.776.260.755.400	Herpes Simplex Virus Protein Vmw65
New Heading	<b>D12.776.260.755.700</b>	<b>Transcription Activator-Like Effectors</b>
New Heading	<b>D12.776.260.755.850</b>	<b>Zinc Finger Protein GLI1</b>
-	D12.776.260.760	Transcription Factor AP-2
-	D12.776.260.775	Transcription Factors, General
-	D12.776.260.775.092	Pol1 Transcription Initiation Complex Proteins
-	D12.776.260.775.186	TATA-Binding Protein Associated Factors
-	D12.776.260.775.374	TATA-Box Binding Protein
-	D12.776.260.775.812	TATA Box Binding Protein-Like Proteins
-	D12.776.260.775.875	Transcription Factors, TFII
-	D12.776.260.775.875.374	Transcription Factor TFIIA
-	D12.776.260.775.875.562	Transcription Factor TFII B
-	D12.776.260.775.875.750	Transcription Factor TFII D
-	D12.776.260.775.875.750.500	TATA-Box Binding Protein
-	D12.776.260.775.875.875	Transcription Factor TFII H
-	D12.776.260.775.875.875.500	Xeroderma Pigmentosum Group D Protein
-	D12.776.260.775.937	Transcription Factors, TFIII
-	D12.776.260.775.937.249	Transcription Factor TFIII A
-	D12.776.260.775.937.500	Transcription Factor TFIII B
-	D12.776.260.790	Tristetraprolin
New Heading	<b>D12.776.260.805</b>	<b>Tumor Suppressor p53-Binding Protein 1</b>
-	D12.776.260.820	Tumor Suppressor Protein p53
New	<b>D12.776.260.885</b>	<b>Tumor Protein p73</b>

## MeSH Tree Changes for 2017

Type	Tree - heading	
Heading		
-	D12.776.260.950	Winged-Helix Transcription Factors
-	D12.776.260.950.249	Forkhead Transcription Factors
New Heading	<b>D12.776.260.950.249.063</b>	<b>Forkhead Box Protein M1</b>
New Heading	<b>D12.776.260.950.249.125</b>	<b>Forkhead Box Protein O3</b>
New Heading	<b>D12.776.260.950.249.250</b>	<b>Forkhead Box Protein O1</b>
-	D12.776.260.950.249.500	Hepatocyte Nuclear Factor 3-alpha
-	D12.776.260.950.249.750	Hepatocyte Nuclear Factor 3-beta
-	D12.776.260.950.249.875	Hepatocyte Nuclear Factor 3-gamma
New Heading	<b>D12.776.260.950.624</b>	<b>Regulatory Factor X Transcription Factors</b>
New Heading	<b>D12.776.260.950.624.500</b>	<b>Regulatory Factor X1</b>
-	D12.776.260.975	Xeroderma Pigmentosum Group A Protein
-	D12.776.290	Egg Proteins
-	D12.776.290.180	Conalbumin
-	D12.776.290.300	Egg Proteins, Dietary
-	D12.776.290.663	Ovalbumin
-	D12.776.290.663.100	Avidin
-	D12.776.290.675	Ovomucin
-	D12.776.290.700	Phosvitin
-	D12.776.290.812	Vitellins
-	D12.776.290.812.500	Vitellogenins
New Heading	<b>D12.776.290.906</b>	<b>Zona Pellucida Glycoproteins</b>
-	D12.776.298	Epididymal Secretory Proteins
-	D12.776.306	Eye Proteins
-	D12.776.306.090	Arrestins
-	D12.776.306.090.050	Arrestin
-	D12.776.306.366	Crystallins
-	D12.776.306.366.100	alpha-Crystallins
-	D12.776.306.366.100.149	alpha-Crystallin A Chain
-	D12.776.306.366.100.300	alpha-Crystallin B Chain
-	D12.776.306.366.300	beta-Crystallins
-	D12.776.306.366.300.100	beta-Crystallin A Chain

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.306.366.300.200 beta-Crystallin B Chain
-	D12.776.306.366.310 delta-Crystallins
-	D12.776.306.366.350 epsilon-Crystallins
-	D12.776.306.366.850 gamma-Crystallins
-	D12.776.306.366.925 omega-Crystallins
-	D12.776.306.366.962 tau-Crystallins
-	D12.776.306.366.981 zeta-Crystallins
-	D12.776.306.399 G-Protein-Coupled Receptor Kinase 1
-	D12.776.306.433 Guanylate Cyclase-Activating Proteins
-	D12.776.306.466 Opsins
-	D12.776.306.466.249 Cone Opsins
-	D12.776.306.466.500 Rod Opsins
-	D12.776.306.750 Recoverin
-	D12.776.313 Fanconi Anemia Complementation Group Proteins
New Tree	<a href="#">D12.776.313.125</a> <a href="#">BRCA1 Protein</a>
-	D12.776.313.249 BRCA2 Protein
-	D12.776.313.500 Fanconi Anemia Complementation Group A Protein
-	D12.776.313.750 Fanconi Anemia Complementation Group C Protein
-	D12.776.313.812 Fanconi Anemia Complementation Group D2 Protein
-	D12.776.313.843 Fanconi Anemia Complementation Group E Protein
-	D12.776.313.875 Fanconi Anemia Complementation Group F Protein
-	D12.776.313.906 Fanconi Anemia Complementation Group G Protein
-	D12.776.313.937 Fanconi Anemia Complementation Group L Protein
New Tree	<a href="#">D12.776.313.968</a> <a href="#">Rad51 Recombinase</a>
-	D12.776.320 Fetal Proteins
-	D12.776.320.525 Fetal Globulins
-	D12.776.320.525.500 alpha-Fetoproteins
-	D12.776.325 Fish Proteins
-	D12.776.325.500 Zebrafish Proteins
-	D12.776.331 Flavoproteins
-	D12.776.331.049 Acetolactate Synthase
-	D12.776.331.099 Acyl-CoA Dehydrogenase
-	D12.776.331.102 Acyl-CoA Dehydrogenase, Long-Chain
-	D12.776.331.149 Acyl-CoA Oxidase

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.331.161 Apoptosis Inducing Factor
-	D12.776.331.174 Butyryl-CoA Dehydrogenase
-	D12.776.331.180 Cryptochromes
-	D12.776.331.186 Cytochrome-B(5) Reductase
-	D12.776.331.192 Dihydrolipoamide Dehydrogenase
-	D12.776.331.199 Electron-Transferring Flavoproteins
-	D12.776.331.199.500 Electron Transport Complex I
-	D12.776.331.199.750 Electron Transport Complex II
-	D12.776.331.199.750.500 Succinate Dehydrogenase
-	D12.776.331.400 Flavodoxin
-	D12.776.331.737 Glutamate Synthase (NADH)
-	D12.776.331.775 Methylene tetrahydrofolate Reductase (NADPH2)
-	D12.776.331.887 NADH Dehydrogenase
-	D12.776.331.894 NADPH Oxidase
-	D12.776.331.899 Nitrate Reductase (NADH)
-	D12.776.331.901 Nitrate Reductase (NAD(P)H)
-	D12.776.331.911 Nitrate Reductase (NADPH)
-	D12.776.331.915 Retinal Dehydrogenase
-	D12.776.331.943 Sarcosine Oxidase
-	D12.776.331.971 Thioredoxin-Disulfide Reductase
-	D12.776.354 Fungal Proteins
-	D12.776.354.374 Killer Factors, Yeast
-	D12.776.354.562 Minichromosome Maintenance 1 Protein
-	D12.776.354.750 Saccharomyces cerevisiae Proteins
-	D12.776.354.750.124 CDC28 Protein Kinase, S cerevisiae
-	D12.776.354.750.249 cdc42 GTP-Binding Protein, Saccharomyces cerevisiae
-	D12.776.354.750.750 Silent Information Regulator Proteins, Saccharomyces cerevisiae
-	D12.776.354.875 Schizosaccharomyces pombe Proteins
-	D12.776.377 Globulins
-	D12.776.377.228 Fetal Globulins
-	D12.776.377.228.500 alpha-Fetoproteins
-	D12.776.377.457 Lactoglobulins
-	D12.776.377.457.507 Lactoferrin
-	D12.776.377.715 Serum Globulins
-	D12.776.377.715.085 Alpha-Globulins



## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.377.715.085.050      alpha 1-Antichymotrypsin
-	D12.776.377.715.085.085      alpha 1-Antitrypsin
-	D12.776.377.715.085.090      alpha-2-Antiplasmin
-	D12.776.377.715.085.092      alpha-Fetoproteins
-	D12.776.377.715.085.100      alpha-Macroglobulins
-	D12.776.377.715.085.125      Antithrombin III
-	D12.776.377.715.085.214      Ceruloplasmin
-	D12.776.377.715.085.304      Fetuins
-	D12.776.377.715.085.304.500      alpha-2-HS-Glycoprotein
-	D12.776.377.715.085.304.750      Fetuin-B
-	D12.776.377.715.085.394      Haptoglobins
-	D12.776.377.715.085.450      Heparin Cofactor II
-	D12.776.377.715.085.640      Orosomuroid
-	D12.776.377.715.085.740      Progesterone-Binding Globulin
-	D12.776.377.715.085.745      Retinol-Binding Proteins, Plasma
-	D12.776.377.715.085.901      Transcortin
-	D12.776.377.715.182      Beta-Globulins
-	D12.776.377.715.182.100      beta 2-Microglobulin
-	D12.776.377.715.182.160      beta-Thromboglobulin
-	D12.776.377.715.182.200      Complement Factor H
-	D12.776.377.715.182.338      Hemopexin
-	D12.776.377.715.182.580      Plasminogen
-	D12.776.377.715.182.580.500      Angiostatins
-	D12.776.377.715.182.624      Properdin
-	D12.776.377.715.182.800      Sex Hormone-Binding Globulin
-	D12.776.377.715.182.839      Transferrin
-	D12.776.377.715.390      Fibronectins
-	D12.776.377.715.548      Immunoglobulins
-	D12.776.377.715.548.114      Antibodies
-	D12.776.377.715.548.114.071      Antibodies, Anti-Idiotypic
-	D12.776.377.715.548.114.071.500      Omalizumab
-	D12.776.377.715.548.114.107      Antibodies, Archaeal
-	D12.776.377.715.548.114.125      Antibodies, Bacterial
-	D12.776.377.715.548.114.125.288      Antistreptolysin
-	D12.776.377.715.548.114.134      Antibodies, Bispecific
-	D12.776.377.715.548.114.143      Antibodies, Blocking

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.377.715.548.114.167      Antibodies, Catalytic
-	D12.776.377.715.548.114.179      Antibodies, Fungal
-	D12.776.377.715.548.114.185      Antibodies, Helminth
-	D12.776.377.715.548.114.191      Antibodies, Heterophile
-	D12.776.377.715.548.114.207      Antibodies, Immobilized
-	D12.776.377.715.548.114.224      Antibodies, Monoclonal
-	D12.776.377.715.548.114.224.200      Antibodies, Monoclonal, Humanized
-	D12.776.377.715.548.114.224.200.250      Adalimumab
-	D12.776.377.715.548.114.224.200.500      Certolizumab Pegol
-	D12.776.377.715.548.114.224.200.750      Cetuximab
-	D12.776.377.715.548.114.224.200.782      Denosumab
-	D12.776.377.715.548.114.224.200.813      Natalizumab
-	D12.776.377.715.548.114.224.200.844      Omalizumab
-	D12.776.377.715.548.114.224.200.860      Palivizumab
-	D12.776.377.715.548.114.224.200.868      Ranibizumab
-	D12.776.377.715.548.114.224.200.875      Trastuzumab
-	D12.776.377.715.548.114.224.200.937      Ustekinumab
-	D12.776.377.715.548.114.224.284      Antibodies, Monoclonal, Murine-Derived
-	D12.776.377.715.548.114.224.284.570      Muromonab-CD3
-	D12.776.377.715.548.114.224.284.785      Rituximab
-	D12.776.377.715.548.114.224.642      Infliximab
-	D12.776.377.715.548.114.240      Antibodies, Neoplasm
-	D12.776.377.715.548.114.244      Antibodies, Neutralizing
-	D12.776.377.715.548.114.248      Antibodies, Phospho-Specific
-	D12.776.377.715.548.114.252      Antibodies, Protozoan
-	D12.776.377.715.548.114.254      Antibodies, Viral
-	D12.776.377.715.548.114.254.150      Deltaretrovirus Antibodies
-	D12.776.377.715.548.114.254.150.440      HIV Antibodies
-	D12.776.377.715.548.114.254.150.500      HTLV-I Antibodies
-	D12.776.377.715.548.114.254.150.510      HTLV-II Antibodies
-	D12.776.377.715.548.114.254.450      Hepatitis Antibodies
-	D12.776.377.715.548.114.254.450.251      Hepatitis A Antibodies
-	D12.776.377.715.548.114.254.450.504      Hepatitis B Antibodies
-	D12.776.377.715.548.114.254.450.510      Hepatitis C Antibodies
-	D12.776.377.715.548.114.257      Antigen-Antibody Complex
-	D12.776.377.715.548.114.323      Autoantibodies

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.377.715.548.114.323.190      Antibodies, Antineutrophil Cytoplasmic
-	D12.776.377.715.548.114.323.204      Antibodies, Antinuclear
-	D12.776.377.715.548.114.323.210      Antibodies, Antiphospholipid
-	D12.776.377.715.548.114.323.210.100      Antibodies, Anticardiolipin
-	D12.776.377.715.548.114.323.210.600      Lupus Coagulation Inhibitor
-	D12.776.377.715.548.114.323.300      Complement C3 Nephritic Factor
-	D12.776.377.715.548.114.323.390      Immunoconglutinins
-	D12.776.377.715.548.114.323.480      Immunoglobulins, Thyroid-Stimulating
-	D12.776.377.715.548.114.323.480.500      Long-Acting Thyroid Stimulator
-	D12.776.377.715.548.114.323.732      Rheumatoid Factor
-	D12.776.377.715.548.114.573      Immune Sera
-	D12.776.377.715.548.114.573.203      Antilymphocyte Serum
-	D12.776.377.715.548.114.573.601      Antitoxins
-	D12.776.377.715.548.114.573.601.138      Antivenins
-	D12.776.377.715.548.114.573.601.268      Botulinum Antitoxin
-	D12.776.377.715.548.114.573.601.438      Diphtheria Antitoxin
-	D12.776.377.715.548.114.573.601.849      Tetanus Antitoxin
-	D12.776.377.715.548.114.580      Immunoconjugates
-	D12.776.377.715.548.114.580.225      Abatacept
-	D12.776.377.715.548.114.580.450      Immunotoxins
-	D12.776.377.715.548.114.606      Immunoglobulin Allotypes
-	D12.776.377.715.548.114.606.586      Immunoglobulin Gm Allotypes
-	D12.776.377.715.548.114.606.587      Immunoglobulin Km Allotypes
-	D12.776.377.715.548.114.619      Immunoglobulin Isotypes
-	D12.776.377.715.548.114.619.026      Immunoglobulin A
-	D12.776.377.715.548.114.619.026.030      Immunoglobulin A, Secretory
-	D12.776.377.715.548.114.619.026.030.500      Secretory Component
-	D12.776.377.715.548.114.619.026.515      Immunoglobulin alpha-Chains
-	D12.776.377.715.548.114.619.251      Immunoglobulin D
-	D12.776.377.715.548.114.619.251.500      Immunoglobulin delta-Chains
-	D12.776.377.715.548.114.619.312      Immunoglobulin E
-	D12.776.377.715.548.114.619.312.500      Immunoglobulin epsilon-Chains
-	D12.776.377.715.548.114.619.393      Immunoglobulin G
-	D12.776.377.715.548.114.619.393.261      Etanercept
-	D12.776.377.715.548.114.619.393.522      Immunoglobulin gamma-Chains
-	D12.776.377.715.548.114.619.393.522.400      Immunoglobulin Gm Allotypes

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.377.715.548.114.619.393.550 Long-Acting Thyroid Stimulator
-	D12.776.377.715.548.114.619.393.570 Muromonab-CD3
-	D12.776.377.715.548.114.619.393.700 Rho(D) Immune Globulin
-	D12.776.377.715.548.114.619.574 Immunoglobulin M
-	D12.776.377.715.548.114.619.574.500 Immunoglobulin mu-Chains
-	D12.776.377.715.548.114.632 Immunoglobulins, Intravenous
-	D12.776.377.715.548.114.656 Insulin Antibodies
-	D12.776.377.715.548.114.664 Isoantibodies
-	D12.776.377.715.548.114.715 Oligoclonal Bands
-	D12.776.377.715.548.114.767 Opsonin Proteins
-	D12.776.377.715.548.114.820 Plantibodies
-	D12.776.377.715.548.114.837 Precipitins
-	D12.776.377.715.548.114.890 Reagins
-	D12.776.377.715.548.397 gamma-Globulins
-	D12.776.377.715.548.397.500 Tuftsin
-	D12.776.377.715.548.538 Immunoglobulin Constant Regions
-	D12.776.377.715.548.538.500 Immunoglobulin Fc Fragments
-	D12.776.377.715.548.538.500.249 CD4 Immunoadhesins
-	D12.776.377.715.548.680 Immunoglobulin Fragments
-	D12.776.377.715.548.680.650 Immunoglobulin Fab Fragments
-	D12.776.377.715.548.680.650.250 Certolizumab Pegol
-	D12.776.377.715.548.680.650.500 Immunoglobulin Variable Region
-	D12.776.377.715.548.680.650.500.180 Complementarity Determining Regions
-	D12.776.377.715.548.680.650.500.590 Immunoglobulin Joining Region
-	D12.776.377.715.548.680.650.500.795 Single-Chain Antibodies
-	D12.776.377.715.548.680.650.500.897 Single-Domain Antibodies
-	D12.776.377.715.548.680.650.750 Tuftsin
-	D12.776.377.715.548.680.660 Immunoglobulin Fc Fragments
-	D12.776.377.715.548.680.660.249 CD4 Immunoadhesins
-	D12.776.377.715.548.680.745 Immunoglobulin Idiotypes
-	D12.776.377.715.548.705 Immunoglobulin Subunits
-	D12.776.377.715.548.705.500 Immunoglobulin Heavy Chains
-	D12.776.377.715.548.705.500.350 Immunoglobulin alpha-Chains
-	D12.776.377.715.548.705.500.360 Immunoglobulin delta-Chains
-	D12.776.377.715.548.705.500.370 Immunoglobulin epsilon-Chains
-	D12.776.377.715.548.705.500.380 Immunoglobulin gamma-Chains

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D12.776.377.715.548.705.500.380.500	Immunoglobulin Gm Allotypes
-	D12.776.377.715.548.705.500.500	Immunoglobulin mu-Chains
-	D12.776.377.715.548.705.625	Immunoglobulin J-Chains
-	D12.776.377.715.548.705.750	Immunoglobulin Light Chains
-	D12.776.377.715.548.705.750.530	Immunoglobulin kappa-Chains
-	D12.776.377.715.548.705.750.530.500	Immunoglobulin Km Allotypes
-	D12.776.377.715.548.705.750.550	Immunoglobulin lambda-Chains
-	D12.776.377.715.548.705.750.775	Immunoglobulin Light Chains, Surrogate
-	D12.776.377.715.548.705.875	Secretory Component
-	D12.776.377.715.548.797	Immunoglobulin Variable Region
-	D12.776.377.715.548.797.180	Complementarity Determining Regions
-	D12.776.377.715.548.797.590	Immunoglobulin Joining Region
-	D12.776.377.715.548.900	Paraproteins
-	D12.776.377.715.548.900.120	Bence Jones Protein
-	D12.776.377.715.548.900.225	Cryoglobulins
-	D12.776.377.715.548.900.500	Myeloma Proteins
-	D12.776.377.715.548.900.700	Pyroglobulins
-	D12.776.377.715.548.950	Receptors, Antigen, B-Cell
-	D12.776.377.715.548.950.500	Antigens, CD79
-	D12.776.377.715.548.950.750	Pre-B Cell Receptors
-	D12.776.377.715.548.950.750.500	Immunoglobulin Light Chains, Surrogate
-	D12.776.377.715.647	Macroglobulins
-	D12.776.377.715.647.100	alpha-Macroglobulins
-	D12.776.377.715.900	Transcobalamins
-	D12.776.377.856	Thyroglobulin
-	D12.776.395	Glycoproteins
-	D12.776.395.022	Activins
-	D12.776.395.022.500	Inhibin-beta Subunits
-	D12.776.395.033	ADAM Proteins
New Heading	<b>D12.776.395.033.125</b>	<b>ADAM12 Protein</b>
New Heading	<b>D12.776.395.033.250</b>	<b>ADAM10 Protein</b>
New Heading	<b>D12.776.395.033.375</b>	<b>ADAM17 Protein</b>
New Heading	<b>D12.776.395.033.500</b>	<b>ADAMTS Proteins</b>

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Heading	<b>D12.776.395.033.500.500</b>	<b>ADAMTS1 Protein</b>
New Heading	<b>D12.776.395.033.500.813</b>	<b>ADAMTS13 Protein</b>
New Heading	<b>D12.776.395.033.500.844</b>	<b>ADAMTS4 Protein</b>
New Heading	<b>D12.776.395.033.500.875</b>	<b>ADAMTS5 Protein</b>
New Heading	<b>D12.776.395.033.500.937</b>	<b>ADAMTS7 Protein</b>
New Heading	<b>D12.776.395.033.500.968</b>	<b>ADAMTS9 Protein</b>
New Heading	<b>D12.776.395.033.750</b>	<b>Fertilins</b>
-	D12.776.395.045	alpha 1-Antichymotrypsin
-	D12.776.395.068	alpha 1-Antitrypsin
-	D12.776.395.080	alpha-2-Antiplasmin
-	D12.776.395.086	alpha-2-HS-Glycoprotein
-	D12.776.395.092	Antigens, CD70
-	D12.776.395.140	Asialoglycoproteins
-	D12.776.395.175	Avidin
-	D12.776.395.191	B-Cell Activating Factor
-	D12.776.395.192	4-1BB Ligand
-	D12.776.395.195	beta 2-Glycoprotein I
-	D12.776.395.199	Cholesterol Ester Transfer Proteins
-	D12.776.395.207	Clusterin
-	D12.776.395.240	Colony-Stimulating Factors
-	D12.776.395.240.150	Erythropoietin
-	D12.776.395.240.150.500	Darbepoetin alfa
-	D12.776.395.240.150.750	Epoetin Alfa
-	D12.776.395.240.200	Granulocyte Colony-Stimulating Factor
-	D12.776.395.240.200.500	Filgrastim
-	D12.776.395.240.300	Granulocyte-Macrophage Colony-Stimulating Factor
-	D12.776.395.240.400	Interleukin-3
-	D12.776.395.240.500	Macrophage Colony-Stimulating Factor
-	D12.776.395.240.750	Thrombopoietin
-	D12.776.395.320	Complement C1 Inhibitor Protein
New Heading	<b>D12.776.395.341</b>	<b>Fibrillins</b>

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Heading	<b>D12.776.395.341.500</b>	<b>Fibrillin-1</b>
New Heading	<b>D12.776.395.341.750</b>	<b>Fibrillin-2</b>
New Heading	<b>D12.776.395.361</b>	<b>Glycodelin</b>
-	D12.776.395.401	Hemopexin
-	D12.776.395.420	HSP47 Heat-Shock Proteins
-	D12.776.395.439	Inhibins
-	D12.776.395.439.500	Inhibin-beta Subunits
-	D12.776.395.507	Lactoferrin
-	D12.776.395.550	Membrane Glycoproteins
-	D12.776.395.550.014	Antigens, CD47
-	D12.776.395.550.016	Antigens, CD82
-	D12.776.395.550.018	Antigens, CD9
-	D12.776.395.550.020	ATP-Binding Cassette Transporters
-	D12.776.395.550.020.304	ATP Binding Cassette Transporter 1
New Heading	<b>D12.776.395.550.020.457 G</b>	<b>ATP Binding Cassette Transporter, Sub-Family G</b>
New Heading	<b>D12.776.395.550.020.457.250 Family G, Member 1</b>	<b>ATP Binding Cassette Transporter, Sub-Family G, Member 1</b>
New Heading	<b>D12.776.395.550.020.457.500 Family G, Member 2</b>	<b>ATP Binding Cassette Transporter, Sub-Family G, Member 2</b>
New Heading	<b>D12.776.395.550.020.457.750 Family G, Member 5</b>	<b>ATP Binding Cassette Transporter, Sub-Family G, Member 5</b>
New Heading	<b>D12.776.395.550.020.457.875 Family G, Member 8</b>	<b>ATP Binding Cassette Transporter, Sub-Family G, Member 8</b>
-	D12.776.395.550.020.610	P-Glycoproteins
New Heading	<b>D12.776.395.550.020.610.305</b>	<b>Antigen Peptide Transporter-1</b>
New Heading	<b>D12.776.395.550.020.610.458</b>	<b>Antigen Peptide Transporter-2</b>
-	D12.776.395.550.020.610.610	P-Glycoprotein
-	D12.776.395.550.034	Antigens, CD58
-	D12.776.395.550.045	Antigens, CD147
-	D12.776.395.550.050	Antigens, Thy-1
New Heading	<b>D12.776.395.550.114</b>	<b>Butyrophilins</b>
New Tree	<a href="#">D12.776.395.550.114.500</a>	<a href="#">Myelin-Oligodendrocyte Glycoprotein</a>

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D12.776.395.550.177	CD30 Ligand
-	D12.776.395.550.185	CD40 Ligand
-	D12.776.395.550.200	Cell Adhesion Molecules
New Heading	<b>D12.776.395.550.200.049</b>	<b>Antigens, CD99</b>
-	D12.776.395.550.200.098	Antigens, CD24
-	D12.776.395.550.200.131	Antigens, CD31
-	D12.776.395.550.200.170	Antigens, CD146
-	D12.776.395.550.200.175	Antigens, CD164
New Heading	<b>D12.776.395.550.200.188</b>	<b>Cadherin Related Proteins</b>
New Tree	<a href="#">D12.776.395.550.200.188.500</a>	<a href="#">Proto-Oncogene Proteins c-ret</a>
-	D12.776.395.550.200.200	Cadherins
-	D12.776.395.550.200.200.500	Desmosomal Cadherins
-	D12.776.395.550.200.200.500.249	Desmocollins
-	D12.776.395.550.200.200.500.500	Desmogleins
-	D12.776.395.550.200.200.500.500.500	Desmoglein 1
-	D12.776.395.550.200.200.500.500.625	Desmoglein 2
-	D12.776.395.550.200.200.500.500.750	Desmoglein 3
-	D12.776.395.550.200.210	Carcinoembryonic Antigen
-	D12.776.395.550.200.230	CD4 Immunoadhesins
-	D12.776.395.550.200.250	Cell Adhesion Molecules, Neuronal
-	D12.776.395.550.200.250.150	Cell Adhesion Molecules, Neuron-Glia
-	D12.776.395.550.200.250.150.050 Molecule	Activated-Leukocyte Cell Adhesion
-	D12.776.395.550.200.250.325	Contactins
-	D12.776.395.550.200.250.325.100	Contactin 1
-	D12.776.395.550.200.250.325.200	Contactin 2
-	D12.776.395.550.200.250.500	Myelin P0 Protein
-	D12.776.395.550.200.250.520	Neural Cell Adhesion Molecules
-	D12.776.395.550.200.250.520.156	Antigens, CD56
-	D12.776.395.550.200.250.520.578	Neural Cell Adhesion Molecule L1
-	D12.776.395.550.200.250.520.789	Neurocan
New Heading	<b>D12.776.395.550.200.263</b>	<b>Epithelial Cell Adhesion Molecule</b>
-	D12.776.395.550.200.275	Integrin alphaXbeta2



## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D12.776.395.550.200.275.500	Antigens, CD11c
-	D12.776.395.550.200.275.750	Antigens, CD18
-	D12.776.395.550.200.450	Intercellular Adhesion Molecule-1
-	D12.776.395.550.200.537	Junctional Adhesion Molecules
-	D12.776.395.550.200.537.500 Membrane Protein	Coxsackie and Adenovirus Receptor-Like
-	D12.776.395.550.200.537.750	Junctional Adhesion Molecule A
-	D12.776.395.550.200.537.875	Junctional Adhesion Molecule B
-	D12.776.395.550.200.537.937	Junctional Adhesion Molecule C
-	D12.776.395.550.200.625	Receptors, Lymphocyte Homing
-	D12.776.395.550.200.625.144	Antigens, CD44
-	D12.776.395.550.200.625.347	Integrin alpha4beta1
-	D12.776.395.550.200.625.550	Lymphocyte Function-Associated Antigen-1
-	D12.776.395.550.200.625.903	L-Selectin
-	D12.776.395.550.200.700	Selectins
-	D12.776.395.550.200.700.300	E-Selectin
-	D12.776.395.550.200.700.510	L-Selectin
-	D12.776.395.550.200.700.775	P-Selectin
-	D12.776.395.550.200.810	Sialic Acid Binding Ig-like Lectin 2
-	D12.776.395.550.200.920	Vascular Cell Adhesion Molecule-1
-	D12.776.395.550.275	Ectodysplasins
-	D12.776.395.550.312	Fas Ligand Protein
New Heading	<b>D12.776.395.550.331</b>	<b>Fertilins</b>
-	D12.776.395.550.350	Fibronectins
-	D12.776.395.550.400	GAP-43 Protein
-	D12.776.395.550.448	GPI-Linked Proteins
-	D12.776.395.550.448.100	Antigens, CD14
-	D12.776.395.550.448.120	Antigens, CD24
-	D12.776.395.550.448.130	Antigens, CD55
-	D12.776.395.550.448.140	Antigens, CD59
-	D12.776.395.550.448.160	Carbonic Anhydrase IV
New Heading	<b>D12.776.395.550.448.180</b>	<b>CD48 Antigen</b>
-	D12.776.395.550.448.200	Ciliary Neurotrophic Factor Receptor alpha Subunit
-	D12.776.395.550.448.250	Contactins
-	D12.776.395.550.448.250.100	Contactin 1

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D12.776.395.550.448.250.200	Contactin 2
-	D12.776.395.550.448.300	Ephrin-A1
-	D12.776.395.550.448.310	Ephrin-A2
-	D12.776.395.550.448.320	Ephrin-A3
-	D12.776.395.550.448.330	Ephrin-A4
-	D12.776.395.550.448.340	Ephrin-A5
-	D12.776.395.550.448.370	Folate Receptors, GPI-Anchored
-	D12.776.395.550.448.370.500	Folate Receptor 1
-	D12.776.395.550.448.370.750	Folate Receptor 2
-	D12.776.395.550.448.400 Receptors	Glial Cell Line-Derived Neurotrophic Factor
-	D12.776.395.550.448.500	Glypicans
New Heading	<b>D12.776.395.550.448.600</b>	<b>Prion Proteins</b>
-	D12.776.395.550.448.700	Matrix Metalloproteinase 17
New Heading	<b>D12.776.395.550.448.738</b>	<b>Nogo Receptors</b>
New Heading	<b>D12.776.395.550.448.738.250</b>	<b>Nogo Receptor 2</b>
New Heading	<b>D12.776.395.550.448.738.500</b>	<b>Nogo Receptor 1</b>
-	D12.776.395.550.448.775	Oligodendrocyte-Myelin Glycoprotein
-	D12.776.395.550.448.850	Uromodulin
New Heading	<b>D12.776.395.550.469</b>	<b>Hepatitis A Virus Cellular Receptor 1</b>
New Heading	<b>D12.776.395.550.479</b>	<b>Hepatitis A Virus Cellular Receptor 2</b>
-	D12.776.395.550.489	Histocompatibility Antigens Class I
New Heading	<b>D12.776.395.550.489.200</b>	<b>Hemochromatosis Protein</b>
-	D12.776.395.550.489.400	HLA-A Antigens
-	D12.776.395.550.489.400.010	HLA-A1 Antigen
-	D12.776.395.550.489.400.020	HLA-A2 Antigen
-	D12.776.395.550.489.400.030	HLA-A3 Antigen
-	D12.776.395.550.489.400.110	HLA-A11 Antigen
-	D12.776.395.550.489.400.240	HLA-A24 Antigen
-	D12.776.395.550.489.500	HLA-B Antigens
-	D12.776.395.550.489.500.070	HLA-B7 Antigen
-	D12.776.395.550.489.500.080	HLA-B8 Antigen

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.395.550.489.500.130 HLA-B13 Antigen
-	D12.776.395.550.489.500.140 HLA-B14 Antigen
-	D12.776.395.550.489.500.150 HLA-B15 Antigen
-	D12.776.395.550.489.500.180 HLA-B18 Antigen
-	D12.776.395.550.489.500.270 HLA-B27 Antigen
-	D12.776.395.550.489.500.350 HLA-B35 Antigen
-	D12.776.395.550.489.500.370 HLA-B37 Antigen
-	D12.776.395.550.489.500.380 HLA-B38 Antigen
-	D12.776.395.550.489.500.390 HLA-B39 Antigen
-	D12.776.395.550.489.500.400 HLA-B40 Antigen
-	D12.776.395.550.489.500.440 HLA-B44 Antigen
-	D12.776.395.550.489.500.510 HLA-B51 Antigen
-	D12.776.395.550.489.500.520 HLA-B52 Antigen
-	D12.776.395.550.489.600 HLA-C Antigens
-	D12.776.395.550.489.700 HLA-G Antigens
-	D12.776.395.550.509 Histocompatibility Antigens Class II
-	D12.776.395.550.509.400 HLA-D Antigens
-	D12.776.395.550.509.400.420 HLA-DP Antigens
-	D12.776.395.550.509.400.420.500 HLA-DP alpha-Chains
-	D12.776.395.550.509.400.420.750 HLA-DP beta-Chains
-	D12.776.395.550.509.400.430 HLA-DQ Antigens
-	D12.776.395.550.509.400.430.500 HLA-DQ alpha-Chains
-	D12.776.395.550.509.400.430.750 HLA-DQ beta-Chains
-	D12.776.395.550.509.400.440 HLA-DR Antigens
-	D12.776.395.550.509.400.440.100 HLA-DR alpha-Chains
-	D12.776.395.550.509.400.440.200 HLA-DR beta-Chains
-	D12.776.395.550.509.400.440.200.010 HLA-DRB1 Chains
-	D12.776.395.550.509.400.440.200.030 HLA-DRB3 Chains
-	D12.776.395.550.509.400.440.200.040 HLA-DRB4 Chains
-	D12.776.395.550.509.400.440.200.050 HLA-DRB5 Chains
-	D12.776.395.550.509.400.440.400 HLA-DR Serological Subtypes
-	D12.776.395.550.509.400.440.400.010 HLA-DR1 Antigen
-	D12.776.395.550.509.400.440.400.020 HLA-DR2 Antigen
-	D12.776.395.550.509.400.440.400.030 HLA-DR3 Antigen
-	D12.776.395.550.509.400.440.400.040 HLA-DR4 Antigen
-	D12.776.395.550.509.400.440.400.050 HLA-DR5 Antigen

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D12.776.395.550.509.400.440.400.060	HLA-DR6 Antigen
-	D12.776.395.550.509.400.440.400.070	HLA-DR7 Antigen
-	D12.776.395.550.530	Laminin
-	D12.776.395.550.550	Lysosome-Associated Membrane Glycoproteins
-	D12.776.395.550.550.249	Antigens, CD63
-	D12.776.395.550.550.500	Lysosomal-Associated Membrane Protein 1
-	D12.776.395.550.550.750	Lysosomal-Associated Membrane Protein 2
-	D12.776.395.550.550.875	Lysosomal-Associated Membrane Protein 3
-	D12.776.395.550.560	Mucin-1
-	D12.776.395.550.565	Mucin-3
-	D12.776.395.550.570	Myelin-Associated Glycoprotein
Old Tree	D12.776.395.550.583	Myelin-Oligodendrocyte Glycoprotein
-	D12.776.395.550.597	OX40 Ligand
-	D12.776.395.550.611	PHEX Phosphate Regulating Neutral Endopeptidase
-	D12.776.395.550.625	Platelet Membrane Glycoproteins
-	D12.776.395.550.625.136	Antigens, CD36
-	D12.776.395.550.625.298	Integrin alpha2beta1
-	D12.776.395.550.625.379	Integrin alpha5beta1
-	D12.776.395.550.625.419	Integrin alpha6beta1
-	D12.776.395.550.625.439	Integrin alphaVbeta3
-	D12.776.395.550.625.449	Lysosomal-Associated Membrane Protein 1
-	D12.776.395.550.625.460	Platelet Glycoprotein GPIb-IX Complex
-	D12.776.395.550.625.785	Platelet Glycoprotein GPIIb-IIIa Complex
-	D12.776.395.550.625.800	Receptors, Thrombin
-	D12.776.395.550.625.800.790	Receptor, PAR-1
-	D12.776.395.550.625.800.800	Thrombomodulin
-	D12.776.395.550.625.905	P-Selectin
New Heading	<b>D12.776.395.550.736</b>	<b>Signaling Lymphocytic Activation Molecule Family</b>
New Heading	<b>D12.776.395.550.736.250</b>	<b>CD48 Antigen</b>
New Heading	<b>D12.776.395.550.736.500</b> <b>Family Member 1</b>	<b>Signaling Lymphocytic Activation Molecule</b>
-	D12.776.395.550.847	Syndecans
-	D12.776.395.550.847.100	Syndecan-1
-	D12.776.395.550.847.200	Syndecan-2
-	D12.776.395.550.847.600	Syndecan-3

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.395.550.847.800      Syndecan-4
-	D12.776.395.550.895      Thrombospondins
-	D12.776.395.550.895.800      Thrombospondin 1
-	D12.776.395.550.942      Uroplakins
-	D12.776.395.550.942.100      Uroplakin Ia
-	D12.776.395.550.942.150      Uroplakin Ib
-	D12.776.395.550.942.200      Uroplakin II
-	D12.776.395.550.942.300      Uroplakin III
-	D12.776.395.550.990      Variant Surface Glycoproteins, Trypanosoma
New Heading	<b>D12.776.395.550.995      Zona Pellucida Glycoproteins</b>
-	D12.776.395.560      Mucoproteins
-	D12.776.395.560.186      Cell Wall Skeleton
-	D12.776.395.560.373      Haptoglobins
-	D12.776.395.560.494      Intrinsic Factor
-	D12.776.395.560.631      Mucins
New Tree	<b>D12.776.395.560.631.050      CA-125 Antigen</b>
-	D12.776.395.560.631.100      Gastric Mucins
-	D12.776.395.560.631.100.500      Mucin 5AC
-	D12.776.395.560.631.100.750      Mucin-6
-	D12.776.395.560.631.115      Mucin-1
-	D12.776.395.560.631.161      Mucin-2
-	D12.776.395.560.631.184      Mucin-3
-	D12.776.395.560.631.195      Mucin-4
-	D12.776.395.560.631.207      Mucin-5B
-	D12.776.395.560.631.650      Sialomucins
-	D12.776.395.560.631.650.143      Antigens, CD43
-	D12.776.395.560.631.650.264      Antigens, CD164
-	D12.776.395.560.742      Orosomucoid
-	D12.776.395.560.760      Ovomucin
-	D12.776.395.560.800      Peptidoglycan
-	D12.776.395.560.825      Phytohemagglutinins
-	D12.776.395.560.912      Uromodulin
-	D12.776.395.570      Myelin P0 Protein
-	D12.776.395.600      Osteonectin

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D12.776.395.635	Protein C
-	D12.776.395.642	Protein S
-	D12.776.395.650	Proteoglycans
-	D12.776.395.650.350	Heparan Sulfate Proteoglycans
-	D12.776.395.650.350.249	Glypicans
-	D12.776.395.650.350.500	Syndecans
-	D12.776.395.650.350.500.100	Syndecan-1
-	D12.776.395.650.350.500.200	Syndecan-2
-	D12.776.395.650.350.500.600	Syndecan-3
-	D12.776.395.650.350.500.800	Syndecan-4
-	D12.776.395.650.750	Chondroitin Sulfate Proteoglycans
New Tree	<a href="#">D12.776.395.650.750.281</a>	<a href="#">Antigens, CD44</a>
-	D12.776.395.650.750.562	Biglycan
-	D12.776.395.650.750.625	Decorin
-	D12.776.395.650.750.687	Hyalectins
-	D12.776.395.650.750.687.100	Aggrecans
-	D12.776.395.650.750.687.275	Brevican
-	D12.776.395.650.750.687.450	Neurocan
-	D12.776.395.650.750.687.800	Versicans
New Heading	<b>D12.776.395.650.875</b>	<b>Small Leucine-Rich Proteoglycans</b>
New Tree	<a href="#">D12.776.395.650.875.250</a>	<a href="#">Biglycan</a>
New Tree	<a href="#">D12.776.395.650.875.500</a>	<a href="#">Decorin</a>
New Heading	<b>D12.776.395.650.875.625</b>	<b>Fibromodulin</b>
New Heading	<b>D12.776.395.650.875.750</b>	<b>Lumican</b>
-	D12.776.395.690	Serum Amyloid P-Component
-	D12.776.395.700	Sialoglycoproteins
-	D12.776.395.700.174	Antigens, CD43
-	D12.776.395.700.350	Glycophorin
-	D12.776.395.700.675	Integrin-Binding Sialoprotein
-	D12.776.395.700.837	Osteopontin
-	D12.776.395.768	Thyroglobulin
-	D12.776.395.832	Thyroxine-Binding Proteins

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.395.832.500 Thyroxine-Binding Globulin
-	D12.776.395.901 Transcortin
-	D12.776.395.930 Tumor Necrosis Factor-alpha
-	D12.776.395.970 Vitronectin
-	D12.776.419 Helminth Proteins
-	D12.776.419.500 Caenorhabditis elegans Proteins
-	D12.776.422 Heme proteins
-	D12.776.422.220 Cytochromes
-	D12.776.422.220.175 Cytochrome a Group
-	D12.776.422.220.175.249 Cytochromes a
-	D12.776.422.220.175.500 Cytochromes a1
-	D12.776.422.220.175.600 Cytochromes a3
-	D12.776.422.220.187 Cytochrome b Group
-	D12.776.422.220.187.249 Cytochromes b
-	D12.776.422.220.187.500 Cytochromes b5
-	D12.776.422.220.187.501 Cytochromes b6
-	D12.776.422.220.286 Cytochrome c Group
-	D12.776.422.220.286.100 Cytochromes c
-	D12.776.422.220.286.150 Cytochromes c'
-	D12.776.422.220.286.200 Cytochromes c1
-	D12.776.422.220.286.300 Cytochromes c2
-	D12.776.422.220.286.600 Cytochromes c6
-	D12.776.422.220.300 Cytochrome d Group
-	D12.776.422.220.453 Cytochrome P-450 Enzyme System
New Tree	<a href="#">D12.776.422.220.453.010</a> Aryl Hydrocarbon Hydroxylases
New Tree	<a href="#">D12.776.422.220.453.010.050</a> Aniline Hydroxylase
New Tree	<a href="#">D12.776.422.220.453.010.110</a> Benzopyrene Hydroxylase
New Tree	<a href="#">D12.776.422.220.453.010.332</a> Cytochrome P-450 CYP1A1
New Tree	<a href="#">D12.776.422.220.453.010.443</a> Cytochrome P-450 CYP1A2
New Tree	<a href="#">D12.776.422.220.453.010.500</a> Cytochrome P-450 CYP1B1
New Tree	<a href="#">D12.776.422.220.453.010.550</a> Cytochrome P-450 CYP2B1

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	D12.776.422.220.453.010.575	Cytochrome P-450 CYP2B6
New Tree	D12.776.422.220.453.010.590	Cytochrome P-450 CYP2C8
New Tree	D12.776.422.220.453.010.600	Cytochrome P-450 CYP2D6
New Tree	D12.776.422.220.453.012	Camphor 5-Monooxygenase
Old Tree	D12.776.422.220.453.040	Aryl Hydrocarbon Hydroxylases
Old Tree	D12.776.422.220.453.040.050	Aniline Hydroxylase
Old Tree	D12.776.422.220.453.040.110	Benzopyrene Hydroxylase
Old Tree	D12.776.422.220.453.040.332	Cytochrome P-450 CYP1A1
Old Tree	D12.776.422.220.453.040.443	Cytochrome P-450 CYP1A2
Old Tree	D12.776.422.220.453.040.500	Cytochrome P-450 CYP1B1
Old Tree	D12.776.422.220.453.040.550	Cytochrome P-450 CYP2B1
Old Tree	D12.776.422.220.453.040.575	Cytochrome P-450 CYP2B6
Old Tree	D12.776.422.220.453.040.590	Cytochrome P-450 CYP2C8
Old Tree	D12.776.422.220.453.040.600	Cytochrome P-450 CYP2D6
Old Tree	D12.776.422.220.453.085	Camphor 5-Monooxygenase
Old Tree	D12.776.422.220.453.100	Cytochrome P-450 CYP2A6
New Heading	<b>D12.776.422.220.453.100</b>	<b>Cytochrome P450 Family 1</b>
New Tree	D12.776.422.220.453.100.500	Cytochrome P-450 CYP1A1
New Tree	D12.776.422.220.453.100.750	Cytochrome P-450 CYP1A2
New Tree	D12.776.422.220.453.100.875	Cytochrome P-450 CYP1B1
Old Tree	D12.776.422.220.453.300	Cytochrome P-450 CYP2E1
Old Tree	D12.776.422.220.453.400	Cytochrome P-450 CYP3A
Old Tree	D12.776.422.220.453.450	Cytochrome P-450 CYP4A
New Heading	<b>D12.776.422.220.453.484</b>	<b>Cytochrome P450 Family 11</b>
New Tree	D12.776.422.220.453.484.250	Cholesterol Side-Chain Cleavage Enzyme
New Tree	D12.776.422.220.453.484.500	Cytochrome P-450 CYP11B2
New Tree	D12.776.422.220.453.484.750	Steroid 11-beta-Hydroxylase
New	<b>D12.776.422.220.453.485</b>	<b>Cytochrome P450 Family 12</b>



## MeSH Tree Changes for 2017

Type	Tree - heading	
Heading		
New Heading	<b>D12.776.422.220.453.487</b>	<b>Cytochrome P450 Family 17</b>
New Tree	D12.776.422.220.453.487.500	Steroid 17-alpha-Hydroxylase
New Heading	<b>D12.776.422.220.453.489</b>	<b>Cytochrome P450 Family 19</b>
New Tree	D12.776.422.220.453.489.500	Aromatase
New Heading	<b>D12.776.422.220.453.491</b>	<b>Cytochrome P450 Family 2</b>
New Tree	D12.776.422.220.453.491.250	Cytochrome P-450 CYP2A6
New Tree	D12.776.422.220.453.491.313	Cytochrome P-450 CYP2B1
New Tree	D12.776.422.220.453.491.344	Cytochrome P-450 CYP2B6
New Tree	D12.776.422.220.453.491.360	Cytochrome P-450 CYP2C8
New Tree	D12.776.422.220.453.491.368	Cytochrome P-450 CYP2D6
New Tree	D12.776.422.220.453.491.375	Cytochrome P-450 CYP2E1
New Tree	D12.776.422.220.453.491.500	Limonene Hydroxylases
New Tree	D12.776.422.220.453.491.500.500	Cytochrome P-450 CYP2C9
New Tree	D12.776.422.220.453.491.500.700	Cytochrome P-450 CYP2C19
New Heading	<b>D12.776.422.220.453.493</b>	<b>Cytochrome P450 Family 21</b>
New Tree	D12.776.422.220.453.493.500	Steroid 21-Hydroxylase
New Heading	<b>D12.776.422.220.453.496</b>	<b>Cytochrome P450 Family 24</b>
New Tree	D12.776.422.220.453.496.500	Vitamin D3 24-Hydroxylase
New Heading	<b>D12.776.422.220.453.498</b>	<b>Cytochrome P450 Family 26</b>
New Heading	<b>D12.776.422.220.453.498.500</b>	<b>Retinoic Acid 4-Hydroxylase</b>
New Heading	<b>D12.776.422.220.453.499</b>	<b>Cytochrome P450 Family 27</b>

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	D12.776.422.220.453.499.500	25-Hydroxyvitamin D3 1-alpha-Hydroxylase
New Tree	D12.776.422.220.453.499.750	Cholestanetriol 26-Monooxygenase
Old Tree	D12.776.422.220.453.500	Limonene Hydroxylases
Old Tree	D12.776.422.220.453.500.500	Cytochrome P-450 CYP2C9
Old Tree	D12.776.422.220.453.500.700	Cytochrome P-450 CYP2C19
New Heading	<b>D12.776.422.220.453.860</b>	<b>Cytochrome P450 Family 3</b>
New Tree	D12.776.422.220.453.860.500	Cytochrome P-450 CYP3A
New Heading	<b>D12.776.422.220.453.870</b>	<b>Cytochrome P450 Family 4</b>
New Tree	D12.776.422.220.453.870.500	Cytochrome P-450 CYP4A
New Heading	<b>D12.776.422.220.453.875</b>	<b>Cytochrome P450 Family 46</b>
New Heading	<b>D12.776.422.220.453.875.500</b>	<b>Cholesterol 24-Hydroxylase</b>
New Heading	<b>D12.776.422.220.453.878</b>	<b>Cytochrome P450 Family 51</b>
New Tree	D12.776.422.220.453.878.500	Sterol 14-Demethylase
New Heading	<b>D12.776.422.220.453.880</b>	<b>Cytochrome P450 Family 6</b>
New Heading	<b>D12.776.422.220.453.890</b>	<b>Cytochrome P450 Family 7</b>
New Tree	D12.776.422.220.453.890.500	Cholesterol 7-alpha-Hydroxylase
New Heading	<b>D12.776.422.220.453.900</b>	<b>Cytochrome P450 Family 8</b>
New Tree	D12.776.422.220.453.900.500	Steroid 12-alpha-Hydroxylase
-	D12.776.422.220.453.915	Steroid Hydroxylases
New Heading	<b>D12.776.422.220.453.915.025</b>	<b>Cholesterol 24-Hydroxylase</b>
-	D12.776.422.220.453.915.050	Cytochrome P-450 CYP11B2
-	D12.776.422.220.453.915.099	Aromatase
-	D12.776.422.220.453.915.150	Cholestanetriol 26-Monooxygenase
-	D12.776.422.220.453.915.200	Cholesterol 7-alpha-Hydroxylase
-	D12.776.422.220.453.915.212	Cholesterol Side-Chain Cleavage Enzyme
-	D12.776.422.220.453.915.400	25-Hydroxyvitamin D3 1-alpha-Hydroxylase

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D12.776.422.220.453.915.730	Steroid 12-alpha-Hydroxylase
-	D12.776.422.220.453.915.737	Steroid 16-alpha-Hydroxylase
-	D12.776.422.220.453.915.748	Steroid 17-alpha-Hydroxylase
-	D12.776.422.220.453.915.750	Steroid 11-beta-Hydroxylase
-	D12.776.422.220.453.915.760	Steroid 21-Hydroxylase
-	D12.776.422.220.453.915.880	Sterol 14-Demethylase
Old Tree	D12.776.422.220.453.957	Vitamin D3 24-Hydroxylase
-	D12.776.422.220.726	Cytochromes f
-	D12.776.422.316	Globins
-	D12.776.422.316.762	Hemoglobins
-	D12.776.422.316.762.149	Carboxyhemoglobin
-	D12.776.422.316.762.260	Erythrocrucorins
-	D12.776.422.316.762.320	Fetal Hemoglobin
-	D12.776.422.316.762.380	Hemoglobin A
-	D12.776.422.316.762.380.440	Hemoglobin A, Glycosylated
-	D12.776.422.316.762.380.450	Hemoglobin A2
-	D12.776.422.316.762.403	Hemoglobin Subunits
-	D12.776.422.316.762.403.320	alpha-Globins
-	D12.776.422.316.762.403.320.500	zeta-Globins
-	D12.776.422.316.762.403.325	beta-Globins
-	D12.776.422.316.762.403.325.249	delta-Globins
-	D12.776.422.316.762.403.325.374	epsilon-Globins
-	D12.776.422.316.762.403.325.500	gamma-Globins
-	D12.776.422.316.762.426	Hemoglobins, Abnormal
-	D12.776.422.316.762.426.338	Hemoglobin C
-	D12.776.422.316.762.426.375	Hemoglobin E
-	D12.776.422.316.762.426.463	Hemoglobin H
-	D12.776.422.316.762.426.480	Hemoglobin J
-	D12.776.422.316.762.426.510	Hemoglobin M
-	D12.776.422.316.762.426.588	Hemoglobin, Sickle
-	D12.776.422.316.762.571	Methemoglobin
-	D12.776.422.316.762.687	Oxyhemoglobins
-	D12.776.422.316.762.865	Sulfhemoglobin
-	D12.776.422.316.762.932	Truncated Hemoglobins
-	D12.776.422.316.881	Leghemoglobin
-	D12.776.422.316.940	Myoglobin

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.422.316.940.500 Metmyoglobin
-	D12.776.422.412 Hemocyanin
-	D12.776.422.680 Methemalbumin
New Heading	<b>D12.776.441 Huntingtin Protein</b>
-	D12.776.460 Immediate-Early Proteins
-	D12.776.460.050 Adenovirus E1 Proteins
-	D12.776.460.050.100 Adenovirus E1A Proteins
-	D12.776.460.050.110 Adenovirus E1B Proteins
-	D12.776.460.287 Butyrate Response Factor 1
-	D12.776.460.525 Early Growth Response Transcription Factors
-	D12.776.460.525.500 Early Growth Response Protein 1
-	D12.776.460.525.750 Early Growth Response Protein 2
-	D12.776.460.525.875 Early Growth Response Protein 3
-	D12.776.460.762 Tristetraprolin
-	D12.776.463 Immobilized Proteins
-	D12.776.463.250 Antibodies, Immobilized
-	D12.776.463.500 Enzymes, Immobilized
-	D12.776.467 Intercellular Signaling Peptides and Proteins
-	D12.776.467.024 Adipokines
-	D12.776.467.024.249 Adiponectin
-	D12.776.467.024.500 Leptin
-	D12.776.467.024.750 Resistin
-	D12.776.467.049 Agouti Signaling Protein
-	D12.776.467.074 Agouti-Related Protein
Old Tree	<b>D12.776.467.087 Alarmins</b>
Old Tree	<b>D12.776.467.087.125 alpha-Defensins</b>
Old Tree	<b>D12.776.467.087.188 beta-Defensins</b>
Old Tree	<b>D12.776.467.087.250 Calgranulin A</b>
Old Tree	<b>D12.776.467.087.375 Calgranulin B</b>
Old Tree	<b>D12.776.467.087.407 Cathelicidins</b>
Old Tree	<b>D12.776.467.087.438 Chaperonin 60</b>
Old Tree	<b>D12.776.467.087.500 HMGB1 Protein</b>
Old Tree	<b>D12.776.467.087.750 HSC70 Heat-Shock Proteins</b>
Old Tree	<b>D12.776.467.087.813 Interleukin-33</b>
Old Tree	<b>D12.776.467.087.875 S100 Calcium Binding Protein beta Subunit</b>

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	<b>D12.776.467.087.937</b> <b>S100A12 Protein</b>
-	D12.776.467.100 Angiogenic Proteins
-	D12.776.467.100.100 Angiopoietins
-	D12.776.467.100.100.100 Angiopoietin-1
-	D12.776.467.100.100.200 Angiopoietin-2
-	D12.776.467.100.450 Angiostatic Proteins
-	D12.776.467.100.450.500 Angiostatins
-	D12.776.467.100.450.750 Endostatins
-	D12.776.467.100.800 Vascular Endothelial Growth Factors
-	D12.776.467.100.800.200 Vascular Endothelial Growth Factor A
-	D12.776.467.100.800.300 Vascular Endothelial Growth Factor B
-	D12.776.467.100.800.400 Vascular Endothelial Growth Factor C
-	D12.776.467.100.800.500 Vascular Endothelial Growth Factor D
-	D12.776.467.100.800.600 Vascular Endothelial Growth Factor, Endocrine-Gland-Derived
-	D12.776.467.150 B7 Antigens
-	D12.776.467.150.100 Antigens, CD80
-	D12.776.467.150.200 Antigens, CD86
-	D12.776.467.150.300 Antigens, CD274
-	D12.776.467.150.500 Inducible T-Cell Co-Stimulator Ligand
-	D12.776.467.150.800 Programmed Cell Death 1 Ligand 2 Protein
-	D12.776.467.150.900 V-Set Domain-Containing T-Cell Activation Inhibitor 1
-	D12.776.467.200 CCN Intercellular Signaling Proteins
-	D12.776.467.200.100 Connective Tissue Growth Factor
-	D12.776.467.200.200 Cysteine-Rich Protein 61
-	D12.776.467.200.500 Nephroblastoma Overexpressed Protein
-	D12.776.467.374 Cytokines
-	D12.776.467.374.200 Chemokines
-	D12.776.467.374.200.070 beta-Thromboglobulin
-	D12.776.467.374.200.100 Chemokines, C
-	D12.776.467.374.200.110 Chemokines, CC
-	D12.776.467.374.200.110.050 Chemokine CCL1
-	D12.776.467.374.200.110.150 Chemokine CCL3
-	D12.776.467.374.200.110.200 Chemokine CCL4
-	D12.776.467.374.200.110.250 Chemokine CCL5
-	D12.776.467.374.200.110.550 Chemokine CCL11

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D12.776.467.374.200.110.850	Chemokine CCL17
-	D12.776.467.374.200.110.870	Chemokine CCL19
-	D12.776.467.374.200.110.880	Chemokine CCL20
-	D12.776.467.374.200.110.890	Chemokine CCL21
-	D12.776.467.374.200.110.900	Chemokine CCL22
-	D12.776.467.374.200.110.910	Chemokine CCL24
-	D12.776.467.374.200.110.930	Chemokine CCL27
-	D12.776.467.374.200.110.990	Monocyte Chemoattractant Proteins
-	D12.776.467.374.200.110.990.600	Chemokine CCL2
-	D12.776.467.374.200.110.990.800	Chemokine CCL7
-	D12.776.467.374.200.110.990.900	Chemokine CCL8
-	D12.776.467.374.200.120	Chemokines, CXC
-	D12.776.467.374.200.120.050	Chemokine CXCL1
-	D12.776.467.374.200.120.100	Chemokine CXCL2
-	D12.776.467.374.200.120.250	Chemokine CXCL5
-	D12.776.467.374.200.120.300	Chemokine CXCL6
-	D12.776.467.374.200.120.450	Chemokine CXCL9
-	D12.776.467.374.200.120.500	Chemokine CXCL10
-	D12.776.467.374.200.120.550	Chemokine CXCL11
-	D12.776.467.374.200.120.600	Chemokine CXCL12
-	D12.776.467.374.200.120.650	Chemokine CXCL13
-	D12.776.467.374.200.120.800	Interleukin-8
-	D12.776.467.374.200.120.900	Platelet Factor 4
-	D12.776.467.374.200.130	Chemokines, CX3C
-	D12.776.467.374.200.130.500	Chemokine CX3CL1
-	D12.776.467.374.200.600	Macrophage Inflammatory Proteins
-	D12.776.467.374.200.600.150	Chemokine CCL3
-	D12.776.467.374.200.600.200	Chemokine CCL4
-	D12.776.467.374.200.600.870	Chemokine CCL19
-	D12.776.467.374.200.600.880	Chemokine CCL20
-	D12.776.467.374.200.600.940	Chemokine CXCL2
-	D12.776.467.374.305	Growth Differentiation Factor 15
-	D12.776.467.374.410	Hematopoietic Cell Growth Factors
-	D12.776.467.374.410.240	Colony-Stimulating Factors
-	D12.776.467.374.410.240.150	Erythropoietin
-	D12.776.467.374.410.240.150.500	Epoetin Alfa

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.467.374.410.240.350 Granulocyte Colony-Stimulating Factor
-	D12.776.467.374.410.240.350.500 Filgrastim
-	D12.776.467.374.410.240.375 Granulocyte-Macrophage Colony-Stimulating Factor
-	D12.776.467.374.410.240.400 Interleukin-3
-	D12.776.467.374.410.240.500 Macrophage Colony-Stimulating Factor
-	D12.776.467.374.410.240.750 Thrombopoietin
-	D12.776.467.374.410.800 Stem Cell Factor
-	D12.776.467.374.420 Hepatocyte Growth Factor
-	D12.776.467.374.440 Interferons
-	D12.776.467.374.440.890 Interferon Type I
-	D12.776.467.374.440.890.250 Interferon-alpha
-	D12.776.467.374.440.890.275 Interferon-beta
-	D12.776.467.374.440.890.275.500 Interferon beta-1a
-	D12.776.467.374.440.890.275.750 Interferon beta-1b
-	D12.776.467.374.440.893 Interferon-gamma
-	D12.776.467.374.460 Interleukin 1 Receptor Antagonist Protein
-	D12.776.467.374.465 Interleukins
-	D12.776.467.374.465.010 Interleukin-1
-	D12.776.467.374.465.010.300 Interleukin-1alpha
-	D12.776.467.374.465.010.600 Interleukin-1beta
-	D12.776.467.374.465.021 Interleukin-2
-	D12.776.467.374.465.032 Interleukin-3
-	D12.776.467.374.465.178 Interleukin-4
-	D12.776.467.374.465.186 Interleukin-5
-	D12.776.467.374.465.202 Interleukin-6
-	D12.776.467.374.465.224 Interleukin-7
-	D12.776.467.374.465.246 Interleukin-8
-	D12.776.467.374.465.312 Interleukin-9
-	D12.776.467.374.465.510 Interleukin-10
-	D12.776.467.374.465.511 Interleukin-11
-	D12.776.467.374.465.512 Interleukin-12
-	D12.776.467.374.465.512.249 Interleukin-12 Subunit p35
-	D12.776.467.374.465.512.500 Interleukin-12 Subunit p40
-	D12.776.467.374.465.513 Interleukin-13
-	D12.776.467.374.465.515 Interleukin-15

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.467.374.465.516 Interleukin-16
-	D12.776.467.374.465.517 Interleukin-17
-	D12.776.467.374.465.518 Interleukin-18
-	D12.776.467.374.465.759 Interleukin-23
-	D12.776.467.374.465.759.249 Interleukin-12 Subunit p40
-	D12.776.467.374.465.759.500 Interleukin-23 Subunit p19
-	D12.776.467.374.465.800 Interleukin-27
-	D12.776.467.374.465.850 Interleukin-33
-	D12.776.467.374.470 Leukemia Inhibitory Factor
-	D12.776.467.374.480 Lymphokines
-	D12.776.467.374.480.372 Interleukin-2
-	D12.776.467.374.480.428 Leukocyte Migration-Inhibitory Factors
-	D12.776.467.374.480.438 Lymphotoxin-alpha
-	D12.776.467.374.480.615 Macrophage-Activating Factors
-	D12.776.467.374.480.615.350 Interferon-gamma
-	D12.776.467.374.480.625 Macrophage Migration-Inhibitory Factors
-	D12.776.467.374.480.700 Suppressor Factors, Immunologic
-	D12.776.467.374.480.750 Transfer Factor
-	D12.776.467.374.500 Monokines
-	D12.776.467.374.500.400 Interleukin-1
-	D12.776.467.374.500.400.300 Interleukin-1alpha
-	D12.776.467.374.500.400.600 Interleukin-1beta
-	D12.776.467.374.500.800 Tumor Necrosis Factor-alpha
-	D12.776.467.374.562 Oncostatin M
-	D12.776.467.374.625 Osteopontin
-	D12.776.467.374.687 Transforming Growth Factor beta
-	D12.776.467.374.687.100 Transforming Growth Factor beta1
-	D12.776.467.374.687.200 Transforming Growth Factor beta2
-	D12.776.467.374.687.300 Transforming Growth Factor beta3
-	D12.776.467.374.750 Tumor Necrosis Factors
-	D12.776.467.374.750.030 Antigens, CD70
-	D12.776.467.374.750.061 B-Cell Activating Factor
-	D12.776.467.374.750.065 4-1BB Ligand
-	D12.776.467.374.750.092 CD30 Ligand
-	D12.776.467.374.750.124 CD40 Ligand
-	D12.776.467.374.750.186 Ectodysplasins



## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.467.374.750.249 Fas Ligand Protein
-	D12.776.467.374.750.500 Lymphotoxin-alpha
-	D12.776.467.374.750.505 Lymphotoxin alpha1, beta2 Heterotrimer
-	D12.776.467.374.750.515 Lymphotoxin-beta
-	D12.776.467.374.750.531 OX40 Ligand
-	D12.776.467.374.750.562 RANK Ligand
-	D12.776.467.374.750.625 TNF-Related Apoptosis-Inducing Ligand
-	D12.776.467.374.750.626 Tumor Necrosis Factor-alpha
-	D12.776.467.374.750.656 Tumor Necrosis Factor Ligand Superfamily Member 13
-	D12.776.467.374.750.690 Tumor Necrosis Factor Ligand Superfamily Member 14
-	D12.776.467.374.750.710 Tumor Necrosis Factor Ligand Superfamily Member 15
-	D12.776.467.382 EGF Family of Proteins
-	D12.776.467.382.249 Amphiregulin
-	D12.776.467.382.374 Betacellulin
-	D12.776.467.382.500 Epidermal Growth Factor
-	D12.776.467.382.531 Epigen
-	D12.776.467.382.562 Epiregulin
-	D12.776.467.382.625 Heparin-binding EGF-like Growth Factor
-	D12.776.467.382.750 Transforming Growth Factor alpha
-	D12.776.467.390 Endothelial Growth Factors
-	D12.776.467.400 Endothelins
-	D12.776.467.400.225 Endothelin-1
-	D12.776.467.400.235 Endothelin-2
-	D12.776.467.400.245 Endothelin-3
-	D12.776.467.500 Ephrins
-	D12.776.467.500.100 Ephrin-A1
-	D12.776.467.500.200 Ephrin-A2
-	D12.776.467.500.300 Ephrin-A3
-	D12.776.467.500.400 Ephrin-A4
-	D12.776.467.500.500 Ephrin-A5
-	D12.776.467.500.600 Ephrin-B1
-	D12.776.467.500.700 Ephrin-B2
-	D12.776.467.500.800 Ephrin-B3
-	D12.776.467.624 Fibroblast Growth Factors

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.467.624.110      Fibroblast Growth Factor 1
-	D12.776.467.624.120      Fibroblast Growth Factor 2
-	D12.776.467.624.130      Fibroblast Growth Factor 3
-	D12.776.467.624.140      Fibroblast Growth Factor 4
-	D12.776.467.624.150      Fibroblast Growth Factor 5
-	D12.776.467.624.160      Fibroblast Growth Factor 6
-	D12.776.467.624.170      Fibroblast Growth Factor 7
-	D12.776.467.624.180      Fibroblast Growth Factor 8
-	D12.776.467.624.190      Fibroblast Growth Factor 9
-	D12.776.467.624.200      Fibroblast Growth Factor 10
-	D12.776.467.671            Hedgehog Proteins
-	D12.776.467.812            Kinins
-	D12.776.467.812.169        Bradykinin
-	D12.776.467.812.169.400    Kallidin
-	D12.776.467.812.654        Kininogens
-	D12.776.467.812.654.350    Kininogen, High-Molecular-Weight
-	D12.776.467.812.654.400    Kininogen, Low-Molecular-Weight
-	D12.776.467.812.900        Tachykinins
-	D12.776.467.812.900.354    Eledoisin
-	D12.776.467.812.900.475    Kassinin
-	D12.776.467.812.900.500    Neurokinin A
-	D12.776.467.812.900.550    Neurokinin B
-	D12.776.467.812.900.800    Physalaemin
-	D12.776.467.812.900.866    Substance P
New Tree	<a href="#">D12.776.467.836</a> <a href="#">Matrix Metalloproteinases, Secreted</a>
New Tree	<a href="#">D12.776.467.836.049</a> <a href="#">Bone Morphogenetic Protein 1</a>
New Tree	<a href="#">D12.776.467.836.100</a> <a href="#">Matrix Metalloproteinase 1</a>
New Tree	<a href="#">D12.776.467.836.150</a> <a href="#">Matrix Metalloproteinase 2</a>
New Tree	<a href="#">D12.776.467.836.200</a> <a href="#">Matrix Metalloproteinase 3</a>
New Tree	<a href="#">D12.776.467.836.250</a> <a href="#">Matrix Metalloproteinase 7</a>
New Tree	<a href="#">D12.776.467.836.300</a> <a href="#">Matrix Metalloproteinase 8</a>

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	<a href="#">D12.776.467.836.350</a>	<a href="#">Matrix Metalloproteinase 9</a>
New Tree	<a href="#">D12.776.467.836.400</a>	<a href="#">Matrix Metalloproteinase 10</a>
New Tree	<a href="#">D12.776.467.836.450</a>	<a href="#">Matrix Metalloproteinase 11</a>
New Tree	<a href="#">D12.776.467.836.500</a>	<a href="#">Matrix Metalloproteinase 12</a>
New Tree	<a href="#">D12.776.467.836.550</a>	<a href="#">Matrix Metalloproteinase 13</a>
New Tree	<a href="#">D12.776.467.836.800</a>	<a href="#">Matrix Metalloproteinase 20</a>
-	D12.776.467.860	Nerve Growth Factors
-	D12.776.467.860.100	Brain-Derived Neurotrophic Factor
-	D12.776.467.860.212	Ciliary Neurotrophic Factor
-	D12.776.467.860.325	Glia Maturation Factor
-	D12.776.467.860.381	Glial Cell Line-Derived Neurotrophic Factors
-	D12.776.467.860.381.500	Glial Cell Line-Derived Neurotrophic Factor
-	D12.776.467.860.381.750	Neurturin
-	D12.776.467.860.437	Nerve Growth Factor
-	D12.776.467.860.550	Neuregulins
-	D12.776.467.860.550.750	Neuregulin-1
-	D12.776.467.860.775	Neurotrophin 3
-	D12.776.467.860.887	Pituitary Adenylate Cyclase-Activating Polypeptide
-	D12.776.467.890	Parathyroid Hormone-Related Protein
-	D12.776.467.910	Platelet-Derived Growth Factor
-	D12.776.467.910.650	Proto-Oncogene Proteins c-sis
-	D12.776.467.923	Semaphorins
-	D12.776.467.923.374	Semaphorin-3A
-	D12.776.467.937	Somatomedins
-	D12.776.467.937.400	Insulin-Like Growth Factor I
-	D12.776.467.937.420	Insulin-Like Growth Factor II
-	D12.776.467.942	TGF-beta Superfamily Proteins
-	D12.776.467.942.200	Bone Morphogenetic Proteins
-	D12.776.467.942.200.100	Bone Morphogenetic Protein 1
-	D12.776.467.942.200.200	Bone Morphogenetic Protein 2
-	D12.776.467.942.200.300	Bone Morphogenetic Protein 3
-	D12.776.467.942.200.400	Bone Morphogenetic Protein 4

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D12.776.467.942.200.500	Bone Morphogenetic Protein 5
-	D12.776.467.942.200.600	Bone Morphogenetic Protein 6
-	D12.776.467.942.200.700	Bone Morphogenetic Protein 7
-	D12.776.467.942.200.900	Bone Morphogenetic Protein 15
-	D12.776.467.942.200.910	Growth Differentiation Factor 2
-	D12.776.467.942.200.950	Growth Differentiation Factor 10
-	D12.776.467.942.300	Growth Differentiation Factors
-	D12.776.467.942.300.049	Bone Morphogenetic Protein 15
-	D12.776.467.942.300.100	Growth Differentiation Factor 1
-	D12.776.467.942.300.200	Growth Differentiation Factor 2
-	D12.776.467.942.300.300	Growth Differentiation Factor 3
-	D12.776.467.942.300.500	Growth Differentiation Factor 5
-	D12.776.467.942.300.600	Growth Differentiation Factor 6
-	D12.776.467.942.300.800	Growth Differentiation Factor 9
-	D12.776.467.942.300.900	Growth Differentiation Factor 10
-	D12.776.467.942.300.915	Growth Differentiation Factor 15
-	D12.776.467.942.300.925	Myostatin
-	D12.776.467.942.550	Nodal Signaling Ligands
-	D12.776.467.942.550.100	Growth Differentiation Factor 1
-	D12.776.467.942.550.300	Growth Differentiation Factor 3
-	D12.776.467.942.550.475	Left-Right Determination Factors
-	D12.776.467.942.550.650	Nodal Protein
-	D12.776.467.942.775	Transforming Growth Factor beta
-	D12.776.467.942.775.100	Transforming Growth Factor beta1
-	D12.776.467.942.775.200	Transforming Growth Factor beta2
-	D12.776.467.942.775.300	Transforming Growth Factor beta3
-	D12.776.467.948	Tolloid-Like Metalloproteinases
-	D12.776.467.948.500	Bone Morphogenetic Protein 1
-	D12.776.467.960	Transforming Growth Factors
-	D12.776.467.960.360	Transforming Growth Factor alpha
-	D12.776.467.984	Wnt Proteins
New Heading	<b>D12.776.467.984.050</b>	<b>Wnt-5a Protein</b>
-	D12.776.467.984.100	Wnt1 Protein
-	D12.776.467.984.200	Wnt2 Protein
-	D12.776.467.984.300	Wnt3 Protein

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.467.984.350 Wnt3A Protein
-	D12.776.467.984.400 Wnt4 Protein
-	D12.776.476 Intracellular Signaling Peptides and Proteins
-	D12.776.476.008 3',5'-Cyclic-AMP Phosphodiesterases
-	D12.776.476.008.100 Cyclic Nucleotide Phosphodiesterases, Type 1
-	D12.776.476.008.200 Cyclic Nucleotide Phosphodiesterases, Type 2
-	D12.776.476.008.300 Cyclic Nucleotide Phosphodiesterases, Type 3
-	D12.776.476.008.400 Cyclic Nucleotide Phosphodiesterases, Type 4
-	D12.776.476.008.500 Cyclic Nucleotide Phosphodiesterases, Type 5
-	D12.776.476.008.600 Cyclic Nucleotide Phosphodiesterases, Type 6
-	D12.776.476.008.700 Cyclic Nucleotide Phosphodiesterases, Type 7
-	D12.776.476.009 3',5'-Cyclic-GMP Phosphodiesterases
-	D12.776.476.009.100 Cyclic Nucleotide Phosphodiesterases, Type 1
-	D12.776.476.009.200 Cyclic Nucleotide Phosphodiesterases, Type 2
-	D12.776.476.009.500 Cyclic Nucleotide Phosphodiesterases, Type 5
-	D12.776.476.009.750 Cyclic Nucleotide Phosphodiesterases, Type 6
-	D12.776.476.011 Activating Transcription Factor 6
-	D12.776.476.024 Adaptor Proteins, Signal Transducing
-	D12.776.476.024.034 5-Lipoxygenase-Activating Proteins
-	D12.776.476.024.050 14-3-3 Proteins
-	D12.776.476.024.069 A Kinase Anchor Proteins
New Tree	<a href="#">D12.776.476.024.104</a> <a href="#">Arrestins</a>
New Tree	<a href="#">D12.776.476.024.104.050</a> <a href="#">Arrestin</a>
New Heading	<b>D12.776.476.024.104.525</b> <b>Beta-Arrestins</b>
New Heading	<b>D12.776.476.024.104.525.500</b> <b>beta-Arrestin 1</b>
New Heading	<b>D12.776.476.024.104.525.750</b> <b>Beta-Arrestin 2</b>
-	D12.776.476.024.139 CARD Signaling Adaptor Proteins
-	D12.776.476.024.139.124 Apoptotic Protease-Activating Factor 1
New Tree	<a href="#">D12.776.476.024.139.155</a> <a href="#">Caspase 9</a>
-	D12.776.476.024.139.186 CRADD Signaling Adaptor Protein
-	D12.776.476.024.139.249 Nod1 Signaling Adaptor Protein
-	D12.776.476.024.139.500 Nod2 Signaling Adaptor Protein

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D12.776.476.024.139.750 Kinase 2	Receptor-Interacting Protein Serine-Threonine
-	D12.776.476.024.280	Caveolin 1
-	D12.776.476.024.295	Caveolin 2
-	D12.776.476.024.311	Cortactin
-	D12.776.476.024.374	Crk-Associated Substrate Protein
-	D12.776.476.024.375	Death Domain Receptor Signaling Adaptor Proteins
New Tree	<a href="#">D12.776.476.024.375.025</a>	<a href="#">Caspase 8</a>
-	D12.776.476.024.375.050	CRADD Signaling Adaptor Protein
-	D12.776.476.024.375.100	Edar-Associated Death Domain Protein
-	D12.776.476.024.375.550	Fas-Associated Death Domain Protein
-	D12.776.476.024.375.662 Kinases	Receptor-Interacting Protein Serine-Threonine
-	D12.776.476.024.375.775	TNF Receptor-Associated Death Domain Protein
New Heading	<b>D12.776.476.024.376</b>	<b>Dishevelled Proteins</b>
-	D12.776.476.024.377	GRB2 Adaptor Protein
-	D12.776.476.024.379	GRB7 Adaptor Protein
-	D12.776.476.024.380	GRB10 Adaptor Protein
New Heading	<b>D12.776.476.024.381</b>	<b>Homer Scaffolding Proteins</b>
-	D12.776.476.024.382	Insulin Receptor Substrate Proteins
-	D12.776.476.024.385	Interferon Regulatory Factors
-	D12.776.476.024.385.124	Interferon Regulatory Factor-1
-	D12.776.476.024.385.249	Interferon Regulatory Factor-2
-	D12.776.476.024.385.374	Interferon Regulatory Factor-3
-	D12.776.476.024.385.437	Interferon Regulatory Factor-7
-	D12.776.476.024.385.500 Subunit	Interferon-Stimulated Gene Factor 3, gamma
-	D12.776.476.024.386	Interferon-Stimulated Gene Factor 3
-	D12.776.476.024.386.500	Interferon-Stimulated Gene Factor 3, alpha Subunit
-	D12.776.476.024.386.500.500	STAT1 Transcription Factor
-	D12.776.476.024.386.500.750	STAT2 Transcription Factor
-	D12.776.476.024.386.750 Subunit	Interferon-Stimulated Gene Factor 3, gamma
New Heading	<b>D12.776.476.024.387</b>	<b>Kelch-Like ECH-Associated Protein 1</b>
-	D12.776.476.024.388	Mediator Complex

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D12.776.476.024.388.249	Cyclin C
-	D12.776.476.024.388.500	Cyclin-Dependent Kinase 8
-	D12.776.476.024.388.750	Mediator Complex Subunit 1
-	D12.776.476.024.390	Myeloid Differentiation Factor 88
-	D12.776.476.024.391	Nod Signaling Adaptor Proteins
-	D12.776.476.024.391.249	Nod1 Signaling Adaptor Protein
-	D12.776.476.024.391.500	Nod2 Signaling Adaptor Protein
-	D12.776.476.024.394	Nuclear Receptor Coactivators
-	D12.776.476.024.394.049	Mediator Complex Subunit 1
-	D12.776.476.024.394.100	Nuclear Receptor Coactivator 1
-	D12.776.476.024.394.200	Nuclear Receptor Coactivator 2
-	D12.776.476.024.394.300	Nuclear Receptor Coactivator 3
New Heading	<b>D12.776.476.024.394.650 Gamma Coactivator 1-alpha</b>	<b>Peroxisome Proliferator-Activated Receptor</b>
-	D12.776.476.024.397	Paxillin
-	D12.776.476.024.400	PII Nitrogen Regulatory Proteins
New Tree	<a href="#">D12.776.476.024.402</a>	<a href="#">Protein Inhibitors of Activated STAT</a>
Old Tree	<del>D12.776.476.024.405</del>	<del>Protein Inhibitors of Activated STAT</del>
New Tree	<a href="#">D12.776.476.024.407</a>	<a href="#">Proto-Oncogene Proteins c-crkl</a>
Old Tree	<del>D12.776.476.024.409</del>	<del>Proto-Oncogene Proteins c-crkl</del>
New Tree	<a href="#">D12.776.476.024.409</a>	<a href="#">Proto-Oncogene Proteins c-vav</a>
New Tree	<a href="#">D12.776.476.024.410</a>	<a href="#">Receptor Activity-Modifying Proteins</a>
New Tree	<a href="#">D12.776.476.024.410.100</a>	<a href="#">Receptor Activity-Modifying Protein 1</a>
New Tree	<a href="#">D12.776.476.024.410.200</a>	<a href="#">Receptor Activity-Modifying Protein 2</a>
New Tree	<a href="#">D12.776.476.024.410.300</a>	<a href="#">Receptor Activity-Modifying Protein 3</a>
Old Tree	<del>D12.776.476.024.411</del>	<del>Proto-Oncogene Proteins c-vav</del>
Old Tree	<del>D12.776.476.024.412</del>	<del>Receptor Activity-Modifying Proteins</del>
Old Tree	<del>D12.776.476.024.412.100</del>	<del>Receptor Activity-Modifying Protein 1</del>
Old Tree	<del>D12.776.476.024.412.200</del>	<del>Receptor Activity-Modifying Protein 2</del>
Old Tree	<del>D12.776.476.024.412.300</del>	<del>Receptor Activity-Modifying Protein 3</del>
Old Tree	<del>D12.776.476.024.414</del>	<del>Retinoblastoma Binding Proteins</del>
Old Tree	<del>D12.776.476.024.414.049</del>	<del>E2F1 Transcription Factor</del>

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	D12.776.476.024.414.100 Retinoblastoma-Binding Protein 1
Old Tree	D12.776.476.024.414.200 Retinoblastoma-Binding Protein 2
Old Tree	D12.776.476.024.414.600 Retinoblastoma-Binding Protein 4
Old Tree	D12.776.476.024.414.700 Retinoblastoma-Binding Protein 7
Old Tree	D12.776.476.024.415 Shc Signaling Adaptor Proteins
Old Tree	D12.776.476.024.417 Smad Proteins
Old Tree	D12.776.476.024.417.249 Smad Proteins, Inhibitory
Old Tree	D12.776.476.024.417.249.600 Smad6 Protein
Old Tree	D12.776.476.024.417.249.700 Smad7 Protein
Old Tree	D12.776.476.024.417.500 Smad Proteins, Receptor-Regulated
Old Tree	D12.776.476.024.417.500.100 Smad1 Protein
Old Tree	D12.776.476.024.417.500.200 Smad2 Protein
Old Tree	D12.776.476.024.417.500.300 Smad3 Protein
Old Tree	D12.776.476.024.417.500.500 Smad5 Protein
Old Tree	D12.776.476.024.417.500.800 Smad8 Protein
Old Tree	D12.776.476.024.417.750 Smad4 Protein
New Tree	D12.776.476.024.420 Retinoblastoma Binding Proteins
New Tree	D12.776.476.024.420.049 E2F1 Transcription Factor
New Tree	D12.776.476.024.420.100 Retinoblastoma-Binding Protein 1
New Tree	D12.776.476.024.420.200 Retinoblastoma-Binding Protein 2
New Tree	D12.776.476.024.420.600 Retinoblastoma-Binding Protein 4
New Tree	D12.776.476.024.420.700 Retinoblastoma-Binding Protein 7
Old Tree	D12.776.476.024.421 STAT Transcription Factors
Old Tree	D12.776.476.024.421.100 STAT1 Transcription Factor
Old Tree	D12.776.476.024.421.200 STAT2 Transcription Factor
Old Tree	D12.776.476.024.421.300 STAT3 Transcription Factor
Old Tree	D12.776.476.024.421.400 STAT4 Transcription Factor
Old Tree	D12.776.476.024.421.500 STAT5 Transcription Factor
Old Tree	D12.776.476.024.421.600 STAT6 Transcription Factor
New Heading	<b>D12.776.476.024.422 Sequestosome-1 Protein</b>
New Tree	D12.776.476.024.424 Shc Signaling Adaptor Proteins



## MeSH Tree Changes for 2017

Type	Tree - heading
New Heading	<b>D12.776.476.024.424.500</b> <b>Src Homology 2 Domain-Containing, Transforming Protein 1</b>
New Heading	<b>D12.776.476.024.424.750</b> <b>Src Homology 2 Domain-Containing, Transforming Protein 2</b>
New Heading	<b>D12.776.476.024.424.875</b> <b>Src Homology 2 Domain-Containing, Transforming Protein 3</b>
New Heading	<b>D12.776.476.024.426</b> <b>Signaling Lymphocytic Activation Molecule Associated Protein</b>
New Tree	D12.776.476.024.428 Smad Proteins
New Tree	D12.776.476.024.428.249 Smad Proteins, Inhibitory
New Tree	D12.776.476.024.428.249.600 Smad6 Protein
New Tree	D12.776.476.024.428.249.700 Smad7 Protein
New Tree	D12.776.476.024.428.500 Smad Proteins, Receptor-Regulated
New Tree	D12.776.476.024.428.500.100 Smad1 Protein
New Tree	D12.776.476.024.428.500.200 Smad2 Protein
New Tree	D12.776.476.024.428.500.300 Smad3 Protein
New Tree	D12.776.476.024.428.500.500 Smad5 Protein
New Tree	D12.776.476.024.428.500.800 Smad8 Protein
New Tree	D12.776.476.024.428.750 Smad4 Protein
New Tree	D12.776.476.024.430 STAT Transcription Factors
New Tree	D12.776.476.024.430.100 STAT1 Transcription Factor
New Tree	D12.776.476.024.430.200 STAT2 Transcription Factor
New Tree	D12.776.476.024.430.300 STAT3 Transcription Factor
New Tree	D12.776.476.024.430.400 STAT4 Transcription Factor
New Tree	D12.776.476.024.430.500 STAT5 Transcription Factor
New Tree	D12.776.476.024.430.600 STAT6 Transcription Factor

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.476.024.437                      Suppressor of Cytokine Signaling Proteins
New Heading	<b>D12.776.476.024.437.500</b> <b>Suppressor of Cytokine Signaling 1 Protein</b>
New Heading	<b>D12.776.476.024.437.750</b> <b>Suppressor of Cytokine Signaling 3 Protein</b>
-	D12.776.476.024.468                      Syntenins
-	D12.776.476.024.500 and Proteins                      Tumor Necrosis Factor Receptor-Associated Peptides
-	D12.776.476.024.500.050                      CRADD Signaling Adaptor Protein
-	D12.776.476.024.500.061                      Edar-Associated Death Domain Protein
-	D12.776.476.024.500.124                      Fas-Associated Death Domain Protein
-	D12.776.476.024.500.186 Kinases                      Receptor-Interacting Protein Serine-Threonine
-	D12.776.476.024.500.186.500 Kinase 2                      Receptor-Interacting Protein Serine-Threonine
-	D12.776.476.024.500.249                      TNF Receptor-Associated Death Domain Protein
-	D12.776.476.024.500.500                      TNF Receptor-Associated Factor 1
-	D12.776.476.024.500.750                      TNF Receptor-Associated Factor 2
-	D12.776.476.024.500.875                      TNF Receptor-Associated Factor 3
-	D12.776.476.024.500.906                      TNF Receptor-Associated Factor 4
-	D12.776.476.024.500.937                      TNF Receptor-Associated Factor 5
-	D12.776.476.024.500.968                      TNF Receptor-Associated Factor 6
-	D12.776.476.050                      Adenylyl Cyclases
-	D12.776.476.062                      AMP-Activated Protein Kinases
-	D12.776.476.075                      Apoptosis Regulatory Proteins
-	D12.776.476.075.311                      Apoptosis Inducing Factor
New Heading	<b>D12.776.476.075.323</b> <b>Bcl-2-Like Protein 11</b>
New Heading	<b>D12.776.476.075.335</b> <b>Beclin-1</b>
-	D12.776.476.075.358                      CARD Signaling Adaptor Proteins
-	D12.776.476.075.358.124                      Apoptotic Protease-Activating Factor 1
New Tree	<a href="#">D12.776.476.075.358.155</a> <a href="#">Caspase 9</a>
-	D12.776.476.075.358.186                      CRADD Signaling Adaptor Protein
-	D12.776.476.075.358.249                      Nod1 Signaling Adaptor Protein
-	D12.776.476.075.358.500                      Nod2 Signaling Adaptor Protein
-	D12.776.476.075.358.750 Kinase 2                      Receptor-Interacting Protein Serine-Threonine

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D12.776.476.075.405	Caspases
-	D12.776.476.075.405.350	Caspases, Effector
-	D12.776.476.075.405.350.300	Caspase 3
-	D12.776.476.075.405.350.600	Caspase 6
-	D12.776.476.075.405.350.700	Caspase 7
-	D12.776.476.075.405.350.900	Caspase 14
-	D12.776.476.075.405.550	Caspases, Initiator
-	D12.776.476.075.405.550.100	Caspase 1
-	D12.776.476.075.405.550.200	Caspase 2
-	D12.776.476.075.405.550.800	Caspase 8
-	D12.776.476.075.405.550.900	Caspase 9
-	D12.776.476.075.405.550.910	Caspase 10
-	D12.776.476.075.405.550.920	Caspase 12
-	D12.776.476.075.421	Death Domain Receptor Signaling Adaptor Proteins
-	D12.776.476.075.421.024 Protein	CASP8 and FADD-Like Apoptosis Regulating
New Tree	<a href="#">D12.776.476.075.421.037</a>	<a href="#">Caspase 8</a>
-	D12.776.476.075.421.050	CRADD Signaling Adaptor Protein
-	D12.776.476.075.421.100	Edar-Associated Death Domain Protein
-	D12.776.476.075.421.200	Fas-Associated Death Domain Protein
-	D12.776.476.075.421.400 Kinases	Receptor-Interacting Protein Serine-Threonine
-	D12.776.476.075.421.600	TNF Receptor-Associated Death Domain Protein
-	D12.776.476.075.437	Inhibitor of Apoptosis Proteins
-	D12.776.476.075.437.500	Neuronal Apoptosis-Inhibitory Protein
-	D12.776.476.075.437.750	X-Linked Inhibitor of Apoptosis Protein
-	D12.776.476.075.718	Proto-Oncogene Proteins c-bcl-2
-	D12.776.476.075.718.100	bcl-Associated Death Protein
-	D12.776.476.075.718.400	bcl-2-Associated X Protein
-	D12.776.476.075.718.425	bcl-2 Homologous Antagonist-Killer Protein
-	D12.776.476.075.718.875	bcl-X Protein
-	D12.776.476.075.718.937	BH3 Interacting Domain Death Agonist Protein
-	D12.776.476.075.718.968	Myeloid Cell Leukemia Sequence 1 Protein
New Heading	<b>D12.776.476.078</b>	<b>Autophagy-Related Protein-1 Homolog</b>
-	D12.776.476.081	Axin Signaling Complex

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.476.081.249      Adenomatous Polyposis Coli Protein
-	D12.776.476.081.500      Axin Protein
-	D12.776.476.081.750      Casein Kinase I
-	D12.776.476.081.750.100      Casein Kinase Ialpha
-	D12.776.476.081.750.200      Casein Kinase Idelta
-	D12.776.476.081.750.300      Casein Kinase Iepsilon
-	D12.776.476.081.875      Glycogen Synthase Kinase 3
New Heading	<b>D12.776.476.081.875.500      Glycogen Synthase Kinase 3 beta</b>
-	D12.776.476.087      Calcineurin
-	D12.776.476.100      Calcium-Calmodulin-Dependent Protein Kinases
-	D12.776.476.100.049      Calcium-Calmodulin-Dependent Protein Kinase Kinase
-	D12.776.476.100.100      Calcium-Calmodulin-Dependent Protein Kinase Type 1
-	D12.776.476.100.200      Calcium-Calmodulin-Dependent Protein Kinase Type 2
-	D12.776.476.100.350      Calcium-Calmodulin-Dependent Protein Kinase Type 4
-	D12.776.476.100.387      Death-Associated Protein Kinases
-	D12.776.476.100.425      Elongation Factor 2 Kinase
-	D12.776.476.100.500      Myosin-Light-Chain Kinase
-	D12.776.476.150      Casein Kinases
-	D12.776.476.150.299      Casein Kinase I
-	D12.776.476.150.299.100      Casein Kinase Ialpha
-	D12.776.476.150.299.200      Casein Kinase Idelta
-	D12.776.476.150.299.300      Casein Kinase Iepsilon
-	D12.776.476.150.600      Casein Kinase II
-	D12.776.476.162      Class I Phosphatidylinositol 3-Kinases
-	D12.776.476.162.100      Class Ia Phosphatidylinositol 3-Kinase
-	D12.776.476.162.200      Class Ib Phosphatidylinositol 3-Kinase
-	D12.776.476.168      Class II Phosphatidylinositol 3-Kinases
-	D12.776.476.171      Class III Phosphatidylinositol 3-Kinases
-	D12.776.476.171.500      Vacuolar Sorting Protein VPS15
-	D12.776.476.175      Cold Shock Proteins and Peptides
-	D12.776.476.200      Cyclic Nucleotide-Regulated Protein Kinases
-	D12.776.476.200.125      Cyclic AMP-Dependent Protein Kinases
-	D12.776.476.200.125.750      Cyclic AMP-Dependent Protein Kinase Type I
-	D12.776.476.200.125.750.124 Subunit      Cyclic AMP-Dependent Protein Kinase RIalpha

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.476.200.125.750.249 Subunit Cyclic AMP-Dependent Protein Kinase RIbeta
-	D12.776.476.200.125.750.500 Subunits Cyclic AMP-Dependent Protein Kinase Catalytic
-	D12.776.476.200.125.875 Cyclic AMP-Dependent Protein Kinase Type II
-	D12.776.476.200.125.875.500 Subunits Cyclic AMP-Dependent Protein Kinase Catalytic
-	D12.776.476.200.125.875.750 Subunit Cyclic AMP-Dependent Protein Kinase RIalpha
-	D12.776.476.200.125.875.875 Subunit Cyclic AMP-Dependent Protein Kinase RIbeta
-	D12.776.476.200.150 Cyclic GMP-Dependent Protein Kinases
-	D12.776.476.200.150.500 Cyclic GMP-Dependent Protein Kinase Type I
-	D12.776.476.200.150.750 Cyclic GMP-Dependent Protein Kinase Type II
-	D12.776.476.200.575 Protamine Kinase
-	D12.776.476.225 Cyclin-Dependent Kinase Inhibitor Proteins
-	D12.776.476.225.100 Cyclin-Dependent Kinase Inhibitor p15
-	D12.776.476.225.200 Cyclin-Dependent Kinase Inhibitor p16
-	D12.776.476.225.300 Cyclin-Dependent Kinase Inhibitor p18
-	D12.776.476.225.400 Cyclin-Dependent Kinase Inhibitor p19
-	D12.776.476.225.500 Cyclin-Dependent Kinase Inhibitor p21
-	D12.776.476.225.600 Cyclin-Dependent Kinase Inhibitor p27
-	D12.776.476.225.700 Cyclin-Dependent Kinase Inhibitor p57
-	D12.776.476.250 Cyclin-Dependent Kinases
-	D12.776.476.250.067 CDC2-CDC28 Kinases
-	D12.776.476.250.067.249 CDC2 Protein Kinase
-	D12.776.476.250.067.500 CDC28 Protein Kinase, S cerevisiae
-	D12.776.476.250.067.875 Cyclin-Dependent Kinase 5
-	D12.776.476.250.067.900 Cyclin-Dependent Kinase 9
-	D12.776.476.250.323 Cyclin-Dependent Kinase 2
-	D12.776.476.250.387 Cyclin-Dependent Kinase 3
-	D12.776.476.250.451 Cyclin-Dependent Kinase 4
-	D12.776.476.250.515 Cyclin-Dependent Kinase 6
-	D12.776.476.250.548 Cyclin-Dependent Kinase 8
-	D12.776.476.250.580 Maturation-Promoting Factor
-	D12.776.476.250.580.500 CDC2 Protein Kinase
-	D12.776.476.262 Cyclins

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.476.262.100 Cyclin A
-	D12.776.476.262.100.100 Cyclin A1
-	D12.776.476.262.100.200 Cyclin A2
-	D12.776.476.262.120 Cyclin B
-	D12.776.476.262.120.100 Cyclin B1
-	D12.776.476.262.120.200 Cyclin B2
-	D12.776.476.262.135 Cyclin C
-	D12.776.476.262.150 Cyclin D
-	D12.776.476.262.150.100 Cyclin D1
-	D12.776.476.262.150.200 Cyclin D2
-	D12.776.476.262.150.300 Cyclin D3
-	D12.776.476.262.180 Cyclin E
-	D12.776.476.262.200 Cyclin G
-	D12.776.476.262.200.100 Cyclin G1
-	D12.776.476.262.200.200 Cyclin G2
-	D12.776.476.262.300 Cyclin H
-	D12.776.476.262.400 Cyclin I
-	D12.776.476.262.700 Cyclin T
-	D12.776.476.268 Dual-Specificity Phosphatases
-	D12.776.476.268.100 cdc25 Phosphatases
-	D12.776.476.268.200 Dual Specificity Phosphatase 1
-	D12.776.476.268.250 Dual Specificity Phosphatase 2
-	D12.776.476.268.300 Dual Specificity Phosphatase 3
-	D12.776.476.268.600 Dual Specificity Phosphatase 6
-	D12.776.476.275 eIF-2 Kinase
-	D12.776.476.287 Focal Adhesion Protein-Tyrosine Kinases
-	D12.776.476.293 G-Protein-Coupled Receptor Kinases
-	D12.776.476.293.249 beta-Adrenergic Receptor Kinases
-	D12.776.476.293.249.200 G-Protein-Coupled Receptor Kinase 2
-	D12.776.476.293.249.300 G-Protein-Coupled Receptor Kinase 3
-	D12.776.476.293.500 G-Protein-Coupled Receptor Kinase 1
-	D12.776.476.293.750 G-Protein-Coupled Receptor Kinase 4
-	D12.776.476.293.875 G-Protein-Coupled Receptor Kinase 5
-	D12.776.476.300 Glycogen Synthase Kinases
-	D12.776.476.300.500 Glycogen Synthase Kinase 3
New	<b>D12.776.476.300.500.500 Glycogen Synthase Kinase 3 beta</b>

## MeSH Tree Changes for 2017

Type	Tree - heading	
Heading		
-	D12.776.476.325	GTP-Binding Protein Regulators
-	D12.776.476.325.150	GTPase-Activating Proteins
-	D12.776.476.325.150.100	Chimerin Proteins
-	D12.776.476.325.150.100.200	Chimerin 1
-	D12.776.476.325.150.300	Eukaryotic Initiation Factor-5
-	D12.776.476.325.150.500	ras GTPase-Activating Proteins
-	D12.776.476.325.150.500.460	Neurofibromin 1
-	D12.776.476.325.150.500.500	p120 GTPase Activating Protein
-	D12.776.476.325.150.750	RGS Proteins
-	D12.776.476.325.225	Guanine Nucleotide Dissociation Inhibitors
-	D12.776.476.325.225.500 Inhibitors	rho-Specific Guanine Nucleotide Dissociation
-	D12.776.476.325.225.500.100 alpha	rho Guanine Nucleotide Dissociation Inhibitor
-	D12.776.476.325.225.500.200 beta	rho Guanine Nucleotide Dissociation Inhibitor
-	D12.776.476.325.225.500.300 gamma	rho Guanine Nucleotide Dissociation Inhibitor
-	D12.776.476.325.300	Guanine Nucleotide Exchange Factors
-	D12.776.476.325.300.050	Eukaryotic Initiation Factor-2B
-	D12.776.476.325.300.099	Rho Guanine Nucleotide Exchange Factors
New Tree	<a href="#">D12.776.476.325.300.099.500</a>	<a href="#">Proto-Oncogene Proteins c-bcr</a>
New Tree	<a href="#">D12.776.476.325.300.099.750</a>	<a href="#">Proto-Oncogene Proteins c-vav</a>
-	D12.776.476.325.300.300	Guanine Nucleotide-Releasing Factor 2
Old Tree	<del>D12.776.476.325.300.450</del>	<del>Proto-Oncogene Proteins c-vav</del>
-	D12.776.476.325.300.600	ral Guanine Nucleotide Exchange Factor
-	D12.776.476.325.300.700	ras Guanine Nucleotide Exchange Factors
-	D12.776.476.325.300.700.500	ras-GRF1
-	D12.776.476.325.300.700.700	Son of Sevenless Proteins
-	D12.776.476.325.300.700.700.600	Son of Sevenless Protein, Drosophila
-	D12.776.476.325.300.700.700.630	SOS1 Protein
-	D12.776.476.350	Guanylate Cyclase
New Heading	<b>D12.776.476.350.500</b>	<b>Soluble Guanylyl Cyclase</b>
-	D12.776.476.375	Heterotrimeric GTP-Binding Proteins

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.476.375.100 GTP-Binding Protein alpha Subunits
-	D12.776.476.375.100.100 GTP-Binding Protein alpha Subunits, G12-G13
-	D12.776.476.375.100.200 GTP-Binding Protein alpha Subunits, Gi-Go
-	D12.776.476.375.100.200.500 GTP-Binding Protein alpha Subunit, Gi2
-	D12.776.476.375.100.300 GTP-Binding Protein alpha Subunits, Gq-G11
-	D12.776.476.375.100.400 GTP-Binding Protein alpha Subunits, Gs
-	D12.776.476.375.520 GTP-Binding Protein beta Subunits
-	D12.776.476.375.730 GTP-Binding Protein gamma Subunits
-	D12.776.476.375.940 Transducin
-	D12.776.476.378 I-kappa B Kinase
-	D12.776.476.381 I-kappa B Proteins
New Heading	<b>D12.776.476.381.500 NF-KappaB Inhibitor alpha</b>
-	D12.776.476.384 Interleukin-1 Receptor-Associated Kinases
-	D12.776.476.387 Intracellular Calcium-Sensing Proteins
-	D12.776.476.387.249 Calmodulin
-	D12.776.476.387.311 Calnexin
-	D12.776.476.387.374 Calreticulin
-	D12.776.476.387.437 Gelsolin
-	D12.776.476.387.500 Neuronal Calcium-Sensor Proteins
-	D12.776.476.387.500.124 Guanylate Cyclase-Activating Proteins
-	D12.776.476.387.500.249 Hippocalcin
-	D12.776.476.387.500.374 Kv Channel-Interacting Proteins
-	D12.776.476.387.500.500 Neurocalcin
-	D12.776.476.387.500.750 Recoverin
-	D12.776.476.393 Janus Kinases
-	D12.776.476.393.100 Janus Kinase 1
-	D12.776.476.393.200 Janus Kinase 2
-	D12.776.476.393.300 Janus Kinase 3
-	D12.776.476.393.650 TYK2 Kinase
-	D12.776.476.396 Lim Kinases
-	D12.776.476.400 MAP Kinase Kinase Kinases
-	D12.776.476.400.100 MAP Kinase Kinase Kinase 1
-	D12.776.476.400.200 MAP Kinase Kinase Kinase 2
-	D12.776.476.400.300 MAP Kinase Kinase Kinase 3
-	D12.776.476.400.400 MAP Kinase Kinase Kinase 4



## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.476.400.500 MAP Kinase Kinase Kinase 5
-	D12.776.476.400.800 Proto-Oncogene Proteins c-mos
-	D12.776.476.400.842 raf Kinases
-	D12.776.476.400.842.249 Oncogene Proteins v-raf
-	D12.776.476.400.842.374 Proto-Oncogene Proteins A-raf
-	D12.776.476.400.842.437 Proto-Oncogene Proteins B-raf
-	D12.776.476.400.842.500 Proto-Oncogene Proteins c-raf
New Heading	<b>D12.776.476.420 Methyl-Accepting Chemotaxis Proteins</b>
-	D12.776.476.440 Mitogen-Activated Protein Kinase Kinases
-	D12.776.476.440.100 MAP Kinase Kinase 1
-	D12.776.476.440.200 MAP Kinase Kinase 2
-	D12.776.476.440.300 MAP Kinase Kinase 3
-	D12.776.476.440.400 MAP Kinase Kinase 4
-	D12.776.476.440.500 MAP Kinase Kinase 5
-	D12.776.476.440.600 MAP Kinase Kinase 6
-	D12.776.476.440.700 MAP Kinase Kinase 7
-	D12.776.476.445 Mitogen-Activated Protein Kinase Phosphatases
-	D12.776.476.445.200 Dual Specificity Phosphatase 1
-	D12.776.476.445.250 Dual Specificity Phosphatase 2
-	D12.776.476.445.300 Dual Specificity Phosphatase 3
-	D12.776.476.445.600 Dual Specificity Phosphatase 6
-	D12.776.476.450 Mitogen-Activated Protein Kinases
-	D12.776.476.450.169 Extracellular Signal-Regulated MAP Kinases
-	D12.776.476.450.169.500 Mitogen-Activated Protein Kinase 1
-	D12.776.476.450.169.750 Mitogen-Activated Protein Kinase 3
-	D12.776.476.450.169.875 Mitogen-Activated Protein Kinase 6
-	D12.776.476.450.169.937 Mitogen-Activated Protein Kinase 7
-	D12.776.476.450.340 JNK Mitogen-Activated Protein Kinases
-	D12.776.476.450.340.500 Mitogen-Activated Protein Kinase 8
-	D12.776.476.450.340.750 Mitogen-Activated Protein Kinase 9
-	D12.776.476.450.340.800 Mitogen-Activated Protein Kinase 10
-	D12.776.476.450.835 p38 Mitogen-Activated Protein Kinases
-	D12.776.476.450.835.500 Mitogen-Activated Protein Kinase 11
-	D12.776.476.450.835.750 Mitogen-Activated Protein Kinase 12
-	D12.776.476.450.835.875 Mitogen-Activated Protein Kinase 13

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.476.450.835.937 Mitogen-Activated Protein Kinase 14
-	D12.776.476.525 Monomeric GTP-Binding Proteins
-	D12.776.476.525.100 ADP-Ribosylation Factors
-	D12.776.476.525.100.100 ADP-Ribosylation Factor 1
-	D12.776.476.525.400 rab GTP-Binding Proteins
-	D12.776.476.525.400.025 rab1 GTP-Binding Proteins
-	D12.776.476.525.400.050 rab2 GTP-Binding Protein
-	D12.776.476.525.400.100 rab3 GTP-Binding Proteins
-	D12.776.476.525.400.100.100 rab3A GTP-Binding Protein
-	D12.776.476.525.400.150 rab4 GTP-Binding Proteins
-	D12.776.476.525.400.200 rab5 GTP-Binding Proteins
-	D12.776.476.525.450 ral GTP-Binding Proteins
-	D12.776.476.525.462 ran GTP-Binding Protein
-	D12.776.476.525.475 rap GTP-Binding Proteins
-	D12.776.476.525.475.100 rap1 GTP-Binding Proteins
-	D12.776.476.525.500 ras Proteins
-	D12.776.476.525.500.300 Oncogene Protein p21(ras)
-	D12.776.476.525.500.600 Proto-Oncogene Proteins p21(ras)
-	D12.776.476.525.700 rho GTP-Binding Proteins
-	D12.776.476.525.700.050 cdc42 GTP-Binding Protein
-	D12.776.476.525.700.050.500 cdc42 GTP-Binding Protein, <i>Saccharomyces cerevisiae</i>
-	D12.776.476.525.700.100 rac GTP-Binding Proteins
-	D12.776.476.525.700.100.100 rac1 GTP-Binding Protein
-	D12.776.476.525.700.200 rhoA GTP-Binding Protein
-	D12.776.476.525.700.300 rhoB GTP-Binding Protein
-	D12.776.476.534 Mucin-4
-	D12.776.476.543 Olfactory Marker Protein
-	D12.776.476.548 p21-Activated Kinases
-	D12.776.476.555 Phosphatidylethanolamine Binding Protein
-	D12.776.476.556 Phosphoinositide Phospholipase C
-	D12.776.476.556.500 Phospholipase C beta
-	D12.776.476.556.625 Phospholipase C delta
-	D12.776.476.556.750 Phospholipase C gamma
-	D12.776.476.561 Protein Phosphatase 2
-	D12.776.476.564 Protein Tyrosine Phosphatases, Non-Receptor

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D12.776.476.564.100	Protein Tyrosine Phosphatase, Non-Receptor Type 1
-	D12.776.476.564.200	Protein Tyrosine Phosphatase, Non-Receptor Type 2
-	D12.776.476.564.300	Protein Tyrosine Phosphatase, Non-Receptor Type 3
-	D12.776.476.564.400	Protein Tyrosine Phosphatase, Non-Receptor Type 4
-	D12.776.476.564.600	Protein Tyrosine Phosphatase, Non-Receptor Type 6
-	D12.776.476.564.800	Protein Tyrosine Phosphatase, Non-Receptor Type 11
-	D12.776.476.564.850	Protein Tyrosine Phosphatase, Non-Receptor Type 12
-	D12.776.476.564.860	Protein Tyrosine Phosphatase, Non-Receptor Type 13
-	D12.776.476.564.930	Protein Tyrosine Phosphatase, Non-Receptor Type 22
-	D12.776.476.565	Proto-Oncogene Proteins c-akt
-	D12.776.476.571	Proto-Oncogene Proteins c-fyn
-	D12.776.476.590	PTEN Phosphohydrolase
-	D12.776.476.592	Receptor-Like Protein Tyrosine Phosphatases
-	D12.776.476.592.100	Receptor-Like Protein Tyrosine Phosphatases, Class 1
-	D12.776.476.592.100.500	Antigens, CD45
-	D12.776.476.592.200	Receptor-Like Protein Tyrosine Phosphatases, Class 2
-	D12.776.476.592.300	Receptor-Like Protein Tyrosine Phosphatases, Class 3
-	D12.776.476.592.400	Receptor-Like Protein Tyrosine Phosphatases, Class 4
-	D12.776.476.592.500	Receptor-Like Protein Tyrosine Phosphatases, Class 5
-	D12.776.476.592.700	Receptor-Like Protein Tyrosine Phosphatases, Class 7
-	D12.776.476.592.800	Receptor-Like Protein Tyrosine Phosphatases, Class 8
-	D12.776.476.595	rho-Associated Kinases
-	D12.776.476.600	Ribosomal Protein S6 Kinases
-	D12.776.476.600.249	Ribosomal Protein S6 Kinases, 70-kDa
-	D12.776.476.600.500	Ribosomal Protein S6 Kinases, 90-kDa
-	D12.776.476.800	SH2 Domain-Containing Protein Tyrosine Phosphatases
-	D12.776.476.800.200	Protein Tyrosine Phosphatase, Non-Receptor Type 6
-	D12.776.476.800.800	Protein Tyrosine Phosphatase, Non-Receptor Type 11
-	D12.776.476.900	Sirtuins
-	D12.776.476.900.100	Sirtuin 1
-	D12.776.476.900.200	Sirtuin 2
-	D12.776.476.900.600	Sirtuin 3
New Heading	<b>D12.776.476.913</b>	<b>Syk Kinase</b>
-	D12.776.476.925	TOR Serine-Threonine Kinases
New	<b>D12.776.476.938</b>	<b>Tumor Necrosis Factor alpha-Induced Protein 3</b>

## MeSH Tree Changes for 2017

Type	Tree - heading	
Heading		
-	D12.776.476.950	ZAP-70 Protein-Tyrosine Kinase
-	D12.776.481	Intrinsically Disordered Proteins
-	D12.776.486	Iodoproteins
-	D12.776.486.706	Thyroglobulin
-	D12.776.494	Iron-Regulatory Proteins
-	D12.776.494.249	Hepcidins
-	D12.776.494.500	Iron Regulatory Protein 1
-	D12.776.494.750	Iron Regulatory Protein 2
-	D12.776.503	Lectins
New Heading	<b>D12.776.503.070</b>	<b>Chitinase-3-Like Protein 1</b>
New Heading	<b>D12.776.503.140</b>	<b>Discoidins</b>
-	D12.776.503.280	Lectins, C-Type
-	D12.776.503.280.124	Asialoglycoprotein Receptor
-	D12.776.503.280.249	Collectins
-	D12.776.503.280.249.500	Mannose-Binding Lectin
-	D12.776.503.280.249.600	Pulmonary Surfactant-Associated Protein A
-	D12.776.503.280.249.625	Pulmonary Surfactant-Associated Protein D
-	D12.776.503.280.437	Hyalectins
-	D12.776.503.280.437.100	Aggrecans
-	D12.776.503.280.437.275	Brevican
-	D12.776.503.280.437.450	Neurocan
-	D12.776.503.280.437.800	Versicans
New Tree	<b>D12.776.503.280.718</b>	<b>Scavenger Receptors, Class E</b>
-	D12.776.503.295	Calnexin
-	D12.776.503.303	Calreticulin
-	D12.776.503.307	Galectins
-	D12.776.503.307.100	Galectin 1
-	D12.776.503.307.200	Galectin 2
-	D12.776.503.307.300	Galectin 3
-	D12.776.503.307.400	Galectin 4
-	D12.776.503.311	Mannose-Binding Lectins
-	D12.776.503.311.500	Mannose-Binding Lectin
-	D12.776.503.499	Plant Lectins

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.503.499.249      Abrin
-	D12.776.503.499.500      Concanavalin A
-	D12.776.503.499.625      Peanut Agglutinin
-	D12.776.503.499.750      Phytohemagglutinins
-	D12.776.503.499.875      Pokeweed Mitogens
-	D12.776.503.499.937      Ricin
-	D12.776.503.499.968      Wheat Germ Agglutinins
-	D12.776.503.499.968.900      Wheat Germ Agglutinin-Horseradish Peroxidase Conjugate
-	D12.776.503.687      Receptors, N-Acetylglucosamine
-	D12.776.503.843      Selectins
-	D12.776.503.843.300      E-Selectin
-	D12.776.503.843.510      L-Selectin
-	D12.776.503.843.775      P-Selectin
-	D12.776.503.921      Sialic Acid Binding Immunoglobulin-like Lectins
-	D12.776.503.921.049      Myelin-Associated Glycoprotein
-	D12.776.503.921.100      Sialic Acid Binding Ig-like Lectin 1
-	D12.776.503.921.200      Sialic Acid Binding Ig-like Lectin 2
-	D12.776.503.921.400      Sialic Acid Binding Ig-like Lectin 3
-	D12.776.512      LIM Domain Proteins
-	D12.776.512.124      Lim Kinases
-	D12.776.512.249      LIM-Homeodomain Proteins
-	D12.776.512.374      Paxillin
-	D12.776.512.500      Zyxin
-	D12.776.521      Lipoproteins
-	D12.776.521.120      Apolipoproteins
-	D12.776.521.120.200      Apolipoproteins A
-	D12.776.521.120.200.100      Apolipoprotein A-I
-	D12.776.521.120.200.150      Apolipoprotein A-II
New Heading	<b>D12.776.521.120.200.575      Apolipoprotein A-V</b>
-	D12.776.521.120.300      Apolipoproteins B
-	D12.776.521.120.300.240      Apolipoprotein B-48
-	D12.776.521.120.300.249      Apolipoprotein B-100
-	D12.776.521.120.400      Apolipoproteins C
-	D12.776.521.120.400.500      Apolipoprotein C-I

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D12.776.521.120.400.750	Apolipoprotein C-II
-	D12.776.521.120.400.875	Apolipoprotein C-III
-	D12.776.521.120.450	Apolipoproteins D
-	D12.776.521.120.500	Apolipoproteins E
-	D12.776.521.120.500.249	Apolipoprotein E2
-	D12.776.521.120.500.500	Apolipoprotein E3
-	D12.776.521.120.500.750	Apolipoprotein E4
New Heading	<b>D12.776.521.181 Member 5</b>	<b>ATP Binding Cassette Transporter, Sub-Family G,</b>
New Heading	<b>D12.776.521.212 Member 8</b>	<b>ATP Binding Cassette Transporter, Sub-Family G,</b>
-	D12.776.521.242	Chylomicrons
-	D12.776.521.242.500	Chylomicron Remnants
-	D12.776.521.400	Lipoprotein(a)
-	D12.776.521.400.500	Apoprotein(a)
-	D12.776.521.450	Lipoprotein-X
-	D12.776.521.479	Lipoproteins, HDL
-	D12.776.521.479.470	Cholesterol, HDL
-	D12.776.521.479.602	High-Density Lipoproteins, Pre-beta
-	D12.776.521.479.735	Lipoproteins, HDL2
-	D12.776.521.479.867	Lipoproteins, HDL3
-	D12.776.521.514	Lipoproteins, IDL
-	D12.776.521.550	Lipoproteins, LDL
-	D12.776.521.550.500	Cholesterol, LDL
-	D12.776.521.622	Lipoproteins, VLDL
-	D12.776.521.622.700	Cholesterol, VLDL
-	D12.776.526	LDL-Receptor Related Proteins
-	D12.776.526.100	Low Density Lipoprotein Receptor-Related Protein-1
-	D12.776.526.200	Low Density Lipoprotein Receptor-Related Protein-2
-	D12.776.526.500	Low Density Lipoprotein Receptor-Related Protein-5
-	D12.776.526.600	Low Density Lipoprotein Receptor-Related Protein-6
-	D12.776.529	Lithostathine
-	D12.776.532	Luminescent Proteins
-	D12.776.532.020	Aequorin
-	D12.776.532.265	Green Fluorescent Proteins
-	D12.776.532.510	Luciferases

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D12.776.532.510.249	Luciferases, Bacterial
-	D12.776.532.510.500	Luciferases, Firefly
-	D12.776.532.510.750	Luciferases, Renilla
-	D12.776.543	Membrane Proteins
-	D12.776.543.019	5-Lipoxygenase-Activating Proteins
-	D12.776.543.029	Amphiregulin
-	D12.776.543.039	Amyloid beta-Protein Precursor
-	D12.776.543.039.500	Amyloid beta-Peptides
-	D12.776.543.080	Ankyrins
-	D12.776.543.090	Arrestins
-	D12.776.543.090.050	Arrestin
New Heading	<b>D12.776.543.090.525</b>	<b>Beta-Arrestins</b>
New Heading	<b>D12.776.543.090.525.500</b>	<b>beta-Arrestin 1</b>
New Heading	<b>D12.776.543.090.525.750</b>	<b>Beta-Arrestin 2</b>
-	D12.776.543.095	B7 Antigens
-	D12.776.543.095.100	Antigens, CD80
-	D12.776.543.095.200	Antigens, CD86
-	D12.776.543.095.300	Antigens, CD274
-	D12.776.543.095.500	Inducible T-Cell Co-Stimulator Ligand
-	D12.776.543.095.800	Programmed Cell Death 1 Ligand 2 Protein
-	D12.776.543.095.900	V-Set Domain-Containing T-Cell Activation Inhibitor 1
-	D12.776.543.100	Bacterial Outer Membrane Proteins
-	D12.776.543.100.050	Adhesins, Bacterial
-	D12.776.543.100.050.040	Adhesins, Escherichia coli
-	D12.776.543.100.300	Fimbriae Proteins
New Heading	<b>D12.776.543.116</b>	<b>Bcl-2-Like Protein 11</b>
New Heading	<b>D12.776.543.131</b>	<b>Cadherin Related Proteins</b>
New Tree	<a href="#">D12.776.543.131.500</a>	<a href="#">Proto-Oncogene Proteins c-ret</a>
-	D12.776.543.162	Calnexin
New Heading	<b>D12.776.543.178</b>	<b>Carbonic Anhydrase IX</b>
-	D12.776.543.193	Chemokine CX3CL1

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.543.200 Cornified Envelope Proline-Rich Proteins
-	D12.776.543.250 Dystrophin
-	D12.776.543.268 Dystrophin-Associated Proteins
-	D12.776.543.268.500 Dystroglycans
-	D12.776.543.268.750 Sarcoglycans
-	D12.776.543.277 Electron Transport Chain Complex Proteins
-	D12.776.543.277.500 Electron-Transferring Flavoproteins
-	D12.776.543.277.500.500 Electron Transport Complex I
-	D12.776.543.277.500.750 Electron Transport Complex II
-	D12.776.543.277.500.750.500 Succinate Dehydrogenase
-	D12.776.543.277.687 Electron Transport Complex IV
-	D12.776.543.277.875 Succinate Cytochrome c Oxidoreductase
-	D12.776.543.277.875.249 Electron Transport Complex II
-	D12.776.543.277.875.249.500 Succinate Dehydrogenase
-	D12.776.543.277.875.500 Electron Transport Complex III
-	D12.776.543.287 Ephrins
-	D12.776.543.287.100 Ephrin-A1
-	D12.776.543.287.200 Ephrin-A2
-	D12.776.543.287.300 Ephrin-A3
-	D12.776.543.287.400 Ephrin-A4
-	D12.776.543.287.500 Ephrin-A5
-	D12.776.543.287.600 Ephrin-B1
-	D12.776.543.287.700 Ephrin-B2
-	D12.776.543.287.800 Ephrin-B3
-	D12.776.543.306 gp100 Melanoma Antigen
-	D12.776.543.325 Heterotrimeric GTP-Binding Proteins
-	D12.776.543.325.100 GTP-Binding Protein alpha Subunits
-	D12.776.543.325.100.100 GTP-Binding Protein alpha Subunits, G12-G13
-	D12.776.543.325.100.200 GTP-Binding Protein alpha Subunits, Gi-Go
-	D12.776.543.325.100.200.500 GTP-Binding Protein alpha Subunit, Gi2
-	D12.776.543.325.100.300 GTP-Binding Protein alpha Subunits, Gq-G11
-	D12.776.543.325.100.400 GTP-Binding Protein alpha Subunits, Gs
-	D12.776.543.325.450 GTP-Binding Protein beta Subunits
-	D12.776.543.325.625 GTP-Binding Protein gamma Subunits
-	D12.776.543.325.800 Transducin
-	D12.776.543.475 LDL-Receptor Related Protein-Associated Protein



## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.543.484 Lipid-Linked Proteins
-	D12.776.543.484.500 GPI-Linked Proteins
-	D12.776.543.484.500.100 Antigens, CD14
-	D12.776.543.484.500.120 Antigens, CD24
-	D12.776.543.484.500.130 Antigens, CD55
-	D12.776.543.484.500.140 Antigens, CD59
-	D12.776.543.484.500.160 Carbonic Anhydrase IV
New Heading	<b>D12.776.543.484.500.180 CD48 Antigen</b>
-	D12.776.543.484.500.200 Ciliary Neurotrophic Factor Receptor alpha Subunit
-	D12.776.543.484.500.250 Contactins
-	D12.776.543.484.500.250.100 Contactin 1
-	D12.776.543.484.500.250.200 Contactin 2
-	D12.776.543.484.500.300 Ephrin-A1
-	D12.776.543.484.500.310 Ephrin-A2
-	D12.776.543.484.500.320 Ephrin-A3
-	D12.776.543.484.500.330 Ephrin-A4
-	D12.776.543.484.500.340 Ephrin-A5
-	D12.776.543.484.500.370 Folate Receptors, GPI-Anchored
-	D12.776.543.484.500.370.500 Folate Receptor 1
-	D12.776.543.484.500.370.750 Folate Receptor 2
-	D12.776.543.484.500.400 Glial Cell Line-Derived Neurotrophic Factor Receptors
-	D12.776.543.484.500.550 Glypicans
New Heading	<b>D12.776.543.484.500.625 Prion Proteins</b>
-	D12.776.543.484.500.700 Matrix Metalloproteinase 17
New Heading	<b>D12.776.543.484.500.738 Nogo Receptors</b>
New Heading	<b>D12.776.543.484.500.738.250 Nogo Receptor 2</b>
New Heading	<b>D12.776.543.484.500.738.500 Nogo Receptor 1</b>
-	D12.776.543.484.500.775 Oligodendrocyte-Myelin Glycoprotein
-	D12.776.543.484.500.850 Uromodulin
-	D12.776.543.488 MARVEL Domain-Containing Proteins
-	D12.776.543.488.249 MARVEL Domain Containing 2 Protein
-	D12.776.543.488.374 Myelin and Lymphocyte-Associated Proteolipid Proteins

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.543.488.500 Occludin
-	D12.776.543.488.750 Synaptogyrins
-	D12.776.543.488.875 Synaptophysin
-	D12.776.543.493 Matrix Metalloproteinases, Membrane-Associated
-	D12.776.543.512 Membrane Fusion Proteins
-	D12.776.543.512.249 SNARE Proteins
-	D12.776.543.512.249.500 Q-SNARE Proteins
-	D12.776.543.512.249.500.500 Qa-SNARE Proteins
-	D12.776.543.512.249.500.500.700 Syntaxin 1
-	D12.776.543.512.249.500.500.750 Syntaxin 16
-	D12.776.543.512.249.500.750 Qb-SNARE Proteins
-	D12.776.543.512.249.500.750.500 Synaptosomal-Associated Protein 25
-	D12.776.543.512.249.500.875 Qc-SNARE Proteins
-	D12.776.543.512.249.500.875.500 Synaptosomal-Associated Protein 25
-	D12.776.543.512.249.750 R-SNARE Proteins
-	D12.776.543.512.249.750.500 Vesicle-Associated Membrane Protein 1
-	D12.776.543.512.249.750.750 Vesicle-Associated Membrane Protein 2
-	D12.776.543.512.249.750.875 Vesicle-Associated Membrane Protein 3
-	D12.776.543.512.500 Viral Fusion Proteins
-	D12.776.543.512.500.330 HIV Envelope Protein gp41
-	D12.776.543.512.500.665 Spike Glycoprotein, Coronavirus
-	D12.776.543.550 Membrane Glycoproteins
-	D12.776.543.550.090 Antigens, CD9
-	D12.776.543.550.147 Antigens, CD47
-	D12.776.543.550.158 Antigens, CD58
-	D12.776.543.550.170 Antigens, CD70
-	D12.776.543.550.182 Antigens, CD82
-	D12.776.543.550.187 Antigens, CD147
-	D12.776.543.550.189 Antigens, Thy-1
-	D12.776.543.550.190 Antiporters
-	D12.776.543.550.190.276 Chloride-Bicarbonate Antiporters
-	D12.776.543.550.190.276.500 Anion Exchange Protein 1, Erythrocyte
-	D12.776.543.550.190.442 Potassium-Hydrogen Antiporters
-	D12.776.543.550.190.608 Reduced Folate Carrier Protein
-	D12.776.543.550.190.775 Sodium-Hydrogen Antiporter
-	D12.776.543.550.192 ATP-Binding Cassette Transporters

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D12.776.543.550.192.304	ATP Binding Cassette Transporter 1
New Heading	<b>D12.776.543.550.192.457 G</b>	<b>ATP Binding Cassette Transporter, Sub-Family G</b>
New Heading	<b>D12.776.543.550.192.457.250 Family G, Member 1</b>	<b>ATP Binding Cassette Transporter, Sub-Family G, Member 1</b>
New Heading	<b>D12.776.543.550.192.457.500 Family G, Member 2</b>	<b>ATP Binding Cassette Transporter, Sub-Family G, Member 2</b>
New Heading	<b>D12.776.543.550.192.457.750 Family G, Member 5</b>	<b>ATP Binding Cassette Transporter, Sub-Family G, Member 5</b>
New Heading	<b>D12.776.543.550.192.457.875 Family G, Member 8</b>	<b>ATP Binding Cassette Transporter, Sub-Family G, Member 8</b>
-	D12.776.543.550.192.610	P-Glycoproteins
New Heading	<b>D12.776.543.550.192.610.305</b>	<b>Antigen Peptide Transporter-1</b>
New Heading	<b>D12.776.543.550.192.610.458</b>	<b>Antigen Peptide Transporter-2</b>
-	D12.776.543.550.192.610.610	P-Glycoprotein
-	D12.776.543.550.193	B-Cell Activating Factor
-	D12.776.543.550.194	4-1BB Ligand
New Heading	<b>D12.776.543.550.195</b>	<b>Butyrophilins</b>
New Tree	<a href="#">D12.776.543.550.195.500</a>	<a href="#">Myelin-Oligodendrocyte Glycoprotein</a>
-	D12.776.543.550.196	CD30 Ligand
-	D12.776.543.550.198	CD40 Ligand
-	D12.776.543.550.200	Cell Adhesion Molecules
New Heading	<b>D12.776.543.550.200.062</b>	<b>Antigens, CD99</b>
-	D12.776.543.550.200.124	Antigens, CD24
-	D12.776.543.550.200.131	Antigens, CD31
-	D12.776.543.550.200.140	Antigens, CD146
-	D12.776.543.550.200.145	Antigens, CD164
-	D12.776.543.550.200.200	Cadherins
-	D12.776.543.550.200.200.500	Desmosomal Cadherins
-	D12.776.543.550.200.200.500.249	Desmocollins
-	D12.776.543.550.200.200.500.500	Desmogleins
-	D12.776.543.550.200.200.500.500.500	Desmoglein 1
-	D12.776.543.550.200.200.500.500.625	Desmoglein 2
-	D12.776.543.550.200.200.500.500.750	Desmoglein 3

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D12.776.543.550.200.210	Carcinoembryonic Antigen
-	D12.776.543.550.200.230	CD4 Immunoadhesins
-	D12.776.543.550.200.250	Cell Adhesion Molecules, Neuronal
-	D12.776.543.550.200.250.150	Cell Adhesion Molecules, Neuron-Glia
-	D12.776.543.550.200.250.150.050 Molecule	Activated-Leukocyte Cell Adhesion
-	D12.776.543.550.200.250.325	Contactins
-	D12.776.543.550.200.250.325.100	Contactin 1
-	D12.776.543.550.200.250.325.200	Contactin 2
-	D12.776.543.550.200.250.500	Myelin P0 Protein
-	D12.776.543.550.200.250.520	Neural Cell Adhesion Molecules
-	D12.776.543.550.200.250.520.156	Antigens, CD56
-	D12.776.543.550.200.250.520.578	Neural Cell Adhesion Molecule L1
-	D12.776.543.550.200.250.520.789	Neurocan
New Heading	<b>D12.776.543.550.200.263</b>	<b>Epithelial Cell Adhesion Molecule</b>
-	D12.776.543.550.200.275	Integrin alphaXbeta2
-	D12.776.543.550.200.275.500	Antigens, CD11c
-	D12.776.543.550.200.275.750	Antigens, CD18
-	D12.776.543.550.200.450	Intercellular Adhesion Molecule-1
-	D12.776.543.550.200.537	Junctional Adhesion Molecules
-	D12.776.543.550.200.537.500 Membrane Protein	Coxsackie and Adenovirus Receptor-Like
-	D12.776.543.550.200.537.750	Junctional Adhesion Molecule A
-	D12.776.543.550.200.537.875	Junctional Adhesion Molecule B
-	D12.776.543.550.200.537.937	Junctional Adhesion Molecule C
-	D12.776.543.550.200.625	Receptors, Lymphocyte Homing
-	D12.776.543.550.200.625.144	Antigens, CD44
-	D12.776.543.550.200.625.347	Integrin alpha4beta1
-	D12.776.543.550.200.625.550	Lymphocyte Function-Associated Antigen-1
-	D12.776.543.550.200.625.903	L-Selectin
-	D12.776.543.550.200.700	Selectins
-	D12.776.543.550.200.700.300	E-Selectin
-	D12.776.543.550.200.700.510	L-Selectin
-	D12.776.543.550.200.700.775	P-Selectin
-	D12.776.543.550.200.810	Sialic Acid Binding Ig-like Lectin 2
-	D12.776.543.550.200.920	Vascular Cell Adhesion Molecule-1

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D12.776.543.550.275	Ectodysplasins
-	D12.776.543.550.312	Fas Ligand Protein
-	D12.776.543.550.350	Fibronectins
-	D12.776.543.550.400	GAP-43 Protein
-	D12.776.543.550.418	GPI-Linked Proteins
-	D12.776.543.550.418.100	Antigens, CD14
-	D12.776.543.550.418.120	Antigens, CD24
-	D12.776.543.550.418.130	Antigens, CD55
-	D12.776.543.550.418.140	Antigens, CD59
-	D12.776.543.550.418.160	Carbonic Anhydrase IV
New Heading	<b>D12.776.543.550.418.180</b>	<b>CD48 Antigen</b>
-	D12.776.543.550.418.200	Ciliary Neurotrophic Factor Receptor alpha Subunit
-	D12.776.543.550.418.250	Contactins
-	D12.776.543.550.418.250.100	Contactin 1
-	D12.776.543.550.418.250.200	Contactin 2
-	D12.776.543.550.418.300	Ephrin-A1
-	D12.776.543.550.418.310	Ephrin-A2
-	D12.776.543.550.418.320	Ephrin-A3
-	D12.776.543.550.418.330	Ephrin-A4
-	D12.776.543.550.418.340	Ephrin-A5
-	D12.776.543.550.418.370	Folate Receptors, GPI-Anchored
-	D12.776.543.550.418.370.500	Folate Receptor 1
-	D12.776.543.550.418.370.750	Folate Receptor 2
-	D12.776.543.550.418.400	Glial Cell Line-Derived Neurotrophic Factor Receptors
-	D12.776.543.550.418.500	Glypicans
New Heading	<b>D12.776.543.550.418.600</b>	<b>Prion Proteins</b>
-	D12.776.543.550.418.700	Matrix Metalloproteinase 17
New Heading	<b>D12.776.543.550.418.738</b>	<b>Nogo Receptors</b>
New Heading	<b>D12.776.543.550.418.738.250</b>	<b>Nogo Receptor 2</b>
New Heading	<b>D12.776.543.550.418.738.500</b>	<b>Nogo Receptor 1</b>
-	D12.776.543.550.418.775	Oligodendrocyte-Myelin Glycoprotein
-	D12.776.543.550.418.850	Uromodulin

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	D12.776.543.550.421                      Histocompatibility Antigens Class I
Old Tree	D12.776.543.550.421.400                HLA-A Antigens
Old Tree	D12.776.543.550.421.400.010            HLA-A1 Antigen
Old Tree	D12.776.543.550.421.400.020            HLA-A2 Antigen
Old Tree	D12.776.543.550.421.400.030            HLA-A3 Antigen
Old Tree	D12.776.543.550.421.400.110            HLA-A11 Antigen
Old Tree	D12.776.543.550.421.400.240            HLA-A24 Antigen
Old Tree	D12.776.543.550.421.500                HLA-B Antigens
Old Tree	D12.776.543.550.421.500.070            HLA-B7 Antigen
Old Tree	D12.776.543.550.421.500.080            HLA-B8 Antigen
Old Tree	D12.776.543.550.421.500.130            HLA-B13 Antigen
Old Tree	D12.776.543.550.421.500.140            HLA-B14 Antigen
Old Tree	D12.776.543.550.421.500.150            HLA-B15 Antigen
Old Tree	D12.776.543.550.421.500.180            HLA-B18 Antigen
Old Tree	D12.776.543.550.421.500.270            HLA-B27 Antigen
Old Tree	D12.776.543.550.421.500.350            HLA-B35 Antigen
Old Tree	D12.776.543.550.421.500.370            HLA-B37 Antigen
Old Tree	D12.776.543.550.421.500.380            HLA-B38 Antigen
Old Tree	D12.776.543.550.421.500.390            HLA-B39 Antigen
Old Tree	D12.776.543.550.421.500.400            HLA-B40 Antigen
Old Tree	D12.776.543.550.421.500.440            HLA-B44 Antigen
Old Tree	D12.776.543.550.421.500.510            HLA-B51 Antigen
Old Tree	D12.776.543.550.421.500.520            HLA-B52 Antigen
Old Tree	D12.776.543.550.421.600                HLA-C Antigens
Old Tree	D12.776.543.550.421.700                HLA-G Antigens
Old Tree	D12.776.543.550.423                    Histocompatibility Antigens Class II
Old Tree	D12.776.543.550.423.400                HLA-D Antigens
Old Tree	D12.776.543.550.423.400.420            HLA-DP Antigens
Old Tree	D12.776.543.550.423.400.420.500        HLA-DP alpha-Chains
Old Tree	D12.776.543.550.423.400.420.750        HLA-DP beta-Chains
Old Tree	D12.776.543.550.423.400.430            HLA-DQ Antigens
Old Tree	D12.776.543.550.423.400.430.500        HLA-DQ alpha-Chains
Old Tree	D12.776.543.550.423.400.430.750        HLA-DQ beta-Chains
Old Tree	D12.776.543.550.423.400.440            HLA-DR Antigens
Old Tree	D12.776.543.550.423.400.440.100        HLA-DR alpha-Chains
Old Tree	D12.776.543.550.423.400.440.200        HLA-DR beta-Chains

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	D12.776.543.550.423.400.440.200.010 HLA-DRB1 Chains
Old Tree	D12.776.543.550.423.400.440.200.030 HLA-DRB3 Chains
Old Tree	D12.776.543.550.423.400.440.200.040 HLA-DRB4 Chains
Old Tree	D12.776.543.550.423.400.440.200.050 HLA-DRB5 Chains
Old Tree	D12.776.543.550.423.400.440.400 HLA-DR Serological Subtypes
Old Tree	D12.776.543.550.423.400.440.400.010 HLA-DR1 Antigen
Old Tree	D12.776.543.550.423.400.440.400.020 HLA-DR2 Antigen
Old Tree	D12.776.543.550.423.400.440.400.030 HLA-DR3 Antigen
Old Tree	D12.776.543.550.423.400.440.400.040 HLA-DR4 Antigen
Old Tree	D12.776.543.550.423.400.440.400.050 HLA-DR5 Antigen
Old Tree	D12.776.543.550.423.400.440.400.060 HLA-DR6 Antigen
Old Tree	D12.776.543.550.423.400.440.400.070 HLA-DR7 Antigen
Old Tree	D12.776.543.550.425 Ion Channels
Old Tree	D12.776.543.550.425.150 Calcium Channels
Old Tree	D12.776.543.550.425.150.400 Calcium Channels, L-Type
Old Tree	D12.776.543.550.425.150.585 Calcium Channels, N-Type
Old Tree	D12.776.543.550.425.150.585.792 Calcium Channels, P-Type
Old Tree	D12.776.543.550.425.150.585.826 Calcium Channels, Q-Type
Old Tree	D12.776.543.550.425.150.585.867 Calcium Channels, R-Type
Old Tree	D12.776.543.550.425.150.720 Calcium Channels, T-Type
Old Tree	D12.776.543.550.425.150.760 Inositol 1,4,5-Trisphosphate Receptors
Old Tree	D12.776.543.550.425.150.800 Ryanodine Receptor Calcium Release Channel
Old Tree	D12.776.543.550.425.175 Chloride Channels
Old Tree	D12.776.543.550.425.175.125 Cystic Fibrosis Transmembrane Conductance Regulator
Old Tree	D12.776.543.550.425.175.562 Receptors, GABA-A
Old Tree	D12.776.543.550.425.175.781 Receptors, Glycine
Old Tree	D12.776.543.550.425.452 Cyclic Nucleotide-Gated Cation Channels
Old Tree	D12.776.543.550.425.476 Hyperpolarization-Activated Cyclic Nucleotide-Gated Channels
Old Tree	D12.776.543.550.425.500 Ligand-Gated Ion Channels
Old Tree	D12.776.543.550.425.500.100 Cysteine Loop Ligand-Gated Ion Channel Receptors
Old Tree	D12.776.543.550.425.500.100.100 Receptors, GABA-A
Old Tree	D12.776.543.550.425.500.100.200 Receptors, Glycine
Old Tree	D12.776.543.550.425.500.100.500 Receptors, Nicotinic
Old Tree	D12.776.543.550.425.500.100.500.500 alpha7 Nicotinic Acetylcholine

## MeSH Tree Changes for 2017

Type	Tree - heading
	<b>Receptor</b>
Old Tree	D12.776.543.550.425.500.100.700                      Receptors, Serotonin, 5-HT3
Old Tree	D12.776.543.550.425.500.200                      Receptors, Ionotropic Glutamate
Old Tree	D12.776.543.550.425.500.200.100                      Receptors, AMPA
Old Tree	D12.776.543.550.425.500.200.200                      Receptors, Kainic Acid
Old Tree	D12.776.543.550.425.500.200.500                      Receptors, N-Methyl-D-Aspartate
Old Tree	D12.776.543.550.425.500.600                      Receptors, Purinergic P2X
Old Tree	D12.776.543.550.425.500.600.100                      Receptors, Purinergic P2X1
Old Tree	D12.776.543.550.425.500.600.200                      Receptors, Purinergic P2X2
Old Tree	D12.776.543.550.425.500.600.300                      Receptors, Purinergic P2X3
Old Tree	D12.776.543.550.425.500.600.400                      Receptors, Purinergic P2X4
Old Tree	D12.776.543.550.425.500.600.500                      Receptors, Purinergic P2X5
Old Tree	D12.776.543.550.425.500.600.700                      Receptors, Purinergic P2X7
Old Tree	D12.776.543.550.425.730                      Porins
Old Tree	D12.776.543.550.425.730.040                      Aquaporins
Old Tree	D12.776.543.550.425.730.040.249                      Aquaglyceroporins
Old Tree	D12.776.543.550.425.730.040.249.500                      Aquaporin 3
Old Tree	D12.776.543.550.425.730.040.249.750                      Aquaporin 6
Old Tree	D12.776.543.550.425.730.040.374                      Aquaporin 1
Old Tree	D12.776.543.550.425.730.040.437                      Aquaporin 2
Old Tree	D12.776.543.550.425.730.040.468                      Aquaporin 4
Old Tree	D12.776.543.550.425.730.040.484                      Aquaporin 5
Old Tree	D12.776.543.550.425.730.520                      Voltage-Dependent Anion Channels
Old Tree	D12.776.543.550.425.730.520.500                      Voltage-Dependent Anion Channel 1
Old Tree	D12.776.543.550.425.730.520.750                      Voltage-Dependent Anion Channel 2
Old Tree	D12.776.543.550.425.750                      Potassium Channels
Old Tree	D12.776.543.550.425.750.150                      Potassium Channels, Calcium-Activated
Old Tree	D12.776.543.550.425.750.150.249 Activated Potassium Channels                      Intermediate-Conductance Calcium-
Old Tree	D12.776.543.550.425.750.150.500 Potassium Channels                      Large-Conductance Calcium-Activated
Old Tree	D12.776.543.550.425.750.150.500.500 Potassium Channel alpha Subunits                      Large-Conductance Calcium-Activated
Old Tree	D12.776.543.550.425.750.150.500.750 Potassium Channel beta Subunits                      Large-Conductance Calcium-Activated
Old Tree	D12.776.543.550.425.750.150.750 Potassium Channels                      Small-Conductance Calcium-Activated



## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	D12.776.543.550.425.750.450 Potassium Channels, Inwardly Rectifying
Old Tree	D12.776.543.550.425.750.450.500 G Protein-Coupled Inwardly-Rectifying Potassium Channels
Old Tree	D12.776.543.550.425.750.450.550 KATP Channels
Old Tree	D12.776.543.550.425.750.450.550.500 Sulfonylurea Receptors
Old Tree	D12.776.543.550.425.750.850 Potassium Channels, Tandem Pore Domain
Old Tree	D12.776.543.550.425.750.900 Potassium Channels, Voltage-Gated
Old Tree	D12.776.543.550.425.750.900.124 Delayed Rectifier Potassium Channels
Old Tree	D12.776.543.550.425.750.900.124.249 KCNQ Potassium Channels
Old Tree	D12.776.543.550.425.750.900.124.249.500 KCNQ1 Potassium Channel
Old Tree	D12.776.543.550.425.750.900.124.249.750 KCNQ2 Potassium Channel
Old Tree	D12.776.543.550.425.750.900.124.249.875 KCNQ3 Potassium Channel
Old Tree	D12.776.543.550.425.750.900.124.311 Kv1.1 Potassium Channel
Old Tree	D12.776.543.550.425.750.900.124.342 Kv1.2 Potassium Channel
Old Tree	D12.776.543.550.425.750.900.124.358 Kv1.3 Potassium Channel
Old Tree	D12.776.543.550.425.750.900.124.374 Kv1.5 Potassium Channel
Old Tree	D12.776.543.550.425.750.900.124.437 Kv1.6 Potassium Channel
Old Tree	D12.776.543.550.425.750.900.124.500 Shab Potassium Channels
Old Tree	D12.776.543.550.425.750.900.249 Ether-A-Go-Go Potassium Channels
Old Tree	D12.776.543.550.425.750.900.500 Shaker Superfamily of Potassium Channels
Old Tree	D12.776.543.550.425.750.900.500.124 Kv1.1 Potassium Channel
Old Tree	D12.776.543.550.425.750.900.500.186 Kv1.2 Potassium Channel
Old Tree	D12.776.543.550.425.750.900.500.233 Kv1.4 Potassium Channel
Old Tree	D12.776.543.550.425.750.900.500.241 Kv1.5 Potassium Channel
Old Tree	D12.776.543.550.425.750.900.500.249 Shab Potassium Channels
Old Tree	D12.776.543.550.425.750.900.500.625 Shal Potassium Channels
Old Tree	D12.776.543.550.425.750.900.500.750 Shaw Potassium Channels
Old Tree	D12.776.543.550.425.875 Sodium Channels
Old Tree	D12.776.543.550.425.875.050 Acid Sensing Ion Channels
Old Tree	D12.776.543.550.425.875.100 Degenerin Sodium Channels
Old Tree	D12.776.543.550.425.875.200 Epithelial Sodium Channels
Old Tree	D12.776.543.550.425.875.750 Voltage-Gated Sodium Channels
Old Tree	D12.776.543.550.425.875.750.100 NAV1.1 Voltage-Gated Sodium Channel
Old Tree	D12.776.543.550.425.875.750.200 NAV1.2 Voltage-Gated Sodium Channel
Old Tree	D12.776.543.550.425.875.750.300 NAV1.3 Voltage-Gated Sodium Channel
Old Tree	D12.776.543.550.425.875.750.400 NAV1.4 Voltage-Gated Sodium Channel

## MeSH Tree Changes for 2017

Type	Tree - heading	
Old Tree	D12.776.543.550.425.875.750.500	NAV1.5 Voltage-Gated Sodium Channel
Old Tree	D12.776.543.550.425.875.750.600	NAV1.6 Voltage-Gated Sodium Channel
Old Tree	D12.776.543.550.425.875.750.700	NAV1.7 Voltage-Gated Sodium Channel
Old Tree	D12.776.543.550.425.875.750.800	NAV1.8 Voltage-Gated Sodium Channel
Old Tree	D12.776.543.550.425.875.750.900	NAV1.9 Voltage-Gated Sodium Channel
Old Tree	D12.776.543.550.425.875.750.970 Subunits	Voltage-Gated Sodium Channel beta
Old Tree	D12.776.543.550.425.875.750.970.100 Subunit	Voltage-Gated Sodium Channel beta-1
Old Tree	D12.776.543.550.425.875.750.970.200 Subunit	Voltage-Gated Sodium Channel beta-2
Old Tree	D12.776.543.550.425.875.750.970.300 Subunit	Voltage-Gated Sodium Channel beta-3
Old Tree	D12.776.543.550.425.875.750.970.400 Subunit	Voltage-Gated Sodium Channel beta-4
New Heading	<b>D12.776.543.550.435</b>	<b>Hepatitis A Virus Cellular Receptor 1</b>
New Heading	<b>D12.776.543.550.437</b>	<b>Hepatitis A Virus Cellular Receptor 2</b>
New Tree	D12.776.543.550.439	Histocompatibility Antigens Class I
New Heading	<b>D12.776.543.550.439.200</b>	<b>Hemochromatosis Protein</b>
New Tree	D12.776.543.550.439.400	HLA-A Antigens
New Tree	D12.776.543.550.439.400.010	HLA-A1 Antigen
New Tree	D12.776.543.550.439.400.020	HLA-A2 Antigen
New Tree	D12.776.543.550.439.400.030	HLA-A3 Antigen
New Tree	D12.776.543.550.439.400.110	HLA-A11 Antigen
New Tree	D12.776.543.550.439.400.240	HLA-A24 Antigen
New Tree	D12.776.543.550.439.500	HLA-B Antigens
New Tree	D12.776.543.550.439.500.070	HLA-B7 Antigen
New Tree	D12.776.543.550.439.500.080	HLA-B8 Antigen
New Tree	D12.776.543.550.439.500.130	HLA-B13 Antigen

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	D12.776.543.550.439.500.140 HLA-B14 Antigen
New Tree	D12.776.543.550.439.500.150 HLA-B15 Antigen
New Tree	D12.776.543.550.439.500.180 HLA-B18 Antigen
New Tree	D12.776.543.550.439.500.270 HLA-B27 Antigen
New Tree	D12.776.543.550.439.500.350 HLA-B35 Antigen
New Tree	D12.776.543.550.439.500.370 HLA-B37 Antigen
New Tree	D12.776.543.550.439.500.380 HLA-B38 Antigen
New Tree	D12.776.543.550.439.500.390 HLA-B39 Antigen
New Tree	D12.776.543.550.439.500.400 HLA-B40 Antigen
New Tree	D12.776.543.550.439.500.440 HLA-B44 Antigen
New Tree	D12.776.543.550.439.500.510 HLA-B51 Antigen
New Tree	D12.776.543.550.439.500.520 HLA-B52 Antigen
New Tree	D12.776.543.550.439.600 HLA-C Antigens
New Tree	D12.776.543.550.439.700 HLA-G Antigens
New Tree	D12.776.543.550.440 Histocompatibility Antigens Class II
New Tree	D12.776.543.550.440.400 HLA-D Antigens
New Tree	D12.776.543.550.440.400.420 HLA-DP Antigens
New Tree	D12.776.543.550.440.400.420.500 HLA-DP alpha-Chains
New Tree	D12.776.543.550.440.400.420.750 HLA-DP beta-Chains
New Tree	D12.776.543.550.440.400.430 HLA-DQ Antigens
New Tree	D12.776.543.550.440.400.430.500 HLA-DQ alpha-Chains
New Tree	D12.776.543.550.440.400.430.750 HLA-DQ beta-Chains

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	D12.776.543.550.440.400.440 HLA-DR Antigens
New Tree	D12.776.543.550.440.400.440.100 HLA-DR alpha-Chains
New Tree	D12.776.543.550.440.400.440.200 HLA-DR beta-Chains
New Tree	D12.776.543.550.440.400.440.200.010 HLA-DRB1 Chains
New Tree	D12.776.543.550.440.400.440.200.030 HLA-DRB3 Chains
New Tree	D12.776.543.550.440.400.440.200.040 HLA-DRB4 Chains
New Tree	D12.776.543.550.440.400.440.200.050 HLA-DRB5 Chains
New Tree	D12.776.543.550.440.400.440.400 HLA-DR Serological Subtypes
New Tree	D12.776.543.550.440.400.440.400.010 HLA-DR1 Antigen
New Tree	D12.776.543.550.440.400.440.400.020 HLA-DR2 Antigen
New Tree	D12.776.543.550.440.400.440.400.030 HLA-DR3 Antigen
New Tree	D12.776.543.550.440.400.440.400.040 HLA-DR4 Antigen
New Tree	D12.776.543.550.440.400.440.400.050 HLA-DR5 Antigen
New Tree	D12.776.543.550.440.400.440.400.060 HLA-DR6 Antigen
New Tree	D12.776.543.550.440.400.440.400.070 HLA-DR7 Antigen
New Tree	D12.776.543.550.450 Ion Channels
New Tree	D12.776.543.550.450.150 Calcium Channels
New Tree	D12.776.543.550.450.150.400 Calcium Channels, L-Type
New Tree	D12.776.543.550.450.150.585 Calcium Channels, N-Type
New Tree	D12.776.543.550.450.150.585.792 Calcium Channels, P-Type
New Tree	D12.776.543.550.450.150.585.826 Calcium Channels, Q-Type
New Tree	D12.776.543.550.450.150.585.867 Calcium Channels, R-Type

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	D12.776.543.550.450.150.720	Calcium Channels, T-Type
New Heading	<b>D12.776.543.550.450.150.740 Channels</b>	<b>Calcium Release Activated Calcium</b>
New Heading	<b>D12.776.543.550.450.150.740.500</b>	<b>ORAI1 Protein</b>
New Heading	<b>D12.776.543.550.450.150.740.750</b>	<b>ORAI2 Protein</b>
New Tree	D12.776.543.550.450.150.760	Inositol 1,4,5-Trisphosphate Receptors
New Tree	D12.776.543.550.450.150.800	Ryanodine Receptor Calcium Release Channel
New Tree	D12.776.543.550.450.175	Chloride Channels
New Tree	D12.776.543.550.450.175.125 Regulator	Cystic Fibrosis Transmembrane Conductance
New Tree	D12.776.543.550.450.175.562	Receptors, GABA-A
New Tree	D12.776.543.550.450.175.781	Receptors, Glycine
New Tree	D12.776.543.550.450.452	Cyclic Nucleotide-Gated Cation Channels
New Tree	D12.776.543.550.450.476 Channels	Hyperpolarization-Activated Cyclic Nucleotide-Gated
New Tree	D12.776.543.550.450.500	Ligand-Gated Ion Channels
New Tree	D12.776.543.550.450.500.100 Receptors	Cysteine Loop Ligand-Gated Ion Channel
New Tree	D12.776.543.550.450.500.100.100	Receptors, GABA-A
New Tree	D12.776.543.550.450.500.100.200	Receptors, Glycine
New Tree	D12.776.543.550.450.500.100.500	Receptors, Nicotinic
New Tree	D12.776.543.550.450.500.100.500.500 Receptor	alpha7 Nicotinic Acetylcholine
New Tree	D12.776.543.550.450.500.100.700	Receptors, Serotonin, 5-HT3
New Tree	D12.776.543.550.450.500.200	Receptors, Ionotropic Glutamate
New Tree	D12.776.543.550.450.500.200.100	Receptors, AMPA
New Tree	D12.776.543.550.450.500.200.200	Receptors, Kainic Acid

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	D12.776.543.550.450.500.200.500      Receptors, N-Methyl-D-Aspartate
New Tree	D12.776.543.550.450.500.600      Receptors, Purinergic P2X
New Tree	D12.776.543.550.450.500.600.100      Receptors, Purinergic P2X1
New Tree	D12.776.543.550.450.500.600.200      Receptors, Purinergic P2X2
New Tree	D12.776.543.550.450.500.600.300      Receptors, Purinergic P2X3
New Tree	D12.776.543.550.450.500.600.400      Receptors, Purinergic P2X4
New Tree	D12.776.543.550.450.500.600.500      Receptors, Purinergic P2X5
New Tree	D12.776.543.550.450.500.600.700      Receptors, Purinergic P2X7
New Tree	D12.776.543.550.450.730      Porins
New Tree	D12.776.543.550.450.730.040      Aquaporins
New Tree	D12.776.543.550.450.730.040.249      Aquaglyceroporins
New Tree	D12.776.543.550.450.730.040.249.500      Aquaporin 3
New Tree	D12.776.543.550.450.730.040.249.750      Aquaporin 6
New Tree	D12.776.543.550.450.730.040.374      Aquaporin 1
New Tree	D12.776.543.550.450.730.040.437      Aquaporin 2
New Tree	D12.776.543.550.450.730.040.468      Aquaporin 4
New Tree	D12.776.543.550.450.730.040.484      Aquaporin 5
New Tree	D12.776.543.550.450.730.520      Voltage-Dependent Anion Channels
New Tree	D12.776.543.550.450.730.520.500      Voltage-Dependent Anion Channel 1
New Tree	D12.776.543.550.450.730.520.750      Voltage-Dependent Anion Channel 2
New Tree	D12.776.543.550.450.750      Potassium Channels
New Tree	D12.776.543.550.450.750.150      Potassium Channels, Calcium-Activated

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	D12.776.543.550.450.750.150.249 Intermediate-Conductance Calcium-Activated Potassium Channels
New Tree	D12.776.543.550.450.750.150.500 Large-Conductance Calcium-Activated Potassium Channels
New Tree	D12.776.543.550.450.750.150.500.500 Large-Conductance Calcium-Activated Potassium Channel alpha Subunits
New Tree	D12.776.543.550.450.750.150.500.750 Large-Conductance Calcium-Activated Potassium Channel beta Subunits
New Tree	D12.776.543.550.450.750.150.750 Small-Conductance Calcium-Activated Potassium Channels
New Tree	D12.776.543.550.450.750.450 Potassium Channels, Inwardly Rectifying
New Tree	D12.776.543.550.450.750.450.500 G Protein-Coupled Inwardly-Rectifying Potassium Channels
New Tree	D12.776.543.550.450.750.450.550 KATP Channels
New Tree	D12.776.543.550.450.750.450.550.500 Sulfonylurea Receptors
New Tree	D12.776.543.550.450.750.850 Potassium Channels, Tandem Pore Domain
New Tree	D12.776.543.550.450.750.900 Potassium Channels, Voltage-Gated
New Tree	D12.776.543.550.450.750.900.124 Delayed Rectifier Potassium Channels
New Tree	D12.776.543.550.450.750.900.124.249 KCNQ Potassium Channels
New Tree	D12.776.543.550.450.750.900.124.249.500 KCNQ1 Potassium Channel
New Tree	D12.776.543.550.450.750.900.124.249.750 KCNQ2 Potassium Channel
New Tree	D12.776.543.550.450.750.900.124.249.875 KCNQ3 Potassium Channel
New Tree	D12.776.543.550.450.750.900.124.311 Kv1.1 Potassium Channel
New Tree	D12.776.543.550.450.750.900.124.342 Kv1.2 Potassium Channel
New Tree	D12.776.543.550.450.750.900.124.358 Kv1.3 Potassium Channel
New Tree	D12.776.543.550.450.750.900.124.374 Kv1.5 Potassium Channel
New Tree	D12.776.543.550.450.750.900.124.437 Kv1.6 Potassium Channel
New Tree	D12.776.543.550.450.750.900.124.500 Shab Potassium Channels

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	D12.776.543.550.450.750.900.249	Ether-A-Go-Go Potassium Channels
New Heading	<b>D12.776.543.550.450.750.900.249.500</b>	<b>ERG1 Potassium Channel</b>
New Tree	D12.776.543.550.450.750.900.500	Shaker Superfamily of Potassium Channels
New Tree	D12.776.543.550.450.750.900.500.124	Kv1.1 Potassium Channel
New Tree	D12.776.543.550.450.750.900.500.186	Kv1.2 Potassium Channel
New Tree	D12.776.543.550.450.750.900.500.233	Kv1.4 Potassium Channel
New Tree	D12.776.543.550.450.750.900.500.241	Kv1.5 Potassium Channel
New Tree	D12.776.543.550.450.750.900.500.249	Shab Potassium Channels
New Tree	D12.776.543.550.450.750.900.500.625	Shal Potassium Channels
New Tree	D12.776.543.550.450.750.900.500.750	Shaw Potassium Channels
New Tree	D12.776.543.550.450.875	Sodium Channels
New Tree	D12.776.543.550.450.875.050	Acid Sensing Ion Channels
New Tree	D12.776.543.550.450.875.100	Degenerin Sodium Channels
New Tree	D12.776.543.550.450.875.200	Epithelial Sodium Channels
New Tree	D12.776.543.550.450.875.750	Voltage-Gated Sodium Channels
New Tree	D12.776.543.550.450.875.750.100	NAV1.1 Voltage-Gated Sodium Channel
New Tree	D12.776.543.550.450.875.750.200	NAV1.2 Voltage-Gated Sodium Channel
New Tree	D12.776.543.550.450.875.750.300	NAV1.3 Voltage-Gated Sodium Channel
New Tree	D12.776.543.550.450.875.750.400	NAV1.4 Voltage-Gated Sodium Channel
New Tree	D12.776.543.550.450.875.750.500	NAV1.5 Voltage-Gated Sodium Channel
New Tree	D12.776.543.550.450.875.750.600	NAV1.6 Voltage-Gated Sodium Channel
New Tree	D12.776.543.550.450.875.750.700	NAV1.7 Voltage-Gated Sodium Channel



## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	D12.776.543.550.450.875.750.800	NAV1.8 Voltage-Gated Sodium Channel
New Tree	D12.776.543.550.450.875.750.900	NAV1.9 Voltage-Gated Sodium Channel
New Tree	D12.776.543.550.450.875.750.970 Subunits	Voltage-Gated Sodium Channel beta
New Tree	D12.776.543.550.450.875.750.970.100 Subunit	Voltage-Gated Sodium Channel beta-1
New Tree	D12.776.543.550.450.875.750.970.200 Subunit	Voltage-Gated Sodium Channel beta-2
New Tree	D12.776.543.550.450.875.750.970.300 Subunit	Voltage-Gated Sodium Channel beta-3
New Tree	D12.776.543.550.450.875.750.970.400 Subunit	Voltage-Gated Sodium Channel beta-4
-	D12.776.543.550.500	Laminin
-	D12.776.543.550.527	Lysosome-Associated Membrane Glycoproteins
-	D12.776.543.550.527.249	Antigens, CD63
-	D12.776.543.550.527.500	Lysosomal-Associated Membrane Protein 1
-	D12.776.543.550.527.750	Lysosomal-Associated Membrane Protein 2
-	D12.776.543.550.527.875	Lysosomal-Associated Membrane Protein 3
-	D12.776.543.550.530	Mucin-1
-	D12.776.543.550.542	Mucin-3
-	D12.776.543.550.555	Myelin-Associated Glycoprotein
Old Tree	<del>D12.776.543.550.572</del>	<del>Myelin-Oligodendrocyte Glycoprotein</del>
-	D12.776.543.550.590	OX40 Ligand
-	D12.776.543.550.607	PHEX Phosphate Regulating Neutral Endopeptidase
-	D12.776.543.550.625	Platelet Membrane Glycoproteins
-	D12.776.543.550.625.136	Antigens, CD36
-	D12.776.543.550.625.298	Integrin alpha2beta1
-	D12.776.543.550.625.379	Integrin alpha5beta1
-	D12.776.543.550.625.419	Integrin alpha6beta1
-	D12.776.543.550.625.439	Integrin alphaVbeta3
-	D12.776.543.550.625.449	Lysosomal-Associated Membrane Protein 1
-	D12.776.543.550.625.460	Platelet Glycoprotein GPIb-IX Complex
-	D12.776.543.550.625.785	Platelet Glycoprotein GPIIb-IIIa Complex
-	D12.776.543.550.625.800	Receptors, Thrombin
-	D12.776.543.550.625.800.790	Receptor, PAR-1
-	D12.776.543.550.625.800.800	Thrombomodulin

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.543.550.625.905 P-Selectin
-	D12.776.543.550.712 Sarcoglycans
New Heading	<b>D12.776.543.550.746 Signaling Lymphocytic Activation Molecule Family</b>
New Heading	<b>D12.776.543.550.746.250 CD48 Antigen</b>
New Heading	<b>D12.776.543.550.746.500 Signaling Lymphocytic Activation Molecule Family Member 1</b>
-	D12.776.543.550.779 SLC4A Proteins
-	D12.776.543.550.779.249 Chloride-Bicarbonate Antiporters
-	D12.776.543.550.779.249.500 Anion Exchange Protein 1, Erythrocyte
-	D12.776.543.550.779.500 Sodium-Bicarbonate Symporters
-	D12.776.543.550.847 Syndecans
-	D12.776.543.550.847.100 Syndecan-1
-	D12.776.543.550.847.200 Syndecan-2
-	D12.776.543.550.847.300 Syndecan-3
-	D12.776.543.550.847.400 Syndecan-4
-	D12.776.543.550.895 Thrombospondins
-	D12.776.543.550.895.800 Thrombospondin 1
-	D12.776.543.550.942 Uroplakins
-	D12.776.543.550.942.100 Uroplakin Ia
-	D12.776.543.550.942.150 Uroplakin Ib
-	D12.776.543.550.942.200 Uroplakin II
-	D12.776.543.550.942.300 Uroplakin III
-	D12.776.543.550.990 Variant Surface Glycoproteins, Trypanosoma
New Heading	<b>D12.776.543.550.995 Zona Pellucida Glycoproteins</b>
-	D12.776.543.585 Membrane Transport Proteins
-	D12.776.543.585.100 ATP-Binding Cassette Transporters
-	D12.776.543.585.100.151 ATP Binding Cassette Transporter 1
New Heading	<b>D12.776.543.585.100.228 ATP Binding Cassette Transporter, Sub-Family G</b>
New Heading	<b>D12.776.543.585.100.228.250 ATP Binding Cassette Transporter, Sub-Family G, Member 1</b>
New Heading	<b>D12.776.543.585.100.228.500 ATP Binding Cassette Transporter, Sub-Family G, Member 2</b>
New Heading	<b>D12.776.543.585.100.228.750 ATP Binding Cassette Transporter, Sub-Family G, Member 5</b>
New	<b>D12.776.543.585.100.228.875 ATP Binding Cassette Transporter, Sub-</b>

## MeSH Tree Changes for 2017

Type	Tree - heading
Heading	<b>Family G, Member 8</b>
-	D12.776.543.585.100.304                      Multidrug Resistance-Associated Proteins
New Tree	<a href="#">D12.776.543.585.100.304.500</a> <a href="#">Cystic Fibrosis Transmembrane Conductance Regulator</a>
-	D12.776.543.585.100.610                      P-Glycoproteins
New Heading	<b>D12.776.543.585.100.610.250</b> <b>Antigen Peptide Transporter-1</b>
New Heading	<b>D12.776.543.585.100.610.375</b> <b>Antigen Peptide Transporter-2</b>
-	D12.776.543.585.100.610.500                      P-Glycoprotein
-	D12.776.543.585.100.805                      Sulfonylurea Receptors
-	D12.776.543.585.200                      Amino Acid Transport Systems
-	D12.776.543.585.200.249                      Amino Acid Transport Systems, Acidic
-	D12.776.543.585.200.249.500                      Amino Acid Transport System X-AG
-	D12.776.543.585.200.249.500.500                      Glutamate Plasma Membrane Transport Proteins
-	D12.776.543.585.200.249.500.500.500                      Excitatory Amino Acid Transporter 1
-	D12.776.543.585.200.249.500.500.750                      Excitatory Amino Acid Transporter 2
-	D12.776.543.585.200.249.500.500.875                      Excitatory Amino Acid Transporter 3
-	D12.776.543.585.200.249.500.500.937                      Excitatory Amino Acid Transporter 4
-	D12.776.543.585.200.249.500.500.968                      Excitatory Amino Acid Transporter 5
-	D12.776.543.585.200.374                      Amino Acid Transport Systems, Basic
-	D12.776.543.585.200.374.600                      Amino Acid Transport System $\gamma$ +
-	D12.776.543.585.200.374.600.200                      Cationic Amino Acid Transporter 1
-	D12.776.543.585.200.374.600.300                      Cationic Amino Acid Transporter 2
-	D12.776.543.585.200.374.750                      Amino Acid Transport System $\gamma$ +L
-	D12.776.543.585.200.374.750.500                      Antigens, CD98
-	D12.776.543.585.200.374.750.500.250                      Antigens, CD98 Heavy Chain
-	D12.776.543.585.200.374.750.500.500                      Antigens, CD98 Light Chains
-	D12.776.543.585.200.500                      Amino Acid Transport Systems, Neutral
-	D12.776.543.585.200.500.100                      Amino Acid Transport System A
-	D12.776.543.585.200.500.200                      Amino Acid Transport System ASC
-	D12.776.543.585.200.500.500                      Amino Acid Transport System L
-	D12.776.543.585.200.500.500.500                      Antigens, CD98
-	D12.776.543.585.200.500.500.500.250                      Antigens, CD98 Heavy Chain
-	D12.776.543.585.200.500.500.500.500                      Antigens, CD98 Light Chains
-	D12.776.543.585.200.500.500.500.500.500                      Large Neutral Amino Acid-

## MeSH Tree Changes for 2017

Type	Tree - heading	
	Transporter 1	
-	D12.776.543.585.200.500.875	Glycine Plasma Membrane Transport Proteins
-	D12.776.543.585.250	Connexins
New Heading	<b>D12.776.543.585.250.100</b>	<b>Connexin 26</b>
-	D12.776.543.585.250.200	Connexin 43
-	D12.776.543.585.300	Fatty Acid Transport Proteins
-	D12.776.543.585.300.500	Antigens, CD36
-	D12.776.543.585.400	Ion Channels
-	D12.776.543.585.400.150	Calcium Channels
-	D12.776.543.585.400.150.400	Calcium Channels, L-Type
-	D12.776.543.585.400.150.585	Calcium Channels, N-Type
-	D12.776.543.585.400.150.585.792	Calcium Channels, P-Type
-	D12.776.543.585.400.150.585.826	Calcium Channels, Q-Type
-	D12.776.543.585.400.150.585.867	Calcium Channels, R-Type
-	D12.776.543.585.400.150.720	Calcium Channels, T-Type
New Heading	<b>D12.776.543.585.400.150.740 Channels</b>	<b>Calcium Release Activated Calcium</b>
New Heading	<b>D12.776.543.585.400.150.740.500</b>	<b>ORAI1 Protein</b>
New Heading	<b>D12.776.543.585.400.150.740.750</b>	<b>ORAI2 Protein</b>
-	D12.776.543.585.400.150.760	Inositol 1,4,5-Trisphosphate Receptors
-	D12.776.543.585.400.150.800	Ryanodine Receptor Calcium Release Channel
-	D12.776.543.585.400.150.900	TRPP Cation Channels
-	D12.776.543.585.400.175	Chloride Channels
-	D12.776.543.585.400.175.125 Regulator	Cystic Fibrosis Transmembrane Conductance
-	D12.776.543.585.400.175.562	Receptors, GABA-A
-	D12.776.543.585.400.175.781	Receptors, Glycine
-	D12.776.543.585.400.452	Cyclic Nucleotide-Gated Cation Channels
-	D12.776.543.585.400.476 Channels	Hyperpolarization-Activated Cyclic Nucleotide-Gated
-	D12.776.543.585.400.500	Ligand-Gated Ion Channels
-	D12.776.543.585.400.500.100 Receptors	Cysteine Loop Ligand-Gated Ion Channel
-	D12.776.543.585.400.500.100.100	Receptors, GABA-A
-	D12.776.543.585.400.500.100.200	Receptors, Glycine

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.543.585.400.500.100.500 Receptors, Nicotinic
-	D12.776.543.585.400.500.100.500.500 Receptor alpha7 Nicotinic Acetylcholine
-	D12.776.543.585.400.500.100.700 Receptors, Serotonin, 5-HT3
-	D12.776.543.585.400.500.200 Receptors, Ionotropic Glutamate
-	D12.776.543.585.400.500.200.100 Receptors, AMPA
-	D12.776.543.585.400.500.200.200 Receptors, Kainic Acid
-	D12.776.543.585.400.500.200.500 Receptors, N-Methyl-D-Aspartate
-	D12.776.543.585.400.500.600 Receptors, Purinergic P2X
-	D12.776.543.585.400.500.600.100 Receptors, Purinergic P2X1
-	D12.776.543.585.400.500.600.200 Receptors, Purinergic P2X2
-	D12.776.543.585.400.500.600.300 Receptors, Purinergic P2X3
-	D12.776.543.585.400.500.600.400 Receptors, Purinergic P2X4
-	D12.776.543.585.400.500.600.500 Receptors, Purinergic P2X5
-	D12.776.543.585.400.500.600.700 Receptors, Purinergic P2X7
-	D12.776.543.585.400.730 Porins
-	D12.776.543.585.400.730.040 Aquaporins
-	D12.776.543.585.400.730.040.249 Aquaglyceroporins
-	D12.776.543.585.400.730.040.249.500 Aquaporin 3
-	D12.776.543.585.400.730.040.249.750 Aquaporin 6
-	D12.776.543.585.400.730.040.436 Aquaporin 1
-	D12.776.543.585.400.730.040.530 Aquaporin 2
-	D12.776.543.585.400.730.040.577 Aquaporin 4
-	D12.776.543.585.400.730.040.600 Aquaporin 5
-	D12.776.543.585.400.730.520 Voltage-Dependent Anion Channels
-	D12.776.543.585.400.730.520.500 Voltage-Dependent Anion Channel 1
-	D12.776.543.585.400.730.520.750 Voltage-Dependent Anion Channel 2
-	D12.776.543.585.400.750 Potassium Channels
-	D12.776.543.585.400.750.150 Potassium Channels, Calcium-Activated
-	D12.776.543.585.400.750.150.249 Intermediate-Conductance Calcium-Activated Potassium Channels
-	D12.776.543.585.400.750.150.500 Large-Conductance Calcium-Activated Potassium Channels
-	D12.776.543.585.400.750.150.500.249 Large-Conductance Calcium-Activated Potassium Channel alpha Subunits
-	D12.776.543.585.400.750.150.500.500 Large-Conductance Calcium-Activated Potassium Channel beta Subunits

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.543.585.400.750.150.750 Small-Conductance Calcium-Activated Potassium Channels
-	D12.776.543.585.400.750.450 Potassium Channels, Inwardly Rectifying
-	D12.776.543.585.400.750.450.500 G Protein-Coupled Inwardly-Rectifying Potassium Channels
-	D12.776.543.585.400.750.450.550 KATP Channels
-	D12.776.543.585.400.750.450.550.500 Sulfonylurea Receptors
-	D12.776.543.585.400.750.850 Potassium Channels, Tandem Pore Domain
-	D12.776.543.585.400.750.900 Potassium Channels, Voltage-Gated
-	D12.776.543.585.400.750.900.124 Delayed Rectifier Potassium Channels
-	D12.776.543.585.400.750.900.124.249 KCNQ Potassium Channels
-	D12.776.543.585.400.750.900.124.249.500 KCNQ1 Potassium Channel
-	D12.776.543.585.400.750.900.124.249.750 KCNQ2 Potassium Channel
-	D12.776.543.585.400.750.900.124.249.875 KCNQ3 Potassium Channel
-	D12.776.543.585.400.750.900.124.374 Kv1.5 Potassium Channel
-	D12.776.543.585.400.750.900.124.500 Shab Potassium Channels
-	D12.776.543.585.400.750.900.249 Ether-A-Go-Go Potassium Channels
New Heading	<b>D12.776.543.585.400.750.900.249.500</b> <b>ERG1 Potassium Channel</b>
-	D12.776.543.585.400.750.900.624 Shaker Superfamily of Potassium Channels
-	D12.776.543.585.400.750.900.624.124 Kv1.1 Potassium Channel
-	D12.776.543.585.400.750.900.624.186 Kv1.2 Potassium Channel
-	D12.776.543.585.400.750.900.624.217 Kv1.3 Potassium Channel
-	D12.776.543.585.400.750.900.624.233 Kv1.4 Potassium Channel
-	D12.776.543.585.400.750.900.624.241 Kv1.5 Potassium Channel
-	D12.776.543.585.400.750.900.624.245 Kv1.6 Potassium Channel
-	D12.776.543.585.400.750.900.624.249 Shab Potassium Channels
-	D12.776.543.585.400.750.900.624.625 Shal Potassium Channels
-	D12.776.543.585.400.750.900.624.750 Shaw Potassium Channels
-	D12.776.543.585.400.875 Sodium Channels
-	D12.776.543.585.400.875.050 Acid Sensing Ion Channels
-	D12.776.543.585.400.875.100 Degenerin Sodium Channels
-	D12.776.543.585.400.875.200 Epithelial Sodium Channels
-	D12.776.543.585.400.875.750 Voltage-Gated Sodium Channels
-	D12.776.543.585.400.875.750.100 NAV1.1 Voltage-Gated Sodium Channel
-	D12.776.543.585.400.875.750.200 NAV1.2 Voltage-Gated Sodium Channel
-	D12.776.543.585.400.875.750.300 NAV1.3 Voltage-Gated Sodium Channel

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D12.776.543.585.400.875.750.400	NAV1.4 Voltage-Gated Sodium Channel
-	D12.776.543.585.400.875.750.500	NAV1.5 Voltage-Gated Sodium Channel
-	D12.776.543.585.400.875.750.600	NAV1.6 Voltage-Gated Sodium Channel
-	D12.776.543.585.400.875.750.700	NAV1.7 Voltage-Gated Sodium Channel
-	D12.776.543.585.400.875.750.800	NAV1.8 Voltage-Gated Sodium Channel
-	D12.776.543.585.400.875.750.900	NAV1.9 Voltage-Gated Sodium Channel
-	D12.776.543.585.400.875.750.970 Subunits	Voltage-Gated Sodium Channel beta
-	D12.776.543.585.400.875.750.970.100 Subunit	Voltage-Gated Sodium Channel beta-1
-	D12.776.543.585.400.875.750.970.200 Subunit	Voltage-Gated Sodium Channel beta-2
-	D12.776.543.585.400.875.750.970.300 Subunit	Voltage-Gated Sodium Channel beta-3
-	D12.776.543.585.400.875.750.970.650 Subunit	Voltage-Gated Sodium Channel beta-4
-	D12.776.543.585.400.901	Transient Receptor Potential Channels
-	D12.776.543.585.400.901.500	TRPC Cation Channels
-	D12.776.543.585.400.901.555	TRPM Cation Channels
-	D12.776.543.585.400.901.777	TRPP Cation Channels
-	D12.776.543.585.450	Ion Pumps
-	D12.776.543.585.450.074	Anion Transport Proteins
-	D12.776.543.585.450.074.249	Halorhodopsins
-	D12.776.543.585.450.074.500	Organic Anion Transporters
-	D12.776.543.585.450.074.500.149	Dicarboxylic Acid Transporters
-	D12.776.543.585.450.074.500.224	Folic Acid Transporters
-	D12.776.543.585.450.074.500.224.500	Folate Receptors, GPI-Anchored
-	D12.776.543.585.450.074.500.224.500.500	Folate Receptor 1
-	D12.776.543.585.450.074.500.224.500.750	Folate Receptor 2
-	D12.776.543.585.450.074.500.224.625	Proton-Coupled Folate Transporter
-	D12.776.543.585.450.074.500.224.750	Reduced Folate Carrier Protein
-	D12.776.543.585.450.074.500.300	Monocarboxylic Acid Transporters
-	D12.776.543.585.450.074.500.500 Dependent	Organic Anion Transporters, ATP-
-	D12.776.543.585.450.074.500.500.500 Proteins	Multidrug Resistance-Associated
New Tree	<a href="#">D12.776.543.585.450.074.500.500.500.500</a> <a href="#">Conductance Regulator</a>	<a href="#">Cystic Fibrosis Transmembrane</a>
Old Tree	<a href="#">D12.776.543.585.450.074.500.500.750</a>	<a href="#">P-Glycoprotein</a>

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D12.776.543.585.450.074.500.500.875	P-Glycoproteins
New Heading	<b>D12.776.543.585.450.074.500.500.875.250</b>	<b>Antigen Peptide Transporter-1</b>
New Heading	<b>D12.776.543.585.450.074.500.500.875.375</b>	<b>Antigen Peptide Transporter-2</b>
New Tree	<b>D12.776.543.585.450.074.500.500.875.500</b>	<b>P-Glycoprotein</b>
-	D12.776.543.585.450.074.500.812 Dependent	Organic Anion Transporters, Sodium-
-	D12.776.543.585.450.074.500.812.500 Transporters	Sodium-Coupled Vitamin C
-	D12.776.543.585.450.074.500.875 Independent	Organic Anion Transporters, Sodium-
-	D12.776.543.585.450.074.500.875.500	Organic Anion Transport Polypeptide C
-	D12.776.543.585.450.074.500.875.500 Transporter Family Member 1b1	Solute Carrier Organic Anion
-	D12.776.543.585.450.074.500.875.750	Organic Anion Transport Protein 1
-	D12.776.543.585.450.074.750	Phosphate Transport Proteins
-	D12.776.543.585.450.074.750.500	Proton-Phosphate Symporters
-	D12.776.543.585.450.074.750.750	Sodium-Phosphate Cotransporter Proteins
-	D12.776.543.585.450.074.750.750.500 Proteins, Type I	Sodium-Phosphate Cotransporter
-	D12.776.543.585.450.074.750.750.750 Proteins, Type II	Sodium-Phosphate Cotransporter
-	D12.776.543.585.450.074.750.750.750.124 Proteins, Type IIa	Sodium-Phosphate Cotransporter
-	D12.776.543.585.450.074.750.750.750.249 Proteins, Type IIb	Sodium-Phosphate Cotransporter
-	D12.776.543.585.450.074.750.750.750.500 Proteins, Type IIc	Sodium-Phosphate Cotransporter
-	D12.776.543.585.450.074.750.750.875 Proteins, Type III	Sodium-Phosphate Cotransporter
-	D12.776.543.585.450.162	Antiporters
-	D12.776.543.585.450.162.193	Chloride-Bicarbonate Antiporters
-	D12.776.543.585.450.162.193.500	Anion Exchange Protein 1, Erythrocyte
-	D12.776.543.585.450.162.225	Mitochondrial ADP, ATP Translocases
-	D12.776.543.585.450.162.225.100	Adenine Nucleotide Translocator 1
-	D12.776.543.585.450.162.225.200	Adenine Nucleotide Translocator 2
-	D12.776.543.585.450.162.225.300	Adenine Nucleotide Translocator 3
-	D12.776.543.585.450.162.234	Organic Anion Transport Protein 1



## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D12.776.543.585.450.162.276	Potassium-Hydrogen Antiporters
-	D12.776.543.585.450.162.359	Reduced Folate Carrier Protein
-	D12.776.543.585.450.162.442	Sodium-Calcium Exchanger
-	D12.776.543.585.450.162.775	Sodium-Hydrogen Antiporter
-	D12.776.543.585.450.162.800	Sodium-Potassium-Exchanging ATPase
-	D12.776.543.585.450.162.887	Vesicular Neurotransmitter Transport Proteins
-	D12.776.543.585.450.162.887.500 Proteins	Vesicular Biogenic Amine Transport
-	D12.776.543.585.450.162.887.500.249 Proteins	Vesicular Acetylcholine Transport
-	D12.776.543.585.450.162.887.500.500 Proteins	Vesicular Monoamine Transport
-	D12.776.543.585.450.162.887.625	Vesicular Glutamate Transport Proteins
-	D12.776.543.585.450.162.887.625.500 1	Vesicular Glutamate Transport Protein
-	D12.776.543.585.450.162.887.625.750 2	Vesicular Glutamate Transport Protein
-	D12.776.543.585.450.162.887.750 Proteins	Vesicular Inhibitory Amino Acid Transport
-	D12.776.543.585.450.250	Cation Transport Proteins
-	D12.776.543.585.450.250.249	Arsenite Transporting ATPases
-	D12.776.543.585.450.250.500	Calcium-Transporting ATPases
-	D12.776.543.585.450.250.500.249 ATPases	Plasma Membrane Calcium-Transporting
-	D12.776.543.585.450.250.500.500 Transporting ATPases	Sarcoplasmic Reticulum Calcium-
-	D12.776.543.585.450.250.812	Organic Cation Transport Proteins
-	D12.776.543.585.450.250.812.500	Organic Cation Transporter 1
-	D12.776.543.585.450.250.875	Proton Pumps
-	D12.776.543.585.450.250.875.249	Bacteriorhodopsins
-	D12.776.543.585.450.250.875.311	Cytochrome b6f Complex
-	D12.776.543.585.450.250.875.311.500	Cytochromes b6
-	D12.776.543.585.450.250.875.311.750	Cytochromes f
-	D12.776.543.585.450.250.875.311.875	Plastoquinol-Plastocyanin Reductase
-	D12.776.543.585.450.250.875.437	Electron Transport Complex I
-	D12.776.543.585.450.250.875.468	Electron Transport Complex III
-	D12.776.543.585.450.250.875.484	Electron Transport Complex IV
-	D12.776.543.585.450.250.875.487	Inorganic Pyrophosphatase
-	D12.776.543.585.450.250.875.492	Photosystem I Protein Complex

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.543.585.450.250.875.500 Proton-Translocating ATPases
-	D12.776.543.585.450.250.875.500.249 Bacterial Proton-Translocating ATPases
-	D12.776.543.585.450.250.875.500.500 Chloroplast Proton-Translocating ATPases
-	D12.776.543.585.450.250.875.500.625 H(+)-K(+)-Exchanging ATPase
-	D12.776.543.585.450.250.875.500.750 Mitochondrial Proton-Translocating ATPases
-	D12.776.543.585.450.250.875.500.875 Vacuolar Proton-Translocating ATPases
-	D12.776.543.585.450.250.890 Sodium-Potassium-Exchanging ATPase
-	D12.776.543.585.450.437 SLC4A Proteins
-	D12.776.543.585.450.437.249 Chloride-Bicarbonate Antiporters
-	D12.776.543.585.450.437.249.500 Anion Exchange Protein 1, Erythrocyte
-	D12.776.543.585.450.437.500 Sodium-Bicarbonate Symporters
-	D12.776.543.585.450.625 Symporters
-	D12.776.543.585.450.625.124 Dopamine Plasma Membrane Transport Proteins
-	D12.776.543.585.450.625.139 GABA Plasma Membrane Transport Proteins
-	D12.776.543.585.450.625.147 Glutamate Plasma Membrane Transport Proteins
-	D12.776.543.585.450.625.147.500 Excitatory Amino Acid Transporter 1
-	D12.776.543.585.450.625.147.750 Excitatory Amino Acid Transporter 2
-	D12.776.543.585.450.625.147.875 Excitatory Amino Acid Transporter 3
-	D12.776.543.585.450.625.147.937 Excitatory Amino Acid Transporter 4
-	D12.776.543.585.450.625.147.968 Excitatory Amino Acid Transporter 5
-	D12.776.543.585.450.625.155 Glycine Plasma Membrane Transport Proteins
-	D12.776.543.585.450.625.186 Norepinephrine Plasma Membrane Transport Proteins
-	D12.776.543.585.450.625.217 Proton-Coupled Folate Transporter
-	D12.776.543.585.450.625.249 Proton-Phosphate Symporters
-	D12.776.543.585.450.625.374 Serotonin Plasma Membrane Transport Proteins
-	D12.776.543.585.450.625.500 Sodium-Bicarbonate Symporters
-	D12.776.543.585.450.625.531 Sodium-Coupled Vitamin C Transporters
-	D12.776.543.585.450.625.562 Sodium-Glucose Transport Proteins
-	D12.776.543.585.450.625.562.500 Sodium-Glucose Transporter 1
-	D12.776.543.585.450.625.562.750 Sodium-Glucose Transporter 2
-	D12.776.543.585.450.625.625 Sodium-Phosphate Cotransporter Proteins

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D12.776.543.585.450.625.625.500 Type I	Sodium-Phosphate Cotransporter Proteins,
-	D12.776.543.585.450.625.625.750 Type II	Sodium-Phosphate Cotransporter Proteins,
-	D12.776.543.585.450.625.625.750.124 Proteins, Type IIa	Sodium-Phosphate Cotransporter
-	D12.776.543.585.450.625.625.750.249 Proteins, Type IIb	Sodium-Phosphate Cotransporter
-	D12.776.543.585.450.625.625.750.500 Proteins, Type IIc	Sodium-Phosphate Cotransporter
-	D12.776.543.585.450.625.625.875 Type III	Sodium-Phosphate Cotransporter Proteins,
-	D12.776.543.585.450.625.750	Sodium-Potassium-Chloride Symporters
-	D12.776.543.585.450.625.750.500	Solute Carrier Family 12, Member 1
-	D12.776.543.585.450.625.750.625	Solute Carrier Family 12, Member 2
-	D12.776.543.585.450.625.750.875	Solute Carrier Family 12, Member 4
-	D12.776.543.585.475	Mitochondrial Membrane Transport Proteins
-	D12.776.543.585.475.500	Mitochondrial ADP, ATP Translocases
-	D12.776.543.585.475.500.100	Adenine Nucleotide Translocator 1
-	D12.776.543.585.475.500.200	Adenine Nucleotide Translocator 2
-	D12.776.543.585.475.500.300	Adenine Nucleotide Translocator 3
-	D12.776.543.585.475.625	Mitochondrial Proton-Translocating ATPases
New Heading	<b>D12.776.543.585.475.688</b>	<b>Mitochondrial Uncoupling Proteins</b>
New Heading	<b>D12.776.543.585.475.688.500</b>	<b>Uncoupling Protein 1</b>
New Heading	<b>D12.776.543.585.475.688.750</b>	<b>Uncoupling Protein 2</b>
New Heading	<b>D12.776.543.585.475.688.875</b>	<b>Uncoupling Protein 3</b>
-	D12.776.543.585.475.750	Voltage-Dependent Anion Channel 1
-	D12.776.543.585.475.875	Voltage-Dependent Anion Channel 2
-	D12.776.543.585.500	Monosaccharide Transport Proteins
-	D12.776.543.585.500.500	Glucose Transport Proteins, Facilitative
-	D12.776.543.585.500.500.500	Glucose Transporter Type 1
-	D12.776.543.585.500.500.750	Glucose Transporter Type 2
-	D12.776.543.585.500.500.875	Glucose Transporter Type 3
-	D12.776.543.585.500.500.937	Glucose Transporter Type 4
-	D12.776.543.585.500.500.968	Glucose Transporter Type 5

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.543.585.500.750 Sodium-Glucose Transport Proteins
-	D12.776.543.585.500.750.500 Sodium-Glucose Transporter 1
-	D12.776.543.585.500.750.750 Sodium-Glucose Transporter 2
-	D12.776.543.585.562 Neurotransmitter Transport Proteins
-	D12.776.543.585.562.374 Plasma Membrane Neurotransmitter Transport Proteins
-	D12.776.543.585.562.374.500 Catecholamine Plasma Membrane Transport Proteins
-	D12.776.543.585.562.374.500.500 Dopamine Plasma Membrane Transport Proteins
-	D12.776.543.585.562.374.500.750 Norepinephrine Plasma Membrane Transport Proteins
-	D12.776.543.585.562.374.750 GABA Plasma Membrane Transport Proteins
-	D12.776.543.585.562.374.781 Glutamate Plasma Membrane Transport Proteins
-	D12.776.543.585.562.374.781.500 Excitatory Amino Acid Transporter 1
-	D12.776.543.585.562.374.781.750 Excitatory Amino Acid Transporter 2
-	D12.776.543.585.562.374.781.781 Excitatory Amino Acid Transporter 3
-	D12.776.543.585.562.374.781.812 Excitatory Amino Acid Transporter 4
-	D12.776.543.585.562.374.781.875 Excitatory Amino Acid Transporter 5
-	D12.776.543.585.562.374.812 Glycine Plasma Membrane Transport Proteins
-	D12.776.543.585.562.374.875 Serotonin Plasma Membrane Transport Proteins
-	D12.776.543.585.562.750 Vesicular Neurotransmitter Transport Proteins
-	D12.776.543.585.562.750.500 Vesicular Biogenic Amine Transport Proteins
-	D12.776.543.585.562.750.500.249 Vesicular Acetylcholine Transport Proteins
-	D12.776.543.585.562.750.500.500 Vesicular Monoamine Transport Proteins
-	D12.776.543.585.562.750.625 Vesicular Glutamate Transport Proteins
-	D12.776.543.585.562.750.625.500 Vesicular Glutamate Transport Protein 1
-	D12.776.543.585.562.750.625.750 Vesicular Glutamate Transport Protein 2
-	D12.776.543.585.562.750.750 Vesicular Inhibitory Amino Acid Transport Proteins
-	D12.776.543.585.625 Nucleobase, Nucleoside, Nucleotide, and Nucleic Acid Transport Proteins
-	D12.776.543.585.625.500 Nucleobase Transport Proteins
-	D12.776.543.585.625.750 Nucleoside Transport Proteins
-	D12.776.543.585.625.750.500 Equilibrative Nucleoside Transport Proteins
-	D12.776.543.585.625.750.500.500 Equilibrative Nucleoside Transporter 1
-	D12.776.543.585.625.750.500.750 Equilibrative-Nucleoside Transporter 2

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.543.585.625.875 Nucleotide Transport Proteins
-	D12.776.543.585.625.875.500 Mitochondrial ADP, ATP Translocases
-	D12.776.543.585.625.875.500.100 Adenine Nucleotide Translocator 1
-	D12.776.543.585.625.875.500.200 Adenine Nucleotide Translocator 2
-	D12.776.543.585.625.875.500.300 Adenine Nucleotide Translocator 3
-	D12.776.543.585.750 Nucleocytoplasmic Transport Proteins
-	D12.776.543.585.750.100 Aryl Hydrocarbon Receptor Nuclear Translocator
-	D12.776.543.585.750.200 Cellular Apoptosis Susceptibility Protein
-	D12.776.543.585.750.500 Karyopherins
-	D12.776.543.585.750.500.100 alpha Karyopherins
-	D12.776.543.585.750.500.249 beta Karyopherins
-	D12.776.543.585.750.625 Nuclear Pore Complex Proteins
-	D12.776.543.585.750.687 Period Circadian Proteins
-	D12.776.543.585.750.750 ran GTP-Binding Protein
New Heading	<b>D12.776.543.585.875 SEC Translocation Channels</b>
New Heading	<b>D12.776.543.585.937 Solute Carrier Proteins</b>
New Tree	<a href="#">D12.776.543.585.937.250 Amino Acid Transport System X-AG</a>
New Tree	<a href="#">D12.776.543.585.937.250.500 Glutamate Plasma Membrane Transport Proteins</a>
New Tree	<a href="#">D12.776.543.585.937.250.500.500 Excitatory Amino Acid Transporter 1</a>
New Tree	<a href="#">D12.776.543.585.937.250.500.750 Excitatory Amino Acid Transporter 2</a>
New Tree	<a href="#">D12.776.543.585.937.250.500.875 Excitatory Amino Acid Transporter 3</a>
New Tree	<a href="#">D12.776.543.585.937.250.500.937 Excitatory Amino Acid Transporter 4</a>
New Tree	<a href="#">D12.776.543.585.937.250.500.968 Excitatory Amino Acid Transporter 5</a>
New Tree	<a href="#">D12.776.543.585.937.313 Amino Acid Transport System y+L</a>
New Tree	<a href="#">D12.776.543.585.937.313.500 Antigens, CD98</a>
New Tree	<a href="#">D12.776.543.585.937.313.500.250 Antigens, CD98 Heavy Chain</a>
New Tree	<a href="#">D12.776.543.585.937.313.500.625 Antigens, CD98 Light Chains</a>

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	D12.776.543.585.937.313.500.625.500 1 Large Neutral Amino Acid-Transporter
New Tree	D12.776.543.585.937.375 Amino Acid Transport System y+
New Tree	D12.776.543.585.937.375.200 Cationic Amino Acid Transporter 1
New Tree	D12.776.543.585.937.375.300 Cationic Amino Acid Transporter 2
New Tree	D12.776.543.585.937.500 Dopamine Plasma Membrane Transport Proteins
New Tree	D12.776.543.585.937.563 Equilibrative Nucleoside Transport Proteins
New Tree	D12.776.543.585.937.563.500 Equilibrative Nucleoside Transporter 1
New Tree	D12.776.543.585.937.563.750 Equilibrative-Nucleoside Transporter 2
New Tree	D12.776.543.585.937.625 Glucose Transport Proteins, Facilitative
New Tree	D12.776.543.585.937.625.500 Glucose Transporter Type 1
New Tree	D12.776.543.585.937.625.750 Glucose Transporter Type 2
New Tree	D12.776.543.585.937.625.875 Glucose Transporter Type 3
New Tree	D12.776.543.585.937.625.937 Glucose Transporter Type 4
New Tree	D12.776.543.585.937.625.968 Glucose Transporter Type 5
New Tree	D12.776.543.585.937.688 Mitochondrial ADP, ATP Translocases
New Tree	D12.776.543.585.937.688.100 Adenine Nucleotide Translocator 1
New Tree	D12.776.543.585.937.688.200 Adenine Nucleotide Translocator 2
New Tree	D12.776.543.585.937.688.300 Adenine Nucleotide Translocator 3
New Heading	<b>D12.776.543.585.937.696 Mitochondrial Uncoupling Proteins</b>
New Heading	<b>D12.776.543.585.937.696.500 Uncoupling Protein 1</b>
New Heading	<b>D12.776.543.585.937.696.750 Uncoupling Protein 2</b>
New Heading	<b>D12.776.543.585.937.696.875 Uncoupling Protein 3</b>

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	D12.776.543.585.937.700 Norepinephrine Plasma Membrane Transport Proteins
New Tree	D12.776.543.585.937.704 Organic Anion Transport Polypeptide C
New Tree	D12.776.543.585.937.704 Solute Carrier Organic Anion Transporter Family Member 1b1
New Tree	D12.776.543.585.937.719 Organic Anion Transport Protein 1
New Tree	D12.776.543.585.937.727 Organic Cation Transporter 1
New Tree	D12.776.543.585.937.735 Proton-Coupled Folate Transporter
New Tree	D12.776.543.585.937.743 Reduced Folate Carrier Protein
New Tree	D12.776.543.585.937.747 Serotonin Plasma Membrane Transport Proteins
New Tree	D12.776.543.585.937.750 SLC4A Proteins
New Tree	D12.776.543.585.937.750.249 Chloride-Bicarbonate Antiporters
New Tree	D12.776.543.585.937.750.249.500 Anion Exchange Protein 1, Erythrocyte
New Tree	D12.776.543.585.937.750.500 Sodium-Bicarbonate Symporters
New Tree	D12.776.543.585.937.813 Sodium-Coupled Vitamin C Transporters
New Tree	D12.776.543.585.937.821 Sodium-Glucose Transporter 1
New Tree	D12.776.543.585.937.825 Sodium-Glucose Transporter 2
New Tree	D12.776.543.585.937.829 Sodium-Phosphate Cotransporter Proteins, Type II
New Tree	D12.776.543.585.937.829.124 Sodium-Phosphate Cotransporter Proteins, Type IIa
New Tree	D12.776.543.585.937.829.249 Sodium-Phosphate Cotransporter Proteins, Type IIb
New Tree	D12.776.543.585.937.829.500 Sodium-Phosphate Cotransporter Proteins, Type IIc
New Tree	D12.776.543.585.937.844 Sodium-Phosphate Cotransporter Proteins, Type III
New Tree	D12.776.543.585.937.875 Sodium-Potassium-Chloride Symporters
New Tree	D12.776.543.585.937.875.500 Solute Carrier Family 12, Member 1

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	<a href="#">D12.776.543.585.937.875.625</a>	<a href="#">Solute Carrier Family 12, Member 2</a>
New Tree	<a href="#">D12.776.543.585.937.875.750</a>	<a href="#">Solute Carrier Family 12, Member 3</a>
New Tree	<a href="#">D12.776.543.585.937.875.875</a>	<a href="#">Solute Carrier Family 12, Member 4</a>
-	D12.776.543.602	Mitochondrial Trifunctional Protein
-	D12.776.543.602.200	Mitochondrial Trifunctional Protein, alpha Subunit
-	D12.776.543.602.500	Mitochondrial Trifunctional Protein, beta Subunit
-	D12.776.543.620	Myelin Proteins
-	D12.776.543.620.264	Myelin and Lymphocyte-Associated Proteolipid Proteins
-	D12.776.543.620.530	Myelin-Associated Glycoprotein
-	D12.776.543.620.540	Myelin Basic Protein
-	D12.776.543.620.550	Myelin-Oligodendrocyte Glycoprotein
-	D12.776.543.620.570	Myelin P0 Protein
-	D12.776.543.620.575	Myelin P2 Protein
-	D12.776.543.620.580	Myelin Proteolipid Protein
New Heading	<b><a href="#">D12.776.543.620.738</a></b>	<b><a href="#">Nogo Proteins</a></b>
-	D12.776.543.620.895	Oligodendrocyte-Myelin Glycoprotein
-	D12.776.543.685	Neurofibromin 2
-	D12.776.543.693	Phospholipid Transfer Proteins
-	D12.776.543.695	Pore Forming Cytotoxic Proteins
-	D12.776.543.695.054	Antimicrobial Cationic Peptides
-	D12.776.543.695.054.099	Cathelicidins
-	D12.776.543.695.054.149	Cecropins
-	D12.776.543.695.054.200	Defensins
-	D12.776.543.695.054.200.050	alpha-Defensins
-	D12.776.543.695.054.200.075	beta-Defensins
-	D12.776.543.695.054.400	Dermcidins
-	D12.776.543.695.054.425	Hepcidins
-	D12.776.543.695.054.450	Histatins
-	D12.776.543.695.054.500	Magainins
-	D12.776.543.695.054.550	Melittin
-	D12.776.543.695.054.600	Polymyxins
-	D12.776.543.695.054.600.110	Colistin
-	D12.776.543.695.054.600.750	Polymyxin B



## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D12.776.543.695.054.800	Thionins
-	D12.776.543.695.110	Bacteriocins
-	D12.776.543.695.110.700	Nisin
New Heading	<b>D12.776.543.695.110.850</b>	<b>Pediocins</b>
-	D12.776.543.695.221	Gramicidin
-	D12.776.543.695.444	Hemolysin Proteins
-	D12.776.543.695.750	Leukocidins
-	D12.776.543.695.875	Perforin
-	D12.776.543.695.937	Streptolysins
-	D12.776.543.696	Presenilins
-	D12.776.543.696.500	Presenilin-1
-	D12.776.543.696.750	Presenilin-2
-	D12.776.543.697	Protoporphyrinogen Oxidase
-	D12.776.543.701	Pulmonary Surfactant-Associated Protein B
-	D12.776.543.717	Pulmonary Surfactant-Associated Protein C
-	D12.776.543.725	Receptor Activity-Modifying Proteins
-	D12.776.543.725.100	Receptor Activity-Modifying Protein 1
-	D12.776.543.725.200	Receptor Activity-Modifying Protein 2
-	D12.776.543.725.300	Receptor Activity-Modifying Protein 3
-	D12.776.543.733	Receptor-Like Protein Tyrosine Phosphatases
-	D12.776.543.733.937	Receptor-Like Protein Tyrosine Phosphatases, Class 1
-	D12.776.543.733.937.500	Antigens, CD45
-	D12.776.543.750	Receptors, Cell Surface
-	D12.776.543.750.006	Advanced Glycosylation End Product-Specific Receptor
-	D12.776.543.750.011	Antigens, CD36
-	D12.776.543.750.024	Asialoglycoprotein Receptor
New Tree	<a href="#">D12.776.543.750.035</a>	<a href="#">Autoreceptors</a>
Old Tree	<a href="#">D12.776.543.750.050</a>	<a href="#">Autoreceptors</a>
Old Tree	<a href="#">D12.776.543.750.051</a>	<a href="#">Cysteine Loop Ligand-Gated Ion Channel Receptors</a>
Old Tree	<a href="#">D12.776.543.750.051.500</a>	<a href="#">Receptors, GABA-A</a>
Old Tree	<a href="#">D12.776.543.750.051.625</a>	<a href="#">Receptors, Glycine</a>
Old Tree	<a href="#">D12.776.543.750.051.687</a>	<a href="#">Receptors, Nicotinic</a>
Old Tree	<a href="#">D12.776.543.750.051.687.500</a>	<a href="#">alpha7 Nicotinic Acetylcholine Receptor</a>
Old Tree	<a href="#">D12.776.543.750.051.750</a>	<a href="#">Receptors, Serotonin, 5-HT3</a>

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	D12.776.543.750.052 Folate Receptors, GPI-Anchored
Old Tree	D12.776.543.750.052.500 Folate Receptor 1
Old Tree	D12.776.543.750.052.750 Folate Receptor 2
Old Tree	D12.776.543.750.053 Junctional Adhesion Molecule A
New Heading	<b>D12.776.543.750.054 Methyl-Accepting Chemotaxis Proteins</b>
Old Tree	D12.776.543.750.055 Neuropilins
Old Tree	D12.776.543.750.055.500 Neuropilin-1
Old Tree	D12.776.543.750.055.750 Neuropilin-2
New Heading	<b>D12.776.543.750.058 Patched Receptors</b>
New Heading	<b>D12.776.543.750.058.500 Patched-1 Receptor</b>
New Heading	<b>D12.776.543.750.058.750 Patched-2 Receptor</b>
Old Tree	D12.776.543.750.060 Receptor Protein-Tyrosine Kinases
Old Tree	D12.776.543.750.060.009 ErbB Receptors
Old Tree	D12.776.543.750.060.009.300 Receptor, Epidermal Growth Factor
Old Tree	D12.776.543.750.060.009.400 Receptor, ErbB-2
Old Tree	D12.776.543.750.060.009.500 Receptor, ErbB-3
Old Tree	D12.776.543.750.060.009.600 Receptor, ErbB-4
Old Tree	D12.776.543.750.060.020 fms-Like Tyrosine Kinase 3
Old Tree	D12.776.543.750.060.124 Proto-Oncogene Proteins c-kit
Old Tree	D12.776.543.750.060.186 Proto-Oncogene Proteins c-met
Old Tree	D12.776.543.750.060.217 Proto-Oncogene Proteins c-ret
Old Tree	D12.776.543.750.060.233 Receptor Tyrosine Kinase-like Orphan Receptors
Old Tree	D12.776.543.750.060.440 Receptor, Fibroblast Growth Factor, Type 1
Old Tree	D12.776.543.750.060.441 Receptor, Fibroblast Growth Factor, Type 2
Old Tree	D12.776.543.750.060.442 Receptor, Fibroblast Growth Factor, Type 3
Old Tree	D12.776.543.750.060.443 Receptor, Fibroblast Growth Factor, Type 4
Old Tree	D12.776.543.750.060.468 Receptor, IGF Type 1
Old Tree	D12.776.543.750.060.484 Receptor, Insulin
Old Tree	D12.776.543.750.060.492 Receptor, Macrophage Colony-Stimulating Factor
Old Tree	D12.776.543.750.060.496 Receptor, trkA
Old Tree	D12.776.543.750.060.498 Receptor, trkB
Old Tree	D12.776.543.750.060.499 Receptor, trkC
Old Tree	D12.776.543.750.060.500 Receptors, Eph Family

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	D12.776.543.750.060.500.050 Receptor, EphA1
Old Tree	D12.776.543.750.060.500.100 Receptor, EphA2
Old Tree	D12.776.543.750.060.500.150 Receptor, EphA3
Old Tree	D12.776.543.750.060.500.200 Receptor, EphA4
Old Tree	D12.776.543.750.060.500.250 Receptor, EphA5
Old Tree	D12.776.543.750.060.500.300 Receptor, EphA6
Old Tree	D12.776.543.750.060.500.350 Receptor, EphA7
Old Tree	D12.776.543.750.060.500.400 Receptor, EphA8
Old Tree	D12.776.543.750.060.500.450 Receptor, EphB1
Old Tree	D12.776.543.750.060.500.500 Receptor, EphB2
Old Tree	D12.776.543.750.060.500.687 Receptor, EphB3
Old Tree	D12.776.543.750.060.500.781 Receptor, EphB4
Old Tree	D12.776.543.750.060.500.875 Receptor, EphB5
Old Tree	D12.776.543.750.060.500.937 Receptor, EphB6
Old Tree	D12.776.543.750.060.625 Receptors, Platelet-Derived Growth Factor
Old Tree	D12.776.543.750.060.625.300 Receptor, Platelet-Derived Growth Factor alpha
Old Tree	D12.776.543.750.060.625.400 Receptor, Platelet-Derived Growth Factor beta
Old Tree	D12.776.543.750.060.687 Receptors, TIE
Old Tree	D12.776.543.750.060.687.249 Receptor, TIE-1
Old Tree	D12.776.543.750.060.687.500 Receptor, TIE-2
Old Tree	D12.776.543.750.060.750 Receptors, Vascular Endothelial Growth Factor
Old Tree	D12.776.543.750.060.750.100 Vascular Endothelial Growth Factor Receptor-1
Old Tree	D12.776.543.750.060.750.200 Vascular Endothelial Growth Factor Receptor-2
Old Tree	D12.776.543.750.060.750.300 Vascular Endothelial Growth Factor Receptor-3
Old Tree	D12.776.543.750.065 Receptors, Adipokine
Old Tree	D12.776.543.750.065.249 Receptors, Adiponectin
Old Tree	D12.776.543.750.065.500 Receptors, Leptin
Old Tree	D12.776.543.750.066 Receptors, Artificial
Old Tree	D12.776.543.750.068 Receptors, Autocrine Motility Factor
Old Tree	D12.776.543.750.069 Receptors, Biogenic Amine
Old Tree	D12.776.543.750.069.300 Receptors, Catecholamine
Old Tree	D12.776.543.750.069.300.300 Receptors, Adrenergic
Old Tree	D12.776.543.750.069.300.300.300 Receptors, Adrenergic, alpha
Old Tree	D12.776.543.750.069.300.300.300.100 Receptors, Adrenergic, alpha-1
Old Tree	D12.776.543.750.069.300.300.300.200 Receptors, Adrenergic, alpha-2
Old Tree	D12.776.543.750.069.300.300.340 Receptors, Adrenergic, beta

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	D12.776.543.750.069.300.300.340.100 Receptors, Adrenergic, beta-1
Old Tree	D12.776.543.750.069.300.300.340.200 Receptors, Adrenergic, beta-2
Old Tree	D12.776.543.750.069.300.300.340.300 Receptors, Adrenergic, beta-3
Old Tree	D12.776.543.750.069.300.400 Receptors, Dopamine
Old Tree	D12.776.543.750.069.300.400.400 Receptors, Dopamine D1
Old Tree	D12.776.543.750.069.300.400.400.500 Receptors, Dopamine D5
Old Tree	D12.776.543.750.069.300.400.500 Receptors, Dopamine D2
Old Tree	D12.776.543.750.069.300.400.500.249 Receptors, Dopamine D3
Old Tree	D12.776.543.750.069.300.400.500.500 Receptors, Dopamine D4
Old Tree	D12.776.543.750.069.450 Receptors, Histamine
Old Tree	D12.776.543.750.069.450.300 Receptors, Histamine H1
Old Tree	D12.776.543.750.069.450.400 Receptors, Histamine H2
Old Tree	D12.776.543.750.069.450.500 Receptors, Histamine H3
Old Tree	D12.776.543.750.069.800 Receptors, Serotonin
Old Tree	D12.776.543.750.069.800.100 Receptors, Serotonin, 5-HT1
Old Tree	D12.776.543.750.069.800.100.100 Receptor, Serotonin, 5-HT1A
Old Tree	D12.776.543.750.069.800.100.150 Receptor, Serotonin, 5-HT1B
Old Tree	D12.776.543.750.069.800.100.200 Receptor, Serotonin, 5-HT1D
Old Tree	D12.776.543.750.069.800.200 Receptors, Serotonin, 5-HT2
Old Tree	D12.776.543.750.069.800.200.100 Receptor, Serotonin, 5-HT2A
Old Tree	D12.776.543.750.069.800.200.150 Receptor, Serotonin, 5-HT2B
Old Tree	D12.776.543.750.069.800.200.200 Receptor, Serotonin, 5-HT2C
Old Tree	D12.776.543.750.069.800.300 Receptors, Serotonin, 5-HT3
Old Tree	D12.776.543.750.069.800.400 Receptors, Serotonin, 5-HT4
Old Tree	D12.776.543.750.070 Receptors, Collagen
Old Tree	D12.776.543.750.070.100 Integrin alpha1beta1
Old Tree	D12.776.543.750.070.200 Integrin alpha2beta1
Old Tree	D12.776.543.750.070.300 Integrin alpha3beta1
Old Tree	D12.776.543.750.073 Receptors, Death Domain
Old Tree	D12.776.543.750.073.500 Antigens, CD95
Old Tree	D12.776.543.750.073.550 Edar Receptor
Old Tree	D12.776.543.750.073.600 Receptors, TNF-Related Apoptosis-Inducing Ligand
Old Tree	D12.776.543.750.073.725 Receptors, Tumor Necrosis Factor, Member 25
Old Tree	D12.776.543.750.073.750 Receptors, Tumor Necrosis Factor, Type I
Old Tree	D12.776.543.750.100 Receptors, G-Protein-Coupled
Old Tree	D12.776.543.750.100.011 Calcitonin Receptor-Like Protein

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	D12.776.543.750.100.017 Frizzled Receptors
Old Tree	D12.776.543.750.100.021 Glucagon-Like Peptide Receptors
Old Tree	D12.776.543.750.100.021.500 Glucagon-Like Peptide-1 Receptor
Old Tree	D12.776.543.750.100.021.600 Glucagon-Like Peptide-2 Receptor
Old Tree	D12.776.543.750.100.023 Orexin Receptors
Old Tree	D12.776.543.750.100.024 Receptor, Anaphylatoxin C5a
Old Tree	D12.776.543.750.100.035 Receptor, PAR-2
Old Tree	D12.776.543.750.100.041 Receptors, Adrenomedullin
Old Tree	D12.776.543.750.100.047 Receptors, Angiotensin
Old Tree	D12.776.543.750.100.047.625 Receptor, Angiotensin, Type 1
Old Tree	D12.776.543.750.100.047.687 Receptor, Angiotensin, Type 2
Old Tree	D12.776.543.750.100.070 Receptors, Bombesin
Old Tree	D12.776.543.750.100.080 Receptors, Bradykinin
Old Tree	D12.776.543.750.100.080.249 Receptor, Bradykinin B1
Old Tree	D12.776.543.750.100.080.500 Receptor, Bradykinin B2
Old Tree	D12.776.543.750.100.100 Receptors, Calcitonin
Old Tree	D12.776.543.750.100.105 Receptors, Calcitonin Gene-Related Peptide
Old Tree	D12.776.543.750.100.115 Receptors, Calcium-Sensing
Old Tree	D12.776.543.750.100.125 Receptors, Cannabinoid
Old Tree	D12.776.543.750.100.125.100 Receptor, Cannabinoid, CB1
Old Tree	D12.776.543.750.100.125.200 Receptor, Cannabinoid, CB2
Old Tree	D12.776.543.750.100.150 Receptors, Catecholamine
Old Tree	D12.776.543.750.100.150.300 Receptors, Adrenergic
Old Tree	D12.776.543.750.100.150.300.300 Receptors, Adrenergic, alpha
Old Tree	D12.776.543.750.100.150.300.300.700 Receptors, Adrenergic, alpha-1
Old Tree	D12.776.543.750.100.150.300.300.725 Receptors, Adrenergic, alpha-2
Old Tree	D12.776.543.750.100.150.300.340 Receptors, Adrenergic, beta
Old Tree	D12.776.543.750.100.150.300.340.700 Receptors, Adrenergic, beta-1
Old Tree	D12.776.543.750.100.150.300.340.725 Receptors, Adrenergic, beta-2
Old Tree	D12.776.543.750.100.150.300.340.900 Receptors, Adrenergic, beta-3
Old Tree	D12.776.543.750.100.150.400 Receptors, Dopamine
Old Tree	D12.776.543.750.100.150.400.400 Receptors, Dopamine D1
Old Tree	D12.776.543.750.100.150.400.400.500 Receptors, Dopamine D5
Old Tree	D12.776.543.750.100.150.400.500 Receptors, Dopamine D2
Old Tree	D12.776.543.750.100.150.400.500.500 Receptors, Dopamine D3
Old Tree	D12.776.543.750.100.150.400.500.750 Receptors, Dopamine D4

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	D12.776.543.750.100.160 Receptors, Chemokine
Old Tree	D12.776.543.750.100.160.150 Receptors, CCR
Old Tree	D12.776.543.750.100.160.150.100 Receptors, CCR1
Old Tree	D12.776.543.750.100.160.150.200 Receptors, CCR2
Old Tree	D12.776.543.750.100.160.150.300 Receptors, CCR3
Old Tree	D12.776.543.750.100.160.150.400 Receptors, CCR4
Old Tree	D12.776.543.750.100.160.150.500 Receptors, CCR5
Old Tree	D12.776.543.750.100.160.150.600 Receptors, CCR6
Old Tree	D12.776.543.750.100.160.150.700 Receptors, CCR7
Old Tree	D12.776.543.750.100.160.150.800 Receptors, CCR8
Old Tree	D12.776.543.750.100.160.150.950 Receptors, CCR10
Old Tree	D12.776.543.750.100.160.500 Receptors, CXCR
Old Tree	D12.776.543.750.100.160.500.300 Receptors, CXCR3
Old Tree	D12.776.543.750.100.160.500.400 Receptors, CXCR4
Old Tree	D12.776.543.750.100.160.500.500 Receptors, CXCR5
Old Tree	D12.776.543.750.100.160.500.750 Receptors, Interleukin-8
Old Tree	D12.776.543.750.100.160.500.750.500 Receptors, Interleukin-8A
Old Tree	D12.776.543.750.100.160.500.750.750 Receptors, Interleukin-8B
Old Tree	D12.776.543.750.100.170 Receptors, Cholecystokinin
Old Tree	D12.776.543.750.100.170.100 Receptor, Cholecystokinin A
Old Tree	D12.776.543.750.100.170.200 Receptor, Cholecystokinin B
Old Tree	D12.776.543.750.100.180 Receptors, Corticotropin-Releasing Hormone
Old Tree	D12.776.543.750.100.200 Receptors, Eicosanoid
Old Tree	D12.776.543.750.100.200.450 Receptors, Leukotriene
Old Tree	D12.776.543.750.100.200.450.300 Receptors, Leukotriene B4
Old Tree	D12.776.543.750.100.200.575 Receptors, Lipoxin
Old Tree	D12.776.543.750.100.200.700 Receptors, Prostaglandin
Old Tree	D12.776.543.750.100.200.700.299 Receptors, Epoprostenol
Old Tree	D12.776.543.750.100.200.700.600 Receptors, Prostaglandin E
Old Tree	D12.776.543.750.100.200.700.600.100 Receptors, Prostaglandin E, EP1 Subtype
Old Tree	D12.776.543.750.100.200.700.600.200 Receptors, Prostaglandin E, EP2 Subtype
Old Tree	D12.776.543.750.100.200.700.600.300 Receptors, Prostaglandin E, EP3 Subtype
Old Tree	D12.776.543.750.100.200.700.600.400 Receptors, Prostaglandin E, EP4 Subtype

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	D12.776.543.750.100.200.700.800 Prostaglandin H2 Receptors, Thromboxane A2,
Old Tree	D12.776.543.750.100.200.800 Receptors, Thromboxane
Old Tree	D12.776.543.750.100.200.800.500 Prostaglandin H2 Receptors, Thromboxane A2,
Old Tree	D12.776.543.750.100.220 Receptors, Endothelin
Old Tree	D12.776.543.750.100.220.100 Receptor, Endothelin A
Old Tree	D12.776.543.750.100.220.200 Receptor, Endothelin B
Old Tree	D12.776.543.750.100.235 Receptors, Formyl Peptide
Old Tree	D12.776.543.750.100.250 Receptors, FSH
Old Tree	D12.776.543.750.100.300 Receptors, GABA-B
Old Tree	D12.776.543.750.100.315 Receptors, Galanin
Old Tree	D12.776.543.750.100.315.100 Receptor, Galanin, Type 1
Old Tree	D12.776.543.750.100.315.200 Receptor, Galanin, Type 2
Old Tree	D12.776.543.750.100.315.300 Receptor, Galanin, Type 3
Old Tree	D12.776.543.750.100.322 Receptors, Ghrelin
Old Tree	D12.776.543.750.100.330 Receptors, Glucagon
Old Tree	D12.776.543.750.100.350 Receptors, Histamine H1
Old Tree	D12.776.543.750.100.355 Receptors, Histamine H2
Old Tree	D12.776.543.750.100.360 Receptors, Islet Amyloid Polypeptide
Old Tree	D12.776.543.750.100.400 Receptors, LH
Old Tree	D12.776.543.750.100.410 Receptors, LHRH
Old Tree	D12.776.543.750.100.420 Receptors, Lysophospholipid
Old Tree	D12.776.543.750.100.420.249 Receptors, Lysophosphatidic Acid
Old Tree	D12.776.543.750.100.420.500 Receptors, Lysosphingolipid
Old Tree	D12.776.543.750.100.430 Receptors, Melanocortin
Old Tree	D12.776.543.750.100.430.500 Receptor, Melanocortin, Type 1
Old Tree	D12.776.543.750.100.430.750 Receptor, Melanocortin, Type 2
Old Tree	D12.776.543.750.100.430.875 Receptor, Melanocortin, Type 3
Old Tree	D12.776.543.750.100.430.937 Receptor, Melanocortin, Type 4
Old Tree	D12.776.543.750.100.440 Receptors, Melatonin
Old Tree	D12.776.543.750.100.440.500 Receptor, Melatonin, MT1
Old Tree	D12.776.543.750.100.440.750 Receptor, Melatonin, MT2
Old Tree	D12.776.543.750.100.450 Receptors, Metabotropic Glutamate
Old Tree	D12.776.543.750.100.450.500 Receptor, Metabotropic Glutamate 5
Old Tree	D12.776.543.750.100.475 Receptors, Muscarinic

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	D12.776.543.750.100.475.100 Receptor, Muscarinic M1
Old Tree	D12.776.543.750.100.475.200 Receptor, Muscarinic M2
Old Tree	D12.776.543.750.100.475.300 Receptor, Muscarinic M3
Old Tree	D12.776.543.750.100.475.400 Receptor, Muscarinic M4
Old Tree	D12.776.543.750.100.475.500 Receptor, Muscarinic M5
Old Tree	D12.776.543.750.100.500 Receptors, Neuropeptide Y
Old Tree	D12.776.543.750.100.550 Receptors, Neurotensin
Old Tree	D12.776.543.750.100.600 Receptors, Odorant
Old Tree	D12.776.543.750.100.620 Receptors, Opioid
Old Tree	D12.776.543.750.100.620.200 Receptors, Opioid, delta
Old Tree	D12.776.543.750.100.620.400 Receptors, Opioid, kappa
Old Tree	D12.776.543.750.100.620.550 Receptors, Opioid, mu
Old Tree	D12.776.543.750.100.620.775 Receptors, sigma
Old Tree	D12.776.543.750.100.630 Receptors, Oxytocin
Old Tree	D12.776.543.750.100.650 Receptors, Parathyroid Hormone
Old Tree	D12.776.543.750.100.650.100 Receptor, Parathyroid Hormone, Type 1
Old Tree	D12.776.543.750.100.650.200 Receptor, Parathyroid Hormone, Type 2
Old Tree	D12.776.543.750.100.660 Receptors, Pheromone
Old Tree	D12.776.543.750.100.660.500 Receptors, Mating Factor
Old Tree	D12.776.543.750.100.665 Receptors, Pituitary Adenylate Cyclase-Activating Polypeptide
Old Tree	D12.776.543.750.100.665.500 Receptors, Pituitary Adenylate Cyclase-Activating Polypeptide, Type I
Old Tree	D12.776.543.750.100.665.750 Receptors, Vasoactive Intestinal Peptide, Type II
Old Tree	D12.776.543.750.100.665.760 Receptors, Vasoactive Intestinal Polypeptide, Type I
Old Tree	D12.776.543.750.100.700 Receptors, Purinergic
Old Tree	D12.776.543.750.100.700.150 Receptors, Cyclic AMP
Old Tree	D12.776.543.750.100.700.700 Receptors, Purinergic P1
Old Tree	D12.776.543.750.100.700.700.100 Receptor, Adenosine A1
Old Tree	D12.776.543.750.100.700.700.200 Receptors, Adenosine A2
Old Tree	D12.776.543.750.100.700.700.200.100 Receptor, Adenosine A2A
Old Tree	D12.776.543.750.100.700.700.200.200 Receptor, Adenosine A2B
Old Tree	D12.776.543.750.100.700.700.300 Receptor, Adenosine A3
Old Tree	D12.776.543.750.100.700.720 Receptors, Purinergic P2
Old Tree	D12.776.543.750.100.700.720.250 Receptors, Purinergic P2X



## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	D12.776.543.750.100.700.720.250.100 Receptors, Purinergic P2X1
Old Tree	D12.776.543.750.100.700.720.250.200 Receptors, Purinergic P2X2
Old Tree	D12.776.543.750.100.700.720.250.300 Receptors, Purinergic P2X3
Old Tree	D12.776.543.750.100.700.720.250.400 Receptors, Purinergic P2X4
Old Tree	D12.776.543.750.100.700.720.250.500 Receptors, Purinergic P2X5
Old Tree	D12.776.543.750.100.700.720.250.700 Receptors, Purinergic P2X7
Old Tree	D12.776.543.750.100.700.720.500 Receptors, Purinergic P2Y
Old Tree	D12.776.543.750.100.700.720.500.100 Receptors, Purinergic P2Y1
Old Tree	D12.776.543.750.100.700.720.500.200 Receptors, Purinergic P2Y2
Old Tree	D12.776.543.750.100.700.720.500.300 Receptors, Purinergic P2Y12
Old Tree	D12.776.543.750.100.800 Receptors, Serotonin
Old Tree	D12.776.543.750.100.800.100 Receptors, Serotonin, 5-HT1
Old Tree	D12.776.543.750.100.800.100.100 Receptor, Serotonin, 5-HT1A
Old Tree	D12.776.543.750.100.800.100.150 Receptor, Serotonin, 5-HT1B
Old Tree	D12.776.543.750.100.800.100.200 Receptor, Serotonin, 5-HT1D
Old Tree	D12.776.543.750.100.800.200 Receptors, Serotonin, 5-HT2
Old Tree	D12.776.543.750.100.800.200.100 Receptor, Serotonin, 5-HT2A
Old Tree	D12.776.543.750.100.800.200.150 Receptor, Serotonin, 5-HT2B
Old Tree	D12.776.543.750.100.800.200.200 Receptor, Serotonin, 5-HT2C
Old Tree	D12.776.543.750.100.800.400 Receptors, Serotonin, 5-HT4
Old Tree	D12.776.543.750.100.850 Receptors, Somatostatin
Old Tree	D12.776.543.750.100.862 Receptors, Tachykinin
Old Tree	D12.776.543.750.100.862.500 Receptors, Neurokinin-1
Old Tree	D12.776.543.750.100.862.540 Receptors, Neurokinin-2
Old Tree	D12.776.543.750.100.862.580 Receptors, Neurokinin-3
Old Tree	D12.776.543.750.100.875 Receptors, Thrombin
Old Tree	D12.776.543.750.100.875.500 Receptor, PAR-1
Old Tree	D12.776.543.750.100.900 Receptors, Vasoactive Intestinal Peptide
Old Tree	D12.776.543.750.100.900.500 II Receptors, Vasoactive Intestinal Peptide, Type II
Old Tree	D12.776.543.750.100.900.520 I Receptors, Vasoactive Intestinal Polypeptide, Type I
Old Tree	D12.776.543.750.100.910 Receptors, Vasopressin
Old Tree	D12.776.543.750.100.955 Rhodopsin
Old Tree	D12.776.543.750.120 Receptors, Guanylate Cyclase-Coupled
Old Tree	D12.776.543.750.120.500 Receptors, Atrial Natriuretic Factor

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	D12.776.543.750.130 Cysteine Loop Ligand-Gated Ion Channel Receptors
New Tree	D12.776.543.750.130.500 Receptors, GABA-A
New Tree	D12.776.543.750.130.625 Receptors, Glycine
New Tree	D12.776.543.750.130.687 Receptors, Nicotinic
New Tree	D12.776.543.750.130.687.500 alpha7 Nicotinic Acetylcholine Receptor
New Tree	D12.776.543.750.130.750 Receptors, Serotonin, 5-HT3
New Heading	<b>D12.776.543.750.250 Endoglin</b>
New Tree	D12.776.543.750.301 Folate Receptors, GPI-Anchored
New Tree	D12.776.543.750.301.500 Folate Receptor 1
New Tree	D12.776.543.750.301.750 Folate Receptor 2
New Tree	D12.776.543.750.400 Junctional Adhesion Molecule A
New Tree	D12.776.543.750.590 Neuropilins
New Tree	D12.776.543.750.590.500 Neuropilin-1
New Tree	D12.776.543.750.590.750 Neuropilin-2
New Heading	<b>D12.776.543.750.600 Nogo Receptors</b>
New Heading	<b>D12.776.543.750.600.250 Nogo Receptor 2</b>
New Heading	<b>D12.776.543.750.600.500 Nogo Receptor 1</b>
New Tree	D12.776.543.750.630 Receptor Protein-Tyrosine Kinases
New Heading	<b>D12.776.543.750.630.005 Discoidin Domain Receptors</b>
New Heading	<b>D12.776.543.750.630.005.500 Discoidin Domain Receptor 1</b>
New Heading	<b>D12.776.543.750.630.005.750 Discoidin Domain Receptor 2</b>
New Tree	D12.776.543.750.630.009 ErbB Receptors

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	D12.776.543.750.630.009.300      Receptor, Epidermal Growth Factor
New Tree	D12.776.543.750.630.009.400      Receptor, ErbB-2
New Tree	D12.776.543.750.630.009.500      Receptor, ErbB-3
New Tree	D12.776.543.750.630.009.600      Receptor, ErbB-4
New Tree	D12.776.543.750.630.020      fms-Like Tyrosine Kinase 3
New Tree	D12.776.543.750.630.124      Proto-Oncogene Proteins c-kit
New Tree	D12.776.543.750.630.186      Proto-Oncogene Proteins c-met
New Tree	D12.776.543.750.630.217      Proto-Oncogene Proteins c-ret
New Tree	D12.776.543.750.630.233      Receptor Tyrosine Kinase-like Orphan Receptors
New Tree	D12.776.543.750.630.440      Receptor, Fibroblast Growth Factor, Type 1
New Tree	D12.776.543.750.630.441      Receptor, Fibroblast Growth Factor, Type 2
New Tree	D12.776.543.750.630.442      Receptor, Fibroblast Growth Factor, Type 3
New Tree	D12.776.543.750.630.443      Receptor, Fibroblast Growth Factor, Type 4
New Tree	D12.776.543.750.630.468      Receptor, IGF Type 1
New Tree	D12.776.543.750.630.484      Receptor, Insulin
New Tree	D12.776.543.750.630.492      Receptor, Macrophage Colony-Stimulating Factor
New Tree	D12.776.543.750.630.496      Receptor, trkA
New Tree	D12.776.543.750.630.498      Receptor, trkB
New Tree	D12.776.543.750.630.499      Receptor, trkC
New Tree	D12.776.543.750.630.500      Receptors, Eph Family
New Tree	D12.776.543.750.630.500.050      Receptor, EphA1
New Tree	D12.776.543.750.630.500.100      Receptor, EphA2

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	D12.776.543.750.630.500.150 Receptor, EphA3
New Tree	D12.776.543.750.630.500.200 Receptor, EphA4
New Tree	D12.776.543.750.630.500.250 Receptor, EphA5
New Tree	D12.776.543.750.630.500.300 Receptor, EphA6
New Tree	D12.776.543.750.630.500.350 Receptor, EphA7
New Tree	D12.776.543.750.630.500.400 Receptor, EphA8
New Tree	D12.776.543.750.630.500.450 Receptor, EphB1
New Tree	D12.776.543.750.630.500.500 Receptor, EphB2
New Tree	D12.776.543.750.630.500.687 Receptor, EphB3
New Tree	D12.776.543.750.630.500.781 Receptor, EphB4
New Tree	D12.776.543.750.630.500.875 Receptor, EphB5
New Tree	D12.776.543.750.630.500.937 Receptor, EphB6
New Tree	D12.776.543.750.630.625 Receptors, Platelet-Derived Growth Factor
New Tree	D12.776.543.750.630.625.300 Receptor, Platelet-Derived Growth Factor alpha
New Tree	D12.776.543.750.630.625.400 Receptor, Platelet-Derived Growth Factor beta
New Tree	D12.776.543.750.630.687 Receptors, TIE
New Tree	D12.776.543.750.630.687.249 Receptor, TIE-1
New Tree	D12.776.543.750.630.687.500 Receptor, TIE-2
New Tree	D12.776.543.750.630.750 Receptors, Vascular Endothelial Growth Factor
New Tree	D12.776.543.750.630.750.100 Vascular Endothelial Growth Factor Receptor-1
New Tree	D12.776.543.750.630.750.200 Vascular Endothelial Growth Factor Receptor-2
New Tree	D12.776.543.750.630.750.300 Vascular Endothelial Growth Factor Receptor-3

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	D12.776.543.750.650 Receptors, Adipokine
New Tree	D12.776.543.750.650.249 Receptors, Adiponectin
New Tree	D12.776.543.750.650.500 Receptors, Leptin
New Tree	D12.776.543.750.655 Receptors, Artificial
New Tree	D12.776.543.750.660 Receptors, Autocrine Motility Factor
New Tree	D12.776.543.750.670 Receptors, Biogenic Amine
New Tree	D12.776.543.750.670.300 Receptors, Catecholamine
New Tree	D12.776.543.750.670.300.300 Receptors, Adrenergic
New Tree	D12.776.543.750.670.300.300.300 Receptors, Adrenergic, alpha
New Tree	D12.776.543.750.670.300.300.300.100 Receptors, Adrenergic, alpha-1
New Tree	D12.776.543.750.670.300.300.300.200 Receptors, Adrenergic, alpha-2
New Tree	D12.776.543.750.670.300.300.340 Receptors, Adrenergic, beta
New Tree	D12.776.543.750.670.300.300.340.100 Receptors, Adrenergic, beta-1
New Tree	D12.776.543.750.670.300.300.340.200 Receptors, Adrenergic, beta-2
New Tree	D12.776.543.750.670.300.300.340.300 Receptors, Adrenergic, beta-3
New Tree	D12.776.543.750.670.300.400 Receptors, Dopamine
New Tree	D12.776.543.750.670.300.400.400 Receptors, Dopamine D1
New Tree	D12.776.543.750.670.300.400.400.500 Receptors, Dopamine D5
New Tree	D12.776.543.750.670.300.400.500 Receptors, Dopamine D2
New Tree	D12.776.543.750.670.300.400.500.249 Receptors, Dopamine D3
New Tree	D12.776.543.750.670.300.400.500.500 Receptors, Dopamine D4
New Tree	D12.776.543.750.670.450 Receptors, Histamine

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	D12.776.543.750.670.450.300 Receptors, Histamine H1
New Tree	D12.776.543.750.670.450.400 Receptors, Histamine H2
New Tree	D12.776.543.750.670.450.500 Receptors, Histamine H3
New Tree	D12.776.543.750.670.800 Receptors, Serotonin
New Tree	D12.776.543.750.670.800.100 Receptors, Serotonin, 5-HT1
New Tree	D12.776.543.750.670.800.100.100 Receptor, Serotonin, 5-HT1A
New Tree	D12.776.543.750.670.800.100.150 Receptor, Serotonin, 5-HT1B
New Tree	D12.776.543.750.670.800.100.200 Receptor, Serotonin, 5-HT1D
New Tree	D12.776.543.750.670.800.200 Receptors, Serotonin, 5-HT2
New Tree	D12.776.543.750.670.800.200.100 Receptor, Serotonin, 5-HT2A
New Tree	D12.776.543.750.670.800.200.150 Receptor, Serotonin, 5-HT2B
New Tree	D12.776.543.750.670.800.200.200 Receptor, Serotonin, 5-HT2C
New Tree	D12.776.543.750.670.800.300 Receptors, Serotonin, 5-HT3
New Tree	D12.776.543.750.670.800.400 Receptors, Serotonin, 5-HT4
New Tree	D12.776.543.750.685 Receptors, Collagen
New Heading	<b>D12.776.543.750.685.050 Discoidin Domain Receptors</b>
New Heading	<b>D12.776.543.750.685.050.500 Discoidin Domain Receptor 1</b>
New Heading	<b>D12.776.543.750.685.050.750 Discoidin Domain Receptor 2</b>
New Tree	D12.776.543.750.685.100 Integrin alpha1beta1
New Tree	D12.776.543.750.685.200 Integrin alpha2beta1
New Tree	D12.776.543.750.685.300 Integrin alpha3beta1
New Tree	D12.776.543.750.690 Receptors, Death Domain

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	D12.776.543.750.690.500      Antigenes, CD95
New Tree	D12.776.543.750.690.550      Edar Receptor
New Tree	D12.776.543.750.690.600      Receptors, TNF-Related Apoptosis-Inducing Ligand
New Tree	D12.776.543.750.690.725      Receptors, Tumor Necrosis Factor, Member 25
New Tree	D12.776.543.750.690.750      Receptors, Tumor Necrosis Factor, Type I
New Tree	D12.776.543.750.695      Receptors, G-Protein-Coupled
New Tree	D12.776.543.750.695.011      Calcitonin Receptor-Like Protein
New Tree	D12.776.543.750.695.017      Frizzled Receptors
New Heading	<b>D12.776.543.750.695.017.500      Smoothened Receptor</b>
New Tree	D12.776.543.750.695.021      Glucagon-Like Peptide Receptors
New Tree	D12.776.543.750.695.021.500      Glucagon-Like Peptide-1 Receptor
New Tree	D12.776.543.750.695.021.600      Glucagon-Like Peptide-2 Receptor
New Tree	D12.776.543.750.695.023      Orexin Receptors
New Tree	D12.776.543.750.695.024      Receptor, Anaphylatoxin C5a
New Tree	D12.776.543.750.695.035      Receptor, PAR-2
New Tree	D12.776.543.750.695.041      Receptors, Adrenomedullin
New Tree	D12.776.543.750.695.047      Receptors, Angiotensin
New Tree	D12.776.543.750.695.047.625      Receptor, Angiotensin, Type 1
New Tree	D12.776.543.750.695.047.687      Receptor, Angiotensin, Type 2
New Tree	D12.776.543.750.695.070      Receptors, Bombesin
New Tree	D12.776.543.750.695.080      Receptors, Bradykinin
New Tree	D12.776.543.750.695.080.249      Receptor, Bradykinin B1

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	D12.776.543.750.695.080.500 Receptor, Bradykinin B2
New Tree	D12.776.543.750.695.100 Receptors, Calcitonin
New Tree	D12.776.543.750.695.105 Receptors, Calcitonin Gene-Related Peptide
New Tree	D12.776.543.750.695.115 Receptors, Calcium-Sensing
New Tree	D12.776.543.750.695.125 Receptors, Cannabinoid
New Tree	D12.776.543.750.695.125.100 Receptor, Cannabinoid, CB1
New Tree	D12.776.543.750.695.125.200 Receptor, Cannabinoid, CB2
New Tree	D12.776.543.750.695.150 Receptors, Catecholamine
New Tree	D12.776.543.750.695.150.300 Receptors, Adrenergic
New Tree	D12.776.543.750.695.150.300.300 Receptors, Adrenergic, alpha
New Tree	D12.776.543.750.695.150.300.300.700 Receptors, Adrenergic, alpha-1
New Tree	D12.776.543.750.695.150.300.300.725 Receptors, Adrenergic, alpha-2
New Tree	D12.776.543.750.695.150.300.340 Receptors, Adrenergic, beta
New Tree	D12.776.543.750.695.150.300.340.700 Receptors, Adrenergic, beta-1
New Tree	D12.776.543.750.695.150.300.340.725 Receptors, Adrenergic, beta-2
New Tree	D12.776.543.750.695.150.300.340.900 Receptors, Adrenergic, beta-3
New Tree	D12.776.543.750.695.150.400 Receptors, Dopamine
New Tree	D12.776.543.750.695.150.400.400 Receptors, Dopamine D1
New Tree	D12.776.543.750.695.150.400.400.500 Receptors, Dopamine D5
New Tree	D12.776.543.750.695.150.400.500 Receptors, Dopamine D2
New Tree	D12.776.543.750.695.150.400.500.500 Receptors, Dopamine D3
New Tree	D12.776.543.750.695.150.400.500.750 Receptors, Dopamine D4



## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	D12.776.543.750.695.160 Receptors, Chemokine
New Tree	D12.776.543.750.695.160.150 Receptors, CCR
New Tree	D12.776.543.750.695.160.150.100 Receptors, CCR1
New Tree	D12.776.543.750.695.160.150.200 Receptors, CCR2
New Tree	D12.776.543.750.695.160.150.300 Receptors, CCR3
New Tree	D12.776.543.750.695.160.150.400 Receptors, CCR4
New Tree	D12.776.543.750.695.160.150.500 Receptors, CCR5
New Tree	D12.776.543.750.695.160.150.600 Receptors, CCR6
New Tree	D12.776.543.750.695.160.150.700 Receptors, CCR7
New Tree	D12.776.543.750.695.160.150.800 Receptors, CCR8
New Tree	D12.776.543.750.695.160.150.950 Receptors, CCR10
New Tree	D12.776.543.750.695.160.500 Receptors, CXCR
New Tree	D12.776.543.750.695.160.500.300 Receptors, CXCR3
New Tree	D12.776.543.750.695.160.500.400 Receptors, CXCR4
New Tree	D12.776.543.750.695.160.500.500 Receptors, CXCR5
New Tree	D12.776.543.750.695.160.500.750 Receptors, Interleukin-8
New Tree	D12.776.543.750.695.160.500.750.500 Receptors, Interleukin-8A
New Tree	D12.776.543.750.695.160.500.750.750 Receptors, Interleukin-8B
New Tree	D12.776.543.750.695.170 Receptors, Cholecystokinin
New Tree	D12.776.543.750.695.170.100 Receptor, Cholecystokinin A
New Tree	D12.776.543.750.695.170.200 Receptor, Cholecystokinin B
New Tree	D12.776.543.750.695.180 Receptors, Corticotropin-Releasing Hormone

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	D12.776.543.750.695.200 Receptors, Eicosanoid
New Tree	D12.776.543.750.695.200.450 Receptors, Leukotriene
New Tree	D12.776.543.750.695.200.450.300 Receptors, Leukotriene B4
New Tree	D12.776.543.750.695.200.575 Receptors, Lipoxin
New Tree	D12.776.543.750.695.200.700 Receptors, Prostaglandin
New Tree	D12.776.543.750.695.200.700.299 Receptors, Epoprostenol
New Tree	D12.776.543.750.695.200.700.600 Receptors, Prostaglandin E
New Tree	D12.776.543.750.695.200.700.600.100 Subtype Receptors, Prostaglandin E, EP1
New Tree	D12.776.543.750.695.200.700.600.200 Subtype Receptors, Prostaglandin E, EP2
New Tree	D12.776.543.750.695.200.700.600.300 Subtype Receptors, Prostaglandin E, EP3
New Tree	D12.776.543.750.695.200.700.600.400 Subtype Receptors, Prostaglandin E, EP4
New Tree	D12.776.543.750.695.200.700.800 Receptors, Thromboxane A2, Prostaglandin H2
New Tree	D12.776.543.750.695.200.800 Receptors, Thromboxane
New Tree	D12.776.543.750.695.200.800.500 Receptors, Thromboxane A2, Prostaglandin H2
New Tree	D12.776.543.750.695.220 Receptors, Endothelin
New Tree	D12.776.543.750.695.220.100 Receptor, Endothelin A
New Tree	D12.776.543.750.695.220.200 Receptor, Endothelin B
New Tree	D12.776.543.750.695.235 Receptors, Formyl Peptide
New Tree	D12.776.543.750.695.250 Receptors, FSH
New Tree	D12.776.543.750.695.300 Receptors, GABA-B
New Tree	D12.776.543.750.695.315 Receptors, Galanin
New Tree	D12.776.543.750.695.315.100 Receptor, Galanin, Type 1

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	D12.776.543.750.695.315.200 Receptor, Galanin, Type 2
New Tree	D12.776.543.750.695.315.300 Receptor, Galanin, Type 3
New Tree	D12.776.543.750.695.322 Receptors, Ghrelin
New Tree	D12.776.543.750.695.330 Receptors, Glucagon
New Tree	D12.776.543.750.695.350 Receptors, Histamine H1
New Tree	D12.776.543.750.695.355 Receptors, Histamine H2
New Tree	D12.776.543.750.695.360 Receptors, Islet Amyloid Polypeptide
New Tree	D12.776.543.750.695.400 Receptors, LH
New Tree	D12.776.543.750.695.410 Receptors, LHRH
New Tree	D12.776.543.750.695.420 Receptors, Lysophospholipid
New Tree	D12.776.543.750.695.420.249 Receptors, Lysophosphatidic Acid
New Tree	D12.776.543.750.695.420.500 Receptors, Lysosphingolipid
New Tree	D12.776.543.750.695.430 Receptors, Melanocortin
New Tree	D12.776.543.750.695.430.500 Receptor, Melanocortin, Type 1
New Tree	D12.776.543.750.695.430.750 Receptor, Melanocortin, Type 2
New Tree	D12.776.543.750.695.430.875 Receptor, Melanocortin, Type 3
New Tree	D12.776.543.750.695.430.937 Receptor, Melanocortin, Type 4
New Tree	D12.776.543.750.695.440 Receptors, Melatonin
New Tree	D12.776.543.750.695.440.500 Receptor, Melatonin, MT1
New Tree	D12.776.543.750.695.440.750 Receptor, Melatonin, MT2
New Tree	D12.776.543.750.695.450 Receptors, Metabotropic Glutamate
New Tree	D12.776.543.750.695.450.500 Receptor, Metabotropic Glutamate 5

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	D12.776.543.750.695.475 Receptors, Muscarinic
New Tree	D12.776.543.750.695.475.100 Receptor, Muscarinic M1
New Tree	D12.776.543.750.695.475.200 Receptor, Muscarinic M2
New Tree	D12.776.543.750.695.475.300 Receptor, Muscarinic M3
New Tree	D12.776.543.750.695.475.400 Receptor, Muscarinic M4
New Tree	D12.776.543.750.695.475.500 Receptor, Muscarinic M5
New Tree	D12.776.543.750.695.500 Receptors, Neuropeptide Y
New Tree	D12.776.543.750.695.550 Receptors, Neurotensin
New Tree	D12.776.543.750.695.600 Receptors, Odorant
New Tree	D12.776.543.750.695.620 Receptors, Opioid
New Tree	D12.776.543.750.695.620.200 Receptors, Opioid, delta
New Tree	D12.776.543.750.695.620.400 Receptors, Opioid, kappa
New Tree	D12.776.543.750.695.620.550 Receptors, Opioid, mu
New Tree	D12.776.543.750.695.620.775 Receptors, sigma
New Tree	D12.776.543.750.695.630 Receptors, Oxytocin
New Tree	D12.776.543.750.695.650 Receptors, Parathyroid Hormone
New Tree	D12.776.543.750.695.650.100 Receptor, Parathyroid Hormone, Type 1
New Tree	D12.776.543.750.695.650.200 Receptor, Parathyroid Hormone, Type 2
New Tree	D12.776.543.750.695.660 Receptors, Pheromone
New Tree	D12.776.543.750.695.660.500 Receptors, Mating Factor
New Tree	D12.776.543.750.695.665 Receptors, Pituitary Adenylate Cyclase-Activating Polypeptide
New Tree	D12.776.543.750.695.665.500 Receptors, Pituitary Adenylate Cyclase-Activating Polypeptide, Type I

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	D12.776.543.750.695.665.750 Receptors, Vasoactive Intestinal Peptide, Type II
New Tree	D12.776.543.750.695.665.760 Receptors, Vasoactive Intestinal Polypeptide, Type I
New Tree	D12.776.543.750.695.700 Receptors, Purinergic
New Tree	D12.776.543.750.695.700.150 Receptors, Cyclic AMP
New Tree	D12.776.543.750.695.700.700 Receptors, Purinergic P1
New Tree	D12.776.543.750.695.700.700.100 Receptor, Adenosine A1
New Tree	D12.776.543.750.695.700.700.200 Receptors, Adenosine A2
New Tree	D12.776.543.750.695.700.700.200.100 Receptor, Adenosine A2A
New Tree	D12.776.543.750.695.700.700.200.200 Receptor, Adenosine A2B
New Tree	D12.776.543.750.695.700.700.300 Receptor, Adenosine A3
New Tree	D12.776.543.750.695.700.720 Receptors, Purinergic P2
New Tree	D12.776.543.750.695.700.720.250 Receptors, Purinergic P2X
New Tree	D12.776.543.750.695.700.720.250.100 Receptors, Purinergic P2X1
New Tree	D12.776.543.750.695.700.720.250.200 Receptors, Purinergic P2X2
New Tree	D12.776.543.750.695.700.720.250.300 Receptors, Purinergic P2X3
New Tree	D12.776.543.750.695.700.720.250.400 Receptors, Purinergic P2X4
New Tree	D12.776.543.750.695.700.720.250.500 Receptors, Purinergic P2X5
New Tree	D12.776.543.750.695.700.720.250.700 Receptors, Purinergic P2X7
New Tree	D12.776.543.750.695.700.720.500 Receptors, Purinergic P2Y
New Tree	D12.776.543.750.695.700.720.500.100 Receptors, Purinergic P2Y1
New Tree	D12.776.543.750.695.700.720.500.200 Receptors, Purinergic P2Y2
New Tree	D12.776.543.750.695.700.720.500.300 Receptors, Purinergic P2Y12

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	D12.776.543.750.695.800 Receptors, Serotonin
New Tree	D12.776.543.750.695.800.100 Receptors, Serotonin, 5-HT1
New Tree	D12.776.543.750.695.800.100.100 Receptor, Serotonin, 5-HT1A
New Tree	D12.776.543.750.695.800.100.150 Receptor, Serotonin, 5-HT1B
New Tree	D12.776.543.750.695.800.100.200 Receptor, Serotonin, 5-HT1D
New Tree	D12.776.543.750.695.800.200 Receptors, Serotonin, 5-HT2
New Tree	D12.776.543.750.695.800.200.100 Receptor, Serotonin, 5-HT2A
New Tree	D12.776.543.750.695.800.200.150 Receptor, Serotonin, 5-HT2B
New Tree	D12.776.543.750.695.800.200.200 Receptor, Serotonin, 5-HT2C
New Tree	D12.776.543.750.695.800.400 Receptors, Serotonin, 5-HT4
New Tree	D12.776.543.750.695.850 Receptors, Somatostatin
New Tree	D12.776.543.750.695.862 Receptors, Tachykinin
New Tree	D12.776.543.750.695.862.500 Receptors, Neurokinin-1
New Tree	D12.776.543.750.695.862.540 Receptors, Neurokinin-2
New Tree	D12.776.543.750.695.862.580 Receptors, Neurokinin-3
New Tree	D12.776.543.750.695.875 Receptors, Thrombin
New Tree	D12.776.543.750.695.875.500 Receptor, PAR-1
New Tree	D12.776.543.750.695.900 Receptors, Vasoactive Intestinal Peptide
New Tree	D12.776.543.750.695.900.500 Receptors, Vasoactive Intestinal Peptide, Type II
New Tree	D12.776.543.750.695.900.520 Receptors, Vasoactive Intestinal Polypeptide, Type I
New Tree	D12.776.543.750.695.910 Receptors, Vasopressin
New Tree	D12.776.543.750.695.955 Rhodopsin

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	<a href="#">D12.776.543.750.700</a>	<a href="#">Receptors, Guanylate Cyclase-Coupled</a>
New Tree	<a href="#">D12.776.543.750.700.500</a>	<a href="#">Receptors, Atrial Natriuretic Factor</a>
-	D12.776.543.750.705	Receptors, Immunologic
-	D12.776.543.750.705.045	Antigens, CD14
-	D12.776.543.750.705.222	Costimulatory and Inhibitory T-Cell Receptors
-	D12.776.543.750.705.222.500	Antigens, CD28
-	D12.776.543.750.705.222.750	CTLA-4 Antigen
-	D12.776.543.750.705.222.812	Inducible T-Cell Co-Stimulator Protein
-	D12.776.543.750.705.222.875	Programmed Cell Death 1 Receptor
-	D12.776.543.750.705.400	Immunophilins
-	D12.776.543.750.705.400.300	Cyclophilins
-	D12.776.543.750.705.400.700	Tacrolimus Binding Proteins
-	D12.776.543.750.705.400.700.500	Tacrolimus Binding Protein 1A
-	D12.776.543.750.705.408	Integrins
-	D12.776.543.750.705.408.100	Integrin alpha Chains
-	D12.776.543.750.705.408.100.100	Antigens, CD11a
-	D12.776.543.750.705.408.100.150	Antigens, CD11b
-	D12.776.543.750.705.408.100.200	Antigens, CD11c
-	D12.776.543.750.705.408.100.300	Integrin alpha1
-	D12.776.543.750.705.408.100.350	Integrin alpha2
-	D12.776.543.750.705.408.100.400	Integrin alpha3
-	D12.776.543.750.705.408.100.450	Integrin alpha4
-	D12.776.543.750.705.408.100.500	Integrin alpha5
-	D12.776.543.750.705.408.100.550	Integrin alpha6
-	D12.776.543.750.705.408.100.900	Integrin alphaV
-	D12.776.543.750.705.408.100.950	Platelet Membrane Glycoprotein IIb
-	D12.776.543.750.705.408.200	Integrin beta Chains
-	D12.776.543.750.705.408.200.249	Antigens, CD18
-	D12.776.543.750.705.408.200.500	Antigens, CD29
-	D12.776.543.750.705.408.200.750	Integrin beta3
-	D12.776.543.750.705.408.200.800	Integrin beta4
-	D12.776.543.750.705.408.460	Receptors, Cytoadhesin
-	D12.776.543.750.705.408.460.170	Integrin alpha6beta4
-	D12.776.543.750.705.408.460.700	Platelet Glycoprotein GPIIb-IIIa Complex

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.543.750.705.408.460.870 Receptors, Vitronectin
-	D12.776.543.750.705.408.460.870.500 Integrin alphaVbeta3
-	D12.776.543.750.705.408.495 Receptors, Fibrinogen
-	D12.776.543.750.705.408.495.500 Macrophage-1 Antigen
-	D12.776.543.750.705.408.530 Receptors, Fibronectin
-	D12.776.543.750.705.408.530.249 Integrin alpha2beta1
-	D12.776.543.750.705.408.530.500 Integrin alpha4beta1
-	D12.776.543.750.705.408.530.750 Integrin alpha5beta1
-	D12.776.543.750.705.408.600 Receptors, Leukocyte-Adhesion
-	D12.776.543.750.705.408.600.100 Integrin alphaXbeta2
-	D12.776.543.750.705.408.600.100.500 Antigens, CD11c
-	D12.776.543.750.705.408.600.100.750 Antigens, CD18
-	D12.776.543.750.705.408.600.400 Lymphocyte Function-Associated Antigen-1
-	D12.776.543.750.705.408.600.500 Macrophage-1 Antigen
-	D12.776.543.750.705.408.850 Receptors, Very Late Antigen
-	D12.776.543.750.705.408.850.099 Integrin alpha1beta1
-	D12.776.543.750.705.408.850.199 Integrin alpha2beta1
-	D12.776.543.750.705.408.850.249 Integrin alpha3beta1
-	D12.776.543.750.705.408.850.299 Integrin alpha4beta1
-	D12.776.543.750.705.408.850.349 Integrin alpha5beta1
-	D12.776.543.750.705.408.850.674 Integrin alpha6beta1
-	D12.776.543.750.705.675 Platelet Membrane Glycoproteins
-	D12.776.543.750.705.675.136 Antigens, CD36
-	D12.776.543.750.705.675.352 Integrin alpha2beta1
-	D12.776.543.750.705.675.460 Integrin alpha5beta1
-	D12.776.543.750.705.675.514 Integrin alpha6beta1
-	D12.776.543.750.705.675.541 Integrin alphaVbeta3
-	D12.776.543.750.705.675.554 Lysosomal-Associated Membrane Protein 1
-	D12.776.543.750.705.675.568 Platelet Glycoprotein GPIb-IX Complex
-	D12.776.543.750.705.675.784 Platelet Glycoprotein GPIIb-IIIa Complex
-	D12.776.543.750.705.675.892 Receptors, Thrombin
-	D12.776.543.750.705.675.892.790 Receptor, PAR-1
-	D12.776.543.750.705.675.892.800 Thrombomodulin
-	D12.776.543.750.705.816 Receptors, Antigen
-	D12.776.543.750.705.816.821 Receptors, Antigen, B-Cell
-	D12.776.543.750.705.816.821.500 Antigens, CD79



## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D12.776.543.750.705.816.821.750	Pre-B Cell Receptors
-	D12.776.543.750.705.816.821.750.500 Surrogate	Immunoglobulin Light Chains,
-	D12.776.543.750.705.816.824	Receptors, Antigen, T-Cell
-	D12.776.543.750.705.816.824.300	Complementarity Determining Regions
-	D12.776.543.750.705.816.824.800	Receptor-CD3 Complex, Antigen, T-Cell
-	D12.776.543.750.705.816.824.825	Receptors, Antigen, T-Cell, alpha-beta
-	D12.776.543.750.705.816.824.830	Receptors, Antigen, T-Cell, gamma-delta
-	D12.776.543.750.705.833	Receptors, Complement
New Tree	<a href="#">D12.776.543.750.705.833.062</a>	<a href="#">Antigens, CD11b</a>
-	D12.776.543.750.705.833.124	Antigens, CD46
-	D12.776.543.750.705.833.249	Integrin alphaXbeta2
-	D12.776.543.750.705.833.249.500	Antigens, CD11c
-	D12.776.543.750.705.833.249.750	Antigens, CD18
-	D12.776.543.750.705.833.500	Macrophage-1 Antigen
-	D12.776.543.750.705.833.550	Receptor, Anaphylatoxin C5a
-	D12.776.543.750.705.833.600	Receptors, Complement 3b
-	D12.776.543.750.705.833.610	Receptors, Complement 3d
-	D12.776.543.750.705.852	Receptors, Cytokine
-	D12.776.543.750.705.852.125	Receptors, Chemokine
-	D12.776.543.750.705.852.125.150	Receptors, CCR
-	D12.776.543.750.705.852.125.150.100	Receptors, CCR1
-	D12.776.543.750.705.852.125.150.200	Receptors, CCR2
-	D12.776.543.750.705.852.125.150.300	Receptors, CCR3
-	D12.776.543.750.705.852.125.150.400	Receptors, CCR4
-	D12.776.543.750.705.852.125.150.500	Receptors, CCR5
-	D12.776.543.750.705.852.125.150.600	Receptors, CCR6
-	D12.776.543.750.705.852.125.150.700	Receptors, CCR7
-	D12.776.543.750.705.852.125.150.800	Receptors, CCR8
-	D12.776.543.750.705.852.125.150.950	Receptors, CCR10
-	D12.776.543.750.705.852.125.500	Receptors, CXCR
-	D12.776.543.750.705.852.125.500.300	Receptors, CXCR3
-	D12.776.543.750.705.852.125.500.400	Receptors, CXCR4
-	D12.776.543.750.705.852.125.500.500	Receptors, CXCR5
-	D12.776.543.750.705.852.125.500.750	Receptors, Interleukin-8

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D12.776.543.750.705.852.125.500.750.500	Receptors, Interleukin-8A
-	D12.776.543.750.705.852.125.500.750.750	Receptors, Interleukin-8B
-	D12.776.543.750.705.852.150	Receptors, Colony-Stimulating Factor
-	D12.776.543.750.705.852.150.100	Proto-Oncogene Proteins c-kit
-	D12.776.543.750.705.852.150.150 Factor	Receptor, Macrophage Colony-Stimulating
-	D12.776.543.750.705.852.150.200	Receptors, Erythropoietin
-	D12.776.543.750.705.852.150.280 Factor	Receptors, Granulocyte Colony-Stimulating
-	D12.776.543.750.705.852.150.310 Colony-Stimulating Factor	Receptors, Granulocyte-Macrophage
-	D12.776.543.750.705.852.150.310.500 Subunit	Cytokine Receptor Common beta
-	D12.776.543.750.705.852.150.360	Receptors, Interleukin-3
-	D12.776.543.750.705.852.400	Receptors, Interferon
-	D12.776.543.750.705.852.400.500	Receptor, Interferon alpha-beta
-	D12.776.543.750.705.852.420	Receptors, Interleukin
-	D12.776.543.750.705.852.420.300	Receptors, Interleukin-1
-	D12.776.543.750.705.852.420.300.249 Protein	Interleukin-1 Receptor Accessory
New Heading	<b>D12.776.543.750.705.852.420.300.375</b>	<b>Interleukin-1 Receptor-Like 1 Protein</b>
-	D12.776.543.750.705.852.420.300.500	Receptors, Interleukin-1 Type I
-	D12.776.543.750.705.852.420.300.750	Receptors, Interleukin-1 Type II
-	D12.776.543.750.705.852.420.320	Receptors, Interleukin-2
-	D12.776.543.750.705.852.420.320.500	Interleukin-2 Receptor alpha Subunit
-	D12.776.543.750.705.852.420.320.750	Interleukin-2 Receptor beta Subunit
-	D12.776.543.750.705.852.420.320.875 Subunit	Interleukin Receptor Common gamma
-	D12.776.543.750.705.852.420.340	Receptors, Interleukin-3
-	D12.776.543.750.705.852.420.340.350 Subunit	Cytokine Receptor Common beta
-	D12.776.543.750.705.852.420.340.500	Interleukin-3 Receptor alpha Subunit
-	D12.776.543.750.705.852.420.360	Receptors, Interleukin-4
-	D12.776.543.750.705.852.420.360.300	Receptors, Interleukin-4, Type I
-	D12.776.543.750.705.852.420.360.300.200 Subunit	Interleukin-4 Receptor alpha
-	D12.776.543.750.705.852.420.360.300.500 gamma Subunit	Interleukin Receptor Common

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.543.750.705.852.420.360.600 Receptors, Interleukin-4, Type II
-	D12.776.543.750.705.852.420.360.600.200 Subunit Interleukin-4 Receptor alpha
-	D12.776.543.750.705.852.420.360.600.500 Subunit Interleukin-13 Receptor alpha1
-	D12.776.543.750.705.852.420.380 Receptors, Interleukin-5
-	D12.776.543.750.705.852.420.380.500 Subunit Cytokine Receptor Common beta
-	D12.776.543.750.705.852.420.380.750 Interleukin-5 Receptor alpha Subunit
-	D12.776.543.750.705.852.420.400 Receptors, Interleukin-6
-	D12.776.543.750.705.852.420.400.500 Cytokine Receptor gp130
-	D12.776.543.750.705.852.420.400.750 Interleukin-6 Receptor alpha Subunit
-	D12.776.543.750.705.852.420.420 Receptors, Interleukin-7
-	D12.776.543.750.705.852.420.420.450 Interleukin-7 Receptor alpha Subunit
-	D12.776.543.750.705.852.420.420.500 Subunit Interleukin Receptor Common gamma
-	D12.776.543.750.705.852.420.421 Receptors, Interleukin-8
-	D12.776.543.750.705.852.420.421.500 Receptors, Interleukin-8A
-	D12.776.543.750.705.852.420.421.750 Receptors, Interleukin-8B
-	D12.776.543.750.705.852.420.500 Receptors, Interleukin-9
-	D12.776.543.750.705.852.420.500.500 Subunit Interleukin Receptor Common gamma
-	D12.776.543.750.705.852.420.550 Receptors, Interleukin-10
-	D12.776.543.750.705.852.420.550.500 Interleukin-10 Receptor alpha Subunit
-	D12.776.543.750.705.852.420.550.750 Interleukin-10 Receptor beta Subunit
-	D12.776.543.750.705.852.420.560 Receptors, Interleukin-11
-	D12.776.543.750.705.852.420.560.249 Cytokine Receptor gp130
-	D12.776.543.750.705.852.420.560.500 Interleukin-11 Receptor alpha Subunit
-	D12.776.543.750.705.852.420.580 Receptors, Interleukin-12
-	D12.776.543.750.705.852.420.580.500 Interleukin-12 Receptor beta 1 Subunit
-	D12.776.543.750.705.852.420.580.750 Interleukin-12 Receptor beta 2 Subunit
-	D12.776.543.750.705.852.420.600 Receptors, Interleukin-13
-	D12.776.543.750.705.852.420.600.300 Interleukin-13 Receptor alpha2 Subunit
-	D12.776.543.750.705.852.420.600.800 Receptors, Interleukin-4, Type II
-	D12.776.543.750.705.852.420.600.800.200 Subunit Interleukin-4 Receptor alpha
-	D12.776.543.750.705.852.420.600.800.500 Subunit Interleukin-13 Receptor alpha1

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.543.750.705.852.420.800 Receptors, Interleukin-15
-	D12.776.543.750.705.852.420.800.500 Interleukin-2 Receptor beta Subunit
-	D12.776.543.750.705.852.420.800.550 Interleukin-15 Receptor alpha Subunit
-	D12.776.543.750.705.852.420.800.750 Interleukin Receptor Common gamma Subunit
-	D12.776.543.750.705.852.420.810 Receptors, Interleukin-16
-	D12.776.543.750.705.852.420.810.500 Antigens, CD4
-	D12.776.543.750.705.852.420.820 Receptors, Interleukin-17
-	D12.776.543.750.705.852.420.830 Receptors, Interleukin-18
-	D12.776.543.750.705.852.420.830.500 Interleukin-18 Receptor alpha Subunit
-	D12.776.543.750.705.852.420.830.750 Interleukin-18 Receptor beta Subunit
-	D12.776.543.750.705.852.420.900 Receptors, Interleukin-21
-	D12.776.543.750.705.852.420.900.450 Interleukin-21 Receptor alpha Subunit
-	D12.776.543.750.705.852.420.900.500 Interleukin Receptor Common gamma Subunit
-	D12.776.543.750.705.852.555 Receptors, Oncostatin M
-	D12.776.543.750.705.852.555.600 Receptors, Oncostatin M, Type II
-	D12.776.543.750.705.852.555.600.500 Cytokine Receptor gp130
-	D12.776.543.750.705.852.555.600.750 Oncostatin M Receptor beta Subunit
-	D12.776.543.750.705.852.583 Receptors, OSM-LIF
-	D12.776.543.750.705.852.583.500 Cytokine Receptor gp130
-	D12.776.543.750.705.852.583.750 Leukemia Inhibitory Factor Receptor alpha Subunit
-	D12.776.543.750.705.852.610 Receptors, Thrombopoietin
-	D12.776.543.750.705.852.720 Receptors, Transforming Growth Factor beta
-	D12.776.543.750.705.852.760 Receptors, Tumor Necrosis Factor
-	D12.776.543.750.705.852.760.048 Antigens, CD27
-	D12.776.543.750.705.852.760.072 Antigens, CD30
-	D12.776.543.750.705.852.760.097 Antigens, CD40
-	D12.776.543.750.705.852.760.195 Antigens, CD95
-	D12.776.543.750.705.852.760.220 Antigens, CD137
-	D12.776.543.750.705.852.760.222 B-Cell Activation Factor Receptor
-	D12.776.543.750.705.852.760.226 B-Cell Maturation Antigen
-	D12.776.543.750.705.852.760.232 Etanercept
-	D12.776.543.750.705.852.760.238 Glucocorticoid-Induced TNFR-Related Protein
-	D12.776.543.750.705.852.760.245 Lymphotoxin beta Receptor

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.543.750.705.852.760.345 B Receptor Activator of Nuclear Factor-kappa B
-	D12.776.543.750.705.852.760.347 Receptors, Ectodysplasin
-	D12.776.543.750.705.852.760.347.500 Edar Receptor
-	D12.776.543.750.705.852.760.347.750 Xedar Receptor
-	D12.776.543.750.705.852.760.350 Receptors, OX40
-	D12.776.543.750.705.852.760.396 Receptors, TNF-Related Apoptosis-Inducing Ligand
-	D12.776.543.750.705.852.760.500 Receptors, Tumor Necrosis Factor, Member 14
-	D12.776.543.750.705.852.760.505 Receptors, Tumor Necrosis Factor, Member 25
-	D12.776.543.750.705.852.760.597 Receptors, Tumor Necrosis Factor, Type I
-	D12.776.543.750.705.852.760.798 Receptors, Tumor Necrosis Factor, Type II
-	D12.776.543.750.705.852.760.899 Transmembrane Activator and CAML Interactor Protein
-	D12.776.543.750.705.852.760.949 Tumor Necrosis Factor Decoy Receptors
-	D12.776.543.750.705.852.760.949.249 Osteoprotegerin
-	D12.776.543.750.705.852.760.949.400 Receptors, Tumor Necrosis Factor, Member 6b
-	D12.776.543.750.705.852.760.949.500 Receptors, Tumor Necrosis Factor, Member 10c
-	D12.776.543.750.705.871 Receptors, Fc
-	D12.776.543.750.705.871.280 Receptors, IgE
-	D12.776.543.750.705.871.300 Receptors, IgG
-	D12.776.543.750.705.871.700 Receptors, Polymeric Immunoglobulin
-	D12.776.543.750.705.873 Receptors, Formyl Peptide
-	D12.776.543.750.705.876 Receptors, Laminin
-	D12.776.543.750.705.876.249 Integrin alpha1beta1
-	D12.776.543.750.705.876.280 Integrin alpha2beta1
-	D12.776.543.750.705.876.311 Integrin alpha3beta1
-	D12.776.543.750.705.876.374 Integrin alpha5beta1
-	D12.776.543.750.705.876.500 Integrin alpha6beta1
-	D12.776.543.750.705.876.750 Integrin alpha6beta4
-	D12.776.543.750.705.877 Receptors, Lymphocyte Homing
-	D12.776.543.750.705.877.144 Antigens, CD44
-	D12.776.543.750.705.877.347 Integrin alpha4beta1
-	D12.776.543.750.705.877.550 Lymphocyte Function-Associated Antigen-1

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.543.750.705.877.903 L-Selectin
-	D12.776.543.750.705.880 Receptors, Mitogen
New Heading	<b>D12.776.543.750.705.880.300 Discoidin Domain Receptors</b>
New Heading	<b>D12.776.543.750.705.880.300.500 Discoidin Domain Receptor 1</b>
New Heading	<b>D12.776.543.750.705.880.300.750 Discoidin Domain Receptor 2</b>
-	D12.776.543.750.705.880.600 Receptors, Concanavalin A
-	D12.776.543.750.705.895 Receptors, Natural Killer Cell
-	D12.776.543.750.705.895.500 Receptors, KIR
-	D12.776.543.750.705.895.500.249 Receptors, KIR2DL1
-	D12.776.543.750.705.895.500.374 Receptors, KIR2DL2
-	D12.776.543.750.705.895.500.437 Receptors, KIR2DL3
-	D12.776.543.750.705.895.500.468 Receptors, KIR2DL4
-	D12.776.543.750.705.895.500.484 Receptors, KIR2DL5
-	D12.776.543.750.705.895.500.500 Receptors, KIR3DL1
-	D12.776.543.750.705.895.500.750 Receptors, KIR3DL2
-	D12.776.543.750.705.895.500.890 Receptors, KIR3DS1
-	D12.776.543.750.705.895.750 Receptors, Natural Cytotoxicity Triggering
-	D12.776.543.750.705.895.750.100 Natural Cytotoxicity Triggering Receptor 1
-	D12.776.543.750.705.895.750.200 Natural Cytotoxicity Triggering Receptor 2
-	D12.776.543.750.705.895.750.300 Natural Cytotoxicity Triggering Receptor 3
-	D12.776.543.750.705.895.800 Receptors, NK Cell Lectin-Like
-	D12.776.543.750.705.895.800.100 NK Cell Lectin-Like Receptor Subfamily A
-	D12.776.543.750.705.895.800.200 NK Cell Lectin-Like Receptor Subfamily B
-	D12.776.543.750.705.895.800.810 NK Cell Lectin-Like Receptor Subfamily C
-	D12.776.543.750.705.895.800.820 NK Cell Lectin-Like Receptor Subfamily D
-	D12.776.543.750.705.895.800.910 NK Cell Lectin-Like Receptor Subfamily K
-	D12.776.543.750.705.910 Receptors, Pattern Recognition
-	D12.776.543.750.705.910.500 Toll-Like Receptors
-	D12.776.543.750.705.910.500.100 Toll-Like Receptor 1
-	D12.776.543.750.705.910.500.200 Toll-Like Receptor 2
-	D12.776.543.750.705.910.500.300 Toll-Like Receptor 3
-	D12.776.543.750.705.910.500.400 Toll-Like Receptor 4
-	D12.776.543.750.705.910.500.500 Toll-Like Receptor 5
-	D12.776.543.750.705.910.500.600 Toll-Like Receptor 6

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D12.776.543.750.705.910.500.700	Toll-Like Receptor 7
-	D12.776.543.750.705.910.500.800	Toll-Like Receptor 8
-	D12.776.543.750.705.910.500.900	Toll-Like Receptor 9
-	D12.776.543.750.705.910.500.910	Toll-Like Receptor 10
-	D12.776.543.750.705.940	Receptors, Scavenger
-	D12.776.543.750.705.940.500	Scavenger Receptors, Class A
-	D12.776.543.750.705.940.625	Scavenger Receptors, Class B
-	D12.776.543.750.705.940.625.249	Antigens, CD36
-	D12.776.543.750.705.940.687	Scavenger Receptors, Class C
-	D12.776.543.750.705.940.718	Scavenger Receptors, Class D
-	D12.776.543.750.705.940.734	Scavenger Receptors, Class E
-	D12.776.543.750.705.940.742	Scavenger Receptors, Class F
New Heading	<b>D12.776.543.750.705.970 Family</b>	<b>Signaling Lymphocytic Activation Molecule</b>
New Heading	<b>D12.776.543.750.705.970.250</b>	<b>CD48 Antigen</b>
New Heading	<b>D12.776.543.750.705.970.500 Family Member 1</b>	<b>Signaling Lymphocytic Activation Molecule</b>
-	D12.776.543.750.710	Receptors, Lipoprotein
-	D12.776.543.750.710.450	Receptors, LDL
-	D12.776.543.750.710.450.500	LDL-Receptor Related Proteins
-	D12.776.543.750.710.450.500.100 Protein-1	Low Density Lipoprotein Receptor-Related
-	D12.776.543.750.710.450.500.200 Protein-2	Low Density Lipoprotein Receptor-Related
-	D12.776.543.750.710.450.500.400 Protein-5	Low Density Lipoprotein Receptor-Related
-	D12.776.543.750.710.450.500.600 Protein-6	Low Density Lipoprotein Receptor-Related
-	D12.776.543.750.710.450.625	Receptors, Oxidized LDL
-	D12.776.543.750.710.450.625.500	Scavenger Receptors, Class E
-	D12.776.543.750.710.450.750	Receptors, Scavenger
-	D12.776.543.750.710.450.750.500	Scavenger Receptors, Class A
-	D12.776.543.750.710.450.750.625	Scavenger Receptors, Class B
-	D12.776.543.750.710.450.750.625.249	Antigens, CD36
-	D12.776.543.750.710.450.750.687	Scavenger Receptors, Class C
-	D12.776.543.750.710.450.750.718	Scavenger Receptors, Class D
-	D12.776.543.750.710.450.750.734	Scavenger Receptors, Class E

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.543.750.710.450.750.742 Scavenger Receptors, Class F
-	D12.776.543.750.715 Receptors, N-Acetylglucosamine
-	D12.776.543.750.720 Receptors, Neurotransmitter
-	D12.776.543.750.720.200 Receptors, Amino Acid
-	D12.776.543.750.720.200.300 Receptors, GABA
-	D12.776.543.750.720.200.300.300 Receptors, GABA-A
-	D12.776.543.750.720.200.300.320 Receptors, GABA-B
-	D12.776.543.750.720.200.450 Receptors, Glutamate
-	D12.776.543.750.720.200.450.400 Receptors, Ionotropic Glutamate
-	D12.776.543.750.720.200.450.400.100 Receptors, AMPA
-	D12.776.543.750.720.200.450.400.200 Receptors, Kainic Acid
-	D12.776.543.750.720.200.450.400.500 Receptors, N-Methyl-D-Aspartate
-	D12.776.543.750.720.200.450.500 Receptors, Metabotropic Glutamate
-	D12.776.543.750.720.200.450.500.500 Receptor, Metabotropic Glutamate 5
-	D12.776.543.750.720.200.470 Receptors, Glycine
-	D12.776.543.750.720.330 Receptors, Catecholamine
-	D12.776.543.750.720.330.300 Receptors, Adrenergic
-	D12.776.543.750.720.330.300.300 Receptors, Adrenergic, alpha
-	D12.776.543.750.720.330.300.300.100 Receptors, Adrenergic, alpha-1
-	D12.776.543.750.720.330.300.300.200 Receptors, Adrenergic, alpha-2
-	D12.776.543.750.720.330.300.340 Receptors, Adrenergic, beta
-	D12.776.543.750.720.330.300.340.100 Receptors, Adrenergic, beta-1
-	D12.776.543.750.720.330.300.340.200 Receptors, Adrenergic, beta-2
-	D12.776.543.750.720.330.300.340.300 Receptors, Adrenergic, beta-3
-	D12.776.543.750.720.330.400 Receptors, Dopamine
-	D12.776.543.750.720.330.400.400 Receptors, Dopamine D1
-	D12.776.543.750.720.330.400.400.500 Receptors, Dopamine D5
-	D12.776.543.750.720.330.400.500 Receptors, Dopamine D2
-	D12.776.543.750.720.330.400.500.249 Receptors, Dopamine D3
-	D12.776.543.750.720.330.400.500.500 Receptors, Dopamine D4
-	D12.776.543.750.720.360 Receptors, Cholinergic
-	D12.776.543.750.720.360.500 Receptors, Muscarinic
-	D12.776.543.750.720.360.500.099 Receptor, Muscarinic M1
-	D12.776.543.750.720.360.500.200 Receptor, Muscarinic M2
-	D12.776.543.750.720.360.500.300 Receptor, Muscarinic M3
-	D12.776.543.750.720.360.500.400 Receptor, Muscarinic M4



## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.543.750.720.360.500.500 Receptor, Muscarinic M5
-	D12.776.543.750.720.360.550 Receptors, Nicotinic
-	D12.776.543.750.720.360.550.500 alpha7 Nicotinic Acetylcholine Receptor
-	D12.776.543.750.720.480 Receptors, Histamine
-	D12.776.543.750.720.480.300 Receptors, Histamine H1
-	D12.776.543.750.720.480.400 Receptors, Histamine H2
-	D12.776.543.750.720.480.500 Receptors, Histamine H3
-	D12.776.543.750.720.600 Receptors, Neuropeptide
-	D12.776.543.750.720.600.074 Orexin Receptors
-	D12.776.543.750.720.600.200 Receptors, Bombesin
-	D12.776.543.750.720.600.220 Receptors, Bradykinin
-	D12.776.543.750.720.600.220.249 Receptor, Bradykinin B1
-	D12.776.543.750.720.600.220.500 Receptor, Bradykinin B2
-	D12.776.543.750.720.600.260 Receptors, Calcitonin Gene-Related Peptide
-	D12.776.543.750.720.600.270 Receptors, Cholecystokinin
-	D12.776.543.750.720.600.270.100 Receptor, Cholecystokinin A
-	D12.776.543.750.720.600.270.200 Receptor, Cholecystokinin B
-	D12.776.543.750.720.600.285 Receptors, Corticotropin
-	D12.776.543.750.720.600.285.500 Receptors, Melanocortin
-	D12.776.543.750.720.600.285.500.500 Receptor, Melanocortin, Type 1
-	D12.776.543.750.720.600.285.500.750 Receptor, Melanocortin, Type 2
-	D12.776.543.750.720.600.285.500.875 Receptor, Melanocortin, Type 3
-	D12.776.543.750.720.600.285.500.937 Receptor, Melanocortin, Type 4
-	D12.776.543.750.720.600.290 Receptors, Corticotropin-Releasing Hormone
-	D12.776.543.750.720.600.370 Receptors, FSH
-	D12.776.543.750.720.600.410 Receptors, Galanin
-	D12.776.543.750.720.600.450 Receptors, LH
-	D12.776.543.750.720.600.460 Receptors, LHRH
-	D12.776.543.750.720.600.540 Receptors, Neuropeptide Y
-	D12.776.543.750.720.600.560 Receptors, Neurotensin
-	D12.776.543.750.720.600.610 Receptors, Opioid
-	D12.776.543.750.720.600.610.200 Receptors, Opioid, delta
-	D12.776.543.750.720.600.610.400 Receptors, Opioid, kappa
-	D12.776.543.750.720.600.610.550 Receptors, Opioid, mu
-	D12.776.543.750.720.600.610.775 Receptors, sigma
-	D12.776.543.750.720.600.630 Receptors, Oxytocin

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.543.750.720.600.710 Receptors, Prolactin
-	D12.776.543.750.720.600.760 Receptors, Somatostatin
-	D12.776.543.750.720.600.830 Receptors, Tachykinin
-	D12.776.543.750.720.600.830.500 Receptors, Neurokinin-1
-	D12.776.543.750.720.600.830.540 Receptors, Neurokinin-2
-	D12.776.543.750.720.600.830.580 Receptors, Neurokinin-3
-	D12.776.543.750.720.600.850 Receptors, Thyrotropin
-	D12.776.543.750.720.600.860 Receptors, Thyrotropin-Releasing Hormone
-	D12.776.543.750.720.600.915 Receptors, Vasoactive Intestinal Peptide
-	D12.776.543.750.720.600.925 Receptors, Vasopressin
-	D12.776.543.750.720.670 Receptors, Presynaptic
-	D12.776.543.750.720.670.100 Autoreceptors
-	D12.776.543.750.720.670.100.500 Receptor, Serotonin, 5-HT1B
-	D12.776.543.750.720.670.100.750 Receptor, Serotonin, 5-HT1D
-	D12.776.543.750.720.700 Receptors, Purinergic
-	D12.776.543.750.720.700.150 Receptors, Cyclic AMP
-	D12.776.543.750.720.700.700 Receptors, Purinergic P1
-	D12.776.543.750.720.700.700.100 Receptor, Adenosine A1
-	D12.776.543.750.720.700.700.200 Receptors, Adenosine A2
-	D12.776.543.750.720.700.700.200.100 Receptor, Adenosine A2A
-	D12.776.543.750.720.700.700.200.200 Receptor, Adenosine A2B
-	D12.776.543.750.720.700.700.300 Receptor, Adenosine A3
-	D12.776.543.750.720.700.720 Receptors, Purinergic P2
-	D12.776.543.750.720.700.720.500 Receptors, Purinergic P2X
-	D12.776.543.750.720.700.720.500.100 Receptors, Purinergic P2X1
-	D12.776.543.750.720.700.720.500.200 Receptors, Purinergic P2X2
-	D12.776.543.750.720.700.720.500.300 Receptors, Purinergic P2X3
-	D12.776.543.750.720.700.720.500.400 Receptors, Purinergic P2X4
-	D12.776.543.750.720.700.720.500.500 Receptors, Purinergic P2X5
-	D12.776.543.750.720.700.720.500.700 Receptors, Purinergic P2X7
-	D12.776.543.750.720.700.720.750 Receptors, Purinergic P2Y
-	D12.776.543.750.720.700.720.750.100 Receptors, Purinergic P2Y1
-	D12.776.543.750.720.700.720.750.200 Receptors, Purinergic P2Y2
-	D12.776.543.750.720.700.720.750.300 Receptors, Purinergic P2Y12
-	D12.776.543.750.720.850 Receptors, Serotonin
-	D12.776.543.750.720.850.100 Receptors, Serotonin, 5-HT1

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D12.776.543.750.720.850.100.100	Receptor, Serotonin, 5-HT1A
-	D12.776.543.750.720.850.100.150	Receptor, Serotonin, 5-HT1B
-	D12.776.543.750.720.850.100.200	Receptor, Serotonin, 5-HT1D
-	D12.776.543.750.720.850.200	Receptors, Serotonin, 5-HT2
-	D12.776.543.750.720.850.200.100	Receptor, Serotonin, 5-HT2A
-	D12.776.543.750.720.850.200.150	Receptor, Serotonin, 5-HT2B
-	D12.776.543.750.720.850.200.200	Receptor, Serotonin, 5-HT2C
-	D12.776.543.750.720.850.300	Receptors, Serotonin, 5-HT3
-	D12.776.543.750.720.850.400	Receptors, Serotonin, 5-HT4
-	D12.776.543.750.725	Receptors, Notch
-	D12.776.543.750.725.500	Receptor, Notch1
-	D12.776.543.750.725.750	Receptor, Notch2
New Heading	<b>D12.776.543.750.725.875</b>	<b>Receptor, Notch3</b>
-	D12.776.543.750.750	Receptors, Peptide
-	D12.776.543.750.750.049	Receptors, Adrenomedullin
-	D12.776.543.750.750.100	Receptors, Albumin
-	D12.776.543.750.750.130	Receptors, Angiotensin
-	D12.776.543.750.750.130.750	Receptor, Angiotensin, Type 1
-	D12.776.543.750.750.130.875	Receptor, Angiotensin, Type 2
-	D12.776.543.750.750.160	Receptors, Atrial Natriuretic Factor
-	D12.776.543.750.750.200	Receptors, Calcitonin
-	D12.776.543.750.750.320	Receptors, Endothelin
-	D12.776.543.750.750.320.100	Receptor, Endothelin A
-	D12.776.543.750.750.320.200	Receptor, Endothelin B
-	D12.776.543.750.750.340	Receptors, Formyl Peptide
-	D12.776.543.750.750.360	Receptors, Gastrointestinal Hormone
-	D12.776.543.750.750.360.100	Glucagon-Like Peptide Receptors
-	D12.776.543.750.750.360.100.500	Glucagon-Like Peptide-1 Receptor
-	D12.776.543.750.750.360.100.600	Glucagon-Like Peptide-2 Receptor
-	D12.776.543.750.750.360.200	Receptors, Cholecystokinin
-	D12.776.543.750.750.360.200.100	Receptor, Cholecystokinin A
-	D12.776.543.750.750.360.200.200	Receptor, Cholecystokinin B
-	D12.776.543.750.750.360.300	Receptor, Epidermal Growth Factor
-	D12.776.543.750.750.360.900	Receptors, Vasoactive Intestinal Peptide
-	D12.776.543.750.750.400	Receptors, Growth Factor

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.543.750.750.400.049 Bone Morphogenetic Protein Receptors
-	D12.776.543.750.750.400.049.500 Bone Morphogenetic Protein Receptors, Type I
-	D12.776.543.750.750.400.049.750 Bone Morphogenetic Protein Receptors, Type II
-	D12.776.543.750.750.400.074 ErbB Receptors
-	D12.776.543.750.750.400.074.300 Receptor, Epidermal Growth Factor
-	D12.776.543.750.750.400.074.400 Receptor, ErbB-2
-	D12.776.543.750.750.400.074.500 Receptor, ErbB-3
-	D12.776.543.750.750.400.074.600 Receptor, ErbB-4
-	D12.776.543.750.750.400.100 Proto-Oncogene Proteins c-met
-	D12.776.543.750.750.400.200 Receptors, Colony-Stimulating Factor
-	D12.776.543.750.750.400.200.170 Proto-Oncogene Proteins c-kit
-	D12.776.543.750.750.400.200.200 Receptor, Macrophage Colony-Stimulating Factor
-	D12.776.543.750.750.400.200.340 Receptors, Erythropoietin
-	D12.776.543.750.750.400.200.400 Receptors, Granulocyte Colony-Stimulating Factor
-	D12.776.543.750.750.400.200.420 Receptors, Granulocyte-Macrophage Colony-Stimulating Factor
-	D12.776.543.750.750.400.200.420.500 Cytokine Receptor Common beta Subunit
-	D12.776.543.750.750.400.200.480 Receptors, Interleukin-3
-	D12.776.543.750.750.400.370 Receptors, Fibroblast Growth Factor
-	D12.776.543.750.750.400.370.500 Receptor, Fibroblast Growth Factor, Type 1
-	D12.776.543.750.750.400.370.750 Receptor, Fibroblast Growth Factor, Type 2
-	D12.776.543.750.750.400.370.875 Receptor, Fibroblast Growth Factor, Type 3
-	D12.776.543.750.750.400.370.937 Receptor, Fibroblast Growth Factor, Type 4
-	D12.776.543.750.750.400.370.968 Receptor, Fibroblast Growth Factor, Type 5
-	D12.776.543.750.750.400.550 Receptors, Nerve Growth Factor
-	D12.776.543.750.750.400.550.124 Glial Cell Line-Derived Neurotrophic Factor Receptors
-	D12.776.543.750.750.400.550.250 Receptor, Ciliary Neurotrophic Factor
-	D12.776.543.750.750.400.550.250.249 Ciliary Neurotrophic Factor Receptor alpha Subunit
-	D12.776.543.750.750.400.550.250.500 Cytokine Receptor gp130
-	D12.776.543.750.750.400.550.250.750 Leukemia Inhibitory Factor Receptor alpha Subunit
-	D12.776.543.750.750.400.550.500 Receptor, Nerve Growth Factor

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.543.750.750.400.550.550 Receptor, trkA
-	D12.776.543.750.750.400.550.600 Receptor, trkB
-	D12.776.543.750.750.400.550.700 Receptor, trkC
-	D12.776.543.750.750.400.630 Receptors, Platelet-Derived Growth Factor
-	D12.776.543.750.750.400.630.300 alpha Receptor, Platelet-Derived Growth Factor
-	D12.776.543.750.750.400.630.400 beta Receptor, Platelet-Derived Growth Factor
-	D12.776.543.750.750.400.780 Receptors, Somatomedin
-	D12.776.543.750.750.400.780.400 Receptor, IGF Type 1
-	D12.776.543.750.750.400.780.410 Receptor, IGF Type 2
-	D12.776.543.750.750.400.820 Receptors, Transforming Growth Factor beta
-	D12.776.543.750.750.400.820.500 Activin Receptors
-	D12.776.543.750.750.400.820.500.500 Activin Receptors, Type I
-	D12.776.543.750.750.400.820.500.750 Activin Receptors, Type II
-	D12.776.543.750.750.400.910 Receptors, Vascular Endothelial Growth Factor
-	D12.776.543.750.750.400.910.100 Vascular Endothelial Growth Factor Receptor-1
-	D12.776.543.750.750.400.910.200 Vascular Endothelial Growth Factor Receptor-2
-	D12.776.543.750.750.400.910.300 Vascular Endothelial Growth Factor Receptor-3
-	D12.776.543.750.750.460 Receptors, Invertebrate Peptide
-	D12.776.543.750.750.555 Receptors, Neuropeptide
-	D12.776.543.750.750.555.074 Orexin Receptors
-	D12.776.543.750.750.555.200 Receptors, Bombesin
-	D12.776.543.750.750.555.220 Receptors, Bradykinin
-	D12.776.543.750.750.555.220.249 Receptor, Bradykinin B1
-	D12.776.543.750.750.555.220.500 Receptor, Bradykinin B2
-	D12.776.543.750.750.555.260 Receptors, Calcitonin Gene-Related Peptide
-	D12.776.543.750.750.555.270 Receptors, Cholecystokinin
-	D12.776.543.750.750.555.270.100 Receptor, Cholecystokinin A
-	D12.776.543.750.750.555.270.200 Receptor, Cholecystokinin B
-	D12.776.543.750.750.555.285 Receptors, Corticotropin
-	D12.776.543.750.750.555.285.500 Receptors, Melanocortin
-	D12.776.543.750.750.555.285.500.500 Receptor, Melanocortin, Type 1
-	D12.776.543.750.750.555.285.500.750 Receptor, Melanocortin, Type 2

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.543.750.750.555.285.500.875 Receptor, Melanocortin, Type 3
-	D12.776.543.750.750.555.285.500.937 Receptor, Melanocortin, Type 4
-	D12.776.543.750.750.555.290 Receptors, Corticotropin-Releasing Hormone
-	D12.776.543.750.750.555.370 Receptors, FSH
-	D12.776.543.750.750.555.410 Receptors, Galanin
-	D12.776.543.750.750.555.450 Receptors, LH
-	D12.776.543.750.750.555.460 Receptors, LHRH
-	D12.776.543.750.750.555.540 Receptors, Neuropeptide Y
-	D12.776.543.750.750.555.560 Receptors, Neurotensin
-	D12.776.543.750.750.555.610 Receptors, Opioid
-	D12.776.543.750.750.555.610.200 Receptors, Opioid, delta
-	D12.776.543.750.750.555.610.400 Receptors, Opioid, kappa
-	D12.776.543.750.750.555.610.550 Receptors, Opioid, mu
-	D12.776.543.750.750.555.610.775 Receptors, sigma
-	D12.776.543.750.750.555.630 Receptors, Oxytocin
-	D12.776.543.750.750.555.710 Receptors, Prolactin
-	D12.776.543.750.750.555.760 Receptors, Somatostatin
-	D12.776.543.750.750.555.770 Receptors, Somatotropin
-	D12.776.543.750.750.555.830 Receptors, Tachykinin
-	D12.776.543.750.750.555.830.500 Receptors, Neurokinin-1
-	D12.776.543.750.750.555.830.540 Receptors, Neurokinin-2
-	D12.776.543.750.750.555.830.580 Receptors, Neurokinin-3
-	D12.776.543.750.750.555.850 Receptors, Thyrotropin
-	D12.776.543.750.750.555.860 Receptors, Thyrotropin-Releasing Hormone
-	D12.776.543.750.750.555.915 Receptors, Vasoactive Intestinal Peptide
-	D12.776.543.750.750.555.925 Receptors, Vasopressin
-	D12.776.543.750.750.580 Receptors, Pancreatic Hormone
-	D12.776.543.750.750.580.300 Receptor, Insulin
-	D12.776.543.750.750.580.350 Receptors, Glucagon
-	D12.776.543.750.750.580.720 Receptors, Somatostatin
-	D12.776.543.750.750.600 Receptors, Parathyroid Hormone
-	D12.776.543.750.750.600.100 Receptor, Parathyroid Hormone, Type 1
-	D12.776.543.750.750.600.200 Receptor, Parathyroid Hormone, Type 2
-	D12.776.543.750.750.660 Receptors, Pituitary Hormone
-	D12.776.543.750.750.660.285 Receptors, Corticotropin
-	D12.776.543.750.750.660.285.500 Receptors, Melanocortin

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D12.776.543.750.750.660.285.500.500	Receptor, Melanocortin, Type 1
-	D12.776.543.750.750.660.285.500.750	Receptor, Melanocortin, Type 2
-	D12.776.543.750.750.660.285.500.875	Receptor, Melanocortin, Type 3
-	D12.776.543.750.750.660.285.500.937	Receptor, Melanocortin, Type 4
-	D12.776.543.750.750.660.350	Receptors, Gonadotropin
-	D12.776.543.750.750.660.350.300	Receptors, FSH
-	D12.776.543.750.750.660.350.450	Receptors, LH
-	D12.776.543.750.750.660.350.725	Receptors, Prolactin
-	D12.776.543.750.750.660.600	Receptors, Oxytocin
-	D12.776.543.750.750.660.750	Receptors, Somatotropin
-	D12.776.543.750.750.660.825	Receptors, Thyrotropin
-	D12.776.543.750.750.660.900	Receptors, Vasopressin
-	D12.776.543.750.750.700	Receptors, Pituitary Hormone-Regulating Hormone
-	D12.776.543.750.750.700.150	Receptors, Corticotropin-Releasing Hormone
-	D12.776.543.750.750.700.460	Receptors, LHRH
-	D12.776.543.750.750.700.800	Receptors, Somatostatin
-	D12.776.543.750.750.700.840	Receptors, Thyrotropin-Releasing Hormone
-	D12.776.543.750.750.850	Receptors, Thrombin
-	D12.776.543.750.750.850.399	Receptor, PAR-1
-	D12.776.543.750.750.850.800	Thrombomodulin
-	D12.776.543.750.783	Receptors, Phospholipase A2
-	D12.776.543.750.792	Receptors, Proteinase-Activated
-	D12.776.543.750.792.249	Receptor, PAR-2
-	D12.776.543.750.792.500	Receptors, Thrombin
-	D12.776.543.750.792.500.500	Receptor, PAR-1
New Tree	<a href="#">D12.776.543.750.800</a>	<a href="#">Receptors, Transferrin</a>
New Tree	<a href="#">D12.776.543.750.800.249</a>	<a href="#">Bacterial Transferrin Receptor Complex</a>
New Tree	<a href="#">D12.776.543.750.800.249.500</a>	<a href="#">Transferrin-Binding Protein A</a>
New Tree	<a href="#">D12.776.543.750.800.249.750</a>	<a href="#">Transferrin-Binding Protein B</a>
New Tree	<a href="#">D12.776.543.750.815</a>	<a href="#">Receptors, Urokinase Plasminogen Activator</a>
New Tree	<a href="#">D12.776.543.750.830</a>	<a href="#">Receptors, Virus</a>
New	<a href="#">D12.776.543.750.830.124</a>	<a href="#">Coxsackie and Adenovirus Receptor-Like</a>

## MeSH Tree Changes for 2017

Type	Tree - heading	
Tree	<a href="#">Membrane Protein</a>	
New Heading	<b>D12.776.543.750.830.187</b>	<b>Hepatitis A Virus Cellular Receptor 1</b>
New Heading	<b>D12.776.543.750.830.219</b>	<b>Hepatitis A Virus Cellular Receptor 2</b>
New Tree	<a href="#">D12.776.543.750.830.250</a>	<a href="#">Receptors, Complement 3d</a>
New Tree	<a href="#">D12.776.543.750.830.700</a>	<a href="#">Receptors, HIV</a>
New Tree	<a href="#">D12.776.543.750.830.700.025</a>	<a href="#">Antigens, CD4</a>
New Tree	<a href="#">D12.776.543.750.830.700.605</a>	<a href="#">Receptors, CCR5</a>
New Tree	<a href="#">D12.776.543.750.830.700.650</a>	<a href="#">Receptors, CXCR4</a>
New Tree	<a href="#">D12.776.543.750.830.850</a>	<a href="#">Receptors, Tumor Necrosis Factor, Member 14</a>
Old Tree	<b>D12.776.543.750.850</b>	<b>Receptors, Transferrin</b>
New Tree	<a href="#">D12.776.543.750.850</a>	<a href="#">Receptors, Wnt</a>
Old Tree	<b>D12.776.543.750.850.249</b>	<b>Bacterial Transferrin Receptor Complex</b>
Old Tree	<b>D12.776.543.750.850.249.500</b>	<b>Transferrin-Binding Protein A</b>
Old Tree	<b>D12.776.543.750.850.249.750</b>	<b>Transferrin-Binding Protein B</b>
New Tree	<a href="#">D12.776.543.750.850.500</a>	<a href="#">Frizzled Receptors</a>
New Heading	<b>D12.776.543.750.850.500.500</b>	<b>Smoothened Receptor</b>
New Tree	<a href="#">D12.776.543.750.850.687</a>	<a href="#">Low Density Lipoprotein Receptor-Related Protein-5</a>
New Tree	<a href="#">D12.776.543.750.850.781</a>	<a href="#">Low Density Lipoprotein Receptor-Related Protein-6</a>
Old Tree	<b>D12.776.543.750.887</b>	<b>Receptors, Urokinase Plasminogen Activator</b>
Old Tree	<b>D12.776.543.750.925</b>	<b>Receptors, Virus</b>
Old Tree	<b>D12.776.543.750.925.124</b> <b>Membrane Protein</b>	<b>Coxsackie and Adenovirus Receptor-Like</b>
Old Tree	<b>D12.776.543.750.925.250</b>	<b>Receptors, Complement 3d</b>
Old Tree	<b>D12.776.543.750.925.700</b>	<b>Receptors, HIV</b>
Old Tree	<b>D12.776.543.750.925.700.025</b>	<b>Antigens, CD4</b>
Old Tree	<b>D12.776.543.750.925.700.605</b>	<b>Receptors, CCR5</b>
Old Tree	<b>D12.776.543.750.925.700.650</b>	<b>Receptors, CXCR4</b>
Old Tree	<b>D12.776.543.750.925.850</b>	<b>Receptors, Tumor Necrosis Factor, Member 14</b>



## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	<a href="#">D12.776.543.750.962</a> Receptors, Wnt
Old Tree	<a href="#">D12.776.543.750.962.500</a> Frizzled Receptors
Old Tree	<a href="#">D12.776.543.750.962.687</a> Low Density Lipoprotein Receptor-Related Protein-5
Old Tree	<a href="#">D12.776.543.750.962.781</a> Low Density Lipoprotein Receptor-Related Protein-6
New Heading	<b>D12.776.543.750.981</b> Zona Pellucida Glycoproteins
New Heading	<b>D12.776.543.800</b> Serrate-Jagged Proteins
New Heading	<b>D12.776.543.800.500</b> Jagged-1 Protein
New Heading	<b>D12.776.543.800.750</b> Jagged-2 Protein
New Tree	<a href="#">D12.776.543.850</a> Spectrin
New Heading	<b>D12.776.543.875</b> Stromal Interaction Molecules
New Heading	<b>D12.776.543.875.500</b> Stromal Interaction Molecule 1
New Heading	<b>D12.776.543.875.750</b> Stromal Interaction Molecule 2
New Tree	<a href="#">D12.776.543.900</a> Tetraspanins
New Tree	<a href="#">D12.776.543.900.109</a> Antigens, CD9
New Tree	<a href="#">D12.776.543.900.153</a> Antigens, CD53
New Tree	<a href="#">D12.776.543.900.163</a> Antigens, CD63
New Tree	<a href="#">D12.776.543.900.181</a> Antigens, CD81
New Tree	<a href="#">D12.776.543.900.182</a> Antigens, CD82
New Tree	<a href="#">D12.776.543.900.251</a> Antigens, CD151
New Tree	<a href="#">D12.776.543.900.750</a> Uroplakin Ia
New Tree	<a href="#">D12.776.543.900.875</a> Uroplakin Ib
New Tree	<a href="#">D12.776.543.930</a> Thylakoid Membrane Proteins
New Tree	<a href="#">D12.776.543.930.249</a> Chloroplast Proton-Translocating ATPases
New	<a href="#">D12.776.543.930.500</a> Photosynthetic Reaction Center Complex Proteins

## MeSH Tree Changes for 2017

Type	Tree - heading
Tree	
New Tree	D12.776.543.930.500.374                      Cytochrome b6f Complex
New Tree	D12.776.543.930.500.374.500                      Cytochromes b6
New Tree	D12.776.543.930.500.374.750                      Cytochromes f
New Tree	D12.776.543.930.500.374.875                      Plastoquinol-Plastocyanin Reductase
New Tree	D12.776.543.930.500.490                      Light-Harvesting Protein Complexes
New Tree	D12.776.543.930.500.490.249                      Chlorophyll Binding Proteins
New Tree	D12.776.543.930.500.500                      Photosystem I Protein Complex
New Tree	D12.776.543.930.500.750                      Photosystem II Protein Complex
New Tree	D12.776.543.940                      Tight Junction Proteins
New Tree	D12.776.543.940.200                      Claudins
New Tree	D12.776.543.940.200.100                      Claudin-1
New Tree	D12.776.543.940.200.200                      Claudin-2
New Tree	D12.776.543.940.200.300                      Claudin-3
New Tree	D12.776.543.940.200.400                      Claudin-4
New Tree	D12.776.543.940.200.500                      Claudin-5
New Tree	D12.776.543.940.600                      Junctional Adhesion Molecules
New Tree	D12.776.543.940.600.500                      Coxsackie and Adenovirus Receptor-Like Membrane Protein
New Tree	D12.776.543.940.600.700                      Junctional Adhesion Molecule A
New Tree	D12.776.543.940.600.800                      Junctional Adhesion Molecule B
New Tree	D12.776.543.940.600.900                      Junctional Adhesion Molecule C
New Tree	D12.776.543.940.675                      MARVEL Domain Containing 2 Protein

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	D12.776.543.940.750 Occludin
New Tree	D12.776.543.940.900 Zonula Occludens Proteins
New Tree	D12.776.543.940.900.500 Zonula Occludens-1 Protein
New Tree	D12.776.543.940.900.750 Zonula Occludens-2 Protein
Old Tree	D12.776.543.980 Spectrin
Old Tree	D12.776.543.982 Tetraspanins
Old Tree	D12.776.543.982.109 Antigens, CD9
Old Tree	D12.776.543.982.153 Antigens, CD53
Old Tree	D12.776.543.982.163 Antigens, CD63
Old Tree	D12.776.543.982.181 Antigens, CD81
Old Tree	D12.776.543.982.182 Antigens, CD82
Old Tree	D12.776.543.982.251 Antigens, CD151
Old Tree	D12.776.543.982.750 Uroplakin Ia
Old Tree	D12.776.543.982.875 Uroplakin Ib
Old Tree	D12.776.543.983 Thylakoid Membrane Proteins
Old Tree	D12.776.543.983.249 Chloroplast Proton-Translocating ATPases
Old Tree	D12.776.543.983.500 Photosynthetic Reaction Center Complex Proteins
Old Tree	D12.776.543.983.500.374 Cytochrome b6f Complex
Old Tree	D12.776.543.983.500.374.500 Cytochromes b6
Old Tree	D12.776.543.983.500.374.750 Cytochromes f
Old Tree	D12.776.543.983.500.374.875 Plastoquinol-Plastocyanin Reductase
Old Tree	D12.776.543.983.500.490 Light-Harvesting Protein Complexes
Old Tree	D12.776.543.983.500.490.249 Chlorophyll Binding Proteins
Old Tree	D12.776.543.983.500.500 Photosystem I Protein Complex
Old Tree	D12.776.543.983.500.750 Photosystem II Protein Complex
Old Tree	D12.776.543.984 Tight Junction Proteins
Old Tree	D12.776.543.984.200 Claudins
Old Tree	D12.776.543.984.200.100 Claudin-1
Old Tree	D12.776.543.984.200.200 Claudin-2
Old Tree	D12.776.543.984.200.300 Claudin-3
Old Tree	D12.776.543.984.200.400 Claudin-4
Old Tree	D12.776.543.984.200.500 Claudin-5
Old Tree	D12.776.543.984.600 Junctional Adhesion Molecules

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	D12.776.543.984.600.500 Coxsackie and Adenovirus Receptor-Like Membrane Protein
Old Tree	D12.776.543.984.600.700 Junctional Adhesion Molecule A
Old Tree	D12.776.543.984.600.800 Junctional Adhesion Molecule B
Old Tree	D12.776.543.984.600.900 Junctional Adhesion Molecule C
Old Tree	D12.776.543.984.675 MARVEL Domain Containing 2 Protein
Old Tree	D12.776.543.984.750 Occludin
Old Tree	D12.776.543.984.900 Zonula Occludens Proteins
Old Tree	D12.776.543.984.900.500 Zonula Occludens-1 Protein
Old Tree	D12.776.543.984.900.750 Zonula Occludens-2 Protein
-	D12.776.543.985 Utrophin
-	D12.776.543.990 Vesicular Transport Proteins
-	D12.776.543.990.024 Antigens, CD63
-	D12.776.543.990.049 Auxilins
-	D12.776.543.990.100 Caveolins
-	D12.776.543.990.100.500 Caveolin 1
-	D12.776.543.990.100.750 Caveolin 2
-	D12.776.543.990.100.875 Caveolin 3
-	D12.776.543.990.150 Adaptor Proteins, Vesicular Transport
-	D12.776.543.990.150.100 Adaptor Protein Complex 1
-	D12.776.543.990.150.200 Adaptor Protein Complex 2
-	D12.776.543.990.150.300 Adaptor Protein Complex 3
-	D12.776.543.990.150.400 Adaptor Protein Complex 4
-	D12.776.543.990.150.500 Adaptor Protein Complex Subunits
-	D12.776.543.990.150.500.100 Adaptor Protein Complex alpha Subunits
-	D12.776.543.990.150.500.200 Adaptor Protein Complex beta Subunits
-	D12.776.543.990.150.500.300 Adaptor Protein Complex delta Subunits
-	D12.776.543.990.150.500.400 Adaptor Protein Complex gamma Subunits
-	D12.776.543.990.150.500.500 Adaptor Protein Complex mu Subunits
-	D12.776.543.990.150.500.750 Adaptor Protein Complex sigma Subunits
-	D12.776.543.990.150.750 Monomeric Clathrin Assembly Proteins
-	D12.776.543.990.150.875 Sorting Nexins
-	D12.776.543.990.175 Class III Phosphatidylinositol 3-Kinases
-	D12.776.543.990.175.500 Vacuolar Sorting Protein VPS15
-	D12.776.543.990.200 Clathrin
-	D12.776.543.990.200.500 Clathrin Heavy Chains

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.543.990.200.750 Clathrin Light Chains
-	D12.776.543.990.300 Coat Protein Complex I
-	D12.776.543.990.300.150 ADP-Ribosylation Factor 1
-	D12.776.543.990.300.300 Coatomer Protein
-	D12.776.543.990.400 Dynamins
-	D12.776.543.990.400.100 Dynamin I
-	D12.776.543.990.400.200 Dynamin II
-	D12.776.543.990.400.300 Dynamin III
-	D12.776.543.990.493 Endosomal Sorting Complexes Required for Transport
-	D12.776.543.990.587 Munc18 Proteins
-	D12.776.543.990.681 Myelin and Lymphocyte-Associated Proteolipid Proteins
-	D12.776.543.990.775 SNARE Proteins
-	D12.776.543.990.775.500 Q-SNARE Proteins
-	D12.776.543.990.775.500.500 Qa-SNARE Proteins
-	D12.776.543.990.775.500.500.700 Syntaxin 1
-	D12.776.543.990.775.500.500.750 Syntaxin 16
-	D12.776.543.990.775.500.750 Qb-SNARE Proteins
-	D12.776.543.990.775.500.750.500 Synaptosomal-Associated Protein 25
-	D12.776.543.990.775.500.875 Qc-SNARE Proteins
-	D12.776.543.990.775.500.875.500 Synaptosomal-Associated Protein 25
-	D12.776.543.990.775.750 R-SNARE Proteins
-	D12.776.543.990.775.750.500 Vesicle-Associated Membrane Protein 1
-	D12.776.543.990.775.750.750 Vesicle-Associated Membrane Protein 2
-	D12.776.543.990.775.750.875 Vesicle-Associated Membrane Protein 3
-	D12.776.543.990.812 Soluble N-Ethylmaleimide-Sensitive Factor Attachment Proteins
-	D12.776.543.990.831 Synaptogyrins
-	D12.776.543.990.840 Synaptophysin
-	D12.776.543.990.850 Synaptotagmins
-	D12.776.543.990.850.249 Synaptotagmin I
-	D12.776.543.990.850.500 Synaptotagmin II
-	D12.776.556 Metalloproteins
-	D12.776.556.080 Azurin
-	D12.776.556.151 Ceruloplasmin
-	D12.776.556.462 Hemocyanin
-	D12.776.556.526 Hemosiderin

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.556.579 Iron-Binding Proteins
-	D12.776.556.579.249 Ferritins
-	D12.776.556.579.249.290 Apoferritins
-	D12.776.556.579.374 Nonheme Iron Proteins
-	D12.776.556.579.374.187 Hemerythrin
-	D12.776.556.579.374.281 Inositol Oxygenase
-	D12.776.556.579.374.375 Iron-Sulfur Proteins
-	D12.776.556.579.374.375.025 Adrenodoxin
-	D12.776.556.579.374.375.140 Electron Transport Complex I
-	D12.776.556.579.374.375.140.500 NADH Dehydrogenase
-	D12.776.556.579.374.375.141 Electron Transport Complex II
-	D12.776.556.579.374.375.141.500 Succinate Dehydrogenase
-	D12.776.556.579.374.375.142 Electron Transport Complex III
-	D12.776.556.579.374.375.150 Ferredoxin-Nitrite Reductase
-	D12.776.556.579.374.375.275 Ferredoxins
-	D12.776.556.579.374.375.275.450 Molybdoferredoxin
-	D12.776.556.579.374.375.275.725 Rubredoxins
-	D12.776.556.579.374.375.617 Ferrochelatase
-	D12.776.556.579.374.375.960 Iron Regulatory Protein 1
-	D12.776.556.579.374.375.962 Iron Regulatory Protein 2
-	D12.776.556.579.374.375.977 Nitrate Reductase (NAD(P)H)
-	D12.776.556.579.374.375.988 Nitrate Reductase (NADPH)
-	D12.776.556.579.374.568 Lipoxygenases
-	D12.776.556.579.374.568.500 Arachidonate Lipoxygenases
-	D12.776.556.579.374.568.500.020 Arachidonate 5-Lipoxygenase
-	D12.776.556.579.374.568.500.025 Arachidonate 12-Lipoxygenase
-	D12.776.556.579.374.568.500.030 Arachidonate 15-Lipoxygenase
-	D12.776.556.579.374.568.750 Lipoxygenase
-	D12.776.556.579.374.687 Retinal Dehydrogenase
-	D12.776.556.579.374.925 Tyrosine 3-Monooxygenase
-	D12.776.556.579.750 Transferrins
-	D12.776.556.579.750.124 Conalbumin
-	D12.776.556.579.750.249 Lactoferrin
-	D12.776.556.579.750.500 Transferrin
-	D12.776.556.670 Metallothionein
-	D12.776.556.760 Plastocyanin

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D12.776.575	Mitochondrial Proteins
New Heading	<b>D12.776.575.093</b>	<b>Aldehyde Dehydrogenase, Mitochondrial</b>
-	D12.776.575.186	Creatine Kinase, Mitochondrial Form
-	D12.776.575.374	Endopeptidase Clp
-	D12.776.575.562	Ferrochelatase
-	D12.776.575.750	Mitochondrial Membrane Transport Proteins
-	D12.776.575.750.500	Mitochondrial ADP, ATP Translocases
-	D12.776.575.750.500.100	Adenine Nucleotide Translocator 1
-	D12.776.575.750.500.200	Adenine Nucleotide Translocator 2
-	D12.776.575.750.500.300	Adenine Nucleotide Translocator 3
-	D12.776.575.750.625	Mitochondrial Proton-Translocating ATPases
New Heading	<b>D12.776.575.750.688</b>	<b>Mitochondrial Uncoupling Proteins</b>
New Heading	<b>D12.776.575.750.688.500</b>	<b>Uncoupling Protein 1</b>
New Heading	<b>D12.776.575.750.688.750</b>	<b>Uncoupling Protein 2</b>
New Heading	<b>D12.776.575.750.688.875</b>	<b>Uncoupling Protein 3</b>
-	D12.776.575.750.750	Voltage-Dependent Anion Channel 1
-	D12.776.575.750.875	Voltage-Dependent Anion Channel 2
-	D12.776.575.765	Mitochondrial Trifunctional Protein
-	D12.776.575.765.200	Mitochondrial Trifunctional Protein, alpha Subunit
-	D12.776.575.765.500	Mitochondrial Trifunctional Protein, beta Subunit
-	D12.776.575.781	Nucleoside Diphosphate Kinase D
-	D12.776.575.812	Peroxiredoxin III
-	D12.776.575.875	Sirtuin 3
-	D12.776.580	Molecular Chaperones
-	D12.776.580.157	alpha-Crystallins
-	D12.776.580.157.149	alpha-Crystallin A Chain
-	D12.776.580.157.300	alpha-Crystallin B Chain
-	D12.776.580.215	Clusterin
-	D12.776.580.216	Heat-Shock Proteins
-	D12.776.580.216.210	Chaperonins
-	D12.776.580.216.210.590	Group I Chaperonins
-	D12.776.580.216.210.590.500	Chaperonin 10

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.580.216.210.590.750 Chaperonin 60
-	D12.776.580.216.210.795 Group II Chaperonins
-	D12.776.580.216.210.795.500 Chaperonin Containing TCP-1
-	D12.776.580.216.210.795.750 Thermosomes
-	D12.776.580.216.270 Heat-Shock Proteins, Small
-	D12.776.580.216.270.500 HSP20 Heat-Shock Proteins
-	D12.776.580.216.270.625 HSP27 Heat-Shock Proteins
-	D12.776.580.216.270.750 HSP30 Heat-Shock Proteins
-	D12.776.580.216.292 HSP40 Heat-Shock Proteins
-	D12.776.580.216.295 HSP47 Heat-Shock Proteins
-	D12.776.580.216.375 HSP70 Heat-Shock Proteins
-	D12.776.580.216.375.200 HSC70 Heat-Shock Proteins
-	D12.776.580.216.375.202 HSP72 Heat-Shock Proteins
-	D12.776.580.216.375.800 HSP110 Heat-Shock Proteins
-	D12.776.580.216.380 HSP90 Heat-Shock Proteins
-	D12.776.580.219 Histone Chaperones
-	D12.776.580.219.200 Chromatin Assembly Factor-1
-	D12.776.580.219.200.500 Retinoblastoma-Binding Protein 4
-	D12.776.580.219.500 Nucleoplasmins
-	D12.776.580.219.550 Nucleosome Assembly Protein 1
-	D12.776.580.612 Metallochaperones
-	D12.776.580.845 Neuroendocrine Secretory Protein 7B2
-	D12.776.580.922 SMN Complex Proteins
-	D12.776.580.922.249 DEAD Box Protein 20
-	D12.776.580.922.500 Survival of Motor Neuron 1 Protein
-	D12.776.580.922.750 Survival of Motor Neuron 2 Protein
-	D12.776.602 Mutant Proteins
-	D12.776.602.500 Mutant Chimeric Proteins
-	D12.776.602.500.500 Oncogene Proteins, Fusion
-	D12.776.602.500.500.100 Fusion Proteins, bcr-abl
-	D12.776.602.500.500.320 Fusion Proteins, gag-onc
-	D12.776.602.500.500.320.700 Oncogene Protein p65(gag-jun)
-	D12.776.602.500.500.660 Oncogene Protein tpr-met
-	D12.776.624 Neoplasm Proteins
-	D12.776.624.301 Melanoma-Specific Antigens
-	D12.776.624.301.249 Antigens, CD146



## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.624.301.500 gp100 Melanoma Antigen
-	D12.776.624.301.750 MART-1 Antigen
-	D12.776.624.553 Myeloma Proteins
-	D12.776.624.664 Oncogene Proteins
-	D12.776.624.664.500 Oncogene Proteins, Fusion
-	D12.776.624.664.500.100 Fusion Proteins, bcr-abl
-	D12.776.624.664.500.320 Fusion Proteins, gag-onc
-	D12.776.624.664.500.320.700 Oncogene Protein p65(gag-jun)
-	D12.776.624.664.500.660 Oncogene Protein tpr-met
-	D12.776.624.664.520 Oncogene Proteins, Viral
-	D12.776.624.664.520.045 Adenovirus Early Proteins
-	D12.776.624.664.520.045.050 Adenovirus E1 Proteins
-	D12.776.624.664.520.045.050.100 Adenovirus E1A Proteins
-	D12.776.624.664.520.045.050.110 Adenovirus E1B Proteins
-	D12.776.624.664.520.045.060 Adenovirus E2 Proteins
-	D12.776.624.664.520.045.070 Adenovirus E3 Proteins
-	D12.776.624.664.520.045.080 Adenovirus E4 Proteins
-	D12.776.624.664.520.090 Antigens, Polyomavirus Transforming
-	D12.776.624.664.520.420 Papillomavirus E7 Proteins
-	D12.776.624.664.520.750 Retroviridae Proteins, Oncogenic
-	D12.776.624.664.520.750.320 Fusion Proteins, gag-onc
-	D12.776.624.664.520.750.320.700 Oncogene Protein p65(gag-jun)
-	D12.776.624.664.520.750.470 Gene Products, rex
-	D12.776.624.664.520.750.480 Gene Products, tax
-	D12.776.624.664.520.750.650 Oncogene Protein gp140(v-fms)
-	D12.776.624.664.520.750.710 Oncogene Protein p21(ras)
-	D12.776.624.664.520.750.750 Oncogene Protein p55(v-myc)
-	D12.776.624.664.520.750.760 Oncogene Protein pp60(v-src)
-	D12.776.624.664.520.750.788 Oncogene Protein v-akt
-	D12.776.624.664.520.750.817 Oncogene Protein v-cbl
-	D12.776.624.664.520.750.846 Oncogene Protein v-crk
-	D12.776.624.664.520.750.860 Oncogene Protein v-maf
-	D12.776.624.664.520.750.875 Oncogene Proteins v-abl
-	D12.776.624.664.520.750.882 Oncogene Proteins v-erbA
-	D12.776.624.664.520.750.883 Oncogene Proteins v-erbB
-	D12.776.624.664.520.750.887 Oncogene Proteins v-fos

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D12.776.624.664.520.750.900	Oncogene Proteins v-mos
-	D12.776.624.664.520.750.903	Oncogene Proteins v-myb
-	D12.776.624.664.520.750.920	Oncogene Proteins v-raf
-	D12.776.624.664.520.750.925	Oncogene Proteins v-rel
-	D12.776.624.664.520.750.935	Oncogene Proteins v-sis
-	D12.776.624.664.700	Proto-Oncogene Proteins
New Heading	<b>D12.776.624.664.700.025</b>	<b>Bcl-2-Like Protein 11</b>
-	D12.776.624.664.700.049	Chemokine CXCL1
-	D12.776.624.664.700.100	Cyclin D1
-	D12.776.624.664.700.110	Fibroblast Growth Factor 4
-	D12.776.624.664.700.112	Fibroblast Growth Factor 6
-	D12.776.624.664.700.114	fms-Like Tyrosine Kinase 3
-	D12.776.624.664.700.117	Janus Kinase 2
New Heading	<b>D12.776.624.664.700.123</b>	<b>Lipocalin-2</b>
-	D12.776.624.664.700.128 p56(lck)	Lymphocyte Specific Protein Tyrosine Kinase
-	D12.776.624.664.700.130	MutS Homolog 2 Protein
-	D12.776.624.664.700.148	Myeloid-Lymphoid Leukemia Protein
New Heading	<b>D12.776.624.664.700.158</b>	<b>N-Myc Proto-Oncogene Protein</b>
-	D12.776.624.664.700.167	Proto-Oncogene Proteins c-abl
-	D12.776.624.664.700.168	Proto-Oncogene Proteins c-akt
-	D12.776.624.664.700.169	Proto-Oncogene Proteins c-bcl-2
-	D12.776.624.664.700.169.500	Myeloid Cell Leukemia Sequence 1 Protein
-	D12.776.624.664.700.170	Proto-Oncogene Proteins c-bcl-6
-	D12.776.624.664.700.171	Proto-Oncogene Proteins c-bcr
-	D12.776.624.664.700.171.500	Fusion Proteins, bcr-abl
-	D12.776.624.664.700.172	Proto-Oncogene Proteins c-cbl
-	D12.776.624.664.700.174	Proto-Oncogene Proteins c-crk
-	D12.776.624.664.700.175	Proto-Oncogene Proteins c-ets
-	D12.776.624.664.700.175.100	Proto-Oncogene Protein c-ets-1
-	D12.776.624.664.700.175.200	Proto-Oncogene Protein c-ets-2
-	D12.776.624.664.700.175.400	Proto-Oncogene Protein c-fli-1
-	D12.776.624.664.700.175.600	Ternary Complex Factors
-	D12.776.624.664.700.175.600.100	ets-Domain Protein Elk-1



## MeSH Tree Changes for 2017

Type	Tree - heading	
Heading		
-	D12.776.624.664.700.967	Wnt1 Protein
-	D12.776.624.664.700.978	Wnt2 Protein
New Heading	<b>D12.776.624.664.700.989</b>	<b>Zinc Finger Protein GLI1</b>
-	D12.776.624.776	Tumor Suppressor Proteins
-	D12.776.624.776.049	Adenomatous Polyposis Coli Protein
-	D12.776.624.776.052	Antigens, CD82
-	D12.776.624.776.100	BRCA1 Protein
-	D12.776.624.776.101	BRCA2 Protein
-	D12.776.624.776.355	Cyclin-Dependent Kinase Inhibitor Proteins
-	D12.776.624.776.355.100	Cyclin-Dependent Kinase Inhibitor p15
-	D12.776.624.776.355.200	Cyclin-Dependent Kinase Inhibitor p16
-	D12.776.624.776.355.300	Cyclin-Dependent Kinase Inhibitor p18
-	D12.776.624.776.355.400	Cyclin-Dependent Kinase Inhibitor p19
-	D12.776.624.776.355.500	Cyclin-Dependent Kinase Inhibitor p21
-	D12.776.624.776.355.600	Cyclin-Dependent Kinase Inhibitor p27
-	D12.776.624.776.355.700	Cyclin-Dependent Kinase Inhibitor p57
-	D12.776.624.776.418	Cystatin M
-	D12.776.624.776.546	Kisspeptins
-	D12.776.624.776.610	Neurofibromin 1
-	D12.776.624.776.612	Neurofibromin 2
New Heading	<b>D12.776.624.776.633</b>	<b>Patched Receptors</b>
New Heading	<b>D12.776.624.776.633.500</b>	<b>Patched-1 Receptor</b>
New Heading	<b>D12.776.624.776.633.750</b>	<b>Patched-2 Receptor</b>
New Heading	<b>D12.776.624.776.654</b>	<b>Promyelocytic Leukemia Protein</b>
-	D12.776.624.776.695	PTEN Phosphohydrolase
-	D12.776.624.776.717	Retinoblastoma-Like Protein p107
-	D12.776.624.776.735	Retinoblastoma-Like Protein p130
-	D12.776.624.776.745	Retinoblastoma Protein
-	D12.776.624.776.760	Smad4 Protein
-	D12.776.624.776.772	Tumor Suppressor Protein p14ARF
-	D12.776.624.776.775	Tumor Suppressor Protein p53

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Heading	<b>D12.776.624.776.820</b>	<b>Tumor Protein p73</b>
-	D12.776.624.776.865	Von Hippel-Lindau Tumor Suppressor Protein
-	D12.776.624.776.960	WT1 Proteins
New Tree	<a href="#">D12.776.631</a>	<a href="#">Nerve Tissue Proteins</a>
New Tree	<a href="#">D12.776.631.024</a>	<a href="#">Acid Sensing Ion Channels</a>
New Tree	<a href="#">D12.776.631.050</a>	<a href="#">Agrin</a>
New Tree	<a href="#">D12.776.631.069</a>	<a href="#">Ataxins</a>
New Tree	<a href="#">D12.776.631.069.500</a>	<a href="#">Ataxin-1</a>
New Tree	<a href="#">D12.776.631.069.750</a>	<a href="#">Ataxin-2</a>
New Tree	<a href="#">D12.776.631.069.875</a>	<a href="#">Ataxin-3</a>
New Tree	<a href="#">D12.776.631.069.901</a>	<a href="#">Ataxin-7</a>
New Tree	<a href="#">D12.776.631.069.950</a>	<a href="#">Ataxin-10</a>
New Tree	<a href="#">D12.776.631.087</a>	<a href="#">Calbindin 1</a>
New Tree	<a href="#">D12.776.631.124</a>	<a href="#">Chimerin Proteins</a>
New Tree	<a href="#">D12.776.631.124.200</a>	<a href="#">Chimerin 1</a>
New Tree	<a href="#">D12.776.631.199</a>	<a href="#">Chromogranins</a>
New Tree	<a href="#">D12.776.631.199.249</a>	<a href="#">Chromogranin A</a>
New Tree	<a href="#">D12.776.631.199.500</a>	<a href="#">Chromogranin B</a>
New Tree	<a href="#">D12.776.631.199.750</a>	<a href="#">Secretogranin II</a>
New Tree	<a href="#">D12.776.631.249</a>	<a href="#">Dopamine and cAMP-Regulated Phosphoprotein 32</a>
New Heading	<b>D12.776.631.274</b>	<b>Fatty Acid-Binding Protein 7</b>
New Tree	<a href="#">D12.776.631.299</a>	<a href="#">Fragile X Mental Retardation Protein</a>
New	<a href="#">D12.776.631.400</a>	<a href="#">GAP-43 Protein</a>

## MeSH Tree Changes for 2017

Type	Tree - heading
Tree	
New Tree	D12.776.631.480      Glucose Transporter Type 3
New Tree	D12.776.631.520      ELAV Proteins
New Tree	D12.776.631.520.500      ELAV-Like Protein 2
New Tree	D12.776.631.520.750      ELAV-Like Protein 3
New Tree	D12.776.631.520.875      ELAV-Like Protein 4
New Tree	D12.776.631.560      Microtubule-Associated Proteins
New Heading	<b>D12.776.631.560.225      Dynactin Complex</b>
New Tree	D12.776.631.560.450      Kinesin
New Tree	D12.776.631.560.510      tau Proteins
New Tree	D12.776.631.580      Myelin Proteins
New Tree	D12.776.631.580.124      2',3'-Cyclic Nucleotide 3'-Phosphodiesterase
New Tree	D12.776.631.580.249      Myelin and Lymphocyte-Associated Proteolipid Proteins
New Tree	D12.776.631.580.500      Myelin-Associated Glycoprotein
New Tree	D12.776.631.580.510      Myelin Basic Protein
New Tree	D12.776.631.580.530      Myelin-Oligodendrocyte Glycoprotein
New Tree	D12.776.631.580.550      Myelin P0 Protein
New Tree	D12.776.631.580.560      Myelin P2 Protein
New Tree	D12.776.631.580.580      Myelin Proteolipid Protein
New Heading	<b>D12.776.631.580.738      Nogo Proteins</b>
New Tree	D12.776.631.580.895      Oligodendrocyte-Myelin Glycoprotein
New Tree	D12.776.631.590      Natriuretic Peptide, Brain

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	D12.776.631.600 Nerve Growth Factors
New Tree	D12.776.631.600.100 Brain-Derived Neurotrophic Factor
New Tree	D12.776.631.600.212 Ciliary Neurotrophic Factor
New Tree	D12.776.631.600.325 Glia Maturation Factor
New Tree	D12.776.631.600.381 Glial Cell Line-Derived Neurotrophic Factors
New Tree	D12.776.631.600.381.500 Glial Cell Line-Derived Neurotrophic Factor
New Tree	D12.776.631.600.381.750 Neurturin
New Tree	D12.776.631.600.437 Nerve Growth Factor
New Tree	D12.776.631.600.550 Neuregulins
New Tree	D12.776.631.600.550.750 Neuregulin-1
New Tree	D12.776.631.600.775 Neurotrophin 3
New Tree	D12.776.631.600.887 Pituitary Adenylate Cyclase-Activating Polypeptide
New Tree	D12.776.631.607 Nestin
New Tree	D12.776.631.615 Neuroendocrine Secretory Protein 7B2
New Tree	D12.776.631.630 Neurofilament Proteins
New Tree	D12.776.631.640 Neurogranin
New Tree	D12.776.631.642 Neuronal Apoptosis-Inhibitory Protein
New Tree	D12.776.631.645 Neuronal Calcium-Sensor Proteins
New Tree	D12.776.631.645.124 Guanylate Cyclase-Activating Proteins
New Tree	D12.776.631.645.249 Hippocalcin
New Tree	D12.776.631.645.374 Kv Channel-Interacting Proteins
New Tree	D12.776.631.645.500 Neurocalcin

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	D12.776.631.645.750 <span style="float: right;">Recoverin</span>
New Tree	D12.776.631.650 <span style="float: right;">Neuropeptides</span>
New Tree	D12.776.631.650.070 <span style="float: right;">Angiotensins</span>
New Tree	D12.776.631.650.070.075 <span style="float: right;">Angiotensin I</span>
New Tree	D12.776.631.650.070.078 <span style="float: right;">Angiotensin II</span>
New Tree	D12.776.631.650.070.080 <span style="float: right;">Angiotensin III</span>
New Tree	D12.776.631.650.085 <span style="float: right;">Bombesin</span>
New Tree	D12.776.631.650.090 <span style="float: right;">Bradykinin</span>
New Tree	D12.776.631.650.095 <span style="float: right;">Calcitonin</span>
New Tree	D12.776.631.650.097 <span style="float: right;">Calcitonin Gene-Related Peptide</span>
New Tree	D12.776.631.650.100 <span style="float: right;">Carnosine</span>
New Tree	D12.776.631.650.200 <span style="float: right;">Delta Sleep-Inducing Peptide</span>
New Tree	D12.776.631.650.235 <span style="float: right;">FMRFamide</span>
New Tree	D12.776.631.650.250 <span style="float: right;">Galanin</span>
New Tree	D12.776.631.650.300 <span style="float: right;">Gastric Inhibitory Polypeptide</span>
New Tree	D12.776.631.650.315 <span style="float: right;">Gastrin-Releasing Peptide</span>
New Tree	D12.776.631.650.320 <span style="float: right;">Gastrins</span>
New Tree	D12.776.631.650.363 <span style="float: right;">Orexins</span>
New Tree	D12.776.631.650.405 <span style="float: right;">Hypothalamic Hormones</span>
New Tree	D12.776.631.650.405.700 <span style="float: right;">Pituitary Hormone Release Inhibiting Hormones</span>
New Tree	D12.776.631.650.405.700.500 <span style="float: right;">MSH Release-Inhibiting Hormone</span>
New Tree	D12.776.631.650.405.700.750 <span style="float: right;">Prolactin Release-Inhibiting Factors</span>



## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	D12.776.631.650.405.700.875 <span style="float: right;">Somatostatin</span>
New Tree	D12.776.631.650.405.700.875.500 <span style="float: right;">Somatostatin-28</span>
New Tree	D12.776.631.650.405.740 <span style="float: right;">Pituitary Hormone-Releasing Hormones</span>
New Tree	D12.776.631.650.405.740.140 <span style="float: right;">Corticotropin-Releasing Hormone</span>
New Tree	D12.776.631.650.405.740.320 <span style="float: right;">Gonadotropin-Releasing Hormone</span>
New Tree	D12.776.631.650.405.740.320.100 <span style="float: right;">Buserelin</span>
New Tree	D12.776.631.650.405.740.320.340 <span style="float: right;">Goserelin</span>
New Tree	D12.776.631.650.405.740.320.400 <span style="float: right;">Leuprolide</span>
New Tree	D12.776.631.650.405.740.320.580 <span style="float: right;">Nafarelin</span>
New Tree	D12.776.631.650.405.740.320.790 <span style="float: right;">Triptorelin Pamoate</span>
New Tree	D12.776.631.650.405.740.860 <span style="float: right;">Growth Hormone-Releasing Hormone</span>
New Tree	D12.776.631.650.405.740.860.780 <span style="float: right;">Sermorelin</span>
New Tree	D12.776.631.650.405.740.880 <span style="float: right;">Thyrotropin-Releasing Hormone</span>
New Tree	D12.776.631.650.405.935 <span style="float: right;">Pro-Opiomelanocortin</span>
New Tree	D12.776.631.650.405.935.119 <span style="float: right;">alpha-Endorphin</span>
New Tree	D12.776.631.650.405.935.179 <span style="float: right;">alpha-MSH</span>
New Tree	D12.776.631.650.405.935.239 <span style="float: right;">beta-Endorphin</span>
New Tree	D12.776.631.650.405.935.480 <span style="float: right;">beta-Lipotropin</span>
New Tree	D12.776.631.650.405.935.492 <span style="float: right;">beta-MSH</span>
New Tree	D12.776.631.650.405.935.498 <span style="float: right;">Corticotropin-Like Intermediate Lobe Peptide</span>
New Tree	D12.776.631.650.405.935.505 <span style="float: right;">gamma-Endorphin</span>
New Tree	D12.776.631.650.405.935.518 <span style="float: right;">gamma-Lipotropin</span>

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	D12.776.631.650.405.935.524 <span style="float: right;">gamma-MSH</span>
New Tree	D12.776.631.650.405.935.531 <span style="float: right;">Melanocortins</span>
New Tree	D12.776.631.650.405.935.531.500 <span style="float: right;">Adrenocorticotrophic Hormone</span>
New Tree	D12.776.631.650.405.935.531.500.200 <span style="float: right;">Cosyntropin</span>
New Tree	D12.776.631.650.405.935.531.750 <span style="float: right;">Melanocyte-Stimulating Hormones</span>
New Tree	D12.776.631.650.405.935.531.750.050 <span style="float: right;">alpha-MSH</span>
New Tree	D12.776.631.650.405.935.531.750.075 <span style="float: right;">beta-MSH</span>
New Tree	D12.776.631.650.405.935.531.750.115 <span style="float: right;">gamma-MSH</span>
New Tree	D12.776.631.650.405.967 <span style="float: right;">Prolactin-Releasing Hormone</span>
New Tree	D12.776.631.650.460 <span style="float: right;">Melanocyte-Stimulating Hormones</span>
New Tree	D12.776.631.650.460.050 <span style="float: right;">alpha-MSH</span>
New Tree	D12.776.631.650.460.075 <span style="float: right;">beta-MSH</span>
New Tree	D12.776.631.650.460.115 <span style="float: right;">gamma-MSH</span>
New Tree	D12.776.631.650.463 <span style="float: right;">Motilin</span>
New Tree	D12.776.631.650.500 <span style="float: right;">Neuropeptide Y</span>
New Tree	D12.776.631.650.525 <span style="float: right;">Neurophysins</span>
New Tree	D12.776.631.650.550 <span style="float: right;">Neurotensin</span>
New Tree	D12.776.631.650.575 <span style="float: right;">Opioid Peptides</span>
New Tree	D12.776.631.650.575.180 <span style="float: right;">Dynorphins</span>
New Tree	D12.776.631.650.575.241 <span style="float: right;">Endorphins</span>
New Tree	D12.776.631.650.575.241.040 <span style="float: right;">alpha-Endorphin</span>
New Tree	D12.776.631.650.575.241.080 <span style="float: right;">beta-Endorphin</span>

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	D12.776.631.650.575.241.540 <span style="float: right;">gamma-Endorphin</span>
New Tree	D12.776.631.650.575.281 <span style="float: right;">Enkephalins</span>
New Tree	D12.776.631.650.575.281.075 <span style="float: right;">Enkephalin, Ala(2)-MePhe(4)-Gly(5)-</span>
New Tree	D12.776.631.650.575.281.231 <span style="float: right;">Enkephalin, Leucine</span>
New Tree	D12.776.631.650.575.281.231.500 <span style="float: right;">Enkephalin, Leucine-2-Alanine</span>
New Tree	D12.776.631.650.575.281.381 <span style="float: right;">Enkephalin, Methionine</span>
New Tree	D12.776.631.650.575.281.381.500 <span style="float: right;">D-Ala(2),MePhe(4),Met(0)-ol-enkephalin</span>
New Tree	D12.776.631.650.575.281.600 <span style="float: right;">Enkephalin, D-Penicillamine (2,5)-</span>
New Tree	D12.776.631.650.600 <span style="float: right;">Pancreatic Polypeptide</span>
New Tree	D12.776.631.650.610 <span style="float: right;">Peptide PHI</span>
New Tree	D12.776.631.650.625 <span style="float: right;">Pituitary Adenylate Cyclase-Activating Polypeptide</span>
New Tree	D12.776.631.650.705 <span style="float: right;">Secretin</span>
New Tree	D12.776.631.650.800 <span style="float: right;">Tachykinins</span>
New Tree	D12.776.631.650.800.354 <span style="float: right;">Eledoisin</span>
New Tree	D12.776.631.650.800.475 <span style="float: right;">Kassinin</span>
New Tree	D12.776.631.650.800.500 <span style="float: right;">Neurokinin A</span>
New Tree	D12.776.631.650.800.550 <span style="float: right;">Neurokinin B</span>
New Tree	D12.776.631.650.800.625 <span style="float: right;">Physalaemin</span>
New Tree	D12.776.631.650.800.750 <span style="float: right;">Substance P</span>
New Tree	D12.776.631.650.810 <span style="float: right;">Thyrotropin-Releasing Hormone</span>
New Tree	D12.776.631.650.875 <span style="float: right;">Vasoactive Intestinal Peptide</span>
New Tree	D12.776.631.650.937 <span style="float: right;">Vasopressins</span>

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	D12.776.631.650.937.100 Arginine Vasopressin
New Tree	D12.776.631.650.937.100.250 Deamino Arginine Vasopressin
New Tree	D12.776.631.650.937.400 Lypressin
New Tree	D12.776.631.650.937.400.350 Felypressin
New Tree	D12.776.631.650.937.700 Ornipressin
New Heading	<b>D12.776.631.651 Nogo Receptors</b>
New Heading	<b>D12.776.631.651.250 Nogo Receptor 2</b>
New Heading	<b>D12.776.631.651.500 Nogo Receptor 1</b>
New Tree	D12.776.631.652 Olfactory Marker Protein
New Tree	D12.776.631.655 S100 Proteins
New Tree	D12.776.631.655.500 Leukocyte L1 Antigen Complex
New Tree	D12.776.631.655.500.100 Calgranulin A
New Tree	D12.776.631.655.500.200 Calgranulin B
New Tree	D12.776.631.655.750 S100 Calcium Binding Protein beta Subunit
New Tree	D12.776.631.750 Synapsins
New Tree	D12.776.631.775 Synaptogyrins
New Tree	D12.776.631.800 Synaptophysin
New Tree	D12.776.631.860 Synucleins
New Tree	D12.776.631.860.500 alpha-Synuclein
New Tree	D12.776.631.860.625 beta-Synuclein
New Tree	D12.776.631.860.750 gamma-Synuclein
New Tree	D12.776.631.920 Tubulin

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	D12.776.631.960 Voltage-Gated Sodium Channels
New Tree	D12.776.631.960.100 NAV1.1 Voltage-Gated Sodium Channel
New Tree	D12.776.631.960.200 NAV1.2 Voltage-Gated Sodium Channel
New Tree	D12.776.631.960.300 NAV1.3 Voltage-Gated Sodium Channel
New Tree	D12.776.631.960.500 NAV1.5 Voltage-Gated Sodium Channel
New Tree	D12.776.631.960.600 NAV1.6 Voltage-Gated Sodium Channel
New Tree	D12.776.631.960.700 NAV1.7 Voltage-Gated Sodium Channel
New Tree	D12.776.631.960.800 NAV1.8 Voltage-Gated Sodium Channel
New Tree	D12.776.631.960.900 NAV1.9 Voltage-Gated Sodium Channel
New Tree	D12.776.631.960.970 Voltage-Gated Sodium Channel beta Subunits
New Tree	D12.776.631.960.970.100 Voltage-Gated Sodium Channel beta-1 Subunit
New Tree	D12.776.631.960.970.200 Voltage-Gated Sodium Channel beta-2 Subunit
New Tree	D12.776.631.960.970.300 Voltage-Gated Sodium Channel beta-3 Subunit
New Tree	D12.776.631.960.970.650 Voltage-Gated Sodium Channel beta-4 Subunit
New Heading	<b>D12.776.637 Parkinson Disease Associated Proteins</b>
New Tree	D12.776.637.500 alpha-Synuclein
New Heading	<b>D12.776.637.750 Leucine-Rich Repeat Serine-Threonine Protein Kinase-2</b>
New Heading	<b>D12.776.637.875 Protein Deglycase DJ-1</b>
New Tree	D12.776.637.937 Ubiquitin Thiolesterase
Old Tree	D12.776.641 Nerve Tissue Proteins
Old Tree	D12.776.641.024 Acid Sensing Ion Channels
Old Tree	D12.776.641.050 Agrin
Old Tree	D12.776.641.069 Ataxins
Old Tree	D12.776.641.069.500 Ataxin-1

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	D12.776.641.069.750 Ataxin-2
Old Tree	D12.776.641.069.875 Ataxin-3
Old Tree	D12.776.641.069.901 Ataxin-7
Old Tree	D12.776.641.069.950 Ataxin-10
Old Tree	D12.776.641.087 Calbindin 1
Old Tree	D12.776.641.124 Chimerin Proteins
Old Tree	D12.776.641.124.200 Chimerin 1
Old Tree	D12.776.641.199 Chromogranins
Old Tree	D12.776.641.199.249 Chromogranin A
Old Tree	D12.776.641.199.500 Chromogranin B
Old Tree	D12.776.641.199.750 Secretogranin II
Old Tree	D12.776.641.249 Dopamine and cAMP-Regulated Phosphoprotein 32
Old Tree	D12.776.641.299 Fragile X Mental Retardation Protein
Old Tree	D12.776.641.400 GAP-43 Protein
Old Tree	D12.776.641.480 Glucose Transporter Type 3
Old Tree	D12.776.641.520 ELAV Proteins
Old Tree	D12.776.641.520.500 ELAV-Like Protein 2
Old Tree	D12.776.641.520.750 ELAV-Like Protein 3
Old Tree	D12.776.641.520.875 ELAV-Like Protein 4
Old Tree	D12.776.641.560 Microtubule-Associated Proteins
Old Tree	D12.776.641.560.450 Kinesin
Old Tree	D12.776.641.560.510 tau Proteins
Old Tree	D12.776.641.580 Myelin Proteins
Old Tree	D12.776.641.580.124 2',3'-Cyclic Nucleotide 3'-Phosphodiesterase
Old Tree	D12.776.641.580.249 Myelin and Lymphocyte-Associated Proteolipid Proteins
Old Tree	D12.776.641.580.500 Myelin-Associated Glycoprotein
Old Tree	D12.776.641.580.510 Myelin Basic Protein
Old Tree	D12.776.641.580.530 Myelin-Oligodendrocyte Glycoprotein
Old Tree	D12.776.641.580.550 Myelin P0 Protein
Old Tree	D12.776.641.580.560 Myelin P2 Protein
Old Tree	D12.776.641.580.580 Myelin Proteolipid Protein
Old Tree	D12.776.641.580.895 Oligodendrocyte-Myelin Glycoprotein
Old Tree	D12.776.641.590 Natriuretic Peptide, Brain
Old Tree	D12.776.641.600 Nerve Growth Factors
Old Tree	D12.776.641.600.100 Brain-Derived Neurotrophic Factor
Old Tree	D12.776.641.600.212 Ciliary Neurotrophic Factor

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	D12.776.641.600.325      Glia Maturation Factor
Old Tree	D12.776.641.600.381      Glial Cell Line-Derived Neurotrophic Factors
Old Tree	D12.776.641.600.381.500      Glial Cell Line-Derived Neurotrophic Factor
Old Tree	D12.776.641.600.381.750      Neurturin
Old Tree	D12.776.641.600.437      Nerve Growth Factor
Old Tree	D12.776.641.600.550      Neuregulins
Old Tree	D12.776.641.600.550.750      Neuregulin-1
Old Tree	D12.776.641.600.775      Neurotrophin 3
Old Tree	D12.776.641.600.887      Pituitary Adenylate Cyclase-Activating Polypeptide
Old Tree	D12.776.641.607      Nestin
Old Tree	D12.776.641.615      Neuroendocrine Secretory Protein 7B2
Old Tree	D12.776.641.630      Neurofilament Proteins
Old Tree	D12.776.641.640      Neurogranin
Old Tree	D12.776.641.642      Neuronal Apoptosis-Inhibitory Protein
Old Tree	D12.776.641.645      Neuronal Calcium-Sensor Proteins
Old Tree	D12.776.641.645.124      Guanylate Cyclase-Activating Proteins
Old Tree	D12.776.641.645.249      Hippocalcin
Old Tree	D12.776.641.645.374      Kv Channel-Interacting Proteins
Old Tree	D12.776.641.645.500      Neurocalcin
Old Tree	D12.776.641.645.750      Recoverin
Old Tree	D12.776.641.650      Neuropeptides
Old Tree	D12.776.641.650.070      Angiotensins
Old Tree	D12.776.641.650.070.075      Angiotensin I
Old Tree	D12.776.641.650.070.078      Angiotensin II
Old Tree	D12.776.641.650.070.080      Angiotensin III
Old Tree	D12.776.641.650.085      Bombesin
Old Tree	D12.776.641.650.090      Bradykinin
Old Tree	D12.776.641.650.095      Calcitonin
Old Tree	D12.776.641.650.097      Calcitonin Gene-Related Peptide
Old Tree	D12.776.641.650.100      Carnosine
Old Tree	D12.776.641.650.200      Delta Sleep-Inducing Peptide
Old Tree	D12.776.641.650.235      FMRFamide
Old Tree	D12.776.641.650.250      Galanin
Old Tree	D12.776.641.650.300      Gastric Inhibitory Polypeptide
Old Tree	D12.776.641.650.315      Gastrin-Releasing Peptide
Old Tree	D12.776.641.650.320      Gastrins

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	D12.776.641.650.363      Orexins
Old Tree	D12.776.641.650.405      Hypothalamic Hormones
Old Tree	D12.776.641.650.405.700      Pituitary Hormone Release Inhibiting Hormones
Old Tree	D12.776.641.650.405.700.500      MSH Release-Inhibiting Hormone
Old Tree	D12.776.641.650.405.700.750      Prolactin Release-Inhibiting Factors
Old Tree	D12.776.641.650.405.700.875      Somatostatin
Old Tree	D12.776.641.650.405.700.875.500      Somatostatin-28
Old Tree	D12.776.641.650.405.740      Pituitary Hormone-Releasing Hormones
Old Tree	D12.776.641.650.405.740.140      Corticotropin-Releasing Hormone
Old Tree	D12.776.641.650.405.740.320      Gonadotropin-Releasing Hormone
Old Tree	D12.776.641.650.405.740.320.100      Buserelin
Old Tree	D12.776.641.650.405.740.320.340      Goserelin
Old Tree	D12.776.641.650.405.740.320.400      Leuprolide
Old Tree	D12.776.641.650.405.740.320.580      Nafarelin
Old Tree	D12.776.641.650.405.740.320.790      Triptorelin Pamoate
Old Tree	D12.776.641.650.405.740.860      Growth Hormone-Releasing Hormone
Old Tree	D12.776.641.650.405.740.860.780      Sermorelin
Old Tree	D12.776.641.650.405.740.880      Thyrotropin-Releasing Hormone
Old Tree	D12.776.641.650.405.935      Pro-Opiomelanocortin
Old Tree	D12.776.641.650.405.935.119      alpha-Endorphin
Old Tree	D12.776.641.650.405.935.179      alpha-MSH
Old Tree	D12.776.641.650.405.935.239      beta-Endorphin
Old Tree	D12.776.641.650.405.935.480      beta-Lipotropin
Old Tree	D12.776.641.650.405.935.492      beta-MSH
Old Tree	D12.776.641.650.405.935.498      Corticotropin-Like Intermediate Lobe Peptide
Old Tree	D12.776.641.650.405.935.505      gamma-Endorphin
Old Tree	D12.776.641.650.405.935.518      gamma-Lipotropin
Old Tree	D12.776.641.650.405.935.524      gamma-MSH
Old Tree	D12.776.641.650.405.935.531      Melanocortins
Old Tree	D12.776.641.650.405.935.531.500      Adrenocorticotrophic Hormone
Old Tree	D12.776.641.650.405.935.531.500.200      Cosyntropin
Old Tree	D12.776.641.650.405.935.531.750      Melanocyte-Stimulating Hormones
Old Tree	D12.776.641.650.405.935.531.750.050      alpha-MSH
Old Tree	D12.776.641.650.405.935.531.750.075      beta-MSH
Old Tree	D12.776.641.650.405.935.531.750.115      gamma-MSH
Old Tree	D12.776.641.650.405.967      Prolactin-Releasing Hormone



## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	D12.776.641.650.460 Melanocyte-Stimulating Hormones
Old Tree	D12.776.641.650.460.050 alpha-MSH
Old Tree	D12.776.641.650.460.075 beta-MSH
Old Tree	D12.776.641.650.460.115 gamma-MSH
Old Tree	D12.776.641.650.463 Motilin
Old Tree	D12.776.641.650.500 Neuropeptide Y
Old Tree	D12.776.641.650.525 Neurophysins
Old Tree	D12.776.641.650.550 Neurotensin
Old Tree	D12.776.641.650.575 Opioid Peptides
Old Tree	D12.776.641.650.575.180 Dynorphins
Old Tree	D12.776.641.650.575.241 Endorphins
Old Tree	D12.776.641.650.575.241.040 alpha-Endorphin
Old Tree	D12.776.641.650.575.241.080 beta-Endorphin
Old Tree	D12.776.641.650.575.241.540 gamma-Endorphin
Old Tree	D12.776.641.650.575.281 Enkephalins
Old Tree	D12.776.641.650.575.281.075 Enkephalin, Ala(2)-MePhe(4)-Gly(5)-
Old Tree	D12.776.641.650.575.281.231 Enkephalin, Leucine
Old Tree	D12.776.641.650.575.281.231.500 Enkephalin, Leucine-2-Alanine
Old Tree	D12.776.641.650.575.281.381 Enkephalin, Methionine
Old Tree	D12.776.641.650.575.281.381.500 D-Ala(2),MePhe(4),Met(0)-ol-enkephalin
Old Tree	D12.776.641.650.575.281.600 Enkephalin, D-Penicillamine (2,5)-
Old Tree	D12.776.641.650.600 Pancreatic Polypeptide
Old Tree	D12.776.641.650.610 Peptide PHI
Old Tree	D12.776.641.650.625 Pituitary Adenylate Cyclase-Activating Polypeptide
Old Tree	D12.776.641.650.705 Secretin
Old Tree	D12.776.641.650.800 Tachykinins
Old Tree	D12.776.641.650.800.354 Eledoisin
Old Tree	D12.776.641.650.800.475 Kassinin
Old Tree	D12.776.641.650.800.500 Neurokinin A
Old Tree	D12.776.641.650.800.550 Neurokinin B
Old Tree	D12.776.641.650.800.625 Physalaemin
Old Tree	D12.776.641.650.800.750 Substance P
Old Tree	D12.776.641.650.810 Thyrotropin-Releasing Hormone
Old Tree	D12.776.641.650.875 Vasoactive Intestinal Peptide
Old Tree	D12.776.641.650.937 Vasopressins
Old Tree	D12.776.641.650.937.100 Arginine Vasopressin

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	D12.776.641.650.937.100.250 Deamino Arginine Vasopressin
Old Tree	D12.776.641.650.937.400 Lypressin
Old Tree	D12.776.641.650.937.400.350 Felypressin
Old Tree	D12.776.641.650.937.700 Ornipressin
Old Tree	D12.776.641.652 Olfactory Marker Protein
Old Tree	D12.776.641.655 S100 Proteins
Old Tree	D12.776.641.655.500 Leukocyte L1 Antigen Complex
Old Tree	D12.776.641.655.500.100 Calgranulin A
Old Tree	D12.776.641.655.500.200 Calgranulin B
Old Tree	D12.776.641.655.750 S100 Calcium Binding Protein beta Subunit
Old Tree	D12.776.641.750 Synapsins
Old Tree	D12.776.641.775 Synaptogyrins
Old Tree	D12.776.641.800 Synaptophysin
Old Tree	D12.776.641.860 Synucleins
Old Tree	D12.776.641.860.500 alpha-Synuclein
Old Tree	D12.776.641.860.625 beta-Synuclein
Old Tree	D12.776.641.860.750 gamma-Synuclein
Old Tree	D12.776.641.920 Tubulin
Old Tree	D12.776.641.960 Voltage-Gated Sodium Channels
Old Tree	D12.776.641.960.100 NAV1.1 Voltage-Gated Sodium Channel
Old Tree	D12.776.641.960.200 NAV1.2 Voltage-Gated Sodium Channel
Old Tree	D12.776.641.960.300 NAV1.3 Voltage-Gated Sodium Channel
Old Tree	D12.776.641.960.500 NAV1.5 Voltage-Gated Sodium Channel
Old Tree	D12.776.641.960.600 NAV1.6 Voltage-Gated Sodium Channel
Old Tree	D12.776.641.960.700 NAV1.7 Voltage-Gated Sodium Channel
Old Tree	D12.776.641.960.800 NAV1.8 Voltage-Gated Sodium Channel
Old Tree	D12.776.641.960.900 NAV1.9 Voltage-Gated Sodium Channel
Old Tree	D12.776.641.960.970 Voltage-Gated Sodium Channel beta Subunits
Old Tree	D12.776.641.960.970.100 Voltage-Gated Sodium Channel beta-1 Subunit
Old Tree	D12.776.641.960.970.200 Voltage-Gated Sodium Channel beta-2 Subunit
Old Tree	D12.776.641.960.970.300 Voltage-Gated Sodium Channel beta-3 Subunit
Old Tree	D12.776.641.960.970.650 Voltage-Gated Sodium Channel beta-4 Subunit
-	D12.776.642 Protein Corona
-	D12.776.643 Protein Degradation End Products
-	D12.776.643.249 Advanced Oxidation Protein Products
-	D12.776.643.500 Glycosylation End Products, Advanced

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.645 Proteinase Inhibitory Proteins, Secretory
-	D12.776.645.437 Elafin
-	D12.776.645.468 Protease Nexins
-	D12.776.645.468.500 Amyloid beta-Protein Precursor
-	D12.776.645.468.750 Serpin E2
-	D12.776.645.500 Secretory Leukocyte Peptidase Inhibitor
-	D12.776.645.875 Tissue Inhibitor of Metalloproteinases
-	D12.776.645.875.450 Tissue Inhibitor of Metalloproteinase-1
-	D12.776.645.875.500 Tissue Inhibitor of Metalloproteinase-2
-	D12.776.645.875.550 Tissue Inhibitor of Metalloproteinase-3
-	D12.776.660 Nuclear Proteins
-	D12.776.660.049 Antennapedia Homeodomain Protein
-	D12.776.660.075 Ataxins
-	D12.776.660.075.500 Ataxin-1
-	D12.776.660.075.750 Ataxin-2
-	D12.776.660.075.875 Ataxin-3
-	D12.776.660.075.900 Ataxin-7
-	D12.776.660.100 BRCA1 Protein
-	D12.776.660.105 BRCA2 Protein
-	D12.776.660.167 CCAAT-Enhancer-Binding Proteins
-	D12.776.660.167.249 CCAAT-Binding Factor
-	D12.776.660.167.500 CCAAT-Enhancer-Binding Protein-alpha
-	D12.776.660.167.750 CCAAT-Enhancer-Binding Protein-beta
-	D12.776.660.167.812 CCAAT-Enhancer-Binding Protein-delta
-	D12.776.660.167.875 Transcription Factor CHOP
-	D12.776.660.167.937 Y-Box-Binding Protein 1
New Heading	<b>D12.776.660.201 CDX2 Transcription Factor</b>
-	D12.776.660.235 Chromosomal Proteins, Non-Histone
-	D12.776.660.235.199 Centromere Protein B
-	D12.776.660.235.400 High Mobility Group Proteins
-	D12.776.660.235.400.400 HMGN Proteins
-	D12.776.660.235.400.400.200 HMGN1 Protein
-	D12.776.660.235.400.400.300 HMGN2 Protein
-	D12.776.660.235.400.500 HMGA Proteins
-	D12.776.660.235.400.500.100 HMGA1a Protein

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.660.235.400.500.200 HMGA1b Protein
-	D12.776.660.235.400.500.400 HMGA1c Protein
-	D12.776.660.235.400.500.600 HMGA2 Protein
-	D12.776.660.235.400.600 HMGB Proteins
-	D12.776.660.235.400.600.300 HMGB1 Protein
-	D12.776.660.235.400.600.600 HMGB2 Protein
-	D12.776.660.235.400.600.800 HMGB3 Protein
-	D12.776.660.235.400.750 SOX Transcription Factors
-	D12.776.660.235.400.750.049 Sex-Determining Region Y Protein
-	D12.776.660.235.400.750.100 SOXB1 Transcription Factors
-	D12.776.660.235.400.750.200 SOXB2 Transcription Factors
-	D12.776.660.235.400.750.300 SOXC Transcription Factors
-	D12.776.660.235.400.750.400 SOXD Transcription Factors
-	D12.776.660.235.400.750.500 SOXE Transcription Factors
-	D12.776.660.235.400.750.500.500 SOX9 Transcription Factor
-	D12.776.660.235.400.750.600 SOXF Transcription Factors
-	D12.776.660.235.400.800 TCF Transcription Factors
-	D12.776.660.235.400.800.500 Lymphoid Enhancer-Binding Factor 1
-	D12.776.660.235.400.800.750 T Cell Transcription Factor 1
-	D12.776.660.235.400.800.812 Transcription Factor 7-Like 1 Protein
-	D12.776.660.235.400.800.875 Transcription Factor 7-Like 2 Protein
-	D12.776.660.235.500 Minichromosome Maintenance Proteins
-	D12.776.660.235.500.100 Minichromosome Maintenance 1 Protein
-	D12.776.660.235.500.200 Minichromosome Maintenance Complex Component 2
-	D12.776.660.235.500.300 Minichromosome Maintenance Complex Component 3
-	D12.776.660.235.500.400 Minichromosome Maintenance Complex Component 4
-	D12.776.660.235.500.500 Minichromosome Maintenance Complex Component 5
-	D12.776.660.235.500.600 Minichromosome Maintenance Complex Component 6
-	D12.776.660.235.500.700 Minichromosome Maintenance Complex Component 7
-	D12.776.660.235.500.800 Minichromosome Maintenance Complex Component 8
-	D12.776.660.235.500.900 Minichromosome Maintenance Complex Component 9

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D12.776.660.235.550	Methyl-CpG-Binding Protein 2
-	D12.776.660.235.600	Polycomb-Group Proteins
-	D12.776.660.235.600.100	Polycomb Repressive Complex 1
-	D12.776.660.235.600.200	Polycomb Repressive Complex 2
New Heading	<b>D12.776.660.235.600.200.250</b>	<b>Enhancer of Zeste Homolog 2 Protein</b>
-	D12.776.660.235.600.200.500	Retinoblastoma-Binding Protein 4
-	D12.776.660.235.600.200.750	Retinoblastoma-Binding Protein 7
New Heading	<b>D12.776.660.235.650</b>	<b>SMARCB1 Protein</b>
-	D12.776.660.235.700	Telomere-Binding Proteins
-	D12.776.660.235.700.500 B	Heterogeneous-Nuclear Ribonucleoprotein Group A-
-	D12.776.660.235.700.750	Telomeric Repeat Binding Protein 1
-	D12.776.660.235.700.875	Telomeric Repeat Binding Protein 2
New Heading	<b>D12.776.660.235.850</b>	<b>Tumor Suppressor p53-Binding Protein 1</b>
-	D12.776.660.264	Fanconi Anemia Complementation Group A Protein
-	D12.776.660.285	Fanconi Anemia Complementation Group D2 Protein
-	D12.776.660.289	Fanconi Anemia Complementation Group E Protein
-	D12.776.660.293	Fanconi Anemia Complementation Group F Protein
-	D12.776.660.352	Hepatocyte Nuclear Factors
-	D12.776.660.352.500	Hepatocyte Nuclear Factor 1
-	D12.776.660.352.500.500	Hepatocyte Nuclear Factor 1-alpha
-	D12.776.660.352.500.750	Hepatocyte Nuclear Factor 1-beta
-	D12.776.660.352.750	Hepatocyte Nuclear Factor 3-alpha
-	D12.776.660.352.875	Hepatocyte Nuclear Factor 3-beta
-	D12.776.660.352.937	Hepatocyte Nuclear Factor 3-gamma
-	D12.776.660.352.968	Hepatocyte Nuclear Factor 4
-	D12.776.660.352.984	Hepatocyte Nuclear Factor 6
-	D12.776.660.470	Histones
New Tree	<b>D12.776.660.478</b>	<b>Host Cell Factor C1</b>
-	D12.776.660.486 Protein	Immunoglobulin J Recombination Signal Sequence-Binding
-	D12.776.660.502	Karyopherins
-	D12.776.660.502.100	alpha Karyopherins
-	D12.776.660.502.500	beta Karyopherins

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D12.776.660.526	Mad2 Proteins
-	D12.776.660.551	Mediator Complex
-	D12.776.660.551.249	Cyclin C
-	D12.776.660.551.500	Cyclin-Dependent Kinase 8
-	D12.776.660.551.750	Mediator Complex Subunit 1
-	D12.776.660.600	NF-kappa B
-	D12.776.660.600.124	NF-kappa B p50 Subunit
-	D12.776.660.600.186	NF-kappa B p52 Subunit
-	D12.776.660.600.249	Transcription Factor RelA
-	D12.776.660.600.500	Transcription Factor RelB
-	D12.776.660.625	Antigens, Nuclear
-	D12.776.660.625.500	Ki-67 Antigen
New Heading	<b>D12.776.660.625.625</b>	<b>Ku Autoantigen</b>
-	D12.776.660.625.750	snRNP Core Proteins
-	D12.776.660.650	Nuclear Matrix-Associated Proteins
-	D12.776.660.650.500	Heterogeneous-Nuclear Ribonucleoprotein U
-	D12.776.660.650.875	Lamins
-	D12.776.660.650.875.500	Lamin Type A
-	D12.776.660.650.875.750	Lamin Type B
-	D12.776.660.675	Nuclear Receptor Coactivators
-	D12.776.660.675.049	Mediator Complex Subunit 1
-	D12.776.660.675.100	Nuclear Receptor Coactivator 1
-	D12.776.660.675.200	Nuclear Receptor Coactivator 2
-	D12.776.660.675.300	Nuclear Receptor Coactivator 3
New Heading	<b>D12.776.660.675.650 Coactivator 1-alpha</b>	<b>Peroxisome Proliferator-Activated Receptor Gamma</b>
-	D12.776.660.700	Oncogene Protein p53(v-myc)
-	D12.776.660.740	Proliferating Cell Nuclear Antigen
New Heading	<b>D12.776.660.745</b>	<b>Promyelocytic Leukemia Protein</b>
-	D12.776.660.750	Protamines
-	D12.776.660.750.270	Clupeine
-	D12.776.660.750.740	Salmine
-	D12.776.660.760	Proto-Oncogene Proteins c-fos
-	D12.776.660.763	Proto-Oncogene Proteins c-jun
-	D12.776.660.764	Proto-Oncogene Proteins c-mdm2

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D12.776.660.765	Proto-Oncogene Proteins c-myc
-	D12.776.660.767	Proto-Oncogene Proteins c-rel
-	D12.776.660.768	ran GTP-Binding Protein
-	D12.776.660.769	Retinoblastoma Binding Proteins
-	D12.776.660.769.049	E2F1 Transcription Factor
-	D12.776.660.769.100	Retinoblastoma-Binding Protein 1
-	D12.776.660.769.200	Retinoblastoma-Binding Protein 2
-	D12.776.660.769.600	Retinoblastoma-Binding Protein 4
-	D12.776.660.769.700	Retinoblastoma-Binding Protein 7
-	D12.776.660.770	Retinoblastoma-Like Protein p107
-	D12.776.660.794	Retinoblastoma-Like Protein p130
-	D12.776.660.807	Retinoblastoma Protein
-	D12.776.660.820 cerevisiae	Silent Information Regulator Proteins, Saccharomyces
-	D12.776.660.825	Tumor Suppressor Protein p53
New Heading	<b>D12.776.660.912</b>	<b>Tumor Protein p73</b>
-	D12.776.664	Nucleoproteins
-	D12.776.664.224	Chromatin
-	D12.776.664.224.270	Euchromatin
-	D12.776.664.224.466	Heterochromatin
-	D12.776.664.224.550	Nucleosomes
-	D12.776.664.235	Chromosomal Proteins, Non-Histone
-	D12.776.664.235.199	Centromere Protein B
-	D12.776.664.235.400	High Mobility Group Proteins
-	D12.776.664.235.400.400	HMGN Proteins
-	D12.776.664.235.400.400.200	HMGN1 Protein
-	D12.776.664.235.400.400.300	HMGN2 Protein
-	D12.776.664.235.400.500	HMGA Proteins
-	D12.776.664.235.400.500.100	HMGA1a Protein
-	D12.776.664.235.400.500.200	HMGA1b Protein
-	D12.776.664.235.400.500.300	HMGA1c Protein
-	D12.776.664.235.400.500.600	HMGA2 Protein
-	D12.776.664.235.400.600	HMGB Proteins
-	D12.776.664.235.400.600.300	HMGB1 Protein
-	D12.776.664.235.400.600.600	HMGB2 Protein

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D12.776.664.235.400.600.800	HMGB3 Protein
-	D12.776.664.235.400.750	SOX Transcription Factors
-	D12.776.664.235.400.750.049	Sex-Determining Region Y Protein
-	D12.776.664.235.400.750.100	SOXB1 Transcription Factors
-	D12.776.664.235.400.750.200	SOXB2 Transcription Factors
-	D12.776.664.235.400.750.300	SOXC Transcription Factors
-	D12.776.664.235.400.750.400	SOXD Transcription Factors
-	D12.776.664.235.400.750.500	SOXE Transcription Factors
-	D12.776.664.235.400.750.500.500	SOX9 Transcription Factor
-	D12.776.664.235.400.750.600	SOXF Transcription Factors
-	D12.776.664.235.400.800	TCF Transcription Factors
-	D12.776.664.235.400.800.500	Lymphoid Enhancer-Binding Factor 1
-	D12.776.664.235.400.800.750	T Cell Transcription Factor 1
-	D12.776.664.235.400.800.812	Transcription Factor 7-Like 1 Protein
-	D12.776.664.235.400.800.875	Transcription Factor 7-Like 2 Protein
-	D12.776.664.235.700	Methyl-CpG-Binding Protein 2
-	D12.776.664.235.750	Minichromosome Maintenance Proteins
-	D12.776.664.235.750.100	Minichromosome Maintenance 1 Protein
-	D12.776.664.235.750.200 2	Minichromosome Maintenance Complex Component
-	D12.776.664.235.750.300 3	Minichromosome Maintenance Complex Component
-	D12.776.664.235.750.400 4	Minichromosome Maintenance Complex Component
-	D12.776.664.235.750.500 5	Minichromosome Maintenance Complex Component
-	D12.776.664.235.750.600 6	Minichromosome Maintenance Complex Component
-	D12.776.664.235.750.700 7	Minichromosome Maintenance Complex Component
-	D12.776.664.235.750.800 8	Minichromosome Maintenance Complex Component
-	D12.776.664.235.750.900 9	Minichromosome Maintenance Complex Component
-	D12.776.664.235.800	Polycomb-Group Proteins
-	D12.776.664.235.800.100	Polycomb Repressive Complex 1
-	D12.776.664.235.800.200	Polycomb Repressive Complex 2
New Heading	<b>D12.776.664.235.800.200.250</b>	<b>Enhancer of Zeste Homolog 2 Protein</b>



## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D12.776.664.235.800.200.500	Retinoblastoma-Binding Protein 4
-	D12.776.664.235.800.200.750	Retinoblastoma-Binding Protein 7
New Heading	<b>D12.776.664.235.900</b>	<b>SMARCB1 Protein</b>
New Heading	<b>D12.776.664.235.950</b>	<b>Tumor Suppressor p53-Binding Protein 1</b>
-	D12.776.664.275	Deoxyribonucleoproteins
-	D12.776.664.469	Histones
-	D12.776.664.750	Protamines
-	D12.776.664.750.270	Clupeine
-	D12.776.664.750.739	Salmine
-	D12.776.664.962	RNA-Binding Proteins
-	D12.776.664.962.061	Butyrate Response Factor 1
-	D12.776.664.962.124	Fragile X Mental Retardation Protein
-	D12.776.664.962.186	Gene Products, rev
-	D12.776.664.962.186.500	rev Gene Products, Human Immunodeficiency Virus
-	D12.776.664.962.249	Host Factor 1 Protein
-	D12.776.664.962.374	Iron Regulatory Protein 1
-	D12.776.664.962.437	Iron Regulatory Protein 2
-	D12.776.664.962.444	mRNA Cleavage and Polyadenylation Factors
-	D12.776.664.962.444.240	Cleavage And Polyadenylation Specificity Factor
-	D12.776.664.962.444.249	Cleavage Stimulation Factor
-	D12.776.664.962.452	Poly(A)-Binding Proteins
-	D12.776.664.962.452.125	Ataxin-2
-	D12.776.664.962.452.249	Poly(A)-Binding Protein I
-	D12.776.664.962.452.500	Poly(A)-Binding Protein II
Old Tree	<b>D12.776.664.962.468</b>	<b>Polypyrimidine Tract-Binding Protein</b>
-	D12.776.664.962.500	Ribonucleoproteins
-	D12.776.664.962.500.625	Ribonuclease P
-	D12.776.664.962.500.750	Ribonucleoproteins, Small Cytoplasmic
-	D12.776.664.962.500.750.800	Signal Recognition Particle
-	D12.776.664.962.500.875	Ribonucleoproteins, Small Nuclear
-	D12.776.664.962.500.875.590	Ribonucleoproteins, Small Nucleolar
-	D12.776.664.962.500.875.600	Ribonucleoprotein, U1 Small Nuclear
-	D12.776.664.962.500.875.605	Ribonucleoprotein, U2 Small Nuclear
-	D12.776.664.962.500.875.615	Ribonucleoprotein, U4-U6 Small Nuclear

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.664.962.500.875.620 Ribonucleoprotein, U5 Small Nuclear
-	D12.776.664.962.500.875.625 Ribonucleoprotein, U7 Small Nuclear
-	D12.776.664.962.500.875.700 snRNP Core Proteins
-	D12.776.664.962.500.906 RNA-Induced Silencing Complex
-	D12.776.664.962.500.906.500 Argonaute Proteins
-	D12.776.664.962.500.921 Telomerase
-	D12.776.664.962.500.937 Vault Ribonucleoprotein Particles
-	D12.776.664.962.750 RNA Cap-Binding Proteins
-	D12.776.664.962.750.374 Eukaryotic Initiation Factor-4F
-	D12.776.664.962.750.750 Nuclear Cap-Binding Protein Complex
-	D12.776.664.962.813 RNA Recognition Motif (RRM) Proteins
-	D12.776.664.962.813 RNA Recognition Motif Proteins
-	D12.776.664.962.813.250 CELF Proteins
-	D12.776.664.962.813.250.500 CELF1 Protein
-	D12.776.664.962.813.500 ELAV Proteins
-	D12.776.664.962.813.500.250 ELAV-Like Protein 1
-	D12.776.664.962.813.500.500 ELAV-Like Protein 2
-	D12.776.664.962.813.500.750 ELAV-Like Protein 3
-	D12.776.664.962.813.500.875 ELAV-Like Protein 4
-	D12.776.664.962.813.750 Heterogeneous-Nuclear Ribonucleoproteins
-	D12.776.664.962.813.750.100 Heterogeneous-Nuclear Ribonucleoprotein Group A-B
-	D12.776.664.962.813.750.200 Heterogeneous-Nuclear Ribonucleoprotein Group C
-	D12.776.664.962.813.750.300 Heterogeneous-Nuclear Ribonucleoprotein D
-	D12.776.664.962.813.750.400 Heterogeneous-Nuclear Ribonucleoprotein Group F-H
-	D12.776.664.962.813.750.500 Heterogeneous-Nuclear Ribonucleoprotein K
-	D12.776.664.962.813.750.600 Heterogeneous-Nuclear Ribonucleoprotein L
-	D12.776.664.962.813.750.700 Heterogeneous-Nuclear Ribonucleoprotein Group M
-	D12.776.664.962.813.750.800 Heterogeneous-Nuclear Ribonucleoprotein U
-	D12.776.664.962.813.750.900 RNA-Binding Protein EWS
-	D12.776.664.962.813.750.902 RNA-Binding Protein FUS
New Heading	<b>D12.776.664.962.813.875 Peroxisome Proliferator-Activated Receptor Gamma Coactivator 1-alpha</b>
New Heading	<b>D12.776.664.962.829 RNA Splicing Factors</b>

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	<a href="#">D12.776.664.962.829.250</a>	<a href="#">Polypyrimidine Tract-Binding Protein</a>
New Heading	<b><a href="#">D12.776.664.962.829.282</a></b>	<b><a href="#">PTB-Associated Splicing Factor</a></b>
New Tree	<a href="#">D12.776.664.962.829.313</a>	<a href="#">Ribonucleoprotein, U1 Small Nuclear</a>
New Tree	<a href="#">D12.776.664.962.829.344</a>	<a href="#">Ribonucleoprotein, U2 Small Nuclear</a>
New Tree	<a href="#">D12.776.664.962.829.375</a>	<a href="#">Ribonucleoprotein, U5 Small Nuclear</a>
New Tree	<a href="#">D12.776.664.962.829.500</a>	<a href="#">Serine-Arginine Splicing Factors</a>
New Heading	<b><a href="#">D12.776.664.962.829.750</a></b>	<b><a href="#">Splicing Factor U2AF</a></b>
Old Tree	<b><a href="#">D12.776.664.962.844</a></b>	<b><a href="#">Serine-Arginine Splicing Factors</a></b>
-	<a href="#">D12.776.664.962.875</a>	<a href="#">SMN Complex Proteins</a>
-	<a href="#">D12.776.664.962.875.249</a>	<a href="#">DEAD Box Protein 20</a>
-	<a href="#">D12.776.664.962.875.500</a>	<a href="#">Survival of Motor Neuron 1 Protein</a>
-	<a href="#">D12.776.664.962.875.750</a>	<a href="#">Survival of Motor Neuron 2 Protein</a>
-	<a href="#">D12.776.691</a>	<a href="#">Oxidative Phosphorylation Coupling Factors</a>
-	<a href="#">D12.776.719</a>	<a href="#">Peptones</a>
-	<a href="#">D12.776.744</a>	<a href="#">Phosphoproteins</a>
-	<a href="#">D12.776.744.049</a>	<a href="#">bcl-Associated Death Protein</a>
-	<a href="#">D12.776.744.100</a>	<a href="#">BRCA1 Protein</a>
-	<a href="#">D12.776.744.150</a>	<a href="#">Caseins</a>
-	<a href="#">D12.776.744.287</a>	<a href="#">Caveolin 1</a>
-	<a href="#">D12.776.744.356</a>	<a href="#">Caveolin 2</a>
-	<a href="#">D12.776.744.360</a>	<a href="#">CDC2 Protein Kinase</a>
-	<a href="#">D12.776.744.390</a>	<a href="#">Cortactin</a>
-	<a href="#">D12.776.744.425</a>	<a href="#">Crk-Associated Substrate Protein</a>
-	<a href="#">D12.776.744.459</a>	<a href="#">Dopamine and cAMP-Regulated Phosphoprotein 32</a>
-	<a href="#">D12.776.744.476</a>	<a href="#">Fanconi Anemia Complementation Group A Protein</a>
-	<a href="#">D12.776.744.484</a>	<a href="#">Fanconi Anemia Complementation Group D2 Protein</a>
-	<a href="#">D12.776.744.488</a>	<a href="#">Fanconi Anemia Complementation Group G Protein</a>
-	<a href="#">D12.776.744.493</a>	<a href="#">Focal Adhesion Kinase 1</a>
-	<a href="#">D12.776.744.527</a>	<a href="#">Integrin-Binding Sialoprotein</a>
-	<a href="#">D12.776.744.562</a>	<a href="#">Interferon Regulatory Factor-3</a>
-	<a href="#">D12.776.744.631</a>	<a href="#">Interferon Regulatory Factor-7</a>

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.744.665 Paxillin
-	D12.776.744.700 Phosvitin
-	D12.776.744.736 Plectin
-	D12.776.744.741 Smad Proteins, Receptor-Regulated
-	D12.776.744.741.500 Smad1 Protein
-	D12.776.744.741.750 Smad2 Protein
-	D12.776.744.741.875 Smad3 Protein
-	D12.776.744.741.937 Smad5 Protein
-	D12.776.744.741.968 Smad8 Protein
-	D12.776.744.747 Retinoblastoma-Like Protein p107
-	D12.776.744.755 Retinoblastoma-Like Protein p130
-	D12.776.744.770 Retinoblastoma Protein
-	D12.776.744.772 Stathmin
-	D12.776.744.840 Synapsins
-	D12.776.744.845 Tumor Suppressor Protein p53
-	D12.776.744.925 Vitellogenins
-	D12.776.744.962 Zyxin
-	D12.776.752 Photoreceptors, Microbial
-	D12.776.752.249 Bacteriochlorophylls
-	D12.776.752.249.500 Bacteriochlorophyll A
-	D12.776.752.812 Rhodopsins, Microbial
-	D12.776.752.812.249 Bacteriorhodopsins
-	D12.776.752.812.500 Halorhodopsins
-	D12.776.752.812.750 Sensory Rhodopsins
-	D12.776.765 Plant Proteins
-	D12.776.765.149 Arabidopsis Proteins
-	D12.776.765.149.500 AGAMOUS Protein, Arabidopsis
-	D12.776.765.199 Chloroplast Proteins
-	D12.776.765.199.186 Chloroplast Thioredoxins
-	D12.776.765.199.249 Endopeptidase Clp
-	D12.776.765.199.499 Ribulose-Bisphosphate Carboxylase
-	D12.776.765.199.750 Thylakoid Membrane Proteins
-	D12.776.765.199.750.374 Chloroplast Proton-Translocating ATPases
-	D12.776.765.199.750.750 Photosynthetic Reaction Center Complex Proteins
-	D12.776.765.199.750.750.374 Cytochrome b6f Complex
-	D12.776.765.199.750.750.374.500 Cytochromes b6

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.765.199.750.750.374.750      Cytochromes f
-	D12.776.765.199.750.750.374.875      Plastoquinol-Plastocyanin Reductase
-	D12.776.765.199.750.750.490      Light-Harvesting Protein Complexes
-	D12.776.765.199.750.750.490.500      Chlorophyll Binding Proteins
-	D12.776.765.199.750.750.500      Photosystem I Protein Complex
-	D12.776.765.199.750.750.750      Photosystem II Protein Complex
-	D12.776.765.249      DEFICIENS Protein
-	D12.776.765.319      Ferredoxins
-	D12.776.765.342      Florigen
-	D12.776.765.365      G-Box Binding Factors
-	D12.776.765.500      Leghemoglobin
-	D12.776.765.537      Periplasmic Proteins
-	D12.776.765.593      Photoreceptors, Plant
-	D12.776.765.593.500      Cryptochromes
-	D12.776.765.593.750      Phototropins
-	D12.776.765.650      Phycocyanin
-	D12.776.765.665      Phycoerythrin
-	D12.776.765.675      Phytochrome
-	D12.776.765.675.249      Phytochrome A
-	D12.776.765.675.500      Phytochrome B
-	D12.776.765.678      Plant Lectins
-	D12.776.765.678.500      Concanavalin A
-	D12.776.765.678.625      Peanut Agglutinin
-	D12.776.765.678.750      Phytohemagglutinins
-	D12.776.765.678.875      Pokeweed Mitogens
-	D12.776.765.678.906      Ribosome Inactivating Proteins, Type 2
-	D12.776.765.678.906.111      Abrin
-	D12.776.765.678.906.750      Ricin
-	D12.776.765.678.968      Wheat Germ Agglutinins
-	D12.776.765.678.968.900      Wheat Germ Agglutinin-Horseradish Peroxidase Conjugate
-	D12.776.765.680      Plastocyanin
-	D12.776.765.710      Ribosome Inactivating Proteins
-	D12.776.765.710.500      Ribosome Inactivating Proteins, Type 1
-	D12.776.765.710.500.500      Trichosanthin
-	D12.776.765.710.750      Ribosome Inactivating Proteins, Type 2

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D12.776.765.725	Seed Storage Proteins
-	D12.776.765.725.249	2S Albumins, Plant
-	D12.776.765.725.500	Prolamins
-	D12.776.765.725.500.500	Glutens
-	D12.776.765.725.500.500.400	Gliadin
-	D12.776.765.725.500.750	Zein
-	D12.776.765.741	Soybean Proteins
-	D12.776.765.741.500	Trypsin Inhibitor, Bowman-Birk Soybean
-	D12.776.765.741.750	Trypsin Inhibitor, Kunitz Soybean
-	D12.776.765.836	Vegetable Proteins
-	D12.776.775	Polyproteins
-	D12.776.775.320	Gene Products, env
-	D12.776.775.330	Gene Products, gag
-	D12.776.775.330.300	Fusion Proteins, gag-pol
-	D12.776.775.330.650	gag Gene Products, Human Immunodeficiency Virus
-	D12.776.775.360	Gene Products, pol
-	D12.776.775.360.300	Fusion Proteins, gag-pol
-	D12.776.775.360.650	pol Gene Products, Human Immunodeficiency Virus
-	D12.776.780	Pregnancy Proteins
-	D12.776.780.400	Chorionic Gonadotropin
-	D12.776.780.400.125	Chorionic Gonadotropin, beta Subunit, Human
New Heading	<b>D12.776.780.426</b>	<b>Glycodelin</b>
-	D12.776.780.451	Gonadotropins, Equine
New Heading	<b>D12.776.780.551</b>	<b>Placenta Growth Factor</b>
-	D12.776.780.650	Placental Lactogen
-	D12.776.780.675	Pregnancy-Associated alpha 2-Macroglobulins
-	D12.776.780.700	Pregnancy-Associated Plasma Protein-A
-	D12.776.780.730	Pregnancy-Specific beta 1-Glycoproteins
-	D12.776.785	Prions
New Heading	<b>D12.776.785.340</b>	<b>Prion Proteins</b>
New Tree	<a href="#">D12.776.785.340.500</a>	<a href="#">PrPC Proteins</a>
New Tree	<a href="#">D12.776.785.340.750</a>	<a href="#">PrPSc Proteins</a>

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	<a href="#">D12.776.785.340.750.700</a>	<a href="#">PrP 27-30 Protein</a>
Old Tree	<a href="#">D12.776.785.680</a>	<a href="#">PrPC Proteins</a>
Old Tree	<a href="#">D12.776.785.700</a>	<a href="#">PrPSc Proteins</a>
Old Tree	<a href="#">D12.776.785.700.700</a>	<a href="#">PrP 27-30 Protein</a>
-	D12.776.796	Protein Hydrolysates
-	D12.776.800	Protein Isoforms
-	D12.776.800.300	Isoenzymes
-	D12.776.811	Protein Precursors
-	D12.776.811.050	Amyloid beta-Protein Precursor
-	D12.776.811.070	Angiotensinogen
-	D12.776.811.185	Chromogranins
-	D12.776.811.300	Fibrinogen
-	D12.776.811.300.290	Fibrin Fibrinogen Degradation Products
-	D12.776.811.300.310	Fibrinopeptide A
-	D12.776.811.300.320	Fibrinopeptide B
-	D12.776.811.420	Kininogens
-	D12.776.811.420.350	Kininogen, High-Molecular-Weight
-	D12.776.811.420.400	Kininogen, Low-Molecular-Weight
-	D12.776.811.690	Procollagen
-	D12.776.811.700	Proglucagon
-	D12.776.811.706	Proinsulin
-	D12.776.811.800	Pro-Opiomelanocortin
-	D12.776.811.850	Tropoelastin
-	D12.776.813	Protein Subunits
-	D12.776.816	Proteolipids
-	D12.776.816.249	Myelin and Lymphocyte-Associated Proteolipid Proteins
-	D12.776.816.500	Myelin Proteolipid Protein
-	D12.776.816.750	Pulmonary Surfactant-Associated Protein C
-	D12.776.817	Proteome
-	D12.776.820	Protozoan Proteins
New Heading	<a href="#">D12.776.820.250</a>	<a href="#">Discoidins</a>
-	D12.776.820.500	Merozoite Surface Protein 1
-	D12.776.823	Pulmonary Surfactant-Associated Proteins
-	D12.776.823.100	Pulmonary Surfactant-Associated Protein A

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D12.776.823.124	Pulmonary Surfactant-Associated Protein B
-	D12.776.823.186	Pulmonary Surfactant-Associated Protein C
-	D12.776.823.249	Pulmonary Surfactant-Associated Protein D
-	D12.776.826	Receptors, Cytoplasmic and Nuclear
-	D12.776.826.119	Hepatocyte Nuclear Factor 4
-	D12.776.826.179	Inositol 1,4,5-Trisphosphate Receptors
New Heading	<b>D12.776.826.194</b>	<b>Liver X Receptors</b>
-	D12.776.826.209	Orphan Nuclear Receptors
-	D12.776.826.209.039	COUP Transcription Factors
-	D12.776.826.209.039.249	COUP Transcription Factor I
-	D12.776.826.209.039.500	COUP Transcription Factor II
-	D12.776.826.209.059	DAX-1 Orphan Nuclear Receptor
-	D12.776.826.209.080	Nuclear Receptor Subfamily 1, Group D, Member 1
-	D12.776.826.209.100	Nuclear Receptor Subfamily 1, Group F, Member 1
-	D12.776.826.209.110	Nuclear Receptor Subfamily 1, Group F, Member 2
-	D12.776.826.209.182	Nuclear Receptor Subfamily 1, Group F, Member 3
-	D12.776.826.209.255	Nuclear Receptor Subfamily 2, Group C, Member 1
-	D12.776.826.209.327	Nuclear Receptor Subfamily 2, Group C, Member 2
-	D12.776.826.209.400	Nuclear Receptor Subfamily 4, Group A, Member 1
-	D12.776.826.209.415	Nuclear Receptor Subfamily 4, Group A, Member 2
-	D12.776.826.209.430	Nuclear Receptor Subfamily 4, Group A, Member 3
-	D12.776.826.209.572	Nuclear Receptor Subfamily 6, Group A, Member 1
-	D12.776.826.209.715	Receptors, Aryl Hydrocarbon
-	D12.776.826.239	Peroxisome Proliferator-Activated Receptors
-	D12.776.826.239.500	PPAR alpha
-	D12.776.826.239.530	PPAR-beta
-	D12.776.826.239.555	PPAR delta
-	D12.776.826.239.588	PPAR gamma
-	D12.776.826.387	Receptors, Artificial
-	D12.776.826.535	Receptors, Calcitriol
-	D12.776.826.590	Receptors, Melatonin
-	D12.776.826.701	Receptors, Retinoic Acid
New Heading	<b>D12.776.826.701.250</b>	<b>Retinoic Acid Receptor alpha</b>
-	D12.776.826.701.500	Retinoid X Receptors



## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.826.701.500.500      Retinoid X Receptor alpha
-	D12.776.826.701.500.625      Retinoid X Receptor beta
-	D12.776.826.701.500.750      Retinoid X Receptor gamma
-	D12.776.826.750                  Receptors, Steroid
-	D12.776.826.750.150              Receptors, Androgen
-	D12.776.826.750.350              Receptors, Estrogen
-	D12.776.826.750.350.174        Estrogen Receptor alpha
-	D12.776.826.750.350.262        Estrogen Receptor beta
-	D12.776.826.750.350.350        Receptors, Estradiol
-	D12.776.826.750.430              Receptors, Glucocorticoid
-	D12.776.826.750.530              Receptors, Mineralocorticoid
-	D12.776.826.750.765              Receptors, Progesterone
-	D12.776.826.850                  Receptors, Thyroid Hormone
-	D12.776.826.850.500              Thyroid Hormone Receptors alpha
-	D12.776.826.850.750              Thyroid Hormone Receptors beta
-	D12.776.826.925                  Steroidogenic Factor 1
-	D12.776.827                        Receptors, Drug
-	D12.776.827.137                    Imidazoline Receptors
-	D12.776.827.275                    Immunophilins
-	D12.776.827.275.300                Cyclophilins
-	D12.776.827.275.700                Tacrolimus Binding Proteins
-	D12.776.827.275.700.500        Tacrolimus Binding Protein 1A
-	D12.776.827.550                    Receptors, Phencyclidine
-	D12.776.827.775                    Sulfonylurea Receptors
-	D12.776.828                        Recombinant Proteins
-	D12.776.828.300                    Recombinant Fusion Proteins
-	D12.776.828.300.200                CD4 Immunoadhesins
-	D12.776.828.868                    Vaccines, Synthetic
-	D12.776.828.868.910                Vaccines, DNA
-	D12.776.828.868.915                Vaccines, Edible
-	D12.776.828.868.940                Vaccines, Virosome
-	D12.776.828.868.955                Vaccines, Virus-Like Particle
-	D12.776.831                        Reptilian Proteins
-	D12.776.831.222                    Cobra Cardiotoxin Proteins
-	D12.776.831.244                    Cobra Neurotoxin Proteins
-	D12.776.835                        Ribosomal Proteins

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.835.700 Peptide Elongation Factors
-	D12.776.835.700.350 GTP Phosphohydrolase-Linked Elongation Factors
-	D12.776.835.700.350.101 Peptide Elongation Factor 1
-	D12.776.835.700.350.102 Peptide Elongation Factor 2
-	D12.776.835.700.350.200 Peptide Elongation Factor G
-	D12.776.835.700.350.700 Peptide Elongation Factor Tu
-	D12.776.835.725 Peptide Initiation Factors
-	D12.776.835.725.868 Eukaryotic Initiation Factors
-	D12.776.835.725.868.124 Eukaryotic Initiation Factor-1
-	D12.776.835.725.868.249 Eukaryotic Initiation Factor-2
-	D12.776.835.725.868.374 Eukaryotic Initiation Factor-2B
-	D12.776.835.725.868.437 Eukaryotic Initiation Factor-3
-	D12.776.835.725.868.500 Eukaryotic Initiation Factor-4F
-	D12.776.835.725.868.500.500 Eukaryotic Initiation Factor-4A
-	D12.776.835.725.868.500.750 Eukaryotic Initiation Factor-4E
-	D12.776.835.725.868.500.875 Eukaryotic Initiation Factor-4G
-	D12.776.835.725.868.750 Eukaryotic Initiation Factor-5
-	D12.776.835.725.934 Prokaryotic Initiation Factors
-	D12.776.835.725.934.374 Prokaryotic Initiation Factor-1
-	D12.776.835.725.934.562 Prokaryotic Initiation Factor-2
-	D12.776.835.725.934.750 Prokaryotic Initiation Factor-3
-	D12.776.835.862 Peptide Termination Factors
-	D12.776.835.931 Ribosomal Protein S6
-	D12.776.850 Salivary Proteins and Peptides
-	D12.776.850.400 Glue Proteins, Drosophila
-	D12.776.850.550 Histatins
-	D12.776.850.700 Lipocalin 1
-	D12.776.850.850 Mucin-5B
-	D12.776.850.887 Salivary alpha-Amylases
-	D12.776.850.906 Salivary Cystatins
-	D12.776.850.925 Salivary Proline-Rich Proteins
-	D12.776.860 Scleroproteins
-	D12.776.860.300 Extracellular Matrix Proteins
-	D12.776.860.300.030 Activated-Leukocyte Cell Adhesion Molecule
New Heading	<b>D12.776.860.300.085 ADAMTS Proteins</b>

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Heading	<b>D12.776.860.300.085.500</b>	<b>ADAMTS1 Protein</b>
New Heading	<b>D12.776.860.300.085.813</b>	<b>ADAMTS13 Protein</b>
New Heading	<b>D12.776.860.300.085.844</b>	<b>ADAMTS4 Protein</b>
New Heading	<b>D12.776.860.300.085.875</b>	<b>ADAMTS5 Protein</b>
New Heading	<b>D12.776.860.300.085.937</b>	<b>ADAMTS7 Protein</b>
New Heading	<b>D12.776.860.300.085.968</b>	<b>ADAMTS9 Protein</b>
-	D12.776.860.300.140	Aggrecans
-	D12.776.860.300.155	Cartilage Oligomeric Matrix Protein
-	D12.776.860.300.200	CCN Intercellular Signaling Proteins
-	D12.776.860.300.200.100	Connective Tissue Growth Factor
-	D12.776.860.300.200.200	Cysteine-Rich Protein 61
-	D12.776.860.300.200.500	Nephroblastoma Overexpressed Protein
-	D12.776.860.300.250	Collagen
-	D12.776.860.300.250.300	Fibrillar Collagens
-	D12.776.860.300.250.300.100	Collagen Type I
-	D12.776.860.300.250.300.200	Collagen Type II
-	D12.776.860.300.250.300.300	Collagen Type III
-	D12.776.860.300.250.300.400	Collagen Type V
-	D12.776.860.300.250.300.500	Collagen Type XI
-	D12.776.860.300.250.400	Non-Fibrillar Collagens
-	D12.776.860.300.250.400.100	Collagen Type IV
-	D12.776.860.300.250.400.200	Collagen Type VI
-	D12.776.860.300.250.400.300	Collagen Type VII
-	D12.776.860.300.250.400.400	Collagen Type VIII
-	D12.776.860.300.250.400.500	Collagen Type X
-	D12.776.860.300.250.400.525	Collagen Type XIII
-	D12.776.860.300.250.400.537	Collagen Type XVIII
-	D12.776.860.300.250.400.537.500	Endostatins
-	D12.776.860.300.250.400.550	Fibril-Associated Collagens
-	D12.776.860.300.250.400.550.200	Collagen Type IX
-	D12.776.860.300.250.400.550.300	Collagen Type XII
-	D12.776.860.300.250.600	Procollagen

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D12.776.860.300.250.700	Tropocollagen
-	D12.776.860.300.350	Elastin
-	D12.776.860.300.350.700	Tropoelastin
New Heading	<b>D12.776.860.300.400</b>	<b>Fibrillins</b>
New Heading	<b>D12.776.860.300.400.500</b>	<b>Fibrillin-1</b>
New Heading	<b>D12.776.860.300.400.750</b>	<b>Fibrillin-2</b>
-	D12.776.860.300.450	Fibronectins
-	D12.776.860.300.562	Integrin-Binding Sialoprotein
-	D12.776.860.300.675	Laminin
New Tree	<a href="#">D12.776.860.300.688</a>	<a href="#">Latent TGF-beta Binding Proteins</a>
-	D12.776.860.300.700	Matrilin Proteins
-	D12.776.860.300.762	Osteopontin
New Heading	<b>D12.776.860.300.806</b>	<b>Small Leucine-Rich Proteoglycans</b>
New Tree	<a href="#">D12.776.860.300.806.500</a>	<a href="#">Biglycan</a>
New Tree	<a href="#">D12.776.860.300.806.750</a>	<a href="#">Decorin</a>
New Heading	<b>D12.776.860.300.806.813</b>	<b>Fibromodulin</b>
New Heading	<b>D12.776.860.300.806.875</b>	<b>Lumican</b>
-	D12.776.860.300.850	Tenascin
-	D12.776.860.300.885	Versicans
-	D12.776.860.300.920	Vitronectin
-	D12.776.860.476	Gelatin
-	D12.776.860.607	Keratins
-	D12.776.860.607.074	beta-Keratins
-	D12.776.860.607.149	Keratins, Hair-Specific
-	D12.776.860.607.300	Keratins, Type I
-	D12.776.860.607.300.100	Keratin-10
-	D12.776.860.607.300.200	Keratin-12
-	D12.776.860.607.300.300	Keratin-13
-	D12.776.860.607.300.400	Keratin-14
-	D12.776.860.607.300.600	Keratin-16

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.860.607.300.700 Keratin-17
-	D12.776.860.607.300.800 Keratin-18
-	D12.776.860.607.300.900 Keratin-19
-	D12.776.860.607.300.950 Keratin-20
-	D12.776.860.607.650 Keratins, Type II
-	D12.776.860.607.650.100 Keratin-1
-	D12.776.860.607.650.200 Keratin-2
-	D12.776.860.607.650.300 Keratin-3
-	D12.776.860.607.650.400 Keratin-4
-	D12.776.860.607.650.500 Keratin-5
-	D12.776.860.607.650.600 Keratin-6
-	D12.776.860.607.650.700 Keratin-7
-	D12.776.860.607.650.800 Keratin-8
-	D12.776.860.607.650.900 Keratin-9
-	D12.776.860.823 Reticulin
-	D12.776.861 Secretoglobins
-	D12.776.861.249 Mammaglobin A
-	D12.776.861.260 Mammaglobin B
-	D12.776.861.311 Prostatein
-	D12.776.861.500 Uteroglobin
-	D12.776.863 Selenium-Binding Proteins
-	D12.776.864 Selenoproteins
-	D12.776.864.100 Methionine Sulfoxide Reductases
-	D12.776.864.124 Selenoprotein P
-	D12.776.864.500 Selenoprotein W
-	D12.776.866 Seminal Plasma Proteins
-	D12.776.866.249 Prostatic Secretory Proteins
-	D12.776.866.249.500 Prostate-Specific Antigen
-	D12.776.866.500 Seminal Vesicle Secretory Proteins
-	D12.776.872 Serpins
-	D12.776.872.020 Angiotensinogen
-	D12.776.872.030 alpha 1-Antichymotrypsin
-	D12.776.872.035 alpha 1-Antitrypsin
-	D12.776.872.050 alpha-2-Antiplasmin
-	D12.776.872.060 Antithrombin Proteins
-	D12.776.872.060.500 Antithrombin III

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D12.776.872.060.750	Heparin Cofactor II
-	D12.776.872.060.875	Hirudins
-	D12.776.872.140	Complement C1 Inactivator Proteins
-	D12.776.872.140.500	Complement C1 Inhibitor Protein
-	D12.776.872.418	HSP47 Heat-Shock Proteins
-	D12.776.872.557	Ovalbumin
-	D12.776.872.695	Plasminogen Inactivators
-	D12.776.872.695.500	Plasminogen Activator Inhibitor 1
-	D12.776.872.695.520	Plasminogen Activator Inhibitor 2
-	D12.776.872.695.700	Protein C Inhibitor
-	D12.776.872.695.850	Serpin E2
-	D12.776.872.847	Thyroxine-Binding Globulin
-	D12.776.872.923	Transcortin
-	D12.776.884	Silver Proteins
-	D12.776.915	Thioredoxins
-	D12.776.915.249	Chloroplast Thioredoxins
-	D12.776.915.624	Thioredoxin h
-	D12.776.922	Thymosin
-	D12.776.930	Transcription Factors
-	D12.776.930.100	Adenovirus E1A Proteins
-	D12.776.930.110	Adenovirus E1B Proteins
-	D12.776.930.120	Antennapedia Homeodomain Protein
-	D12.776.930.125	Basic Helix-Loop-Helix Transcription Factors
-	D12.776.930.125.249	ARNTL Transcription Factors
-	D12.776.930.125.500	Basic Helix-Loop-Helix Leucine Zipper Transcription Factors
-	D12.776.930.125.500.500	Microphthalmia-Associated Transcription Factor
New Heading	<b>D12.776.930.125.500.563</b>	<b>N-Myc Proto-Oncogene Protein</b>
New Tree	<b>D12.776.930.125.500.625</b>	<b>Nuclear Receptor Coactivator 1</b>
-	D12.776.930.125.500.750	Sterol Regulatory Element Binding Proteins
-	D12.776.930.125.500.750.500	Sterol Regulatory Element Binding Protein 1
-	D12.776.930.125.500.750.750	Sterol Regulatory Element Binding Protein 2
-	D12.776.930.125.500.875	Transcription Factor 3
-	D12.776.930.125.562	CLOCK Proteins
-	D12.776.930.125.625	Hypoxia-Inducible Factor 1

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D12.776.930.125.625.500	Aryl Hydrocarbon Receptor Nuclear Translocator
-	D12.776.930.125.625.750	Hypoxia-Inducible Factor 1, alpha Subunit
-	D12.776.930.125.750	Myogenic Regulatory Factors
-	D12.776.930.125.750.294	MEF2 Transcription Factors
-	D12.776.930.125.750.590	MyoD Protein
-	D12.776.930.125.750.595	Myogenic Regulatory Factor 5
-	D12.776.930.125.750.600	Myogenin
New Tree	<a href="#">D12.776.930.125.813</a>	<a href="#">Proto-Oncogene Proteins c-myc</a>
New Heading	<b>D12.776.930.125.844</b>	<b>Transcription Factor HES-1</b>
Old Tree	<del>D12.776.930.125.875</del>	<del>Twist Transcription Factor</del>
Old Tree	<del>D12.776.930.125.875</del>	<del>Twist-Related Protein 1</del>
New Heading	<b>D12.776.930.125.906</b>	<b>Twist Transcription Factors</b>
New Tree	<a href="#">D12.776.930.125.906.250</a>	<a href="#">Twist Transcription Factor</a>
New Tree	<a href="#">D12.776.930.125.906.250</a>	<a href="#">Twist-Related Protein 1</a>
New Heading	<b>D12.776.930.125.906.500</b>	<b>Twist-Related Protein 2</b>
-	D12.776.930.125.937	Upstream Stimulatory Factors
-	D12.776.930.127	Basic-Leucine Zipper Transcription Factors
-	D12.776.930.127.061	Activating Transcription Factors
-	D12.776.930.127.061.500	Activating Transcription Factor 1
-	D12.776.930.127.061.750	Activating Transcription Factor 2
-	D12.776.930.127.061.875	Activating Transcription Factor 3
-	D12.776.930.127.061.937	Activating Transcription Factor 4
-	D12.776.930.127.061.968	Activating Transcription Factor 6
-	D12.776.930.127.092	Basic Helix-Loop-Helix Leucine Zipper Transcription Factors
-	D12.776.930.127.092.500	Microphthalmia-Associated Transcription Factor
New Heading	<b>D12.776.930.127.092.625</b>	<b>N-Myc Proto-Oncogene Protein</b>
-	D12.776.930.127.092.750	Sterol Regulatory Element Binding Proteins
-	D12.776.930.127.092.750.500	Sterol Regulatory Element Binding Protein 1
-	D12.776.930.127.092.750.750	Sterol Regulatory Element Binding Protein 2
-	D12.776.930.127.092.875	Transcription Factor 3
-	D12.776.930.127.124	CCAAT-Enhancer-Binding Proteins

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D12.776.930.127.124.249	CCAAT-Binding Factor
-	D12.776.930.127.124.500	CCAAT-Enhancer-Binding Protein-alpha
-	D12.776.930.127.124.750	CCAAT-Enhancer-Binding Protein-beta
-	D12.776.930.127.124.812	CCAAT-Enhancer-Binding Protein-delta
-	D12.776.930.127.124.875	Transcription Factor CHOP
-	D12.776.930.127.124.937	Y-Box-Binding Protein 1
-	D12.776.930.127.184	Cyclic AMP Response Element-Binding Protein
-	D12.776.930.127.186	Cyclic AMP Response Element-Binding Protein A
-	D12.776.930.127.217	Cyclic AMP Response Element Modulator
-	D12.776.930.127.311	Fos-Related Antigen-2
-	D12.776.930.127.374	Interferon Regulatory Factor-1
-	D12.776.930.127.500	Maf Transcription Factors
-	D12.776.930.127.500.061	Maf Transcription Factors, Large
-	D12.776.930.127.500.061.500	MafB Transcription Factor
-	D12.776.930.127.500.061.625	Oncogene Protein v-maf
-	D12.776.930.127.500.061.750	Proto-Oncogene Proteins c-maf
-	D12.776.930.127.500.500	Maf Transcription Factors, Small
-	D12.776.930.127.500.500.249	MafF Transcription Factor
-	D12.776.930.127.500.500.500	MafG Transcription Factor
-	D12.776.930.127.500.500.750	MafK Transcription Factor
-	D12.776.930.127.656	NF-E2 Transcription Factor
-	D12.776.930.127.656.750	Maf Transcription Factors, Small
-	D12.776.930.127.656.750.249	MafF Transcription Factor
-	D12.776.930.127.656.750.500	MafG Transcription Factor
-	D12.776.930.127.656.750.750	MafK Transcription Factor
-	D12.776.930.127.656.770	NF-E2 Transcription Factor, p45 Subunit
-	D12.776.930.127.710	NF-E2-Related Factor 1
-	D12.776.930.127.737	NF-E2-Related Factor 2
-	D12.776.930.127.765	Proto-Oncogene Proteins c-fos
-	D12.776.930.127.820	Proto-Oncogene Proteins c-jun
-	D12.776.930.127.875	Transcription Factor AP-1
New Heading	<b>D12.776.930.127.937</b>	<b>X-Box Binding Protein 1</b>
-	D12.776.930.130	beta Catenin
-	D12.776.930.137	BRCA1 Protein
New	<b>D12.776.930.146</b>	<b>CDX2 Transcription Factor</b>



## MeSH Tree Changes for 2017

Type	Tree - heading
Heading	
-	D12.776.930.155 Core Binding Factors
-	D12.776.930.155.200 Core Binding Factor alpha Subunits
-	D12.776.930.155.200.100 Core Binding Factor Alpha 1 Subunit
-	D12.776.930.155.200.200 Core Binding Factor Alpha 2 Subunit
-	D12.776.930.155.200.300 Core Binding Factor Alpha 3 Subunit
-	D12.776.930.155.400 Core Binding Factor beta Subunit
-	D12.776.930.160 COUP Transcription Factors
-	D12.776.930.165 Cyclic AMP Receptor Protein
New Tree	<a href="#">D12.776.930.211</a> E2F Transcription Factors
New Tree	<a href="#">D12.776.930.211.500</a> E2F1 Transcription Factor
New Tree	<a href="#">D12.776.930.211.625</a> E2F2 Transcription Factor
New Tree	<a href="#">D12.776.930.211.750</a> E2F3 Transcription Factor
New Tree	<a href="#">D12.776.930.211.812</a> E2F4 Transcription Factor
New Tree	<a href="#">D12.776.930.211.875</a> E2F5 Transcription Factor
New Tree	<a href="#">D12.776.930.211.937</a> E2F6 Transcription Factor
New Tree	<a href="#">D12.776.930.211.968</a> E2F7 Transcription Factor
New Tree	<a href="#">D12.776.930.213</a> Early Growth Response Transcription Factors
New Tree	<a href="#">D12.776.930.213.500</a> Early Growth Response Protein 1
New Tree	<a href="#">D12.776.930.213.750</a> Early Growth Response Protein 2
New Tree	<a href="#">D12.776.930.213.875</a> Early Growth Response Protein 3
New Tree	<a href="#">D12.776.930.216</a> Erythroid-Specific DNA-Binding Factors
New Tree	<a href="#">D12.776.930.216.500</a> GATA1 Transcription Factor
New Tree	<a href="#">D12.776.930.216.625</a> GATA2 Transcription Factor
New Tree	<a href="#">D12.776.930.216.687</a> GATA3 Transcription Factor
New	<a href="#">D12.776.930.216.750</a> NF-E2 Transcription Factor

## MeSH Tree Changes for 2017

Type	Tree - heading	
Tree		
New Tree	D12.776.930.216.750.750	Maf Transcription Factors, Small
New Tree	D12.776.930.216.750.750.249	MafF Transcription Factor
New Tree	D12.776.930.216.750.750.500	MafG Transcription Factor
New Tree	D12.776.930.216.750.750.750	MafK Transcription Factor
New Tree	D12.776.930.216.750.770	NF-E2 Transcription Factor, p45 Subunit
New Tree	D12.776.930.216.875	YY1 Transcription Factor
Old Tree	D12.776.930.311	E2F Transcription Factors
Old Tree	D12.776.930.311.500	E2F1 Transcription Factor
Old Tree	D12.776.930.311.625	E2F2 Transcription Factor
Old Tree	D12.776.930.311.750	E2F3 Transcription Factor
Old Tree	D12.776.930.311.812	E2F4 Transcription Factor
Old Tree	D12.776.930.311.875	E2F5 Transcription Factor
Old Tree	D12.776.930.311.937	E2F6 Transcription Factor
Old Tree	D12.776.930.311.968	E2F7 Transcription Factor
Old Tree	D12.776.930.313	Early Growth Response Transcription Factors
Old Tree	D12.776.930.313.500	Early Growth Response Protein 1
Old Tree	D12.776.930.313.750	Early Growth Response Protein 2
Old Tree	D12.776.930.313.875	Early Growth Response Protein 3
Old Tree	D12.776.930.316	Erythroid-Specific DNA-Binding Factors
Old Tree	D12.776.930.316.500	GATA1 Transcription Factor
Old Tree	D12.776.930.316.625	GATA2 Transcription Factor
Old Tree	D12.776.930.316.687	GATA3 Transcription Factor
Old Tree	D12.776.930.316.750	NF-E2 Transcription Factor
Old Tree	D12.776.930.316.750.750	Maf Transcription Factors, Small
Old Tree	D12.776.930.316.750.750.249	MafF Transcription Factor
Old Tree	D12.776.930.316.750.750.500	MafG Transcription Factor
Old Tree	D12.776.930.316.750.750.750	MafK Transcription Factor
Old Tree	D12.776.930.316.750.770	NF-E2 Transcription Factor, p45 Subunit
Old Tree	D12.776.930.316.875	YY1 Transcription Factor
-	D12.776.930.318	Fushi Tarazu Transcription Factors
-	D12.776.930.319	G-Box Binding Factors

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.930.321 GATA Transcription Factors
-	D12.776.930.321.100 GATA1 Transcription Factor
-	D12.776.930.321.200 GATA2 Transcription Factor
-	D12.776.930.321.300 GATA3 Transcription Factor
-	D12.776.930.321.400 GATA4 Transcription Factor
-	D12.776.930.321.500 GATA5 Transcription Factor
-	D12.776.930.321.600 GATA6 Transcription Factor
-	D12.776.930.322 Goosecoid Protein
-	D12.776.930.323 Hepatocyte Nuclear Factors
-	D12.776.930.323.500 Hepatocyte Nuclear Factor 1
-	D12.776.930.323.500.500 Hepatocyte Nuclear Factor 1-alpha
-	D12.776.930.323.500.750 Hepatocyte Nuclear Factor 1-beta
-	D12.776.930.323.750 Hepatocyte Nuclear Factor 3-alpha
-	D12.776.930.323.875 Hepatocyte Nuclear Factor 3-beta
-	D12.776.930.323.937 Hepatocyte Nuclear Factor 3-gamma
-	D12.776.930.323.968 Hepatocyte Nuclear Factor 4
-	D12.776.930.323.984 Hepatocyte Nuclear Factor 6
New Heading	<b>D12.776.930.324 Homeobox Protein Nkx-2.5</b>
-	D12.776.930.325 Host Cell Factor C1
-	D12.776.930.326 I-kappa B Proteins
New Heading	<b>D12.776.930.326.500 NF-KappaB Inhibitor alpha</b>
-	D12.776.930.327 Immunoglobulin J Recombination Signal Sequence-Binding Protein
-	D12.776.930.329 Inhibitor of Differentiation Proteins
-	D12.776.930.329.249 Inhibitor of Differentiation Protein 1
-	D12.776.930.329.500 Inhibitor of Differentiation Protein 2
-	D12.776.930.332 Interferon Regulatory Factors
-	D12.776.930.332.124 Interferon Regulatory Factor-1
-	D12.776.930.332.249 Interferon Regulatory Factor-2
-	D12.776.930.332.374 Interferon Regulatory Factor-3
-	D12.776.930.332.437 Interferon Regulatory Factor-7
-	D12.776.930.332.500 Interferon-Stimulated Gene Factor 3, gamma Subunit
-	D12.776.930.354 Interferon-Stimulated Gene Factor 3
-	D12.776.930.354.249 Interferon-Stimulated Gene Factor 3, alpha Subunit
-	D12.776.930.354.249.500 STAT1 Transcription Factor

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.930.354.249.750                      STAT2 Transcription Factor
-	D12.776.930.354.500                      Interferon-Stimulated Gene Factor 3, gamma Subunit
-	D12.776.930.375                              Kruppel-Like Transcription Factors
-	D12.776.930.375.500                      Ikaros Transcription Factor
-	D12.776.930.375.750                      Sp Transcription Factors
-	D12.776.930.375.750.249                      Sp1 Transcription Factor
-	D12.776.930.375.750.500                      Sp2 Transcription Factor
-	D12.776.930.375.750.750                      Sp3 Transcription Factor
-	D12.776.930.375.750.875                      Sp4 Transcription Factor
-	D12.776.930.386                              Leucine-Responsive Regulatory Protein
-	D12.776.930.397                              MADS Domain Proteins
-	D12.776.930.397.249                      AGAMOUS Protein, Arabidopsis
-	D12.776.930.397.374                      DEFICIENS Protein
-	D12.776.930.397.700                      MEF2 Transcription Factors
-	D12.776.930.397.750                      Minichromosome Maintenance 1 Protein
-	D12.776.930.397.875                      Serum Response Factor
-	D12.776.930.440                              MSX1 Transcription Factor
-	D12.776.930.483                              Myeloid-Lymphoid Leukemia Protein
New Heading	<b>D12.776.930.542                      Nanog Homeobox Protein</b>
-	D12.776.930.600                              NF-kappa B
-	D12.776.930.600.124                      NF-kappa B p50 Subunit
-	D12.776.930.600.186                      NF-kappa B p52 Subunit
-	D12.776.930.600.249                      Transcription Factor RelA
-	D12.776.930.600.500                      Transcription Factor RelB
-	D12.776.930.608                              NFATC Transcription Factors
-	D12.776.930.612                              NFI Transcription Factors
-	D12.776.930.616                              Nuclear Factor 90 Proteins
-	D12.776.930.616.500                      Nuclear Factor 45 Protein
-	D12.776.930.617                              Nuclear Receptor Coactivators
-	D12.776.930.617.049                      Mediator Complex Subunit 1
-	D12.776.930.617.100                      Nuclear Receptor Coactivator 1
-	D12.776.930.617.200                      Nuclear Receptor Coactivator 2
-	D12.776.930.617.300                      Nuclear Receptor Coactivator 3
New Heading	<b>D12.776.930.617.650                      Peroxisome Proliferator-Activated Receptor Gamma Coactivator 1-alpha</b>

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D12.776.930.618	Nuclear Respiratory Factors
-	D12.776.930.618.249	GA-Binding Protein Transcription Factor
-	D12.776.930.618.500	Nuclear Respiratory Factor 1
-	D12.776.930.640	Onecut Transcription Factors
-	D12.776.930.640.500	Hepatocyte Nuclear Factor 6
-	D12.776.930.645	Orphan Nuclear Receptors
-	D12.776.930.650	Otx Transcription Factors
-	D12.776.930.680	p300-CBP Transcription Factors
New Tree	<a href="#">D12.776.930.680.300</a>	<a href="#">CREB-Binding Protein</a>
-	D12.776.930.680.600	E1A-Associated p300 Protein
-	D12.776.930.700	Paired Box Transcription Factors
-	D12.776.930.700.500	B-Cell-Specific Activator Protein
-	D12.776.930.700.500	PAX5 Transcription Factor
-	D12.776.930.700.750	PAX2 Transcription Factor
New Heading	<b>D12.776.930.700.782</b>	<b>PAX3 Transcription Factor</b>
New Heading	<b>D12.776.930.700.813</b>	<b>PAX6 Transcription Factor</b>
-	D12.776.930.700.875	PAX7 Transcription Factor
New Heading	<b>D12.776.930.700.906</b>	<b>PAX8 Transcription Factor</b>
-	D12.776.930.700.937	PAX9 Transcription Factor
-	D12.776.930.705	Peroxisome Proliferator-Activated Receptors
-	D12.776.930.710	POU Domain Factors
-	D12.776.930.710.500	Octamer Transcription Factors
-	D12.776.930.710.500.100	Octamer Transcription Factor-1
-	D12.776.930.710.500.200	Octamer Transcription Factor-2
-	D12.776.930.710.500.300	Octamer Transcription Factor-3
-	D12.776.930.710.500.600	Octamer Transcription Factor-6
-	D12.776.930.710.625	Transcription Factor Brn-3
-	D12.776.930.710.625.500	Transcription Factor Brn-3A
-	D12.776.930.710.625.750	Transcription Factor Brn-3B
-	D12.776.930.710.625.875	Transcription Factor Brn-3C
-	D12.776.930.710.750	Transcription Factor Pit-1
New Heading	<b>D12.776.930.713</b>	<b>Promyelocytic Leukemia Protein</b>

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D12.776.930.715	Proto-Oncogene Proteins c-bcl-6
-	D12.776.930.720	Proto-Oncogene Proteins c-ets
-	D12.776.930.720.100	Proto-Oncogene Protein c-ets-1
-	D12.776.930.720.200	Proto-Oncogene Protein c-ets-2
-	D12.776.930.720.400	Proto-Oncogene Protein c-fli-1
-	D12.776.930.720.600	Ternary Complex Factors
-	D12.776.930.720.600.100	ets-Domain Protein Elk-1
-	D12.776.930.720.600.300	ets-Domain Protein Elk-4
-	D12.776.930.725	Proto-Oncogene Proteins c-myb
-	D12.776.930.730	Proto-Oncogene Proteins c-rel
-	D12.776.930.760	Receptors, Aryl Hydrocarbon
-	D12.776.930.770	Receptors, Notch
-	D12.776.930.770.500	Receptor, Notch1
-	D12.776.930.770.750	Receptor, Notch2
New Heading	<b>D12.776.930.770.875</b>	<b>Receptor, Notch3</b>
-	D12.776.930.775	Receptors, Retinoic Acid
New Heading	<b>D12.776.930.775.250</b>	<b>Retinoic Acid Receptor alpha</b>
-	D12.776.930.775.500	Retinoid X Receptors
-	D12.776.930.778	Receptors, Steroid
-	D12.776.930.778.350	Receptors, Estrogen
-	D12.776.930.780	Repressor Proteins
-	D12.776.930.780.500	AraC Transcription Factor
-	D12.776.930.780.625	Co-Repressor Proteins
-	D12.776.930.780.625.186	DAX-1 Orphan Nuclear Receptor
-	D12.776.930.780.625.374	Nuclear Receptor Co-Repressor 1
-	D12.776.930.780.625.400	Nuclear Receptor Co-Repressor 2
-	D12.776.930.780.625.750 Complex	Sin3 Histone Deacetylase and Corepressor
-	D12.776.930.780.750	COUP Transcription Factor II
-	D12.776.930.780.781	Lac Repressors
-	D12.776.930.780.890	Polycomb-Group Proteins
-	D12.776.930.780.890.100	Polycomb Repressive Complex 1
-	D12.776.930.780.890.200	Polycomb Repressive Complex 2
New Heading	<b>D12.776.930.780.890.200.250</b>	<b>Enhancer of Zeste Homolog 2 Protein</b>

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D12.776.930.780.890.200.500	Retinoblastoma-Binding Protein 4
-	D12.776.930.780.890.200.750	Retinoblastoma-Binding Protein 7
New Heading	<b>D12.776.930.780.945</b>	<b>Zinc Finger E-box-Binding Homeobox 1</b>
-	D12.776.930.785	Rho Factor
-	D12.776.930.800	Sigma Factor
-	D12.776.930.806	Smad Proteins
-	D12.776.930.806.500	Smad Proteins, Receptor-Regulated
-	D12.776.930.806.500.100	Smad1 Protein
-	D12.776.930.806.500.200	Smad2 Protein
-	D12.776.930.806.500.300	Smad3 Protein
-	D12.776.930.806.500.500	Smad5 Protein
-	D12.776.930.806.500.800	Smad8 Protein
-	D12.776.930.806.750	Smad4 Protein
-	D12.776.930.806.875	Smad6 Protein
New Heading	<b>D12.776.930.809</b>	<b>SMARCB1 Protein</b>
New Heading	<b>D12.776.930.815</b>	<b>Snail Family Transcription Factors</b>
-	D12.776.930.823	SOX Transcription Factors
-	D12.776.930.823.049	Sex-Determining Region Y Protein
-	D12.776.930.823.100	SOXB1 Transcription Factors
-	D12.776.930.823.200	SOXB2 Transcription Factors
-	D12.776.930.823.300	SOXC Transcription Factors
-	D12.776.930.823.400	SOXD Transcription Factors
-	D12.776.930.823.500	SOXE Transcription Factors
-	D12.776.930.823.500.500	SOX9 Transcription Factor
-	D12.776.930.823.600	SOXF Transcription Factors
-	D12.776.930.840	STAT Transcription Factors
-	D12.776.930.840.100	STAT1 Transcription Factor
-	D12.776.930.840.200	STAT2 Transcription Factor
-	D12.776.930.840.300	STAT3 Transcription Factor
-	D12.776.930.840.400	STAT4 Transcription Factor
-	D12.776.930.840.500	STAT5 Transcription Factor
-	D12.776.930.840.600	STAT6 Transcription Factor
-	D12.776.930.850	T-Box Domain Proteins
-	D12.776.930.875	TCF Transcription Factors

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D12.776.930.875.500	Lymphoid Enhancer-Binding Factor 1
-	D12.776.930.875.750	T Cell Transcription Factor 1
-	D12.776.930.875.812	Transcription Factor 7-Like 1 Protein
-	D12.776.930.875.875	Transcription Factor 7-Like 2 Protein
-	D12.776.930.900	Trans-Activators
-	D12.776.930.900.199	Gene Products, tat
-	D12.776.930.900.199.500	tat Gene Products, Human Immunodeficiency Virus
-	D12.776.930.900.400	Herpes Simplex Virus Protein Vmw65
New Heading	<b>D12.776.930.900.550</b>	<b>Transcription Activator-Like Effectors</b>
New Heading	<b>D12.776.930.900.625</b>	<b>Transcriptional Regulator ERG</b>
New Heading	<b>D12.776.930.900.700</b>	<b>Zinc Finger Protein GLI1</b>
-	D12.776.930.907	Transcription Factor AP-2
-	D12.776.930.908	Transcription Factor DP1
-	D12.776.930.930	Transcription Factors, General
-	D12.776.930.930.092	PoI1 Transcription Initiation Complex Proteins
-	D12.776.930.930.186	TATA-Binding Protein Associated Factors
-	D12.776.930.930.374	TATA-Box Binding Protein
-	D12.776.930.930.812	TATA Box Binding Protein-Like Proteins
-	D12.776.930.930.875	Transcription Factors, TFII
-	D12.776.930.930.875.374	Transcription Factor TFIIA
-	D12.776.930.930.875.562	Transcription Factor TFIIIB
-	D12.776.930.930.875.750	Transcription Factor TFIIID
-	D12.776.930.930.875.750.500	TATA-Box Binding Protein
-	D12.776.930.930.875.875	Transcription Factor TFIIH
-	D12.776.930.930.875.875.500	Xeroderma Pigmentosum Group D Protein
-	D12.776.930.930.937	Transcription Factors, TFIII
-	D12.776.930.930.937.249	Transcription Factor TFIIIA
-	D12.776.930.930.937.500	Transcription Factor TFIIIB
-	D12.776.930.955	Transcriptional Elongation Factors
-	D12.776.930.955.500	Positive Transcriptional Elongation Factor B
-	D12.776.930.955.500.249	Cyclin T
-	D12.776.930.955.500.500	Cyclin-Dependent Kinase 9
-	D12.776.930.960	Tristetraprolin
New	<b>D12.776.930.969</b>	<b>Tumor Protein p73</b>



## MeSH Tree Changes for 2017

Type	Tree - heading	
Heading		
-	D12.776.930.977	Winged-Helix Transcription Factors
-	D12.776.930.977.249	Forkhead Transcription Factors
New Heading	<b>D12.776.930.977.249.063</b>	<b>Forkhead Box Protein M1</b>
New Heading	<b>D12.776.930.977.249.125</b>	<b>Forkhead Box Protein O3</b>
New Heading	<b>D12.776.930.977.249.250</b>	<b>Forkhead Box Protein O1</b>
-	D12.776.930.977.249.500	Hepatocyte Nuclear Factor 3-alpha
-	D12.776.930.977.249.750	Hepatocyte Nuclear Factor 3-beta
-	D12.776.930.977.249.875	Hepatocyte Nuclear Factor 3-gamma
New Heading	<b>D12.776.930.977.624</b>	<b>Regulatory Factor X Transcription Factors</b>
New Heading	<b>D12.776.930.977.624.500</b>	<b>Regulatory Factor X1</b>
New Heading	<b>D12.776.934</b>	<b>Tripartite Motif Proteins</b>
New Heading	<b>D12.776.934.500</b>	<b>Promyelocytic Leukemia Protein</b>
New Heading	<b>D12.776.934.750</b>	<b>Pyrin</b>
-	D12.776.938	Ubiquitinated Proteins
-	D12.776.947	Ubiquitins
New Heading	<b>D12.776.947.125</b>	<b>Autophagy-Related Protein 12</b>
New Heading	<b>D12.776.947.187</b>	<b>Autophagy-Related Protein 8 Family</b>
-	D12.776.947.249	Small Ubiquitin-Related Modifier Proteins
-	D12.776.947.249.500	SUMO-1 Protein
-	D12.776.947.500	Ubiquitin
-	D12.776.947.500.500	Polyubiquitin
-	D12.776.947.500.750	Ubiquitin C
-	D12.776.964	Viral Proteins
-	D12.776.964.700	Oncogene Proteins, Viral
-	D12.776.964.700.045	Adenovirus Early Proteins
-	D12.776.964.700.045.050	Adenovirus E1 Proteins
-	D12.776.964.700.045.050.100	Adenovirus E1A Proteins
-	D12.776.964.700.045.050.110	Adenovirus E1B Proteins

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.964.700.045.060      Adenovirus E2 Proteins
-	D12.776.964.700.045.070      Adenovirus E3 Proteins
-	D12.776.964.700.045.080      Adenovirus E4 Proteins
-	D12.776.964.700.090      Antigens, Polyomavirus Transforming
-	D12.776.964.700.750      Retroviridae Proteins, Oncogenic
-	D12.776.964.700.750.320      Fusion Proteins, gag-onc
-	D12.776.964.700.750.320.700      Oncogene Protein p65(gag-jun)
-	D12.776.964.700.750.470      Gene Products, rex
-	D12.776.964.700.750.480      Gene Products, tax
-	D12.776.964.700.750.650      Oncogene Protein gp140(v-fms)
-	D12.776.964.700.750.710      Oncogene Protein p21(ras)
-	D12.776.964.700.750.750      Oncogene Protein p55(v-myc)
-	D12.776.964.700.750.760      Oncogene Protein pp60(v-src)
-	D12.776.964.700.750.817      Oncogene Protein v-maf
-	D12.776.964.700.750.875      Oncogene Proteins v-abl
-	D12.776.964.700.750.882      Oncogene Proteins v-erbA
-	D12.776.964.700.750.883      Oncogene Proteins v-erbB
-	D12.776.964.700.750.887      Oncogene Proteins v-fos
-	D12.776.964.700.750.900      Oncogene Proteins v-mos
-	D12.776.964.700.750.903      Oncogene Proteins v-myb
-	D12.776.964.700.750.920      Oncogene Proteins v-raf
-	D12.776.964.700.750.925      Oncogene Proteins v-rel
-	D12.776.964.700.750.935      Oncogene Proteins v-sis
-	D12.776.964.775      Retroviridae Proteins
-	D12.776.964.775.325      Gene Products, env
-	D12.776.964.775.325.164      env Gene Products, Human Immunodeficiency Virus
-	D12.776.964.775.325.164.200      HIV Envelope Protein gp41
-	D12.776.964.775.325.164.249      HIV Envelope Protein gp120
-	D12.776.964.775.325.164.374      HIV Envelope Protein gp160
-	D12.776.964.775.350      Gene Products, gag
-	D12.776.964.775.350.320      Fusion Proteins, gag-onc
-	D12.776.964.775.350.320.700      Oncogene Protein p65(gag-jun)
-	D12.776.964.775.350.325      Fusion Proteins, gag-pol
-	D12.776.964.775.350.362      gag Gene Products, Human Immunodeficiency Virus
-	D12.776.964.775.350.362.500      HIV Core Protein p24
-	D12.776.964.775.362      Gene Products, nef

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.964.775.362.500                      nef Gene Products, Human Immunodeficiency Virus
-	D12.776.964.775.375                              Gene Products, pol
-	D12.776.964.775.375.325                      Fusion Proteins, gag-pol
-	D12.776.964.775.375.545                      pol Gene Products, Human Immunodeficiency Virus
-	D12.776.964.775.375.545.500                      HIV Integrase
-	D12.776.964.775.375.545.750                      HIV Protease
-	D12.776.964.775.375.545.875                      HIV Reverse Transcriptase
-	D12.776.964.775.375.545.875.500                      Ribonuclease H, Human Immunodeficiency Virus
-	D12.776.964.775.375.750                              RNA-Directed DNA Polymerase
-	D12.776.964.775.375.750.187                              HIV Reverse Transcriptase
-	D12.776.964.775.468                              Gene Products, vif
-	D12.776.964.775.468.500                              vif Gene Products, Human Immunodeficiency Virus
-	D12.776.964.775.562                              Human Immunodeficiency Virus Proteins
-	D12.776.964.775.562.500                              env Gene Products, Human Immunodeficiency Virus
-	D12.776.964.775.562.500.200                              HIV Envelope Protein gp41
-	D12.776.964.775.562.500.500                              HIV Envelope Protein gp120
-	D12.776.964.775.562.500.750                              HIV Envelope Protein gp160
-	D12.776.964.775.562.750                              gag Gene Products, Human Immunodeficiency Virus
-	D12.776.964.775.562.750.500                              HIV Core Protein p24
-	D12.776.964.775.562.760                              nef Gene Products, Human Immunodeficiency Virus
-	D12.776.964.775.562.764                              pol Gene Products, Human Immunodeficiency Virus
-	D12.776.964.775.562.764.500                              HIV Integrase
-	D12.776.964.775.562.764.875                              HIV Reverse Transcriptase
-	D12.776.964.775.562.764.875.500                              Ribonuclease H, Human Immunodeficiency Virus
-	D12.776.964.775.562.770                              rev Gene Products, Human Immunodeficiency Virus
-	D12.776.964.775.562.773                              tat Gene Products, Human Immunodeficiency Virus
-	D12.776.964.775.562.781                              vif Gene Products, Human Immunodeficiency Virus
-	D12.776.964.775.562.937                              vpr Gene Products, Human Immunodeficiency Virus
-	D12.776.964.775.750                              Retroviridae Proteins, Oncogenic
-	D12.776.964.775.750.320                              Fusion Proteins, gag-onc
-	D12.776.964.775.750.320.700                              Oncogene Protein p65(gag-jun)
-	D12.776.964.775.750.470                              Gene Products, rex
-	D12.776.964.775.750.480                              Gene Products, tax
-	D12.776.964.775.750.650                              Oncogene Protein gp140(v-fms)

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.964.775.750.710 Oncogene Protein p21(ras)
-	D12.776.964.775.750.750 Oncogene Protein p55(v-myc)
-	D12.776.964.775.750.760 Oncogene Protein pp60(v-src)
-	D12.776.964.775.750.817 Oncogene Protein v-maf
-	D12.776.964.775.750.875 Oncogene Proteins v-abl
-	D12.776.964.775.750.882 Oncogene Proteins v-erbA
-	D12.776.964.775.750.883 Oncogene Proteins v-erbB
-	D12.776.964.775.750.887 Oncogene Proteins v-fos
-	D12.776.964.775.750.900 Oncogene Proteins v-mos
-	D12.776.964.775.750.903 Oncogene Proteins v-myb
-	D12.776.964.775.750.920 Oncogene Proteins v-raf
-	D12.776.964.775.750.925 Oncogene Proteins v-rel
-	D12.776.964.775.750.935 Oncogene Proteins v-sis
-	D12.776.964.900 Viral Nonstructural Proteins
-	D12.776.964.925 Viral Regulatory and Accessory Proteins
-	D12.776.964.925.500 Gene Products, nef
-	D12.776.964.925.500.500 nef Gene Products, Human Immunodeficiency Virus
-	D12.776.964.925.750 Gene Products, rex
-	D12.776.964.925.875 Gene Products, vif
-	D12.776.964.925.875.500 vif Gene Products, Human Immunodeficiency Virus
-	D12.776.964.925.968 Immediate-Early Proteins
-	D12.776.964.925.984 Trans-Activators
-	D12.776.964.925.984.385 Gene Products, rev
-	D12.776.964.925.984.385.500 rev Gene Products, Human Immunodeficiency Virus
-	D12.776.964.925.984.400 Gene Products, tat
-	D12.776.964.925.984.400.500 tat Gene Products, Human Immunodeficiency Virus
-	D12.776.964.925.984.410 Gene Products, tax
-	D12.776.964.925.984.430 Gene Products, vpr
-	D12.776.964.925.984.430.500 vpr Gene Products, Human Immunodeficiency Virus
-	D12.776.964.925.984.445 Herpes Simplex Virus Protein Vmw65
-	D12.776.964.970 Viral Structural Proteins
-	D12.776.964.970.600 Nucleocapsid Proteins
-	D12.776.964.970.600.550 Capsid Proteins
-	D12.776.964.970.600.850 Viral Core Proteins

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D12.776.964.970.600.850.350 Gene Products, gag
-	D12.776.964.970.600.850.350.325 Fusion Proteins, gag-pol
-	D12.776.964.970.600.850.350.362 gag Gene Products, Human Immunodeficiency Virus
-	D12.776.964.970.600.850.350.362.500 HIV Core Protein p24
-	D12.776.964.970.600.850.375 Gene Products, pol
-	D12.776.964.970.600.850.375.325 Fusion Proteins, gag-pol
-	D12.776.964.970.600.850.375.545 pol Gene Products, Human Immunodeficiency Virus
-	D12.776.964.970.600.850.375.545.500 HIV Integrase
-	D12.776.964.970.600.850.375.545.875 HIV Reverse Transcriptase
-	D12.776.964.970.600.850.375.545.875.500 Ribonuclease H, Human Immunodeficiency Virus
-	D12.776.964.970.600.850.375.750 RNA-Directed DNA Polymerase
-	D12.776.964.970.600.850.375.750.187 HIV Reverse Transcriptase
-	D12.776.964.970.600.850.375.750.187.500 Ribonuclease H, Human Immunodeficiency Virus
-	D12.776.964.970.700 Plant Viral Movement Proteins
-	D12.776.964.970.880 Viral Envelope Proteins
-	D12.776.964.970.880.325 Gene Products, env
-	D12.776.964.970.880.325.164 env Gene Products, Human Immunodeficiency Virus
-	D12.776.964.970.880.325.164.200 HIV Envelope Protein gp41
-	D12.776.964.970.880.325.164.249 HIV Envelope Protein gp120
-	D12.776.964.970.880.325.164.374 HIV Envelope Protein gp160
-	D12.776.964.970.880.345 Hemagglutinins, Viral
-	D12.776.964.970.880.345.500 Hemagglutinin Glycoproteins, Influenza Virus
-	D12.776.964.970.880.350 HN Protein
-	D12.776.964.970.880.910 Viral Fusion Proteins
-	D12.776.964.970.880.910.330 HIV Envelope Protein gp41
-	D12.776.964.970.880.910.665 Spike Glycoprotein, Coronavirus
-	D12.776.964.970.880.940 Viral Matrix Proteins
-	D12.776.964.970.910 Viral Tail Proteins
-	D13 Nucleic Acids, Nucleotides, and Nucleosides
-	D13.150 Antisense Elements (Genetics)
-	D13.150.200 DNA, Antisense
-	D13.150.200.640 Oligodeoxyribonucleotides, Antisense

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D13.150.480	Oligonucleotides, Antisense
-	D13.150.480.640	Oligodeoxyribonucleotides, Antisense
-	D13.150.480.645	Oligoribonucleotides, Antisense
-	D13.150.650	RNA, Antisense
-	D13.150.650.319	MicroRNAs
-	D13.150.650.640	Oligoribonucleotides, Antisense
-	D13.150.650.700	RNA, Small Interfering
-	D13.400	Nucleic Acid Precursors
-	D13.400.730	RNA Precursors
-	D13.444	Nucleic Acids
-	D13.444.308	DNA
New Heading	<b>D13.444.308.065</b>	<b>DNA, Ancient</b>
-	D13.444.308.130	DNA, A-Form
-	D13.444.308.135	DNA Adducts
-	D13.444.308.148	DNA, Algal
-	D13.444.308.150	DNA, Antisense
-	D13.444.308.150.640	Oligodeoxyribonucleotides, Antisense
-	D13.444.308.180	DNA, Archaeal
-	D13.444.308.196	DNA, B-Form
-	D13.444.308.212	DNA, Bacterial
-	D13.444.308.227	DNA, C-Form
-	D13.444.308.243	DNA, Catalytic
-	D13.444.308.283	DNA, Circular
-	D13.444.308.283.084	DNA, Catenated
-	D13.444.308.283.170	DNA, Chloroplast
-	D13.444.308.283.225	DNA, Mitochondrial
-	D13.444.308.283.225.200	DNA, Kinetoplast
-	D13.444.308.283.250	DNA, Superhelical
-	D13.444.308.291	DNA, Concatenated
-	D13.444.308.295	DNA, Cruciform
-	D13.444.308.300	DNA, Fungal
-	D13.444.308.315	DNA, Helminth
-	D13.444.308.324	DNA, Intergenic
-	D13.444.308.324.230	DNA, Ribosomal Spacer
-	D13.444.308.425	DNA, Neoplasm

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D13.444.308.435 DNA, Plant
-	D13.444.308.435.275 DNA, Chloroplast
-	D13.444.308.442 DNA, Protozoan
-	D13.444.308.442.200 DNA, Kinetoplast
-	D13.444.308.460 DNA, Recombinant
-	D13.444.308.475 DNA, Ribosomal
-	D13.444.308.475.230 DNA, Ribosomal Spacer
-	D13.444.308.480 DNA, Satellite
-	D13.444.308.497 DNA, Single-Stranded
-	D13.444.308.497.220 DNA, Complementary
-	D13.444.308.520 DNA Transposable Elements
-	D13.444.308.568 DNA, Viral
-	D13.444.308.574 DNA, Z-Form
-	D13.444.308.580 Isochores
-	D13.444.308.760 Retroelements
-	D13.444.404 Immobilized Nucleic Acids
-	D13.444.500 Nucleic Acid Heteroduplexes
-	D13.444.600 Nucleic Acid Probes
-	D13.444.600.150 Antisense Elements (Genetics)
-	D13.444.600.150.200 DNA, Antisense
-	D13.444.600.150.200.640 Oligodeoxyribonucleotides, Antisense
-	D13.444.600.150.640 Oligonucleotides, Antisense
-	D13.444.600.150.640.640 Oligodeoxyribonucleotides, Antisense
-	D13.444.600.150.640.645 Oligoribonucleotides, Antisense
-	D13.444.600.150.760 RNA, Antisense
-	D13.444.600.150.760.640 Oligoribonucleotides, Antisense
-	D13.444.600.223 DNA Probes
-	D13.444.600.223.500 DNA, Complementary
-	D13.444.600.223.550 DNA Probes, HLA
-	D13.444.600.223.555 DNA Probes, HPV
-	D13.444.600.601 Oligonucleotide Probes
-	D13.444.600.723 RNA Probes
-	D13.444.600.723.480 RNA, Complementary
-	D13.444.735 RNA
-	D13.444.735.064 RNA Isoforms
-	D13.444.735.130 RNA, Algal

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D13.444.735.150 RNA, Antisense
-	D13.444.735.150.319 MicroRNAs
-	D13.444.735.150.640 Oligoribonucleotides, Antisense
-	D13.444.735.150.700 RNA, Small Interfering
-	D13.444.735.300 RNA, Archaeal
-	D13.444.735.473 RNA, Bacterial
-	D13.444.735.480 RNA, Complementary
-	D13.444.735.490 RNA, Double-Stranded
-	D13.444.735.500 RNA, Fungal
-	D13.444.735.520 RNA, Helminth
-	D13.444.735.544 RNA, Messenger
-	D13.444.735.544.355 Codon
-	D13.444.735.544.355.225 Codon, Initiator
-	D13.444.735.544.355.250 Codon, Terminator
-	D13.444.735.544.355.250.235 Codon, Nonsense
-	D13.444.735.544.427 Riboswitch
-	D13.444.735.544.500 RNA Caps
-	D13.444.735.544.500.710 RNA Cap Analogs
-	D13.444.735.544.527 RNA, Messenger, Stored
-	D13.444.735.544.550 RNA Splice Sites
-	D13.444.735.544.875 Untranslated Regions
-	D13.444.735.544.875.880 3' Untranslated Regions
-	D13.444.735.544.875.885 5' Untranslated Regions
-	D13.444.735.615 RNA, Neoplasm
-	D13.444.735.628 RNA, Nuclear
-	D13.444.735.628.806 RNA, Heterogeneous Nuclear
-	D13.444.735.628.818 RNA, Small Nuclear
-	D13.444.735.628.818.800 RNA, Small Nucleolar
-	D13.444.735.635 RNA, Plant
-	D13.444.735.635.575 RNA, Chloroplast
-	D13.444.735.640 RNA Precursors
-	D13.444.735.650 RNA, Protozoan
-	D13.444.735.686 RNA, Ribosomal
-	D13.444.735.686.650 RNA, Ribosomal, 5S
-	D13.444.735.686.660 RNA, Ribosomal, 5.8S
-	D13.444.735.686.670 RNA, Ribosomal, 16S



## MeSH Tree Changes for 2017

Type	Tree - heading
-	D13.444.735.686.675 RNA, Ribosomal, 18S
-	D13.444.735.686.680 RNA, Ribosomal, 23S
-	D13.444.735.686.690 RNA, Ribosomal, 28S
-	D13.444.735.686.845 RNA, Ribosomal, Self-Splicing
-	D13.444.735.721 RNA, Satellite
-	D13.444.735.721.250 Cucumber Mosaic Virus Satellite
-	D13.444.735.757 RNA, Transfer
-	D13.444.735.757.286 Anticodon
-	D13.444.735.757.700 RNA, Transfer, Amino Acid-Specific
-	D13.444.735.757.700.050 RNA, Transfer, Ala
-	D13.444.735.757.700.075 RNA, Transfer, Arg
-	D13.444.735.757.700.085 RNA, Transfer, Asn
-	D13.444.735.757.700.090 RNA, Transfer, Asp
-	D13.444.735.757.700.200 RNA, Transfer, Cys
-	D13.444.735.757.700.400 RNA, Transfer, Gln
-	D13.444.735.757.700.410 RNA, Transfer, Glu
-	D13.444.735.757.700.420 RNA, Transfer, Gly
-	D13.444.735.757.700.450 RNA, Transfer, His
-	D13.444.735.757.700.480 RNA, Transfer, Ile
-	D13.444.735.757.700.500 RNA, Transfer, Leu
-	D13.444.735.757.700.510 RNA, Transfer, Lys
-	D13.444.735.757.700.525 RNA, Transfer, Met
-	D13.444.735.757.700.650 RNA, Transfer, Phe
-	D13.444.735.757.700.660 RNA, Transfer, Pro
-	D13.444.735.757.700.700 RNA, Transfer, Ser
-	D13.444.735.757.700.725 RNA, Transfer, Thr
-	D13.444.735.757.700.740 RNA, Transfer, Trp
-	D13.444.735.757.700.750 RNA, Transfer, Tyr
-	D13.444.735.757.700.900 RNA, Transfer, Val
-	D13.444.735.757.715 RNA, Transfer, Amino Acyl
-	D13.444.735.790 RNA, Untranslated
-	D13.444.735.790.199 RNA, Catalytic
-	D13.444.735.790.375 RNA, Long Noncoding
-	D13.444.735.790.552 RNA, Small Untranslated
-	D13.444.735.790.552.500 MicroRNAs
-	D13.444.735.790.552.625 RNA, Guide

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D13.444.735.790.552.750 RNA, Small Cytoplasmic
-	D13.444.735.790.552.875 RNA, Small Interfering
-	D13.444.735.790.552.937 RNA, Small Nuclear
-	D13.444.735.790.552.937.800 RNA, Small Nucleolar
-	D13.444.735.790.552.968 RNA, Spliced Leader
-	D13.444.735.790.878 Untranslated Regions
-	D13.444.735.790.878.880 3' Untranslated Regions
-	D13.444.735.790.878.885 5' Untranslated Regions
-	D13.444.735.790.878.942 Regulatory Sequences, Ribonucleic Acid
-	D13.444.735.790.878.942.250 Internal Ribosome Entry Sites
-	D13.444.735.790.878.942.500 Riboswitch
-	D13.444.735.828 RNA, Viral
-	D13.570 Nucleosides
-	D13.570.065 Arabinonucleosides
-	D13.570.065.090 Arabinofuranosyluracil
-	D13.570.065.300 Cytarabine
-	D13.570.065.300.050 Ancitabine
-	D13.570.065.950 Vidarabine
-	D13.570.230 Deoxyribonucleosides
-	D13.570.230.229 Deoxyadenosines
-	D13.570.230.229.075 Cladribine
-	D13.570.230.229.105 Dideoxyadenosine
-	D13.570.230.229.650 Puromycin Aminonucleoside
-	D13.570.230.329 Deoxycytidine
-	D13.570.230.329.100 Bromodeoxycytidine
-	D13.570.230.329.313 Capecitabine
-	D13.570.230.329.525 Emtricitabine
-	D13.570.230.329.525.250 Elvitegravir, Cobicistat, Emtricitabine, Tenofovir Disoproxil Fumarate Drug Combination
-	D13.570.230.329.525.500 Emtricitabine, Tenofovir Disoproxil Fumarate Drug Combination
-	D13.570.230.329.950 Zalcitabine
-	D13.570.230.329.950.500 Lamivudine
-	D13.570.230.360 Deoxyguanosine
-	D13.570.230.430 Deoxyuridine
-	D13.570.230.430.196 Bromodeoxyuridine

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D13.570.230.430.432                      Floxuridine
-	D13.570.230.430.609                      Idoxuridine
-	D13.570.230.500                            Dideoxynucleosides
-	D13.570.230.500.090                      Didanosine
-	D13.570.230.500.105                      Dideoxyadenosine
-	D13.570.230.500.850                      Stavudine
-	D13.570.230.500.925                      Zalcitabine
-	D13.570.230.500.925.500                      Lamivudine
-	D13.570.230.500.950                      Zidovudine
-	D13.570.230.677                            Pentostatin
-	D13.570.230.855                            Thymidine
-	D13.570.230.855.875                      Stavudine
-	D13.570.230.855.900                      Trifluridine
-	D13.570.230.855.950                      Zidovudine
-	D13.570.583                                Purine Nucleosides
-	D13.570.583.138                            Adenosine
-	D13.570.583.138.025                      Adenosine-5'-(N-ethylcarboxamide)
-	D13.570.583.138.240                      S-Adenosylhomocysteine
-	D13.570.583.138.264                      S-Adenosylmethionine
-	D13.570.583.138.300                      2-Chloroadenosine
-	D13.570.583.138.300.200                      Cladribine
-	D13.570.583.138.325                      Deoxyadenosines
-	D13.570.583.138.325.075                      Cladribine
-	D13.570.583.138.325.105                      Dideoxyadenosine
-	D13.570.583.138.325.800                      Puromycin Aminonucleoside
-	D13.570.583.138.500                      Isopentenyladenosine
-	D13.570.583.138.630                      Phenylisopropyladenosine
-	D13.570.583.138.711                      Puromycin
-	D13.570.583.138.711.650                      Puromycin Aminonucleoside
-	D13.570.583.138.900                      Vidarabine
-	D13.570.583.454                            Guanosine
-	D13.570.583.454.240                      Deoxyguanosine
-	D13.570.583.454.500                      Nucleoside Q
-	D13.570.583.616                            Inosine
-	D13.570.583.616.130                      Didanosine
-	D13.570.583.616.450                      Inosine Pranobex

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D13.570.583.616.900 Thioinosine
-	D13.570.583.616.900.500 Methylthioinosine
-	D13.570.583.910 Tubercidin
-	D13.570.685 Pyrimidine Nucleosides
-	D13.570.685.245 Cytidine
-	D13.570.685.245.217 Azacitidine
-	D13.570.685.245.453 Cytarabine
-	D13.570.685.245.453.050 Ancitabine
-	D13.570.685.245.500 Deoxycytidine
-	D13.570.685.245.500.250 Bromodeoxycytidine
-	D13.570.685.245.500.425 Capecitabine
-	D13.570.685.245.500.600 Emtricitabine
-	D13.570.685.245.500.600.250 Elvitegravir, Cobicistat, Emtricitabine, Tenofovir Disoproxil Fumarate Drug Combination
-	D13.570.685.245.500.600.500 Emtricitabine, Tenofovir Disoproxil Fumarate Drug Combination
-	D13.570.685.245.500.950 Zalcitabine
-	D13.570.685.245.500.950.500 Lamivudine
-	D13.570.685.350 Formycins
-	D13.570.685.350.200 Coformycin
-	D13.570.685.350.200.700 Pentostatin
-	D13.570.685.705 Thymidine
-	D13.570.685.705.875 Stavudine
-	D13.570.685.705.900 Trifluridine
-	D13.570.685.705.950 Zidovudine
-	D13.570.685.725 Tunicamycin
-	D13.570.685.852 Uridine
-	D13.570.685.852.150 Arabinofuranosyluracil
-	D13.570.685.852.176 Azauridine
-	D13.570.685.852.250 3-Deazauridine
-	D13.570.685.852.300 Deoxyuridine
-	D13.570.685.852.300.150 Bromodeoxyuridine
-	D13.570.685.852.300.350 Floxuridine
-	D13.570.685.852.300.400 Idoxuridine
-	D13.570.685.852.628 Pseudouridine
-	D13.570.685.852.800 Tetrahydrouridine

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D13.570.685.852.829 Thiouridine
-	D13.570.800 Ribonucleosides
-	D13.570.800.096 Adenosine
-	D13.570.800.096.250 Adenosine-5'-(N-ethylcarboxamide)
-	D13.570.800.096.262 S-Adenosylhomocysteine
-	D13.570.800.096.264 S-Adenosylmethionine
-	D13.570.800.096.300 2-Chloroadenosine
-	D13.570.800.096.300.200 Cladribine
-	D13.570.800.096.500 Isopentenyladenosine
-	D13.570.800.096.630 Phenylisopropyladenosine
-	D13.570.800.286 Cytidine
-	D13.570.800.286.300 Azacitidine
-	D13.570.800.330 Dichlororibofuranosylbenzimidazole
-	D13.570.800.410 Formycins
-	D13.570.800.410.200 Coformycin
-	D13.570.800.453 Guanosine
-	D13.570.800.453.500 Nucleoside Q
-	D13.570.800.573 Inosine
-	D13.570.800.573.130 Didanosine
-	D13.570.800.573.450 Inosine Pranobex
-	D13.570.800.573.900 Thioinosine
-	D13.570.800.573.900.500 Methylthioinosine
-	D13.570.800.790 Ribavirin
-	D13.570.800.810 Showdomycin
-	D13.570.800.840 Toyocamycin
-	D13.570.800.850 Tubercidin
-	D13.570.800.892 Uridine
-	D13.570.800.892.176 Azauridine
-	D13.570.800.892.250 3-Deazauridine
-	D13.570.800.892.628 Pseudouridine
-	D13.570.800.892.800 Tetrahydrouridine
-	D13.570.800.892.829 Thiouridine
-	D13.570.900 Thionucleosides
-	D13.570.900.111 Azathioprine
-	D13.570.900.800 Thioinosine
-	D13.570.900.800.500 Methylthioinosine

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D13.570.900.829 Thiouridine
-	D13.695 Nucleotides
-	D13.695.065 Arabinonucleotides
-	D13.695.065.200 Arabinofuranosylcytosine Triphosphate
-	D13.695.065.900 Vidarabine Phosphate
-	D13.695.201 Deoxyribonucleotides
-	D13.695.201.100 Deoxyadenine Nucleotides
-	D13.695.201.150 Deoxycytosine Nucleotides
-	D13.695.201.150.200 Deoxycytidine Monophosphate
-	D13.695.201.175 Deoxyguanine Nucleotides
-	D13.695.201.200 Deoxyuracil Nucleotides
-	D13.695.201.200.270 Fluorodeoxyuridylate
-	D13.695.201.486 Nucleoside Diphosphate Sugars
-	D13.695.201.789 Thymine Nucleotides
-	D13.695.201.789.788 Thymidine Monophosphate
-	D13.695.225 Dideoxynucleotides
-	D13.695.250 Dinucleoside Phosphates
-	D13.695.462 Nucleotides, Cyclic
-	D13.695.462.200 Cyclic AMP
-	D13.695.462.200.225 8-Bromo Cyclic Adenosine Monophosphate
-	D13.695.462.200.250 Bucladesine
-	D13.695.462.250 Cyclic CMP
-	D13.695.462.275 Cyclic GMP
-	D13.695.462.275.325 Dibutyryl Cyclic GMP
-	D13.695.462.300 Cyclic IMP
-	D13.695.578 Polynucleotides
-	D13.695.578.424 Oligonucleotides
New Heading	<b>D13.695.578.424.112 Antagomirs</b>
-	D13.695.578.424.224 Aptamers, Nucleotide
-	D13.695.578.424.337 Morpholinos
-	D13.695.578.424.450 Oligodeoxyribonucleotides
-	D13.695.578.424.450.275 DNA Primers
-	D13.695.578.424.480 Oligonucleotides, Antisense
-	D13.695.578.424.480.640 Oligodeoxyribonucleotides, Antisense
-	D13.695.578.424.480.645 Oligoribonucleotides, Antisense

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D13.695.578.424.500 Oligoribonucleotides
-	D13.695.578.424.550 Peptide Nucleic Acids
-	D13.695.578.424.575 Phosphorothioate Oligonucleotides
-	D13.695.578.424.600 Pyrimidine Dimers
-	D13.695.578.500 Polydeoxyribonucleotides
-	D13.695.578.500.050 Apurinic Acid
-	D13.695.578.500.300 Poly dA-dT
-	D13.695.578.500.600 Poly T
-	D13.695.578.550 Polyribonucleotides
-	D13.695.578.550.050 Apurinic Acid
-	D13.695.578.550.500 Poly A
-	D13.695.578.550.500.510 Poly A-U
-	D13.695.578.550.530 Poly Adenosine Diphosphate Ribose
-	D13.695.578.550.560 Poly C
-	D13.695.578.550.560.600 Poly I-C
-	D13.695.578.550.600 Poly G
-	D13.695.578.550.650 Poly I
-	D13.695.578.550.650.600 Poly I-C
-	D13.695.578.550.750 Poly U
-	D13.695.578.550.750.510 Poly A-U
-	D13.695.667 Purine Nucleotides
-	D13.695.667.138 Adenine Nucleotides
-	D13.695.667.138.124 Adenosine Diphosphate
-	D13.695.667.138.124.070 Adenosine Diphosphate Sugars
-	D13.695.667.138.124.070.075 Adenosine Diphosphate Glucose
-	D13.695.667.138.124.070.125 Adenosine Diphosphate Ribose
-	D13.695.667.138.124.070.125.040 O-Acetyl-ADP-Ribose
-	D13.695.667.138.124.070.125.195 Cyclic ADP-Ribose
-	D13.695.667.138.180 Adenosine Monophosphate
-	D13.695.667.138.180.080 Adenosine Phosphosulfate
-	D13.695.667.138.236 Adenosine Triphosphate
-	D13.695.667.138.236.050 Adenylyl Imidodiphosphate
-	D13.695.667.138.236.250 Ethenoadenosine Triphosphate
-	D13.695.667.138.382 Coenzyme A
-	D13.695.667.138.382.300 Acyl Coenzyme A
-	D13.695.667.138.382.300.020 Acetyl Coenzyme A

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D13.695.667.138.382.300.500 Malonyl Coenzyme A
-	D13.695.667.138.382.300.700 Palmitoyl Coenzyme A
-	D13.695.667.138.395 Cyclic AMP
-	D13.695.667.138.395.225 8-Bromo Cyclic Adenosine Monophosphate
-	D13.695.667.138.395.250 Bucladesine
-	D13.695.667.138.410 Deoxyadenine Nucleotides
-	D13.695.667.138.506 Flavin-Adenine Dinucleotide
-	D13.695.667.138.694 NAD
-	D13.695.667.138.749 NADP
-	D13.695.667.138.850 Phosphoadenosine Phosphosulfate
-	D13.695.667.138.925 Vidarabine Phosphate
-	D13.695.667.454 Guanine Nucleotides
-	D13.695.667.454.160 Cyclic GMP
-	D13.695.667.454.160.325 Dibutyryl Cyclic GMP
-	D13.695.667.454.200 Deoxyguanine Nucleotides
-	D13.695.667.454.340 Guanosine Diphosphate
-	D13.695.667.454.340.350 Guanosine Diphosphate Sugars
-	D13.695.667.454.340.350.400 Guanosine Diphosphate Fucose
-	D13.695.667.454.340.350.500 Guanosine Diphosphate Mannose
-	D13.695.667.454.440 Guanosine Pentaphosphate
-	D13.695.667.454.480 Guanosine Tetraphosphate
-	D13.695.667.454.504 Guanosine Triphosphate
-	D13.695.667.454.504.380 Guanosine 5'-O-(3-Thiotriphosphate)
-	D13.695.667.454.504.400 Guanylyl Imidodiphosphate
-	D13.695.667.454.525 Guanosine Monophosphate
-	D13.695.667.454.700 RNA Caps
-	D13.695.667.454.700.710 RNA Cap Analogs
-	D13.695.667.616 Inosine Nucleotides
-	D13.695.667.616.300 Cyclic IMP
-	D13.695.667.616.400 Inosine Diphosphate
-	D13.695.667.616.500 Inosine Monophosphate
-	D13.695.667.616.800 Inosine Triphosphate
-	D13.695.740 Pyrimidine Nucleotides
-	D13.695.740.050 Apurinic Acid
-	D13.695.740.246 Cytosine Nucleotides
-	D13.695.740.246.050 Arabinofuranosylcytosine Triphosphate



## MeSH Tree Changes for 2017

Type	Tree - heading
-	D13.695.740.246.115      Cyclic CMP
-	D13.695.740.246.150      Cytidine Diphosphate
-	D13.695.740.246.150.180      Cytidine Diphosphate Choline
-	D13.695.740.246.150.210      Cytidine Diphosphate Diglycerides
-	D13.695.740.246.370      Cytidine Monophosphate
-	D13.695.740.246.370.250      Cytidine Monophosphate N-Acetylneuraminic Acid
-	D13.695.740.246.400      Cytidine Triphosphate
-	D13.695.740.246.425      Deoxycytosine Nucleotides
-	D13.695.740.246.425.300      Deoxycytidine Monophosphate
-	D13.695.740.600      Pyrimidine Dimers
-	D13.695.740.706      Thymine Nucleotides
-	D13.695.740.706.788      Thymidine Monophosphate
-	D13.695.740.850      Uracil Nucleotides
-	D13.695.740.850.210      Deoxyuracil Nucleotides
-	D13.695.740.850.210.200      Fluorodeoxyuridylate
-	D13.695.740.850.600      Uridine Diphosphate
-	D13.695.740.850.600.677      Uridine Diphosphate Sugars
-	D13.695.740.850.600.677.100      Uridine Diphosphate N-Acetylgalactosamine
-	D13.695.740.850.600.677.120      Uridine Diphosphate N-Acetylglucosamine
-	D13.695.740.850.600.677.150      Uridine Diphosphate N-Acetylmuramic Acid
-	D13.695.740.850.600.677.300      Uridine Diphosphate Galactose
-	D13.695.740.850.600.677.350      Uridine Diphosphate Glucose
-	D13.695.740.850.600.677.375      Uridine Diphosphate Glucuronic Acid
-	D13.695.740.850.600.677.800      Uridine Diphosphate Xylose
-	D13.695.740.850.877      Uridine Monophosphate
-	D13.695.740.850.877.500      Sofosbuvir
-	D13.695.740.850.950      Uridine Triphosphate
-	D13.695.827      Ribonucleotides
-	D13.695.827.068      Adenine Nucleotides
-	D13.695.827.068.124      Adenosine Diphosphate
-	D13.695.827.068.124.070      Adenosine Diphosphate Sugars
-	D13.695.827.068.124.070.075      Adenosine Diphosphate Glucose
-	D13.695.827.068.124.070.125      Adenosine Diphosphate Ribose
-	D13.695.827.068.124.070.125.040      O-Acetyl-ADP-Ribose
-	D13.695.827.068.124.070.125.195      Cyclic ADP-Ribose
-	D13.695.827.068.180      Adenosine Monophosphate

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D13.695.827.068.180.080 Adenosine Phosphosulfate
-	D13.695.827.068.236 Adenosine Triphosphate
-	D13.695.827.068.236.050 Adenylyl Imidodiphosphate
-	D13.695.827.068.236.250 Ethenoadenosine Triphosphate
-	D13.695.827.068.382 Coenzyme A
-	D13.695.827.068.382.300 Acyl Coenzyme A
-	D13.695.827.068.382.300.020 Acetyl Coenzyme A
-	D13.695.827.068.382.300.500 Malonyl Coenzyme A
-	D13.695.827.068.382.300.700 Palmitoyl Coenzyme A
-	D13.695.827.068.395 Cyclic AMP
-	D13.695.827.068.395.225 8-Bromo Cyclic Adenosine Monophosphate
-	D13.695.827.068.395.250 Bucladesine
-	D13.695.827.068.506 Flavin-Adenine Dinucleotide
-	D13.695.827.068.694 NAD
-	D13.695.827.068.749 NADP
-	D13.695.827.068.850 Phosphoadenosine Phosphosulfate
-	D13.695.827.232 Cytosine Nucleotides
-	D13.695.827.232.115 Cyclic CMP
-	D13.695.827.232.150 Cytidine Diphosphate
-	D13.695.827.232.150.180 Cytidine Diphosphate Choline
-	D13.695.827.232.150.210 Cytidine Diphosphate Diglycerides
-	D13.695.827.232.370 Cytidine Monophosphate
-	D13.695.827.232.370.250 Cytidine Monophosphate N-Acetylneuraminic Acid
-	D13.695.827.232.400 Cytidine Triphosphate
-	D13.695.827.349 Flavin Mononucleotide
-	D13.695.827.426 Guanine Nucleotides
-	D13.695.827.426.160 Cyclic GMP
-	D13.695.827.426.160.325 Dibutyryl Cyclic GMP
-	D13.695.827.426.340 Guanosine Diphosphate
-	D13.695.827.426.340.350 Guanosine Diphosphate Sugars
-	D13.695.827.426.340.350.400 Guanosine Diphosphate Fucose
-	D13.695.827.426.340.350.500 Guanosine Diphosphate Mannose
-	D13.695.827.426.440 Guanosine Pentaphosphate
-	D13.695.827.426.480 Guanosine Tetraphosphate
-	D13.695.827.426.504 Guanosine Triphosphate
-	D13.695.827.426.504.380 Guanosine 5'-O-(3-Thiotriphosphate)

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D13.695.827.426.504.400 Guanylyl Imidodiphosphate
-	D13.695.827.426.525 Guanosine Monophosphate
-	D13.695.827.426.700 RNA Caps
-	D13.695.827.426.700.710 RNA Cap Analogs
-	D13.695.827.519 Inosine Nucleotides
-	D13.695.827.519.300 Cyclic IMP
-	D13.695.827.519.400 Inosine Diphosphate
-	D13.695.827.519.500 Inosine Monophosphate
-	D13.695.827.519.800 Inosine Triphosphate
-	D13.695.827.648 Nicotinamide Mononucleotide
-	D13.695.827.708 Nucleoside Diphosphate Sugars
-	D13.695.827.708.070 Adenosine Diphosphate Sugars
-	D13.695.827.708.070.075 Adenosine Diphosphate Glucose
-	D13.695.827.708.070.125 Adenosine Diphosphate Ribose
-	D13.695.827.708.070.125.040 O-Acetyl-ADP-Ribose
-	D13.695.827.708.070.125.195 Cyclic ADP-Ribose
-	D13.695.827.708.070.125.600 Poly Adenosine Diphosphate Ribose
-	D13.695.827.708.260 Cytidine Diphosphate Diglycerides
-	D13.695.827.708.400 Guanosine Diphosphate Sugars
-	D13.695.827.708.400.410 Guanosine Diphosphate Fucose
-	D13.695.827.708.400.500 Guanosine Diphosphate Mannose
-	D13.695.827.708.727 Uridine Diphosphate Sugars
-	D13.695.827.708.727.100 Uridine Diphosphate N-Acetylgalactosamine
-	D13.695.827.708.727.120 Uridine Diphosphate N-Acetylglucosamine
-	D13.695.827.708.727.150 Uridine Diphosphate N-Acetylmuramic Acid
-	D13.695.827.708.727.300 Uridine Diphosphate Galactose
-	D13.695.827.708.727.350 Uridine Diphosphate Glucose
-	D13.695.827.708.727.375 Uridine Diphosphate Glucuronic Acid
-	D13.695.827.708.727.800 Uridine Diphosphate Xylose
-	D13.695.827.919 Uracil Nucleotides
-	D13.695.827.919.600 Uridine Diphosphate
-	D13.695.827.919.600.677 Uridine Diphosphate Sugars
-	D13.695.827.919.600.677.100 Uridine Diphosphate N-Acetylgalactosamine
-	D13.695.827.919.600.677.120 Uridine Diphosphate N-Acetylglucosamine
-	D13.695.827.919.600.677.150 Uridine Diphosphate N-Acetylmuramic Acid
-	D13.695.827.919.600.677.300 Uridine Diphosphate Galactose

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D13.695.827.919.600.677.350 Uridine Diphosphate Glucose
-	D13.695.827.919.600.677.375 Uridine Diphosphate Glucuronic Acid
-	D13.695.827.919.600.677.800 Uridine Diphosphate Xylose
-	D13.695.827.919.877 Uridine Monophosphate
-	D13.695.827.919.877.500 Sofosbuvir
-	D13.695.827.919.950 Uridine Triphosphate
-	D13.695.900 Thionucleotides
-	D13.695.900.380 Guanosine 5'-O-(3-Thiotriphosphate)
-	D20 Complex Mixtures
-	D20.050 Acid Rain
-	D20.080 Ambergris
-	D20.147 Biofuels
-	D20.215 Biological Products
-	D20.215.113 Biological Control Agents
-	D20.215.226 Biological Warfare Agents
-	D20.215.261 Biosimilar Pharmaceuticals
-	D20.215.401 Immune Sera
-	D20.215.401.203 Antilymphocyte Serum
-	D20.215.401.601 Antitoxins
-	D20.215.401.601.163 Antivenins
-	D20.215.401.601.276 Botulinum Antitoxin
-	D20.215.401.601.388 Diphtheria Antitoxin
-	D20.215.401.601.824 Tetanus Antitoxin
-	D20.215.535 Menotropins
-	D20.215.659 Picibanil
-	D20.215.721 Plant Exudates
-	D20.215.721.061 Gutta-Percha
-	D20.215.721.124 Latex
-	D20.215.721.249 Plant Gums
-	D20.215.721.249.249 Chewing Gum
-	D20.215.721.249.500 Gum Arabic
-	D20.215.721.249.625 Karaya Gum
-	D20.215.721.249.750 Tragacanth
-	D20.215.721.500 Resins, Plant
-	D20.215.721.500.109 Amber
-	D20.215.721.500.219 Balsams

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D20.215.721.500.354	Frankincense
-	D20.215.721.500.490	Guaiac
New Heading	<b>D20.215.721.500.626</b>	<b>Mastic Resin</b>
-	D20.215.721.500.762	Propolis
-	D20.215.721.500.881	Turpentine
-	D20.215.780	Plant Nectar
-	D20.215.784	Plant Preparations
-	D20.215.784.249	Coffee
-	D20.215.784.500	Plant Extracts
-	D20.215.784.500.087	Flower Essences
-	D20.215.784.500.175	Cascara
-	D20.215.784.500.200	Curare
-	D20.215.784.500.350	Drugs, Chinese Herbal
-	D20.215.784.500.400	Grape Seed Extract
-	D20.215.784.500.450	Ipecac
-	D20.215.784.500.492	Lecithins
-	D20.215.784.500.534	Opium
-	D20.215.784.500.618	Pectins
-	D20.215.784.500.675	Podophyllin
-	D20.215.784.500.720	Psyllium
-	D20.215.784.500.860	Senna Extract
-	D20.215.784.750	Plant Oils
-	D20.215.784.750.132	Castor Oil
-	D20.215.784.750.186	Clove Oil
-	D20.215.784.750.240	Corn Oil
-	D20.215.784.750.263	Cottonseed Oil
-	D20.215.784.750.315	Croton Oil
-	D20.215.784.750.550	Iodized Oil
-	D20.215.784.750.550.300	Ethiodized Oil
-	D20.215.784.750.615	Linseed Oil
-	D20.215.784.750.840	Safflower Oil
-	D20.215.784.750.865	Sesame Oil
-	D20.215.784.750.880	Soybean Oil
-	D20.215.784.750.940	Tea Tree Oil
-	D20.215.784.844	Tea

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D20.215.784.844.500 Kombucha Tea
-	D20.215.784.937 Teas, Herbal
-	D20.215.894 Vaccines
-	D20.215.894.067 Alzheimer Vaccines
-	D20.215.894.135 Bacterial Vaccines
-	D20.215.894.135.063 Anthrax Vaccines
-	D20.215.894.135.098 Autovaccines
-	D20.215.894.135.134 Brucella Vaccine
-	D20.215.894.135.225 Cholera Vaccines
-	D20.215.894.135.310 Diphtheria-Tetanus Vaccine
-	D20.215.894.135.390 Escherichia coli Vaccines
-	D20.215.894.135.450 Haemophilus Vaccines
-	D20.215.894.135.492 Lyme Disease Vaccines
-	D20.215.894.135.500 Meningococcal Vaccines
-	D20.215.894.135.535 Pertussis Vaccine
-	D20.215.894.135.535.295 Diphtheria-Tetanus-acellular Pertussis Vaccines
-	D20.215.894.135.535.300 Diphtheria-Tetanus-Pertussis Vaccine
-	D20.215.894.135.609 Plague Vaccine
-	D20.215.894.135.620 Pseudomonas Vaccines
-	D20.215.894.135.650 Rickettsial Vaccines
-	D20.215.894.135.685 Salmonella Vaccines
-	D20.215.894.135.685.910 Typhoid-Paratyphoid Vaccines
-	D20.215.894.135.720 Shigella Vaccines
-	D20.215.894.135.744 Staphylococcal Vaccines
-	D20.215.894.135.750 Streptococcal Vaccines
-	D20.215.894.135.750.600 Pneumococcal Vaccines
-	D20.215.894.135.750.600.500 Heptavalent Pneumococcal Conjugate Vaccine
-	D20.215.894.135.825 Tuberculosis Vaccines
-	D20.215.894.135.825.100 BCG Vaccine
-	D20.215.894.200 Cancer Vaccines
-	D20.215.894.354 Fungal Vaccines
-	D20.215.894.582 Protozoan Vaccines
-	D20.215.894.582.480 Leishmaniasis Vaccines
-	D20.215.894.582.500 Malaria Vaccines
-	D20.215.894.691 Toxoids
-	D20.215.894.691.263 Diphtheria Toxoid

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D20.215.894.691.263.290 Diphtheria-Tetanus-acellular Pertussis Vaccines
-	D20.215.894.691.263.300 Diphtheria-Tetanus-Pertussis Vaccine
-	D20.215.894.691.263.310 Diphtheria-Tetanus Vaccine
-	D20.215.894.691.681 Staphylococcal Toxoid
-	D20.215.894.691.824 Tetanus Toxoid
-	D20.215.894.691.824.290 Diphtheria-Tetanus-acellular Pertussis Vaccines
-	D20.215.894.691.824.300 Diphtheria-Tetanus-Pertussis Vaccine
-	D20.215.894.691.824.310 Diphtheria-Tetanus Vaccine
-	D20.215.894.811 Vaccines, Attenuated
-	D20.215.894.815 Vaccines, Combined
-	D20.215.894.815.149 Diphtheria-Tetanus-acellular Pertussis Vaccines
-	D20.215.894.815.300 Diphtheria-Tetanus-Pertussis Vaccine
-	D20.215.894.815.320 Diphtheria-Tetanus Vaccine
-	D20.215.894.815.500 Measles-Mumps-Rubella Vaccine
-	D20.215.894.818 Vaccines, Contraceptive
-	D20.215.894.830 Vaccines, Inactivated
-	D20.215.894.830.614 Poliovirus Vaccine, Inactivated
-	D20.215.894.845 Vaccines, Marker
-	D20.215.894.860 Vaccines, Subunit
-	D20.215.894.860.449 ISCOMs
-	D20.215.894.860.900 Vaccines, Acellular
-	D20.215.894.860.900.267 Diphtheria-Tetanus-acellular Pertussis Vaccines
-	D20.215.894.860.915 Vaccines, Edible
-	D20.215.894.865 Vaccines, Synthetic
-	D20.215.894.865.900 Vaccines, Conjugate
-	D20.215.894.865.910 Vaccines, DNA
-	D20.215.894.865.915 Vaccines, Edible
-	D20.215.894.865.940 Vaccines, Virosome
-	D20.215.894.865.955 Vaccines, Virus-Like Particle
-	D20.215.894.882 Vaccines, Live, Unattenuated
-	D20.215.894.899 Viral Vaccines
-	D20.215.894.899.024 Adenovirus Vaccines
-	D20.215.894.899.050 AIDS Vaccines
-	D20.215.894.899.120 Cytomegalovirus Vaccines
-	D20.215.894.899.162 Dengue Vaccines
-	D20.215.894.899.205 Ebola Vaccines

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D20.215.894.899.290 Herpesvirus Vaccines
-	D20.215.894.899.290.130 Chickenpox Vaccine
-	D20.215.894.899.290.130.400 Herpes Zoster Vaccine
-	D20.215.894.899.290.400 Herpes Simplex Virus Vaccines
-	D20.215.894.899.290.500 Marek Disease Vaccines
-	D20.215.894.899.302 Influenza Vaccines
-	D20.215.894.899.320 Japanese Encephalitis Vaccines
-	D20.215.894.899.404 Measles Vaccine
-	D20.215.894.899.404.500 Measles-Mumps-Rubella Vaccine
-	D20.215.894.899.488 Mumps Vaccine
-	D20.215.894.899.488.500 Measles-Mumps-Rubella Vaccine
-	D20.215.894.899.498 Papillomavirus Vaccines
-	D20.215.894.899.498.500 Human Papillomavirus Recombinant Vaccine Quadrivalent, Types 6, 11, 16, 18
-	D20.215.894.899.500 Parainfluenza Vaccines
-	D20.215.894.899.623 Poliovirus Vaccines
-	D20.215.894.899.623.500 Poliovirus Vaccine, Inactivated
-	D20.215.894.899.623.750 Poliovirus Vaccine, Oral
-	D20.215.894.899.632 Pseudorabies Vaccines
-	D20.215.894.899.700 Rabies Vaccines
-	D20.215.894.899.730 Respiratory Syncytial Virus Vaccines
-	D20.215.894.899.760 Rotavirus Vaccines
-	D20.215.894.899.779 Rubella Vaccine
-	D20.215.894.899.779.500 Measles-Mumps-Rubella Vaccine
-	D20.215.894.899.790 SAIDS Vaccines
-	D20.215.894.899.859 Smallpox Vaccine
-	D20.215.894.899.955 Viral Hepatitis Vaccines
-	D20.215.894.899.955.395 Hepatitis A Vaccines
-	D20.215.894.899.955.400 Hepatitis B Vaccines
-	D20.215.894.899.962 West Nile Virus Vaccines
-	D20.215.894.899.970 Yellow Fever Vaccine
-	D20.280 Colloids
-	D20.280.055 Aerosols
-	D20.280.055.500 Nasal Sprays
-	D20.280.055.750 Oral Sprays
-	D20.280.260 Emulsions



## MeSH Tree Changes for 2017

Type	Tree - heading
-	D20.280.320 Gels
-	D20.280.320.375 Hydrogels
-	D20.280.320.375.500 Cryogels
-	D20.280.320.687 Silica Gel
-	D20.280.810 Suspensions
-	D20.345 Fossil Fuels
-	D20.345.108 Coal
-	D20.345.108.110 Coke
-	D20.345.369 Natural Gas
-	D20.345.630 Petroleum
-	D20.345.630.500 Fuel Oils
-	D20.345.630.540 Gasoline
-	D20.345.630.600 Kerosene
-	D20.475 Freund's Adjuvant
-	D20.538 Lignin
-	D20.601 Manure
-	D20.633 Particulate Matter
-	D20.633.110 Coal Ash
-	D20.633.222 Dust
-	D20.633.222.200 Cosmic Dust
-	D20.633.875 Smog
-	D20.633.937 Smoke
-	D20.633.937.339 Soot
-	D20.633.937.680 Tobacco Smoke Pollution
Old Tree	<b>D20.663 Phytochemicals</b>
-	D20.693 Radioactive Pollutants
-	D20.693.101 Air Pollutants, Radioactive
-	D20.693.555 Radioactive Fallout
-	D20.693.638 Radioactive Waste
-	D20.693.756 Soil Pollutants, Radioactive
-	D20.693.903 Water Pollutants, Radioactive
-	D20.721 Soil
-	D20.721.500 Humic Substances
-	D20.749 Tars
-	D20.749.354 Coal Tar
-	D20.749.850 Tar-Water

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D20.777 Tissue Extracts
-	D20.777.050 Actihaemyl
-	D20.777.162 Cell Extracts
-	D20.777.351 Liver Extracts
-	D20.777.500 Pancreatic Extracts
-	D20.777.500.720 Pancreatin
-	D20.777.500.745 Pancrelipase
-	D20.777.608 Placental Extracts
-	D20.777.871 Thymus Extracts
-	D20.832 Vehicle Emissions
-	D20.888 Venoms
-	D20.888.033 Amphibian Venoms
-	D20.888.033.112 Batrachotoxins
-	D20.888.033.137 Bombesin
-	D20.888.033.163 Bufotenin
-	D20.888.033.728 Physalaemin
-	D20.888.065 Arthropod Venoms
-	D20.888.065.055 Ant Venoms
-	D20.888.065.115 Bee Venoms
-	D20.888.065.115.060 Apamin
-	D20.888.065.115.580 Melitten
-	D20.888.065.830 Scorpion Venoms
-	D20.888.065.830.150 Charybdotoxin
-	D20.888.065.870 Spider Venoms
-	D20.888.065.870.324 Agatoxins
-	D20.888.065.870.324.500 omega-Agatoxin IVA
-	D20.888.065.970 Wasp Venoms
-	D20.888.230 Cnidarian Venoms
-	D20.888.370 Fish Venoms
-	D20.888.590 Mollusk Venoms
-	D20.888.590.162 Conotoxins
-	D20.888.590.162.720 omega-Conotoxins
-	D20.888.590.162.720.700 omega-Conotoxin GVIA
-	D20.888.590.325 Eledoisin
-	D20.888.850 Snake Venoms
-	D20.888.850.325 Elapid Venoms

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D20.888.850.325.139	Bungarotoxins
-	D20.888.850.325.220	Cobra Venoms
-	D20.888.850.325.220.222	Cobra Cardiotoxin Proteins
-	D20.888.850.325.220.244	Cobra Neurotoxin Proteins
-	D20.888.850.960	Viper Venoms
-	D20.888.850.960.200	Crotalid Venoms
-	D20.888.850.960.200.050	Ancrod
-	D20.888.850.960.200.105	Batroxobin
-	D20.888.850.960.200.210	Crotoxin
-	D20.944	Waste Products
-	D20.944.380	Hazardous Waste
-	D20.944.380.638	Radioactive Waste
-	D20.944.420	Industrial Waste
-	D20.944.460	Medical Waste
-	D20.944.460.150	Dental Waste
-	D20.944.460.300	Medical Waste Disposal
-	D20.944.730	Solid Waste
-	D20.944.932	Waste Water
-	D20.944.932.500	Sewage
-	D23	Biological Factors
New Tree	<a href="#">D23.035</a>	<a href="#">Alarmins</a>
-	D23.050	Antigens
-	D23.050.063	Allergens
-	D23.050.101	Antigen-Antibody Complex
-	D23.050.140	Antigens, Archaeal
-	D23.050.161	Antigens, Bacterial
-	D23.050.161.050	Adhesins, Bacterial
-	D23.050.161.050.040	Adhesins, Escherichia coli
-	D23.050.161.386	Lepromin
-	D23.050.161.616	Polysaccharides, Bacterial
-	D23.050.161.616.525	Lipopolysaccharides
-	D23.050.161.616.525.500	Lipid A
-	D23.050.161.616.525.600	O Antigens
-	D23.050.161.616.594	Peptidoglycan
-	D23.050.161.616.797	Teichoic Acids

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D23.050.161.821 Staphylococcal Protein A
-	D23.050.161.845 Tuberculin
-	D23.050.181 Antigens, Dermatophagoides
-	D23.050.202 Antigens, Fungal
-	D23.050.202.180 Coccidioidin
-	D23.050.202.283 Fungal Polysaccharides
-	D23.050.202.386 Histoplasmin
-	D23.050.202.800 Trichophytin
-	D23.050.223 Antigens, Helminth
-	D23.050.244 Antigens, Heterophile
-	D23.050.244.436 Forssman Antigen
-	D23.050.285 Antigens, Neoplasm
-	D23.050.285.018 Antigens, CD24
-	D23.050.285.025 Antigens, CD30
-	D23.050.285.040 Antigens, CD147
-	D23.050.285.050 Antigens, Tumor-Associated, Carbohydrate
-	D23.050.285.050.050 Antigens, CD15
-	D23.050.285.050.119 CA-19-9 Antigen
-	D23.050.285.050.225 CA-125 Antigen
-	D23.050.285.050.300 Mucin-1
-	D23.050.285.062 Antigens, Viral, Tumor
-	D23.050.285.062.045 Adenovirus E1A Proteins
-	D23.050.285.062.050 Adenovirus E1B Proteins
-	D23.050.285.062.090 Antigens, Polyomavirus Transforming
New Heading	<b>D23.050.285.196 Carbonic Anhydrase IX</b>
-	D23.050.285.329 Carcinoembryonic Antigen
New Heading	<b>D23.050.285.357 Epithelial Cell Adhesion Molecule</b>
-	D23.050.285.384 Folate Receptor 1
-	D23.050.285.439 Melanoma-Specific Antigens
-	D23.050.285.439.249 Antigens, CD146
-	D23.050.285.439.500 gp100 Melanoma Antigen
-	D23.050.285.439.750 MART-1 Antigen
-	D23.050.285.550 Neprilysin
-	D23.050.285.625 Prostate-Specific Antigen

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D23.050.285.840 Tissue Polypeptide Antigen
-	D23.050.290 Antigens, Nuclear
-	D23.050.290.249 Epstein-Barr Virus Nuclear Antigens
-	D23.050.290.500 Ki-67 Antigen
New Heading	<b>D23.050.290.625 Ku Autoantigen</b>
-	D23.050.290.750 Proliferating Cell Nuclear Antigen
-	D23.050.290.875 snRNP Core Proteins
-	D23.050.291 Antigens, Plant
-	D23.050.293 Antigens, Protozoan
-	D23.050.293.500 Merozoite Surface Protein 1
-	D23.050.293.900 Variant Surface Glycoproteins, Trypanosoma
-	D23.050.301 Antigens, Surface
-	D23.050.301.264 Antigens, Differentiation
-	D23.050.301.264.035 Antigens, CD
New Heading	<b>D23.050.301.264.035.015 AC133 Antigen</b>
-	D23.050.301.264.035.030 Activated-Leukocyte Cell Adhesion Molecule
New Heading	<b>D23.050.301.264.035.048 ADAM10 Protein</b>
New Heading	<b>D23.050.301.264.035.057 ADAM17 Protein</b>
New Heading	<b>D23.050.301.264.035.065 Antigens, CD99</b>
-	D23.050.301.264.035.100 Antigens, CD1
-	D23.050.301.264.035.100.500 Antigens, CD1d
-	D23.050.301.264.035.102 Antigens, CD2
-	D23.050.301.264.035.103 Antigens, CD3
-	D23.050.301.264.035.103.800 Receptor-CD3 Complex, Antigen, T-Cell
-	D23.050.301.264.035.104 Antigens, CD4
-	D23.050.301.264.035.105 Antigens, CD5
-	D23.050.301.264.035.107 Antigens, CD7
-	D23.050.301.264.035.108 Antigens, CD8
-	D23.050.301.264.035.109 Antigens, CD9
-	D23.050.301.264.035.111 Antigens, CD11
-	D23.050.301.264.035.111.024 Antigens, CD11a
-	D23.050.301.264.035.111.049 Antigens, CD11b

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D23.050.301.264.035.111.074                      Antigen, CD11c
-	D23.050.301.264.035.111.400                      Lymphocyte Function-Associated Antigen-1
-	D23.050.301.264.035.113                          Antigen, CD13
-	D23.050.301.264.035.114                          Antigen, CD14
-	D23.050.301.264.035.115                          Antigen, CD15
-	D23.050.301.264.035.118                          Antigen, CD18
-	D23.050.301.264.035.119                          Antigen, CD19
-	D23.050.301.264.035.120                          Antigen, CD20
-	D23.050.301.264.035.124                          Antigen, CD24
-	D23.050.301.264.035.127                          Antigen, CD27
-	D23.050.301.264.035.128                          Antigen, CD28
-	D23.050.301.264.035.129                          Antigen, CD29
-	D23.050.301.264.035.130                          Antigen, CD30
-	D23.050.301.264.035.131                          Antigen, CD31
-	D23.050.301.264.035.134                          Antigen, CD34
-	D23.050.301.264.035.136                          Antigen, CD36
-	D23.050.301.264.035.138                          Antigen, CD38
-	D23.050.301.264.035.140                          Antigen, CD40
-	D23.050.301.264.035.143                          Antigen, CD43
-	D23.050.301.264.035.145                          Antigen, CD45
-	D23.050.301.264.035.146                          Antigen, CD46
-	D23.050.301.264.035.147                          Antigen, CD47
-	D23.050.301.264.035.153                          Antigen, CD53
-	D23.050.301.264.035.155                          Antigen, CD55
-	D23.050.301.264.035.156                          Antigen, CD56
-	D23.050.301.264.035.157                          Antigen, CD57
-	D23.050.301.264.035.158                          Antigen, CD58
-	D23.050.301.264.035.159                          Antigen, CD59
-	D23.050.301.264.035.163                          Antigen, CD63
-	D23.050.301.264.035.170                          Antigen, CD70
-	D23.050.301.264.035.179                          Antigen, CD79
-	D23.050.301.264.035.181                          Antigen, CD81
-	D23.050.301.264.035.182                          Antigen, CD82
-	D23.050.301.264.035.195                          Antigen, CD95
-	D23.050.301.264.035.198                          Antigen, CD98
-	D23.050.301.264.035.198.500                      Antigen, CD98 Heavy Chain

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D23.050.301.264.035.198.625	Antigens, CD98 Light Chains
-	D23.050.301.264.035.198.625.500	Large Neutral Amino Acid-Transporter 1
-	D23.050.301.264.035.237	Antigens, CD137
-	D23.050.301.264.035.246	Antigens, CD146
-	D23.050.301.264.035.247	Antigens, CD147
-	D23.050.301.264.035.251	Antigens, CD151
-	D23.050.301.264.035.264	Antigens, CD164
-	D23.050.301.264.035.270	Antigens, Thy-1
New Heading	<b>D23.050.301.264.035.272 G, Member 2</b>	<b>ATP Binding Cassette Transporter, Sub-Family</b>
New Tree	<a href="#">D23.050.301.264.035.273</a>	<a href="#">B-Cell Activating Factor</a>
New Tree	<a href="#">D23.050.301.264.035.274</a>	<a href="#">B-Cell Activation Factor Receptor</a>
Old Tree	<del>D23.050.301.264.035.275</del>	<del>B-Cell Activation Factor Receptor</del>
-	D23.050.301.264.035.276	B7 Antigens
-	D23.050.301.264.035.276.100	Antigens, CD80
-	D23.050.301.264.035.276.200	Antigens, CD86
-	D23.050.301.264.035.276.300	Antigens, CD274
-	D23.050.301.264.035.276.500	Inducible T-Cell Co-Stimulator Ligand
-	D23.050.301.264.035.276.800	Programmed Cell Death 1 Ligand 2 Protein
-	D23.050.301.264.035.276.900 Inhibitor 1	V-Set Domain-Containing T-Cell Activation
-	D23.050.301.264.035.277	CD30 Ligand
-	D23.050.301.264.035.278	CD40 Ligand
-	D23.050.301.264.035.279	CTLA-4 Antigen
-	D23.050.301.264.035.280	Cytokine Receptor Common beta Subunit
New Tree	<a href="#">D23.050.301.264.035.281</a>	<a href="#">Cytokine Receptor gp130</a>
Old Tree	<del>D23.050.301.264.035.282</del>	<del>Cytokine Receptor gp130</del>
New Tree	<a href="#">D23.050.301.264.035.282</a>	<a href="#">Dipeptidyl Peptidase 4</a>
Old Tree	<del>D23.050.301.264.035.283</del>	<del>Dipeptidyl Peptidase 4</del>
Old Tree	<del>D23.050.301.264.035.284</del>	<del>Fas Ligand Protein</del>
Old Tree	<del>D23.050.301.264.035.285</del>	<del>fms-Like Tyrosine Kinase 3</del>
Old Tree	<del>D23.050.301.264.035.286</del>	<del>Inducible T-Cell Co-Stimulator Protein</del>
Old Tree	<del>D23.050.301.264.035.287</del>	<del>Integrin alpha1</del>
Old Tree	<del>D23.050.301.264.035.288</del>	<del>Integrin alpha2</del>

## MeSH Tree Changes for 2017

Type	Tree - heading	
Old Tree	D23.050.301.264.035.289	Integrin alpha3
Old Tree	D23.050.301.264.035.291	Integrin alpha4
Old Tree	D23.050.301.264.035.293	Integrin alpha5
New Heading	<b>D23.050.301.264.035.294</b>	<b>Endoglin</b>
Old Tree	D23.050.301.264.035.296	Integrin alpha6
Old Tree	D23.050.301.264.035.298	Integrin alphaV
New Heading	<b>D23.050.301.264.035.300</b>	<b>Epithelial Cell Adhesion Molecule</b>
Old Tree	D23.050.301.264.035.302	Integrin beta3
Old Tree	D23.050.301.264.035.303	Integrin beta4
New Tree	D23.050.301.264.035.305	Fas Ligand Protein
New Tree	D23.050.301.264.035.310	fms-Like Tyrosine Kinase 3
New Heading	<b>D23.050.301.264.035.320</b>	<b>Hepatitis A Virus Cellular Receptor 1</b>
New Heading	<b>D23.050.301.264.035.325</b>	<b>Hepatitis A Virus Cellular Receptor 2</b>
New Tree	D23.050.301.264.035.330	Inducible T-Cell Co-Stimulator Protein
New Tree	D23.050.301.264.035.350	Integrin alpha1
New Tree	D23.050.301.264.035.351	Integrin alpha2
New Tree	D23.050.301.264.035.352	Integrin alpha3
New Tree	D23.050.301.264.035.353	Integrin alpha4
New Tree	D23.050.301.264.035.354	Integrin alpha5
New Tree	D23.050.301.264.035.355	Integrin alpha6
New Tree	D23.050.301.264.035.356	Integrin alphaV
New Tree	D23.050.301.264.035.357	Integrin beta3
New Tree	D23.050.301.264.035.358	Integrin beta4
-	D23.050.301.264.035.400	Intercellular Adhesion Molecule-1
-	D23.050.301.264.035.401	Interleukin-2 Receptor alpha Subunit
-	D23.050.301.264.035.402	Interleukin-2 Receptor beta Subunit



## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D23.050.301.264.035.403	Interleukin-3 Receptor alpha Subunit
-	D23.050.301.264.035.404	Interleukin-4 Receptor alpha Subunit
-	D23.050.301.264.035.405	Interleukin-5 Receptor alpha Subunit
-	D23.050.301.264.035.406	Interleukin-6 Receptor alpha Subunit
-	D23.050.301.264.035.407	Interleukin-7 Receptor alpha Subunit
-	D23.050.301.264.035.409	Interleukin-10 Receptor alpha Subunit
-	D23.050.301.264.035.410	Interleukin-10 Receptor beta Subunit
-	D23.050.301.264.035.412	Interleukin-13 Receptor alpha1 Subunit
-	D23.050.301.264.035.413	Interleukin-13 Receptor alpha2 Subunit
-	D23.050.301.264.035.417	Interleukin-18 Receptor alpha Subunit
-	D23.050.301.264.035.418	Interleukin-18 Receptor beta Subunit
-	D23.050.301.264.035.420	Interleukin Receptor Common gamma Subunit
New Heading	<b>D23.050.301.264.035.438</b>	<b>Jagged-1 Protein</b>
-	D23.050.301.264.035.456	Junctional Adhesion Molecule A
-	D23.050.301.264.035.474	Junctional Adhesion Molecule B
-	D23.050.301.264.035.493	Leukemia Inhibitory Factor Receptor alpha Subunit
-	D23.050.301.264.035.502	Low Density Lipoprotein Receptor-Related Protein-1
-	D23.050.301.264.035.512	Lysosomal-Associated Membrane Protein 1
-	D23.050.301.264.035.531	Lysosomal-Associated Membrane Protein 2
-	D23.050.301.264.035.533	Lysosomal-Associated Membrane Protein 3
New Heading	<b>D23.050.301.264.035.534</b>	<b>Prion Proteins</b>
-	D23.050.301.264.035.535	Mucin-1
-	D23.050.301.264.035.540	Natural Cytotoxicity Triggering Receptor 1
-	D23.050.301.264.035.545	Natural Cytotoxicity Triggering Receptor 2
-	D23.050.301.264.035.547	Natural Cytotoxicity Triggering Receptor 3
-	D23.050.301.264.035.550	Neprilysin
-	D23.050.301.264.035.570	5'-Nucleotidase
-	D23.050.301.264.035.577	NK Cell Lectin-Like Receptor Subfamily B
-	D23.050.301.264.035.579	NK Cell Lectin-Like Receptor Subfamily D
-	D23.050.301.264.035.582	NK Cell Lectin-Like Receptor Subfamily K
New Tree	<b>D23.050.301.264.035.584</b>	<b>P-Glycoprotein</b>
-	D23.050.301.264.035.585	Peptidyl-Dipeptidase A
-	D23.050.301.264.035.587	Platelet Membrane Glycoprotein IIb
-	D23.050.301.264.035.588	Programmed Cell Death 1 Receptor

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D23.050.301.264.035.590	Proto-Oncogene Proteins c-kit
New Tree	<a href="#">D23.050.301.264.035.593</a>	<a href="#">RANK Ligand</a>
New Tree	<a href="#">D23.050.301.264.035.594</a>	<a href="#">Receptor Activator of Nuclear Factor-kappa B</a>
-	D23.050.301.264.035.595	Receptor, Anaphylatoxin C5a
-	D23.050.301.264.035.596	Receptor, ErbB-2
Old Tree	<a href="#">D23.050.301.264.035.597</a>	<a href="#">Receptor, Macrophage Colony-Stimulating Factor</a>
New Tree	<a href="#">D23.050.301.264.035.598</a>	<a href="#">Receptor, Fibroblast Growth Factor, Type 1</a>
New Tree	<a href="#">D23.050.301.264.035.602</a>	<a href="#">Receptor, Fibroblast Growth Factor, Type 2</a>
New Tree	<a href="#">D23.050.301.264.035.605</a>	<a href="#">Receptor, Fibroblast Growth Factor, Type 3</a>
Old Tree	<a href="#">D23.050.301.264.035.605</a>	<a href="#">Receptors, Chemokine</a>
Old Tree	<a href="#">D23.050.301.264.035.605.150</a>	<a href="#">Receptors, CCR</a>
Old Tree	<a href="#">D23.050.301.264.035.605.150.100</a>	<a href="#">Receptors, CCR1</a>
Old Tree	<a href="#">D23.050.301.264.035.605.150.200</a>	<a href="#">Receptors, CCR2</a>
Old Tree	<a href="#">D23.050.301.264.035.605.150.300</a>	<a href="#">Receptors, CCR3</a>
Old Tree	<a href="#">D23.050.301.264.035.605.150.400</a>	<a href="#">Receptors, CCR4</a>
Old Tree	<a href="#">D23.050.301.264.035.605.150.500</a>	<a href="#">Receptors, CCR5</a>
Old Tree	<a href="#">D23.050.301.264.035.605.150.600</a>	<a href="#">Receptors, CCR6</a>
Old Tree	<a href="#">D23.050.301.264.035.605.150.700</a>	<a href="#">Receptors, CCR7</a>
Old Tree	<a href="#">D23.050.301.264.035.605.150.800</a>	<a href="#">Receptors, CCR8</a>
Old Tree	<a href="#">D23.050.301.264.035.605.150.950</a>	<a href="#">Receptors, CCR10</a>
Old Tree	<a href="#">D23.050.301.264.035.605.500</a>	<a href="#">Receptors, CXCR</a>
Old Tree	<a href="#">D23.050.301.264.035.605.500.300</a>	<a href="#">Receptors, CXCR3</a>
Old Tree	<a href="#">D23.050.301.264.035.605.500.400</a>	<a href="#">Receptors, CXCR4</a>
Old Tree	<a href="#">D23.050.301.264.035.605.500.500</a>	<a href="#">Receptors, CXCR5</a>
Old Tree	<a href="#">D23.050.301.264.035.605.500.750</a>	<a href="#">Receptors, Interleukin-8</a>
Old Tree	<a href="#">D23.050.301.264.035.605.500.750.500</a>	<a href="#">Receptors, Interleukin-8A</a>
Old Tree	<a href="#">D23.050.301.264.035.605.500.750.750</a>	<a href="#">Receptors, Interleukin-8B</a>
Old Tree	<a href="#">D23.050.301.264.035.608</a>	<a href="#">Receptors, Complement 3b</a>
Old Tree	<a href="#">D23.050.301.264.035.610</a>	<a href="#">Receptors, Complement 3d</a>
New Tree	<a href="#">D23.050.301.264.035.611</a>	<a href="#">Receptor, Macrophage Colony-Stimulating Factor</a>
New Tree	<a href="#">D23.050.301.264.035.615</a>	<a href="#">Receptors, Chemokine</a>

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	D23.050.301.264.035.615.150	Receptors, CCR
New Tree	D23.050.301.264.035.615.150.100	Receptors, CCR1
New Tree	D23.050.301.264.035.615.150.200	Receptors, CCR2
New Tree	D23.050.301.264.035.615.150.300	Receptors, CCR3
New Tree	D23.050.301.264.035.615.150.400	Receptors, CCR4
New Tree	D23.050.301.264.035.615.150.500	Receptors, CCR5
New Tree	D23.050.301.264.035.615.150.600	Receptors, CCR6
New Tree	D23.050.301.264.035.615.150.700	Receptors, CCR7
New Tree	D23.050.301.264.035.615.150.800	Receptors, CCR8
New Tree	D23.050.301.264.035.615.150.950	Receptors, CCR10
New Tree	D23.050.301.264.035.615.500	Receptors, CXCR
New Tree	D23.050.301.264.035.615.500.300	Receptors, CXCR3
New Tree	D23.050.301.264.035.615.500.400	Receptors, CXCR4
New Tree	D23.050.301.264.035.615.500.500	Receptors, CXCR5
New Tree	D23.050.301.264.035.615.500.750	Receptors, Interleukin-8
New Tree	D23.050.301.264.035.615.500.750.500	Receptors, Interleukin-8A
New Tree	D23.050.301.264.035.615.500.750.750	Receptors, Interleukin-8B
New Tree	D23.050.301.264.035.618	Receptors, Complement 3b
New Tree	D23.050.301.264.035.620	Receptors, Complement 3d
-	D23.050.301.264.035.690	Receptors, IgE
-	D23.050.301.264.035.695	Receptors, IgG
-	D23.050.301.264.035.697	Receptors, Interleukin-1 Type I
-	D23.050.301.264.035.699	Receptors, Interleukin-1 Type II
-	D23.050.301.264.035.715	Receptors, Interleukin-7

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D23.050.301.264.035.782	Receptors, Interleukin-8A
-	D23.050.301.264.035.816	Receptors, Interleukin-9
-	D23.050.301.264.035.820	Receptors, Interleukin-17
-	D23.050.301.264.035.835	Receptors, KIR2DL1
-	D23.050.301.264.035.838	Receptors, KIR2DL2
-	D23.050.301.264.035.842	Receptors, KIR2DL3
-	D23.050.301.264.035.846	Receptors, KIR2DL4
-	D23.050.301.264.035.848	Receptors, KIR3DL1
-	D23.050.301.264.035.849	Receptors, KIR3DL2
-	D23.050.301.264.035.850	Receptors, Lymphocyte Homing
-	D23.050.301.264.035.850.144	Antigens, CD44
-	D23.050.301.264.035.850.347	Integrin alpha4beta1
-	D23.050.301.264.035.850.550	Lymphocyte Function-Associated Antigen-1
-	D23.050.301.264.035.850.903	L-Selectin
-	D23.050.301.264.035.855	Receptors, OX40
-	D23.050.301.264.035.881	Receptors, Thrombopoietin
New Tree	<a href="#">D23.050.301.264.035.895</a>	<a href="#">Receptors, TNF-Related Apoptosis-Inducing Ligand</a>
-	D23.050.301.264.035.908	Receptors, Tumor Necrosis Factor, Type I
-	D23.050.301.264.035.910	Receptors, Tumor Necrosis Factor, Type II
-	D23.050.301.264.035.911	Receptors, Urokinase Plasminogen Activator
-	D23.050.301.264.035.912	Selectins
-	D23.050.301.264.035.912.300	E-Selectin
-	D23.050.301.264.035.912.510	L-Selectin
-	D23.050.301.264.035.912.775	P-Selectin
-	D23.050.301.264.035.913	Sialic Acid Binding Immunoglobulin-like Lectins
-	D23.050.301.264.035.913.049	Myelin-Associated Glycoprotein
-	D23.050.301.264.035.913.100	Sialic Acid Binding Ig-like Lectin 1
-	D23.050.301.264.035.913.200	Sialic Acid Binding Ig-like Lectin 2
-	D23.050.301.264.035.913.300	Sialic Acid Binding Ig-like Lectin 3
New Heading	<b>D23.050.301.264.035.915</b> <b>Family</b>	<b>Signaling Lymphocytic Activation Molecule</b>
New Heading	<b>D23.050.301.264.035.915.250</b>	<b>CD48 Antigen</b>
New Heading	<b>D23.050.301.264.035.915.500</b> <b>Family Member 1</b>	<b>Signaling Lymphocytic Activation Molecule</b>
-	D23.050.301.264.035.916	Syndecan-1

## MeSH Tree Changes for 2017

Type	Tree - heading	
Old Tree	D23.050.301.264.035.917 Protein	Transmembrane Activator and CAML Interactor
New Tree	D23.050.301.264.035.918	TNF-Related Apoptosis-Inducing Ligand
New Tree	D23.050.301.264.035.919	Toll-Like Receptor 9
Old Tree	D23.050.301.264.035.920	Vascular Cell Adhesion Molecule-1
New Tree	D23.050.301.264.035.935 Protein	Transmembrane Activator and CAML Interactor
New Tree	D23.050.301.264.035.945	Vascular Cell Adhesion Molecule-1
-	D23.050.301.264.051	Antigens, Differentiation, B-Lymphocyte
-	D23.050.301.264.051.101	Antigens, CD5
-	D23.050.301.264.051.119	Antigens, CD19
-	D23.050.301.264.051.120	Antigens, CD20
-	D23.050.301.264.051.140	Antigens, CD40
-	D23.050.301.264.894	Antigens, Differentiation, T-Lymphocyte
-	D23.050.301.264.894.080	Antigens, CD1
-	D23.050.301.264.894.080.500	Antigens, CD1d
-	D23.050.301.264.894.090	Antigens, CD2
-	D23.050.301.264.894.095	Antigens, CD3
-	D23.050.301.264.894.095.800	Receptor-CD3 Complex, Antigen, T-Cell
-	D23.050.301.264.894.100	Antigens, CD4
-	D23.050.301.264.894.101	Antigens, CD5
-	D23.050.301.264.894.107	Antigens, CD7
-	D23.050.301.264.894.108	Antigens, CD8
-	D23.050.301.264.894.113	Antigens, CD13
-	D23.050.301.264.894.118	Antigens, CD18
-	D23.050.301.264.894.127	Antigens, CD27
-	D23.050.301.264.894.128	Antigens, CD28
-	D23.050.301.264.894.156	Antigens, CD56
-	D23.050.301.264.894.157	Antigens, CD57
-	D23.050.301.264.894.158	CTLA-4 Antigen
-	D23.050.301.264.894.160	Dipeptidyl Peptidase 4
-	D23.050.301.264.894.580	Inducible T-Cell Co-Stimulator Protein
-	D23.050.301.264.894.790	Programmed Cell Death 1 Receptor
-	D23.050.301.264.900	Antigens, Differentiation, Myelomonocytic

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D23.050.301.264.900.045	Antigens, CD14
-	D23.050.301.264.900.050	Antigens, CD15
-	D23.050.301.264.900.131	Antigens, CD31
-	D23.050.301.264.900.565	Sialic Acid Binding Ig-like Lectin 3
-	D23.050.301.264.920	Antigens, Ly
-	D23.050.301.264.920.500	NK Cell Lectin-Like Receptor Subfamily A
-	D23.050.301.264.965	Stage-Specific Embryonic Antigens
-	D23.050.301.264.965.500	Antigens, CD15
-	D23.050.301.280	Arrestin
-	D23.050.301.285	B7 Antigens
-	D23.050.301.285.100	Antigens, CD80
-	D23.050.301.285.200	Antigens, CD86
-	D23.050.301.285.400	Antigens, CD274
-	D23.050.301.285.500	Inducible T-Cell Co-Stimulator Ligand
-	D23.050.301.285.800	Programmed Cell Death 1 Ligand 2 Protein
-	D23.050.301.285.900	V-Set Domain-Containing T-Cell Activation Inhibitor 1
-	D23.050.301.290	Blood Group Antigens
-	D23.050.301.290.031	ABO Blood-Group System
-	D23.050.301.290.301	Duffy Blood-Group System
-	D23.050.301.290.501	I Blood-Group System
-	D23.050.301.290.533	Kell Blood-Group System
-	D23.050.301.290.538	Kidd Blood-Group System
-	D23.050.301.290.544	Lewis Blood-Group System
-	D23.050.301.290.544.059	Antigens, CD15
-	D23.050.301.290.544.119	CA-19-9 Antigen
-	D23.050.301.290.552	Lutheran Blood-Group System
-	D23.050.301.290.606	MNSs Blood-Group System
-	D23.050.301.290.691	P Blood-Group System
-	D23.050.301.290.775	Rh-Hr Blood-Group System
-	D23.050.301.350	Cell Adhesion Molecules
New Heading	<b>D23.050.301.350.049</b>	<b>Antigens, CD99</b>
-	D23.050.301.350.098	Antigens, CD24
-	D23.050.301.350.131	Antigens, CD31
-	D23.050.301.350.150	Antigens, CD146
-	D23.050.301.350.154	Antigens, CD164

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D23.050.301.350.200                      Cadherins
-	D23.050.301.350.200.500                      Desmosomal Cadherins
-	D23.050.301.350.200.500.249                      Desmocollins
-	D23.050.301.350.200.500.500                      Desmogleins
-	D23.050.301.350.200.500.500.500                      Desmoglein 1
-	D23.050.301.350.200.500.500.625                      Desmoglein 2
-	D23.050.301.350.200.500.500.750                      Desmoglein 3
-	D23.050.301.350.210                      Carcinoembryonic Antigen
-	D23.050.301.350.230                      CD4 Immunoadhesins
-	D23.050.301.350.250                      Cell Adhesion Molecules, Neuronal
-	D23.050.301.350.250.150                      Cell Adhesion Molecules, Neuron-Glia
-	D23.050.301.350.250.150.050                      Activated-Leukocyte Cell Adhesion Molecule
-	D23.050.301.350.250.325                      Contactins
-	D23.050.301.350.250.325.100                      Contactin 1
-	D23.050.301.350.250.325.200                      Contactin 2
-	D23.050.301.350.250.500                      Myelin P0 Protein
-	D23.050.301.350.250.520                      Neural Cell Adhesion Molecules
-	D23.050.301.350.250.520.156                      Antigens, CD56
-	D23.050.301.350.250.520.578                      Neural Cell Adhesion Molecule L1
-	D23.050.301.350.250.520.789                      Neurocan
New Heading	<b>D23.050.301.350.263                      Epithelial Cell Adhesion Molecule</b>
-	D23.050.301.350.275                      Integrin alphaXbeta2
-	D23.050.301.350.275.500                      Antigens, CD11c
-	D23.050.301.350.275.750                      Antigens, CD18
-	D23.050.301.350.450                      Intercellular Adhesion Molecule-1
-	D23.050.301.350.537                      Junctional Adhesion Molecules
-	D23.050.301.350.537.500                      Coxsackie and Adenovirus Receptor-Like Membrane Protein
-	D23.050.301.350.537.750                      Junctional Adhesion Molecule A
-	D23.050.301.350.537.875                      Junctional Adhesion Molecule B
-	D23.050.301.350.537.937                      Junctional Adhesion Molecule C
-	D23.050.301.350.625                      Receptors, Lymphocyte Homing
-	D23.050.301.350.625.144                      Antigens, CD44
-	D23.050.301.350.625.347                      Integrin alpha4beta1
-	D23.050.301.350.625.550                      Lymphocyte Function-Associated Antigen-1

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D23.050.301.350.625.903 L-Selectin
-	D23.050.301.350.700 Selectins
-	D23.050.301.350.700.300 E-Selectin
-	D23.050.301.350.700.510 L-Selectin
-	D23.050.301.350.700.775 P-Selectin
-	D23.050.301.350.720 Sialic Acid Binding Ig-like Lectin 2
-	D23.050.301.350.920 Vascular Cell Adhesion Molecule-1
-	D23.050.301.500 Histocompatibility Antigens
-	D23.050.301.500.100 Histocompatibility Antigens Class I
-	D23.050.301.500.100.350 H-2 Antigens
New Heading	<b>D23.050.301.500.100.363 Hemochromatosis Protein</b>
-	D23.050.301.500.100.375 Histocompatibility Antigen H-2D
-	D23.050.301.500.100.400 HLA-A Antigens
-	D23.050.301.500.100.400.010 HLA-A1 Antigen
-	D23.050.301.500.100.400.020 HLA-A2 Antigen
-	D23.050.301.500.100.400.030 HLA-A3 Antigen
-	D23.050.301.500.100.400.110 HLA-A11 Antigen
-	D23.050.301.500.100.400.240 HLA-A24 Antigen
-	D23.050.301.500.100.500 HLA-B Antigens
-	D23.050.301.500.100.500.070 HLA-B7 Antigen
-	D23.050.301.500.100.500.080 HLA-B8 Antigen
-	D23.050.301.500.100.500.130 HLA-B13 Antigen
-	D23.050.301.500.100.500.140 HLA-B14 Antigen
-	D23.050.301.500.100.500.150 HLA-B15 Antigen
-	D23.050.301.500.100.500.180 HLA-B18 Antigen
-	D23.050.301.500.100.500.270 HLA-B27 Antigen
-	D23.050.301.500.100.500.350 HLA-B35 Antigen
-	D23.050.301.500.100.500.370 HLA-B37 Antigen
-	D23.050.301.500.100.500.380 HLA-B38 Antigen
-	D23.050.301.500.100.500.390 HLA-B39 Antigen
-	D23.050.301.500.100.500.400 HLA-B40 Antigen
-	D23.050.301.500.100.500.440 HLA-B44 Antigen
-	D23.050.301.500.100.500.510 HLA-B51 Antigen
-	D23.050.301.500.100.500.520 HLA-B52 Antigen
-	D23.050.301.500.100.600 HLA-C Antigens



## MeSH Tree Changes for 2017

Type	Tree - heading
-	D23.050.301.500.100.700 HLA-G Antigens
-	D23.050.301.500.400 Histocompatibility Antigens Class II
-	D23.050.301.500.400.199 H-2 Antigens
-	D23.050.301.500.400.400 HLA-D Antigens
-	D23.050.301.500.400.400.420 HLA-DP Antigens
-	D23.050.301.500.400.400.420.500 HLA-DP alpha-Chains
-	D23.050.301.500.400.400.420.750 HLA-DP beta-Chains
-	D23.050.301.500.400.400.430 HLA-DQ Antigens
-	D23.050.301.500.400.400.430.500 HLA-DQ alpha-Chains
-	D23.050.301.500.400.400.430.750 HLA-DQ beta-Chains
-	D23.050.301.500.400.400.440 HLA-DR Antigens
-	D23.050.301.500.400.400.440.100 HLA-DR alpha-Chains
-	D23.050.301.500.400.400.440.200 HLA-DR beta-Chains
-	D23.050.301.500.400.400.440.200.010 HLA-DRB1 Chains
-	D23.050.301.500.400.400.440.200.030 HLA-DRB3 Chains
-	D23.050.301.500.400.400.440.200.040 HLA-DRB4 Chains
-	D23.050.301.500.400.400.440.200.050 HLA-DRB5 Chains
-	D23.050.301.500.400.400.440.400 HLA-DR Serological Subtypes
-	D23.050.301.500.400.400.440.400.010 HLA-DR1 Antigen
-	D23.050.301.500.400.400.440.400.020 HLA-DR2 Antigen
-	D23.050.301.500.400.400.440.400.030 HLA-DR3 Antigen
-	D23.050.301.500.400.400.440.400.040 HLA-DR4 Antigen
-	D23.050.301.500.400.400.440.400.050 HLA-DR5 Antigen
-	D23.050.301.500.400.400.440.400.060 HLA-DR6 Antigen
-	D23.050.301.500.400.400.440.400.070 HLA-DR7 Antigen
-	D23.050.301.500.450 HLA Antigens
-	D23.050.301.500.450.370 HLA-A Antigens
-	D23.050.301.500.450.370.010 HLA-A1 Antigen
-	D23.050.301.500.450.370.020 HLA-A2 Antigen
-	D23.050.301.500.450.370.030 HLA-A3 Antigen
-	D23.050.301.500.450.370.110 HLA-A11 Antigen
-	D23.050.301.500.450.370.240 HLA-A24 Antigen
-	D23.050.301.500.450.380 HLA-B Antigens
-	D23.050.301.500.450.380.070 HLA-B7 Antigen
-	D23.050.301.500.450.380.080 HLA-B8 Antigen
-	D23.050.301.500.450.380.130 HLA-B13 Antigen

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D23.050.301.500.450.380.140 HLA-B14 Antigen
-	D23.050.301.500.450.380.150 HLA-B15 Antigen
-	D23.050.301.500.450.380.180 HLA-B18 Antigen
-	D23.050.301.500.450.380.270 HLA-B27 Antigen
-	D23.050.301.500.450.380.350 HLA-B35 Antigen
-	D23.050.301.500.450.380.370 HLA-B37 Antigen
-	D23.050.301.500.450.380.380 HLA-B38 Antigen
-	D23.050.301.500.450.380.390 HLA-B39 Antigen
-	D23.050.301.500.450.380.400 HLA-B40 Antigen
-	D23.050.301.500.450.380.440 HLA-B44 Antigen
-	D23.050.301.500.450.380.510 HLA-B51 Antigen
-	D23.050.301.500.450.380.520 HLA-B52 Antigen
-	D23.050.301.500.450.390 HLA-C Antigens
-	D23.050.301.500.450.400 HLA-D Antigens
-	D23.050.301.500.450.400.420 HLA-DP Antigens
-	D23.050.301.500.450.400.420.500 HLA-DP alpha-Chains
-	D23.050.301.500.450.400.420.750 HLA-DP beta-Chains
-	D23.050.301.500.450.400.430 HLA-DQ Antigens
-	D23.050.301.500.450.400.430.500 HLA-DQ alpha-Chains
-	D23.050.301.500.450.400.430.750 HLA-DQ beta-Chains
-	D23.050.301.500.450.400.440 HLA-DR Antigens
-	D23.050.301.500.450.400.440.222 HLA-DR alpha-Chains
-	D23.050.301.500.450.400.440.333 HLA-DR beta-Chains
-	D23.050.301.500.450.400.440.333.500 HLA-DRB1 Chains
-	D23.050.301.500.450.400.440.333.750 HLA-DRB3 Chains
-	D23.050.301.500.450.400.440.333.875 HLA-DRB4 Chains
-	D23.050.301.500.450.400.440.333.937 HLA-DRB5 Chains
-	D23.050.301.500.450.400.440.389 HLA-DR Serological Subtypes
-	D23.050.301.500.450.400.440.389.500 HLA-DR1 Antigen
-	D23.050.301.500.450.400.440.389.750 HLA-DR2 Antigen
-	D23.050.301.500.450.400.440.389.875 HLA-DR3 Antigen
-	D23.050.301.500.450.400.440.389.937 HLA-DR4 Antigen
-	D23.050.301.500.450.400.440.389.968 HLA-DR5 Antigen
-	D23.050.301.500.450.400.440.389.984 HLA-DR6 Antigen
-	D23.050.301.500.450.400.440.389.992 HLA-DR7 Antigen
-	D23.050.301.500.450.700 HLA-G Antigens

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D23.050.301.500.600	Minor Histocompatibility Antigens
New Tree	<a href="#">D23.050.301.500.600.200</a>	<a href="#">Antigens, CD19</a>
New Tree	<a href="#">D23.050.301.500.600.300</a>	<a href="#">Cathepsin H</a>
New Tree	<a href="#">D23.050.301.500.600.350</a>	<a href="#">Focal Adhesion Kinase 2</a>
-	D23.050.301.500.600.400	H-Y Antigen
New Tree	<a href="#">D23.050.301.500.600.700</a>	<a href="#">Receptor, ErbB-2</a>
New Tree	<a href="#">D23.050.301.500.600.775</a>	<a href="#">Receptors, Purinergic P2X5</a>
New Tree	<a href="#">D23.050.301.500.600.813</a>	<a href="#">Reduced Folate Carrier Protein</a>
New Tree	<a href="#">D23.050.301.500.600.850</a>	<a href="#">rho Guanine Nucleotide Dissociation Inhibitor beta</a>
New Tree	<a href="#">D23.050.301.500.600.925</a>	<a href="#">Thymidine Phosphorylase</a>
-	D23.050.301.562	Leukocyte L1 Antigen Complex
-	D23.050.301.562.100	Calgranulin A
-	D23.050.301.562.200	Calgranulin B
-	D23.050.301.593	Lymphocyte Antigen 96
-	D23.050.301.625	Minor Lymphocyte Stimulatory Antigens
-	D23.050.301.900	Variant Surface Glycoproteins, Trypanosoma
-	D23.050.324	Antigens, T-Independent
-	D23.050.327	Antigens, Viral
-	D23.050.327.045	Adenovirus Early Proteins
-	D23.050.327.045.050	Adenovirus E1 Proteins
-	D23.050.327.045.060	Adenovirus E2 Proteins
-	D23.050.327.045.070	Adenovirus E3 Proteins
-	D23.050.327.045.080	Adenovirus E4 Proteins
-	D23.050.327.062	Antigens, Viral, Tumor
-	D23.050.327.062.045	Adenovirus E1A Proteins
-	D23.050.327.062.050	Adenovirus E1B Proteins
-	D23.050.327.062.090	Antigens, Polyomavirus Transforming
-	D23.050.327.150	Deltaretrovirus Antigens
-	D23.050.327.150.500	HTLV-I Antigens
-	D23.050.327.150.510	HTLV-II Antigens
-	D23.050.327.300	Epstein-Barr Virus Nuclear Antigens

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D23.050.327.461 Hemagglutinins, Viral
-	D23.050.327.461.350 HN Protein
-	D23.050.327.495 Hepatitis Antigens
-	D23.050.327.495.249 Hepatitis A Antigens
-	D23.050.327.495.500 Hepatitis B Antigens
-	D23.050.327.495.500.450 Hepatitis B Core Antigens
-	D23.050.327.495.500.469 Hepatitis B e Antigens
-	D23.050.327.495.500.475 Hepatitis B Surface Antigens
-	D23.050.327.495.505 Hepatitis C Antigens
-	D23.050.327.495.752 Hepatitis delta Antigens
-	D23.050.327.520 HIV Antigens
-	D23.050.327.520.300 HIV Core Protein p24
-	D23.050.327.520.330 HIV Envelope Protein gp41
-	D23.050.327.520.350 HIV Envelope Protein gp120
-	D23.050.422 Autoantigens
-	D23.050.422.061 Centromere Protein B
-	D23.050.422.124 Desmoglein 1
-	D23.050.422.249 Desmoglein 3
-	D23.050.422.500 Heymann Nephritis Antigenic Complex
-	D23.050.422.500.500 Low Density Lipoprotein Receptor-Related Protein-2
-	D23.050.422.500.750 LDL-Receptor Related Protein-Associated Protein
-	D23.050.422.625 Myelin-Oligodendrocyte Glycoprotein
-	D23.050.422.750 snRNP Core Proteins
-	D23.050.550 Epitopes
-	D23.050.550.325 Antigens, Tumor-Associated, Carbohydrate
-	D23.050.550.325.050 Antigens, CD15
-	D23.050.550.325.119 CA-19-9 Antigen
-	D23.050.550.325.225 CA-125 Antigen
-	D23.050.550.325.300 Mucin-1
-	D23.050.550.395 Epitopes, B-Lymphocyte
-	D23.050.550.402 Epitopes, T-Lymphocyte
-	D23.050.550.480 Haptens
-	D23.050.550.480.097 p-Azobenzene arsonate
-	D23.050.550.480.330 Cardiolipins
-	D23.050.550.480.350 Dinitrochlorobenzene
-	D23.050.550.480.641 Nitrohydroxyiodophenylacetate

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D23.050.550.480.820                      Picryl Chloride
-	D23.050.550.500                              Immunodominant Epitopes
-	D23.050.550.750                              Immunoglobulin Idiotypes
-	D23.050.705                                    Isoantigens
-	D23.050.705.115                              Antigen, Human Platelet
-	D23.050.705.230                              Blood Group Antigens
-	D23.050.705.230.031                          ABO Blood-Group System
-	D23.050.705.230.301                          Duffy Blood-Group System
-	D23.050.705.230.501                          I Blood-Group System
-	D23.050.705.230.533                          Kell Blood-Group System
-	D23.050.705.230.538                          Kidd Blood-Group System
-	D23.050.705.230.544                          Lewis Blood-Group System
-	D23.050.705.230.544.059                      Antigen, CD15
-	D23.050.705.230.544.119                      CA-19-9 Antigen
-	D23.050.705.230.552                          Lutheran Blood-Group System
-	D23.050.705.230.606                          MNSs Blood-Group System
-	D23.050.705.230.691                          P Blood-Group System
-	D23.050.705.230.775                          Rh-Hr Blood-Group System
-	D23.050.705.552                              Histocompatibility Antigens
-	D23.050.705.552.100                          Histocompatibility Antigens Class I
-	D23.050.705.552.100.350                      H-2 Antigens
New Heading	<b>D23.050.705.552.100.375                      Hemochromatosis Protein</b>
-	D23.050.705.552.100.400                      HLA-A Antigens
-	D23.050.705.552.100.400.010                      HLA-A1 Antigen
-	D23.050.705.552.100.400.020                      HLA-A2 Antigen
-	D23.050.705.552.100.400.030                      HLA-A3 Antigen
-	D23.050.705.552.100.400.110                      HLA-A11 Antigen
-	D23.050.705.552.100.400.240                      HLA-A24 Antigen
-	D23.050.705.552.100.500                      HLA-B Antigens
-	D23.050.705.552.100.500.070                      HLA-B7 Antigen
-	D23.050.705.552.100.500.080                      HLA-B8 Antigen
-	D23.050.705.552.100.500.130                      HLA-B13 Antigen
-	D23.050.705.552.100.500.140                      HLA-B14 Antigen
-	D23.050.705.552.100.500.150                      HLA-B15 Antigen
-	D23.050.705.552.100.500.180                      HLA-B18 Antigen

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D23.050.705.552.100.500.270 HLA-B27 Antigen
-	D23.050.705.552.100.500.350 HLA-B35 Antigen
-	D23.050.705.552.100.500.370 HLA-B37 Antigen
-	D23.050.705.552.100.500.380 HLA-B38 Antigen
-	D23.050.705.552.100.500.390 HLA-B39 Antigen
-	D23.050.705.552.100.500.400 HLA-B40 Antigen
-	D23.050.705.552.100.500.440 HLA-B44 Antigen
-	D23.050.705.552.100.500.510 HLA-B51 Antigen
-	D23.050.705.552.100.500.520 HLA-B52 Antigen
-	D23.050.705.552.100.600 HLA-C Antigens
-	D23.050.705.552.100.700 HLA-G Antigens
-	D23.050.705.552.410 Histocompatibility Antigens Class II
-	D23.050.705.552.410.199 H-2 Antigens
-	D23.050.705.552.410.400 HLA-D Antigens
-	D23.050.705.552.410.400.420 HLA-DP Antigens
-	D23.050.705.552.410.400.420.500 HLA-DP alpha-Chains
-	D23.050.705.552.410.400.420.750 HLA-DP beta-Chains
-	D23.050.705.552.410.400.430 HLA-DQ Antigens
-	D23.050.705.552.410.400.430.500 HLA-DQ alpha-Chains
-	D23.050.705.552.410.400.430.750 HLA-DQ beta-Chains
-	D23.050.705.552.410.400.440 HLA-DR Antigens
-	D23.050.705.552.410.400.440.100 HLA-DR alpha-Chains
-	D23.050.705.552.410.400.440.200 HLA-DR beta-Chains
-	D23.050.705.552.410.400.440.200.010 HLA-DRB1 Chains
-	D23.050.705.552.410.400.440.200.030 HLA-DRB3 Chains
-	D23.050.705.552.410.400.440.200.040 HLA-DRB4 Chains
-	D23.050.705.552.410.400.440.200.050 HLA-DRB5 Chains
-	D23.050.705.552.410.400.440.389 HLA-DR Serological Subtypes
-	D23.050.705.552.410.400.440.389.010 HLA-DR1 Antigen
-	D23.050.705.552.410.400.440.389.020 HLA-DR2 Antigen
-	D23.050.705.552.410.400.440.389.030 HLA-DR3 Antigen
-	D23.050.705.552.410.400.440.389.040 HLA-DR4 Antigen
-	D23.050.705.552.410.400.440.389.050 HLA-DR5 Antigen
-	D23.050.705.552.410.400.440.389.060 HLA-DR6 Antigen
-	D23.050.705.552.410.400.440.389.070 HLA-DR7 Antigen
-	D23.050.705.552.450 HLA Antigens

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D23.050.705.552.450.370 HLA-A Antigens
-	D23.050.705.552.450.370.372 HLA-A1 Antigen
-	D23.050.705.552.450.370.374 HLA-A2 Antigen
-	D23.050.705.552.450.370.376 HLA-A3 Antigen
-	D23.050.705.552.450.370.500 HLA-A11 Antigen
-	D23.050.705.552.450.370.600 HLA-A24 Antigen
-	D23.050.705.552.450.380 HLA-B Antigens
-	D23.050.705.552.450.380.070 HLA-B7 Antigen
-	D23.050.705.552.450.380.080 HLA-B8 Antigen
-	D23.050.705.552.450.380.130 HLA-B13 Antigen
-	D23.050.705.552.450.380.140 HLA-B14 Antigen
-	D23.050.705.552.450.380.150 HLA-B15 Antigen
-	D23.050.705.552.450.380.180 HLA-B18 Antigen
-	D23.050.705.552.450.380.270 HLA-B27 Antigen
-	D23.050.705.552.450.380.350 HLA-B35 Antigen
-	D23.050.705.552.450.380.370 HLA-B37 Antigen
-	D23.050.705.552.450.380.380 HLA-B38 Antigen
-	D23.050.705.552.450.380.390 HLA-B39 Antigen
-	D23.050.705.552.450.380.400 HLA-B40 Antigen
-	D23.050.705.552.450.380.440 HLA-B44 Antigen
-	D23.050.705.552.450.380.510 HLA-B51 Antigen
-	D23.050.705.552.450.380.520 HLA-B52 Antigen
-	D23.050.705.552.450.390 HLA-C Antigens
-	D23.050.705.552.450.400 HLA-D Antigens
-	D23.050.705.552.450.400.420 HLA-DP Antigens
-	D23.050.705.552.450.400.420.500 HLA-DP alpha-Chains
-	D23.050.705.552.450.400.420.750 HLA-DP beta-Chains
-	D23.050.705.552.450.400.430 HLA-DQ Antigens
-	D23.050.705.552.450.400.430.500 HLA-DQ alpha-Chains
-	D23.050.705.552.450.400.430.750 HLA-DQ beta-Chains
-	D23.050.705.552.450.400.440 HLA-DR Antigens
-	D23.050.705.552.450.400.440.222 HLA-DR alpha-Chains
-	D23.050.705.552.450.400.440.333 HLA-DR beta-Chains
-	D23.050.705.552.450.400.440.333.500 HLA-DRB1 Chains
-	D23.050.705.552.450.400.440.333.750 HLA-DRB3 Chains
-	D23.050.705.552.450.400.440.333.875 HLA-DRB4 Chains

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D23.050.705.552.450.400.440.333.937	HLA-DRB5 Chains
-	D23.050.705.552.450.400.440.389	HLA-DR Serological Subtypes
-	D23.050.705.552.450.400.440.389.500	HLA-DR1 Antigen
-	D23.050.705.552.450.400.440.389.750	HLA-DR2 Antigen
-	D23.050.705.552.450.400.440.389.875	HLA-DR3 Antigen
-	D23.050.705.552.450.400.440.389.937	HLA-DR4 Antigen
-	D23.050.705.552.450.400.440.389.968	HLA-DR5 Antigen
-	D23.050.705.552.450.400.440.389.984	HLA-DR6 Antigen
-	D23.050.705.552.450.400.440.389.992	HLA-DR7 Antigen
-	D23.050.705.552.450.700	HLA-G Antigens
-	D23.050.705.552.600	Minor Histocompatibility Antigens
New Tree	<a href="#">D23.050.705.552.600.200</a>	<a href="#">Antigens, CD19</a>
New Tree	<a href="#">D23.050.705.552.600.300</a>	<a href="#">Cathepsin H</a>
New Tree	<a href="#">D23.050.705.552.600.350</a>	<a href="#">Focal Adhesion Kinase 2</a>
-	D23.050.705.552.600.400	H-Y Antigen
New Tree	<a href="#">D23.050.705.552.600.550</a>	<a href="#">Receptor, ErbB-2</a>
New Tree	<a href="#">D23.050.705.552.600.625</a>	<a href="#">Receptors, Purinergic P2X5</a>
New Tree	<a href="#">D23.050.705.552.600.663</a>	<a href="#">Reduced Folate Carrier Protein</a>
New Tree	<a href="#">D23.050.705.552.600.700</a>	<a href="#">rho Guanine Nucleotide Dissociation Inhibitor beta</a>
New Tree	<a href="#">D23.050.705.552.600.850</a>	<a href="#">Thymidine Phosphorylase</a>
-	D23.050.820	Superantigens
-	D23.050.820.625	Minor Lymphocyte Stimulatory Antigens
-	D23.050.865	Vaccines, Synthetic
-	D23.050.865.900	Vaccines, Conjugate
-	D23.050.865.910	Vaccines, DNA
-	D23.050.865.915	Vaccines, Edible
-	D23.050.865.940	Vaccines, Virosome
-	D23.050.865.955	Vaccines, Virus-Like Particle
-	D23.101	Biomarkers
-	D23.101.050	Antibodies, Antineutrophil Cytoplasmic
-	D23.101.100	Antigens, Differentiation



## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D23.101.100.110	Antigens, CD
New Heading	<b>D23.101.100.110.015</b>	<b>AC133 Antigen</b>
-	D23.101.100.110.030	Activated-Leukocyte Cell Adhesion Molecule
New Heading	<b>D23.101.100.110.048</b>	<b>ADAM10 Protein</b>
New Heading	<b>D23.101.100.110.057</b>	<b>ADAM17 Protein</b>
New Heading	<b>D23.101.100.110.065</b>	<b>Antigens, CD99</b>
-	D23.101.100.110.100	Antigens, CD1
-	D23.101.100.110.100.500	Antigens, CD1d
-	D23.101.100.110.102	Antigens, CD2
-	D23.101.100.110.103	Antigens, CD3
-	D23.101.100.110.103.800	Receptor-CD3 Complex, Antigen, T-Cell
-	D23.101.100.110.104	Antigens, CD4
-	D23.101.100.110.105	Antigens, CD5
-	D23.101.100.110.106	Antigens, CD53
-	D23.101.100.110.107	Antigens, CD7
-	D23.101.100.110.109	Antigens, CD8
-	D23.101.100.110.110	Antigens, CD9
-	D23.101.100.110.111	Antigens, CD11
-	D23.101.100.110.111.024	Antigens, CD11a
-	D23.101.100.110.111.049	Antigens, CD11b
-	D23.101.100.110.111.074	Antigens, CD11c
-	D23.101.100.110.111.400	Lymphocyte Function-Associated Antigen-1
-	D23.101.100.110.113	Antigens, CD13
-	D23.101.100.110.114	Antigens, CD14
-	D23.101.100.110.115	Antigens, CD15
-	D23.101.100.110.118	Antigens, CD18
-	D23.101.100.110.119	Antigens, CD19
-	D23.101.100.110.120	Antigens, CD20
-	D23.101.100.110.124	Antigens, CD24
-	D23.101.100.110.127	Antigens, CD27
-	D23.101.100.110.128	Antigens, CD28
-	D23.101.100.110.129	Antigens, CD29
-	D23.101.100.110.130	Antigens, CD30

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D23.101.100.110.131	Antigens, CD31
-	D23.101.100.110.134	Antigens, CD34
-	D23.101.100.110.136	Antigens, CD36
-	D23.101.100.110.138	Antigens, CD38
-	D23.101.100.110.140	Antigens, CD40
-	D23.101.100.110.143	Antigens, CD43
-	D23.101.100.110.145	Antigens, CD45
-	D23.101.100.110.146	Antigens, CD46
-	D23.101.100.110.147	Antigens, CD47
-	D23.101.100.110.155	Antigens, CD55
-	D23.101.100.110.156	Antigens, CD56
-	D23.101.100.110.157	Antigens, CD57
-	D23.101.100.110.158	Antigens, CD58
-	D23.101.100.110.159	Antigens, CD59
-	D23.101.100.110.163	Antigens, CD63
-	D23.101.100.110.170	Antigens, CD70
-	D23.101.100.110.179	Antigens, CD79
-	D23.101.100.110.181	Antigens, CD81
-	D23.101.100.110.182	Antigens, CD82
-	D23.101.100.110.195	Antigens, CD95
-	D23.101.100.110.198	Antigens, CD98
-	D23.101.100.110.198.500	Antigens, CD98 Heavy Chain
-	D23.101.100.110.198.625	Antigens, CD98 Light Chains
-	D23.101.100.110.198.625.500	Large Neutral Amino Acid-Transporter 1
-	D23.101.100.110.237	Antigens, CD137
-	D23.101.100.110.246	Antigens, CD146
-	D23.101.100.110.247	Antigens, CD147
-	D23.101.100.110.251	Antigens, CD151
-	D23.101.100.110.264	Antigens, CD164
-	D23.101.100.110.270	Antigens, Thy-1
New Heading	<b>D23.101.100.110.272 Member 2</b>	<b>ATP Binding Cassette Transporter, Sub-Family G,</b>
New Tree	<a href="#">D23.101.100.110.273</a>	<a href="#">B-Cell Activating Factor</a>
New Tree	<a href="#">D23.101.100.110.274</a>	<a href="#">B-Cell Activation Factor Receptor</a>
Old Tree	<a href="#">D23.101.100.110.275</a>	<a href="#">B-Cell Activation Factor Receptor</a>

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D23.101.100.110.276	B7 Antigens
-	D23.101.100.110.276.100	Antigens, CD80
-	D23.101.100.110.276.200	Antigens, CD86
-	D23.101.100.110.276.300	Antigens, CD274
-	D23.101.100.110.276.500	Inducible T-Cell Co-Stimulator Ligand
-	D23.101.100.110.276.800	Programmed Cell Death 1 Ligand 2 Protein
-	D23.101.100.110.276.900 1	V-Set Domain-Containing T-Cell Activation Inhibitor
-	D23.101.100.110.277	CD30 Ligand
-	D23.101.100.110.278	CD40 Ligand
-	D23.101.100.110.279	CTLA-4 Antigen
-	D23.101.100.110.280	Cytokine Receptor gp130
-	D23.101.100.110.281	Cytokine Receptor Common beta Subunit
Old Tree	<del>D23.101.100.110.282</del>	<del>Dipeptidyl Peptidase 4</del>
Old Tree	<del>D23.101.100.110.283</del>	<del>Fas Ligand Protein</del>
Old Tree	<del>D23.101.100.110.284</del>	<del>fms-Like Tyrosine Kinase 3</del>
Old Tree	<del>D23.101.100.110.285</del>	<del>Inducible T-Cell Co-Stimulator Protein</del>
Old Tree	<del>D23.101.100.110.286</del>	<del>Integrin alpha1</del>
New Tree	D23.101.100.110.287	Dipeptidyl Peptidase 4
Old Tree	<del>D23.101.100.110.288</del>	<del>Integrin alpha2</del>
Old Tree	<del>D23.101.100.110.290</del>	<del>Integrin alpha3</del>
Old Tree	<del>D23.101.100.110.292</del>	<del>Integrin alpha4</del>
Old Tree	<del>D23.101.100.110.294</del>	<del>Integrin alpha5</del>
New Heading	<b>D23.101.100.110.296</b>	<b>Endoglin</b>
Old Tree	<del>D23.101.100.110.297</del>	<del>Integrin alpha6</del>
Old Tree	<del>D23.101.100.110.298</del>	<del>Integrin alphaV</del>
Old Tree	<del>D23.101.100.110.299</del>	<del>Integrin beta3</del>
New Heading	<b>D23.101.100.110.301</b>	<b>Epithelial Cell Adhesion Molecule</b>
Old Tree	<del>D23.101.100.110.302</del>	<del>Integrin beta4</del>
New Tree	D23.101.100.110.305	Fas Ligand Protein
New Tree	D23.101.100.110.310	fms-Like Tyrosine Kinase 3
New Heading	<b>D23.101.100.110.320</b>	<b>Hepatitis A Virus Cellular Receptor 1</b>

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Heading	<b>D23.101.100.110.325</b>	<b>Hepatitis A Virus Cellular Receptor 2</b>
New Tree	D23.101.100.110.330	Inducible T-Cell Co-Stimulator Protein
New Tree	D23.101.100.110.350	Integrin alpha1
New Tree	D23.101.100.110.351	Integrin alpha2
New Tree	D23.101.100.110.352	Integrin alpha3
New Tree	D23.101.100.110.353	Integrin alpha4
New Tree	D23.101.100.110.354	Integrin alpha5
New Tree	D23.101.100.110.355	Integrin alpha6
New Tree	D23.101.100.110.356	Integrin alphaV
New Tree	D23.101.100.110.357	Integrin beta3
New Tree	D23.101.100.110.358	Integrin beta4
-	D23.101.100.110.400	Intercellular Adhesion Molecule-1
-	D23.101.100.110.401	Interleukin-2 Receptor alpha Subunit
-	D23.101.100.110.402	Interleukin-2 Receptor beta Subunit
-	D23.101.100.110.403	Interleukin-3 Receptor alpha Subunit
-	D23.101.100.110.404	Interleukin-4 Receptor alpha Subunit
-	D23.101.100.110.405	Interleukin-5 Receptor alpha Subunit
-	D23.101.100.110.406	Interleukin-6 Receptor alpha Subunit
-	D23.101.100.110.407	Interleukin-7 Receptor alpha Subunit
-	D23.101.100.110.409	Interleukin-10 Receptor alpha Subunit
-	D23.101.100.110.410	Interleukin-10 Receptor beta Subunit
-	D23.101.100.110.412	Interleukin-13 Receptor alpha1 Subunit
-	D23.101.100.110.413	Interleukin-13 Receptor alpha2 Subunit
-	D23.101.100.110.417	Interleukin-18 Receptor alpha Subunit
-	D23.101.100.110.418	Interleukin-18 Receptor beta Subunit
-	D23.101.100.110.420	Interleukin Receptor Common gamma Subunit
New Heading	<b>D23.101.100.110.438</b>	<b>Jagged-1 Protein</b>
-	D23.101.100.110.456	Junctional Adhesion Molecule A

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D23.101.100.110.474	Junctional Adhesion Molecule B
-	D23.101.100.110.493	Leukemia Inhibitory Factor Receptor alpha Subunit
-	D23.101.100.110.498	Low Density Lipoprotein Receptor-Related Protein-1
-	D23.101.100.110.512	Lysosomal-Associated Membrane Protein 1
-	D23.101.100.110.531	Lysosomal-Associated Membrane Protein 2
-	D23.101.100.110.533	Lysosomal-Associated Membrane Protein 3
New Heading	<b>D23.101.100.110.534</b>	<b>Prion Proteins</b>
-	D23.101.100.110.535	Mucin-1
-	D23.101.100.110.540	Natural Cytotoxicity Triggering Receptor 1
-	D23.101.100.110.545	Natural Cytotoxicity Triggering Receptor 2
-	D23.101.100.110.547	Natural Cytotoxicity Triggering Receptor 3
-	D23.101.100.110.550	Nepriylsin
-	D23.101.100.110.570	5'-Nucleotidase
-	D23.101.100.110.577	NK Cell Lectin-Like Receptor Subfamily B
-	D23.101.100.110.579	NK Cell Lectin-Like Receptor Subfamily D
-	D23.101.100.110.582	NK Cell Lectin-Like Receptor Subfamily K
New Tree	<a href="#">D23.101.100.110.584</a>	<a href="#">P-Glycoprotein</a>
-	D23.101.100.110.585	Peptidyl-Dipeptidase A
-	D23.101.100.110.587	Platelet Membrane Glycoprotein IIb
-	D23.101.100.110.588	Programmed Cell Death 1 Receptor
-	D23.101.100.110.590	Proto-Oncogene Proteins c-kit
New Tree	<a href="#">D23.101.100.110.593</a>	<a href="#">RANK Ligand</a>
New Tree	<a href="#">D23.101.100.110.594</a>	<a href="#">Receptor Activator of Nuclear Factor-kappa B</a>
-	D23.101.100.110.595	Receptor, Anaphylatoxin C5a
-	D23.101.100.110.596	Receptor, ErbB-2
Old Tree	<del>D23.101.100.110.597</del>	<del>Receptor, Macrophage Colony-Stimulating Factor</del>
New Tree	<a href="#">D23.101.100.110.598</a>	<a href="#">Receptor, Fibroblast Growth Factor, Type 1</a>
New Tree	<a href="#">D23.101.100.110.602</a>	<a href="#">Receptor, Fibroblast Growth Factor, Type 2</a>
New Tree	<a href="#">D23.101.100.110.605</a>	<a href="#">Receptor, Fibroblast Growth Factor, Type 3</a>
Old Tree	<del>D23.101.100.110.605</del>	<del>Receptors, Chemokine</del>
Old Tree	<del>D23.101.100.110.605.150</del>	<del>Receptors, CCR</del>

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	D23.101.100.110.605.150.100      Receptors, CCR1
Old Tree	D23.101.100.110.605.150.200      Receptors, CCR2
Old Tree	D23.101.100.110.605.150.300      Receptors, CCR3
Old Tree	D23.101.100.110.605.150.400      Receptors, CCR4
Old Tree	D23.101.100.110.605.150.500      Receptors, CCR5
Old Tree	D23.101.100.110.605.150.600      Receptors, CCR6
Old Tree	D23.101.100.110.605.150.700      Receptors, CCR7
Old Tree	D23.101.100.110.605.150.800      Receptors, CCR8
Old Tree	D23.101.100.110.605.150.950      Receptors, CCR10
Old Tree	D23.101.100.110.605.500              Receptors, CXCR
Old Tree	D23.101.100.110.605.500.300      Receptors, CXCR3
Old Tree	D23.101.100.110.605.500.400      Receptors, CXCR4
Old Tree	D23.101.100.110.605.500.500      Receptors, CXCR5
Old Tree	D23.101.100.110.605.500.750      Receptors, Interleukin-8
Old Tree	D23.101.100.110.605.500.750.500      Receptors, Interleukin-8A
Old Tree	D23.101.100.110.605.500.750.750      Receptors, Interleukin-8B
Old Tree	D23.101.100.110.608                  Receptors, Complement 3b
Old Tree	D23.101.100.110.610                  Receptors, Complement 3d
New Tree	D23.101.100.110.611                  Receptor, Macrophage Colony-Stimulating Factor
New Tree	D23.101.100.110.615                  Receptors, Chemokine
New Tree	D23.101.100.110.615.150              Receptors, CCR
New Tree	D23.101.100.110.615.150.100      Receptors, CCR1
New Tree	D23.101.100.110.615.150.200      Receptors, CCR2
New Tree	D23.101.100.110.615.150.300      Receptors, CCR3
New Tree	D23.101.100.110.615.150.400      Receptors, CCR4
New Tree	D23.101.100.110.615.150.500      Receptors, CCR5
New Tree	D23.101.100.110.615.150.600      Receptors, CCR6
New Tree	D23.101.100.110.615.150.700      Receptors, CCR7
New Tree	D23.101.100.110.615.150.800      Receptors, CCR8

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	D23.101.100.110.615.150.950	Receptors, CCR10
New Tree	D23.101.100.110.615.500	Receptors, CXCR
New Tree	D23.101.100.110.615.500.300	Receptors, CXCR3
New Tree	D23.101.100.110.615.500.400	Receptors, CXCR4
New Tree	D23.101.100.110.615.500.500	Receptors, CXCR5
New Tree	D23.101.100.110.615.500.750	Receptors, Interleukin-8
New Tree	D23.101.100.110.615.500.750.500	Receptors, Interleukin-8A
New Tree	D23.101.100.110.615.500.750.750	Receptors, Interleukin-8B
New Tree	D23.101.100.110.618	Receptors, Complement 3b
New Tree	D23.101.100.110.620	Receptors, Complement 3d
-	D23.101.100.110.690	Receptors, IgE
-	D23.101.100.110.695	Receptors, IgG
-	D23.101.100.110.697	Receptors, Interleukin-1 Type I
-	D23.101.100.110.699	Receptors, Interleukin-1 Type II
-	D23.101.100.110.715	Receptors, Interleukin-7
-	D23.101.100.110.782	Receptors, Interleukin-8A
-	D23.101.100.110.816	Receptors, Interleukin-9
-	D23.101.100.110.820	Receptors, Interleukin-17
-	D23.101.100.110.835	Receptors, KIR2DL1
-	D23.101.100.110.838	Receptors, KIR2DL2
-	D23.101.100.110.842	Receptors, KIR2DL3
-	D23.101.100.110.846	Receptors, KIR2DL4
-	D23.101.100.110.848	Receptors, KIR3DL1
-	D23.101.100.110.849	Receptors, KIR3DL2
-	D23.101.100.110.850	Receptors, Lymphocyte Homing
-	D23.101.100.110.850.144	Antigens, CD44
-	D23.101.100.110.850.347	Integrin alpha4beta1
-	D23.101.100.110.850.550	Lymphocyte Function-Associated Antigen-1
-	D23.101.100.110.850.903	L-Selectin

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D23.101.100.110.855      Receptors, OX40
-	D23.101.100.110.883      Receptors, Thrombopoietin
New Tree	<a href="#">D23.101.100.110.898</a> <a href="#">Receptors, TNF-Related Apoptosis-Inducing Ligand</a>
-	D23.101.100.110.912      Receptors, Tumor Necrosis Factor, Type I
-	D23.101.100.110.916      Receptors, Tumor Necrosis Factor, Type II
-	D23.101.100.110.917      Receptors, Urokinase Plasminogen Activator
-	D23.101.100.110.918      Selectins
-	D23.101.100.110.918.300      E-Selectin
-	D23.101.100.110.918.510      L-Selectin
-	D23.101.100.110.918.775      P-Selectin
-	D23.101.100.110.929      Sialic Acid Binding Immunoglobulin-like Lectins
-	D23.101.100.110.929.049      Myelin-Associated Glycoprotein
-	D23.101.100.110.929.100      Sialic Acid Binding Ig-like Lectin 1
-	D23.101.100.110.929.200      Sialic Acid Binding Ig-like Lectin 2
-	D23.101.100.110.929.300      Sialic Acid Binding Ig-like Lectin 3
New Heading	<b>D23.101.100.110.935</b> <b>Signaling Lymphocytic Activation Molecule Family</b>
New Heading	<b>D23.101.100.110.935.250</b> <b>CD48 Antigen</b>
New Heading	<b>D23.101.100.110.935.500</b> <b>Signaling Lymphocytic Activation Molecule Family Member 1</b>
-	D23.101.100.110.940      Syndecan-1
New Tree	<a href="#">D23.101.100.110.943</a> <a href="#">TNF-Related Apoptosis-Inducing Ligand</a>
New Tree	<a href="#">D23.101.100.110.945</a> <a href="#">Toll-Like Receptor 9</a>
-	D23.101.100.110.950      Transmembrane Activator and CAML Interactor Protein
-	D23.101.100.110.970      Vascular Cell Adhesion Molecule-1
-	D23.101.100.150      Antigens, Differentiation, B-Lymphocyte
-	D23.101.100.150.101      Antigens, CD5
-	D23.101.100.150.119      Antigens, CD19
-	D23.101.100.150.120      Antigens, CD20
-	D23.101.100.150.140      Antigens, CD40
-	D23.101.100.894      Antigens, Differentiation, T-Lymphocyte
-	D23.101.100.894.080      Antigens, CD1
-	D23.101.100.894.080.500      Antigens, CD1d
-	D23.101.100.894.090      Antigens, CD2



## MeSH Tree Changes for 2017

Type	Tree - heading
-	D23.101.100.894.095                      Antigens, CD3
-	D23.101.100.894.095.800                      Receptor-CD3 Complex, Antigen, T-Cell
-	D23.101.100.894.100                      Antigens, CD4
-	D23.101.100.894.101                      Antigens, CD5
-	D23.101.100.894.107                      Antigens, CD7
-	D23.101.100.894.108                      Antigens, CD8
-	D23.101.100.894.113                      Antigens, CD13
-	D23.101.100.894.118                      Antigens, CD18
-	D23.101.100.894.127                      Antigens, CD27
-	D23.101.100.894.128                      Antigens, CD28
-	D23.101.100.894.156                      Antigens, CD56
-	D23.101.100.894.157                      Antigens, CD57
-	D23.101.100.894.158                      CTLA-4 Antigen
-	D23.101.100.894.160                      Dipeptidyl Peptidase 4
-	D23.101.100.894.580                      Inducible T-Cell Co-Stimulator Protein
-	D23.101.100.894.790                      Programmed Cell Death 1 Receptor
-	D23.101.100.900                      Antigens, Differentiation, Myelomonocytic
-	D23.101.100.900.045                      Antigens, CD14
-	D23.101.100.900.050                      Antigens, CD15
-	D23.101.100.900.131                      Antigens, CD31
-	D23.101.100.900.565                      Sialic Acid Binding Ig-like Lectin 3
-	D23.101.100.920                      Antigens, Ly
-	D23.101.100.920.500                      NK Cell Lectin-Like Receptor Subfamily A
-	D23.101.100.965                      Stage-Specific Embryonic Antigens
-	D23.101.137                      Biomarkers, Pharmacological
-	D23.101.140                      Biomarkers, Tumor
-	D23.101.140.050                      alpha-Fetoproteins
-	D23.101.140.055                      Antigens, CD30
-	D23.101.140.075                      Antigens, Tumor-Associated, Carbohydrate
-	D23.101.140.075.050                      Antigens, CD15
-	D23.101.140.075.119                      CA-19-9 Antigen
-	D23.101.140.075.225                      CA-125 Antigen
-	D23.101.140.075.300                      Mucin-1
-	D23.101.140.300                      Carcinoembryonic Antigen
-	D23.101.140.325                      Chorionic Gonadotropin, beta Subunit, Human
-	D23.101.140.375                      Hormones, Ectopic

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D23.101.140.400 Ki-67 Antigen
-	D23.101.140.425 Mammaglobin A
-	D23.101.140.450 Nephroblastoma Overexpressed Protein
-	D23.101.140.500 Neprilysin
-	D23.101.140.540 Normetanephrine
-	D23.101.140.600 Proliferating Cell Nuclear Antigen
-	D23.101.140.625 Prostate-Specific Antigen
-	D23.101.140.642 Receptor, ErbB-2
-	D23.101.140.721 Receptor, ErbB-3
-	D23.101.140.760 Receptor, ErbB-4
-	D23.101.140.800 Synaptophysin
-	D23.101.140.870 Tissue Kallikreins
-	D23.101.140.880 Tissue Polypeptide Antigen
-	D23.101.175 Chorionic Gonadotropin, beta Subunit, Human
-	D23.101.340 Fibrinopeptide A
-	D23.101.387 Genetic Markers
-	D23.101.613 Oligoclonal Bands
-	D23.113 Blood Coagulation Factor Inhibitors
-	D23.113.025 Antithrombin III
-	D23.113.475 Lupus Coagulation Inhibitor
-	D23.113.700 Protein C
-	D23.113.725 Protein S
-	D23.119 Blood Coagulation Factors
-	D23.119.050 beta-Thromboglobulin
-	D23.119.100 Calcium
-	D23.119.300 Factor V
-	D23.119.300.300 Factor Va
-	D23.119.325 Factor VII
-	D23.119.325.300 Factor VIIa
-	D23.119.350 Factor VIII
-	D23.119.350.300 Factor VIIIa
-	D23.119.375 Factor IX
-	D23.119.375.310 Factor IXa
-	D23.119.400 Factor X
-	D23.119.400.315 Factor Xa
-	D23.119.425 Factor XI

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D23.119.425.320 Factor XIa
-	D23.119.450 Factor XII
-	D23.119.450.324 Factor XIIIa
-	D23.119.475 Factor XIII
-	D23.119.475.500 Factor XIIIa
-	D23.119.490 Fibrinogen
-	D23.119.490.265 Fibrinogens, Abnormal
-	D23.119.500 Fibrinopeptide A
-	D23.119.510 Fibrinopeptide B
-	D23.119.597 Kallikreins
-	D23.119.597.712 Prekallikrein
-	D23.119.630 Kininogens
-	D23.119.630.480 Kininogen, High-Molecular-Weight
-	D23.119.630.490 Kininogen, Low-Molecular-Weight
-	D23.119.832 Plasminogen Inactivators
-	D23.119.832.500 Plasminogen Activator Inhibitor 1
-	D23.119.832.750 Plasminogen Activator Inhibitor 2
-	D23.119.865 Platelet Activating Factor
-	D23.119.932 Platelet Factor 3
-	D23.119.940 Platelet Factor 4
-	D23.119.945 Prothrombin
-	D23.119.960 Thrombin
-	D23.119.965 Thromboplastin
-	D23.119.970 Tissue Plasminogen Activator
-	D23.119.985 von Willebrand Factor
-	D23.125 Chemotactic Factors
-	D23.125.300 Chemokines
-	D23.125.300.070 beta-Thromboglobulin
-	D23.125.300.100 Chemokines, C
-	D23.125.300.110 Chemokines, CC
-	D23.125.300.110.050 Chemokine CCL1
-	D23.125.300.110.150 Chemokine CCL3
-	D23.125.300.110.200 Chemokine CCL4
-	D23.125.300.110.250 Chemokine CCL5
-	D23.125.300.110.550 Chemokine CCL11
-	D23.125.300.110.850 Chemokine CCL17

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D23.125.300.110.870 Chemokine CCL19
-	D23.125.300.110.880 Chemokine CCL20
-	D23.125.300.110.890 Chemokine CCL21
-	D23.125.300.110.900 Chemokine CCL22
-	D23.125.300.110.910 Chemokine CCL24
-	D23.125.300.110.930 Chemokine CCL27
-	D23.125.300.110.990 Monocyte Chemoattractant Proteins
-	D23.125.300.110.990.600 Chemokine CCL2
-	D23.125.300.110.990.800 Chemokine CCL7
-	D23.125.300.110.990.900 Chemokine CCL8
-	D23.125.300.120 Chemokines, CXC
-	D23.125.300.120.050 Chemokine CXCL1
-	D23.125.300.120.100 Chemokine CXCL2
-	D23.125.300.120.250 Chemokine CXCL5
-	D23.125.300.120.300 Chemokine CXCL6
-	D23.125.300.120.450 Chemokine CXCL9
-	D23.125.300.120.500 Chemokine CXCL10
-	D23.125.300.120.550 Chemokine CXCL11
-	D23.125.300.120.600 Chemokine CXCL12
-	D23.125.300.120.650 Chemokine CXCL13
-	D23.125.300.120.800 Interleukin-8
-	D23.125.300.120.900 Platelet Factor 4
-	D23.125.300.130 Chemokines, CX3C
-	D23.125.300.130.500 Chemokine CX3CL1
-	D23.125.300.600 Macrophage Inflammatory Proteins
-	D23.125.300.600.500 Chemokine CCL3
-	D23.125.300.600.750 Chemokine CCL4
-	D23.125.300.600.870 Chemokine CCL19
-	D23.125.300.600.880 Chemokine CCL20
-	D23.125.300.600.940 Chemokine CXCL2
-	D23.125.320 Chemotactic Factors, Eosinophil
-	D23.125.477 Leukocyte Migration-Inhibitory Factors
-	D23.125.477.500 Macrophage Migration-Inhibitory Factors
-	D23.125.685 N-Formylmethionine Leucyl-Phenylalanine
-	D23.297 Host-Derived Cellular Factors
-	D23.297.500 Integration Host Factors

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D23.469 Inflammation Mediators
-	D23.469.050 Autacoids
-	D23.469.050.050 Angiotensins
-	D23.469.050.050.025 Angiotensin I
-	D23.469.050.050.050 Angiotensin II
-	D23.469.050.050.050.050 Angiotensin Amide
-	D23.469.050.050.050.700 Saralasin
-	D23.469.050.050.050.725 1-Sarcosine-8-Isoleucine Angiotensin II
-	D23.469.050.050.075 Angiotensin III
-	D23.469.050.175 Eicosanoids
-	D23.469.050.175.450 Leukotrienes
-	D23.469.050.175.450.400 Leukotriene A4
-	D23.469.050.175.450.415 Leukotriene B4
-	D23.469.050.175.450.725 SRS-A
-	D23.469.050.175.450.725.400 Leukotriene C4
-	D23.469.050.175.450.725.415 Leukotriene D4
-	D23.469.050.175.450.725.425 Leukotriene E4
-	D23.469.050.175.725 Prostaglandins
-	D23.469.050.175.725.025 Prostaglandin Endoperoxides
-	D23.469.050.175.725.025.600 Prostaglandins G
-	D23.469.050.175.725.025.650 Prostaglandins H
-	D23.469.050.175.725.025.650.500 Prostaglandin H2
-	D23.469.050.175.725.025.650.500.500 (epoxymethano)prosta-5,13-dienoic Acid 15-Hydroxy-11 alpha,9 alpha-
-	D23.469.050.175.725.100 Prostaglandins A
-	D23.469.050.175.725.150 Prostaglandins B
-	D23.469.050.175.725.200 Prostaglandins D
-	D23.469.050.175.725.200.200 Prostaglandin D2
-	D23.469.050.175.725.250 Prostaglandins E
-	D23.469.050.175.725.250.100 Alprostadil
-	D23.469.050.175.725.250.200 Dinoprostone
-	D23.469.050.175.725.400 Prostaglandins F
-	D23.469.050.175.725.400.200 Dinoprost
-	D23.469.050.175.725.400.350 6-Ketoprostaglandin F1 alpha
-	D23.469.050.175.725.550 Prostaglandins I
-	D23.469.050.175.725.550.500 Epoprostenol

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D23.469.050.175.725.775 Prostaglandins, Synthetic
-	D23.469.050.175.725.775.125 Iloprost
-	D23.469.050.175.725.775.250 Prostaglandin Endoperoxides, Synthetic
-	D23.469.050.175.725.775.350 Prostaglandins A, Synthetic
-	D23.469.050.175.725.775.450 Prostaglandins E, Synthetic
-	D23.469.050.175.725.775.450.050 Arbutoprost
-	D23.469.050.175.725.775.450.300 16,16-Dimethylprostaglandin E2
-	D23.469.050.175.725.775.450.350 Enprostil
-	D23.469.050.175.725.775.450.500 Misoprostol
-	D23.469.050.175.725.775.450.750 Rioprostil
-	D23.469.050.175.725.775.500 Prostaglandins F, Synthetic
-	D23.469.050.175.725.775.500.150 Carboprost
-	D23.469.050.175.725.775.500.175 Cloprostenol
-	D23.469.050.175.725.775.500.175.250 Bimatoprost
-	D23.469.050.175.725.775.500.175.500 Travoprost
-	D23.469.050.300 Histamine
-	D23.469.050.375 Kinins
-	D23.469.050.375.110 Bradykinin
-	D23.469.050.375.110.400 Kallidin
-	D23.469.050.375.425 Kininogens
-	D23.469.050.375.425.350 Kininogen, High-Molecular-Weight
-	D23.469.050.375.425.400 Kininogen, Low-Molecular-Weight
-	D23.469.050.375.850 Tachykinins
-	D23.469.050.375.850.275 Eledoisin
-	D23.469.050.375.850.400 Kassinin
-	D23.469.050.375.850.550 Neurokinin A
-	D23.469.050.375.850.570 Neurokinin B
-	D23.469.050.375.850.780 Physalaemin
-	D23.469.050.375.850.890 Substance P
-	D23.469.050.375.925 Urotensins
-	D23.469.050.600 Platelet Activating Factor
-	D23.469.050.650 Serotonin
-	D23.469.200 Chemokines
-	D23.469.200.070 beta-Thromboglobulin
-	D23.469.200.100 Chemokines, C
-	D23.469.200.110 Chemokines, CC

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D23.469.200.110.050 Chemokine CCL1
-	D23.469.200.110.150 Chemokine CCL3
-	D23.469.200.110.200 Chemokine CCL4
-	D23.469.200.110.250 Chemokine CCL5
-	D23.469.200.110.550 Chemokine CCL11
-	D23.469.200.110.850 Chemokine CCL17
-	D23.469.200.110.870 Chemokine CCL19
-	D23.469.200.110.880 Chemokine CCL20
-	D23.469.200.110.890 Chemokine CCL21
-	D23.469.200.110.900 Chemokine CCL22
-	D23.469.200.110.910 Chemokine CCL24
-	D23.469.200.110.930 Chemokine CCL27
-	D23.469.200.110.990 Monocyte Chemoattractant Proteins
-	D23.469.200.110.990.600 Chemokine CCL2
-	D23.469.200.110.990.800 Chemokine CCL7
-	D23.469.200.110.990.900 Chemokine CCL8
-	D23.469.200.120 Chemokines, CXC
-	D23.469.200.120.050 Chemokine CXCL1
-	D23.469.200.120.100 Chemokine CXCL2
-	D23.469.200.120.250 Chemokine CXCL5
-	D23.469.200.120.300 Chemokine CXCL6
-	D23.469.200.120.450 Chemokine CXCL9
-	D23.469.200.120.500 Chemokine CXCL10
-	D23.469.200.120.550 Chemokine CXCL11
-	D23.469.200.120.600 Chemokine CXCL12
-	D23.469.200.120.650 Chemokine CXCL13
-	D23.469.200.120.800 Interleukin-8
-	D23.469.200.120.900 Platelet Factor 4
-	D23.469.200.130 Chemokines, CX3C
-	D23.469.200.130.500 Chemokine CX3CL1
-	D23.469.200.600 Macrophage Inflammatory Proteins
-	D23.469.200.600.150 Chemokine CCL3
-	D23.469.200.600.870 Chemokine CCL19
-	D23.469.200.600.880 Chemokine CCL20
-	D23.469.200.600.940 Chemokine CXCL2
-	D23.469.700 Prostaglandins, Synthetic

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D23.469.700.275 lloprost
-	D23.469.700.630 Prostaglandin Endoperoxides, Synthetic
-	D23.469.700.630.500 15-Hydroxy-11 alpha,9 alpha- (epoxymethano)prosta-5,13-dienoic Acid
-	D23.469.700.645 Prostaglandins A, Synthetic
-	D23.469.700.660 Prostaglandins E, Synthetic
-	D23.469.700.660.050 Arbaprostil
-	D23.469.700.660.200 16,16-Dimethylprostaglandin E2
-	D23.469.700.660.250 Enprostil
-	D23.469.700.660.500 Misoprostol
-	D23.469.700.660.750 Rioprostil
-	D23.469.700.670 Prostaglandins F, Synthetic
-	D23.469.700.670.150 Carboprost
-	D23.469.700.670.175 Cloprostenol
-	D23.469.700.670.175.250 Bimatoprost
-	D23.469.700.670.175.500 Travoprost
-	D23.529 Intercellular Signaling Peptides and Proteins
-	D23.529.024 Adipokines
-	D23.529.024.249 Adiponectin
-	D23.529.024.500 Leptin
-	D23.529.024.750 Resistin
-	D23.529.049 Agouti Signaling Protein
-	D23.529.074 Agouti-Related Protein
Old Tree	<b>D23.529.087 Alarmins</b>
Old Tree	<b>D23.529.087.048 alpha-Defensins</b>
Old Tree	<b>D23.529.087.233 beta-Defensins</b>
Old Tree	<b>D23.529.087.416 Calgranulin A</b>
Old Tree	<b>D23.529.087.452 Calgranulin B</b>
Old Tree	<b>D23.529.087.470 Cathelicidins</b>
Old Tree	<b>D23.529.087.488 Chaperonin 60</b>
Old Tree	<b>D23.529.087.612 HMGB1 Protein</b>
Old Tree	<b>D23.529.087.672 HSC70 Heat-Shock Proteins</b>
Old Tree	<b>D23.529.087.780 Interleukin-33</b>
Old Tree	<b>D23.529.087.887 S100 Calcium Binding Protein beta Subunit</b>
Old Tree	<b>D23.529.087.943 S100A12 Protein</b>
-	D23.529.100 Angiogenic Proteins



## MeSH Tree Changes for 2017

Type	Tree - heading
-	D23.529.100.100 Angiopoietins
-	D23.529.100.100.100 Angiopoietin-1
-	D23.529.100.100.200 Angiopoietin-2
-	D23.529.100.450 Angiostatic Proteins
-	D23.529.100.450.500 Angiostatins
-	D23.529.100.450.750 Endostatins
-	D23.529.100.800 Vascular Endothelial Growth Factors
-	D23.529.100.800.200 Vascular Endothelial Growth Factor A
-	D23.529.100.800.300 Vascular Endothelial Growth Factor B
-	D23.529.100.800.400 Vascular Endothelial Growth Factor C
-	D23.529.100.800.500 Vascular Endothelial Growth Factor D
-	D23.529.100.800.600 Vascular Endothelial Growth Factor, Endocrine-Gland-Derived
-	D23.529.168 B7 Antigens
-	D23.529.168.100 Antigens, CD80
-	D23.529.168.200 Antigens, CD86
-	D23.529.168.300 Antigens, CD274
-	D23.529.168.500 Inducible T-Cell Co-Stimulator Ligand
-	D23.529.168.800 Programmed Cell Death 1 Ligand 2 Protein
-	D23.529.168.900 V-Set Domain-Containing T-Cell Activation Inhibitor 1
-	D23.529.237 CCN Intercellular Signaling Proteins
-	D23.529.237.100 Connective Tissue Growth Factor
-	D23.529.237.200 Cysteine-Rich Protein 61
-	D23.529.237.600 Nephroblastoma Overexpressed Protein
-	D23.529.374 Cytokines
-	D23.529.374.200 Chemokines
-	D23.529.374.200.070 beta-Thromboglobulin
-	D23.529.374.200.100 Chemokines, C
-	D23.529.374.200.110 Chemokines, CC
-	D23.529.374.200.110.050 Chemokine CCL1
-	D23.529.374.200.110.150 Chemokine CCL3
-	D23.529.374.200.110.200 Chemokine CCL4
-	D23.529.374.200.110.250 Chemokine CCL5
-	D23.529.374.200.110.550 Chemokine CCL11
-	D23.529.374.200.110.850 Chemokine CCL17
-	D23.529.374.200.110.870 Chemokine CCL19

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D23.529.374.200.110.880 Chemokine CCL20
-	D23.529.374.200.110.890 Chemokine CCL21
-	D23.529.374.200.110.900 Chemokine CCL22
-	D23.529.374.200.110.910 Chemokine CCL24
-	D23.529.374.200.110.930 Chemokine CCL27
-	D23.529.374.200.110.990 Monocyte Chemoattractant Proteins
-	D23.529.374.200.110.990.500 Chemokine CCL2
-	D23.529.374.200.110.990.750 Chemokine CCL7
-	D23.529.374.200.110.990.875 Chemokine CCL8
-	D23.529.374.200.120 Chemokines, CXC
-	D23.529.374.200.120.050 Chemokine CXCL1
-	D23.529.374.200.120.100 Chemokine CXCL2
-	D23.529.374.200.120.250 Chemokine CXCL5
-	D23.529.374.200.120.300 Chemokine CXCL6
-	D23.529.374.200.120.450 Chemokine CXCL9
-	D23.529.374.200.120.500 Chemokine CXCL10
-	D23.529.374.200.120.550 Chemokine CXCL11
-	D23.529.374.200.120.600 Chemokine CXCL12
-	D23.529.374.200.120.650 Chemokine CXCL13
-	D23.529.374.200.120.800 Interleukin-8
-	D23.529.374.200.120.900 Platelet Factor 4
-	D23.529.374.200.130 Chemokines, CX3C
-	D23.529.374.200.130.500 Chemokine CX3CL1
-	D23.529.374.200.600 Macrophage Inflammatory Proteins
-	D23.529.374.200.600.150 Chemokine CCL3
-	D23.529.374.200.600.200 Chemokine CCL4
-	D23.529.374.200.600.870 Chemokine CCL19
-	D23.529.374.200.600.880 Chemokine CCL20
-	D23.529.374.200.600.940 Chemokine CXCL2
-	D23.529.374.305 Growth Differentiation Factor 15
-	D23.529.374.410 Hematopoietic Cell Growth Factors
-	D23.529.374.410.240 Colony-Stimulating Factors
-	D23.529.374.410.240.150 Erythropoietin
-	D23.529.374.410.240.150.750 Epoetin Alfa
-	D23.529.374.410.240.350 Granulocyte Colony-Stimulating Factor
-	D23.529.374.410.240.350.500 Filgrastim

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D23.529.374.410.240.375 Granulocyte-Macrophage Colony-Stimulating Factor
-	D23.529.374.410.240.400 Interleukin-3
-	D23.529.374.410.240.500 Macrophage Colony-Stimulating Factor
-	D23.529.374.410.240.750 Thrombopoietin
-	D23.529.374.410.800 Stem Cell Factor
-	D23.529.374.420 Hepatocyte Growth Factor
-	D23.529.374.440 Interferons
-	D23.529.374.440.890 Interferon Type I
-	D23.529.374.440.890.250 Interferon-alpha
-	D23.529.374.440.890.275 Interferon-beta
-	D23.529.374.440.890.275.500 Interferon beta-1a
-	D23.529.374.440.890.275.750 Interferon beta-1b
-	D23.529.374.440.893 Interferon-gamma
-	D23.529.374.460 Interleukin 1 Receptor Antagonist Protein
-	D23.529.374.465 Interleukins
-	D23.529.374.465.131 Interleukin-1
-	D23.529.374.465.131.300 Interleukin-1alpha
-	D23.529.374.465.131.600 Interleukin-1beta
-	D23.529.374.465.155 Interleukin-2
-	D23.529.374.465.169 Interleukin-3
-	D23.529.374.465.186 Interleukin-4
-	D23.529.374.465.202 Interleukin-5
-	D23.529.374.465.224 Interleukin-6
-	D23.529.374.465.246 Interleukin-7
-	D23.529.374.465.312 Interleukin-8
-	D23.529.374.465.377 Interleukin-9
-	D23.529.374.465.510 Interleukin-10
-	D23.529.374.465.511 Interleukin-11
-	D23.529.374.465.512 Interleukin-12
-	D23.529.374.465.512.249 Interleukin-12 Subunit p35
-	D23.529.374.465.512.500 Interleukin-12 Subunit p40
-	D23.529.374.465.513 Interleukin-13
-	D23.529.374.465.515 Interleukin-15
-	D23.529.374.465.516 Interleukin-16
-	D23.529.374.465.517 Interleukin-17
-	D23.529.374.465.518 Interleukin-18

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D23.529.374.465.550 Interleukin-23
-	D23.529.374.465.550.249 Interleukin-12 Subunit p40
-	D23.529.374.465.550.500 Interleukin-23 Subunit p19
-	D23.529.374.465.600 Interleukin-27
-	D23.529.374.465.850 Interleukin-33
-	D23.529.374.470 Leukemia Inhibitory Factor
-	D23.529.374.480 Lymphokines
-	D23.529.374.480.372 Interleukin-2
-	D23.529.374.480.428 Leukocyte Migration-Inhibitory Factors
-	D23.529.374.480.438 Lymphotoxin-alpha
-	D23.529.374.480.615 Macrophage-Activating Factors
-	D23.529.374.480.615.350 Interferon-gamma
-	D23.529.374.480.625 Macrophage Migration-Inhibitory Factors
-	D23.529.374.480.700 Suppressor Factors, Immunologic
-	D23.529.374.480.750 Transfer Factor
-	D23.529.374.500 Monokines
-	D23.529.374.500.400 Interleukin-1
-	D23.529.374.500.400.300 Interleukin-1alpha
-	D23.529.374.500.400.600 Interleukin-1beta
-	D23.529.374.500.800 Tumor Necrosis Factor-alpha
-	D23.529.374.562 Oncostatin M
-	D23.529.374.625 Osteopontin
-	D23.529.374.687 Transforming Growth Factor beta
-	D23.529.374.687.100 Transforming Growth Factor beta1
-	D23.529.374.687.200 Transforming Growth Factor beta2
-	D23.529.374.750 Tumor Necrosis Factors
-	D23.529.374.750.030 Antigens, CD70
-	D23.529.374.750.061 B-Cell Activating Factor
-	D23.529.374.750.065 4-1BB Ligand
-	D23.529.374.750.092 CD30 Ligand
-	D23.529.374.750.124 CD40 Ligand
-	D23.529.374.750.186 Ectodysplasins
-	D23.529.374.750.249 Fas Ligand Protein
-	D23.529.374.750.500 Lymphotoxin-alpha
-	D23.529.374.750.505 Lymphotoxin alpha1, beta2 Heterotrimer
-	D23.529.374.750.515 Lymphotoxin-beta

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D23.529.374.750.531 OX40 Ligand
-	D23.529.374.750.562 RANK Ligand
-	D23.529.374.750.625 TNF-Related Apoptosis-Inducing Ligand
-	D23.529.374.750.626 Tumor Necrosis Factor-alpha
-	D23.529.374.750.656 Tumor Necrosis Factor Ligand Superfamily Member 13
-	D23.529.374.750.690 Tumor Necrosis Factor Ligand Superfamily Member 14
-	D23.529.374.750.710 Tumor Necrosis Factor Ligand Superfamily Member 15
-	D23.529.382 EGF Family of Proteins
-	D23.529.382.249 Amphiregulin
-	D23.529.382.374 Betacellulin
-	D23.529.382.500 Epidermal Growth Factor
-	D23.529.382.531 Epigen
-	D23.529.382.562 Epiregulin
-	D23.529.382.625 Heparin-binding EGF-like Growth Factor
-	D23.529.382.750 Transforming Growth Factor alpha
-	D23.529.390 Endothelial Growth Factors
-	D23.529.400 Endothelins
-	D23.529.400.225 Endothelin-1
-	D23.529.400.235 Endothelin-2
-	D23.529.400.245 Endothelin-3
-	D23.529.500 Ephrins
-	D23.529.500.100 Ephrin-A1
-	D23.529.500.200 Ephrin-A2
-	D23.529.500.300 Ephrin-A3
-	D23.529.500.400 Ephrin-A4
-	D23.529.500.500 Ephrin-A5
-	D23.529.500.600 Ephrin-B1
-	D23.529.500.700 Ephrin-B2
-	D23.529.500.800 Ephrin-B3
-	D23.529.624 Fibroblast Growth Factors
-	D23.529.624.110 Fibroblast Growth Factor 1
-	D23.529.624.120 Fibroblast Growth Factor 2
-	D23.529.624.130 Fibroblast Growth Factor 3
-	D23.529.624.140 Fibroblast Growth Factor 4
-	D23.529.624.150 Fibroblast Growth Factor 5
-	D23.529.624.160 Fibroblast Growth Factor 6

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D23.529.624.170 Fibroblast Growth Factor 7
-	D23.529.624.180 Fibroblast Growth Factor 8
-	D23.529.624.190 Fibroblast Growth Factor 9
-	D23.529.624.200 Fibroblast Growth Factor 10
-	D23.529.671 Hedgehog Proteins
-	D23.529.812 Kinins
-	D23.529.812.169 Bradykinin
-	D23.529.812.169.400 Kallidin
-	D23.529.812.654 Kininogens
-	D23.529.812.654.350 Kininogen, High-Molecular-Weight
-	D23.529.812.654.400 Kininogen, Low-Molecular-Weight
-	D23.529.812.900 Tachykinins
-	D23.529.812.900.354 Eledoisin
-	D23.529.812.900.475 Kassinin
-	D23.529.812.900.500 Neurokinin A
-	D23.529.812.900.550 Neurokinin B
-	D23.529.812.900.800 Physalaemin
-	D23.529.812.900.866 Substance P
-	D23.529.850 Nerve Growth Factors
-	D23.529.850.100 Brain-Derived Neurotrophic Factor
-	D23.529.850.212 Ciliary Neurotrophic Factor
-	D23.529.850.325 Glia Maturation Factor
-	D23.529.850.381 Glial Cell Line-Derived Neurotrophic Factors
-	D23.529.850.381.500 Glial Cell Line-Derived Neurotrophic Factor
-	D23.529.850.381.750 Neurturin
-	D23.529.850.437 Nerve Growth Factor
-	D23.529.850.550 Neuregulins
-	D23.529.850.550.750 Neuregulin-1
-	D23.529.850.775 Neurotrophin 3
-	D23.529.850.887 Pituitary Adenylate Cyclase-Activating Polypeptide
-	D23.529.890 Parathyroid Hormone-Related Protein
-	D23.529.910 Platelet-Derived Growth Factor
-	D23.529.910.650 Proto-Oncogene Proteins c-sis
-	D23.529.923 Semaphorins
-	D23.529.923.374 Semaphorin-3A
New	<b>D23.529.930 Serrate-Jagged Proteins</b>

## MeSH Tree Changes for 2017

Type	Tree - heading	
Heading		
New Heading	<b>D23.529.930.500</b>	<b>Jagged-1 Protein</b>
New Heading	<b>D23.529.930.750</b>	<b>Jagged-2 Protein</b>
-	D23.529.937	Somatomedins
-	D23.529.937.400	Insulin-Like Growth Factor I
-	D23.529.937.420	Insulin-Like Growth Factor II
-	D23.529.942	TGF-beta Superfamily Proteins
-	D23.529.942.200	Bone Morphogenetic Proteins
-	D23.529.942.200.100	Bone Morphogenetic Protein 1
-	D23.529.942.200.200	Bone Morphogenetic Protein 2
-	D23.529.942.200.300	Bone Morphogenetic Protein 3
-	D23.529.942.200.400	Bone Morphogenetic Protein 4
-	D23.529.942.200.500	Bone Morphogenetic Protein 5
-	D23.529.942.200.600	Bone Morphogenetic Protein 6
-	D23.529.942.200.700	Bone Morphogenetic Protein 7
-	D23.529.942.200.900	Bone Morphogenetic Protein 15
-	D23.529.942.200.910	Growth Differentiation Factor 2
-	D23.529.942.200.950	Growth Differentiation Factor 10
-	D23.529.942.300	Growth Differentiation Factors
-	D23.529.942.300.049	Bone Morphogenetic Protein 15
-	D23.529.942.300.100	Growth Differentiation Factor 1
-	D23.529.942.300.200	Growth Differentiation Factor 2
-	D23.529.942.300.300	Growth Differentiation Factor 3
-	D23.529.942.300.500	Growth Differentiation Factor 5
-	D23.529.942.300.600	Growth Differentiation Factor 6
-	D23.529.942.300.800	Growth Differentiation Factor 9
-	D23.529.942.300.900	Growth Differentiation Factor 10
-	D23.529.942.300.915	Growth Differentiation Factor 15
-	D23.529.942.300.925	Myostatin
-	D23.529.942.550	Nodal Signaling Ligands
-	D23.529.942.550.100	Growth Differentiation Factor 1
-	D23.529.942.550.300	Growth Differentiation Factor 3
-	D23.529.942.550.475	Left-Right Determination Factors
-	D23.529.942.550.650	Nodal Protein

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D23.529.942.775	Transforming Growth Factor beta
-	D23.529.942.775.100	Transforming Growth Factor beta1
-	D23.529.942.775.200	Transforming Growth Factor beta2
-	D23.529.942.775.300	Transforming Growth Factor beta3
-	D23.529.948	Tolloid-Like Metalloproteinases
-	D23.529.948.500	Bone Morphogenetic Protein 1
-	D23.529.960	Transforming Growth Factors
-	D23.529.960.360	Transforming Growth Factor alpha
-	D23.529.984	Wnt Proteins
New Heading	<b>D23.529.984.050</b>	<b>Wnt-5a Protein</b>
-	D23.529.984.100	Wnt1 Protein
-	D23.529.984.200	Wnt2 Protein
-	D23.529.984.300	Wnt3 Protein
-	D23.529.984.350	Wnt3A Protein
-	D23.529.984.400	Wnt4 Protein
-	D23.585	Pathogen-Associated Molecular Pattern Molecules
-	D23.641	Pheromones
New Heading	<b>D23.641.200</b>	<b>Mating Factor</b>
-	D23.641.399	Pheromones, Human
-	D23.641.800	Sex Attractants
New Tree	<b>D23.704</b>	<b>Phytochemicals</b>
New Tree	<b>D23.704.500</b>	<b>Phytosterols</b>
New Tree	<b>D23.704.500.071</b>	<b>Brassinosteroids</b>
New Tree	<b>D23.704.500.143</b>	<b>Ecdysteroids</b>
New Tree	<b>D23.704.500.669</b>	<b>Sitosterols</b>
New Tree	<b>D23.704.500.808</b>	<b>Stigmasterol</b>
New Tree	<b>D23.704.500.904</b>	<b>Withanolides</b>
-	D23.767	Pigments, Biological
-	D23.767.061	Adrenochrome
-	D23.767.124	Anthocyanins



## MeSH Tree Changes for 2017

Type	Tree - heading
-	D23.767.193 Bile Pigments
-	D23.767.193.184 Bilirubin
-	D23.767.193.184.200 Biliverdine
-	D23.767.193.727 Urobilin
-	D23.767.193.852 Urobilinogen
-	D23.767.261 Carotenoids
-	D23.767.261.050 beta Carotene
-	D23.767.261.700 Retinoids
-	D23.767.261.700.050 Acitretin
-	D23.767.261.700.250 Etretnate
-	D23.767.261.700.270 Fenretinide
-	D23.767.261.700.325 Isotretinoin
-	D23.767.261.700.690 Retinaldehyde
-	D23.767.261.700.780 Tretinoin
-	D23.767.261.700.860 Vitamin A
-	D23.767.261.887 Xanthophylls
-	D23.767.261.887.249 Canthaxanthin
-	D23.767.261.887.374 Cryptoxanthins
New Heading	<b>D23.767.261.887.374.500 Beta-Cryptoxanthin</b>
-	D23.767.261.887.500 Lutein
-	D23.767.261.887.625 Macular Pigment
-	D23.767.261.887.750 Zeaxanthins
-	D23.767.261.925 zeta Carotene
-	D23.767.300 Ceroid
-	D23.767.405 Flavins
-	D23.767.405.650 Riboflavin
-	D23.767.405.650.249 Flavin-Adenine Dinucleotide
-	D23.767.405.650.500 Flavin Mononucleotide
-	D23.767.482 Hemocyanin
-	D23.767.550 Lipofuscin
-	D23.767.620 Melanins
-	D23.767.690 Phycocyanin
-	D23.767.705 Phycoerythrin
-	D23.767.712 Phytochrome
-	D23.767.712.249 Phytochrome A

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D23.767.712.500                      Phytochrome B
-	D23.767.727                              Porphyrins
-	D23.767.727.250                      Coproporphyrins
-	D23.767.727.280                      Deuteroporphyrins
-	D23.767.727.340                      Etioporphyrins
-	D23.767.727.462                      Hematoporphyrins
-	D23.767.727.462.400                      Hematoporphyrin Derivative
-	D23.767.727.462.400.200                      Dihematoporphyrin Ether
-	D23.767.727.620                      Mesoporphyrins
-	D23.767.727.640                      Metalloporphyrins
-	D23.767.727.640.220                      Chlorophyll
-	D23.767.727.640.220.100                      Bacteriochlorophylls
-	D23.767.727.640.220.180                      Chlorophyllides
-	D23.767.727.640.220.453                      Pheophytins
-	D23.767.727.640.220.725                      Protochlorophyllide
-	D23.767.727.640.587                      Heme
-	D23.767.727.640.587.462                      Hemin
-	D23.767.727.725                      Protoporphyrins
-	D23.767.727.880                      Uroporphyrins
-	D23.767.778                              Prodigiosin
-	D23.767.831                              Pterins
-	D23.767.831.742                      Xanthopterin
-	D23.767.860                              Pyocyanine
-	D23.767.930                              Retinal Pigments
-	D23.767.930.750                      Opsins
-	D23.767.930.750.249                      Cone Opsins
-	D23.767.930.750.500                      Rod Opsins
-	D23.767.930.750.500.500                      Rhodopsin
-	D23.946                                      Toxins, Biological
-	D23.946.123                              Bacterial Toxins
-	D23.946.123.179                      Botulinum Toxins
-	D23.946.123.179.050                      Botulinum Toxins, Type A
-	D23.946.123.194                      Cholera Toxin
-	D23.946.123.208                      Cord Factors
-	D23.946.123.305                      Diphtheria Toxin
-	D23.946.123.329                      Endotoxins

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D23.946.123.329.500 Lipopolysaccharides
-	D23.946.123.329.500.500 Lipid A
-	D23.946.123.329.500.600 O Antigens
-	D23.946.123.353 Exfoliatins
-	D23.946.123.794 Shiga Toxins
-	D23.946.123.794.095 Shiga Toxin
-	D23.946.123.794.100 Shiga Toxin 1
-	D23.946.123.794.124 Shiga Toxin 2
-	D23.946.123.868 Streptolysins
-	D23.946.123.893 Tetanus Toxin
-	D23.946.123.946 Virulence Factors, Bordetella
-	D23.946.123.946.040 Adenylate Cyclase Toxin
-	D23.946.123.946.690 Pertussis Toxin
-	D23.946.330 Enterotoxins
-	D23.946.330.150 Cholera Toxin
-	D23.946.330.575 Shiga Toxins
-	D23.946.330.575.100 Shiga Toxin
-	D23.946.330.575.120 Shiga Toxin 1
-	D23.946.330.575.124 Shiga Toxin 2
-	D23.946.350 Exotoxins
-	D23.946.350.500 Exfoliatins
-	D23.946.350.750 Streptolysins
-	D23.946.412 Hypoglycins
-	D23.946.580 Marine Toxins
-	D23.946.580.160 Ciguatoxins
-	D23.946.580.230 Cnidarian Venoms
-	D23.946.580.370 Fish Venoms
-	D23.946.580.450 Holothurin
-	D23.946.580.565 Lyngbya Toxins
-	D23.946.580.590 Mollusk Venoms
-	D23.946.580.590.162 Conotoxins
-	D23.946.580.590.162.720 omega-Conotoxins
-	D23.946.580.590.162.720.700 omega-Conotoxin GVIA
-	D23.946.580.590.325 Eledoisin
-	D23.946.580.830 Saxitoxin
-	D23.946.580.910 Tetrodotoxin

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D23.946.587 Mycotoxins
-	D23.946.587.142 Aflatoxins
-	D23.946.587.142.075 Aflatoxin B1
-	D23.946.587.142.100 Aflatoxin M1
-	D23.946.587.175 Amanitins
-	D23.946.587.175.111 Alpha-Amanitin
-	D23.946.587.272 Citrinin
-	D23.946.587.370 Cytochalasins
-	D23.946.587.370.370 Cytochalasin B
-	D23.946.587.370.450 Cytochalasin D
-	D23.946.587.385 Fumonisin
-	D23.946.587.401 Gliotoxin
-	D23.946.587.475 Ibotenic Acid
-	D23.946.587.531 Killer Factors, Yeast
-	D23.946.587.587 Muscimol
-	D23.946.587.697 Ochratoxins
-	D23.946.587.725 Patulin
-	D23.946.587.735 Penicillic Acid
-	D23.946.587.755 Phalloidine
-	D23.946.587.800 Sporidesmins
-	D23.946.587.850 Sterigmatocystin
-	D23.946.587.915 Tenuazonic Acid
-	D23.946.587.933 Trichothecenes
-	D23.946.587.933.870 T-2 Toxin
-	D23.946.587.933.900 Trichodermin
-	D23.946.587.989 Zearalenone
-	D23.946.833 Venoms
-	D23.946.833.033 Amphibian Venoms
-	D23.946.833.033.112 Batrachotoxins
-	D23.946.833.033.137 Bombesin
-	D23.946.833.033.163 Bufotenin
-	D23.946.833.033.728 Physalaemin
-	D23.946.833.065 Arthropod Venoms
-	D23.946.833.065.055 Ant Venoms
-	D23.946.833.065.115 Bee Venoms
-	D23.946.833.065.115.060 Apamin

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D23.946.833.065.115.580 Melitten
-	D23.946.833.065.830 Scorpion Venoms
-	D23.946.833.065.830.150 Charybdotoxin
-	D23.946.833.065.870 Spider Venoms
-	D23.946.833.065.870.324 Agatoxins
-	D23.946.833.065.870.324.500 omega-Agatoxin IVA
-	D23.946.833.065.970 Wasp Venoms
-	D23.946.833.230 Cnidarian Venoms
-	D23.946.833.370 Fish Venoms
-	D23.946.833.590 Mollusk Venoms
-	D23.946.833.590.162 Conotoxins
-	D23.946.833.590.162.720 omega-Conotoxins
-	D23.946.833.590.162.720.700 omega-Conotoxin GVIA
-	D23.946.833.590.325 Eledoisin
-	D23.946.833.850 Snake Venoms
-	D23.946.833.850.325 Elapid Venoms
-	D23.946.833.850.325.139 Bungarotoxins
-	D23.946.833.850.325.220 Cobra Venoms
-	D23.946.833.850.325.220.222 Cobra Cardiotoxin Proteins
-	D23.946.833.850.325.220.244 Cobra Neurotoxin Proteins
-	D23.946.833.850.960 Viper Venoms
-	D23.946.833.850.960.200 Crotalid Venoms
-	D23.946.833.850.960.200.050 Ancrod
-	D23.946.833.850.960.200.105 Batroxobin
-	D23.946.833.850.960.200.210 Crotoxin
-	D23.946.896 Virulence Factors
New Heading	<b>D23.946.896.490 Transcription Activator-Like Effectors</b>
-	D23.946.896.980 Virulence Factors, Bordetella
-	D23.946.896.980.040 Adenylate Cyclase Toxin
-	D23.946.896.980.690 Pertussis Toxin
-	D25 Biomedical and Dental Materials
-	D25.058 Alloys
-	D25.058.224 Chromium Alloys
-	D25.058.224.735 Vitallium
-	D25.058.451 Gold Alloys

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D25.058.520 Metal Ceramic Alloys
-	D25.058.520.250 Cermet Cements
-	D25.058.807 Steel
-	D25.058.807.681 Stainless Steel
-	D25.130 Biocompatible Materials
-	D25.130.325 Bone Substitutes
-	D25.130.420 Coated Materials, Biocompatible
-	D25.130.650 Polydioxanone
-	D25.187 Cariogenic Agents
-	D25.223 Cariostatic Agents
-	D25.223.015 Acidulated Phosphate Fluoride
-	D25.223.432 Fluorides, Topical
-	D25.223.716 Sodium Fluoride
-	D25.223.800 Tin Fluorides
-	D25.339 Dental Materials
-	D25.339.208 Dental Alloys
-	D25.339.208.224 Chromium Alloys
-	D25.339.208.224.959 Vitallium
-	D25.339.208.291 Dental Amalgam
-	D25.339.208.534 Gold Alloys
-	D25.339.208.720 Metal Ceramic Alloys
-	D25.339.208.720.250 Cermet Cements
-	D25.339.250 Dental Casting Investment
-	D25.339.291 Dental Cements
-	D25.339.291.150 Compomers
-	D25.339.291.300 Dentin-Bonding Agents
-	D25.339.291.300.500 Silorane Resins
-	D25.339.291.402 Glass Ionomer Cements
-	D25.339.291.402.120 Cermet Cements
-	D25.339.291.551 Organically Modified Ceramics
-	D25.339.291.700 Polycarboxylate Cement
-	D25.339.291.750 Resin Cements
-	D25.339.291.800 Silicate Cement
-	D25.339.291.925 Zinc Oxide-Eugenol Cement
-	D25.339.291.950 Zinc Phosphate Cement
-	D25.339.312 Dental Implants

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D25.339.334 Dental Impression Materials
-	D25.339.334.574 Inlay Casting Wax
-	D25.339.376 Dental Porcelain
-	D25.339.574 Photoinitiators, Dental
-	D25.339.773 Pit and Fissure Sealants
-	D25.339.788 Pulp Capping and Pulpectomy Agents
-	D25.339.816 Resins, Synthetic
-	D25.339.816.500 Composite Resins
-	D25.339.816.500.200 Bisphenol A-Glycidyl Methacrylate
-	D25.339.816.500.300 Compomers
-	D25.339.859 Root Canal Filling Materials
-	D25.339.859.495 Gutta-Percha
-	D25.376 Dentifrices
-	D25.376.262 Denture Cleansers
-	D25.376.711 Toothpastes
-	D25.479 Membranes, Artificial
-	D25.479.517 Liposomes
-	D25.479.517.500 Unilamellar Liposomes
-	D25.479.900 Virosomes
-	D25.583 Mouthwashes
-	D25.583.820 Saliva, Artificial
-	D25.651 Pharmaceutical Preparations, Dental
-	D25.720 Polymers
-	D25.720.099 Biopolymers
-	D25.720.099.500 Cellulose
-	D25.720.099.500.252 Cellophane
-	D25.720.099.500.439 Collodion
-	D25.720.099.500.719 Hypromellose Derivatives
-	D25.720.099.625 Latex
-	D25.720.099.687 Lignin
-	D25.720.099.718 Plant Mucilage
-	D25.720.099.750 Rubber
-	D25.720.099.750.500 Latex
-	D25.720.200 Colestipol
-	D25.720.259 Cyanoacrylates
-	D25.720.259.341 Enbucrilate

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D25.720.259.341.110 Bucrylate
-	D25.720.327 Elastomers
-	D25.720.327.390 Neoprene
-	D25.720.327.782 Polyurethanes
-	D25.720.327.811 Polyvinyl Chloride
-	D25.720.327.840 Rubber
-	D25.720.327.840.119 Gutta-Percha
-	D25.720.327.840.239 Latex
-	D25.720.327.900 Silicone Elastomers
-	D25.720.395 Fluorocarbon Polymers
-	D25.720.395.616 Polytetrafluoroethylene
-	D25.720.395.616.755 Proplast
-	D25.720.470 Hexadimethrine Bromide
-	D25.720.593 Organically Modified Ceramics
-	D25.720.716 Plastics
-	D25.720.716.195 Biodegradable Plastics
-	D25.720.716.392 Nylons
-	D25.720.716.507 Polyethylenes
-	D25.720.716.507.500 Polyethylene
-	D25.720.716.507.600 Polyethyleneimine
-	D25.720.716.550 Polypropylenes
-	D25.720.716.579 Polystyrenes
-	D25.720.716.579.159 Cholestyramine Resin
-	D25.720.716.650 Polyurethanes
-	D25.720.716.721 Polyvinyls
-	D25.720.716.721.616 Polyvinyl Alcohol
-	D25.720.716.721.812 Polyvinyl Chloride
-	D25.720.716.721.838 Povidone
-	D25.720.716.721.838.745 Povidone-Iodine
-	D25.720.716.822 Resins, Synthetic
-	D25.720.716.822.111 Acrylic Resins
-	D25.720.716.822.111.650 Polymethacrylic Acids
-	D25.720.716.822.111.650.605 Methylmethacrylates
-	D25.720.716.822.111.650.605.450 Methylmethacrylate
-	D25.720.716.822.111.650.605.500 Polymethyl Methacrylate
-	D25.720.716.822.111.650.750 Polyhydroxyethyl Methacrylate



## MeSH Tree Changes for 2017

Type	Tree - heading
-	D25.720.716.822.300 Bone Cements
-	D25.720.716.822.308 Composite Resins
-	D25.720.716.822.308.200 Bisphenol A-Glycidyl Methacrylate
-	D25.720.716.822.308.300 Compomers
-	D25.720.716.822.461 Epoxy Resins
-	D25.720.716.822.730 Resin Cements
-	D25.720.722 Polyanetholesulfonate
-	D25.720.728 Polyesters
-	D25.720.728.700 Polydioxanone
-	D25.720.728.764 Polyethylene Terephthalates
-	D25.720.728.772 Polyglactin 910
-	D25.720.728.780 Polyglycolic Acid
-	D25.720.741 Polyethylene Glycols
-	D25.720.741.125 Certolizumab Pegol
-	D25.720.741.250 Cetomacrogol
-	D25.720.741.485 Hydrogel
-	D25.720.741.575 Nonoxynol
-	D25.720.741.610 Octoxynol
-	D25.720.741.650 Poloxalene
-	D25.720.741.667 Poloxamer
-	D25.720.741.685 Polyhydroxyethyl Methacrylate
-	D25.720.741.700 Polysorbates
-	D25.720.780 Polygeline
-	D25.720.795 Polyphloreitin Phosphate
-	D25.720.830 Pyran Copolymer
-	D25.720.900 Siloxanes
-	D25.720.900.850 Silicones
-	D25.720.900.850.150 Dimethylpolysiloxanes
-	D25.720.900.850.150.750 Simethicone
-	D25.720.900.850.900 Silicone Elastomers
-	D25.720.900.850.905 Silicone Gels
-	D25.720.900.850.950 Silicone Oils
-	D25.800 Root Canal Irrigants
-	D25.919 Tissue Adhesives
-	D25.919.367 Cyanoacrylates
-	D25.919.367.341 Enbucrilate

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D25.919.367.341.110 Bucrylate
-	D26 Pharmaceutical Preparations
-	D26.049 Controlled Substances
-	D26.074 Cosmeceuticals
-	D26.099 Counterfeit Drugs
-	D26.200 Designer Drugs
-	D26.255 Dosage Forms
-	D26.255.150 Capsules
-	D26.255.165 Colloids
-	D26.255.165.055 Aerosols
-	D26.255.165.055.055 Aerosol Propellants
-	D26.255.165.055.291 Nasal Sprays
-	D26.255.165.055.527 Oral Sprays
-	D26.255.165.260 Emulsions
-	D26.255.165.260.270 Fat Emulsions, Intravenous
-	D26.255.165.320 Gels
-	D26.255.165.320.375 Hydrogels
-	D26.255.165.320.375.187 Cryogels
-	D26.255.165.320.375.375 Hydrogel
-	D26.255.165.320.687 Silica Gel
-	D26.255.165.810 Suspensions
-	D26.255.210 Delayed-Action Preparations
-	D26.255.210.315 Drug Implants
-	D26.255.210.860 Tablets, Enteric-Coated
-	D26.255.260 Drug Carriers
-	D26.255.260.517 Liposomes
-	D26.255.260.575 Nanocapsules
-	D26.255.260.600 Nanoconjugates
-	D26.255.260.900 Virosomes
-	D26.255.480 Liniments
-	D26.255.560 Micelles
-	D26.255.640 Ointments
-	D26.255.779 Powders
-	D26.255.785 Suppositories
-	D26.255.830 Tablets
-	D26.255.830.860 Tablets, Enteric-Coated

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D26.255.955 Vaginal Creams, Foams, and Jellies
-	D26.310 Drug Combinations
-	D26.310.032 Adapalene, Benzoyl Peroxide Drug Combination
-	D26.310.048 Aminophylline
-	D26.310.063 Amlodipine Besylate, Olmesartan Medoxomil Drug Combination
-	D26.310.094 Amlodipine, Valsartan Drug Combination
-	D26.310.125 Aspirin, Dipyridamole Drug Combination
-	D26.310.249 Biphasic Insulins
-	D26.310.312 Brimonidine Tartrate, Timolol Maleate Drug Combination
-	D26.310.344 Albuterol, Ipratropium Drug Combination
-	D26.310.348 Budesonide, Formoterol Fumarate Drug Combination
-	D26.310.352 Buprenorphine, Naloxone Drug Combination
-	D26.310.360 Contraceptives, Oral, Combined
-	D26.310.368 Dimenhydrinate
-	D26.310.375 Efavirenz, Emtricitabine, Tenofovir Disoproxil Fumarate Drug Combination
-	D26.310.391 Elvitegravir, Cobicistat, Emtricitabine, Tenofovir Disoproxil Fumarate Drug Combination
-	D26.310.407 Emtricitabine, Rilpivirine, Tenofovir Drug Combination
-	D26.310.423 Emtricitabine, Tenofovir Disoproxil Fumarate Drug Combination
-	D26.310.431 Ezetimibe, Simvastatin Drug Combination
-	D26.310.438 Fluticasone Propionate, Salmeterol Xinafoate Drug Combination
-	D26.310.469 Mometasone Furoate, Formoterol Fumarate Drug Combination
-	D26.310.500 Multi-Ingredient Cold, Flu, and Allergy Medications
-	D26.310.750 Sitagliptin Phosphate, Metformin Hydrochloride Drug Combination
-	D26.310.875 Trimethoprim, Sulfamethoxazole Drug Combination
-	D26.335 Drugs, Chinese Herbal
-	D26.355 Drugs, Essential
-	D26.360 Drugs, Generic
-	D26.371 Drugs, Investigational
-	D26.526 Materia Medica
-	D26.528 Medical Marijuana
-	D26.530 Nonprescription Drugs
-	D26.530.500 Behind-the-Counter Drugs
-	D26.593 Nostrums
-	D26.650 Pharmaceutic Aids

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D26.650.064                      Adjuvants, Pharmaceutic
-	D26.650.523                      Ointment Bases
-	D26.650.700                      Pharmaceutical Vehicles
-	D26.650.700.419                  Excipients
-	D26.650.702                      Preservatives, Pharmaceutical
-	D26.655                            Pharmaceutical Preparations, Dental
-	D26.660                            Placebos
-	D26.667                            Plant Extracts
-	D26.667.249                      Flower Essences
-	D26.667.500                      Grape Seed Extract
-	D26.670                            Prescription Drugs
-	D26.675                            Prodrugs
-	D26.776                            Solutions
-	D26.776.210                      Contact Lens Solutions
-	D26.776.314                      Hypertonic Solutions
-	D26.776.314.420                  Glucose Solution, Hypertonic
-	D26.776.314.890                  Saline Solution, Hypertonic
-	D26.776.399                      Hypotonic Solutions
-	D26.776.498                      Isotonic Solutions
-	D26.776.675                      Organ Preservation Solutions
-	D26.776.708                      Pharmaceutical Solutions
-	D26.776.708.160                  Cardioplegic Solutions
-	D26.776.708.322                  Dialysis Solutions
-	D26.776.708.322.651              Hemodialysis Solutions
-	D26.776.708.645                  Ophthalmic Solutions
-	D26.776.708.645.500              Lubricant Eye Drops
-	D26.776.708.733                  Parenteral Nutrition Solutions
-	D26.776.708.733.500              Fat Emulsions, Intravenous
-	D26.776.708.822                  Sclerosing Solutions
-	D26.776.741                      Rehydration Solutions
-	D26.878                            Street Drugs
-	D26.878.250                      Crack Cocaine
-	D26.939                            Veterinary Drugs
-	D26.969                            Xenobiotics
-	D27                                  Chemical Actions and Uses
-	D27.505                            Pharmacologic Actions

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D27.505.259 Diagnostic Uses of Chemicals
-	D27.505.259.500 Contrast Media
-	D27.505.259.750 Molecular Probes
-	D27.505.259.750.600 Nucleic Acid Probes
-	D27.505.259.750.600.223 DNA Probes
-	D27.505.259.750.600.223.615 DNA Probes, HLA
-	D27.505.259.750.600.223.620 DNA Probes, HPV
-	D27.505.259.750.600.650 Oligonucleotide Probes
-	D27.505.259.750.600.825 RNA Probes
-	D27.505.259.812 Neuronal Tract-Tracers
-	D27.505.259.843 Radiopharmaceuticals
-	D27.505.259.875 Reagent Kits, Diagnostic
-	D27.505.259.875.680 Reagent Strips
-	D27.505.389 Metabolic Side Effects of Drugs and Substances
-	D27.505.389.249 Cytochrome P-450 Enzyme Inducers
-	D27.505.389.249.120 Cytochrome P-450 CYP1A2 Inducers
-	D27.505.389.249.260 Cytochrome P-450 CYP2B6 Inducers
-	D27.505.389.249.308 Cytochrome P-450 CYP2C8 Inducers
-	D27.505.389.249.320 Cytochrome P-450 CYP2C9 Inducers
-	D27.505.389.249.350 Cytochrome P-450 CYP2C19 Inducers
-	D27.505.389.249.500 Cytochrome P-450 CYP2D6 Inducers
-	D27.505.389.249.600 Cytochrome P-450 CYP2E1 Inducers
-	D27.505.389.249.700 Cytochrome P-450 CYP3A Inducers
-	D27.505.389.500 Cytochrome P-450 Enzyme Inhibitors
-	D27.505.389.500.059 14-alpha Demethylase Inhibitors
-	D27.505.389.500.120 Cytochrome P-450 CYP1A2 Inhibitors
-	D27.505.389.500.260 Cytochrome P-450 CYP2B6 Inhibitors
-	D27.505.389.500.308 Cytochrome P-450 CYP2C8 Inhibitors
-	D27.505.389.500.315 Cytochrome P-450 CYP2C9 Inhibitors
-	D27.505.389.500.319 Cytochrome P-450 CYP2C19 Inhibitors
-	D27.505.389.500.368 Cytochrome P-450 CYP2D6 Inhibitors
-	D27.505.389.500.421 Cytochrome P-450 CYP2E1 Inhibitors
-	D27.505.389.500.503 Cytochrome P-450 CYP3A Inhibitors
-	D27.505.519 Molecular Mechanisms of Pharmacological Action
-	D27.505.519.124 Alkylating Agents
-	D27.505.519.124.035 Antineoplastic Agents, Alkylating

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D27.505.519.143 Amylin Receptor Agonists
-	D27.505.519.162 Angiotensin Receptor Antagonists
-	D27.505.519.162.500 Angiotensin II Type 1 Receptor Blockers
-	D27.505.519.162.750 Angiotensin II Type 2 Receptor Blockers
-	D27.505.519.170 Antacids
-	D27.505.519.174 Antidiuretic Hormone Receptor Antagonists
-	D27.505.519.178 Antifoaming Agents
-	D27.505.519.186 Antimetabolites
-	D27.505.519.186.071 Hypolipidemic Agents
-	D27.505.519.186.071.202 Anticholesteremic Agents
-	D27.505.519.186.071.202.370 Hydroxymethylglutaryl-CoA Reductase Inhibitors
-	D27.505.519.186.071.401 Fatty Acid Synthesis Inhibitors
-	D27.505.519.186.071.601 Lipotropic Agents
-	D27.505.519.186.144 Antimetabolites, Antineoplastic
-	D27.505.519.217 Antioxidants
-	D27.505.519.217.500 Free Radical Scavengers
-	D27.505.519.265 Bradykinin Receptor Antagonists
-	D27.505.519.265.249 Bradykinin B1 Receptor Antagonists
-	D27.505.519.265.500 Bradykinin B2 Receptor Antagonists
-	D27.505.519.275 CCR5 Receptor Antagonists
-	D27.505.519.295 Cerumenolytic Agents
-	D27.505.519.334 Cystine Depleting Agents
-	D27.505.519.349 Cytochrome P-450 Enzyme Inducers
-	D27.505.519.349.120 Cytochrome P-450 CYP1A2 Inducers
-	D27.505.519.349.260 Cytochrome P-450 CYP2B6 Inducers
-	D27.505.519.349.308 Cytochrome P-450 CYP2C8 Inducers
-	D27.505.519.349.320 Cytochrome P-450 CYP2C9 Inducers
-	D27.505.519.349.350 Cytochrome P-450 CYP2C19 Inducers
-	D27.505.519.349.500 Cytochrome P-450 CYP2D6 Inducers
-	D27.505.519.349.600 Cytochrome P-450 CYP2E1 Inducers
-	D27.505.519.349.700 Cytochrome P-450 CYP3A Inducers
-	D27.505.519.364 Endothelin Receptor Antagonists
-	D27.505.519.364.500 Endothelin A Receptor Antagonists
-	D27.505.519.364.750 Endothelin B Receptor Antagonists
-	D27.505.519.374 Enzyme Activators

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D27.505.519.374.099 Carbamoyl Phosphate Synthetase I Activators
-	D27.505.519.374.200 GTP Phosphohydrolase Activators
-	D27.505.519.374.600 Lipoprotein Lipase Activators
-	D27.505.519.389 Enzyme Inhibitors
-	D27.505.519.389.086 5-Lipoxygenase-Activating Protein Inhibitors
-	D27.505.519.389.089 Acetaldehyde Dehydrogenase Inhibitors
-	D27.505.519.389.092 Adenosine Deaminase Inhibitors
-	D27.505.519.389.108 Adenylyl Cyclase Inhibitors
-	D27.505.519.389.124 Aromatic Amino Acid Decarboxylase Inhibitors
-	D27.505.519.389.174 Calcineurin Inhibitors
-	D27.505.519.389.200 Carbonic Anhydrase Inhibitors
-	D27.505.519.389.237 Catechol O-Methyltransferase Inhibitors
-	D27.505.519.389.275 Cholinesterase Inhibitors
-	D27.505.519.389.310 Cyclooxygenase Inhibitors
-	D27.505.519.389.310.500 Cyclooxygenase 2 Inhibitors
-	D27.505.519.389.320 Glycoside Hydrolase Inhibitors
-	D27.505.519.389.335 Cytochrome P-450 Enzyme Inhibitors
-	D27.505.519.389.335.059 14-alpha Demethylase Inhibitors
-	D27.505.519.389.335.120 Cytochrome P-450 CYP1A2 Inhibitors
-	D27.505.519.389.335.260 Cytochrome P-450 CYP2B6 Inhibitors
-	D27.505.519.389.335.308 Cytochrome P-450 CYP2C8 Inhibitors
-	D27.505.519.389.335.319 Cytochrome P-450 CYP2C19 Inhibitors
-	D27.505.519.389.335.329 Cytochrome P-450 CYP2C9 Inhibitors
-	D27.505.519.389.335.368 Cytochrome P-450 CYP2D6 Inhibitors
-	D27.505.519.389.335.421 Cytochrome P-450 CYP2E1 Inhibitors
-	D27.505.519.389.335.503 Cytochrome P-450 CYP3A Inhibitors
-	D27.505.519.389.350 Folic Acid Antagonists
-	D27.505.519.389.360 Histone Deacetylase Inhibitors
-	D27.505.519.389.370 Hydroxymethylglutaryl-CoA Reductase Inhibitors
-	D27.505.519.389.375 Integrase Inhibitors
-	D27.505.519.389.375.400 HIV Integrase Inhibitors
-	D27.505.519.389.400 beta-Lactamase Inhibitors
-	D27.505.519.389.480 Lipoxygenase Inhibitors
-	D27.505.519.389.616 Monoamine Oxidase Inhibitors
-	D27.505.519.389.675 Nucleic Acid Synthesis Inhibitors
-	D27.505.519.389.675.850 Reverse Transcriptase Inhibitors

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D27.505.519.389.705 Ornithine Decarboxylase Inhibitors
-	D27.505.519.389.735 Phosphodiesterase Inhibitors
-	D27.505.519.389.735.249 Phosphodiesterase 3 Inhibitors
-	D27.505.519.389.735.374 Phosphodiesterase 4 Inhibitors
-	D27.505.519.389.735.500 Phosphodiesterase 5 Inhibitors
-	D27.505.519.389.737 Phospholipase A2 Inhibitors
-	D27.505.519.389.739 Poly(ADP-ribose) Polymerase Inhibitors
-	D27.505.519.389.740 Prolyl-Hydroxylase Inhibitors
-	D27.505.519.389.745 Protease Inhibitors
-	D27.505.519.389.745.085 Angiotensin-Converting Enzyme Inhibitors
-	D27.505.519.389.745.325 Cysteine Proteinase Inhibitors
-	D27.505.519.389.745.325.500 Caspase Inhibitors
-	D27.505.519.389.745.335 Dipeptidyl-Peptidase IV Inhibitors
-	D27.505.519.389.745.420 HIV Protease Inhibitors
-	D27.505.519.389.745.610 Matrix Metalloproteinase Inhibitors
-	D27.505.519.389.745.705 Proteasome Inhibitors
-	D27.505.519.389.745.800 Serine Proteinase Inhibitors
-	D27.505.519.389.745.800.449 Antithrombins
-	D27.505.519.389.745.800.449.500 Factor Xa Inhibitors
-	D27.505.519.389.745.800.900 Trypsin Inhibitors
-	D27.505.519.389.755 Protein Kinase Inhibitors
-	D27.505.519.389.760 Protein Synthesis Inhibitors
-	D27.505.519.389.848 Proton Pump Inhibitors
-	D27.505.519.389.870 Steroid Synthesis Inhibitors
-	D27.505.519.389.870.100 14-alpha Demethylase Inhibitors
-	D27.505.519.389.870.200 5-alpha Reductase Inhibitors
-	D27.505.519.389.870.300 Aromatase Inhibitors
-	D27.505.519.389.892 Topoisomerase Inhibitors
-	D27.505.519.389.892.500 Topoisomerase I Inhibitors
-	D27.505.519.389.892.750 Topoisomerase II Inhibitors
-	D27.505.519.389.936 Uncoupling Agents
-	D27.505.519.389.936.500 Proton Ionophores
-	D27.505.519.405 Enzyme Reactivators
-	D27.505.519.405.347 Cholinesterase Reactivators
-	D27.505.519.421 Fibrin Modulating Agents
-	D27.505.519.421.500 Antifibrinolytic Agents



## MeSH Tree Changes for 2017

Type	Tree - heading
-	D27.505.519.421.750 Fibrinolytic Agents
-	D27.505.519.452 Heparin Antagonists
-	D27.505.519.562 Membrane Transport Modulators
-	D27.505.519.562.061 Acetylcholine Release Inhibitors
-	D27.505.519.562.124 Calcium Channel Agonists
-	D27.505.519.562.249 Calcium Channel Blockers
-	D27.505.519.562.311 Chloride Channel Agonists
-	D27.505.519.562.374 Ionophores
-	D27.505.519.562.374.100 Calcium Ionophores
-	D27.505.519.562.374.400 Potassium Ionophores
-	D27.505.519.562.374.500 Proton Ionophores
-	D27.505.519.562.374.750 Sodium Ionophores
-	D27.505.519.562.437 Neurotransmitter Uptake Inhibitors
-	D27.505.519.562.437.050 Adrenergic Uptake Inhibitors
-	D27.505.519.562.437.220 Dopamine Uptake Inhibitors
-	D27.505.519.562.437.535 GABA Uptake Inhibitors
-	D27.505.519.562.437.693 Serotonin and Noradrenaline Reuptake Inhibitors
-	D27.505.519.562.437.850 Serotonin Uptake Inhibitors
-	D27.505.519.562.500 Potassium Channel Blockers
-	D27.505.519.562.625 Sodium Channel Agonists
-	D27.505.519.562.625.249 Epithelial Sodium Channel Agonists
-	D27.505.519.562.625.500 Voltage-Gated Sodium Channel Agonists
-	D27.505.519.562.750 Sodium Channel Blockers
-	D27.505.519.562.750.100 Acid Sensing Ion Channel Blockers
-	D27.505.519.562.750.374 Epithelial Sodium Channel Blockers
-	D27.505.519.562.750.500 Voltage-Gated Sodium Channel Blockers
-	D27.505.519.562.812 Sodium Chloride Symporter Inhibitors
-	D27.505.519.562.906 Sodium Potassium Chloride Symporter Inhibitors
-	D27.505.519.593 Mitosis Modulators
-	D27.505.519.593.249 Antimitotic Agents
-	D27.505.519.593.249.500 Tubulin Modulators
-	D27.505.519.593.624 Mitogens
-	D27.505.519.625 Neurotransmitter Agents
-	D27.505.519.625.050 Adrenergic Agents
-	D27.505.519.625.050.100 Adrenergic Agonists
-	D27.505.519.625.050.100.100 Adrenergic alpha-Agonists

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D27.505.519.625.050.100.100.100 Adrenergic alpha-1 Receptor Agonists
-	D27.505.519.625.050.100.100.200 Adrenergic alpha-2 Receptor Agonists
-	D27.505.519.625.050.100.200 Adrenergic beta-Agonists
-	D27.505.519.625.050.100.200.100 Adrenergic beta-1 Receptor Agonists
-	D27.505.519.625.050.100.200.200 Adrenergic beta-2 Receptor Agonists
-	D27.505.519.625.050.100.200.300 Adrenergic beta-3 Receptor Agonists
-	D27.505.519.625.050.200 Adrenergic Antagonists
-	D27.505.519.625.050.200.100 Adrenergic alpha-Antagonists
-	D27.505.519.625.050.200.100.100 Adrenergic alpha-1 Receptor Antagonists
-	D27.505.519.625.050.200.100.200 Adrenergic alpha-2 Receptor Antagonists
-	D27.505.519.625.050.200.200 Adrenergic beta-Antagonists
-	D27.505.519.625.050.200.200.100 Adrenergic beta-1 Receptor Antagonists
-	D27.505.519.625.050.200.200.200 Adrenergic beta-2 Receptor Antagonists
-	D27.505.519.625.050.200.200.300 Adrenergic beta-3 Receptor Antagonists
-	D27.505.519.625.050.601 Adrenergic Uptake Inhibitors
-	D27.505.519.625.085 Cannabinoid Receptor Modulators
-	D27.505.519.625.085.500 Cannabinoid Receptor Agonists
-	D27.505.519.625.085.750 Cannabinoid Receptor Antagonists
-	D27.505.519.625.120 Cholinergic Agents
-	D27.505.519.625.120.069 Acetylcholine Release Inhibitors
-	D27.505.519.625.120.140 Cholinergic Agonists
-	D27.505.519.625.120.140.500 Muscarinic Agonists
-	D27.505.519.625.120.140.700 Nicotinic Agonists
-	D27.505.519.625.120.200 Cholinergic Antagonists
-	D27.505.519.625.120.200.500 Muscarinic Antagonists
-	D27.505.519.625.120.200.700 Nicotinic Antagonists
-	D27.505.519.625.120.300 Cholinesterase Inhibitors
-	D27.505.519.625.120.400 Cholinesterase Reactivators
-	D27.505.519.625.150 Dopamine Agents
-	D27.505.519.625.150.151 Dopamine Agonists
-	D27.505.519.625.150.175 Dopamine Antagonists
-	D27.505.519.625.150.175.500 Dopamine D2 Receptor Antagonists
-	D27.505.519.625.150.800 Dopamine Uptake Inhibitors
-	D27.505.519.625.190 Excitatory Amino Acid Agents
-	D27.505.519.625.190.200 Excitatory Amino Acid Agonists
-	D27.505.519.625.190.300 Excitatory Amino Acid Antagonists

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D27.505.519.625.240 GABA Agents
-	D27.505.519.625.240.200 GABA Agonists
-	D27.505.519.625.240.200.500 GABA-A Receptor Agonists
-	D27.505.519.625.240.200.750 GABA-B Receptor Agonists
-	D27.505.519.625.240.300 GABA Antagonists
-	D27.505.519.625.240.300.500 GABA-A Receptor Antagonists
-	D27.505.519.625.240.300.750 GABA-B Receptor Antagonists
-	D27.505.519.625.240.500 GABA Modulators
-	D27.505.519.625.240.750 GABA Uptake Inhibitors
-	D27.505.519.625.270 Gasotransmitters
-	D27.505.519.625.300 Glycine Agents
-	D27.505.519.625.375 Histamine Agents
-	D27.505.519.625.375.400 Histamine Agonists
-	D27.505.519.625.375.425 Histamine Antagonists
-	D27.505.519.625.375.425.400 Histamine H1 Antagonists
-	D27.505.519.625.375.425.400.500 Histamine H1 Antagonists, Non-Sedating
-	D27.505.519.625.375.425.425 Histamine H2 Antagonists
-	D27.505.519.625.375.425.712 Histamine H3 Antagonists
-	D27.505.519.625.487 Neurokinin-1 Receptor Antagonists
-	D27.505.519.625.600 Neurotransmitter Uptake Inhibitors
-	D27.505.519.625.600.050 Adrenergic Uptake Inhibitors
-	D27.505.519.625.600.220 Dopamine Uptake Inhibitors
-	D27.505.519.625.600.535 GABA Uptake Inhibitors
-	D27.505.519.625.600.693 Serotonin and Noradrenaline Reuptake Inhibitors
-	D27.505.519.625.600.850 Serotonin Uptake Inhibitors
-	D27.505.519.625.663 Orexin Receptor Antagonists
-	D27.505.519.625.725 Purinergic Agents
-	D27.505.519.625.725.200 Purinergic Agonists
-	D27.505.519.625.725.200.100 Purinergic P1 Receptor Agonists
-	D27.505.519.625.725.200.100.100 Adenosine A1 Receptor Agonists
-	D27.505.519.625.725.200.100.200 Adenosine A2 Receptor Agonists
-	D27.505.519.625.725.200.100.300 Adenosine A3 Receptor Agonists
-	D27.505.519.625.725.200.200 Purinergic P2 Receptor Agonists
-	D27.505.519.625.725.200.200.100 Purinergic P2X Receptor Agonists
-	D27.505.519.625.725.200.200.200 Purinergic P2Y Receptor Agonists
-	D27.505.519.625.725.400 Purinergic Antagonists

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D27.505.519.625.725.400.100 Purinergic P1 Receptor Antagonists
-	D27.505.519.625.725.400.100.100 Adenosine A1 Receptor Antagonists
-	D27.505.519.625.725.400.100.200 Adenosine A2 Receptor Antagonists
-	D27.505.519.625.725.400.100.300 Adenosine A3 Receptor Antagonists
-	D27.505.519.625.725.400.200 Purinergic P2 Receptor Antagonists
-	D27.505.519.625.725.400.200.100 Purinergic P2X Receptor Antagonists
-	D27.505.519.625.725.400.200.200 Purinergic P2Y Receptor Antagonists
-	D27.505.519.625.850 Serotonin Agents
-	D27.505.519.625.850.800 Serotonin Receptor Agonists
-	D27.505.519.625.850.800.100 Serotonin 5-HT1 Receptor Agonists
-	D27.505.519.625.850.800.200 Serotonin 5-HT2 Receptor Agonists
-	D27.505.519.625.850.800.300 Serotonin 5-HT3 Receptor Agonists
-	D27.505.519.625.850.800.400 Serotonin 5-HT4 Receptor Agonists
-	D27.505.519.625.850.850 Serotonin Antagonists
-	D27.505.519.625.850.850.100 Serotonin 5-HT1 Receptor Antagonists
-	D27.505.519.625.850.850.200 Serotonin 5-HT2 Receptor Antagonists
-	D27.505.519.625.850.850.300 Serotonin 5-HT3 Receptor Antagonists
-	D27.505.519.625.850.850.400 Serotonin 5-HT4 Receptor Antagonists
-	D27.505.519.625.850.900 Serotonin Uptake Inhibitors
-	D27.505.519.656 Nitric Oxide Donors
-	D27.505.519.828 Peptidomimetics
-	D27.505.519.871 Radiopharmaceuticals
-	D27.505.519.914 Sequestering Agents
-	D27.505.519.914.500 Chelating Agents
-	D27.505.519.914.500.204 Calcium Chelating Agents
-	D27.505.519.914.500.410 Iron Chelating Agents
-	D27.505.519.914.500.410.750 Siderophores
-	D27.505.519.957 Viral Fusion Protein Inhibitors
-	D27.505.519.957.500 HIV Fusion Inhibitors
-	D27.505.696 Physiological Effects of Drugs
-	D27.505.696.068 Antipyretics
-	D27.505.696.138 Antispermatic Agents
-	D27.505.696.138.379 Sperm Immobilizing Agents
-	D27.505.696.138.569 Spermaticidal Agents
-	D27.505.696.138.760 Spermaticogenesis-Blocking Agents
-	D27.505.696.207 Astringents

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	D27.505.696.242	Bone Density Conservation Agents
-	D27.505.696.277	Central Nervous System Depressants
-	D27.505.696.277.100	Anesthetics
-	D27.505.696.277.100.017	Anesthetics, Combined
-	D27.505.696.277.100.035	Anesthetics, General
-	D27.505.696.277.100.035.060	Anesthetics, Inhalation
-	D27.505.696.277.100.035.075	Anesthetics, Intravenous
-	D27.505.696.277.100.035.075.035	Anesthetics, Dissociative
-	D27.505.696.277.100.200	Anesthetics, Local
-	D27.505.696.277.350	Hypnotics and Sedatives
-	D27.505.696.277.600	Narcotics
-	D27.505.696.277.600.500	Analgesics, Opioid
-	D27.505.696.277.950	Tranquilizing Agents
-	D27.505.696.277.950.015	Anti-Anxiety Agents
-	D27.505.696.277.950.025	Antimanic Agents
-	D27.505.696.277.950.040	Antipsychotic Agents
-	D27.505.696.282	Central Nervous System Stimulants
-	D27.505.696.282.045	Aphrodisiacs
-	D27.505.696.282.050	Appetite Stimulants
-	D27.505.696.282.224	Convulsants
-	D27.505.696.305	Cerumenolytic Agents
-	D27.505.696.329	Emetics
-	D27.505.696.353	Endocrine Disruptors
-	D27.505.696.365	Galactogogues
-	D27.505.696.377	Growth Substances
-	D27.505.696.377.077	Angiogenesis Modulating Agents
-	D27.505.696.377.077.077	Angiogenesis Inducing Agents
-	D27.505.696.377.077.099	Angiogenesis Inhibitors
-	D27.505.696.377.450	Growth Inhibitors
-	D27.505.696.377.450.100	Angiogenesis Inhibitors
-	D27.505.696.377.605	Micronutrients
-	D27.505.696.377.605.555	Trace Elements
-	D27.505.696.377.605.600	Vitamins
New Heading	<b>D27.505.696.377.605.600.354</b>	<b>Provitamins</b>
-	D27.505.696.377.605.600.708	Vitamin B Complex

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D27.505.696.377.760 Plant Growth Regulators
-	D27.505.696.388 Hallucinogens
-	D27.505.696.399 Hormones, Hormone Substitutes, and Hormone Antagonists
-	D27.505.696.399.450 Hormone Antagonists
-	D27.505.696.399.450.065 Androgen Antagonists
-	D27.505.696.399.450.065.249 Androgen Receptor Antagonists
-	D27.505.696.399.450.065.500 Nonsteroidal Anti-Androgens
-	D27.505.696.399.450.100 Antithyroid Agents
-	D27.505.696.399.450.230 Calcimimetic Agents
-	D27.505.696.399.450.327 Estrogen Antagonists
-	D27.505.696.399.450.327.149 Aromatase Inhibitors
-	D27.505.696.399.450.327.650 Estrogen Receptor Antagonists
-	D27.505.696.399.450.360 Estrogen Receptor Modulators
-	D27.505.696.399.450.360.827 Selective Estrogen Receptor Modulators
-	D27.505.696.399.450.420 Insulin Antagonists
-	D27.505.696.399.450.565 Leukotriene Antagonists
-	D27.505.696.399.450.600 Mineralocorticoid Receptor Antagonists
-	D27.505.696.399.450.710 Prostaglandin Antagonists
-	D27.505.696.399.450.855 Steroid Synthesis Inhibitors
-	D27.505.696.399.450.855.100 14-alpha Demethylase Inhibitors
-	D27.505.696.399.450.855.200 5-alpha Reductase Inhibitors
-	D27.505.696.399.450.855.300 Aromatase Inhibitors
-	D27.505.696.399.472 Hormones
-	D27.505.696.399.472.080 Anabolic Agents
-	D27.505.696.399.472.161 Androgens
-	D27.505.696.399.472.188 Cannabinoid Receptor Modulators
-	D27.505.696.399.472.188.500 Cannabinoid Receptor Agonists
-	D27.505.696.399.472.188.750 Cannabinoid Receptor Antagonists
-	D27.505.696.399.472.277 Estrogens
-	D27.505.696.399.472.277.540 Estrogens, Non-Steroidal
-	D27.505.696.399.472.277.540.500 Phytoestrogens
-	D27.505.696.399.472.488 Glucocorticoids
-	D27.505.696.399.472.580 Incretins
-	D27.505.696.399.472.673 Mineralocorticoids
-	D27.505.696.399.472.858 Progestins
-	D27.505.696.422 Hypoglycemic Agents

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D27.505.696.422.249 Glycoside Hydrolase Inhibitors
-	D27.505.696.422.374 Amylin Receptor Agonists
-	D27.505.696.422.500 Dipeptidyl-Peptidase IV Inhibitors
-	D27.505.696.477 Immunologic Factors
-	D27.505.696.477.067 Adjuvants, Immunologic
-	D27.505.696.477.136 Agglutinins
-	D27.505.696.477.136.377 Hemagglutinins
-	D27.505.696.477.656 Immunosuppressive Agents
-	D27.505.696.477.656.500 Complement Inactivating Agents
-	D27.505.696.477.656.500.500 Bradykinin B2 Receptor Antagonists
-	D27.505.696.477.656.750 Myeloablative Agonists
-	D27.505.696.477.828 Interferon Inducers
-	D27.505.696.510 Muscle Relaxants, Central
-	D27.505.696.543 Narcotic Antagonists
-	D27.505.696.560 Natriuretic Agents
-	D27.505.696.560.249 Antidiuretic Agents
-	D27.505.696.560.311 Antidiuretic Hormone Receptor Antagonists
-	D27.505.696.560.374 Antihyperkalemic Agents
-	D27.505.696.560.500 Diuretics
-	D27.505.696.560.500.453 Diuretics, Osmotic
-	D27.505.696.560.500.726 Diuretics, Potassium Sparing
-	D27.505.696.560.500.726.100 Epithelial Sodium Channel Blockers
-	D27.505.696.560.500.726.249 Mineralocorticoid Receptor Antagonists
-	D27.505.696.560.500.863 Sodium Chloride Symporter Inhibitors
-	D27.505.696.560.500.931 Sodium Potassium Chloride Symporter Inhibitors
-	D27.505.696.577 Neurotransmitter Agents
-	D27.505.696.577.050 Adrenergic Agents
-	D27.505.696.577.050.100 Adrenergic Agonists
-	D27.505.696.577.050.100.100 Adrenergic alpha-Agonists
-	D27.505.696.577.050.100.100.100 Adrenergic alpha-1 Receptor Agonists
-	D27.505.696.577.050.100.100.200 Adrenergic alpha-2 Receptor Agonists
-	D27.505.696.577.050.100.200 Adrenergic beta-Agonists
-	D27.505.696.577.050.100.200.100 Adrenergic beta-1 Receptor Agonists
-	D27.505.696.577.050.100.200.200 Adrenergic beta-2 Receptor Agonists
-	D27.505.696.577.050.100.200.300 Adrenergic beta-3 Receptor Agonists
-	D27.505.696.577.050.200 Adrenergic Antagonists

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D27.505.696.577.050.200.100                      Adrenergic alpha-Antagonists
-	D27.505.696.577.050.200.100.100                      Adrenergic alpha-1 Receptor Antagonists
-	D27.505.696.577.050.200.100.200                      Adrenergic alpha-2 Receptor Antagonists
-	D27.505.696.577.050.200.200                      Adrenergic beta-Antagonists
-	D27.505.696.577.050.200.200.100                      Adrenergic beta-1 Receptor Antagonists
-	D27.505.696.577.050.200.200.200                      Adrenergic beta-2 Receptor Antagonists
-	D27.505.696.577.050.200.200.300                      Adrenergic beta-3 Receptor Antagonists
-	D27.505.696.577.050.601                      Adrenergic Uptake Inhibitors
-	D27.505.696.577.120                      Cholinergic Agents
-	D27.505.696.577.120.069                      Acetylcholine Release Inhibitors
-	D27.505.696.577.120.140                      Cholinergic Agonists
-	D27.505.696.577.120.140.500                      Muscarinic Agonists
-	D27.505.696.577.120.140.700                      Nicotinic Agonists
-	D27.505.696.577.120.200                      Cholinergic Antagonists
-	D27.505.696.577.120.200.500                      Muscarinic Antagonists
-	D27.505.696.577.120.200.700                      Nicotinic Antagonists
-	D27.505.696.577.120.300                      Cholinesterase Inhibitors
-	D27.505.696.577.120.400                      Cholinesterase Reactivators
-	D27.505.696.577.150                      Dopamine Agents
-	D27.505.696.577.150.151                      Dopamine Agonists
-	D27.505.696.577.150.175                      Dopamine Antagonists
-	D27.505.696.577.150.175.500                      Dopamine D2 Receptor Antagonists
-	D27.505.696.577.150.800                      Dopamine Uptake Inhibitors
-	D27.505.696.577.190                      Excitatory Amino Acid Agents
-	D27.505.696.577.190.200                      Excitatory Amino Acid Agonists
-	D27.505.696.577.190.300                      Excitatory Amino Acid Antagonists
-	D27.505.696.577.240                      GABA Agents
-	D27.505.696.577.240.200                      GABA Agonists
-	D27.505.696.577.240.200.500                      GABA-A Receptor Agonists
-	D27.505.696.577.240.200.750                      GABA-B Receptor Agonists
-	D27.505.696.577.240.300                      GABA Antagonists
-	D27.505.696.577.240.300.500                      GABA-A Receptor Antagonists
-	D27.505.696.577.240.300.750                      GABA-B Receptor Antagonists
-	D27.505.696.577.240.500                      GABA Modulators
-	D27.505.696.577.240.750                      GABA Uptake Inhibitors
-	D27.505.696.577.300                      Glycine Agents



## MeSH Tree Changes for 2017

Type	Tree - heading
-	D27.505.696.577.375 Histamine Agents
-	D27.505.696.577.375.400 Histamine Agonists
-	D27.505.696.577.375.425 Histamine Antagonists
-	D27.505.696.577.375.425.400 Histamine H1 Antagonists
-	D27.505.696.577.375.425.400.500 Histamine H1 Antagonists, Non-Sedating
-	D27.505.696.577.375.425.425 Histamine H2 Antagonists
-	D27.505.696.577.375.425.712 Histamine H3 Antagonists
-	D27.505.696.577.487 Neurokinin-1 Receptor Antagonists
-	D27.505.696.577.600 Neurotransmitter Uptake Inhibitors
-	D27.505.696.577.600.050 Adrenergic Uptake Inhibitors
-	D27.505.696.577.600.220 Dopamine Uptake Inhibitors
-	D27.505.696.577.600.535 GABA Uptake Inhibitors
-	D27.505.696.577.600.693 Serotonin and Noradrenaline Reuptake Inhibitors
-	D27.505.696.577.600.850 Serotonin Uptake Inhibitors
-	D27.505.696.577.663 Orexin Receptor Antagonists
-	D27.505.696.577.725 Purinergic Agents
-	D27.505.696.577.725.200 Purinergic Agonists
-	D27.505.696.577.725.200.100 Purinergic P1 Receptor Agonists
-	D27.505.696.577.725.200.100.100 Adenosine A1 Receptor Agonists
-	D27.505.696.577.725.200.100.200 Adenosine A2 Receptor Agonists
-	D27.505.696.577.725.200.100.300 Adenosine A3 Receptor Agonists
-	D27.505.696.577.725.200.200 Purinergic P2 Receptor Agonists
-	D27.505.696.577.725.200.200.100 Purinergic P2X Receptor Agonists
-	D27.505.696.577.725.200.200.200 Purinergic P2Y Receptor Agonists
-	D27.505.696.577.725.400 Purinergic Antagonists
-	D27.505.696.577.725.400.100 Purinergic P1 Receptor Antagonists
-	D27.505.696.577.725.400.100.100 Adenosine A1 Receptor Antagonists
-	D27.505.696.577.725.400.100.200 Adenosine A2 Receptor Antagonists
-	D27.505.696.577.725.400.100.300 Adenosine A3 Receptor Antagonists
-	D27.505.696.577.725.400.200 Purinergic P2 Receptor Antagonists
-	D27.505.696.577.725.400.200.100 Purinergic P2X Receptor Antagonists
-	D27.505.696.577.725.400.200.200 Purinergic P2Y Receptor Antagonists
-	D27.505.696.577.850 Serotonin Agents
-	D27.505.696.577.850.800 Serotonin Receptor Agonists
-	D27.505.696.577.850.800.100 Serotonin 5-HT1 Receptor Agonists
-	D27.505.696.577.850.800.200 Serotonin 5-HT2 Receptor Agonists

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D27.505.696.577.850.800.300 Serotonin 5-HT3 Receptor Agonists
-	D27.505.696.577.850.800.400 Serotonin 5-HT4 Receptor Agonists
-	D27.505.696.577.850.850 Serotonin Antagonists
-	D27.505.696.577.850.850.100 Serotonin 5-HT1 Receptor Antagonists
-	D27.505.696.577.850.850.200 Serotonin 5-HT2 Receptor Antagonists
-	D27.505.696.577.850.850.300 Serotonin 5-HT3 Receptor Antagonists
-	D27.505.696.577.850.850.400 Serotonin 5-HT4 Receptor Antagonists
-	D27.505.696.577.850.900 Serotonin Uptake Inhibitors
-	D27.505.696.620 Performance-Enhancing Substances
-	D27.505.696.663 Peripheral Nervous System Agents
-	D27.505.696.663.050 Autonomic Agents
-	D27.505.696.663.050.030 Antiemetics
-	D27.505.696.663.050.100 Bronchoconstrictor Agents
-	D27.505.696.663.050.110 Bronchodilator Agents
-	D27.505.696.663.050.225 Emetics
-	D27.505.696.663.050.340 Ganglionic Blockers
-	D27.505.696.663.050.400 Ganglionic Stimulants
-	D27.505.696.663.050.495 Miotics
-	D27.505.696.663.050.500 Mydriatics
-	D27.505.696.663.050.650 Parasympatholytics
-	D27.505.696.663.050.675 Parasympathomimetics
-	D27.505.696.663.050.850 Sympatholytics
-	D27.505.696.663.050.870 Sympathomimetics
-	D27.505.696.663.700 Neuromuscular Agents
-	D27.505.696.663.700.600 Muscle Relaxants, Central
-	D27.505.696.663.700.710 Neuromuscular Blocking Agents
-	D27.505.696.663.700.710.550 Neuromuscular Depolarizing Agents
-	D27.505.696.663.700.710.575 Neuromuscular Nondepolarizing Agents
-	D27.505.696.663.850 Sensory System Agents
-	D27.505.696.663.850.014 Analgesics
-	D27.505.696.663.850.014.040 Analgesics, Non-Narcotic
-	D27.505.696.663.850.014.040.500 Anti-Inflammatory Agents, Non-Steroidal
-	D27.505.696.663.850.014.040.500.500 Cyclooxygenase Inhibitors
-	D27.505.696.663.850.014.040.500.500.500 Cyclooxygenase 2 Inhibitors
-	D27.505.696.663.850.014.580 Analgesics, Short-Acting
-	D27.505.696.663.850.014.640 Dentin Desensitizing Agents

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D27.505.696.663.850.014.760 Narcotics
-	D27.505.696.663.850.014.760.500 Analgesics, Opioid
-	D27.505.696.663.850.025 Anesthetics, Local
-	D27.505.696.663.850.512 Narcotic Antagonists
-	D27.505.696.706 Protective Agents
-	D27.505.696.706.018 Anticarcinogenic Agents
-	D27.505.696.706.037 Antidotes
-	D27.505.696.706.080 Antimutagenic Agents
-	D27.505.696.706.125 Antioxidants
-	D27.505.696.706.222 Cariostatic Agents
-	D27.505.696.706.320 Cryoprotective Agents
-	D27.505.696.706.434 Insect Repellents
-	D27.505.696.706.548 Neuroprotective Agents
-	D27.505.696.706.776 Radiation-Protective Agents
-	D27.505.696.706.776.800 Sunscreening Agents
-	D27.505.696.706.888 Viscosupplements
-	D27.505.696.875 Reproductive Control Agents
-	D27.505.696.875.131 Abortifacient Agents
-	D27.505.696.875.131.100 Abortifacient Agents, Nonsteroidal
-	D27.505.696.875.131.200 Abortifacient Agents, Steroidal
-	D27.505.696.875.360 Contraceptive Agents
-	D27.505.696.875.360.276 Contraceptive Agents, Female
-	D27.505.696.875.360.276.210 Contraceptives, Oral
-	D27.505.696.875.360.276.210.100 Contraceptives, Oral, Combined
-	D27.505.696.875.360.276.210.277 Contraceptives, Oral, Hormonal
-	D27.505.696.875.360.276.210.400 Contraceptives, Oral, Sequential
-	D27.505.696.875.360.276.210.443 Contraceptives, Oral, Synthetic
-	D27.505.696.875.360.276.310 Contraceptives, Postcoital
-	D27.505.696.875.360.276.310.235 Contraceptives, Postcoital, Hormonal
-	D27.505.696.875.360.276.310.360 Contraceptives, Postcoital, Synthetic
-	D27.505.696.875.360.276.450 Luteolytic Agents
-	D27.505.696.875.360.276.500 Menstruation-Inducing Agents
-	D27.505.696.875.360.276.727 Sperm Immobilizing Agents
-	D27.505.696.875.360.276.827 Spermaticidal Agents
-	D27.505.696.875.360.443 Contraceptive Agents, Male
-	D27.505.696.875.360.443.068 Antispermaticidal Agents

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D27.505.696.875.360.443.068.760 Spermatogenesis-Blocking Agents
-	D27.505.696.875.552 Fertility Agents
-	D27.505.696.875.552.344 Fertility Agents, Female
-	D27.505.696.875.552.510 Fertility Agents, Male
-	D27.505.696.875.610 Luteolytic Agents
-	D27.505.696.875.650 Menstruation-Inducing Agents
-	D27.505.696.875.737 Oxytocics
-	D27.505.696.875.825 Tocolytic Agents
-	D27.505.954 Therapeutic Uses
-	D27.505.954.016 Anti-Allergic Agents
-	D27.505.954.122 Anti-Infective Agents
-	D27.505.954.122.085 Anti-Bacterial Agents
-	D27.505.954.122.085.222 Antitreponemal Agents
-	D27.505.954.122.085.255 Antitubercular Agents
-	D27.505.954.122.085.255.135 Antibiotics, Antitubercular
-	D27.505.954.122.085.516 beta-Lactamase Inhibitors
-	D27.505.954.122.085.777 Leprostatic Agents
-	D27.505.954.122.136 Antifungal Agents
-	D27.505.954.122.187 Anti-Infective Agents, Local
-	D27.505.954.122.187.500 Hand Sanitizers
-	D27.505.954.122.237 Anti-Infective Agents, Urinary
-	D27.505.954.122.250 Antiparasitic Agents
-	D27.505.954.122.250.075 Anthelmintics
-	D27.505.954.122.250.075.080 Antinematodal Agents
-	D27.505.954.122.250.075.080.275 Filaricides
-	D27.505.954.122.250.075.100 Antiplatyhelminthic Agents
-	D27.505.954.122.250.075.100.040 Anticestodal Agents
-	D27.505.954.122.250.075.100.750 Schistosomicides
-	D27.505.954.122.250.100 Antiprotozoal Agents
-	D27.505.954.122.250.100.055 Amebicides
-	D27.505.954.122.250.100.085 Antimalarials
-	D27.505.954.122.250.100.115 Antitrichomonal Agents
-	D27.505.954.122.250.100.170 Coccidiostats
-	D27.505.954.122.250.100.875 Trypanocidal Agents
-	D27.505.954.122.388 Antiviral Agents
-	D27.505.954.122.388.077 Anti-Retroviral Agents

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D27.505.954.122.388.077.088                      Anti-HIV Agents
-	D27.505.954.122.388.077.088.104                      CCR5 Receptor Antagonists
-	D27.505.954.122.388.077.088.209                      HIV Fusion Inhibitors
-	D27.505.954.122.388.077.088.314                      HIV Integrase Inhibitors
-	D27.505.954.122.388.077.088.420                      HIV Protease Inhibitors
-	D27.505.954.122.388.077.750                      Reverse Transcriptase Inhibitors
-	D27.505.954.122.388.538                      Viral Fusion Protein Inhibitors
-	D27.505.954.122.388.538.500                      HIV Fusion Inhibitors
-	D27.505.954.122.425                      Disinfectants
-	D27.505.954.122.425.150                      Contact Lens Solutions
-	D27.505.954.122.425.300                      Dental Disinfectants
-	D27.505.954.122.425.300.500                      Root Canal Irrigants
-	D27.505.954.122.425.650                      Hand Sanitizers
-	D27.505.954.158                      Anti-Inflammatory Agents
-	D27.505.954.158.030                      Anti-Inflammatory Agents, Non-Steroidal
-	D27.505.954.158.030.500                      Cyclooxygenase Inhibitors
-	D27.505.954.158.030.500.500                      Cyclooxygenase 2 Inhibitors
-	D27.505.954.158.515                      Demulcents
-	D27.505.954.248                      Antineoplastic Agents
-	D27.505.954.248.025                      Angiogenesis Inhibitors
-	D27.505.954.248.106                      Antibiotics, Antineoplastic
-	D27.505.954.248.125                      Anticarcinogenic Agents
-	D27.505.954.248.144                      Antimetabolites, Antineoplastic
-	D27.505.954.248.147                      Antimitotic Agents
-	D27.505.954.248.150                      Antineoplastic Agents, Alkylating
-	D27.505.954.248.169                      Antineoplastic Agents, Hormonal
-	D27.505.954.248.179                      Antineoplastic Agents, Phytogenic
-	D27.505.954.248.589                      Myeloablative Agonists
-	D27.505.954.248.692                      Poly(ADP-ribose) Polymerase Inhibitors
-	D27.505.954.248.794                      Topoisomerase Inhibitors
-	D27.505.954.248.794.500                      Topoisomerase I Inhibitors
-	D27.505.954.248.794.750                      Topoisomerase II Inhibitors
-	D27.505.954.329                      Antirheumatic Agents
-	D27.505.954.329.030                      Anti-Inflammatory Agents, Non-Steroidal
-	D27.505.954.329.030.500                      Cyclooxygenase Inhibitors
-	D27.505.954.329.030.500.500                      Cyclooxygenase 2 Inhibitors

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D27.505.954.329.337 Gout Suppressants
-	D27.505.954.329.337.900 Uricosuric Agents
-	D27.505.954.411 Cardiovascular Agents
-	D27.505.954.411.097 Anti-Arrhythmia Agents
-	D27.505.954.411.162 Antihypertensive Agents
-	D27.505.954.411.192 Calcium Channel Blockers
-	D27.505.954.411.207 Cardioplegic Solutions
-	D27.505.954.411.222 Cardiotonic Agents
-	D27.505.954.411.320 Fibrinolytic Agents
-	D27.505.954.411.455 Natriuretic Agents
-	D27.505.954.411.590 Nitric Oxide Donors
-	D27.505.954.411.645 Potassium Channel Blockers
-	D27.505.954.411.700 Sclerosing Solutions
-	D27.505.954.411.720 Sodium Channel Blockers
-	D27.505.954.411.720.500 Voltage-Gated Sodium Channel Blockers
-	D27.505.954.411.793 Vasoconstrictor Agents
-	D27.505.954.411.793.205 Calcium Channel Agonists
-	D27.505.954.411.793.610 Nasal Decongestants
-	D27.505.954.411.918 Vasodilator Agents
-	D27.505.954.411.918.500 Endothelium-Dependent Relaxing Factors
-	D27.505.954.427 Central Nervous System Agents
-	D27.505.954.427.010 Adjuvants, Anesthesia
-	D27.505.954.427.020 Alcohol Deterrents
-	D27.505.954.427.040 Analgesics
-	D27.505.954.427.040.100 Analgesics, Non-Narcotic
-	D27.505.954.427.040.381 Analgesics, Short-Acting
-	D27.505.954.427.040.437 Dentin Desensitizing Agents
-	D27.505.954.427.040.550 Narcotics
-	D27.505.954.427.040.550.500 Analgesics, Opioid
-	D27.505.954.427.080 Anticonvulsants
-	D27.505.954.427.090 Anti-Dyskinesia Agents
-	D27.505.954.427.090.050 Antiparkinson Agents
-	D27.505.954.427.090.050.249 Aromatic Amino Acid Decarboxylase Inhibitors
-	D27.505.954.427.090.050.500 Catechol O-Methyltransferase Inhibitors
-	D27.505.954.427.095 Antiemetics
-	D27.505.954.427.140 Anti-Obesity Agents

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D27.505.954.427.140.155                      Appetite Depressants
-	D27.505.954.427.153                            Antitussive Agents
-	D27.505.954.427.210                          Central Nervous System Depressants
-	D27.505.954.427.210.100                    Anesthetics
-	D27.505.954.427.210.100.017              Anesthetics, Combined
-	D27.505.954.427.210.100.035              Anesthetics, General
-	D27.505.954.427.210.100.035.060           Anesthetics, Inhalation
-	D27.505.954.427.210.100.035.075           Anesthetics, Intravenous
-	D27.505.954.427.210.100.035.075.035      Anesthetics, Dissociative
-	D27.505.954.427.210.100.200              Anesthetics, Local
-	D27.505.954.427.210.350                    Hypnotics and Sedatives
-	D27.505.954.427.210.600                    Narcotics
-	D27.505.954.427.210.600.500              Analgesics, Opioid
-	D27.505.954.427.210.950                    Tranquilizing Agents
-	D27.505.954.427.210.950.015              Anti-Anxiety Agents
-	D27.505.954.427.210.950.025              Antimanic Agents
-	D27.505.954.427.210.950.040              Antipsychotic Agents
-	D27.505.954.427.220                          Central Nervous System Stimulants
-	D27.505.954.427.220.045                    Aphrodisiacs
-	D27.505.954.427.220.050                    Appetite Stimulants
-	D27.505.954.427.220.224                    Convulsants
-	D27.505.954.427.220.612                    Wakefulness-Promoting Agents
-	D27.505.954.427.270                          Emetics
-	D27.505.954.427.525                          Muscle Relaxants, Central
-	D27.505.954.427.550                          Narcotic Antagonists
-	D27.505.954.427.575                          Neuroprotective Agents
-	D27.505.954.427.637                          Nootropic Agents
-	D27.505.954.427.700                          Psychotropic Drugs
-	D27.505.954.427.700.122                    Antidepressive Agents
-	D27.505.954.427.700.122.050              Antidepressive Agents, Second-Generation
-	D27.505.954.427.700.122.055              Antidepressive Agents, Tricyclic
-	D27.505.954.427.700.372                    Hallucinogens
-	D27.505.954.427.700.872                    Tranquilizing Agents
-	D27.505.954.427.700.872.015              Anti-Anxiety Agents
-	D27.505.954.427.700.872.025              Antimanic Agents
-	D27.505.954.427.700.872.331              Antipsychotic Agents

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D27.505.954.444 Dermatologic Agents
-	D27.505.954.444.075 Antipruritics
-	D27.505.954.444.100 Astringents
-	D27.505.954.444.200 Emollients
-	D27.505.954.444.400 Keratolytic Agents
-	D27.505.954.444.600 Photosensitizing Agents
-	D27.505.954.444.695 Sunscreening Agents
-	D27.505.954.483 Gastrointestinal Agents
-	D27.505.954.483.080 Antacids
-	D27.505.954.483.161 Antidiarrheals
-	D27.505.954.483.200 Antiemetics
-	D27.505.954.483.203 Anti-Ulcer Agents
-	D27.505.954.483.396 Cathartics
-	D27.505.954.483.508 Cholagogues and Cholaretics
-	D27.505.954.483.534 Demulcents
-	D27.505.954.483.560 Emetics
-	D27.505.954.483.620 Laxatives
-	D27.505.954.483.680 Lipotropic Agents
-	D27.505.954.502 Hematologic Agents
-	D27.505.954.502.119 Anticoagulants
-	D27.505.954.502.119.500 Antithrombins
-	D27.505.954.502.119.500.500 Factor Xa Inhibitors
-	D27.505.954.502.135 Antisickling Agents
-	D27.505.954.502.140 Blood Substitutes
-	D27.505.954.502.140.500 Plasma Substitutes
-	D27.505.954.502.270 Coagulants
-	D27.505.954.502.270.463 Hemostatics
-	D27.505.954.502.270.463.091 Antifibrinolytic Agents
-	D27.505.954.502.270.546 Heparin Antagonists
-	D27.505.954.502.427 Fibrinolytic Agents
-	D27.505.954.502.543 Hematinics
-	D27.505.954.502.780 Platelet Aggregation Inhibitors
-	D27.505.954.557 Lipid Regulating Agents
-	D27.505.954.557.500 Hypolipidemic Agents
-	D27.505.954.557.500.202 Anticholesteremic Agents
-	D27.505.954.557.500.202.370 Hydroxymethylglutaryl-CoA Reductase



## MeSH Tree Changes for 2017

Type	Tree - heading
	Inhibitors
-	D27.505.954.557.500.601 Lipotropic Agents
-	D27.505.954.557.500.601.500 Lipoprotein Lipase Activators
-	D27.505.954.578 Pharmaceutical Solutions
-	D27.505.954.578.322 Cardioplegic Solutions
-	D27.505.954.578.483 Dialysis Solutions
-	D27.505.954.578.483.651 Hemodialysis Solutions
-	D27.505.954.578.645 Ophthalmic Solutions
-	D27.505.954.578.645.500 Lubricant Eye Drops
-	D27.505.954.578.733 Parenteral Nutrition Solutions
-	D27.505.954.578.733.500 Fat Emulsions, Intravenous
-	D27.505.954.578.822 Sclerosing Solutions
-	D27.505.954.600 Radiation-Sensitizing Agents
-	D27.505.954.600.710 Photosensitizing Agents
-	D27.505.954.613 Renal Agents
-	D27.505.954.613.056 Anti-Infective Agents, Urinary
-	D27.505.954.613.860 Uricosuric Agents
-	D27.505.954.705 Reproductive Control Agents
-	D27.505.954.705.131 Abortifacient Agents
-	D27.505.954.705.131.100 Abortifacient Agents, Nonsteroidal
-	D27.505.954.705.131.200 Abortifacient Agents, Steroidal
-	D27.505.954.705.360 Contraceptive Agents
-	D27.505.954.705.360.276 Contraceptive Agents, Female
-	D27.505.954.705.360.276.210 Contraceptives, Oral
-	D27.505.954.705.360.276.210.100 Contraceptives, Oral, Combined
-	D27.505.954.705.360.276.210.277 Contraceptives, Oral, Hormonal
-	D27.505.954.705.360.276.210.400 Contraceptives, Oral, Sequential
-	D27.505.954.705.360.276.210.443 Contraceptives, Oral, Synthetic
-	D27.505.954.705.360.276.310 Contraceptives, Postcoital
-	D27.505.954.705.360.276.310.235 Contraceptives, Postcoital, Hormonal
-	D27.505.954.705.360.276.310.360 Contraceptives, Postcoital, Synthetic
-	D27.505.954.705.360.276.450 Luteolytic Agents
-	D27.505.954.705.360.276.500 Menstruation-Inducing Agents
-	D27.505.954.705.360.276.727 Sperm Immobilizing Agents
-	D27.505.954.705.360.276.827 Spermaticidal Agents
-	D27.505.954.705.360.443 Contraceptive Agents, Male



## MeSH Tree Changes for 2017

Type	Tree - heading
-	D27.720.031.700.748 Pesticide Synergists
-	D27.720.031.700.853 Rodenticides
-	D27.720.066 Antistatic Agents
-	D27.720.102 Biomedical and Dental Materials
-	D27.720.102.130 Biocompatible Materials
-	D27.720.102.158 Bone Cements
-	D27.720.102.187 Cariogenic Agents
-	D27.720.102.223 Cariostatic Agents
-	D27.720.102.339 Dental Materials
-	D27.720.102.339.249 Photoinitiators, Dental
-	D27.720.102.339.500 Pulp Capping and Pulpectomy Agents
-	D27.720.102.461 Dermal Fillers
-	D27.720.102.583 Mouthwashes
-	D27.720.102.919 Tissue Adhesives
-	D27.720.185 Caustics
-	D27.720.233 Coloring Agents
-	D27.720.233.174 Chromogenic Compounds
-	D27.720.233.348 Fluorescent Dyes
-	D27.720.233.674 Food Coloring Agents
-	D27.720.233.837 Hair Dyes
-	D27.720.259 Contrast Media
-	D27.720.269 Cosmetics
-	D27.720.269.189 Antiperspirants
-	D27.720.269.285 Cosmeceuticals
-	D27.720.269.380 Dentifrices
-	D27.720.269.385 Deodorants
-	D27.720.269.430 Hair Preparations
-	D27.720.269.430.214 Hair Bleaching Agents
-	D27.720.269.430.430 Hair Dyes
-	D27.720.269.583 Mouthwashes
-	D27.720.269.700 Perfume
-	D27.720.269.750 Skin Lightening Preparations
-	D27.720.269.800 Sunscreening Agents
-	D27.720.274 Disinfectants
-	D27.720.274.150 Contact Lens Solutions
-	D27.720.274.300 Dental Disinfectants

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D27.720.274.300.500                      Root Canal Irrigants
-	D27.720.274.650                              Hand Sanitizers
-	D27.720.317                                   Explosive Agents
-	D27.720.355                                   Fixatives
-	D27.720.361                                   Flame Retardants
-	D27.720.364                                   Flavoring Agents
-	D27.720.364.609                            Sweetening Agents
-	D27.720.364.609.500                      Non-Nutritive Sweeteners
-	D27.720.364.609.750                      Nutritive Sweeteners
-	D27.720.368                                  Food Additives
-	D27.720.368.350                            Fat Substitutes
-	D27.720.368.355                            Food Coloring Agents
-	D27.720.368.385                            Food Preservatives
-	D27.720.369                                  Hygroscopic Agents
-	D27.720.395                                  Ionophores
-	D27.720.395.100                            Calcium Ionophores
-	D27.720.395.400                            Potassium Ionophores
-	D27.720.395.500                            Proton Ionophores
-	D27.720.395.750                            Sodium Ionophores
-	D27.720.400                                  Irritants
-	D27.720.470                                  Laboratory Chemicals
-	D27.720.470.280                            Buffers
-	D27.720.470.280.060                      Ampholyte Mixtures
-	D27.720.470.305                            Culture Media
-	D27.720.470.305.250                      Culture Media, Conditioned
-	D27.720.470.305.255                      Culture Media, Serum-Free
-	D27.720.470.410                            Indicators and Reagents
-	D27.720.470.410.080                      Affinity Labels
-	D27.720.470.410.080.600                   Photoaffinity Labels
-	D27.720.470.410.200                      Chromogenic Compounds
-	D27.720.470.410.210                      Cross-Linking Reagents
-	D27.720.470.410.360                      Intercalating Agents
-	D27.720.470.410.505                      Luminescent Agents
-	D27.720.470.410.505.500                   Fluorescent Dyes
-	D27.720.470.410.577                      Neuronal Tract-Tracers
-	D27.720.470.410.650                      Radiopharmaceuticals

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D27.720.470.410.680 Reagent Kits, Diagnostic
-	D27.720.470.410.680.680 Reagent Strips
-	D27.720.470.410.690 Reducing Agents
-	D27.720.470.410.700 Sulfhydryl Reagents
-	D27.720.470.410.750 Thiobarbituric Acid Reactive Substances
-	D27.720.470.420 Ion Exchange Resins
-	D27.720.470.420.050 Anion Exchange Resins
-	D27.720.470.420.275 Cation Exchange Resins
-	D27.720.470.480 Ligands
-	D27.720.470.530 Molecular Probes
-	D27.720.470.530.600 Nucleic Acid Probes
-	D27.720.470.530.600.150 Antisense Elements (Genetics)
-	D27.720.470.530.600.150.200 DNA, Antisense
-	D27.720.470.530.600.150.200.640 Oligodeoxyribonucleotides, Antisense
-	D27.720.470.530.600.150.640 Oligonucleotides, Antisense
-	D27.720.470.530.600.150.640.640 Oligodeoxyribonucleotides, Antisense
-	D27.720.470.530.600.150.640.645 Oligoribonucleotides, Antisense
-	D27.720.470.530.600.150.760 RNA, Antisense
-	D27.720.470.530.600.150.760.645 Oligoribonucleotides, Antisense
-	D27.720.470.530.600.223 DNA Probes
-	D27.720.470.530.600.223.260 DNA, Complementary
-	D27.720.470.530.600.223.600 DNA Primers
-	D27.720.470.530.600.223.615 DNA Probes, HLA
-	D27.720.470.530.600.223.620 DNA Probes, HPV
-	D27.720.470.530.600.650 Oligonucleotide Probes
-	D27.720.470.530.600.825 RNA Probes
-	D27.720.470.530.600.825.840 RNA, Complementary
-	D27.720.470.765 Small Molecule Libraries
-	D27.720.556 Lubricants
-	D27.720.556.249 Lubricant Eye Drops
-	D27.720.556.500 Viscosupplements
-	D27.720.599 Nerve Agents
-	D27.720.642 Oxidants
-	D27.720.642.315 Bleaching Agents
-	D27.720.642.315.249 Hair Bleaching Agents
-	D27.720.642.315.500 Tooth Bleaching Agents

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D27.720.642.631                      Oxidants, Photochemical
-	D27.720.744                              Pharmaceutic Aids
-	D27.720.744.064                        Adjuvants, Pharmaceutic
-	D27.720.744.523                        Ointment Bases
-	D27.720.744.770                        Pharmaceutical Vehicles
-	D27.720.744.770.419                    Excipients
-	D27.720.744.771                        Preservatives, Pharmaceutical
-	D27.720.752                              Pharmaceutical Solutions
-	D27.720.752.322                        Cardioplegic Solutions
-	D27.720.752.483                        Dialysis Solutions
-	D27.720.752.483.651                    Hemodialysis Solutions
-	D27.720.752.608                        Ophthalmic Solutions
-	D27.720.752.608.500                    Lubricant Eye Drops
-	D27.720.752.733                        Parenteral Nutrition Solutions
-	D27.720.752.733.500                    Fat Emulsions, Intravenous
-	D27.720.752.822                        Sclerosing Solutions
-	D27.720.760                              Plasticizers
-	D27.720.777                              Poisons
-	D27.720.777.300                        Chemical Warfare Agents
-	D27.720.777.300.500                    Nerve Agents
-	D27.720.799                              Protective Agents
-	D27.720.799.018                        Anticarcinogenic Agents
-	D27.720.799.037                        Antidotes
-	D27.720.799.042                        Antimutagenic Agents
-	D27.720.799.047                        Antioxidants
-	D27.720.799.080                        Cardiotonic Agents
-	D27.720.799.113                        Cariostatic Agents
-	D27.720.799.180                        Cryoprotective Agents
-	D27.720.799.763                        Radiation-Protective Agents
-	D27.720.799.763.764                    Sunscreening Agents
-	D27.720.821                              Riot Control Agents, Chemical
-	D27.720.821.500                        Tear Gases
-	D27.720.832                              Sequestering Agents
-	D27.720.832.500                        Chelating Agents
-	D27.720.832.500.204                    Calcium Chelating Agents
-	D27.720.832.500.410                    Iron Chelating Agents

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D27.720.832.500.410.750 Siderophores
-	D27.720.844 Solvents
-	D27.720.844.500 Ionic Liquids
-	D27.720.877 Surface-Active Agents
-	D27.720.877.048 Antifoaming Agents
-	D27.720.877.156 Demulcents
-	D27.720.877.265 Detergents
-	D27.720.877.265.861 Soaps
-	D27.720.877.383 Emulsifying Agents
-	D27.720.877.383.500 Cerumenolytic Agents
-	D27.720.877.501 Immunosorbents
-	D27.720.877.974 Wetting Agents
-	D27.720.888 Sweetening Agents
-	D27.720.888.500 Non-Nutritive Sweeteners
-	D27.720.888.750 Nutritive Sweeteners
-	D27.720.944 Viscoelastic Substances
-	D27.720.944.500 Viscosupplements
-	D27.888 Toxic Actions
-	D27.888.141 Endocrine Disruptors
-	D27.888.284 Environmental Pollutants
-	D27.888.284.101 Air Pollutants
-	D27.888.284.101.268 Air Pollutants, Occupational
-	D27.888.284.101.393 Air Pollutants, Radioactive
-	D27.888.284.756 Soil Pollutants
-	D27.888.284.756.674 Soil Pollutants, Radioactive
-	D27.888.284.903 Water Pollutants
-	D27.888.284.903.655 Water Pollutants, Chemical
-	D27.888.284.903.821 Water Pollutants, Radioactive
-	D27.888.426 Hazardous Substances
-	D27.888.426.500 Hazardous Waste
-	D27.888.426.500.638 Radioactive Waste
-	D27.888.569 Noxae
-	D27.888.569.035 Alkylating Agents
-	D27.888.569.035.035 Antineoplastic Agents, Alkylating
-	D27.888.569.042 Antimetabolites
-	D27.888.569.042.030 Antimetabolites, Antineoplastic

## MeSH Tree Changes for 2017

Type	Tree - heading
-	D27.888.569.071 Antispermato-genic Agents
-	D27.888.569.071.379 Sperm Immobilizing Agents
-	D27.888.569.071.569 Spermato-cidal Agents
-	D27.888.569.071.760 Spermato-genesis-Blocking Agents
-	D27.888.569.100 Carcinogens
-	D27.888.569.100.125 Carcinogens, Environmental
-	D27.888.569.100.675 Peroxisome Proliferators
-	D27.888.569.142 Cardiotoxins
-	D27.888.569.185 Caustics
-	D27.888.569.199 Cytostatic Agents
-	D27.888.569.213 Cytotoxins
-	D27.888.569.213.444 Hemolytic Agents
-	D27.888.569.242 Dermotoxins
-	D27.888.569.271 Immunotoxins
-	D27.888.569.300 Irritants
-	D27.888.569.468 Mutagens
-	D27.888.569.468.060 Aneugens
-	D27.888.569.504 Neurotoxins
-	D27.888.569.540 Oxidants
-	D27.888.569.540.631 Oxidants, Photochemical
-	D27.888.569.612 Poisons
-	D27.888.569.612.150 Chemical Warfare Agents
-	D27.888.569.673 Pyrogens
-	D27.888.569.734 Riot Control Agents, Chemical
-	D27.888.569.734.500 Tear Gases
-	D27.888.569.864 Teratogens
-	D27.888.723 Pesticides
-	D27.888.723.071 Acaricides
-	D27.888.723.141 Chemosterilants
-	D27.888.723.288 Fungicides, Industrial
-	D27.888.723.366 Herbicides
-	D27.888.723.366.181 Defoliant-s, Chemical
-	D27.888.723.441 Insect Repellents
-	D27.888.723.491 Insecticides
-	D27.888.723.596 Molluscacides
-	D27.888.723.697 Pesticide Residues



## MeSH Tree Changes for 2017

Type	Tree - heading
-	D27.888.723.748 Pesticide Synergists
-	D27.888.723.853 Rodenticides
-	E01 Diagnosis
-	E01.055 Clinical Decision-Making
-	E01.110 Delayed Diagnosis
-	E01.158 Diagnosis, Computer-Assisted
-	E01.158.600 Image Interpretation, Computer-Assisted
-	E01.158.600.339 Neuronavigation
-	E01.158.600.680 Radiographic Image Interpretation, Computer-Assisted
-	E01.171 Diagnosis, Differential
-	E01.190 Diagnosis, Dual (Psychiatry)
-	E01.354 Diagnostic Errors
-	E01.354.340 False Negative Reactions
-	E01.354.506 False Positive Reactions
-	E01.354.753 Observer Variation
-	E01.370 Diagnostic Techniques and Procedures
-	E01.370.049 Age Determination by Skeleton
-	E01.370.060 Autopsy
-	E01.370.100 Breath Tests
-	E01.370.225 Clinical Laboratory Techniques
-	E01.370.225.124 Clinical Chemistry Tests
-	E01.370.225.124.100 Blood Chemical Analysis
-	E01.370.225.124.100.100 Blood Gas Analysis
-	E01.370.225.124.100.100.600 Oximetry
-	E01.370.225.124.100.100.600.100 Blood Gas Monitoring, Transcutaneous
-	E01.370.225.124.100.105 Blood Glucose Self-Monitoring
-	E01.370.225.124.100.110 Blood Protein Electrophoresis
-	E01.370.225.124.100.115 Blood Urea Nitrogen
-	E01.370.225.124.100.232 Dried Blood Spot Testing
-	E01.370.225.124.100.350 Glucose Clamp Technique
-	E01.370.225.124.100.355 Glucose Tolerance Test
-	E01.370.225.124.100.450 Lactose Tolerance Test
-	E01.370.225.124.100.600 Petrosal Sinus Sampling
-	E01.370.225.124.200 Clinical Enzyme Tests
-	E01.370.225.124.300 Gastric Acidity Determination
-	E01.370.225.124.470 Limulus Test

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E01.370.225.124.810                      Urinalysis
-	E01.370.225.500                            Cytological Techniques
-	E01.370.225.500.195                      Cell Count
-	E01.370.225.500.195.107                Blood Cell Count
-	E01.370.225.500.195.107.330            Erythrocyte Count
-	E01.370.225.500.195.107.330.725      Reticulocyte Count
-	E01.370.225.500.195.107.595            Leukocyte Count
-	E01.370.225.500.195.107.595.500      Lymphocyte Count
-	E01.370.225.500.195.107.595.500.150    CD4 Lymphocyte Count
-	E01.370.225.500.195.107.595.500.150.160    CD4-CD8 Ratio
-	E01.370.225.500.195.107.740            Platelet Count
-	E01.370.225.500.195.870                Sperm Count
-	E01.370.225.500.223                      Cell Culture Techniques
-	E01.370.225.500.223.500                Primary Cell Culture
-	E01.370.225.500.335                      Cell Migration Assays
-	E01.370.225.500.335.500                Cell Migration Assays, Leukocyte
-	E01.370.225.500.335.750                Cell Migration Assays, Macrophage
-	E01.370.225.500.335.875                Skin Window Technique
-	E01.370.225.500.363                      Cell Separation
-	E01.370.225.500.363.342                Flow Cytometry
-	E01.370.225.500.373                      Cell Tracking
-	E01.370.225.500.383                      Colony-Forming Units Assay
-	E01.370.225.500.383.910                Tumor Stem Cell Assay
-	E01.370.225.500.384                      Cytodiagnosis
-	E01.370.225.500.384.050                Amniocentesis
-	E01.370.225.500.384.100                Biopsy
-	E01.370.225.500.384.100.119            Biopsy, Needle
-	E01.370.225.500.384.100.119.500      Biopsy, Fine-Needle
-	E01.370.225.500.384.100.119.500.500    Endoscopic Ultrasound-Guided Fine Needle Aspiration
-	E01.370.225.500.384.100.119.750      Biopsy, Large-Core Needle
-	E01.370.225.500.384.100.149            Chorionic Villi Sampling
-	E01.370.225.500.384.100.160            Conization
-	E01.370.225.500.384.100.370            Image-Guided Biopsy
-	E01.370.225.500.384.100.370.500      Endoscopic Ultrasound-Guided Fine Needle Aspiration

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E01.370.225.500.384.100.422 Papanicolaou Test
-	E01.370.225.500.384.100.580 Sentinel Lymph Node Biopsy
-	E01.370.225.500.384.100.800 Vaginal Smears
-	E01.370.225.500.384.235 Cytopathogenic Effect, Viral
-	E01.370.225.500.384.617 Karyometry
-	E01.370.225.500.385 Cytogenetic Analysis
-	E01.370.225.500.385.130 Chromosome Banding
-	E01.370.225.500.385.315 Karyotyping
-	E01.370.225.500.385.315.800 Spectral Karyotyping
-	E01.370.225.500.385.500 Mitotic Index
-	E01.370.225.500.386 Cytophotometry
-	E01.370.225.500.386.350 Flow Cytometry
-	E01.370.225.500.386.400 Image Cytometry
-	E01.370.225.500.386.400.500 Laser Scanning Cytometry
-	E01.370.225.500.388 Drug Screening Assays, Antitumor
-	E01.370.225.500.388.930 Tumor Stem Cell Assay
-	E01.370.225.500.508 Enzyme-Linked Immunospot Assay
-	E01.370.225.500.607 Histochemistry
-	E01.370.225.500.607.512 Immunohistochemistry
-	E01.370.225.500.607.512.240 Fluorescent Antibody Technique
-	E01.370.225.500.607.512.240.149 Antibody-Coated Bacteria Test, Urinary
-	E01.370.225.500.607.512.240.300 Fluorescent Antibody Technique, Direct
-	E01.370.225.500.607.512.240.310 Fluorescent Antibody Technique, Indirect
-	E01.370.225.500.607.512.240.655 Fluoroimmunoassay
-	E01.370.225.500.607.512.240.655.350 Fluorescence Polarization Immunoassay
-	E01.370.225.500.607.790 Periodic Acid-Schiff Reaction
-	E01.370.225.500.607.810 Prussian Blue Reaction
-	E01.370.225.500.620 Histochemical Preparation Techniques
-	E01.370.225.500.620.520 Microdissection
-	E01.370.225.500.620.530 Microtomy
-	E01.370.225.500.620.530.160 Cryoultramicrotomy
-	E01.370.225.500.620.530.160.260 Frozen Sections
-	E01.370.225.500.620.620 Replica Techniques
-	E01.370.225.500.620.620.150 Corrosion Casting
-	E01.370.225.500.620.620.260 Freeze Fracturing

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E01.370.225.500.620.620.260.400 Freeze Etching
-	E01.370.225.500.620.670 Staining and Labeling
-	E01.370.225.500.620.670.130 Chromosome Banding
-	E01.370.225.500.620.670.325 In Situ Hybridization
-	E01.370.225.500.620.670.325.350 In Situ Hybridization, Fluorescence
-	E01.370.225.500.620.670.325.350.125 Chromosome Painting
-	E01.370.225.500.620.670.325.680 Primed In Situ Labeling
-	E01.370.225.500.620.670.520 Negative Staining
-	E01.370.225.500.620.670.570 Neuroanatomical Tract-Tracing Techniques
-	E01.370.225.500.620.670.615 Periodic Acid-Schiff Reaction
-	E01.370.225.500.620.670.660 Prussian Blue Reaction
-	E01.370.225.500.620.670.770 Shadowing (Histology)
-	E01.370.225.500.620.670.780 Silver Staining
-	E01.370.225.500.620.720 Tissue Embedding
-	E01.370.225.500.620.720.610 Paraffin Embedding
-	E01.370.225.500.620.720.640 Plastic Embedding
-	E01.370.225.500.620.760 Tissue Preservation
-	E01.370.225.500.620.760.160 Cryopreservation
-	E01.370.225.500.620.760.160.260 Freeze Drying
-	E01.370.225.500.620.760.160.260.270 Freeze Substitution
-	E01.370.225.500.620.760.720 Tissue Fixation
-	E01.370.225.500.810 Metabolic Flux Analysis
-	E01.370.225.562 Genetic Testing
New Heading	<b>E01.370.225.562.500 Pharmacogenomic Testing</b>
-	E01.370.225.625 Hematologic Tests
-	E01.370.225.625.107 Blood Cell Count
-	E01.370.225.625.107.330 Erythrocyte Count
-	E01.370.225.625.107.330.725 Reticulocyte Count
-	E01.370.225.625.107.595 Leukocyte Count
-	E01.370.225.625.107.595.500 Lymphocyte Count
-	E01.370.225.625.107.595.500.150 CD4 Lymphocyte Count
-	E01.370.225.625.107.595.500.150.160 CD4-CD8 Ratio
-	E01.370.225.625.107.700 Platelet Count
-	E01.370.225.625.115 Blood Coagulation Tests
-	E01.370.225.625.115.320 International Normalized Ratio

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E01.370.225.625.115.600 Partial Thromboplastin Time
-	E01.370.225.625.115.610 Prothrombin Time
-	E01.370.225.625.115.830 Thrombelastography
-	E01.370.225.625.115.870 Thrombin Time
-	E01.370.225.625.115.950 Whole Blood Coagulation Time
-	E01.370.225.625.120 Blood Grouping and Crossmatching
-	E01.370.225.625.125 Blood Sedimentation
-	E01.370.225.625.135 Bone Marrow Examination
-	E01.370.225.625.225 Erythrocyte Aggregation
-	E01.370.225.625.230 Erythrocyte Indices
-	E01.370.225.625.265 Fibrin Clot Lysis Time
-	E01.370.225.625.300 FIGLU Test
-	E01.370.225.625.400 Hematocrit
-	E01.370.225.625.410 Hemoglobinometry
-	E01.370.225.625.550 Osmotic Fragility
-	E01.370.225.625.625 Platelet Function Tests
-	E01.370.225.625.625.100 Bleeding Time
-	E01.370.225.625.625.160 Clot Retraction
-	E01.370.225.625.625.392 Mean Platelet Volume
-	E01.370.225.625.625.625 Platelet Count
-	E01.370.225.625.750 Schilling Test
-	E01.370.225.750 Histological Techniques
-	E01.370.225.750.132 Autoradiography
-	E01.370.225.750.288 Decalcification Technique
-	E01.370.225.750.551 Histochemistry
-	E01.370.225.750.551.512 Immunohistochemistry
-	E01.370.225.750.551.512.240 Fluorescent Antibody Technique
-	E01.370.225.750.551.512.240.149 Antibody-Coated Bacteria Test, Urinary
-	E01.370.225.750.551.512.240.300 Fluorescent Antibody Technique, Direct
-	E01.370.225.750.551.512.240.310 Fluorescent Antibody Technique, Indirect
-	E01.370.225.750.551.512.240.655 Fluoroimmunoassay
-	E01.370.225.750.551.512.240.655.350 Immunoassay Fluorescence Polarization
-	E01.370.225.750.551.790 Periodic Acid-Schiff Reaction
-	E01.370.225.750.551.810 Prussian Blue Reaction
-	E01.370.225.750.600 Histochemical Preparation Techniques

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E01.370.225.750.600.520                      Microdissection
-	E01.370.225.750.600.530                      Microtomy
-	E01.370.225.750.600.530.160                      Cryoultramicrotomy
-	E01.370.225.750.600.530.160.260                      Frozen Sections
-	E01.370.225.750.600.620                      Replica Techniques
-	E01.370.225.750.600.620.150                      Corrosion Casting
-	E01.370.225.750.600.620.260                      Freeze Fracturing
-	E01.370.225.750.600.620.260.400                      Freeze Etching
-	E01.370.225.750.600.670                      Staining and Labeling
-	E01.370.225.750.600.670.130                      Chromosome Banding
-	E01.370.225.750.600.670.325                      In Situ Hybridization
-	E01.370.225.750.600.670.325.350                      In Situ Hybridization, Fluorescence
-	E01.370.225.750.600.670.325.350.125                      Chromosome Painting
-	E01.370.225.750.600.670.325.680                      Primed In Situ Labeling
-	E01.370.225.750.600.670.520                      Negative Staining
-	E01.370.225.750.600.670.570                      Neuroanatomical Tract-Tracing Techniques
-	E01.370.225.750.600.670.660                      Prussian Blue Reaction
-	E01.370.225.750.600.670.770                      Shadowing (Histology)
-	E01.370.225.750.600.670.780                      Silver Staining
-	E01.370.225.750.600.720                      Tissue Embedding
-	E01.370.225.750.600.720.610                      Paraffin Embedding
-	E01.370.225.750.600.720.640                      Plastic Embedding
-	E01.370.225.750.600.760                      Tissue Preservation
-	E01.370.225.750.600.760.160                      Cryopreservation
-	E01.370.225.750.600.760.160.260                      Freeze Drying
-	E01.370.225.750.600.760.160.260.270                      Freeze Substitution
-	E01.370.225.750.600.760.720                      Tissue Fixation
-	E01.370.225.812                      Immunologic Tests
-	E01.370.225.812.049                      Antibody-Coated Bacteria Test, Urinary
-	E01.370.225.812.100                      Basophil Degranulation Test
-	E01.370.225.812.125                      Cell Migration Assays
-	E01.370.225.812.125.500                      Cell Migration Assays, Leukocyte
-	E01.370.225.812.125.750                      Cell Migration Assays, Macrophage
-	E01.370.225.812.125.875                      Skin Window Technique
-	E01.370.225.812.160                      Cytotoxicity Tests, Immunologic
-	E01.370.225.812.160.155                      Complement Hemolytic Activity Assay

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E01.370.225.812.375 Hemolytic Plaque Technique
-	E01.370.225.812.385 Histocompatibility Testing
-	E01.370.225.812.385.120 Blood Grouping and Crossmatching
-	E01.370.225.812.385.475 Lymphocyte Culture Test, Mixed
-	E01.370.225.812.400 Immune Adherence Reaction
-	E01.370.225.812.447 Immunophenotyping
-	E01.370.225.812.453 Interferon-gamma Release Tests
-	E01.370.225.812.460 Leukocyte Adherence Inhibition Test
-	E01.370.225.812.482 Lymphocyte Activation
-	E01.370.225.812.505 Monitoring, Immunologic
-	E01.370.225.812.620 Pregnancy Tests, Immunologic
-	E01.370.225.812.706 Rosette Formation
-	E01.370.225.812.735 Serologic Tests
-	E01.370.225.812.735.050 Agglutination Tests
-	E01.370.225.812.735.050.375 Hemagglutination Tests
-	E01.370.225.812.735.050.375.150 Coombs Test
-	E01.370.225.812.735.050.450 Latex Fixation Tests
-	E01.370.225.812.735.060 AIDS Serodiagnosis
-	E01.370.225.812.735.150 Complement Fixation Tests
-	E01.370.225.812.735.155 Complement Hemolytic Activity Assay
-	E01.370.225.812.735.360 Hemadsorption Inhibition Tests
-	E01.370.225.812.735.370 Hemagglutination Inhibition Tests
-	E01.370.225.812.735.550 Neutralization Tests
-	E01.370.225.812.735.645 Precipitin Tests
-	E01.370.225.812.735.645.300 Flocculation Tests
-	E01.370.225.812.735.645.350 Immunodiffusion
-	E01.370.225.812.735.645.350.350 Immunoelectrophoresis
-	E01.370.225.812.735.645.350.350.150 Counterimmunoelectrophoresis
-	E01.370.225.812.735.645.350.350.350 Immunoelectrophoresis, Two-Dimensional
-	E01.370.225.812.735.830 Radioallergosorbent Test
-	E01.370.225.812.735.840 Radioimmunoprecipitation Assay
-	E01.370.225.812.735.845 Serum Bactericidal Antibody Assay
-	E01.370.225.812.735.850 Syphilis Serodiagnosis
-	E01.370.225.812.735.850.200 Fluorescent Treponemal Antibody-Absorption Test

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	E01.370.225.812.735.850.800	Treponema Immobilization Test
-	E01.370.225.812.742	Serotyping
-	E01.370.225.812.871	Skin Tests
-	E01.370.225.812.871.300	Intradermal Tests
-	E01.370.225.812.871.300.540	Kveim Test
-	E01.370.225.812.871.300.750	Skin Test End-Point Titration
-	E01.370.225.812.871.450	Local Lymph Node Assay
-	E01.370.225.812.871.600	Passive Cutaneous Anaphylaxis
-	E01.370.225.812.871.610	Patch Tests
-	E01.370.225.812.871.705	Skin Window Technique
-	E01.370.225.812.871.800	Tuberculin Test
New Heading	<b>E01.370.225.828</b>	<b>Laboratory Critical Values</b>
-	E01.370.225.843	Metabolic Clearance Rate
-	E01.370.225.875	Microbiological Techniques
-	E01.370.225.875.074	Axenic Culture
-	E01.370.225.875.150	Bacteriological Techniques
-	E01.370.225.875.150.115	Bacterial Load
-	E01.370.225.875.150.125	Bacterial Typing Techniques
-	E01.370.225.875.150.125.150	Bacteriophage Typing
-	E01.370.225.875.150.125.457	Molecular Typing
-	E01.370.225.875.150.125.457.500	Multilocus Sequence Typing
-	E01.370.225.875.150.125.765	Ribotyping
-	E01.370.225.875.150.125.890	Serotyping
-	E01.370.225.875.150.570	Limulus Test
New Heading	<b>E01.370.225.875.185</b>	<b>Blood Culture</b>
-	E01.370.225.875.220	Colony Count, Microbial
-	E01.370.225.875.220.115	Bacterial Load
-	E01.370.225.875.595	Microbial Sensitivity Tests
-	E01.370.225.875.595.399	Disk Diffusion Antimicrobial Tests
-	E01.370.225.875.595.800	Serum Bactericidal Test
-	E01.370.225.875.610	Mycological Typing Techniques
-	E01.370.225.875.837	Serial Passage
-	E01.370.225.875.950	Viral Load
-	E01.370.225.875.970	Virus Cultivation



## MeSH Tree Changes for 2017

Type	Tree - heading
-	E01.370.225.875.970.790                      Viral Plaque Assay
-	E01.370.225.875.977                              Virus Inactivation
-	E01.370.225.875.985                              Xenodiagnosis
-	E01.370.225.880                                    Molecular Diagnostic Techniques
-	E01.370.225.880.500                              Human Papillomavirus DNA Tests
-	E01.370.225.910                                    Neonatal Screening
-	E01.370.225.925                                    Occult Blood
-	E01.370.225.932                                    Parasite Load
-	E01.370.225.932.600                              Parasite Egg Count
-	E01.370.225.940                                    Parasitic Sensitivity Tests
-	E01.370.225.970                                    Pregnancy Tests
-	E01.370.225.970.620                              Pregnancy Tests, Immunologic
-	E01.370.225.985                                    Radioligand Assay
-	E01.370.225.992                                    Semen Analysis
-	E01.370.225.992.624                              Sperm Count
-	E01.370.225.992.812                              Sperm Motility
-	E01.370.225.996                                    Sex Determination Analysis
-	E01.370.225.996.500                              Sex Determination by Skeleton
-	E01.370.225.998                                    Specimen Handling
-	E01.370.225.998.054                              Biopsy
-	E01.370.225.998.054.119                         Biopsy, Needle
-	E01.370.225.998.054.119.500                    Biopsy, Fine-Needle
-	E01.370.225.998.054.119.500.500              Endoscopic Ultrasound-Guided Fine Needle Aspiration
-	E01.370.225.998.054.119.750                    Biopsy, Large-Core Needle
-	E01.370.225.998.054.149                         Chorionic Villi Sampling
-	E01.370.225.998.054.160                         Conization
-	E01.370.225.998.054.370                         Image-Guided Biopsy
-	E01.370.225.998.054.370.500                    Endoscopic Ultrasound-Guided Fine Needle Aspiration
-	E01.370.225.998.054.422                         Papanicolaou Test
-	E01.370.225.998.054.580                         Sentinel Lymph Node Biopsy
-	E01.370.225.998.054.790                         Spinal Puncture
-	E01.370.225.998.054.800                         Vaginal Smears
-	E01.370.225.998.110                              Blood Specimen Collection
-	E01.370.225.998.110.150                         Cordocentesis

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E01.370.225.998.110.600                      Petrosal Sinus Sampling
-	E01.370.225.998.110.625                      Phlebotomy
-	E01.370.225.998.221                            Dissection
-	E01.370.225.998.221.580                      Microdissection
-	E01.370.225.998.221.580.500                Laser Capture Microdissection
-	E01.370.225.998.293                           DNA Contamination
-	E01.370.225.998.329                           Paracentesis
-	E01.370.225.998.329.309                      Amniocentesis
-	E01.370.225.998.329.465                      Arthrocentesis
-	E01.370.225.998.329.543                      Cordocentesis
-	E01.370.225.998.329.620                      Pericardiocentesis
-	E01.370.225.998.329.810                      Thoracentesis
-	E01.370.225.998.329.905                      Tympanocentesis
-	E01.370.225.998.762                           Urine Specimen Collection
-	E01.370.350                                      Diagnostic Imaging
-	E01.370.350.130                                Cardiac Imaging Techniques
-	E01.370.350.130.249                           Angiocardiology
-	E01.370.350.130.500                           Cardiac-Gated Imaging Techniques
-	E01.370.350.130.500.500                      Cardiac-Gated Single-Photon Emission Computer-Assisted Tomography
-	E01.370.350.130.500.750                      Gated Blood-Pool Imaging
-	E01.370.350.130.625                           Coronary Angiography
-	E01.370.350.130.750                           Echocardiography
-	E01.370.350.130.750.220                      Echocardiography, Doppler
-	E01.370.350.130.750.220.220                Echocardiography, Doppler, Color
-	E01.370.350.130.750.220.225                Echocardiography, Doppler, Pulsed
-	E01.370.350.130.750.228                      Echocardiography, Stress
-	E01.370.350.130.750.230                      Echocardiography, Three-Dimensional
-	E01.370.350.130.750.230.230                Echocardiography, Four-Dimensional
-	E01.370.350.130.750.235                      Echocardiography, Transesophageal
-	E01.370.350.130.875                           Myocardial Perfusion Imaging
-	E01.370.350.130.937                           Radionuclide Ventriculography
-	E01.370.350.130.937.350                      Gated Blood-Pool Imaging
-	E01.370.350.130.937.950                      Ventriculography, First-Pass
-	E01.370.350.350                                Image Interpretation, Computer-Assisted
-	E01.370.350.350.349                           Neuronavigation

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E01.370.350.350.700 Radiographic Image Interpretation, Computer-Assisted
-	E01.370.350.350.800 Tomography, Emission-Computed
-	E01.370.350.350.800.700 Positron-Emission Tomography
New Heading	<b>E01.370.350.350.800.700.500 Positron Emission Tomography Computed Tomography</b>
-	E01.370.350.350.800.800 Tomography, Emission-Computed, Single-Photon
-	E01.370.350.350.800.800.500 Cardiac-Gated Single-Photon Emission Computer-Assisted Tomography
New Heading	<b>E01.370.350.350.800.800.750 Single Photon Emission Computed Tomography Computed Tomography</b>
-	E01.370.350.350.810 Tomography, X-Ray Computed
-	E01.370.350.350.810.180 Colonography, Computed Tomographic
New Heading	<b>E01.370.350.350.810.335 Computed Tomography Angiography</b>
-	E01.370.350.350.810.490 Four-Dimensional Computed Tomography
New Heading	<b>E01.370.350.350.810.645 Positron Emission Tomography Computed Tomography</b>
New Heading	<b>E01.370.350.350.810.723 Single Photon Emission Computed Tomography Computed Tomography</b>
-	E01.370.350.350.810.800 Tomography, Spiral Computed
-	E01.370.350.350.810.800.500 Multidetector Computed Tomography
-	E01.370.350.400 Imaging, Three-Dimensional
-	E01.370.350.400.200 Echocardiography, Three-Dimensional
-	E01.370.350.400.200.230 Echocardiography, Four-Dimensional
-	E01.370.350.400.350 Four-Dimensional Computed Tomography
-	E01.370.350.400.500 Holography
-	E01.370.350.515 Microscopy
-	E01.370.350.515.277 Intravital Microscopy
-	E01.370.350.515.277.250 Dermoscopy
-	E01.370.350.515.277.500 Microscopic Angioscopy
-	E01.370.350.515.370 Microscopy, Acoustic
-	E01.370.350.515.395 Microscopy, Confocal
-	E01.370.350.515.395.500 Laser Scanning Cytometry
-	E01.370.350.515.402 Microscopy, Electron
-	E01.370.350.515.402.150 Cryoelectron Microscopy
-	E01.370.350.515.402.250 Electron Probe Microanalysis
-	E01.370.350.515.402.541 Microscopy, Electron, Scanning
-	E01.370.350.515.402.580 Microscopy, Electron, Transmission



## MeSH Tree Changes for 2017

Type	Tree - heading	
-	E01.370.350.578.937.260	Echoencephalography
-	E01.370.350.578.937.260.850	Ultrasonography, Doppler, Transcranial
-	E01.370.350.578.937.505	Myelography
-	E01.370.350.578.937.620	Pneumoencephalography
-	E01.370.350.589	Optical Imaging
-	E01.370.350.589.124	Narrow Band Imaging
-	E01.370.350.589.249	Tomography, Optical
-	E01.370.350.589.249.500	Tomography, Optical Coherence
-	E01.370.350.589.500	Transillumination
-	E01.370.350.600	Photography
-	E01.370.350.600.300	Holography
-	E01.370.350.600.350	Image Enhancement
-	E01.370.350.600.350.700	Radiographic Image Enhancement
-	E01.370.350.600.350.700.060	Angiography, Digital Subtraction
-	E01.370.350.600.350.700.690	Radiography, Dental, Digital
-	E01.370.350.600.350.700.700	Radiography, Dual-Energy Scanned Projection
-	E01.370.350.600.350.700.810	Tomography, X-Ray Computed
-	E01.370.350.600.350.700.810.180	Colonography, Computed Tomographic
New Heading	<b>E01.370.350.600.350.700.810.335</b>	<b>Computed Tomography Angiography</b>
New Heading	<b>E01.370.350.600.350.700.810.490 Computed Tomography</b>	<b>Positron Emission Tomography</b>
New Heading	<b>E01.370.350.600.350.700.810.645 Tomography Computed Tomography</b>	<b>Single Photon Emission Computed</b>
-	E01.370.350.600.350.700.810.800	Tomography, Spiral Computed
-	E01.370.350.600.350.700.810.800.500	Multidetector Computed Tomography
-	E01.370.350.600.350.800	Tomography, Emission-Computed
-	E01.370.350.600.350.800.399	Positron-Emission Tomography
New Heading	<b>E01.370.350.600.350.800.399.500 Computed Tomography</b>	<b>Positron Emission Tomography</b>
-	E01.370.350.600.350.800.800	Tomography, Emission-Computed, Single-Photon
New Heading	<b>E01.370.350.600.350.800.800.500 Tomography Computed Tomography</b>	<b>Single Photon Emission Computed</b>
-	E01.370.350.600.625	Photofluorography
-	E01.370.350.600.630	Photogrammetry
-	E01.370.350.600.630.500	Moire Topography
-	E01.370.350.600.630.750	Radiostereometric Analysis

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	E01.370.350.600.631	Photography, Dental
-	E01.370.350.600.635	Photomicrography
-	E01.370.350.600.817	Time-Lapse Imaging
-	E01.370.350.700	Radiography
-	E01.370.350.700.024	Absorptiometry, Photon
-	E01.370.350.700.050	Age Determination by Skeleton
-	E01.370.350.700.060	Angiography
-	E01.370.350.700.060.050	Angiocardiography
-	E01.370.350.700.060.060	Angiography, Digital Subtraction
-	E01.370.350.700.060.070	Aortography
-	E01.370.350.700.060.180	Cerebral Angiography
-	E01.370.350.700.060.190	Cineangiography
-	E01.370.350.700.060.200	Coronary Angiography
-	E01.370.350.700.060.600	Phlebography
-	E01.370.350.700.060.610	Portography
-	E01.370.350.700.070	Arthrography
-	E01.370.350.700.150	Cineradiography
-	E01.370.350.700.150.190	Cineangiography
-	E01.370.350.700.200	Electrokymography
-	E01.370.350.700.225	Fluoroscopy
New Heading	<b>E01.370.350.700.225.313</b>	<b>Barium Enema</b>
-	E01.370.350.700.225.625	Photofluorography
-	E01.370.350.700.325	Hysterosalpingography
-	E01.370.350.700.475	Lymphography
-	E01.370.350.700.500	Mammography
-	E01.370.350.700.500.975	Xeromammography
-	E01.370.350.700.510	Microradiography
-	E01.370.350.700.560	Neuroradiography
-	E01.370.350.700.560.180	Cerebral Angiography
-	E01.370.350.700.560.190	Cerebral Ventriculography
-	E01.370.350.700.560.260	Echoencephalography
-	E01.370.350.700.560.260.850	Ultrasonography, Doppler, Transcranial
-	E01.370.350.700.560.505	Myelography
-	E01.370.350.700.560.620	Pneumoencephalography
-	E01.370.350.700.625	Pneumoradiography

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E01.370.350.700.625.620 Pneumoencephalography
-	E01.370.350.700.700 Radiographic Image Enhancement
-	E01.370.350.700.700.060 Angiography, Digital Subtraction
-	E01.370.350.700.700.690 Radiography, Dental, Digital
-	E01.370.350.700.700.700 Radiography, Dual-Energy Scanned Projection
-	E01.370.350.700.700.810 Tomography, X-Ray Computed
-	E01.370.350.700.700.810.180 Colonography, Computed Tomographic
New Heading	<b>E01.370.350.700.700.810.335 Computed Tomography Angiography</b>
-	E01.370.350.700.700.810.490 Four-Dimensional Computed Tomography
New Heading	<b>E01.370.350.700.700.810.645 Positron Emission Tomography Computed Tomography</b>
New Heading	<b>E01.370.350.700.700.810.723 Single Photon Emission Computed Tomography Computed Tomography</b>
-	E01.370.350.700.700.810.800 Tomography, Spiral Computed
-	E01.370.350.700.700.810.800.500 Multidetector Computed Tomography
-	E01.370.350.700.705 Radiographic Image Interpretation, Computer-Assisted
-	E01.370.350.700.710 Radiographic Magnification
-	E01.370.350.700.715 Radiography, Abdominal
-	E01.370.350.700.715.200 Cholangiography
-	E01.370.350.700.715.200.200 Retrograde Cholangiopancreatography, Endoscopic
-	E01.370.350.700.715.210 Cholecystography
-	E01.370.350.700.715.250 Defecography
-	E01.370.350.700.715.610 Portography
-	E01.370.350.700.720 Radiography, Dental
-	E01.370.350.700.720.050 Age Determination by Teeth
-	E01.370.350.700.720.700 Radiography, Bitewing
-	E01.370.350.700.720.720 Radiography, Dental, Digital
-	E01.370.350.700.720.750 Radiography, Panoramic
-	E01.370.350.700.720.793 Sialography
-	E01.370.350.700.725 Radiography, Interventional
-	E01.370.350.700.730 Radiography, Thoracic
-	E01.370.350.700.730.100 Bronchography
-	E01.370.350.700.730.500 Mass Chest X-Ray
-	E01.370.350.700.750 Radiostereometric Analysis
-	E01.370.350.700.770 Sex Determination by Skeleton

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E01.370.350.700.810 Tomography, X-Ray
-	E01.370.350.700.810.810 Tomography, X-Ray Computed
-	E01.370.350.700.810.810.180 Colonography, Computed Tomographic
-	E01.370.350.700.810.810.490 Cone-Beam Computed Tomography
-	E01.370.350.700.810.810.490.500 Spiral Cone-Beam Computed Tomography
New Heading	<b>E01.370.350.700.810.810.568 Computed Tomography Angiography</b>
-	E01.370.350.700.810.810.645 Four-Dimensional Computed Tomography
New Heading	<b>E01.370.350.700.810.810.723 Positron Emission Tomography Computed Tomography</b>
New Heading	<b>E01.370.350.700.810.810.762 Single Photon Emission Computed Tomography Computed Tomography</b>
-	E01.370.350.700.810.810.800 Tomography, Spiral Computed
-	E01.370.350.700.810.810.800.249 Multidetector Computed Tomography
-	E01.370.350.700.810.810.800.500 Spiral Cone-Beam Computed Tomography
-	E01.370.350.700.810.810.900 X-Ray Microtomography
-	E01.370.350.700.830 Urography
New Heading	<b>E01.370.350.700.830.500 Cystography</b>
-	E01.370.350.700.975 Xeroradiography
-	E01.370.350.700.975.975 Xeromammography
-	E01.370.350.710 Radionuclide Imaging
-	E01.370.350.710.299 Lymphoscintigraphy
-	E01.370.350.710.600 Perfusion Imaging
-	E01.370.350.710.600.500 Myocardial Perfusion Imaging
-	E01.370.350.710.710 Radioimmunodetection
-	E01.370.350.710.715 Radionuclide Angiography
-	E01.370.350.710.715.700 Radioisotope Renography
-	E01.370.350.710.715.710 Radionuclide Ventriculography
-	E01.370.350.710.715.710.350 Gated Blood-Pool Imaging
-	E01.370.350.710.715.710.950 Ventriculography, First-Pass
-	E01.370.350.710.800 Tomography, Emission-Computed
-	E01.370.350.710.800.399 Positron-Emission Tomography
New Heading	<b>E01.370.350.710.800.399.500 Positron Emission Tomography Computed Tomography</b>
-	E01.370.350.710.800.800 Tomography, Emission-Computed, Single-Photon
-	E01.370.350.710.800.800.500 Cardiac-Gated Single-Photon Emission Computer-Assisted Tomography



## MeSH Tree Changes for 2017

Type	Tree - heading	
New Heading	<b>E01.370.350.710.800.800.750</b>	<b>Single Photon Emission Computed Tomography Computed Tomography</b>
-	E01.370.350.730	Respiratory-Gated Imaging Techniques
-	E01.370.350.750	Spectroscopy, Near-Infrared
-	E01.370.350.755	Stroboscopy
-	E01.370.350.760	Subtraction Technique
-	E01.370.350.760.060	Angiography, Digital Subtraction
-	E01.370.350.760.700	Radiography, Dual-Energy Scanned Projection
-	E01.370.350.780	Terahertz Imaging
-	E01.370.350.800	Thermography
-	E01.370.350.825	Tomography
-	E01.370.350.825.249	Electron Microscope Tomography
-	E01.370.350.825.500	Magnetic Resonance Imaging
-	E01.370.350.825.500.100	Cholangiopancreatography, Magnetic Resonance
-	E01.370.350.825.500.150	Diffusion Magnetic Resonance Imaging
-	E01.370.350.825.500.200	Echo-Planar Imaging
-	E01.370.350.825.500.350	Fluorine-19 Magnetic Resonance Imaging
-	E01.370.350.825.500.500	Magnetic Resonance Angiography
-	E01.370.350.825.500.510	Magnetic Resonance Imaging, Cine
-	E01.370.350.825.800	Tomography, Emission-Computed
-	E01.370.350.825.800.399	Positron-Emission Tomography
New Heading	<b>E01.370.350.825.800.399.500</b>	<b>Positron Emission Tomography Computed Tomography</b>
-	E01.370.350.825.800.800	Tomography, Emission-Computed, Single-Photon
-	E01.370.350.825.800.800.500 Computer-Assisted Tomography	Cardiac-Gated Single-Photon Emission
New Heading	<b>E01.370.350.825.800.800.750</b>	<b>Single Photon Emission Computed Tomography Computed Tomography</b>
-	E01.370.350.825.805	Tomography, Optical
-	E01.370.350.825.805.500	Tomography, Optical Coherence
-	E01.370.350.825.810	Tomography, X-Ray
-	E01.370.350.825.810.810	Tomography, X-Ray Computed
-	E01.370.350.825.810.810.399	Cone-Beam Computed Tomography
-	E01.370.350.825.810.810.399.500	Spiral Cone-Beam Computed Tomography
New Heading	<b>E01.370.350.825.810.810.499</b>	<b>Computed Tomography Angiography</b>
-	E01.370.350.825.810.810.599	Four-Dimensional Computed Tomography

## MeSH Tree Changes for 2017

Type	Tree - heading
New Heading	<b>E01.370.350.825.810.810.700</b> <b>Positron Emission Tomography Computed Tomography</b>
New Heading	<b>E01.370.350.825.810.810.750</b> <b>Single Photon Emission Computed Tomography Computed Tomography</b>
-	E01.370.350.825.810.810.800 Tomography, Spiral Computed
-	E01.370.350.825.810.810.800.249 Multidetector Computed Tomography
-	E01.370.350.825.810.810.800.500 Spiral Cone-Beam Computed Tomography
-	E01.370.350.825.810.810.900 X-Ray Microtomography
-	E01.370.350.850 Ultrasonography
-	E01.370.350.850.150 Carotid Intima-Media Thickness
-	E01.370.350.850.220 Echocardiography
-	E01.370.350.850.220.220 Echocardiography, Doppler
-	E01.370.350.850.220.220.220 Echocardiography, Doppler, Color
-	E01.370.350.850.220.220.225 Echocardiography, Doppler, Pulsed
-	E01.370.350.850.220.228 Echocardiography, Stress
-	E01.370.350.850.220.230 Echocardiography, Three-Dimensional
-	E01.370.350.850.220.230.230 Echocardiography, Four-Dimensional
-	E01.370.350.850.220.235 Echocardiography, Transesophageal
-	E01.370.350.850.260 Echoencephalography
-	E01.370.350.850.260.850 Ultrasonography, Doppler, Transcranial
-	E01.370.350.850.270 Elasticity Imaging Techniques
-	E01.370.350.850.280 Endosonography
-	E01.370.350.850.565 Microscopy, Acoustic
-	E01.370.350.850.850 Ultrasonography, Doppler
-	E01.370.350.850.850.220 Echocardiography, Doppler
-	E01.370.350.850.850.220.220 Echocardiography, Doppler, Color
-	E01.370.350.850.850.220.225 Echocardiography, Doppler, Pulsed
-	E01.370.350.850.850.850 Ultrasonography, Doppler, Duplex
-	E01.370.350.850.850.850.850 Ultrasonography, Doppler, Color
-	E01.370.350.850.850.850.850.220 Echocardiography, Doppler, Color
-	E01.370.350.850.850.860 Ultrasonography, Doppler, Pulsed
-	E01.370.350.850.850.860.225 Echocardiography, Doppler, Pulsed
-	E01.370.350.850.850.870 Ultrasonography, Doppler, Transcranial
-	E01.370.350.850.855 Ultrasonography, Interventional
-	E01.370.350.850.855.500 Endoscopic Ultrasound-Guided Fine Needle Aspiration
-	E01.370.350.850.860 Ultrasonography, Mammary

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E01.370.350.850.865                      Ultrasonography, Prenatal
-	E01.370.350.850.865.249                      Cervical Length Measurement
-	E01.370.350.850.865.500                      Nuchal Translucency Measurement
-	E01.370.350.887                      Voltage-Sensitive Dye Imaging
-	E01.370.350.925                      Whole Body Imaging
-	E01.370.360                      Diagnostic Self Evaluation
-	E01.370.370                      Diagnostic Techniques, Cardiovascular
-	E01.370.370.050                      Angiography
-	E01.370.370.050.050                      Angiocardiology
-	E01.370.370.050.060                      Angiography, Digital Subtraction
-	E01.370.370.050.070                      Aortography
-	E01.370.370.050.180                      Cerebral Angiography
-	E01.370.370.050.190                      Cineangiography
-	E01.370.370.050.200                      Coronary Angiography
-	E01.370.370.050.350                      Fluorescein Angiography
-	E01.370.370.050.500                      Magnetic Resonance Angiography
-	E01.370.370.050.600                      Phlebography
-	E01.370.370.050.610                      Portography
-	E01.370.370.050.650                      Radionuclide Angiography
-	E01.370.370.050.650.650                      Radionuclide Ventriculography
-	E01.370.370.050.650.650.350                      Gated Blood-Pool Imaging
-	E01.370.370.050.650.650.950                      Ventriculography, First-Pass
-	E01.370.370.065                      Angioscopy
-	E01.370.370.120                      Blood Circulation Time
-	E01.370.370.130                      Blood Flow Velocity
-	E01.370.370.140                      Blood Pressure Determination
-	E01.370.370.140.049                      Ankle Brachial Index
-	E01.370.370.140.100                      Blood Pressure Monitoring, Ambulatory
-	E01.370.370.150                      Blood Volume Determination
-	E01.370.370.170                      Capillary Fragility
-	E01.370.370.180                      Carotid Intima-Media Thickness
-	E01.370.370.380                      Heart Function Tests
-	E01.370.370.380.050                      Angiocardiology
-	E01.370.370.380.100                      Ballistocardiography
-	E01.370.370.380.140                      Cardiac Catheterization
-	E01.370.370.380.140.210                      Catheterization, Swan-Ganz

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	E01.370.370.380.150	Cardiac Output
-	E01.370.370.380.150.700	Stroke Volume
-	E01.370.370.380.160	Cardiography, Impedance
-	E01.370.370.380.170	Cardiotocography
-	E01.370.370.380.200	Coronary Angiography
-	E01.370.370.380.220	Echocardiography
-	E01.370.370.380.220.220	Echocardiography, Doppler
-	E01.370.370.380.220.220.220	Echocardiography, Doppler, Color
-	E01.370.370.380.220.220.225	Echocardiography, Doppler, Pulsed
-	E01.370.370.380.220.228	Echocardiography, Stress
-	E01.370.370.380.220.230	Echocardiography, Three-Dimensional
-	E01.370.370.380.220.230.230	Echocardiography, Four-Dimensional
-	E01.370.370.380.220.235	Echocardiography, Transesophageal
-	E01.370.370.380.240	Electrocardiography
-	E01.370.370.380.240.230	Electrocardiography, Ambulatory
-	E01.370.370.380.240.850	Vectorcardiography
-	E01.370.370.380.240.850.100	Body Surface Potential Mapping
-	E01.370.370.380.245	Electrophysiologic Techniques, Cardiac
-	E01.370.370.380.245.500	Epicardial Mapping
-	E01.370.370.380.250	Exercise Test
New Heading	<b>E01.370.370.380.250.500</b>	<b>Walk Test</b>
-	E01.370.370.380.400	Heart Auscultation
-	E01.370.370.380.400.500	Phonocardiography
New Heading	<b>E01.370.370.380.425</b>	<b>Heart Rate Determination</b>
-	E01.370.370.380.450	Kinetocardiography
-	E01.370.370.380.487	Magnetocardiography
-	E01.370.370.380.500	Myocardial Perfusion Imaging
-	E01.370.370.380.600	Oximetry
-	E01.370.370.380.600.100	Blood Gas Monitoring, Transcutaneous
-	E01.370.370.380.650	Pulse
-	E01.370.370.380.710	Radionuclide Ventriculography
-	E01.370.370.380.710.350	Gated Blood-Pool Imaging
-	E01.370.370.380.710.950	Ventriculography, First-Pass
-	E01.370.370.380.950	Valsalva Maneuver

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	E01.370.370.475	Laser-Doppler Flowmetry
-	E01.370.370.560	Microscopic Angioscopy
-	E01.370.370.610	Plethysmography
-	E01.370.370.610.600	Photoplethysmography
-	E01.370.370.610.610	Plethysmography, Impedance
-	E01.370.370.610.610.200	Cardiography, Impedance
-	E01.370.370.610.805	Plethysmography, Whole Body
-	E01.370.370.680	Pulse Wave Analysis
-	E01.370.370.750	Tilt-Table Test
-	E01.370.372	Diagnostic Techniques, Digestive System
-	E01.370.372.200	Cholangiography
-	E01.370.372.200.200	Cholangiopancreatography, Endoscopic Retrograde
-	E01.370.372.207	Cholangiopancreatography, Magnetic Resonance
-	E01.370.372.210	Cholecystography
-	E01.370.372.230	Colonography, Computed Tomographic
-	E01.370.372.250	Endoscopy, Digestive System
-	E01.370.372.250.200	Cholangiopancreatography, Endoscopic Retrograde
-	E01.370.372.250.250	Endoscopy, Gastrointestinal
New Heading	<b>E01.370.372.250.250.100</b>	<b>Balloon Enteroscopy</b>
New Tree	<a href="#">E01.370.372.250.250.100.500</a>	<a href="#">Double-Balloon Enteroscopy</a>
New Heading	<b>E01.370.372.250.250.100.750</b>	<b>Single-Balloon Enteroscopy</b>
-	E01.370.372.250.250.200	Colonoscopy
-	E01.370.372.250.250.200.700	Sigmoidoscopy
-	E01.370.372.250.250.225	Duodenoscopy
New Heading	<b>E01.370.372.250.250.250</b>	<b>Endoscopic Mucosal Resection</b>
-	E01.370.372.250.250.275	Esophagoscopy
-	E01.370.372.250.250.325	Gastrosopy
-	E01.370.372.250.250.600	Proctoscopy
-	E01.370.372.255	Esophageal pH Monitoring
-	E01.370.372.300	Gastric Acidity Determination
-	E01.370.372.310	Gastrointestinal Transit
-	E01.370.372.450	Lactose Tolerance Test
-	E01.370.372.460	Liver Function Tests

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E01.370.372.600 Pancreatic Function Tests
-	E01.370.372.610 Portography
-	E01.370.372.700 Sialography
-	E01.370.374 Diagnostic Techniques, Endocrine
-	E01.370.374.050 Adrenal Cortex Function Tests
-	E01.370.374.100 Blood Glucose Self-Monitoring
-	E01.370.374.355 Glucose Tolerance Test
-	E01.370.374.530 Ovarian Function Tests
-	E01.370.374.600 Pituitary-Adrenal Function Tests
-	E01.370.374.605 Pituitary Function Tests
-	E01.370.374.650 Radioligand Assay
-	E01.370.374.750 Thyroid Function Tests
-	E01.370.374.750.100 Basal Metabolism
-	E01.370.376 Diagnostic Techniques, Neurological
-	E01.370.376.300 Electroencephalography
-	E01.370.376.300.150 Brain Waves
-	E01.370.376.300.150.374 Alpha Rhythm
-	E01.370.376.300.150.750 Beta Rhythm
-	E01.370.376.300.150.875 Delta Rhythm
-	E01.370.376.300.150.906 Gamma Rhythm
-	E01.370.376.300.150.937 Theta Rhythm
-	E01.370.376.300.294 Electrocorticography
-	E01.370.376.300.437 Electroencephalography Phase Synchronization
-	E01.370.376.300.437.500 Cortical Synchronization
-	E01.370.376.500 Magnetoencephalography
-	E01.370.376.525 Neuroendoscopy
-	E01.370.376.537 Neuroimaging
-	E01.370.376.537.500 Diffusion Tensor Imaging
-	E01.370.376.537.625 Functional Neuroimaging
-	E01.370.376.537.625.500 Brain Mapping
-	E01.370.376.537.625.500.249 Connectome
-	E01.370.376.537.750 Neuroradiography
-	E01.370.376.537.750.180 Cerebral Angiography
-	E01.370.376.537.750.190 Cerebral Ventriculography
-	E01.370.376.537.750.260 Echoencephalography
-	E01.370.376.537.750.260.850 Ultrasonography, Doppler, Transcranial

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E01.370.376.537.750.505                      Myelography
-	E01.370.376.537.750.620                      Pneumoencephalography
-	E01.370.376.550                                  Neurologic Examination
-	E01.370.376.550.650                              Reflex
-	E01.370.376.550.650.324                          Diving Reflex
-	E01.370.376.550.650.650                          Reflex, Abdominal
-	E01.370.376.550.650.655                          Reflex, Abnormal
-	E01.370.376.550.650.670                          Reflex, Acoustic
-	E01.370.376.550.650.680                          Reflex, Babinski
-	E01.370.376.550.650.690                          Reflex, Pupillary
-	E01.370.376.550.650.695                          Reflex, Righting
-	E01.370.376.550.650.700                          Reflex, Stretch
-	E01.370.376.550.650.800                          Reflex, Startle
-	E01.370.376.625                                  Olfactometry
-	E01.370.376.700                                  Spinal Puncture
-	E01.370.378    Diagnostic Techniques, Obstetrical and Gynecological
-	E01.370.378.150                                      Colposcopy
-	E01.370.378.155                                      Culdoscopy
-	E01.370.378.225                                      Fallopian Tube Patency Tests
-	E01.370.378.230                                      Fetal Monitoring
-	E01.370.378.230.150                                  Cardiotocography
-	E01.370.378.325                                      Hysterosalpingography
-	E01.370.378.330                                      Hysteroscopy
-	E01.370.378.530                                      Ovarian Function Tests
-	E01.370.378.530.550                                  Ovulation Detection
-	E01.370.378.530.775                                  Ovulation Prediction
-	E01.370.378.605                                      Pelvimetry
-	E01.370.378.610                                      Placental Function Tests
-	E01.370.378.620                                      Pregnancy Tests
-	E01.370.378.620.620                                  Pregnancy Tests, Immunologic
-	E01.370.378.625                                      Preimplantation Diagnosis
-	E01.370.378.630                                      Prenatal Diagnosis
-	E01.370.378.630.050                                  Amniocentesis
-	E01.370.378.630.150                                  Chorionic Villi Sampling
-	E01.370.378.630.300                                  Fetoscopy
-	E01.370.378.630.582                                  Maternal Serum Screening Tests

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	E01.370.378.630.865	Ultrasonography, Prenatal
-	E01.370.378.630.865.249	Cervical Length Measurement
-	E01.370.378.630.865.500	Nuchal Translucency Measurement
-	E01.370.378.850	Ultrasonography, Mammary
-	E01.370.378.860	Uterine Monitoring
-	E01.370.378.900	Vaginal Smears
-	E01.370.380	Diagnostic Techniques, Ophthalmological
-	E01.370.380.074	Aberrometry
-	E01.370.380.112	Corneal Pachymetry
-	E01.370.380.150	Corneal Topography
-	E01.370.380.225	Electroretinography
-	E01.370.380.230	Eye Movement Measurements
-	E01.370.380.230.280	Electronystagmography
-	E01.370.380.230.285	Electrooculography
-	E01.370.380.245	Flicker Fusion
-	E01.370.380.250	Fluorescein Angiography
-	E01.370.380.255	Fluorophotometry
-	E01.370.380.300	Gonioscopy
-	E01.370.380.550	Ophthalmodynamometry
-	E01.370.380.560	Ophthalmoscopy
-	E01.370.380.560.500	Retinoscopy
-	E01.370.380.655	Scanning Laser Polarimetry
Old Tree	E01.370.380.703	Slit Lamp
New Heading	E01.370.380.727	Slit Lamp Microscopy
-	E01.370.380.750	Tonometry, Ocular
-	E01.370.380.850	Vision Tests
-	E01.370.380.850.150	Color Perception Tests
-	E01.370.380.850.700	Refraction, Ocular
-	E01.370.380.850.900	Vision Screening
-	E01.370.380.850.950	Visual Acuity
-	E01.370.380.850.950.500	Contrast Sensitivity
-	E01.370.380.850.950.750	Emmetropia
-	E01.370.380.850.962	Visual Field Tests
-	E01.370.382	Diagnostic Techniques, Otological
-	E01.370.382.375	Hearing Tests



## MeSH Tree Changes for 2017

Type	Tree - heading	
-	E01.370.382.375.050	Acoustic Impedance Tests
-	E01.370.382.375.060	Audiometry
-	E01.370.382.375.060.050	Audiometry, Evoked Response
-	E01.370.382.375.060.055	Audiometry, Pure-Tone
-	E01.370.382.375.060.060	Audiometry, Speech
-	E01.370.382.375.060.060.750	Speech Discrimination Tests
-	E01.370.382.375.060.060.760	Speech Reception Threshold Test
-	E01.370.382.375.060.530	Psychoacoustics
-	E01.370.382.375.200	Dichotic Listening Tests
-	E01.370.382.375.650	Recruitment Detection, Audiologic
-	E01.370.382.637	Otoscopy
-	E01.370.382.900	Vestibular Function Tests
-	E01.370.382.900.150	Caloric Tests
-	E01.370.382.900.280	Electronystagmography
-	E01.370.382.900.640	Head Impulse Test
-	E01.370.384	Diagnostic Techniques, Radioisotope
-	E01.370.384.700	Radioimmunoassay
-	E01.370.384.710	Radioisotope Dilution Technique
-	E01.370.384.720	Radioligand Assay
-	E01.370.384.730	Radionuclide Imaging
-	E01.370.384.730.176	Lymphoscintigraphy
-	E01.370.384.730.354	Perfusion Imaging
-	E01.370.384.730.354.500	Myocardial Perfusion Imaging
-	E01.370.384.730.710	Radioimmunodetection
-	E01.370.384.730.715	Radionuclide Angiography
-	E01.370.384.730.715.700	Radioisotope Renography
-	E01.370.384.730.715.710	Radionuclide Ventriculography
-	E01.370.384.730.715.710.350	Gated Blood-Pool Imaging
-	E01.370.384.730.715.710.950	Ventriculography, First-Pass
-	E01.370.384.730.800	Tomography, Emission-Computed
-	E01.370.384.730.800.399	Positron-Emission Tomography
New Heading	<b>E01.370.384.730.800.399.500</b> <b>Tomography</b>	<b>Positron Emission Tomography Computed</b>
-	E01.370.384.730.800.800	Tomography, Emission-Computed, Single-Photon
-	E01.370.384.730.800.800.500 Computer-Assisted Tomography	Cardiac-Gated Single-Photon Emission

## MeSH Tree Changes for 2017

Type	Tree - heading
New Heading	<b>E01.370.384.730.800.800.750</b> <span style="float: right;"><b>Single Photon Emission Computed Tomography Computed Tomography</b></span>
-	E01.370.384.750 Schilling Test
-	E01.370.386 Diagnostic Techniques, Respiratory System
-	E01.370.386.100 Bronchography
-	E01.370.386.105 Bronchoscopy
-	E01.370.386.460 Laryngoscopy
-	E01.370.386.500 Mass Chest X-Ray
-	E01.370.386.520 Mucociliary Clearance
-	E01.370.386.550 Nasal Provocation Tests
-	E01.370.386.700 Respiratory Function Tests
-	E01.370.386.700.050 Airway Resistance
-	E01.370.386.700.100 Blood Gas Analysis
-	E01.370.386.700.100.600 Oximetry
-	E01.370.386.700.100.600.100 Blood Gas Monitoring, Transcutaneous
-	E01.370.386.700.125 Bronchial Provocation Tests
-	E01.370.386.700.150 Capnography
-	E01.370.386.700.250 Exercise Test
-	E01.370.386.700.475 Lung Compliance
-	E01.370.386.700.485 Lung Volume Measurements
-	E01.370.386.700.485.750 Total Lung Capacity
-	E01.370.386.700.485.750.150 Closing Volume
-	E01.370.386.700.485.750.275 Functional Residual Capacity
-	E01.370.386.700.485.750.275.260 Expiratory Reserve Volume
-	E01.370.386.700.485.750.275.650 Residual Volume
-	E01.370.386.700.485.750.900 Vital Capacity
-	E01.370.386.700.485.750.900.260 Expiratory Reserve Volume
-	E01.370.386.700.485.750.900.350 Inspiratory Capacity
-	E01.370.386.700.485.750.900.350.350 Inspiratory Reserve Volume
-	E01.370.386.700.485.750.900.350.750 Tidal Volume
New Heading	<b>E01.370.386.700.550</b> <span style="float: right;"><b>Maximal Respiratory Pressures</b></span>
-	E01.370.386.700.615 Plethysmography, Whole Body
-	E01.370.386.700.650 Pulmonary Gas Exchange
-	E01.370.386.700.650.650 Pulmonary Diffusing Capacity
-	E01.370.386.700.650.900 Ventilation-Perfusion Ratio

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E01.370.386.700.660 Pulmonary Ventilation
-	E01.370.386.700.660.225 Forced Expiratory Flow Rates
-	E01.370.386.700.660.225.500 Maximal Expiratory Flow Rate
-	E01.370.386.700.660.225.505 Maximal Expiratory Flow-Volume Curves
-	E01.370.386.700.660.225.510 Maximal Midexpiratory Flow Rate
-	E01.370.386.700.660.225.600 Peak Expiratory Flow Rate
-	E01.370.386.700.660.230 Forced Expiratory Volume
-	E01.370.386.700.660.500 Maximal Voluntary Ventilation
-	E01.370.386.700.750 Spirometry
-	E01.370.386.700.750.100 Bronchosprometry
-	E01.370.386.700.950 Valsalva Maneuver
-	E01.370.386.700.975 Work of Breathing
-	E01.370.386.720 Respiratory Sounds
-	E01.370.386.730 Respiratory-Gated Imaging Techniques
-	E01.370.386.740 Rhinomanometry
-	E01.370.386.870 Rhinometry, Acoustic
-	E01.370.388 Diagnostic Techniques, Surgical
-	E01.370.388.100 Biopsy
-	E01.370.388.100.100 Biopsy, Needle
-	E01.370.388.100.100.500 Biopsy, Fine-Needle
-	E01.370.388.100.100.500.500 Endoscopic Ultrasound-Guided Fine Needle Aspiration
-	E01.370.388.100.100.750 Biopsy, Large-Core Needle
-	E01.370.388.100.150 Chorionic Villi Sampling
-	E01.370.388.100.160 Conization
-	E01.370.388.100.370 Image-Guided Biopsy
-	E01.370.388.100.370.500 Endoscopic Ultrasound-Guided Fine Needle Aspiration
-	E01.370.388.100.580 Sentinel Lymph Node Biopsy
-	E01.370.388.250 Endoscopy
-	E01.370.388.250.035 Angioscopy
-	E01.370.388.250.070 Arthroscopy
-	E01.370.388.250.100 Bronchoscopy
-	E01.370.388.250.150 Colposcopy
-	E01.370.388.250.160 Culdoscopy
-	E01.370.388.250.180 Cystoscopy

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	E01.370.388.250.250	Endoscopy, Digestive System
-	E01.370.388.250.250.160	Cholangiopancreatography, Endoscopic Retrograde
-	E01.370.388.250.250.250	Endoscopy, Gastrointestinal
New Heading	<b>E01.370.388.250.250.250.070</b>	<b>Balloon Enteroscopy</b>
New Tree	<a href="#">E01.370.388.250.250.250.070.500</a>	<a href="#">Double-Balloon Enteroscopy</a>
New Heading	<b>E01.370.388.250.250.250.070.750</b>	<b>Single-Balloon Enteroscopy</b>
-	E01.370.388.250.250.250.140	Capsule Endoscopy
-	E01.370.388.250.250.250.160	Colonoscopy
-	E01.370.388.250.250.250.160.800	Sigmoidoscopy
Old Tree	<del>E01.370.388.250.250.250.180</del>	<del>Double-Balloon Enteroscopy</del>
-	E01.370.388.250.250.250.200	Duodenoscopy
New Heading	<b>E01.370.388.250.250.250.230</b>	<b>Endoscopic Mucosal Resection</b>
-	E01.370.388.250.250.250.260	Esophagoscopy
-	E01.370.388.250.250.250.320	Gastrosocopy
-	E01.370.388.250.250.250.680	Proctoscopy
-	E01.370.388.250.280	Fetoscopy
-	E01.370.388.250.360	Hysteroscopy
-	E01.370.388.250.520	Laparoscopy
-	E01.370.388.250.520.500	Hand-Assisted Laparoscopy
-	E01.370.388.250.525	Laryngoscopy
-	E01.370.388.250.560	Mediastinoscopy
-	E01.370.388.250.630	Natural Orifice Endoscopic Surgery
-	E01.370.388.250.630.500	Transanal Endoscopic Surgery
-	E01.370.388.250.630.500.500	Transanal Endoscopic Microsurgery
-	E01.370.388.250.700	Neuroendoscopy
-	E01.370.388.250.840	Thoracoscopy
-	E01.370.388.250.840.830	Thoracic Surgery, Video-Assisted
-	E01.370.388.250.920	Ureteroscopy
-	E01.370.388.250.950	Video-Assisted Surgery
-	E01.370.388.250.950.830	Thoracic Surgery, Video-Assisted
-	E01.370.388.600	Pneumomediastinum, Diagnostic
-	E01.370.388.605	Pneumoperitoneum, Artificial
-	E01.370.390	Diagnostic Techniques, Urological

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	E01.370.390.050	Antibody-Coated Bacteria Test, Urinary
-	E01.370.390.175	Cystoscopy
-	E01.370.390.400	Kidney Function Tests
-	E01.370.390.400.100	Blood Urea Nitrogen
-	E01.370.390.400.300	Glomerular Filtration Rate
-	E01.370.390.400.700	Radioisotope Renography
-	E01.370.390.550	Nephrostomy, Percutaneous
-	E01.370.390.800	Ureteroscopy
-	E01.370.390.810	Urinalysis
-	E01.370.390.820	Urinary Catheterization
-	E01.370.390.830	Urography
New Heading	<b>E01.370.390.830.500</b>	<b>Cystography</b>
-	E01.370.395	Diagnostic Tests, Routine
-	E01.370.398	Direct-To-Consumer Screening and Testing
-	E01.370.400	Disability Evaluation
-	E01.370.400.925	Work Capacity Evaluation
-	E01.370.405	Electrodiagnosis
-	E01.370.405.240	Electrocardiography
-	E01.370.405.240.230	Electrocardiography, Ambulatory
-	E01.370.405.240.850	Vectorcardiography
-	E01.370.405.240.850.100	Body Surface Potential Mapping
-	E01.370.405.245	Electroencephalography
-	E01.370.405.245.287	Brain Waves
-	E01.370.405.245.287.374	Alpha Rhythm
-	E01.370.405.245.287.750	Beta Rhythm
-	E01.370.405.245.287.875	Delta Rhythm
-	E01.370.405.245.287.906	Gamma Rhythm
-	E01.370.405.245.287.937	Theta Rhythm
-	E01.370.405.245.431	Electrocorticography
-	E01.370.405.245.575	Electroencephalography Phase Synchronization
-	E01.370.405.245.575.500	Cortical Synchronization
-	E01.370.405.250	Electrokymography
-	E01.370.405.255	Electromyography
-	E01.370.405.260	Electronystagmography
-	E01.370.405.265	Electrooculography

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	E01.370.405.267	Electrophysiologic Techniques, Cardiac
-	E01.370.405.270	Electroretinography
-	E01.370.405.435	Magnetocardiography
-	E01.370.405.440	Magnetoencephalography
-	E01.370.405.610	Plethysmography, Impedance
-	E01.370.450	Insufflation
-	E01.370.475	Kymography
-	E01.370.475.200	Electrokymography
-	E01.370.500	Mass Screening
-	E01.370.500.174	Anonymous Testing
-	E01.370.500.500	Mass Chest X-Ray
-	E01.370.500.540	Multiphasic Screening
-	E01.370.500.580	Neonatal Screening
New Heading	<b>E01.370.503</b>	<b>Traditional Pulse Diagnosis</b>
-	E01.370.505	Surgical Clearance
-	E01.370.510	Medical History Taking
-	E01.370.510.150	Cornell Medical Index
-	E01.370.510.700	Reproductive History
-	E01.370.520	Monitoring, Physiologic
-	E01.370.520.049	Actigraphy
-	E01.370.520.100	Blood Glucose Self-Monitoring
-	E01.370.520.200	Drug Monitoring
-	E01.370.520.215	Esophageal pH Monitoring
-	E01.370.520.230	Fetal Monitoring
-	E01.370.520.230.150	Cardiotocography
-	E01.370.520.500	Monitoring, Ambulatory
-	E01.370.520.500.100	Blood Pressure Monitoring, Ambulatory
-	E01.370.520.500.230	Electrocardiography, Ambulatory
-	E01.370.520.505	Monitoring, Immunologic
-	E01.370.520.510	Monitoring, Intraoperative
-	E01.370.520.510.500	Intraoperative Neurophysiological Monitoring
-	E01.370.520.567	Neuromuscular Monitoring
-	E01.370.520.596	Neurophysiological Monitoring
-	E01.370.520.596.500	Intraoperative Neurophysiological Monitoring
-	E01.370.520.625	Polysomnography

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E01.370.520.750                      Telemetry
-	E01.370.520.750.500                      Remote Sensing Technology
-	E01.370.520.860                      Uterine Monitoring
-	E01.370.530                      Myography
-	E01.370.530.255                      Electromyography
-	E01.370.565                      Photoacoustic Techniques
-	E01.370.600                      Physical Examination
-	E01.370.600.024                      Anthropometry
-	E01.370.600.024.250                      Cephalometry
New Heading	<b>E01.370.600.024.450                      Kinanthropometry</b>
-	E01.370.600.024.650                      Odontometry
-	E01.370.600.024.650.500                      Age Determination by Teeth
-	E01.370.600.024.690                      Pelvimetry
-	E01.370.600.050                      Apgar Score
-	E01.370.600.060                      Auscultation
-	E01.370.600.060.400                      Heart Auscultation
-	E01.370.600.100                      Blood Pressure Determination
-	E01.370.600.115                      Body Constitution
-	E01.370.600.115.100                      Body Weights and Measures
-	E01.370.600.115.100.062                      Body Fat Distribution
-	E01.370.600.115.100.062.500                      Adiposity
-	E01.370.600.115.100.125                      Body Mass Index
-	E01.370.600.115.100.160                      Body Size
-	E01.370.600.115.100.160.100                      Body Height
-	E01.370.600.115.100.160.100.500                      Crown-Rump Length
-	E01.370.600.115.100.160.120                      Body Weight
-	E01.370.600.115.100.160.120.186                      Birth Weight
-	E01.370.600.115.100.160.120.300                      Fetal Weight
-	E01.370.600.115.100.160.120.499                      Ideal Body Weight
-	E01.370.600.115.100.160.120.699                      Overweight
-	E01.370.600.115.100.160.120.699.500                      Obesity
-	E01.370.600.115.100.160.120.699.500.249                      Obesity, Abdominal
-	E01.370.600.115.100.160.120.699.500.375                      Obesity, Metabolically Benign
-	E01.370.600.115.100.160.120.699.500.500                      Obesity, Morbid
-	E01.370.600.115.100.160.120.699.500.750                      Pediatric Obesity

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	E01.370.600.115.100.160.120.828	Thinness
-	E01.370.600.115.100.160.340	Sagittal Abdominal Diameter
-	E01.370.600.115.100.160.560	Waist Circumference
-	E01.370.600.115.100.160.560.500	Lipid Accumulation Product
-	E01.370.600.115.100.160.780	Waist-Height Ratio
-	E01.370.600.115.100.231	Body Surface Area
-	E01.370.600.115.100.660	Organ Size
-	E01.370.600.115.100.660.500	Axial Length, Eye
-	E01.370.600.115.100.803	Skinfold Thickness
-	E01.370.600.115.100.960	Waist-Hip Ratio
New Heading	<b>E01.370.600.115.275</b>	<b>Breast Density</b>
-	E01.370.600.115.450	Body, Physical Appearance
-	E01.370.600.115.450	Physical Appearance, Body
-	E01.370.600.115.450.500	Skin Pigmentation
-	E01.370.600.115.800	Somatotypes
-	E01.370.600.225	Facial Expression
-	E01.370.600.230	Facies
-	E01.370.600.250	Gait
New Heading	<b>E01.370.600.250.500</b>	<b>Walking Speed</b>
-	E01.370.600.293	Gynecological Examination
New Heading	<b>E01.370.600.315</b>	<b>Heart Rate Determination</b>
-	E01.370.600.337	Metabolic Equivalent
-	E01.370.600.425	Muscle Strength
-	E01.370.600.425.500	Hand Strength
-	E01.370.600.425.500.500	Pinch Strength
-	E01.370.600.550	Neurologic Examination
-	E01.370.600.550.162	Abnormal Involuntary Movement Scale
-	E01.370.600.550.324	Pain Measurement
-	E01.370.600.550.650	Reflex
-	E01.370.600.550.650.324	Diving Reflex
-	E01.370.600.550.650.650	Reflex, Abdominal
-	E01.370.600.550.650.655	Reflex, Abnormal
-	E01.370.600.550.650.670	Reflex, Acoustic
-	E01.370.600.550.650.680	Reflex, Babinski



## MeSH Tree Changes for 2017

Type	Tree - heading	
-	E01.370.600.550.650.690	Reflex, Pupillary
-	E01.370.600.550.650.695	Reflex, Righting
-	E01.370.600.550.650.700	Reflex, Stretch
-	E01.370.600.550.650.800	Reflex, Startle
-	E01.370.600.600	Palpation
-	E01.370.600.600.500	Digital Rectal Examination
-	E01.370.600.610	Percussion
-	E01.370.600.620	Pigmentation
-	E01.370.600.620.750	Skin Pigmentation
Old Tree	E01.370.600.650	Pulse
-	E01.370.600.700	Range of Motion, Articular
-	E01.370.600.700.500	Arthrometry, Articular
-	E01.370.600.750	Self-Examination
-	E01.370.600.750.100	Breast Self-Examination
-	E01.370.600.875	Vital Signs
-	E01.370.600.875.249	Blood Pressure
-	E01.370.600.875.374	Body Temperature
-	E01.370.600.875.500	Heart Rate
-	E01.370.600.875.875	Respiratory Rate
-	E01.370.620	Premarital Examinations
-	E01.370.685	Psychophysics
-	E01.370.685.628	Psychoacoustics
-	E01.370.685.814	Signal Detection, Psychological
-	E01.370.760	Speech Production Measurement
-	E01.370.760.760	Speech Articulation Tests
-	E01.370.872	Symptom Assessment
-	E01.370.928	Visual Analog Scale
-	E01.370.985	Xenodiagnosis
-	E01.390	Early Diagnosis
-	E01.390.500	Early Detection of Cancer
-	E01.410	Incidental Findings
-	E01.599	Prodromal Symptoms
-	E01.789	Prognosis
-	E01.789.200	Disease-Free Survival
-	E01.789.600	Medical Futility
-	E01.789.612	Neoplasm Grading

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	E01.789.625	Neoplasm Staging
-	E01.789.650	Nomograms
-	E01.789.700	Pregnancy Outcome
-	E01.789.800	Treatment Outcome
-	E01.789.800.379	Response Evaluation Criteria in Solid Tumors
New Heading	<b>E01.789.800.570</b>	<b>Sustained Virologic Response</b>
-	E01.789.800.760	Treatment Failure
-	E01.789.800.760.500	Failure to Rescue, Health Care
-	E01.894	Theranostic Nanomedicine
-	E02	Therapeutics
-	E02.037	Acoustic Stimulation
-	E02.041	Airway Management
-	E02.041.249	Airway Extubation
-	E02.041.500	Intubation, Intratracheal
-	E02.041.500.475	Laryngeal Masks
-	E02.041.625	Respiration, Artificial
-	E02.041.625.508	High-Frequency Ventilation
-	E02.041.625.508.510	High-Frequency Jet Ventilation
-	E02.041.625.516	Interactive Ventilatory Support
-	E02.041.625.525	Liquid Ventilation
-	E02.041.625.591	Noninvasive Ventilation
-	E02.041.625.657	One-Lung Ventilation
-	E02.041.625.790	Positive-Pressure Respiration
-	E02.041.625.790.259	Continuous Positive Airway Pressure
-	E02.041.625.790.520	Intermittent Positive-Pressure Breathing
-	E02.041.625.790.550	Intermittent Positive-Pressure Ventilation
-	E02.041.625.950	Ventilator Weaning
-	E02.041.750	Tracheostomy
-	E02.046	Apitherapy
-	E02.056	Balneology
-	E02.056.020	Ammotherapy
-	E02.056.110	Baths
-	E02.056.610	Mud Therapy
-	E02.056.902	Steam Bath
-	E02.065	Bariatrics

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E02.065.062                      Bariatric Surgery
-	E02.065.062.249                  Gastric Bypass
-	E02.065.062.750                  Gastroplasty
-	E02.065.062.875                  Jejunoileal Bypass
-	E02.065.062.937                  Lipectomy
-	E02.075                              Bed Rest
-	E02.085                              Behavior Control
-	E02.085.700                        Restraint, Physical
-	E02.095                              Biological Therapy
-	E02.095.125                        Blood Patch, Epidural
-	E02.095.135                        Blood Transfusion
-	E02.095.135.140                  Blood Component Transfusion
-	E02.095.135.140.275              Erythrocyte Transfusion
-	E02.095.135.140.425              Leukocyte Transfusion
-	E02.095.135.140.425.445        Lymphocyte Transfusion
-	E02.095.135.140.650              Platelet Transfusion
-	E02.095.135.164                  Blood Transfusion, Autologous
-	E02.095.135.264                  Blood Transfusion, Intrauterine
-	E02.095.135.469                  Exchange Transfusion, Whole Blood
-	E02.095.135.750                  Plasma Exchange
-	E02.095.147                        Cell- and Tissue-Based Therapy
-	E02.095.147.500                  Cell Transplantation
-	E02.095.147.750                  Tissue Transplantation
-	E02.095.160                        Cytapheresis
-	E02.095.160.570                  Leukapheresis
-	E02.095.160.790                  Plateletpheresis
-	E02.095.231                        Fecal Microbiota Transplantation
-	E02.095.301                        Genetic Therapy
-	E02.095.301.250                  RNAi Therapeutics
-	E02.095.301.500                  Targeted Gene Repair
-	E02.095.410                        Hematopoietic Stem Cell Mobilization
-	E02.095.437                        Immunomagnetic Separation
-	E02.095.465                        Immunomodulation
-	E02.095.465.425                  Immunotherapy
-	E02.095.465.425.400              Immunization
-	E02.095.465.425.400.330        Immunization, Passive

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	E02.095.465.425.400.330.050	Adoptive Transfer
-	E02.095.465.425.400.330.050.400	Immunotherapy, Adoptive
-	E02.095.465.425.400.470	Immunization Schedule
-	E02.095.465.425.400.485	Immunization, Secondary
-	E02.095.465.425.400.530	Immunotherapy, Active
-	E02.095.465.425.400.530.890	Vaccination
-	E02.095.465.425.400.530.890.500	Mass Vaccination
-	E02.095.465.425.450	Immunosuppression
-	E02.095.465.425.450.310	Desensitization, Immunologic
-	E02.095.465.425.450.310.500	Sublingual Immunotherapy
-	E02.095.465.425.450.440	Graft Enhancement, Immunologic
-	E02.095.465.425.450.521	Lymphocyte Depletion
-	E02.095.465.425.450.800	Transplantation Conditioning
-	E02.095.465.425.750	Radioimmunotherapy
-	E02.095.601	Oncolytic Virotherapy
-	E02.095.682	Organotherapy
-	E02.095.682.884	Tissue Therapy, Historical
New Heading	<b>E02.095.841</b>	<b>Phage Therapy</b>
-	E02.120	Blood Component Removal
-	E02.120.142	Bone Marrow Purging
-	E02.120.285	Cytapheresis
-	E02.120.285.570	Leukapheresis
-	E02.120.285.790	Plateletpheresis
-	E02.120.527	Leukocyte Reduction Procedures
-	E02.120.527.570	Leukapheresis
-	E02.120.770	Plasmapheresis
-	E02.148	Catheterization
-	E02.148.050	Angioplasty
-	E02.148.050.060	Angioplasty, Balloon
-	E02.148.050.060.100	Angioplasty, Balloon, Coronary
-	E02.148.050.060.105	Angioplasty, Balloon, Laser-Assisted
-	E02.148.050.075	Angioplasty, Laser
-	E02.148.050.075.080	Angioplasty, Balloon, Laser-Assisted
-	E02.148.050.120	Atherectomy
-	E02.148.050.120.125	Atherectomy, Coronary

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	E02.148.093	Balloon Embolectomy
-	E02.148.106	Balloon Occlusion
-	E02.148.106.500	Uterine Balloon Tamponade
-	E02.148.108	Balloon Valvuloplasty
-	E02.148.167	Catheterization, Central Venous
-	E02.148.224	Catheterization, Peripheral
-	E02.148.224.165	Catheterization, Swan-Ganz
-	E02.148.442	Cardiac Catheterization
-	E02.148.442.165	Catheterization, Swan-Ganz
-	E02.148.947	Urinary Catheterization
-	E02.148.947.500	Intermittent Urethral Catheterization
-	E02.154	Cautery
-	E02.154.402	Electrocoagulation
-	E02.154.402.054	Argon Plasma Coagulation
-	E02.168	Chronotherapy
-	E02.168.500	Drug Chronotherapy
-	E02.175	Climatotherapy
-	E02.183	Clinical Protocols
-	E02.183.750	Antineoplastic Protocols
-	E02.183.750.500	Antineoplastic Combined Chemotherapy Protocols
New Heading	<b>E02.183.875</b>	<b>Standing Orders</b>
-	E02.186	Combined Modality Therapy
-	E02.186.079	Chemoradiotherapy
-	E02.186.079.500	Chemoradiotherapy, Adjuvant
-	E02.186.170	Chemotherapy, Adjuvant
-	E02.186.250	Electroacupuncture
-	E02.186.450	Neoadjuvant Therapy
-	E02.186.500	Photochemotherapy
-	E02.186.750	Radioimmunotherapy
-	E02.186.775	Radiotherapy, Adjuvant
-	E02.190	Complementary Therapies
-	E02.190.044	Acupuncture Therapy
-	E02.190.044.105	Acupuncture Analgesia
-	E02.190.044.133	Acupuncture, Ear
-	E02.190.044.244	Electroacupuncture

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E02.190.044.555 Meridians
-	E02.190.044.555.035 Acupuncture Points
-	E02.190.044.588 Moxibustion
-	E02.190.088 Anthroposophy
-	E02.190.204 Auriculotherapy
-	E02.190.204.500 Acupuncture, Ear
-	E02.190.262 Diffuse Noxious Inhibitory Control
-	E02.190.321 Holistic Health
-	E02.190.321.500 Bioresonance Therapy
-	E02.190.388 Homeopathy
-	E02.190.438 Horticultural Therapy
-	E02.190.488 Medicine, Traditional
-	E02.190.488.505 Medicine, African Traditional
-	E02.190.488.510 Medicine, Arabic
-	E02.190.488.510.500 Medicine, Unani
-	E02.190.488.515 Medicine, Ayurvedic
-	E02.190.488.585 Medicine, East Asian Traditional
-	E02.190.488.585.520 Medicine, Chinese Traditional
-	E02.190.488.585.600 Medicine, Kampo
-	E02.190.488.585.700 Medicine, Korean Traditional
-	E02.190.488.585.850 Medicine, Mongolian Traditional
New Tree	<a href="#">E02.190.488.585.875</a> <a href="#">Medicine, Tibetan Traditional</a>
-	E02.190.488.830 Shamanism
-	E02.190.506 Mesotherapy
-	E02.190.525 Mind-Body Therapies
-	E02.190.525.061 Aromatherapy
-	E02.190.525.123 Biofeedback, Psychology
-	E02.190.525.123.500 Neurofeedback
-	E02.190.525.186 Breathing Exercises
-	E02.190.525.186.500 Qigong
-	E02.190.525.217 Hypnosis
-	E02.190.525.217.100 Autogenic Training
-	E02.190.525.217.771 Suggestion
-	E02.190.525.217.771.100 Autosuggestion
-	E02.190.525.249 Imagery (Psychotherapy)

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E02.190.525.311 Laughter Therapy
-	E02.190.525.374 Meditation
-	E02.190.525.500 Mental Healing
-	E02.190.525.781 Psychodrama
-	E02.190.525.781.653 Role Playing
-	E02.190.525.812 Psychophysiology
-	E02.190.525.875 Relaxation Therapy
-	E02.190.525.890 Tai Ji
-	E02.190.525.906 Therapeutic Touch
-	E02.190.525.937 Yoga
-	E02.190.599 Musculoskeletal Manipulations
-	E02.190.599.186 Kinesiology, Applied
-	E02.190.599.233 Manipulation, Chiropractic
-	E02.190.599.280 Manipulation, Osteopathic
-	E02.190.599.750 Therapy, Soft Tissue
-	E02.190.599.750.500 Acupressure
-	E02.190.599.750.750 Massage
-	E02.190.655 Naturopathy
-	E02.190.701 Organotherapy
-	E02.190.701.884 Tissue Therapy, Historical
-	E02.190.755 Phytotherapy
-	E02.190.755.100 Aromatherapy
-	E02.190.755.624 Eclecticism, Historical
-	E02.190.799 Reflexotherapy
-	E02.190.888 Sensory Art Therapies
-	E02.190.888.030 Acoustic Stimulation
-	E02.190.888.061 Aromatherapy
-	E02.190.888.124 Art Therapy
-	E02.190.888.249 Color Therapy
-	E02.190.888.374 Dance Therapy
-	E02.190.888.500 Music Therapy
-	E02.190.888.625 Play Therapy
-	E02.190.894 Speleotherapy
-	E02.190.901 Spiritual Therapies
-	E02.190.901.155 Faith Healing
-	E02.190.901.411 Magic

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	E02.190.901.433	Medicine, African Traditional
-	E02.190.901.455	Meditation
-	E02.190.901.500	Mental Healing
-	E02.190.901.740	Radiesthesia
-	E02.190.901.788	Shamanism
-	E02.190.901.830	Therapeutic Touch
-	E02.190.901.968	Witchcraft
-	E02.190.901.984	Yoga
New Heading	<b>E02.197</b>	<b>Conservative Treatment</b>
Old Tree	<b>E02.204</b>	<b>Continuity of Patient Care</b>
Old Tree	<b>E02.204.125</b>	<b>Patient Discharge</b>
Old Tree	<b>E02.204.249</b>	<b>Patient Handoff</b>
Old Tree	<b>E02.204.624</b>	<b>Patient Transfer</b>
Old Tree	<b>E02.204.718</b>	<b>Transition to Adult Care</b>
Old Tree	<b>E02.204.812</b>	<b>Transitional Care</b>
-	E02.218	Cosmetic Techniques
-	E02.218.042	Abdominoplasty
-	E02.218.085	Body Modification, Non-Therapeutic
-	E02.218.085.090	Body Piercing
-	E02.218.085.165	Circumcision, Female
-	E02.218.085.170	Circumcision, Male
-	E02.218.085.840	Tattooing
-	E02.218.170	Chemexfoliation
-	E02.218.210	Dermabrasion
-	E02.218.372	Hair Removal
-	E02.218.530	Lipectomy
-	E02.218.565	Mammoplasty
-	E02.218.565.210	Breast Implantation
-	E02.218.660	Mesotherapy
-	E02.218.707	Plasma Skin Regeneration
-	E02.218.755	Rhinoplasty
-	E02.218.765	Rhytidoplasty
-	E02.258	Cryotherapy
-	E02.258.750	Hypothermia, Induced
-	E02.258.750.500	Gastric Hypothermia



## MeSH Tree Changes for 2017

Type	Tree - heading	
-	E02.278	Decompression
-	E02.278.500	Lower Body Negative Pressure
-	E02.288	Delayed Diagnosis
-	E02.299	Directly Observed Therapy
-	E02.309	Drainage
-	E02.309.221	Drainage, Postural
-	E02.309.610	Negative-Pressure Wound Therapy
-	E02.309.805	Paracentesis
-	E02.319	Drug Therapy
-	E02.319.077	Antineoplastic Protocols
-	E02.319.077.500	Antineoplastic Combined Chemotherapy Protocols
-	E02.319.155	Chelation Therapy
-	E02.319.162	Chemoprevention
-	E02.319.162.150	Antibiotic Prophylaxis
-	E02.319.164	Chemoradiotherapy
-	E02.319.164.500	Chemoradiotherapy, Adjuvant
-	E02.319.170	Chemotherapy, Adjuvant
-	E02.319.218	Consolidation Chemotherapy
-	E02.319.267	Drug Administration Routes
-	E02.319.267.050	Administration, Inhalation
-	E02.319.267.082	Administration, Intravenous
-	E02.319.267.082.500	Infusions, Intravenous
-	E02.319.267.082.750	Injections, Intravenous
-	E02.319.267.100	Administration, Oral
-	E02.319.267.100.028	Administration, Buccal
-	E02.319.267.100.878	Administration, Sublingual
-	E02.319.267.120	Administration, Topical
-	E02.319.267.120.040	Administration, Buccal
-	E02.319.267.120.060	Administration, Cutaneous
-	E02.319.267.120.500	Administration, Intravaginal
-	E02.319.267.120.505	Administration, Intravesical
-	E02.319.267.120.655	Administration, Mucosal
-	E02.319.267.120.655.500	Administration, Intranasal
-	E02.319.267.120.655.750	Administration, Rectal
-	E02.319.267.120.805	Administration, Ophthalmic
-	E02.319.267.200	Chemotherapy, Cancer, Regional Perfusion

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E02.319.267.510                      Infusions, Parenteral
-	E02.319.267.510.520                      Infusions, Intra-Arterial
-	E02.319.267.510.555                      Infusions, Intralesional
-	E02.319.267.510.560                      Infusions, Intraosseous
-	E02.319.267.510.590                      Infusions, Intravenous
-	E02.319.267.510.692                      Infusions, Intraventricular
-	E02.319.267.510.794                      Infusions, Spinal
-	E02.319.267.510.795                      Infusions, Subcutaneous
-	E02.319.267.510.795.500                      Hypodermoclysis
-	E02.319.267.530                      Injections
-	E02.319.267.530.370                      Injections, Intra-Arterial
-	E02.319.267.530.380                      Injections, Intra-Articular
-	E02.319.267.530.380.500                      Viscosupplementation
-	E02.319.267.530.430                      Injections, Intralesional
-	E02.319.267.530.440                      Injections, Intralymphatic
-	E02.319.267.530.460                      Injections, Intramuscular
-	E02.319.267.530.475                      Injections, Intraocular
-	E02.319.267.530.475.500                      Intravitreal Injections
-	E02.319.267.530.490                      Injections, Intraperitoneal
-	E02.319.267.530.540                      Injections, Intravenous
-	E02.319.267.530.550                      Injections, Intraventricular
-	E02.319.267.530.580                      Injections, Spinal
-	E02.319.267.530.580.300                      Injections, Epidural
-	E02.319.267.530.580.300.145                      Blood Patch, Epidural
-	E02.319.267.530.620                      Injections, Subcutaneous
-	E02.319.267.530.620.410                      Injections, Intradermal
-	E02.319.267.530.620.570                      Injections, Jet
-	E02.319.267.530.620.570.100                      Biolistics
-	E02.319.267.530.620.785                      Mesotherapy
-	E02.319.267.530.655                      Injection, Intra-tympanic
-	E02.319.267.530.690                      Microinjections
-	E02.319.267.641                      Instillation, Drug
-	E02.319.267.650                      Iontophoresis
-	E02.319.267.700                      Phonophoresis
-	E02.319.283                      Drug Administration Schedule
-	E02.319.283.199                      Administration, Metronomic

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	E02.319.283.400	Drug Chronotherapy
-	E02.319.283.600	Pulse Therapy, Drug
-	E02.319.300	Drug Delivery Systems
-	E02.319.300.253	Delayed-Action Preparations
-	E02.319.300.380	Drug Carriers
-	E02.319.300.380.200	Dendrimers
-	E02.319.300.380.575	Nanocapsules
-	E02.319.300.380.600	Nanoconjugates
-	E02.319.300.508	Insulin Infusion Systems
-	E02.319.300.754	Pharmaceutical Vehicles
-	E02.319.305	Drug Dosage Calculations
-	E02.319.307	Drug Prescriptions
-	E02.319.307.312	Drug Substitution
-	E02.319.307.500	Off-Label Use
-	E02.319.310	Drug Therapy, Combination
-	E02.319.310.037	Antineoplastic Combined Chemotherapy Protocols
-	E02.319.310.075	Antiretroviral Therapy, Highly Active
-	E02.319.335	Drug Therapy, Computer-Assisted
-	E02.319.341	Electrochemotherapy
-	E02.319.347	Enema
New Heading	<b>E02.319.347.500</b>	<b>Barium Enema</b>
-	E02.319.353	Enzyme Therapy
-	E02.319.353.500	Enzyme Replacement Therapy
-	E02.319.360	Fluid Therapy
-	E02.319.360.500	Hypodermoclysis
-	E02.319.444	Home Infusion Therapy
-	E02.319.452	Hormone Replacement Therapy
-	E02.319.452.150	Estrogen Replacement Therapy
-	E02.319.490	Inappropriate Prescribing
-	E02.319.499	Induction Chemotherapy
-	E02.319.509	Maintenance Chemotherapy
-	E02.319.529	Medication Errors
-	E02.319.529.500	Medication Reconciliation
-	E02.319.529.750	Near Miss, Healthcare
-	E02.319.574	Molecular Targeted Therapy

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	E02.319.620	Opiate Substitution Treatment
-	E02.319.630	Orthomolecular Therapy
-	E02.319.685	Photochemotherapy
-	E02.319.694	Pleurodesis
-	E02.319.698	Polypharmacy
-	E02.319.698.500	Deprescriptions
-	E02.319.703	Premedication
-	E02.319.703.150	Antibiotic Prophylaxis
-	E02.319.754	Prescription Drug Misuse
-	E02.319.754.500	Drug Overdose
-	E02.319.754.750	Prescription Drug Overuse
-	E02.319.805	Sclerotherapy
-	E02.319.890	Self Administration
-	E02.319.900	Self Medication
-	E02.319.913	Thrombolytic Therapy
-	E02.319.913.500	Hirudin Therapy
-	E02.331	Electric Stimulation Therapy
-	E02.331.200	Cardiac Pacing, Artificial
-	E02.331.200.500	Cardiac Resynchronization Therapy
-	E02.331.300	Deep Brain Stimulation
-	E02.331.350	Electric Countershock
-	E02.331.399	Electroacupuncture
-	E02.331.599	Pulsed Radiofrequency Treatment
-	E02.331.699	Spinal Cord Stimulation
-	E02.331.750	Transcranial Direct Current Stimulation
-	E02.331.800	Transcutaneous Electric Nerve Stimulation
-	E02.331.900	Vagus Nerve Stimulation
-	E02.365	Emergency Treatment
-	E02.365.152	Advanced Trauma Life Support Care
-	E02.365.305	First Aid
-	E02.365.476	Heimlich Maneuver
-	E02.365.647	Resuscitation
-	E02.365.647.110	Cardiopulmonary Resuscitation
-	E02.365.647.110.500	Advanced Cardiac Life Support
-	E02.365.647.375	Heart Massage
-	E02.365.647.729	Respiration, Artificial

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E02.365.647.740                      Resuscitation Orders
-	E02.365.839                              Transportation of Patients
-	E02.421                                      Feeding Methods
-	E02.421.150                              Bottle Feeding
-	E02.421.360                              Enteral Nutrition
-	E02.421.505                              Parenteral Nutrition
-	E02.421.505.550                        Parenteral Nutrition, Home
-	E02.421.505.550.565                    Parenteral Nutrition, Home Total
-	E02.421.505.575                        Parenteral Nutrition, Total
-	E02.421.505.575.565                    Parenteral Nutrition, Home Total
-	E02.467                                      Fetal Therapies
-	E02.467.500                              Blood Transfusion, Intrauterine
-	E02.467.750                              Fetoscopy
-	E02.514                                      Hemodilution
-	E02.520                                      Hemostatic Techniques
-	E02.520.089                              Argon Plasma Coagulation
-	E02.520.360                              Embolization, Therapeutic
-	E02.520.360.150                        Chemoembolization, Therapeutic
-	E02.520.360.575                        Uterine Artery Embolization
-	E02.520.392                              Endotamponade
-	E02.520.392.500                        Balloon Occlusion
-	E02.520.392.500.500                    Uterine Balloon Tamponade
-	E02.520.425                              Hemostasis, Endoscopic
-	E02.520.490                              Hemostasis, Surgical
-	E02.520.745                              Light Coagulation
-	E02.520.745.410                        Laser Coagulation
-	E02.547                                      Hygiene
New Tree	<a href="#">E02.547.300</a> <a href="#">Hand Hygiene</a>
-	E02.547.600                              Oral Hygiene
-	E02.547.800                              Skin Care
-	E02.547.800.500                        Skin Cream
-	E02.565                                      Hyperthermia, Induced
-	E02.565.020                              Ammotherapy
-	E02.565.280                              Diathermy
-	E02.565.280.853                        Short-Wave Therapy

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E02.565.280.945                      Ultrasonic Therapy
-	E02.565.280.945.399                      High-Intensity Focused Ultrasound Ablation
-	E02.565.280.945.399.500                      Ultrasound, High-Intensity Focused, Transrectal
-	E02.565.640                      Steam Bath
-	E02.574                      Precision Medicine
-	E02.583                      Insufflation
-	E02.585                      Intubation
-	E02.585.412                      Intubation, Gastrointestinal
-	E02.585.578                      Intubation, Intratracheal
-	E02.585.578.475                      Laryngeal Masks
-	E02.587                      Ischemic Postconditioning
-	E02.592                      Ischemic Preconditioning
-	E02.592.325                      Ischemic Preconditioning, Myocardial
-	E02.594                      Laser Therapy
-	E02.594.060                      Angioplasty, Laser
-	E02.594.060.080                      Angioplasty, Balloon, Laser-Assisted
-	E02.594.480                      Corneal Surgery, Laser
-	E02.594.480.500                      Keratectomy, Subepithelial, Laser-Assisted
-	E02.594.480.750                      Keratomileusis, Laser In Situ
-	E02.594.480.875                      Photorefractive Keratectomy
-	E02.594.530                      Laser Coagulation
-	E02.594.540                      Low-Level Light Therapy
-	E02.594.550                      Lithotripsy, Laser
-	E02.596                      Leeching
-	E02.600                      Lithotripsy
-	E02.600.500                      Lithotripsy, Laser
-	E02.621                      Magnetic Field Therapy
-	E02.621.820                      Transcranial Magnetic Stimulation
-	E02.631                      Mechanical Thrombolysis
-	E02.642                      Nutrition Therapy
-	E02.642.249                      Diet Therapy
-	E02.642.249.200                      Caloric Restriction
New Heading	<b>E02.642.249.220                      Diet, Carbohydrate Loading</b>
-	E02.642.249.240                      Diet, Diabetic
-	E02.642.249.245                      Diet, Carbohydrate-Restricted

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	E02.642.249.260	Diet, Fat-Restricted
-	E02.642.249.265	Diet, Gluten-Free
-	E02.642.249.270	Diet, Mediterranean
-	E02.642.249.275	Diet, Paleolithic
-	E02.642.249.280	Diet, Protein-Restricted
-	E02.642.249.285	Diet, Reducing
-	E02.642.249.290	Diet, Sodium-Restricted
-	E02.642.249.300	Diet, Vegetarian
-	E02.642.249.300.500	Diet, Macrobiotic
-	E02.642.249.300.750	Diet, Vegan
-	E02.642.249.650	Ketogenic Diet
-	E02.642.500	Nutritional Support
-	E02.642.500.360	Enteral Nutrition
-	E02.642.500.505	Parenteral Nutrition
-	E02.642.500.505.550	Parenteral Nutrition, Home
-	E02.642.500.505.550.565	Parenteral Nutrition, Home Total
-	E02.642.500.505.750	Parenteral Nutrition, Total
-	E02.642.500.505.750.565	Parenteral Nutrition, Home Total
-	E02.674	Organ Sparing Treatments
-	E02.706	Orthokeratologic Procedures
-	E02.718	Orthopedic Procedures
-	E02.718.275	Cementoplasty
-	E02.718.275.500	Vertebroplasty
-	E02.718.275.500.500	Kyphoplasty
-	E02.718.500	Intervertebral Disc Chemolysis
-	E02.718.625	Manipulation, Orthopedic
New Heading	<b>E02.718.688</b>	<b>Posterior Cruciate Ligament Reconstruction</b>
New Heading	<b>E02.718.750</b>	<b>Ulnar Collateral Ligament Reconstruction</b>
-	E02.718.875	Viscosupplementation
-	E02.730	Orthoptics
-	E02.745	Pain Management
-	E02.760	Patient Care
Old Tree	<b>E02.760.058</b>	<b>Aftercare</b>
-	E02.760.106	Ambulatory Care

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	E02.760.106.500	Peritoneal Dialysis, Continuous Ambulatory
-	E02.760.148	Bloodless Medical and Surgical Procedures
New Tree	<a href="#">E02.760.169</a>	<a href="#">Continuity of Patient Care</a>
New Tree	<a href="#">E02.760.169.063</a>	<a href="#">Aftercare</a>
New Tree	<a href="#">E02.760.169.063.500</a>	<a href="#">Rehabilitation</a>
New Tree	<a href="#">E02.760.169.063.500.067</a>	<a href="#">Activities of Daily Living</a>
New Tree	<a href="#">E02.760.169.063.500.083</a>	<a href="#">Animal Assisted Therapy</a>
New Tree	<a href="#">E02.760.169.063.500.083.500</a>	<a href="#">Equine-Assisted Therapy</a>
New Tree	<a href="#">E02.760.169.063.500.100</a>	<a href="#">Art Therapy</a>
New Tree	<a href="#">E02.760.169.063.500.169</a>	<a href="#">Bibliotherapy</a>
New Heading	<b><a href="#">E02.760.169.063.500.185</a></b>	<b><a href="#">Cardiac Rehabilitation</a></b>
New Tree	<a href="#">E02.760.169.063.500.200</a>	<a href="#">Correction of Hearing Impairment</a>
New Tree	<a href="#">E02.760.169.063.500.200.221</a>	<a href="#">Communication Methods, Total</a>
New Tree	<a href="#">E02.760.169.063.500.200.432</a>	<a href="#">Lipreading</a>
New Tree	<a href="#">E02.760.169.063.500.200.609</a>	<a href="#">Manual Communication</a>
New Tree	<a href="#">E02.760.169.063.500.200.609.668</a>	<a href="#">Sign Language</a>
New Tree	<a href="#">E02.760.169.063.500.230</a>	<a href="#">Dance Therapy</a>
New Tree	<a href="#">E02.760.169.063.500.335</a>	<a href="#">Early Ambulation</a>
New Tree	<a href="#">E02.760.169.063.500.387</a>	<a href="#">Exercise Therapy</a>
New Tree	<a href="#">E02.760.169.063.500.387.500</a>	<a href="#">Motion Therapy, Continuous Passive</a>
New Tree	<a href="#">E02.760.169.063.500.387.750</a>	<a href="#">Muscle Stretching Exercises</a>
New Tree	<a href="#">E02.760.169.063.500.387.812</a>	<a href="#">Plyometric Exercise</a>
New	<a href="#">E02.760.169.063.500.387.875</a>	<a href="#">Resistance Training</a>



## MeSH Tree Changes for 2017

Type	Tree - heading	
Tree		
New Tree	<a href="#">E02.760.169.063.500.440</a>	Music Therapy
New Tree	<a href="#">E02.760.169.063.500.477</a>	Neurological Rehabilitation
New Heading	<b><a href="#">E02.760.169.063.500.477.500</a></b>	<b>Stroke Rehabilitation</b>
New Tree	<a href="#">E02.760.169.063.500.489</a>	Occupational Therapy
New Tree	<a href="#">E02.760.169.063.500.580</a>	Recreation Therapy
New Tree	<a href="#">E02.760.169.063.500.727</a>	Rehabilitation of Speech and Language Disorders
New Tree	<a href="#">E02.760.169.063.500.727.344</a>	Language Therapy
New Tree	<a href="#">E02.760.169.063.500.727.442</a>	Myofunctional Therapy
New Tree	<a href="#">E02.760.169.063.500.727.541</a>	Speech, Alaryngeal
New Tree	<a href="#">E02.760.169.063.500.727.541.677</a>	Speech, Esophageal
New Tree	<a href="#">E02.760.169.063.500.727.552</a>	Speech Therapy
New Tree	<a href="#">E02.760.169.063.500.727.862</a>	Voice Training
New Tree	<a href="#">E02.760.169.063.500.782</a>	Rehabilitation, Vocational
New Tree	<a href="#">E02.760.169.063.500.891</a>	Telerehabilitation
New Tree	<a href="#">E02.760.169.125</a>	Patient Discharge
New Tree	<a href="#">E02.760.169.249</a>	Patient Handoff
New Tree	<a href="#">E02.760.169.624</a>	Patient Transfer
New Tree	<a href="#">E02.760.169.718</a>	Transition to Adult Care
New Tree	<a href="#">E02.760.169.812</a>	Transitional Care
-	<a href="#">E02.760.190</a>	Critical Care
-	<a href="#">E02.760.190.405</a>	Intensive Care, Neonatal
-	<a href="#">E02.760.215</a>	Custodial Care
-	<a href="#">E02.760.246</a>	Day Care, Medical

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	E02.760.300	Episode of Care
-	E02.760.352	Foster Home Care
-	E02.760.400	Hospitalization
-	E02.760.400.480	Length of Stay
-	E02.760.400.600	Patient Admission
-	E02.760.400.610	Patient Discharge
-	E02.760.400.615	Patient Handoff
-	E02.760.400.620	Patient Readmission
-	E02.760.400.630	Patient Transfer
-	E02.760.415	Institutionalization
-	E02.760.415.280	Deinstitutionalization
-	E02.760.440	Life Support Care
-	E02.760.440.028	Advanced Cardiac Life Support
-	E02.760.440.040	Advanced Trauma Life Support Care
-	E02.760.476	Long-Term Care
-	E02.760.564	Night Care
-	E02.760.611	Nursing Care
-	E02.760.611.470	Home Nursing
-	E02.760.611.470.610	Respite Care
-	E02.760.611.735	Primary Care Nursing
-	E02.760.666	Palliative Care
-	E02.760.670	Patient Positioning
-	E02.760.670.500	Kangaroo-Mother Care Method
-	E02.760.703	Perinatal Care
-	E02.760.703.500	Postnatal Care
-	E02.760.731	Perioperative Care
-	E02.760.731.400	Intraoperative Care
-	E02.760.731.500	Perioperative Nursing
-	E02.760.731.700	Postoperative Care
-	E02.760.775	Preconception Care
-	E02.760.786	Prenatal Care
-	E02.760.795	Preoperative Care
-	E02.760.850	Subacute Care
-	E02.760.905	Terminal Care
-	E02.760.905.199	Euthanasia
-	E02.760.905.199.249	Euthanasia, Active

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E02.760.905.199.249.200 Euthanasia, Active, Voluntary
-	E02.760.905.199.249.750 Euthanasia, Animal
-	E02.760.905.199.500 Euthanasia, Passive
-	E02.760.905.400 Hospice Care
-	E02.760.905.700 Resuscitation Orders
-	E02.760.905.850 Suicide, Assisted
-	E02.760.928 Time-to-Treatment
-	E02.760.940 Transitional Care
-	E02.760.952 Withholding Treatment
-	E02.760.952.500 Euthanasia, Passive
-	E02.765 Patient Care Bundles
-	E02.770 Patient Isolation
-	E02.774 Phototherapy
-	E02.774.215 Color Therapy
-	E02.774.430 Heliotherapy
-	E02.774.465 Intense Pulsed Light Therapy
-	E02.774.500 Low-Level Light Therapy
-	E02.774.722 Photochemotherapy
-	E02.774.722.435 Hematoporphyrin Photoradiation
-	E02.774.945 Ultraviolet Therapy
-	E02.774.945.500 PUVA Therapy
-	E02.774.945.500.500 Photopheresis
-	E02.779 Physical Therapy Modalities
-	E02.779.124 Animal Assisted Therapy
-	E02.779.124.500 Equine-Assisted Therapy
-	E02.779.358 Drainage, Postural
-	E02.779.468 Electric Stimulation Therapy
-	E02.779.468.399 Electroacupuncture
-	E02.779.468.599 Pulsed Radiofrequency Treatment
-	E02.779.468.699 Spinal Cord Stimulation
-	E02.779.468.800 Transcutaneous Electric Nerve Stimulation
-	E02.779.474 Exercise Movement Techniques
-	E02.779.474.124 Breathing Exercises
-	E02.779.474.124.500 Qigong
-	E02.779.474.186 Dance Therapy
-	E02.779.474.913 Tai Ji

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E02.779.474.937 Yoga
-	E02.779.483 Exercise Therapy
-	E02.779.483.500 Motion Therapy, Continuous Passive
-	E02.779.483.750 Muscle Stretching Exercises
-	E02.779.483.812 Plyometric Exercise
-	E02.779.483.875 Resistance Training
-	E02.779.492 Hydrotherapy
-	E02.779.492.500 Therapeutic Irrigation
-	E02.779.867 Musculoskeletal Manipulations
-	E02.779.867.344 Kinesiology, Applied
-	E02.779.867.433 Manipulation, Orthopedic
-	E02.779.867.444 Manipulation, Osteopathic
-	E02.779.867.466 Manipulation, Spinal
-	E02.779.867.761 Motion Therapy, Continuous Passive
-	E02.779.867.880 Therapy, Soft Tissue
-	E02.779.867.880.500 Acupressure
-	E02.779.867.880.750 Massage
-	E02.779.933 Myofunctional Therapy
-	E02.785 Placebos
-	E02.792 Preservation, Biological
-	E02.792.156 Cryopreservation
-	E02.792.156.260 Freeze Drying
-	E02.792.156.260.270 Freeze Substitution
-	E02.792.643 Refrigeration
-	E02.792.833 Tissue Preservation
-	E02.792.833.230 Blood Preservation
-	E02.792.833.230.500 Blood Safety
-	E02.792.833.660 Organ Preservation
-	E02.792.833.890 Semen Preservation
-	E02.794 Prosthesis Fitting
-	E02.794.500 Prosthesis Retention
-	E02.800 Punctures
-	E02.800.550 Paracentesis
-	E02.800.550.310 Arthrocentesis
-	E02.800.550.465 Cordocentesis
-	E02.800.550.620 Pericardiocentesis

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E02.800.550.810 Thoracentesis
-	E02.800.550.905 Tympanocentesis
-	E02.800.558 Phlebotomy
-	E02.800.558.500 Bloodletting
-	E02.800.779 Spinal Puncture
-	E02.815 Radiotherapy
-	E02.815.150 Brachytherapy
-	E02.815.160 Chemoradiotherapy
-	E02.815.160.500 Chemoradiotherapy, Adjuvant
-	E02.815.190 Cranial Irradiation
-	E02.815.190.600 Pituitary Irradiation
-	E02.815.230 Craniospinal Irradiation
-	E02.815.250 Heavy Ion Radiotherapy
-	E02.815.270 Hemibody Irradiation
-	E02.815.350 Lymphatic Irradiation
-	E02.815.435 Proton Therapy
-	E02.815.520 Radioimmunotherapy
-	E02.815.530 Radiosurgery
-	E02.815.565 Radiotherapy Setup Errors
-	E02.815.600 Radiotherapy, Adjuvant
-	E02.815.635 Radiotherapy, Computer-Assisted
-	E02.815.635.700 Radiotherapy, Conformal
-	E02.815.635.700.700 Radiotherapy, Intensity-Modulated
-	E02.815.639 Radiotherapy Dosage
-	E02.815.639.200 Dose Fractionation
-	E02.815.639.200.500 Dose Hypofractionation
-	E02.815.639.600 Dose Hypofractionation
-	E02.815.722 Radiotherapy, High-Energy
-	E02.815.722.500 Neutron Capture Therapy
-	E02.815.722.500.100 Boron Neutron Capture Therapy
-	E02.815.722.820 Radioisotope Teletherapy
-	E02.815.768 Radiotherapy, Image-Guided
-	E02.815.791 Re-Irradiation
-	E02.815.814 Whole-Body Irradiation
-	E02.815.907 X-Ray Therapy
-	E02.831 Rehabilitation

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	E02.831.067	Activities of Daily Living
Old Tree	<b>E02.831.083</b>	<b>Animal Assisted Therapy</b>
Old Tree	<b>E02.831.083.500</b>	<b>Equine-Assisted Therapy</b>
-	E02.831.100	Art Therapy
-	E02.831.169	Bibliotherapy
New Heading	<b>E02.831.185</b>	<b>Cardiac Rehabilitation</b>
-	E02.831.200	Correction of Hearing Impairment
-	E02.831.200.221	Communication Methods, Total
-	E02.831.200.432	Lipreading
-	E02.831.200.609	Manual Communication
-	E02.831.200.609.668	Sign Language
-	E02.831.230	Dance Therapy
-	E02.831.335	Early Ambulation
Old Tree	<b>E02.831.387</b>	<b>Exercise Therapy</b>
Old Tree	<b>E02.831.387.500</b>	<b>Motion Therapy, Continuous Passive</b>
Old Tree	<b>E02.831.387.750</b>	<b>Muscle Stretching Exercises</b>
Old Tree	<b>E02.831.387.812</b>	<b>Plyometric Exercise</b>
Old Tree	<b>E02.831.387.875</b>	<b>Resistance Training</b>
-	E02.831.440	Music Therapy
-	E02.831.477	Neurological Rehabilitation
New Heading	<b>E02.831.477.500</b>	<b>Stroke Rehabilitation</b>
-	E02.831.489	Occupational Therapy
New Tree	<b>E02.831.535</b>	<b>Physical Therapy Modalities</b>
New Tree	<b>E02.831.535.358</b>	<b>Drainage, Postural</b>
New Tree	<b>E02.831.535.468</b>	<b>Electric Stimulation Therapy</b>
New Tree	<b>E02.831.535.468.399</b>	<b>Electroacupuncture</b>
New Tree	<b>E02.831.535.468.599</b>	<b>Pulsed Radiofrequency Treatment</b>
New Tree	<b>E02.831.535.468.699</b>	<b>Spinal Cord Stimulation</b>
New Tree	<b>E02.831.535.468.800</b>	<b>Transcutaneous Electric Nerve Stimulation</b>
New	<b>E02.831.535.483</b>	<b>Exercise Therapy</b>

## MeSH Tree Changes for 2017

Type	Tree - heading	
Tree		
New Tree	<a href="#">E02.831.535.483.500</a>	Motion Therapy, Continuous Passive
New Tree	<a href="#">E02.831.535.483.750</a>	Muscle Stretching Exercises
New Tree	<a href="#">E02.831.535.483.812</a>	Plyometric Exercise
New Tree	<a href="#">E02.831.535.483.875</a>	Resistance Training
New Tree	<a href="#">E02.831.535.492</a>	Hydrotherapy
New Tree	<a href="#">E02.831.535.492.500</a>	Therapeutic Irrigation
New Tree	<a href="#">E02.831.535.867</a>	Musculoskeletal Manipulations
New Tree	<a href="#">E02.831.535.867.344</a>	Kinesiology, Applied
New Tree	<a href="#">E02.831.535.867.433</a>	Manipulation, Orthopedic
New Tree	<a href="#">E02.831.535.867.444</a>	Manipulation, Osteopathic
New Tree	<a href="#">E02.831.535.867.466</a>	Manipulation, Spinal
New Tree	<a href="#">E02.831.535.867.761</a>	Motion Therapy, Continuous Passive
New Tree	<a href="#">E02.831.535.867.880</a>	Therapy, Soft Tissue
New Tree	<a href="#">E02.831.535.867.880.500</a>	Acupressure
New Tree	<a href="#">E02.831.535.867.880.750</a>	Massage
New Tree	<a href="#">E02.831.535.933</a>	Myofunctional Therapy
-	<a href="#">E02.831.580</a>	Recreation Therapy
-	<a href="#">E02.831.727</a>	Rehabilitation of Speech and Language Disorders
-	<a href="#">E02.831.727.344</a>	Language Therapy
-	<a href="#">E02.831.727.442</a>	Myofunctional Therapy
-	<a href="#">E02.831.727.541</a>	Speech, Alaryngeal
-	<a href="#">E02.831.727.541.677</a>	Speech, Esophageal
-	<a href="#">E02.831.727.552</a>	Speech Therapy
-	<a href="#">E02.831.727.862</a>	Voice Training
-	<a href="#">E02.831.782</a>	Rehabilitation, Vocational

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	E02.831.891	Telerehabilitation
-	E02.849	Rejuvenation
-	E02.860	Remission Induction
-	E02.860.500	Induction Chemotherapy
-	E02.870	Renal Replacement Therapy
-	E02.870.300	Renal Dialysis
-	E02.870.300.200	Hemodiafiltration
-	E02.870.300.300	Hemodialysis, Home
-	E02.870.300.650	Peritoneal Dialysis
-	E02.870.300.650.500	Peritoneal Dialysis, Continuous Ambulatory
-	E02.870.500	Kidney Transplantation
-	E02.875	Reproductive Techniques
-	E02.875.194	Contraception
-	E02.875.194.250	Coitus Interruptus
-	E02.875.194.285	Contraception, Barrier
-	E02.875.194.300	Contraception, Immunologic
-	E02.875.194.540	Contraception, Postcoital
-	E02.875.194.605	Natural Family Planning Methods
-	E02.875.194.670	Ovulation Inhibition
-	E02.875.194.910	Sterilization, Reproductive
-	E02.875.390	Fallopian Tube Patency Tests
-	E02.875.690	Ovulation Detection
-	E02.875.745	Ovulation Prediction
-	E02.875.800	Reproductive Techniques, Assisted
New Heading	<b>E02.875.800.250</b>	<b>Donor Conception</b>
-	E02.875.800.500	Embryo Transfer
-	E02.875.800.500.500	Single Embryo Transfer
-	E02.875.800.625	Fertility Preservation
-	E02.875.800.750	Fertilization in Vitro
-	E02.875.800.750.350	Mitochondrial Replacement Therapy
-	E02.875.800.750.700	Sperm Injections, Intracytoplasmic
-	E02.875.800.875	Gamete Intrafallopian Transfer
-	E02.875.800.906	In Vitro Oocyte Maturation Techniques
-	E02.875.800.937	Insemination, Artificial
-	E02.875.800.937.515	Insemination, Artificial, Heterologous



## MeSH Tree Changes for 2017

Type	Tree - heading
-	E02.875.800.937.525 Insemination, Artificial, Homologous
-	E02.875.800.968 Oocyte Donation
-	E02.875.800.976 Oocyte Retrieval
-	E02.875.800.984 Ovulation Induction
-	E02.875.800.984.500 Superovulation
-	E02.875.800.986 Posthumous Conception
-	E02.875.800.988 Sperm Retrieval
-	E02.875.800.992 Zygote Intrafallopian Transfer
-	E02.875.900 Tocolysis
-	E02.880 Respiratory Therapy
-	E02.880.130 Chest Wall Oscillation
-	E02.880.150 Drainage, Postural
-	E02.880.301 Extracorporeal Membrane Oxygenation
-	E02.880.690 Oxygen Inhalation Therapy
-	E02.880.690.490 Hyperbaric Oxygenation
-	E02.880.820 Respiration, Artificial
-	E02.880.820.508 High-Frequency Ventilation
-	E02.880.820.508.510 High-Frequency Jet Ventilation
-	E02.880.820.516 Interactive Ventilatory Support
-	E02.880.820.525 Liquid Ventilation
-	E02.880.820.657 Noninvasive Ventilation
-	E02.880.820.790 Positive-Pressure Respiration
-	E02.880.820.790.259 Continuous Positive Airway Pressure
-	E02.880.820.790.520 Intermittent Positive-Pressure Breathing
-	E02.880.820.790.550 Intermittent Positive-Pressure Ventilation
-	E02.880.820.950 Ventilator Weaning
-	E02.887 Retreatment
-	E02.887.500 Re-Irradiation
-	E02.891 Rewarming
-	E02.895 Salvage Therapy
-	E02.897 Secondary Prevention
-	E02.900 Self Care
-	E02.900.100 Blood Glucose Self-Monitoring
-	E02.900.890 Self Administration
-	E02.900.900 Self Medication
-	E02.906 Sex Reassignment Procedures

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E02.906.500                      Sex Reassignment Surgery
-	E02.912                              Sorption Detoxification
-	E02.912.300                      Enterosorption
-	E02.912.430                      Hemoperfusion
-	E02.912.715                      Plasmapheresis
-	E02.912.800                      Renal Dialysis
-	E02.912.800.200                  Hemodiafiltration
-	E02.912.800.300                  Hemodialysis, Home
-	E02.912.800.650                  Peritoneal Dialysis
-	E02.912.800.650.500              Peritoneal Dialysis, Continuous Ambulatory
-	E02.919                              Tertiary Prevention
-	E02.921                              Theranostic Nanomedicine
-	E02.926                              Therapeutic Occlusion
-	E02.926.500                      Embolization, Therapeutic
-	E02.926.500.074                  Balloon Occlusion
-	E02.926.500.074.500              Uterine Balloon Tamponade
-	E02.926.500.150                  Chemoembolization, Therapeutic
-	E02.926.500.575                  Uterine Artery Embolization
-	E02.931                              Therapies, Investigational
-	E02.931.500                      Compassionate Use Trials
-	E02.950                              Therapy, Computer-Assisted
-	E02.950.825                      Radiotherapy Planning, Computer-Assisted
-	E02.950.875                      Surgery, Computer-Assisted
-	E02.950.875.500                  Robotic Surgical Procedures
-	E02.960                              Therapy with Helminths
-	E03                                      Anesthesia and Analgesia
-	E03.091                              Analgesia
-	E03.091.048                      Acupuncture Analgesia
-	E03.091.080                      Analgesia, Epidural
-	E03.091.110                      Analgesia, Obstetrical
-	E03.091.120                      Analgesia, Patient-Controlled
-	E03.091.214                      Audioanalgesia
-	E03.091.322                      Diffuse Noxious Inhibitory Control
-	E03.091.430                      Interpleural Analgesia
-	E03.091.646                      Neuroleptanalgesia
-	E03.091.823                      Transcutaneous Electric Nerve Stimulation

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E03.091.823.500 Electroacupuncture
-	E03.155 Anesthesia
-	E03.155.086 Anesthesia, Conduction
-	E03.155.086.131 Anesthesia, Epidural
-	E03.155.086.131.100 Anesthesia, Caudal
-	E03.155.086.231 Anesthesia, Local
-	E03.155.086.331 Anesthesia, Spinal
-	E03.155.086.711 Nerve Block
-	E03.155.086.711.299 Autonomic Nerve Block
-	E03.155.086.711.299.500 Sphenopalatine Ganglion Block
-	E03.155.086.711.649 Brachial Plexus Block
-	E03.155.086.711.824 Cervical Plexus Block
-	E03.155.141 Anesthesia, Dental
-	E03.155.141.481 Hypnosis, Dental
-	E03.155.197 Anesthesia, General
-	E03.155.197.197 Anesthesia, Inhalation
-	E03.155.197.197.280 Anesthesia, Closed-Circuit
-	E03.155.197.364 Anesthesia, Rectal
-	E03.155.197.682 Balanced Anesthesia
-	E03.155.253 Anesthesia, Endotracheal
-	E03.155.308 Anesthesia, Intravenous
-	E03.155.364 Anesthesia, Obstetrical
-	E03.155.441 Cryoanesthesia
-	E03.155.519 Electroacupuncture
-	E03.155.675 Hypnosis, Anesthetic
-	E03.155.675.481 Hypnosis, Dental
-	E03.160 Anesthesia Recovery Period
-	E03.250 Conscious Sedation
-	E03.295 Deep Sedation
-	E03.340 Electronarcosis
-	E03.545 Hypotension, Controlled
-	E03.706 Neuromuscular Blockade
-	E03.806 Preanesthetic Medication
-	E04 Surgical Procedures, Operative
-	E04.014 Ablation Techniques
-	E04.014.085 Catheter Ablation

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E04.014.170                      Cautery
-	E04.014.170.402                  Electrocoagulation
-	E04.014.170.402.054              Argon Plasma Coagulation
-	E04.014.180                      Cryosurgery
-	E04.014.240                      Endometrial Ablation Techniques
-	E04.014.380                      High-Intensity Focused Ultrasound Ablation
-	E04.014.380.500                  Ultrasound, High-Intensity Focused, Transrectal
-	E04.014.520                      Laser Therapy
-	E04.014.520.060                  Angioplasty, Laser
-	E04.014.520.060.080              Angioplasty, Balloon, Laser-Assisted
-	E04.014.520.480                  Corneal Surgery, Laser
-	E04.014.520.480.500              Keratectomy, Subepithelial, Laser-Assisted
-	E04.014.520.480.750              Keratomileusis, Laser In Situ
-	E04.014.520.480.875              Photorefractive Keratectomy
-	E04.014.520.530                  Laser Coagulation
-	E04.014.520.550                  Lithotripsy, Laser
-	E04.030                          Ambulatory Surgical Procedures
-	E04.035                          Anastomosis, Surgical
-	E04.035.070                      Anastomosis, Roux-en-Y
-	E04.035.087                      Arteriovenous Shunt, Surgical
-	E04.035.112                      Axillofemoral Bypass Grafting
-	E04.035.137                      Blalock-Taussig Procedure
-	E04.035.188                      Cerebrospinal Fluid Shunts
-	E04.035.188.850                  Ventriculoperitoneal Shunt
-	E04.035.188.957                  Ventriculostomy
-	E04.035.195                      Cholecystostomy
-	E04.035.200                      Choledochostomy
-	E04.035.335                      Endolymphatic Shunt
-	E04.035.398                      Gastroenterostomy
-	E04.035.398.385                  Gastric Bypass
-	E04.035.410                      Heart Bypass, Right
-	E04.035.410.295                  Fontan Procedure
-	E04.035.415                      Jejunioileal Bypass
-	E04.035.703                      Pancreaticojejunostomy
-	E04.035.720                      Pericardial Window Techniques
-	E04.035.735                      Peritoneovenous Shunt

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E04.035.760 Portasystemic Shunt, Surgical
-	E04.035.760.755 Portacaval Shunt, Surgical
-	E04.035.760.795 Portasystemic Shunt, Transjugular Intrahepatic
-	E04.035.760.835 Splenorenal Shunt, Surgical
-	E04.035.775 Portoenterostomy, Hepatic
-	E04.035.832 Salpingostomy
-	E04.035.956 Vasovasostomy
-	E04.050 Assisted Circulation
-	E04.050.215 Counterpulsation
-	E04.050.215.400 Intra-Aortic Balloon Pumping
-	E04.050.430 Heart-Assist Devices
-	E04.062 Bariatric Surgery
-	E04.062.249 Gastric Bypass
-	E04.062.750 Gastroplasty
-	E04.062.875 Jejunioileal Bypass
-	E04.062.937 Lipectomy
-	E04.074 Biopsy
-	E04.074.119 Biopsy, Needle
-	E04.074.119.500 Biopsy, Fine-Needle
-	E04.074.119.500.500 Endoscopic Ultrasound-Guided Fine Needle Aspiration
-	E04.074.119.750 Biopsy, Large-Core Needle
-	E04.074.149 Chorionic Villi Sampling
-	E04.074.160 Conization
-	E04.074.370 Image-Guided Biopsy
-	E04.074.370.500 Endoscopic Ultrasound-Guided Fine Needle Aspiration
-	E04.074.422 Papanicolaou Test
-	E04.074.580 Sentinel Lymph Node Biopsy
-	E04.074.790 Spinal Puncture
-	E04.074.800 Vaginal Smears
-	E04.079 Bloodless Medical and Surgical Procedures
-	E04.085 Body Modification, Non-Therapeutic
-	E04.085.090 Body Piercing
-	E04.085.165 Circumcision, Female
-	E04.085.170 Circumcision, Male
-	E04.085.840 Tattooing
-	E04.100 Cardiovascular Surgical Procedures

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E04.100.376                      Cardiac Surgical Procedures
-	E04.100.376.031                  Arterial Switch Operation
-	E04.100.376.062                  Cardiac Valve Annuloplasty
-	E04.100.376.062.500              Mitral Valve Annuloplasty
-	E04.100.376.125                  Cardiomyoplasty
-	E04.100.376.374                  Heart Arrest, Induced
-	E04.100.376.374.500              Circulatory Arrest, Deep Hypothermia Induced
-	E04.100.376.410                  Heart Bypass, Right
-	E04.100.376.410.295              Fontan Procedure
-	E04.100.376.458                  Heart Massage
-	E04.100.376.475                  Heart Transplantation
-	E04.100.376.475.450              Heart-Lung Transplantation
-	E04.100.376.485                  Heart Valve Prosthesis Implantation
-	E04.100.376.485.500              Transcatheter Aortic Valve Replacement
-	E04.100.376.719                  Myocardial Revascularization
-	E04.100.376.719.100              Angioplasty, Balloon, Coronary
-	E04.100.376.719.125              Atherectomy, Coronary
-	E04.100.376.719.332              Coronary Artery Bypass
-	E04.100.376.719.332.199           Coronary Artery Bypass, Off-Pump
-	E04.100.376.719.332.400           Internal Mammary-Coronary Artery Anastomosis
-	E04.100.376.719.666              Transmyocardial Laser Revascularization
-	E04.100.376.724                  Norwood Procedures
-	E04.100.376.724.500              Fontan Procedure
-	E04.100.376.730                  Pericardial Window Techniques
-	E04.100.376.735                  Pericardiectomy
-	E04.100.376.745                  Pericardiocentesis
-	E04.100.700                      Reperfusion
-	E04.100.700.600                  Myocardial Reperfusion
-	E04.100.814                      Vascular Surgical Procedures
-	E04.100.814.334                  Axillofemoral Bypass Grafting
-	E04.100.814.445                  Embolectomy
-	E04.100.814.445.500              Balloon Embolectomy
-	E04.100.814.456                  Endarterectomy
-	E04.100.814.456.250              Endarterectomy, Carotid
-	E04.100.814.529                  Endovascular Procedures
-	E04.100.814.529.124              Angioplasty

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E04.100.814.529.124.060                      Angioplasty, Balloon
-	E04.100.814.529.124.060.100                      Angioplasty, Balloon, Coronary
-	E04.100.814.529.124.060.105                      Angioplasty, Balloon, Laser-Assisted
-	E04.100.814.529.124.075                      Angioplasty, Laser
-	E04.100.814.529.124.075.080                      Angioplasty, Balloon, Laser-Assisted
-	E04.100.814.529.124.120                      Atherectomy
-	E04.100.814.529.124.120.125                      Atherectomy, Coronary
-	E04.100.814.529.500                      Angioscopy
-	E04.100.814.529.875                      Catheterization, Central Venous
-	E04.100.814.529.937                      Catheterization, Peripheral
-	E04.100.814.529.937.165                      Catheterization, Swan-Ganz
-	E04.100.814.529.968                      Percutaneous Coronary Intervention
-	E04.100.814.529.968.050                      Angioplasty, Balloon, Coronary
-	E04.100.814.529.968.060                      Atherectomy, Coronary
-	E04.100.814.603                      Limb Salvage
-	E04.100.814.750                      Peritoneovenous Shunt
-	E04.100.814.842                      Thrombectomy
-	E04.100.814.842.500                      Mechanical Thrombolysis
-	E04.100.814.868                      Vascular Grafting
-	E04.100.814.868.249                      Arteriovenous Shunt, Surgical
-	E04.100.814.868.374                      Blalock-Taussig Procedure
-	E04.100.814.868.500                      Blood Vessel Prosthesis Implantation
-	E04.100.814.868.625                      Cerebral Revascularization
-	E04.100.814.868.750                      Coronary Artery Bypass
-	E04.100.814.868.750.199                      Coronary Artery Bypass, Off-Pump
-	E04.100.814.868.750.400                      Internal Mammary-Coronary Artery Anastomosis
-	E04.100.814.868.875                      Heart Bypass, Right
-	E04.100.814.868.875.295                      Fontan Procedure
-	E04.100.814.868.937                      Portasystemic Shunt, Surgical
-	E04.100.814.868.937.790                      Portacaval Shunt, Surgical
-	E04.100.814.868.937.830                      Portasystemic Shunt, Transjugular Intrahepatic
-	E04.100.814.868.937.870                      Splenorenal Shunt, Surgical
-	E04.100.814.895                      Venous Cutdown
-	E04.157                      Curettage
-	E04.157.310                      Dilatation and Curettage
-	E04.157.310.970                      Vacuum Curettage

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	E04.166	Cytoreduction Surgical Procedures
-	E04.176	Debridement
-	E04.188	Decompression, Surgical
-	E04.188.200	Decompressive Craniectomy
-	E04.188.600	Microvascular Decompression Surgery
-	E04.190	Deep Brain Stimulation
-	E04.199	Device Removal
-	E04.210	Digestive System Surgical Procedures
-	E04.210.070	Anastomosis, Roux-en-Y
-	E04.210.078	Appendectomy
-	E04.210.120	Biliary Tract Surgical Procedures
-	E04.210.120.086	Biliopancreatic Diversion
-	E04.210.120.172	Cholecystectomy
-	E04.210.120.172.140	Cholecystectomy, Laparoscopic
-	E04.210.120.195	Cholecystostomy
-	E04.210.120.200	Choledochostomy
-	E04.210.120.775	Portoenterostomy, Hepatic
-	E04.210.120.850	Sphincterotomy, Endoscopic
-	E04.210.120.860	Sphincterotomy, Transhepatic
-	E04.210.169	Biliopancreatic Diversion
-	E04.210.219	Colectomy
-	E04.210.219.620	Proctocolectomy, Restorative
-	E04.210.240	Endoscopy, Digestive System
-	E04.210.240.160	Cholangiopancreatography, Endoscopic Retrograde
-	E04.210.240.250	Endoscopy, Gastrointestinal
New Heading	<b>E04.210.240.250.080</b>	<b>Balloon Enteroscopy</b>
New Tree	<a href="#">E04.210.240.250.080.500</a>	<a href="#">Double-Balloon Enteroscopy</a>
New Heading	<b>E04.210.240.250.080.750</b>	<b>Single-Balloon Enteroscopy</b>
-	E04.210.240.250.160	Colonoscopy
-	E04.210.240.250.160.800	Sigmoidoscopy
-	E04.210.240.250.200	Duodenoscopy
New Heading	<b>E04.210.240.250.230</b>	<b>Endoscopic Mucosal Resection</b>
-	E04.210.240.250.260	Esophagoscopy



## MeSH Tree Changes for 2017

Type	Tree - heading
-	E04.210.240.250.320                      Gastroscopy
-	E04.210.240.250.680                      Proctoscopy
-	E04.210.240.250.680.500                      Transanal Endoscopic Surgery
-	E04.210.240.250.680.500.500                      Transanal Endoscopic Microsurgery
-	E04.210.240.250.840                      Sphincterotomy, Endoscopic
-	E04.210.338                      Enterostomy
-	E04.210.338.175                      Cecostomy
-	E04.210.338.225                      Colostomy
-	E04.210.338.303                      Duodenostomy
-	E04.210.338.508                      Ileostomy
-	E04.210.338.523                      Jejunostomy
-	E04.210.346                      Esophagectomy
-	E04.210.355                      Esophagoplasty
-	E04.210.358                      Esophagostomy
-	E04.210.390                      Fundoplication
-	E04.210.419                      Gastrectomy
-	E04.210.457                      Gastroenterostomy
-	E04.210.457.430                      Gastric Bypass
-	E04.210.471                      Gastropexy
-	E04.210.485                      Gastroplasty
-	E04.210.496                      Gastrostomy
-	E04.210.526                      Hemorrhoidectomy
-	E04.210.556                      Hepatectomy
-	E04.210.626                      Jejunioileal Bypass
-	E04.210.650                      Liver Transplantation
-	E04.210.725                      Pancreas Transplantation
-	E04.210.752                      Pancreatectomy
-	E04.210.760                      Pancreaticoduodenectomy
-	E04.210.762                      Pancreaticojejunostomy
-	E04.210.790                      Peritoneovenous Shunt
-	E04.221                      Dissection
-	E04.221.580                      Microdissection
-	E04.237                      Drainage
-	E04.237.444                      Negative-Pressure Wound Therapy
-	E04.237.667                      Paracentesis
-	E04.237.890                      Suction

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	E04.249	Elective Surgical Procedures
-	E04.262	Electrosurgery
-	E04.270	Endocrine Surgical Procedures
-	E04.270.115	Adrenalectomy
-	E04.270.282	Castration
-	E04.270.282.679	Orchiectomy
-	E04.270.282.685	Ovariectomy
-	E04.270.532	Hypophysectomy
-	E04.270.532.490	Hypophysectomy, Chemical
-	E04.270.550	Islets of Langerhans Transplantation
-	E04.270.694	Parathyroidectomy
-	E04.270.856	Thyroidectomy
-	E04.292	Extracorporeal Circulation
-	E04.292.413	Cardiopulmonary Bypass
-	E04.292.425	Chemotherapy, Cancer, Regional Perfusion
-	E04.292.451	Extracorporeal Membrane Oxygenation
-	E04.292.465	Heart Bypass, Left
-	E04.292.471	Hemofiltration
-	E04.292.471.350	Hemodiafiltration
-	E04.292.510	Hemoperfusion
-	E04.292.762	Photopheresis
-	E04.292.975	Ultrafiltration
New Heading	<b>E04.321</b>	<b>Fasciotomy</b>
-	E04.350	Hemostasis, Surgical
-	E04.350.500	Argon Plasma Coagulation
-	E04.350.750	Light Coagulation
-	E04.350.750.410	Laser Coagulation
-	E04.406	Laparotomy
-	E04.426	Ligation
-	E04.446	Lymph Node Excision
-	E04.446.318	Neck Dissection
-	E04.446.819	Sentinel Lymph Node Biopsy
-	E04.466	Mastectomy
-	E04.466.678	Mastectomy, Radical
-	E04.466.678.354	Mastectomy, Extended Radical

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	E04.466.678.476	Mastectomy, Modified Radical
-	E04.466.701	Mastectomy, Segmental
-	E04.466.754	Mastectomy, Simple
-	E04.466.823	Mastectomy, Subcutaneous
New Heading	<b>E04.466.911</b>	<b>Prophylactic Mastectomy</b>
-	E04.480	Metastasectomy
-	E04.494	Microsurgery
-	E04.494.150	Cerebral Revascularization
-	E04.494.575	Mohs Surgery
-	E04.494.787	Transanal Endoscopic Microsurgery
-	E04.502	Minimally Invasive Surgical Procedures
-	E04.502.250	Endoscopy
-	E04.502.250.035	Angioscopy
-	E04.502.250.070	Arthroscopy
-	E04.502.250.070.500	Joint Capsule Release
-	E04.502.250.100	Bronchoscopy
-	E04.502.250.150	Colposcopy
-	E04.502.250.155	Conversion to Open Surgery
-	E04.502.250.160	Culdoscopy
-	E04.502.250.180	Cystoscopy
-	E04.502.250.250	Endoscopy, Digestive System
-	E04.502.250.250.160	Cholangiopancreatography, Endoscopic Retrograde
-	E04.502.250.250.250	Endoscopy, Gastrointestinal
New Heading	<b>E04.502.250.250.250.080</b>	<b>Balloon Enteroscopy</b>
New Tree	<a href="#">E04.502.250.250.250.080.250</a>	<a href="#">Double-Balloon Enteroscopy</a>
New Heading	<b>E04.502.250.250.250.080.500</b>	<b>Single-Balloon Enteroscopy</b>
-	E04.502.250.250.250.160	Colonoscopy
-	E04.502.250.250.250.160.800	Sigmoidoscopy
Old Tree	<del>E04.502.250.250.250.180</del>	<del>Double-Balloon Enteroscopy</del>
-	E04.502.250.250.250.200	Duodenoscopy
New Heading	<b>E04.502.250.250.250.230</b>	<b>Endoscopic Mucosal Resection</b>
-	E04.502.250.250.250.260	Esophagoscopy



## MeSH Tree Changes for 2017

Type	Tree - heading
-	E04.502.382.968.060                      Atherectomy, Coronary
-	E04.502.515                                      Magnetic Resonance Imaging, Interventional
-	E04.502.648                                      Morcellation
-	E04.502.780                                      Radiography, Interventional
-	E04.502.890                                      Ultrasonography, Interventional
-	E04.502.890.500                                  Endoscopic Ultrasound-Guided Fine Needle Aspiration
-	E04.506    Minor Surgical Procedures
-	E04.510    Monitoring, Intraoperative
-	E04.510.500    Intraoperative Neurophysiological Monitoring
-	E04.520    Obstetric Surgical Procedures
-	E04.520.050    Abortion, Induced
-	E04.520.050.050                                      Abortion, Eugenic
-	E04.520.050.055                                      Abortion, Legal
-	E04.520.050.060                                      Abortion, Therapeutic
-	E04.520.050.600    Pregnancy Reduction, Multifetal
-	E04.520.100    Cerclage, Cervical
-	E04.520.150    Colposcopy
-	E04.520.155    Colpotomy
-	E04.520.160    Culdoscopy
-	E04.520.252    Delivery, Obstetric
-	E04.520.252.500    Cesarean Section
-	E04.520.252.500.150                                      Cesarean Section, Repeat
-	E04.520.252.750    Episiotomy
-	E04.520.252.875    Extraction, Obstetrical
-	E04.520.252.875.970    Vacuum Extraction, Obstetrical
-	E04.520.252.968    Labor, Induced
-	E04.520.252.992    Vaginal Birth after Cesarean
-	E04.520.252.996    Version, Fetal
-	E04.520.280    Fetoscopy
-	E04.520.360    Hysteroscopy
-	E04.520.365    Hysterotomy
-	E04.525    Neurosurgical Procedures
-	E04.525.044    Anterior Temporal Lobectomy
-	E04.525.090    Brain Tissue Transplantation
-	E04.525.160    Cerebral Decortication
-	E04.525.160.500    Hemispherectomy

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E04.525.170 Cerebrospinal Fluid Shunts
-	E04.525.170.850 Ventriculoperitoneal Shunt
-	E04.525.170.860 Ventriculostomy
-	E04.525.190 Craniotomy
-	E04.525.190.200 Decompressive Craniectomy
-	E04.525.190.840 Trephining
-	E04.525.210 Denervation
-	E04.525.210.080 Autonomic Denervation
-	E04.525.210.080.600 Parasympathectomy
-	E04.525.210.080.600.850 Vagotomy
-	E04.525.210.080.600.850.850 Vagotomy, Proximal Gastric
-	E04.525.210.080.600.850.860 Vagotomy, Truncal
-	E04.525.210.080.800 Sympathectomy
-	E04.525.210.080.800.380 Ganglionectomy
-	E04.525.210.080.800.800 Sympathectomy, Chemical
-	E04.525.210.145 Axotomy
-	E04.525.210.210 Cordotomy
-	E04.525.210.380 Ganglionectomy
-	E04.525.210.500 Muscle Denervation
-	E04.525.210.550 Nerve Block
-	E04.525.210.550.500 Autonomic Nerve Block
-	E04.525.210.550.500.500 Sphenopalatine Ganglion Block
-	E04.525.210.560 Nerve Crush
-	E04.525.210.700 Rhizotomy
-	E04.525.210.850 Vagotomy
-	E04.525.210.850.850 Vagotomy, Proximal Gastric
-	E04.525.210.850.860 Vagotomy, Truncal
-	E04.525.305 Foraminotomy
-	E04.525.400 Hypophysectomy
-	E04.525.450 Laminectomy
-	E04.525.475 Laminoplasty
-	E04.525.500 Microvascular Decompression Surgery
-	E04.525.550 Nerve Transfer
-	E04.525.562 Neuroendoscopy
-	E04.525.575 Pallidotomy
-	E04.525.600 Psychosurgery

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E04.525.770                      Split-Brain Procedure
-	E04.525.800                      Stereotaxic Techniques
-	E04.525.800.324                      Neuronavigation
-	E04.525.800.650                      Radiosurgery
-	E04.540                      Ophthalmologic Surgical Procedures
-	E04.540.104                      Blepharoplasty
-	E04.540.255                      Dacryocystorhinostomy
-	E04.540.429                      Eye Enucleation
-	E04.540.431                      Eye Evisceration
-	E04.540.450                      Filtering Surgery
-	E04.540.450.600                      Sclerostomy
-	E04.540.450.700                      Trabeculectomy
-	E04.540.480                      Iridectomy
-	E04.540.630                      Light Coagulation
-	E04.540.630.410                      Laser Coagulation
-	E04.540.760                      Orbit Evisceration
-	E04.540.825                      Refractive Surgical Procedures
-	E04.540.825.249                      Cataract Extraction
-	E04.540.825.249.352                      Capsulorhexis
-	E04.540.825.249.704                      Phacoemulsification
-	E04.540.825.374                      Corneal Transplantation
-	E04.540.825.374.112                      Descemet Stripping Endothelial Keratoplasty
-	E04.540.825.374.225                      Epikeratophakia
-	E04.540.825.374.350                      Keratoplasty, Penetrating
-	E04.540.825.437                      Corneal Surgery, Laser
-	E04.540.825.437.249                      Keratectomy, Subepithelial, Laser-Assisted
-	E04.540.825.437.374                      Keratomileusis, Laser In Situ
-	E04.540.825.437.500                      Photorefractive Keratectomy
-	E04.540.825.550                      Keratotomy, Radial
-	E04.540.825.600                      Lens Implantation, Intraocular
-	E04.540.825.784                      Posterior Capsulotomy
-	E04.540.825.968                      Scleroplasty
-	E04.540.890                      Scleral Buckling
-	E04.540.960                      Vitrectomy
-	E04.540.980                      Vitreoretinal Surgery
-	E04.545                      Oral Surgical Procedures

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E04.545.100 Apicoectomy
-	E04.545.350 Gingivectomy
-	E04.545.355 Gingivoplasty
-	E04.545.380 Glossectomy
-	E04.545.440 Jaw Fixation Techniques
-	E04.545.500 Mandibular Advancement
-	E04.545.510 Maxillofacial Prosthesis Implantation
-	E04.545.510.500 Mandibular Prosthesis Implantation
-	E04.545.550 Oral Surgical Procedures, Preprosthetic
-	E04.545.550.100 Alveolar Ridge Augmentation
-	E04.545.550.110 Alveolectomy
-	E04.545.550.120 Alveoloplasty
-	E04.545.550.280 Dental Implantation
-	E04.545.550.280.280 Dental Implantation, Endosseous
-	E04.545.550.280.280.150 Blade Implantation
-	E04.545.550.280.280.280 Dental Implantation, Endosseous, Endodontic
-	E04.545.550.280.280.350 Immediate Dental Implant Loading
-	E04.545.550.280.290 Dental Implantation, Subperiosteal
-	E04.545.550.800 Vestibuloplasty
-	E04.545.562 Orthognathic Surgical Procedures
-	E04.545.562.500 Alveolar Bone Grafting
-	E04.545.575 Osteotomy, Le Fort
-	E04.545.637 Osteotomy, Sagittal Split Ramus
-	E04.545.668 Sinus Floor Augmentation
-	E04.545.700 Tooth Extraction
-	E04.545.700.680 Serial Extraction
-	E04.545.710 Tooth Replantation
-	E04.555 Orthopedic Procedures
-	E04.555.039 Acetabuloplasty
-	E04.555.080 Amputation
-	E04.555.080.250 Disarticulation
-	E04.555.080.380 Hemipelvectomy
-	E04.555.100 Arthrodesis
-	E04.555.100.700 Spinal Fusion
-	E04.555.110 Arthroplasty
-	E04.555.110.026 Anterior Cruciate Ligament Reconstruction



## MeSH Tree Changes for 2017

Type	Tree - heading	
-	E04.555.110.026.500	Bone-Patellar Tendon-Bone Grafting
-	E04.555.110.110	Arthroplasty, Replacement
-	E04.555.110.110.010	Arthroplasty, Replacement, Ankle
-	E04.555.110.110.045	Arthroplasty, Replacement, Elbow
-	E04.555.110.110.054	Arthroplasty, Replacement, Finger
-	E04.555.110.110.110	Arthroplasty, Replacement, Hip
-	E04.555.110.110.115	Arthroplasty, Replacement, Knee
New Heading	<b>E04.555.110.110.299</b>	<b>Arthroplasty, Replacement, Shoulder</b>
-	E04.555.110.110.482	Hemiarthroplasty
-	E04.555.110.110.850	Total Disc Replacement
-	E04.555.110.115	Arthroplasty, Subchondral
New Heading	<b>E04.555.110.557</b>	<b>Posterior Cruciate Ligament Reconstruction</b>
-	E04.555.113	Arthroscopy
-	E04.555.113.500	Joint Capsule Release
-	E04.555.120	Bone Lengthening
-	E04.555.120.380	Ilizarov Technique
-	E04.555.120.690	Osteogenesis, Distraction
-	E04.555.130	Bone Transplantation
-	E04.555.130.550	Sinus Floor Augmentation
-	E04.555.165	Cementoplasty
-	E04.555.165.500	Vertebroplasty
-	E04.555.165.500.500	Kyphoplasty
-	E04.555.200	Discectomy
-	E04.555.200.200	Discectomy, Percutaneous
-	E04.555.300	Fracture Fixation
New Heading	<b>E04.555.300.150</b>	<b>Closed Fracture Reduction</b>
-	E04.555.300.300	Fracture Fixation, Internal
-	E04.555.300.300.300	Fracture Fixation, Intramedullary
-	E04.555.300.380	Ilizarov Technique
New Heading	<b>E04.555.300.690</b>	<b>Open Fracture Reduction</b>
-	E04.555.400	Limb Salvage
-	E04.555.580	Osteotomy
-	E04.555.580.289	Orthognathic Surgical Procedures

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E04.555.580.289.500 Alveolar Bone Grafting
-	E04.555.580.580 Osteotomy, Le Fort
-	E04.555.580.790 Osteotomy, Sagittal Split Ramus
-	E04.555.700 Tendon Transfer
-	E04.555.710 Tenodesis
-	E04.555.712 Tenotomy
-	E04.555.720 Traction
New Heading	<b>E04.555.860 Ulnar Collateral Ligament Reconstruction</b>
-	E04.579 Ostomy
-	E04.579.240 Cystostomy
-	E04.579.255 Dacryocystorhinostomy
-	E04.579.338 Enterostomy
-	E04.579.338.175 Cecostomy
-	E04.579.338.225 Colostomy
-	E04.579.338.303 Duodenostomy
-	E04.579.338.508 Ileostomy
-	E04.579.338.523 Jejunostomy
-	E04.579.358 Esophagostomy
-	E04.579.408 Gastrostomy
-	E04.579.592 Middle Ear Ventilation
-	E04.579.642 Nephrostomy, Percutaneous
-	E04.579.740 Pharyngostomy
-	E04.579.895 Sclerostomy
-	E04.579.918 Thoracostomy
-	E04.579.935 Tracheostomy
-	E04.579.947 Ureterostomy
-	E04.580 Otorhinolaryngologic Surgical Procedures
-	E04.580.068 Adenoidectomy
-	E04.580.369 Laryngectomy
-	E04.580.371 Laryngoplasty
-	E04.580.373 Laryngoscopy
-	E04.580.392 Nasal Surgical Procedures
-	E04.580.392.500 Rhinoplasty
-	E04.580.411 Neck Dissection
-	E04.580.450 Otologic Surgical Procedures

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	E04.580.450.109	Auditory Brain Stem Implantation
-	E04.580.450.220	Cochlear Implantation
-	E04.580.450.230	Endolymphatic Shunt
-	E04.580.450.277	Fenestration, Labyrinth
-	E04.580.450.472	Middle Ear Ventilation
-	E04.580.450.505	Myringoplasty
-	E04.580.450.600	Ossicular Replacement
-	E04.580.450.731	Stapes Surgery
-	E04.580.450.731.681	Stapes Mobilization
-	E04.580.450.802	Transtympanic Micropressure Treatment
-	E04.580.450.873	Tympanoplasty
-	E04.580.515	Pharyngectomy
-	E04.580.540	Pharyngostomy
-	E04.580.848	Tonsillectomy
-	E04.580.900	Tracheostomy
-	E04.580.907	Tracheotomy
-	E04.584	Pelvic Exenteration
-	E04.604	Perioperative Care
-	E04.604.249	Intraoperative Care
-	E04.604.500	Postoperative Care
-	E04.604.750	Preoperative Care
-	E04.604.750.619	Preanesthetic Medication
-	E04.614	Perioperative Period
-	E04.614.374	Intraoperative Period
-	E04.614.374.500	Operative Time
-	E04.614.750	Postoperative Period
-	E04.614.750.055	Anesthesia Recovery Period
-	E04.614.937	Preoperative Period
-	E04.625	Prophylactic Surgical Procedures
New Heading	<b>E04.625.500</b>	<b>Prophylactic Mastectomy</b>
-	E04.650	Prosthesis Implantation
-	E04.650.110	Arthroplasty, Replacement
-	E04.650.110.010	Arthroplasty, Replacement, Ankle
-	E04.650.110.045	Arthroplasty, Replacement, Elbow
-	E04.650.110.054	Arthroplasty, Replacement, Finger

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	E04.650.110.110	Arthroplasty, Replacement, Hip
-	E04.650.110.115	Arthroplasty, Replacement, Knee
New Heading	<b>E04.650.110.299</b>	<b>Arthroplasty, Replacement, Shoulder</b>
-	E04.650.110.482	Hemiarthroplasty
-	E04.650.110.850	Total Disc Replacement
-	E04.650.155	Auditory Brain Stem Implantation
-	E04.650.200	Blood Vessel Prosthesis Implantation
-	E04.650.210	Breast Implantation
-	E04.650.220	Cochlear Implantation
-	E04.650.230	Dental Implantation
-	E04.650.230.500	Dental Implantation, Endosseous
-	E04.650.230.500.150	Blade Implantation
-	E04.650.230.500.280	Dental Implantation, Endosseous, Endodontic
-	E04.650.230.500.350	Immediate Dental Implant Loading
-	E04.650.230.750	Dental Implantation, Subperiosteal
-	E04.650.410	Heart Valve Prosthesis Implantation
-	E04.650.410.500	Transcatheter Aortic Valve Replacement
-	E04.650.510	Maxillofacial Prosthesis Implantation
-	E04.650.510.500	Mandibular Prosthesis Implantation
-	E04.650.600	Ossicular Replacement
-	E04.650.620	Penile Implantation
-	E04.665	Punctures
-	E04.665.100	Biopsy, Needle
-	E04.665.100.500	Biopsy, Fine-Needle
-	E04.665.100.750	Biopsy, Large-Core Needle
-	E04.665.150	Blood Specimen Collection
-	E04.665.150.150	Cordocentesis
-	E04.665.150.600	Petrosal Sinus Sampling
-	E04.665.150.625	Phlebotomy
-	E04.665.600	Paracentesis
-	E04.665.600.309	Amniocentesis
-	E04.665.600.465	Arthrocentesis
-	E04.665.600.543	Cordocentesis
-	E04.665.600.620	Pericardiocentesis
-	E04.665.600.810	Thoracentesis

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	E04.665.600.905	Tympanocentesis
-	E04.665.700	Spinal Puncture
-	E04.680	Reconstructive Surgical Procedures
-	E04.680.025	Abdominoplasty
-	E04.680.038	Acetabuloplasty
-	E04.680.101	Arthroplasty
-	E04.680.101.026	Anterior Cruciate Ligament Reconstruction
-	E04.680.101.026.500	Bone-Patellar Tendon-Bone Grafting
-	E04.680.101.110	Arthroplasty, Replacement
-	E04.680.101.110.010	Arthroplasty, Replacement, Ankle
-	E04.680.101.110.045	Arthroplasty, Replacement, Elbow
-	E04.680.101.110.054	Arthroplasty, Replacement, Finger
-	E04.680.101.110.110	Arthroplasty, Replacement, Hip
-	E04.680.101.110.115	Arthroplasty, Replacement, Knee
New Heading	<b>E04.680.101.110.299</b>	<b>Arthroplasty, Replacement, Shoulder</b>
-	E04.680.101.110.482	Hemiarthroplasty
-	E04.680.101.110.850	Total Disc Replacement
-	E04.680.101.115	Arthroplasty, Subchondral
New Heading	<b>E04.680.101.557</b>	<b>Posterior Cruciate Ligament Reconstruction</b>
-	E04.680.152	Cervicoplasty
-	E04.680.275	Dermatologic Surgical Procedures
-	E04.680.275.090	Blepharoplasty
-	E04.680.275.200	Chemexfoliation
-	E04.680.275.250	Dermabrasion
-	E04.680.275.580	Mohs Surgery
-	E04.680.275.700	Rhytidoplasty
-	E04.680.275.850	Skin Transplantation
-	E04.680.300	Guided Tissue Regeneration
-	E04.680.300.500	Guided Tissue Regeneration, Periodontal
-	E04.680.325	Herniorrhaphy
-	E04.680.350	Limb Salvage
-	E04.680.450	Lipectomy
-	E04.680.500	Mammoplasty
-	E04.680.500.210	Breast Implantation

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	E04.680.650	Rhinoplasty
-	E04.680.700	Scleroplasty
-	E04.680.750	Sex Reassignment Surgery
-	E04.680.800	Tissue Expansion
-	E04.680.800.500	Nerve Expansion
New Heading	<b>E04.680.900</b>	<b>Ulnar Collateral Ligament Reconstruction</b>
-	E04.690	Reoperation
-	E04.708	Second-Look Surgery
-	E04.726	Splenectomy
-	E04.749	Surgery, Computer-Assisted
-	E04.749.500	Robotic Surgical Procedures
-	E04.910	Symphysiotomy
-	E04.928	Thoracic Surgical Procedures
-	E04.928.220	Cardiac Surgical Procedures
-	E04.928.220.055	Arterial Switch Operation
-	E04.928.220.109	Cardiac Valve Annuloplasty
-	E04.928.220.109.500	Mitral Valve Annuloplasty
-	E04.928.220.220	Cardiomyoplasty
-	E04.928.220.360	Heart Arrest, Induced
-	E04.928.220.360.500	Circulatory Arrest, Deep Hypothermia Induced
-	E04.928.220.370	Heart Bypass, Right
-	E04.928.220.370.295	Fontan Procedure
-	E04.928.220.380	Heart Massage
-	E04.928.220.390	Heart Transplantation
-	E04.928.220.390.450	Heart-Lung Transplantation
-	E04.928.220.410	Heart Valve Prosthesis Implantation
-	E04.928.220.410.500	Transcatheter Aortic Valve Replacement
-	E04.928.220.520	Myocardial Revascularization
-	E04.928.220.520.100	Angioplasty, Balloon, Coronary
-	E04.928.220.520.110	Atherectomy, Coronary
-	E04.928.220.520.220	Coronary Artery Bypass
-	E04.928.220.520.220.189	Coronary Artery Bypass, Off-Pump
-	E04.928.220.520.220.380	Internal Mammary-Coronary Artery Anastomosis
-	E04.928.220.520.610	Transmyocardial Laser Revascularization
-	E04.928.220.560	Norwood Procedures

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E04.928.220.560.500                      Fontan Procedure
-	E04.928.220.600                      Pericardial Window Techniques
-	E04.928.220.605                      Pericardiectomy
-	E04.928.220.620                      Pericardiocentesis
-	E04.928.490                              Mediastinoscopy
-	E04.928.600                              Pulmonary Surgical Procedures
-	E04.928.600.080                      Bronchoscopy
-	E04.928.600.220                      Collapse Therapy
-	E04.928.600.220.610                      Pneumonolysis
-	E04.928.600.220.620                      Pneumothorax, Artificial
-	E04.928.600.495                      Lung Transplantation
-	E04.928.600.495.450                      Heart-Lung Transplantation
-	E04.928.600.600                      Pneumonectomy
-	E04.928.710                              Sternotomy
-	E04.928.750                              Thoracoplasty
-	E04.928.752                              Thoracoscopy
-	E04.928.752.830                      Thoracic Surgery, Video-Assisted
-	E04.928.755                              Thoracostomy
-	E04.928.760                              Thoracotomy
-	E04.928.770                              Thymectomy
-	E04.928.780                              Tracheostomy
-	E04.928.790                              Tracheotomy
-	E04.936                                      Transplantation
-	E04.936.225                              Cell Transplantation
-	E04.936.225.375                      Islets of Langerhans Transplantation
-	E04.936.225.687                      Stem Cell Transplantation
-	E04.936.225.687.312                      Cord Blood Stem Cell Transplantation
-	E04.936.225.687.500                      Hematopoietic Stem Cell Transplantation
New Tree	<a href="#">E04.936.225.687.500.500</a> <a href="#">Peripheral Blood Stem Cell Transplantation</a>
-	E04.936.225.687.625                      Mesenchymal Stem Cell Transplantation
Old Tree	<del><a href="#">E04.936.225.687.750</a></del> <del><a href="#">Peripheral Blood Stem Cell Transplantation</a></del>
-	E04.936.337                              Cold Ischemia
-	E04.936.450                              Organ Transplantation
-	E04.936.450.475                      Heart Transplantation
-	E04.936.450.475.450                      Heart-Lung Transplantation

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E04.936.450.485 Kidney Transplantation
-	E04.936.450.490 Liver Transplantation
-	E04.936.450.495 Lung Transplantation
-	E04.936.450.495.450 Heart-Lung Transplantation
-	E04.936.450.650 Pancreas Transplantation
-	E04.936.450.825 Vascularized Composite Allotransplantation
-	E04.936.450.825.249 Facial Transplantation
-	E04.936.450.825.500 Hand Transplantation
-	E04.936.494 Replantation
-	E04.936.494.711 Tooth Replantation
-	E04.936.537 Tissue and Organ Harvesting
-	E04.936.537.500 Donor Selection
-	E04.936.537.562 Fertility Preservation
-	E04.936.537.625 Oocyte Retrieval
-	E04.936.537.687 Operative Blood Salvage
-	E04.936.537.750 Sperm Retrieval
-	E04.936.580 Tissue Transplantation
-	E04.936.580.040 Bone Marrow Transplantation
-	E04.936.580.052 Bone Transplantation
-	E04.936.580.065 Bone-Patellar Tendon-Bone Grafting
-	E04.936.580.090 Brain Tissue Transplantation
-	E04.936.580.225 Corneal Transplantation
-	E04.936.580.225.225 Epikeratophakia
-	E04.936.580.225.350 Keratoplasty, Penetrating
-	E04.936.580.262 Descemet Stripping Endothelial Keratoplasty
-	E04.936.580.300 Fetal Tissue Transplantation
-	E04.936.580.490 Liver Transplantation
-	E04.936.580.700 Skin Transplantation
-	E04.936.664 Transplantation, Autologous
-	E04.936.764 Transplantation, Heterologous
-	E04.936.800 Transplantation, Heterotopic
-	E04.936.864 Transplantation, Homologous
-	E04.936.864.700 Transplantation, Isogeneic
-	E04.936.932 Warm Ischemia
-	E04.943 Ultrasonic Surgical Procedures
-	E04.943.500 Lithotripsy



## MeSH Tree Changes for 2017

Type	Tree - heading
-	E04.943.500.500 Lithotripsy, Laser
-	E04.943.875 Phacoemulsification
-	E04.943.937 Piezosurgery
-	E04.950 Urogenital Surgical Procedures
-	E04.950.165 Castration
-	E04.950.165.679 Orchiectomy
-	E04.950.165.685 Ovariectomy
-	E04.950.300 Gynecologic Surgical Procedures
-	E04.950.300.200 Circumcision, Female
-	E04.950.300.210 Colposcopy
-	E04.950.300.220 Colpotomy
-	E04.950.300.259 Culdoscopy
-	E04.950.300.299 Dilatation and Curettage
-	E04.950.300.299.970 Vacuum Curettage
-	E04.950.300.320 Endometrial Ablation Techniques
-	E04.950.300.399 Hysterectomy
-	E04.950.300.399.380 Hysterectomy, Vaginal
-	E04.950.300.399.690 Trachelectomy
-	E04.950.300.539 Hysteroscopy
-	E04.950.300.680 Ovariectomy
-	E04.950.300.715 Salpingectomy
-	E04.950.300.750 Salpingostomy
-	E04.950.300.766 Sterilization, Tubal
-	E04.950.300.883 Uterine Artery Embolization
-	E04.950.300.941 Uterine Myomectomy
-	E04.950.449 Sex Reassignment Surgery
-	E04.950.599 Sterilization, Reproductive
-	E04.950.599.450 Sterilization, Involuntary
-	E04.950.599.500 Sterilization Reversal
-	E04.950.599.500.800 Vasovasostomy
-	E04.950.599.683 Sterilization, Tubal
-	E04.950.599.900 Vasectomy
-	E04.950.774 Urologic Surgical Procedures
-	E04.950.774.150 Cystectomy
-	E04.950.774.155 Cystoscopy
-	E04.950.774.162 Cystotomy

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	E04.950.774.400	Kidney Transplantation
-	E04.950.774.435	Nephrectomy
-	E04.950.774.840	Ureteroscopy
-	E04.950.774.852	Urinary Diversion
-	E04.950.774.852.240	Cystostomy
-	E04.950.774.852.642	Nephrostomy, Percutaneous
-	E04.950.774.852.947	Ureterostomy
-	E04.950.774.860	Urologic Surgical Procedures, Male
-	E04.950.774.860.226	Circumcision, Male
-	E04.950.774.860.618	Orchiectomy
-	E04.950.774.860.619	Orchiopexy
-	E04.950.774.860.620	Penile Implantation
-	E04.950.774.860.625	Prostatectomy
-	E04.950.774.860.625.750	Transurethral Resection of Prostate
-	E04.950.774.860.625.875	Ultrasound, High-Intensity Focused, Transrectal
-	E04.950.774.860.856	Vasectomy
-	E04.950.774.860.956	Vasovasostomy
-	E04.987	Wound Closure Techniques
-	E04.987.100	Abdominal Wound Closure Techniques
-	E04.987.550	Negative-Pressure Wound Therapy
-	E04.987.775	Suture Techniques
-	E04.987.775.800	Surgical Stapling
-	E04.987.775.900	Tenodesis
New Heading	<b>E04.987.887</b>	<b>Sutureless Surgical Procedures</b>
-	E05	Investigative Techniques
-	E05.003	Accelerometry
-	E05.003.500	Actigraphy
-	E05.008	Airway Extubation
-	E05.017	Animal Experimentation
-	E05.017.080	Animal Use Alternatives
-	E05.017.080.100	Animal Testing Alternatives
-	E05.017.449	Rotarod Performance Test
-	E05.017.900	Vivisection
-	E05.035	Animal Identification Systems
-	E05.041	Anthropometry

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E05.041.124                      Body Weights and Measures
-	E05.041.124.125                  Body Mass Index
-	E05.041.124.160                  Body Size
-	E05.041.124.160.500              Body Height
-	E05.041.124.160.500.500        Crown-Rump Length
-	E05.041.124.160.750              Body Weight
-	E05.041.124.160.750.149        Birth Weight
-	E05.041.124.160.750.300        Fetal Weight
-	E05.041.124.160.750.650        Ideal Body Weight
-	E05.041.124.160.812              Sagittal Abdominal Diameter
-	E05.041.124.160.875              Waist Circumference
-	E05.041.124.160.875.500        Lipid Accumulation Product
-	E05.041.124.160.937              Waist-Height Ratio
-	E05.041.124.231                  Body Surface Area
-	E05.041.124.715                  Organ Size
-	E05.041.124.803                  Skinfold Thickness
-	E05.041.124.892                  Tumor Burden
-	E05.041.124.946                  Waist-Hip Ratio
-	E05.041.250                      Cephalometry
-	E05.041.650                      Odontometry
-	E05.041.650.500                  Age Determination by Teeth
-	E05.041.690                      Pelvimetry
-	E05.047                          Artifacts
-	E05.059                          Autoanalysis
-	E05.064                          Automation, Laboratory
-	E05.070                          Autopsy
-	E05.091                          Biological Assay
-	E05.091.570                      Limulus Test
-	E05.104                          Biomedical Enhancement
-	E05.104.500                      Genetic Enhancement
-	E05.111                          Bioprinting
-	E05.114                          Bioprospecting
-	E05.118                          Bone Demineralization Technique
-	E05.157                          Catheterization
-	E05.157.016                      Angioplasty
-	E05.157.016.060                  Angioplasty, Balloon

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E05.157.016.060.100      Angioplasty, Balloon, Coronary
-	E05.157.016.060.105      Angioplasty, Balloon, Laser-Assisted
-	E05.157.016.075            Angioplasty, Laser
-	E05.157.016.075.080      Angioplasty, Balloon, Laser-Assisted
-	E05.157.016.120            Atherectomy
-	E05.157.016.120.125      Atherectomy, Coronary
-	E05.157.032                Balloon Embolectomy
-	E05.157.063                Balloon Occlusion
-	E05.157.063.500          Uterine Balloon Tamponade
-	E05.157.125                Balloon Valvuloplasty
-	E05.157.250                Cardiac Catheterization
-	E05.157.250.165          Catheterization, Swan-Ganz
-	E05.157.313                Catheterization, Central Venous
-	E05.157.375                Catheterization, Peripheral
-	E05.157.375.165          Catheterization, Swan-Ganz
-	E05.157.500                Urinary Catheterization
-	E05.157.500.500          Intermittent Urethral Catheterization
-	E05.170                    Cementation
-	E05.181                    Centrifugation
-	E05.181.724                Ultracentrifugation
-	E05.181.724.336          Centrifugation, Density Gradient
-	E05.181.724.336.253      Centrifugation, Isopycnic
-	E05.181.724.336.336      Centrifugation, Zonal
-	E05.196                    Chemistry Techniques, Analytical
-	E05.196.039                Activation Analysis
-	E05.196.039.564          Neutron Activation Analysis
-	E05.196.059                Analytic Sample Preparation Methods
-	E05.196.080                Binding, Competitive
-	E05.196.085                Biuret Reaction
-	E05.196.131                Calorimetry
-	E05.196.131.310          Calorimetry, Differential Scanning
-	E05.196.131.655          Calorimetry, Indirect
-	E05.196.150                Chemical Precipitation
-	E05.196.150.347          Flocculation
-	E05.196.150.347.500      Flocculation Tests
-	E05.196.150.519          Fractional Precipitation

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E05.196.150.639 Immunoprecipitation
-	E05.196.150.639.500 Precipitin Tests
-	E05.196.155 Chemical Fractionation
-	E05.196.155.249 Distillation
-	E05.196.155.500 Fractionation, Field Flow
-	E05.196.155.650 Liquid-Liquid Extraction
-	E05.196.155.650.500 Liquid Phase Microextraction
-	E05.196.155.800 Solid Phase Extraction
-	E05.196.155.800.500 Solid Phase Microextraction
-	E05.196.181 Chromatography
-	E05.196.181.349 Chromatography, Gas
-	E05.196.181.349.390 Flame Ionization
-	E05.196.181.349.500 Gas Chromatography-Mass Spectrometry
-	E05.196.181.400 Chromatography, Liquid
-	E05.196.181.400.150 Capillary Electrochromatography
-	E05.196.181.400.170 Chromatography, Affinity
-	E05.196.181.400.170.500 Immunochromatography
-	E05.196.181.400.250 Chromatography, Gel
-	E05.196.181.400.250.200 Chromatography, Agarose
-	E05.196.181.400.300 Chromatography, High Pressure Liquid
-	E05.196.181.400.383 Chromatography, Ion Exchange
-	E05.196.181.400.383.349 Chromatography, DEAE-Cellulose
-	E05.196.181.400.454 Chromatography, Paper
-	E05.196.181.400.454.655 Nucleotide Mapping
-	E05.196.181.400.454.720 Peptide Mapping
-	E05.196.181.400.495 Chromatography, Reverse-Phase
-	E05.196.181.400.537 Chromatography, Thin Layer
-	E05.196.181.400.555 Countercurrent Distribution
-	E05.196.181.500 Chromatography, Micellar Electrokinetic Capillary
-	E05.196.181.750 Chromatography, Supercritical Fluid
-	E05.196.278 Countercurrent Distribution
-	E05.196.300 Crystallization
-	E05.196.309 Crystallography
-	E05.196.309.555 Neutron Diffraction
-	E05.196.309.711 Powder Diffraction
-	E05.196.309.742 X-Ray Diffraction

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E05.196.309.742.225 Crystallography, X-Ray
-	E05.196.335 Desiccation
-	E05.196.344 Deuterium Exchange Measurement
-	E05.196.353 Dialysis
-	E05.196.353.500 Microdialysis
-	E05.196.370 Differential Thermal Analysis
-	E05.196.370.310 Calorimetry, Differential Scanning
-	E05.196.385 Electroosmosis
-	E05.196.401 Electrophoresis
-	E05.196.401.076 Blood Protein Electrophoresis
-	E05.196.401.095 Blotting, Northern
-	E05.196.401.114 Blotting, Southern
-	E05.196.401.133 Blotting, Southwestern
-	E05.196.401.143 Blotting, Western
-	E05.196.401.143.500 Blotting, Far-Western
-	E05.196.401.153 Electrophoresis, Agar Gel
-	E05.196.401.153.150 Comet Assay
-	E05.196.401.190 Electrophoresis, Capillary
-	E05.196.401.190.249 Capillary Electrochromatography
-	E05.196.401.190.500 Electrophoresis, Microchip
-	E05.196.401.200 Electrophoresis, Cellulose Acetate
-	E05.196.401.220 Electrophoresis, Gel, Pulsed-Field
-	E05.196.401.250 Electrophoresis, Gel, Two-Dimensional
-	E05.196.401.250.500 Two-Dimensional Difference Gel Electrophoresis
-	E05.196.401.319 Electrophoresis, Paper
-	E05.196.401.319.670 Nucleotide Mapping
-	E05.196.401.319.720 Peptide Mapping
-	E05.196.401.402 Electrophoresis, Polyacrylamide Gel
-	E05.196.401.402.117 Denaturing Gradient Gel Electrophoresis
-	E05.196.401.402.236 Electrophoresis, Disc
-	E05.196.401.402.625 Native Polyacrylamide Gel Electrophoresis
-	E05.196.401.485 Electrophoresis, Starch Gel
-	E05.196.401.500 Electrophoretic Mobility Shift Assay
-	E05.196.401.568 Immunoelectrophoresis
-	E05.196.401.568.250 Counterimmunoelectrophoresis
-	E05.196.401.568.520 Immunoelectrophoresis, Two-Dimensional

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E05.196.401.663 Isoelectric Focusing
-	E05.196.401.831 Isotachopheresis
-	E05.196.427 Enzyme Assays
-	E05.196.427.200 Clinical Enzyme Tests
-	E05.196.454 Filtration
-	E05.196.454.201 Air Filters
-	E05.196.454.403 Micropore Filters
-	E05.196.454.807 Ultrafiltration
-	E05.196.460 Flow Injection Analysis
-	E05.196.500 Glucose Clamp Technique
-	E05.196.566 Mass Spectrometry
-	E05.196.566.500 Gas Chromatography-Mass Spectrometry
-	E05.196.566.600 Spectrometry, Mass, Electrospray Ionization
-	E05.196.566.750 Spectrometry, Mass, Fast Atom Bombardment
-	E05.196.566.755 Spectrometry, Mass, Matrix-Assisted Laser Desorption-Ionization
-	E05.196.566.760 Spectrometry, Mass, Secondary Ion
-	E05.196.566.880 Tandem Mass Spectrometry
-	E05.196.620 Microchemistry
-	E05.196.655 Molecular Imprinting
-	E05.196.680 Nucleotide Mapping
-	E05.196.690 Osmometry
-	E05.196.700 Peptide Mapping
-	E05.196.700.690 Protein Footprinting
-	E05.196.712 Photometry
-	E05.196.712.224 Densitometry
-	E05.196.712.224.187 Absorptiometry, Photon
-	E05.196.712.516 Luminescent Measurements
-	E05.196.712.516.099 Bioluminescence Resonance Energy Transfer Techniques
-	E05.196.712.516.600 Fluorometry
-	E05.196.712.516.600.240 Cytophotometry
-	E05.196.712.516.600.240.350 Flow Cytometry
-	E05.196.712.516.600.240.400 Image Cytometry
-	E05.196.712.516.600.240.400.500 Laser Scanning Cytometry
-	E05.196.712.516.600.390 Fluorescence Polarization

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E05.196.712.516.600.390.350 Fluorescence Polarization Immunoassay
-	E05.196.712.516.600.393 Fluorescence Recovery After Photobleaching
-	E05.196.712.516.600.410 Fluorophotometry
-	E05.196.712.516.600.676 Spectrometry, Fluorescence
-	E05.196.712.516.600.676.500 Fluorescence Resonance Energy Transfer
-	E05.196.712.650 Nephelometry and Turbidimetry
-	E05.196.712.726 Spectrophotometry
-	E05.196.712.726.300 Microspectrophotometry
-	E05.196.712.726.551 Spectrophotometry, Atomic
-	E05.196.712.726.676 Spectrophotometry, Infrared
-	E05.196.712.726.676.700 Spectroscopy, Fourier Transform Infrared
-	E05.196.712.726.802 Spectrophotometry, Ultraviolet
-	E05.196.749 Polarography
-	E05.196.778 Quartz Crystal Microbalance Techniques
-	E05.196.808 Refractometry
-	E05.196.822 Scattering, Radiation
-	E05.196.822.325 Dynamic Light Scattering
-	E05.196.822.650 Neutron Diffraction
-	E05.196.822.830 Scattering, Small Angle
-	E05.196.822.860 Spectrum Analysis, Raman
-	E05.196.822.950 X-Ray Diffraction
-	E05.196.867 Spectrum Analysis
-	E05.196.867.151 Circular Dichroism
-	E05.196.867.335 Dielectric Spectroscopy
-	E05.196.867.519 Magnetic Resonance Spectroscopy
-	E05.196.867.519.136 Carbon-13 Magnetic Resonance Spectroscopy
-	E05.196.867.519.274 Electron Spin Resonance Spectroscopy
-	E05.196.867.519.550 Nuclear Magnetic Resonance, Biomolecular
-	E05.196.867.519.775 Proton Magnetic Resonance Spectroscopy
-	E05.196.867.576 Optical Rotatory Dispersion
-	E05.196.867.618 Photoelectron Spectroscopy
-	E05.196.867.660 Pulse Radiolysis
-	E05.196.867.726 Spectrometry, Fluorescence
-	E05.196.867.776 Spectrometry, Gamma
-	E05.196.867.776.751 Spectroscopy, Mossbauer
-	E05.196.867.800 Spectrometry, X-Ray Emission



## MeSH Tree Changes for 2017

Type	Tree - heading
-	E05.196.867.800.360 Electron Probe Microanalysis
-	E05.196.867.826 Spectrophotometry
-	E05.196.867.826.150 Action Spectrum
-	E05.196.867.826.300 Microspectrophotometry
-	E05.196.867.826.551 Spectrophotometry, Atomic
-	E05.196.867.826.676 Spectrophotometry, Infrared
-	E05.196.867.826.676.700 Spectroscopy, Fourier Transform Infrared
-	E05.196.867.826.802 Spectrophotometry, Ultraviolet
-	E05.196.867.838 Spectroscopy, Electron Energy-Loss
-	E05.196.867.838.500 Microscopy, Energy-Filtering Transmission Electron
-	E05.196.867.851 Spectroscopy, Near-Infrared
-	E05.196.867.890 Spectrum Analysis, Raman
-	E05.196.867.917 Sun Protection Factor
-	E05.196.867.945 Terahertz Spectroscopy
-	E05.196.867.972 X-Ray Absorption Spectroscopy
-	E05.196.885 Spin Trapping
-	E05.196.890 Surface Plasmon Resonance
-	E05.196.904 Thermogravimetry
-	E05.196.922 Titrimetry
-	E05.196.922.250 Colorimetry
-	E05.196.922.500 Conductometry
-	E05.196.922.750 Potentiometry
-	E05.196.941 Ultracentrifugation
-	E05.196.941.336 Centrifugation, Density Gradient
-	E05.196.941.336.253 Centrifugation, Isopycnic
-	E05.196.941.336.419 Centrifugation, Zonal
New Tree	<a href="#">E05.197</a> <a href="#">Chemistry Techniques, Synthetic</a>
New Tree	<a href="#">E05.197.124</a> <a href="#">Click Chemistry</a>
New Tree	<a href="#">E05.197.312</a> <a href="#">Combinatorial Chemistry Techniques</a>
New Tree	<a href="#">E05.197.312.500</a> <a href="#">SELEX Aptamer Technique</a>
New Tree	<a href="#">E05.197.406</a> <a href="#">Cycloaddition Reaction</a>
New Tree	<a href="#">E05.197.500</a> <a href="#">Solid-Phase Synthesis Techniques</a>

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	E05.198	Chromophore-Assisted Light Inactivation
-	E05.200	Clinical Laboratory Techniques
-	E05.200.124	Clinical Chemistry Tests
-	E05.200.124.100	Blood Chemical Analysis
-	E05.200.124.100.100	Blood Gas Analysis
-	E05.200.124.100.100.600	Oximetry
-	E05.200.124.100.100.600.100	Blood Gas Monitoring, Transcutaneous
-	E05.200.124.100.105	Blood Glucose Self-Monitoring
-	E05.200.124.100.110	Blood Protein Electrophoresis
-	E05.200.124.100.115	Blood Urea Nitrogen
-	E05.200.124.100.232	Dried Blood Spot Testing
-	E05.200.124.100.350	Glucose Clamp Technique
-	E05.200.124.100.355	Glucose Tolerance Test
-	E05.200.124.100.450	Lactose Tolerance Test
-	E05.200.124.100.600	Petrosal Sinus Sampling
-	E05.200.124.200	Clinical Enzyme Tests
-	E05.200.124.300	Gastric Acidity Determination
-	E05.200.124.470	Limulus Test
-	E05.200.124.810	Urinalysis
-	E05.200.500	Cytological Techniques
-	E05.200.500.195	Cell Count
-	E05.200.500.195.107	Blood Cell Count
-	E05.200.500.195.107.330	Erythrocyte Count
-	E05.200.500.195.107.330.725	Reticulocyte Count
-	E05.200.500.195.107.595	Leukocyte Count
-	E05.200.500.195.107.595.500	Lymphocyte Count
-	E05.200.500.195.107.595.500.150	CD4 Lymphocyte Count
-	E05.200.500.195.107.595.500.150.160	CD4-CD8 Ratio
-	E05.200.500.195.107.740	Platelet Count
-	E05.200.500.195.870	Sperm Count
-	E05.200.500.265	Cell Culture Techniques
-	E05.200.500.265.500	Primary Cell Culture
-	E05.200.500.335	Cell Migration Assays
-	E05.200.500.335.500	Cell Migration Assays, Leukocyte
-	E05.200.500.335.750	Cell Migration Assays, Macrophage
-	E05.200.500.335.875	Skin Window Technique

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E05.200.500.363 Cell Separation
-	E05.200.500.363.171 Cytapheresis
-	E05.200.500.363.171.570 Leukapheresis
-	E05.200.500.363.171.790 Plateletpheresis
-	E05.200.500.363.342 Flow Cytometry
-	E05.200.500.363.441 Immunomagnetic Separation
-	E05.200.500.363.835 Leukocyte Reduction Procedures
-	E05.200.500.373 Cell Tracking
-	E05.200.500.380 Cellular Reprogramming Techniques
-	E05.200.500.380.250 Mitochondrial Replacement Therapy
-	E05.200.500.380.500 Nuclear Transfer Techniques
-	E05.200.500.383 Colony-Forming Units Assay
-	E05.200.500.383.910 Tumor Stem Cell Assay
-	E05.200.500.384 Cytodiagnosis
-	E05.200.500.384.050 Amniocentesis
-	E05.200.500.384.100 Biopsy
-	E05.200.500.384.100.119 Biopsy, Needle
-	E05.200.500.384.100.119.500 Biopsy, Fine-Needle
-	E05.200.500.384.100.119.500.500 Endoscopic Ultrasound-Guided Fine Needle Aspiration
-	E05.200.500.384.100.119.750 Biopsy, Large-Core Needle
-	E05.200.500.384.100.149 Chorionic Villi Sampling
-	E05.200.500.384.100.160 Conization
-	E05.200.500.384.100.370 Image-Guided Biopsy
-	E05.200.500.384.100.370.500 Endoscopic Ultrasound-Guided Fine Needle Aspiration
-	E05.200.500.384.100.422 Papanicolaou Test
-	E05.200.500.384.100.580 Sentinel Lymph Node Biopsy
-	E05.200.500.384.100.800 Vaginal Smears
-	E05.200.500.384.235 Cytopathogenic Effect, Viral
-	E05.200.500.384.467 Karyometry
-	E05.200.500.385 Cytogenetic Analysis
-	E05.200.500.385.130 Chromosome Banding
-	E05.200.500.385.315 Karyotyping
-	E05.200.500.385.315.800 Spectral Karyotyping
-	E05.200.500.385.500 Mitotic Index

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E05.200.500.386 Cytophotometry
-	E05.200.500.386.350 Flow Cytometry
-	E05.200.500.386.400 Image Cytometry
-	E05.200.500.386.400.500 Laser Scanning Cytometry
-	E05.200.500.388 Drug Screening Assays, Antitumor
-	E05.200.500.388.930 Tumor Stem Cell Assay
-	E05.200.500.454 Electroporation
-	E05.200.500.454.500 Electrochemotherapy
-	E05.200.500.508 Enzyme-Linked Immunospot Assay
-	E05.200.500.607 Histocytochemistry
-	E05.200.500.607.512 Immunohistochemistry
-	E05.200.500.607.512.240 Fluorescent Antibody Technique
-	E05.200.500.607.512.240.149 Antibody-Coated Bacteria Test, Urinary
-	E05.200.500.607.512.240.300 Fluorescent Antibody Technique, Direct
-	E05.200.500.607.512.240.310 Fluorescent Antibody Technique, Indirect
-	E05.200.500.607.512.240.655 Fluoroimmunoassay
-	E05.200.500.607.512.240.655.350 Fluorescence Polarization Immunoassay
-	E05.200.500.607.790 Periodic Acid-Schiff Reaction
-	E05.200.500.607.810 Prussian Blue Reaction
-	E05.200.500.620 Histocytological Preparation Techniques
-	E05.200.500.620.265 Microdissection
-	E05.200.500.620.530 Microtomy
-	E05.200.500.620.530.160 Cryoultramicrotomy
-	E05.200.500.620.530.160.260 Frozen Sections
-	E05.200.500.620.620 Replica Techniques
-	E05.200.500.620.620.150 Corrosion Casting
-	E05.200.500.620.620.260 Freeze Fracturing
-	E05.200.500.620.620.260.400 Freeze Etching
-	E05.200.500.620.670 Staining and Labeling
-	E05.200.500.620.670.130 Chromosome Banding
-	E05.200.500.620.670.325 In Situ Hybridization
-	E05.200.500.620.670.325.350 In Situ Hybridization, Fluorescence
-	E05.200.500.620.670.325.350.125 Chromosome Painting
-	E05.200.500.620.670.325.680 Primed In Situ Labeling
-	E05.200.500.620.670.520 Negative Staining
-	E05.200.500.620.670.570 Neuroanatomical Tract-Tracing Techniques

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E05.200.500.620.670.620      Periodic Acid-Schiff Reaction
-	E05.200.500.620.670.660      Prussian Blue Reaction
-	E05.200.500.620.670.770      Shadowing (Histology)
-	E05.200.500.620.670.780      Silver Staining
-	E05.200.500.620.720      Tissue Embedding
-	E05.200.500.620.720.610      Paraffin Embedding
-	E05.200.500.620.720.640      Plastic Embedding
-	E05.200.500.620.760      Tissue Preservation
-	E05.200.500.620.760.160      Cryopreservation
-	E05.200.500.620.760.160.260      Freeze Drying
-	E05.200.500.620.760.160.260.270      Freeze Substitution
-	E05.200.500.620.760.720      Tissue Fixation
-	E05.200.500.810      Metabolic Flux Analysis
-	E05.200.500.905      Patch-Clamp Techniques
-	E05.200.562      Genetic Testing
New Heading	<b>E05.200.562.500      Pharmacogenomic Testing</b>
-	E05.200.625      Hematologic Tests
-	E05.200.625.107      Blood Cell Count
-	E05.200.625.107.330      Erythrocyte Count
-	E05.200.625.107.330.725      Reticulocyte Count
-	E05.200.625.107.595      Leukocyte Count
-	E05.200.625.107.595.500      Lymphocyte Count
-	E05.200.625.107.595.500.150      CD4 Lymphocyte Count
-	E05.200.625.107.595.500.150.160      CD4-CD8 Ratio
-	E05.200.625.107.700      Platelet Count
-	E05.200.625.115      Blood Coagulation Tests
-	E05.200.625.115.320      International Normalized Ratio
-	E05.200.625.115.600      Partial Thromboplastin Time
-	E05.200.625.115.610      Prothrombin Time
-	E05.200.625.115.830      Thrombelastography
-	E05.200.625.115.870      Thrombin Time
-	E05.200.625.115.950      Whole Blood Coagulation Time
-	E05.200.625.120      Blood Grouping and Crossmatching
-	E05.200.625.125      Blood Sedimentation
-	E05.200.625.135      Bone Marrow Examination

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E05.200.625.225 Erythrocyte Aggregation
-	E05.200.625.230 Erythrocyte Indices
-	E05.200.625.265 Fibrin Clot Lysis Time
-	E05.200.625.300 FIGLU Test
-	E05.200.625.400 Hematocrit
-	E05.200.625.410 Hemoglobinometry
-	E05.200.625.550 Osmotic Fragility
-	E05.200.625.625 Platelet Function Tests
-	E05.200.625.625.100 Bleeding Time
-	E05.200.625.625.160 Clot Retraction
-	E05.200.625.625.392 Mean Platelet Volume
-	E05.200.625.625.625 Platelet Count
-	E05.200.625.750 Schilling Test
-	E05.200.750 Histological Techniques
-	E05.200.750.132 Autoradiography
-	E05.200.750.288 Decalcification Technique
-	E05.200.750.551 Histocytochemistry
-	E05.200.750.551.512 Immunohistochemistry
-	E05.200.750.551.512.240 Fluorescent Antibody Technique
-	E05.200.750.551.512.240.149 Antibody-Coated Bacteria Test, Urinary
-	E05.200.750.551.512.240.300 Fluorescent Antibody Technique, Direct
-	E05.200.750.551.512.240.310 Fluorescent Antibody Technique, Indirect
-	E05.200.750.551.512.240.655 Fluoroimmunoassay
-	E05.200.750.551.512.240.655.350 Fluorescence Polarization Immunoassay
-	E05.200.750.551.790 Periodic Acid-Schiff Reaction
-	E05.200.750.551.810 Prussian Blue Reaction
-	E05.200.750.600 Histocytological Preparation Techniques
-	E05.200.750.600.520 Microdissection
-	E05.200.750.600.530 Microtomy
-	E05.200.750.600.530.160 Cryoultramicrotomy
-	E05.200.750.600.530.160.260 Frozen Sections
-	E05.200.750.600.620 Replica Techniques
-	E05.200.750.600.620.150 Corrosion Casting
-	E05.200.750.600.620.260 Freeze Fracturing
-	E05.200.750.600.620.260.400 Freeze Etching
-	E05.200.750.600.670 Staining and Labeling

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E05.200.750.600.670.130 Chromosome Banding
-	E05.200.750.600.670.325 In Situ Hybridization
-	E05.200.750.600.670.325.350 In Situ Hybridization, Fluorescence
-	E05.200.750.600.670.325.350.125 Chromosome Painting
-	E05.200.750.600.670.325.680 Primed In Situ Labeling
-	E05.200.750.600.670.520 Negative Staining
-	E05.200.750.600.670.570 Neuroanatomical Tract-Tracing Techniques
-	E05.200.750.600.670.660 Prussian Blue Reaction
-	E05.200.750.600.670.770 Shadowing (Histology)
-	E05.200.750.600.670.780 Silver Staining
-	E05.200.750.600.720 Tissue Embedding
-	E05.200.750.600.720.610 Paraffin Embedding
-	E05.200.750.600.720.640 Plastic Embedding
-	E05.200.750.600.760 Tissue Preservation
-	E05.200.750.600.760.160 Cryopreservation
-	E05.200.750.600.760.160.260 Freeze Drying
-	E05.200.750.600.760.160.260.270 Freeze Substitution
-	E05.200.750.600.760.720 Tissue Fixation
-	E05.200.812 Immunologic Tests
-	E05.200.812.049 Antibody-Coated Bacteria Test, Urinary
-	E05.200.812.100 Basophil Degranulation Test
-	E05.200.812.125 Cell Migration Assays
-	E05.200.812.125.500 Cell Migration Assays, Leukocyte
-	E05.200.812.125.750 Cell Migration Assays, Macrophage
-	E05.200.812.125.875 Skin Window Technique
-	E05.200.812.160 Cytotoxicity Tests, Immunologic
-	E05.200.812.160.155 Complement Hemolytic Activity Assay
-	E05.200.812.375 Hemolytic Plaque Technique
-	E05.200.812.385 Histocompatibility Testing
-	E05.200.812.385.120 Blood Grouping and Crossmatching
-	E05.200.812.385.475 Lymphocyte Culture Test, Mixed
-	E05.200.812.400 Immune Adherence Reaction
-	E05.200.812.447 Immunophenotyping
-	E05.200.812.453 Interferon-gamma Release Tests
-	E05.200.812.460 Leukocyte Adherence Inhibition Test
-	E05.200.812.482 Lymphocyte Activation

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E05.200.812.505 Monitoring, Immunologic
-	E05.200.812.620 Pregnancy Tests, Immunologic
-	E05.200.812.706 Rosette Formation
-	E05.200.812.735 Serologic Tests
-	E05.200.812.735.050 Agglutination Tests
-	E05.200.812.735.050.375 Hemagglutination Tests
-	E05.200.812.735.050.375.150 Coombs Test
-	E05.200.812.735.050.450 Latex Fixation Tests
-	E05.200.812.735.060 AIDS Serodiagnosis
-	E05.200.812.735.150 Complement Fixation Tests
-	E05.200.812.735.155 Complement Hemolytic Activity Assay
-	E05.200.812.735.360 Hemadsorption Inhibition Tests
-	E05.200.812.735.370 Hemagglutination Inhibition Tests
-	E05.200.812.735.550 Neutralization Tests
-	E05.200.812.735.645 Precipitin Tests
-	E05.200.812.735.645.300 Flocculation Tests
-	E05.200.812.735.645.350 Immunodiffusion
-	E05.200.812.735.645.350.350 Immunoelectrophoresis
-	E05.200.812.735.645.350.350.150 Counterimmunoelectrophoresis
-	E05.200.812.735.645.350.350.350 Immunoelectrophoresis, Two-Dimensional
-	E05.200.812.735.830 Radioallergosorbent Test
-	E05.200.812.735.840 Radioimmunoprecipitation Assay
-	E05.200.812.735.845 Serum Bactericidal Antibody Assay
-	E05.200.812.735.850 Syphilis Serodiagnosis
-	E05.200.812.735.850.200 Fluorescent Treponemal Antibody-Absorption Test
-	E05.200.812.735.850.800 Treponema Immobilization Test
-	E05.200.812.742 Serotyping
-	E05.200.812.871 Skin Tests
-	E05.200.812.871.300 Intradermal Tests
-	E05.200.812.871.300.540 Kveim Test
-	E05.200.812.871.300.750 Skin Test End-Point Titration
-	E05.200.812.871.450 Local Lymph Node Assay
-	E05.200.812.871.600 Passive Cutaneous Anaphylaxis
-	E05.200.812.871.610 Patch Tests
-	E05.200.812.871.705 Skin Window Technique
-	E05.200.812.871.800 Tuberculin Test



## MeSH Tree Changes for 2017

Type	Tree - heading	
New Heading	<b>E05.200.828</b>	<b>Laboratory Critical Values</b>
-	E05.200.843	Metabolic Clearance Rate
-	E05.200.875	Microbiological Techniques
-	E05.200.875.074	Axenic Culture
-	E05.200.875.150	Bacteriological Techniques
-	E05.200.875.150.115	Bacterial Load
-	E05.200.875.150.125	Bacterial Typing Techniques
-	E05.200.875.150.125.150	Bacteriophage Typing
-	E05.200.875.150.125.457	Molecular Typing
-	E05.200.875.150.125.457.500	Multilocus Sequence Typing
-	E05.200.875.150.125.765	Ribotyping
-	E05.200.875.150.125.890	Serotyping
-	E05.200.875.150.570	Limulus Test
New Heading	<b>E05.200.875.185</b>	<b>Blood Culture</b>
-	E05.200.875.220	Colony Count, Microbial
-	E05.200.875.220.115	Bacterial Load
-	E05.200.875.595	Microbial Sensitivity Tests
-	E05.200.875.595.399	Disk Diffusion Antimicrobial Tests
-	E05.200.875.595.800	Serum Bactericidal Test
-	E05.200.875.610	Mycological Typing Techniques
-	E05.200.875.837	Serial Passage
-	E05.200.875.950	Viral Load
-	E05.200.875.970	Virus Cultivation
-	E05.200.875.970.790	Viral Plaque Assay
-	E05.200.875.977	Virus Inactivation
-	E05.200.875.985	Xenodiagnosis
-	E05.200.880	Molecular Diagnostic Techniques
-	E05.200.880.500	Human Papillomavirus DNA Tests
-	E05.200.910	Neonatal Screening
-	E05.200.925	Occult Blood
-	E05.200.932	Parasite Load
-	E05.200.932.600	Parasite Egg Count
-	E05.200.940	Parasitic Sensitivity Tests
-	E05.200.970	Pregnancy Tests

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E05.200.970.620      Pregnancy Tests, Immunologic
-	E05.200.985      Radioligand Assay
-	E05.200.992      Semen Analysis
-	E05.200.992.624      Sperm Count
-	E05.200.992.812      Sperm Motility
-	E05.200.996      Sex Determination Analysis
-	E05.200.996.500      Sex Determination by Skeleton
-	E05.200.998      Specimen Handling
-	E05.200.998.054      Biopsy
-	E05.200.998.054.119      Biopsy, Needle
-	E05.200.998.054.119.500      Biopsy, Fine-Needle
-	E05.200.998.054.119.500.500      Endoscopic Ultrasound-Guided Fine Needle Aspiration
-	E05.200.998.054.119.750      Biopsy, Large-Core Needle
-	E05.200.998.054.149      Chorionic Villi Sampling
-	E05.200.998.054.160      Conization
-	E05.200.998.054.370      Image-Guided Biopsy
-	E05.200.998.054.370.500      Endoscopic Ultrasound-Guided Fine Needle Aspiration
-	E05.200.998.054.422      Papanicolaou Test
-	E05.200.998.054.580      Sentinel Lymph Node Biopsy
-	E05.200.998.054.790      Spinal Puncture
-	E05.200.998.054.800      Vaginal Smears
-	E05.200.998.110      Blood Specimen Collection
-	E05.200.998.110.150      Cordocentesis
-	E05.200.998.110.600      Petrosal Sinus Sampling
-	E05.200.998.110.625      Phlebotomy
-	E05.200.998.221      Dissection
-	E05.200.998.221.580      Microdissection
-	E05.200.998.221.580.500      Laser Capture Microdissection
-	E05.200.998.293      DNA Contamination
-	E05.200.998.329      Paracentesis
-	E05.200.998.329.309      Amniocentesis
-	E05.200.998.329.465      Arthrocentesis
-	E05.200.998.329.543      Cordocentesis
-	E05.200.998.329.620      Pericardiocentesis

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E05.200.998.329.810                      Thoracentesis
-	E05.200.998.329.905                      Tympanocentesis
-	E05.200.998.762                            Urine Specimen Collection
-	E05.225                                        Constriction
-	E05.235                                        Containment of Biohazards
-	E05.242                                        Cytological Techniques
-	E05.242.195                                  Cell Count
-	E05.242.195.107                            Blood Cell Count
-	E05.242.195.107.330                      Erythrocyte Count
-	E05.242.195.107.330.725                Reticulocyte Count
-	E05.242.195.107.595                      Leukocyte Count
-	E05.242.195.107.595.500                Lymphocyte Count
-	E05.242.195.107.595.500.150            CD4 Lymphocyte Count
-	E05.242.195.107.595.500.150.160      CD4-CD8 Ratio
-	E05.242.195.107.740                      Platelet Count
-	E05.242.195.870                            Sperm Count
-	E05.242.223                                  Cell Culture Techniques
-	E05.242.223.500                            Primary Cell Culture
-	E05.242.251                                  Cell Fractionation
-	E05.242.307                                  Cell Fusion
-	E05.242.335                                  Cell Migration Assays
-	E05.242.335.500                            Cell Migration Assays, Leukocyte
-	E05.242.335.750                            Cell Migration Assays, Macrophage
-	E05.242.335.875                            Skin Window Technique
-	E05.242.363                                  Cell Separation
-	E05.242.363.171                            Cytapheresis
-	E05.242.363.171.570                      Leukapheresis
-	E05.242.363.171.790                      Plateletpheresis
-	E05.242.363.342                            Flow Cytometry
-	E05.242.363.441                            Immunomagnetic Separation
-	E05.242.363.540                            Leukocyte Reduction Procedures
-	E05.242.373                                  Cell Tracking
-	E05.242.378                                  Cellular Reprogramming Techniques
-	E05.242.383                                  Colony-Forming Units Assay
-	E05.242.383.910                            Tumor Stem Cell Assay
-	E05.242.384                                  Cytodiagnosis

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E05.242.384.050 Amniocentesis
-	E05.242.384.100 Biopsy
-	E05.242.384.100.119 Biopsy, Needle
-	E05.242.384.100.119.500 Biopsy, Fine-Needle
-	E05.242.384.100.119.500.500 Endoscopic Ultrasound-Guided Fine Needle Aspiration
-	E05.242.384.100.119.750 Biopsy, Large-Core Needle
-	E05.242.384.100.149 Chorionic Villi Sampling
-	E05.242.384.100.160 Conization
-	E05.242.384.100.370 Image-Guided Biopsy
-	E05.242.384.100.370.500 Endoscopic Ultrasound-Guided Fine Needle Aspiration
-	E05.242.384.100.422 Papanicolaou Test
-	E05.242.384.100.580 Sentinel Lymph Node Biopsy
-	E05.242.384.100.800 Vaginal Smears
-	E05.242.384.235 Cytopathogenic Effect, Viral
-	E05.242.384.467 Karyometry
-	E05.242.385 Cytogenetic Analysis
-	E05.242.385.130 Chromosome Banding
-	E05.242.385.315 Karyotyping
-	E05.242.385.315.800 Spectral Karyotyping
-	E05.242.385.500 Mitotic Index
-	E05.242.386 Cytophotometry
-	E05.242.386.350 Flow Cytometry
-	E05.242.386.400 Image Cytometry
-	E05.242.386.400.500 Laser Scanning Cytometry
-	E05.242.417 Drug Screening Assays, Antitumor
-	E05.242.417.500 Tumor Stem Cell Assay
-	E05.242.448 Electroporation
-	E05.242.448.500 Electrochemotherapy
-	E05.242.551 Enzyme-Linked Immunospot Assay
-	E05.242.654 Metabolic Flux Analysis
-	E05.242.800 Patch-Clamp Techniques
-	E05.242.900 Single-Cell Analysis
-	E05.242.950 Stem Cell Research
-	E05.245 Decision Support Techniques

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	E05.245.380	Data Interpretation, Statistical
-	E05.256	Dermatoglyphics
Old Tree	<b>E05.263</b>	<b>Diet, Atherogenic</b>
Old Tree	<b>E05.270</b>	<b>Diet, Cariogenic</b>
-	E05.284	Dilatation
-	E05.295	Drug Discovery
-	E05.295.500	Drug Design
-	E05.295.500.500	Polypharmacology
-	E05.295.750	Drug Evaluation, Preclinical
-	E05.295.875	Drug Repositioning
-	E05.298	Ecological Parameter Monitoring
-	E05.301	Electrochemical Techniques
-	E05.301.160	Conductometry
-	E05.301.250	Electrolysis
-	E05.301.250.348	Electroplating
-	E05.301.270	Electroosmosis
-	E05.301.300	Electrophoresis
-	E05.301.300.049	Blood Protein Electrophoresis
-	E05.301.300.074	Blotting, Northern
-	E05.301.300.087	Blotting, Southern
-	E05.301.300.093	Blotting, Southwestern
-	E05.301.300.096	Blotting, Western
-	E05.301.300.096.500	Blotting, Far-Western
-	E05.301.300.100	Electrophoresis, Agar Gel
-	E05.301.300.100.150	Comet Assay
-	E05.301.300.190	Electrophoresis, Capillary
-	E05.301.300.190.500	Capillary Electrochromatography
-	E05.301.300.200	Electrophoresis, Cellulose Acetate
-	E05.301.300.220	Electrophoresis, Gel, Pulsed-Field
-	E05.301.300.230	Electrophoresis, Gel, Two-Dimensional
-	E05.301.300.230.500	Two-Dimensional Difference Gel Electrophoresis
-	E05.301.300.236	Electrophoresis, Paper
-	E05.301.300.319	Electrophoresis, Polyacrylamide Gel
-	E05.301.300.319.201	Denaturing Gradient Gel Electrophoresis
-	E05.301.300.319.403	Electrophoresis, Disc
-	E05.301.300.319.725	Native Polyacrylamide Gel Electrophoresis

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E05.301.300.402 Electrophoresis, Starch Gel
-	E05.301.300.568 Immunoelectrophoresis
-	E05.301.300.568.250 Counterimmunoelectrophoresis
-	E05.301.300.568.520 Immunoelectrophoresis, Two-Dimensional
-	E05.301.300.575 Iontophoresis
-	E05.301.300.663 Isoelectric Focusing
-	E05.301.300.663.500 Isoelectric Point
-	E05.301.300.831 Isotachopheresis
-	E05.301.500 Electroporation
-	E05.301.500.500 Electrochemotherapy
-	E05.301.700 Polarography
-	E05.301.710 Potentiometry
-	E05.313 Embryo Research
-	E05.313.750 Research Embryo Creation
-	E05.315 Endpoint Determination
-	E05.318 Epidemiologic Methods
-	E05.318.270 Contact Tracing
-	E05.318.308 Data Collection
-	E05.318.308.028 Data Accuracy
-	E05.318.308.056 Datasets as Topic
-	E05.318.308.112 Focus Groups
-	E05.318.308.225 Geriatric Assessment
-	E05.318.308.237 Health Impact Assessment
-	E05.318.308.420 Interviews as Topic
-	E05.318.308.461 Lot Quality Assurance Sampling
-	E05.318.308.502 Narration
-	E05.318.308.585 Nutrition Assessment
-	E05.318.308.940 Records as Topic
-	E05.318.308.940.250 Birth Certificates
-	E05.318.308.940.350 Death Certificates
-	E05.318.308.940.375 Dental Records
-	E05.318.308.940.425 Hospital Records
-	E05.318.308.940.968 Medical Records
-	E05.318.308.940.968.249 Health Records, Personal
-	E05.318.308.940.968.249.500 Electronic Health Records
New	<b>E05.318.308.940.968.249.500.500 Patient Portals</b>

## MeSH Tree Changes for 2017

Type	Tree - heading	
Heading		
New Heading	<b>E05.318.308.940.968.249.750</b>	<b>Patient Portals</b>
-	E05.318.308.940.968.500	Medical Record Linkage
-	E05.318.308.940.968.550	Medical Records, Problem-Oriented
-	E05.318.308.940.968.625	Medical Records Systems, Computerized
-	E05.318.308.940.968.625.500	Electronic Health Records
-	E05.318.308.940.968.625.500.500	Health Information Exchange
-	E05.318.308.940.968.625.750	Health Smart Cards
-	E05.318.308.940.968.875	Trauma Severity Indices
-	E05.318.308.940.968.875.125	Abbreviated Injury Scale
-	E05.318.308.940.968.875.250	Glasgow Coma Scale
-	E05.318.308.940.968.875.260	Glasgow Outcome Scale
-	E05.318.308.940.968.875.500	Injury Severity Score
-	E05.318.308.940.984	Nursing Records
-	E05.318.308.970	Registries
-	E05.318.308.970.725	SEER Program
-	E05.318.308.980	Surveys and Questionnaires
-	E05.318.308.980.313	Contraceptive Prevalence Surveys
-	E05.318.308.980.344	Health Care Surveys
New Heading	<b>E05.318.308.980.344.500</b>	<b>Patient Reported Outcome Measures</b>
-	E05.318.308.980.438	Health Surveys
-	E05.318.308.980.438.149	Behavioral Risk Factor Surveillance System
-	E05.318.308.980.438.300	Dental Health Surveys
-	E05.318.308.980.438.300.300	Dental Plaque Index
-	E05.318.308.980.438.300.350	DMF Index
-	E05.318.308.980.438.300.512	Index of Orthodontic Treatment Need
-	E05.318.308.980.438.300.675	Oral Hygiene Index
-	E05.318.308.980.438.300.725	Periodontal Index
-	E05.318.308.980.438.475	Health Status Indicators
-	E05.318.308.980.438.475.365	APACHE
Old Tree	<b>E05.318.308.980.438.475.410</b>	<b>Organ Dysfunction Scores</b>
-	E05.318.308.980.438.475.456	Patient Acuity
-	E05.318.308.980.438.475.456.500	Severity of Illness Index
New Tree	<b>E05.318.308.980.438.475.456.500.250</b>	<b>APACHE</b>

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E05.318.308.980.438.475.456.500.500      Karnofsky Performance Status
New Tree	<a href="#">E05.318.308.980.438.475.456.500.625</a> <a href="#">Organ Dysfunction Scores</a>
New Heading	<b>E05.318.308.980.438.475.456.500.750</b> <b>Simplified Acute Physiology Score</b>
-	E05.318.308.980.438.475.730      Sickness Impact Profile
-	E05.318.308.980.438.580      Mass Screening
-	E05.318.308.980.438.580.174      Anonymous Testing
-	E05.318.308.980.438.580.510      Mass Chest X-Ray
-	E05.318.308.980.438.580.560      Multiphasic Screening
-	E05.318.308.980.438.580.580      Neonatal Screening
-	E05.318.308.980.438.580.925      Vision Screening
-	E05.318.308.980.438.700      Population Surveillance
-	E05.318.308.980.438.700.324      Public Health Surveillance
-	E05.318.308.980.438.700.650      Sentinel Surveillance
-	E05.318.308.980.485      Nutrition Surveys
-	E05.318.308.980.485.350      Diet Surveys
-	E05.318.308.980.500      Self Report
-	E05.318.308.985      Vital Statistics
-	E05.318.308.985.450      Life Expectancy
-	E05.318.308.985.475      Life Tables
-	E05.318.308.985.525      Morbidity
-	E05.318.308.985.525.080      Basic Reproduction Number
-	E05.318.308.985.525.375      Incidence
-	E05.318.308.985.525.750      Prevalence
-	E05.318.308.985.550      Mortality
-	E05.318.308.985.550.250      Cause of Death
-	E05.318.308.985.550.287      Child Mortality
-	E05.318.308.985.550.325      Fatal Outcome
-	E05.318.308.985.550.362      Fetal Mortality
-	E05.318.308.985.550.400      Hospital Mortality
-	E05.318.308.985.550.475      Infant Mortality
New Tree	<a href="#">E05.318.308.985.550.475.500</a> <a href="#">Perinatal Mortality</a>
-	E05.318.308.985.550.500      Maternal Mortality
-	E05.318.308.985.550.550      Mortality, Premature
Old Tree	<b>E05.318.308.985.550.700</b> <b>Perinatal Mortality</b>



## MeSH Tree Changes for 2017

Type	Tree - heading
-	E05.318.308.985.550.900 Survival Rate
-	E05.318.308.985.775 Pregnancy Rate
-	E05.318.308.985.775.500 Birth Rate
-	E05.318.362 Disease Notification
-	E05.318.375 Epidemiological Monitoring
-	E05.318.389 Geographic Mapping
-	E05.318.416 Molecular Epidemiology
-	E05.318.416.249 Genome-Wide Association Study
-	E05.318.416.500 Mendelian Randomization Analysis
-	E05.318.650 Sentinel Surveillance
-	E05.318.740 Statistics as Topic
-	E05.318.740.100 Actuarial Analysis
-	E05.318.740.100.500 Life Tables
-	E05.318.740.100.500.700 Quality-Adjusted Life Years
-	E05.318.740.150 Analysis of Variance
-	E05.318.740.150.500 Multivariate Analysis
-	E05.318.740.200 Area Under Curve
-	E05.318.740.225 Biometry
-	E05.318.740.225.500 Biometric Identification
-	E05.318.740.225.500.500 DNA Fingerprinting
-	E05.318.740.237 Biostatistics
-	E05.318.740.250 Cluster Analysis
-	E05.318.740.250.675 Small-Area Analysis
-	E05.318.740.250.700 Space-Time Clustering
-	E05.318.740.275 Confidence Intervals
-	E05.318.740.300 Data Interpretation, Statistical
-	E05.318.740.350 Discriminant Analysis
-	E05.318.740.400 Factor Analysis, Statistical
-	E05.318.740.475 Matched-Pair Analysis
-	E05.318.740.500 Models, Statistical
-	E05.318.740.500.475 Likelihood Functions
-	E05.318.740.500.500 Linear Models
-	E05.318.740.500.525 Logistic Models
-	E05.318.740.500.600 Models, Economic
-	E05.318.740.500.600.500 Models, Econometric
-	E05.318.740.500.625 Nomograms

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E05.318.740.500.700 Proportional Hazards Models
-	E05.318.740.525 Monte Carlo Method
-	E05.318.740.562 Principal Component Analysis
-	E05.318.740.600 Probability
-	E05.318.740.600.200 Bayes Theorem
-	E05.318.740.600.400 Likelihood Functions
-	E05.318.740.600.500 Markov Chains
-	E05.318.740.600.600 Odds Ratio
-	E05.318.740.600.650 Position-Specific Scoring Matrices
-	E05.318.740.600.675 Propensity Score
-	E05.318.740.600.700 Proportional Hazards Models
-	E05.318.740.600.800 Risk
-	E05.318.740.600.800.450 Logistic Models
-	E05.318.740.600.800.582 Protective Factors
-	E05.318.740.600.800.715 Risk Assessment
-	E05.318.740.600.800.715.500 Healthcare Failure Mode and Effect Analysis
-	E05.318.740.600.800.725 Risk Factors
-	E05.318.740.600.900 Uncertainty
-	E05.318.740.750 Regression Analysis
-	E05.318.740.750.400 Least-Squares Analysis
-	E05.318.740.750.425 Linear Models
-	E05.318.740.750.450 Logistic Models
-	E05.318.740.750.725 Proportional Hazards Models
-	E05.318.740.750.862 Spatial Regression
-	E05.318.740.872 Sensitivity and Specificity
-	E05.318.740.872.374 Limit of Detection
-	E05.318.740.872.750 ROC Curve
-	E05.318.740.872.875 Signal-To-Noise Ratio
-	E05.318.740.933 Spatial Analysis
-	E05.318.740.933.249 Geographic Mapping
-	E05.318.740.933.375 Spatial Regression
-	E05.318.740.933.500 Spatio-Temporal Analysis
-	E05.318.740.994 Statistical Distributions
-	E05.318.740.994.250 Binomial Distribution
-	E05.318.740.994.300 Chi-Square Distribution
-	E05.318.740.994.500 Normal Distribution

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E05.318.740.994.750 Poisson Distribution
-	E05.318.740.995 Statistics, Nonparametric
-	E05.318.740.996 Stochastic Processes
-	E05.318.740.996.500 Markov Chains
-	E05.318.740.998 Survival Analysis
-	E05.318.740.998.300 Disease-Free Survival
-	E05.318.740.998.650 Kaplan-Meier Estimate
-	E05.318.740.998.825 Proportional Hazards Models
-	E05.318.760 Epidemiologic Study Characteristics as Topic
-	E05.318.760.250 Clinical Studies as Topic
-	E05.318.760.250.500 Clinical Trials as Topic
-	E05.318.760.250.500.200 Clinical Trials, Phase I as Topic
-	E05.318.760.250.500.210 Clinical Trials, Phase II as Topic
-	E05.318.760.250.500.220 Clinical Trials, Phase III as Topic
-	E05.318.760.250.500.230 Clinical Trials, Phase IV as Topic
-	E05.318.760.250.500.365 Controlled Clinical Trials as Topic
-	E05.318.760.250.500.365.250 Non-Randomized Controlled Trials as Topic
-	E05.318.760.250.500.365.500 Randomized Controlled Trials as Topic
-	E05.318.760.250.500.365.500.250 Intention to Treat Analysis
-	E05.318.760.250.500.365.500.500 Pragmatic Clinical Trials as Topic
-	E05.318.760.250.750 Observational Studies as Topic
-	E05.318.760.500 Epidemiologic Studies
-	E05.318.760.500.500 Case-Control Studies
-	E05.318.760.500.500.500 Retrospective Studies
-	E05.318.760.500.750 Cohort Studies
-	E05.318.760.500.750.249 Follow-Up Studies
-	E05.318.760.500.750.500 Longitudinal Studies
-	E05.318.760.500.750.500.500 National Longitudinal Study of Adolescent Health
-	E05.318.760.500.750.625 Prospective Studies
-	E05.318.760.500.750.750 Retrospective Studies
-	E05.318.760.500.812 Controlled Before-After Studies
-	E05.318.760.500.875 Cross-Sectional Studies
-	E05.318.760.500.912 Historically Controlled Study
-	E05.318.760.500.931 Interrupted Time Series Analysis
-	E05.318.760.500.950 Seroepidemiologic Studies

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E05.318.760.500.950.375 HIV Seroprevalence
-	E05.318.760.550 Feasibility Studies
-	E05.318.760.658 Multicenter Studies as Topic
-	E05.318.760.750 Pilot Projects
-	E05.318.760.875 Sampling Studies
-	E05.318.760.900 Twin Studies as Topic
-	E05.318.780 Epidemiologic Research Design
-	E05.318.780.074 Control Groups
-	E05.318.780.150 Cross-Over Studies
-	E05.318.780.300 Double-Blind Method
-	E05.318.780.392 Genome-Wide Association Study
-	E05.318.780.438 Lost to Follow-Up
-	E05.318.780.485 Matched-Pair Analysis
-	E05.318.780.500 Meta-Analysis as Topic
New Heading	<b>E05.318.780.500.500 Network Meta-Analysis</b>
-	E05.318.780.700 Random Allocation
-	E05.318.780.725 Reproducibility of Results
-	E05.318.780.725.250 Data Accuracy
-	E05.318.780.725.500 Dimensional Measurement Accuracy
-	E05.318.780.762 Sample Size
-	E05.318.780.800 Sensitivity and Specificity
-	E05.318.780.800.650 Predictive Value of Tests
-	E05.318.780.800.750 ROC Curve
-	E05.318.780.800.875 Signal-To-Noise Ratio
-	E05.318.780.850 Single-Blind Method
-	E05.320 Equipment Design
-	E05.320.274 Orthodontic Appliance Design
-	E05.320.550 Prosthesis Design
-	E05.320.550.550 Prosthesis Coloring
-	E05.325 Equipment Failure
-	E05.325.095 Catheter Obstruction
-	E05.325.192 Equipment Failure Analysis
-	E05.325.578 Intrauterine Device Migration
-	E05.325.578.500 Intrauterine Device Expulsion
-	E05.325.771 Prosthesis Failure

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	E05.328	Equipment Reuse
-	E05.330	Equipment Safety
-	E05.333	Ergometry
-	E05.333.250	Exercise Test
-	E05.335	Euthanasia, Animal
-	E05.337	Evaluation Studies as Topic
-	E05.337.275	Device Approval
-	E05.337.275.500	Diagnostic Test Approval
-	E05.337.300	Drug Approval
-	E05.337.300.199	Compassionate Use Trials
-	E05.337.300.400	Investigational New Drug Application
-	E05.337.425	Drug Evaluation
-	E05.337.550	Drug Evaluation, Preclinical
-	E05.337.550.200	Drug Screening Assays, Antitumor
-	E05.337.550.200.800	Tumor Stem Cell Assay
-	E05.337.550.200.900	Xenograft Model Antitumor Assays
-	E05.337.550.200.900.830	Subrenal Capsule Assay
-	E05.337.550.400	Microbial Sensitivity Tests
-	E05.337.550.400.800	Serum Bactericidal Test
-	E05.337.550.700	Parasitic Sensitivity Tests
-	E05.337.675	Feasibility Studies
-	E05.337.737	Pilot Projects
-	E05.337.800	Product Surveillance, Postmarketing
-	E05.337.800.120	Adverse Drug Reaction Reporting Systems
-	E05.337.800.200	Clinical Trials, Phase IV as Topic
-	E05.337.800.600	Pharmacovigilance
-	E05.337.820	Program Evaluation
-	E05.337.851	Reproducibility of Results
-	E05.337.851.500	Dimensional Measurement Accuracy
-	E05.337.925	Validation Studies as Topic
-	E05.349	Fetal Research
-	E05.355	Finite Element Analysis
-	E05.362	Food Analysis
-	E05.377	Fourier Analysis
-	E05.385	Games, Experimental
-	E05.385.500	Prisoner Dilemma

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E05.393 Genetic Techniques
-	E05.393.085 Cellular Reprogramming Techniques
-	E05.393.085.250 Mitochondrial Replacement Therapy
-	E05.393.085.500 Nuclear Transfer Techniques
-	E05.393.170 Chromatin Immunoprecipitation
-	E05.393.183 Chromosome Mapping
-	E05.393.183.250 Chromosome Walking
-	E05.393.183.620 Physical Chromosome Mapping
-	E05.393.183.620.160 Contig Mapping
-	E05.393.183.620.405 Radiation Hybrid Mapping
-	E05.393.183.620.650 Restriction Mapping
-	E05.393.183.620.650.500 Optical Restriction Mapping
-	E05.393.220 Cloning, Molecular
-	E05.393.220.250 Artificial Gene Fusion
-	E05.393.220.250.500 Cell Surface Display Techniques
-	E05.393.220.870 Two-Hybrid System Techniques
-	E05.393.240 Cloning, Organism
-	E05.393.260 Contig Mapping
-	E05.393.281 Crosses, Genetic
-	E05.393.281.526 Genetic Complementation Test
-	E05.393.285 Cytogenetic Analysis
-	E05.393.285.130 Chromosome Banding
-	E05.393.285.240 Comparative Genomic Hybridization
-	E05.393.285.350 In Situ Hybridization, Fluorescence
-	E05.393.285.350.125 Chromosome Painting
-	E05.393.285.350.125.800 Spectral Karyotyping
-	E05.393.285.475 Karyotyping
-	E05.393.285.475.800 Spectral Karyotyping
-	E05.393.285.830 Sex Determination Analysis
-	E05.393.285.830.500 Sex Determination by Skeleton
-	E05.393.290 DNA Fingerprinting
-	E05.393.290.382 Amplified Fragment Length Polymorphism Analysis
-	E05.393.290.573 DNA Contamination
-	E05.393.290.765 Ribotyping
-	E05.393.300 DNA Footprinting
-	E05.393.332 Gene Expression Profiling

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E05.393.332.500 Subtractive Hybridization Techniques
-	E05.393.335 Gene Targeting
-	E05.393.335.249 Gene Knock-In Techniques
-	E05.393.335.500 Gene Knockdown Techniques
-	E05.393.335.750 Gene Knockout Techniques
-	E05.393.350 Gene Transfer Techniques
-	E05.393.350.100 Biolistics
-	E05.393.350.800 Transduction, Genetic
-	E05.393.350.810 Transfection
-	E05.393.350.810.500 Transformation, Bacterial
-	E05.393.385 Genetic Association Studies
-	E05.393.385.500 Genome-Wide Association Study
-	E05.393.420 Genetic Engineering
-	E05.393.420.175 Directed Molecular Evolution
-	E05.393.420.238 DNA Shuffling
New Heading	<b>E05.393.420.270 Gene Editing</b>
-	E05.393.420.301 Genetic Therapy
-	E05.393.420.301.500 Targeted Gene Repair
-	E05.393.420.451 Genetic Enhancement
-	E05.393.420.526 Metabolic Engineering
-	E05.393.420.601 Protein Engineering
-	E05.393.420.601.035 Amino Acid Substitution
-	E05.393.420.601.292 Cell Surface Display Techniques
-	E05.393.420.601.550 Mutagenesis, Insertional
-	E05.393.420.601.575 Mutagenesis, Site-Directed
-	E05.393.420.601.575.500 Targeted Gene Repair
-	E05.393.420.890 Sex Preselection
-	E05.393.435 Genetic Testing
New Heading	<b>E05.393.435.500 Pharmacogenomic Testing</b>
-	E05.393.442 Genotyping Techniques
-	E05.393.450 Heterozygote Detection
-	E05.393.475 In Situ Nick-End Labeling
-	E05.393.520 Molecular Diagnostic Techniques
-	E05.393.520.500 Comparative Genomic Hybridization

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E05.393.520.750 Human Papillomavirus DNA Tests
-	E05.393.522 Molecular Epidemiology
-	E05.393.522.500 Genome-Wide Association Study
-	E05.393.542 Molecular Typing
-	E05.393.542.249 DNA Barcoding, Taxonomic
-	E05.393.542.500 Multilocus Sequence Typing
-	E05.393.560 Mutagenicity Tests
-	E05.393.560.150 Comet Assay
-	E05.393.560.598 Micronucleus Tests
-	E05.393.600 Nuclease Protection Assays
-	E05.393.620 Nucleic Acid Amplification Techniques
-	E05.393.620.311 Ligase Chain Reaction
-	E05.393.620.500 Polymerase Chain Reaction
-	E05.393.620.500.324 Amplified Fragment Length Polymorphism Analysis
-	E05.393.620.500.487 Multiplex Polymerase Chain Reaction
-	E05.393.620.500.650 Primed In Situ Labeling
-	E05.393.620.500.687 Random Amplified Polymorphic DNA Technique
-	E05.393.620.500.706 Real-Time Polymerase Chain Reaction
-	E05.393.620.500.725 Reverse Transcriptase Polymerase Chain Reaction
-	E05.393.620.700 Self-Sustained Sequence Replication
-	E05.393.640 Nucleic Acid Denaturation
-	E05.393.661 Nucleic Acid Hybridization
-	E05.393.661.124 Branched DNA Signal Amplification Assay
-	E05.393.661.187 Comparative Genomic Hybridization
-	E05.393.661.250 Heteroduplex Analysis
-	E05.393.661.475 In Situ Hybridization
-	E05.393.661.475.350 In Situ Hybridization, Fluorescence
-	E05.393.661.475.350.125 Chromosome Painting
-	E05.393.661.475.350.125.800 Spectral Karyotyping
-	E05.393.661.475.680 Primed In Situ Labeling
-	E05.393.661.640 Oligonucleotide Array Sequence Analysis
-	E05.393.661.820 Subtractive Hybridization Techniques
-	E05.393.667 Optogenetics
-	E05.393.673 Pedigree
-	E05.393.712 Restriction Mapping
-	E05.393.712.500 Optical Restriction Mapping



## MeSH Tree Changes for 2017

Type	Tree - heading
-	E05.393.731 Reverse Genetics
-	E05.393.751 Sequence Alignment
-	E05.393.760 Sequence Analysis
-	E05.393.760.319 High-Throughput Nucleotide Sequencing
-	E05.393.760.479 Molecular Sequence Annotation
-	E05.393.760.640 Oligonucleotide Array Sequence Analysis
-	E05.393.760.640.500 Genome-Wide Association Study
-	E05.393.760.670 Position-Specific Scoring Matrices
-	E05.393.760.700 Sequence Analysis, DNA
-	E05.393.760.700.149 DNA Barcoding, Taxonomic
-	E05.393.760.700.224 DNA Contamination
-	E05.393.760.700.300 DNA Mutational Analysis
-	E05.393.760.700.650 Multilocus Sequence Typing
-	E05.393.760.705 Sequence Analysis, Protein
-	E05.393.760.705.685 Peptide Mapping
-	E05.393.760.705.685.690 Protein Footprinting
-	E05.393.760.710 Sequence Analysis, RNA
-	E05.417 Hardness Tests
-	E05.445 Human Experimentation
-	E05.445.100 Autoexperimentation
-	E05.445.750 Nontherapeutic Human Experimentation
-	E05.445.875 Therapeutic Human Experimentation
-	E05.466 Immersion
-	E05.472 Immobilization
-	E05.472.760 Restraint, Physical
-	E05.472.760.370 Hindlimb Suspension
-	E05.478 Immunologic Techniques
-	E05.478.274 Epitope Mapping
-	E05.478.550 Immunization
-	E05.478.550.520 Immunization, Passive
-	E05.478.550.520.050 Adoptive Transfer
-	E05.478.550.520.050.400 Immunotherapy, Adoptive
-	E05.478.550.545 Immunization Schedule
-	E05.478.550.550 Immunization, Secondary
-	E05.478.550.600 Immunotherapy, Active
-	E05.478.550.600.890 Vaccination

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E05.478.550.600.890.500 Mass Vaccination
-	E05.478.566 Immunoassay
-	E05.478.566.159 Fluoroimmunoassay
-	E05.478.566.159.350 Fluorescence Polarization Immunoassay
-	E05.478.566.320 Immunoblotting
-	E05.478.566.320.200 Blotting, Western
-	E05.478.566.320.200.200 Blotting, Far-Western
-	E05.478.566.335 Immunochromatography
-	E05.478.566.350 Immunoenzyme Techniques
-	E05.478.566.350.170 Enzyme-Linked Immunosorbent Assay
-	E05.478.566.350.170.500 Enzyme-Linked Immunospot Assay
-	E05.478.566.350.180 Enzyme Multiplied Immunoassay Technique
-	E05.478.566.380 Immunosorbent Techniques
-	E05.478.566.380.360 Enzyme-Linked Immunosorbent Assay
-	E05.478.566.380.360.500 Enzyme-Linked Immunospot Assay
-	E05.478.566.380.810 Radioallergosorbent Test
-	E05.478.566.380.825 Radioimmunoprecipitation Assay
-	E05.478.566.380.830 Radioimmunosorbent Test
-	E05.478.566.639 Radioimmunoassay
-	E05.478.566.639.405 Immunoradiometric Assay
-	E05.478.566.639.810 Radioallergosorbent Test
-	E05.478.566.639.825 Radioimmunoprecipitation Assay
-	E05.478.566.639.830 Radioimmunosorbent Test
-	E05.478.583 Immunohistochemistry
-	E05.478.583.375 Fluorescent Antibody Technique
-	E05.478.583.375.050 Antibody-Coated Bacteria Test, Urinary
-	E05.478.583.375.300 Fluorescent Antibody Technique, Direct
-	E05.478.583.375.310 Fluorescent Antibody Technique, Indirect
-	E05.478.583.375.655 Fluoroimmunoassay
-	E05.478.583.375.655.350 Fluorescence Polarization Immunoassay
-	E05.478.583.400 Immunoenzyme Techniques
-	E05.478.583.400.170 Enzyme-Linked Immunosorbent Assay
-	E05.478.583.400.170.500 Enzyme-Linked Immunospot Assay
-	E05.478.583.400.180 Enzyme Multiplied Immunoassay Technique
-	E05.478.594 Immunologic Tests
-	E05.478.594.049 Antibody-Coated Bacteria Test, Urinary

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E05.478.594.100 Basophil Degranulation Test
-	E05.478.594.122 Cell Migration Assays
-	E05.478.594.122.500 Cell Migration Assays, Leukocyte
-	E05.478.594.122.750 Cell Migration Assays, Macrophage
-	E05.478.594.122.875 Skin Window Technique
-	E05.478.594.160 Cytotoxicity Tests, Immunologic
-	E05.478.594.160.150 Complement Hemolytic Activity Assay
-	E05.478.594.375 Hemolytic Plaque Technique
-	E05.478.594.385 Histocompatibility Testing
-	E05.478.594.385.120 Blood Grouping and Crossmatching
-	E05.478.594.385.429 Lymphocyte Culture Test, Mixed
-	E05.478.594.400 Immune Adherence Reaction
-	E05.478.594.450 Immunophenotyping
-	E05.478.594.475 Interferon-gamma Release Tests
-	E05.478.594.500 Leukocyte Adherence Inhibition Test
-	E05.478.594.530 Lymphocyte Activation
-	E05.478.594.550 Monitoring, Immunologic
-	E05.478.594.662 Pregnancy Tests, Immunologic
-	E05.478.594.730 Rosette Formation
-	E05.478.594.760 Serologic Tests
-	E05.478.594.760.050 Agglutination Tests
-	E05.478.594.760.050.375 Hemagglutination Tests
-	E05.478.594.760.050.375.150 Coombs Test
-	E05.478.594.760.050.450 Latex Fixation Tests
-	E05.478.594.760.060 AIDS Serodiagnosis
-	E05.478.594.760.150 Complement Fixation Tests
-	E05.478.594.760.155 Complement Hemolytic Activity Assay
-	E05.478.594.760.360 Hemadsorption Inhibition Tests
-	E05.478.594.760.370 Hemagglutination Inhibition Tests
-	E05.478.594.760.550 Neutralization Tests
-	E05.478.594.760.645 Precipitin Tests
-	E05.478.594.760.645.300 Flocculation Tests
-	E05.478.594.760.645.350 Immunodiffusion
-	E05.478.594.760.645.350.350 Immunelectrophoresis
-	E05.478.594.760.645.350.350.150 Counterimmunolectrophoresis
-	E05.478.594.760.645.350.350.350 Immunelectrophoresis, Two-Dimensional

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E05.478.594.760.830 Radioallergosorbent Test
-	E05.478.594.760.840 Radioimmunoprecipitation Assay
-	E05.478.594.760.845 Serum Bactericidal Antibody Assay
-	E05.478.594.760.850 Syphilis Serodiagnosis
-	E05.478.594.760.850.200 Fluorescent Treponemal Antibody-Absorption Test
-	E05.478.594.760.850.800 Treponema Immobilization Test
-	E05.478.594.780 Serotyping
-	E05.478.594.890 Skin Tests
-	E05.478.594.890.300 Intradermal Tests
-	E05.478.594.890.300.540 Kveim Test
-	E05.478.594.890.300.750 Skin Test End-Point Titration
-	E05.478.594.890.450 Local Lymph Node Assay
-	E05.478.594.890.600 Passive Cutaneous Anaphylaxis
-	E05.478.594.890.610 Patch Tests
-	E05.478.594.890.705 Skin Window Technique
-	E05.478.594.890.800 Tuberculin Test
-	E05.478.605 Immunoprecipitation
-	E05.478.605.160 Chromatin Immunoprecipitation
-	E05.478.605.492 Precipitin Tests
-	E05.478.605.492.300 Flocculation Tests
-	E05.478.605.492.350 Immunodiffusion
-	E05.478.605.492.350.350 Immunelectrophoresis
-	E05.478.605.492.350.350.150 Counterimmunelectrophoresis
-	E05.478.605.492.350.350.350 Immunelectrophoresis, Two-Dimensional
-	E05.478.605.825 Radioimmunoprecipitation Assay
-	E05.478.610 Immunosuppression
-	E05.478.610.310 Desensitization, Immunologic
-	E05.478.610.310.500 Sublingual Immunotherapy
-	E05.478.610.500 Graft Enhancement, Immunologic
-	E05.478.610.570 Lymphocyte Depletion
-	E05.478.610.800 Transplantation Conditioning
-	E05.478.805 Radioimmunodetection
-	E05.481 In Vitro Techniques
-	E05.481.500 Culture Techniques
-	E05.481.500.124 Axenic Culture
-	E05.481.500.249 Cell Culture Techniques

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E05.481.500.249.124 Batch Cell Culture Techniques
-	E05.481.500.249.500 Primary Cell Culture
-	E05.481.500.311 Cell Engineering
-	E05.481.500.311.249 Metabolic Engineering
-	E05.481.500.311.500 Tissue Engineering
-	E05.481.500.374 Coculture Techniques
-	E05.481.500.468 Embryo Culture Techniques
-	E05.481.500.484 Organ Culture Techniques
-	E05.481.500.550 Plant Somatic Embryogenesis Techniques
-	E05.481.500.617 Tissue Culture Techniques
-	E05.484 Indicator Dilution Techniques
-	E05.484.250 Dye Dilution Technique
-	E05.484.650 Radioisotope Dilution Technique
-	E05.484.750 Thermodilution
-	E05.490 Interferometry
-	E05.490.630 Microscopy, Interference
-	E05.490.630.569 Microscopy, Phase-Contrast
-	E05.490.815 Moire Topography
-	E05.497 Intubation
-	E05.497.412 Intubation, Gastrointestinal
-	E05.497.578 Intubation, Intratracheal
-	E05.497.578.475 Laryngeal Masks
-	E05.516 Ischemic Preconditioning
-	E05.516.325 Ischemic Preconditioning, Myocardial
-	E05.519 Isolated Heart Preparation
-	E05.522 Isotope Labeling
-	E05.540 Magnetometry
-	E05.540.249 Magnetocardiography
-	E05.540.500 Magnetoencephalography
-	E05.559 Manometry
-	E05.559.740 Rhinomanometry
-	E05.570 Materials Testing
-	E05.581 Methods
-	E05.581.249 Observation
-	E05.581.500 Research Design
-	E05.581.500.149 Control Groups

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	E05.581.500.300	Double-Blind Method
-	E05.581.500.400	Early Termination of Clinical Trials
-	E05.581.500.501	Meta-Analysis as Topic
New Heading	<b>E05.581.500.501.500</b>	<b>Network Meta-Analysis</b>
-	E05.581.500.653	Patient Selection
-	E05.581.500.805	Random Allocation
-	E05.581.500.902	Sample Size
-	E05.581.500.902.500	Numbers Needed To Treat
-	E05.588	Microchip Analytical Procedures
-	E05.588.465	Microfluidic Analytical Techniques
-	E05.588.465.340	Electrophoresis, Microchip
-	E05.588.570	Microarray Analysis
-	E05.588.570.660	Oligonucleotide Array Sequence Analysis
-	E05.588.570.700	Protein Array Analysis
-	E05.588.570.850	Tissue Array Analysis
-	E05.591	Micromanipulation
-	E05.591.560	Microdissection
-	E05.591.560.500	Laser Capture Microdissection
-	E05.591.570	Microinjections
-	E05.591.580	Microsurgery
-	E05.595	Microscopy
-	E05.595.185	Intravital Microscopy
-	E05.595.185.250	Dermoscopy
-	E05.595.185.500	Microscopic Angioscopy
-	E05.595.370	Microscopy, Acoustic
-	E05.595.395	Microscopy, Confocal
-	E05.595.395.500	Laser Scanning Cytometry
-	E05.595.402	Microscopy, Electron
-	E05.595.402.150	Cryoelectron Microscopy
-	E05.595.402.250	Electron Probe Microanalysis
-	E05.595.402.541	Microscopy, Electron, Scanning
-	E05.595.402.580	Microscopy, Electron, Transmission
-	E05.595.402.580.239	Electron Microscope Tomography
-	E05.595.402.580.480	Microscopy, Electron, Scanning Transmission
-	E05.595.402.580.500	Microscopy, Energy-Filtering Transmission Electron

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E05.595.402.625 Microscopy, Immunoelectron
-	E05.595.458 Microscopy, Fluorescence
-	E05.595.458.500 Microscopy, Fluorescence, Multiphoton
-	E05.595.513 Microscopy, Interference
-	E05.595.513.569 Microscopy, Phase-Contrast
-	E05.595.624 Microscopy, Polarization
-	E05.595.666 Microscopy, Scanning Probe
-	E05.595.666.400 Microscopy, Atomic Force
-	E05.595.666.450 Microscopy, Electrochemical, Scanning
-	E05.595.666.500 Microscopy, Scanning Tunneling
-	E05.595.680 Microscopy, Ultraviolet
-	E05.595.690 Microscopy, Video
-	E05.595.744 Nuclear Microscopy
-	E05.595.799 Photomicrography
New Heading	<b>E05.595.899 Single Molecule Imaging</b>
-	E05.598 Models, Animal
-	E05.598.500 Disease Models, Animal
-	E05.598.500.249 Arthritis, Experimental
-	E05.598.500.374 Diabetes Mellitus, Experimental
-	E05.598.500.468 Liver Cirrhosis, Experimental
-	E05.598.500.496 Neoplasms, Experimental
-	E05.598.500.496.500 Leukemia, Experimental
-	E05.598.500.496.750 Liver Neoplasms, Experimental
-	E05.598.500.496.843 Mammary Neoplasms, Experimental
-	E05.598.500.496.937 Melanoma, Experimental
-	E05.598.500.496.968 Sarcoma, Experimental
-	E05.598.500.500 Nervous System Autoimmune Disease, Experimental
-	E05.598.500.500.500 Encephalomyelitis, Autoimmune, Experimental
-	E05.598.500.500.750 Myasthenia Gravis, Autoimmune, Experimental
-	E05.598.500.500.875 Neuritis, Autoimmune, Experimental
-	E05.598.500.750 Radiation Injuries, Experimental
-	E05.598.750 Isolated Heart Preparation
-	E05.599 Models, Theoretical
-	E05.599.125 Fractals
-	E05.599.250 Fuzzy Logic

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E05.599.395 Models, Biological
-	E05.599.395.080 Disease Models, Animal
-	E05.599.395.161 Models, Cardiovascular
-	E05.599.395.397 Models, Genetic
-	E05.599.395.500 Models, Immunological
-	E05.599.395.642 Models, Neurological
-	E05.599.395.642.550 Neurolinguistic Programming
-	E05.599.395.821 Patient-Specific Modeling
-	E05.599.495 Models, Chemical
-	E05.599.545 Models, Educational
-	E05.599.595 Models, Molecular
-	E05.599.595.249 Molecular Docking Simulation
-	E05.599.595.500 Molecular Dynamics Simulation
-	E05.599.645 Models, Nursing
-	E05.599.670 Models, Organizational
-	E05.599.695 Models, Psychological
-	E05.599.695.550 Neurolinguistic Programming
-	E05.599.835 Models, Statistical
-	E05.599.835.500 Likelihood Functions
-	E05.599.835.750 Linear Models
-	E05.599.835.875 Logistic Models
-	E05.599.835.890 Models, Economic
-	E05.599.835.890.500 Models, Econometric
-	E05.599.835.895 Nomograms
-	E05.599.835.900 Proportional Hazards Models
-	E05.599.850 Nonlinear Dynamics
-	E05.601 Molecular Probe Techniques
-	E05.601.043 Biosensing Techniques
-	E05.601.043.700 Surface Plasmon Resonance
-	E05.601.085 Biotinylation
-	E05.601.100 Blotting, Northern
-	E05.601.150 Blotting, Southern
-	E05.601.225 Blotting, Southwestern
-	E05.601.262 Blotting, Western
-	E05.601.262.500 Blotting, Far-Western
-	E05.601.300 Branched DNA Signal Amplification Assay



## MeSH Tree Changes for 2017

Type	Tree - heading
-	E05.601.470                      Immunoassay
-	E05.601.470.320                      Immunoblotting
-	E05.601.470.320.200                      Blotting, Western
-	E05.601.470.320.200.500                      Blotting, Far-Western
-	E05.601.470.350                      Immunoenzyme Techniques
-	E05.601.470.350.170                      Enzyme-Linked Immunosorbent Assay
-	E05.601.470.350.170.500                      Enzyme-Linked Immunospot Assay
-	E05.601.470.350.180                      Enzyme Multiplied Immunoassay Technique
-	E05.601.470.380                      Immunosorbent Techniques
-	E05.601.470.380.360                      Enzyme-Linked Immunosorbent Assay
-	E05.601.470.380.360.500                      Enzyme-Linked Immunospot Assay
-	E05.601.470.380.810                      Radioallergosorbent Test
-	E05.601.470.380.825                      Radioimmunoprecipitation Assay
-	E05.601.470.380.830                      Radioimmunosorbent Test
-	E05.601.470.639                      Radioimmunoassay
-	E05.601.470.639.405                      Immunoradiometric Assay
-	E05.601.470.639.810                      Radioallergosorbent Test
-	E05.601.470.639.825                      Radioimmunoprecipitation Assay
-	E05.601.470.639.830                      Radioimmunosorbent Test
-	E05.601.555                      Molecular Imaging
New Heading	<b>E05.601.555.500                      Single Molecule Imaging</b>
-	E05.601.640                      Oligonucleotide Array Sequence Analysis
-	E05.601.680                      Protein Array Analysis
-	E05.601.690                      Protein Interaction Mapping
-	E05.601.690.149                      Blotting, Far-Western
-	E05.601.690.300                      Epitope Mapping
-	E05.601.690.650                      Two-Hybrid System Techniques
-	E05.601.700                      Random Amplified Polymorphic DNA Technique
-	E05.601.870                      Two-Hybrid System Techniques
-	E05.624                      Neoplasm Transplantation
-	E05.624.850                      Xenograft Model Antitumor Assays
-	E05.624.850.830                      Subrenal Capsule Assay
-	E05.629                      Neuroimaging
-	E05.629.750                      Diffusion Tensor Imaging
-	E05.629.875                      Functional Neuroimaging

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	E05.629.875.500	Brain Mapping
New Tree	E05.629.875.500.500	Connectome
-	E05.629.937	Neuroradiography
-	E05.629.937.180	Cerebral Angiography
-	E05.629.937.190	Cerebral Ventriculography
-	E05.629.937.260	Echoencephalography
-	E05.629.937.260.850	Ultrasonography, Doppler, Transcranial
-	E05.629.937.505	Myelography
-	E05.629.937.620	Pneumoencephalography
-	E05.635	Neuromuscular Blockade
-	E05.638	Olfactometry
-	E05.642	Optical Imaging
-	E05.642.124	Narrow Band Imaging
-	E05.642.249	Tomography, Optical
-	E05.642.249.500	Tomography, Optical Coherence
-	E05.642.500	Transillumination
-	E05.650	Optical Tweezers
-	E05.654	Oscillometry
New Heading	<b>E05.657</b>	<b>Oxygen Radical Absorbance Capacity</b>
-	E05.660	Parabiosis
-	E05.660.365	Cross Circulation
-	E05.680	Perfusion
-	E05.680.730	Reperfusion
-	E05.680.730.600	Myocardial Reperfusion
-	E05.696	Photoacoustic Techniques
-	E05.712	Photography
-	E05.712.315	Photography, Dental
-	E05.712.657	Time-Lapse Imaging
-	E05.723	Physical Stimulation
-	E05.723.136	Acoustic Stimulation
-	E05.723.402	Electric Stimulation
-	E05.723.402.201	Chronaxy
-	E05.723.402.403	Electroshock
-	E05.723.729	Photic Stimulation

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	E05.734	Physiognomy
-	E05.760	Preservation, Biological
-	E05.760.156	Cryopreservation
-	E05.760.156.260	Freeze Drying
-	E05.760.156.260.270	Freeze Substitution
-	E05.760.643	Refrigeration
-	E05.760.833	Tissue Preservation
-	E05.760.833.230	Blood Preservation
-	E05.760.833.230.500	Blood Safety
-	E05.760.833.445	Cold Ischemia
-	E05.760.833.660	Organ Preservation
-	E05.760.833.890	Semen Preservation
-	E05.790	Protein Refolding
-	E05.793	Protein Unfolding
-	E05.793.500	Protein Denaturation
-	E05.796	Psychological Techniques
-	E05.796.112	Behavior Observation Techniques
-	E05.796.332	Galvanic Skin Response
-	E05.796.817	Reaction Time
-	E05.796.817.559	Refractory Period, Psychological
-	E05.796.908	Signal Detection, Psychological
-	E05.799	Radiometry
-	E05.799.256	Autoradiography
-	E05.799.513	Radiation Dosage
-	E05.799.513.500	Dose-Response Relationship, Radiation
-	E05.799.638	Radiation Monitoring
-	E05.799.638.231	Body Burden
-	E05.799.638.420	Film Dosimetry
New Heading	<b>E05.799.638.511</b>	<b>In Vivo Dosimetry</b>
-	E05.799.638.602	Optically Stimulated Luminescence Dosimetry
-	E05.799.638.785	Thermoluminescent Dosimetry
-	E05.799.669	Radiometric Dating
-	E05.799.700	Scintillation Counting
-	E05.799.801	Spectrometry, Gamma
-	E05.799.830	Spectrometry, X-Ray Emission

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E05.799.830.360                      Electron Probe Microanalysis
-	E05.799.890                              Sun Protection Factor
-	E05.799.950                              Whole-Body Counting
-	E05.820                                    Reproductive Techniques
-	E05.820.150                              Breeding
-	E05.820.150.360                        Estrus Detection
-	E05.820.150.370                        Estrus Synchronization
-	E05.820.150.390                        Hybridization, Genetic
-	E05.820.150.520                        Inbreeding
-	E05.820.150.760                        Selective Breeding
-	E05.820.180                              Cloning, Organism
-	E05.820.240                              Embryo Disposition
-	E05.820.390                              Fallopian Tube Patency Tests
-	E05.820.540                              Nuclear Transfer Techniques
-	E05.820.690                              Ovulation Detection
-	E05.820.745                              Ovulation Prediction
-	E05.820.800                              Reproductive Techniques, Assisted
New Heading	<b>E05.820.800.250                        Donor Conception</b>
-	E05.820.800.500                        Embryo Transfer
-	E05.820.800.500.500                    Single Embryo Transfer
-	E05.820.800.625                        Fertility Preservation
-	E05.820.800.750                        Fertilization in Vitro
-	E05.820.800.750.350                    Mitochondrial Replacement Therapy
-	E05.820.800.750.700                    Sperm Injections, Intracytoplasmic
-	E05.820.800.875                        Gamete Intrafallopian Transfer
-	E05.820.800.906                        In Vitro Oocyte Maturation Techniques
-	E05.820.800.937                        Insemination, Artificial
-	E05.820.800.937.515                    Insemination, Artificial, Heterologous
-	E05.820.800.937.525                    Insemination, Artificial, Homologous
-	E05.820.800.968                        Oocyte Donation
-	E05.820.800.976                        Oocyte Retrieval
-	E05.820.800.984                        Ovulation Induction
-	E05.820.800.984.500                    Superovulation
-	E05.820.800.986                        Posthumous Conception
-	E05.820.800.988                        Sperm Retrieval

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E05.820.800.992 Zygote Intrafallopian Transfer
-	E05.830 Rheology
-	E05.830.250 Hemorheology
-	E05.830.500 Laser-Doppler Flowmetry
-	E05.830.583 Lubrication
-	E05.830.666 Microfluidics
-	E05.830.666.500 Electrowetting
-	E05.848 Sonication
-	E05.855 Sound Spectrography
-	E05.873 Stereotaxic Techniques
-	E05.873.249 Neuronavigation
-	E05.873.500 Radiosurgery
-	E05.885 Substance Abuse Detection
-	E05.916 Technology, Pharmaceutical
Old Tree	<b>E05.916.039 Chemistry Techniques, Synthetic</b>
Old Tree	<b>E05.916.039.124 Click Chemistry</b>
Old Tree	<b>E05.916.039.312 Combinatorial Chemistry Techniques</b>
Old Tree	<b>E05.916.039.312.500 SELEX Aptamer Technique</b>
Old Tree	<b>E05.916.039.406 Cycloaddition Reaction</b>
Old Tree	<b>E05.916.039.500 Solid-Phase Synthesis Techniques</b>
-	E05.916.250 Dosage Forms
-	E05.916.270 Drug Compounding
-	E05.916.290 Drug Incompatibility
-	E05.916.310 Drug Labeling
-	E05.916.320 Drug Packaging
-	E05.916.330 Drug Stability
-	E05.916.350 Drug Storage
-	E05.916.360 Enzyme Stability
-	E05.916.680 High-Throughput Screening Assays
-	E05.920 Technology, Radiologic
-	E05.920.700 Teleradiology
-	E05.925 Telemetry
-	E05.925.500 Remote Sensing Technology
-	E05.925.500.500 Satellite Imagery
-	E05.927 Therapeutic Irrigation
-	E05.927.100 Bronchoalveolar Lavage

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E05.927.100.500 Bronchoalveolar Lavage Fluid
-	E05.927.441 Gastric Lavage
-	E05.927.573 Nasal Lavage
-	E05.927.573.500 Nasal Lavage Fluid
-	E05.927.705 Peritoneal Lavage
-	E05.927.852 Vaginal Douching
-	E05.933 Thermometry
-	E05.933.500 Thermography
-	E05.940 Toxicity Tests
-	E05.940.150 Carcinogenicity Tests
-	E05.940.245 Cytotoxicity Tests, Immunologic
-	E05.940.350 Inhibitory Concentration 50
-	E05.940.402 Lethal Dose 50
-	E05.940.481 Maximum Tolerated Dose
-	E05.940.560 Mutagenicity Tests
-	E05.940.560.150 Comet Assay
-	E05.940.600 No-Observed-Adverse-Effect Level
-	E05.940.790 Toxicity Tests, Acute
-	E05.940.790.500 Skin Irritancy Tests
-	E05.940.800 Toxicity Tests, Chronic
-	E05.940.850 Toxicity Tests, Subacute
-	E05.940.900 Toxicity Tests, Subchronic
-	E05.944 Trauma Severity Indices
-	E05.944.250 Abbreviated Injury Scale
-	E05.944.500 Glasgow Coma Scale
-	E05.944.510 Glasgow Outcome Scale
-	E05.944.600 Injury Severity Score
-	E05.959 Wavelet Analysis
-	E05.974 Weightlessness Countermeasures
-	E05.977 Weightlessness Simulation
-	E05.977.400 Hindlimb Suspension
-	E05.978 Weights and Measures
-	E05.978.155 Calibration
-	E05.978.507 International System of Units
-	E05.978.539 Metric System
-	E05.978.673 Quartz Crystal Microbalance Techniques

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E05.978.808 Reference Standards
-	E05.978.808.249 Fiducial Markers
-	E05.978.808.500 Growth Charts
-	E05.978.810 Reference Values
-	E05.979 Whole Body Imaging
-	E05.980 Whole-Body Irradiation
-	E06 Dentistry
-	E06.020 Air Abrasion, Dental
-	E06.045 Anesthesia, Dental
-	E06.045.481 Hypnosis, Dental
-	E06.070 Dental Atraumatic Restorative Treatment
-	E06.095 Dental Bonding
-	E06.095.170 Cementation
-	E06.095.585 Light-Curing of Dental Adhesives
-	E06.095.850 Self-Curing of Dental Resins
-	E06.170 Dental Care
-	E06.170.100 Dental Care for Aged
-	E06.170.152 Dental Care for Children
-	E06.170.205 Dental Care for Chronically Ill
-	E06.170.310 Dental Care for Disabled
-	E06.178 Dental Debonding
-	E06.186 Dental Equipment
-	E06.186.104 Curing Lights, Dental
-	E06.186.210 Dental Articulators
-	E06.186.250 Dental Devices, Home Care
-	E06.186.376 Dental High-Speed Equipment
-	E06.186.501 Dental Instruments
-	E06.186.501.630 Matrix Bands
-	E06.186.670 Rubber Dams
-	E06.208 Dental Health Surveys
-	E06.208.250 Dental Plaque Index
-	E06.208.266 DMF Index
-	E06.208.421 Index of Orthodontic Treatment Need
-	E06.208.576 Oral Hygiene Index
-	E06.208.720 Periodontal Index
-	E06.216 Dental High-Speed Technique

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	E06.261	Dental Models
-	E06.276	Dental Occlusion
-	E06.276.125	Bite Force
-	E06.276.293	Dental Occlusion, Balanced
-	E06.276.459	Dental Occlusion, Centric
-	E06.276.459.733	Vertical Dimension
-	E06.292	Dental Pins
-	E06.298	Dental Polishing
-	E06.308	Dental Stress Analysis
-	E06.323	Dentistry, Operative
-	E06.323.115	Crown Lengthening
-	E06.323.225	Dental Cavity Lining
-	E06.323.400	Dental Restoration Failure
-	E06.323.428	Dental Restoration, Permanent
-	E06.323.428.187	Dental Restoration Repair
-	E06.323.428.275	Inlays
-	E06.323.528	Dental Restoration, Temporary
-	E06.323.764	Dental Marginal Adaptation
-	E06.342	Diagnosis, Oral
-	E06.342.250	Dental Caries Activity Tests
-	E06.342.377	Dental Pulp Test
-	E06.342.488	Photography, Dental
-	E06.342.764	Radiography, Dental
-	E06.342.764.142	Age Determination by Teeth
-	E06.342.764.600	Radiography, Bitewing
-	E06.342.764.716	Radiography, Dental, Digital
-	E06.342.764.750	Radiography, Panoramic
-	E06.342.764.793	Sialography
-	E06.377	Electrogalvanism, Intraoral
-	E06.397	Endodontics
-	E06.397.102	Apicoectomy
-	E06.397.345	Dental Implantation, Endosseous, Endodontic
-	E06.397.370	Dental Pulp Capping
-	E06.397.633	Pulpectomy
-	E06.397.695	Pulpotomy
-	E06.397.778	Root Canal Therapy



## MeSH Tree Changes for 2017

Type	Tree - heading
-	E06.397.778.188 Apexification
-	E06.397.778.377 Dental Pulp Devitalization
-	E06.397.778.778 Root Canal Obturation
-	E06.397.778.778.648 Retrograde Obturation
-	E06.397.778.889 Root Canal Preparation
-	E06.397.898 Tooth Replantation
-	E06.420 Esthetics, Dental
-	E06.420.500 Enamel Microabrasion
-	E06.420.750 Tooth Bleaching
-	E06.470 Infection Control, Dental
-	E06.520 Jaw Relation Record
-	E06.520.200 Centric Relation
-	E06.584 Mouth Rehabilitation
-	E06.603 Myofunctional Therapy
-	E06.623 Odontometry
-	E06.623.500 Age Determination by Teeth
-	E06.640 Oral Medicine
-	E06.645 Oral Surgical Procedures
-	E06.645.100 Apicoectomy
-	E06.645.350 Gingivectomy
-	E06.645.355 Gingivoplasty
-	E06.645.380 Glossectomy
-	E06.645.410 Guided Tissue Regeneration, Periodontal
-	E06.645.440 Jaw Fixation Techniques
-	E06.645.500 Mandibular Advancement
-	E06.645.510 Maxillofacial Prosthesis Implantation
-	E06.645.510.500 Mandibular Prosthesis Implantation
-	E06.645.550 Oral Surgical Procedures, Preprosthetic
-	E06.645.550.100 Alveolar Ridge Augmentation
-	E06.645.550.110 Alveolectomy
-	E06.645.550.120 Alveoloplasty
-	E06.645.550.280 Dental Implantation
-	E06.645.550.280.280 Dental Implantation, Endosseous
-	E06.645.550.280.280.150 Blade Implantation
-	E06.645.550.280.280.280 Dental Implantation, Endosseous, Endodontic
-	E06.645.550.280.280.450 Immediate Dental Implant Loading

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E06.645.550.280.290                      Dental Implantation, Subperiosteal
-	E06.645.550.800                              Vestibuloplasty
-	E06.645.562                                    Orthognathic Surgical Procedures
-	E06.645.562.124                              Alveolar Bone Grafting
-	E06.645.562.249                              Genioplasty
-	E06.645.562.374                              Mandibular Osteotomy
-	E06.645.562.500                              Mandibular Reconstruction
-	E06.645.562.750                              Maxillary Osteotomy
-	E06.645.575                                    Osteotomy, Le Fort
-	E06.645.637                                    Osteotomy, Sagittal Split Ramus
-	E06.645.668                                    Sinus Floor Augmentation
-	E06.645.700                                    Tooth Extraction
-	E06.645.700.680                              Serial Extraction
-	E06.645.710                                    Tooth Replantation
-	E06.658    Orthodontics
-	E06.658.224                                    Dental Marginal Adaptation
-	E06.658.280                                    Mandibular Advancement
-	E06.658.337                                    Orthodontic Anchorage Procedures
-	E06.658.450                                    Orthodontic Appliance Design
-	E06.658.453                                    Orthodontic Appliances
-	E06.658.453.510                              Occlusal Splints
-	E06.658.453.560                              Orthodontic Appliances, Functional
-	E06.658.453.560.100                        Activator Appliances
-	E06.658.453.578                              Orthodontic Appliances, Removable
-	E06.658.453.578.100                        Activator Appliances
-	E06.658.453.578.360                        Extraoral Traction Appliances
-	E06.658.453.590                              Orthodontic Brackets
-	E06.658.453.637                              Orthodontic Retainers
-	E06.658.453.684                              Orthodontic Wires
-	E06.658.578                                    Orthodontics, Corrective
-	E06.658.578.099                              Crown Lengthening
-	E06.658.578.200                              Occlusal Adjustment
-	E06.658.578.225                              Orthodontic Extrusion
-	E06.658.578.229                              Orthodontic Space Closure
-	E06.658.578.458                              Palatal Expansion Technique
-	E06.658.578.836                              Tooth Movement

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E06.658.578.836 Tooth Movement Techniques
-	E06.658.641 Orthodontics, Interceptive
-	E06.658.641.890 Serial Extraction
-	E06.658.703 Orthodontics, Preventive
-	E06.658.703.675 Space Maintenance, Orthodontic
-	E06.685 Pathology, Oral
-	E06.721 Periodontics
-	E06.721.189 Dental Prophylaxis
-	E06.721.189.350 Dental Scaling
-	E06.721.189.350.650 Root Planing
-	E06.721.189.675 Periodontal Debridement
-	E06.721.321 Gingivectomy
-	E06.721.384 Gingivoplasty
-	E06.721.485 Guided Tissue Regeneration, Periodontal
-	E06.721.595 Periodontal Dressings
-	E06.721.658 Periodontal Index
-	E06.721.721 Periodontal Prosthesis
-	E06.721.721.680 Periodontal Splints
-	E06.721.874 Subgingival Curettage
-	E06.721.874.650 Root Planing
-	E06.761 Preventive Dentistry
-	E06.761.227 Dental Prophylaxis
-	E06.761.227.350 Dental Scaling
-	E06.761.227.350.650 Root Planing
-	E06.761.382 Fluoridation
-	E06.761.599 Mouth Protectors
-	E06.761.726 Oral Hygiene
-	E06.761.726.292 Dental Devices, Home Care
-	E06.761.726.794 Toothbrushing
-	E06.780 Prosthodontics
-	E06.780.250 Dental Casting Technique
-	E06.780.314 Dental Implantation
-	E06.780.314.310 Dental Implantation, Endosseous
-	E06.780.314.310.340 Blade Implantation
-	E06.780.314.310.360 Dental Implantation, Endosseous, Endodontic
-	E06.780.314.310.450 Immediate Dental Implant Loading

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E06.780.314.482                      Dental Implantation, Subperiosteal
-	E06.780.346                              Dental Prosthesis
-	E06.780.346.250                        Crowns
-	E06.780.346.250.500                    Post and Core Technique
-	E06.780.346.500                        Dental Abutments
-	E06.780.346.562                        Dental Clasps
-	E06.780.346.593                        Dental Implants
-	E06.780.346.593.185                    Dental Implants, Single-Tooth
-	E06.780.346.625                        Dental Prosthesis Design
-	E06.780.346.625.500                    Dental Implant-Abutment Design
-	E06.780.346.630                        Dental Prosthesis, Implant-Supported
-	E06.780.346.656                        Dental Prosthesis Repair
-	E06.780.346.687                        Dental Prosthesis Retention
-	E06.780.346.725                        Dental Restoration Failure
-	E06.780.346.737                        Dental Restoration, Permanent
-	E06.780.346.737.112                    Dental Restoration Repair
-	E06.780.346.737.125                    Dental Restoration Wear
-	E06.780.346.737.500                    Inlays
-	E06.780.346.740                        Dental Restoration, Temporary
-	E06.780.346.750                        Dental Veneers
-	E06.780.346.760                        Dentures
-	E06.780.346.760.281                    Denture Bases
-	E06.780.346.760.290                    Denture, Complete
-	E06.780.346.760.290.254                Denture, Complete, Immediate
-	E06.780.346.760.290.379                Denture, Complete, Lower
-	E06.780.346.760.290.504                Denture, Complete, Upper
-	E06.780.346.760.300                    Denture Design
-	E06.780.346.760.357                    Denture Liners
-	E06.780.346.760.395                    Denture, Overlay
-	E06.780.346.760.433                    Denture, Partial
-	E06.780.346.760.433.271                Denture, Partial, Fixed
-	E06.780.346.760.433.271.260            Denture, Partial, Fixed, Resin-Bonded
-	E06.780.346.760.433.342                Denture, Partial, Immediate
-	E06.780.346.760.433.413                Denture, Partial, Removable
-	E06.780.346.760.433.484                Denture, Partial, Temporary
-	E06.780.346.760.433.555                Denture Precision Attachment

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E06.780.346.760.471 Denture Rebasing
-	E06.780.346.760.500 Denture Repair
-	E06.780.346.760.550 Denture Retention
-	E06.780.346.850 Palatal Obturators
-	E06.780.346.875 Tooth, Artificial
-	E06.780.620 Dental Marginal Adaptation
-	E06.780.895 Tissue Conditioning (Dental)
-	E06.892 Surgery, Oral
-	E06.892.500 Orthognathic Surgery
-	E06.912 Technology, Dental
-	E06.912.115 Dental Casting Technique
-	E06.912.130 Dental Impression Technique
-	E06.912.145 Dental Prosthesis Design
-	E06.912.145.750 Dental Implant-Abutment Design
-	E06.912.160 Dental Prosthesis Repair
-	E06.912.190 Dental Soldering
-	E06.912.250 Denture Design
-	E06.912.275 Denture Identification Marking
-	E06.912.280 Denture Rebasing
-	E06.912.285 Denture Repair
-	E06.912.480 Gingival Retraction Techniques
-	E06.912.675 Orthodontic Appliance Design
-	E06.931 Tooth Preparation
-	E06.931.325 Dental Cavity Preparation
-	E06.931.475 Dental Etching
-	E06.931.475.111 Acid Etching, Dental
-	E06.931.625 Root Canal Preparation
-	E06.931.750 Tooth Preparation, Prosthodontic
-	E06.950 Tooth Remineralization
-	E07 Equipment and Supplies
-	E07.039 Air Filters
-	E07.079 Atmosphere Exposure Chambers
-	E07.101 Bandages
-	E07.101.036 Athletic Tape
-	E07.101.074 Bandages, Hydrocolloid
-	E07.101.150 Biological Dressings

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	E07.101.400	Compression Bandages
-	E07.101.400.500	Stockings, Compression
-	E07.101.650	Occlusive Dressings
New Heading	<b>E07.108</b>	<b>Bathroom Equipment</b>
-	E07.115	Bioreactors
-	E07.115.600	Photobioreactors
-	E07.123	Capillary Tubing
-	E07.132	Catheters
New Heading	<b>E07.132.125</b>	<b>Cannula</b>
-	E07.132.249	Catheter Obstruction
-	E07.132.500	Catheters, Indwelling
-	E07.132.625	Urinary Catheters
-	E07.132.750	Vascular Access Devices
-	E07.132.750.249	Cardiac Catheters
-	E07.132.750.500	Central Venous Catheters
-	E07.161	Compressed Air
-	E07.190	Contraceptive Devices
-	E07.190.250	Contraceptive Devices, Female
-	E07.190.250.260	Condoms, Female
-	E07.190.250.520	Intrauterine Devices
-	E07.190.250.520.520	Intrauterine Devices, Medicated
-	E07.190.250.520.520.300	Intrauterine Devices, Copper
-	E07.190.270	Contraceptive Devices, Male
-	E07.190.270.150	Condoms
-	E07.206	Culture Media
-	E07.206.250	Culture Media, Conditioned
-	E07.206.255	Culture Media, Serum-Free
-	E07.206.627	Tissue Scaffolds
-	E07.222	Dental Equipment
-	E07.222.104	Curing Lights, Dental
-	E07.222.210	Dental Articulators
-	E07.222.250	Dental Devices, Home Care
-	E07.222.376	Dental High-Speed Equipment
-	E07.222.501	Dental Instruments

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E07.222.501.630 Matrix Bands
-	E07.222.670 Rubber Dams
-	E07.230 Diagnostic Equipment
-	E07.230.200 Clinical Alarms
-	E07.230.210 Electronic Nose
-	E07.230.220 Endoscopes
-	E07.230.220.040 Angioscopes
-	E07.230.220.060 Arthroscopes
-	E07.230.220.090 Bronchoscopes
-	E07.230.220.170 Colposcopes
-	E07.230.220.180 Culdoscopes
-	E07.230.220.190 Cystoscopes
-	E07.230.220.260 Endoscopes, Gastrointestinal
-	E07.230.220.260.140 Capsule Endoscopes
-	E07.230.220.260.160 Colonoscopes
-	E07.230.220.260.160.800 Sigmoidoscopes
-	E07.230.220.260.200 Duodenoscopes
-	E07.230.220.260.260 Esophagoscopes
-	E07.230.220.260.320 Gastrosopes
-	E07.230.220.260.680 Proctoscopes
-	E07.230.220.280 Fetoscopes
-	E07.230.220.390 Hysteroscopes
-	E07.230.220.520 Laparoscopes
-	E07.230.220.525 Laryngoscopes
-	E07.230.220.565 Mediastinoscopes
-	E07.230.220.707 Neuroendoscopes
-	E07.230.220.850 Thoracoscopes
-	E07.230.220.895 Ureteroscopes
New Heading	<b>E07.230.300 Fitness Trackers</b>
-	E07.230.380 Flowmeters
-	E07.230.460 Muscle Strength Dynamometer
-	E07.230.540 Ophthalmoscopes
-	E07.230.540.500 Retinoscopes
New Tree	<b>E07.230.540.750 Slit Lamp</b>

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	E07.230.550	Otosopes
-	E07.230.740	Sphygmomanometers
-	E07.230.740.100	Blood Pressure Monitors
-	E07.230.750	Stethoscopes
-	E07.241	Diffusion Chambers, Culture
-	E07.252	Disposable Equipment
-	E07.278	Durable Medical Equipment
-	E07.305	Electrical Equipment and Supplies
-	E07.305.061	Amplifiers, Electronic
-	E07.305.076	Brain-Computer Interfaces
-	E07.305.092	Cathode Ray Tube
-	E07.305.124	Electric Power Supplies
-	E07.305.124.150	Bioelectric Energy Sources
-	E07.305.245	Electric Wiring
-	E07.305.250	Electrodes
-	E07.305.250.159	Defibrillators
-	E07.305.250.159.175	Defibrillators, Implantable
-	E07.305.250.319	Electrodes, Implanted
-	E07.305.250.319.175	Defibrillators, Implantable
-	E07.305.250.319.381	Implantable Neurostimulators
-	E07.305.250.319.381.500	Neural Prostheses
-	E07.305.250.319.381.500.249	Auditory Brain Stem Implants
-	E07.305.250.319.381.500.500	Cochlear Implants
-	E07.305.250.471	Ion-Selective Electrodes
-	E07.305.250.500	Microelectrodes
-	E07.305.250.750	Pacemaker, Artificial
-	E07.305.250.750.500	Cardiac Resynchronization Therapy Devices
-	E07.305.296	Electronic Nose
-	E07.305.343	Micro-Electrical-Mechanical Systems
-	E07.305.343.500	Lab-On-A-Chip Devices
Old Tree	<b>E07.305.437</b>	<b>Power Plants</b>
Old Tree	<b>E07.305.437.600</b>	<b>Nuclear Power Plants</b>
-	E07.305.625	Semiconductors
-	E07.305.625.714	Transistors, Electronic
-	E07.305.812	Transducers
-	E07.305.812.901	Transducers, Pressure



## MeSH Tree Changes for 2017

Type	Tree - heading	
-	E07.315	Emergency Medical Tags
-	E07.325	Equipment and Supplies, Hospital
-	E07.325.068	Absorbent Pads
-	E07.325.068.124	Diapers, Adult
-	E07.325.068.249	Diapers, Infant
-	E07.325.068.500	Incontinence Pads
-	E07.325.137	Bedding and Linens
-	E07.325.137.750	Surgical Drapes
-	E07.325.220	Beds
-	E07.325.220.500	Stretchers
-	E07.325.409	Examination Tables
-	E07.325.473	Hospitals, Packaged
-	E07.325.569	Incubators, Infant
-	E07.325.662	Operating Tables
-	E07.325.755	Patient Isolators
-	E07.325.877	Surgical Attire
-	E07.325.877.500	Masks
-	E07.341	Exoskeleton Device
-	E07.357	Feminine Hygiene Products
-	E07.357.500	Menstrual Hygiene Products
-	E07.357.750	Vaginal Creams, Foams, and Jellies
-	E07.420	Gamma Cameras
-	E07.427	Gas Scavengers
-	E07.430	Gastric Balloon
-	E07.440	Gravity Suits
-	E07.451	Humidifiers
-	E07.461	Incubators
-	E07.461.487	Incubators, Infant
-	E07.490	Infant Equipment
-	E07.490.249	Child Restraint Systems
-	E07.490.500	Pacifiers
-	E07.505	Infusion Pumps
-	E07.505.254	Infusion Pumps, Implantable
-	E07.505.508	Insulin Infusion Systems
-	E07.515	Intermittent Pneumatic Compression Devices
-	E07.546	Medicine Chests

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	E07.553	Microbubbles
-	E07.560	Micropore Filters
-	E07.565	Microspheres
-	E07.595	Nanospheres
-	E07.605	Nebulizers and Vaporizers
-	E07.605.500	Dry Powder Inhalers
-	E07.605.750	Metered Dose Inhalers
-	E07.605.750.500	Inhalation Spacers
-	E07.612	Needles
-	E07.632	Optical Devices
-	E07.632.490	Lasers
-	E07.632.490.230	Lasers, Dye
-	E07.632.490.244	Lasers, Excimer
-	E07.632.490.367	Lasers, Gas
-	E07.632.490.480	Lasers, Semiconductor
-	E07.632.490.490	Lasers, Solid-State
-	E07.632.500	Lenses
-	E07.632.500.276	Contact Lenses
-	E07.632.500.276.360	Contact Lenses, Hydrophilic
-	E07.632.500.276.360.220	Contact Lenses, Extended-Wear
-	E07.632.500.300	Eyeglasses
-	E07.632.500.460	Lenses, Intraocular
-	E07.632.500.460.500	Phakic Intraocular Lenses
-	E07.632.750	Optical Fibers
-	E07.632.875	Telescopes
-	E07.652	Oxygenators
-	E07.652.582	Oxygenators, Membrane
Old Tree	<b>E07.662</b>	<b>Personal Protective Equipment</b>
Old Tree	<b>E07.662.250</b>	<b>Eye Protective Devices</b>
Old Tree	<b>E07.662.375</b>	<b>Head Protective Devices</b>
Old Tree	<b>E07.662.438</b>	<b>Masks</b>
Old Tree	<b>E07.662.500</b>	<b>Protective Clothing</b>
Old Tree	<b>E07.662.500.400</b>	<b>Gloves, Protective</b>
Old Tree	<b>E07.662.500.400.400</b>	<b>Gloves, Surgical</b>
Old Tree	<b>E07.662.500.700</b>	<b>Space Suits</b>
Old Tree	<b>E07.662.750</b>	<b>Respiratory Protective Devices</b>

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E07.671                      Phantoms, Imaging
-	E07.695                      Protheses and Implants
-	E07.695.025                  Absorbable Implants
-	E07.695.050                  Artificial Limbs
-	E07.695.075                  Auditory Brain Stem Implants
-	E07.695.100                  Bioprosthesis
-	E07.695.110                  Blood Vessel Prosthesis
-	E07.695.125                  Bone-Implant Interface
-	E07.695.140                  Breast Implants
-	E07.695.150                  Cochlear Implants
-	E07.695.185                  Dental Implants
-	E07.695.185.185              Dental Implants, Single-Tooth
-	E07.695.190                  Dental Prosthesis
-	E07.695.190.088              Crowns
-	E07.695.190.088.500          Post and Core Technique
-	E07.695.190.175              Dental Abutments
-	E07.695.190.180              Dental Clasps
-	E07.695.190.185              Dental Prosthesis, Implant-Supported
-	E07.695.190.190              Dental Restoration, Permanent
-	E07.695.190.190.250          Dental Restoration Repair
-	E07.695.190.190.350          Inlays
-	E07.695.190.192              Dental Restoration, Temporary
-	E07.695.190.196              Dental Veneers
-	E07.695.190.200              Dentures
-	E07.695.190.200.200          Denture Bases
-	E07.695.190.200.205          Denture, Complete
-	E07.695.190.200.205.205      Denture, Complete, Immediate
-	E07.695.190.200.205.210      Denture, Complete, Lower
-	E07.695.190.200.205.215      Denture, Complete, Upper
-	E07.695.190.200.210          Denture Liners
-	E07.695.190.200.215          Denture, Overlay
-	E07.695.190.200.220          Denture, Partial
-	E07.695.190.200.220.220      Denture, Partial, Fixed
-	E07.695.190.200.220.220.220      Denture, Partial, Fixed, Resin-Bonded
-	E07.695.190.200.220.225      Denture, Partial, Immediate
-	E07.695.190.200.220.230      Denture, Partial, Removable

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E07.695.190.200.220.235 Denture, Partial, Temporary
-	E07.695.190.200.220.245 Denture Precision Attachment
-	E07.695.190.600 Palatal Obturators
-	E07.695.190.610 Periodontal Prosthesis
-	E07.695.190.800 Tooth, Artificial
-	E07.695.202 Electrodes, Implanted
-	E07.695.202.175 Defibrillators, Implantable
-	E07.695.202.381 Implantable Neurostimulators
-	E07.695.202.381.500 Neural Prostheses
-	E07.695.202.381.500.249 Auditory Brain Stem Implants
-	E07.695.202.381.500.500 Cochlear Implants
-	E07.695.207 Embolic Protection Devices
-	E07.695.207.500 Vena Cava Filters
-	E07.695.225 Eye, Artificial
-	E07.695.237 Fiducial Markers
-	E07.695.250 Glaucoma Drainage Implants
-	E07.695.250.500 Molteno Implants
-	E07.695.300 Heart, Artificial
-	E07.695.300.300 Heart-Assist Devices
-	E07.695.310 Heart Valve Prosthesis
-	E07.695.340 Implants, Experimental
-	E07.695.370 Internal Fixators
-	E07.695.370.249 Bone Nails
-	E07.695.370.374 Bone Plates
-	E07.695.370.437 Bone Screws
-	E07.695.370.437.500 Pedicle Screws
-	E07.695.370.468 Bone Wires
-	E07.695.370.734 Suture Anchors
-	E07.695.400 Joint Prosthesis
-	E07.695.400.300 Elbow Prosthesis
-	E07.695.400.400 Hip Prosthesis
-	E07.695.400.410 Knee Prosthesis
-	E07.695.400.705 Metal-on-Metal Joint Prostheses
New Heading	<b>E07.695.400.852</b> <b>Shoulder Prosthesis</b>
-	E07.695.450 Larynx, Artificial

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	E07.695.460	Lenses, Intraocular
-	E07.695.510	Maxillofacial Prosthesis
-	E07.695.510.500	Mandibular Prosthesis
-	E07.695.540	Orbital Implants
-	E07.695.550	Ossicular Prosthesis
-	E07.695.610	Penile Prosthesis
-	E07.695.680	Prosthesis Design
-	E07.695.680.550	Prosthesis Coloring
New Heading	<b>E07.695.698</b>	<b>Punctal Plugs</b>
-	E07.695.715	Septal Occluder Device
-	E07.695.750	Stents
-	E07.695.750.500	Drug-Eluting Stents
-	E07.695.750.750	Self Expandable Metallic Stents
-	E07.695.752	Suburethral Slings
-	E07.695.800	Tissue Expansion Devices
-	E07.695.825	Tissue Scaffolds
-	E07.695.850	Urinary Sphincter, Artificial
-	E07.695.950	Visual Prosthesis
-	E07.700	Protective Devices
-	E07.700.100	Air Bags
-	E07.700.175	Child Restraint Systems
-	E07.700.250	Ear Protective Devices
Old Tree	<b>E07.700.260</b>	<b>Eye Protective Devices</b>
-	E07.700.380	Head Protective Devices
-	E07.700.500	Masks
-	E07.700.500.450	Laryngeal Masks
-	E07.700.510	Mosquito Nets
-	E07.700.510.500	Insecticide-Treated Bednets
-	E07.700.520	Mouth Protectors
New Tree	<b>E07.700.560</b>	<b>Personal Protective Equipment</b>
-	E07.700.600	Protective Clothing
-	E07.700.600.400	Gloves, Protective
-	E07.700.600.400.400	Gloves, Surgical
-	E07.700.600.700	Space Suits

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E07.700.700                      Respiratory Protective Devices
-	E07.700.800                      Seat Belts
-	E07.705                              Quantum Dots
-	E07.710                              Radiation Equipment and Supplies
-	E07.710.259                      Fiducial Markers
-	E07.710.520                      Lasers
-	E07.710.520.230                  Lasers, Dye
-	E07.710.520.244                  Lasers, Excimer
-	E07.710.520.367                  Lasers, Gas
-	E07.710.520.480                  Lasers, Semiconductor
-	E07.710.520.490                  Lasers, Solid-State
-	E07.710.600                      Nuclear Reactors
-	E07.710.600.500                  Nuclear Power Plants
-	E07.710.680                      Particle Accelerators
-	E07.710.680.250                  Cyclotrons
-	E07.710.680.700                  Synchrotrons
New Heading	<b>E07.710.725                      Radiation Dosimeters</b>
-	E07.710.770                      Radionuclide Generators
-	E07.715                              Radio Frequency Identification Device
-	E07.720                              Reagent Kits, Diagnostic
-	E07.720.720                      Reagent Strips
-	E07.796                              Self-Help Devices
-	E07.796.250                      Communication Aids for Disabled
-	E07.796.980                      Wheelchairs
-	E07.814                              Sensory Aids
-	E07.814.458                      Hearing Aids
-	E07.814.458.074                  Auditory Brain Stem Implants
-	E07.814.458.150                  Cochlear Implants
-	E07.858                              Surgical Equipment
-	E07.858.082                      Artificial Organs
-	E07.858.082.050                  Artificial Limbs
-	E07.858.082.212                  Bioartificial Organs
-	E07.858.082.374                  Heart, Artificial
-	E07.858.082.374.300              Heart-Assist Devices
-	E07.858.082.458                  Heart-Lung Machine

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E07.858.082.505 Infusion Pumps
-	E07.858.082.505.254 Infusion Pumps, Implantable
-	E07.858.082.505.508 Insulin Infusion Systems
-	E07.858.082.585 Kidneys, Artificial
-	E07.858.082.595 Larynx, Artificial
-	E07.858.082.620 Liver, Artificial
-	E07.858.082.760 Pancreas, Artificial
-	E07.858.082.800 Skin, Artificial
-	E07.858.082.900 Urinary Sphincter, Artificial
-	E07.858.150 Chest Tubes
-	E07.858.195 Consciousness Monitors
-	E07.858.240 Endoscopes
-	E07.858.240.040 Angioscopes
-	E07.858.240.060 Arthroscopes
-	E07.858.240.090 Bronchoscopes
-	E07.858.240.170 Colposcopes
-	E07.858.240.180 Culdoscopes
-	E07.858.240.190 Cystoscopes
-	E07.858.240.260 Endoscopes, Gastrointestinal
-	E07.858.240.260.160 Colonoscopes
-	E07.858.240.260.160.800 Sigmoidoscopes
-	E07.858.240.260.200 Duodenoscopes
-	E07.858.240.260.260 Esophagoscopes
-	E07.858.240.260.320 Gastrosopes
-	E07.858.240.260.680 Proctoscopes
-	E07.858.240.280 Fetoscopes
-	E07.858.240.390 Hysteroscopes
-	E07.858.240.520 Laparoscopes
-	E07.858.240.525 Laryngoscopes
-	E07.858.240.565 Mediastinoscopes
-	E07.858.240.707 Neuroendoscopes
-	E07.858.240.850 Thoracoscopes
-	E07.858.240.895 Ureteroscopes
-	E07.858.421 Operating Tables
-	E07.858.442 Orthopedic Equipment
-	E07.858.442.050 Artificial Limbs

## MeSH Tree Changes for 2017

Type	Tree - heading
-	E07.858.442.250 Canes
-	E07.858.442.281 Crutches
-	E07.858.442.660 Orthopedic Fixation Devices
-	E07.858.442.660.430 External Fixators
-	E07.858.442.660.430.500 Casts, Surgical
-	E07.858.442.660.430.750 Splints
-	E07.858.442.660.460 Internal Fixators
-	E07.858.442.660.460.249 Bone Nails
-	E07.858.442.660.460.374 Bone Plates
-	E07.858.442.660.460.437 Bone Screws
-	E07.858.442.660.460.437.500 Pedicle Screws
-	E07.858.442.660.460.468 Bone Wires
-	E07.858.442.660.460.734 Suture Anchors
-	E07.858.442.743 Orthotic Devices
-	E07.858.442.743.159 Athletic Tape
-	E07.858.442.743.319 Braces
-	E07.858.442.743.659 Foot Orthoses
-	E07.858.442.980 Walkers
-	E07.858.499 Pessaries
-	E07.858.594 Surgical Attire
-	E07.858.594.500 Gloves, Surgical
-	E07.858.594.750 Masks
-	E07.858.642 Surgical Drapes
-	E07.858.690 Surgical Fixation Devices
-	E07.858.690.725 Orthopedic Fixation Devices
-	E07.858.690.725.430 External Fixators
-	E07.858.690.725.430.500 Casts, Surgical
-	E07.858.690.725.430.750 Splints
-	E07.858.690.725.460 Internal Fixators
-	E07.858.690.725.460.249 Bone Nails
-	E07.858.690.725.460.374 Bone Plates
-	E07.858.690.725.460.437 Bone Screws
-	E07.858.690.725.460.437.500 Pedicle Screws
-	E07.858.690.725.460.468 Bone Wires
-	E07.858.690.725.460.734 Suture Anchors
-	E07.858.690.800 Surgical Tape



## MeSH Tree Changes for 2017

Type	Tree - heading
-	E07.858.690.820 Sutures
-	E07.858.690.820.250 Catgut
-	E07.858.690.860 Tissue Adhesives
-	E07.858.690.930 Vascular Closure Devices
-	E07.858.700 Surgical Instruments
-	E07.858.700.650 Obstetrical Forceps
-	E07.858.700.750 Surgical Staplers
-	E07.858.708 Surgical Mesh
-	E07.858.740 Surgical Sponges
-	E07.858.740.300 Gelatin Sponge, Absorbable
-	E07.858.826 Tampons, Surgical
-	E07.858.886 Tissue Expansion Devices
-	E07.858.925 Trusses
-	E07.862 Surgically-Created Structures
-	E07.862.200 Colonic Pouches
-	E07.862.700 Skeletal Muscle Ventricle
-	E07.862.710 Surgical Flaps
-	E07.862.710.500 Free Tissue Flaps
-	E07.862.710.625 Myocutaneous Flap
-	E07.862.710.750 Perforator Flap
-	E07.862.750 Urinary Reservoirs, Continent
-	E07.877 Syringes
-	E07.900 Thermometers
-	E07.913 Tomography Scanners, X-Ray Computed
-	E07.926 Tourniquets
-	E07.935 Transdermal Patch
-	E07.950 Ventilators, Mechanical
-	E07.950.500 Ventilators, Negative-Pressure
-	E07.960 X-Ray Film
-	E07.970 X-Ray Intensifying Screens
-	F01 Behavior and Behavior Mechanisms
-	F01.058 Adaptation, Psychological
-	F01.058.144 Emotional Adjustment
-	F01.058.288 Feedback, Psychological
-	F01.058.577 Orientation
New	<b>F01.058.577.500 Orientation, Spatial</b>

## MeSH Tree Changes for 2017

Type	Tree - heading
Heading	
-	F01.058.788 Sense of Coherence
-	F01.100 Attitude
-	F01.100.050 Attitude of Health Personnel
New Heading	<b>F01.100.050.500 Alert Fatigue, Health Personnel</b>
-	F01.100.100 Attitude to Computers
-	F01.100.125 Attitude to Death
-	F01.100.150 Attitude to Health
-	F01.100.150.500 Health Knowledge, Attitudes, Practice
-	F01.100.575 Catastrophization
-	F01.100.787 Optimism
-	F01.100.893 Pessimism
-	F01.145 Behavior
-	F01.145.015 Accident Proneness
-	F01.145.022 Adolescent Behavior
-	F01.145.022.750 Underage Drinking
-	F01.145.113 Behavior, Animal
-	F01.145.113.055 Animal Communication
-	F01.145.113.055.400 Echolocation
-	F01.145.113.055.800 Vocalization, Animal
-	F01.145.113.069 Animal Distribution
-	F01.145.113.069.500 Animal Migration
-	F01.145.113.111 Appetitive Behavior
-	F01.145.113.111.453 Grooming
New Heading	<b>F01.145.113.111.527 Host-Seeking Behavior</b>
-	F01.145.113.111.600 Predatory Behavior
-	F01.145.113.252 Consummatory Behavior
-	F01.145.113.252.478 Nesting Behavior
-	F01.145.113.252.520 Predatory Behavior
-	F01.145.113.252.748 Sexual Behavior, Animal
-	F01.145.113.252.748.200 Copulation
-	F01.145.113.252.748.300 Mating Preference, Animal
-	F01.145.113.252.748.400 Pair Bond
-	F01.145.113.367 Eliminative Behavior, Animal

## MeSH Tree Changes for 2017

Type	Tree - heading	
Old Tree	F01.145.113.448	Escape Reaction
-	F01.145.113.547	Feeding Behavior
-	F01.145.113.547.200	Cannibalism
-	F01.145.113.547.400	Carnivory
-	F01.145.113.547.500	Coprophagia
-	F01.145.113.547.600	Herbivory
-	F01.145.113.555	Freezing Reaction, Cataleptic
-	F01.145.113.646	Homing Behavior
-	F01.145.113.680	Immobility Response, Tonic
New Tree	F01.145.113.700	Kinesis
New Heading	F01.145.113.780	Taxis Response
New Tree	F01.145.113.780.500	Chemotaxis
New Tree	F01.145.113.780.688	Escape Reaction
New Heading	F01.145.113.780.875	Phototaxis
-	F01.145.113.840	Tool Use Behavior
-	F01.145.126	Behavioral Symptoms
-	F01.145.126.100	Affective Symptoms
-	F01.145.126.125	Aggression
-	F01.145.126.125.100	Agonistic Behavior
-	F01.145.126.125.550	Bullying
-	F01.145.126.156	Catatonia
-	F01.145.126.159	Child Reactive Disorders
-	F01.145.126.200	Delusions
-	F01.145.126.300	Depersonalization
-	F01.145.126.350	Depression
-	F01.145.126.837	Encopresis
-	F01.145.126.856	Enuresis
-	F01.145.126.856.249	Diurnal Enuresis
-	F01.145.126.856.500	Nocturnal Enuresis
-	F01.145.126.875	Hearing Loss, Functional
-	F01.145.126.900	Human Coprophagia
-	F01.145.126.925	Malingering

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	F01.145.126.937	Mental Fatigue
New Heading	<b>F01.145.126.937.250</b>	<b>Alert Fatigue, Health Personnel</b>
-	F01.145.126.937.500	Compassion Fatigue
-	F01.145.126.950	Obsessive Behavior
-	F01.145.126.950.500	Stalking
-	F01.145.126.962	Paranoid Behavior
-	F01.145.126.968	Polydipsia, Psychogenic
-	F01.145.126.972	Problem Behavior
-	F01.145.126.975	Schizophrenic Language
-	F01.145.126.980	Self-Injurious Behavior
-	F01.145.126.980.750	Self Mutilation
-	F01.145.126.980.875	Suicide
-	F01.145.126.980.875.149	Suicidal Ideation
-	F01.145.126.980.875.300	Suicide, Assisted
-	F01.145.126.980.875.600	Suicide, Attempted
-	F01.145.126.990	Stress, Psychological
-	F01.145.126.990.469	Compassion Fatigue
-	F01.145.126.995	Wandering Behavior
-	F01.145.179	Child Behavior
-	F01.145.179.500	Infant Behavior
-	F01.145.179.750	Problem Behavior
-	F01.145.194	Codependency (Psychology)
-	F01.145.209	Communication
-	F01.145.209.372	Information Seeking Behavior
-	F01.145.209.399	Language
-	F01.145.209.429	Literacy
-	F01.145.209.459	Narration
-	F01.145.209.520	Negotiating
-	F01.145.209.530	Nonverbal Communication
-	F01.145.209.530.136	Blushing
-	F01.145.209.530.258	Crying
-	F01.145.209.530.385	Facial Expression
-	F01.145.209.530.385.671	Smiling
-	F01.145.209.530.538	Kinesics
-	F01.145.209.530.538.445	Gestures

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	F01.145.209.530.614	Laughter
-	F01.145.209.530.702	Manual Communication
-	F01.145.209.530.702.668	Sign Language
-	F01.145.209.719	Teach-Back Communication
-	F01.145.209.908	Verbal Behavior
-	F01.145.209.908.677	Speech
-	F01.145.209.908.677.610	Speech Intelligibility
-	F01.145.236	Consumer Behavior
-	F01.145.250	Criminal Behavior
-	F01.145.250.250	Driving Under the Influence
-	F01.145.263	Dangerous Behavior
-	F01.145.263.500	Driving Under the Influence
-	F01.145.317	Drinking Behavior
-	F01.145.317.134	Alcohol Abstinence
-	F01.145.317.269	Alcohol Drinking
-	F01.145.317.269.500	Binge Drinking
-	F01.145.317.269.625	Alcohol Drinking in College
-	F01.145.317.269.875	Underage Drinking
-	F01.145.342	Drug-Seeking Behavior
-	F01.145.367	Escape Reaction
-	F01.145.387	Exploratory Behavior
-	F01.145.407	Feeding Behavior
-	F01.145.407.099	Bottle Feeding
-	F01.145.407.199	Breast Feeding
-	F01.145.407.199.500	Breast Milk Expression
-	F01.145.407.299	Carnivory
-	F01.145.407.400	Fasting
Old Tree	F01.145.407.432	Food Habits
Old Tree	F01.145.407.432.500	Meals
Old Tree	F01.145.407.432.500.100	Breakfast
Old Tree	F01.145.407.432.500.550	Lunch
Old Tree	F01.145.407.432.500.775	Snacks
-	F01.145.407.516	Food Preferences
-	F01.145.407.758	Herbivory
Old Tree	F01.145.437	Functional Behavior, Psychology
-	F01.145.466	Habits

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	F01.145.466.263	Fingersucking
Old Tree	<b>F01.145.466.349</b>	<b>Food Habits</b>
Old Tree	<b>F01.145.466.349.500</b>	<b>Meals</b>
Old Tree	<b>F01.145.466.349.500.100</b>	<b>Breakfast</b>
Old Tree	<b>F01.145.466.349.500.550</b>	<b>Lunch</b>
Old Tree	<b>F01.145.466.349.500.775</b>	<b>Snacks</b>
-	F01.145.466.554	Nail Biting
-	F01.145.466.874	Tongue Habits
-	F01.145.477	Harm Reduction
-	F01.145.488	Health Behavior
-	F01.145.488.500	Patient Compliance
-	F01.145.488.500.500	Medication Adherence
-	F01.145.488.500.750	No-Show Patients
-	F01.145.488.600	Patient Dropouts
-	F01.145.488.700	Self-Examination
-	F01.145.488.700.100	Breast Self-Examination
New Heading	<b>F01.145.488.725</b>	<b>Sleep Hygiene</b>
-	F01.145.488.750	Tobacco Use Cessation
-	F01.145.488.750.700	Smoking Cessation
New Heading	<b>F01.145.488.750.850</b>	<b>Vaping</b>
-	F01.145.488.800	Treatment Refusal
New Heading	<b>F01.145.488.800.500</b>	<b>Vaccination Refusal</b>
-	F01.145.499	Illness Behavior
-	F01.145.510	Imitative Behavior
-	F01.145.527	Impulsive Behavior
-	F01.145.527.100	Compulsive Behavior
-	F01.145.527.100.120	Behavior, Addictive
-	F01.145.535	Information Seeking Behavior
-	F01.145.544	Inhibition (Psychology)
-	F01.145.544.538	Proactive Inhibition
-	F01.145.544.724	Reactive Inhibition
-	F01.145.588	Marijuana Smoking
-	F01.145.632	Motor Activity
-	F01.145.632.555	Freezing Reaction, Cataleptic

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	F01.145.632.680	Immobility Response, Tonic
New Tree	<a href="#">F01.145.655</a>	<a href="#">Hoarding</a>
New Tree	<a href="#">F01.145.655</a>	<a href="#">Obsessive Hoarding</a>
-	F01.145.677	Personal Satisfaction
-	F01.145.688	Reproductive Behavior
-	F01.145.688.500	Contraception Behavior
-	F01.145.699	Risk Reduction Behavior
-	F01.145.699.400	HIV Serosorting
-	F01.145.722	Risk-Taking
-	F01.145.722.408	Gambling
-	F01.145.775	Self Stimulation
-	F01.145.802	Sexual Behavior
-	F01.145.802.188	Coitus
-	F01.145.802.188.300	Coitus Interruptus
-	F01.145.802.279	Courtship
-	F01.145.802.295	Extramarital Relations
-	F01.145.802.400	HIV Serosorting
-	F01.145.802.526	Masturbation
-	F01.145.802.790	Prostitution
-	F01.145.802.790	Sex work
-	F01.145.802.845	Safe Sex
-	F01.145.802.900	Sexual Abstinence
-	F01.145.802.950	Sexual Harassment
-	F01.145.802.975	Sexuality
-	F01.145.802.975.200	Bisexuality
-	F01.145.802.975.400	Heterosexuality
-	F01.145.802.975.500	Homosexuality
-	F01.145.802.975.500.400	Homosexuality, Female
-	F01.145.802.975.500.600	Homosexuality, Male
New Tree	<a href="#">F01.145.802.975.750</a>	<a href="#">Transsexualism</a>
-	F01.145.802.987	Unsafe Sex
-	F01.145.813	Social Behavior
-	F01.145.813.045	Aggression
-	F01.145.813.090	Altruism

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	F01.145.813.093	Anomie
-	F01.145.813.097	Ceremonial Behavior
-	F01.145.813.105	Competitive Behavior
-	F01.145.813.115	Cooperative Behavior
-	F01.145.813.157	Deception
-	F01.145.813.191	Dehumanization
-	F01.145.813.191.250	Commodification
-	F01.145.813.199	Femininity
-	F01.145.813.208	Gift Giving
New Heading	<b>F01.145.813.213</b>	<b>Harassment, Non-Sexual</b>
New Tree	<a href="#">F01.145.813.213.500</a>	<a href="#">Bullying</a>
-	F01.145.813.217	Help-Seeking Behavior
-	F01.145.813.225	Helping Behavior
-	F01.145.813.376	Masculinity
-	F01.145.813.527	Mass Behavior
-	F01.145.813.546	Permissiveness
-	F01.145.813.550	Prejudice
-	F01.145.813.550.249	Ageism
-	F01.145.813.550.374	Homophobia
-	F01.145.813.550.500	Racism
-	F01.145.813.550.750	Sexism
-	F01.145.813.550.875	Xenophobia
-	F01.145.813.565	Rejection (Psychology)
-	F01.145.813.572	Scapegoating
-	F01.145.813.595	Self-Control
New Tree	<a href="#">F01.145.813.606</a>	<a href="#">Sexual Harassment</a>
-	F01.145.813.617	Shyness
-	F01.145.813.621	Social Adjustment
-	F01.145.813.625	Social Conformity
-	F01.145.813.628	Social Desirability
-	F01.145.813.629	Social Discrimination
-	F01.145.813.629.500	Ageism
-	F01.145.813.629.562	Homophobia
-	F01.145.813.629.625	Racism



## MeSH Tree Changes for 2017

Type	Tree - heading	
-	F01.145.813.629.750	Sexism
-	F01.145.813.629.875	Xenophobia
-	F01.145.813.630	Social Distance
-	F01.145.813.650	Social Dominance
-	F01.145.813.650.400	Dominance-Subordination
-	F01.145.813.655	Social Facilitation
-	F01.145.813.708	Social Identification
-	F01.145.813.781	Social Isolation
-	F01.145.813.817	Social Marginalization
-	F01.145.813.828	Social Skills
-	F01.145.813.840	Social Stigma
-	F01.145.813.854	Stereotyping
-	F01.145.813.963	Whistleblowing
-	F01.145.875	Spatial Behavior
-	F01.145.875.281	Crowding
New Heading	<b>F01.145.875.439</b>	<b>Orientation, Spatial</b>
New Heading	<b>F01.145.875.439.500</b>	<b>Taxis Response</b>
New Tree	<a href="#">F01.145.875.439.500.500</a>	<a href="#">Chemotaxis</a>
New Tree	<a href="#">F01.145.875.439.500.688</a>	<a href="#">Escape Reaction</a>
New Heading	<b>F01.145.875.439.500.875</b>	<b>Phototaxis</b>
-	F01.145.875.596	Personal Space
-	F01.145.875.730	Spatial Learning
-	F01.145.875.797	Spatial Navigation
-	F01.145.875.830	Spatial Processing
-	F01.145.875.864	Territoriality
-	F01.145.896	Stereotyped Behavior
-	F01.145.916	Sucking Behavior
-	F01.145.958	Tobacco Use
-	F01.145.958.500	Smoking
New Heading	<b>F01.145.979</b>	<b>Vaping</b>
-	F01.318	Child Rearing
-	F01.318.709	Toilet Training

## MeSH Tree Changes for 2017

Type	Tree - heading
-	F01.393 Defense Mechanisms
-	F01.393.047 Acting Out
-	F01.393.200 Denial (Psychology)
-	F01.393.246 Displacement (Psychology)
-	F01.393.246.250 Cathexis
-	F01.393.246.500 Scapegoating
-	F01.393.351 Fantasy
-	F01.393.398 Helplessness, Learned
-	F01.393.446 Identification (Psychology)
-	F01.393.446.250 Gender Identity
-	F01.393.446.250.500 Femininity
-	F01.393.446.250.750 Masculinity
-	F01.393.628 Perceptual Defense
-	F01.393.693 Projection
-	F01.393.693.500 Scapegoating
-	F01.393.746 Rationalization
-	F01.393.784 Regression (Psychology)
-	F01.393.821 Repression, Psychology
-	F01.393.821.645 Repression-Sensitization
-	F01.393.898 Sublimation
-	F01.470 Emotions
-	F01.470.047 Affect
-	F01.470.047.110 Irritable Mood
-	F01.470.093 Anger
-	F01.470.093.640 Rage
-	F01.470.132 Anxiety
-	F01.470.132.150 Anxiety, Castration
-	F01.470.132.150.500 Koro
-	F01.470.132.225 Catastrophization
-	F01.470.132.300 Dental Anxiety
-	F01.470.132.650 Performance Anxiety
-	F01.470.137 Apathy
-	F01.470.142 Bereavement
-	F01.470.142.110 Grief
-	F01.470.192 Boredom
-	F01.470.257 Courage

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	F01.470.323	Euphoria
-	F01.470.342	Expressed Emotion
-	F01.470.361	Fear
-	F01.470.361.300	Dental Anxiety
-	F01.470.361.585	Panic
-	F01.470.383	Forgiveness
-	F01.470.405	Frustration
-	F01.470.483	Guilt
-	F01.470.483.666	Shame
-	F01.470.516	Happiness
-	F01.470.548	Hate
-	F01.470.572	Hope
-	F01.470.596	Hostility
-	F01.470.653	Jealousy
-	F01.470.693	Laughter
-	F01.470.713	Loneliness
-	F01.470.734	Love
-	F01.470.867	Pleasure
-	F01.510	Human Characteristics
-	F01.525	Human Development
-	F01.525.049	Adolescent Development
-	F01.525.200	Child Development
-	F01.525.200.310	Language Development
-	F01.525.200.310.250	Child Language
-	F01.525.200.310.300	Crying
-	F01.590	Mental Competency
-	F01.658	Motivation
-	F01.658.059	Achievement
-	F01.658.100	Aspirations (Psychology)
-	F01.658.209	Conflict (Psychology)
-	F01.658.293	Drive
-	F01.658.293.195	Craving
-	F01.658.293.391	Hunger
-	F01.658.293.787	Thirst
-	F01.658.370	Exploratory Behavior
-	F01.658.433	Food Deprivation

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	F01.658.500	Goals
-	F01.658.556	Handling (Psychology)
-	F01.658.642	Instinct
-	F01.658.650	Intention
-	F01.658.780	Power (Psychology)
-	F01.658.938	Water Deprivation
-	F01.700	Neurobehavioral Manifestations
-	F01.700.039	Anhedonia
-	F01.700.165	Catatonia
-	F01.700.250	Confusion
-	F01.700.250.500	Delirium
New Heading	<b>F01.700.250.500.500</b>	<b>Emergence Delirium</b>
-	F01.700.315	Consciousness Disorders
-	F01.700.470	Lethargy
-	F01.700.625	Memory Disorders
-	F01.700.625.100	Amnesia
-	F01.700.625.100.075	Amnesia, Anterograde
-	F01.700.625.100.150	Amnesia, Retrograde
-	F01.700.625.100.800	Amnesia, Transient Global
-	F01.700.625.400	Korsakoff Syndrome
-	F01.700.687	Intellectual Disability
-	F01.700.750	Perceptual Disorders
-	F01.700.750.100	Agnosia
-	F01.700.750.100.300	Gerstmann Syndrome
-	F01.700.750.100.650	Prosopagnosia
-	F01.700.750.150	Alice in Wonderland Syndrome
-	F01.700.750.175	Allesthesia
-	F01.700.750.200	Auditory Perceptual Disorders
-	F01.700.750.300	Hallucinations
-	F01.700.750.400	Illusions
-	F01.700.750.700	Phantom Limb
-	F01.700.812	Polydipsia, Psychogenic
-	F01.700.875	Psychomotor Disorders
-	F01.700.875.350	Apraxias
-	F01.700.875.350.099	Alien Hand Syndrome

## MeSH Tree Changes for 2017

Type	Tree - heading
-	F01.700.875.350.200 Apraxia, Ideomotor
-	F01.700.875.350.600 Gait Apraxia
-	F01.700.875.700 Psychomotor Agitation
-	F01.752 Personality
-	F01.752.049 Assertiveness
-	F01.752.098 Authoritarianism
-	F01.752.190 Character
-	F01.752.264 Creativity
-	F01.752.330 Dependency (Psychology)
-	F01.752.355 Empathy
-	F01.752.488 Individuality
-	F01.752.543 Intelligence
-	F01.752.543.500 Emotional Intelligence
-	F01.752.543.500.250 Emotional Adjustment
-	F01.752.543.500.500 Empathy
-	F01.752.543.500.750 Motivation
-	F01.752.609 Leadership
-	F01.752.650 Machiavellianism
-	F01.752.698 Negativism
-	F01.752.747 Personality Development
-	F01.752.747.189 Ego
-	F01.752.747.189.508 Reality Testing
-	F01.752.747.246 Extraversion (Psychology)
-	F01.752.747.347 Id
-	F01.752.747.385 Identification (Psychology)
-	F01.752.747.385.200 Gender Identity
-	F01.752.747.385.200.500 Femininity
-	F01.752.747.385.200.750 Masculinity
-	F01.752.747.423 Identity Crisis
-	F01.752.747.466 Individuation
-	F01.752.747.511 Introversion (Psychology)
-	F01.752.747.616 Moral Development
-	F01.752.747.722 Psychosexual Development
-	F01.752.747.722.200 Gender Identity
-	F01.752.747.722.458 Latency Period (Psychology)
-	F01.752.747.722.626 Oral Stage

## MeSH Tree Changes for 2017

Type	Tree - heading
-	F01.752.747.792 Self Concept
-	F01.752.747.792.110 Body Image
-	F01.752.747.792.220 Diagnostic Self Evaluation
-	F01.752.747.792.537 Self-Assessment
-	F01.752.747.792.662 Self Disclosure
-	F01.752.747.792.700 Self Efficacy
-	F01.752.747.792.925 Sense of Coherence
-	F01.752.747.859 Superego
-	F01.752.747.880 Type A Personality
-	F01.752.747.905 Type B Personality
-	F01.752.747.917 Type D Personality
-	F01.752.747.930 Unconscious (Psychology)
New Heading	<b>F01.752.823 Perfectionism</b>
-	F01.752.898 Temperament
-	F01.829 Psychology, Social
-	F01.829.131 Double Bind Interaction
-	F01.829.263 Family
-	F01.829.263.065 Adult Children
-	F01.829.263.132 Birth Order
-	F01.829.263.315 Family Characteristics
-	F01.829.263.315.500 Marital Status
-	F01.829.263.315.500.300 Divorce
-	F01.829.263.315.500.500 Marriage
-	F01.829.263.315.500.725 Single Person
-	F01.829.263.315.500.725.700 Single Parent
-	F01.829.263.315.500.862 Widowhood
-	F01.829.263.370 Family Relations
-	F01.829.263.370.054 Family Conflict
-	F01.829.263.370.110 Intergenerational Relations
-	F01.829.263.370.215 Maternal Behavior
-	F01.829.263.370.215.374 Maternal-Fetal Relations
-	F01.829.263.370.240 Maternal Deprivation
-	F01.829.263.370.290 Parent-Child Relations
-	F01.829.263.370.290.110 Father-Child Relations
-	F01.829.263.370.290.170 Mother-Child Relations

## MeSH Tree Changes for 2017

Type	Tree - heading
-	F01.829.263.370.290.170.500 Maternal-Fetal Relations
-	F01.829.263.370.310 Parenting
-	F01.829.263.370.330 Paternal Behavior
-	F01.829.263.370.380 Paternal Deprivation
-	F01.829.263.370.450 Sibling Relations
-	F01.829.263.403 Grandparents
-	F01.829.263.435 Military Family
-	F01.829.263.500 Nuclear Family
-	F01.829.263.500.280 Only Child
-	F01.829.263.500.320 Parents
-	F01.829.263.500.320.100 Fathers
-	F01.829.263.500.320.200 Mothers
-	F01.829.263.500.320.785 Single Parent
-	F01.829.263.500.320.892 Surrogate Mothers
-	F01.829.263.500.490 Siblings
-	F01.829.263.500.660 Spouses
-	F01.829.263.750 Single-Parent Family
-	F01.829.316 Group Processes
-	F01.829.316.068 Consensus
-	F01.829.316.171 Diplomacy
-	F01.829.316.274 Group Structure
-	F01.829.316.483 Peer Group
-	F01.829.316.483.750 Peer Influence
-	F01.829.316.549 Peer Review
-	F01.829.316.549.690 Peer Review, Health Care
-	F01.829.316.549.700 Peer Review, Research
-	F01.829.316.616 Role
-	F01.829.316.616.625 Professional Role
-	F01.829.316.616.625.450 Nurse's Role
-	F01.829.316.616.625.600 Physician's Role
-	F01.829.316.616.751 Sick Role
-	F01.829.316.697 Sensitivity Training Groups
-	F01.829.316.777 Social Distance
-	F01.829.379 Internal-External Control
-	F01.829.401 Interpersonal Relations
-	F01.829.401.046 Disclosure

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	F01.829.401.046.800	Truth Disclosure
-	F01.829.401.046.800.200	Duty to Warn
-	F01.829.401.094	Dissent and Disputes
-	F01.829.401.142	Family Conflict
-	F01.829.401.166	Forgiveness
-	F01.829.401.190	Intergenerational Relations
-	F01.829.401.205	Interprofessional Relations
-	F01.829.401.205.249	Interdisciplinary Communication
-	F01.829.401.205.500	Physician-Nurse Relations
-	F01.829.401.520	Negotiating
-	F01.829.401.550	Professional-Family Relations
-	F01.829.401.650	Professional-Patient Relations
-	F01.829.401.650.410	Dentist-Patient Relations
-	F01.829.401.650.505	Duty to Recontact
-	F01.829.401.650.600	Nurse-Patient Relations
-	F01.829.401.650.675	Physician-Patient Relations
-	F01.829.401.650.837	Researcher-Subject Relations
-	F01.829.401.737	Social Skills
-	F01.829.401.825	Trust
-	F01.829.458	Life Style
New Heading	<b>F01.829.458.205</b>	<b>Healthy Lifestyle</b>
New Heading	<b>F01.829.458.205.500</b>	<b>Healthy Diet</b>
-	F01.829.458.410	Life Change Events
-	F01.829.458.705	Sedentary Lifestyle
-	F01.829.477	Morale
-	F01.829.500	Morals
-	F01.829.500.359	Conscience
-	F01.829.500.679	Moral Development
New Heading	<b>F01.829.500.720</b>	<b>Moral Status</b>
-	F01.829.500.760	Social Responsibility
-	F01.829.500.760.500	Moral Obligations
-	F01.829.500.840	Virtues
-	F01.829.547	Paternalism
-	F01.829.595	Prejudice



## MeSH Tree Changes for 2017

Type	Tree - heading
-	F01.829.595.249 Ageism
-	F01.829.595.374 Homophobia
-	F01.829.595.500 Racism
-	F01.829.595.750 Sexism
-	F01.829.595.875 Xenophobia
-	F01.829.628 Psychosocial Deprivation
-	F01.829.750 Social Norms
-	F01.829.873 Social Values
-	F01.914 Temperance
-	F02 Psychological Phenomena and Processes
-	F02.410 Mental Competency
-	F02.418 Mental Health
-	F02.463 Mental Processes
-	F02.463.093 Anticipation, Psychological
-	F02.463.188 Cognition
-	F02.463.188.150 Awareness
-	F02.463.188.305 Cognitive Dissonance
-	F02.463.188.331 Cognitive Reserve
-	F02.463.188.357 Comprehension
-	F02.463.188.409 Consciousness
-	F02.463.188.634 Imagination
-	F02.463.188.634.309 Dreams
-	F02.463.188.634.507 Fantasy
-	F02.463.188.675 Intuition
-	F02.463.188.756 Metacognition
-	F02.463.217 Executive Function
-	F02.463.247 Higher Nervous Activity
-	F02.463.306 Intention
-	F02.463.425 Learning
-	F02.463.425.069 Association
-	F02.463.425.069.296 Association Learning
-	F02.463.425.097 Avoidance Learning
-	F02.463.425.179 Conditioning (Psychology)
-	F02.463.425.179.149 Automatism
-	F02.463.425.179.308 Conditioning, Classical
-	F02.463.425.179.408 Conditioning, Eyelid

## MeSH Tree Changes for 2017

Type	Tree - heading
-	F02.463.425.179.509                      Conditioning, Operant
-	F02.463.425.209                            Critical Period (Psychology)
-	F02.463.425.234                            Cues
-	F02.463.425.280                            Discrimination Learning
-	F02.463.425.357                            Generalization (Psychology)
-	F02.463.425.357.359                      Generalization, Response
-	F02.463.425.357.526                      Generalization, Stimulus
-	F02.463.425.393                            Habituation, Psychophysiologic
-	F02.463.425.420                            Helplessness, Learned
-	F02.463.425.448                            Imprinting (Psychology)
-	F02.463.425.475                            Inhibition (Psychology)
-	F02.463.425.475.538                      Proactive Inhibition
-	F02.463.425.475.724                      Reactive Inhibition
-	F02.463.425.540                            Memory
-	F02.463.425.540.203                      Deja Vu
-	F02.463.425.540.254                      Memory, Episodic
-	F02.463.425.540.305                      Memory, Long-Term
-	F02.463.425.540.305.500                Memory Consolidation
-	F02.463.425.540.407                      Memory, Short-Term
-	F02.463.425.540.641                      Mental Recall
-	F02.463.425.540.706                      Recognition (Psychology)
-	F02.463.425.540.739                      Repetition Priming
-	F02.463.425.540.772                      Retention (Psychology)
-	F02.463.425.540.886                      Spatial Memory
-	F02.463.425.575                            Neurolinguistic Programming
-	F02.463.425.612                            Overlearning
-	F02.463.425.674                            Practice (Psychology)
-	F02.463.425.701                            Probability Learning
-	F02.463.425.720                            Problem-Based Learning
-	F02.463.425.725                            Problem Solving
-	F02.463.425.725.500                      Heuristics
-	F02.463.425.770                            Reinforcement (Psychology)
-	F02.463.425.770.232                      Extinction, Psychological
-	F02.463.425.770.379                      Knowledge of Results (Psychology)
-	F02.463.425.770.571                      Punishment
-	F02.463.425.770.644                      Reinforcement Schedule

## MeSH Tree Changes for 2017

Type	Tree - heading
-	F02.463.425.770.706 Reinforcement, Social
-	F02.463.425.770.769 Reinforcement, Verbal
-	F02.463.425.770.836 Reward
-	F02.463.425.770.836.500 Token Economy
-	F02.463.425.798 Reversal Learning
-	F02.463.425.838 Set (Psychology)
-	F02.463.425.874 Spatial Learning
-	F02.463.425.874.500 Maze Learning
-	F02.463.425.910 Transfer (Psychology)
-	F02.463.425.952 Verbal Learning
-	F02.463.425.952.500 Paired-Associate Learning
-	F02.463.425.952.747 Serial Learning
-	F02.463.509 Mind-Body Relations, Metaphysical
-	F02.463.551 Mindfulness
-	F02.463.593 Perception
-	F02.463.593.071 Auditory Perception
-	F02.463.593.071.173 Auditory Threshold
-	F02.463.593.071.173.095 Auditory Fatigue
-	F02.463.593.071.576 Loudness Perception
-	F02.463.593.071.594 Perceptual Masking
-	F02.463.593.071.700 Pitch Perception
-	F02.463.593.071.700.408 Pitch Discrimination
-	F02.463.593.071.869 Sound Localization
-	F02.463.593.071.875 Speech Perception
-	F02.463.593.112 Body Image
-	F02.463.593.200 Depth Perception
-	F02.463.593.200.390 Distance Perception
-	F02.463.593.257 Discrimination (Psychology)
-	F02.463.593.257.800 Signal Detection, Psychological
-	F02.463.593.292 Eidetic Imagery
-	F02.463.593.343 Field Dependence-Independence
-	F02.463.593.373 Form Perception
-	F02.463.593.373.765 Stereognosis
-	F02.463.593.446 Illusions
-	F02.463.593.446.659 Optical Illusions
-	F02.463.593.465 Interoception

## MeSH Tree Changes for 2017

Type	Tree - heading
-	F02.463.593.485 Olfactory Perception
-	F02.463.593.504 Pain Perception
-	F02.463.593.504.500 Nociception
-	F02.463.593.524 Pattern Recognition, Physiological
-	F02.463.593.524.500 Pattern Recognition, Visual
-	F02.463.593.524.500.500 Facial Recognition
-	F02.463.593.603 Perceptual Distortion
-	F02.463.593.696 Sensory Deprivation
-	F02.463.593.710 Sensory Thresholds
-	F02.463.593.710.190 Auditory Threshold
-	F02.463.593.710.370 Differential Threshold
-	F02.463.593.710.560 Pain Threshold
-	F02.463.593.710.725 Signal Detection, Psychological
-	F02.463.593.710.750 Subliminal Stimulation
-	F02.463.593.710.790 Taste Threshold
-	F02.463.593.725 Size Perception
-	F02.463.593.752 Social Perception
-	F02.463.593.778 Space Perception
-	F02.463.593.778.255 Depth Perception
-	F02.463.593.778.255.390 Distance Perception
-	F02.463.593.778.255.780 Vision Disparity
-	F02.463.593.778.435 Form Perception
-	F02.463.593.778.435.110 Contrast Sensitivity
-	F02.463.593.778.794 Size Perception
-	F02.463.593.817 Taste Perception
-	F02.463.593.857 Time Perception
-	F02.463.593.894 Touch Perception
-	F02.463.593.894.500 Stereognosis
-	F02.463.593.932 Visual Perception
-	F02.463.593.932.073 Afterimage
-	F02.463.593.932.145 Attentional Blink
-	F02.463.593.932.217 Color Perception
-	F02.463.593.932.281 Contrast Sensitivity
-	F02.463.593.932.401 Figural Aftereffect
-	F02.463.593.932.458 Flicker Fusion
-	F02.463.593.932.567 Motion Perception

## MeSH Tree Changes for 2017

Type	Tree - heading
-	F02.463.593.932.594 Optic Flow
-	F02.463.593.932.622 Pattern Recognition, Visual
-	F02.463.593.932.622.500 Facial Recognition
-	F02.463.593.932.677 Perceptual Closure
-	F02.463.593.932.733 Perceptual Masking
-	F02.463.593.932.877 Vision Disparity
-	F02.463.593.932.885 Vision, Binocular
-	F02.463.593.932.893 Vision, Monocular
-	F02.463.593.932.901 Visual Acuity
-	F02.463.593.932.901.500 Contrast Sensitivity
-	F02.463.593.932.901.750 Emmetropia
-	F02.463.593.932.934 Visual Fields
-	F02.463.593.959 Weight Perception
-	F02.463.641 Spatial Navigation
-	F02.463.689 Theory of Mind
-	F02.463.785 Thinking
-	F02.463.785.233 Concept Formation
-	F02.463.785.302 Creativity
-	F02.463.785.373 Decision Making
-	F02.463.785.373.173 Avoidance Learning
-	F02.463.785.373.346 Choice Behavior
-	F02.463.785.373.346.400 Career Choice
-	F02.463.785.373.346.700 Delay Discounting
-	F02.463.785.373.433 Consensus
-	F02.463.785.373.476 Dissent and Disputes
-	F02.463.785.373.476.850 Refusal to Participate
-	F02.463.785.373.520 Negotiating
-	F02.463.785.373.520.500 Diplomacy
-	F02.463.785.373.820 Uncertainty
-	F02.463.785.477 Esthetics
-	F02.463.785.477.304 Beauty
-	F02.463.785.626 Judgment
-	F02.463.785.810 Problem Solving
-	F02.463.785.810.500 Heuristics
-	F02.463.902 Volition
-	F02.550 Parapsychology

## MeSH Tree Changes for 2017

Type	Tree - heading
-	F02.550.847 Telepathy
-	F02.600 Personal Autonomy
-	F02.694 Psycholinguistics
-	F02.694.550 Neurolinguistic Programming
-	F02.694.663 Semantic Differential
-	F02.739 Psychological Theory
-	F02.739.138 Behaviorism
-	F02.739.418 Existentialism
-	F02.739.527 Gestalt Theory
-	F02.739.660 Personal Construct Theory
-	F02.739.794 Psychoanalytic Theory
-	F02.739.794.206 Ego
-	F02.739.794.206.508 Reality Testing
-	F02.739.794.253 Extraversion (Psychology)
-	F02.739.794.297 Freudian Theory
-	F02.739.794.371 Id
-	F02.739.794.405 Inhibition (Psychology)
-	F02.739.794.438 Introversion (Psychology)
-	F02.739.794.471 Jungian Theory
-	F02.739.794.511 Libido
-	F02.739.794.582 Narcissism
-	F02.739.794.624 Object Attachment
-	F02.739.794.624.120 Bonding, Human-Pet
-	F02.739.794.653 Oedipus Complex
-	F02.739.794.746 Pleasure-Pain Principle
-	F02.739.794.793 Psychosexual Development
-	F02.739.794.793.200 Gender Identity
-	F02.739.794.793.458 Latency Period (Psychology)
-	F02.739.794.793.626 Oral Stage
-	F02.739.794.837 Self Psychology
-	F02.739.794.881 Superego
-	F02.739.794.942 Unconscious (Psychology)
-	F02.739.897 Theory of Mind
-	F02.784 Psychology, Applied
-	F02.784.176 Counseling
-	F02.784.176.279 Directive Counseling

## MeSH Tree Changes for 2017

Type	Tree - heading
-	F02.784.176.279.500 Motivational Interviewing
-	F02.784.176.350 Distance Counseling
-	F02.784.176.560 Pastoral Care
-	F02.784.176.700 Sex Counseling
-	F02.784.240 Criminal Psychology
-	F02.784.240.514 Lie Detection
-	F02.784.412 Human Engineering
-	F02.784.412.221 Data Display
-	F02.784.412.575 Man-Machine Systems
-	F02.784.412.846 Task Performance and Analysis
-	F02.784.412.846.707 Time and Motion Studies
-	F02.784.412.846.853 Work Simplification
-	F02.784.412.923 Time Management
-	F02.784.629 Psychology, Educational
-	F02.784.629.054 Achievement
-	F02.784.629.131 Aptitude
-	F02.784.629.155 Aspirations (Psychology)
-	F02.784.629.228 Child, Exceptional
-	F02.784.629.228.350 Child, Gifted
-	F02.784.629.272 Child Guidance
-	F02.784.629.375 Education of Intellectually Disabled
-	F02.784.629.529 Learning
-	F02.784.629.529.223 Formative Feedback
-	F02.784.629.529.274 Learning Curve
-	F02.784.629.529.550 Neurolinguistic Programming
-	F02.784.629.529.663 Social Learning
-	F02.784.629.529.775 Spatial Learning
-	F02.784.629.709 Remedial Teaching
-	F02.784.629.796 Student Dropouts
-	F02.784.629.880 Underachievement
-	F02.784.629.937 Vocational Guidance
-	F02.784.692 Psychology, Industrial
-	F02.784.692.107 Absenteeism
-	F02.784.692.351 Efficiency
-	F02.784.692.425 Job Satisfaction
-	F02.784.692.586 Presenteeism

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	F02.784.692.746	Task Performance and Analysis
-	F02.784.692.746.707	Time and Motion Studies
-	F02.784.692.746.853	Work Simplification
-	F02.784.692.816	Time Management
-	F02.784.692.887	Vocational Guidance
-	F02.784.754	Psychology, Military
-	F02.808	Psychomotor Performance
-	F02.808.260	Motor Skills
-	F02.808.600	Task Performance and Analysis
-	F02.830	Psychophysiology
-	F02.830.071	Appetite
-	F02.830.104	Arousal
-	F02.830.104.214	Attention
New Heading	<b>F02.830.104.214.500</b>	<b>Attentional Bias</b>
-	F02.830.104.821	Wakefulness
-	F02.830.131	Biofeedback, Psychology
-	F02.830.131.500	Neurofeedback
-	F02.830.158	Blushing
-	F02.830.233	Consciousness
-	F02.830.297	Dominance, Cerebral
-	F02.830.297.425	Functional Laterality
-	F02.830.422	Habituation, Psychophysiological
-	F02.830.512	Lie Detection
-	F02.830.606	Orientation
Old Tree	<b>F02.830.606.587</b>	<b>Kinesis</b>
New Heading	<b>F02.830.606.793</b>	<b>Orientation, Spatial</b>
-	F02.830.650	Reaction Time
-	F02.830.650.400	Refractory Period, Psychological
-	F02.830.702	Reflex
-	F02.830.702.157	Gagging
-	F02.830.702.315	Galvanic Skin Response
-	F02.830.702.607	Piloerection
-	F02.830.702.807	Reflex, Startle
-	F02.830.749	Satiation



## MeSH Tree Changes for 2017

Type	Tree - heading
-	F02.830.749.658 Satiety Response
-	F02.830.784 Self Stimulation
-	F02.830.816 Sensation
-	F02.830.816.131 Breakthrough Pain
-	F02.830.816.263 Hearing
-	F02.830.816.263.500 Bone Conduction
-	F02.830.816.308 Mastodynia
-	F02.830.816.353 Musculoskeletal Pain
-	F02.830.816.353.500 Myalgia
-	F02.830.816.444 Pain
-	F02.830.816.444.350 Arthralgia
-	F02.830.816.444.350.500 Shoulder Pain
-	F02.830.816.444.700 Pain Threshold
-	F02.830.816.444.850 Renal Colic
-	F02.830.816.492 Pleasure
-	F02.830.816.541 Proprioception
-	F02.830.816.541.504 Kinesthesia
-	F02.830.816.643 Smell
-	F02.830.816.724 Taste
-	F02.830.816.781 Thermosensing
-	F02.830.816.850 Touch
-	F02.830.816.964 Vision, Ocular
-	F02.830.816.964.124 Color Vision
-	F02.830.816.964.186 Mesopic Vision
-	F02.830.816.964.186.624 Rod-Cone Interaction
-	F02.830.816.964.249 Night Vision
-	F02.830.816.964.500 Phosphenes
-	F02.830.816.964.730 Vision, Entoptic
-	F02.830.855 Sleep
-	F02.830.855.268 Dreams
-	F02.830.855.671 Sleep Deprivation
New Heading	<b>F02.830.855.734 Sleep Hygiene</b>
-	F02.830.855.796 Sleep Stages
-	F02.830.855.796.671 Sleep, REM
-	F02.830.900 Stress, Psychological

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	F02.830.900.110	Burnout, Professional
-	F02.830.900.333	Compassion Fatigue
-	F02.880	Religion and Psychology
-	F02.880.410	Pastoral Care
-	F02.880.705	Spirituality
-	F02.940	Resilience, Psychological
-	F02.970	Social Theory
-	F03	Mental Disorders
-	F03.080	Anxiety Disorders
-	F03.080.100	Agoraphobia
-	F03.080.300	Anxiety, Separation
-	F03.080.500	Neurocirculatory Asthenia
-	F03.080.550	Neurotic Disorders
-	F03.080.600	Obsessive-Compulsive Disorder
-	F03.080.600.250	Hoarding Disorder
Old Tree	<b>F03.080.600.500</b>	<b>Hoarding</b>
Old Tree	<b>F03.080.600.500</b>	<b>Obsessive Hoarding</b>
-	F03.080.700	Panic Disorder
-	F03.080.725	Phobic Disorders
New Heading	<b>F03.080.725.500</b>	<b>Phobia, Social</b>
-	F03.084	Bipolar and Related Disorders
-	F03.084.500	Bipolar Disorder
-	F03.250	Disruptive, Impulse Control, and Conduct Disorders
-	F03.250.300	Firesetting Behavior
-	F03.250.400	Gambling
-	F03.250.800	Trichotillomania
-	F03.300	Dissociative Disorders
-	F03.300.500	Multiple Personality Disorder
-	F03.388	Elimination Disorders
-	F03.388.300	Encopresis
-	F03.388.400	Enuresis
-	F03.388.400.249	Diurnal Enuresis
-	F03.388.400.500	Nocturnal Enuresis
-	F03.400	Feeding and Eating Disorders
-	F03.400.125	Anorexia Nervosa

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	F03.400.188	Binge-Eating Disorder
-	F03.400.250	Bulimia Nervosa
-	F03.400.500	Feeding and Eating Disorders of Childhood
-	F03.400.750	Female Athlete Triad Syndrome
-	F03.400.875	Pica
-	F03.600	Mood Disorders
-	F03.600.300	Depressive Disorder
-	F03.600.300.350	Depression, Postpartum
-	F03.600.300.375	Depressive Disorder, Major
-	F03.600.300.388	Depressive Disorder, Treatment-Resistant
-	F03.600.300.400	Dysthymic Disorder
-	F03.600.300.550	Premenstrual Dysphoric Disorder
-	F03.600.300.775	Seasonal Affective Disorder
-	F03.600.500	Cyclothymic Disorder
-	F03.608	Motor Disorders
-	F03.615	Neurocognitive Disorders
-	F03.615.200	Amnesia
-	F03.615.200.125	Alcohol Amnestic Disorder
-	F03.615.200.125.500	Korsakoff Syndrome
-	F03.615.200.137	Amnesia, Anterograde
-	F03.615.200.150	Amnesia, Retrograde
-	F03.615.200.800	Amnesia, Transient Global
-	F03.615.250	Cognition Disorders
-	F03.615.250.100	Auditory Perceptual Disorders
-	F03.615.250.400	Huntington Disease
-	F03.615.250.700	Cognitive Dysfunction
-	F03.615.250.700	Mild Cognitive Impairment
-	F03.615.300	Consciousness Disorders
-	F03.615.350	Delirium
New Heading	<b>F03.615.350.500</b>	<b>Emergence Delirium</b>
-	F03.615.400	Dementia
-	F03.615.400.050	AIDS Dementia Complex
-	F03.615.400.100	Alzheimer Disease
-	F03.615.400.125	Aphasia, Primary Progressive
-	F03.615.400.125.600	Primary Progressive Nonfluent Aphasia

## MeSH Tree Changes for 2017

Type	Tree - heading
-	F03.615.400.300 Creutzfeldt-Jakob Syndrome
-	F03.615.400.350 Dementia, Vascular
-	F03.615.400.350.400 Dementia, Multi-Infarct
-	F03.615.400.370 Diffuse Neurofibrillary Tangles with Calcification
-	F03.615.400.380 Frontotemporal Lobar Degeneration
-	F03.615.400.380.299 Frontotemporal Dementia
-	F03.615.400.380.299.500 Pick Disease of the Brain
-	F03.615.400.380.600 Primary Progressive Nonfluent Aphasia
-	F03.615.400.390 Huntington Disease
-	F03.615.400.431 Kluver-Bucy Syndrome
-	F03.615.400.512 Lewy Body Disease
-	F03.615.700 Dyslexia, Acquired
-	F03.615.700.500 Alexia, Pure
-	F03.625 Neurodevelopmental Disorders
-	F03.625.047 Anxiety, Separation
-	F03.625.094 Attention Deficit and Disruptive Behavior Disorders
-	F03.625.094.150 Attention Deficit Disorder with Hyperactivity
-	F03.625.094.300 Conduct Disorder
-	F03.625.141 Child Behavior Disorders
-	F03.625.164 Child Development Disorders, Pervasive
-	F03.625.164.100 Asperger Syndrome
-	F03.625.164.113 Autism Spectrum Disorder
-	F03.625.164.125 Autistic Disorder
-	F03.625.374 Communication Disorders
-	F03.625.374.125 Childhood-Onset Fluency Disorder
-	F03.625.374.250 Social Communication Disorder
-	F03.625.374.500 Speech Sound Disorder
-	F03.625.421 Developmental Disabilities
-	F03.625.539 Intellectual Disability
-	F03.625.562 Learning Disorders
-	F03.625.562.199 Dyscalculia
-	F03.625.562.400 Dyslexia
-	F03.625.562.400.500 Dyslexia, Acquired
-	F03.625.562.700 Specific Learning Disorder
-	F03.625.813 Motor Skills Disorders
-	F03.625.875 Mutism

## MeSH Tree Changes for 2017

Type	Tree - heading
-	F03.625.937 Reactive Attachment Disorder
-	F03.625.968 Schizophrenia, Childhood
-	F03.625.984 Stereotypic Movement Disorder
-	F03.625.992 Tic Disorders
-	F03.625.992.850 Tourette Syndrome
-	F03.650 Neurotic Disorders
-	F03.657 Paraphilic Disorders
-	F03.657.300 Exhibitionism
-	F03.657.350 Fetishism (Psychiatric)
-	F03.657.500 Masochism
-	F03.657.600 Pedophilia
-	F03.657.700 Sadism
-	F03.657.800 Transvestism
-	F03.657.900 Voyeurism
-	F03.675 Personality Disorders
-	F03.675.050 Antisocial Personality Disorder
-	F03.675.100 Borderline Personality Disorder
-	F03.675.150 Compulsive Personality Disorder
-	F03.675.200 Dependent Personality Disorder
-	F03.675.400 Histrionic Personality Disorder
-	F03.675.400.500 Hysteria
-	F03.675.600 Paranoid Personality Disorder
-	F03.675.625 Passive-Aggressive Personality Disorder
-	F03.675.700 Schizoid Personality Disorder
-	F03.675.725 Schizotypal Personality Disorder
-	F03.700 Schizophrenia Spectrum and Other Psychotic Disorders
-	F03.700.150 Affective Disorders, Psychotic
-	F03.700.300 Capgras Syndrome
-	F03.700.356 Delusional Parasitosis
-	F03.700.412 Morgellons Disease
-	F03.700.450 Paranoid Disorders
-	F03.700.675 Psychotic Disorders
-	F03.700.675.600 Psychoses, Substance-Induced
-	F03.700.675.600.750 Psychoses, Alcoholic
-	F03.700.750 Schizophrenia
-	F03.700.750.300 Schizophrenia, Catatonic

## MeSH Tree Changes for 2017

Type	Tree - heading
-	F03.700.750.350 Schizophrenia, Disorganized
-	F03.700.750.600 Schizophrenia, Paranoid
-	F03.700.750.700 Shared Paranoid Disorder
-	F03.835 Sexual Dysfunctions, Psychological
-	F03.835.199 Dyspareunia
-	F03.835.400 Erectile Dysfunction
-	F03.835.550 Gender Dysphoria
-	F03.835.700 Premature Ejaculation
-	F03.835.750 Sexual and Gender Disorders
Old Tree	<b>F03.835.800 Transsexualism</b>
-	F03.835.900 Vaginismus
-	F03.870 Sleep Wake Disorders
-	F03.870.400 Dyssomnias
-	F03.870.400.099 Sleep Deprivation
-	F03.870.400.200 Sleep Disorders, Circadian Rhythm
-	F03.870.400.200.500 Jet Lag Syndrome
-	F03.870.400.800 Sleep Disorders, Intrinsic
-	F03.870.400.800.200 Disorders of Excessive Somnolence
-	F03.870.400.800.200.400 Hypersomnolence, Idiopathic
-	F03.870.400.800.200.500 Kleine-Levin Syndrome
-	F03.870.400.800.200.750 Narcolepsy
-	F03.870.400.800.200.750.500 Cataplexy
-	F03.870.400.800.700 Restless Legs Syndrome
-	F03.870.400.800.800 Sleep Initiation and Maintenance Disorders
-	F03.870.664 Parasomnias
-	F03.870.664.627 Nocturnal Paroxysmal Dystonia
-	F03.870.664.633 REM Sleep Parasomnias
-	F03.870.664.633.700 REM Sleep Behavior Disorder
-	F03.870.664.633.800 Sleep Paralysis
-	F03.870.664.634 Restless Legs Syndrome
-	F03.870.664.635 Sleep Arousal Disorders
-	F03.870.664.635.600 Night Terrors
-	F03.870.664.635.700 Somnambulism
-	F03.870.664.637 Sleep Bruxism
-	F03.870.664.700 Sleep-Wake Transition Disorders
-	F03.875 Somatoform Disorders

## MeSH Tree Changes for 2017

Type	Tree - heading
-	F03.875.149                      Body Dysmorphic Disorders
-	F03.875.300                      Conversion Disorder
-	F03.875.375                      Factitious Disorders
-	F03.875.375.600                  Munchausen Syndrome
-	F03.875.375.800                  Munchausen Syndrome by Proxy
-	F03.875.450                      Hypochondriasis
-	F03.875.600                      Neurasthenia
-	F03.900                            Substance-Related Disorders
-	F03.900.100                      Alcohol-Related Disorders
-	F03.900.100.050                  Alcohol Amnestic Disorder
-	F03.900.100.050.500              Korsakoff Syndrome
-	F03.900.100.100                  Alcohol Withdrawal Delirium
-	F03.900.100.300                  Alcoholic Intoxication
-	F03.900.100.350                  Alcoholism
-	F03.900.100.550                  Binge Drinking
-	F03.900.100.750                  Psychoses, Alcoholic
-	F03.900.100.875                  Wernicke Encephalopathy
-	F03.900.225                      Amphetamine-Related Disorders
-	F03.900.300                      Cocaine-Related Disorders
-	F03.900.467                      Inhalant Abuse
-	F03.900.635                      Marijuana Abuse
-	F03.900.650                      Neonatal Abstinence Syndrome
-	F03.900.675                      Opioid-Related Disorders
-	F03.900.675.400                  Heroin Dependence
-	F03.900.675.600                  Morphine Dependence
-	F03.900.700                      Phencyclidine Abuse
-	F03.900.746                      Psychoses, Substance-Induced
-	F03.900.793                      Substance Abuse, Intravenous
-	F03.900.825                      Substance Withdrawal Syndrome
-	F03.900.825.500                  Alcohol Withdrawal Delirium
-	F03.900.912                      Tobacco Use Disorder
-	F03.950                            Trauma and Stressor Related Disorders
-	F03.950.500                      Adjustment Disorders
-	F03.950.750                      Stress Disorders, Traumatic
-	F03.950.750.124                  Battered Child Syndrome
-	F03.950.750.249                  Combat Disorders

## MeSH Tree Changes for 2017

Type	Tree - heading
-	F03.950.750.375 Psychological Trauma
-	F03.950.750.500 Stress Disorders, Post-Traumatic
-	F03.950.750.550 Stress Disorders, Traumatic, Acute
-	F04 Behavioral Disciplines and Activities
-	F04.047 Behavior Control
-	F04.096 Behavioral Sciences
-	F04.096.080 Behavioral Medicine
-	F04.096.144 Behavioral Research
-	F04.096.208 Ethology
-	F04.096.276 Genetics, Behavioral
-	F04.096.276.400 Genetic Determinism
-	F04.096.462 Parapsychology
-	F04.096.544 Psychiatry
-	F04.096.544.065 Adolescent Psychiatry
-	F04.096.544.090 Biological Psychiatry
-	F04.096.544.193 Child Psychiatry
-	F04.096.544.215 Community Psychiatry
-	F04.096.544.215.508 Preventive Psychiatry
-	F04.096.544.275 Ethnopsychology
-	F04.096.544.335 Forensic Psychiatry
-	F04.096.544.335.200 Commitment of Mentally Ill
-	F04.096.544.335.240 Confidentiality
-	F04.096.544.335.240.270 Duty to Warn
-	F04.096.544.335.528 Insanity Defense
-	F04.096.544.380 Geriatric Psychiatry
-	F04.096.544.480 Military Psychiatry
-	F04.096.544.504 Neuropsychiatry
-	F04.096.544.528 Orthopsychiatry
-	F04.096.544.779 Psychoanalysis
-	F04.096.544.830 Psychosomatic Medicine
-	F04.096.586 Psycholinguistics
-	F04.096.586.550 Neurolinguistic Programming
-	F04.096.628 Psychology
-	F04.096.628.033 Environmental Psychology
-	F04.096.628.065 Psychology, Adolescent
-	F04.096.628.193 Psychology, Child



## MeSH Tree Changes for 2017

Type	Tree - heading
-	F04.096.628.255 Cognitive Science
-	F04.096.628.255.500 Cognitive Neuroscience
-	F04.096.628.270 Psychology, Developmental
-	F04.096.628.286 Economics, Behavioral
-	F04.096.628.317 Ethnopsychology
-	F04.096.628.579 Psychology, Clinical
-	F04.096.628.629 Psychology, Comparative
-	F04.096.628.679 Psychology, Educational
-	F04.096.628.729 Psychology, Experimental
-	F04.096.628.779 Psychology, Industrial
-	F04.096.628.779.450 Presenteeism
-	F04.096.628.779.900 Time Management
-	F04.096.628.808 Psychology, Medical
-	F04.096.628.829 Psychology, Social
-	F04.096.628.914 Psychology, Sports
-	F04.096.670 Psychopathology
-	F04.096.712 Psychopharmacology
-	F04.096.753 Psychophysics
-	F04.096.753.628 Psychoacoustics
-	F04.096.753.814 Signal Detection, Psychological
-	F04.096.795 Psychophysiology
-	F04.096.795.600 Neuropsychology
-	F04.096.795.650 Psychoneuroimmunology
-	F04.096.837 Sexology
-	F04.096.837.249 Sex Counseling
-	F04.096.837.500 Sex Education
-	F04.096.879 Social Sciences
-	F04.096.879.201 Anthropology
-	F04.096.879.201.500 Anthropology, Medical
-	F04.096.879.757 Sociology
-	F04.096.879.757.400 Sociology, Medical
-	F04.096.939 Sociobiology
-	F04.408 Mental Health Services
-	F04.408.192 Child Guidance
-	F04.408.307 Community Mental Health Services
-	F04.408.413 Counseling

## MeSH Tree Changes for 2017

Type	Tree - heading
-	F04.408.413.349 Directive Counseling
-	F04.408.413.349.500 Motivational Interviewing
-	F04.408.413.437 Distance Counseling
-	F04.408.413.700 Sex Counseling
-	F04.408.525 Emergency Services, Psychiatric
-	F04.408.823 Social Work, Psychiatric
-	F04.513 Personality Assessment
-	F04.513.636 Q-Sort
-	F04.570 Psychiatric Somatic Therapies
-	F04.570.200 Convulsive Therapy
-	F04.570.200.583 Electroconvulsive Therapy
-	F04.570.200.791 Transcranial Direct Current Stimulation
-	F04.570.400 Orthomolecular Therapy
-	F04.570.505 Psychopharmacology
-	F04.570.505.557 Narcotherapy
-	F04.570.630 Psychosurgery
-	F04.586 Psychiatric Status Rating Scales
-	F04.586.083 Brief Psychiatric Rating Scale
-	F04.586.574 Mental Status Schedule
-	F04.628 Psychoanalytic Interpretation
-	F04.669 Psychological Techniques
-	F04.669.112 Behavior Observation Techniques
-	F04.669.224 Electroshock
-	F04.669.224.300 Electroconvulsive Therapy
-	F04.669.224.650 Transcranial Direct Current Stimulation
-	F04.669.332 Galvanic Skin Response
-	F04.669.599 Interview, Psychological
-	F04.669.817 Reaction Time
-	F04.669.817.559 Refractory Period, Psychological
-	F04.669.908 Signal Detection, Psychological
-	F04.711 Psychological Tests
-	F04.711.141 Aptitude Tests
-	F04.711.141.493 Intelligence Tests
-	F04.711.141.493.225 Stanford-Binet Test
-	F04.711.141.493.822 Wechsler Scales
-	F04.711.271 Behavior Rating Scale

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Heading	<b>F04.711.336</b>	<b>Ecological Momentary Assessment</b>
-	F04.711.400	Language Tests
-	F04.711.513	Neuropsychological Tests
-	F04.711.513.180	Bender-Gestalt Test
-	F04.711.513.300	Luria-Nebraska Neuropsychological Battery
New Heading	<b>F04.711.513.502</b>	<b>Mental Navigation Tests</b>
-	F04.711.513.703	Stroop Test
-	F04.711.513.838	Trail Making Test
-	F04.711.647	Personality Tests
-	F04.711.647.138	Bender-Gestalt Test
-	F04.711.647.513	Personality Inventory
-	F04.711.647.513.146	Cattell Personality Factor Questionnaire
-	F04.711.647.513.463	Manifest Anxiety Scale
-	F04.711.647.513.535	Millon Clinical Multiaxial Inventory
-	F04.711.647.513.607	MMPI
-	F04.711.647.513.886	Test Anxiety Scale
-	F04.711.647.622	Projective Techniques
-	F04.711.647.622.341	Ink Blot Tests
-	F04.711.647.622.341.387	Holtzman Inkblot Test
-	F04.711.647.622.341.736	Rorschach Test
-	F04.711.647.622.851	Thematic Apperception Test
-	F04.711.647.745	Semantic Differential
-	F04.711.647.905	Word Association Tests
-	F04.711.780	Psychometrics
-	F04.754	Psychotherapy
-	F04.754.017	Animal Assisted Therapy
-	F04.754.017.500	Equine-Assisted Therapy
-	F04.754.035	Aromatherapy
-	F04.754.070	Art Therapy
-	F04.754.137	Behavior Therapy
-	F04.754.137.087	Anger Management Therapy
New Heading	<b>F04.754.137.131</b>	<b>Applied Behavior Analysis</b>
-	F04.754.137.174	Aversive Therapy
-	F04.754.137.301	Biofeedback, Psychology

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	F04.754.137.301.500	Feedback, Sensory
-	F04.754.137.301.750	Neurofeedback
New Heading	<b>F04.754.137.365</b>	<b>Cognitive Remediation</b>
-	F04.754.137.428	Cognitive Therapy
-	F04.754.137.428.249	Acceptance and Commitment Therapy
-	F04.754.137.428.500	Mindfulness
-	F04.754.137.506	Desensitization, Psychologic
-	F04.754.137.506.162	Eye Movement Desensitization Reprocessing
-	F04.754.137.506.325	Implosive Therapy
-	F04.754.137.506.662	Virtual Reality Exposure Therapy
-	F04.754.137.750	Relaxation Therapy
-	F04.754.137.750.500	Meditation
-	F04.754.137.875	Sleep Phase Chronotherapy
-	F04.754.168	Bibliotherapy
-	F04.754.215	Color Therapy
-	F04.754.252	Crisis Intervention
-	F04.754.278	Dance Therapy
New Heading	<b>F04.754.293</b>	<b>Emotion-Focused Therapy</b>
-	F04.754.308	Feedback, Psychological
-	F04.754.308.500	Biofeedback, Psychology
-	F04.754.308.500.500	Feedback, Sensory
-	F04.754.308.500.750	Neurofeedback
-	F04.754.339	Feedback, Sensory
-	F04.754.360	Gestalt Therapy
-	F04.754.392	Horticultural Therapy
-	F04.754.424	Hypnosis
-	F04.754.424.386	Autogenic Training
-	F04.754.424.771	Suggestion
-	F04.754.424.771.299	Autosuggestion
-	F04.754.462	Imagery (Psychotherapy)
-	F04.754.549	Music Therapy
-	F04.754.570	Narrative Therapy
-	F04.754.592	Person-Centered Therapy
-	F04.754.664	Play Therapy

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	F04.754.709	Psychoanalytic Therapy
-	F04.754.709.437	Free Association
-	F04.754.709.838	Transactional Analysis
-	F04.754.720	Psychotherapeutic Processes
-	F04.754.720.107	Abreaction
-	F04.754.720.107.333	Catharsis
-	F04.754.720.346	Association
-	F04.754.720.864	Transference (Psychology)
-	F04.754.720.864.363	Countertransference (Psychology)
-	F04.754.738	Psychotherapy, Brief
-	F04.754.766	Psychotherapy, Multiple
-	F04.754.775	Psychotherapy, Psychodynamic
-	F04.754.785	Psychotherapy, Rational-Emotive
-	F04.754.804	Reality Therapy
-	F04.754.864	Socioenvironmental Therapy
-	F04.754.864.392	Milieu Therapy
-	F04.754.864.392.701	Therapeutic Community
-	F04.754.864.581	Psychotherapy, Group
-	F04.754.864.581.136	Couples Therapy
-	F04.754.864.581.273	Family Therapy
-	F04.754.864.581.550	Marital Therapy
-	F04.754.864.581.679	Psychodrama
-	F04.754.864.581.679.653	Role Playing
-	F04.754.864.581.813	Sensitivity Training Groups
-	F04.754.864.696	Residential Treatment
-	F04.824	Schizophrenic Psychology
-	G01	Physical Phenomena
New Tree	<a href="#">G01.015</a>	<a href="#">Absorption</a>
New Tree	<a href="#">G01.015.249</a>	<a href="#">Absorption, Physicochemical</a>
New Tree	<a href="#">G01.015.249.500</a>	<a href="#">Absorption, Radiation</a>
New Tree	<a href="#">G01.030</a>	<a href="#">Adsorption</a>
-	G01.060	Astronomical Phenomena
New	<a href="#">G01.060.075</a>	<a href="#">Astronomical Objects</a>

## MeSH Tree Changes for 2017

Type	Tree - heading	
Tree		
New Tree	<a href="#">G01.060.075.159</a>	Extraterrestrial Environment
New Tree	<a href="#">G01.060.075.159.200</a>	Cosmic Dust
New Tree	<a href="#">G01.060.075.320</a>	Galaxies
New Tree	<a href="#">G01.060.075.500</a>	Meteoroids
New Tree	<a href="#">G01.060.075.680</a>	Planets
New Tree	<a href="#">G01.060.075.730</a>	Solar System
New Tree	<a href="#">G01.060.075.730.600</a>	Minor Planets
New Tree	<a href="#">G01.060.075.730.600.500</a>	Pluto
New Tree	<a href="#">G01.060.075.730.700</a>	Planets
New Tree	<a href="#">G01.060.075.730.700.200</a>	Earth (Planet)
New Tree	<a href="#">G01.060.075.730.700.200.500</a>	Moon
New Tree	<a href="#">G01.060.075.730.700.412</a>	Jupiter
New Tree	<a href="#">G01.060.075.730.700.625</a>	Mars
New Tree	<a href="#">G01.060.075.730.700.762</a>	Mercury (Planet)
New Tree	<a href="#">G01.060.075.730.700.796</a>	Neptune
New Tree	<a href="#">G01.060.075.730.700.831</a>	Saturn
New Tree	<a href="#">G01.060.075.730.700.865</a>	Uranus
New Tree	<a href="#">G01.060.075.730.700.900</a>	Venus
New Tree	<a href="#">G01.060.075.750</a>	Stars, Celestial
New Tree	<a href="#">G01.060.185</a>	Cosmic Radiation
Old Tree	<a href="#">G01.060.249</a>	Astronomical Objects
Old Tree	<a href="#">G01.060.249.159</a>	Extraterrestrial Environment

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	G01.060.249.159.200 Cosmic Dust
Old Tree	G01.060.249.320 Galaxies
Old Tree	G01.060.249.500 Meteoroids
Old Tree	G01.060.249.680 Planets
Old Tree	G01.060.249.730 Solar System
Old Tree	G01.060.249.730.600 Minor Planets
Old Tree	G01.060.249.730.600.500 Pluto
Old Tree	G01.060.249.730.700 Planets
Old Tree	G01.060.249.730.700.200 Earth (Planet)
Old Tree	G01.060.249.730.700.200.500 Moon
Old Tree	G01.060.249.730.700.412 Jupiter
Old Tree	G01.060.249.730.700.625 Mars
Old Tree	G01.060.249.730.700.762 Mercury (Planet)
Old Tree	G01.060.249.730.700.796 Neptune
Old Tree	G01.060.249.730.700.831 Saturn
Old Tree	G01.060.249.730.700.865 Uranus
Old Tree	G01.060.249.730.700.900 Venus
Old Tree	G01.060.249.750 Stars, Celestial
New Tree	G01.060.275 Evolution, Planetary
New Tree	G01.060.275.500 Evolution, Chemical
New Tree	G01.060.350 Gravitation
New Tree	G01.060.350.369 Gravity, Altered
New Tree	G01.060.350.369.300 Hypergravity
New Tree	G01.060.350.369.400 Hypogravity
New Tree	G01.060.350.369.400.900 Weightlessness
Old Tree	G01.060.374 Astronomical Processes
Old Tree	G01.060.374.134 Cosmic Radiation
Old Tree	G01.060.374.270 Evolution, Planetary
Old Tree	G01.060.374.270.500 Evolution, Chemical
Old Tree	G01.060.374.800 Solar Activity
New Tree	G01.060.800 Solar Activity

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	G01.154	Biophysical Phenomena
-	G01.154.090	Biomechanical Phenomena
New Tree	<a href="#">G01.154.090.500</a>	<a href="#">Mechanotransduction, Cellular</a>
New Heading	<b>G01.154.090.625</b>	<b>Orthodontic Friction</b>
New Tree	<a href="#">G01.154.090.750</a>	<a href="#">Osmosis</a>
New Tree	<a href="#">G01.154.090.750.500</a>	<a href="#">Electroosmosis</a>
Old Tree	<del>G01.154.100</del>	<del>Biophysical Processes</del>
Old Tree	<del>G01.154.100.240</del>	<del>Energy Transfer</del>
Old Tree	<del>G01.154.100.240.280</del>	<del>Fluorescence Resonance Energy Transfer</del>
Old Tree	<del>G01.154.100.240.400</del>	<del>Linear Energy Transfer</del>
Old Tree	<del>G01.154.100.280</del>	<del>Facilitated Diffusion</del>
Old Tree	<del>G01.154.100.440</del>	<del>Mechanotransduction, Cellular</del>
Old Tree	<del>G01.154.100.600</del>	<del>Osmosis</del>
Old Tree	<del>G01.154.100.600.500</del>	<del>Electroosmosis</del>
Old Tree	<del>G01.154.100.640</del>	<del>Protein Folding</del>
Old Tree	<del>G01.154.100.640.501</del>	<del>Protein Refolding</del>
Old Tree	<del>G01.154.100.640.750</del>	<del>Protein Unfolding</del>
Old Tree	<del>G01.154.100.640.750.500</del>	<del>Protein Denaturation</del>
Old Tree	<del>G01.154.100.820</del>	<del>RNA Folding</del>
New Tree	<a href="#">G01.154.240</a>	<a href="#">Energy Transfer</a>
New Tree	<a href="#">G01.154.240.280</a>	<a href="#">Fluorescence Resonance Energy Transfer</a>
New Tree	<a href="#">G01.154.240.400</a>	<a href="#">Linear Energy Transfer</a>
New Tree	<a href="#">G01.154.280</a>	<a href="#">Facilitated Diffusion</a>
-	G01.154.500	Membrane Fluidity
-	G01.154.535	Membrane Potentials
New Tree	<a href="#">G01.154.651</a>	<a href="#">Protein Folding</a>
New Tree	<a href="#">G01.154.651.501</a>	<a href="#">Protein Refolding</a>
New Tree	<a href="#">G01.154.651.750</a>	<a href="#">Protein Unfolding</a>
New	<a href="#">G01.154.651.750.500</a>	<a href="#">Protein Denaturation</a>



## MeSH Tree Changes for 2017

Type	Tree - heading	
Tree		
New Tree	<a href="#">G01.154.709</a>	<a href="#">RNA Folding</a>
-	<a href="#">G01.154.767</a>	<a href="#">Seed Dispersal</a>
New Tree	<a href="#">G01.202</a>	<a href="#">Diffusion</a>
New Tree	<a href="#">G01.202.374</a>	<a href="#">Facilitated Diffusion</a>
New Tree	<a href="#">G01.202.750</a>	<a href="#">Thermal Diffusion</a>
New Tree	<a href="#">G01.226</a>	<a href="#">Doppler Effect</a>
New Tree	<a href="#">G01.238</a>	<a href="#">Elementary Particle Interactions</a>
-	<a href="#">G01.249</a>	<a href="#">Elementary Particles</a>
-	<a href="#">G01.249.335</a>	<a href="#">Electrons</a>
-	<a href="#">G01.249.467</a>	<a href="#">Heavy Ions</a>
-	<a href="#">G01.249.600</a>	<a href="#">Mesons</a>
Old Tree	<a href="#">G01.249.645</a>	<a href="#">Neutrons</a>
Old Tree	<a href="#">G01.249.645.368</a>	<a href="#">Fast Neutrons</a>
New Heading	<b><a href="#">G01.249.660</a></b>	<b><a href="#">Nucleons</a></b>
New Tree	<a href="#">G01.249.660.250</a>	<a href="#">Neutrons</a>
New Tree	<a href="#">G01.249.660.250.368</a>	<a href="#">Fast Neutrons</a>
New Tree	<a href="#">G01.249.660.500</a>	<a href="#">Protons</a>
-	<a href="#">G01.249.675</a>	<a href="#">Phonons</a>
-	<a href="#">G01.249.705</a>	<a href="#">Photons</a>
Old Tree	<a href="#">G01.249.765</a>	<a href="#">Protons</a>
New Tree	<a href="#">G01.280</a>	<a href="#">Filtration</a>
New Tree	<a href="#">G01.280.807</a>	<a href="#">Ultrafiltration</a>
-	<a href="#">G01.311</a>	<a href="#">Geological Phenomena</a>
New Tree	<a href="#">G01.311.085</a>	<a href="#">Avalanches</a>
-	<a href="#">G01.311.169</a>	<a href="#">Caves</a>
New Tree	<a href="#">G01.311.250</a>	<a href="#">Earthquakes</a>

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	<a href="#">G01.311.290</a>	<a href="#">Evolution, Planetary</a>
-	G01.311.330	Geologic Sediments
Old Tree	<a href="#">G01.311.340</a>	<a href="#">Geological Processes</a>
Old Tree	<a href="#">G01.311.340.111</a>	<a href="#">Avalanches</a>
Old Tree	<a href="#">G01.311.340.224</a>	<a href="#">Earthquakes</a>
Old Tree	<a href="#">G01.311.340.449</a>	<a href="#">Evolution, Planetary</a>
Old Tree	<a href="#">G01.311.340.674</a>	<a href="#">Landslides</a>
Old Tree	<a href="#">G01.311.340.787</a>	<a href="#">Tidal Waves</a>
Old Tree	<a href="#">G01.311.340.843</a>	<a href="#">Tsunamis</a>
Old Tree	<a href="#">G01.311.340.900</a>	<a href="#">Volcanic Eruptions</a>
Old Tree	<a href="#">G01.311.340.950</a>	<a href="#">Water Cycle</a>
-	G01.311.355	Groundwater
-	G01.311.355.750	Natural Springs
-	G01.311.355.750.500	Hot Springs
-	G01.311.355.750.500.400	Hydrothermal Vents
-	G01.311.400	Ice Cover
-	G01.311.580	Lakes
New Tree	<a href="#">G01.311.603</a>	<a href="#">Landslides</a>
-	G01.311.625	Oceans and Seas
-	G01.311.625.080	Bays
-	G01.311.625.540	Estuaries
-	G01.311.687	Oil and Gas Fields
-	G01.311.718	Ponds
-	G01.311.750	Rivers
-	G01.311.820	Soil
-	G01.311.820.500	Permafrost
New Tree	<a href="#">G01.311.910</a>	<a href="#">Tidal Waves</a>
New Tree	<a href="#">G01.311.933</a>	<a href="#">Tsunamis</a>
New Tree	<a href="#">G01.311.955</a>	<a href="#">Volcanic Eruptions</a>
New Tree	<a href="#">G01.311.977</a>	<a href="#">Water Cycle</a>
-	G01.342	Hydrodynamics
-	G01.358	Magnetic Phenomena

## MeSH Tree Changes for 2017

Type	Tree - heading
-	G01.358.500 Electromagnetic Phenomena
-	G01.358.500.249 Electricity
-	G01.358.500.249.270 Electric Capacitance
-	G01.358.500.249.277 Electric Conductivity
-	G01.358.500.249.277.350 Electric Impedance
-	G01.358.500.249.820 Static Electricity
-	G01.358.500.260 Electromagnetic Fields
-	G01.358.500.505 Electromagnetic Radiation
-	G01.358.500.505.300 Gamma Rays
-	G01.358.500.505.650 Light
-	G01.358.500.505.650.440 Incandescence
-	G01.358.500.505.650.552 Infrared Rays
-	G01.358.500.505.650.665 Luminescence
-	G01.358.500.505.650.665.500 Fluorescence
-	G01.358.500.505.650.782 Photons
-	G01.358.500.505.650.836 Sunlight
-	G01.358.500.505.650.891 Ultraviolet Rays
-	G01.358.500.505.810 Radio Waves
-	G01.358.500.505.810.500 Microwaves
-	G01.358.500.505.890 Terahertz Radiation
-	G01.358.500.505.970 X-Rays
-	G01.358.500.750 Electrons
New Heading	<b>G01.358.500.875 Superconductivity</b>
-	G01.358.750 Magnetic Fields
-	G01.358.750.500 Electromagnetic Fields
-	G01.374 Mechanical Phenomena
-	G01.374.089 Biomechanical Phenomena
-	G01.374.180 Compressive Strength
-	G01.374.590 Elasticity
-	G01.374.590.210 Compliance
-	G01.374.590.605 Elastic Modulus
-	G01.374.618 Friction
-	G01.374.647 Hardness
-	G01.374.647.457 Hardness Tests
-	G01.374.661 Kinetics

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	<a href="#">G01.374.669</a>	<a href="#">Lifting</a>
Old Tree	<a href="#">G01.374.676</a>	<a href="#">Mechanical Processes</a>
Old Tree	<a href="#">G01.374.676.530</a>	<a href="#">Lifting</a>
Old Tree	<a href="#">G01.374.676.830</a>	<a href="#">Stress, Mechanical</a>
Old Tree	<a href="#">G01.374.676.860</a>	<a href="#">Torsion, Mechanical</a>
Old Tree	<a href="#">G01.374.676.860.500</a>	<a href="#">Torque</a>
Old Tree	<a href="#">G01.374.676.930</a>	<a href="#">Vibration</a>
Old Tree	<a href="#">G01.374.676.965</a>	<a href="#">Weight-Bearing</a>
-	G01.374.705	Pliability
-	G01.374.710	Porosity
-	G01.374.715	Pressure
-	G01.374.715.250	Decompression
-	G01.374.715.250.500	Decompression, Explosive
-	G01.374.715.352	Hydrostatic Pressure
-	G01.374.715.578	Osmotic Pressure
-	G01.374.715.714	Partial Pressure
-	G01.374.715.902	Vacuum
-	G01.374.715.951	Vapor Pressure
-	G01.374.820	Shear Strength
New Tree	<a href="#">G01.374.835</a>	<a href="#">Stress, Mechanical</a>
-	G01.374.850	Tensile Strength
New Tree	<a href="#">G01.374.860</a>	<a href="#">Torsion, Mechanical</a>
New Tree	<a href="#">G01.374.860.500</a>	<a href="#">Torque</a>
New Tree	<a href="#">G01.374.930</a>	<a href="#">Vibration</a>
New Tree	<a href="#">G01.374.965</a>	<a href="#">Weight-Bearing</a>
New Tree	<a href="#">G01.482</a>	<a href="#">Motion</a>
New Tree	<a href="#">G01.482.107</a>	<a href="#">Acceleration</a>
New Tree	<a href="#">G01.482.107.373</a>	<a href="#">Deceleration</a>
New Tree	<a href="#">G01.482.311</a>	<a href="#">Coriolis Force</a>

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	<a href="#">G01.482.620</a>	<a href="#">Pulsatile Flow</a>
New Tree	<a href="#">G01.482.703</a>	<a href="#">Rotation</a>
New Tree	<a href="#">G01.536</a>	<a href="#">Nuclear Energy</a>
New Tree	<a href="#">G01.536.485</a>	<a href="#">Nuclear Fission</a>
New Tree	<a href="#">G01.536.652</a>	<a href="#">Nuclear Fusion</a>
-	G01.590	Optical Phenomena
-	G01.590.040	Anisotropy
-	G01.590.080	Birefringence
-	G01.590.310	Iridescence
-	G01.590.540	Light
-	G01.590.540.199	Color
-	G01.590.540.233	Darkness
-	G01.590.540.333	Glare
-	G01.590.540.440	Incandescence
-	G01.590.540.552	Infrared Rays
-	G01.590.540.665	Luminescence
-	G01.590.540.665.500	Fluorescence
-	G01.590.540.782	Photons
-	G01.590.540.891	Ultraviolet Rays
Old Tree	<a href="#">G01.590.770</a>	<a href="#">Optical Processes</a>
Old Tree	<a href="#">G01.590.770.640</a>	<a href="#">Optical Rotation</a>
Old Tree	<a href="#">G01.590.770.760</a>	<a href="#">Refraction, Ocular</a>
New Tree	<a href="#">G01.590.773</a>	<a href="#">Optical Rotation</a>
New Tree	<a href="#">G01.590.775</a>	<a href="#">Refraction, Ocular</a>
Old Tree	<a href="#">G01.595</a>	<a href="#">Physical Processes</a>
Old Tree	<a href="#">G01.595.014</a>	<a href="#">Absorption</a>
Old Tree	<a href="#">G01.595.014.249</a>	<a href="#">Absorption, Physicochemical</a>
Old Tree	<a href="#">G01.595.014.249.500</a>	<a href="#">Absorption, Radiation</a>
Old Tree	<a href="#">G01.595.037</a>	<a href="#">Adsorption</a>
Old Tree	<a href="#">G01.595.060</a>	<a href="#">Astronomical Processes</a>
Old Tree	<a href="#">G01.595.060.134</a>	<a href="#">Cosmic Radiation</a>

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	G01.595.060.270 Evolution, Planetary
Old Tree	G01.595.060.270.500 Evolution, Chemical
Old Tree	G01.595.060.535 Gravitation
Old Tree	G01.595.060.535.369 Gravity, Altered
Old Tree	G01.595.060.535.369.300 Hypergravity
Old Tree	G01.595.060.535.369.400 Hypogravity
Old Tree	G01.595.060.535.369.400.900 Weightlessness
Old Tree	G01.595.060.800 Solar Activity
Old Tree	G01.595.100 Biophysical Processes
Old Tree	G01.595.100.240 Energy Transfer
Old Tree	G01.595.100.240.280 Fluorescence Resonance Energy Transfer
Old Tree	G01.595.100.240.400 Linear Energy Transfer
Old Tree	G01.595.100.280 Facilitated Diffusion
Old Tree	G01.595.100.440 Mechanotransduction, Cellular
Old Tree	G01.595.100.600 Osmosis
Old Tree	G01.595.100.600.500 Electroosmosis
Old Tree	G01.595.100.640 Protein Folding
Old Tree	G01.595.100.640.501 Protein Refolding
Old Tree	G01.595.100.640.750 Protein Unfolding
Old Tree	G01.595.100.640.750.500 Protein Denaturation
Old Tree	G01.595.100.820 RNA Folding
Old Tree	G01.595.200 Diffusion
Old Tree	G01.595.200.374 Facilitated Diffusion
Old Tree	G01.595.200.750 Thermal Diffusion
Old Tree	G01.595.210 Doppler Effect
Old Tree	G01.595.240 Elementary Particle Interactions
Old Tree	G01.595.280 Filtration
Old Tree	G01.595.280.807 Ultrafiltration
Old Tree	G01.595.340 Geological Processes
Old Tree	G01.595.340.111 Avalanches
Old Tree	G01.595.340.224 Earthquakes
Old Tree	G01.595.340.449 Evolution, Planetary
Old Tree	G01.595.340.674 Landslides
Old Tree	G01.595.340.787 Tsunamis
Old Tree	G01.595.340.900 Volcanic Eruptions
Old Tree	G01.595.540 Mechanical Processes

## MeSH Tree Changes for 2017

Type	Tree - heading	
Old Tree	G01.595.540.530	Lifting
Old Tree	G01.595.540.830	Stress, Mechanical
Old Tree	G01.595.540.860	Torsion, Mechanical
Old Tree	G01.595.540.860.500	Torque
Old Tree	G01.595.540.930	Vibration
Old Tree	G01.595.540.965	Weight-Bearing
Old Tree	G01.595.560	Motion
Old Tree	G01.595.560.107	Acceleration
Old Tree	G01.595.560.107.373	Deceleration
Old Tree	G01.595.560.311	Coriolis Force
Old Tree	G01.595.560.620	Pulsatile Flow
Old Tree	G01.595.560.703	Rotation
Old Tree	G01.595.630	Nuclear Energy
Old Tree	G01.595.630.485	Nuclear Fission
Old Tree	G01.595.630.652	Nuclear Fusion
Old Tree	G01.595.640	Optical Processes
Old Tree	G01.595.640.640	Optical Rotation
Old Tree	G01.595.640.760	Refraction, Ocular
Old Tree	G01.595.680	Phase Transition
Old Tree	G01.595.680.500	Freezing
Old Tree	G01.595.680.625	Vitrification
Old Tree	G01.595.680.750	Volatilization
Old Tree	G01.595.815	Radioactivity
Old Tree	G01.595.861	Scattering, Radiation
Old Tree	G01.595.861.650	Neutron Diffraction
Old Tree	G01.595.861.755	Scattering, Small Angle
Old Tree	G01.595.861.950	X-Ray Diffraction
New Tree	G01.645	Phase Transition
New Tree	G01.645.500	Freezing
New Tree	G01.645.625	Vitrification
New Tree	G01.645.750	Volatilization
-	G01.750	Radiation
-	G01.750.124	Absorption, Radiation

## MeSH Tree Changes for 2017

Type	Tree - heading
-	G01.750.250 Electromagnetic Radiation
-	G01.750.250.300 Gamma Rays
-	G01.750.250.650 Light
-	G01.750.250.650.440 Incandescence
-	G01.750.250.650.552 Infrared Rays
-	G01.750.250.650.665 Luminescence
-	G01.750.250.650.782 Photons
-	G01.750.250.650.836 Sunlight
-	G01.750.250.650.891 Ultraviolet Rays
-	G01.750.250.810 Radio Waves
-	G01.750.250.810.500 Microwaves
-	G01.750.250.890 Terahertz Radiation
-	G01.750.250.970 X-Rays
-	G01.750.740 Radiation Dosage
-	G01.750.740.500 Dose-Response Relationship, Radiation
-	G01.750.745 Radiation Effects
-	G01.750.748 Radiation Exposure
-	G01.750.748.500 Radiation Injuries
-	G01.750.748.500.031 Abnormalities, Radiation-Induced
-	G01.750.748.500.188 Acute Radiation Syndrome
-	G01.750.748.500.266 Cardiotoxicity
-	G01.750.748.500.345 Leukemia, Radiation-Induced
-	G01.750.748.500.476 Neoplasms, Radiation-Induced
-	G01.750.748.500.579 Osteoradionecrosis
-	G01.750.748.500.720 Radiation Injuries, Experimental
-	G01.750.748.500.762 Radiation Pneumonitis
-	G01.750.748.500.804 Radiodermatitis
-	G01.750.750 Radiation, Ionizing
-	G01.750.750.055 Alpha Particles
-	G01.750.750.115 Background Radiation
-	G01.750.750.125 Beta Particles
-	G01.750.750.235 Cosmic Radiation
-	G01.750.750.400 Gamma Rays
-	G01.750.750.659 Ultraviolet Rays
-	G01.750.750.918 X-Rays
-	G01.750.770 Radiation, Nonionizing



## MeSH Tree Changes for 2017

Type	Tree - heading
-	G01.750.770.578                      Light
-	G01.750.770.578.440                      Incandescence
-	G01.750.770.578.552                      Infrared Rays
-	G01.750.770.578.665                      Luminescence
-	G01.750.770.578.782                      Photons
-	G01.750.770.578.836                      Sunlight
-	G01.750.770.578.891                      Ultraviolet Rays
-	G01.750.770.721                      Radio Waves
-	G01.750.770.721.500                      Microwaves
-	G01.750.770.776                      Sound
-	G01.750.770.776.567                      Noise
-	G01.750.770.776.783                      Phonons
-	G01.750.770.776.891                      Ultrasonic Waves
-	G01.750.770.776.891.500                      High-Energy Shock Waves
-	G01.750.770.832                      Terahertz Radiation
-	G01.750.897                      Solar Energy
New Tree	<a href="#">G01.828</a> <a href="#">Radioactivity</a>
New Tree	<a href="#">G01.867</a> <a href="#">Scattering, Radiation</a>
New Tree	<a href="#">G01.867.650</a> <a href="#">Neutron Diffraction</a>
New Tree	<a href="#">G01.867.755</a> <a href="#">Scattering, Small Angle</a>
New Tree	<a href="#">G01.867.950</a> <a href="#">X-Ray Diffraction</a>
-	G01.906                      Thermodynamics
-	G01.906.230                      Convection
-	G01.906.345                      Entropy
-	G01.906.595                      Temperature
-	G01.906.595.272                      Cold Temperature
-	G01.906.595.272.437                      Freezing
-	G01.906.595.543                      Hot Temperature
-	G01.906.595.850                      Transition Temperature
-	G01.906.730                      Thermal Conductivity
-	G01.910                      Time
-	G01.910.200                      Chronology as Topic

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	G01.910.405	Half-Life
-	G01.910.645	Periodicity
-	G01.910.645.661	Seasons
-	G01.910.675	Photoperiod
-	G01.910.857	Time Factors
-	G02	Chemical Phenomena
New Tree	<a href="#">G02.010</a>	<a href="#">Absorption</a>
New Tree	<a href="#">G02.010.500</a>	<a href="#">Absorption, Physicochemical</a>
New Tree	<a href="#">G02.010.500.500</a>	<a href="#">Absorption, Radiation</a>
New Tree	<a href="#">G02.020</a>	<a href="#">Adsorption</a>
New Tree	<a href="#">G02.035</a>	<a href="#">Air Ionization</a>
New Tree	<a href="#">G02.050</a>	<a href="#">Anisotropy</a>
-	G02.111	Biochemical Phenomena
New Tree	<a href="#">G02.111.007</a>	<a href="#">Acid-Base Equilibrium</a>
New Tree	<a href="#">G02.111.012</a>	<a href="#">Acylation</a>
New Tree	<a href="#">G02.111.012.052</a>	<a href="#">Acetylation</a>
New Tree	<a href="#">G02.111.012.055</a>	<a href="#">Aminoacylation</a>
New Tree	<a href="#">G02.111.012.055.860</a>	<a href="#">Transfer RNA Aminoacylation</a>
New Tree	<a href="#">G02.111.017</a>	<a href="#">Aerobiosis</a>
New Tree	<a href="#">G02.111.026</a>	<a href="#">Agglutination</a>
New Tree	<a href="#">G02.111.026.500</a>	<a href="#">Hemagglutination</a>
New Tree	<a href="#">G02.111.026.500.500</a>	<a href="#">Hemagglutination, Viral</a>
Old Tree	<a href="#">G02.111.029</a>	<a href="#">Acid-Base Equilibrium</a>
New Tree	<a href="#">G02.111.035</a>	<a href="#">Alkylation</a>
New Tree	<a href="#">G02.111.035.538</a>	<a href="#">Methylation</a>

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	<a href="#">G02.111.035.538.161</a>	DNA Methylation
New Tree	<a href="#">G02.111.044</a>	Allosteric Regulation
New Tree	<a href="#">G02.111.053</a>	Amination
New Tree	<a href="#">G02.111.062</a>	Anaerobiosis
New Tree	<a href="#">G02.111.071</a>	Autotrophic Processes
New Tree	<a href="#">G02.111.071.314</a>	Chemoautotrophic Growth
New Tree	<a href="#">G02.111.071.630</a>	Nitrogen Fixation
-	<a href="#">G02.111.080</a>	Base Composition
New Tree	<a href="#">G02.111.084</a>	Binding, Competitive
New Tree	<a href="#">G02.111.086</a>	Biocatalysis
Old Tree	<a href="#">G02.111.087</a>	Biochemical Processes
Old Tree	<a href="#">G02.111.087.019</a>	Acylation
Old Tree	<a href="#">G02.111.087.019.052</a>	Acetylation
Old Tree	<a href="#">G02.111.087.019.055</a>	Aminoacylation
Old Tree	<a href="#">G02.111.087.019.055.860</a>	Transfer RNA Aminoacylation
Old Tree	<a href="#">G02.111.087.024</a>	Aerobiosis
Old Tree	<a href="#">G02.111.087.026</a>	Agglutination
Old Tree	<a href="#">G02.111.087.026.500</a>	Hemagglutination
Old Tree	<a href="#">G02.111.087.026.500.500</a>	Hemagglutination, Viral
Old Tree	<a href="#">G02.111.087.029</a>	Alkylation
Old Tree	<a href="#">G02.111.087.029.538</a>	Methylation
Old Tree	<a href="#">G02.111.087.029.538.161</a>	DNA Methylation
Old Tree	<a href="#">G02.111.087.040</a>	Allosteric Regulation
Old Tree	<a href="#">G02.111.087.045</a>	Amination
Old Tree	<a href="#">G02.111.087.050</a>	Anaerobiosis
Old Tree	<a href="#">G02.111.087.070</a>	Autotrophic Processes
Old Tree	<a href="#">G02.111.087.070.314</a>	Chemoautotrophic Growth
Old Tree	<a href="#">G02.111.087.070.630</a>	Nitrogen Fixation
Old Tree	<a href="#">G02.111.087.090</a>	Binding, Competitive
Old Tree	<a href="#">G02.111.087.095</a>	Biocatalysis

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	G02.111.087.126 Biosynthetic Pathways
Old Tree	G02.111.087.157 Biotinylation
Old Tree	G02.111.087.160 Carbohydrate Metabolism
Old Tree	G02.111.087.160.249 Fermentation
Old Tree	G02.111.087.160.500 Gluconeogenesis
Old Tree	G02.111.087.160.625 Glycogenolysis
Old Tree	G02.111.087.160.750 Glycolysis
Old Tree	G02.111.087.160.812 Glycosylation
Old Tree	G02.111.087.160.875 Pentose Phosphate Pathway
Old Tree	G02.111.087.160.937 Photosynthesis
Old Tree	G02.111.087.160.937.700 Photophosphorylation
Old Tree	G02.111.087.165 Citric Acid Cycle
Old Tree	G02.111.087.170 Cyclization
Old Tree	G02.111.087.200 Dealkylation
Old Tree	G02.111.087.205 Deamination
Old Tree	G02.111.087.210 Decarboxylation
Old Tree	G02.111.087.217 DNA Cleavage
Old Tree	G02.111.087.218 DNA Methylation
Old Tree	G02.111.087.219 DNA Repair
Old Tree	G02.111.087.219.200 DNA End-Joining Repair
Old Tree	G02.111.087.219.220 DNA Mismatch Repair
Old Tree	G02.111.087.219.700 Recombinational DNA Repair
Old Tree	G02.111.087.219.830 SOS Response (Genetics)
Old Tree	G02.111.087.222 DNA Replication
Old Tree	G02.111.087.222.760 DNA Replication Timing
Old Tree	G02.111.087.222.880 S Phase
Old Tree	G02.111.087.222.940 Telomere Shortening
Old Tree	G02.111.087.225 Down-Regulation
Old Tree	G02.111.087.240 Electron Transport
Old Tree	G02.111.087.242 Energy Transfer
Old Tree	G02.111.087.242.280 Fluorescence Resonance Energy Transfer
Old Tree	G02.111.087.242.400 Linear Energy Transfer
Old Tree	G02.111.087.245 Enzyme Activation
Old Tree	G02.111.087.451 Esterification
Old Tree	G02.111.087.485 Halogenation
Old Tree	G02.111.087.490 Heterotrophic Processes

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	G02.111.087.497 Hydrogenation
Old Tree	G02.111.087.512 Hydroxylation
Old Tree	G02.111.087.520 Lipid Peroxidation
Old Tree	G02.111.087.525 Lipogenesis
Old Tree	G02.111.087.530 Lipolysis
Old Tree	G02.111.087.535 Lipoylation
Old Tree	G02.111.087.560 Molecular Mimicry
Old Tree	G02.111.087.599 Nitrogen Cycle
Old Tree	G02.111.087.612 Nitrosation
Old Tree	G02.111.087.615 Nucleic Acid Denaturation
Old Tree	G02.111.087.620 Nucleic Acid Hybridization
Old Tree	G02.111.087.620.500 Base Pairing
Old Tree	G02.111.087.625 Nucleic Acid Renaturation
Old Tree	G02.111.087.640 Osmosis
Old Tree	G02.111.087.640.500 Electroosmosis
Old Tree	G02.111.087.675 Peptide Biosynthesis
Old Tree	G02.111.087.675.050 Aminoacylation
Old Tree	G02.111.087.675.050.860 Transfer RNA Aminoacylation
Old Tree	G02.111.087.675.333 Peptide Biosynthesis, Nucleic Acid-Independent
Old Tree	G02.111.087.675.871 Protein Biosynthesis
Old Tree	G02.111.087.675.871.200 Frameshifting, Ribosomal
Old Tree	G02.111.087.675.871.640 Peptide Chain Elongation, Translational
Old Tree	G02.111.087.675.871.650 Peptide Chain Initiation, Translational
Old Tree	G02.111.087.675.871.720 Peptide Chain Termination, Translational
Old Tree	G02.111.087.675.871.790 Protein Modification, Translational
Old Tree	G02.111.087.675.871.790.600 Protein Processing, Post-Translational
Old Tree	G02.111.087.675.871.790.600.400 Protein Prenylation
Old Tree	G02.111.087.675.871.790.600.700 Protein Splicing
Old Tree	G02.111.087.675.871.790.600.925 Ubiquitination
Old Tree	G02.111.087.675.871.790.600.925.500 Sumoylation
Old Tree	G02.111.087.675.871.790.600.962 Unfolded Protein Response
Old Tree	G02.111.087.675.871.790.600.962.500 Endoplasmic Reticulum-Associated Degradation
Old Tree	G02.111.087.675.871.850 Transfer RNA Aminoacylation
Old Tree	G02.111.087.677 Phosphorylation
Old Tree	G02.111.087.677.550 Oxidative Phosphorylation

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	G02.111.087.677.605 Photophosphorylation
Old Tree	G02.111.087.678 Phototrophic Processes
Old Tree	G02.111.087.678.700 Photosynthesis
Old Tree	G02.111.087.678.700.700 Photophosphorylation
Old Tree	G02.111.087.679 Prenylation
Old Tree	G02.111.087.679.500 Protein Prenylation
Old Tree	G02.111.087.680 Protein Binding
Old Tree	G02.111.087.682 Protein Carbonylation
Old Tree	G02.111.087.690 Protein Folding
Old Tree	G02.111.087.690.501 Protein Refolding
Old Tree	G02.111.087.690.501.500 Protein Renaturation
Old Tree	G02.111.087.690.750 Protein Unfolding
Old Tree	G02.111.087.690.750.500 Protein Denaturation
Old Tree	G02.111.087.693 Protein Modification, Translational
Old Tree	G02.111.087.693.600 Protein Processing, Post-Translational
Old Tree	G02.111.087.693.600.400 Protein Prenylation
Old Tree	G02.111.087.693.600.700 Protein Splicing
Old Tree	G02.111.087.693.600.775 Ubiquitination
Old Tree	G02.111.087.693.600.775.500 Sumoylation
Old Tree	G02.111.087.693.600.850 Unfolded Protein Response
Old Tree	G02.111.087.693.600.850.500 Endoplasmic Reticulum-Associated Degradation
Old Tree	G02.111.087.697 Protein Multimerization
Old Tree	G02.111.087.710 Proteolysis
Old Tree	G02.111.087.723 RNA Cleavage
Old Tree	G02.111.087.736 RNA Folding
Old Tree	G02.111.087.750 RNA Processing, Post-Transcriptional
Old Tree	G02.111.087.750.112 Nonsense Mediated mRNA Decay
Old Tree	G02.111.087.750.225 RNA 3' End Processing
Old Tree	G02.111.087.750.225.710 Polyadenylation
Old Tree	G02.111.087.750.250 RNA Editing
Old Tree	G02.111.087.750.700 RNA Splicing
Old Tree	G02.111.087.750.700.100 Alternative Splicing
Old Tree	G02.111.087.750.700.750 Trans-Splicing
Old Tree	G02.111.087.800 Signal Transduction
Old Tree	G02.111.087.800.400 Ion Channel Gating

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	G02.111.087.800.480 Light Signal Transduction
Old Tree	G02.111.087.800.480.900 Vision, Ocular
Old Tree	G02.111.087.800.560 MAP Kinase Signaling System
Old Tree	G02.111.087.800.580 Mechanotransduction, Cellular
Old Tree	G02.111.087.800.800 Second Messenger Systems
Old Tree	G02.111.087.800.800.100 Calcium Signaling
Old Tree	G02.111.087.800.800.100.500 Excitation Contraction Coupling
Old Tree	G02.111.087.800.850 Synaptic Transmission
Old Tree	G02.111.087.800.850.500 Postsynaptic Potential Summation
Old Tree	G02.111.087.820 Substrate Cycling
Old Tree	G02.111.087.847 Transcription, Genetic
Old Tree	G02.111.087.847.500 Reverse Transcription
Old Tree	G02.111.087.847.562 Transcription Elongation, Genetic
Old Tree	G02.111.087.847.625 Transcription Initiation, Genetic
Old Tree	G02.111.087.847.687 Transcription Termination, Genetic
Old Tree	G02.111.087.847.750 Transcriptome
Old Tree	G02.111.087.880 Up-Regulation
Old Tree	G02.111.097 Body Composition
Old Tree	G02.111.097.134 Body Fat Distribution
Old Tree	G02.111.097.134.500 Adiposity
New Tree	G02.111.098 Biosynthetic Pathways
Old Tree	G02.111.100 Brain Chemistry
New Tree	G02.111.109 Biotinylation
New Tree	G02.111.130 Body Composition
New Tree	G02.111.130.134 Body Fat Distribution
New Tree	G02.111.130.134.500 Adiposity
New Tree	G02.111.150 Brain Chemistry
New Tree	G02.111.158 Carbohydrate Metabolism
New Tree	G02.111.158.249 Fermentation
New Tree	G02.111.158.500 Gluconeogenesis

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	G02.111.158.625 Glycogenolysis
New Tree	G02.111.158.750 Glycolysis
New Tree	G02.111.158.812 Glycosylation
New Tree	G02.111.158.875 Pentose Phosphate Pathway
New Tree	G02.111.158.937 Photosynthesis
New Tree	G02.111.158.937.700 Photophosphorylation
New Tree	G02.111.165 Citric Acid Cycle
New Tree	G02.111.180 Cyclization
New Tree	G02.111.188 Dealkylation
New Tree	G02.111.192 Deamination
New Tree	G02.111.195 Decarboxylation
New Tree	G02.111.210 DNA Cleavage
New Tree	G02.111.218 DNA Methylation
New Tree	G02.111.222 DNA Repair
New Tree	G02.111.222.200 DNA End-Joining Repair
New Tree	G02.111.222.220 DNA Mismatch Repair
New Tree	G02.111.222.700 Recombinational DNA Repair
New Tree	G02.111.222.830 SOS Response (Genetics)
New Tree	G02.111.225 DNA Replication
New Tree	G02.111.225.760 DNA Replication Timing
New Tree	G02.111.225.880 S Phase
New Tree	G02.111.225.940 Telomere Shortening



## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	G02.111.240	Down-Regulation
New Tree	G02.111.248	Electron Transport
New Tree	G02.111.255	Energy Transfer
New Tree	G02.111.255.280	Fluorescence Resonance Energy Transfer
New Tree	G02.111.255.400	Linear Energy Transfer
New Tree	G02.111.263	Enzyme Activation
New Tree	G02.111.270	Esterification
New Tree	G02.111.323	Halogenation
Old Tree	G02.111.325	Kinetics
New Tree	G02.111.375	Heterotrophic Processes
New Tree	G02.111.380	Hydrogenation
New Tree	G02.111.385	Hydroxylation
New Tree	G02.111.490	Kinetics
New Tree	G02.111.515	Lipid Peroxidation
New Tree	G02.111.528	Lipogenesis
New Tree	G02.111.534	Lipolysis
New Tree	G02.111.540	Lipoylation
-	G02.111.550	Membrane Fluidity
New Tree	G02.111.560	Molecular Mimicry
-	G02.111.570	Molecular Structure
-	G02.111.570.060	Amino Acid Sequence
Old Tree	G02.111.570.060.040	Amino Acid Motifs
Old Tree	G02.111.570.060.040.500	F-Box Motifs
Old Tree	G02.111.570.060.040.750	Immunoreceptor Tyrosine-Based Activation Motif
Old Tree	G02.111.570.060.040.875	Immunoreceptor Tyrosine-Based Inhibition Motif

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	G02.111.570.060.270	Exteins
-	G02.111.570.060.360	Histone Code
-	G02.111.570.060.425	Immunoglobulin Variable Region
-	G02.111.570.060.425.079	Binding Sites, Antibody
-	G02.111.570.060.425.160	Complementarity Determining Regions
-	G02.111.570.060.425.580	Immunoglobulin Idiotypes
-	G02.111.570.060.440	Inteins
-	G02.111.570.060.620	Peptide Library
-	G02.111.570.060.670	Protein Sorting Signals
-	G02.111.570.060.670.600	Nuclear Export Signals
-	G02.111.570.060.670.610	Nuclear Localization Signals
-	G02.111.570.060.720	Repetitive Sequences, Amino Acid
-	G02.111.570.060.720.030	Ankyrin Repeat
New Heading	<b>G02.111.570.060.720.273</b>	<b>Kelch Repeat</b>
New Heading	<b>G02.111.570.060.720.515</b>	<b>WD40 Repeats</b>
-	G02.111.570.080	Base Sequence
-	G02.111.570.080.040	AT Rich Sequence
-	G02.111.570.080.380	GC Rich Sequence
-	G02.111.570.080.380.160	CpG Islands
-	G02.111.570.080.534	Matrix Attachment Regions
-	G02.111.570.080.611	Nucleotide Motifs
-	G02.111.570.080.689	Regulatory Sequences, Nucleic Acid
-	G02.111.570.080.689.330	Enhancer Elements, Genetic
-	G02.111.570.080.689.330.240	E-Box Elements
-	G02.111.570.080.689.330.400	HIV Enhancer
-	G02.111.570.080.689.330.700	Response Elements
-	G02.111.570.080.689.330.700.800	Serum Response Element
-	G02.111.570.080.689.330.700.920	Vitamin D Response Element
-	G02.111.570.080.689.390	Insulator Elements
-	G02.111.570.080.689.450	Locus Control Region
-	G02.111.570.080.689.650	Operator Regions, Genetic
-	G02.111.570.080.689.675	Promoter Regions, Genetic
-	G02.111.570.080.689.675.700	Response Elements
-	G02.111.570.080.689.675.700.040	Antioxidant Response Elements

## MeSH Tree Changes for 2017

Type	Tree - heading
-	G02.111.570.080.689.675.700.800 Serum Response Element
-	G02.111.570.080.689.675.700.920 Vitamin D Response Element
-	G02.111.570.080.689.675.850 TATA Box
-	G02.111.570.080.689.687 Regulatory Sequences, Ribonucleic Acid
-	G02.111.570.080.689.687.061 AU Rich Elements
-	G02.111.570.080.689.687.093 Internal Ribosome Entry Sites
-	G02.111.570.080.689.687.124 Riboswitch
-	G02.111.570.080.689.687.249 RNA 3' Polyadenylation Signals
-	G02.111.570.080.689.687.275 RNA 5' Terminal Oligopyrimidine Sequence
-	G02.111.570.080.689.687.490 RNA Splice Sites
-	G02.111.570.080.689.755 Silencer Elements, Transcriptional
-	G02.111.570.080.689.810 Terminator Regions, Genetic
-	G02.111.570.080.708 Repetitive Sequences, Nucleic Acid
-	G02.111.570.080.708.330 Interspersed Repetitive Sequences
-	G02.111.570.080.708.330.200 DNA Transposable Elements
-	G02.111.570.080.708.330.200.500 Integrons
-	G02.111.570.080.708.330.330 Genomic Islands
-	G02.111.570.080.708.330.800 Retroelements
-	G02.111.570.080.708.330.800.175 Endogenous Retroviruses
-	G02.111.570.080.708.330.800.200 Genes, Intracisternal A-Particle
-	G02.111.570.080.708.330.800.400 Long Interspersed Nucleotide Elements
-	G02.111.570.080.708.330.800.800 Short Interspersed Nucleotide Elements
-	G02.111.570.080.708.330.800.800.050 Alu Elements
-	G02.111.570.080.708.565 Segmental Duplications, Genomic
-	G02.111.570.080.708.800 Tandem Repeat Sequences
-	G02.111.570.080.708.800.140 DNA Repeat Expansion
-	G02.111.570.080.708.800.140.865 Trinucleotide Repeat Expansion
-	G02.111.570.080.708.800.150 DNA, Satellite
-	G02.111.570.080.708.800.325 Inverted Repeat Sequences
-	G02.111.570.080.708.800.325.500 Clustered Regularly Interspaced Short Palindromic Repeats
-	G02.111.570.080.708.800.500 Microsatellite Repeats
-	G02.111.570.080.708.800.500.150 Dinucleotide Repeats
-	G02.111.570.080.708.800.500.850 Trinucleotide Repeats
-	G02.111.570.080.708.800.500.850.200 Trinucleotide Repeat Expansion
-	G02.111.570.080.708.800.550 Minisatellite Repeats

## MeSH Tree Changes for 2017

Type	Tree - heading
-	G02.111.570.080.708.850 Terminal Repeat Sequences
-	G02.111.570.080.708.850.400 HIV Long Terminal Repeat
-	G02.111.570.080.708.850.400.400 HIV Enhancer
-	G02.111.570.120 Binding Sites
-	G02.111.570.120.147 Allosteric Site
-	G02.111.570.120.290 Bay-Region, Polycyclic Aromatic Hydrocarbon
-	G02.111.570.120.309 Binding, Competitive
-	G02.111.570.120.408 Binding Sites, Antibody
-	G02.111.570.120.704 Catalytic Domain
-	G02.111.570.160 Carbohydrate Sequence
-	G02.111.570.580 Conserved Sequence
-	G02.111.570.580.175 Consensus Sequence
-	G02.111.570.580.175.500 Position-Specific Scoring Matrices
-	G02.111.570.685 Isomerism
-	G02.111.570.820 Molecular Conformation
-	G02.111.570.820.117 Bay-Region, Polycyclic Aromatic Hydrocarbon
-	G02.111.570.820.235 Carbohydrate Conformation
-	G02.111.570.820.486 Nucleic Acid Conformation
-	G02.111.570.820.486.100 Base Pairing
-	G02.111.570.820.486.128 DNA, A-Form
-	G02.111.570.820.486.142 DNA, B-Form
-	G02.111.570.820.486.156 DNA, C-Form
-	G02.111.570.820.486.212 DNA, Circular
-	G02.111.570.820.486.212.084 DNA, Catenated
-	G02.111.570.820.486.212.250 DNA, Superhelical
-	G02.111.570.820.486.268 DNA, Concatenated
-	G02.111.570.820.486.325 DNA, Cruciform
-	G02.111.570.820.486.437 DNA, Single-Stranded
-	G02.111.570.820.486.493 DNA, Z-Form
-	G02.111.570.820.486.550 G-Quadruplexes
-	G02.111.570.820.486.662 Nucleotide Motifs
-	G02.111.570.820.486.718 RNA Folding
-	G02.111.570.820.486.775 RNA, Double-Stranded
-	G02.111.570.820.709 Protein Conformation
New Heading	<b>G02.111.570.820.709.275 Protein Structural Elements</b>

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	G02.111.570.820.709.275.500	Amino Acid Motifs
New Tree	G02.111.570.820.709.275.500.030	Ankyrin Repeat
New Tree	G02.111.570.820.709.275.500.050	AT-Hook Motifs
New Tree	G02.111.570.820.709.275.500.127	Cystine Knot Motifs
New Tree	G02.111.570.820.709.275.500.205	F-Box Motifs
New Tree	G02.111.570.820.709.275.500.360	Helix-Loop-Helix Motifs
New Tree	G02.111.570.820.709.275.500.360.240	EF Hand Motifs
New Tree	G02.111.570.820.709.275.500.360.360	Helix-Turn-Helix Motifs
New Heading	<b>G02.111.570.820.709.275.500.360.360.500</b>	<b>ETS Motif</b>
New Tree	G02.111.570.820.709.275.500.440 Motif	Immunoreceptor Tyrosine-Based Activation
New Tree	G02.111.570.820.709.275.500.480 Motif	Immunoreceptor Tyrosine-Based Inhibition
New Heading	<b>G02.111.570.820.709.275.500.500</b>	<b>Kelch Repeat</b>
New Tree	G02.111.570.820.709.275.500.520	Leucine Zippers
New Heading	<b>G02.111.570.820.709.275.500.869</b>	<b>RNA-Binding Motifs</b>
New Heading	<b>G02.111.570.820.709.275.500.869.250</b>	<b>Double-Stranded RNA Binding Motif</b>
New Heading	<b>G02.111.570.820.709.275.500.869.500</b>	<b>RNA Recognition Motif</b>
New Heading	<b>G02.111.570.820.709.275.500.898</b>	<b>Sterile Alpha Motif</b>
New Heading	<b>G02.111.570.820.709.275.500.927</b>	<b>WD40 Repeats</b>
New Tree	G02.111.570.820.709.275.500.985	Zinc Fingers
New Heading	<b>G02.111.570.820.709.275.500.985.250</b>	<b>CYS2-HIS2 Zinc Fingers</b>
New Tree	G02.111.570.820.709.275.500.985.500	RING Finger Domains
New Heading	<b>G02.111.570.820.709.275.750</b>	<b>Protein Domains</b>

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Heading	<b>G02.111.570.820.709.275.750.125</b>	<b>C2 Domains</b>
New Tree	<a href="#">G02.111.570.820.709.275.750.188</a>	<a href="#">Catalytic Domain</a>
New Heading	<b>G02.111.570.820.709.275.750.219</b>	<b>Discoidin Domain</b>
New Tree	<a href="#">G02.111.570.820.709.275.750.235</a>	<a href="#">HMG-Box Domains</a>
New Heading	<b>G02.111.570.820.709.275.750.250</b>	<b>Immunoglobulin Domains</b>
New Heading	<b>G02.111.570.820.709.275.750.250.500</b>	<b>Fibronectin Type III Domain</b>
New Tree	<a href="#">G02.111.570.820.709.275.750.375</a>	<a href="#">Kringles</a>
New Heading	<b>G02.111.570.820.709.275.750.438</b>	<b>Methyl CpG Binding Domain</b>
New Heading	<b>G02.111.570.820.709.275.750.469</b>	<b>Pleckstrin Homology Domains</b>
New Tree	<a href="#">G02.111.570.820.709.275.750.485</a>	<a href="#">Proline-Rich Protein Domains</a>
New Tree	<a href="#">G02.111.570.820.709.275.750.500</a>	<a href="#">Protein Interaction Domains and Motifs</a>
New Tree	<a href="#">G02.111.570.820.709.275.750.500.080</a>	<a href="#">Ankyrin Repeat</a>
New Heading	<b>G02.111.570.820.709.275.750.500.290</b>	<b>B30.2-SPRY Domain</b>
New Heading	<b>G02.111.570.820.709.275.750.500.343</b>	<b>BTB-POZ Domain</b>
New Heading	<b>G02.111.570.820.709.275.750.500.395</b>	<b>Death Domain Superfamily</b>
New Heading	<b>G02.111.570.820.709.275.750.500.395.250</b> <b>Recruitment Domain</b>	<b>Caspase Activation and</b>
New Heading	<b>G02.111.570.820.709.275.750.500.395.500</b>	<b>Death Domain</b>
New Heading	<b>G02.111.570.820.709.275.750.500.395.750</b>	<b>Death Effector Domain</b>
New Heading	<b>G02.111.570.820.709.275.750.500.395.875</b>	<b>Pyrin Domain</b>
New Heading	<b>G02.111.570.820.709.275.750.500.448</b>	<b>Kelch Repeat</b>
New Tree	<a href="#">G02.111.570.820.709.275.750.500.500</a>	<a href="#">PDZ Domains</a>
New Tree	<a href="#">G02.111.570.820.709.275.750.500.625</a>	<a href="#">RING Finger Domains</a>

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	<a href="#">G02.111.570.820.709.275.750.500.750</a>	<a href="#">src Homology Domains</a>
New Heading	<b>G02.111.570.820.709.275.750.500.813</b>	<b>Sterile Alpha Motif</b>
New Heading	<b>G02.111.570.820.709.275.750.500.844</b>	<b>Tudor Domain</b>
New Heading	<b>G02.111.570.820.709.275.750.500.875</b>	<b>WD40 Repeats</b>
New Tree	<a href="#">G02.111.570.820.709.275.875</a>	<a href="#">Repetitive Sequences, Amino Acid</a>
New Tree	<a href="#">G02.111.570.820.709.275.875.030</a>	<a href="#">Ankyrin Repeat</a>
New Heading	<b>G02.111.570.820.709.275.875.273</b>	<b>Kelch Repeat</b>
New Heading	<b>G02.111.570.820.709.275.875.515</b>	<b>WD40 Repeats</b>
-	G02.111.570.820.709.550	Protein Structure, Quaternary
-	G02.111.570.820.709.600	Protein Structure, Secondary
New Heading	<b>G02.111.570.820.709.600.020</b>	<b>Protein Conformation, alpha-Helical</b>
Old Tree	<a href="#">G02.111.570.820.709.600.040</a>	<a href="#">Amino Acid Motifs</a>
Old Tree	<a href="#">G02.111.570.820.709.600.040.030</a>	<a href="#">Ankyrin Repeat</a>
Old Tree	<a href="#">G02.111.570.820.709.600.040.050</a>	<a href="#">AT-Hook Motifs</a>
Old Tree	<a href="#">G02.111.570.820.709.600.040.127</a>	<a href="#">Cystine Knot Motifs</a>
Old Tree	<a href="#">G02.111.570.820.709.600.040.205</a>	<a href="#">F-Box Motifs</a>
Old Tree	<a href="#">G02.111.570.820.709.600.040.360</a>	<a href="#">Helix-Loop-Helix Motifs</a>
Old Tree	<a href="#">G02.111.570.820.709.600.040.360.240</a>	<a href="#">EF Hand Motifs</a>
Old Tree	<a href="#">G02.111.570.820.709.600.040.360.360</a>	<a href="#">Helix-Turn-Helix Motifs</a>
Old Tree	<a href="#">G02.111.570.820.709.600.040.440 Motif</a>	<a href="#">Immunoreceptor Tyrosine-Based Activation</a>
Old Tree	<a href="#">G02.111.570.820.709.600.040.480 Motif</a>	<a href="#">Immunoreceptor Tyrosine-Based Inhibition</a>
Old Tree	<a href="#">G02.111.570.820.709.600.040.520</a>	<a href="#">Leucine Zippers</a>
Old Tree	<a href="#">G02.111.570.820.709.600.040.752</a>	<a href="#">Proline-Rich Protein Domains</a>
Old Tree	<a href="#">G02.111.570.820.709.600.040.985</a>	<a href="#">Zinc Fingers</a>
Old Tree	<a href="#">G02.111.570.820.709.600.040.985.500</a>	<a href="#">RING Finger Domains</a>
New Tree	<a href="#">G02.111.570.820.709.600.500</a>	<a href="#">Amino Acid Motifs</a>
New Heading	<b>G02.111.570.820.709.600.750</b>	<b>Protein Conformation, beta-Strand</b>

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	G02.111.570.820.709.610	Protein Structure, Tertiary
Old Tree	<del>G02.111.570.820.709.610.189</del>	<del>Catalytic Domain</del>
Old Tree	<del>G02.111.570.820.709.610.380</del>	<del>HMG-Box Domains</del>
Old Tree	<del>G02.111.570.820.709.610.480</del>	<del>Kringles</del>
New Heading	<b>G02.111.570.820.709.610.500</b>	<b>Protein Domains</b>
Old Tree	<del>G02.111.570.820.709.610.640</del>	<del>Protein Interaction Domains and Motifs</del>
Old Tree	<del>G02.111.570.820.709.610.640.080</del>	<del>Ankyrin Repeat</del>
Old Tree	<del>G02.111.570.820.709.610.640.500</del>	<del>PDZ Domains</del>
Old Tree	<del>G02.111.570.820.709.610.640.625</del>	<del>RING Finger Domains</del>
Old Tree	<del>G02.111.570.820.709.610.640.750</del>	<del>src Homology Domains</del>
-	G02.111.570.820.709.805	Structural Homology, Protein
-	G02.111.570.895	Molecular Dynamics Simulation
New Tree	<a href="#">G02.111.587</a>	<a href="#">Nitrogen Cycle</a>
New Tree	<a href="#">G02.111.587.250</a>	<a href="#">Denitrification</a>
New Tree	<a href="#">G02.111.587.500</a>	<a href="#">Nitrification</a>
New Tree	<a href="#">G02.111.587.750</a>	<a href="#">Nitrogen Fixation</a>
New Tree	<a href="#">G02.111.595</a>	<a href="#">Nitrosation</a>
New Tree	<a href="#">G02.111.603</a>	<a href="#">Nucleic Acid Denaturation</a>
New Tree	<a href="#">G02.111.611</a>	<a href="#">Nucleic Acid Hybridization</a>
New Tree	<a href="#">G02.111.611.500</a>	<a href="#">Base Pairing</a>
New Tree	<a href="#">G02.111.619</a>	<a href="#">Nucleic Acid Renaturation</a>
-	G02.111.635	Osmoregulation
-	G02.111.635.500	Water-Electrolyte Balance
-	G02.111.635.500.500	Kallikrein-Kinin System
-	G02.111.635.500.750	Water Loss, Insensible
New Tree	<a href="#">G02.111.655</a>	<a href="#">Osmosis</a>
New Tree	<a href="#">G02.111.655.500</a>	<a href="#">Electroosmosis</a>
New	<a href="#">G02.111.660</a>	<a href="#">Peptide Biosynthesis</a>



## MeSH Tree Changes for 2017

Type	Tree - heading
Tree	
New Tree	G02.111.660.050 Aminoacylation
New Tree	G02.111.660.050.860 Transfer RNA Aminoacylation
New Tree	G02.111.660.333 Peptide Biosynthesis, Nucleic Acid-Independent
New Tree	G02.111.660.871 Protein Biosynthesis
New Tree	G02.111.660.871.200 Frameshifting, Ribosomal
New Tree	G02.111.660.871.640 Peptide Chain Elongation, Translational
New Tree	G02.111.660.871.650 Peptide Chain Initiation, Translational
New Tree	G02.111.660.871.720 Peptide Chain Termination, Translational
New Tree	G02.111.660.871.790 Protein Modification, Translational
New Tree	G02.111.660.871.790.600 Protein Processing, Post-Translational
New Tree	G02.111.660.871.790.600.400 Protein Prenylation
New Tree	G02.111.660.871.790.600.700 Protein Splicing
New Tree	G02.111.660.871.790.600.925 Ubiquitination
New Tree	G02.111.660.871.790.600.925.500 Sumoylation
New Tree	G02.111.660.871.790.600.962 Unfolded Protein Response
New Tree	G02.111.660.871.790.600.962.500 Endoplasmic Reticulum-Associated Degradation
New Tree	G02.111.660.871.850 Transfer RNA Aminoacylation
New Tree	G02.111.665 Phosphorylation
New Tree	G02.111.665.550 Oxidative Phosphorylation
New Tree	G02.111.665.605 Photophosphorylation
Old Tree	G02.111.668 Protein Aggregation, Pathological
New	G02.111.669 Phototrophic Processes

## MeSH Tree Changes for 2017

Type	Tree - heading
Tree	
New Tree	G02.111.669.700                      Photosynthesis
New Tree	G02.111.669.700.700                      Photophosphorylation
New Tree	G02.111.672                      Prenylation
New Tree	G02.111.672.500                      Protein Prenylation
New Tree	G02.111.675                      Protein Aggregation, Pathological
New Tree	G02.111.679                      Protein Binding
New Tree	G02.111.682                      Protein Carbonylation
New Tree	G02.111.688                      Protein Folding
New Tree	G02.111.688.501                      Protein Refolding
New Tree	G02.111.688.501.500                      Protein Renaturation
New Tree	G02.111.688.750                      Protein Unfolding
New Tree	G02.111.688.750.500                      Protein Denaturation
New Tree	G02.111.691                      Protein Modification, Translational
New Tree	G02.111.691.600                      Protein Processing, Post-Translational
New Tree	G02.111.691.600.400                      Protein Prenylation
New Tree	G02.111.691.600.700                      Protein Splicing
New Tree	G02.111.691.600.775                      Ubiquitination
New Tree	G02.111.691.600.775.500                      Sumoylation
New Tree	G02.111.691.600.850                      Unfolded Protein Response
New Tree	G02.111.691.600.850.500                      Endoplasmic Reticulum-Associated Degradation
New Tree	G02.111.694                      Protein Multimerization

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	G02.111.700	Protein Stability
-	G02.111.700.500	Enzyme Stability
New Tree	<a href="#">G02.111.720</a>	<a href="#">Proteolysis</a>
New Tree	<a href="#">G02.111.740</a>	<a href="#">RNA Cleavage</a>
New Tree	<a href="#">G02.111.750</a>	<a href="#">RNA Folding</a>
New Tree	<a href="#">G02.111.760</a>	<a href="#">RNA Processing, Post-Transcriptional</a>
New Tree	<a href="#">G02.111.760.112</a>	<a href="#">Nonsense Mediated mRNA Decay</a>
New Tree	<a href="#">G02.111.760.225</a>	<a href="#">RNA 3' End Processing</a>
New Tree	<a href="#">G02.111.760.225.710</a>	<a href="#">Polyadenylation</a>
New Tree	<a href="#">G02.111.760.250</a>	<a href="#">RNA Editing</a>
New Tree	<a href="#">G02.111.760.700</a>	<a href="#">RNA Splicing</a>
New Tree	<a href="#">G02.111.760.700.100</a>	<a href="#">Alternative Splicing</a>
New Tree	<a href="#">G02.111.760.700.750</a>	<a href="#">Trans-Splicing</a>
-	G02.111.780	RNA Stability
-	G02.111.780.500	Nonsense Mediated mRNA Decay
-	G02.111.810	Sequence Homology
-	G02.111.810.200	Sequence Homology, Amino Acid
-	G02.111.810.200.820	Structural Homology, Protein
-	G02.111.810.550	Sequence Homology, Nucleic Acid
-	G02.111.810.550.830	Synteny
New Tree	<a href="#">G02.111.820</a>	<a href="#">Signal Transduction</a>
New Tree	<a href="#">G02.111.820.400</a>	<a href="#">Ion Channel Gating</a>
New Tree	<a href="#">G02.111.820.480</a>	<a href="#">Light Signal Transduction</a>
New Tree	<a href="#">G02.111.820.480.900</a>	<a href="#">Vision, Ocular</a>
New Tree	<a href="#">G02.111.820.560</a>	<a href="#">MAP Kinase Signaling System</a>
New	<a href="#">G02.111.820.580</a>	<a href="#">Mechanotransduction, Cellular</a>

## MeSH Tree Changes for 2017

Type	Tree - heading	
Tree		
New Heading	<b>G02.111.820.690</b>	<b>Non-Neuronal Cholinergic System</b>
New Tree	G02.111.820.800	Second Messenger Systems
New Tree	G02.111.820.800.100	Calcium Signaling
New Tree	G02.111.820.800.100.500	Excitation Contraction Coupling
New Tree	G02.111.820.850	Synaptic Transmission
New Tree	G02.111.820.850.500	Postsynaptic Potential Summation
New Tree	G02.111.820.925	Wnt Signaling Pathway
-	G02.111.830	Structure-Activity Relationship
-	G02.111.830.500	Quantitative Structure-Activity Relationship
New Tree	G02.111.833	Substrate Cycling
-	G02.111.835	Substrate Specificity
-	G02.111.840	Th1-Th2 Balance
New Tree	G02.111.873	Transcription, Genetic
New Tree	G02.111.873.500	Reverse Transcription
New Tree	G02.111.873.562	Transcription Elongation, Genetic
New Tree	G02.111.873.625	Transcription Initiation, Genetic
New Tree	G02.111.873.687	Transcription Termination, Genetic
New Tree	G02.111.873.750	Transcriptome
New Tree	G02.111.905	Up-Regulation
New Tree	G02.130	Catalysis
New Tree	G02.130.500	Biocatalysis
Old Tree	G02.149	Chemical Processes
Old Tree	G02.149.115	Biochemical Processes
Old Tree	G02.149.115.019	Acylation

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	G02.149.115.019.052 Acetylation
Old Tree	G02.149.115.019.055 Aminoacylation
Old Tree	G02.149.115.019.055.860 Transfer RNA Aminoacylation
Old Tree	G02.149.115.024 Aerobiosis
Old Tree	G02.149.115.026 Agglutination
Old Tree	G02.149.115.026.500 Hemagglutination
Old Tree	G02.149.115.026.500.500 Hemagglutination, Viral
Old Tree	G02.149.115.029 Alkylation
Old Tree	G02.149.115.029.538 Methylation
Old Tree	G02.149.115.029.538.161 DNA Methylation
Old Tree	G02.149.115.040 Allosteric Regulation
Old Tree	G02.149.115.045 Amination
Old Tree	G02.149.115.050 Anaerobiosis
Old Tree	G02.149.115.070 Autotrophic Processes
Old Tree	G02.149.115.070.314 Chemoautotrophic Growth
Old Tree	G02.149.115.070.630 Nitrogen Fixation
Old Tree	G02.149.115.090 Binding, Competitive
Old Tree	G02.149.115.095 Biocatalysis
Old Tree	G02.149.115.126 Biosynthetic Pathways
Old Tree	G02.149.115.157 Biotinylation
Old Tree	G02.149.115.160 Carbohydrate Metabolism
Old Tree	G02.149.115.160.249 Fermentation
Old Tree	G02.149.115.160.500 Gluconeogenesis
Old Tree	G02.149.115.160.625 Glycogenolysis
Old Tree	G02.149.115.160.750 Glycolysis
Old Tree	G02.149.115.160.812 Glycosylation
Old Tree	G02.149.115.160.875 Pentose Phosphate Pathway
Old Tree	G02.149.115.160.937 Photosynthesis
Old Tree	G02.149.115.160.937.700 Photophosphorylation
Old Tree	G02.149.115.165 Citric Acid Cycle
Old Tree	G02.149.115.170 Cyclization
Old Tree	G02.149.115.200 Dealkylation
Old Tree	G02.149.115.205 Deamination
Old Tree	G02.149.115.210 Decarboxylation
Old Tree	G02.149.115.217 DNA Cleavage
Old Tree	G02.149.115.218 DNA Methylation

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	G02.149.115.219 DNA Repair
Old Tree	G02.149.115.219.200 DNA End-Joining Repair
Old Tree	G02.149.115.219.220 DNA Mismatch Repair
Old Tree	G02.149.115.219.700 Recombinational DNA Repair
Old Tree	G02.149.115.219.830 SOS Response (Genetics)
Old Tree	G02.149.115.222 DNA Replication
Old Tree	G02.149.115.222.760 DNA Replication Timing
Old Tree	G02.149.115.222.880 S Phase
Old Tree	G02.149.115.222.940 Telomere Shortening
Old Tree	G02.149.115.225 Down-Regulation
Old Tree	G02.149.115.240 Electron Transport
Old Tree	G02.149.115.242 Energy Transfer
Old Tree	G02.149.115.242.280 Fluorescence Resonance Energy Transfer
Old Tree	G02.149.115.242.400 Linear Energy Transfer
Old Tree	G02.149.115.245 Enzyme Activation
Old Tree	G02.149.115.451 Esterification
Old Tree	G02.149.115.485 Halogenation
Old Tree	G02.149.115.490 Heterotrophic Processes
Old Tree	G02.149.115.497 Hydrogenation
Old Tree	G02.149.115.512 Hydroxylation
Old Tree	G02.149.115.520 Lipid Peroxidation
Old Tree	G02.149.115.525 Lipogenesis
Old Tree	G02.149.115.530 Lipolysis
Old Tree	G02.149.115.535 Lipoylation
Old Tree	G02.149.115.560 Molecular Mimicry
Old Tree	G02.149.115.599 Nitrogen Cycle
Old Tree	G02.149.115.612 Nitrosation
Old Tree	G02.149.115.615 Nucleic Acid Denaturation
Old Tree	G02.149.115.620 Nucleic Acid Hybridization
Old Tree	G02.149.115.620.500 Base Pairing
Old Tree	G02.149.115.625 Nucleic Acid Renaturation
Old Tree	G02.149.115.640 Osmosis
Old Tree	G02.149.115.640.500 Electroosmosis
Old Tree	G02.149.115.675 Peptide Biosynthesis
Old Tree	G02.149.115.675.050 Aminoacylation
Old Tree	G02.149.115.675.050.860 Transfer RNA Aminoacylation

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	G02.149.115.675.333 Peptide Biosynthesis, Nucleic Acid-Independent
Old Tree	G02.149.115.675.871 Protein Biosynthesis
Old Tree	G02.149.115.675.871.200 Frameshifting, Ribosomal
Old Tree	G02.149.115.675.871.640 Peptide Chain Elongation, Translational
Old Tree	G02.149.115.675.871.650 Peptide Chain Initiation, Translational
Old Tree	G02.149.115.675.871.720 Peptide Chain Termination, Translational
Old Tree	G02.149.115.675.871.790 Protein Modification, Translational
Old Tree	G02.149.115.675.871.790.600 Protein Processing, Post-Translational
Old Tree	G02.149.115.675.871.790.600.400 Protein Prenylation
Old Tree	G02.149.115.675.871.790.600.700 Protein Splicing
Old Tree	G02.149.115.675.871.790.600.831 Ubiquitination
Old Tree	G02.149.115.675.871.790.600.831.500 Sumoylation
Old Tree	G02.149.115.675.871.790.600.850 Unfolded Protein Response
Old Tree	G02.149.115.675.871.790.600.850.500 Endoplasmic Reticulum-Associated Degradation
Old Tree	G02.149.115.675.871.850 Transfer RNA Aminoacylation
Old Tree	G02.149.115.677 Phosphorylation
Old Tree	G02.149.115.677.550 Oxidative Phosphorylation
Old Tree	G02.149.115.677.605 Photophosphorylation
Old Tree	G02.149.115.678 Phototrophic Processes
Old Tree	G02.149.115.678.700 Photosynthesis
Old Tree	G02.149.115.678.700.700 Photophosphorylation
Old Tree	G02.149.115.679 Prenylation
Old Tree	G02.149.115.679.500 Protein Prenylation
Old Tree	G02.149.115.680 Protein Binding
Old Tree	G02.149.115.682 Protein Carbonylation
Old Tree	G02.149.115.690 Protein Folding
Old Tree	G02.149.115.690.501 Protein Refolding
Old Tree	G02.149.115.690.501.500 Protein Renaturation
Old Tree	G02.149.115.690.750 Protein Unfolding
Old Tree	G02.149.115.690.750.500 Protein Denaturation
Old Tree	G02.149.115.693 Protein Modification, Translational
Old Tree	G02.149.115.693.600 Protein Processing, Post-Translational
Old Tree	G02.149.115.693.600.400 Protein Prenylation
Old Tree	G02.149.115.693.600.700 Protein Splicing
Old Tree	G02.149.115.693.600.850 Ubiquitination

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	G02.149.115.693.600.850.500 Sumoylation
Old Tree	G02.149.115.693.600.925 Unfolded Protein Response
Old Tree	G02.149.115.693.600.925.500 Endoplasmic Reticulum-Associated Degradation
Old Tree	G02.149.115.697 Protein Multimerization
Old Tree	G02.149.115.710 Proteolysis
Old Tree	G02.149.115.723 RNA Cleavage
Old Tree	G02.149.115.736 RNA Folding
Old Tree	G02.149.115.750 RNA Processing, Post-Transcriptional
Old Tree	G02.149.115.750.112 Nonsense Mediated mRNA Decay
Old Tree	G02.149.115.750.225 RNA 3' End Processing
Old Tree	G02.149.115.750.225.710 Polyadenylation
Old Tree	G02.149.115.750.250 RNA Editing
Old Tree	G02.149.115.750.700 RNA Splicing
Old Tree	G02.149.115.750.700.100 Alternative Splicing
Old Tree	G02.149.115.750.700.750 Trans-Splicing
Old Tree	G02.149.115.800 Signal Transduction
Old Tree	G02.149.115.800.400 Ion Channel Gating
Old Tree	G02.149.115.800.480 Light Signal Transduction
Old Tree	G02.149.115.800.480.900 Vision, Ocular
Old Tree	G02.149.115.800.560 MAP Kinase Signaling System
Old Tree	G02.149.115.800.580 Mechanotransduction, Cellular
Old Tree	G02.149.115.800.800 Second Messenger Systems
Old Tree	G02.149.115.800.800.100 Calcium Signaling
Old Tree	G02.149.115.800.850 Synaptic Transmission
Old Tree	G02.149.115.800.850.500 Postsynaptic Potential Summation
Old Tree	G02.149.115.800.925 Wnt Signaling Pathway
Old Tree	G02.149.115.820 Substrate Cycling
Old Tree	G02.149.115.847 Transcription, Genetic
Old Tree	G02.149.115.847.500 Reverse Transcription
Old Tree	G02.149.115.847.562 Transcription Elongation, Genetic
Old Tree	G02.149.115.847.625 Transcription Initiation, Genetic
Old Tree	G02.149.115.847.687 Transcription Termination, Genetic
Old Tree	G02.149.115.847.750 Transcriptome
Old Tree	G02.149.115.880 Up-Regulation
Old Tree	G02.149.170 Catalysis



## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	G02.149.170.500 Biocatalysis
Old Tree	G02.149.200 Corrosion
Old Tree	G02.149.250 Dimerization
Old Tree	G02.149.260 Evolution, Chemical
Old Tree	G02.149.395 Hydrolysis
Old Tree	G02.149.465 Organic Chemistry Processes
Old Tree	G02.149.465.030 Acylation
Old Tree	G02.149.465.030.052 Acetylation
Old Tree	G02.149.465.030.526 Aminoacylation
Old Tree	G02.149.465.040 Alkylation
Old Tree	G02.149.465.040.538 Methylation
Old Tree	G02.149.465.050 Amination
Old Tree	G02.149.465.190 Cyclization
Old Tree	G02.149.465.200 Dealkylation
Old Tree	G02.149.465.210 Deamination
Old Tree	G02.149.465.220 Decarboxylation
Old Tree	G02.149.465.270 Esterification
Old Tree	G02.149.465.350 Glycosylation
Old Tree	G02.149.465.400 Hydrogenation
Old Tree	G02.149.465.410 Hydroxylation
Old Tree	G02.149.465.500 Maillard Reaction
Old Tree	G02.149.465.620 Nitrosation
Old Tree	G02.149.465.700 Phosphorylation
Old Tree	G02.149.767 Physicochemical Processes
Old Tree	G02.149.767.019 Absorption
Old Tree	G02.149.767.019.500 Absorption, Physicochemical
Old Tree	G02.149.767.019.500.500 Absorption, Radiation
Old Tree	G02.149.767.029 Adsorption
Old Tree	G02.149.767.040 Air Ionization
Old Tree	G02.149.767.500 Catalysis
Old Tree	G02.149.767.500.500 Biocatalysis
Old Tree	G02.149.767.502 Chemical Precipitation
Old Tree	G02.149.767.502.347 Flocculation
Old Tree	G02.149.767.505 Crystallization
Old Tree	G02.149.767.530 Desiccation
Old Tree	G02.149.767.540 Dialysis

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	G02.149.767.560 Diffusion
Old Tree	G02.149.767.560.500 Facilitated Diffusion
Old Tree	G02.149.767.560.750 Thermal Diffusion
Old Tree	G02.149.767.580 Energy Transfer
Old Tree	G02.149.767.580.400 Linear Energy Transfer
Old Tree	G02.149.767.600 Filtration
Old Tree	G02.149.767.600.807 Ultrafiltration
Old Tree	G02.149.767.620 Hydrogen Bonding
Old Tree	G02.149.767.640 Ion Exchange
Old Tree	G02.149.767.642 Neutron Diffraction
Old Tree	G02.149.767.645 Osmosis
Old Tree	G02.149.767.645.250 Electroosmosis
Old Tree	G02.149.767.650 Oxidation-Reduction
Old Tree	G02.149.767.680 Phase Transition
Old Tree	G02.149.767.680.466 Freezing
Old Tree	G02.149.767.680.933 Volatilization
Old Tree	G02.149.767.690 Photochemical Processes
Old Tree	G02.149.767.690.680 Photobleaching
Old Tree	G02.149.767.690.685 Photolysis
Old Tree	G02.149.767.690.842 Photophosphorylation
Old Tree	G02.149.767.690.921 Photosynthesis
Old Tree	G02.149.767.710 Proton-Motive Force
Old Tree	G02.149.767.830 Spontaneous Combustion
Old Tree	G02.149.767.915 X-Ray Diffraction
Old Tree	G02.149.883 Polymerization
New Tree	G02.159 Chemical Precipitation
New Tree	G02.159.347 Flocculation
New Tree	G02.165 Corrosion
New Tree	G02.171 Crystallization
New Tree	G02.176 Desiccation
New Tree	G02.186 Dialysis
New	G02.196 Diffusion

## MeSH Tree Changes for 2017

Type	Tree - heading	
Tree		
New Tree	<a href="#">G02.196.500</a>	<a href="#">Facilitated Diffusion</a>
New Tree	<a href="#">G02.196.750</a>	<a href="#">Thermal Diffusion</a>
New Tree	<a href="#">G02.206</a>	<a href="#">Dimerization</a>
New Tree	<a href="#">G02.211</a>	<a href="#">Drug Liberation</a>
New Tree	<a href="#">G02.216</a>	<a href="#">Energy Transfer</a>
New Tree	<a href="#">G02.216.400</a>	<a href="#">Linear Energy Transfer</a>
New Tree	<a href="#">G02.225</a>	<a href="#">Evolution, Chemical</a>
New Tree	<a href="#">G02.263</a>	<a href="#">Filtration</a>
New Tree	<a href="#">G02.263.807</a>	<a href="#">Ultrafiltration</a>
New Tree	<a href="#">G02.282</a>	<a href="#">Hydrogen Bonding</a>
-	<a href="#">G02.300</a>	<a href="#">Hydrogen-Ion Concentration</a>
-	<a href="#">G02.300.176</a>	<a href="#">Acid-Base Equilibrium</a>
-	<a href="#">G02.300.500</a>	<a href="#">Isoelectric Point</a>
New Tree	<a href="#">G02.380</a>	<a href="#">Hydrolysis</a>
New Tree	<a href="#">G02.409</a>	<a href="#">Hydrophobic and Hydrophilic Interactions</a>
New Tree	<a href="#">G02.409.500</a>	<a href="#">Wettability</a>
New Tree	<a href="#">G02.437</a>	<a href="#">Ion Exchange</a>
New Tree	<a href="#">G02.466</a>	<a href="#">Molecular Structure</a>
New Tree	<a href="#">G02.494</a>	<a href="#">Molecular Weight</a>
New Tree	<a href="#">G02.551</a>	<a href="#">Neutron Diffraction</a>
-	<a href="#">G02.607</a>	<a href="#">Organic Chemistry Phenomena</a>
New Tree	<a href="#">G02.607.063</a>	<a href="#">Acylation</a>
New Tree	<a href="#">G02.607.063.052</a>	<a href="#">Acetylation</a>

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	<a href="#">G02.607.063.526</a>	<a href="#">Aminoacylation</a>
New Tree	<a href="#">G02.607.094</a>	<a href="#">Alkylation</a>
New Tree	<a href="#">G02.607.094.538</a>	<a href="#">Methylation</a>
New Tree	<a href="#">G02.607.110</a>	<a href="#">Amination</a>
New Tree	<a href="#">G02.607.125</a>	<a href="#">Carbon Cycle</a>
New Tree	<a href="#">G02.607.125.500</a>	<a href="#">Carbon Sequestration</a>
New Tree	<a href="#">G02.607.133</a>	<a href="#">Cyclization</a>
New Tree	<a href="#">G02.607.141</a>	<a href="#">Dealkylation</a>
New Tree	<a href="#">G02.607.157</a>	<a href="#">Deamination</a>
New Tree	<a href="#">G02.607.188</a>	<a href="#">Decarboxylation</a>
New Tree	<a href="#">G02.607.250</a>	<a href="#">Esterification</a>
New Tree	<a href="#">G02.607.299</a>	<a href="#">Glycosylation</a>
New Tree	<a href="#">G02.607.324</a>	<a href="#">Hydrogenation</a>
New Tree	<a href="#">G02.607.348</a>	<a href="#">Hydroxylation</a>
New Tree	<a href="#">G02.607.445</a>	<a href="#">Isomerism</a>
New Tree	<a href="#">G02.607.445.682</a>	<a href="#">Stereoisomerism</a>
Old Tree	<a href="#">G02.607.500</a>	<a href="#">Isomerism</a>
Old Tree	<a href="#">G02.607.500.682</a>	<a href="#">Stereoisomerism</a>
New Tree	<a href="#">G02.607.522</a>	<a href="#">Maillard Reaction</a>
New Tree	<a href="#">G02.607.560</a>	<a href="#">Nitrogen Cycle</a>
New Tree	<a href="#">G02.607.560.250</a>	<a href="#">Denitrification</a>
New Tree	<a href="#">G02.607.560.500</a>	<a href="#">Nitrification</a>
New	<a href="#">G02.607.560.750</a>	<a href="#">Nitrogen Fixation</a>

## MeSH Tree Changes for 2017

Type	Tree - heading	
Tree		
New Tree	<a href="#">G02.607.598</a>	Nitrosation
Old Tree	<a href="#">G02.607.750</a>	Organic Chemistry Processes
Old Tree	<a href="#">G02.607.750.030</a>	Acylation
Old Tree	<a href="#">G02.607.750.030.052</a>	Acetylation
Old Tree	<a href="#">G02.607.750.030.526</a>	Aminoacylation
Old Tree	<a href="#">G02.607.750.040</a>	Alkylation
Old Tree	<a href="#">G02.607.750.040.538</a>	Methylation
Old Tree	<a href="#">G02.607.750.050</a>	Amination
Old Tree	<a href="#">G02.607.750.120</a>	Carbon Cycle
Old Tree	<a href="#">G02.607.750.155</a>	Chemistry Techniques, Synthetic
Old Tree	<a href="#">G02.607.750.155.124</a>	Click Chemistry
Old Tree	<a href="#">G02.607.750.155.249</a>	Combinatorial Chemistry Techniques
Old Tree	<a href="#">G02.607.750.155.249.500</a>	SELEX Aptamer Technique
Old Tree	<a href="#">G02.607.750.155.374</a>	Cycloaddition Reaction
Old Tree	<a href="#">G02.607.750.155.500</a>	Solid-Phase Synthesis Techniques
Old Tree	<a href="#">G02.607.750.190</a>	Cyclization
Old Tree	<a href="#">G02.607.750.200</a>	Dealkylation
Old Tree	<a href="#">G02.607.750.210</a>	Deamination
Old Tree	<a href="#">G02.607.750.220</a>	Decarboxylation
Old Tree	<a href="#">G02.607.750.270</a>	Esterification
Old Tree	<a href="#">G02.607.750.350</a>	Glycosylation
Old Tree	<a href="#">G02.607.750.400</a>	Hydrogenation
Old Tree	<a href="#">G02.607.750.410</a>	Hydroxylation
Old Tree	<a href="#">G02.607.750.500</a>	Maillard Reaction
Old Tree	<a href="#">G02.607.750.560</a>	Nitrogen Cycle
Old Tree	<a href="#">G02.607.750.620</a>	Nitrosation
Old Tree	<a href="#">G02.607.750.660</a>	Oxidative Coupling
Old Tree	<a href="#">G02.607.750.700</a>	Phosphorylation
New Tree	<a href="#">G02.607.765</a>	Oxidative Coupling
New Tree	<a href="#">G02.607.780</a>	Phosphorylation
-	G02.640	Osmolar Concentration
-	G02.640.249	Osmotic Pressure
-	G02.640.500	Salinity

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	G02.691	Osmosis
New Tree	G02.691.250	Electroosmosis
New Tree	G02.700	Oxidation-Reduction
New Tree	G02.700.500	Oxidative Coupling
New Tree	G02.712	Particle Size
New Tree	G02.723	Permeability
New Tree	G02.723.495	Osmosis
New Tree	G02.723.495.250	Electroosmosis
New Tree	G02.723.661	Osmotic Pressure
New Tree	G02.734	Phase Transition
New Tree	G02.734.466	Freezing
New Tree	G02.734.933	Volatilization
New Tree	G02.740	Photochemical Processes
New Tree	G02.740.680	Photobleaching
New Tree	G02.740.685	Photolysis
New Tree	G02.740.842	Photophosphorylation
New Tree	G02.740.921	Photosynthesis
New Tree	G02.750	Polymerization
New Tree	G02.765	Proton-Motive Force
New Tree	G02.805	Solubility
New Tree	G02.819	Specific Gravity
New Tree	G02.833	Spontaneous Combustion

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	G02.842 Physicochemical Phenomena
Old Tree	G02.842.050 Anisotropy
Old Tree	G02.842.312 Hydrophobic and Hydrophilic Interactions
Old Tree	G02.842.312.500 Wettability
Old Tree	G02.842.575 Molecular Structure
Old Tree	G02.842.580 Molecular Weight
Old Tree	G02.842.680 Particle Size
Old Tree	G02.842.700 Permeability
Old Tree	G02.842.700.495 Osmosis
Old Tree	G02.842.700.495.250 Electroosmosis
Old Tree	G02.842.700.661 Osmotic Pressure
Old Tree	G02.842.750 Physicochemical Processes
Old Tree	G02.842.750.019 Absorption
Old Tree	G02.842.750.019.500 Absorption, Physicochemical
Old Tree	G02.842.750.019.500.500 Absorption, Radiation
Old Tree	G02.842.750.029 Adsorption
Old Tree	G02.842.750.040 Air Ionization
Old Tree	G02.842.750.270 Carbon Cycle
Old Tree	G02.842.750.270.500 Carbon Sequestration
Old Tree	G02.842.750.500 Catalysis
Old Tree	G02.842.750.500.500 Biocatalysis
Old Tree	G02.842.750.502 Chemical Precipitation
Old Tree	G02.842.750.502.347 Flocculation
Old Tree	G02.842.750.505 Crystallization
Old Tree	G02.842.750.530 Desiccation
Old Tree	G02.842.750.540 Dialysis
Old Tree	G02.842.750.560 Diffusion
Old Tree	G02.842.750.560.500 Facilitated Diffusion
Old Tree	G02.842.750.560.750 Thermal Diffusion
Old Tree	G02.842.750.570 Drug Liberation
Old Tree	G02.842.750.580 Energy Transfer
Old Tree	G02.842.750.580.400 Linear Energy Transfer
Old Tree	G02.842.750.600 Filtration
Old Tree	G02.842.750.600.807 Ultrafiltration
Old Tree	G02.842.750.620 Hydrogen Bonding
Old Tree	G02.842.750.640 Ion Exchange

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	G02.842.750.642 Neutron Diffraction
Old Tree	G02.842.750.643 Nitrogen Cycle
Old Tree	G02.842.750.643.500 Denitrification
Old Tree	G02.842.750.645 Osmosis
Old Tree	G02.842.750.645.250 Electroosmosis
Old Tree	G02.842.750.650 Oxidation-Reduction
Old Tree	G02.842.750.650.500 Oxidative Coupling
Old Tree	G02.842.750.680 Phase Transition
Old Tree	G02.842.750.680.466 Freezing
Old Tree	G02.842.750.680.933 Volatilization
Old Tree	G02.842.750.690 Photochemical Processes
Old Tree	G02.842.750.690.680 Photobleaching
Old Tree	G02.842.750.690.685 Photolysis
Old Tree	G02.842.750.690.842 Photophosphorylation
Old Tree	G02.842.750.690.921 Photosynthesis
Old Tree	G02.842.750.710 Proton-Motive Force
Old Tree	G02.842.750.830 Spontaneous Combustion
Old Tree	G02.842.750.915 X-Ray Diffraction
Old Tree	G02.842.800 Solubility
Old Tree	G02.842.810 Specific Gravity
Old Tree	G02.842.850 Surface Properties
Old Tree	G02.842.850.139 Adhesiveness
Old Tree	G02.842.850.327 Capillary Action
Old Tree	G02.842.850.816 Surface Tension
Old Tree	G02.842.850.908 Wettability
Old Tree	G02.842.925 Viscosity
New Tree	G02.860 Surface Properties
New Tree	G02.860.139 Adhesiveness
New Tree	G02.860.327 Capillary Action
New Tree	G02.860.816 Surface Tension
New Tree	G02.860.908 Wettability
New Tree	G02.930 Viscosity



## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	G02.965	X-Ray Diffraction
Old Tree	G03	Metabolic Phenomena
New Tree	G03	Metabolism
New Tree	G03.015	Absorption
New Tree	G03.015.500	Absorption, Physiological
New Tree	G03.015.500.374	Gastrointestinal Absorption
New Tree	G03.015.500.374.249	Gastric Absorption
New Tree	G03.015.500.374.500	Intestinal Absorption
New Tree	G03.015.500.374.500.500	Intestinal Reabsorption
New Tree	G03.015.500.374.650	Oral Mucosal Absorption
New Tree	G03.015.500.374.825	Rectal Absorption
New Tree	G03.015.500.562	Intramuscular Absorption
New Tree	G03.015.500.656	Ocular Absorption
New Tree	G03.015.500.667	Peritoneal Absorption
New Tree	G03.015.500.679	Renal Reabsorption
New Tree	G03.015.500.703	Respiratory Tract Absorption
New Tree	G03.015.500.703.500	Nasal Absorption
New Tree	G03.015.500.750	Skin Absorption
New Tree	G03.015.500.875	Subcutaneous Absorption
New Tree	G03.015.500.937	Vaginal Absorption
-	G03.030	Acid-Base Equilibrium
New Tree	G03.040	Acylation
New	G03.040.052	Acetylation

## MeSH Tree Changes for 2017

Type	Tree - heading
Tree	
New Tree	G03.040.055 Aminoacylation
New Tree	G03.040.055.860 Transfer RNA Aminoacylation
New Tree	G03.049 Aerobiosis
New Tree	G03.059 Alkylation
New Tree	G03.059.538 Methylation
New Tree	G03.059.538.161 DNA Methylation
New Tree	G03.068 Amination
New Tree	G03.078 Anaerobiosis
New Tree	G03.087 Autotrophic Processes
New Tree	G03.087.314 Chemoautotrophic Growth
New Tree	G03.087.630 Nitrogen Fixation
New Tree	G03.105 Biocatalysis
New Tree	G03.143 Biological Transport
New Tree	G03.143.310 Biological Transport, Active
New Tree	G03.143.310.100 Active Transport, Cell Nucleus
New Tree	G03.143.330 Capillary Permeability
New Tree	G03.143.335 Cell Membrane Permeability
New Tree	G03.143.355 Cytoplasmic Streaming
New Tree	G03.143.355.040 Axonal Transport
New Tree	G03.143.427 Facilitated Diffusion
New Tree	G03.143.500 Ion Transport

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	G03.143.500.100	Calcium Signaling
New Tree	G03.143.700	Protein Transport
New Tree	G03.143.700.100	Active Transport, Cell Nucleus
New Tree	G03.143.775	Respiratory Transport
New Tree	G03.143.775.602	Pulmonary Gas Exchange
New Tree	G03.143.850	RNA Transport
New Tree	G03.143.925	Secretory Pathway
New Tree	G03.143.962	Transcytosis
New Tree	G03.143.962.500	Transcellular Cell Migration
New Tree	G03.162	Biotinylation
New Tree	G03.171	Biotransformation
New Tree	G03.171.224	Activation, Metabolic
New Tree	G03.171.450	Inactivation, Metabolic
New Tree	G03.171.450.500	Metabolic Detoxication, Phase I
New Tree	G03.171.450.750	Metabolic Detoxication, Phase II
-	G03.180	Body Composition
-	G03.180.134	Body Fat Distribution
-	G03.180.134.500	Adiposity
New Tree	G03.185	Brain Chemistry
New Tree	G03.191	Carbohydrate Metabolism
New Tree	G03.191.249	Fermentation
New Tree	G03.191.500	Gluconeogenesis
New Tree	G03.191.625	Glycogenolysis

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	G03.191.750	Glycolysis
New Tree	G03.191.812	Glycosylation
New Tree	G03.191.875	Pentose Phosphate Pathway
New Tree	G03.191.937	Photosynthesis
New Tree	G03.191.937.700	Photophosphorylation
New Tree	G03.194	Catabolite Repression
New Tree	G03.197	Cell Respiration
New Tree	G03.197.300	Cell Hypoxia
New Heading	<b>G03.197.300.500</b>	<b>Tumor Hypoxia</b>
New Tree	G03.197.620	Respiratory Burst
Old Tree	<b>G03.200</b>	<b>Brain Chemistry</b>
New Tree	G03.208	Cyclization
New Tree	G03.219	Dealkylation
New Tree	G03.222	Deamination
New Tree	G03.225	Decarboxylation
New Tree	G03.230	Dimerization
New Tree	G03.295	Energy Metabolism
New Tree	G03.295.154	Basal Metabolism
New Tree	G03.295.342	Citric Acid Cycle
New Tree	G03.295.436	Glycolysis
New Tree	G03.295.531	Oxidation-Reduction
New Tree	G03.295.531.403	Electron Transport

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	G03.295.531.587 Lipid Peroxidation
New Tree	G03.295.631 Oxidative Phosphorylation
New Tree	G03.295.693 Pentose Phosphate Pathway
New Tree	G03.295.724 Photophosphorylation
New Tree	G03.295.770 Proton-Motive Force
New Tree	G03.295.770.500 Membrane Potential, Mitochondrial
New Tree	G03.295.835 Substrate Cycling
New Tree	G03.312 Enterohepatic Circulation
New Tree	G03.312.500 Intestinal Reabsorption
New Tree	G03.328 Enzyme Activation
New Tree	G03.344 Esterification
New Tree	G03.360 Halogenation
New Tree	G03.393 Heterotrophic Processes
New Tree	G03.409 Hydrogenation
New Tree	G03.425 Hydroxylation
New Tree	G03.458 Lipid Metabolism
New Tree	G03.458.249 Lipogenesis
New Tree	G03.458.500 Lipolysis
New Tree	G03.458.500.500 Lipid Mobilization
New Tree	G03.458.875 Lipoylation
-	G03.490 Metabolic Clearance Rate
New Tree	G03.493 Metabolic Networks and Pathways

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	G03.493.100	Biosynthetic Pathways
New Tree	G03.493.170	Citric Acid Cycle
New Tree	G03.493.350	Electron Transport
New Tree	G03.493.360	Glycolysis
New Tree	G03.493.485	Kallikrein-Kinin System
New Tree	G03.493.560	MAP Kinase Signaling System
New Heading	<b>G03.493.628</b>	<b>Non-Neuronal Cholinergic System</b>
New Tree	G03.493.695	Pentose Phosphate Pathway
New Tree	G03.493.700	Photosynthesis
New Tree	G03.493.750	Protein Interaction Maps
New Tree	G03.493.875	Secondary Metabolism
Old Tree	G03.495	Metabolism
Old Tree	G03.495.023	Absorption
Old Tree	G03.495.023.500	Absorption, Physiological
Old Tree	G03.495.023.500.374	Gastrointestinal Absorption
Old Tree	G03.495.023.500.374.249	Gastric Absorption
Old Tree	G03.495.023.500.374.500	Intestinal Absorption
Old Tree	G03.495.023.500.374.500.500	Intestinal Reabsorption
Old Tree	G03.495.023.500.374.650	Oral Mucosal Absorption
Old Tree	G03.495.023.500.374.825	Rectal Absorption
Old Tree	G03.495.023.500.562	Intramuscular Absorption
Old Tree	G03.495.023.500.656	Ocular Absorption
Old Tree	G03.495.023.500.667	Peritoneal Absorption
Old Tree	G03.495.023.500.679	Renal Reabsorption
Old Tree	G03.495.023.500.703	Respiratory Tract Absorption
Old Tree	G03.495.023.500.703.500	Nasal Absorption
Old Tree	G03.495.023.500.750	Skin Absorption
Old Tree	G03.495.023.500.875	Subcutaneous Absorption
Old Tree	G03.495.023.500.937	Vaginal Absorption

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	G03.495.095 Acylation
Old Tree	G03.495.095.052 Acetylation
Old Tree	G03.495.095.055 Aminoacylation
Old Tree	G03.495.095.055.860 Transfer RNA Aminoacylation
Old Tree	G03.495.112 Aerobiosis
Old Tree	G03.495.130 Alkylation
Old Tree	G03.495.130.538 Methylation
Old Tree	G03.495.130.538.161 DNA Methylation
Old Tree	G03.495.140 Amination
Old Tree	G03.495.146 Anaerobiosis
Old Tree	G03.495.153 Autotrophic Processes
Old Tree	G03.495.153.314 Chemoautotrophic Growth
Old Tree	G03.495.153.630 Nitrogen Fixation
Old Tree	G03.495.159 Biocatalysis
Old Tree	G03.495.166 Biological Transport
Old Tree	G03.495.166.310 Biological Transport, Active
Old Tree	G03.495.166.310.100 Active Transport, Cell Nucleus
Old Tree	G03.495.166.330 Capillary Permeability
Old Tree	G03.495.166.335 Cell Membrane Permeability
Old Tree	G03.495.166.355 Cytoplasmic Streaming
Old Tree	G03.495.166.355.040 Axonal Transport
Old Tree	G03.495.166.427 Facilitated Diffusion
Old Tree	G03.495.166.500 Ion Transport
Old Tree	G03.495.166.500.100 Calcium Signaling
Old Tree	G03.495.166.700 Protein Transport
Old Tree	G03.495.166.700.100 Active Transport, Cell Nucleus
Old Tree	G03.495.166.775 Respiratory Transport
Old Tree	G03.495.166.775.602 Pulmonary Gas Exchange
Old Tree	G03.495.166.850 RNA Transport
Old Tree	G03.495.166.925 Secretory Pathway
Old Tree	G03.495.166.962 Transcytosis
Old Tree	G03.495.166.962.500 Transcellular Cell Migration
Old Tree	G03.495.233 Biotinylation
Old Tree	G03.495.244 Biotransformation
Old Tree	G03.495.244.224 Activation, Metabolic
Old Tree	G03.495.244.450 Inactivation, Metabolic

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	G03.495.244.450.500                      Metabolic Detoxication, Phase I
Old Tree	G03.495.244.450.750                      Metabolic Detoxication, Phase II
Old Tree	G03.495.256                                    Carbohydrate Metabolism
Old Tree	G03.495.256.249                            Fermentation
Old Tree	G03.495.256.500                            Gluconeogenesis
Old Tree	G03.495.256.625                            Glycogenolysis
Old Tree	G03.495.256.750                            Glycolysis
Old Tree	G03.495.256.812                            Glycosylation
Old Tree	G03.495.256.875                            Pentose Phosphate Pathway
Old Tree	G03.495.256.937                            Photosynthesis
Old Tree	G03.495.256.937.700                      Photophosphorylation
Old Tree	G03.495.278                                   Cell Respiration
Old Tree	G03.495.278.300                            Cell Hypoxia
Old Tree	G03.495.278.620                            Respiratory Burst
Old Tree	G03.495.300                                   Cyclization
Old Tree	G03.495.305                                   Dealkylation
Old Tree	G03.495.310                                   Deamination
Old Tree	G03.495.315                                   Decarboxylation
Old Tree	G03.495.320                                   Dimerization
Old Tree	G03.495.335                                   Energy Metabolism
Old Tree	G03.495.335.154                            Basal Metabolism
Old Tree	G03.495.335.342                            Citric Acid Cycle
Old Tree	G03.495.335.436                            Glycolysis
Old Tree	G03.495.335.531                            Oxidation-Reduction
Old Tree	G03.495.335.531.403                      Electron Transport
Old Tree	G03.495.335.531.587                      Lipid Peroxidation
Old Tree	G03.495.335.631                            Oxidative Phosphorylation
Old Tree	G03.495.335.693                            Pentose Phosphate Pathway
Old Tree	G03.495.335.724                            Photophosphorylation
Old Tree	G03.495.335.770                            Proton-Motive Force
Old Tree	G03.495.335.770.500                      Membrane Potential, Mitochondrial
Old Tree	G03.495.335.835                            Substrate Cycling
Old Tree	G03.495.355                                   Enterohepatic Circulation
Old Tree	G03.495.355.500                            Intestinal Reabsorption
Old Tree	G03.495.368                                   Enzyme Activation
Old Tree	G03.495.475                                   Esterification



## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	G03.495.496 Halogenation
Old Tree	G03.495.505 Heterotrophic Processes
Old Tree	G03.495.506 Hydrogenation
Old Tree	G03.495.510 Hydroxylation
Old Tree	G03.495.520 Lipid Metabolism
Old Tree	G03.495.520.249 Lipogenesis
Old Tree	G03.495.520.500 Lipolysis
Old Tree	G03.495.520.500.500 Lipid Mobilization
Old Tree	G03.495.520.875 Lipoylation
Old Tree	G03.495.553 Metabolic Networks and Pathways
Old Tree	G03.495.553.100 Biosynthetic Pathways
Old Tree	G03.495.553.170 Citric Acid Cycle
Old Tree	G03.495.553.350 Electron Transport
Old Tree	G03.495.553.360 Glycolysis
Old Tree	G03.495.553.485 Kallikrein-Kinin System
Old Tree	G03.495.553.560 MAP Kinase Signaling System
Old Tree	G03.495.553.695 Pentose Phosphate Pathway
Old Tree	G03.495.553.700 Photosynthesis
Old Tree	G03.495.553.750 Protein Interaction Maps
Old Tree	G03.495.553.875 Secondary Metabolism
Old Tree	G03.495.655 Nitrosation
Old Tree	G03.495.710 Oxidative Stress
Old Tree	G03.495.710.690 Protein Carbonylation
Old Tree	G03.495.770 Peptide Biosynthesis
Old Tree	G03.495.770.050 Aminoacylation
Old Tree	G03.495.770.050.860 Transfer RNA Aminoacylation
Old Tree	G03.495.770.333 Peptide Biosynthesis, Nucleic Acid-Independent
Old Tree	G03.495.770.871 Protein Biosynthesis
Old Tree	G03.495.770.871.200 Frameshifting, Ribosomal
Old Tree	G03.495.770.871.640 Peptide Chain Elongation, Translational
Old Tree	G03.495.770.871.650 Peptide Chain Initiation, Translational
Old Tree	G03.495.770.871.720 Peptide Chain Termination, Translational
Old Tree	G03.495.770.871.790 Protein Modification, Translational
Old Tree	G03.495.770.871.790.600 Protein Processing, Post-Translational
Old Tree	G03.495.770.871.790.600.400 Protein Prenylation
Old Tree	G03.495.770.871.790.600.700 Protein Splicing

## MeSH Tree Changes for 2017

Type	Tree - heading	
Old Tree	G03.495.770.871.790.600.831	Ubiquitination
Old Tree	G03.495.770.871.790.600.831.500	Sumoylation
Old Tree	G03.495.770.871.790.600.850	Unfolded Protein Response
Old Tree	G03.495.770.871.790.600.850.500 Degradation	Endoplasmic Reticulum-Associated
Old Tree	G03.495.770.871.850	Transfer RNA Aminoacylation
Old Tree	G03.495.790	Phosphorylation
Old Tree	G03.495.790.550	Oxidative Phosphorylation
Old Tree	G03.495.790.605	Photophosphorylation
Old Tree	G03.495.795	Phototrophic Processes
Old Tree	G03.495.795.700	Photosynthesis
Old Tree	G03.495.795.700.700	Photophosphorylation
Old Tree	G03.495.797	Prenylation
Old Tree	G03.495.797.500	Protein Prenylation
Old Tree	G03.495.800	Protein Binding
Old Tree	G03.495.802	Proteolysis
Old Tree	G03.495.805	Renin-Angiotensin System
Old Tree	G03.495.839	RNA Processing, Post-Transcriptional
Old Tree	G03.495.839.112	Nonsense Mediated mRNA Decay
Old Tree	G03.495.839.225	RNA 3' End Processing
Old Tree	G03.495.839.225.710	Polyadenylation
Old Tree	G03.495.839.250	RNA Editing
Old Tree	G03.495.839.700	RNA Splicing
Old Tree	G03.495.839.700.100	Alternative Splicing
Old Tree	G03.495.839.700.750	Trans-Splicing
Old Tree	G03.495.873	Secretory Rate
-	G03.500	Metabolome
New Tree	G03.558	Nitrosation
-	G03.615	Osmoregulation
-	G03.615.500	Water-Electrolyte Balance
-	G03.615.500.500	Kallikrein-Kinin System
-	G03.615.500.750	Water Loss, Insensible
New Tree	G03.673	Oxidative Stress
New Tree	G03.673.690	Protein Carbonylation

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	G03.680	Oxygen Consumption
New Tree	G03.680.110	Anaerobic Threshold
New Tree	G03.680.365	Metabolic Equivalent
Old Tree	G03.730	Oxygen Consumption
Old Tree	G03.730.110	Anaerobic Threshold
Old Tree	G03.730.365	Metabolic Equivalent
New Tree	G03.734	Peptide Biosynthesis
New Tree	G03.734.050	Aminoacylation
New Tree	G03.734.050.860	Transfer RNA Aminoacylation
New Tree	G03.734.333	Peptide Biosynthesis, Nucleic Acid-Independent
New Tree	G03.734.871	Protein Biosynthesis
New Tree	G03.734.871.200	Frameshifting, Ribosomal
New Tree	G03.734.871.640	Peptide Chain Elongation, Translational
New Tree	G03.734.871.650	Peptide Chain Initiation, Translational
New Tree	G03.734.871.720	Peptide Chain Termination, Translational
New Tree	G03.734.871.790	Protein Modification, Translational
New Tree	G03.734.871.790.600	Protein Processing, Post-Translational
New Tree	G03.734.871.790.600.400	Protein Prenylation
New Tree	G03.734.871.790.600.700	Protein Splicing
New Tree	G03.734.871.790.600.831	Ubiquitination
New Tree	G03.734.871.790.600.831.500	Sumoylation
New Tree	G03.734.871.790.600.850	Unfolded Protein Response
New Tree	G03.734.871.790.600.850.500	Endoplasmic Reticulum-Associated Degradation

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	<a href="#">G03.734.871.850</a> <a href="#">Transfer RNA Aminoacylation</a>
-	G03.787 Pharmacokinetics
-	G03.787.024 Absorption
-	G03.787.024.500 Absorption, Physiological
-	G03.787.024.500.374 Gastrointestinal Absorption
-	G03.787.024.500.374.249 Gastric Absorption
-	G03.787.024.500.374.500 Intestinal Absorption
-	G03.787.024.500.374.500.500 Intestinal Reabsorption
-	G03.787.024.500.374.650 Oral Mucosal Absorption
-	G03.787.024.500.374.825 Rectal Absorption
-	G03.787.024.500.562 Intramuscular Absorption
-	G03.787.024.500.656 Ocular Absorption
-	G03.787.024.500.667 Peritoneal Absorption
-	G03.787.024.500.679 Renal Reabsorption
-	G03.787.024.500.703 Respiratory Tract Absorption
-	G03.787.024.500.703.500 Nasal Absorption
-	G03.787.024.500.750 Skin Absorption
-	G03.787.024.500.875 Subcutaneous Absorption
-	G03.787.024.500.937 Vaginal Absorption
-	G03.787.101 Area Under Curve
-	G03.787.151 Biological Availability
-	G03.787.225 Biotransformation
-	G03.787.225.224 Activation, Metabolic
-	G03.787.225.450 Inactivation, Metabolic
-	G03.787.225.450.500 Metabolic Detoxication, Phase I
-	G03.787.225.450.750 Metabolic Detoxication, Phase II
-	G03.787.273 Cutaneous Elimination
-	G03.787.321 Drug Liberation
-	G03.787.380 Hepatobiliary Elimination
-	G03.787.383 Intestinal Elimination
-	G03.787.387 Lacrimal Elimination
-	G03.787.391 Lacteal Elimination
-	G03.787.395 Pulmonary Elimination
-	G03.787.410 Renal Elimination
-	G03.787.440 Salivary Elimination

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	G03.787.559	Therapeutic Equivalency
-	G03.787.917	Tissue Distribution
New Tree	<a href="#">G03.796</a>	<a href="#">Phosphorylation</a>
New Tree	<a href="#">G03.796.550</a>	<a href="#">Oxidative Phosphorylation</a>
New Tree	<a href="#">G03.796.605</a>	<a href="#">Photophosphorylation</a>
New Tree	<a href="#">G03.800</a>	<a href="#">Phototrophic Processes</a>
New Tree	<a href="#">G03.800.700</a>	<a href="#">Photosynthesis</a>
New Tree	<a href="#">G03.800.700.700</a>	<a href="#">Photophosphorylation</a>
New Tree	<a href="#">G03.804</a>	<a href="#">Prenylation</a>
New Tree	<a href="#">G03.804.500</a>	<a href="#">Protein Prenylation</a>
New Tree	<a href="#">G03.808</a>	<a href="#">Protein Binding</a>
New Tree	<a href="#">G03.812</a>	<a href="#">Proteolysis</a>
New Tree	<a href="#">G03.820</a>	<a href="#">Renin-Angiotensin System</a>
New Tree	<a href="#">G03.839</a>	<a href="#">RNA Processing, Post-Transcriptional</a>
New Tree	<a href="#">G03.839.112</a>	<a href="#">Nonsense Mediated mRNA Decay</a>
New Tree	<a href="#">G03.839.225</a>	<a href="#">RNA 3' End Processing</a>
New Tree	<a href="#">G03.839.225.710</a>	<a href="#">Polyadenylation</a>
New Tree	<a href="#">G03.839.250</a>	<a href="#">RNA Editing</a>
New Tree	<a href="#">G03.839.700</a>	<a href="#">RNA Splicing</a>
New Tree	<a href="#">G03.839.700.100</a>	<a href="#">Alternative Splicing</a>
New Tree	<a href="#">G03.839.700.750</a>	<a href="#">Trans-Splicing</a>
New Tree	<a href="#">G03.857</a>	<a href="#">Secretory Rate</a>
-	G03.893	Toxicokinetics

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	G04	Cell Physiological Phenomena
New Tree	G04.022	Cell Adhesion
New Tree	G04.043	Cell Aging
New Tree	G04.043.260	Erythrocyte Aging
New Tree	G04.043.630	Telomere Shortening
New Tree	G04.085	Cell Communication
New Tree	G04.085.100	Autocrine Communication
New Tree	G04.085.155	Bystander Effect
New Tree	G04.085.300	Embryonic Induction
New Tree	G04.085.600	Paracrine Communication
New Tree	G04.085.700	Quorum Sensing
New Tree	G04.128	Cell Compartmentation
New Tree	G04.128.180	Chromosome Positioning
New Tree	G04.140	Cell Count
New Tree	G04.140.107	Blood Cell Count
New Tree	G04.140.107.330	Erythrocyte Count
New Tree	G04.140.107.330.725	Reticulocyte Count
New Tree	G04.140.107.595	Leukocyte Count
New Tree	G04.140.107.595.500	Lymphocyte Count
New Tree	G04.140.107.595.500.150	CD4 Lymphocyte Count
New Tree	G04.140.107.595.500.150.160	CD4-CD8 Ratio
New Tree	G04.140.107.740	Platelet Count

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	G04.140.870 Sperm Count
New Tree	G04.144 Cell Cycle
New Tree	G04.144.109 Cell Cycle Checkpoints
New Tree	G04.144.109.249 G1 Phase Cell Cycle Checkpoints
New Tree	G04.144.109.500 G2 Phase Cell Cycle Checkpoints
New Tree	G04.144.109.750 M Phase Cell Cycle Checkpoints
New Tree	G04.144.109.875 S Phase Cell Cycle Checkpoints
New Tree	G04.144.220 Cell Division
New Tree	G04.144.220.109 Asymmetric Cell Division
New Tree	G04.144.220.220 Cell Nucleus Division
New Tree	G04.144.220.220.625 Chromosome Segregation
New Tree	G04.144.220.220.687 Meiosis
New Tree	G04.144.220.220.687.222 Anaphase
New Tree	G04.144.220.220.687.444 Meiotic Prophase I
New Tree	G04.144.220.220.687.625 Metaphase
New Tree	G04.144.220.220.687.812 Prometaphase
New Tree	G04.144.220.220.687.883 Prophase
New Tree	G04.144.220.220.687.883.250 Chromosome Pairing
New Tree	G04.144.220.220.687.883.250.500 Synaptonemal Complex
New Tree	G04.144.220.220.687.883.750 Pachytene Stage
New Tree	G04.144.220.220.687.953 Telophase
New Tree	G04.144.220.220.781 Mitosis

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	G04.144.220.220.781.050 Anaphase
New Tree	G04.144.220.220.781.337 M Phase Cell Cycle Checkpoints
New Tree	G04.144.220.220.781.625 Metaphase
New Tree	G04.144.220.220.781.812 Prometaphase
New Tree	G04.144.220.220.781.906 Prophase
New Tree	G04.144.220.220.781.906.250 Chromosome Pairing
New Tree	G04.144.220.220.781.906.250.500 Synaptonemal Complex
New Tree	G04.144.220.220.781.906.750 Pachytene Stage
New Tree	G04.144.220.220.781.953 Telophase
New Tree	G04.144.220.235 Cell Self Renewal
New Tree	G04.144.220.250 Cytokinesis
New Tree	G04.144.220.625 Telomere Homeostasis
New Tree	G04.144.500 Interphase
New Tree	G04.144.500.300 G0 Phase
New Tree	G04.144.500.300 Resting Phase, Cell Cycle
New Tree	G04.144.500.320 G1 Phase
New Tree	G04.144.500.320.500 G1 Phase Cell Cycle Checkpoints
New Tree	G04.144.500.340 G2 Phase
New Tree	G04.144.500.340.500 G2 Phase Cell Cycle Checkpoints
New Tree	G04.144.500.800 S Phase
New Tree	G04.144.500.800.500 S Phase Cell Cycle Checkpoints
New Tree	G04.146 Cell Death



## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	G04.146.160 Apoptosis
New Tree	G04.146.160.060 Anoikis
New Heading	<b>G04.146.160.295 Eryptosis</b>
New Tree	G04.146.160.530 Pyroptosis
New Tree	G04.146.399 Autophagy
New Tree	G04.146.638 Necrosis
New Tree	G04.146.638.200 DNA Degradation, Necrotic
New Tree	G04.148 Cell Dedifferentiation
New Tree	G04.152 Cell Differentiation
New Tree	G04.152.149 Adipogenesis
New Tree	G04.152.224 Asymmetric Cell Division
New Tree	G04.152.262 Cellular Reprogramming
New Tree	G04.152.300 Embryonic Induction
New Tree	G04.152.650 Gametogenesis
New Tree	G04.152.650.240 Gametogenesis, Plant
New Tree	G04.152.650.249 Oogenesis
New Tree	G04.152.650.249.880 Vitellogenesis
New Tree	G04.152.650.624 Spermatogenesis
New Tree	G04.152.650.624.700 Sperm Maturation
New Tree	G04.152.825 Hematopoiesis
New Tree	G04.152.825.414 Erythropoiesis
New Tree	G04.152.825.463 Hematopoiesis, Extramedullary

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	G04.152.825.597	Leukopoiesis
New Tree	G04.152.825.597.500	Lymphopoiesis
New Tree	G04.152.825.597.750	Myelopoiesis
New Tree	G04.152.825.798	Thrombopoiesis
New Tree	G04.152.912	Neurogenesis
New Heading	<b>G04.152.912.750</b>	<b>Neuronal Outgrowth</b>
New Heading	<b>G04.152.912.750.250</b>	<b>Axon Fasciculation</b>
New Heading	<b>G04.152.912.750.500</b>	<b>Axon Guidance</b>
New Tree	G04.155	Cell Fusion
New Tree	G04.161	Cell Growth Processes
New Tree	G04.161.500	Cell Enlargement
New Tree	G04.161.750	Cell Proliferation
New Tree	G04.161.750.500	Cell Division
New Tree	G04.161.750.500.249	Asymmetric Cell Division
New Tree	G04.161.750.500.375	Cell Self Renewal
New Tree	G04.161.750.500.500	Telomere Homeostasis
New Tree	G04.165	Cell-in-Cell Formation
New Tree	G04.165.249	Cytophagocytosis
New Tree	G04.165.500	Emperipoiesis
New Tree	G04.165.500.500	Transcellular Cell Migration
New Tree	G04.165.750	Entosis
Old Tree	<b>G04.170</b>	<b>Cell Count</b>

## MeSH Tree Changes for 2017

Type	Tree - heading	
Old Tree	<b>G04.170.107</b>	<b>Blood Cell Count</b>
Old Tree	<b>G04.170.107.330</b>	<b>Erythrocyte Count</b>
Old Tree	<b>G04.170.107.330.725</b>	<b>Reticulocyte Count</b>
Old Tree	<b>G04.170.107.595</b>	<b>Leukocyte Count</b>
Old Tree	<b>G04.170.107.595.500</b>	<b>Lymphocyte Count</b>
Old Tree	<b>G04.170.107.595.500.150</b>	<b>CD4 Lymphocyte Count</b>
Old Tree	<b>G04.170.107.595.500.150.160</b>	<b>CD4-CD8 Ratio</b>
Old Tree	<b>G04.170.107.740</b>	<b>Platelet Count</b>
Old Tree	<b>G04.170.870</b>	<b>Sperm Count</b>
-	G04.172	Cell Lineage
-	G04.175	Cell Membrane Permeability
New Tree	<b>G04.198</b>	<b>Cell Movement</b>
New Tree	<b>G04.198.251</b>	<b>Cell Aggregation</b>
New Tree	<b>G04.198.337</b>	<b>Cell Migration Inhibition</b>
New Tree	<b>G04.198.424</b>	<b>Chemotaxis</b>
New Tree	<b>G04.198.424.233</b>	<b>Chemotaxis, Leukocyte</b>
New Tree	<b>G04.198.424.233.500</b>	<b>Leukocyte Rolling</b>
New Tree	<b>G04.198.700</b>	<b>Ovum Transport</b>
New Tree	<b>G04.198.750</b>	<b>Sperm Motility</b>
New Tree	<b>G04.198.800</b>	<b>Sperm Transport</b>
New Tree	<b>G04.198.900</b>	<b>Transcellular Cell Migration</b>
New Tree	<b>G04.198.950</b>	<b>Transendothelial and Transepithelial Migration</b>
New Tree	<b>G04.250</b>	<b>Cell Polarity</b>
New Tree	<b>G04.270</b>	<b>Cell Respiration</b>
New Tree	<b>G04.270.300</b>	<b>Cell Hypoxia</b>
New Heading	<b>G04.270.300.500</b>	<b>Tumor Hypoxia</b>

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	<a href="#">G04.270.620</a> <a href="#">Respiratory Burst</a>
Old Tree	<a href="#">G04.299</a> <a href="#">Cell Physiological Processes</a>
Old Tree	<a href="#">G04.299.117</a> <a href="#">Cell Adhesion</a>
Old Tree	<a href="#">G04.299.119</a> <a href="#">Cell Aging</a>
Old Tree	<a href="#">G04.299.119.260</a> <a href="#">Erythrocyte Aging</a>
Old Tree	<a href="#">G04.299.119.630</a> <a href="#">Telomere Shortening</a>
Old Tree	<a href="#">G04.299.122</a> <a href="#">Cell Communication</a>
Old Tree	<a href="#">G04.299.122.100</a> <a href="#">Autocrine Communication</a>
Old Tree	<a href="#">G04.299.122.155</a> <a href="#">Bystander Effect</a>
Old Tree	<a href="#">G04.299.122.300</a> <a href="#">Embryonic Induction</a>
Old Tree	<a href="#">G04.299.122.600</a> <a href="#">Paracrine Communication</a>
Old Tree	<a href="#">G04.299.122.700</a> <a href="#">Quorum Sensing</a>
Old Tree	<a href="#">G04.299.125</a> <a href="#">Cell Compartmentation</a>
Old Tree	<a href="#">G04.299.125.180</a> <a href="#">Chromosome Positioning</a>
Old Tree	<a href="#">G04.299.134</a> <a href="#">Cell Cycle</a>
Old Tree	<a href="#">G04.299.134.109</a> <a href="#">Cell Cycle Checkpoints</a>
Old Tree	<a href="#">G04.299.134.109.249</a> <a href="#">G1 Phase Cell Cycle Checkpoints</a>
Old Tree	<a href="#">G04.299.134.109.500</a> <a href="#">G2 Phase Cell Cycle Checkpoints</a>
Old Tree	<a href="#">G04.299.134.109.750</a> <a href="#">M Phase Cell Cycle Checkpoints</a>
Old Tree	<a href="#">G04.299.134.109.875</a> <a href="#">S Phase Cell Cycle Checkpoints</a>
Old Tree	<a href="#">G04.299.134.220</a> <a href="#">Cell Division</a>
Old Tree	<a href="#">G04.299.134.220.109</a> <a href="#">Asymmetric Cell Division</a>
Old Tree	<a href="#">G04.299.134.220.220</a> <a href="#">Cell Nucleus Division</a>
Old Tree	<a href="#">G04.299.134.220.220.625</a> <a href="#">Chromosome Segregation</a>
Old Tree	<a href="#">G04.299.134.220.220.687</a> <a href="#">Meiosis</a>
Old Tree	<a href="#">G04.299.134.220.220.687.222</a> <a href="#">Anaphase</a>
Old Tree	<a href="#">G04.299.134.220.220.687.444</a> <a href="#">Meiotic Prophase I</a>
Old Tree	<a href="#">G04.299.134.220.220.687.625</a> <a href="#">Metaphase</a>
Old Tree	<a href="#">G04.299.134.220.220.687.812</a> <a href="#">Prometaphase</a>
Old Tree	<a href="#">G04.299.134.220.220.687.883</a> <a href="#">Prophase</a>
Old Tree	<a href="#">G04.299.134.220.220.687.883.250</a> <a href="#">Chromosome Pairing</a>
Old Tree	<a href="#">G04.299.134.220.220.687.883.250.500</a> <a href="#">Synaptonemal Complex</a>
Old Tree	<a href="#">G04.299.134.220.220.687.883.750</a> <a href="#">Pachytene Stage</a>
Old Tree	<a href="#">G04.299.134.220.220.687.953</a> <a href="#">Telophase</a>
Old Tree	<a href="#">G04.299.134.220.220.781</a> <a href="#">Mitosis</a>

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	G04.299.134.220.220.781.050 Anaphase
Old Tree	G04.299.134.220.220.781.337 M Phase Cell Cycle Checkpoints
Old Tree	G04.299.134.220.220.781.625 Metaphase
Old Tree	G04.299.134.220.220.781.812 Prometaphase
Old Tree	G04.299.134.220.220.781.906 Prophase
Old Tree	G04.299.134.220.220.781.906.250 Chromosome Pairing
Old Tree	G04.299.134.220.220.781.906.250.500 Synaptonemal Complex
Old Tree	G04.299.134.220.220.781.906.750 Pachytene Stage
Old Tree	G04.299.134.220.220.781.953 Telophase
Old Tree	G04.299.134.220.235 Cell Self Renewal
Old Tree	G04.299.134.220.250 Cytokinesis
Old Tree	G04.299.134.220.625 Telomere Homeostasis
Old Tree	G04.299.134.500 Interphase
Old Tree	G04.299.134.500.300 G0 Phase
Old Tree	G04.299.134.500.300 Resting Phase, Cell Cycle
Old Tree	G04.299.134.500.320 G1 Phase
Old Tree	G04.299.134.500.320.500 G1 Phase Cell Cycle Checkpoints
Old Tree	G04.299.134.500.340 G2 Phase
Old Tree	G04.299.134.500.340.500 G2 Phase Cell Cycle Checkpoints
Old Tree	G04.299.134.500.800 S Phase
Old Tree	G04.299.134.500.800.500 S Phase Cell Cycle Checkpoints
Old Tree	G04.299.139 Cell Death
Old Tree	G04.299.139.160 Apoptosis
Old Tree	G04.299.139.160.060 Anoikis
Old Tree	G04.299.139.160.530 Pyroptosis
Old Tree	G04.299.139.399 Autophagy
Old Tree	G04.299.139.638 Necrosis
Old Tree	G04.299.139.638.200 DNA Degradation, Necrotic
Old Tree	G04.299.145 Cell Dedifferentiation
Old Tree	G04.299.151 Cell Differentiation
Old Tree	G04.299.151.149 Adipogenesis
Old Tree	G04.299.151.224 Asymmetric Cell Division
Old Tree	G04.299.151.262 Cellular Reprogramming
Old Tree	G04.299.151.300 Embryonic Induction
Old Tree	G04.299.151.650 Gametogenesis
Old Tree	G04.299.151.650.240 Gametogenesis, Plant

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	G04.299.151.650.249 Oogenesis
Old Tree	G04.299.151.650.249.880 Vitellogenesis
Old Tree	G04.299.151.650.624 Spermatogenesis
Old Tree	G04.299.151.650.624.700 Sperm Maturation
Old Tree	G04.299.151.825 Hematopoiesis
Old Tree	G04.299.151.825.414 Erythropoiesis
Old Tree	G04.299.151.825.463 Hematopoiesis, Extramedullary
Old Tree	G04.299.151.825.597 Leukopoiesis
Old Tree	G04.299.151.825.597.500 Lymphopoiesis
Old Tree	G04.299.151.825.597.750 Myelopoiesis
Old Tree	G04.299.151.825.798 Thrombopoiesis
Old Tree	G04.299.151.912 Neurogenesis
Old Tree	G04.299.217 Cell Fusion
Old Tree	G04.299.233 Cell Growth Processes
Old Tree	G04.299.233.500 Cell Enlargement
Old Tree	G04.299.233.750 Cell Proliferation
Old Tree	G04.299.233.750.500 Cell Division
Old Tree	G04.299.233.750.500.249 Asymmetric Cell Division
Old Tree	G04.299.233.750.500.375 Cell Self Renewal
Old Tree	G04.299.233.750.500.500 Telomere Homeostasis
Old Tree	G04.299.283 Cell Movement
Old Tree	G04.299.283.251 Cell Aggregation
Old Tree	G04.299.283.337 Cell Migration Inhibition
Old Tree	G04.299.283.424 Chemotaxis
Old Tree	G04.299.283.424.233 Chemotaxis, Leukocyte
Old Tree	G04.299.283.424.233.500 Leukocyte Rolling
Old Tree	G04.299.283.700 Ovum Transport
Old Tree	G04.299.283.750 Sperm Motility
Old Tree	G04.299.283.800 Sperm Transport
Old Tree	G04.299.283.900 Transcellular Cell Migration
Old Tree	G04.299.283.950 Transendothelial and Transepithelial Migration
Old Tree	G04.299.305 Cell Respiration
Old Tree	G04.299.305.300 Cell Hypoxia
Old Tree	G04.299.305.620 Respiratory Burst
Old Tree	G04.299.316 Cell Survival
Old Tree	G04.299.335 Cell Transdifferentiation

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	G04.299.335.250 Cell Plasticity
Old Tree	G04.299.335.500 Epithelial-Mesenchymal Transition
Old Tree	G04.299.345 Cell-in-Cell Formation
Old Tree	G04.299.345.249 Cytophagocytosis
Old Tree	G04.299.345.500 Emperipolesis
Old Tree	G04.299.345.500.500 Transcellular Cell Migration
Old Tree	G04.299.345.750 Entosis
Old Tree	G04.299.350 Clonal Evolution
Old Tree	G04.299.355 Contact Inhibition
Old Tree	G04.299.418 Cytoplasmic Streaming
Old Tree	G04.299.418.040 Axonal Transport
Old Tree	G04.299.452 DNA Packaging
Old Tree	G04.299.452.095 Chromatin Assembly and Disassembly
Old Tree	G04.299.482 Endocytosis
Old Tree	G04.299.482.350 Phagocytosis
Old Tree	G04.299.482.350.091 Autophagy
Old Tree	G04.299.482.350.545 Cytophagocytosis
Old Tree	G04.299.482.370 Pinocytosis
Old Tree	G04.299.482.685 Transcytosis
Old Tree	G04.299.482.685.500 Transcellular Cell Migration
Old Tree	G04.299.490 Exocytosis
Old Tree	G04.299.490.160 Cell Degranulation
Old Tree	G04.299.490.580 Secretory Pathway
Old Tree	G04.299.490.790 Transcytosis
Old Tree	G04.299.490.790.500 Transcellular Cell Migration
Old Tree	G04.299.540 Lymphocyte Cooperation
Old Tree	G04.299.570 Membrane Fusion
Old Tree	G04.299.580 Mitochondrial Swelling
Old Tree	G04.299.630 Mitochondrial Turnover
Old Tree	G04.299.630.500 Mitochondrial Degradation
Old Tree	G04.299.630.750 Mitochondrial Dynamics
Old Tree	G04.299.680 Radiation Tolerance
Old Tree	G04.299.680.500 Dose-Response Relationship, Radiation
Old Tree	G04.299.730 Re-Epithelialization
Old Tree	G04.299.780 Receptor Aggregation
Old Tree	G04.299.780.612 Immunologic Capping

## MeSH Tree Changes for 2017

Type	Tree - heading	
Old Tree	G04.299.785	Receptor Cross-Talk
Old Tree	G04.299.880	Signal Transduction
Old Tree	G04.299.880.199	Excitation Contraction Coupling
Old Tree	G04.299.880.400	Ion Channel Gating
Old Tree	G04.299.880.480	Light Signal Transduction
Old Tree	G04.299.880.480.900	Vision, Ocular
Old Tree	G04.299.880.560	MAP Kinase Signaling System
Old Tree	G04.299.880.580	Mechanotransduction, Cellular
Old Tree	G04.299.880.800	Second Messenger Systems
Old Tree	G04.299.880.800.100	Calcium Signaling
Old Tree	G04.299.880.850	Synaptic Transmission
Old Tree	G04.299.880.850.500	Postsynaptic Potential Summation
Old Tree	G04.310	Cell Polarity
-	G04.320	Cell Shape
-	G04.325	Cell Size
-	G04.325.500	Mean Platelet Volume
New Tree	G04.346	Cell Survival
New Tree	G04.356	Cell Transdifferentiation
New Tree	G04.356.250	Cell Plasticity
New Tree	G04.356.500	Epithelial-Mesenchymal Transition
-	G04.366	Cellular Microenvironment
-	G04.366.249	Stem Cell Niche
-	G04.366.500	Tumor Microenvironment
New Tree	G04.375	Clonal Evolution
New Tree	G04.383	Contact Inhibition
New Tree	G04.392	Cytoplasmic Streaming
New Tree	G04.392.040	Axonal Transport
New Tree	G04.400	DNA Packaging
New Tree	G04.400.095	Chromatin Assembly and Disassembly



## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	G04.417	Endocytosis
New Tree	G04.417.350	Phagocytosis
New Tree	G04.417.350.091	Autophagy
New Tree	G04.417.350.545	Cytophagocytosis
New Tree	G04.417.370	Pinocytosis
New Tree	G04.417.685	Transcytosis
New Tree	G04.417.685.500	Transcellular Cell Migration
-	G04.434	Endoplasmic Reticulum Stress
New Tree	G04.468	Exocytosis
New Tree	G04.468.160	Cell Degranulation
New Tree	G04.468.580	Secretory Pathway
New Tree	G04.468.790	Transcytosis
New Tree	G04.468.790.500	Transcellular Cell Migration
New Tree	G04.502	Lymphocyte Cooperation
-	G04.570	Membrane Fluidity
New Tree	G04.575	Membrane Fusion
-	G04.580	Membrane Potentials
-	G04.580.100	Action Potentials
-	G04.580.550	Membrane Potential, Mitochondrial
-	G04.580.887	Synaptic Potentials
-	G04.580.887.249	Excitatory Postsynaptic Potentials
-	G04.580.887.374	Inhibitory Postsynaptic Potentials
-	G04.580.887.500	Miniature Postsynaptic Potentials
-	G04.580.887.750	Postsynaptic Potential Summation
New Tree	G04.590	Mitochondrial Swelling
New Tree	G04.599	Mitochondrial Turnover

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	<a href="#">G04.599.500</a>	<a href="#">Mitochondrial Degradation</a>
New Tree	<a href="#">G04.599.750</a>	<a href="#">Mitochondrial Dynamics</a>
-	G04.618	Organelle Biogenesis
-	G04.655	Organelle Shape
-	G04.655.270	Cell Nucleus Shape
-	G04.670	Organelle Size
-	G04.670.160	Cell Nucleus Size
-	G04.670.560	Mitochondrial Size
New Tree	<a href="#">G04.712</a>	<a href="#">Radiation Tolerance</a>
New Tree	<a href="#">G04.712.500</a>	<a href="#">Dose-Response Relationship, Radiation</a>
New Tree	<a href="#">G04.753</a>	<a href="#">Re-Epithelialization</a>
New Tree	<a href="#">G04.774</a>	<a href="#">Receptor Aggregation</a>
New Tree	<a href="#">G04.774.612</a>	<a href="#">Immunologic Capping</a>
New Tree	<a href="#">G04.794</a>	<a href="#">Receptor Cross-Talk</a>
New Tree	<a href="#">G04.835</a>	<a href="#">Signal Transduction</a>
New Tree	<a href="#">G04.835.199</a>	<a href="#">Excitation Contraction Coupling</a>
New Tree	<a href="#">G04.835.400</a>	<a href="#">Ion Channel Gating</a>
New Tree	<a href="#">G04.835.480</a>	<a href="#">Light Signal Transduction</a>
New Tree	<a href="#">G04.835.480.900</a>	<a href="#">Vision, Ocular</a>
New Tree	<a href="#">G04.835.560</a>	<a href="#">MAP Kinase Signaling System</a>
New Tree	<a href="#">G04.835.580</a>	<a href="#">Mechanotransduction, Cellular</a>
New Heading	<b><a href="#">G04.835.690</a></b>	<b><a href="#">Non-Neuronal Cholinergic System</a></b>
New Tree	<a href="#">G04.835.800</a>	<a href="#">Second Messenger Systems</a>
New Tree	<a href="#">G04.835.800.100</a>	<a href="#">Calcium Signaling</a>

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	G04.835.850	Synaptic Transmission
New Tree	G04.835.850.500	Postsynaptic Potential Summation
-	G05	Genetic Phenomena
New Tree	G05.045	Biological Evolution
New Tree	G05.045.250	Evolution, Molecular
New Tree	G05.045.250.750	Mutation Rate
New Tree	G05.045.300	Genetic Drift
New Tree	G05.045.350	Genetic Speciation
New Heading	<b>G05.045.513</b>	<b>Life History Traits</b>
New Tree	G05.045.675	Reproductive Isolation
New Tree	G05.090	Breeding
New Tree	G05.090.390	Hybridization, Genetic
New Tree	G05.090.403	Inbreeding
New Tree	G05.090.403.180	Consanguinity
New Tree	G05.090.403.590	Self-Fertilization
New Tree	G05.090.701	Selective Breeding
New Tree	G05.113	Cell Division
New Tree	G05.113.109	Asymmetric Cell Division
New Tree	G05.113.220	Cell Nucleus Division
New Tree	G05.113.220.625	Chromosome Segregation
New Tree	G05.113.220.625.620	Nondisjunction, Genetic
New Tree	G05.113.220.687	Meiosis

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	<a href="#">G05.113.220.687.250</a>	<a href="#">Anaphase</a>
New Tree	<a href="#">G05.113.220.687.500</a>	<a href="#">Meiotic Prophase I</a>
New Tree	<a href="#">G05.113.220.687.625</a>	<a href="#">Metaphase</a>
New Tree	<a href="#">G05.113.220.687.812</a>	<a href="#">Prometaphase</a>
New Tree	<a href="#">G05.113.220.687.883</a>	<a href="#">Prophase</a>
New Tree	<a href="#">G05.113.220.687.883.250</a>	<a href="#">Chromosome Pairing</a>
New Tree	<a href="#">G05.113.220.687.883.250.500</a>	<a href="#">Synaptonemal Complex</a>
New Tree	<a href="#">G05.113.220.687.883.750</a>	<a href="#">Pachytene Stage</a>
New Tree	<a href="#">G05.113.220.687.953</a>	<a href="#">Telophase</a>
New Tree	<a href="#">G05.113.220.781</a>	<a href="#">Mitosis</a>
New Tree	<a href="#">G05.113.220.781.050</a>	<a href="#">Anaphase</a>
New Tree	<a href="#">G05.113.220.781.338</a>	<a href="#">M Phase Cell Cycle Checkpoints</a>
New Tree	<a href="#">G05.113.220.781.625</a>	<a href="#">Metaphase</a>
New Tree	<a href="#">G05.113.220.781.812</a>	<a href="#">Prometaphase</a>
New Tree	<a href="#">G05.113.220.781.906</a>	<a href="#">Prophase</a>
New Tree	<a href="#">G05.113.220.781.906.250</a>	<a href="#">Chromosome Pairing</a>
New Tree	<a href="#">G05.113.220.781.906.250.500</a>	<a href="#">Synaptonemal Complex</a>
New Tree	<a href="#">G05.113.220.781.906.750</a>	<a href="#">Pachytene Stage</a>
New Tree	<a href="#">G05.113.220.781.953</a>	<a href="#">Telophase</a>
New Tree	<a href="#">G05.113.415</a>	<a href="#">Cell Self Renewal</a>
New Tree	<a href="#">G05.113.610</a>	<a href="#">Telomere Homeostasis</a>
New Tree	<a href="#">G05.135</a>	<a href="#">Cellular Reprogramming</a>

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	G05.158	Clonal Evolution
-	G05.180	Consanguinity
New Tree	G05.193	DNA Cleavage
New Tree	G05.200	DNA Damage
New Tree	G05.200.104	DNA Adducts
New Tree	G05.200.210	DNA Breaks
New Tree	G05.200.210.170	Chromosome Breakage
New Tree	G05.200.210.170.500	Chromosome Breakpoints
New Tree	G05.200.210.220	DNA Breaks, Double-Stranded
New Tree	G05.200.210.230	DNA Breaks, Single-Stranded
New Tree	G05.200.220	DNA Degradation, Necrotic
New Tree	G05.200.230	DNA Fragmentation
New Tree	G05.206	DNA Methylation
New Tree	G05.213	DNA Packaging
New Tree	G05.213.095	Chromatin Assembly and Disassembly
New Tree	G05.219	DNA Repair
New Tree	G05.219.200	DNA End-Joining Repair
New Tree	G05.219.220	DNA Mismatch Repair
New Tree	G05.219.700	Recombinational DNA Repair
New Tree	G05.219.830	SOS Response (Genetics)
New Tree	G05.226	DNA Replication
New Tree	G05.226.760	DNA Replication Timing

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	<a href="#">G05.226.820</a>	<a href="#">Endoreduplication</a>
New Tree	<a href="#">G05.226.880</a>	<a href="#">S Phase</a>
New Tree	<a href="#">G05.226.940</a>	<a href="#">Telomere Shortening</a>
-	G05.232	DNA Transformation Competence
-	G05.285	Founder Effect
New Tree	<a href="#">G05.297</a>	<a href="#">Gene Expression</a>
New Tree	<a href="#">G05.297.670</a>	<a href="#">Protein Biosynthesis</a>
New Tree	<a href="#">G05.297.700</a>	<a href="#">Transcription, Genetic</a>
New Tree	<a href="#">G05.297.700.500</a>	<a href="#">Reverse Transcription</a>
New Tree	<a href="#">G05.297.700.562</a>	<a href="#">Transcription Elongation, Genetic</a>
New Tree	<a href="#">G05.297.700.625</a>	<a href="#">Transcription Initiation, Genetic</a>
New Tree	<a href="#">G05.297.700.687</a>	<a href="#">Transcription Termination, Genetic</a>
New Tree	<a href="#">G05.297.700.750</a>	<a href="#">Transcriptome</a>
New Tree	<a href="#">G05.308</a>	<a href="#">Gene Expression Regulation</a>
New Tree	<a href="#">G05.308.048</a>	<a href="#">Catabolite Repression</a>
New Tree	<a href="#">G05.308.095</a>	<a href="#">Chromatin Assembly and Disassembly</a>
New Tree	<a href="#">G05.308.200</a>	<a href="#">Down-Regulation</a>
New Tree	<a href="#">G05.308.202</a>	<a href="#">Ectopic Gene Expression</a>
New Tree	<a href="#">G05.308.203</a>	<a href="#">Epigenesis, Genetic</a>
New Tree	<a href="#">G05.308.203.124</a>	<a href="#">Chromosomal Position Effects</a>
New Tree	<a href="#">G05.308.203.249</a>	<a href="#">Dosage Compensation, Genetic</a>
New Tree	<a href="#">G05.308.203.249.970</a>	<a href="#">X Chromosome Inactivation</a>
New	<a href="#">G05.308.203.311</a>	<a href="#">Epigenetic Repression</a>

## MeSH Tree Changes for 2017

Type	Tree - heading
Tree	
New Tree	G05.308.203.374 Gene Silencing
New Tree	G05.308.203.374.394 CRISPR-Cas Systems
New Tree	G05.308.203.374.790 RNA Interference
New Tree	G05.308.203.500 Genomic Imprinting
New Tree	G05.308.207 Epistasis, Genetic
New Tree	G05.308.215 Frameshifting, Ribosomal
New Tree	G05.308.250 Gene Amplification
New Tree	G05.308.290 Gene Expression Regulation, Archaeal
New Tree	G05.308.300 Gene Expression Regulation, Bacterial
New Tree	G05.308.310 Gene Expression Regulation, Developmental
New Tree	G05.308.320 Gene Expression Regulation, Enzymologic
New Tree	G05.308.320.200 Enzyme Induction
New Tree	G05.308.320.300 Enzyme Repression
New Tree	G05.308.330 Gene Expression Regulation, Fungal
New Tree	G05.308.370 Gene Expression Regulation, Neoplastic
New Tree	G05.308.370.500 Gene Expression Regulation, Leukemic
New Tree	G05.308.375 Gene Expression Regulation, Plant
New Tree	G05.308.385 Gene Expression Regulation, Viral
New Tree	G05.308.670 Protein Modification, Translational
New Tree	G05.308.670.600 Protein Processing, Post-Translational
New Tree	G05.308.670.600.400 Protein Prenylation

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	G05.308.670.600.700	Protein Splicing
New Tree	G05.308.670.600.831	Ubiquitination
New Tree	G05.308.670.600.831.500	Sumoylation
New Tree	G05.308.670.600.850	Unfolded Protein Response
New Tree	G05.308.670.600.850.500	Endoplasmic Reticulum-Associated Degradation
New Tree	G05.308.700	RNA Processing, Post-Transcriptional
New Tree	G05.308.700.112	Nonsense Mediated mRNA Decay
New Tree	G05.308.700.225	RNA 3' End Processing
New Tree	G05.308.700.225.710	Polyadenylation
New Tree	G05.308.700.250	RNA Editing
New Tree	G05.308.700.700	RNA Splicing
New Tree	G05.308.700.700.100	Alternative Splicing
New Tree	G05.308.700.700.750	Trans-Splicing
New Tree	G05.308.800	Transcriptional Activation
New Tree	G05.308.850	Up-Regulation
-	G05.330	Gene Frequency
-	G05.330.159	Gene Flow
-	G05.330.320	Genetic Drift
-	G05.340	Gene Order
New Tree	G05.342	Gene Pool
New Tree	G05.344	Gene Rearrangement
New Tree	G05.344.401	Gene Rearrangement, B-Lymphocyte
New Tree	G05.344.401.501	Gene Rearrangement, B-Lymphocyte, Heavy Chain
New	G05.344.401.501.450	Immunoglobulin Class Switching



## MeSH Tree Changes for 2017

Type	Tree - heading	
Tree		
New Tree	<a href="#">G05.344.401.601</a>	<a href="#">Gene Rearrangement, B-Lymphocyte, Light Chain</a>
New Tree	<a href="#">G05.344.801</a>	<a href="#">Gene Rearrangement, T-Lymphocyte</a>
New Tree	<a href="#">G05.344.801.111</a>	<a href="#">Gene Rearrangement, alpha-Chain T-Cell Antigen Receptor</a>
New Tree	<a href="#">G05.344.801.211</a>	<a href="#">Gene Rearrangement, beta-Chain T-Cell Antigen Receptor</a>
New Tree	<a href="#">G05.344.801.261</a>	<a href="#">Gene Rearrangement, delta-Chain T-Cell Antigen Receptor</a>
New Tree	<a href="#">G05.344.801.311</a>	<a href="#">Gene Rearrangement, gamma-Chain T-Cell Antigen Receptor</a>
New Tree	<a href="#">G05.344.900</a>	<a href="#">V(D)J Recombination</a>
Old Tree	<del>G05.345</del>	<del>Gene Pool</del>
-	G05.346	Genetic Background
-	G05.347	Genetic Fitness
New Tree	<a href="#">G05.348</a>	<a href="#">Genetic Linkage</a>
New Tree	<a href="#">G05.348.500</a>	<a href="#">Linkage Disequilibrium</a>
New Tree	<a href="#">G05.348.750</a>	<a href="#">Lod Score</a>
-	G05.350	Genetic Load
Old Tree	<del>G05.355</del>	<del>Genetic Processes</del>
Old Tree	<del>G05.355.044</del>	<del>Biological Evolution</del>
Old Tree	<del>G05.355.044.250</del>	<del>Evolution, Molecular</del>
Old Tree	<del>G05.355.044.250.750</del>	<del>Mutation Rate</del>
Old Tree	<del>G05.355.044.300</del>	<del>Genetic Drift</del>
Old Tree	<del>G05.355.044.350</del>	<del>Genetic Speciation</del>
Old Tree	<del>G05.355.044.675</del>	<del>Reproductive Isolation</del>
Old Tree	<del>G05.355.090</del>	<del>Breeding</del>
Old Tree	<del>G05.355.090.390</del>	<del>Hybridization, Genetic</del>
Old Tree	<del>G05.355.090.403</del>	<del>Inbreeding</del>
Old Tree	<del>G05.355.090.403.180</del>	<del>Consanguinity</del>
Old Tree	<del>G05.355.090.403.590</del>	<del>Self-Fertilization</del>
Old Tree	<del>G05.355.090.701</del>	<del>Selective Breeding</del>
Old Tree	<del>G05.355.105</del>	<del>Cell Division</del>

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	G05.355.105.109 Asymmetric Cell Division
Old Tree	G05.355.105.220 Cell Nucleus Division
Old Tree	G05.355.105.220.625 Chromosome Segregation
Old Tree	G05.355.105.220.625.620 Nondisjunction, Genetic
Old Tree	G05.355.105.220.687 Meiosis
Old Tree	G05.355.105.220.687.250 Anaphase
Old Tree	G05.355.105.220.687.500 Meiotic Prophase I
Old Tree	G05.355.105.220.687.625 Metaphase
Old Tree	G05.355.105.220.687.812 Prometaphase
Old Tree	G05.355.105.220.687.883 Prophase
Old Tree	G05.355.105.220.687.883.250 Chromosome Pairing
Old Tree	G05.355.105.220.687.883.250.500 Synaptonemal Complex
Old Tree	G05.355.105.220.687.883.750 Pachytene Stage
Old Tree	G05.355.105.220.687.953 Telophase
Old Tree	G05.355.105.220.781 Mitosis
Old Tree	G05.355.105.220.781.050 Anaphase
Old Tree	G05.355.105.220.781.338 M Phase Cell Cycle Checkpoints
Old Tree	G05.355.105.220.781.625 Metaphase
Old Tree	G05.355.105.220.781.812 Prometaphase
Old Tree	G05.355.105.220.781.906 Prophase
Old Tree	G05.355.105.220.781.906.250 Chromosome Pairing
Old Tree	G05.355.105.220.781.906.250.500 Synaptonemal Complex
Old Tree	G05.355.105.220.781.906.750 Pachytene Stage
Old Tree	G05.355.105.220.781.953 Telophase
Old Tree	G05.355.105.415 Cell Self Renewal
Old Tree	G05.355.105.610 Telomere Homeostasis
Old Tree	G05.355.114 Cellular Reprogramming
Old Tree	G05.355.123 Clonal Evolution
Old Tree	G05.355.142 DNA Cleavage
Old Tree	G05.355.180 DNA Damage
Old Tree	G05.355.180.104 DNA Adducts
Old Tree	G05.355.180.210 DNA Breaks
Old Tree	G05.355.180.210.170 Chromosome Breakage
Old Tree	G05.355.180.210.170.500 Chromosome Breakpoints
Old Tree	G05.355.180.210.220 DNA Breaks, Double-Stranded
Old Tree	G05.355.180.210.230 DNA Breaks, Single-Stranded

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	G05.355.180.220 DNA Degradation, Necrotic
Old Tree	G05.355.180.230 DNA Fragmentation
Old Tree	G05.355.190 DNA Methylation
Old Tree	G05.355.192 DNA Packaging
Old Tree	G05.355.192.095 Chromatin Assembly and Disassembly
Old Tree	G05.355.195 DNA Repair
Old Tree	G05.355.195.200 DNA End-Joining Repair
Old Tree	G05.355.195.220 DNA Mismatch Repair
Old Tree	G05.355.195.700 Recombinational DNA Repair
Old Tree	G05.355.195.830 SOS Response (Genetics)
Old Tree	G05.355.200 DNA Replication
Old Tree	G05.355.200.760 DNA Replication Timing
Old Tree	G05.355.200.820 Endoreduplication
Old Tree	G05.355.200.880 S Phase
Old Tree	G05.355.200.940 Telomere Shortening
Old Tree	G05.355.310 Gene Expression
Old Tree	G05.355.310.670 Protein Biosynthesis
Old Tree	G05.355.310.700 Transcription, Genetic
Old Tree	G05.355.310.700.500 Reverse Transcription
Old Tree	G05.355.310.700.562 Transcription Elongation, Genetic
Old Tree	G05.355.310.700.625 Transcription Initiation, Genetic
Old Tree	G05.355.310.700.687 Transcription Termination, Genetic
Old Tree	G05.355.310.700.750 Transcriptome
Old Tree	G05.355.315 Gene Expression Regulation
Old Tree	G05.355.315.095 Chromatin Assembly and Disassembly
Old Tree	G05.355.315.200 Down-Regulation
Old Tree	G05.355.315.202 Ectopic Gene Expression
Old Tree	G05.355.315.203 Epigenesis, Genetic
Old Tree	G05.355.315.203.124 Chromosomal Position Effects
Old Tree	G05.355.315.203.249 Dosage Compensation, Genetic
Old Tree	G05.355.315.203.249.970 X Chromosome Inactivation
Old Tree	G05.355.315.203.311 Epigenetic Repression
Old Tree	G05.355.315.203.374 Gene Silencing
Old Tree	G05.355.315.203.374.394 CRISPR-Cas Systems
Old Tree	G05.355.315.203.374.790 RNA Interference
Old Tree	G05.355.315.203.500 Genomic Imprinting

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	G05.355.315.207 Epistasis, Genetic
Old Tree	G05.355.315.215 Frameshifting, Ribosomal
Old Tree	G05.355.315.250 Gene Amplification
Old Tree	G05.355.315.290 Gene Expression Regulation, Archaeal
Old Tree	G05.355.315.300 Gene Expression Regulation, Bacterial
Old Tree	G05.355.315.310 Gene Expression Regulation, Developmental
Old Tree	G05.355.315.320 Gene Expression Regulation, Enzymologic
Old Tree	G05.355.315.320.200 Enzyme Induction
Old Tree	G05.355.315.320.300 Enzyme Repression
Old Tree	G05.355.315.330 Gene Expression Regulation, Fungal
Old Tree	G05.355.315.370 Gene Expression Regulation, Neoplastic
Old Tree	G05.355.315.370.500 Gene Expression Regulation, Leukemic
Old Tree	G05.355.315.375 Gene Expression Regulation, Plant
Old Tree	G05.355.315.385 Gene Expression Regulation, Viral
Old Tree	G05.355.315.670 Protein Modification, Translational
Old Tree	G05.355.315.670.600 Protein Processing, Post-Translational
Old Tree	G05.355.315.670.600.400 Protein Prenylation
Old Tree	G05.355.315.670.600.700 Protein Splicing
Old Tree	G05.355.315.670.600.831 Ubiquitination
Old Tree	G05.355.315.670.600.831.500 Sumoylation
Old Tree	G05.355.315.670.600.850 Unfolded Protein Response
Old Tree	G05.355.315.670.600.850.500 Endoplasmic Reticulum-Associated Degradation
Old Tree	G05.355.315.700 RNA Processing, Post-Transcriptional
Old Tree	G05.355.315.700.112 Nonsense Mediated mRNA Decay
Old Tree	G05.355.315.700.225 RNA 3' End Processing
Old Tree	G05.355.315.700.225.710 Polyadenylation
Old Tree	G05.355.315.700.250 RNA Editing
Old Tree	G05.355.315.700.700 RNA Splicing
Old Tree	G05.355.315.700.700.100 Alternative Splicing
Old Tree	G05.355.315.700.700.750 Trans-Splicing
Old Tree	G05.355.315.800 Transcriptional Activation
Old Tree	G05.355.315.850 Up-Regulation
Old Tree	G05.355.330 Gene Rearrangement
Old Tree	G05.355.330.401 Gene Rearrangement, B-Lymphocyte
Old Tree	G05.355.330.401.501 Gene Rearrangement, B-Lymphocyte, Heavy Chain

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	G05.355.330.401.501.450 Immunoglobulin Class Switching
Old Tree	G05.355.330.401.601 Gene Rearrangement, B-Lymphocyte, Light Chain
Old Tree	G05.355.330.801 Gene Rearrangement, T-Lymphocyte
Old Tree	G05.355.330.801.111 Receptor Gene Rearrangement, alpha-Chain T-Cell Antigen
Old Tree	G05.355.330.801.211 Receptor Gene Rearrangement, beta-Chain T-Cell Antigen
Old Tree	G05.355.330.801.261 Receptor Gene Rearrangement, delta-Chain T-Cell Antigen
Old Tree	G05.355.330.801.311 Receptor Gene Rearrangement, gamma-Chain T-Cell Antigen
Old Tree	G05.355.330.900 V(D)J Recombination
Old Tree	G05.355.380 Heredity
Old Tree	G05.355.590 Molecular Mimicry
Old Tree	G05.355.600 Mutagenesis
Old Tree	G05.355.600.109 Amino Acid Substitution
Old Tree	G05.355.600.164 Chromosome Duplication
Old Tree	G05.355.600.220 DNA Repeat Expansion
Old Tree	G05.355.600.220.865 Trinucleotide Repeat Expansion
Old Tree	G05.355.600.315 Gene Amplification
Old Tree	G05.355.600.320 Gene Duplication
Old Tree	G05.355.600.370 INDEL Mutation
Old Tree	G05.355.600.550 Mutagenesis, Insertional
Old Tree	G05.355.600.620 Nondisjunction, Genetic
Old Tree	G05.355.600.800 Sequence Deletion
Old Tree	G05.355.600.800.180 Chromosome Deletion
Old Tree	G05.355.600.800.320 Gene Deletion
Old Tree	G05.355.600.805 Sequence Inversion
Old Tree	G05.355.600.805.500 Chromosome Inversion
Old Tree	G05.355.600.810 Somatic Hypermutation, Immunoglobulin
Old Tree	G05.355.600.835 Suppression, Genetic
Old Tree	G05.355.600.860 Translocation, Genetic
Old Tree	G05.355.720 Nucleic Acid Denaturation
Old Tree	G05.355.760 Recombination, Genetic
Old Tree	G05.355.760.200 Conjugation, Genetic
Old Tree	G05.355.760.385 Gene Fusion
Old Tree	G05.355.760.385.500 Oncogene Fusion

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	G05.355.760.390 Gene Transfer, Horizontal
Old Tree	G05.355.760.615 Homologous Recombination
Old Tree	G05.355.760.615.200 Crossing Over, Genetic
Old Tree	G05.355.760.615.475 Gene Conversion
Old Tree	G05.355.760.615.612 Recombinational DNA Repair
Old Tree	G05.355.760.615.750 Sister Chromatid Exchange
Old Tree	G05.355.760.850 Transduction, Genetic
Old Tree	G05.355.760.860 Transfection
Old Tree	G05.355.760.860.500 Transformation, Bacterial
Old Tree	G05.355.760.865 Transformation, Genetic
Old Tree	G05.355.760.865.820 Transformation, Bacterial
Old Tree	G05.355.760.932 V(D)J Recombination
Old Tree	G05.355.780 RNA Cleavage
Old Tree	G05.355.800 Selection, Genetic
Old Tree	G05.355.865 Sex Determination Processes
Old Tree	G05.355.930 Virus Integration
Old Tree	G05.355.930.500 Lysogeny
-	G05.360 Genetic Structures
-	G05.360.080 Base Sequence
-	G05.360.080.040 AT Rich Sequence
-	G05.360.080.380 GC Rich Sequence
-	G05.360.080.380.160 CpG Islands
-	G05.360.080.534 Matrix Attachment Regions
-	G05.360.080.611 Nucleotide Motifs
-	G05.360.080.689 Regulatory Sequences, Nucleic Acid
-	G05.360.080.689.330 Enhancer Elements, Genetic
-	G05.360.080.689.330.240 E-Box Elements
-	G05.360.080.689.330.400 HIV Enhancer
-	G05.360.080.689.330.700 Response Elements
-	G05.360.080.689.330.700.800 Serum Response Element
-	G05.360.080.689.330.700.920 Vitamin D Response Element
-	G05.360.080.689.360 Gene Regulatory Networks
-	G05.360.080.689.390 Insulator Elements
-	G05.360.080.689.450 Locus Control Region
-	G05.360.080.689.650 Operator Regions, Genetic
-	G05.360.080.689.675 Promoter Regions, Genetic

## MeSH Tree Changes for 2017

Type	Tree - heading
-	G05.360.080.689.675.700 Response Elements
-	G05.360.080.689.675.700.040 Antioxidant Response Elements
-	G05.360.080.689.675.700.800 Serum Response Element
-	G05.360.080.689.675.700.920 Vitamin D Response Element
-	G05.360.080.689.675.850 TATA Box
-	G05.360.080.689.687 Regulatory Sequences, Ribonucleic Acid
-	G05.360.080.689.687.061 AU Rich Elements
-	G05.360.080.689.687.093 Internal Ribosome Entry Sites
-	G05.360.080.689.687.124 Riboswitch
-	G05.360.080.689.687.249 RNA 3' Polyadenylation Signals
-	G05.360.080.689.687.275 RNA 5' Terminal Oligopyrimidine Sequence
-	G05.360.080.689.687.490 RNA Splice Sites
-	G05.360.080.689.755 Silencer Elements, Transcriptional
-	G05.360.080.689.810 Terminator Regions, Genetic
-	G05.360.080.708 Repetitive Sequences, Nucleic Acid
-	G05.360.080.708.330 Interspersed Repetitive Sequences
-	G05.360.080.708.330.200 DNA Transposable Elements
-	G05.360.080.708.330.330 Genomic Islands
-	G05.360.080.708.330.800 Retroelements
-	G05.360.080.708.330.800.175 Endogenous Retroviruses
-	G05.360.080.708.330.800.200 Genes, Intracisternal A-Particle
-	G05.360.080.708.330.800.400 Long Interspersed Nucleotide Elements
-	G05.360.080.708.330.800.800 Short Interspersed Nucleotide Elements
-	G05.360.080.708.330.800.800.050 Alu Elements
-	G05.360.080.708.565 Segmental Duplications, Genomic
-	G05.360.080.708.800 Tandem Repeat Sequences
-	G05.360.080.708.800.074 DNA Repeat Expansion
-	G05.360.080.708.800.074.865 Trinucleotide Repeat Expansion
-	G05.360.080.708.800.150 DNA, Satellite
-	G05.360.080.708.800.325 Inverted Repeat Sequences
-	G05.360.080.708.800.325.500 Clustered Regularly Interspaced Short Palindromic Repeats
-	G05.360.080.708.800.500 Microsatellite Repeats
-	G05.360.080.708.800.500.150 Dinucleotide Repeats
-	G05.360.080.708.800.500.850 Trinucleotide Repeats
-	G05.360.080.708.800.500.850.200 Trinucleotide Repeat Expansion

## MeSH Tree Changes for 2017

Type	Tree - heading
-	G05.360.080.708.800.550                      Minisatellite Repeats
-	G05.360.080.708.850                              Terminal Repeat Sequences
-	G05.360.080.708.850.400                        HIV Long Terminal Repeat
-	G05.360.080.708.850.400.400                 HIV Enhancer
-	G05.360.160                                        Chromosome Structures
-	G05.360.160.165                                 Centromere
-	G05.360.160.165.500                            Kinetochores
-	G05.360.160.175                                 Chromatids
-	G05.360.160.180                                 Chromatin
-	G05.360.160.180.270                            Euchromatin
-	G05.360.160.180.383                            Heterochromatin
-	G05.360.160.180.383.800                      Sex Chromatin
-	G05.360.160.180.625                            Nucleosomes
-	G05.360.160.650                                 Nucleolus Organizer Region
-	G05.360.160.830                                 Synaptonemal Complex
-	G05.360.160.845                                 Telomere
-	G05.360.162                                        Chromosomes
-	G05.360.162.167                                 Chromosomes, Archaeal
-	G05.360.162.178                                 Chromosomes, Artificial
-	G05.360.162.178.170                            Chromosomes, Artificial, Bacterial
-	G05.360.162.178.190                            Chromosomes, Artificial, Mammalian
-	G05.360.162.178.190.117                      Chromosomes, Artificial, Human
-	G05.360.162.178.195                            Chromosomes, Artificial, P1 Bacteriophage
-	G05.360.162.178.200                            Chromosomes, Artificial, Yeast
-	G05.360.162.190                                 Chromosomes, Bacterial
-	G05.360.162.190.170                            Chromosomes, Artificial, Bacterial
-	G05.360.162.360                                 Chromosomes, Fungal
-	G05.360.162.360.800                            Chromosomes, Artificial, Yeast
-	G05.360.162.440                                 Chromosomes, Insect
-	G05.360.162.440.500                            Polytene Chromosomes
-	G05.360.162.520                                 Chromosomes, Mammalian
-	G05.360.162.520.190                            Chromosomes, Artificial, Mammalian
-	G05.360.162.520.190.117                      Chromosomes, Artificial, Human
-	G05.360.162.520.300                            Chromosomes, Human
-	G05.360.162.520.300.117                      Chromosomes, Artificial, Human
-	G05.360.162.520.300.235                      Chromosomes, Human, 1-3



## MeSH Tree Changes for 2017

Type	Tree - heading
-	G05.360.162.520.300.235.240 Chromosomes, Human, Pair 1
-	G05.360.162.520.300.235.245 Chromosomes, Human, Pair 2
-	G05.360.162.520.300.235.250 Chromosomes, Human, Pair 3
-	G05.360.162.520.300.280 Chromosomes, Human, 4-5
-	G05.360.162.520.300.280.285 Chromosomes, Human, Pair 4
-	G05.360.162.520.300.280.290 Chromosomes, Human, Pair 5
-	G05.360.162.520.300.325 Chromosomes, Human, 6-12 and X
-	G05.360.162.520.300.325.330 Chromosomes, Human, Pair 6
-	G05.360.162.520.300.325.335 Chromosomes, Human, Pair 7
-	G05.360.162.520.300.325.340 Chromosomes, Human, Pair 8
-	G05.360.162.520.300.325.345 Chromosomes, Human, Pair 9
-	G05.360.162.520.300.325.345.700 Philadelphia Chromosome
-	G05.360.162.520.300.325.350 Chromosomes, Human, Pair 10
-	G05.360.162.520.300.325.355 Chromosomes, Human, Pair 11
-	G05.360.162.520.300.325.360 Chromosomes, Human, Pair 12
-	G05.360.162.520.300.325.680 Chromosomes, Human, X
-	G05.360.162.520.300.370 Chromosomes, Human, 13-15
-	G05.360.162.520.300.370.375 Chromosomes, Human, Pair 13
-	G05.360.162.520.300.370.380 Chromosomes, Human, Pair 14
-	G05.360.162.520.300.370.385 Chromosomes, Human, Pair 15
-	G05.360.162.520.300.415 Chromosomes, Human, 16-18
-	G05.360.162.520.300.415.420 Chromosomes, Human, Pair 16
-	G05.360.162.520.300.415.425 Chromosomes, Human, Pair 17
-	G05.360.162.520.300.415.430 Chromosomes, Human, Pair 18
-	G05.360.162.520.300.460 Chromosomes, Human, 19-20
-	G05.360.162.520.300.460.465 Chromosomes, Human, Pair 19
-	G05.360.162.520.300.460.470 Chromosomes, Human, Pair 20
-	G05.360.162.520.300.505 Chromosomes, Human, 21-22 and Y
-	G05.360.162.520.300.505.510 Chromosomes, Human, Pair 21
-	G05.360.162.520.300.505.515 Chromosomes, Human, Pair 22
-	G05.360.162.520.300.505.515.700 Philadelphia Chromosome
-	G05.360.162.520.300.505.757 Chromosomes, Human, Y
-	G05.360.162.560 Chromosomes, Plant
-	G05.360.162.570 Isochromosomes
-	G05.360.162.679 Karyotype
-	G05.360.162.679.500 Abnormal Karyotype

## MeSH Tree Changes for 2017

Type	Tree - heading
-	G05.360.162.679.500.500 XYY Karyotype
-	G05.360.162.788 Ring Chromosomes
-	G05.360.162.865 Sex Chromosomes
New Heading	<b>G05.360.162.865.400</b> <b>Pseudoautosomal Regions</b>
-	G05.360.162.865.800 Sex Chromatin
-	G05.360.162.865.982 X Chromosome
-	G05.360.162.865.982.500 Chromosomes, Human, X
-	G05.360.162.865.983 Y Chromosome
-	G05.360.162.865.983.500 Chromosomes, Human, Y
-	G05.360.325 Gene Library
-	G05.360.325.425 Genomic Library
-	G05.360.325.640 Peptide Library
-	G05.360.335 Genetic Code
-	G05.360.335.060 Anticodon
-	G05.360.335.355 Codon
-	G05.360.335.355.225 Codon, Initiator
-	G05.360.335.355.250 Codon, Terminator
-	G05.360.335.355.250.235 Codon, Nonsense
-	G05.360.335.760 Reading Frames
-	G05.360.335.760.640 Open Reading Frames
-	G05.360.337 Genetic Vectors
-	G05.360.337.249 Chromosomes, Artificial
-	G05.360.337.249.170 Chromosomes, Artificial, Bacterial
-	G05.360.337.249.190 Chromosomes, Artificial, Mammalian
-	G05.360.337.249.190.117 Chromosomes, Artificial, Human
-	G05.360.337.249.195 Chromosomes, Artificial, P1 Bacteriophage
-	G05.360.337.249.200 Chromosomes, Artificial, Yeast
-	G05.360.337.500 Cosmids
-	G05.360.340 Genome
-	G05.360.340.011 Exome
-	G05.360.340.024 Genome Components
-	G05.360.340.024.079 Attachment Sites, Microbiological
-	G05.360.340.024.159 CpG Islands
-	G05.360.340.024.189 DNA Sequence, Unstable
-	G05.360.340.024.189.220 DNA Repeat Expansion

## MeSH Tree Changes for 2017

Type	Tree - heading
-	G05.360.340.024.189.220.865 Trinucleotide Repeat Expansion
-	G05.360.340.024.189.610 Chromosome Fragile Sites
-	G05.360.340.024.220 DNA, Intergenic
-	G05.360.340.024.220.150 DNA, Satellite
-	G05.360.340.024.220.280 3' Flanking Region
-	G05.360.340.024.220.282 5' Flanking Region
-	G05.360.340.024.220.400 Introns
-	G05.360.340.024.220.760 Replication Origin
-	G05.360.340.024.220.880 Untranslated Regions
-	G05.360.340.024.220.880.880 3' Untranslated Regions
-	G05.360.340.024.220.880.880.500 AU Rich Elements
-	G05.360.340.024.220.880.885 5' Untranslated Regions
-	G05.360.340.024.340 Genes
-	G05.360.340.024.340.030 Alleles
-	G05.360.340.024.340.137 Gene Components
-	G05.360.340.024.340.137.190 Codon
-	G05.360.340.024.340.137.190.225 Codon, Initiator
-	G05.360.340.024.340.137.190.250 Codon, Terminator
-	G05.360.340.024.340.137.232 Exons
-	G05.360.340.024.340.137.232.459 Hinge Exons
-	G05.360.340.024.340.137.232.920 VDJ Exons
-	G05.360.340.024.340.137.275 Expressed Sequence Tags
-	G05.360.340.024.340.137.290 3' Flanking Region
-	G05.360.340.024.340.137.295 5' Flanking Region
-	G05.360.340.024.340.137.430 Immunoglobulin Switch Region
-	G05.360.340.024.340.137.515 Introns
-	G05.360.340.024.340.137.650 Open Reading Frames
-	G05.360.340.024.340.137.750 Regulatory Elements, Transcriptional
-	G05.360.340.024.340.137.750.249 Enhancer Elements, Genetic
-	G05.360.340.024.340.137.750.249.240 E-Box Elements
-	G05.360.340.024.340.137.750.249.400 HIV Enhancer
-	G05.360.340.024.340.137.750.249.765 Response Elements
-	G05.360.340.024.340.137.750.249.765.800 Serum Response Element
-	G05.360.340.024.340.137.750.249.765.920 Vitamin D Response Element
-	G05.360.340.024.340.137.750.680 Promoter Regions, Genetic
-	G05.360.340.024.340.137.750.680.765 Response Elements

## MeSH Tree Changes for 2017

Type	Tree - heading
-	G05.360.340.024.340.137.750.680.765.040      Antioxidant Response Elements
-	G05.360.340.024.340.137.750.680.765.800      Serum Response Element
-	G05.360.340.024.340.137.750.680.765.920      Vitamin D Response Element
-	G05.360.340.024.340.137.750.680.850      TATA Box
-	G05.360.340.024.340.137.750.830      Terminator Regions, Genetic
-	G05.360.340.024.340.137.750.840      Transcription Initiation Site
-	G05.360.340.024.340.137.775      RNA 3' Polyadenylation Signals
-	G05.360.340.024.340.137.785      RNA 5' Terminal Oligopyrimidine Sequence
-	G05.360.340.024.340.137.800      RNA Splice Sites
-	G05.360.340.024.340.137.910      Untranslated Regions
-	G05.360.340.024.340.137.910.880      3' Untranslated Regions
-	G05.360.340.024.340.137.910.880.500      AU Rich Elements
-	G05.360.340.024.340.137.910.885      5' Untranslated Regions
-	G05.360.340.024.340.220      Genes, cdc
-	G05.360.340.024.340.225      Genes, Chloroplast
-	G05.360.340.024.340.230      Genes, Developmental
-	G05.360.340.024.340.230.500      Genes, Homeobox
-	G05.360.340.024.340.240      Genes, Dominant
-	G05.360.340.024.340.250      Genes, Duplicate
-	G05.360.340.024.340.270      Genes, Essential
-	G05.360.340.024.340.310      Genes, Helminth
-	G05.360.340.024.340.330      Genes, Immediate-Early
-	G05.360.340.024.340.335      Genes, Immunoglobulin
-	G05.360.340.024.340.335.300      Genes, Immunoglobulin Heavy Chain
-	G05.360.340.024.340.335.300.249      Hinge Exons
-	G05.360.340.024.340.335.300.500      Immunoglobulin Switch Region
-	G05.360.340.024.340.335.310      Genes, Immunoglobulin Light Chain
-	G05.360.340.024.340.335.655      VDJ Exons
-	G05.360.340.024.340.340      Genes, Insect
-	G05.360.340.024.340.350      Genes, Lethal
-	G05.360.340.024.340.361      Genes, MDR
-	G05.360.340.024.340.364      Genes, Microbial
-	G05.360.340.024.340.364.124      Genes, Archaeal
-	G05.360.340.024.340.364.249      Genes, Bacterial
-	G05.360.340.024.340.364.500      Genes, Fungal
-	G05.360.340.024.340.364.500.089      Genes, Mating Type, Fungal

## MeSH Tree Changes for 2017

Type	Tree - heading
-	G05.360.340.024.340.364.875 Genes, Viral
-	G05.360.340.024.340.364.875.172 Genes, env
-	G05.360.340.024.340.364.875.258 Genes, gag
-	G05.360.340.024.340.364.875.345 Genes, Immediate-Early
-	G05.360.340.024.340.364.875.360 Genes, Intracisternal A-Particle
-	G05.360.340.024.340.364.875.600 Genes, nef
-	G05.360.340.024.340.364.875.667 Genes, pol
-	G05.360.340.024.340.364.875.735 Genes, pX
-	G05.360.340.024.340.364.875.775 Genes, rev
-	G05.360.340.024.340.364.875.850 Genes, tat
-	G05.360.340.024.340.364.875.890 Genes, vif
-	G05.360.340.024.340.364.875.897 Genes, vpr
-	G05.360.340.024.340.364.875.900 Genes, vpu
-	G05.360.340.024.340.365 Genes, Mitochondrial
-	G05.360.340.024.340.370 Genes, Modifier
-	G05.360.340.024.340.375 Genes, Neoplasm
-	G05.360.340.024.340.375.249 Genes, Tumor Suppressor
-	G05.360.340.024.340.375.249.050 Genes, APC
-	G05.360.340.024.340.375.249.100 Genes, BRCA1
-	G05.360.340.024.340.375.249.105 Genes, BRCA2
-	G05.360.340.024.340.375.249.200 Genes, DCC
-	G05.360.340.024.340.375.249.320 Genes, MCC
-	G05.360.340.024.340.375.249.340 Genes, Neurofibromatosis 1
-	G05.360.340.024.340.375.249.345 Genes, Neurofibromatosis 2
-	G05.360.340.024.340.375.249.375 Genes, p16
-	G05.360.340.024.340.375.249.385 Genes, p53
-	G05.360.340.024.340.375.249.400 Genes, Retinoblastoma
-	G05.360.340.024.340.375.249.420 Genes, Wilms Tumor
-	G05.360.340.024.340.375.500 Oncogenes
-	G05.360.340.024.340.375.500.791 Proto-Oncogenes
-	G05.360.340.024.340.375.500.791.100 Genes, abl
-	G05.360.340.024.340.375.500.791.148 Genes, bcl-1
-	G05.360.340.024.340.375.500.791.150 Genes, bcl-2
-	G05.360.340.024.340.375.500.791.290 Genes, erbA
-	G05.360.340.024.340.375.500.791.295 Genes, erbB
-	G05.360.340.024.340.375.500.791.295.300 Genes, erbB-1

## MeSH Tree Changes for 2017

Type	Tree - heading
-	G05.360.340.024.340.375.500.791.295.305 Genes, erbB-2
-	G05.360.340.024.340.375.500.791.325 Genes, fms
-	G05.360.340.024.340.375.500.791.330 Genes, fos
-	G05.360.340.024.340.375.500.791.365 Genes, jun
-	G05.360.340.024.340.375.500.791.400 Genes, mos
-	G05.360.340.024.340.375.500.791.418 Genes, myb
-	G05.360.340.024.340.375.500.791.420 Genes, myc
-	G05.360.340.024.340.375.500.791.550 Genes, ras
-	G05.360.340.024.340.375.500.791.552 Genes, rel
-	G05.360.340.024.340.375.500.791.560 Genes, sis
-	G05.360.340.024.340.375.500.791.570 Genes, src
-	G05.360.340.024.340.385 Genes, Overlapping
-	G05.360.340.024.340.385.600 Nested Genes
-	G05.360.340.024.340.393 Genes, Plant
-	G05.360.340.024.340.396 Genes, Protozoan
-	G05.360.340.024.340.400 Genes, RAG-1
-	G05.360.340.024.340.415 Genes, Recessive
-	G05.360.340.024.340.415.400 Genes, Tumor Suppressor
-	G05.360.340.024.340.415.400.050 Genes, APC
-	G05.360.340.024.340.415.400.100 Genes, BRCA1
-	G05.360.340.024.340.415.400.105 Genes, BRCA2
-	G05.360.340.024.340.415.400.200 Genes, DCC
-	G05.360.340.024.340.415.400.320 Genes, MCC
-	G05.360.340.024.340.415.400.340 Genes, Neurofibromatosis 1
-	G05.360.340.024.340.415.400.345 Genes, Neurofibromatosis 2
-	G05.360.340.024.340.415.400.375 Genes, p16
-	G05.360.340.024.340.415.400.385 Genes, p53
-	G05.360.340.024.340.415.400.400 Genes, Retinoblastoma
-	G05.360.340.024.340.415.400.420 Genes, Wilms Tumor
-	G05.360.340.024.340.425 Genes, Regulator
-	G05.360.340.024.340.425.412 Genes, araC
-	G05.360.340.024.340.425.413 Genes, nef
-	G05.360.340.024.340.425.416 Genes, pX
-	G05.360.340.024.340.425.418 Genes, rev
-	G05.360.340.024.340.425.420 Genes, Switch
-	G05.360.340.024.340.425.560 Genes, tat

## MeSH Tree Changes for 2017

Type	Tree - heading
-	G05.360.340.024.340.425.575 Genes, vif
-	G05.360.340.024.340.425.578 Genes, vpr
-	G05.360.340.024.340.425.580 Genes, vpu
-	G05.360.340.024.340.435 Genes, Reporter
-	G05.360.340.024.340.450 Genes, sry
-	G05.360.340.024.340.460 Genes, Suppressor
-	G05.360.340.024.340.465 Genes, Synthetic
-	G05.360.340.024.340.475 Genes, T-Cell Receptor
-	G05.360.340.024.340.475.050 Genes, T-Cell Receptor alpha
-	G05.360.340.024.340.475.080 Genes, T-Cell Receptor beta
-	G05.360.340.024.340.475.240 Genes, T-Cell Receptor delta
-	G05.360.340.024.340.475.400 Genes, T-Cell Receptor gamma
-	G05.360.340.024.340.500 Genes, X-Linked
-	G05.360.340.024.340.515 Genes, Y-Linked
-	G05.360.340.024.340.610 Major Histocompatibility Complex
-	G05.360.340.024.340.610.595 Genes, MHC Class I
-	G05.360.340.024.340.610.600 Genes, MHC Class II
-	G05.360.340.024.340.645 Multigene Family
-	G05.360.340.024.340.645.500 Genes, MDR
-	G05.360.340.024.340.645.750 Genes, rRNA
-	G05.360.340.024.340.700 Pseudogenes
-	G05.360.340.024.340.825 Transgenes
-	G05.360.340.024.340.825.500 Genes, Transgenic, Suicide
-	G05.360.340.024.380 Genetic Loci
-	G05.360.340.024.380.124 Achaete-Scute Complex Genome Region
-	G05.360.340.024.380.249 Locus Control Region
-	G05.360.340.024.380.500 Major Histocompatibility Complex
-	G05.360.340.024.380.500.595 Genes, MHC Class I
-	G05.360.340.024.380.500.600 Genes, MHC Class II
-	G05.360.340.024.380.625 Minor Histocompatibility Loci
-	G05.360.340.024.380.750 Minor Lymphocyte Stimulatory Loci
-	G05.360.340.024.380.875 Nucleolus Organizer Region
New Heading	<b>G05.360.340.024.380.906 Pseudoautosomal Regions</b>
-	G05.360.340.024.380.937 Quantitative Trait Loci
-	G05.360.340.024.380.968 t-Complex Genome Region

## MeSH Tree Changes for 2017

Type	Tree - heading
-	G05.360.340.024.420 Insulator Elements
-	G05.360.340.024.425 Interspersed Repetitive Sequences
-	G05.360.340.024.425.200 DNA Transposable Elements
-	G05.360.340.024.425.500 Genomic Islands
-	G05.360.340.024.425.800 Retroelements
-	G05.360.340.024.425.800.175 Endogenous Retroviruses
-	G05.360.340.024.425.800.200 Genes, Intracisternal A-Particle
-	G05.360.340.024.425.800.400 Long Interspersed Nucleotide Elements
-	G05.360.340.024.425.800.800 Short Interspersed Nucleotide Elements
-	G05.360.340.024.425.800.800.050 Alu Elements
-	G05.360.340.024.430 Isochores
-	G05.360.340.024.686 Operon
-	G05.360.340.024.686.545 Lac Operon
-	G05.360.340.024.686.645 Operator Regions, Genetic
-	G05.360.340.024.686.817 rRNA Operon
-	G05.360.340.024.742 Regulon
-	G05.360.340.024.745 Replicon
-	G05.360.340.024.745.725 Replication Origin
-	G05.360.340.024.810 Sequence Tagged Sites
-	G05.360.340.024.815 Silencer Elements, Transcriptional
-	G05.360.340.024.850 Tandem Repeat Sequences
-	G05.360.340.024.850.069 Clustered Regularly Interspaced Short Palindromic Repeats
-	G05.360.340.024.850.140 DNA Repeat Expansion
-	G05.360.340.024.850.150 DNA, Satellite
-	G05.360.340.024.850.500 Microsatellite Repeats
-	G05.360.340.024.850.500.150 Dinucleotide Repeats
-	G05.360.340.024.850.500.850 Trinucleotide Repeats
-	G05.360.340.024.850.500.850.200 Trinucleotide Repeat Expansion
-	G05.360.340.024.850.550 Minisatellite Repeats
-	G05.360.340.037 Genome Size
-	G05.360.340.337 Genome, Helminth
-	G05.360.340.337.500 Genes, Helminth
-	G05.360.340.350 Genome, Human
-	G05.360.340.357 Genome, Insect
-	G05.360.340.357.500 Genes, Insect



## MeSH Tree Changes for 2017

Type	Tree - heading
-	G05.360.340.358 Genome, Microbial
-	G05.360.340.358.024 Genes, Microbial
-	G05.360.340.358.024.124 Genes, Archaeal
-	G05.360.340.358.024.249 Genes, Bacterial
-	G05.360.340.358.024.500 Genes, Fungal
-	G05.360.340.358.024.500.089 Genes, Mating Type, Fungal
-	G05.360.340.358.024.875 Genes, Viral
-	G05.360.340.358.024.875.172 Genes, env
-	G05.360.340.358.024.875.258 Genes, gag
-	G05.360.340.358.024.875.345 Genes, Immediate-Early
-	G05.360.340.358.024.875.360 Genes, Intracisternal A-Particle
-	G05.360.340.358.024.875.600 Genes, nef
-	G05.360.340.358.024.875.667 Genes, pol
-	G05.360.340.358.024.875.735 Genes, pX
-	G05.360.340.358.024.875.775 Genes, rev
-	G05.360.340.358.024.875.850 Genes, tat
-	G05.360.340.358.024.875.890 Genes, vif
-	G05.360.340.358.024.875.897 Genes, vpr
-	G05.360.340.358.024.875.900 Genes, vpu
-	G05.360.340.358.050 Genome, Archaeal
-	G05.360.340.358.050.500 Genes, Archaeal
-	G05.360.340.358.207 Genome, Bacterial
-	G05.360.340.358.207.249 Genes, Bacterial
-	G05.360.340.358.207.500 Operon
-	G05.360.340.358.207.500.545 Lac Operon
-	G05.360.340.358.207.500.645 Operator Regions, Genetic
-	G05.360.340.358.207.500.817 rRNA Operon
-	G05.360.340.358.365 Genome, Fungal
-	G05.360.340.358.365.500 Genes, Fungal
-	G05.360.340.358.365.500.089 Genes, Mating Type, Fungal
-	G05.360.340.358.840 Genome, Viral
-	G05.360.340.358.840.500 Genes, Viral
-	G05.360.340.358.840.500.172 Genes, env
-	G05.360.340.358.840.500.258 Genes, gag
-	G05.360.340.358.840.500.345 Genes, Immediate-Early
-	G05.360.340.358.840.500.360 Genes, Intracisternal A-Particle

## MeSH Tree Changes for 2017

Type	Tree - heading
-	G05.360.340.358.840.500.600 Genes, nef
-	G05.360.340.358.840.500.667 Genes, pol
-	G05.360.340.358.840.500.735 Genes, pX
-	G05.360.340.358.840.500.775 Genes, rev
-	G05.360.340.358.840.500.850 Genes, tat
-	G05.360.340.358.840.500.890 Genes, vif
-	G05.360.340.358.840.500.897 Genes, vpr
-	G05.360.340.358.840.500.900 Genes, vpu
-	G05.360.340.360 Genome, Mitochondrial
-	G05.360.340.365 Genome, Plant
-	G05.360.340.365.500 Genes, Plant
-	G05.360.340.370 Genome, Plastid
-	G05.360.340.370.200 Genome, Chloroplast
-	G05.360.340.397 Genome, Protozoan
-	G05.360.340.397.500 Genes, Protozoan
-	G05.360.340.425 Genomic Library
-	G05.360.340.550 Metagenome
-	G05.360.360 Histone Code
-	G05.360.580 Nucleic Acid Conformation
-	G05.360.580.100 Base Pairing
-	G05.360.580.114 DNA, A-Form
-	G05.360.580.121 DNA, B-Form
-	G05.360.580.128 DNA, C-Form
-	G05.360.580.156 DNA, Circular
-	G05.360.580.156.084 DNA, Catenated
-	G05.360.580.156.250 DNA, Superhelical
-	G05.360.580.212 DNA, Concatenated
-	G05.360.580.325 DNA, Cruciform
-	G05.360.580.437 DNA, Single-Stranded
-	G05.360.580.493 DNA, Z-Form
-	G05.360.580.550 G-Quadruplexes
-	G05.360.580.662 Nucleotide Motifs
-	G05.360.580.718 RNA Folding
-	G05.360.580.775 RNA, Double-Stranded
-	G05.360.600 Plasmids
-	G05.360.600.080 Bacteriocin Plasmids

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	G05.360.600.250	Cosmids
-	G05.360.600.300	F Factor
-	G05.360.600.430	Hemolysin Factors
-	G05.360.600.500	Lactose Factors
-	G05.360.600.550	Plant Tumor-Inducing Plasmids
-	G05.360.600.600	R Factors
-	G05.360.840	Templates, Genetic
-	G05.360.920	Transcriptome
-	G05.365	Genetic Variation
-	G05.365.036	Antibody Diversity
-	G05.365.073	Antigenic Variation
-	G05.365.331	Genetic Heterogeneity
-	G05.365.590	Mutation
-	G05.365.590.029	Allelic Imbalance
-	G05.365.590.029.530	Loss of Heterozygosity
-	G05.365.590.029.530.175	Chromosome Deletion
-	G05.365.590.029.530.587	Haploinsufficiency
-	G05.365.590.060	Base Pair Mismatch
-	G05.365.590.175	Chromosome Aberrations
-	G05.365.590.175.024	Abnormal Karyotype
-	G05.365.590.175.024.500	XYY Karyotype
-	G05.365.590.175.050	Aneuploidy
-	G05.365.590.175.050.500	Monosomy
-	G05.365.590.175.050.500.500	Chromosome Deletion
-	G05.365.590.175.050.625	Tetrasomy
-	G05.365.590.175.050.750	Trisomy
-	G05.365.590.175.125	Chimerism
-	G05.365.590.175.165	Chromosomal Instability
-	G05.365.590.175.165.180	Chromosome Fragility
-	G05.365.590.175.175	Chromosome Breakage
-	G05.365.590.175.183	Chromosome Duplication
-	G05.365.590.175.183.249	Tetrasomy
-	G05.365.590.175.183.500	Trisomy
-	G05.365.590.175.190	Chromosome Inversion
New Heading	<b>G05.365.590.175.310</b>	<b>Chromothripsis</b>

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	G05.365.590.175.430	Isochromosomes
-	G05.365.590.175.570	Micronuclei, Chromosome-Defective
-	G05.365.590.175.595	Mosaicism
-	G05.365.590.175.677	Polyploidy
-	G05.365.590.175.677.249	Tetraploidy
-	G05.365.590.175.677.500	Triploidy
-	G05.365.590.175.760	Ring Chromosomes
-	G05.365.590.175.815	Sex Chromosome Aberrations
-	G05.365.590.175.815.970	XYY Karyotype
-	G05.365.590.175.870	Translocation, Genetic
-	G05.365.590.175.870.680	Philadelphia Chromosome
-	G05.365.590.175.935	Uniparental Disomy
-	G05.365.590.195	Codon, Nonsense
-	G05.365.590.220	DNA Repeat Expansion
-	G05.365.590.220.865	Trinucleotide Repeat Expansion
-	G05.365.590.265	Frameshift Mutation
-	G05.365.590.310	Gene Amplification
-	G05.365.590.320	Gene Duplication
-	G05.365.590.335	Genomic Instability
-	G05.365.590.335.590	Microsatellite Instability
-	G05.365.590.350	Germ-Line Mutation
-	G05.365.590.500	INDEL Mutation
-	G05.365.590.575	Mutagenesis, Insertional
-	G05.365.590.594	Mutation Accumulation
-	G05.365.590.612	Mutation Rate
-	G05.365.590.650	Mutation, Missense
-	G05.365.590.675	Point Mutation
-	G05.365.590.762	Sequence Deletion
-	G05.365.590.762.180	Chromosome Deletion
-	G05.365.590.762.320	Gene Deletion
-	G05.365.590.770	Sequence Inversion
-	G05.365.590.770.500	Chromosome Inversion
-	G05.365.590.803	Silent Mutation
-	G05.365.590.835	Suppression, Genetic
New Heading	<b>G05.365.590.917</b>	<b>Synthetic Lethal Mutations</b>

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	G05.365.795	Polymorphism, Genetic
-	G05.365.795.297	Genomic Structural Variation
-	G05.365.795.297.500	DNA Copy Number Variations
New Heading	<b>G05.365.795.446</b>	<b>Pharmacogenomic Variants</b>
-	G05.365.795.595	Polymorphism, Restriction Fragment Length
-	G05.365.795.598	Polymorphism, Single Nucleotide
-	G05.365.795.600	Polymorphism, Single-Stranded Conformational
-	G05.370	Genomic Instability
-	G05.370.180	Chromosomal Instability
-	G05.370.180.180	Chromosome Fragility
-	G05.370.590	Microsatellite Instability
-	G05.380	Genotype
-	G05.380.350	Gene Dosage
-	G05.380.350.500	Haploinsufficiency
-	G05.380.355	Genetic Predisposition to Disease
-	G05.380.360	Haplotypes
-	G05.380.371	Hemizygote
-	G05.380.383	Heterozygote
-	G05.380.554	Homozygote
New Tree	<b>G05.390</b>	<b>Heredity</b>
-	G05.400	Hybrid Vigor
New Heading	<b>G05.410</b>	<b>Inbreeding Depression</b>
-	G05.420	Inheritance Patterns
-	G05.420.040	Anticipation, Genetic
-	G05.420.275	Extrachromosomal Inheritance
-	G05.420.275.249	Genes, Chloroplast
-	G05.420.275.500	Genes, Mitochondrial
New Heading	<b>G05.420.275.750</b>	<b>Maternal Inheritance</b>
-	G05.420.320	Genes, Dominant
-	G05.420.325	Genes, Recessive
-	G05.420.457	Genes, X-Linked
-	G05.420.523	Genes, Y-Linked
-	G05.420.556	Genetic Pleiotropy

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	G05.420.590	Multifactorial Inheritance
New Heading	<b>G05.420.623</b>	<b>Paternal Inheritance</b>
-	G05.420.655	Penetrance
-	G05.420.720	Quantitative Trait, Heritable
Old Tree	G05.540	Genetic Linkage
Old Tree	G05.540.500	Linkage Disequilibrium
Old Tree	G05.540.750	Lod Score
New Tree	G05.545	Molecular Mimicry
New Tree	G05.558	Mutagenesis
New Tree	G05.558.109	Amino Acid Substitution
New Tree	G05.558.164	Chromosome Duplication
New Tree	G05.558.220	DNA Repeat Expansion
New Tree	G05.558.220.865	Trinucleotide Repeat Expansion
New Tree	G05.558.315	Gene Amplification
New Tree	G05.558.320	Gene Duplication
New Tree	G05.558.370	INDEL Mutation
New Tree	G05.558.550	Mutagenesis, Insertional
New Tree	G05.558.620	Nondisjunction, Genetic
New Tree	G05.558.800	Sequence Deletion
New Tree	G05.558.800.180	Chromosome Deletion
New Tree	G05.558.800.320	Gene Deletion
New Tree	G05.558.805	Sequence Inversion
New Tree	G05.558.805.500	Chromosome Inversion
New Tree	G05.558.810	Somatic Hypermutation, Immunoglobulin

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	<a href="#">G05.558.835</a>	<a href="#">Suppression, Genetic</a>
New Tree	<a href="#">G05.558.860</a>	<a href="#">Translocation, Genetic</a>
New Tree	<a href="#">G05.627</a>	<a href="#">Nucleic Acid Denaturation</a>
-	G05.695	Phenotype
-	G05.695.200	Ecotype
-	G05.695.224	Endophenotypes
-	G05.695.337	Gene-Environment Interaction
-	G05.695.450	Genetic Markers
-	G05.695.550	Genetic Pleiotropy
-	G05.695.650	Penetrance
-	G05.695.825	Serogroup
-	G05.697	Phylogeny
-	G05.700	Ploidies
-	G05.700.131	Aneuploidy
-	G05.700.131.500	Monosomy
-	G05.700.131.500.500	Chromosome Deletion
-	G05.700.131.625	Tetrasomy
-	G05.700.131.750	Trisomy
-	G05.700.264	Diploidy
-	G05.700.456	Haploidy
-	G05.700.740	Polyploidy
-	G05.700.740.249	Tetraploidy
-	G05.700.740.500	Triploidy
New Tree	<a href="#">G05.728</a>	<a href="#">Recombination, Genetic</a>
New Tree	<a href="#">G05.728.200</a>	<a href="#">Conjugation, Genetic</a>
New Tree	<a href="#">G05.728.385</a>	<a href="#">Gene Fusion</a>
New Tree	<a href="#">G05.728.385.500</a>	<a href="#">Oncogene Fusion</a>
New Tree	<a href="#">G05.728.390</a>	<a href="#">Gene Transfer, Horizontal</a>
New Tree	<a href="#">G05.728.615</a>	<a href="#">Homologous Recombination</a>
New	<a href="#">G05.728.615.200</a>	<a href="#">Crossing Over, Genetic</a>

## MeSH Tree Changes for 2017

Type	Tree - heading	
Tree		
New Tree	<a href="#">G05.728.615.475</a>	<a href="#">Gene Conversion</a>
New Tree	<a href="#">G05.728.615.612</a>	<a href="#">Recombinational DNA Repair</a>
New Tree	<a href="#">G05.728.615.750</a>	<a href="#">Sister Chromatid Exchange</a>
New Tree	<a href="#">G05.728.850</a>	<a href="#">Transduction, Genetic</a>
New Tree	<a href="#">G05.728.860</a>	<a href="#">Transfection</a>
New Tree	<a href="#">G05.728.860.500</a>	<a href="#">Transformation, Bacterial</a>
New Tree	<a href="#">G05.728.865</a>	<a href="#">Transformation, Genetic</a>
New Tree	<a href="#">G05.728.865.820</a>	<a href="#">Transformation, Bacterial</a>
New Tree	<a href="#">G05.728.932</a>	<a href="#">V(D)J Recombination</a>
New Tree	<a href="#">G05.755</a>	<a href="#">RNA Cleavage</a>
New Tree	<a href="#">G05.783</a>	<a href="#">Selection, Genetic</a>
-	<a href="#">G05.810</a>	<a href="#">Sequence Homology</a>
-	<a href="#">G05.810.200</a>	<a href="#">Sequence Homology, Amino Acid</a>
-	<a href="#">G05.810.550</a>	<a href="#">Sequence Homology, Nucleic Acid</a>
-	<a href="#">G05.810.550.830</a>	<a href="#">Synteny</a>
New Tree	<a href="#">G05.813</a>	<a href="#">Sex Determination Processes</a>
-	<a href="#">G05.815</a>	<a href="#">Sex Ratio</a>
-	<a href="#">G05.820</a>	<a href="#">Structural Homology, Protein</a>
-	<a href="#">G05.910</a>	<a href="#">Sympatry</a>
New Tree	<a href="#">G05.935</a>	<a href="#">Virus Integration</a>
New Tree	<a href="#">G05.935.500</a>	<a href="#">Lysogeny</a>
-	<a href="#">G06</a>	<a href="#">Microbiological Phenomena</a>
-	<a href="#">G06.099</a>	<a href="#">Bacterial Physiological Phenomena</a>
New Tree	<a href="#">G06.099.050</a>	<a href="#">Bacterial Adhesion</a>
-	<a href="#">G06.099.100</a>	<a href="#">Bacterial Load</a>



## MeSH Tree Changes for 2017

Type	Tree - heading	
Old Tree	G06.099.112	Bacterial Processes
Old Tree	G06.099.112.100	Bacterial Adhesion
Old Tree	G06.099.112.105	Bacterial Translocation
Old Tree	G06.099.112.120	Bacteriolysis
Old Tree	G06.099.112.805	Transformation, Bacterial
Old Tree	G06.099.112.805.500	DNA Transformation Competence
New Tree	G06.099.114	Bacterial Translocation
New Tree	G06.099.115	Bacteriolysis
-	G06.099.225	Drug Resistance, Bacterial
-	G06.099.225.500	beta-Lactam Resistance
-	G06.099.225.500.175	Cephalosporin Resistance
-	G06.099.225.500.600	Penicillin Resistance
-	G06.099.225.500.600.050	Ampicillin Resistance
-	G06.099.225.500.600.525	Methicillin Resistance
-	G06.099.225.750	Chloramphenicol Resistance
-	G06.099.225.812	Drug Resistance, Multiple, Bacterial
-	G06.099.225.875	Kanamycin Resistance
-	G06.099.225.937	Tetracycline Resistance
-	G06.099.225.968	Trimethoprim Resistance
-	G06.099.225.984	Vancomycin Resistance
New Tree	G06.099.850	Transformation, Bacterial
New Tree	G06.099.850.500	DNA Transformation Competence
New Tree	G06.099.925	Viral Tropism
-	G06.120	Biofilms
New Tree	G06.173	Catabolite Repression
-	G06.225	Drug Resistance, Microbial
-	G06.225.347	Drug Resistance, Bacterial
-	G06.225.347.500	beta-Lactam Resistance
-	G06.225.347.500.175	Cephalosporin Resistance
-	G06.225.347.500.600	Penicillin Resistance
-	G06.225.347.500.600.050	Ampicillin Resistance
-	G06.225.347.500.600.525	Methicillin Resistance

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	G06.225.347.750	Chloramphenicol Resistance
-	G06.225.347.812	Drug Resistance, Multiple, Bacterial
-	G06.225.347.875	Kanamycin Resistance
-	G06.225.347.937	Tetracycline Resistance
-	G06.225.347.968	Trimethoprim Resistance
-	G06.225.347.984	Vancomycin Resistance
-	G06.225.383	Drug Resistance, Fungal
-	G06.225.383.500	Drug Resistance, Multiple, Fungal
-	G06.225.420	Drug Resistance, Viral
-	G06.225.420.500	Drug Resistance, Multiple, Viral
-	G06.320	Germ-Free Life
-	G06.320.676	Specific Pathogen-Free Organisms
New Tree	<a href="#">G06.365</a>	<a href="#">Hemadsorption</a>
New Tree	<a href="#">G06.380</a>	<a href="#">Host-Pathogen Interactions</a>
New Tree	<a href="#">G06.380.380</a>	<a href="#">Host Specificity</a>
New Tree	<a href="#">G06.380.400</a>	<a href="#">Immune Evasion</a>
New Tree	<a href="#">G06.550</a>	<a href="#">Microbial Interactions</a>
New Tree	<a href="#">G06.550.050</a>	<a href="#">Antibiosis</a>
New Tree	<a href="#">G06.550.700</a>	<a href="#">Quorum Sensing</a>
New Tree	<a href="#">G06.550.800</a>	<a href="#">Symbiosis</a>
-	G06.580	Microbial Viability
Old Tree	<a href="#">G06.590</a>	<a href="#">Microbiological Processes</a>
Old Tree	<a href="#">G06.590.110</a>	<a href="#">Bacterial Processes</a>
Old Tree	<a href="#">G06.590.110.100</a>	<a href="#">Bacterial Adhesion</a>
Old Tree	<a href="#">G06.590.110.105</a>	<a href="#">Bacterial Translocation</a>
Old Tree	<a href="#">G06.590.110.120</a>	<a href="#">Bacteriolysis</a>
Old Tree	<a href="#">G06.590.110.805</a>	<a href="#">Transformation, Bacterial</a>
Old Tree	<a href="#">G06.590.235</a>	<a href="#">Catabolite Repression</a>
Old Tree	<a href="#">G06.590.360</a>	<a href="#">Hemadsorption</a>
Old Tree	<a href="#">G06.590.470</a>	<a href="#">Host-Pathogen Interactions</a>
Old Tree	<a href="#">G06.590.470.380</a>	<a href="#">Host Specificity</a>

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	G06.590.470.400 Immune Evasion
Old Tree	G06.590.580 Microbial Interactions
Old Tree	G06.590.580.050 Antibiosis
Old Tree	G06.590.580.700 Quorum Sensing
Old Tree	G06.590.580.800 Symbiosis
Old Tree	G06.590.620 Nitrogen Fixation
Old Tree	G06.590.875 Virus Physiological Processes
Old Tree	G06.590.875.040 Antibody-Dependent Enhancement
Old Tree	G06.590.875.210 Cell Transformation, Viral
Old Tree	G06.590.875.380 Hemagglutination, Viral
Old Tree	G06.590.875.710 Viral Interference
Old Tree	G06.590.875.712 Viral Tropism
Old Tree	G06.590.875.715 Virus Attachment
Old Tree	G06.590.875.730 Virus Inactivation
Old Tree	G06.590.875.750 Virus Integration
Old Tree	G06.590.875.750.500 Lysogeny
Old Tree	G06.590.875.755 Virus Internalization
Old Tree	G06.590.875.776 Virus Release
Old Tree	G06.590.875.780 Virus Replication
Old Tree	G06.590.875.780.940 Virus Activation
Old Tree	G06.590.875.780.950 Virus Assembly
New Tree	G06.591 Microbiota
New Tree	G06.591.375 Gastrointestinal Microbiome
New Tree	G06.591.750 Microbial Consortia
New Heading	<b>G06.591.875 Mycobiome</b>
New Tree	G06.625 Nitrogen Fixation
Old Tree	G06.755 Microbiota
Old Tree	G06.755.375 Gastrointestinal Microbiome
Old Tree	G06.755.750 Microbial Consortia
-	G06.920 Virus Physiological Phenomena
New Tree	G06.920.095 Antibody-Dependent Enhancement
New	G06.920.143 Cell Transformation, Viral

## MeSH Tree Changes for 2017

Type	Tree - heading	
Tree		
-	G06.920.190	Cytopathogenic Effect, Viral
-	G06.920.225	Drug Resistance, Viral
-	G06.920.225.500	Drug Resistance, Multiple, Viral
New Tree	<a href="#">G06.920.370</a>	<a href="#">Hemagglutination, Viral</a>
-	G06.920.400	Inclusion Bodies, Viral
New Tree	<a href="#">G06.920.800</a>	<a href="#">Viral Interference</a>
-	G06.920.850	Viral Load
New Tree	<a href="#">G06.920.863</a>	<a href="#">Viral Tropism</a>
New Tree	<a href="#">G06.920.868</a>	<a href="#">Virus Attachment</a>
New Tree	<a href="#">G06.920.872</a>	<a href="#">Virus Inactivation</a>
Old Tree	<a href="#">G06.920.875</a>	<a href="#">Virus Physiological Processes</a>
Old Tree	<a href="#">G06.920.875.040</a>	<a href="#">Antibody-Dependent Enhancement</a>
Old Tree	<a href="#">G06.920.875.210</a>	<a href="#">Cell Transformation, Viral</a>
Old Tree	<a href="#">G06.920.875.380</a>	<a href="#">Hemagglutination, Viral</a>
Old Tree	<a href="#">G06.920.875.710</a>	<a href="#">Viral Interference</a>
Old Tree	<a href="#">G06.920.875.712</a>	<a href="#">Viral Tropism</a>
Old Tree	<a href="#">G06.920.875.715</a>	<a href="#">Virus Attachment</a>
Old Tree	<a href="#">G06.920.875.730</a>	<a href="#">Virus Inactivation</a>
Old Tree	<a href="#">G06.920.875.750</a>	<a href="#">Virus Integration</a>
Old Tree	<a href="#">G06.920.875.750.500</a>	<a href="#">Lysogeny</a>
Old Tree	<a href="#">G06.920.875.755</a>	<a href="#">Virus Internalization</a>
Old Tree	<a href="#">G06.920.875.776</a>	<a href="#">Virus Release</a>
Old Tree	<a href="#">G06.920.875.780</a>	<a href="#">Virus Replication</a>
Old Tree	<a href="#">G06.920.875.780.940</a>	<a href="#">Virus Activation</a>
Old Tree	<a href="#">G06.920.875.780.950</a>	<a href="#">Virus Assembly</a>
Old Tree	<a href="#">G06.920.875.890</a>	<a href="#">Virus Uncoating</a>
New Tree	<a href="#">G06.920.877</a>	<a href="#">Virus Integration</a>
New Tree	<a href="#">G06.920.877.500</a>	<a href="#">Lysogeny</a>
New Tree	<a href="#">G06.920.881</a>	<a href="#">Virus Internalization</a>
-	G06.920.900	Virus Latency

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	G06.920.913	Virus Release
New Tree	G06.920.925	Virus Replication
New Tree	G06.920.925.940	Virus Activation
New Tree	G06.920.925.950	Virus Assembly
New Tree	G06.920.950	Virus Uncoating
-	G06.930	Virulence
-	G07	Physiological Phenomena
New Tree	G07.025	Adaptation, Physiological
New Tree	G07.025.133	Acclimatization
New Tree	G07.025.133.250	Salt-Tolerance
New Heading	<b>G07.025.133.500</b>	<b>Thermotolerance</b>
-	G07.049	Anatomic Variation
New Tree	G07.075	Bacterial Shedding
-	G07.100	Body Constitution
-	G07.100.049	Body Composition
-	G07.100.049.134	Body Fat Distribution
-	G07.100.049.134.500	Adiposity
-	G07.100.100	Body Weights and Measures
-	G07.100.100.125	Body Mass Index
-	G07.100.100.160	Body Size
-	G07.100.100.160.100	Body Height
-	G07.100.100.160.100.500	Crown-Rump Length
-	G07.100.100.160.120	Body Weight
-	G07.100.100.160.120.186	Birth Weight
-	G07.100.100.160.120.300	Fetal Weight
-	G07.100.100.160.120.499	Ideal Body Weight
-	G07.100.100.160.120.699	Overweight
-	G07.100.100.160.120.699.500	Obesity
-	G07.100.100.160.120.699.500.249	Obesity, Abdominal

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	G07.100.100.160.120.699.500.375	Obesity, Metabolically Benign
-	G07.100.100.160.120.699.500.500	Obesity, Morbid
-	G07.100.100.160.120.699.500.750	Pediatric Obesity
-	G07.100.100.160.120.828	Thinness
-	G07.100.100.160.340	Sagittal Abdominal Diameter
-	G07.100.100.160.560	Waist Circumference
-	G07.100.100.231	Body Surface Area
-	G07.100.100.250	Crown-Rump Length
-	G07.100.100.660	Organ Size
-	G07.100.100.803	Skinfold Thickness
-	G07.100.100.960	Waist-Hip Ratio
New Heading	<b>G07.100.138</b>	<b>Breast Density</b>
-	G07.100.175	Body, Physical Appearance
-	G07.100.175	Physical Appearance, Body
-	G07.100.175.500	Skin Pigmentation
-	G07.100.250	Disease Susceptibility
-	G07.100.800	Somatotypes
-	G07.110	Body Temperature
-	G07.110.232	Body Temperature Regulation
-	G07.110.232.693	Sweating
-	G07.110.232.778	Thermogenesis
-	G07.110.232.778.500	Shivering
-	G07.110.232.889	Torpor
-	G07.110.232.889.249	Estivation
-	G07.110.232.889.500	Hibernation
-	G07.110.753	Skin Temperature
-	G07.180	Chronobiology Phenomena
-	G07.180.562	Periodicity
-	G07.180.562.094	Biological Clocks
-	G07.180.562.094.500	Circadian Clocks
-	G07.180.562.190	Circadian Rhythm
Old Tree	<b>G07.180.562.190.025</b>	<b>Activity Cycles</b>
New Heading	<b>G07.180.562.595</b>	<b>Infradian Rhythm</b>
New Heading	<b>G07.180.562.797</b>	<b>Ultradian Rhythm</b>

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	<a href="#">G07.180.562.797.500</a>	<a href="#">Activity Cycles</a>
-	G07.203	Diet, Food, and Nutrition
-	G07.203.100	Beverages
-	G07.203.100.100	Alcoholic Beverages
-	G07.203.100.100.100	Absinthe
-	G07.203.100.100.200	Beer
-	G07.203.100.100.900	Wine
-	G07.203.100.300	Carbonated Beverages
-	G07.203.100.300.500	Carbonated Water
-	G07.203.100.325	Coffee
-	G07.203.100.418	Drinking Water
-	G07.203.100.418.500	Carbonated Water
-	G07.203.100.512	Energy Drinks
-	G07.203.100.606	Fruit and Vegetable Juices
-	G07.203.100.700	Milk
Old Tree	<del>G07.203.100.700.124</del>	<del>Cultured Milk Products</del>
Old Tree	<del>G07.203.100.700.249</del>	<del>Infant Formula</del>
New Heading	<b>G07.203.100.700.250</b>	<b>Buttermilk</b>
New Heading	<b>G07.203.100.700.375</b>	<b>Kefir</b>
New Heading	<b>G07.203.100.700.438</b>	<b>Koumiss</b>
-	G07.203.100.700.500	Milk, Human
-	G07.203.100.700.750	Whey
-	G07.203.100.712	Milk Substitutes
-	G07.203.100.712.249	Infant Formula
-	G07.203.100.712.500	Soy Milk
-	G07.203.100.831	Tea
-	G07.203.100.831.500	Kombucha Tea
-	G07.203.100.950	Teas, Herbal
-	G07.203.100.975	Teas, Medicinal
-	G07.203.300	Food
-	G07.203.300.100	Bread
-	G07.203.300.140	Candy
-	G07.203.300.140.200	Chewing Gum

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Heading	<b>G07.203.300.195</b>	<b>Chocolate</b>
-	G07.203.300.250	Condiments
-	G07.203.300.250.725	Spices
-	G07.203.300.300	Crops, Agricultural
-	G07.203.300.300.100	Animal Feed
-	G07.203.300.300.100.800	Silage
-	G07.203.300.300.550	Edible Grain
-	G07.203.300.300.550.500	Whole Grains
-	G07.203.300.350	Dairy Products
-	G07.203.300.350.100	Butter
-	G07.203.300.350.100.500	Ghee
Old Tree	<b>G07.203.300.350.200</b>	<b>Cheese</b>
New Tree	<b>G07.203.300.350.300</b>	<b>Cultured Milk Products</b>
New Heading	<b>G07.203.300.350.300.222</b>	<b>Buttermilk</b>
New Tree	<b>G07.203.300.350.300.444</b>	<b>Cheese</b>
New Heading	<b>G07.203.300.350.300.666</b>	<b>Kefir</b>
New Heading	<b>G07.203.300.350.300.777</b>	<b>Koumiss</b>
New Tree	<b>G07.203.300.350.300.888</b>	<b>Yogurt</b>
-	G07.203.300.350.400	Ice Cream
-	G07.203.300.350.500	Margarine
-	G07.203.300.350.525	Milk
Old Tree	<b>G07.203.300.350.525.221</b>	<b>Cultured Milk Products</b>
Old Tree	<b>G07.203.300.350.525.221.888</b>	<b>Yogurt</b>
New Heading	<b>G07.203.300.350.525.250</b>	<b>Buttermilk</b>
Old Tree	<b>G07.203.300.350.525.332</b>	<b>Infant Formula</b>
New Heading	<b>G07.203.300.350.525.375</b>	<b>Kefir</b>
New Heading	<b>G07.203.300.350.525.438</b>	<b>Koumiss</b>
-	G07.203.300.350.525.500	Milk, Human
-	G07.203.300.350.525.520	Milk Proteins



## MeSH Tree Changes for 2017

Type	Tree - heading
-	G07.203.300.350.525.520.500 Whey Proteins
-	G07.203.300.350.525.760 Whey
-	G07.203.300.350.525.760.500 Whey Proteins
-	G07.203.300.362 Dietary Carbohydrates
-	G07.203.300.362.325 Dietary Sucrose
-	G07.203.300.362.662 High Fructose Corn Syrup
-	G07.203.300.375 Dietary Fats
-	G07.203.300.375.200 Butter
-	G07.203.300.375.200.500 Ghee
-	G07.203.300.375.400 Dietary Fats, Unsaturated
-	G07.203.300.375.400.250 Corn Oil
-	G07.203.300.375.400.300 Cottonseed Oil
-	G07.203.300.375.400.500 Olive Oil
-	G07.203.300.375.400.700 Safflower Oil
-	G07.203.300.375.400.725 Sesame Oil
-	G07.203.300.375.400.750 Soybean Oil
-	G07.203.300.375.650 Margarine
-	G07.203.300.400 Dietary Fiber
-	G07.203.300.400.500 Prebiotics
-	G07.203.300.428 Dietary Proteins
-	G07.203.300.428.317 Egg Proteins, Dietary
-	G07.203.300.428.626 Milk Proteins
-	G07.203.300.428.626.500 Whey Proteins
-	G07.203.300.428.920 Vegetable Proteins
-	G07.203.300.456 Dietary Supplements
-	G07.203.300.456.249 Prebiotics
-	G07.203.300.456.500 Probiotics
-	G07.203.300.456.716 Synbiotics
-	G07.203.300.456.933 Yeast, Dried
-	G07.203.300.470 Eggs
-	G07.203.300.470.349 Egg Proteins, Dietary
-	G07.203.300.470.700 Egg White
-	G07.203.300.470.800 Egg Yolk
-	G07.203.300.477 Fast Foods
-	G07.203.300.484 Flour
-	G07.203.300.512 Food Additives

## MeSH Tree Changes for 2017

Type	Tree - heading
-	G07.203.300.512.350                      Fat Substitutes
-	G07.203.300.512.400                      Flavoring Agents
-	G07.203.300.512.400.700                      Sweetening Agents
-	G07.203.300.512.400.700.500                      Non-Nutritive Sweeteners
-	G07.203.300.512.400.700.750                      Nutritive Sweeteners
-	G07.203.300.512.400.700.750.250                      Dietary Sucrose
-	G07.203.300.512.400.700.750.500                      High Fructose Corn Syrup
-	G07.203.300.512.700                      Food Preservatives
-	G07.203.300.515                      Food, Fortified
-	G07.203.300.518                      Food, Genetically Modified
-	G07.203.300.519                      Food, Organic
-	G07.203.300.521                      Food, Preserved
-	G07.203.300.521.500                      Frozen Foods
-	G07.203.300.525                      Foods, Specialized
-	G07.203.300.525.350                      Food, Formulated
-	G07.203.300.525.350.500                      Infant Formula
-	G07.203.300.525.500                      Infant Food
-	G07.203.300.525.500.500                      Infant Formula
-	G07.203.300.562                      Fruit
-	G07.203.300.572                      Functional Food
-	G07.203.300.581                      Honey
-	G07.203.300.590                      Meals
-	G07.203.300.590.120                      Breakfast
-	G07.203.300.590.560                      Lunch
-	G07.203.300.590.780                      Snacks
-	G07.203.300.600                      Meat
-	G07.203.300.600.500                      Meat Products
-	G07.203.300.600.750                      Poultry
-	G07.203.300.600.750.500                      Poultry Products
-	G07.203.300.600.813                      Red Meat
-	G07.203.300.600.875                      Seafood
-	G07.203.300.600.875.400                      Fish Products
-	G07.203.300.600.875.400.400                      Fish Flour
-	G07.203.300.600.875.700                      Shellfish
-	G07.203.300.631                      Micronutrients
-	G07.203.300.631.555                      Trace Elements

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	G07.203.300.631.600	Vitamins
New Heading	<b>G07.203.300.631.600.500</b>	<b>Provitamins</b>
-	G07.203.300.662	Molasses
-	G07.203.300.700	Nuts
-	G07.203.300.737	Raw Foods
-	G07.203.300.775	Seeds
-	G07.203.300.775.500	Edible Grain
-	G07.203.300.775.500.500	Whole Grains
-	G07.203.300.850	Vegetables
-	G07.203.300.850.450	Vegetable Products
-	G07.203.300.850.450.500	Soy Foods
-	G07.203.300.850.450.500.500	Soy Milk
-	G07.203.300.850.450.500.750	Soybean Proteins
-	G07.203.300.850.900	Vegetable Proteins
-	G07.203.650	Nutritional Physiological Phenomena
-	G07.203.650.161	Animal Nutritional Physiological Phenomena
New Tree	<a href="#">G07.203.650.170</a>	<a href="#">Appetite Regulation</a>
New Tree	<a href="#">G07.203.650.195</a>	<a href="#">Breast Feeding</a>
-	G07.203.650.220	Child Nutritional Physiological Phenomena
-	G07.203.650.220.060	Adolescent Nutritional Physiological Phenomena
-	G07.203.650.220.500	Infant Nutritional Physiological Phenomena
-	G07.203.650.220.500.250	Bottle Feeding
-	G07.203.650.220.500.500	Breast Feeding
-	G07.203.650.220.500.500.500	Breast Milk Expression
-	G07.203.650.220.500.750	Weaning
-	G07.203.650.240	Diet
New Heading	<b>G07.203.650.240.120</b>	<b>Diet, Carbohydrate Loading</b>
-	G07.203.650.240.240	Diet, Diabetic
-	G07.203.650.240.242	Diet, Atherogenic
-	G07.203.650.240.245	Diet, Carbohydrate-Restricted
-	G07.203.650.240.250	Diet Fads
-	G07.203.650.240.255	Diet, Cariogenic
-	G07.203.650.240.260	Diet, Fat-Restricted

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	G07.203.650.240.265	Diet, Gluten-Free
-	G07.203.650.240.267	Diet, High-Fat
-	G07.203.650.240.270	Diet, Mediterranean
-	G07.203.650.240.275	Diet, Paleolithic
-	G07.203.650.240.280	Diet, Protein-Restricted
-	G07.203.650.240.285	Diet, Reducing
-	G07.203.650.240.290	Diet, Sodium-Restricted
-	G07.203.650.240.300	Diet, Vegetarian
-	G07.203.650.240.300.375	Diet, Macrobiotic
-	G07.203.650.240.300.750	Diet, Vegan
-	G07.203.650.240.310	Diet, Western
-	G07.203.650.240.340	Energy Intake
-	G07.203.650.240.340.150	Caloric Restriction
-	G07.203.650.240.587	Fasting
New Heading	<b>G07.203.650.240.629</b>	<b>Healthy Diet</b>
-	G07.203.650.240.670	Ketogenic Diet
-	G07.203.650.240.835	Portion Size
-	G07.203.650.240.917	Serving Size
New Tree	<b>G07.203.650.250</b>	<b>Digestion</b>
New Tree	<b>G07.203.650.250.800</b>	<b>Salivation</b>
New Tree	<b>G07.203.650.283</b>	<b>Eating</b>
New Tree	<b>G07.203.650.283.249</b>	<b>Drinking</b>
New Tree	<b>G07.203.650.283.500</b>	<b>Mastication</b>
-	G07.203.650.315	Elder Nutritional Physiological Phenomena
-	G07.203.650.353	Feeding Behavior
-	G07.203.650.353.099	Bottle Feeding
-	G07.203.650.353.199	Breast Feeding
-	G07.203.650.353.199.500	Breast Milk Expression
-	G07.203.650.353.299	Carnivory
-	G07.203.650.353.400	Fasting
Old Tree	<b>G07.203.650.353.432</b>	<b>Food Habits</b>
Old Tree	<b>G07.203.650.353.432.500</b>	<b>Meals</b>

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	G07.203.650.353.432.500.100 <span style="float: right;">Breakfast</span>
Old Tree	G07.203.650.353.432.500.550 <span style="float: right;">Lunch</span>
Old Tree	G07.203.650.353.432.500.775 <span style="float: right;">Snacks</span>
-	G07.203.650.353.516 <span style="float: right;">Food Preferences</span>
-	G07.203.650.353.758 <span style="float: right;">Herbivory</span>
New Tree	G07.203.650.372 <span style="float: right;">Gastrointestinal Absorption</span>
New Tree	G07.203.650.372.249 <span style="float: right;">Gastric Absorption</span>
New Tree	G07.203.650.372.500 <span style="float: right;">Intestinal Absorption</span>
New Tree	G07.203.650.372.500.500 <span style="float: right;">Intestinal Reabsorption</span>
New Tree	G07.203.650.372.650 <span style="float: right;">Oral Mucosal Absorption</span>
-	G07.203.650.390 <span style="float: right;">Hunger</span>
-	G07.203.650.390.070 <span style="float: right;">Appetite</span>
-	G07.203.650.390.070.290 <span style="float: right;">Appetite Regulation</span>
-	G07.203.650.566 <span style="float: right;">Maternal Nutritional Physiological Phenomena</span>
-	G07.203.650.566.624 <span style="float: right;">Prenatal Nutritional Physiological Phenomena</span>
Old Tree	G07.203.650.593 <span style="float: right;">Nutrition Processes</span>
Old Tree	G07.203.650.593.080 <span style="float: right;">Appetite Regulation</span>
Old Tree	G07.203.650.593.185 <span style="float: right;">Breast Feeding</span>
Old Tree	G07.203.650.593.220 <span style="float: right;">Digestion</span>
Old Tree	G07.203.650.593.220.800 <span style="float: right;">Salivation</span>
Old Tree	G07.203.650.593.260 <span style="float: right;">Eating</span>
Old Tree	G07.203.650.593.260.249 <span style="float: right;">Drinking</span>
Old Tree	G07.203.650.593.260.500 <span style="float: right;">Mastication</span>
Old Tree	G07.203.650.593.345 <span style="float: right;">Gastrointestinal Absorption</span>
Old Tree	G07.203.650.593.345.249 <span style="float: right;">Gastric Absorption</span>
Old Tree	G07.203.650.593.345.500 <span style="float: right;">Intestinal Absorption</span>
Old Tree	G07.203.650.593.345.500.500 <span style="float: right;">Intestinal Reabsorption</span>
Old Tree	G07.203.650.593.345.650 <span style="float: right;">Oral Mucosal Absorption</span>
Old Tree	G07.203.650.593.762 <span style="float: right;">Weaning</span>
-	G07.203.650.620 <span style="float: right;">Nutritional Requirements</span>
-	G07.203.650.620.500 <span style="float: right;">Recommended Dietary Allowances</span>
-	G07.203.650.650 <span style="float: right;">Nutritional Status</span>

## MeSH Tree Changes for 2017

Type	Tree - heading
-	G07.203.650.660                      Nutritive Value
-	G07.203.650.660.500                      Glycemic Index
-	G07.203.650.660.750                      Glycemic Load
-	G07.203.650.830                      Sports Nutritional Physiological Phenomena
New Tree	<a href="#">G07.203.650.915</a> <a href="#">Weaning</a>
-	G07.225                      Dose-Response Relationship, Radiation
-	G07.225.500                      Lethal Dose 50
-	G07.265                      Electrophysiological Phenomena
-	G07.265.087                      Brain Waves
-	G07.265.087.500                      Alpha Rhythm
-	G07.265.087.750                      Beta Rhythm
-	G07.265.087.875                      Delta Rhythm
-	G07.265.087.906                      Gamma Rhythm
-	G07.265.087.937                      Theta Rhythm
-	G07.265.175                      Chronaxy
New Heading	<b>G07.265.216</b> <b>Cortical Excitability</b>
New Tree	<a href="#">G07.265.216.500</a> <a href="#">Evoked Potentials</a>
New Tree	<a href="#">G07.265.216.500.250</a> <a href="#">Contingent Negative Variation</a>
New Tree	<a href="#">G07.265.216.500.350</a> <a href="#">Event-Related Potentials, P300</a>
New Tree	<a href="#">G07.265.216.500.370</a> <a href="#">Evoked Potentials, Auditory</a>
New Tree	<a href="#">G07.265.216.500.370.223</a> <a href="#">Cochlear Microphonic Potentials</a>
New Tree	<a href="#">G07.265.216.500.370.300</a> <a href="#">Evoked Potentials, Auditory, Brain Stem</a>
New Tree	<a href="#">G07.265.216.500.385</a> <a href="#">Evoked Potentials, Motor</a>
New Tree	<a href="#">G07.265.216.500.385.500</a> <a href="#">Vestibular Evoked Myogenic Potentials</a>
New Tree	<a href="#">G07.265.216.500.400</a> <a href="#">Evoked Potentials, Somatosensory</a>
New Tree	<a href="#">G07.265.216.500.400.500</a> <a href="#">Laser-Evoked Potentials</a>
New Tree	<a href="#">G07.265.216.500.425</a> <a href="#">Evoked Potentials, Visual</a>

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	<a href="#">G07.265.256</a> <a href="#">Electroencephalography Phase Synchronization</a>
New Tree	<a href="#">G07.265.256.249</a> <a href="#">Cortical Synchronization</a>
Old Tree	<a href="#">G07.265.337</a> <a href="#">Electrophysiological Processes</a>
Old Tree	<a href="#">G07.265.337.200</a> <a href="#">Electroencephalography Phase Synchronization</a>
Old Tree	<a href="#">G07.265.337.200.249</a> <a href="#">Cortical Synchronization</a>
Old Tree	<a href="#">G07.265.337.249</a> <a href="#">Galvanic Skin Response</a>
Old Tree	<a href="#">G07.265.337.500</a> <a href="#">Ion Channel Gating</a>
Old Tree	<a href="#">G07.265.337.600</a> <a href="#">Neural Conduction</a>
Old Tree	<a href="#">G07.265.337.600.760</a> <a href="#">Recruitment, Neurophysiological</a>
Old Tree	<a href="#">G07.265.337.600.770</a> <a href="#">Refractory Period, Electrophysiological</a>
Old Tree	<a href="#">G07.265.337.615</a> <a href="#">Neural Inhibition</a>
Old Tree	<a href="#">G07.265.337.615.500</a> <a href="#">Inhibitory Postsynaptic Potentials</a>
Old Tree	<a href="#">G07.265.337.620</a> <a href="#">Neuroimmunomodulation</a>
Old Tree	<a href="#">G07.265.337.900</a> <a href="#">Synaptic Transmission</a>
Old Tree	<a href="#">G07.265.337.900.750</a> <a href="#">Synaptic Potentials</a>
Old Tree	<a href="#">G07.265.337.900.750.199</a> <a href="#">Excitatory Postsynaptic Potentials</a>
Old Tree	<a href="#">G07.265.337.900.750.400</a> <a href="#">Inhibitory Postsynaptic Potentials</a>
Old Tree	<a href="#">G07.265.337.900.750.560</a> <a href="#">Miniature Postsynaptic Potentials</a>
Old Tree	<a href="#">G07.265.337.900.750.780</a> <a href="#">Postsynaptic Potential Summation</a>
Old Tree	<a href="#">G07.265.500</a> <a href="#">Evoked Potentials</a>
Old Tree	<a href="#">G07.265.500.250</a> <a href="#">Contingent Negative Variation</a>
Old Tree	<a href="#">G07.265.500.350</a> <a href="#">Event-Related Potentials, P300</a>
Old Tree	<a href="#">G07.265.500.370</a> <a href="#">Evoked Potentials, Auditory</a>
Old Tree	<a href="#">G07.265.500.370.223</a> <a href="#">Cochlear Microphonic Potentials</a>
Old Tree	<a href="#">G07.265.500.370.300</a> <a href="#">Evoked Potentials, Auditory, Brain Stem</a>
Old Tree	<a href="#">G07.265.500.385</a> <a href="#">Evoked Potentials, Motor</a>
Old Tree	<a href="#">G07.265.500.385.500</a> <a href="#">Vestibular Evoked Myogenic Potentials</a>
Old Tree	<a href="#">G07.265.500.400</a> <a href="#">Evoked Potentials, Somatosensory</a>
Old Tree	<a href="#">G07.265.500.400.500</a> <a href="#">Laser-Evoked Potentials</a>
Old Tree	<a href="#">G07.265.500.425</a> <a href="#">Evoked Potentials, Visual</a>
New Tree	<a href="#">G07.265.563</a> <a href="#">Galvanic Skin Response</a>
New Tree	<a href="#">G07.265.625</a> <a href="#">Ion Channel Gating</a>
New	<a href="#">G07.265.675</a> <a href="#">Membrane Potentials</a>

## MeSH Tree Changes for 2017

Type	Tree - heading	
Tree		
New Tree	<a href="#">G07.265.675.100</a>	Action Potentials
New Tree	<a href="#">G07.265.675.550</a>	Membrane Potential, Mitochondrial
New Tree	<a href="#">G07.265.675.775</a>	Myoelectric Complex, Migrating
New Tree	<a href="#">G07.265.675.887</a>	Synaptic Potentials
New Tree	<a href="#">G07.265.675.887.249</a>	Excitatory Postsynaptic Potentials
New Tree	<a href="#">G07.265.675.887.374</a>	Inhibitory Postsynaptic Potentials
New Tree	<a href="#">G07.265.675.887.500</a>	Miniature Postsynaptic Potentials
New Tree	<a href="#">G07.265.675.887.750</a>	Postsynaptic Potential Summation
Old Tree	<a href="#">G07.265.750</a>	Membrane Potentials
Old Tree	<a href="#">G07.265.750.100</a>	Action Potentials
Old Tree	<a href="#">G07.265.750.550</a>	Membrane Potential, Mitochondrial
Old Tree	<a href="#">G07.265.750.775</a>	Myoelectric Complex, Migrating
Old Tree	<a href="#">G07.265.750.887</a>	Synaptic Potentials
Old Tree	<a href="#">G07.265.750.887.249</a>	Excitatory Postsynaptic Potentials
Old Tree	<a href="#">G07.265.750.887.374</a>	Inhibitory Postsynaptic Potentials
Old Tree	<a href="#">G07.265.750.887.500</a>	Miniature Postsynaptic Potentials
Old Tree	<a href="#">G07.265.750.887.750</a>	Postsynaptic Potential Summation
New Tree	<a href="#">G07.265.753</a>	Neural Conduction
New Tree	<a href="#">G07.265.753.760</a>	Recruitment, Neurophysiological
New Tree	<a href="#">G07.265.753.770</a>	Refractory Period, Electrophysiological
New Tree	<a href="#">G07.265.755</a>	Neural Inhibition
New Tree	<a href="#">G07.265.755.500</a>	Inhibitory Postsynaptic Potentials
New Tree	<a href="#">G07.265.758</a>	Neuroimmunomodulation
-	<a href="#">G07.265.760</a>	Refractory Period, Electrophysiological
New Tree	<a href="#">G07.265.880</a>	Synaptic Transmission



## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	G07.265.880.750	Synaptic Potentials
New Tree	G07.265.880.750.199	Excitatory Postsynaptic Potentials
New Tree	G07.265.880.750.400	Inhibitory Postsynaptic Potentials
New Tree	G07.265.880.750.560	Miniature Postsynaptic Potentials
New Tree	G07.265.880.750.780	Postsynaptic Potential Summation
-	G07.280	Fetal Viability
New Tree	G07.313	Fluid Shifts
New Tree	G07.345	Growth and Development
New Tree	G07.345.124	Aging
New Tree	G07.345.124.260	Cognitive Aging
New Tree	G07.345.124.390	Immunosenescence
New Tree	G07.345.124.519	Longevity
New Tree	G07.345.186	Calcification, Physiologic
New Tree	G07.345.186.710	Tooth Calcification
New Tree	G07.345.217	Diapause, Insect
New Tree	G07.345.249	Growth
New Tree	G07.345.249.314	Body Size
New Tree	G07.345.249.314.100	Body Height
New Tree	G07.345.249.314.120	Body Weight
New Tree	G07.345.249.314.120.186	Birth Weight
New Tree	G07.345.249.314.120.200	Body Weight Changes
New Tree	G07.345.249.314.120.200.926	Weight Gain

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	G07.345.249.314.120.200.963 Weight Loss
New Tree	G07.345.249.314.120.250 Body Weight Maintenance
New Tree	G07.345.249.314.120.300 Fetal Weight
New Tree	G07.345.249.314.120.650 Ideal Body Weight
New Tree	G07.345.249.410 Cell Growth Processes
New Tree	G07.345.249.410.500 Cell Enlargement
New Tree	G07.345.249.410.750 Cell Proliferation
New Tree	G07.345.249.410.750.500 Cell Division
New Tree	G07.345.249.410.750.500.500 Asymmetric Cell Division
New Tree	G07.345.249.410.750.500.625 Cell Self Renewal
New Tree	G07.345.249.410.750.500.750 Telomere Homeostasis
New Tree	G07.345.249.690 Organ Size
New Tree	G07.345.249.845 Tropism
New Tree	G07.345.249.845.400 Gravitropism
New Tree	G07.345.249.845.700 Phototropism
New Tree	G07.345.374 Human Development
New Tree	G07.345.374.500 Adolescent Development
New Tree	G07.345.374.750 Child Development
New Tree	G07.345.500 Morphogenesis
New Tree	G07.345.500.100 Body Patterning
New Tree	G07.345.500.100.250 Embryonic Induction
New Tree	G07.345.500.325 Embryonic and Fetal Development

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	G07.345.500.325.089 Ectogenesis
New Tree	G07.345.500.325.180 Embryonic Development
New Tree	G07.345.500.325.180.500 Cell Lineage
New Tree	G07.345.500.325.180.750 Embryonic Induction
New Tree	G07.345.500.325.180.812 Gastrulation
New Tree	G07.345.500.325.180.875 Neurulation
New Tree	G07.345.500.325.235 Fetal Development
New Tree	G07.345.500.325.235.374 Fetal Movement
New Tree	G07.345.500.325.235.750 Fetal Organ Maturity
New Tree	G07.345.500.325.235.875 Fetal Viability
New Tree	G07.345.500.325.235.937 Fetal Weight
New Tree	G07.345.500.325.235.968 Gestational Age
New Tree	G07.345.500.325.377 Organogenesis
New Tree	G07.345.500.325.377.124 Cementogenesis
New Tree	G07.345.500.325.377.186 Dentinogenesis
New Tree	G07.345.500.325.377.249 Fetal Organ Maturity
New Tree	G07.345.500.325.377.437 Lymphangiogenesis
New Tree	G07.345.500.325.377.625 Musculoskeletal Development
New Tree	G07.345.500.325.377.625.100 Bone Development
New Tree	G07.345.500.325.377.625.100.175 Calcification, Physiologic
New Tree	G07.345.500.325.377.625.100.478 Maxillofacial Development
New Tree	G07.345.500.325.377.625.100.729 Osteogenesis

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	G07.345.500.325.377.625.180 Chondrogenesis
New Tree	G07.345.500.325.377.625.590 Muscle Development
New Tree	G07.345.500.325.377.687 Neurogenesis
New Heading	<b>G07.345.500.325.377.687.750 Neuronal Outgrowth</b>
New Heading	<b>G07.345.500.325.377.687.750.500 Axon Fasciculation</b>
New Heading	<b>G07.345.500.325.377.687.750.750 Axon Guidance</b>
New Tree	G07.345.500.325.377.750 Odontogenesis
New Tree	G07.345.500.325.377.750.190 Amelogenesis
New Tree	G07.345.500.325.377.765 Organogenesis, Plant
New Tree	G07.345.500.325.377.812 Sex Determination Processes
New Tree	G07.345.500.325.377.843 Sex Differentiation
New Tree	G07.345.500.550 Metamorphosis, Biological
New Tree	G07.345.500.550.500 Life Cycle Stages
New Tree	G07.345.500.550.500.500 Larva
New Tree	G07.345.500.550.500.500.150 Cercaria
New Tree	G07.345.500.550.500.500.150.500 Metacercariae
New Tree	G07.345.500.550.500.650 Nymph
New Tree	G07.345.500.550.500.675 Oocysts
New Tree	G07.345.500.550.500.675.800 Sporozoites
New Tree	G07.345.500.550.500.700 Pupa
New Tree	G07.345.500.550.500.800 Schizonts
New Tree	G07.345.500.550.500.800.500 Merozoites

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	G07.345.500.550.500.950 Trophozoites
New Tree	G07.345.500.550.750 Molting
New Tree	G07.345.500.550.875 Parasite Encystment
New Tree	G07.345.625 Plant Development
New Tree	G07.345.625.124 Etiolation
New Tree	G07.345.625.249 Germination
New Tree	G07.345.625.875 Plant Dormancy
New Tree	G07.345.750 Sexual Development
New Tree	G07.345.750.437 Sex Determination Processes
New Tree	G07.345.750.500 Sex Differentiation
New Tree	G07.345.750.750 Sexual Maturation
New Tree	G07.410 Homeostasis
New Tree	G07.410.110 Acid-Base Equilibrium
New Tree	G07.410.421 Body Temperature Regulation
New Tree	G07.410.421.693 Sweating
New Tree	G07.410.421.778 Thermogenesis
New Tree	G07.410.421.778.500 Shivering
New Tree	G07.410.421.889 Torpor
New Tree	G07.410.421.889.249 Estivation
New Tree	G07.410.421.889.500 Hibernation
New Tree	G07.410.732 Feedback, Physiological
New Tree	G07.410.732.500 Feedback, Sensory

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	G07.410.810	Osmoregulation
New Tree	G07.410.810.500	Water-Electrolyte Balance
New Tree	G07.410.810.500.500	Kallikrein-Kinin System
New Tree	G07.410.810.500.750	Water Loss, Insensible
-	G07.540	Longevity
New Tree	G07.568	Movement
New Tree	G07.568.500	Locomotion
New Tree	G07.568.500.180	Cell Movement
New Tree	G07.568.500.180.500	Transendothelial and Transepithelial Migration
New Heading	<b>G07.568.500.590</b>	<b>Taxis Response</b>
New Tree	G07.568.500.590.500	Chemotaxis
New Tree	G07.568.500.590.688	Escape Reaction
New Heading	<b>G07.568.500.590.875</b>	<b>Phototaxis</b>
-	G07.650	Organ Specificity
-	G07.690	Pharmacological and Toxicological Phenomena and Processes
-	G07.690.595	Metabolic Clearance Rate
-	G07.690.725	Pharmacokinetics
-	G07.690.725.015	Absorption
-	G07.690.725.015.500	Absorption, Physiological
-	G07.690.725.015.500.374	Gastrointestinal Absorption
-	G07.690.725.015.500.374.249	Gastric Absorption
-	G07.690.725.015.500.374.500	Intestinal Absorption
-	G07.690.725.015.500.374.500.500	Intestinal Reabsorption
-	G07.690.725.015.500.374.650	Oral Mucosal Absorption
-	G07.690.725.015.500.374.825	Rectal Absorption
-	G07.690.725.015.500.562	Intramuscular Absorption
-	G07.690.725.015.500.656	Ocular Absorption
-	G07.690.725.015.500.667	Peritoneal Absorption

## MeSH Tree Changes for 2017

Type	Tree - heading
-	G07.690.725.015.500.679                      Renal Reabsorption
-	G07.690.725.015.500.703                      Respiratory Tract Absorption
-	G07.690.725.015.500.703.500                      Nasal Absorption
-	G07.690.725.015.500.750                      Skin Absorption
-	G07.690.725.015.500.875                      Subcutaneous Absorption
-	G07.690.725.015.500.937                      Vaginal Absorption
-	G07.690.725.064                      Area Under Curve
-	G07.690.725.129                      Biological Availability
-	G07.690.725.225                      Biotransformation
-	G07.690.725.225.224                      Activation, Metabolic
-	G07.690.725.225.450                      Inactivation, Metabolic
-	G07.690.725.225.450.500                      Metabolic Detoxication, Phase I
-	G07.690.725.225.450.750                      Metabolic Detoxication, Phase II
-	G07.690.725.249                      Chronopharmacokinetics
-	G07.690.725.273                      Cutaneous Elimination
-	G07.690.725.321                      Drug Liberation
-	G07.690.725.417                      Hepatobiliary Elimination
-	G07.690.725.465                      Intestinal Elimination
-	G07.690.725.477                      Lacrimal Elimination
-	G07.690.725.489                      Lacteal Elimination
-	G07.690.725.513                      Metabolic Clearance Rate
-	G07.690.725.561                      Pulmonary Elimination
-	G07.690.725.609                      Renal Elimination
-	G07.690.725.705                      Salivary Elimination
-	G07.690.725.898                      Therapeutic Equivalency
-	G07.690.725.949                      Tissue Distribution
-	G07.690.773                      Pharmacological Phenomena
-	G07.690.773.500                      Cytoprotection
-	G07.690.773.750                      Depression, Chemical
-	G07.690.773.875                      Dose-Response Relationship, Drug
-	G07.690.773.875.500                      Hormesis
-	G07.690.773.875.750                      Lethal Dose 50
-	G07.690.773.937                      Down-Regulation
-	G07.690.773.968                      Drug Interactions
-	G07.690.773.968.154                      Drug Agonism
-	G07.690.773.968.154.500                      Drug Partial Agonism

## MeSH Tree Changes for 2017

Type	Tree - heading
-	G07.690.773.968.310 Drug Antagonism
-	G07.690.773.968.393 Drug Inverse Agonism
-	G07.690.773.968.477 Drug Synergism
-	G07.690.773.968.511 Food-Drug Interactions
-	G07.690.773.968.755 Herb-Drug Interactions
-	G07.690.773.984 Drug Resistance
-	G07.690.773.984.269 Drug Resistance, Microbial
-	G07.690.773.984.269.347 Drug Resistance, Bacterial
-	G07.690.773.984.269.347.500 beta-Lactam Resistance
-	G07.690.773.984.269.347.500.175 Cephalosporin Resistance
-	G07.690.773.984.269.347.500.600 Penicillin Resistance
-	G07.690.773.984.269.347.500.600.050 Ampicillin Resistance
-	G07.690.773.984.269.347.500.600.525 Methicillin Resistance
-	G07.690.773.984.269.347.750 Chloramphenicol Resistance
-	G07.690.773.984.269.347.812 Drug Resistance, Multiple, Bacterial
-	G07.690.773.984.269.347.875 Kanamycin Resistance
-	G07.690.773.984.269.347.937 Tetracycline Resistance
-	G07.690.773.984.269.347.968 Trimethoprim Resistance
-	G07.690.773.984.269.347.984 Vancomycin Resistance
-	G07.690.773.984.269.383 Drug Resistance, Fungal
-	G07.690.773.984.269.383.500 Drug Resistance, Multiple, Fungal
-	G07.690.773.984.269.420 Drug Resistance, Viral
-	G07.690.773.984.269.420.500 Drug Resistance, Multiple, Viral
-	G07.690.773.984.300 Drug Resistance, Multiple
-	G07.690.773.984.300.500 Drug Resistance, Multiple, Bacterial
-	G07.690.773.984.300.625 Drug Resistance, Multiple, Fungal
-	G07.690.773.984.300.750 Drug Resistance, Multiple, Viral
-	G07.690.773.984.395 Drug Resistance, Neoplasm
-	G07.690.773.984.443 Herbicide Resistance
-	G07.690.773.984.491 Insecticide Resistance
-	G07.690.773.984.617 Insulin Resistance
-	G07.690.773.992 Drug Tolerance
-	G07.690.773.992.910 Tachyphylaxis
-	G07.690.773.996 Stimulation, Chemical
-	G07.690.773.997 Structure-Activity Relationship
-	G07.690.773.997.500 Quantitative Structure-Activity Relationship



## MeSH Tree Changes for 2017

Type	Tree - heading
-	G07.690.773.998 Up-Regulation
-	G07.690.915 Toxicokinetics
-	G07.690.936 Toxicological Phenomena
-	G07.690.936.500 Dose-Response Relationship, Drug
-	G07.690.936.500.500 Hormesis
-	G07.690.936.500.750 Lethal Dose 50
-	G07.690.936.563 Inhibitory Concentration 50
-	G07.690.936.625 Maximum Tolerated Dose
-	G07.690.936.750 No-Observed-Adverse-Effect Level
Old Tree	<b>G07.700 Physiological Processes</b>
Old Tree	<b>G07.700.062 Adaptation, Physiological</b>
Old Tree	<b>G07.700.062.133 Acclimatization</b>
Old Tree	<b>G07.700.062.796 Salt-Tolerance</b>
Old Tree	<b>G07.700.085 Bacterial Shedding</b>
Old Tree	<b>G07.700.240 Electrophysiological Processes</b>
Old Tree	<b>G07.700.240.249 Galvanic Skin Response</b>
Old Tree	<b>G07.700.240.500 Ion Channel Gating</b>
Old Tree	<b>G07.700.240.600 Neural Conduction</b>
Old Tree	<b>G07.700.240.600.760 Recruitment, Neurophysiological</b>
Old Tree	<b>G07.700.240.600.770 Refractory Period, Electrophysiological</b>
Old Tree	<b>G07.700.240.615 Neural Inhibition</b>
Old Tree	<b>G07.700.240.615.500 Inhibitory Postsynaptic Potentials</b>
Old Tree	<b>G07.700.240.620 Neuroimmunomodulation</b>
Old Tree	<b>G07.700.240.900 Synaptic Transmission</b>
Old Tree	<b>G07.700.240.900.750 Synaptic Potentials</b>
Old Tree	<b>G07.700.240.900.750.199 Excitatory Postsynaptic Potentials</b>
Old Tree	<b>G07.700.240.900.750.400 Inhibitory Postsynaptic Potentials</b>
Old Tree	<b>G07.700.240.900.750.560 Miniature Postsynaptic Potentials</b>
Old Tree	<b>G07.700.240.900.750.780 Postsynaptic Potential Summation</b>
Old Tree	<b>G07.700.280 Fluid Shifts</b>
Old Tree	<b>G07.700.320 Growth and Development</b>
Old Tree	<b>G07.700.320.124 Aging</b>
Old Tree	<b>G07.700.320.124.260 Cognitive Aging</b>
Old Tree	<b>G07.700.320.124.390 Immunosenescence</b>
Old Tree	<b>G07.700.320.124.519 Longevity</b>
Old Tree	<b>G07.700.320.186 Calcification, Physiologic</b>

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	G07.700.320.186.710 Tooth Calcification
Old Tree	G07.700.320.217 Diapause, Insect
Old Tree	G07.700.320.249 Growth
Old Tree	G07.700.320.249.314 Body Size
Old Tree	G07.700.320.249.314.100 Body Height
Old Tree	G07.700.320.249.314.120 Body Weight
Old Tree	G07.700.320.249.314.120.186 Birth Weight
Old Tree	G07.700.320.249.314.120.200 Body Weight Changes
Old Tree	G07.700.320.249.314.120.200.926 Weight Gain
Old Tree	G07.700.320.249.314.120.200.963 Weight Loss
Old Tree	G07.700.320.249.314.120.250 Body Weight Maintenance
Old Tree	G07.700.320.249.314.120.300 Fetal Weight
Old Tree	G07.700.320.249.314.120.650 Ideal Body Weight
Old Tree	G07.700.320.249.410 Cell Growth Processes
Old Tree	G07.700.320.249.410.500 Cell Enlargement
Old Tree	G07.700.320.249.410.750 Cell Proliferation
Old Tree	G07.700.320.249.410.750.500 Cell Division
Old Tree	G07.700.320.249.410.750.500.500 Asymmetric Cell Division
Old Tree	G07.700.320.249.410.750.500.625 Cell Self Renewal
Old Tree	G07.700.320.249.410.750.500.750 Telomere Homeostasis
Old Tree	G07.700.320.249.690 Organ Size
Old Tree	G07.700.320.249.845 Tropism
Old Tree	G07.700.320.249.845.400 Gravitropism
Old Tree	G07.700.320.249.845.700 Phototropism
Old Tree	G07.700.320.374 Human Development
Old Tree	G07.700.320.374.500 Adolescent Development
Old Tree	G07.700.320.374.750 Child Development
Old Tree	G07.700.320.500 Morphogenesis
Old Tree	G07.700.320.500.100 Body Patterning
Old Tree	G07.700.320.500.100.250 Embryonic Induction
Old Tree	G07.700.320.500.325 Embryonic and Fetal Development
Old Tree	G07.700.320.500.325.089 Ectogenesis
Old Tree	G07.700.320.500.325.180 Embryonic Development
Old Tree	G07.700.320.500.325.180.500 Cell Lineage
Old Tree	G07.700.320.500.325.180.750 Embryonic Induction
Old Tree	G07.700.320.500.325.180.781 Epithelial-Mesenchymal Transition

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	G07.700.320.500.325.180.812      Gastrulation
Old Tree	G07.700.320.500.325.180.875      Neurulation
Old Tree	G07.700.320.500.325.235      Fetal Development
Old Tree	G07.700.320.500.325.235.374      Fetal Movement
Old Tree	G07.700.320.500.325.235.750      Fetal Organ Maturity
Old Tree	G07.700.320.500.325.235.875      Fetal Viability
Old Tree	G07.700.320.500.325.235.937      Fetal Weight
Old Tree	G07.700.320.500.325.235.968      Gestational Age
Old Tree	G07.700.320.500.325.377      Organogenesis
Old Tree	G07.700.320.500.325.377.124      Cementogenesis
Old Tree	G07.700.320.500.325.377.186      Dentinogenesis
Old Tree	G07.700.320.500.325.377.249      Fetal Organ Maturity
Old Tree	G07.700.320.500.325.377.437      Lymphangiogenesis
Old Tree	G07.700.320.500.325.377.625      Musculoskeletal Development
Old Tree	G07.700.320.500.325.377.625.100      Bone Development
Old Tree	G07.700.320.500.325.377.625.100.175      Calcification, Physiologic
Old Tree	G07.700.320.500.325.377.625.100.478      Maxillofacial Development
Old Tree	G07.700.320.500.325.377.625.100.729      Osteogenesis
Old Tree	G07.700.320.500.325.377.625.180      Chondrogenesis
Old Tree	G07.700.320.500.325.377.625.590      Muscle Development
Old Tree	G07.700.320.500.325.377.687      Neurogenesis
Old Tree	G07.700.320.500.325.377.750      Odontogenesis
Old Tree	G07.700.320.500.325.377.750.190      Amelogenesis
Old Tree	G07.700.320.500.325.377.765      Organogenesis, Plant
Old Tree	G07.700.320.500.325.377.812      Sex Determination Processes
Old Tree	G07.700.320.500.325.377.843      Sex Differentiation
Old Tree	G07.700.320.500.550      Metamorphosis, Biological
Old Tree	G07.700.320.500.550.500      Life Cycle Stages
Old Tree	G07.700.320.500.550.500.500      Larva
Old Tree	G07.700.320.500.550.500.500.150      Cercaria
Old Tree	G07.700.320.500.550.500.500.150.500      Metacercariae
Old Tree	G07.700.320.500.550.500.650      Nymph
Old Tree	G07.700.320.500.550.500.675      Oocysts
Old Tree	G07.700.320.500.550.500.675.800      Sporozoites
Old Tree	G07.700.320.500.550.500.700      Pupa
Old Tree	G07.700.320.500.550.500.800      Schizonts

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	G07.700.320.500.550.500.800.500 Merozoites
Old Tree	G07.700.320.500.550.500.950 Trophozoites
Old Tree	G07.700.320.500.550.750 Molting
Old Tree	G07.700.320.500.550.875 Parasite Encystment
Old Tree	G07.700.320.625 Plant Development
Old Tree	G07.700.320.625.124 Etiolation
Old Tree	G07.700.320.625.249 Germination
Old Tree	G07.700.320.625.875 Plant Dormancy
Old Tree	G07.700.320.750 Sexual Development
Old Tree	G07.700.320.750.437 Sex Determination Processes
Old Tree	G07.700.320.750.500 Sex Differentiation
Old Tree	G07.700.320.750.750 Sexual Maturation
Old Tree	G07.700.345 Homeostasis
Old Tree	G07.700.345.110 Acid-Base Equilibrium
Old Tree	G07.700.345.421 Body Temperature Regulation
Old Tree	G07.700.345.421.693 Sweating
Old Tree	G07.700.345.421.778 Thermogenesis
Old Tree	G07.700.345.421.778.500 Shivering
Old Tree	G07.700.345.421.889 Torpor
Old Tree	G07.700.345.421.889.249 Estivation
Old Tree	G07.700.345.421.889.500 Hibernation
Old Tree	G07.700.345.732 Feedback, Physiological
Old Tree	G07.700.345.732.500 Feedback, Sensory
Old Tree	G07.700.345.810 Osmoregulation
Old Tree	G07.700.345.810.500 Water-Electrolyte Balance
Old Tree	G07.700.345.810.500.500 Kallikrein-Kinin System
Old Tree	G07.700.345.810.500.750 Water Loss, Insensible
Old Tree	G07.700.560 Movement
Old Tree	G07.700.560.500 Locomotion
Old Tree	G07.700.560.500.180 Cell Movement
Old Tree	G07.700.560.500.180.500 Transendothelial and Transepithelial Migration
Old Tree	G07.700.620 Nutrition Processes
Old Tree	G07.700.620.080 Appetite Regulation
Old Tree	G07.700.620.185 Breast Feeding
Old Tree	G07.700.620.220 Digestion
Old Tree	G07.700.620.220.800 Salivation

## MeSH Tree Changes for 2017

Type	Tree - heading	
Old Tree	G07.700.620.260	Eating
Old Tree	G07.700.620.260.249	Drinking
Old Tree	G07.700.620.260.500	Mastication
Old Tree	G07.700.620.431	Intestinal Absorption
Old Tree	G07.700.620.762	Weaning
Old Tree	G07.700.760	Radiation Tolerance
Old Tree	G07.700.760.500	Dose-Response Relationship, Radiation
Old Tree	G07.700.830	Stress, Physiological
Old Tree	G07.700.830.249	Cold-Shock Response
Old Tree	G07.700.830.500	Heat-Shock Response
Old Tree	G07.700.830.750	Oxidative Stress
Old Tree	G07.700.850	Thermosensing
Old Tree	G07.700.915	Virus Shedding
New Tree	G07.738	Radiation Tolerance
New Tree	G07.738.500	Dose-Response Relationship, Radiation
New Tree	G07.775	Stress, Physiological
New Tree	G07.775.249	Cold-Shock Response
New Tree	G07.775.500	Heat-Shock Response
New Tree	G07.775.750	Oxidative Stress
New Tree	G07.850	Thermosensing
New Tree	G07.925	Virus Shedding
-	G08	Reproductive and Urinary Physiological Phenomena
-	G08.686	Reproductive Physiological Phenomena
-	G08.686.155	Cell Lineage
-	G08.686.157	Climacteric
-	G08.686.157.249	Andropause
-	G08.686.157.500	Menopause
-	G08.686.157.500.500	Menopause, Premature
-	G08.686.157.500.562	Perimenopause
-	G08.686.157.500.625	Postmenopause
-	G08.686.157.500.812	Premenopause

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	G08.686.160	Clutch Size
-	G08.686.195	Estrous Cycle
-	G08.686.195.249	Anestrus
-	G08.686.195.374	Diestrus
-	G08.686.195.500	Estrus
-	G08.686.195.500.500	Estrus Synchronization
-	G08.686.195.750	Metestrus
-	G08.686.195.875	Proestrus
-	G08.686.210	Fertility
-	G08.686.210.249	Ovarian Reserve
-	G08.686.210.500	Time-to-Pregnancy
-	G08.686.220	Fetal Viability
-	G08.686.280	Fetal Weight
New Tree	<a href="#">G08.686.290</a>	<a href="#">Follicular Atresia</a>
-	G08.686.320	Gestational Age
-	G08.686.340	Gravidity
-	G08.686.520	Labor Presentation
-	G08.686.520.150	Breech Presentation
New Tree	<a href="#">G08.686.523</a>	<a href="#">Lactation</a>
New Tree	<a href="#">G08.686.523.460</a>	<a href="#">Milk Ejection</a>
New Tree	<a href="#">G08.686.525</a>	<a href="#">Lacteal Elimination</a>
-	G08.686.530	Litter Size
-	G08.686.560	Maternal Age
-	G08.686.605	Menstrual Cycle
-	G08.686.605.154	Fertile Period
-	G08.686.605.310	Follicular Phase
-	G08.686.605.410	Luteal Phase
-	G08.686.605.428	Menstruation
New Tree	<a href="#">G08.686.617</a>	<a href="#">Menstruation</a>
New Tree	<a href="#">G08.686.628</a>	<a href="#">Ovarian Reserve</a>
-	G08.686.650	Oviparity
-	G08.686.655	Ovoviviparity

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	G08.686.677	Parity
-	G08.686.700	Paternal Age
-	G08.686.701	Peripartum Period
-	G08.686.702	Postpartum Period
-	G08.686.702.500	Lactation
-	G08.686.702.500.460	Milk Ejection
-	G08.686.705	Pregnancy Rate
-	G08.686.707	Pregnancy Trimesters
-	G08.686.707.408	Pregnancy Trimester, First
-	G08.686.707.490	Pregnancy Trimester, Second
-	G08.686.707.520	Pregnancy Trimester, Third
-	G08.686.710	Pregnancy, Multiple
-	G08.686.710.030	Pregnancy, Quadruplet
-	G08.686.710.062	Pregnancy, Quintuplet
-	G08.686.710.126	Pregnancy, Triplet
-	G08.686.710.253	Pregnancy, Twin
-	G08.686.710.608	Superfetation
-	G08.686.760	Puberty
-	G08.686.760.204	Adrenarche
-	G08.686.760.410	Menarche
New Tree	<a href="#">G08.686.784</a>	<a href="#">Reproduction</a>
New Tree	<a href="#">G08.686.784.041</a>	<a href="#">Coitus</a>
New Tree	<a href="#">G08.686.784.084</a>	<a href="#">Ejaculation</a>
New Tree	<a href="#">G08.686.784.170</a>	<a href="#">Embryonic and Fetal Development</a>
New Tree	<a href="#">G08.686.784.170.104</a>	<a href="#">Embryonic Development</a>
New Tree	<a href="#">G08.686.784.170.104.249</a>	<a href="#">Cell Lineage</a>
New Tree	<a href="#">G08.686.784.170.104.500</a>	<a href="#">Embryo Implantation</a>
New Tree	<a href="#">G08.686.784.170.104.500.100</a>	<a href="#">Embryo Implantation, Delayed</a>
New Tree	<a href="#">G08.686.784.170.104.750</a>	<a href="#">Embryonic Induction</a>
New	<a href="#">G08.686.784.170.104.875</a>	<a href="#">Twinning, Monozygotic</a>

## MeSH Tree Changes for 2017

Type	Tree - heading	
Tree		
New Tree	<a href="#">G08.686.784.170.157</a>	Fetal Development
New Tree	<a href="#">G08.686.784.170.157.374</a>	Fetal Movement
New Tree	<a href="#">G08.686.784.170.157.750</a>	Fetal Organ Maturity
New Tree	<a href="#">G08.686.784.170.450</a>	Organogenesis
New Tree	<a href="#">G08.686.784.170.450.500</a>	Neurogenesis
New Heading	<b><a href="#">G08.686.784.170.450.500.750</a></b>	<b>Neuronal Outgrowth</b>
New Heading	<b><a href="#">G08.686.784.170.450.500.750.500</a></b>	<b>Axon Fasciculation</b>
New Heading	<b><a href="#">G08.686.784.170.450.500.750.750</a></b>	<b>Axon Guidance</b>
New Tree	<a href="#">G08.686.784.277</a>	Fertilization
New Tree	<a href="#">G08.686.784.277.360</a>	Ovum Transport
New Tree	<a href="#">G08.686.784.277.560</a>	Self-Fertilization
New Tree	<a href="#">G08.686.784.277.760</a>	Sperm Capacitation
New Tree	<a href="#">G08.686.784.277.800</a>	Sperm-Ovum Interactions
New Tree	<a href="#">G08.686.784.277.800.100</a>	Acrosome Reaction
New Tree	<a href="#">G08.686.784.277.820</a>	Sperm Transport
New Tree	<a href="#">G08.686.784.310</a>	Gametogenesis
New Tree	<a href="#">G08.686.784.310.249</a>	Gametogenesis, Plant
New Tree	<a href="#">G08.686.784.310.500</a>	Oogenesis
New Tree	<a href="#">G08.686.784.310.500.880</a>	Vitellogenesis
New Tree	<a href="#">G08.686.784.310.760</a>	Spermatogenesis
New Tree	<a href="#">G08.686.784.310.760.700</a>	Sperm Maturation



## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	G08.686.784.363 Insemination
New Tree	G08.686.784.363.492 Insemination, Artificial
New Tree	G08.686.784.363.492.500 Insemination, Artificial, Heterologous
New Tree	G08.686.784.363.492.750 Insemination, Artificial, Homologous
New Tree	G08.686.784.460 Orgasm
New Tree	G08.686.784.480 Oviposition
New Tree	G08.686.784.690 Ovulation
New Tree	G08.686.784.690.080 Anovulation
New Tree	G08.686.784.690.355 Luteinization
New Tree	G08.686.784.690.355.500 Corpus Luteum Maintenance
New Tree	G08.686.784.690.380 Luteolysis
New Tree	G08.686.784.690.493 Ovulation Inhibition
New Tree	G08.686.784.690.768 Superovulation
New Tree	G08.686.784.717 Penile Erection
New Tree	G08.686.784.743 Pollination
New Tree	G08.686.784.769 Pregnancy
New Tree	G08.686.784.769.213 Gravidity
New Tree	G08.686.784.769.326 Labor, Obstetric
New Tree	G08.686.784.769.326.100 Cervical Ripening
New Tree	G08.686.784.769.326.500 Labor Onset
New Tree	G08.686.784.769.326.500.080 Labor Stage, First
New Tree	G08.686.784.769.326.500.090 Labor Stage, Second

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	G08.686.784.769.326.500.110 Labor Stage, Third
New Tree	G08.686.784.769.326.520 Labor Presentation
New Tree	G08.686.784.769.326.520.150 Breech Presentation
New Tree	G08.686.784.769.326.660 Trial of Labor
New Tree	G08.686.784.769.326.700 Uterine Contraction
New Tree	G08.686.784.769.455 Maternal-Fetal Exchange
New Tree	G08.686.784.769.472 Parity
New Tree	G08.686.784.769.490 Parturition
New Tree	G08.686.784.769.490.124 Home Childbirth
New Tree	G08.686.784.769.490.249 Natural Childbirth
New Tree	G08.686.784.769.490.500 Term Birth
New Tree	G08.686.784.769.491 Placentation
New Tree	G08.686.784.769.494 Pregnancy in Adolescence
New Tree	G08.686.784.769.498 Pregnancy, Animal
New Tree	G08.686.784.769.498.300 Litter Size
New Tree	G08.686.784.769.500 Pregnancy, High-Risk
New Tree	G08.686.784.769.520 Pregnancy Maintenance
New Tree	G08.686.784.769.520.500 Corpus Luteum Maintenance
New Tree	G08.686.784.769.525 Pregnancy, Multiple
New Tree	G08.686.784.769.525.608 Superfetation
New Tree	G08.686.784.769.530 Pregnancy Outcome
New Tree	G08.686.784.769.530.249 Live Birth

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	G08.686.784.769.530.500 Stillbirth
New Tree	G08.686.784.769.570 Pregnancy, Unplanned
New Tree	G08.686.784.769.580 Pregnancy, Unwanted
New Tree	G08.686.784.769.600 Prenatal Nutritional Physiological Phenomena
New Tree	G08.686.784.769.887 Pseudopregnancy
New Tree	G08.686.784.830 Reproduction, Asexual
New Tree	G08.686.784.830.500 Parthenogenesis
New Tree	G08.686.784.830.500.500 Apomixis
New Tree	G08.686.784.891 Reproductive Behavior
New Tree	G08.686.784.891.500 Contraception Behavior
Old Tree	G08.686.785 Reproductive Physiological Processes
Old Tree	G08.686.785.280 Follicular Atresia
Old Tree	G08.686.785.420 Lactation
Old Tree	G08.686.785.420.460 Milk Ejection
Old Tree	G08.686.785.490 Lacteal Elimination
Old Tree	G08.686.785.560 Menstruation
Old Tree	G08.686.785.660 Ovarian Reserve
Old Tree	G08.686.785.760 Reproduction
Old Tree	G08.686.785.760.041 Coitus
Old Tree	G08.686.785.760.084 Ejaculation
Old Tree	G08.686.785.760.170 Embryonic and Fetal Development
Old Tree	G08.686.785.760.170.104 Embryonic Development
Old Tree	G08.686.785.760.170.104.249 Cell Lineage
Old Tree	G08.686.785.760.170.104.500 Embryo Implantation
Old Tree	G08.686.785.760.170.104.500.100 Embryo Implantation, Delayed
Old Tree	G08.686.785.760.170.104.750 Embryonic Induction
Old Tree	G08.686.785.760.170.104.812 Epithelial-Mesenchymal Transition
Old Tree	G08.686.785.760.170.104.875 Twinning, Monozygotic
Old Tree	G08.686.785.760.170.157 Fetal Development

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	G08.686.785.760.170.157.374      Fetal Movement
Old Tree	G08.686.785.760.170.157.750      Fetal Organ Maturity
Old Tree	G08.686.785.760.170.450      Organogenesis
Old Tree	G08.686.785.760.170.450.500      Neurogenesis
Old Tree	G08.686.785.760.277      Fertilization
Old Tree	G08.686.785.760.277.360      Ovum Transport
Old Tree	G08.686.785.760.277.560      Self-Fertilization
Old Tree	G08.686.785.760.277.760      Sperm Capacitation
Old Tree	G08.686.785.760.277.800      Sperm-Ovum Interactions
Old Tree	G08.686.785.760.277.800.100      Acrosome Reaction
Old Tree	G08.686.785.760.277.820      Sperm Transport
Old Tree	G08.686.785.760.310      Gametogenesis
Old Tree	G08.686.785.760.310.249      Gametogenesis, Plant
Old Tree	G08.686.785.760.310.500      Oogenesis
Old Tree	G08.686.785.760.310.500.880      Vitellogenesis
Old Tree	G08.686.785.760.310.760      Spermatogenesis
Old Tree	G08.686.785.760.310.760.700      Sperm Maturation
Old Tree	G08.686.785.760.363      Insemination
Old Tree	G08.686.785.760.363.492      Insemination, Artificial
Old Tree	G08.686.785.760.363.492.500      Insemination, Artificial, Heterologous
Old Tree	G08.686.785.760.363.492.750      Insemination, Artificial, Homologous
Old Tree	G08.686.785.760.460      Orgasm
Old Tree	G08.686.785.760.480      Oviposition
Old Tree	G08.686.785.760.690      Ovulation
Old Tree	G08.686.785.760.690.080      Anovulation
Old Tree	G08.686.785.760.690.355      Luteinization
Old Tree	G08.686.785.760.690.355.500      Corpus Luteum Maintenance
Old Tree	G08.686.785.760.690.380      Luteolysis
Old Tree	G08.686.785.760.690.493      Ovulation Inhibition
Old Tree	G08.686.785.760.690.768      Superovulation
Old Tree	G08.686.785.760.717      Penile Erection
Old Tree	G08.686.785.760.743      Pollination
Old Tree	G08.686.785.760.769      Pregnancy
Old Tree	G08.686.785.760.769.213      Gravity
Old Tree	G08.686.785.760.769.326      Labor, Obstetric
Old Tree	G08.686.785.760.769.326.100      Cervical Ripening

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	G08.686.785.760.769.326.500 Labor Onset
Old Tree	G08.686.785.760.769.326.500.080 Labor Stage, First
Old Tree	G08.686.785.760.769.326.500.090 Labor Stage, Second
Old Tree	G08.686.785.760.769.326.500.110 Labor Stage, Third
Old Tree	G08.686.785.760.769.326.520 Labor Presentation
Old Tree	G08.686.785.760.769.326.520.150 Breech Presentation
Old Tree	G08.686.785.760.769.326.660 Trial of Labor
Old Tree	G08.686.785.760.769.326.700 Uterine Contraction
Old Tree	G08.686.785.760.769.455 Maternal-Fetal Exchange
Old Tree	G08.686.785.760.769.472 Parity
Old Tree	G08.686.785.760.769.490 Parturition
Old Tree	G08.686.785.760.769.490.124 Home Childbirth
Old Tree	G08.686.785.760.769.490.249 Natural Childbirth
Old Tree	G08.686.785.760.769.490.500 Term Birth
Old Tree	G08.686.785.760.769.491 Placentation
Old Tree	G08.686.785.760.769.494 Pregnancy in Adolescence
Old Tree	G08.686.785.760.769.498 Pregnancy, Animal
Old Tree	G08.686.785.760.769.498.300 Litter Size
Old Tree	G08.686.785.760.769.500 Pregnancy, High-Risk
Old Tree	G08.686.785.760.769.520 Pregnancy Maintenance
Old Tree	G08.686.785.760.769.520.500 Corpus Luteum Maintenance
Old Tree	G08.686.785.760.769.525 Pregnancy, Multiple
Old Tree	G08.686.785.760.769.525.608 Superfetation
Old Tree	G08.686.785.760.769.530 Pregnancy Outcome
Old Tree	G08.686.785.760.769.530.249 Live Birth
Old Tree	G08.686.785.760.769.530.500 Stillbirth
Old Tree	G08.686.785.760.769.570 Pregnancy, Unplanned
Old Tree	G08.686.785.760.769.580 Pregnancy, Unwanted
Old Tree	G08.686.785.760.769.600 Prenatal Nutritional Physiological Phenomena
Old Tree	G08.686.785.760.769.887 Pseudopregnancy
Old Tree	G08.686.785.760.830 Reproduction, Asexual
Old Tree	G08.686.785.760.830.249 Apomixis
Old Tree	G08.686.785.760.830.500 Parthenogenesis
Old Tree	G08.686.785.760.891 Reproductive Behavior
Old Tree	G08.686.785.760.891.500 Contraception Behavior
Old Tree	G08.686.785.880 Sexual Development

## MeSH Tree Changes for 2017

Type	Tree - heading	
Old Tree	G08.686.785.880.249	Climacteric
Old Tree	G08.686.785.880.249.249	Andropause
Old Tree	G08.686.785.880.249.500	Menopause
Old Tree	G08.686.785.880.249.500.500	Menopause, Premature
Old Tree	G08.686.785.880.249.500.562	Perimenopause
Old Tree	G08.686.785.880.249.500.625	Postmenopause
Old Tree	G08.686.785.880.249.500.812	Premenopause
Old Tree	G08.686.785.880.374	Puberty
Old Tree	G08.686.785.880.374.204	Adrenarche
Old Tree	G08.686.785.880.374.410	Menarche
Old Tree	G08.686.785.880.437	Sex Determination Processes
Old Tree	G08.686.785.880.500	Sex Differentiation
Old Tree	G08.686.785.880.750	Sexual Maturation
-	G08.686.810	Sex
-	G08.686.815	Sex Characteristics
New Tree	G08.686.841	Sexual Development
New Tree	G08.686.841.249	Climacteric
New Tree	G08.686.841.249.249	Andropause
New Tree	G08.686.841.249.500	Menopause
New Tree	G08.686.841.249.500.500	Menopause, Premature
New Tree	G08.686.841.249.500.562	Perimenopause
New Tree	G08.686.841.249.500.625	Postmenopause
New Tree	G08.686.841.249.500.812	Premenopause
New Tree	G08.686.841.374	Puberty
New Tree	G08.686.841.374.204	Adrenarche
New Tree	G08.686.841.374.410	Menarche
New Tree	G08.686.841.437	Sex Determination Processes
New	G08.686.841.500	Sex Differentiation

## MeSH Tree Changes for 2017

Type	Tree - heading	
Tree		
New Tree	<a href="#">G08.686.841.750</a>	<a href="#">Sexual Maturation</a>
-	G08.686.867	Sexuality
-	G08.686.867.200	Bisexuality
-	G08.686.867.400	Heterosexuality
-	G08.686.867.500	Homosexuality
-	G08.686.867.500.400	Homosexuality, Female
-	G08.686.867.500.600	Homosexuality, Male
-	G08.686.920	Viviparity, Nonmammalian
-	G08.852	Urinary Tract Physiological Phenomena
New Tree	<a href="#">G08.852.179</a>	<a href="#">Diuresis</a>
New Tree	<a href="#">G08.852.179.557</a>	<a href="#">Natriuresis</a>
-	G08.852.357	Glomerular Filtration Rate
-	G08.852.536	Kidney Concentrating Ability
-	G08.852.536.500	Renal Reabsorption
New Tree	<a href="#">G08.852.725</a>	<a href="#">Renal Circulation</a>
New Tree	<a href="#">G08.852.725.700</a>	<a href="#">Renal Blood Flow, Effective</a>
New Tree	<a href="#">G08.852.725.740</a>	<a href="#">Renal Plasma Flow</a>
New Tree	<a href="#">G08.852.725.750</a>	<a href="#">Renal Plasma Flow, Effective</a>
New Tree	<a href="#">G08.852.761</a>	<a href="#">Renal Elimination</a>
Old Tree	<a href="#">G08.852.796</a>	<a href="#">Urinary Tract Physiological Processes</a>
Old Tree	<a href="#">G08.852.796.241</a>	<a href="#">Diuresis</a>
Old Tree	<a href="#">G08.852.796.241.557</a>	<a href="#">Natriuresis</a>
Old Tree	<a href="#">G08.852.796.700</a>	<a href="#">Renal Circulation</a>
Old Tree	<a href="#">G08.852.796.700.700</a>	<a href="#">Renal Blood Flow, Effective</a>
Old Tree	<a href="#">G08.852.796.700.740</a>	<a href="#">Renal Plasma Flow</a>
Old Tree	<a href="#">G08.852.796.700.750</a>	<a href="#">Renal Plasma Flow, Effective</a>
Old Tree	<a href="#">G08.852.796.788</a>	<a href="#">Renal Elimination</a>
Old Tree	<a href="#">G08.852.796.877</a>	<a href="#">Urination</a>
New Tree	<a href="#">G08.852.880</a>	<a href="#">Urination</a>

## MeSH Tree Changes for 2017

Type	Tree - heading
-	G08.852.898 Urodynamics
-	G09 Circulatory and Respiratory Physiological Phenomena
-	G09.188 Blood Physiological Phenomena
-	G09.188.050 Acid-Base Equilibrium
-	G09.188.087 Bleeding Time
-	G09.188.100 Blood Bactericidal Activity
-	G09.188.105 Blood Cell Count
-	G09.188.105.330 Erythrocyte Count
-	G09.188.105.330.725 Reticulocyte Count
-	G09.188.105.595 Leukocyte Count
-	G09.188.105.595.500 Lymphocyte Count
-	G09.188.105.595.500.150 CD4 Lymphocyte Count
-	G09.188.105.595.500.150.160 CD4-CD8 Ratio
-	G09.188.105.700 Platelet Count
-	G09.188.114 Blood Group Incompatibility
-	G09.188.114.750 Rh Isoimmunization
Old Tree	<b>G09.188.124 Blood Physiological Processes</b>
Old Tree	<b>G09.188.124.454 Erythrocyte Aging</b>
Old Tree	<b>G09.188.124.544 Hematopoiesis</b>
Old Tree	<b>G09.188.124.544.414 Erythropoiesis</b>
Old Tree	<b>G09.188.124.544.463 Hematopoiesis, Extramedullary</b>
Old Tree	<b>G09.188.124.544.597 Leukopoiesis</b>
Old Tree	<b>G09.188.124.544.597.500 Lymphopoiesis</b>
Old Tree	<b>G09.188.124.544.597.750 Myelopoiesis</b>
Old Tree	<b>G09.188.124.544.798 Thrombopoiesis</b>
Old Tree	<b>G09.188.124.552 Hemorheology</b>
Old Tree	<b>G09.188.124.552.249 Erythrocyte Aggregation</b>
Old Tree	<b>G09.188.124.552.624 Platelet Aggregation</b>
Old Tree	<b>G09.188.124.560 Hemostasis</b>
Old Tree	<b>G09.188.124.560.150 Blood Coagulation</b>
Old Tree	<b>G09.188.124.560.150.390 Fibrinolysis</b>
Old Tree	<b>G09.188.124.560.600 Platelet Activation</b>
Old Tree	<b>G09.188.124.560.600.180 Clot Retraction</b>
Old Tree	<b>G09.188.124.560.600.500 Platelet Adhesiveness</b>
Old Tree	<b>G09.188.124.560.600.640 Platelet Aggregation</b>
Old Tree	<b>G09.188.124.780 Phagocytosis</b>



## MeSH Tree Changes for 2017

Type	Tree - heading	
Old Tree	<del>G09.188.124.780.500</del>	<del>Cytophagocytosis</del>
-	G09.188.130	Blood Volume
New Heading	<b>G09.188.130.185</b>	<b>Cerebral Blood Volume</b>
-	G09.188.130.370	Erythrocyte Volume
-	G09.188.130.610	Plasma Volume
New Tree	<a href="#">G09.188.230</a>	<a href="#">Erythrocyte Aging</a>
-	G09.188.260	Erythrocyte Indices
-	G09.188.315	Fibrin Clot Lysis Time
New Tree	<a href="#">G09.188.343</a>	<a href="#">Hematopoiesis</a>
New Tree	<a href="#">G09.188.343.414</a>	<a href="#">Erythropoiesis</a>
New Tree	<a href="#">G09.188.343.463</a>	<a href="#">Hematopoiesis, Extramedullary</a>
New Tree	<a href="#">G09.188.343.597</a>	<a href="#">Leukopoiesis</a>
New Tree	<a href="#">G09.188.343.597.500</a>	<a href="#">Lymphopoiesis</a>
New Tree	<a href="#">G09.188.343.597.750</a>	<a href="#">Myelopoiesis</a>
New Tree	<a href="#">G09.188.343.798</a>	<a href="#">Thrombopoiesis</a>
-	G09.188.370	Hemorheology
-	G09.188.370.124	Blood Viscosity
New Tree	<a href="#">G09.188.370.187</a>	<a href="#">Erythrocyte Aggregation</a>
-	G09.188.370.249	Erythrocyte Deformability
-	G09.188.370.374	Hematocrit
New Tree	<a href="#">G09.188.370.687</a>	<a href="#">Platelet Aggregation</a>
New Tree	<a href="#">G09.188.390</a>	<a href="#">Hemostasis</a>
New Tree	<a href="#">G09.188.390.150</a>	<a href="#">Blood Coagulation</a>
New Tree	<a href="#">G09.188.390.150.390</a>	<a href="#">Fibrinolysis</a>
New Tree	<a href="#">G09.188.390.600</a>	<a href="#">Platelet Activation</a>
New Tree	<a href="#">G09.188.390.600.180</a>	<a href="#">Clot Retraction</a>

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	<a href="#">G09.188.390.600.500</a>	<a href="#">Platelet Adhesiveness</a>
New Tree	<a href="#">G09.188.390.600.640</a>	<a href="#">Platelet Aggregation</a>
-	G09.188.640	Osmotic Fragility
-	G09.188.660	Partial Thromboplastin Time
New Tree	<a href="#">G09.188.665</a>	<a href="#">Phagocytosis</a>
New Tree	<a href="#">G09.188.665.500</a>	<a href="#">Cytophagocytosis</a>
-	G09.188.670	Platelet Adhesiveness
-	G09.188.680	Prothrombin Time
-	G09.188.760	Reticulocytosis
-	G09.188.840	Thrombin Time
-	G09.188.960	Whole Blood Coagulation Time
-	G09.330	Cardiovascular Physiological Phenomena
New Tree	<a href="#">G09.330.040</a>	<a href="#">Atrial Function</a>
New Tree	<a href="#">G09.330.040.100</a>	<a href="#">Atrial Function, Left</a>
New Tree	<a href="#">G09.330.040.200</a>	<a href="#">Atrial Function, Right</a>
New Tree	<a href="#">G09.330.040.600</a>	<a href="#">Atrial Pressure</a>
New Tree	<a href="#">G09.330.040.800</a>	<a href="#">Atrial Remodeling</a>
New Tree	<a href="#">G09.330.100</a>	<a href="#">Blood Circulation</a>
New Tree	<a href="#">G09.330.100.159</a>	<a href="#">Cerebrovascular Circulation</a>
New Tree	<a href="#">G09.330.100.159.500</a>	<a href="#">Neurovascular Coupling</a>
New Tree	<a href="#">G09.330.100.248</a>	<a href="#">Collateral Circulation</a>
New Tree	<a href="#">G09.330.100.324</a>	<a href="#">Coronary Circulation</a>
New Tree	<a href="#">G09.330.100.324.500</a>	<a href="#">Fractional Flow Reserve, Myocardial</a>
New Tree	<a href="#">G09.330.100.645</a>	<a href="#">Microcirculation</a>
New Tree	<a href="#">G09.330.100.749</a>	<a href="#">Placental Circulation</a>

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	<a href="#">G09.330.100.770</a>	<a href="#">Pulmonary Circulation</a>
New Tree	<a href="#">G09.330.100.780</a>	<a href="#">Regional Blood Flow</a>
New Tree	<a href="#">G09.330.100.812</a>	<a href="#">Renal Circulation</a>
New Tree	<a href="#">G09.330.100.812.700</a>	<a href="#">Renal Blood Flow, Effective</a>
New Tree	<a href="#">G09.330.100.812.740</a>	<a href="#">Renal Plasma Flow</a>
New Tree	<a href="#">G09.330.100.812.750</a>	<a href="#">Renal Plasma Flow, Effective</a>
New Tree	<a href="#">G09.330.100.881</a>	<a href="#">Splanchnic Circulation</a>
New Tree	<a href="#">G09.330.100.881.552</a>	<a href="#">Liver Circulation</a>
-	<a href="#">G09.330.160</a>	<a href="#">Capillary Fragility</a>
-	<a href="#">G09.330.165</a>	<a href="#">Capillary Permeability</a>
New Tree	<a href="#">G09.330.170</a>	<a href="#">Cardiomegaly, Exercise-Induced</a>
-	<a href="#">G09.330.175</a>	<a href="#">Cardiovascular Deconditioning</a>
Old Tree	<a href="#">G09.330.190</a>	<a href="#">Cardiovascular Physiological Processes</a>
Old Tree	<a href="#">G09.330.190.075</a>	<a href="#">Atrial Function</a>
Old Tree	<a href="#">G09.330.190.075.100</a>	<a href="#">Atrial Function, Left</a>
Old Tree	<a href="#">G09.330.190.075.200</a>	<a href="#">Atrial Function, Right</a>
Old Tree	<a href="#">G09.330.190.075.600</a>	<a href="#">Atrial Pressure</a>
Old Tree	<a href="#">G09.330.190.075.800</a>	<a href="#">Atrial Remodeling</a>
Old Tree	<a href="#">G09.330.190.163</a>	<a href="#">Blood Circulation</a>
Old Tree	<a href="#">G09.330.190.163.159</a>	<a href="#">Cerebrovascular Circulation</a>
Old Tree	<a href="#">G09.330.190.163.159.500</a>	<a href="#">Neurovascular Coupling</a>
Old Tree	<a href="#">G09.330.190.163.248</a>	<a href="#">Collateral Circulation</a>
Old Tree	<a href="#">G09.330.190.163.324</a>	<a href="#">Coronary Circulation</a>
Old Tree	<a href="#">G09.330.190.163.324.500</a>	<a href="#">Fractional Flow Reserve, Myocardial</a>
Old Tree	<a href="#">G09.330.190.163.645</a>	<a href="#">Microcirculation</a>
Old Tree	<a href="#">G09.330.190.163.749</a>	<a href="#">Placental Circulation</a>
Old Tree	<a href="#">G09.330.190.163.770</a>	<a href="#">Pulmonary Circulation</a>
Old Tree	<a href="#">G09.330.190.163.780</a>	<a href="#">Regional Blood Flow</a>
Old Tree	<a href="#">G09.330.190.163.812</a>	<a href="#">Renal Circulation</a>
Old Tree	<a href="#">G09.330.190.163.812.700</a>	<a href="#">Renal Blood Flow, Effective</a>

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	G09.330.190.163.812.740 Renal Plasma Flow
Old Tree	G09.330.190.163.812.750 Renal Plasma Flow, Effective
Old Tree	G09.330.190.163.881 Splanchnic Circulation
Old Tree	G09.330.190.163.881.552 Liver Circulation
Old Tree	G09.330.190.281 Cardiomegaly, Exercise-Induced
Old Tree	G09.330.190.340 Diving Reflex
Old Tree	G09.330.190.400 Hemodynamics
Old Tree	G09.330.190.400.090 Baroreflex
Old Tree	G09.330.190.400.295 Hemorheology
Old Tree	G09.330.190.400.295.750 Pulsatile Flow
Old Tree	G09.330.190.400.500 Kallikrein-Kinin System
Old Tree	G09.330.190.400.750 Renin-Angiotensin System
Old Tree	G09.330.190.400.920 Vasoconstriction
Old Tree	G09.330.190.400.925 Vasodilation
Old Tree	G09.330.190.541 Myocardial Contraction
Old Tree	G09.330.190.541.295 Diastole
Old Tree	G09.330.190.541.880 Systole
Old Tree	G09.330.190.751 Neovascularization, Physiologic
Old Tree	G09.330.190.962 Ventricular Function
Old Tree	G09.330.190.962.800 Ventricular Function, Left
Old Tree	G09.330.190.962.900 Ventricular Function, Right
Old Tree	G09.330.190.962.950 Ventricular Pressure
Old Tree	G09.330.190.962.975 Ventricular Remodeling
-	G09.330.210 Carotid Intima-Media Thickness
New Tree	G09.330.260 Diving Reflex
-	G09.330.380 Hemodynamics
-	G09.330.380.037 Atrial Pressure
New Tree	G09.330.380.057 Baroreflex
-	G09.330.380.076 Blood Pressure
-	G09.330.380.076.347 Arterial Pressure
-	G09.330.380.076.695 Pulmonary Wedge Pressure
-	G09.330.380.076.732 Venous Pressure
-	G09.330.380.076.732.336 Central Venous Pressure
-	G09.330.380.076.732.650 Portal Pressure

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	G09.330.380.092	Blood Volume
New Heading	<b>G09.330.380.092.185</b>	<b>Cerebral Blood Volume</b>
-	G09.330.380.092.370	Erythrocyte Volume
-	G09.330.380.092.610	Plasma Volume
-	G09.330.380.124	Cardiac Output
-	G09.330.380.124.882	Stroke Volume
-	G09.330.380.249	Cardiac Volume
-	G09.330.380.500	Heart Rate
-	G09.330.380.500.430	Heart Rate, Fetal
-	G09.330.380.500.715	Respiratory Sinus Arrhythmia
-	G09.330.380.510	Heart Sounds
-	G09.330.380.630	Hemorheology
-	G09.330.380.630.080	Blood Flow Velocity
-	G09.330.380.630.110	Blood Viscosity
New Tree	<a href="#">G09.330.380.630.555</a>	<a href="#">Pulsatile Flow</a>
New Tree	<a href="#">G09.330.380.690</a>	<a href="#">Kallikrein-Kinin System</a>
-	G09.330.380.750	Pulse
New Tree	<a href="#">G09.330.380.813</a>	<a href="#">Renin-Angiotensin System</a>
-	G09.330.380.875	Valsalva Maneuver
-	G09.330.380.906	Vascular Capacitance
-	G09.330.380.921	Vascular Resistance
-	G09.330.380.921.327	Capillary Resistance
New Tree	<a href="#">G09.330.380.925</a>	<a href="#">Vasoconstriction</a>
New Tree	<a href="#">G09.330.380.928</a>	<a href="#">Vasodilation</a>
-	G09.330.380.937	Ventricular Pressure
New Tree	<a href="#">G09.330.580</a>	<a href="#">Myocardial Contraction</a>
New Tree	<a href="#">G09.330.580.295</a>	<a href="#">Diastole</a>
New Tree	<a href="#">G09.330.580.880</a>	<a href="#">Systole</a>
New Tree	<a href="#">G09.330.630</a>	<a href="#">Neovascularization, Physiologic</a>

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	G09.330.920	Vascular Patency
-	G09.330.930	Vascular Remodeling
-	G09.330.940	Vascular Stiffness
New Tree	<a href="#">G09.330.955</a>	<a href="#">Ventricular Function</a>
New Tree	<a href="#">G09.330.955.800</a>	<a href="#">Ventricular Function, Left</a>
New Tree	<a href="#">G09.330.955.900</a>	<a href="#">Ventricular Function, Right</a>
New Tree	<a href="#">G09.330.955.950</a>	<a href="#">Ventricular Pressure</a>
New Tree	<a href="#">G09.330.955.975</a>	<a href="#">Ventricular Remodeling</a>
-	G09.772	Respiratory Physiological Phenomena
-	G09.772.029	Airway Remodeling
-	G09.772.060	Airway Resistance
-	G09.772.540	Lung Compliance
New Tree	<a href="#">G09.772.555</a>	<a href="#">Mucociliary Clearance</a>
New Tree	<a href="#">G09.772.585</a>	<a href="#">Phonation</a>
New Tree	<a href="#">G09.772.585.500</a>	<a href="#">Singing</a>
New Tree	<a href="#">G09.772.593</a>	<a href="#">Pulmonary Circulation</a>
-	G09.772.600	Pulmonary Diffusing Capacity
New Tree	<a href="#">G09.772.625</a>	<a href="#">Pulmonary Elimination</a>
-	G09.772.650	Pulmonary Ventilation
-	G09.772.650.300	Forced Expiratory Flow Rates
-	G09.772.650.300.590	Maximal Expiratory Flow Rate
-	G09.772.650.300.630	Maximal Expiratory Flow-Volume Curves
-	G09.772.650.300.670	Maximal Midexpiratory Flow Rate
-	G09.772.650.300.790	Peak Expiratory Flow Rate
-	G09.772.650.430	Forced Expiratory Volume
-	G09.772.650.630	Maximal Voluntary Ventilation
New Tree	<a href="#">G09.772.705</a>	<a href="#">Respiration</a>
New Tree	<a href="#">G09.772.705.349</a>	<a href="#">Breath Holding</a>

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	<a href="#">G09.772.705.700</a>	<a href="#">Respiratory Mechanics</a>
New Tree	<a href="#">G09.772.705.700.080</a>	<a href="#">Bronchoconstriction</a>
New Tree	<a href="#">G09.772.705.700.275</a>	<a href="#">Exhalation</a>
New Tree	<a href="#">G09.772.705.700.390</a>	<a href="#">Inhalation</a>
New Tree	<a href="#">G09.772.705.730</a>	<a href="#">Respiratory Rate</a>
New Tree	<a href="#">G09.772.705.760</a>	<a href="#">Respiratory Transport</a>
New Tree	<a href="#">G09.772.705.760.602</a>	<a href="#">Pulmonary Gas Exchange</a>
-	<a href="#">G09.772.760</a>	<a href="#">Respiratory Dead Space</a>
Old Tree	<a href="#">G09.772.770</a>	<a href="#">Respiratory Physiological Processes</a>
Old Tree	<a href="#">G09.772.770.580</a>	<a href="#">Mucociliary Clearance</a>
Old Tree	<a href="#">G09.772.770.680</a>	<a href="#">Phonation</a>
Old Tree	<a href="#">G09.772.770.680.500</a>	<a href="#">Singing</a>
Old Tree	<a href="#">G09.772.770.690</a>	<a href="#">Pulmonary Circulation</a>
Old Tree	<a href="#">G09.772.770.723</a>	<a href="#">Pulmonary Elimination</a>
Old Tree	<a href="#">G09.772.770.755</a>	<a href="#">Respiration</a>
Old Tree	<a href="#">G09.772.770.755.349</a>	<a href="#">Breath Holding</a>
Old Tree	<a href="#">G09.772.770.755.700</a>	<a href="#">Respiratory Mechanics</a>
Old Tree	<a href="#">G09.772.770.755.700.080</a>	<a href="#">Bronchoconstriction</a>
Old Tree	<a href="#">G09.772.770.755.700.275</a>	<a href="#">Exhalation</a>
Old Tree	<a href="#">G09.772.770.755.700.390</a>	<a href="#">Inhalation</a>
Old Tree	<a href="#">G09.772.770.755.730</a>	<a href="#">Respiratory Rate</a>
Old Tree	<a href="#">G09.772.770.755.760</a>	<a href="#">Respiratory Transport</a>
Old Tree	<a href="#">G09.772.770.755.760.602</a>	<a href="#">Pulmonary Gas Exchange</a>
Old Tree	<a href="#">G09.772.770.787</a>	<a href="#">Respiratory Tract Absorption</a>
Old Tree	<a href="#">G09.772.770.787.500</a>	<a href="#">Nasal Absorption</a>
Old Tree	<a href="#">G09.772.770.820</a>	<a href="#">Sneezing</a>
Old Tree	<a href="#">G09.772.770.980</a>	<a href="#">Yawning</a>
-	<a href="#">G09.772.775</a>	<a href="#">Respiratory Sounds</a>
New Tree	<a href="#">G09.772.813</a>	<a href="#">Respiratory Tract Absorption</a>
New Tree	<a href="#">G09.772.813.500</a>	<a href="#">Nasal Absorption</a>

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	<a href="#">G09.772.832</a>	<a href="#">Sneezing</a>
-	G09.772.850	Total Lung Capacity
-	G09.772.850.250	Closing Volume
-	G09.772.850.390	Functional Residual Capacity
-	G09.772.850.390.360	Expiratory Reserve Volume
-	G09.772.850.390.820	Residual Volume
-	G09.772.850.970	Vital Capacity
-	G09.772.850.970.360	Expiratory Reserve Volume
-	G09.772.850.970.500	Inspiratory Capacity
-	G09.772.850.970.500.375	Inspiratory Reserve Volume
-	G09.772.850.970.500.700	Tidal Volume
-	G09.772.910	Valsalva Maneuver
-	G09.772.920	Ventilation-Perfusion Ratio
-	G09.772.925	Voice
-	G09.772.925.960	Voice Quality
-	G09.772.965	Work of Breathing
New Tree	<a href="#">G09.772.982</a>	<a href="#">Yawning</a>
-	G10	Digestive System and Oral Physiological Phenomena
-	G10.261	Digestive System Physiological Phenomena
New Tree	<a href="#">G10.261.165</a>	<a href="#">Defecation</a>
New Tree	<a href="#">G10.261.178</a>	<a href="#">Deglutition</a>
New Tree	<a href="#">G10.261.190</a>	<a href="#">Digestion</a>
New Tree	<a href="#">G10.261.190.800</a>	<a href="#">Salivation</a>
Old Tree	<a href="#">G10.261.326</a>	<a href="#">Digestive System Processes</a>
Old Tree	<a href="#">G10.261.326.210</a>	<a href="#">Defecation</a>
Old Tree	<a href="#">G10.261.326.215</a>	<a href="#">Deglutition</a>
Old Tree	<a href="#">G10.261.326.222</a>	<a href="#">Digestion</a>
Old Tree	<a href="#">G10.261.326.222.800</a>	<a href="#">Salivation</a>
Old Tree	<a href="#">G10.261.326.240</a>	<a href="#">Eating</a>
Old Tree	<a href="#">G10.261.326.240.249</a>	<a href="#">Drinking</a>
Old Tree	<a href="#">G10.261.326.240.500</a>	<a href="#">Mastication</a>
Old Tree	<a href="#">G10.261.326.270</a>	<a href="#">Gagging</a>



## MeSH Tree Changes for 2017

Type	Tree - heading	
Old Tree	G10.261.326.300	Gallbladder Emptying
Old Tree	G10.261.326.305	Gastrointestinal Absorption
Old Tree	G10.261.326.305.249	Gastric Absorption
Old Tree	G10.261.326.305.500	Intestinal Absorption
Old Tree	G10.261.326.305.500.500	Intestinal Reabsorption
Old Tree	G10.261.326.305.650	Oral Mucosal Absorption
Old Tree	G10.261.326.305.825	Rectal Absorption
Old Tree	G10.261.326.310	Gastrointestinal Motility
Old Tree	G10.261.326.310.400	Gastric Emptying
Old Tree	G10.261.326.310.525	Gastrointestinal Transit
Old Tree	G10.261.326.310.570	Myoelectric Complex, Migrating
Old Tree	G10.261.326.310.596	Peristalsis
Old Tree	G10.261.326.415	Hepatobiliary Elimination
Old Tree	G10.261.326.467	Intestinal Elimination
Old Tree	G10.261.326.520	Liver Regeneration
New Tree	G10.261.330	Eating
New Tree	G10.261.330.249	Drinking
New Tree	G10.261.330.500	Mastication
New Tree	G10.261.338	Gagging
New Tree	G10.261.345	Gallbladder Emptying
New Tree	G10.261.353	Gastrointestinal Absorption
New Tree	G10.261.353.249	Gastric Absorption
New Tree	G10.261.353.500	Intestinal Absorption
New Tree	G10.261.353.500.500	Intestinal Reabsorption
New Tree	G10.261.353.650	Oral Mucosal Absorption
New Tree	G10.261.353.825	Rectal Absorption
New Tree	G10.261.360	Gastrointestinal Motility
New	G10.261.360.400	Gastric Emptying

## MeSH Tree Changes for 2017

Type	Tree - heading	
Tree		
New Tree	<a href="#">G10.261.360.525</a>	<a href="#">Gastrointestinal Transit</a>
New Tree	<a href="#">G10.261.360.570</a>	<a href="#">Myoelectric Complex, Migrating</a>
New Tree	<a href="#">G10.261.360.596</a>	<a href="#">Peristalsis</a>
New Tree	<a href="#">G10.261.375</a>	<a href="#">Hepatobiliary Elimination</a>
-	G10.261.390	Hunger
-	G10.261.390.070	Appetite
-	G10.261.390.070.290	Appetite Regulation
New Tree	<a href="#">G10.261.433</a>	<a href="#">Intestinal Elimination</a>
New Tree	<a href="#">G10.261.475</a>	<a href="#">Liver Regeneration</a>
-	G10.261.700	Postprandial Period
-	G10.549	Dental Physiological Phenomena
-	G10.549.140	Dental Caries Susceptibility
-	G10.549.181	Dental Enamel Solubility
New Tree	<a href="#">G10.549.208</a>	<a href="#">Dental Occlusion</a>
Old Tree	<a href="#">G10.549.235</a>	<a href="#">Dental Physiological Processes</a>
Old Tree	<a href="#">G10.549.235.202</a>	<a href="#">Dental Occlusion</a>
Old Tree	<a href="#">G10.549.235.688</a>	<a href="#">Tooth Calcification</a>
Old Tree	<a href="#">G10.549.235.726</a>	<a href="#">Tooth Eruption</a>
Old Tree	<a href="#">G10.549.235.764</a>	<a href="#">Tooth Exfoliation</a>
Old Tree	<a href="#">G10.549.235.803</a>	<a href="#">Tooth Migration</a>
Old Tree	<a href="#">G10.549.235.803.535</a>	<a href="#">Mesial Movement of Teeth</a>
Old Tree	<a href="#">G10.549.235.917</a>	<a href="#">Tooth Resorption</a>
Old Tree	<a href="#">G10.549.235.917.653</a>	<a href="#">Root Resorption</a>
-	G10.549.245	Dentin Solubility
New Tree	<a href="#">G10.549.800</a>	<a href="#">Tooth Calcification</a>
New Tree	<a href="#">G10.549.810</a>	<a href="#">Tooth Eruption</a>
New Tree	<a href="#">G10.549.820</a>	<a href="#">Tooth Exfoliation</a>
New Tree	<a href="#">G10.549.830</a>	<a href="#">Tooth Migration</a>

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	<a href="#">G10.549.830.535</a> <b>Mesial Movement of Teeth</b>
-	G10.549.840 Tooth Mobility
-	G10.549.845 Tooth Permeability
-	G10.549.845.292 Dental Enamel Permeability
-	G10.549.845.461 Dentin Permeability
New Tree	<a href="#">G10.549.855</a> <b>Tooth Resorption</b>
New Tree	<a href="#">G10.549.855.653</a> <b>Root Resorption</b>
-	G11 Musculoskeletal and Neural Physiological Phenomena
-	G11.427 Musculoskeletal Physiological Phenomena
-	G11.427.100 Bone Density
New Tree	<a href="#">G11.427.213</a> <b>Bone Remodeling</b>
New Tree	<a href="#">G11.427.213.140</a> <b>Bone Regeneration</b>
New Tree	<a href="#">G11.427.213.140.570</a> <b>Osseointegration</b>
New Tree	<a href="#">G11.427.213.150</a> <b>Bone Resorption</b>
New Tree	<a href="#">G11.427.213.150.570</a> <b>Osteolysis</b>
New Tree	<a href="#">G11.427.325</a> <b>Intramuscular Absorption</b>
New Tree	<a href="#">G11.427.410</a> <b>Movement</b>
New Tree	<a href="#">G11.427.410.140</a> <b>Eye Movements</b>
New Tree	<a href="#">G11.427.410.478</a> <b>Head Movements</b>
New Tree	<a href="#">G11.427.410.568</a> <b>Locomotion</b>
New Tree	<a href="#">G11.427.410.568.304</a> <b>Flight, Animal</b>
New Tree	<a href="#">G11.427.410.568.610</a> <b>Running</b>
New Tree	<a href="#">G11.427.410.568.610.320</a> <b>Jogging</b>
New Tree	<a href="#">G11.427.410.568.800</a> <b>Swimming</b>
New	<b><a href="#">G11.427.410.568.850</a> Taxis Response</b>

## MeSH Tree Changes for 2017

Type	Tree - heading	
Heading		
New Tree	<a href="#">G11.427.410.568.850.500</a>	<a href="#">Chemotaxis</a>
New Tree	<a href="#">G11.427.410.568.850.688</a>	<a href="#">Escape Reaction</a>
New Heading	<b><a href="#">G11.427.410.568.850.875</a></b>	<b><a href="#">Phototaxis</a></b>
New Tree	<a href="#">G11.427.410.568.900</a>	<a href="#">Walking</a>
New Tree	<a href="#">G11.427.410.568.900.500</a>	<a href="#">Dependent Ambulation</a>
New Tree	<a href="#">G11.427.410.568.900.750</a>	<a href="#">Gait</a>
New Heading	<b><a href="#">G11.427.410.568.900.750.500</a></b>	<b><a href="#">Walking Speed</a></b>
New Heading	<b><a href="#">G11.427.410.568.900.875</a></b>	<b><a href="#">Stair Climbing</a></b>
New Tree	<a href="#">G11.427.410.698</a>	<a href="#">Motor Activity</a>
New Tree	<a href="#">G11.427.410.698.277</a>	<a href="#">Exercise</a>
New Tree	<a href="#">G11.427.410.698.277.061</a>	<a href="#">Circuit-Based Exercise</a>
New Tree	<a href="#">G11.427.410.698.277.124</a>	<a href="#">Cool-Down Exercise</a>
New Tree	<a href="#">G11.427.410.698.277.156</a>	<a href="#">Gymnastics</a>
New Heading	<b><a href="#">G11.427.410.698.277.187</a></b>	<b><a href="#">High-Intensity Interval Training</a></b>
New Tree	<a href="#">G11.427.410.698.277.249</a>	<a href="#">Muscle Stretching Exercises</a>
New Tree	<a href="#">G11.427.410.698.277.280</a>	<a href="#">Physical Conditioning, Animal</a>
New Tree	<a href="#">G11.427.410.698.277.311</a>	<a href="#">Physical Conditioning, Human</a>
New Tree	<a href="#">G11.427.410.698.277.374</a>	<a href="#">Plyometric Exercise</a>
New Tree	<a href="#">G11.427.410.698.277.500</a>	<a href="#">Resistance Training</a>
New Tree	<a href="#">G11.427.410.698.277.750</a>	<a href="#">Running</a>
New Tree	<a href="#">G11.427.410.698.277.750.320</a>	<a href="#">Jogging</a>

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	G11.427.410.698.277.875	Swimming
New Tree	G11.427.410.698.277.937	Walking
New Heading	<b>G11.427.410.698.277.937.500</b>	<b>Stair Climbing</b>
New Tree	G11.427.410.698.277.968	Warm-Up Exercise
New Tree	G11.427.410.698.416	Flight, Animal
New Tree	G11.427.410.698.555	Freezing Reaction, Cataleptic
New Tree	G11.427.410.698.680	Immobility Response, Tonic
New Tree	G11.427.410.698.840	Pronation
New Tree	G11.427.410.698.920	Supination
New Tree	G11.427.494	Muscle Contraction
New Tree	G11.427.494.235	Excitation Contraction Coupling
New Tree	G11.427.494.472	Isometric Contraction
New Tree	G11.427.494.493	Isotonic Contraction
New Tree	G11.427.494.554	Muscle Relaxation
New Tree	G11.427.494.554.250	Diastole
New Tree	G11.427.494.570	Myocardial Contraction
New Tree	G11.427.494.570.295	Diastole
New Tree	G11.427.494.570.880	Systole
New Tree	G11.427.494.730	Myoelectric Complex, Migrating
New Tree	G11.427.494.890	Uterine Contraction
-	G11.427.550	Muscle Fatigue
-	G11.427.560	Muscle Strength
-	G11.427.560.500	Hand Strength

## MeSH Tree Changes for 2017

Type	Tree - heading
-	G11.427.560.500.500 Pinch Strength
-	G11.427.565 Muscle Tonus
New Tree	<a href="#">G11.427.578</a> <a href="#">Musculoskeletal Development</a>
New Tree	<a href="#">G11.427.578.100</a> <a href="#">Bone Development</a>
New Tree	<a href="#">G11.427.578.100.175</a> <a href="#">Calcification, Physiologic</a>
New Tree	<a href="#">G11.427.578.100.478</a> <a href="#">Maxillofacial Development</a>
New Tree	<a href="#">G11.427.578.100.729</a> <a href="#">Osteogenesis</a>
New Tree	<a href="#">G11.427.578.180</a> <a href="#">Chondrogenesis</a>
New Tree	<a href="#">G11.427.578.590</a> <a href="#">Muscle Development</a>
Old Tree	<a href="#">G11.427.590</a> <a href="#">Musculoskeletal Physiological Processes</a>
Old Tree	<a href="#">G11.427.590.195</a> <a href="#">Bone Remodeling</a>
Old Tree	<a href="#">G11.427.590.195.140</a> <a href="#">Bone Regeneration</a>
Old Tree	<a href="#">G11.427.590.195.140.570</a> <a href="#">Osseointegration</a>
Old Tree	<a href="#">G11.427.590.195.150</a> <a href="#">Bone Resorption</a>
Old Tree	<a href="#">G11.427.590.195.150.570</a> <a href="#">Osteolysis</a>
Old Tree	<a href="#">G11.427.590.362</a> <a href="#">Intramuscular Absorption</a>
Old Tree	<a href="#">G11.427.590.530</a> <a href="#">Movement</a>
Old Tree	<a href="#">G11.427.590.530.140</a> <a href="#">Eye Movements</a>
Old Tree	<a href="#">G11.427.590.530.389</a> <a href="#">Gait</a>
Old Tree	<a href="#">G11.427.590.530.478</a> <a href="#">Head Movements</a>
Old Tree	<a href="#">G11.427.590.530.568</a> <a href="#">Locomotion</a>
Old Tree	<a href="#">G11.427.590.530.568.304</a> <a href="#">Flight, Animal</a>
Old Tree	<a href="#">G11.427.590.530.568.610</a> <a href="#">Running</a>
Old Tree	<a href="#">G11.427.590.530.568.610.320</a> <a href="#">Jogging</a>
Old Tree	<a href="#">G11.427.590.530.568.800</a> <a href="#">Swimming</a>
Old Tree	<a href="#">G11.427.590.530.568.900</a> <a href="#">Walking</a>
Old Tree	<a href="#">G11.427.590.530.568.900.500</a> <a href="#">Dependent Ambulation</a>
Old Tree	<a href="#">G11.427.590.530.698</a> <a href="#">Motor Activity</a>
Old Tree	<a href="#">G11.427.590.530.698.277</a> <a href="#">Exercise</a>
Old Tree	<a href="#">G11.427.590.530.698.277.061</a> <a href="#">Circuit-Based Exercise</a>
Old Tree	<a href="#">G11.427.590.530.698.277.124</a> <a href="#">Cool-Down Exercise</a>

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	G11.427.590.530.698.277.249 Muscle Stretching Exercises
Old Tree	G11.427.590.530.698.277.280 Physical Conditioning, Animal
Old Tree	G11.427.590.530.698.277.311 Physical Conditioning, Human
Old Tree	G11.427.590.530.698.277.374 Plyometric Exercise
Old Tree	G11.427.590.530.698.277.500 Resistance Training
Old Tree	G11.427.590.530.698.277.750 Running
Old Tree	G11.427.590.530.698.277.750.320 Jogging
Old Tree	G11.427.590.530.698.277.875 Swimming
Old Tree	G11.427.590.530.698.277.937 Walking
Old Tree	G11.427.590.530.698.277.968 Warm-Up Exercise
Old Tree	G11.427.590.530.698.416 Flight, Animal
Old Tree	G11.427.590.530.698.555 Freezing Reaction, Cataleptic
Old Tree	G11.427.590.530.698.680 Immobility Response, Tonic
Old Tree	G11.427.590.530.698.840 Pronation
Old Tree	G11.427.590.530.698.920 Supination
Old Tree	G11.427.590.540 Muscle Contraction
Old Tree	G11.427.590.540.235 Excitation Contraction Coupling
Old Tree	G11.427.590.540.472 Isometric Contraction
Old Tree	G11.427.590.540.493 Isotonic Contraction
Old Tree	G11.427.590.540.554 Muscle Relaxation
Old Tree	G11.427.590.540.554.250 Diastole
Old Tree	G11.427.590.540.570 Myocardial Contraction
Old Tree	G11.427.590.540.570.295 Diastole
Old Tree	G11.427.590.540.570.880 Systole
Old Tree	G11.427.590.540.730 Myoelectric Complex, Migrating
Old Tree	G11.427.590.540.890 Uterine Contraction
Old Tree	G11.427.590.560 Musculoskeletal Development
Old Tree	G11.427.590.560.100 Bone Development
Old Tree	G11.427.590.560.100.175 Calcification, Physiologic
Old Tree	G11.427.590.560.100.478 Maxillofacial Development
Old Tree	G11.427.590.560.100.729 Osteogenesis
Old Tree	G11.427.590.560.180 Chondrogenesis
Old Tree	G11.427.590.560.590 Muscle Development
Old Tree	G11.427.590.780 Physical Exertion
-	G11.427.680 Physical Endurance
-	G11.427.680.134 Anaerobic Threshold

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	G11.427.680.270	Exercise Tolerance
New Tree	<a href="#">G11.427.683</a>	<a href="#">Physical Exertion</a>
-	G11.427.685	Physical Fitness
New Heading	<b>G11.427.685.500</b>	<b>Cardiorespiratory Fitness</b>
-	G11.427.690	Postural Balance
-	G11.427.695	Posture
-	G11.427.695.300	Head-Down Tilt
-	G11.427.695.380	Knee-Chest Position
-	G11.427.695.525	Prone Position
-	G11.427.695.625	Supine Position
-	G11.427.700	Psychomotor Performance
-	G11.427.760	Range of Motion, Articular
-	G11.427.760.500	Arthrometry, Articular
-	G11.561	Nervous System Physiological Phenomena
New Tree	<a href="#">G11.561.035</a>	<a href="#">Arousal</a>
New Tree	<a href="#">G11.561.035.738</a>	<a href="#">Wakefulness</a>
New Tree	<a href="#">G11.561.050</a>	<a href="#">Axonal Transport</a>
-	G11.561.127	Brain Waves
-	G11.561.127.500	Alpha Rhythm
-	G11.561.127.750	Beta Rhythm
-	G11.561.127.875	Delta Rhythm
-	G11.561.127.906	Gamma Rhythm
-	G11.561.127.937	Theta Rhythm
-	G11.561.148	Central Nervous System Sensitization
-	G11.561.170	Cerebrospinal Fluid Pressure
-	G11.561.170.505	Intracranial Pressure
-	G11.561.175	Chronaxy
New Heading	<b>G11.561.200</b>	<b>Cortical Excitability</b>
New Tree	<a href="#">G11.561.200.500</a>	<a href="#">Evoked Potentials</a>
New Tree	<a href="#">G11.561.200.500.250</a>	<a href="#">Contingent Negative Variation</a>
New	<a href="#">G11.561.200.500.300</a>	<a href="#">Cortical Spreading Depression</a>



## MeSH Tree Changes for 2017

Type	Tree - heading	
Tree		
New Tree	<a href="#">G11.561.200.500.350</a>	<a href="#">Event-Related Potentials, P300</a>
New Tree	<a href="#">G11.561.200.500.370</a>	<a href="#">Evoked Potentials, Auditory</a>
New Tree	<a href="#">G11.561.200.500.370.223</a>	<a href="#">Cochlear Microphonic Potentials</a>
New Tree	<a href="#">G11.561.200.500.370.300</a>	<a href="#">Evoked Potentials, Auditory, Brain Stem</a>
New Tree	<a href="#">G11.561.200.500.385</a>	<a href="#">Evoked Potentials, Motor</a>
New Tree	<a href="#">G11.561.200.500.400</a>	<a href="#">Evoked Potentials, Somatosensory</a>
New Tree	<a href="#">G11.561.200.500.400.500</a>	<a href="#">Laser-Evoked Potentials</a>
New Tree	<a href="#">G11.561.200.500.425</a>	<a href="#">Evoked Potentials, Visual</a>
-	<a href="#">G11.561.225</a>	<a href="#">Dominance, Cerebral</a>
-	<a href="#">G11.561.225.425</a>	<a href="#">Functional Laterality</a>
-	<a href="#">G11.561.225.425.500</a>	<a href="#">Dominance, Ocular</a>
Old Tree	<a href="#">G11.561.270</a>	<a href="#">Evoked Potentials</a>
Old Tree	<a href="#">G11.561.270.250</a>	<a href="#">Contingent Negative Variation</a>
Old Tree	<a href="#">G11.561.270.300</a>	<a href="#">Cortical Spreading Depression</a>
Old Tree	<a href="#">G11.561.270.350</a>	<a href="#">Event-Related Potentials, P300</a>
Old Tree	<a href="#">G11.561.270.370</a>	<a href="#">Evoked Potentials, Auditory</a>
Old Tree	<a href="#">G11.561.270.370.223</a>	<a href="#">Cochlear Microphonic Potentials</a>
Old Tree	<a href="#">G11.561.270.370.300</a>	<a href="#">Evoked Potentials, Auditory, Brain Stem</a>
Old Tree	<a href="#">G11.561.270.385</a>	<a href="#">Evoked Potentials, Motor</a>
Old Tree	<a href="#">G11.561.270.400</a>	<a href="#">Evoked Potentials, Somatosensory</a>
Old Tree	<a href="#">G11.561.270.400.500</a>	<a href="#">Laser-Evoked Potentials</a>
Old Tree	<a href="#">G11.561.270.425</a>	<a href="#">Evoked Potentials, Visual</a>
New Tree	<a href="#">G11.561.312</a>	<a href="#">Habituation, Psychophysiologic</a>
New Tree	<a href="#">G11.561.398</a>	<a href="#">Higher Nervous Activity</a>
New Tree	<a href="#">G11.561.484</a>	<a href="#">Kindling, Neurologic</a>
-	<a href="#">G11.561.570</a>	<a href="#">Membrane Potentials</a>
-	<a href="#">G11.561.570.100</a>	<a href="#">Action Potentials</a>
-	<a href="#">G11.561.570.837</a>	<a href="#">Myoelectric Complex, Migrating</a>

## MeSH Tree Changes for 2017

Type	Tree - heading
-	G11.561.570.918 Synaptic Potentials
-	G11.561.570.918.249 Excitatory Postsynaptic Potentials
-	G11.561.570.918.374 Inhibitory Postsynaptic Potentials
-	G11.561.570.918.500 Miniature Postsynaptic Potentials
-	G11.561.570.918.750 Postsynaptic Potential Summation
New Tree	<a href="#">G11.561.585</a> Nerve Regeneration
New Tree	<a href="#">G11.561.585.500</a> Spinal Cord Regeneration
Old Tree	<a href="#">G11.561.600</a> Nervous System Physiological Processes
Old Tree	<a href="#">G11.561.600.042</a> Arousal
Old Tree	<a href="#">G11.561.600.042.738</a> Wakefulness
Old Tree	<a href="#">G11.561.600.085</a> Axonal Transport
Old Tree	<a href="#">G11.561.600.227</a> Habituation, Psychophysiologic
Old Tree	<a href="#">G11.561.600.370</a> Higher Nervous Activity
Old Tree	<a href="#">G11.561.600.480</a> Kindling, Neurologic
Old Tree	<a href="#">G11.561.600.615</a> Nerve Regeneration
Old Tree	<a href="#">G11.561.600.615.500</a> Spinal Cord Regeneration
Old Tree	<a href="#">G11.561.600.620</a> Neural Conduction
Old Tree	<a href="#">G11.561.600.620.760</a> Recruitment, Neurophysiological
Old Tree	<a href="#">G11.561.600.620.770</a> Refractory Period, Electrophysiological
Old Tree	<a href="#">G11.561.600.625</a> Neural Inhibition
Old Tree	<a href="#">G11.561.600.625.500</a> Inhibitory Postsynaptic Potentials
Old Tree	<a href="#">G11.561.600.630</a> Neuroimmunomodulation
Old Tree	<a href="#">G11.561.600.635</a> Neuronal Plasticity
Old Tree	<a href="#">G11.561.600.635.350</a> Long-Term Potentiation
Old Tree	<a href="#">G11.561.600.635.355</a> Long-Term Synaptic Depression
Old Tree	<a href="#">G11.561.600.638</a> Neuroprotection
Old Tree	<a href="#">G11.561.600.640</a> Neurosecretion
Old Tree	<a href="#">G11.561.600.765</a> Reflex
Old Tree	<a href="#">G11.561.600.765.063</a> Baroreflex
Old Tree	<a href="#">G11.561.600.765.127</a> Blinking
Old Tree	<a href="#">G11.561.600.765.189</a> Diving Reflex
Old Tree	<a href="#">G11.561.600.765.251</a> Gagging
Old Tree	<a href="#">G11.561.600.765.482</a> Piloerection
Old Tree	<a href="#">G11.561.600.765.580</a> Reflex, Abdominal

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	G11.561.600.765.587 Reflex, Abnormal
Old Tree	G11.561.600.765.594 Reflex, Acoustic
Old Tree	G11.561.600.765.618 Reflex, Babinski
Old Tree	G11.561.600.765.643 Reflex, Monosynaptic
Old Tree	G11.561.600.765.643.474 H-Reflex
Old Tree	G11.561.600.765.685 Reflex, Oculocardiac
Old Tree	G11.561.600.765.705 Reflex, Pupillary
Old Tree	G11.561.600.765.745 Reflex, Righting
Old Tree	G11.561.600.765.768 Reflex, Stretch
Old Tree	G11.561.600.765.781 Reflex, Trigemino-cardiac
Old Tree	G11.561.600.765.795 Reflex, Vestibulo-Ocular
Old Tree	G11.561.600.765.869 Reflex, Startle
Old Tree	G11.561.600.810 Sensation
Old Tree	G11.561.600.810.255 Gravity Sensing
Old Tree	G11.561.600.810.263 Hearing
Old Tree	G11.561.600.810.263.398 Bone Conduction
Old Tree	G11.561.600.810.263.570 Otoacoustic Emissions, Spontaneous
Old Tree	G11.561.600.810.308 Mastodynia
Old Tree	G11.561.600.810.353 Musculoskeletal Pain
Old Tree	G11.561.600.810.353.500 Myalgia
Old Tree	G11.561.600.810.444 Pain
Old Tree	G11.561.600.810.444.350 Arthralgia
Old Tree	G11.561.600.810.444.350.500 Shoulder Pain
Old Tree	G11.561.600.810.444.393 Breakthrough Pain
Old Tree	G11.561.600.810.444.700 Pain Threshold
Old Tree	G11.561.600.810.444.850 Renal Colic
Old Tree	G11.561.600.810.541 Proprioception
Old Tree	G11.561.600.810.541.587 Kinesthesia
Old Tree	G11.561.600.810.541.595 Postural Balance
Old Tree	G11.561.600.810.643 Smell
Old Tree	G11.561.600.810.724 Taste
Old Tree	G11.561.600.810.724.880 Taste Threshold
Old Tree	G11.561.600.810.781 Thermosensing
Old Tree	G11.561.600.810.850 Touch
Old Tree	G11.561.600.810.964 Vision, Ocular
Old Tree	G11.561.600.810.964.124 Color Vision

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	G11.561.600.810.964.186                      Mesopic Vision
Old Tree	G11.561.600.810.964.186.624                      Rod-Cone Interaction
Old Tree	G11.561.600.810.964.249                      Night Vision
Old Tree	G11.561.600.810.964.500                      Phosphenes
Old Tree	G11.561.600.810.964.730                      Vision, Entoptic
Old Tree	G11.561.600.812                      Sensory Gating
Old Tree	G11.561.600.812.500                      Prepulse Inhibition
Old Tree	G11.561.600.815                      Sleep
Old Tree	G11.561.600.815.587                      Sleep Deprivation
Old Tree	G11.561.600.815.754                      Sleep Stages
Old Tree	G11.561.600.815.754.671                      Sleep, REM
Old Tree	G11.561.600.825                      Speech
Old Tree	G11.561.600.825.650                      Speech Acoustics
Old Tree	G11.561.600.825.686                      Speech Intelligibility
Old Tree	G11.561.600.835                      Synaptic Transmission
Old Tree	G11.561.600.835.750                      Synaptic Potentials
Old Tree	G11.561.600.835.750.199                      Excitatory Postsynaptic Potentials
Old Tree	G11.561.600.835.750.400                      Inhibitory Postsynaptic Potentials
Old Tree	G11.561.600.835.750.560                      Miniature Postsynaptic Potentials
Old Tree	G11.561.600.835.750.780                      Postsynaptic Potential Summation
New Tree	G11.561.601                      Neural Conduction
New Tree	G11.561.601.760                      Recruitment, Neurophysiological
New Tree	G11.561.601.770                      Refractory Period, Electrophysiological
New Tree	G11.561.616                      Neural Inhibition
New Tree	G11.561.616.500                      Inhibitory Postsynaptic Potentials
New Tree	G11.561.620                      Neurogenesis
New Heading	<b>G11.561.620.750                      Neuronal Outgrowth</b>
New Heading	<b>G11.561.620.750.250                      Axon Fasciculation</b>
New Heading	<b>G11.561.620.750.500                      Axon Guidance</b>
Old Tree	G11.561.623                      Psychomotor Performance

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	G11.561.630	Neuroimmunomodulation
New Tree	G11.561.638	Neuronal Plasticity
New Tree	G11.561.638.350	Long-Term Potentiation
New Tree	G11.561.638.355	Long-Term Synaptic Depression
New Tree	G11.561.645	Neuroprotection
New Tree	G11.561.653	Neurosecretion
New Tree	G11.561.660	Psychomotor Performance
-	G11.561.677	Reaction Time
New Tree	G11.561.731	Reflex
New Tree	G11.561.731.063	Baroreflex
New Tree	G11.561.731.127	Blinking
New Tree	G11.561.731.189	Diving Reflex
New Tree	G11.561.731.251	Gagging
New Tree	G11.561.731.482	Piloerection
New Tree	G11.561.731.580	Reflex, Abdominal
New Tree	G11.561.731.587	Reflex, Abnormal
New Tree	G11.561.731.594	Reflex, Acoustic
New Tree	G11.561.731.618	Reflex, Babinski
New Tree	G11.561.731.643	Reflex, Monosynaptic
New Tree	G11.561.731.643.474	H-Reflex
New Tree	G11.561.731.685	Reflex, Oculocardiac
New Tree	G11.561.731.705	Reflex, Pupillary

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	G11.561.731.745	Reflex, Righting
New Tree	G11.561.731.768	Reflex, Stretch
New Tree	G11.561.731.781	Reflex, Trigemino-cardiac
New Tree	G11.561.731.795	Reflex, Vestibulo-Ocular
New Tree	G11.561.731.869	Reflex, Startle
-	G11.561.785	Refractory Period, Electrophysiological
New Tree	G11.561.790	Sensation
New Tree	G11.561.790.255	Gravity Sensing
New Tree	G11.561.790.263	Hearing
New Tree	G11.561.790.263.398	Bone Conduction
New Tree	G11.561.790.263.570	Otoacoustic Emissions, Spontaneous
New Tree	G11.561.790.308	Mastodynia
New Tree	G11.561.790.353	Musculoskeletal Pain
New Tree	G11.561.790.353.500	Myalgia
New Tree	G11.561.790.444	Pain
New Tree	G11.561.790.444.350	Arthralgia
New Tree	G11.561.790.444.350.500	Shoulder Pain
New Tree	G11.561.790.444.393	Breakthrough Pain
New Tree	G11.561.790.444.700	Pain Threshold
New Tree	G11.561.790.444.850	Renal Colic
New Tree	G11.561.790.541	Proprioception
New Tree	G11.561.790.541.587	Kinesthesia

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	G11.561.790.541.595 Postural Balance
New Tree	G11.561.790.643 Smell
New Tree	G11.561.790.724 Taste
New Tree	G11.561.790.724.880 Taste Threshold
New Tree	G11.561.790.781 Thermosensing
New Tree	G11.561.790.850 Touch
New Tree	G11.561.790.964 Vision, Ocular
New Tree	G11.561.790.964.124 Color Vision
New Tree	G11.561.790.964.186 Mesopic Vision
New Tree	G11.561.790.964.186.624 Rod-Cone Interaction
New Tree	G11.561.790.964.249 Night Vision
New Tree	G11.561.790.964.500 Phosphenes
New Tree	G11.561.790.964.730 Vision, Entoptic
New Tree	G11.561.794 Sensory Gating
New Tree	G11.561.794.500 Prepulse Inhibition
New Tree	G11.561.803 Sleep
New Tree	G11.561.803.754 Sleep Stages
New Tree	G11.561.803.754.671 Sleep, REM
New Tree	G11.561.812 Speech
New Tree	G11.561.812.650 Speech Acoustics
New Tree	G11.561.812.686 Speech Intelligibility
-	G11.561.820 Speech Acoustics

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	G11.561.830	Synaptic Transmission
New Tree	G11.561.830.750	Synaptic Potentials
New Tree	G11.561.830.750.199	Excitatory Postsynaptic Potentials
New Tree	G11.561.830.750.400	Inhibitory Postsynaptic Potentials
New Tree	G11.561.830.750.560	Miniature Postsynaptic Potentials
New Tree	G11.561.830.750.780	Postsynaptic Potential Summation
-	G11.561.840	Taste Threshold
-	G12	Immune System Phenomena
-	G12.040	Antibody Affinity
New Tree	G12.070	Antibody Formation
New Heading	<b>G12.070.500</b>	<b>Immunogenicity, Vaccine</b>
-	G12.100	Antibody Specificity
New Tree	G12.113	Antibody-Dependent Enhancement
New Tree	G12.119	Antigen Presentation
New Tree	G12.122	Antigen-Antibody Reactions
New Tree	G12.122.100	Agglutination
New Tree	G12.122.100.200	Hemagglutination
New Tree	G12.122.100.700	Sperm Agglutination
New Tree	G12.122.125	Antibody Affinity
New Tree	G12.122.140	Antigenic Modulation
New Tree	G12.122.232	Binding Sites, Antibody
New Tree	G12.122.281	Cross Reactions
New Tree	G12.122.545	Hemolysis
New	G12.122.612	Immunologic Capping



## MeSH Tree Changes for 2017

Type	Tree - heading	
Tree		
New Tree	<a href="#">G12.122.754</a>	<a href="#">Passive Cutaneous Anaphylaxis</a>
New Tree	<a href="#">G12.122.780</a>	<a href="#">Rh Isoimmunization</a>
-	G12.125	Binding Sites, Antibody
-	G12.186	Blood Group Incompatibility
-	G12.186.750	Rh Isoimmunization
-	G12.248	CD4-CD8 Ratio
New Tree	<a href="#">G12.261</a>	<a href="#">Clonal Selection, Antigen-Mediated</a>
New Tree	<a href="#">G12.274</a>	<a href="#">Complement Activation</a>
New Tree	<a href="#">G12.274.695</a>	<a href="#">Complement Pathway, Alternative</a>
New Tree	<a href="#">G12.274.698</a>	<a href="#">Complement Pathway, Classical</a>
New Tree	<a href="#">G12.274.849</a>	<a href="#">Complement Pathway, Mannose-Binding Lectin</a>
New Tree	<a href="#">G12.287</a>	<a href="#">Cytotoxicity, Immunologic</a>
New Tree	<a href="#">G12.287.070</a>	<a href="#">Antibody-Dependent Cell Cytotoxicity</a>
New Tree	<a href="#">G12.287.500</a>	<a href="#">Macrophage Activation</a>
-	G12.300	Dose-Response Relationship, Immunologic
-	G12.300.500	Vaccine Potency
New Tree	<a href="#">G12.350</a>	<a href="#">Histamine Release</a>
-	G12.400	HIV Seronegativity
New Tree	<a href="#">G12.413</a>	<a href="#">Immune Evasion</a>
Old Tree	<a href="#">G12.425</a>	<a href="#">Immune System Processes</a>
Old Tree	<a href="#">G12.425.040</a>	<a href="#">Antibody-Dependent Enhancement</a>
Old Tree	<a href="#">G12.425.071</a>	<a href="#">Antibody Formation</a>
Old Tree	<a href="#">G12.425.107</a>	<a href="#">Antigen Presentation</a>
Old Tree	<a href="#">G12.425.143</a>	<a href="#">Antigen-Antibody Reactions</a>
Old Tree	<a href="#">G12.425.143.100</a>	<a href="#">Agglutination</a>
Old Tree	<a href="#">G12.425.143.100.200</a>	<a href="#">Hemagglutination</a>
Old Tree	<a href="#">G12.425.143.100.700</a>	<a href="#">Sperm Agglutination</a>

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	G12.425.143.125 Antibody Affinity
Old Tree	G12.425.143.140 Antigenic Modulation
Old Tree	G12.425.143.232 Binding Sites, Antibody
Old Tree	G12.425.143.281 Cross Reactions
Old Tree	G12.425.143.545 Hemolysis
Old Tree	G12.425.143.612 Immunologic Capping
Old Tree	G12.425.143.754 Passive Cutaneous Anaphylaxis
Old Tree	G12.425.143.780 Rh Isoimmunization
Old Tree	G12.425.199 Clonal Selection, Antigen-Mediated
Old Tree	G12.425.255 Complement Activation
Old Tree	G12.425.255.695 Complement Pathway, Alternative
Old Tree	G12.425.255.698 Complement Pathway, Classical
Old Tree	G12.425.255.849 Complement Pathway, Mannose-Binding Lectin
Old Tree	G12.425.270 Cytotoxicity, Immunologic
Old Tree	G12.425.270.070 Antibody-Dependent Cell Cytotoxicity
Old Tree	G12.425.270.500 Macrophage Activation
Old Tree	G12.425.431 Histamine Release
Old Tree	G12.425.700 Immune Evasion
Old Tree	G12.425.742 Immunogenetic Processes
Old Tree	G12.425.742.500 Gene Rearrangement, B-Lymphocyte
Old Tree	G12.425.742.500.501 Gene Rearrangement, B-Lymphocyte, Heavy Chain
Old Tree	G12.425.742.500.501.450 Immunoglobulin Class Switching
Old Tree	G12.425.742.500.601 Gene Rearrangement, B-Lymphocyte, Light Chain
Old Tree	G12.425.742.530 Gene Rearrangement, T-Lymphocyte
Old Tree	G12.425.742.530.111 Gene Rearrangement, alpha-Chain T-Cell Antigen Receptor
Old Tree	G12.425.742.530.211 Gene Rearrangement, beta-Chain T-Cell Antigen Receptor
Old Tree	G12.425.742.530.261 Gene Rearrangement, delta-Chain T-Cell Antigen Receptor
Old Tree	G12.425.742.530.311 Gene Rearrangement, gamma-Chain T-Cell Antigen Receptor
Old Tree	G12.425.742.765 Somatic Hypermutation, Immunoglobulin
Old Tree	G12.425.745 Immunologic Surveillance
Old Tree	G12.425.746 Immunomodulation
Old Tree	G12.425.746.425 Immune Tolerance
Old Tree	G12.425.746.425.100 Clonal Anergy

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	G12.425.746.425.120 Clonal Deletion
Old Tree	G12.425.746.425.800 Self Tolerance
Old Tree	G12.425.746.425.800.200 Central Tolerance
Old Tree	G12.425.746.425.800.600 Peripheral Tolerance
Old Tree	G12.425.746.425.910 Tachyphylaxis
Old Tree	G12.425.746.425.955 Transplantation Tolerance
Old Tree	G12.425.746.575 Neuroimmunomodulation
Old Tree	G12.425.747 Lymphocyte Activation
Old Tree	G12.425.747.150 Cross-Priming
Old Tree	G12.425.749 Lymphocyte Cooperation
Old Tree	G12.425.897 Neutrophil Activation
Old Tree	G12.425.898 Neutrophil Infiltration
Old Tree	G12.425.899 Phagocytosis
Old Tree	G12.425.899.500 Cytophagocytosis
Old Tree	G12.425.900 Seroconversion
Old Tree	G12.425.901 Transplantation Immunology
Old Tree	G12.425.901.402 Graft vs Host Reaction
Old Tree	G12.425.901.402.320 Graft vs Tumor Effect
Old Tree	G12.425.901.402.320.320 Graft vs Leukemia Effect
Old Tree	G12.425.901.519 Histocompatibility
Old Tree	G12.425.901.519.500 Histocompatibility, Maternal-Fetal
Old Tree	G12.425.901.545 Host vs Graft Reaction
Old Tree	G12.425.901.545.328 Graft Rejection
Old Tree	G12.425.901.545.340 Graft Survival
Old Tree	G12.425.905 Tumor Escape
-	G12.450 Immunity
-	G12.450.050 Adaptive Immunity
-	G12.450.050.184 Clonal Selection, Antigen-Mediated
-	G12.450.050.370 Immunity, Active
New Heading	<b>G12.450.050.370.500 Immunogenicity, Vaccine</b>
-	G12.450.050.400 Immunity, Cellular
-	G12.450.050.400.070 Antigen Presentation
-	G12.450.050.400.412 Immunologic Surveillance
-	G12.450.050.400.545 Lymphocyte Activation
-	G12.450.050.400.545.150 Cross-Priming

## MeSH Tree Changes for 2017

Type	Tree - heading
-	G12.450.050.420 Immunity, Humoral
New Heading	<b>G12.450.050.460 Immunogenicity, Vaccine</b>
-	G12.450.050.500 Immunologic Memory
-	G12.450.192 Autoimmunity
-	G12.450.275 Cross Protection
-	G12.450.400 Immunity, Herd
-	G12.450.418 Immunity, Heterologous
-	G12.450.564 Immunity, Innate
-	G12.450.564.204 Blood Bactericidal Activity
-	G12.450.564.250 Disease Resistance
-	G12.450.564.809 Phagocytosis
-	G12.450.564.809.500 Cytophagocytosis
-	G12.450.570 Immunity, Maternally-Acquired
-	G12.450.573 Immunity, Mucosal
-	G12.450.800 Plant Immunity
-	G12.450.800.250 Disease Resistance
-	G12.450.900 T-Cell Antigen Receptor Specificity
-	G12.460 Immunocompetence
-	G12.460.500 Immunosenescence
-	G12.470 Immunocompromised Host
-	G12.470.500 Radiation Chimera
-	G12.500 Immunogenetic Phenomena
-	G12.500.199 Antibody Diversity
-	G12.500.249 Antigenic Variation
New Tree	<b>G12.500.274 Gene Rearrangement, B-Lymphocyte</b>
New Tree	<b>G12.500.274.501 Gene Rearrangement, B-Lymphocyte, Heavy Chain</b>
New Tree	<b>G12.500.274.501.450 Immunoglobulin Class Switching</b>
New Tree	<b>G12.500.274.601 Gene Rearrangement, B-Lymphocyte, Light Chain</b>
New Tree	<b>G12.500.287 Gene Rearrangement, T-Lymphocyte</b>
New Tree	<b>G12.500.287.111 Gene Rearrangement, alpha-Chain T-Cell Antigen Receptor</b>
New Tree	<b>G12.500.287.211 Gene Rearrangement, beta-Chain T-Cell Antigen Receptor</b>

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	<a href="#">G12.500.287.261</a>	<a href="#">Gene Rearrangement, delta-Chain T-Cell Antigen Receptor</a>
New Tree	<a href="#">G12.500.287.311 Receptor</a>	<a href="#">Gene Rearrangement, gamma-Chain T-Cell Antigen</a>
-	G12.500.299	Genes, Immunoglobulin
Old Tree	<a href="#">G12.500.349</a>	<a href="#">Immunogenetic Processes</a>
Old Tree	<a href="#">G12.500.349.500</a>	<a href="#">Gene Rearrangement, B-Lymphocyte</a>
Old Tree	<a href="#">G12.500.349.500.501</a>	<a href="#">Gene Rearrangement, B-Lymphocyte, Heavy Chain</a>
Old Tree	<a href="#">G12.500.349.500.501.450</a>	<a href="#">Immunoglobulin Class Switching</a>
Old Tree	<a href="#">G12.500.349.500.601</a>	<a href="#">Gene Rearrangement, B-Lymphocyte, Light Chain</a>
Old Tree	<a href="#">G12.500.349.530</a>	<a href="#">Gene Rearrangement, T-Lymphocyte</a>
Old Tree	<a href="#">G12.500.349.530.111 Receptor</a>	<a href="#">Gene Rearrangement, alpha-Chain T-Cell Antigen</a>
Old Tree	<a href="#">G12.500.349.530.211 Receptor</a>	<a href="#">Gene Rearrangement, beta-Chain T-Cell Antigen</a>
Old Tree	<a href="#">G12.500.349.530.261 Receptor</a>	<a href="#">Gene Rearrangement, delta-Chain T-Cell Antigen</a>
Old Tree	<a href="#">G12.500.349.530.311 Receptor</a>	<a href="#">Gene Rearrangement, gamma-Chain T-Cell Antigen</a>
Old Tree	<a href="#">G12.500.349.765</a>	<a href="#">Somatic Hypermutation, Immunoglobulin</a>
-	G12.500.400	Immunoglobulin Allotypes
-	G12.500.400.586	Immunoglobulin Gm Allotypes
-	G12.500.400.587	Immunoglobulin Km Allotypes
-	G12.500.450	Immunoglobulin Idiotypes
-	G12.500.500	Major Histocompatibility Complex
-	G12.500.500.595	Genes, MHC Class I
-	G12.500.500.600	Genes, MHC Class II
-	G12.500.530	Minor Histocompatibility Loci
-	G12.500.560	Minor Lymphocyte Stimulatory Loci
New Tree	<a href="#">G12.500.780</a>	<a href="#">Somatic Hypermutation, Immunoglobulin</a>
New Heading	<b>G12.513</b>	<b>Immunogenicity, Vaccine</b>
New Tree	<a href="#">G12.525</a>	<a href="#">Immunologic Surveillance</a>
New Tree	<a href="#">G12.535</a>	<a href="#">Immunomodulation</a>
New Tree	<a href="#">G12.535.425</a>	<a href="#">Immune Tolerance</a>
New	<a href="#">G12.535.425.100</a>	<a href="#">Clonal Anergy</a>

## MeSH Tree Changes for 2017

Type	Tree - heading
Tree	
New Tree	G12.535.425.120 Clonal Deletion
New Tree	G12.535.425.800 Self Tolerance
New Tree	G12.535.425.800.200 Central Tolerance
New Tree	G12.535.425.800.600 Peripheral Tolerance
New Tree	G12.535.425.910 Tachyphylaxis
New Tree	G12.535.425.955 Transplantation Tolerance
New Tree	G12.535.575 Neuroimmunomodulation
New Tree	G12.565 Lymphocyte Activation
New Tree	G12.565.150 Cross-Priming
New Tree	G12.575 Lymphocyte Cooperation
New Tree	G12.604 Neutrophil Activation
New Tree	G12.632 Neutrophil Infiltration
New Tree	G12.688 Phagocytosis
New Tree	G12.688.500 Cytophagocytosis
New Tree	G12.800 Seroconversion
New Tree	G12.875 Transplantation Immunology
New Tree	G12.875.402 Graft vs Host Reaction
New Tree	G12.875.402.320 Graft vs Tumor Effect
New Tree	G12.875.402.320.320 Graft vs Leukemia Effect
New Tree	G12.875.519 Histocompatibility
New Tree	G12.875.519.500 Histocompatibility, Maternal-Fetal

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	<a href="#">G12.875.545</a>	<a href="#">Host vs Graft Reaction</a>
New Tree	<a href="#">G12.875.545.328</a>	<a href="#">Graft Rejection</a>
New Tree	<a href="#">G12.875.545.340</a>	<a href="#">Graft Survival</a>
New Tree	<a href="#">G12.900</a>	<a href="#">Tumor Escape</a>
-	G13	Integumentary System Physiological Phenomena
-	G13.500	Hair Color
-	G13.750	Skin Physiological Phenomena
New Tree	<a href="#">G13.750.208</a>	<a href="#">Cutaneous Elimination</a>
New Tree	<a href="#">G13.750.415</a>	<a href="#">Galvanic Skin Response</a>
New Tree	<a href="#">G13.750.622</a>	<a href="#">Piloerection</a>
New Tree	<a href="#">G13.750.726</a>	<a href="#">Re-Epithelialization</a>
New Tree	<a href="#">G13.750.778</a>	<a href="#">Skin Absorption</a>
New Tree	<a href="#">G13.750.804</a>	<a href="#">Skin Aging</a>
Old Tree	<a href="#">G13.750.829</a>	<a href="#">Skin Physiological Processes</a>
Old Tree	<a href="#">G13.750.829.159</a>	<a href="#">Cutaneous Elimination</a>
Old Tree	<a href="#">G13.750.829.320</a>	<a href="#">Galvanic Skin Response</a>
Old Tree	<a href="#">G13.750.829.690</a>	<a href="#">Piloerection</a>
Old Tree	<a href="#">G13.750.829.755</a>	<a href="#">Re-Epithelialization</a>
Old Tree	<a href="#">G13.750.829.820</a>	<a href="#">Skin Absorption</a>
Old Tree	<a href="#">G13.750.829.826</a>	<a href="#">Skin Aging</a>
Old Tree	<a href="#">G13.750.829.833</a>	<a href="#">Skin Pigmentation</a>
Old Tree	<a href="#">G13.750.829.833.500</a>	<a href="#">Suntan</a>
Old Tree	<a href="#">G13.750.829.855</a>	<a href="#">Sweating</a>
New Tree	<a href="#">G13.750.837</a>	<a href="#">Skin Pigmentation</a>
New Tree	<a href="#">G13.750.837.500</a>	<a href="#">Suntan</a>
-	G13.750.844	Skin Temperature
New Tree	<a href="#">G13.750.860</a>	<a href="#">Sweating</a>

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	G14	Ocular Physiological Phenomena
New Tree	<a href="#">G14.010</a>	<a href="#">Accommodation, Ocular</a>
New Tree	<a href="#">G14.020</a>	<a href="#">Adaptation, Ocular</a>
New Tree	<a href="#">G14.020.371</a>	<a href="#">Dark Adaptation</a>
-	G14.040	Afterimage
New Tree	<a href="#">G14.152</a>	<a href="#">Blinking</a>
-	G14.264	Dominance, Ocular
-	G14.330	Evoked Potentials, Visual
-	G14.340	Eye Color
New Tree	<a href="#">G14.350</a>	<a href="#">Eye Movements</a>
New Tree	<a href="#">G14.350.217</a>	<a href="#">Convergence, Ocular</a>
New Tree	<a href="#">G14.350.253</a>	<a href="#">Fixation, Ocular</a>
New Tree	<a href="#">G14.350.378</a>	<a href="#">Nystagmus, Physiologic</a>
New Tree	<a href="#">G14.350.378.500</a>	<a href="#">Nystagmus, Optokinetic</a>
New Tree	<a href="#">G14.350.453</a>	<a href="#">Pursuit, Smooth</a>
New Tree	<a href="#">G14.350.500</a>	<a href="#">Saccades</a>
-	G14.360	Figural Aftereffect
-	G14.370	Flicker Fusion
-	G14.400	Glare
-	G14.440	Intraocular Pressure
New Tree	<a href="#">G14.540</a>	<a href="#">Lacrimal Elimination</a>
New Tree	<a href="#">G14.590</a>	<a href="#">Ocular Absorption</a>
Old Tree	<a href="#">G14.640</a>	<a href="#">Ocular Physiological Processes</a>
Old Tree	<a href="#">G14.640.079</a>	<a href="#">Accommodation, Ocular</a>
Old Tree	<a href="#">G14.640.154</a>	<a href="#">Adaptation, Ocular</a>
Old Tree	<a href="#">G14.640.154.371</a>	<a href="#">Dark Adaptation</a>
Old Tree	<a href="#">G14.640.182</a>	<a href="#">Blinking</a>



## MeSH Tree Changes for 2017

Type	Tree - heading	
Old Tree	G14.640.260	Eye Movements
Old Tree	G14.640.260.217	Convergence, Ocular
Old Tree	G14.640.260.253	Fixation, Ocular
Old Tree	G14.640.260.378	Nystagmus, Physiologic
Old Tree	G14.640.260.378.500	Nystagmus, Optokinetic
Old Tree	G14.640.260.453	Pursuit, Smooth
Old Tree	G14.640.260.500	Saccades
Old Tree	G14.640.399	Lacrimal Elimination
Old Tree	G14.640.538	Ocular Absorption
Old Tree	G14.640.816	Vision, Ocular
Old Tree	G14.640.816.124	Color Vision
Old Tree	G14.640.816.186	Mesopic Vision
Old Tree	G14.640.816.186.624	Rod-Cone Interaction
Old Tree	G14.640.816.249	Night Vision
Old Tree	G14.640.816.500	Phosphenes
Old Tree	G14.640.816.730	Vision, Entoptic
-	G14.700	Optic Flow
-	G14.760	Refraction, Ocular
-	G14.930	Vision Disparity
New Tree	G14.935	Vision, Ocular
New Tree	G14.935.124	Color Vision
New Tree	G14.935.186	Mesopic Vision
New Tree	G14.935.186.624	Rod-Cone Interaction
New Tree	G14.935.249	Night Vision
New Tree	G14.935.500	Phosphenes
New Tree	G14.935.730	Vision, Entoptic
-	G14.940	Visual Acuity
-	G14.940.500	Contrast Sensitivity
-	G14.940.750	Emmetropia
-	G14.950	Visual Fields
-	G15	Plant Physiological Phenomena

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	G15.150	Etiolation
Old Tree	G15.277	Organogenesis, Plant
New Tree	G15.300	Gametogenesis, Plant
New Tree	G15.357	Germination
New Tree	G15.357.500	Plant Dormancy
New Tree	G15.375	Gravity Sensing
New Tree	G15.525	Organogenesis, Plant
New Tree	G15.547	Parthenogenesis
New Tree	G15.547.500	Apomixis
Old Tree	G15.556	Plant Diseases
Old Tree	G15.556.700	Plant Tumors
New Tree	G15.568	Photosynthesis
New Tree	G15.589	Plant Development
Old Tree	G15.602	Plant Immunity
Old Tree	G15.602.250	Disease Resistance
New Tree	G15.610	Plant Diseases
New Tree	G15.610.700	Plant Tumors
New Tree	G15.620	Plant Dispersal
New Tree	G15.620.500	Seed Dispersal
New Tree	G15.630	Plant Immunity
New Tree	G15.630.250	Disease Resistance
-	G15.649	Plant Infertility
New Tree	G15.682	Plant Root Nodulation
New Tree	G15.698	Plant Somatic Embryogenesis Techniques

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	G15.713	Plant Transpiration
Old Tree	G15.744	Plant Physiological Processes
Old Tree	G15.744.074	Apomixis
Old Tree	G15.744.111	Etiolation
Old Tree	G15.744.149	Gametogenesis, Plant
Old Tree	G15.744.300	Germination
Old Tree	G15.744.300.500	Plant Dormancy
Old Tree	G15.744.450	Gravity Sensing
Old Tree	G15.744.700	Photosynthesis
Old Tree	G15.744.723	Plant Development
Old Tree	G15.744.735	Plant Dispersal
Old Tree	G15.744.735.500	Seed Dispersal
Old Tree	G15.744.747	Plant Root Nodulation
Old Tree	G15.744.749	Plant Somatic Embryogenesis Techniques
Old Tree	G15.744.750	Plant Transpiration
Old Tree	G15.744.800	Pollination
Old Tree	G15.744.850	Self-Fertilization
Old Tree	G15.744.875	Self-Incompatibility in Flowering Plants
Old Tree	G15.744.900	Tropism
Old Tree	G15.744.900.400	Gravitropism
Old Tree	G15.744.900.700	Phototropism
New Tree	G15.776	Pollination
-	G15.808	Seed Dispersal
New Tree	G15.830	Self-Fertilization
New Tree	G15.840	Self-Incompatibility in Flowering Plants
New Tree	G15.920	Tropism
New Tree	G15.920.400	Gravitropism
New Tree	G15.920.700	Phototropism
-	G16	Biological Phenomena
New Tree	G16.012	Adaptation, Biological
New	G16.012.500	Adaptation, Physiological

## MeSH Tree Changes for 2017

Type	Tree - heading	
Tree		
New Tree	<a href="#">G16.012.500.133</a>	Acclimatization
New Tree	<a href="#">G16.012.500.133.250</a>	Salt-Tolerance
New Heading	<b><a href="#">G16.012.500.133.500</a></b>	<b>Thermotolerance</b>
New Tree	<a href="#">G16.012.500.274</a>	Allostasis
New Tree	<a href="#">G16.012.500.535</a>	Body Temperature Regulation
New Tree	<a href="#">G16.012.500.535.693</a>	Sweating
New Tree	<a href="#">G16.012.500.535.778</a>	Thermogenesis
New Tree	<a href="#">G16.012.500.535.778.500</a>	Shivering
New Tree	<a href="#">G16.012.500.535.889</a>	Torpor
New Tree	<a href="#">G16.012.500.535.889.249</a>	Estivation
New Tree	<a href="#">G16.012.500.535.889.500</a>	Hibernation
New Tree	<a href="#">G16.012.750</a>	Biological Mimicry
New Tree	<a href="#">G16.012.750.500</a>	Molecular Mimicry
-	G16.024	Allelopathy
-	G16.049	Animal Distribution
New Tree	<a href="#">G16.062</a>	Antibiosis
New Tree	<a href="#">G16.075</a>	Biological Evolution
New Tree	<a href="#">G16.075.250</a>	Evolution, Molecular
New Tree	<a href="#">G16.075.250.750</a>	Mutation Rate
New Tree	<a href="#">G16.075.350</a>	Genetic Speciation
New Tree	<a href="#">G16.075.605</a>	Phylogeny
New Tree	<a href="#">G16.075.802</a>	Reproductive Isolation

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	G16.100 Biological Processes
Old Tree	G16.100.057 Adaptation, Biological
Old Tree	G16.100.057.500 Adaptation, Physiological
Old Tree	G16.100.057.500.133 Acclimatization
Old Tree	G16.100.057.500.274 Allostasis
Old Tree	G16.100.057.500.535 Body Temperature Regulation
Old Tree	G16.100.057.500.535.693 Sweating
Old Tree	G16.100.057.500.535.778 Thermogenesis
Old Tree	G16.100.057.500.535.778.500 Shivering
Old Tree	G16.100.057.500.535.889 Torpor
Old Tree	G16.100.057.500.535.889.249 Estivation
Old Tree	G16.100.057.500.535.889.500 Hibernation
Old Tree	G16.100.057.750 Biological Mimicry
Old Tree	G16.100.057.750.500 Molecular Mimicry
Old Tree	G16.100.065 Antibiosis
Old Tree	G16.100.178 Biological Evolution
Old Tree	G16.100.178.250 Evolution, Molecular
Old Tree	G16.100.178.250.750 Mutation Rate
Old Tree	G16.100.178.350 Genetic Speciation
Old Tree	G16.100.178.605 Phylogeny
Old Tree	G16.100.178.802 Reproductive Isolation
Old Tree	G16.100.200 Colony Collapse
Old Tree	G16.100.277 Denitrification
Old Tree	G16.100.280 Extinction, Biological
Old Tree	G16.100.380 Host-Pathogen Interactions
Old Tree	G16.100.380.380 Host Specificity
Old Tree	G16.100.380.400 Host-Parasite Interactions
Old Tree	G16.100.380.700 Immune Evasion
Old Tree	G16.100.400 Organelle Biogenesis
Old Tree	G16.100.856 Regeneration
Old Tree	G16.100.856.150 Bone Remodeling
Old Tree	G16.100.856.150.150 Bone Regeneration
Old Tree	G16.100.856.150.150.570 Osseointegration
Old Tree	G16.100.856.468 Liver Regeneration
Old Tree	G16.100.856.611 Nerve Regeneration
Old Tree	G16.100.856.611.500 Spinal Cord Regeneration

## MeSH Tree Changes for 2017

Type	Tree - heading	
Old Tree	G16.100.856.891	Wound Healing
Old Tree	G16.100.856.891.249	Cicatrix
Old Tree	G16.100.856.891.500	Fracture Healing
Old Tree	G16.100.856.891.750	Re-Epithelialization
Old Tree	G16.100.870	Remission, Spontaneous
Old Tree	G16.100.870.500	Neoplasm Regression, Spontaneous
Old Tree	G16.100.900	Symbiosis
New Tree	G16.150	Colony Collapse
-	G16.500	Ecological and Environmental Phenomena
New Tree	G16.500.050	Air Ionization
New Tree	G16.500.100	Biofouling
New Tree	G16.500.150	Carbon Cycle
New Tree	G16.500.150.500	Carbon Sequestration
New Tree	G16.500.175	Climatic Processes
New Tree	G16.500.175.249	Air Movements
New Tree	G16.500.175.249.200	Wind
New Tree	G16.500.175.374	Climate Change
New Tree	G16.500.175.374.500	Global Warming
New Tree	G16.500.175.500	Cyclonic Storms
New Tree	G16.500.175.781	Droughts
New Tree	G16.500.175.812	Floods
New Tree	G16.500.175.827	Greenhouse Effect
New Tree	G16.500.175.843	Lightning
New Tree	G16.500.175.859	Rain
New Tree	G16.500.175.859.050	Acid Rain

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	G16.500.175.867 Snow
New Tree	G16.500.175.875 Tidal Waves
New Tree	G16.500.200 Cosmic Radiation
Old Tree	G16.500.240 Ecological and Environmental Processes
Old Tree	G16.500.240.070 Air Ionization
Old Tree	G16.500.240.102 Biofouling
Old Tree	G16.500.240.118 Carbon Cycle
Old Tree	G16.500.240.135 Climatic Processes
Old Tree	G16.500.240.135.249 Air Movements
Old Tree	G16.500.240.135.249.200 Wind
Old Tree	G16.500.240.135.374 Climate Change
Old Tree	G16.500.240.135.374.500 Global Warming
Old Tree	G16.500.240.135.500 Cyclonic Storms
Old Tree	G16.500.240.135.781 Droughts
Old Tree	G16.500.240.135.812 Floods
Old Tree	G16.500.240.135.827 Greenhouse Effect
Old Tree	G16.500.240.135.843 Lightning
Old Tree	G16.500.240.135.859 Rain
Old Tree	G16.500.240.135.859.050 Acid Rain
Old Tree	G16.500.240.135.867 Snow
Old Tree	G16.500.240.135.875 Tidal Waves
Old Tree	G16.500.240.200 Cosmic Radiation
Old Tree	G16.500.240.300 Eutrophication
Old Tree	G16.500.240.300.400 Harmful Algal Bloom
Old Tree	G16.500.240.465 Nitrogen Cycle
Old Tree	G16.500.240.465.249 Denitrification
Old Tree	G16.500.240.465.500 Nitrification
Old Tree	G16.500.240.547 Ozone Depletion
Old Tree	G16.500.240.630 Rhizosphere
Old Tree	G16.500.240.795 Water Cycle
Old Tree	G16.500.240.960 Water Movements
-	G16.500.275 Environment
-	G16.500.275.058 Altitude
-	G16.500.275.063 Atmosphere

## MeSH Tree Changes for 2017

Type	Tree - heading
-	G16.500.275.063.150 Air
-	G16.500.275.063.150.500 Compressed Air
-	G16.500.275.063.700 Stratospheric Ozone
-	G16.500.275.063.725 Weather
-	G16.500.275.063.725.154 Air Movements
-	G16.500.275.063.725.154.200 Wind
-	G16.500.275.063.725.310 Humidity
-	G16.500.275.063.725.395 Rain
-	G16.500.275.063.725.395.050 Acid Rain
-	G16.500.275.063.725.480 Snow
-	G16.500.275.063.725.525 Sunlight
-	G16.500.275.063.725.525.400 Infrared Rays
-	G16.500.275.063.725.525.600 Ultraviolet Rays
-	G16.500.275.063.725.710 Temperature
-	G16.500.275.063.725.710.300 Cold Temperature
-	G16.500.275.063.725.710.380 Hot Temperature
-	G16.500.275.066 Carbon Footprint
-	G16.500.275.067 Caves
-	G16.500.275.069 Cities
-	G16.500.275.071 Climate
-	G16.500.275.071.275 Cold Climate
-	G16.500.275.071.325 Desert Climate
-	G16.500.275.071.387 El Nino-Southern Oscillation
-	G16.500.275.071.450 Microclimate
-	G16.500.275.071.590 Seasons
-	G16.500.275.071.600 Tropical Climate
-	G16.500.275.074 Confined Spaces
-	G16.500.275.157 Ecosystem
-	G16.500.275.157.049 Biodiversity
-	G16.500.275.157.049.100 Biota
-	G16.500.275.157.049.100.500 Microbiota
-	G16.500.275.157.049.100.500.375 Gastrointestinal Microbiome
-	G16.500.275.157.049.100.500.750 Microbial Consortia
New Heading	<b>G16.500.275.157.049.100.500.875 Mycobiome</b>
-	G16.500.275.157.049.230 Ecotype



## MeSH Tree Changes for 2017

Type	Tree - heading
-	G16.500.275.157.049.250                      Endangered Species
-	G16.500.275.157.049.400                      Introduced Species
-	G16.500.275.157.100                          Biomass
-	G16.500.275.157.130                          Coral Reefs
-	G16.500.275.157.240                          Ecological Systems, Closed
-	G16.500.275.157.250                          Food Chain
-	G16.500.275.157.437                          Forests
-	G16.500.275.157.437.500                      Rainforest
-	G16.500.275.157.437.750                      Taiga
-	G16.500.275.157.531                          Grassland
-	G16.500.275.157.625                          Rhizosphere
-	G16.500.275.157.718                          Tundra
-	G16.500.275.157.718.500                      Permafrost
-	G16.500.275.157.812                          Wetlands
-	G16.500.275.240                              Extraterrestrial Environment
-	G16.500.275.240.200                          Cosmic Dust
New Heading	<b>G16.500.275.260                              Extreme Environments</b>
New Tree	<b>G16.500.275.260.500                          Hot Springs</b>
New Tree	<b>G16.500.275.260.750                          Hydrothermal Vents</b>
-	G16.500.275.280                              Fresh Water
-	G16.500.275.280.500                          Lakes
-	G16.500.275.280.625                          Ponds
-	G16.500.275.280.650                          Rivers
-	G16.500.275.410                              Ice
-	G16.500.275.410.500                          Ice Cover
-	G16.500.275.505                              Islands
-	G16.500.275.553                              Natural Resources
-	G16.500.275.553.500                          Water Resources
-	G16.500.275.600                              Noise
-	G16.500.275.640                              Odorants
-	G16.500.275.640                              Odors
-	G16.500.275.683                              Refugium
-	G16.500.275.725                              Saline Waters
-	G16.500.275.725.500                          Seawater

## MeSH Tree Changes for 2017

Type	Tree - heading
-	G16.500.275.725.500.650 Oceans and Seas
-	G16.500.275.725.500.650.075 Aquatic Organisms
-	G16.500.275.815 Soil
-	G16.500.275.815.500 Permafrost
-	G16.500.275.965 Wilderness
New Tree	<a href="#">G16.500.285</a> Eutrophication
New Tree	<a href="#">G16.500.285.400</a> Harmful Algal Bloom
-	G16.500.320 Geologic Sediments
-	G16.500.750 Meteorological Concepts
-	G16.500.750.274 Atmospheric Pressure
-	G16.500.750.274.100 Air Pressure
-	G16.500.750.274.902 Vacuum
-	G16.500.750.775 Weather
-	G16.500.750.775.154 Droughts
-	G16.500.750.775.232 Extreme Cold
-	G16.500.750.775.271 Extreme Heat
-	G16.500.750.775.310 Humidity
-	G16.500.750.775.342 Ice
-	G16.500.750.775.375 Lightning
-	G16.500.750.775.450 Rain
-	G16.500.750.775.450.050 Acid Rain
-	G16.500.750.775.480 Snow
-	G16.500.750.775.525 Sunlight
-	G16.500.750.775.525.400 Infrared Rays
-	G16.500.750.775.525.600 Ultraviolet Rays
-	G16.500.750.775.710 Temperature
-	G16.500.750.775.710.300 Cold Temperature
-	G16.500.750.775.710.380 Hot Temperature
-	G16.500.750.775.745 Tornadoes
-	G16.500.750.775.780 Wind
New Tree	<a href="#">G16.500.768</a> Nitrogen Cycle
New Tree	<a href="#">G16.500.768.249</a> Denitrification
New	<a href="#">G16.500.768.500</a> Nitrification

## MeSH Tree Changes for 2017

Type	Tree - heading	
Tree		
New Tree	<a href="#">G16.500.768.600</a>	<a href="#">Nitrogen Fixation</a>
New Tree	<a href="#">G16.500.785</a>	<a href="#">Ozone Depletion</a>
-	G16.500.819	Refugium
New Tree	<a href="#">G16.500.853</a>	<a href="#">Rhizosphere</a>
-	G16.500.887	Steam
New Tree	<a href="#">G16.500.943</a>	<a href="#">Water Cycle</a>
New Tree	<a href="#">G16.500.971</a>	<a href="#">Water Movements</a>
New Tree	<a href="#">G16.510</a>	<a href="#">Extinction, Biological</a>
New Tree	<a href="#">G16.543</a>	<a href="#">Host-Pathogen Interactions</a>
New Tree	<a href="#">G16.543.380</a>	<a href="#">Host Specificity</a>
New Tree	<a href="#">G16.543.400</a>	<a href="#">Host-Parasite Interactions</a>
New Tree	<a href="#">G16.543.700</a>	<a href="#">Immune Evasion</a>
New Heading	<b>G16.559</b>	<b>Host-Seeking Behavior</b>
New Heading	<b>G16.575</b>	<b>Life History Traits</b>
Old Tree	<b>G16.595</b>	<b>Origin of Life</b>
New Tree	<a href="#">G16.645</a>	<a href="#">Organelle Biogenesis</a>
New Tree	<a href="#">G16.650</a>	<a href="#">Origin of Life</a>
-	G16.690	Pigmentation
-	G16.690.360	Eye Color
-	G16.690.490	Hair Color
-	G16.690.890	Skin Pigmentation
-	G16.757	Recovery of Function
New Tree	<a href="#">G16.762</a>	<a href="#">Regeneration</a>
New Tree	<a href="#">G16.762.150</a>	<a href="#">Bone Remodeling</a>
New	<a href="#">G16.762.150.150</a>	<a href="#">Bone Regeneration</a>

## MeSH Tree Changes for 2017

Type	Tree - heading	
Tree		
New Tree	<a href="#">G16.762.150.150.570</a>	<a href="#">Osseointegration</a>
New Tree	<a href="#">G16.762.468</a>	<a href="#">Liver Regeneration</a>
New Tree	<a href="#">G16.762.611</a>	<a href="#">Nerve Regeneration</a>
New Tree	<a href="#">G16.762.611.500</a>	<a href="#">Spinal Cord Regeneration</a>
New Tree	<a href="#">G16.762.891</a>	<a href="#">Wound Healing</a>
New Tree	<a href="#">G16.762.891.249</a>	<a href="#">Cicatrix</a>
New Tree	<a href="#">G16.762.891.500</a>	<a href="#">Fracture Healing</a>
New Tree	<a href="#">G16.762.891.750</a>	<a href="#">Re-Epithelialization</a>
New Tree	<a href="#">G16.767</a>	<a href="#">Remission, Spontaneous</a>
New Tree	<a href="#">G16.767.500</a>	<a href="#">Neoplasm Regression, Spontaneous</a>
-	<a href="#">G16.824</a>	<a href="#">Species Specificity</a>
New Tree	<a href="#">G16.840</a>	<a href="#">Symbiosis</a>
-	<a href="#">G16.920</a>	<a href="#">Tissue Survival</a>
-	<a href="#">G17</a>	<a href="#">Mathematical Concepts</a>
-	<a href="#">G17.035</a>	<a href="#">Algorithms</a>
-	<a href="#">G17.035.250</a>	<a href="#">Artificial Intelligence</a>
-	<a href="#">G17.035.250.500</a>	<a href="#">Machine Learning</a>
-	<a href="#">G17.035.250.500.500</a>	<a href="#">Supervised Machine Learning</a>
-	<a href="#">G17.035.250.500.500.500</a>	<a href="#">Support Vector Machine</a>
-	<a href="#">G17.035.250.500.750</a>	<a href="#">Unsupervised Machine Learning</a>
-	<a href="#">G17.162</a>	<a href="#">Decision Theory</a>
-	<a href="#">G17.162.500</a>	<a href="#">Decision Trees</a>
-	<a href="#">G17.226</a>	<a href="#">Fourier Analysis</a>
-	<a href="#">G17.290</a>	<a href="#">Fractals</a>
-	<a href="#">G17.388</a>	<a href="#">Game Theory</a>
-	<a href="#">G17.485</a>	<a href="#">Neural Networks (Computer)</a>
-	<a href="#">G17.582</a>	<a href="#">Nomograms</a>
-	<a href="#">G17.680</a>	<a href="#">Probability</a>

## MeSH Tree Changes for 2017

Type	Tree - heading
-	G17.680.500 Odds Ratio
-	G17.680.750 Risk
-	G17.680.875 Uncertainty
-	G17.800 Sensitivity and Specificity
-	G17.800.500 Signal-To-Noise Ratio
-	G17.820 Statistical Distributions
-	G17.820.250 Binomial Distribution
-	G17.820.300 Chi-Square Distribution
-	G17.820.500 Normal Distribution
-	G17.820.750 Poisson Distribution
-	G17.830 Stochastic Processes
-	G17.830.500 Markov Chains
-	G17.915 Wavelet Analysis
-	H01 Natural Science Disciplines
-	H01.158 Biological Science Disciplines
-	H01.158.100 Anatomy
-	H01.158.100.091 Anatomy, Artistic
-	H01.158.100.154 Anatomy, Comparative
-	H01.158.100.185 Anatomy, Cross-Sectional
-	H01.158.100.185.905 Visible Human Projects
-	H01.158.100.216 Anatomy, Regional
-	H01.158.100.279 Anatomy, Veterinary
-	H01.158.100.433 Cell Biology
-	H01.158.100.529 Embryology
-	H01.158.100.529.500 Teratology
-	H01.158.100.656 Histology
-	H01.158.100.656.234 Histochemistry
-	H01.158.100.656.234.512 Immunohistochemistry
-	H01.158.100.656.469 Histology, Comparative
-	H01.158.100.700 Neuroanatomy
-	H01.158.100.720 Osteology
-	H01.158.201 Biochemistry
-	H01.158.201.085 Carbohydrate Biochemistry
-	H01.158.201.085.320 Glycomics
-	H01.158.201.214 Chemistry, Bioinorganic
-	H01.158.201.344 Histochemistry

## MeSH Tree Changes for 2017

Type	Tree - heading
-	H01.158.201.344.512 Immunohistochemistry
-	H01.158.201.486 Immunochemistry
-	H01.158.201.486.512 Immunohistochemistry
-	H01.158.201.586 Metabolomics
-	H01.158.201.636 Molecular Biology
-	H01.158.201.636.475 Molecular Medicine
-	H01.158.201.636.475.500 Molecular Epidemiology
-	H01.158.201.636.475.750 Pathology, Molecular
-	H01.158.201.687 Neurochemistry
-	H01.158.201.843 Proteomics
New Heading	<b>H01.158.201.843.500 Proteogenomics</b>
-	H01.158.273 Biology
-	H01.158.273.118 Botany
-	H01.158.273.118.299 Ethnobotany
-	H01.158.273.118.598 Pharmacognosy
-	H01.158.273.118.598.500 Herbal Medicine
-	H01.158.273.180 Computational Biology
-	H01.158.273.180.350 Genomics
-	H01.158.273.180.350.074 Epigenomics
-	H01.158.273.180.350.150 Glycomics
-	H01.158.273.180.350.162 HapMap Project
-	H01.158.273.180.350.174 Human Genome Project
-	H01.158.273.180.350.349 Nutrigenomics
-	H01.158.273.180.350.700 Proteomics
New Heading	<b>H01.158.273.180.350.700.500 Proteogenomics</b>
-	H01.158.273.180.599 Metabolomics
-	H01.158.273.180.800 Systems Biology
-	H01.158.273.190 Cell Biology
-	H01.158.273.200 Developmental Biology
-	H01.158.273.200.350 Embryology
-	H01.158.273.200.350.500 Teratology
-	H01.158.273.200.675 Plant Pathology
-	H01.158.273.248 Ecology
-	H01.158.273.248.500 Ecotoxicology

## MeSH Tree Changes for 2017

Type	Tree - heading
-	H01.158.273.248.750 Hydrobiology
-	H01.158.273.248.750.249 Freshwater Biology
-	H01.158.273.248.750.500 Marine Biology
-	H01.158.273.248.875 Hydrology
-	H01.158.273.295 Exobiology
-	H01.158.273.343 Genetics
-	H01.158.273.343.180 Cytogenetics
-	H01.158.273.343.249 Genetic Research
-	H01.158.273.343.249.099 Gene Ontology
-	H01.158.273.343.249.199 HapMap Project
-	H01.158.273.343.249.400 Human Genome Project
-	H01.158.273.343.290 Genetics, Behavioral
-	H01.158.273.343.315 Genetics, Medical
-	H01.158.273.343.315.384 Genetic Counseling
-	H01.158.273.343.330 Genetics, Microbial
-	H01.158.273.343.335 Genetics, Population
-	H01.158.273.343.335.500 Phylogeography
-	H01.158.273.343.350 Genomics
-	H01.158.273.343.350.042 Epigenomics
-	H01.158.273.343.350.086 Glycomics
-	H01.158.273.343.350.130 HapMap Project
-	H01.158.273.343.350.174 Human Genome Project
-	H01.158.273.343.350.261 Metagenomics
-	H01.158.273.343.350.349 Nutrigenomics
-	H01.158.273.343.350.700 Proteomics
New Heading	<b>H01.158.273.343.350.700.500 Proteogenomics</b>
-	H01.158.273.343.420 Immunogenetics
-	H01.158.273.343.595 Molecular Biology
-	H01.158.273.343.595.475 Molecular Medicine
-	H01.158.273.343.595.475.500 Molecular Epidemiology
-	H01.158.273.343.595.475.750 Pathology, Molecular
-	H01.158.273.343.750 Pharmacogenetics
-	H01.158.273.343.800 Radiation Genetics
-	H01.158.273.343.900 Toxicogenetics
-	H01.158.273.368 Laboratory Animal Science

## MeSH Tree Changes for 2017

Type	Tree - heading
-	H01.158.273.540 Microbiology
-	H01.158.273.540.177 Bacteriology
-	H01.158.273.540.274 Environmental Microbiology
-	H01.158.273.540.274.110 Air Microbiology
-	H01.158.273.540.274.332 Food Microbiology
-	H01.158.273.540.274.555 Soil Microbiology
-	H01.158.273.540.274.777 Water Microbiology
-	H01.158.273.540.367 Genetics, Microbial
-	H01.158.273.540.460 Industrial Microbiology
-	H01.158.273.540.553 Mycology
-	H01.158.273.540.706 Plant Pathology
-	H01.158.273.540.859 Virology
-	H01.158.273.602 Natural History
-	H01.158.273.610 Neurobiology
-	H01.158.273.688 Parasitology
-	H01.158.273.688.500 Food Parasitology
-	H01.158.273.738 Photobiology
-	H01.158.273.789 Radiobiology
-	H01.158.273.866 Sociobiology
-	H01.158.273.904 Synthetic Biology
-	H01.158.273.943 Zoology
-	H01.158.273.943.409 Entomology
-	H01.158.344 Biophysics
-	H01.158.344.310 Bionics
-	H01.158.344.528 Electrophysiology
-	H01.158.344.528.500 Cardiac Electrophysiology
-	H01.158.550 Biotechnology
-	H01.158.550.100 Biomimetics
-	H01.158.580 Chronobiology Discipline
-	H01.158.610 Neurosciences
-	H01.158.610.030 Cognitive Neuroscience
-	H01.158.610.060 Neuroanatomy
-	H01.158.610.080 Neurobiology
-	H01.158.610.110 Neurochemistry
-	H01.158.610.160 Neuroendocrinology
-	H01.158.610.184 Neuropathology



## MeSH Tree Changes for 2017

Type	Tree - heading
-	H01.158.610.208      Neuropharmacology
-	H01.158.610.268      Neurophysiology
-	H01.158.703            Pharmacology
-	H01.158.703.003      Biopharmaceutics
-	H01.158.703.007      Chemistry, Pharmaceutical
-	H01.158.703.007.675      Drug Discovery
-	H01.158.703.007.675.500      Drug Design
-	H01.158.703.015      Ethnopharmacology
-	H01.158.703.030      Neuropharmacology
-	H01.158.703.045      Pharmacoepidemiology
-	H01.158.703.052      Pharmacogenetics
-	H01.158.703.060      Pharmacognosy
-	H01.158.703.060.500      Herbal Medicine
-	H01.158.703.152      Pharmacology, Clinical
-	H01.158.703.546      Psychopharmacology
-	H01.158.782            Physiology
-	H01.158.782.236      Electrophysiology
-	H01.158.782.236.500      Cardiac Electrophysiology
-	H01.158.782.323      Endocrinology
-	H01.158.782.323.200      Neuroendocrinology
-	H01.158.782.562      Neurophysiology
-	H01.158.782.688      Physiology, Comparative
-	H01.158.782.795      Psychophysiology
-	H01.158.782.795.110      Neuropsychology
-	H01.158.782.795.180      Psychoneuroimmunology
-	H01.158.891            Toxicology
-	H01.158.891.211      Ecotoxicology
-	H01.158.891.424      Forensic Toxicology
-	H01.158.891.850      Toxicogenetics
-	H01.181                Chemistry
-	H01.181.122            Biochemistry
-	H01.181.122.253      Chemistry, Bioinorganic
-	H01.181.122.508      Glycomics
-	H01.181.122.573      Histochemistry
-	H01.181.122.573.512      Immunohistochemistry
-	H01.181.122.605      Immunochemistry

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	H01.181.122.605.512	Immunohistochemistry
-	H01.181.122.638	Metabolomics
-	H01.181.122.650	Molecular Biology
-	H01.181.122.650.475	Molecular Medicine
-	H01.181.122.650.475.550	Molecular Epidemiology
-	H01.181.122.650.475.680	Pathology, Molecular
-	H01.181.122.709	Neurochemistry
-	H01.181.122.738	Proteomics
New Heading	<b>H01.181.122.738.500</b>	<b>Proteogenomics</b>
-	H01.181.216	Chemistry, Agricultural
-	H01.181.309	Chemistry, Analytic
-	H01.181.341	Chemistry, Clinical
-	H01.181.370	Chemistry, Inorganic
-	H01.181.370.310	Chemistry, Bioinorganic
-	H01.181.404	Chemistry, Organic
-	H01.181.466	Chemistry, Pharmaceutical
-	H01.181.466.675	Drug Discovery
-	H01.181.466.675.500	Drug Design
-	H01.181.529	Chemistry, Physical
-	H01.181.529.240	Crystallography
-	H01.181.529.307	Electrochemistry
-	H01.181.529.711	Photochemistry
-	H01.181.529.776	Radiochemistry
-	H01.181.650	Microchemistry
-	H01.277	Earth Sciences
-	H01.277.249	Ecology
-	H01.277.249.500	Ecotoxicology
-	H01.277.249.750	Hydrobiology
-	H01.277.249.750.249	Freshwater Biology
-	H01.277.249.750.500	Marine Biology
-	H01.277.249.875	Hydrology
-	H01.277.500	Geography
-	H01.277.500.097	Geography, Medical
-	H01.277.500.097.500	Topography, Medical
-	H01.277.500.589	Phylogeography

## MeSH Tree Changes for 2017

Type	Tree - heading
-	H01.277.562 Geology
-	H01.277.625 Limnology
-	H01.277.625.500 Freshwater Biology
-	H01.277.687 Meteorology
-	H01.277.750 Oceanography
-	H01.277.750.500 Marine Biology
-	H01.277.875 Paleontology
-	H01.548 Mathematics
-	H01.548.675 Nonlinear Dynamics
-	H01.548.832 Statistics as Topic
-	H01.570 Microtechnology
-	H01.603 Nanotechnology
-	H01.603.600 Nanomedicine
-	H01.603.600.500 Theranostic Nanomedicine
-	H01.671 Physics
-	H01.671.031 Acoustics
-	H01.671.031.849 Ultrasonics
-	H01.671.065 Astronomy
-	H01.671.100 Biophysics
-	H01.671.100.202 Bionics
-	H01.671.293 Electronics
-	H01.671.293.159 Bionics
-	H01.671.293.319 Electronics, Medical
-	H01.671.293.643 Robotics
-	H01.671.368 Health Physics
-	H01.671.493 Magnetism
-	H01.671.515 Mechanics
-	H01.671.547 Meteorology
-	H01.671.579 Nuclear Physics
-	H01.671.579.800 Quantum Theory
-	H01.671.617 Optics and Photonics
-	H01.671.617.249 Fiber Optic Technology
-	H01.671.617.562 Microscopy
-	H01.671.617.755 Refractometry
-	H01.671.808 Rheology
-	H01.671.808.500 Microfluidics

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	H01.770	Science
-	H01.770.552	Natural History
-	H01.770.644	Research
-	H01.770.644.053	Animal Experimentation
-	H01.770.644.108	Behavioral Research
-	H01.770.644.108.687	Biobehavioral Sciences
-	H01.770.644.145	Biomedical Research
-	H01.770.644.145.124	Dental Research
-	H01.770.644.145.250	Embryo Research
-	H01.770.644.145.300	Fetal Research
-	H01.770.644.145.350	Genetic Research
-	H01.770.644.145.350.124	Gene Ontology
-	H01.770.644.145.350.249	HapMap Project
-	H01.770.644.145.350.500	Human Genome Project
-	H01.770.644.145.360	Health Services Research
-	H01.770.644.145.360.500	Comparative Effectiveness Research
-	H01.770.644.145.365	Human Experimentation
-	H01.770.644.145.365.100	Autoexperimentation
-	H01.770.644.145.365.750	Nontherapeutic Human Experimentation
-	H01.770.644.145.365.875	Therapeutic Human Experimentation
-	H01.770.644.145.390	Nursing Research
-	H01.770.644.145.390.234	Clinical Nursing Research
-	H01.770.644.145.390.385	Nursing Administration Research
-	H01.770.644.145.390.413	Nursing Education Research
-	H01.770.644.145.390.432	Nursing Evaluation Research
-	H01.770.644.145.390.634	Nursing Methodology Research
-	H01.770.644.145.431	Outcome Assessment (Health Care)
-	H01.770.644.145.431.500	Failure to Rescue, Health Care
New Heading	<b>H01.770.644.145.442</b>	<b>Pharmaceutical Research</b>
New Heading	<b>H01.770.644.145.452</b>	<b>Pharmacy Research</b>
-	H01.770.644.145.472	Rehabilitation Research
-	H01.770.644.145.512	Stem Cell Research
-	H01.770.644.145.675	Translational Medical Research
-	H01.770.644.193	Community-Based Participatory Research

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	H01.770.644.205	Data Anonymization
-	H01.770.644.217	Dual Use Research
-	H01.770.644.241	Empirical Research
-	H01.770.644.241.424	Grounded Theory
-	H01.770.644.241.850	Qualitative Research
-	H01.770.644.241.850.750	Hermeneutics
-	H01.770.644.333	Operations Research
-	H01.770.644.364	Parapsychology
-	H01.770.644.487	Peer Review, Research
New Heading	<b>H01.770.644.608</b>	<b>Public Health Systems Research</b>
-	H01.770.644.728	Research Design
-	H01.770.644.864	Research Report
-	H01.770.644.932	Social Validity, Research
-	H01.770.787	Systems Integration
-	H01.770.808	Systems Theory
-	H02	Health Occupations
-	H02.004	Acupuncture
-	H02.010	Allied Health Occupations
-	H02.010.150	Audiology
-	H02.010.450	Medical Laboratory Science
-	H02.010.500	Occupational Therapy
-	H02.010.625	Physical Therapy Specialty
-	H02.010.750	Speech-Language Pathology
-	H02.010.800	Technology, Dental
-	H02.010.850	Technology, Radiologic
-	H02.010.850.700	Teleradiology
-	H02.070	Biomedical Engineering
-	H02.110	Chiropractic
-	H02.163	Dentistry
-	H02.163.090	Dental Research
-	H02.163.180	Dentistry, Operative
-	H02.163.232	Evidence-Based Dentistry
-	H02.163.285	Forensic Dentistry
-	H02.163.342	General Practice, Dental
-	H02.163.394	Geriatric Dentistry

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	H02.163.592	Military Dentistry
-	H02.163.640	Occupational Dentistry
-	H02.163.670	Oral Medicine
-	H02.163.721	Preventive Dentistry
-	H02.163.809	School Dentistry
-	H02.163.876	Specialties, Dental
-	H02.163.876.213	Endodontics
-	H02.163.876.439	Orthodontics
-	H02.163.876.506	Pathology, Oral
-	H02.163.876.600	Pediatric Dentistry
-	H02.163.876.623	Periodontics
-	H02.163.876.708	Prosthodontics
-	H02.163.876.770	Public Health Dentistry
-	H02.163.876.770.357	Community Dentistry
-	H02.163.876.886	Surgery, Oral
-	H02.163.876.886.500	Orthognathic Surgery
-	H02.229	Environmental Health
-	H02.229.333	Health Physics
-	H02.229.656	Sanitary Engineering
-	H02.229.782	Sanitation
-	H02.249	Evidence-Based Practice
-	H02.249.500	Evidence-Based Dentistry
-	H02.249.750	Evidence-Based Medicine
-	H02.249.750.500	Evidence-Based Emergency Medicine
-	H02.249.875	Evidence-Based Nursing
-	H02.269	Health Services Administration
-	H02.309	Hospital Administration
-	H02.385	Medical Illustration
-	H02.403	Medicine
-	H02.403.014	Adolescent Medicine
-	H02.403.029	Aerospace Medicine
-	H02.403.044	Allergy and Immunology
-	H02.403.044.500	Immunochemistry
Old Tree	<b>H02.403.059</b>	<b>Andrology</b>
-	H02.403.066	Anesthesiology
-	H02.403.074	Bariatric Medicine

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	H02.403.090	Behavioral Medicine
-	H02.403.200	Clinical Medicine
-	H02.403.200.400	Evidence-Based Medicine
-	H02.403.200.400.500	Evidence-Based Emergency Medicine
-	H02.403.200.700	Precision Medicine
-	H02.403.220	Community Medicine
-	H02.403.225	Dermatology
-	H02.403.230	Disaster Medicine
-	H02.403.250	Emergency Medicine
New Heading	<b>H02.403.250.500</b>	<b>Pediatric Emergency Medicine</b>
-	H02.403.330	Forensic Medicine
-	H02.403.330.149	Forensic Genetics
-	H02.403.330.300	Forensic Pathology
-	H02.403.340	General Practice
-	H02.403.340.500	Family Practice
-	H02.403.350	Genetics, Medical
-	H02.403.352	Geography, Medical
-	H02.403.352.500	Topography, Medical
-	H02.403.355	Geriatrics
-	H02.403.371	Global Health
-	H02.403.377	Hospital Medicine
-	H02.403.400	Integrative Medicine
-	H02.403.429	Internal Medicine
-	H02.403.429.163	Cardiology
-	H02.403.429.163.300	Cardiac Electrophysiology
-	H02.403.429.323	Endocrinology
-	H02.403.429.405	Gastroenterology
-	H02.403.429.445	Hematology
-	H02.403.429.445.500	Transfusion Medicine
-	H02.403.429.480	Infectious Disease Medicine
-	H02.403.429.515	Medical Oncology
-	H02.403.429.515.500	Radiation Oncology
New Heading	<b>H02.403.429.515.750</b>	<b>Surgical Oncology</b>
-	H02.403.429.580	Nephrology

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	H02.403.429.675	Pulmonary Medicine
-	H02.403.429.730	Rheumatology
-	H02.403.429.865	Sleep Medicine Specialty
-	H02.403.500	Military Medicine
-	H02.403.530	Molecular Medicine
-	H02.403.560	Naval Medicine
-	H02.403.560.508	Submarine Medicine
-	H02.403.600	Neurology
-	H02.403.600.250	Neuropathology
-	H02.403.600.500	Neurotology
-	H02.403.640	Osteopathic Medicine
-	H02.403.645	Palliative Medicine
-	H02.403.650	Pathology
-	H02.403.650.249	Forensic Pathology
-	H02.403.650.375	Neuropathology
-	H02.403.650.500	Pathology, Clinical
-	H02.403.650.505	Pathology, Molecular
-	H02.403.650.510	Pathology, Surgical
-	H02.403.650.600	Telepathology
-	H02.403.670	Pediatrics
-	H02.403.670.400	Neonatology
New Heading	<b>H02.403.670.450</b>	<b>Pediatric Emergency Medicine</b>
-	H02.403.670.500	Perinatology
-	H02.403.680	Physical and Rehabilitation Medicine
-	H02.403.680.600	Rehabilitation
New Heading	<b>H02.403.680.600.250</b>	<b>Cardiac Rehabilitation</b>
-	H02.403.680.600.500	Correction of Hearing Impairment
-	H02.403.680.600.750	Neurological Rehabilitation
New Heading	<b>H02.403.680.600.750.500</b>	<b>Stroke Rehabilitation</b>
-	H02.403.680.600.875	Psychiatric Rehabilitation
-	H02.403.680.600.937	Telerehabilitation
-	H02.403.690	Psychiatry
-	H02.403.690.080	Adolescent Psychiatry
-	H02.403.690.100	Biological Psychiatry



## MeSH Tree Changes for 2017

Type	Tree - heading	
-	H02.403.690.130	Child Psychiatry
-	H02.403.690.150	Community Psychiatry
-	H02.403.690.150.580	Preventive Psychiatry
-	H02.403.690.208	Forensic Psychiatry
-	H02.403.690.260	Geriatric Psychiatry
-	H02.403.690.508	Military Psychiatry
-	H02.403.690.754	Neuropsychiatry
-	H02.403.720	Public Health
-	H02.403.720.500	Epidemiology
-	H02.403.720.500.300	Molecular Epidemiology
-	H02.403.720.500.650	Pharmacoepidemiology
-	H02.403.720.750	Preventive Medicine
-	H02.403.720.750.250	Environmental Medicine
-	H02.403.720.750.510	Occupational Medicine
-	H02.403.720.750.550	Preventive Psychiatry
-	H02.403.740	Radiology
-	H02.403.740.500	Nuclear Medicine
-	H02.403.740.650	Radiation Oncology
-	H02.403.740.675	Radiology, Interventional
-	H02.403.750	Regenerative Medicine
-	H02.403.763	Reproductive Medicine
New Tree	<a href="#">H02.403.763.500</a>	<a href="#">Andrology</a>
New Tree	<a href="#">H02.403.763.750</a>	<a href="#">Gynecology</a>
-	H02.403.800	Social Medicine
-	H02.403.810	Specialties, Surgical
-	H02.403.810.208	Colorectal Surgery
-	H02.403.810.300	General Surgery
-	H02.403.810.310	Gynecology
-	H02.403.810.425	Neurosurgery
-	H02.403.810.450	Obstetrics
-	H02.403.810.468	Ophthalmology
-	H02.403.810.481	Orthognathic Surgery
-	H02.403.810.494	Orthopedics
-	H02.403.810.526	Otolaryngology

## MeSH Tree Changes for 2017

Type	Tree - heading
-	H02.403.810.526.500                      Neurotology
-	H02.403.810.788                            Surgery, Plastic
New Heading	<b>H02.403.810.796                            Surgical Oncology</b>
-	H02.403.810.803                            Thoracic Surgery
-	H02.403.810.850                            Traumatology
-	H02.403.810.860                            Urology
-	H02.403.830                                Sports Medicine
-	H02.403.830.500                            Sports Nutritional Sciences
-	H02.403.830.750                            Veterinary Sports Medicine
-	H02.403.840                                Telemedicine
-	H02.403.840.600                            Telepathology
-	H02.403.840.700                            Teleradiology
-	H02.403.840.850                            Telerehabilitation
-	H02.403.845                                Theranostic Nanomedicine
-	H02.403.850                                Travel Medicine
-	H02.403.879                                Tropical Medicine
-	H02.403.909                                Venereology
-	H02.403.959                                Wilderness Medicine
-	H02.438                                      Mortuary Practice
-	H02.478                                      Nursing
-	H02.478.197                                Evidence-Based Nursing
-	H02.478.395                                Nursing Research
-	H02.478.395.234                            Clinical Nursing Research
-	H02.478.395.385                            Nursing Administration Research
-	H02.478.395.413                            Nursing Education Research
-	H02.478.395.432                            Nursing Evaluation Research
-	H02.478.395.634                            Nursing Methodology Research
-	H02.478.408                                Nursing Theory
-	H02.478.676                                Specialties, Nursing
-	H02.478.676.074                            Advanced Practice Nursing
-	H02.478.676.112                            Cardiovascular Nursing
-	H02.478.676.150                            Community Health Nursing
-	H02.478.676.150.500                      Home Health Nursing
-	H02.478.676.150.750                      Parish Nursing
-	H02.478.676.175                            Critical Care Nursing

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	H02.478.676.200	Emergency Nursing
-	H02.478.676.218	Family Nursing
-	H02.478.676.227	Forensic Nursing
-	H02.478.676.236	Geriatric Nursing
-	H02.478.676.313	Holistic Nursing
-	H02.478.676.350	Hospice and Palliative Care Nursing
-	H02.478.676.390	Maternal-Child Nursing
-	H02.478.676.390.600	Neonatal Nursing
-	H02.478.676.403	Medical-Surgical Nursing
-	H02.478.676.416	Midwifery
-	H02.478.676.458	Military Nursing
-	H02.478.676.514	Nephrology Nursing
-	H02.478.676.542	Neuroscience Nursing
-	H02.478.676.570	Obstetric Nursing
-	H02.478.676.590	Occupational Health Nursing
-	H02.478.676.605	Oncology Nursing
-	H02.478.676.615	Orthopedic Nursing
-	H02.478.676.631	Pediatric Nursing
-	H02.478.676.631.600	Neonatal Nursing
-	H02.478.676.650	Perioperative Nursing
-	H02.478.676.650.625	Operating Room Nursing
-	H02.478.676.650.650	Postanesthesia Nursing
-	H02.478.676.710	Psychiatric Nursing
-	H02.478.676.755	Public Health Nursing
New Heading	<b>H02.478.676.772</b>	<b>Radiologic and Imaging Nursing</b>
-	H02.478.676.789	Rehabilitation Nursing
-	H02.478.676.806	Rural Nursing
-	H02.478.676.824	School Nursing
-	H02.478.676.920	Transcultural Nursing
-	H02.495	Nursing, Practical
-	H02.533	Nutritional Sciences
-	H02.533.124	Animal Nutrition Sciences
-	H02.533.252	Child Nutrition Sciences
-	H02.533.290	Dietetics
-	H02.533.468	Nutrigenomics

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	H02.533.645	Sports Nutritional Sciences
-	H02.553	Optometry
-	H02.573	Orthoptics
-	H02.628	Pharmacology
-	H02.628.049	Biopharmaceutics
-	H02.628.100	Ethnopharmacology
-	H02.628.190	Herbal Medicine
-	H02.628.280	Neuropharmacology
-	H02.628.413	Pharmacoepidemiology
-	H02.628.479	Pharmacogenetics
-	H02.628.512	Pharmacology, Clinical
-	H02.628.546	Psychopharmacology
-	H02.646	Pharmacy
New Heading	<b>H02.646.250</b>	<b>Nuclear Pharmacy</b>
New Heading	<b>H02.646.375</b>	<b>Pharmacy Research</b>
Old Tree	<b>H02.646.500</b>	<b>Technology, Pharmaceutical</b>
-	H02.696	Podiatry
-	H02.720	Psychology, Medical
-	H02.781	Serology
-	H02.790	Sociology, Medical
-	H02.811	Specialization
-	H02.884	Toxicology
-	H02.884.211	Ecotoxicology
-	H02.884.424	Forensic Toxicology
-	H02.884.850	Toxicogenetics
-	H02.956	Veterinary Medicine
-	H02.956.465	Pathology, Veterinary
-	H02.956.692	Surgery, Veterinary
-	H02.956.858	Veterinary Service, Military
-	H02.956.929	Veterinary Sports Medicine
-	I01	Social Sciences
-	I01.076	Anthropology
-	I01.076.201	Anthropology, Cultural
-	I01.076.201.208	Archaeology

## MeSH Tree Changes for 2017

Type	Tree - heading
-	I01.076.201.450 Culture
-	I01.076.201.450.050 Acculturation
-	I01.076.201.450.170 Ceremonial Behavior
-	I01.076.201.450.199 Circumcision, Female
-	I01.076.201.450.226 Civilization
-	I01.076.201.450.226.200 Arab World
-	I01.076.201.450.226.800 Western World
-	I01.076.201.450.226.800.250 Greek World
-	I01.076.201.450.226.800.500 Roman World
-	I01.076.201.450.281 Cross-Cultural Comparison
-	I01.076.201.450.324 Cultural Characteristics
-	I01.076.201.450.350 Cultural Diversity
-	I01.076.201.450.370 Cultural Evolution
-	I01.076.201.450.457 Ethnology
-	I01.076.201.450.515 Folklore
-	I01.076.201.450.515.500 Legendary Creatures
-	I01.076.201.450.550 Funeral Rites
-	I01.076.201.450.550.150 Burial
-	I01.076.201.450.550.150.500 Cemeteries
-	I01.076.201.450.550.175 Cremation
-	I01.076.201.450.550.250 Embalming
-	I01.076.201.450.560 Human Body
-	I01.076.201.450.654 Medicine, Traditional
-	I01.076.201.450.654.252 Ethnopharmacology
-	I01.076.201.450.654.505 Medicine, African Traditional
-	I01.076.201.450.654.510 Medicine, Arabic
-	I01.076.201.450.654.510.500 Medicine, Unani
-	I01.076.201.450.654.515 Medicine, Ayurvedic
-	I01.076.201.450.654.558 Medicine, East Asian Traditional
-	I01.076.201.450.654.558.520 Medicine, Chinese Traditional
-	I01.076.201.450.654.558.520.300 Qi
-	I01.076.201.450.654.558.520.300.500 Meridians
-	I01.076.201.450.654.558.520.967 Yin-Yang
-	I01.076.201.450.654.558.600 Medicine, Kampo
-	I01.076.201.450.654.558.700 Medicine, Korean Traditional
-	I01.076.201.450.654.558.750 Medicine, Mongolian Traditional

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	<b>I01.076.201.450.654.558.800</b> <span style="float: right;"><b>Medicine, Tibetan Traditional</b></span>
New Tree	<b>I01.076.201.450.654.558.875</b> <span style="float: right;"><b>Medicine, Tibetan Traditional</b></span>
-	I01.076.201.450.654.830 <span style="float: right;">Shamanism</span>
New Tree	<b>I01.076.201.450.776</b> <span style="float: right;"><b>Social Change</b></span>
-	I01.076.201.450.897 <span style="float: right;">Superstitions</span>
-	I01.076.201.450.897.439 <span style="float: right;">Magic</span>
-	I01.076.201.450.897.439.925 <span style="float: right;">Witchcraft</span>
-	I01.076.201.450.948 <span style="float: right;">Taboo</span>
-	I01.076.284 <span style="float: right;">Anthropology, Medical</span>
-	I01.076.368 <span style="float: right;">Anthropology, Physical</span>
-	I01.076.368.239 <span style="float: right;">Craniology</span>
-	I01.076.368.400 <span style="float: right;">Forensic Anthropology</span>
-	I01.076.368.584 <span style="float: right;">Paleontology</span>
-	I01.076.368.584.311 <span style="float: right;">Fossils</span>
-	I01.076.368.584.500 <span style="float: right;">Mummies</span>
-	I01.076.368.584.542 <span style="float: right;">Origin of Life</span>
-	I01.076.368.584.583 <span style="float: right;">Paleodontology</span>
-	I01.076.368.584.709 <span style="float: right;">Paleopathology</span>
New Heading	<b>I01.076.368.792</b> <span style="float: right;"><b>Body Remains</b></span>
-	I01.198 <span style="float: right;">Criminology</span>
-	I01.198.240 <span style="float: right;">Crime</span>
-	I01.198.240.089 <span style="float: right;">Abortion, Criminal</span>
-	I01.198.240.194 <span style="float: right;">Corpse Dismemberment</span>
-	I01.198.240.240 <span style="float: right;">Defamation</span>
-	I01.198.240.247 <span style="float: right;">Drug Trafficking</span>
-	I01.198.240.300 <span style="float: right;">Fraud</span>
-	I01.198.240.300.249 <span style="float: right;">Identity Theft</span>
-	I01.198.240.300.249.500 <span style="float: right;">Medical Identity Theft</span>
-	I01.198.240.385 <span style="float: right;">Grave Robbing</span>
-	I01.198.240.470 <span style="float: right;">Homicide</span>
-	I01.198.240.470.572 <span style="float: right;">Infanticide</span>
-	I01.198.240.609 <span style="float: right;">Prescription Drug Diversion</span>
-	I01.198.240.748 <span style="float: right;">Sex Offenses</span>
-	I01.198.240.748.300 <span style="float: right;">Child Abuse, Sexual</span>

## MeSH Tree Changes for 2017

Type	Tree - heading
-	I01.198.240.748.470 Human Trafficking
-	I01.198.240.748.640 Rape
-	I01.198.240.810 Theft
-	I01.198.240.856 Violence
-	I01.198.240.856.350 Domestic Violence
-	I01.198.240.856.350.250 Child Abuse
-	I01.198.240.856.350.250.255 Child Abuse, Sexual
-	I01.198.240.856.350.390 Elder Abuse
-	I01.198.240.856.350.750 Spouse Abuse
-	I01.198.240.856.575 Intimate Partner Violence
-	I01.198.240.856.575.500 Spouse Abuse
-	I01.198.240.856.688 Physical Abuse
New Tree	<a href="#">I01.198.240.856.744</a> Rape
-	I01.198.240.856.800 Terrorism
-	I01.198.240.856.800.100 Bioterrorism
-	I01.198.240.856.800.325 Chemical Terrorism
-	I01.198.240.856.800.537 Mass Casualty Incidents
-	I01.198.240.856.800.750 September 11 Terrorist Attacks
-	I01.198.240.856.825 Torture
-	I01.198.240.856.912 Workplace Violence
-	I01.198.240.903 War Crimes
-	I01.198.240.903.074 Ethnic Cleansing
-	I01.198.240.903.149 Genocide
-	I01.198.240.903.149.500 Holocaust
-	I01.198.290 Criminal Law
-	I01.198.780 Forensic Sciences
-	I01.198.780.750 Forensic Anthropology
-	I01.198.780.760 Forensic Ballistics
-	I01.198.780.875 Forensic Dentistry
-	I01.198.780.937 Forensic Medicine
-	I01.198.780.937.120 Autopsy
-	I01.198.780.937.206 Blood Stains
-	I01.198.780.937.343 Dermatoglyphics
-	I01.198.780.937.375 DNA Fingerprinting
-	I01.198.780.937.375.500 DNA Contamination

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	I01.198.780.937.422	Exhumation
-	I01.198.780.937.441	Forensic Genetics
-	I01.198.780.937.460	Forensic Pathology
-	I01.198.780.937.469	Forensic Psychiatry
-	I01.198.780.937.469.528	Insanity Defense
-	I01.198.780.937.619	Lie Detection
-	I01.198.780.937.766	Paternity
New Heading	<b>I01.198.780.937.883</b>	<b>Body Remains</b>
-	I01.198.780.952	Forensic Nursing
-	I01.198.780.968	Forensic Toxicology
-	I01.240	Demography
-	I01.240.050	Age Distribution
-	I01.240.205	Censuses
-	I01.240.361	Family Characteristics
-	I01.240.361.150	Birth Intervals
-	I01.240.361.160	Birth Order
-	I01.240.361.500	Marital Status
-	I01.240.361.500.300	Divorce
-	I01.240.361.500.500	Marriage
-	I01.240.361.500.725	Single Person
-	I01.240.361.500.725.700	Single Parent
-	I01.240.361.500.862	Widowhood
-	I01.240.425	Health Status
-	I01.240.425.350	Geriatric Assessment
-	I01.240.425.675	Health Status Disparities
-	I01.240.600	Population Dynamics
-	I01.240.600.400	Health Transition
-	I01.240.600.525	Human Migration
-	I01.240.600.525.500	Emigration and Immigration
-	I01.240.600.650	Population Control
-	I01.240.600.660	Population Growth
-	I01.240.800	Sex Distribution
-	I01.240.800.815	Sex Ratio
-	I01.261	Economics
-	I01.261.100	Capitalism



## MeSH Tree Changes for 2017

Type	Tree - heading	
-	I01.261.262	Economic Development
-	I01.261.262.500	Industrial Development
-	I01.261.425	Economic Recession
New Heading	<b>I01.261.506</b>	<b>Expropriation</b>
-	I01.261.587	Gross Domestic Product
-	I01.261.750	Resource Allocation
-	I01.261.750.500	Health Care Rationing
-	I01.283	Environment Design
-	I01.320	Forecasting
-	I01.320.500	Population Forecast
-	I01.409	Government
-	I01.409.137	Federal Government
-	I01.409.418	Government Agencies
-	I01.409.418.500	Child Protective Services
-	I01.409.418.625	Federal Government
-	I01.409.418.750	United States Government Agencies
-	I01.409.418.750.099	Peace Corps
-	I01.409.418.750.199	United States Agency for International Development
-	I01.409.418.750.400	United States Department of Agriculture
-	I01.409.418.750.500	United States Department of Defense
-	I01.409.418.750.600	United States Dept. of Health and Human Services
-	I01.409.418.750.600.310	Centers for Medicare and Medicaid Services (U.S.)
Old Tree	<b>I01.409.418.750.600.480</b>	<b>National Institutes of Health (U.S.)</b>
Old Tree	<b>I01.409.418.750.600.480.150</b>	<b>National Cancer Institute (U.S.)</b>
Old Tree	<b>I01.409.418.750.600.480.200</b>	<b>National Eye Institute (U.S.)</b>
Old Tree	<b>I01.409.418.750.600.480.300</b>	<b>National Heart, Lung, and Blood Institute (U.S.)</b>
Old Tree	<b>I01.409.418.750.600.480.325 (U.S.)</b>	<b>National Human Genome Research Institute</b>
Old Tree	<b>I01.409.418.750.600.480.400 Diseases (U.S.)</b>	<b>National Institute of Allergy and Infectious</b>
Old Tree	<b>I01.409.418.750.600.480.425 and Skin Diseases (U.S.)</b>	<b>National Institute of Arthritis and Musculoskeletal</b>
Old Tree	<b>I01.409.418.750.600.480.435 Bioengineering (U.S.)</b>	<b>National Institute of Biomedical Imaging and</b>
Old Tree	<b>I01.409.418.750.600.480.440 Development (U.S.)</b>	<b>National Institute of Child Health and Human</b>
Old Tree	<b>I01.409.418.750.600.480.445</b>	<b>National Institute of Dental and Craniofacial</b>

## MeSH Tree Changes for 2017

Type	Tree - heading	
	<b>Research (U.S.)</b>	
Old Tree	<b>I01.409.418.750.600.480.450 Kidney Diseases (U.S.)</b>	National Institute of Diabetes and Digestive and
Old Tree	<b>I01.409.418.750.600.480.455 Sciences (U.S.)</b>	National Institute of Environmental Health
Old Tree	<b>I01.409.418.750.600.480.457 (U.S.)</b>	National Institute of General Medical Sciences
Old Tree	<b>I01.409.418.750.600.480.460</b>	National Institute of Mental Health (U.S.)
Old Tree	<b>I01.409.418.750.600.480.461 Stroke</b>	National Institute of Neurological Disorders and
Old Tree	<b>I01.409.418.750.600.480.462</b>	National Institute of Nursing Research (U.S.)
Old Tree	<b>I01.409.418.750.600.480.465</b>	National Institute on Aging (U.S.)
Old Tree	<b>I01.409.418.750.600.480.467 Alcoholism (U.S.)</b>	National Institute on Alcohol Abuse and
Old Tree	<b>I01.409.418.750.600.480.470 Communication Disorders (U.S.)</b>	National Institute on Deafness and Other
Old Tree	<b>I01.409.418.750.600.480.485</b>	National Institute on Drug Abuse (U.S.)
Old Tree	<b>I01.409.418.750.600.480.490</b>	National Library of Medicine (U.S.)
-	<b>I01.409.418.750.600.650</b>	United States Public Health Service
New Tree	<b>I01.409.418.750.600.650.200 (U.S.)</b>	Centers for Disease Control and Prevention
New Tree	<b>I01.409.418.750.600.650.200.260</b>	National Center for Health Statistics (U.S.)
New Tree	<b>I01.409.418.750.600.650.200.520 and Health (U.S.)</b>	National Institute for Occupational Safety
Old Tree	<b>I01.409.418.750.600.650.225 (U.S.)</b>	Centers for Disease Control and Prevention
Old Tree	<b>I01.409.418.750.600.650.225.520 and Health (U.S.)</b>	National Institute for Occupational Safety
-	<b>I01.409.418.750.600.650.400 (U.S.)</b>	National Center for Health Care Technology
Old Tree	<b>I01.409.418.750.600.650.425</b>	National Center for Health Statistics (U.S.)
New Tree	<b>I01.409.418.750.600.650.496</b>	National Institutes of Health (U.S.)
New Tree	<b>I01.409.418.750.600.650.496.150</b>	National Cancer Institute (U.S.)
New Heading	<b>I01.409.418.750.600.650.496.175 Translational Sciences (U.S.)</b>	National Center for Advancing
New Heading	<b>I01.409.418.750.600.650.496.188 Integrative Health (U.S.)</b>	National Center for Complementary and
New Tree	<b>I01.409.418.750.600.650.496.200</b>	National Eye Institute (U.S.)

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	<a href="#">I01.409.418.750.600.650.496.300 (U.S.)</a>	<a href="#">National Heart, Lung, and Blood Institute</a>
New Tree	<a href="#">I01.409.418.750.600.650.496.325 (U.S.)</a>	<a href="#">National Human Genome Research Institute</a>
New Tree	<a href="#">I01.409.418.750.600.650.496.400 Diseases (U.S.)</a>	<a href="#">National Institute of Allergy and Infectious</a>
New Tree	<a href="#">I01.409.418.750.600.650.496.425 Musculoskeletal and Skin Diseases (U.S.)</a>	<a href="#">National Institute of Arthritis and</a>
New Tree	<a href="#">I01.409.418.750.600.650.496.435 Bioengineering (U.S.)</a>	<a href="#">National Institute of Biomedical Imaging and</a>
New Tree	<a href="#">I01.409.418.750.600.650.496.440 Development (U.S.)</a>	<a href="#">National Institute of Child Health and Human</a>
New Tree	<a href="#">I01.409.418.750.600.650.496.445 Research (U.S.)</a>	<a href="#">National Institute of Dental and Craniofacial</a>
New Tree	<a href="#">I01.409.418.750.600.650.496.450 and Kidney Diseases (U.S.)</a>	<a href="#">National Institute of Diabetes and Digestive</a>
New Tree	<a href="#">I01.409.418.750.600.650.496.455 Sciences (U.S.)</a>	<a href="#">National Institute of Environmental Health</a>
New Tree	<a href="#">I01.409.418.750.600.650.496.457 Sciences (U.S.)</a>	<a href="#">National Institute of General Medical</a>
New Tree	<a href="#">I01.409.418.750.600.650.496.460</a>	<a href="#">National Institute of Mental Health (U.S.)</a>
New Tree	<a href="#">I01.409.418.750.600.650.496.461 and Stroke</a>	<a href="#">National Institute of Neurological Disorders</a>
New Tree	<a href="#">I01.409.418.750.600.650.496.462</a>	<a href="#">National Institute of Nursing Research (U.S.)</a>
New Tree	<a href="#">I01.409.418.750.600.650.496.465</a>	<a href="#">National Institute on Aging (U.S.)</a>
New Tree	<a href="#">I01.409.418.750.600.650.496.467 Alcoholism (U.S.)</a>	<a href="#">National Institute on Alcohol Abuse and</a>
New Tree	<a href="#">I01.409.418.750.600.650.496.470 Communication Disorders (U.S.)</a>	<a href="#">National Institute on Deafness and Other</a>
New Tree	<a href="#">I01.409.418.750.600.650.496.485</a>	<a href="#">National Institute on Drug Abuse (U.S.)</a>
New Tree	<a href="#">I01.409.418.750.600.650.496.490</a>	<a href="#">National Library of Medicine (U.S.)</a>
-	<a href="#">I01.409.418.750.600.650.592 and Quality</a>	<a href="#">United States Agency for Healthcare Research</a>
-	<a href="#">I01.409.418.750.600.650.760</a>	<a href="#">United States Food and Drug Administration</a>
-	<a href="#">I01.409.418.750.600.650.790 Administration</a>	<a href="#">United States Health Resources and Services</a>
-	<a href="#">I01.409.418.750.600.650.790.525 (U.S.)</a>	<a href="#">National Health Planning Information Center</a>

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	I01.409.418.750.600.650.825	United States Indian Health Service
-	I01.409.418.750.600.650.912	United States Office of Research Integrity
-	I01.409.418.750.600.650.920 Health Services Administration	United States Substance Abuse and Mental
-	I01.409.418.750.650	United States Department of Homeland Security
-	I01.409.418.750.700	United States Department of Veterans Affairs
-	I01.409.418.750.937	United States Environmental Protection Agency
-	I01.409.418.750.968	United States Federal Trade Commission
-	I01.409.418.750.984 Administration	United States National Aeronautics and Space
-	I01.409.418.750.992 Administration	United States Occupational Safety and Health
-	I01.409.418.750.996	United States Office of Economic Opportunity
-	I01.409.418.750.997	United States Office of National Drug Control Policy
-	I01.409.418.750.998	United States Office of Technology Assessment
-	I01.409.418.750.999	United States Social Security Administration
-	I01.409.700	Local Government
-	I01.409.775	State Government
-	I01.451	Government Programs
-	I01.451.227	Civil Defense
-	I01.451.617	Postal Service
-	I01.615	Internationality
-	I01.615.500	International Cooperation
-	I01.615.500.125	African Union
-	I01.615.500.250	Developed Countries
-	I01.615.500.300	Developing Countries
-	I01.615.500.475	European Union
-	I01.615.500.650	International Educational Exchange
-	I01.615.500.750	Medical Missions, Official
-	I01.655	Policy
New Heading	<b>I01.655.125</b>	<b>Fiscal Policy</b>
-	I01.655.249	Smoke-Free Policy
-	I01.655.500	Social Control Policies
-	I01.655.500.550	Organizational Policy
-	I01.655.500.608	Public Policy
-	I01.655.500.608.180	Environmental Policy

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	I01.655.500.608.200	Family Planning Policy
-	I01.655.500.608.400	Health Policy
-	I01.655.500.608.400.285	Health Care Reform
-	I01.655.500.608.400.650	Nutrition Policy
-	I01.655.500.608.400.650.500	Recommended Dietary Allowances
-	I01.696	Political Systems
New Heading	<b>I01.696.050</b>	<b>Apartheid</b>
-	I01.696.100	Capitalism
-	I01.696.116	Colonialism
-	I01.696.232	Communism
-	I01.696.374	Democracy
-	I01.696.586	National Socialism
-	I01.696.798	Socialism
-	I01.738	Politics
-	I01.738.305	Diplomacy
-	I01.738.610	Lobbying
-	I01.791	Private Sector
New Heading	<b>I01.791.500</b>	<b>Informal Sector</b>
-	I01.795	Public Sector
-	I01.800	Quality of Life
-	I01.880	Sociology
-	I01.880.604	Social Control, Formal
-	I01.880.604.100	Animal Welfare
-	I01.880.604.100.049	Animal Care Committees
-	I01.880.604.100.100	Animal Rights
-	I01.880.604.160	Capital Punishment
-	I01.880.604.238	Censorship, Research
-	I01.880.604.316	Coercion
-	I01.880.604.394	Government Regulation
-	I01.880.604.473	Human Rights
-	I01.880.604.473.300	Child Advocacy
-	I01.880.604.473.352	Civil Rights
-	I01.880.604.473.352.500	Privacy
-	I01.880.604.473.352.500.030	Access to Information

## MeSH Tree Changes for 2017

Type	Tree - heading
-	I01.880.604.473.352.500.030.500 Patient Access to Records
-	I01.880.604.473.352.500.320 Genetic Privacy
-	I01.880.604.473.368 Consumer Advocacy
New Tree	<b>I01.880.604.473.374</b> <b>Feminism</b>
-	I01.880.604.473.380 Freedom
-	I01.880.604.473.380.500 Personal Autonomy
-	I01.880.604.473.650 Patient Rights
-	I01.880.604.473.650.500 Confidentiality
-	I01.880.604.473.650.500.320 Genetic Privacy
-	I01.880.604.473.650.718 Informed Consent
-	I01.880.604.473.650.718.500 Informed Consent By Minors
-	I01.880.604.473.650.827 Patient Access to Records
-	I01.880.604.473.650.952 Right to Die
-	I01.880.604.473.650.960 Therapeutic Misconception
-	I01.880.604.473.650.968 Treatment Refusal
New Heading	<b>I01.880.604.473.650.968.500</b> <b>Vaccination Refusal</b>
-	I01.880.604.473.675 Reproductive Rights
-	I01.880.604.473.700 Social Justice
-	I01.880.604.473.850 Women's Rights
-	I01.880.604.583 Jurisprudence
-	I01.880.604.583.020 Advance Directives
-	I01.880.604.583.020.500 Living Wills
-	I01.880.604.583.050 Compensation and Redress
-	I01.880.604.583.080 Confidentiality
-	I01.880.604.583.080.134 Disclosure
-	I01.880.604.583.080.134.300 Mandatory Reporting
-	I01.880.604.583.080.134.399 Parental Notification
-	I01.880.604.583.080.134.800 Truth Disclosure
-	I01.880.604.583.080.134.800.200 Duty to Warn
-	I01.880.604.583.080.134.900 Whistleblowing
-	I01.880.604.583.080.270 Duty to Warn
-	I01.880.604.583.080.320 Genetic Privacy
-	I01.880.604.583.080.660 Personally Identifiable Information
-	I01.880.604.583.090 Contracts

## MeSH Tree Changes for 2017

Type	Tree - heading
-	I01.880.604.583.100 Criminal Law
-	I01.880.604.583.166 Duty to Recontact
-	I01.880.604.583.232 Expert Testimony
-	I01.880.604.583.310 Forensic Psychiatry
-	I01.880.604.583.310.528 Insanity Defense
-	I01.880.604.583.427 Informed Consent
-	I01.880.604.583.427.134 Consent Forms
-	I01.880.604.583.427.384 Informed Consent By Minors
-	I01.880.604.583.427.635 Third-Party Consent
-	I01.880.604.583.427.635.500 Parental Consent
-	I01.880.604.583.458 Intellectual Property
-	I01.880.604.583.458.300 Copyright
-	I01.880.604.583.458.650 Patents as Topic
-	I01.880.604.583.466 International Law
-	I01.880.604.583.474 Judicial Role
-	I01.880.604.583.482 Legal Services
-	I01.880.604.583.490 Liability, Legal
-	I01.880.604.583.524 Malpractice
-	I01.880.604.583.524.300 Defensive Medicine
-	I01.880.604.583.524.528 Professional Impairment
-	I01.880.604.583.524.528.500 Physician Impairment
-	I01.880.604.583.527 Mandatory Reporting
-	I01.880.604.583.530 Mental Competency
-	I01.880.604.583.594 Ownership
-	I01.880.604.583.659 Presumed Consent
-	I01.880.604.583.789 Resuscitation Orders
-	I01.880.604.583.858 Supreme Court Decisions
-	I01.880.604.583.927 Wills
-	I01.880.604.583.927.500 Living Wills
-	I01.880.604.583.963 Wrongful Life
-	I01.880.604.594 Law Enforcement
-	I01.880.604.605 Legislation, Drug
-	I01.880.604.605.250 Drug and Narcotic Control
-	I01.880.604.605.250.250 Drug Approval
-	I01.880.604.605.250.625 Drug Recalls
-	I01.880.604.605.250.812 Safety-Based Drug Withdrawals

## MeSH Tree Changes for 2017

Type	Tree - heading
-	I01.880.604.622 Mandatory Programs
-	I01.880.604.622.249 Mandatory Reporting
-	I01.880.604.622.500 Mandatory Testing
-	I01.880.604.631 Patient Advocacy
-	I01.880.604.640 Peer Review
-	I01.880.604.640.690 Peer Review, Health Care
-	I01.880.604.640.700 Peer Review, Research
-	I01.880.604.787 Prisons
-	I01.880.604.787.358 Concentration Camps
-	I01.880.604.825 Social Control Policies
-	I01.880.604.825.550 Organizational Policy
-	I01.880.604.825.608 Public Policy
-	I01.880.604.825.608.180 Environmental Policy
-	I01.880.604.825.608.200 Family Planning Policy
-	I01.880.604.825.608.400 Health Policy
-	I01.880.604.825.608.400.285 Health Care Reform
-	I01.880.604.825.608.400.650 Nutrition Policy
-	I01.880.604.825.608.400.650.500 Recommended Dietary Allowances
-	I01.880.630 Social Control, Informal
-	I01.880.630.099 Behavior Control
-	I01.880.630.200 Coercion
-	I01.880.630.548 Public Opinion
-	I01.880.630.716 Punishment
-	I01.880.709 Social Planning
-	I01.880.709.202 City Planning
-	I01.880.709.359 Environment Design
-	I01.880.709.876 Urban Renewal
-	I01.880.735 Social Problems
-	I01.880.735.070 Bullying
-	I01.880.735.140 Civil Disorders
-	I01.880.735.140.652 Riots
-	I01.880.735.191 Crime
-	I01.880.735.191.051 Defamation
-	I01.880.735.191.101 Drug Trafficking
-	I01.880.735.191.152 Organ Trafficking
-	I01.880.735.191.204 Prescription Drug Diversion



## MeSH Tree Changes for 2017

Type	Tree - heading	
-	I01.880.735.191.410	Theft
-	I01.880.735.223	Dangerous Behavior
-	I01.880.735.223.250	Driving Under the Influence
-	I01.880.735.223.500	Stalking
-	I01.880.735.255	Divorce
-	I01.880.735.265	Doping in Sports
-	I01.880.735.344	Homicide
-	I01.880.735.344.500	Euthanasia
-	I01.880.735.344.500.249	Euthanasia, Active
-	I01.880.735.344.500.249.200	Euthanasia, Active, Voluntary
-	I01.880.735.384	Human Rights Abuses
-	I01.880.735.384.399	Human Trafficking
-	I01.880.735.384.599	Organ Trafficking
-	I01.880.735.384.800	Slavery
-	I01.880.735.424	Illegitimacy
-	I01.880.735.442	Incest
-	I01.880.735.479	Juvenile Delinquency
-	I01.880.735.580	Needle Sharing
-	I01.880.735.607	Parental Death
-	I01.880.735.607.500	Maternal Death
-	I01.880.735.634	Poverty
-	I01.880.735.679	Prostitution
-	I01.880.735.679	Sex work
-	I01.880.735.728	Quackery
Old Tree	<b>I01.880.735.768</b>	<b>Race Relations</b>
Old Tree	<b>I01.880.735.768.500</b>	<b>Racism</b>
-	I01.880.735.779	Runaway Behavior
-	I01.880.735.784	Social Behavior Disorders
-	I01.880.735.784.350	General Adaptation Syndrome
New Heading	<b>I01.880.735.820</b>	<b>Social Segregation</b>
New Tree	<b>I01.880.735.820.500</b>	<b>Race Relations</b>
New Heading	<b>I01.880.735.820.500.125</b>	<b>Apartheid</b>
New Heading	<b>I01.880.735.820.500.250</b>	<b>Desegregation</b>

## MeSH Tree Changes for 2017

Type	Tree - heading
New Tree	<a href="#">I01.880.735.820.500.500</a> <b>Racism</b>
-	I01.880.735.856 Suicide
-	I01.880.735.856.149 Suicidal Ideation
-	I01.880.735.856.300 Suicide, Assisted
-	I01.880.735.856.600 Suicide, Attempted
-	I01.880.735.878 Underage Drinking
-	I01.880.735.900 Violence
-	I01.880.735.900.350 Domestic Violence
-	I01.880.735.900.350.250 Child Abuse
-	I01.880.735.900.350.250.255 Child Abuse, Sexual
-	I01.880.735.900.350.390 Elder Abuse
-	I01.880.735.900.350.750 Spouse Abuse
-	I01.880.735.900.575 Ethnic Violence
-	I01.880.735.900.575.500 Ethnic Cleansing
-	I01.880.735.900.688 Intimate Partner Violence
-	I01.880.735.900.688.500 Spouse Abuse
-	I01.880.735.900.744 Physical Abuse
New Tree	<a href="#">I01.880.735.900.772</a> <b>Rape</b>
-	I01.880.735.900.800 Terrorism
-	I01.880.735.900.800.100 Bioterrorism
-	I01.880.735.900.800.150 Chemical Terrorism
-	I01.880.735.900.800.575 Mass Casualty Incidents
-	I01.880.735.900.825 Torture
-	I01.880.735.900.869 Exposure to Violence
-	I01.880.735.900.912 Workplace Violence
-	I01.880.735.950 Warfare and Armed Conflicts
-	I01.880.735.950.250 Armed Conflicts
-	I01.880.735.950.250.094 Afghan Campaign 2001-
-	I01.880.735.950.250.188 American Civil War
-	I01.880.735.950.250.375 American Revolution
-	I01.880.735.950.250.469 Crimean War
-	I01.880.735.950.250.563 French Revolution
-	I01.880.735.950.250.657 Gulf War
-	I01.880.735.950.250.782 Iraq War, 2003-2011

## MeSH Tree Changes for 2017

Type	Tree - heading
-	I01.880.735.950.250.813 Korean War
-	I01.880.735.950.250.844 Russian-Japanese War
-	I01.880.735.950.250.860 September 11 Terrorist Attacks
-	I01.880.735.950.250.875 Spanish-American War, 1898
-	I01.880.735.950.250.937 Vietnam Conflict
-	I01.880.735.950.250.968 World War I
-	I01.880.735.950.250.984 World War II
-	I01.880.735.950.500 Warfare
-	I01.880.735.950.500.226 Biological Warfare
-	I01.880.735.950.500.226.100 Bioterrorism
-	I01.880.735.950.500.340 Chemical Warfare
-	I01.880.735.950.500.340.150 Chemical Terrorism
-	I01.880.735.950.500.600 Nuclear Warfare
-	I01.880.735.950.500.711 Psychological Warfare
-	I01.880.735.950.500.903 War Crimes
-	I01.880.735.950.500.903.074 Ethnic Cleansing
-	I01.880.735.950.500.903.149 Genocide
-	I01.880.735.950.500.903.149.500 Holocaust
-	I01.880.735.950.500.951 War Exposure
-	I01.880.735.950.500.951.500 War-Related Injuries
-	I01.880.787 Social Welfare
-	I01.880.787.190 Charities
-	I01.880.787.190.500 Almshouses
-	I01.880.787.293 Child Welfare
-	I01.880.787.293.350 Child Advocacy
-	I01.880.787.293.360 Child Care
-	I01.880.787.293.370 Child Custody
-	I01.880.787.354 Community Integration
-	I01.880.787.416 Foster Home Care
-	I01.880.787.539 Infant Welfare
-	I01.880.787.678 Maternal Welfare
-	I01.880.787.839 Relief Work
-	I01.880.787.839.500 Food Assistance
-	I01.880.792 Social Work
-	I01.880.792.410 Social Work, Psychiatric
-	I01.880.853 Sociological Factors

## MeSH Tree Changes for 2017

Type	Tree - heading
-	I01.880.853.100 Culture
-	I01.880.853.100.079 Acculturation
-	I01.880.853.100.257 Cross-Cultural Comparison
-	I01.880.853.100.329 Cultural Characteristics
-	I01.880.853.100.364 Cultural Competency
-	I01.880.853.100.400 Cultural Deprivation
-	I01.880.853.100.450 Cultural Diversity
-	I01.880.853.100.806 Psychosocial Deprivation
-	I01.880.853.150 Family
-	I01.880.853.150.140 Adoption
-	I01.880.853.150.281 Adult Children
-	I01.880.853.150.423 Family Characteristics
-	I01.880.853.150.423.500 Marital Status
-	I01.880.853.150.423.500.300 Divorce
-	I01.880.853.150.423.500.500 Marriage
-	I01.880.853.150.423.500.725 Single Person
-	I01.880.853.150.423.500.725.700 Single Parent
-	I01.880.853.150.423.500.862 Widowhood
-	I01.880.853.150.439 Family Relations
-	I01.880.853.150.452 Grandparents
-	I01.880.853.150.465 Illegitimacy
-	I01.880.853.150.482 Military Family
-	I01.880.853.150.500 Nuclear Family
-	I01.880.853.150.500.280 Only Child
-	I01.880.853.150.500.340 Parents
-	I01.880.853.150.500.340.210 Fathers
-	I01.880.853.150.500.340.270 Mothers
-	I01.880.853.150.500.340.785 Single Parent
-	I01.880.853.150.500.340.892 Surrogate Mothers
-	I01.880.853.150.500.505 Siblings
-	I01.880.853.150.500.670 Spouses
-	I01.880.853.150.750 Single-Parent Family
-	I01.880.853.200 Hierarchy, Social
-	I01.880.853.250 Medicalization
-	I01.880.853.300 Minority Groups
-	I01.880.853.350 Secularism

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	I01.880.853.375	Social Capital
-	I01.880.853.400	Social Change
-	I01.880.853.400.726	Urbanization
-	I01.880.853.450	Social Conditions
-	I01.880.853.450.283	Anomie
-	I01.880.853.500	Social Environment
-	I01.880.853.500.300	Community Networks
-	I01.880.853.500.600	Social Support
New Heading	<b>I01.880.853.500.600.500</b>	<b>Psychosocial Support Systems</b>
-	I01.880.853.748	Social Isolation
-	I01.880.853.748.435	Loneliness
-	I01.880.853.748.755	Social Alienation
-	I01.880.853.872	Social Marginalization
-	I01.880.853.903	Social Norms
-	I01.880.853.934	Socialization
-	I01.880.853.996	Socioeconomic Factors
-	I01.880.853.996.535	Poverty
-	I01.880.853.996.535.550	Poverty Areas
-	I01.880.853.996.755	Social Class
-	I01.880.853.996.755.673	Social Mobility
-	I01.880.859	Sociology, Medical
-	I01.880.866	Sociometric Techniques
New Heading	<b>I01.940</b>	<b>Work-Life Balance</b>
-	I02	Education
-	I02.158	Curriculum
-	I02.158.210	Competency-Based Education
-	I02.158.405	Interdisciplinary Studies
-	I02.158.600	Mainstreaming (Education)
-	I02.158.660	Problem-Based Learning
-	I02.195	Education, Distance
-	I02.233	Education, Nonprofessional
-	I02.233.213	Education, Special
-	I02.233.213.199	Education of Hearing Disabled
-	I02.233.213.400	Education of Intellectually Disabled

## MeSH Tree Changes for 2017

Type	Tree - heading
-	I02.233.213.500 Education of Visually Disabled
-	I02.233.213.600 Mainstreaming (Education)
-	I02.233.332 Health Education
-	I02.233.332.186 Consumer Health Information
-	I02.233.332.186.500 Health Literacy
-	I02.233.332.374 Health Education, Dental
-	I02.233.332.390 Health Fairs
-	I02.233.332.500 Patient Education as Topic
-	I02.233.332.500.500 Prenatal Education
-	I02.233.332.749 Sex Education
-	I02.233.332.874 Teach-Back Communication
-	I02.233.543 Physical Education and Training
Old Tree	I02.233.543.454 <b>Gymnastics</b>
-	I02.233.862 Vocational Education
-	I02.275 Education, Predental
-	I02.316 Education, Premedical
-	I02.358 Education, Professional
-	I02.358.105 Clinical Clerkship
-	I02.358.212 Education, Continuing
-	I02.358.212.249 Education, Dental, Continuing
-	I02.358.212.350 Education, Medical, Continuing
-	I02.358.212.450 Education, Nursing, Continuing
-	I02.358.212.550 Education, Pharmacy, Continuing
-	I02.358.212.590 Education, Professional, Retraining
-	I02.358.274 Education, Dental
-	I02.358.274.316 Education, Dental, Continuing
-	I02.358.274.482 Education, Dental, Graduate
-	I02.358.337 Education, Graduate
-	I02.358.337.249 Education, Dental, Graduate
-	I02.358.337.350 Education, Medical, Graduate
-	I02.358.337.450 Education, Nursing, Graduate
-	I02.358.337.550 Education, Pharmacy, Graduate
-	I02.358.399 Education, Medical
-	I02.358.399.250 Education, Medical, Continuing
-	I02.358.399.350 Education, Medical, Graduate
-	I02.358.399.450 Education, Medical, Undergraduate

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	I02.358.399.644	Internship and Residency
-	I02.358.399.644.500	Teaching Rounds
-	I02.358.462	Education, Nursing
-	I02.358.462.233	Education, Nursing, Associate
-	I02.358.462.316	Education, Nursing, Baccalaureate
-	I02.358.462.399	Education, Nursing, Continuing
-	I02.358.462.482	Education, Nursing, Diploma Programs
-	I02.358.462.565	Education, Nursing, Graduate
-	I02.358.462.612	Nursing Education Research
-	I02.358.525	Education, Pharmacy
-	I02.358.525.317	Education, Pharmacy, Continuing
-	I02.358.525.483	Education, Pharmacy, Graduate
-	I02.358.525.741	Pharmacy Residencies
-	I02.358.556	Education, Public Health Professional
-	I02.358.588	Education, Veterinary
-	I02.358.750	Internship, Nonmedical
-	I02.358.750.500	Pharmacy Residencies
New Heading	<b>I02.358.859</b>	<b>Mentoring</b>
-	I02.358.968	Preceptorship
New Heading	<b>I02.358.984</b>	<b>Teacher Training</b>
-	I02.399	Educational Measurement
-	I02.399.272	College Admission Test
-	I02.399.630	Professional Competence
-	I02.399.630.210	Clinical Competence
-	I02.399.750	School Admission Criteria
-	I02.399.780	Self-Evaluation Programs
-	I02.399.890	Test Taking Skills
-	I02.574	Inservice Training
-	I02.574.700	Staff Development
-	I02.581	International Educational Exchange
New Heading	<b>I02.588</b>	<b>Mentoring</b>
-	I02.594	Needs Assessment
-	I02.783	Schools
-	I02.783.330	Library Schools

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	I02.783.495	Schools, Health Occupations
-	I02.783.495.125	Area Health Education Centers
-	I02.783.495.481	Schools, Dental
-	I02.783.495.552	Schools, Medical
-	I02.783.495.623	Schools, Nursing
-	I02.783.495.694	Schools, Pharmacy
-	I02.783.495.765	Schools, Public Health
-	I02.783.495.836	Schools, Veterinary
-	I02.783.660	Schools, Nursery
-	I02.783.830	Universities
-	I02.903	Teaching
-	I02.903.080	Computer User Training
-	I02.903.302	Models, Educational
-	I02.903.565	Problem-Based Learning
-	I02.903.573	Programmed Instruction as Topic
-	I02.903.573.208	Computer-Assisted Instruction
-	I02.903.694	Remedial Teaching
-	I02.903.847	Simulation Training
New Heading	<b>I02.903.847.250</b>	<b>High Fidelity Simulation Training</b>
-	I02.903.847.500	Patient Simulation
New Heading	<b>I02.903.923</b>	<b>Study Guide as Topic</b>
-	I03	Human Activities
-	I03.050	Activities of Daily Living
-	I03.050.500	Independent Living
-	I03.050.750	Social Participation
New Heading	<b>I03.088</b>	<b>Anti-Vaccination Movement</b>
-	I03.125	Automobile Driving
-	I03.125.299	Automobile Driver Examination
New Heading	<b>I03.125.474</b>	<b>Distracted Driving</b>
-	I03.125.649	Driving Under the Influence
-	I03.287	Diet Fads
-	I03.350	Exercise
-	I03.350.061	Circuit-Based Exercise



## MeSH Tree Changes for 2017

Type	Tree - heading	
-	I03.350.124	Cool-Down Exercise
New Tree	<a href="#">I03.350.156</a>	<a href="#">Gymnastics</a>
New Heading	<b>I03.350.187</b>	<b>High-Intensity Interval Training</b>
-	I03.350.249	Muscle Stretching Exercises
-	I03.350.311	Physical Conditioning, Human
-	I03.350.374	Plyometric Exercise
-	I03.350.500	Resistance Training
-	I03.350.750	Running
-	I03.350.750.320	Jogging
-	I03.350.875	Swimming
-	I03.350.937	Walking
New Heading	<b>I03.350.937.500</b>	<b>Stair Climbing</b>
-	I03.350.968	Warm-Up Exercise
-	I03.450	Leisure Activities
-	I03.450.345	Holidays
-	I03.450.642	Recreation
-	I03.450.642.159	Camping
-	I03.450.642.287	Dancing
-	I03.450.642.378	Gardening
-	I03.450.642.469	Hobbies
-	I03.450.642.693	Play and Playthings
-	I03.450.642.693.465	Games, Recreational
-	I03.450.642.693.930	Video Games
-	I03.450.642.845	Sports
-	I03.450.642.845.054	Athletic Performance
New Heading	<b>I03.450.642.845.054.300</b>	<b>Cardiorespiratory Fitness</b>
-	I03.450.642.845.054.600	Physical Endurance
-	I03.450.642.845.054.800	Physical Fitness
New Heading	<b>I03.450.642.845.054.800.500</b>	<b>Cardiorespiratory Fitness</b>
-	I03.450.642.845.110	Baseball
-	I03.450.642.845.117	Basketball
-	I03.450.642.845.140	Bicycling

## MeSH Tree Changes for 2017

Type	Tree - heading
-	I03.450.642.845.210 Boxing
-	I03.450.642.845.300 Football
-	I03.450.642.845.400 Golf
-	I03.450.642.845.417 Gymnastics
-	I03.450.642.845.475 Hockey
-	I03.450.642.845.560 Martial Arts
-	I03.450.642.845.560.500 Tai Ji
-	I03.450.642.845.582 Mountaineering
-	I03.450.642.845.600 Racquet Sports
-	I03.450.642.845.600.900 Tennis
-	I03.450.642.845.605 Return to Sport
-	I03.450.642.845.610 Running
-	I03.450.642.845.610.320 Jogging
-	I03.450.642.845.700 Skating
-	I03.450.642.845.787 Snow Sports
-	I03.450.642.845.787.500 Skiing
-	I03.450.642.845.800 Soccer
-	I03.450.642.845.834 Sports for Persons with Disabilities
-	I03.450.642.845.869 Swimming
-	I03.450.642.845.869.110 Diving
-	I03.450.642.845.925 Track and Field
-	I03.450.642.845.932 Volleyball
-	I03.450.642.845.940 Walking
-	I03.450.642.845.950 Weight Lifting
-	I03.450.642.845.975 Wrestling
-	I03.450.642.845.987 Youth Sports
-	I03.450.642.922 Sunbathing
-	I03.450.769 Relaxation
-	I03.450.769.647 Rest
-	I03.548 Nudism
-	I03.702 Retirement
-	I03.784 Survival
-	I03.833 Temperance Movement
-	I03.883 Travel
-	I03.883.209 Air Travel
-	I03.883.420 Expeditions

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	I03.883.710	Medical Tourism
-	I03.946	Work
-	I03.946.449	Return to Work
-	I03.946.675	Work Performance
-	I03.946.900	Work Schedule Tolerance
New Heading	<b>I03.946.950</b>	<b>Work-Life Balance</b>
-	J01	Technology, Industry, and Agriculture
-	J01.040	Agriculture
-	J01.040.044	Agricultural Irrigation
-	J01.040.090	Animal Husbandry
-	J01.040.090.500	Animal Culling
-	J01.040.168	Aquaculture
-	J01.040.168.300	Fisheries
-	J01.040.207	Beekeeping
-	J01.040.227	Crop Production
New Heading	<b>J01.040.227.250</b>	<b>Biofortification</b>
-	J01.040.227.500	Plant Breeding
-	J01.040.237	Crop Protection
-	J01.040.246	Dairying
New Heading	<b>J01.040.330</b>	<b>Domestication</b>
-	J01.040.413	Gardening
New Heading	<b>J01.040.415</b>	<b>Gardens</b>
-	J01.040.581	Hydroponics
-	J01.040.895	Organic Agriculture
-	J01.040.947	Weed Control
-	J01.086	Architecture as Topic
-	J01.086.339	Facility Design and Construction
Old Tree	<b>J01.086.339.034</b>	<b>Airports</b>
-	J01.086.339.070	Architectural Accessibility
-	J01.086.339.115	Elevators and Escalators
-	J01.086.339.140	Floors and Floorcoverings
-	J01.086.339.250	Hospital Design and Construction
-	J01.086.339.290	Interior Design and Furnishings

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	J01.086.339.400	Location Directories and Signs
-	J01.086.339.445	Military Facilities
Old Tree	<b>J01.086.339.445.500</b>	<b>Hospitals, Military</b>
Old Tree	<b>J01.086.339.490</b>	<b>Parking Facilities</b>
-	J01.219	Commerce
-	J01.219.187	Commodification
-	J01.219.375	Entrepreneurship
-	J01.219.687	Marketing
-	J01.219.687.274	Advertising as Topic
-	J01.219.687.274.500	Direct-to-Consumer Advertising
-	J01.219.687.550	Marketing of Health Services
-	J01.219.687.750	Social Marketing
-	J01.219.843	Public-Private Sector Partnerships
-	J01.219.921	Small Business
-	J01.256	Conservation of Natural Resources
-	J01.256.078	Conservation of Energy Resources
-	J01.256.214	Endangered Species
-	J01.256.282	Environmental Restoration and Remediation
-	J01.256.350	Green Chemistry Technology
-	J01.256.675	Recycling
-	J01.293	Engineering
-	J01.293.069	Bioengineering
-	J01.293.069.124	Bioprinting
-	J01.293.069.249	Cell Engineering
-	J01.293.069.249.249	Metabolic Engineering
-	J01.293.069.249.500	Tissue Engineering
-	J01.293.069.500	Synthetic Biology
-	J01.293.140	Biomedical Engineering
-	J01.293.257	Chemical Engineering
-	J01.293.556	Human Engineering
-	J01.293.556.441	Man-Machine Systems
-	J01.293.622	Hydrology
-	J01.293.688	Optics and Photonics
-	J01.293.754	Quality Improvement
-	J01.293.821	Sanitary Engineering
-	J01.293.821.500	Water Supply

## MeSH Tree Changes for 2017

Type	Tree - heading
-	J01.293.821.500.500 Water Wells
-	J01.494 Household Articles
-	J01.494.221 Bedding and Linens
-	J01.494.221.750 Mosquito Nets
-	J01.494.300 Cooking and Eating Utensils
-	J01.494.650 Humidifiers
-	J01.516 Household Products
-	J01.516.213 Cosmetics
-	J01.516.213.249 Skin Cream
-	J01.516.213.500 Skin Lightening Preparations
-	J01.516.381 Detergents
-	J01.516.872 Soaps
-	J01.576 Industry
-	J01.576.082 Barbering
-	J01.576.112 Beauty Culture
-	J01.576.423 Food Industry
-	J01.576.423.200 Food Handling
-	J01.576.423.200.200 Cooking
-	J01.576.423.200.200.500 Cookbooks as Topic
-	J01.576.423.200.300 Cooking and Eating Utensils
-	J01.576.423.200.350 Food Dispensers, Automatic
-	J01.576.423.200.375 Food Packaging
-	J01.576.423.200.387 Food Storage
-	J01.576.423.200.400 Food-Processing Industry
-	J01.576.423.200.700 Meat-Packing Industry
-	J01.576.423.200.700.100 Abattoirs
-	J01.576.423.200.850 Pasteurization
-	J01.576.423.500 Food Services
-	J01.576.423.500.300 Food Service, Hospital
-	J01.576.423.500.500 Menu Planning
-	J01.576.423.500.700 Restaurants
-	J01.576.423.750 Food Supply
-	J01.576.423.850 Food Technology
-	J01.576.423.850.100 Food Analysis
-	J01.576.423.850.600 Food Packaging
-	J01.576.423.850.600.400 Food Labeling

## MeSH Tree Changes for 2017

Type	Tree - heading
-	J01.576.423.850.700 Food Preservation
-	J01.576.423.850.700.700 Pasteurization
-	J01.576.423.850.700.700.500 Food Irradiation
-	J01.576.423.850.730 Food Quality
-	J01.576.423.850.730.500 Food Safety
-	J01.576.423.850.730.500.249 Food Contamination
-	J01.576.423.850.730.500.249.250 Food Contamination, Radioactive
-	J01.576.423.850.730.500.249.300 Food Microbiology
-	J01.576.423.850.730.500.249.650 Food Parasitology
-	J01.576.423.850.730.500.500 Food Inspection
-	J01.576.423.850.730.500.750 Hazard Analysis and Critical Control Points
-	J01.576.423.850.730.750 Nutritive Value
-	J01.576.423.850.730.750.500 Glycemic Index
-	J01.576.423.850.730.750.750 Glycemic Load
-	J01.576.430 Forestry
-	J01.576.489 Health Care Sector
-	J01.576.519 Industrial Development
-	J01.576.549 Laundering
-	J01.576.655 Manufacturing Industry
-	J01.576.655.125 Construction Industry
-	J01.576.655.374 Book Industry
-	J01.576.655.437 Chemical Industry
-	J01.576.655.750 Drug Industry
-	J01.576.655.750.321 Drug Packaging
-	J01.576.655.750.321.400 Drug Labeling
-	J01.576.655.750.600 Orphan Drug Production
-	J01.576.655.875 Extraction and Processing Industry
-	J01.576.655.875.200 Coal Industry
-	J01.576.655.875.400 Metallurgy
-	J01.576.655.875.400.900 Welding
-	J01.576.655.875.500 Mining
-	J01.576.655.875.500.500 Coal Mining
-	J01.576.655.875.750 Oil and Gas Industry
-	J01.576.655.875.750.500 Hydraulic Fracking
-	J01.576.655.937 Textile Industry
-	J01.576.655.937.800 Textiles

## MeSH Tree Changes for 2017

Type	Tree - heading
-	J01.576.655.937.800.299 Cotton Fiber
-	J01.576.655.937.800.600 Nylons
-	J01.576.655.937.800.800 Wool
-	J01.576.655.968 Tobacco Industry
-	J01.576.761 Product Packaging
-	J01.576.761.300 Drug Packaging
-	J01.576.761.300.400 Drug Labeling
-	J01.576.761.400 Food Packaging
-	J01.576.761.400.450 Food Labeling
-	J01.576.761.700 Product Labeling
-	J01.576.852 Tanning
-	J01.637 Manufactured Materials
-	J01.637.016 Adhesives
-	J01.637.051 Biomedical and Dental Materials
-	J01.637.051.058 Alloys
-	J01.637.051.058.224 Chromium Alloys
-	J01.637.051.058.224.735 Vitallium
-	J01.637.051.058.451 Gold Alloys
-	J01.637.051.058.520 Metal Ceramic Alloys
-	J01.637.051.058.520.250 Cermet Cements
-	J01.637.051.058.807 Steel
-	J01.637.051.058.807.681 Stainless Steel
-	J01.637.051.130 Biocompatible Materials
-	J01.637.051.130.325 Bone Substitutes
-	J01.637.051.130.420 Coated Materials, Biocompatible
-	J01.637.051.130.650 Polydioxanone
-	J01.637.051.187 Cariogenic Agents
-	J01.637.051.223 Cariostatic Agents
-	J01.637.051.223.015 Acidulated Phosphate Fluoride
-	J01.637.051.223.432 Fluorides, Topical
-	J01.637.051.223.716 Sodium Fluoride
-	J01.637.051.223.800 Tin Fluorides
-	J01.637.051.339 Dental Materials
-	J01.637.051.339.208 Dental Alloys
-	J01.637.051.339.208.224 Chromium Alloys
-	J01.637.051.339.208.224.959 Vitallium

## MeSH Tree Changes for 2017

Type	Tree - heading
-	J01.637.051.339.208.291 Dental Amalgam
-	J01.637.051.339.208.534 Gold Alloys
-	J01.637.051.339.208.720 Metal Ceramic Alloys
-	J01.637.051.339.208.720.250 Cermet Cements
-	J01.637.051.339.250 Dental Casting Investment
-	J01.637.051.339.291 Dental Cements
-	J01.637.051.339.291.150 Compomers
-	J01.637.051.339.291.300 Dentin-Bonding Agents
-	J01.637.051.339.291.300.500 Silorane Resins
-	J01.637.051.339.291.402 Glass Ionomer Cements
-	J01.637.051.339.291.402.120 Cermet Cements
-	J01.637.051.339.291.476 Organically Modified Ceramics
-	J01.637.051.339.291.700 Polycarboxylate Cement
-	J01.637.051.339.291.750 Resin Cements
-	J01.637.051.339.291.800 Silicate Cement
-	J01.637.051.339.291.925 Zinc Oxide-Eugenol Cement
-	J01.637.051.339.291.950 Zinc Phosphate Cement
-	J01.637.051.339.312 Dental Implants
-	J01.637.051.339.334 Dental Impression Materials
-	J01.637.051.339.334.574 Inlay Casting Wax
-	J01.637.051.339.376 Dental Porcelain
-	J01.637.051.339.574 Photoinitiators, Dental
-	J01.637.051.339.773 Pit and Fissure Sealants
-	J01.637.051.339.788 Pulp Capping and Pulpectomy Agents
-	J01.637.051.339.816 Resins, Synthetic
-	J01.637.051.339.816.500 Composite Resins
-	J01.637.051.339.816.500.200 Bisphenol A-Glycidyl Methacrylate
-	J01.637.051.339.816.500.300 Compomers
-	J01.637.051.339.859 Root Canal Filling Materials
-	J01.637.051.339.859.495 Gutta-Percha
-	J01.637.051.376 Dentifrices
-	J01.637.051.376.262 Denture Cleansers
-	J01.637.051.376.711 Toothpastes
-	J01.637.051.479 Membranes, Artificial
-	J01.637.051.479.258 Artificial Cells
-	J01.637.051.479.517 Liposomes



## MeSH Tree Changes for 2017

Type	Tree - heading
-	J01.637.051.479.517.500                      Unilamellar Liposomes
-	J01.637.051.479.900                              Virosomes
-	J01.637.051.583                                  Mouthwashes
-	J01.637.051.583.820                            Saliva, Artificial
-	J01.637.051.720                                 Polymers
-	J01.637.051.720.099                            Biopolymers
-	J01.637.051.720.099.500                      Cellulose
-	J01.637.051.720.099.500.252                Cellophane
-	J01.637.051.720.099.500.439                Collodion
-	J01.637.051.720.099.500.719                Hypromellose Derivatives
-	J01.637.051.720.099.625                      Latex
-	J01.637.051.720.099.687                      Lignin
-	J01.637.051.720.099.750                      Rubber
-	J01.637.051.720.200                            Colestipol
-	J01.637.051.720.259                            Cyanoacrylates
-	J01.637.051.720.259.341                      Enbucrilate
-	J01.637.051.720.259.341.110                Bucrylate
-	J01.637.051.720.327                            Elastomers
-	J01.637.051.720.327.782                      Polyurethanes
-	J01.637.051.720.327.840                      Rubber
-	J01.637.051.720.327.840.480                Neoprene
-	J01.637.051.720.327.900                      Silicone Elastomers
-	J01.637.051.720.395                            Fluorocarbon Polymers
-	J01.637.051.720.395.616                      Polytetrafluoroethylene
-	J01.637.051.720.395.616.755                Proplast
-	J01.637.051.720.470                            Hexadimethrine Bromide
-	J01.637.051.720.540                            Latex
-	J01.637.051.720.628                            Organically Modified Ceramics
-	J01.637.051.720.716                            Plastics
-	J01.637.051.720.716.195                      Biodegradable Plastics
-	J01.637.051.720.716.392                      Nylons
-	J01.637.051.720.716.507                      Polyethylenes
-	J01.637.051.720.716.507.500                Polyethylene
-	J01.637.051.720.716.507.600                Polyethyleneimine
-	J01.637.051.720.716.550                      Polypropylenes
-	J01.637.051.720.716.579                      Polystyrenes

## MeSH Tree Changes for 2017

Type	Tree - heading
-	J01.637.051.720.716.579.159 Cholestyramine Resin
-	J01.637.051.720.716.650 Polyurethanes
-	J01.637.051.720.716.721 Polyvinyls
-	J01.637.051.720.716.721.616 Polyvinyl Alcohol
-	J01.637.051.720.716.721.812 Polyvinyl Chloride
-	J01.637.051.720.716.721.838 Povidone
-	J01.637.051.720.716.721.838.745 Povidone-Iodine
-	J01.637.051.720.716.822 Resins, Synthetic
-	J01.637.051.720.716.822.111 Acrylic Resins
-	J01.637.051.720.716.822.111.650 Polymethacrylic Acids
-	J01.637.051.720.716.822.111.650.605 Methylmethacrylates
-	J01.637.051.720.716.822.111.650.605.450 Methylmethacrylate
-	J01.637.051.720.716.822.111.650.605.500 Polymethyl Methacrylate
-	J01.637.051.720.716.822.111.650.750 Polyhydroxyethyl Methacrylate
-	J01.637.051.720.716.822.300 Bone Cements
-	J01.637.051.720.716.822.308 Composite Resins
-	J01.637.051.720.716.822.308.200 Bisphenol A-Glycidyl Methacrylate
-	J01.637.051.720.716.822.308.300 Compomers
-	J01.637.051.720.716.822.461 Epoxy Resins
-	J01.637.051.720.716.822.730 Resin Cements
-	J01.637.051.720.722 Polyanetholesulfonate
-	J01.637.051.720.728 Polyesters
-	J01.637.051.720.728.700 Polydioxanone
-	J01.637.051.720.728.764 Polyethylene Terephthalates
-	J01.637.051.720.728.772 Polyglactin 910
-	J01.637.051.720.728.780 Polyglycolic Acid
-	J01.637.051.720.741 Polyethylene Glycols
-	J01.637.051.720.741.125 Certolizumab Pegol
-	J01.637.051.720.741.250 Cetomacrogol
-	J01.637.051.720.741.485 Hydrogel
-	J01.637.051.720.741.575 Nonoxynol
-	J01.637.051.720.741.610 Octoxynol
-	J01.637.051.720.741.650 Poloxalene
-	J01.637.051.720.741.667 Poloxamer
-	J01.637.051.720.741.685 Polyhydroxyethyl Methacrylate
-	J01.637.051.720.741.700 Polysorbates

## MeSH Tree Changes for 2017

Type	Tree - heading
-	J01.637.051.720.780 Polygeline
-	J01.637.051.720.795 Polyphlorethin Phosphate
-	J01.637.051.720.830 Pyran Copolymer
-	J01.637.051.720.900 Siloxanes
-	J01.637.051.720.900.850 Silicones
-	J01.637.051.720.900.850.150 Dimethylpolysiloxanes
-	J01.637.051.720.900.850.150.750 Simethicone
-	J01.637.051.720.900.850.900 Silicone Elastomers
-	J01.637.051.720.900.850.905 Silicone Gels
-	J01.637.051.720.900.850.950 Silicone Oils
-	J01.637.051.800 Root Canal Irrigants
-	J01.637.051.919 Tissue Adhesives
-	J01.637.051.919.367 Cyanoacrylates
-	J01.637.051.919.367.341 Enbucrilate
-	J01.637.051.919.367.341.110 Bucrylate
-	J01.637.087 Biomimetic Materials
-	J01.637.087.249 Blood Substitutes
-	J01.637.087.500 Membranes, Artificial
-	J01.637.087.500.254 Artificial Cells
-	J01.637.087.500.510 Lipid Bilayers
-	J01.637.087.500.517 Liposomes
-	J01.637.087.500.517.500 Unilamellar Liposomes
-	J01.637.087.500.900 Virosomes
-	J01.637.087.750 Receptors, Artificial
-	J01.637.118 Cellophane
-	J01.637.153 Ceramics
-	J01.637.153.377 Dental Porcelain
-	J01.637.153.688 Organically Modified Ceramics
-	J01.637.215 Clothing
-	J01.637.215.600 Protective Clothing
-	J01.637.215.600.199 Ear Protective Devices
-	J01.637.215.600.400 Gloves, Protective
-	J01.637.215.600.400.400 Gloves, Surgical
-	J01.637.215.600.700 Space Suits
-	J01.637.215.800 Shoes
-	J01.637.241 Construction Materials

## MeSH Tree Changes for 2017

Type	Tree - heading
-	J01.637.241.900 Wood
-	J01.637.370 Cooking and Eating Utensils
-	J01.637.412 Elastomers
-	J01.637.412.700 Polyurethanes
-	J01.637.412.767 Rubber
-	J01.637.412.767.480 Neoprene
-	J01.637.412.900 Silicone Elastomers
-	J01.637.437 Glass
-	J01.637.450 Industrial Oils
-	J01.637.500 Ink
-	J01.637.502 Jewelry
-	J01.637.506 Liquid Crystals
-	J01.637.507 Magnets
-	J01.637.507.500 Magnetite Nanoparticles
-	J01.637.509 Micro-Electrical-Mechanical Systems
-	J01.637.512 Nanostructures
-	J01.637.512.150 Nanocomposites
-	J01.637.512.300 Nanofibers
-	J01.637.512.600 Nanoparticles
-	J01.637.512.600.150 Calcifying Nanoparticles
-	J01.637.512.600.200 Dendrimers
-	J01.637.512.600.500 Metal Nanoparticles
-	J01.637.512.600.500.287 Magnetite Nanoparticles
-	J01.637.512.600.500.575 Nanoshells
-	J01.637.512.600.575 Nanocapsules
-	J01.637.512.600.577 Nanoconjugates
-	J01.637.512.600.580 Nanodiamonds
-	J01.637.512.600.612 Nanospheres
-	J01.637.512.600.612.350 Fullerenes
-	J01.637.512.600.650 Quantum Dots
-	J01.637.512.650 Nanopores
-	J01.637.512.850 Nanotubes
-	J01.637.512.850.500 Nanotubes, Carbon
-	J01.637.512.850.750 Nanotubes, Peptide
-	J01.637.512.925 Nanowires
-	J01.637.548 Nylons

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	J01.637.597	Paint
-	J01.637.597.422	Lacquer
-	J01.637.650	Paper
-	J01.637.708	Protective Devices
-	J01.637.708.100	Air Bags
-	J01.637.708.175	Child Restraint Systems
Old Tree	J01.637.708.250	Ear Protective Devices
Old Tree	J01.637.708.260	Eye Protective Devices
Old Tree	J01.637.708.380	Head Protective Devices
Old Tree	J01.637.708.500	Masks
Old Tree	J01.637.708.500.450	Laryngeal Masks
Old Tree	J01.637.708.520	Mouth Protectors
New Tree	J01.637.708.560	Personal Protective Equipment
New Tree	J01.637.708.560.250	Ear Protective Devices
New Tree	J01.637.708.560.500	Eye Protective Devices
New Tree	J01.637.708.560.750	Head Protective Devices
New Tree	J01.637.708.560.782	Masks
New Tree	J01.637.708.560.782.450	Laryngeal Masks
New Tree	J01.637.708.560.813	Mouth Protectors
New Tree	J01.637.708.560.875	Protective Clothing
New Tree	J01.637.708.560.875.400	Gloves, Protective
New Tree	J01.637.708.560.875.400.400	Gloves, Surgical
New Tree	J01.637.708.560.875.700	Space Suits
New Tree	J01.637.708.560.937	Respiratory Protective Devices
Old Tree	J01.637.708.600	Protective Clothing
Old Tree	J01.637.708.600.400	Gloves, Protective
Old Tree	J01.637.708.600.400.400	Gloves, Surgical
Old Tree	J01.637.708.600.700	Space Suits

## MeSH Tree Changes for 2017

Type	Tree - heading
Old Tree	<b>J01.637.708.700</b> <b>Respiratory Protective Devices</b>
Old Tree	<b>J01.637.708.800</b> <b>Seat Belts</b>
-	J01.637.825 Sports Equipment
-	J01.637.836 Tobacco Products
-	J01.637.836.500 Tobacco, Smokeless
-	J01.637.847 Tobacco Use Cessation Products
-	J01.637.847.500 Electronic Cigarettes
-	J01.637.870 Weapons
-	J01.637.870.175 Bombs
-	J01.637.870.175.500 Nuclear Weapons
-	J01.637.870.350 Firearms
-	J01.637.870.800 Tear Gases
-	J01.637.870.900 Weapons of Mass Destruction
-	J01.637.870.900.100 Biological Warfare Agents
-	J01.637.870.900.200 Chemical Warfare Agents
-	J01.637.870.900.575 Nuclear Weapons
-	J01.637.883 Wool
-	J01.675 Military Science
-	J01.728 Natural Resources
-	J01.728.500 Water Resources
-	J01.780 Power Plants
-	J01.780.600 Nuclear Power Plants
-	J01.897 Technology
-	J01.897.052 Culturally Appropriate Technology
-	J01.897.104 Automation
-	J01.897.104.416 Automation, Laboratory
-	J01.897.104.834 Robotics
-	J01.897.115 Biomedical Technology
-	J01.897.115.250 Biomedical Enhancement
-	J01.897.115.250.500 Genetic Enhancement
-	J01.897.115.625 Nanomedicine
-	J01.897.120 Biotechnology
-	J01.897.120.100 Biomimetics
-	J01.897.120.115 Bioreactors
-	J01.897.120.115.600 Photobioreactors
-	J01.897.120.460 Industrial Microbiology

## MeSH Tree Changes for 2017

Type	Tree - heading
-	J01.897.280 Educational Technology
-	J01.897.280.500 Audiovisual Aids
-	J01.897.280.500.117 Books, Illustrated
-	J01.897.280.500.269 Exhibits as Topic
-	J01.897.280.500.426 Maps as Topic
-	J01.897.280.500.480 Medical Illustration
-	J01.897.280.500.545 Models, Structural
-	J01.897.280.500.545.129 Models, Anatomic
-	J01.897.280.500.545.129.200 Dental Models
-	J01.897.280.500.545.129.400 Manikins
-	J01.897.280.500.545.129.805 Visible Human Projects
-	J01.897.280.500.598 Motion Pictures as Topic
-	J01.897.280.500.633 Multimedia
-	J01.897.280.500.668 Optical Storage Devices
-	J01.897.280.500.668.800 Videodisc Recording
-	J01.897.280.500.668.800.250 Compact Disks
-	J01.897.280.500.668.800.250.100 CD-I
-	J01.897.280.500.668.800.250.200 CD-ROM
-	J01.897.280.500.739 Radio
-	J01.897.280.500.846 Tape Recording
-	J01.897.280.500.846.734 Videotape Recording
-	J01.897.280.500.898 Television
-	J01.897.280.500.898.475 Microscopy, Video
-	J01.897.280.500.898.800 Videodisc Recording
-	J01.897.280.500.898.800.250 Compact Disks
-	J01.897.280.500.898.800.250.100 CD-I
-	J01.897.280.500.898.800.250.200 CD-ROM
-	J01.897.280.500.898.840 Videotape Recording
-	J01.897.360 Green Chemistry Technology
-	J01.897.400 Inventions
-	J01.897.441 Man-Machine Systems
-	J01.897.480 Medical Laboratory Science
-	J01.897.520 Miniaturization
-	J01.897.520.500 Microtechnology
-	J01.897.520.500.500 Microfluidics
-	J01.897.520.600 Nanotechnology

## MeSH Tree Changes for 2017

Type	Tree - heading
-	J01.897.520.600.600                      Nanomedicine
-	J01.897.520.600.600.500                      Theranostic Nanomedicine
-	J01.897.564                      Printing, Three-Dimensional
-	J01.897.608                      Quality Control
-	J01.897.608.500                      Laboratory Proficiency Testing
-	J01.897.724                      Technology, Dental
-	J01.897.836                      Technology, Pharmaceutical
-	J01.897.836.249                      Chemistry Techniques, Synthetic
-	J01.897.836.249.124                      Click Chemistry
-	J01.897.836.249.249                      Combinatorial Chemistry Techniques
-	J01.897.836.249.249.500                      SELEX Aptamer Technique
-	J01.897.836.249.374                      Cycloaddition Reaction
-	J01.897.836.249.500                      Solid-Phase Synthesis Techniques
-	J01.897.836.750                      Molecular Farming
-	J01.897.891                      Technology, Radiologic
-	J01.897.900                      Technology Transfer
-	J01.937                      Transportation
-	J01.937.285                      Aviation
-	J01.937.285.100                      Aircraft
-	J01.937.285.100.100                      Air Ambulances
-	J01.937.285.850                      Space Flight
-	J01.937.285.850.400                      Extravehicular Activity
-	J01.937.285.850.900                      Spacecraft
-	J01.937.500                      Motor Vehicles
-	J01.937.500.050                      Ambulances
-	J01.937.500.100                      Automobiles
-	J01.937.500.500                      Motorcycles
-	J01.937.500.600                      Off-Road Motor Vehicles
-	J01.937.690                      Railroads
-	J01.937.817                      Ships
-	J02                      Food and Beverages
-	J02.200                      Beverages
-	J02.200.100                      Alcoholic Beverages
-	J02.200.100.100                      Absinthe
-	J02.200.100.200                      Beer
-	J02.200.100.900                      Wine



## MeSH Tree Changes for 2017

Type	Tree - heading	
-	J02.200.300	Carbonated Beverages
-	J02.200.300.500	Carbonated Water
-	J02.200.325	Coffee
-	J02.200.418	Drinking Water
-	J02.200.418.500	Carbonated Water
-	J02.200.512	Energy Drinks
-	J02.200.606	Fruit and Vegetable Juices
-	J02.200.700	Milk
Old Tree	J02.200.700.124	Cultured Milk Products
Old Tree	J02.200.700.249	Infant Formula
New Heading	J02.200.700.250	Buttermilk
New Heading	J02.200.700.375	Kefir
New Heading	J02.200.700.438	Koumiss
-	J02.200.700.500	Milk, Human
-	J02.200.700.750	Whey
-	J02.200.712	Milk Substitutes
-	J02.200.712.249	Infant Formula
-	J02.200.712.500	Soy Milk
-	J02.200.831	Tea
-	J02.200.831.500	Kombucha Tea
-	J02.200.950	Teas, Herbal
-	J02.200.975	Teas, Medicinal
-	J02.500	Food
-	J02.500.100	Bread
-	J02.500.140	Candy
-	J02.500.140.200	Chewing Gum
New Heading	J02.500.195	Chocolate
-	J02.500.250	Condiments
-	J02.500.250.725	Spices
-	J02.500.300	Crops, Agricultural
-	J02.500.300.100	Animal Feed
-	J02.500.300.100.800	Silage
-	J02.500.300.550	Edible Grain

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	J02.500.300.550.500	Whole Grains
-	J02.500.350	Dairy Products
-	J02.500.350.100	Butter
-	J02.500.350.100.500	Ghee
Old Tree	<b>J02.500.350.200</b>	<b>Cheese</b>
New Tree	<b>J02.500.350.300</b>	<b>Cultured Milk Products</b>
New Heading	<b>J02.500.350.300.222</b>	<b>Buttermilk</b>
New Tree	<b>J02.500.350.300.444</b>	<b>Cheese</b>
New Heading	<b>J02.500.350.300.666</b>	<b>Kefir</b>
New Heading	<b>J02.500.350.300.777</b>	<b>Koumiss</b>
New Tree	<b>J02.500.350.300.888</b>	<b>Yogurt</b>
-	J02.500.350.400	Ice Cream
-	J02.500.350.500	Margarine
-	J02.500.350.525	Milk
Old Tree	<b>J02.500.350.525.221</b>	<b>Cultured Milk Products</b>
Old Tree	<b>J02.500.350.525.221.888</b>	<b>Yogurt</b>
New Heading	<b>J02.500.350.525.250</b>	<b>Buttermilk</b>
Old Tree	<b>J02.500.350.525.332</b>	<b>Infant Formula</b>
New Heading	<b>J02.500.350.525.375</b>	<b>Kefir</b>
New Heading	<b>J02.500.350.525.438</b>	<b>Koumiss</b>
-	J02.500.350.525.500	Milk, Human
-	J02.500.350.525.520	Milk Proteins
-	J02.500.350.525.520.500	Whey Proteins
-	J02.500.350.525.760	Whey
-	J02.500.350.525.760.500	Whey Proteins
-	J02.500.362	Dietary Carbohydrates
-	J02.500.362.325	Dietary Sucrose
-	J02.500.362.662	High Fructose Corn Syrup
-	J02.500.375	Dietary Fats
-	J02.500.375.200	Butter

## MeSH Tree Changes for 2017

Type	Tree - heading
-	J02.500.375.200.500 Ghee
-	J02.500.375.400 Dietary Fats, Unsaturated
-	J02.500.375.400.250 Corn Oil
-	J02.500.375.400.300 Cottonseed Oil
-	J02.500.375.400.500 Olive Oil
-	J02.500.375.400.700 Safflower Oil
-	J02.500.375.400.725 Sesame Oil
-	J02.500.375.400.750 Soybean Oil
-	J02.500.375.650 Margarine
-	J02.500.400 Dietary Fiber
-	J02.500.400.500 Prebiotics
-	J02.500.428 Dietary Proteins
-	J02.500.428.317 Egg Proteins, Dietary
-	J02.500.428.626 Milk Proteins
-	J02.500.428.626.500 Whey Proteins
-	J02.500.428.920 Vegetable Proteins
-	J02.500.456 Dietary Supplements
-	J02.500.456.249 Prebiotics
-	J02.500.456.500 Probiotics
-	J02.500.456.716 Synbiotics
-	J02.500.456.933 Yeast, Dried
-	J02.500.470 Eggs
-	J02.500.470.349 Egg Proteins, Dietary
-	J02.500.470.700 Egg White
-	J02.500.470.800 Egg Yolk
-	J02.500.477 Fast Foods
-	J02.500.484 Flour
-	J02.500.512 Food Additives
-	J02.500.512.350 Fat Substitutes
-	J02.500.512.400 Flavoring Agents
-	J02.500.512.400.700 Sweetening Agents
-	J02.500.512.400.700.500 Non-Nutritive Sweeteners
-	J02.500.512.400.700.750 Nutritive Sweeteners
-	J02.500.512.400.700.750.250 Dietary Sucrose
-	J02.500.512.400.700.750.500 High Fructose Corn Syrup
-	J02.500.512.700 Food Preservatives

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	J02.500.515	Food, Fortified
-	J02.500.518	Food, Genetically Modified
-	J02.500.519	Food, Organic
-	J02.500.521	Food, Preserved
-	J02.500.521.500	Frozen Foods
-	J02.500.525	Foods, Specialized
-	J02.500.525.350	Food, Formulated
-	J02.500.525.350.500	Infant Formula
-	J02.500.525.500	Infant Food
-	J02.500.525.500.500	Infant Formula
-	J02.500.562	Fruit
-	J02.500.572	Functional Food
-	J02.500.581	Honey
-	J02.500.590	Meals
-	J02.500.590.120	Breakfast
-	J02.500.590.560	Lunch
-	J02.500.590.780	Snacks
-	J02.500.600	Meat
-	J02.500.600.500	Meat Products
-	J02.500.600.750	Poultry
-	J02.500.600.750.500	Poultry Products
-	J02.500.600.813	Red Meat
-	J02.500.600.875	Seafood
-	J02.500.600.875.400	Fish Products
-	J02.500.600.875.400.400	Fish Flour
-	J02.500.600.875.700	Shellfish
-	J02.500.631	Micronutrients
-	J02.500.631.555	Trace Elements
-	J02.500.631.600	Vitamins
New Heading	<b>J02.500.631.600.500</b>	<b>Provitamins</b>
-	J02.500.662	Molasses
-	J02.500.700	Nuts
-	J02.500.737	Raw Foods
-	J02.500.775	Seeds
-	J02.500.775.500	Edible Grain

## MeSH Tree Changes for 2017

Type	Tree - heading
-	J02.500.775.500.500 Whole Grains
-	J02.500.850 Vegetables
-	J02.500.850.800 Vegetable Products
-	J02.500.850.800.500 Soy Foods
-	J02.500.850.800.500.500 Soy Milk
-	J02.500.850.800.500.750 Soybean Proteins
-	J02.500.850.900 Vegetable Proteins
New Heading	<b>J03 Non-Medical Public and Private Facilities</b>
New Tree	J03.040 Adult Day Care Centers
New Tree	J03.150 Cemeteries
New Tree	J03.160 Child Day Care Centers
New Heading	<b>J03.320 Gardens</b>
New Tree	J03.340 Housing
New Tree	J03.340.030 Almshouses
New Tree	J03.340.250 Housing, Animal
New Tree	J03.340.260 Housing for the Elderly
New Tree	J03.340.650 Public Housing
New Tree	J03.400 Information Centers
New Tree	J03.400.208 Archives
New Tree	J03.400.596 Libraries
New Tree	J03.400.596.363 Libraries, Dental
New Tree	J03.400.596.413 Libraries, Digital
New Tree	J03.400.596.463 Libraries, Hospital
New Tree	J03.400.596.563 Libraries, Medical
New	J03.400.596.563.725 National Library of Medicine (U.S.)

## MeSH Tree Changes for 2017

Type	Tree - heading
Tree	
New Tree	J03.400.596.663 Libraries, Nursing
New Tree	J03.400.596.831 Libraries, Special
New Tree	J03.520 Laboratories
New Heading	<b>J03.540 Manufacturing and Industrial Facilities</b>
New Tree	J03.540.020 Abattoirs
New Heading	<b>J03.540.150 Farms</b>
New Tree	J03.540.280 Fisheries
New Tree	J03.540.680 Power Plants
New Tree	J03.540.680.600 Nuclear Power Plants
New Tree	J03.560 Military Facilities
New Tree	J03.570 Museums
New Tree	J03.693 Prisons
New Tree	J03.693.358 Concentration Camps
New Heading	<b>J03.700 Private Facilities</b>
New Tree	J03.720 Public Facilities
New Tree	J03.748 Religious Missions
New Tree	J03.775 Residential Facilities
New Tree	J03.775.187 Assisted Living Facilities
New Tree	J03.775.375 Group Homes
New Tree	J03.775.450 Halfway Houses
New Tree	J03.775.462 Homes for the Aged

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	J03.775.640	Orphanages
New Tree	J03.813	Restaurants
New Tree	J03.832	Schools
New Tree	J03.832.330	Library Schools
New Tree	J03.832.660	Schools, Nursery
New Tree	J03.832.830	Universities
New Tree	J03.850	Senior Centers
New Tree	J03.860	Sheltered Workshops
New Heading	<b>J03.925</b>	<b>Sports and Recreational Facilities</b>
New Tree	J03.925.080	Bathing Beaches
New Tree	J03.925.680	Parks, Recreational
New Tree	J03.925.830	Swimming Pools
New Tree	J03.962	Toilet Facilities
New Heading	<b>J03.970</b>	<b>Transportation Facilities</b>
New Tree	J03.970.100	Airports
New Tree	J03.970.680	Parking Facilities
-	K01	Humanities
-	K01.093	Art
-	K01.093.206	Caricatures as Topic
-	K01.093.206.332	Cartoons as Topic
-	K01.093.346	Engraving and Engravings
-	K01.093.378	Human Body
-	K01.093.410	Medical Illustration
-	K01.093.410.100	Anatomy, Artistic
-	K01.093.530	Medicine in Art
-	K01.093.560	Numismatics

## MeSH Tree Changes for 2017

Type	Tree - heading
-	K01.093.646 Paintings
-	K01.093.660 Philately
-	K01.093.742 Portraits as Topic
-	K01.093.849 Sculpture
-	K01.150 Awards and Prizes
-	K01.150.567 Nobel Prize
-	K01.400 History
-	K01.400.075 Alchemy
-	K01.400.095 Anniversaries and Special Events
-	K01.400.152 Archives
-	K01.400.152.552 Museums
-	K01.400.255 Emblems and Insignia
-	K01.400.307 Eugenics
-	K01.400.359 Genealogy and Heraldry
-	K01.400.441 Historiography
-	K01.400.470 History, Ancient
-	K01.400.475 History, Early Modern 1451-1600
-	K01.400.475.500 History, 15th Century
-	K01.400.475.750 History, 16th Century
-	K01.400.500 History, Medieval
-	K01.400.504 History, Modern 1601-
-	K01.400.504.750 History, 17th Century
-	K01.400.504.875 History, 18th Century
-	K01.400.504.875.100 American Revolution
-	K01.400.504.875.374 French Revolution
-	K01.400.504.937 History, 19th Century
-	K01.400.504.937.100 American Civil War
-	K01.400.504.937.500 Crimean War
-	K01.400.504.937.750 Spanish-American War, 1898
-	K01.400.504.968 History, 20th Century
-	K01.400.504.968.074 Bhopal Accidental Release
-	K01.400.504.968.150 Chernobyl Nuclear Accident
-	K01.400.504.968.400 Gulf War
-	K01.400.504.968.450 Influenza Pandemic, 1918-1919
-	K01.400.504.968.500 Korean War
-	K01.400.504.968.650 Russian-Japanese War



## MeSH Tree Changes for 2017

Type	Tree - heading
-	K01.400.504.968.700 Seveso Accidental Release
-	K01.400.504.968.750 Sverdlovsk Accidental Release
-	K01.400.504.968.850 Vietnam Conflict
-	K01.400.504.968.900 World War I
-	K01.400.504.968.901 World War II
-	K01.400.504.984 History, 21st Century
-	K01.400.504.984.124 Afghan Campaign 2001-
-	K01.400.504.984.186 Fukushima Nuclear Accident
-	K01.400.504.984.249 Iraq War, 2003-2011
-	K01.400.504.984.500 September 11 Terrorist Attacks
-	K01.400.551 History of Dentistry
-	K01.400.552 History of Medicine
-	K01.400.608 History of Nursing
-	K01.400.703 Natural History
-	K01.400.799 Phrenology
-	K01.400.899 Symbolism
-	K01.400.899.500 Metaphor
-	K01.468 Knowledge
-	K01.517 Literature
-	K01.517.067 Anecdotes as Topic
-	K01.517.116 Aphorisms and Proverbs as Topic
-	K01.517.172 Bible
-	K01.517.211 Biography as Topic
-	K01.517.211.215 Autobiography as Topic
-	K01.517.211.506 Famous Persons
-	K01.517.211.753 Personal Narratives as Topic
-	K01.517.333 Drama
-	K01.517.495 Literature, Medieval
-	K01.517.533 Literature, Modern
-	K01.517.584 Medicine in Literature
-	K01.517.584.500 Psychiatry in Literature
-	K01.517.647 Mythology
-	K01.517.732 Philology
-	K01.517.732.477 Philology, Classical
-	K01.517.732.602 Philology, Oriental
-	K01.517.732.727 Philology, Romance

## MeSH Tree Changes for 2017

Type	Tree - heading
-	K01.517.781 Poetry as Topic
-	K01.517.946 Wit and Humor as Topic
-	K01.602 Music
-	K01.637 Nature
-	K01.672 Occultism
-	K01.672.296 Astrology
-	K01.752 Philosophy
-	K01.752.210 Esthetics
-	K01.752.210.304 Beauty
-	K01.752.304 Existentialism
-	K01.752.400 Life
-	K01.752.400.100 Beginning of Human Life
-	K01.752.400.750 Quality of Life
-	K01.752.400.900 Value of Life
-	K01.752.448 Logic
-	K01.752.448.250 Fuzzy Logic
-	K01.752.511 Metaphysics
-	K01.752.566 Morals
-	K01.752.566.359 Conscience
-	K01.752.566.479 Ethics
-	K01.752.566.479.023 Bioethical Issues
-	K01.752.566.479.045 Bioethics
-	K01.752.566.479.045.500 Ethics, Clinical
-	K01.752.566.479.068 Codes of Ethics
-	K01.752.566.479.068.249 Helsinki Declaration
-	K01.752.566.479.068.400 Hippocratic Oath
-	K01.752.566.479.090 Complicity
-	K01.752.566.479.095 Conflict of Interest
-	K01.752.566.479.101 Double Effect Principle
-	K01.752.566.479.113 Ethical Analysis
-	K01.752.566.479.113.150 Casuistry
-	K01.752.566.479.113.750 Retrospective Moral Judgment
-	K01.752.566.479.113.875 Wedge Argument
-	K01.752.566.479.116 Ethical Relativism
-	K01.752.566.479.117 Ethical Review
-	K01.752.566.479.117.200 Ethics Consultation

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	K01.752.566.479.118	Ethical Theory
-	K01.752.566.479.119	Ethicists
-	K01.752.566.479.147	Ethics Committees
-	K01.752.566.479.147.500	Ethics Committees, Clinical
-	K01.752.566.479.147.750	Ethics Committees, Research
-	K01.752.566.479.161	Ethics, Business
-	K01.752.566.479.168	Ethics, Institutional
-	K01.752.566.479.171	Ethics, Professional
-	K01.752.566.479.171.066	Codes of Ethics
-	K01.752.566.479.171.066.249	Helsinki Declaration
-	K01.752.566.479.171.132	Ethics, Clinical
-	K01.752.566.479.171.132.500	Ethics, Dental
-	K01.752.566.479.171.132.750	Ethics, Medical
-	K01.752.566.479.171.132.750.551	Hippocratic Oath
-	K01.752.566.479.171.132.750.775	Therapeutic Equipoise
-	K01.752.566.479.171.132.875	Ethics, Nursing
-	K01.752.566.479.171.132.937	Ethics, Pharmacy
-	K01.752.566.479.171.566	Professionalism
-	K01.752.566.479.173	Ethics, Research
-	K01.752.566.479.173.200	Censorship, Research
-	K01.752.566.479.173.400	Helsinki Declaration
-	K01.752.566.479.173.700	Therapeutic Equipoise
-	K01.752.566.479.173.850	Therapeutic Misconception
-	K01.752.566.479.174	Humanism
Old Tree	K01.752.566.479.174.400	Feminism
New Tree	K01.752.566.479.174.700	Feminism
New Heading	<b>K01.752.566.479.417</b>	<b>Moral Status</b>
-	K01.752.566.479.660	Personhood
-	K01.752.566.479.830	Principle-Based Ethics
-	K01.752.566.479.830.500	Beneficence
-	K01.752.566.479.830.650	Personal Autonomy
-	K01.752.566.479.830.750	Social Justice
-	K01.752.566.479.915	Professional Misconduct
-	K01.752.566.479.915.500	Scientific Misconduct

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	K01.752.566.739	Moral Development
-	K01.752.566.869	Social Responsibility
-	K01.752.566.869.500	Moral Obligations
-	K01.752.566.934	Virtues
-	K01.752.622	Philosophy, Dental
-	K01.752.622.416	Ethics, Dental
-	K01.752.667	Philosophy, Medical
-	K01.752.667.400	Empiricism
-	K01.752.667.555	Germ Theory of Disease
-	K01.752.667.710	Holistic Health
-	K01.752.667.750	Humoralism
-	K01.752.667.875	Hygiene Hypothesis
-	K01.752.712	Philosophy, Nursing
-	K01.752.712.416	Ethics, Nursing
-	K01.752.712.708	Holistic Nursing
-	K01.752.720	Postmodernism
-	K01.752.730	Qi
-	K01.752.798	Symbolism
-	K01.752.798.500	Metaphor
-	K01.752.827	Thanatology
-	K01.752.884	Utopias
-	K01.752.935	Vitalism
-	K01.752.967	Yin-Yang
-	K01.844	Religion
-	K01.844.058	Anthroposophy
-	K01.844.117	Buddhism
-	K01.844.188	Christianity
-	K01.844.188.250	Catholicism
-	K01.844.188.431	Christian Science
-	K01.844.188.466	Church of Jesus Christ of Latter-day Saints
-	K01.844.188.502	Eastern Orthodoxy
-	K01.844.188.573	Jehovah's Witnesses
-	K01.844.188.644	Protestantism
-	K01.844.188.715	Saints
-	K01.844.231	Hinduism
-	K01.844.275	Islam

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	K01.844.385	Judaism
-	K01.844.619	Religion and Medicine
-	K01.844.619.500	Spirit Possession
-	K01.844.664	Religion and Psychology
-	K01.844.664.500	Spirituality
-	K01.844.709	Religion and Science
-	K01.844.754	Religion and Sex
Old Tree	<b>K01.844.776</b>	<b>Religious Missions</b>
New Tree	<b>K01.844.777</b>	<b>Religious Missions</b>
-	K01.844.799	Religious Philosophies
-	K01.844.799.214	Confucianism
-	K01.844.799.430	Mysticism
-	K01.844.799.679	Spiritualism
-	K01.844.799.867	Yoga
-	K01.844.949	Theology
-	K01.883	Romanticism
-	K01.922	Secularism
-	L01	Information Science
-	L01.040	Book Collecting
New Heading	<b>L01.060</b>	<b>Calendars as Topic</b>
-	L01.080	Chronology as Topic
-	L01.100	Classification
-	L01.100.697	Phylogeny
-	L01.143	Communication
-	L01.143.024	Access to Information
-	L01.143.050	Advertising as Topic
-	L01.143.140	Answering Services
-	L01.143.230	Communication Barriers
-	L01.143.230.500	Digital Divide
-	L01.143.256	Computer Literacy
-	L01.143.283	Cybernetics
-	L01.143.283.425	Feedback
-	L01.143.283.425.156	Formative Feedback
-	L01.143.283.425.311	Feedback, Physiological

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	L01.143.283.425.311.500	Feedback, Sensory
-	L01.143.283.425.624	Feedback, Psychological
-	L01.143.283.425.624.500	Biofeedback, Psychology
-	L01.143.283.425.624.500.500	Neurofeedback
-	L01.143.283.425.624.750	Feedback, Sensory
-	L01.143.320	Diffusion of Innovation
-	L01.143.320.800	Technology Transfer
-	L01.143.335	Disclosure
-	L01.143.350	Health Communication
Old Tree	L01.143.380	Hotlines
-	L01.143.443	Information Dissemination
-	L01.143.450	Information Literacy
-	L01.143.450.500	Health Literacy
-	L01.143.458	Information Seeking Behavior
-	L01.143.474	Interdisciplinary Communication
-	L01.143.506	Language
-	L01.143.506.423	Language Arts
-	L01.143.506.423.348	Lipreading
-	L01.143.506.423.452	Multilingualism
-	L01.143.506.423.557	Reading
-	L01.143.506.423.676	Speech
-	L01.143.506.423.796	Translating
-	L01.143.506.423.906	Writing
-	L01.143.506.423.906.215	Authorship
-	L01.143.506.423.906.377	Correspondence as Topic
-	L01.143.506.423.906.377.333	Electronic Mail
New Heading	L01.143.506.423.906.377.500	Postcards as Topic
-	L01.143.506.423.906.377.666	Text Messaging
-	L01.143.506.423.906.539	Handwriting
-	L01.143.506.423.906.539.500	Paleography
-	L01.143.506.423.906.539.750	Shorthand
-	L01.143.506.423.906.654	Medical Writing
-	L01.143.506.423.906.769	Research Report
-	L01.143.506.598	Linguistics
-	L01.143.506.598.400	Terminology as Topic

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	L01.143.506.598.400.556	Names
-	L01.143.506.598.400.556.131	Abbreviations as Topic
-	L01.143.506.598.400.556.283	Anonyms and Pseudonyms
-	L01.143.506.598.400.556.536	Eponyms
-	L01.143.506.598.400.778	RxNorm
New Heading	<b>L01.143.506.598.400.889</b>	<b>Standardized Nursing Terminology</b>
-	L01.143.506.598.518	Phonetics
-	L01.143.506.598.628	Psycholinguistics
-	L01.143.506.598.628.550	Neurolinguistic Programming
-	L01.143.506.598.745	Semantics
-	L01.143.506.598.901	Vocabulary
-	L01.143.563	Literacy
-	L01.143.620	Negotiating
-	L01.143.649	Nonverbal Communication
-	L01.143.649.526	Manual Communication
-	L01.143.649.526.668	Sign Language
-	L01.143.762	Persuasive Communication
-	L01.143.790	Propaganda
-	L01.143.805	Public Service Announcements as Topic
-	L01.143.820	Reminder Systems
-	L01.143.910	Social Networking
-	L01.178	Communications Media
New Heading	<b>L01.178.181</b>	<b>Call Centers</b>
New Tree	<a href="#">L01.178.181.500</a>	<a href="#">Hotlines</a>
New Heading	<b>L01.178.272</b>	<b>Emergency Police Dispatcher</b>
-	L01.178.362	Erotica
-	L01.178.476	Library Materials
-	L01.178.590	Mass Media
-	L01.178.590.500	Motion Pictures as Topic
-	L01.178.590.700	Radio
-	L01.178.590.875	Television
-	L01.178.590.875.800	Videodisc Recording
-	L01.178.590.875.800.250	Compact Disks

## MeSH Tree Changes for 2017

Type	Tree - heading
-	L01.178.590.875.800.250.100 CD-I
-	L01.178.590.875.800.250.200 CD-ROM
-	L01.178.590.875.840 Videotape Recording
-	L01.178.682 Publications
-	L01.178.682.099 Bibliography as Topic
-	L01.178.682.099.183 Bibliography, National
-	L01.178.682.099.308 Bibliography of Medicine
-	L01.178.682.099.325 Bibliometrics
-	L01.178.682.099.325.500 Journal Impact Factor
-	L01.178.682.099.325.750 Literature Based Discovery
-	L01.178.682.099.434 Biobibliography as Topic
-	L01.178.682.150 Book Reviews as Topic
-	L01.178.682.192 Books
-	L01.178.682.192.122 Book Imprints
-	L01.178.682.192.122.622 Printers' Marks
-	L01.178.682.192.178 Book Ornamentation
-	L01.178.682.192.234 Bookplates as Topic
-	L01.178.682.192.289 Books, Illustrated
-	L01.178.682.192.402 Cookbooks as Topic
-	L01.178.682.192.515 Incunabula as Topic
-	L01.178.682.192.609 Manuals as Topic
-	L01.178.682.192.609.666 Sex Manuals
-	L01.178.682.192.779 Rare Books
-	L01.178.682.192.836 Reference Books
-	L01.178.682.192.836.072 Almanacs as Topic
-	L01.178.682.192.836.147 Atlases as Topic
-	L01.178.682.192.836.285 Dictionaries as Topic
-	L01.178.682.192.836.285.206 Dictionaries, Chemical
-	L01.178.682.192.836.285.277 Dictionaries, Classical
-	L01.178.682.192.836.285.348 Dictionaries, Dental
-	L01.178.682.192.836.285.419 Dictionaries, Medical
-	L01.178.682.192.836.285.490 Dictionaries, Pharmaceutic
-	L01.178.682.192.836.285.561 Dictionaries, Polyglot
-	L01.178.682.192.836.339 Directories as Topic
-	L01.178.682.192.836.390 Dispensatories
-	L01.178.682.192.836.456 Encyclopedias as Topic



## MeSH Tree Changes for 2017

Type	Tree - heading
-	L01.178.682.192.836.535                      Formularies as Topic
-	L01.178.682.192.836.535.310                      Formularies, Dental
-	L01.178.682.192.836.535.435                      Formularies, Homeopathic
-	L01.178.682.192.836.535.561                      Formularies, Hospital
-	L01.178.682.192.836.642                      Herbals as Topic
-	L01.178.682.192.836.749                      Pharmacopoeias as Topic
-	L01.178.682.192.836.749.599                      Pharmacopoeias, Homeopathic
-	L01.178.682.192.836.842                      Reference Books, Medical
-	L01.178.682.192.836.842.385                      Dictionaries, Medical
-	L01.178.682.192.900                      Textbooks as Topic
-	L01.178.682.236                      Broadsides as Topic
-	L01.178.682.290                      Catalogs as Topic
-	L01.178.682.290.183                      Catalogs, Commercial
-	L01.178.682.290.183.249                      Catalogs, Booksellers'
-	L01.178.682.290.183.416                      Catalogs, Publishers'
-	L01.178.682.290.283                      Catalogs, Drug
-	L01.178.682.290.383                      Catalogs, Library
-	L01.178.682.290.483                      Catalogs, Union
-	L01.178.682.389                      Dissertations, Academic as Topic
-	L01.178.682.492                      Government Publications as Topic
-	L01.178.682.608                      Manuscripts as Topic
-	L01.178.682.608.526                      Manuscripts, Medical
-	L01.178.682.707                      Pamphlets
-	L01.178.682.759                      Review Literature as Topic
-	L01.178.682.759.150                      Consensus Development Conferences as Topic
-	L01.178.682.759.150.500                      Consensus Development Conferences, NIH as Topic
-	L01.178.682.829                      Serial Publications
-	L01.178.682.829.481                      Newspapers as Topic
-	L01.178.682.829.678                      Periodicals as Topic
-	L01.178.682.920                      Translations
-	L01.178.820                      Teaching Materials
-	L01.178.820.090                      Audiovisual Aids
-	L01.178.820.090.269                      Exhibits as Topic
-	L01.178.820.090.426                      Maps as Topic
-	L01.178.820.090.480                      Medical Illustration
-	L01.178.820.090.545                      Models, Structural

## MeSH Tree Changes for 2017

Type	Tree - heading
-	L01.178.820.090.545.129 Models, Anatomic
-	L01.178.820.090.545.129.200 Dental Models
-	L01.178.820.090.545.129.400 Manikins
-	L01.178.820.090.598 Motion Pictures as Topic
-	L01.178.820.090.615 MP3-Player
-	L01.178.820.090.633 Multimedia
-	L01.178.820.090.668 Optical Storage Devices
-	L01.178.820.090.668.800 Videodisc Recording
-	L01.178.820.090.668.800.250 Compact Disks
-	L01.178.820.090.668.800.250.100 CD-I
-	L01.178.820.090.668.800.250.200 CD-ROM
-	L01.178.820.090.703 Posters as Topic
-	L01.178.820.090.739 Radio
-	L01.178.820.090.846 Tape Recording
-	L01.178.820.090.846.734 Videotape Recording
-	L01.178.820.090.898 Television
-	L01.178.820.090.898.475 Microscopy, Video
-	L01.178.820.090.898.800 Videodisc Recording
-	L01.178.820.090.898.800.250 Compact Disks
-	L01.178.820.090.898.800.250.100 CD-I
-	L01.178.820.090.898.800.250.200 CD-ROM
-	L01.178.820.090.898.840 Videotape Recording
-	L01.178.820.500 Manuals as Topic
-	L01.178.820.500.666 Sex Manuals
-	L01.178.820.900 Textbooks as Topic
-	L01.178.847 Telecommunications
-	L01.178.847.249 Electronic Mail
-	L01.178.847.500 Radar
-	L01.178.847.514 Radio
-	L01.178.847.550 Satellite Communications
-	L01.178.847.550.500 Satellite Imagery
-	L01.178.847.606 Telefacsimile
-	L01.178.847.652 Telemedicine
-	L01.178.847.652.550 Remote Consultation
-	L01.178.847.652.600 Telepathology
-	L01.178.847.652.700 Teleradiology

## MeSH Tree Changes for 2017

Type	Tree - heading
-	L01.178.847.652.850                      Telerehabilitation
-	L01.178.847.675                              Telemetry
-	L01.178.847.675.500                      Remote Sensing Technology
-	L01.178.847.675.500.500                      Satellite Imagery
-	L01.178.847.698                              Telephone
-	L01.178.847.698.249                      Answering Services
-	L01.178.847.698.300                      Cell Phones
-	L01.178.847.698.300.250                      Smartphone
-	L01.178.847.698.300.500                      Text Messaging
-	L01.178.847.698.500                      Modems
-	L01.178.847.823                              Television
-	L01.178.847.823.475                      Microscopy, Video
-	L01.178.847.900                              Videoconferencing
-	L01.178.847.900.500                      Webcasts as Topic
-	L01.178.847.950                              Wireless Technology
-	L01.209    Computer Security
-	L01.209.500                                      Data Anonymization
-	L01.224    Computing Methodologies
-	L01.224.050                                      Algorithms
-	L01.224.050.375                              Artificial Intelligence
-	L01.224.050.375.095                      Computer Heuristics
-	L01.224.050.375.190                      Expert Systems
-	L01.224.050.375.250                      Fuzzy Logic
-	L01.224.050.375.480                      Knowledge Bases
-	L01.224.050.375.480.500                      Biological Ontologies
-	L01.224.050.375.480.500.500                      Gene Ontology
-	L01.224.050.375.530                      Machine Learning
-	L01.224.050.375.530.500                      Supervised Machine Learning
-	L01.224.050.375.530.500.500                      Support Vector Machine
-	L01.224.050.375.530.750                      Unsupervised Machine Learning
-	L01.224.050.375.580                      Natural Language Processing
-	L01.224.050.375.605                      Neural Networks (Computer)
-	L01.224.050.375.630                      Robotics
-	L01.224.085                                      Automatic Data Processing
-	L01.224.085.800                              Punched-Card Systems
-	L01.224.097                                      Cloud Computing

## MeSH Tree Changes for 2017

Type	Tree - heading
-	L01.224.108 Computer Graphics
-	L01.224.108.150 Computer-Aided Design
-	L01.224.108.150.500 Printing, Three-Dimensional
-	L01.224.160 Computer Simulation
-	L01.224.160.249 Molecular Docking Simulation
-	L01.224.160.500 Molecular Dynamics Simulation
-	L01.224.160.750 Patient-Specific Modeling
-	L01.224.230 Computer Systems
-	L01.224.230.110 Computer Communication Networks
-	L01.224.230.110.500 Internet
-	L01.224.230.110.500.500 Blogging
-	L01.224.230.110.500.750 Social Media
-	L01.224.230.110.600 Local Area Networks
-	L01.224.230.260 Computers
-	L01.224.230.260.115 Computer Peripherals
-	L01.224.230.260.115.475 Computer Storage Devices
-	L01.224.230.260.115.475.600 Optical Storage Devices
-	L01.224.230.260.115.475.600.325 Compact Disks
-	L01.224.230.260.115.475.600.325.500 CD-I
-	L01.224.230.260.115.475.600.325.750 CD-ROM
-	L01.224.230.260.115.500 Computer Terminals
-	L01.224.230.260.115.750 Modems
-	L01.224.230.260.230 Computers, Analog
-	L01.224.230.260.280 Computers, Hybrid
-	L01.224.230.260.280.080 Analog-Digital Conversion
-	L01.224.230.260.308 Computers, Mainframe
-	L01.224.230.260.315 Computers, Molecular
-	L01.224.230.260.550 Microcomputers
-	L01.224.230.260.550.500 Computers, Handheld
-	L01.224.230.260.550.500.500 Smartphone
-	L01.224.230.260.580 Minicomputers
-	L01.224.300 Computers, Molecular
-	L01.224.308 Image Processing, Computer-Assisted
-	L01.224.308.189 Data Compression
-	L01.224.308.380 Image Enhancement
-	L01.224.308.380.600 Radiographic Image Enhancement

## MeSH Tree Changes for 2017

Type	Tree - heading
-	L01.224.308.380.800                      Radiography, Dual-Energy Scanned Projection
-	L01.224.308.410                              Imaging, Three-Dimensional
-	L01.224.680                                  Mathematical Computing
-	L01.224.680.700                              Numerical Analysis, Computer-Assisted
-	L01.224.800                                  Signal Processing, Computer-Assisted
-	L01.224.800.500                              Data Compression
-	L01.224.800.625                              Fourier Analysis
-	L01.224.800.750                              Wavelet Analysis
-	L01.224.900                                  Software
-	L01.224.900.280                              Database Management Systems
-	L01.224.900.400                              Grateful Med
-	L01.224.900.590                              Hypermedia
-	L01.224.900.685                              Mobile Applications
-	L01.224.900.780                              Programming Languages
-	L01.224.900.820                              Software Design
-	L01.224.900.868                              Software Validation
-	L01.224.900.889                              Speech Recognition Software
-	L01.224.900.910                              User-Computer Interface
-	L01.224.900.930                              Video Games
-	L01.224.900.940                              Web Browser
-	L01.224.900.950                              Word Processing
-	L01.240    Copying Processes
-	L01.240.630                                  Microfilming
-	L01.240.940                                  Tape Recording
-	L01.240.940.840                              Videotape Recording
-	L01.240.950                                  Telefacsimile
-	L01.240.960                                  Video Recording
-	L01.240.960.399                              Time-Lapse Imaging
-	L01.240.960.800                              Videodisc Recording
-	L01.240.960.800.250                          Compact Disks
-	L01.240.960.800.250.100                      CD-I
-	L01.240.960.800.250.200                      CD-ROM
-	L01.240.960.880                              Videotape Recording
-	L01.280    Data Collection
-	L01.280.090                                  Common Data Elements
-	L01.280.179                                  Crowdsourcing

## MeSH Tree Changes for 2017

Type	Tree - heading
-	L01.280.202 Data Accuracy
-	L01.280.224 Datasets as Topic
-	L01.280.269 Focus Groups
-	L01.280.360 Geriatric Assessment
-	L01.280.520 Interviews as Topic
-	L01.280.660 Narration
-	L01.280.900 Records as Topic
-	L01.280.900.250 Birth Certificates
-	L01.280.900.350 Death Certificates
-	L01.280.900.375 Dental Records
-	L01.280.900.425 Hospital Records
-	L01.280.900.968 Medical Records
-	L01.280.900.968.500 Medical Record Linkage
-	L01.280.900.968.550 Medical Records, Problem-Oriented
-	L01.280.900.968.775 Medical Records Systems, Computerized
-	L01.280.900.968.775.500 Health Smart Cards
-	L01.280.900.968.875 Trauma Severity Indices
-	L01.280.900.968.875.125 Abbreviated Injury Scale
-	L01.280.900.968.875.250 Glasgow Coma Scale
-	L01.280.900.968.875.260 Glasgow Outcome Scale
-	L01.280.900.968.875.500 Injury Severity Score
-	L01.280.900.984 Nursing Records
-	L01.280.950 Registries
-	L01.280.950.725 SEER Program
-	L01.280.960 Surveys and Questionnaires
-	L01.280.960.233 Contraceptive Prevalence Surveys
-	L01.280.960.310 Delphi Technique
-	L01.280.960.500 Health Surveys
-	L01.280.960.500.149 Behavioral Risk Factor Surveillance System
-	L01.280.960.500.300 Dental Health Surveys
-	L01.280.960.500.300.300 Dental Plaque Index
-	L01.280.960.500.300.350 DMF Index
-	L01.280.960.500.300.512 Index of Orthodontic Treatment Need
-	L01.280.960.500.300.675 Oral Hygiene Index
-	L01.280.960.500.300.725 Periodontal Index
-	L01.280.960.500.475 Health Status Indicators

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	L01.280.960.500.475.365	APACHE
Old Tree	<b>L01.280.960.500.475.410</b>	<b>Organ Dysfunction Scores</b>
-	L01.280.960.500.475.456	Patient Acuity
-	L01.280.960.500.475.456.500	Severity of Illness Index
New Tree	<b>L01.280.960.500.475.456.500.250</b>	<b>APACHE</b>
-	L01.280.960.500.475.456.500.500	Karnofsky Performance Status
New Tree	<b>L01.280.960.500.475.456.500.625</b>	<b>Organ Dysfunction Scores</b>
New Heading	<b>L01.280.960.500.475.456.500.750</b>	<b>Simplified Acute Physiology Score</b>
-	L01.280.960.500.475.730	Sickness Impact Profile
-	L01.280.960.500.580	Mass Screening
-	L01.280.960.500.580.174	Anonymous Testing
-	L01.280.960.500.580.510	Mass Chest X-Ray
-	L01.280.960.500.580.560	Multiphasic Screening
-	L01.280.960.500.580.580	Neonatal Screening
-	L01.280.960.500.580.925	Vision Screening
-	L01.280.960.500.700	Population Surveillance
-	L01.280.960.500.700.324	Public Health Surveillance
-	L01.280.960.500.700.650	Sentinel Surveillance
-	L01.280.960.578	Nutrition Surveys
-	L01.280.960.578.350	Diet Surveys
-	L01.280.960.655	Self Report
-	L01.280.975	Vital Statistics
-	L01.280.975.450	Life Expectancy
-	L01.280.975.475	Life Tables
-	L01.280.975.475.700	Quality-Adjusted Life Years
-	L01.280.975.525	Morbidity
-	L01.280.975.525.080	Basic Reproduction Number
-	L01.280.975.525.375	Incidence
-	L01.280.975.525.750	Prevalence
-	L01.280.975.550	Mortality
-	L01.280.975.550.250	Cause of Death
-	L01.280.975.550.287	Child Mortality
-	L01.280.975.550.325	Fatal Outcome
-	L01.280.975.550.362	Fetal Mortality

## MeSH Tree Changes for 2017

Type	Tree - heading
-	L01.280.975.550.400 Hospital Mortality
-	L01.280.975.550.475 Infant Mortality
New Tree	<a href="#">L01.280.975.550.475.500</a> Perinatal Mortality
-	L01.280.975.550.500 Maternal Mortality
-	L01.280.975.550.550 Mortality, Premature
Old Tree	<del>L01.280.975.550.700</del> Perinatal Mortality
-	L01.280.975.550.900 Survival Rate
-	L01.280.975.775 Pregnancy Rate
-	L01.280.975.775.500 Birth Rate
-	L01.296 Data Display
-	L01.296.110 Computer Graphics
-	L01.296.110.150 Computer-Aided Design
-	L01.296.110.150.500 Printing, Three-Dimensional
-	L01.313 Informatics
-	L01.313.124 Computational Biology
-	L01.313.249 Dental Informatics
-	L01.313.500 Medical Informatics
-	L01.313.500.500 Health Information Exchange
-	L01.313.500.750 Medical Informatics Applications
-	L01.313.500.750.100 Decision Making, Computer-Assisted
-	L01.313.500.750.100.158 Diagnosis, Computer-Assisted
-	L01.313.500.750.100.158.600 Image Interpretation, Computer-Assisted
-	L01.313.500.750.100.158.600.680 Radiographic Image Interpretation, Computer-Assisted
-	L01.313.500.750.100.710 Therapy, Computer-Assisted
-	L01.313.500.750.100.710.180 Drug Therapy, Computer-Assisted
-	L01.313.500.750.100.710.600 Radiotherapy, Computer-Assisted
-	L01.313.500.750.100.710.600.550 Radiotherapy, Conformal
-	L01.313.500.750.100.710.600.550.700 Radiotherapy, Intensity-Modulated
-	L01.313.500.750.100.710.600.608 Radiotherapy Planning, Computer-Assisted
-	L01.313.500.750.100.710.800 Surgery, Computer-Assisted
-	L01.313.500.750.100.710.800.500 Robotic Surgical Procedures
-	L01.313.500.750.190 Decision Support Techniques
-	L01.313.500.750.190.380 Data Interpretation, Statistical
-	L01.313.500.750.280 Information Storage and Retrieval



## MeSH Tree Changes for 2017

Type	Tree - heading
-	L01.313.500.750.280.199 Data Mining
-	L01.313.500.750.280.199.500 Multifactor Dimensionality Reduction
-	L01.313.500.750.280.400 Grateful Med
-	L01.313.500.750.280.710 MEDLARS
-	L01.313.500.750.280.710.500 MEDLINE
-	L01.313.500.750.280.730 MedlinePlus
-	L01.313.500.750.280.750 PubMed
-	L01.313.500.750.280.750.500 MEDLINE
-	L01.313.500.750.280.875 RxNorm
-	L01.313.500.750.300 Information Systems
-	L01.313.500.750.300.184 Community Networks
-	L01.313.500.750.300.188 Databases as Topic
-	L01.313.500.750.300.188.300 Databases, Bibliographic
-	L01.313.500.750.300.188.300.650 PubMed
-	L01.313.500.750.300.188.300.650.500 MEDLINE
-	L01.313.500.750.300.188.400 Databases, Factual
-	L01.313.500.750.300.188.400.300 Databases, Chemical
-	L01.313.500.750.300.188.400.300.500 Databases, Nucleic Acid
-	L01.313.500.750.300.188.400.300.750 Databases, Protein
-	L01.313.500.750.300.188.400.325 Databases, Genetic
-	L01.313.500.750.300.188.400.325.630 Databases, Nucleic Acid
-	L01.313.500.750.300.188.400.325.710 Databases, Protein
-	L01.313.500.750.300.188.400.400 Databases, Pharmaceutical
-	L01.313.500.750.300.188.400.500 Datasets as Topic
-	L01.313.500.750.300.188.400.600 National Practitioner Data Bank
-	L01.313.500.750.300.188.400.905 Visible Human Projects
-	L01.313.500.750.300.190 Decision Support Systems, Clinical
-	L01.313.500.750.300.314 Geographic Information Systems
-	L01.313.500.750.300.361 Health Information Systems
-	L01.313.500.750.300.420 Integrated Advanced Information Management Systems
-	L01.313.500.750.300.550 Knowledge Bases
-	L01.313.500.750.300.550.500 Biological Ontologies
-	L01.313.500.750.300.550.500.500 Gene Ontology
-	L01.313.500.750.300.680 Management Information Systems
-	L01.313.500.750.300.680.055 Clinical Laboratory Information Systems

## MeSH Tree Changes for 2017

Type	Tree - heading
-	L01.313.500.750.300.680.083 Clinical Pharmacy Information Systems
-	L01.313.500.750.300.680.110 Database Management Systems
-	L01.313.500.750.300.680.135 Decision Support Systems, Management
-	L01.313.500.750.300.680.350 Healthcare Common Procedure Coding System
-	L01.313.500.750.300.680.360 Hospital Information Systems
-	L01.313.500.750.300.680.360.113 Ambulatory Care Information Systems
-	L01.313.500.750.300.680.360.450 Medical Order Entry Systems
-	L01.313.500.750.300.680.360.500 Operating Room Information Systems
-	L01.313.500.750.300.680.580 Office Automation
-	L01.313.500.750.300.680.580.950 Word Processing
-	L01.313.500.750.300.680.800 Personnel Staffing and Scheduling Information Systems
-	L01.313.500.750.300.680.900 Radiology Information Systems
-	L01.313.500.750.300.680.900.500 Teleradiology
-	L01.313.500.750.300.695 Medical Records Systems, Computerized
-	L01.313.500.750.300.695.300 Health Smart Cards
-	L01.313.500.750.300.695.600 Medical Order Entry Systems
-	L01.313.500.750.300.710 MEDLARS
-	L01.313.500.750.300.710.500 MEDLINE
-	L01.313.500.750.300.742 Online Systems
-	L01.313.500.750.300.742.470 Libraries, Digital
-	L01.313.500.750.300.742.650 PubMed
-	L01.313.500.750.300.742.650.500 MEDLINE
-	L01.313.500.750.300.790 Reminder Systems
-	L01.313.500.875 Medical Informatics Computing
-	L01.313.500.875.500 Data Curation
-	L01.313.650 Nursing Informatics
-	L01.313.750 Public Health Informatics
-	L01.346 Information Centers
-	L01.346.208 Archives
-	L01.346.596 Libraries
-	L01.346.596.363 Libraries, Dental
-	L01.346.596.413 Libraries, Digital
-	L01.346.596.463 Libraries, Hospital
-	L01.346.596.563 Libraries, Medical
-	L01.346.596.563.725 National Library of Medicine (U.S.)

## MeSH Tree Changes for 2017

Type	Tree - heading
-	L01.346.596.663 Libraries, Nursing
-	L01.346.596.831 Libraries, Special
-	L01.399 Information Management
-	L01.399.500 Health Information Management
-	L01.399.500.500 Health Information Exchange
-	L01.453 Information Services
-	L01.453.183 Bibliography as Topic
-	L01.453.183.190 Bibliography, Descriptive
-	L01.453.183.224 Bibliography of Medicine
-	L01.453.183.291 Bibliometrics
-	L01.453.183.291.500 Journal Impact Factor
-	L01.453.183.392 Biobibliography as Topic
-	L01.453.220 Book Selection
-	L01.453.245 Documentation
-	L01.453.245.100 Abstracting and Indexing as Topic
-	L01.453.245.250 Cataloging
-	L01.453.245.250.150 Book Classification
-	L01.453.245.275 Classification
-	L01.453.245.332 Data Curation
-	L01.453.245.390 Filing
-	L01.453.245.667 Molecular Sequence Data
-	L01.453.245.667.060 Amino Acid Sequence
-	L01.453.245.667.080 Base Sequence
-	L01.453.245.667.160 Carbohydrate Sequence
-	L01.453.245.667.580 Molecular Sequence Annotation
-	L01.453.245.945 Vocabulary, Controlled
-	L01.453.245.945.079 Biological Ontologies
-	L01.453.245.945.079.500 Gene Ontology
-	L01.453.245.945.160 Current Procedural Terminology
-	L01.453.245.945.200 Diagnostic and Statistical Manual of Mental Disorders
-	L01.453.245.945.380 Healthcare Common Procedure Coding System
-	L01.453.245.945.400 International Classification of Diseases
-	L01.453.245.945.450 International Classification of Functioning, Disability and Health
-	L01.453.245.945.500 Logical Observation Identifiers Names and Codes
New	<b>L01.453.245.945.600 Standardized Nursing Terminology</b>

## MeSH Tree Changes for 2017

Type	Tree - heading
Heading	
-	L01.453.245.945.700 Subject Headings
-	L01.453.245.945.700.500 Medical Subject Headings
-	L01.453.245.945.720 Systematized Nomenclature of Medicine
-	L01.453.245.945.800 Unified Medical Language System
-	L01.453.270 Drug Information Services
-	L01.453.270.120 Adverse Drug Reaction Reporting Systems
-	L01.453.270.200 Clinical Pharmacy Information Systems
-	L01.453.270.600 Pharmacovigilance
-	L01.453.450 Human Genome Project
-	L01.453.639 Library Services
-	L01.453.639.493 Interlibrary Loans
-	L01.453.819 Library Technical Services
-	L01.453.819.250 Cataloging
-	L01.453.819.250.150 Book Classification
-	L01.470 Information Storage and Retrieval
-	L01.470.500 Data Compression
-	L01.470.625 Data Mining
-	L01.470.625.500 Multifactor Dimensionality Reduction
-	L01.470.750 Databases as Topic
-	L01.470.750.500 Databases, Bibliographic
-	L01.470.750.500.650 PubMed
-	L01.470.750.500.650.500 MEDLINE
-	L01.470.750.750 Databases, Factual
-	L01.470.750.750.300 Databases, Chemical
-	L01.470.750.750.300.500 Databases, Nucleic Acid
-	L01.470.750.750.300.750 Databases, Protein
-	L01.470.750.750.325 Databases, Genetic
-	L01.470.750.750.325.630 Databases, Nucleic Acid
-	L01.470.750.750.325.710 Databases, Protein
-	L01.470.750.750.400 Databases, Pharmaceutical
-	L01.470.750.750.431 Datasets as Topic
-	L01.470.750.750.462 Geographic Information Systems
-	L01.470.750.750.600 National Practitioner Data Bank
-	L01.470.750.750.905 Visible Human Projects
-	L01.470.875 Search Engine

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	L01.470.937	Web Browser
-	L01.488	Information Theory
-	L01.535	Knowledge
-	L01.583	Library Science
-	L01.583.363	Library Administration
-	L01.583.390	Library Associations
-	L01.583.400	Library Automation
-	L01.583.410	Library Collection Development
-	L01.583.430	Library Schools
-	L01.583.464	Library Services
-	L01.583.464.493	Interlibrary Loans
-	L01.583.564	Library Surveys
-	L01.583.782	Library Technical Services
New Heading	<b>L01.654</b>	<b>Metadata</b>
-	L01.725	Pattern Recognition, Automated
-	L01.725.249	Biometric Identification
-	L01.725.500	Neural Networks (Computer)
-	L01.725.750	Support Vector Machine
New Heading	<b>L01.731</b>	<b>Postcards as Topic</b>
-	L01.737	Publishing
-	L01.737.030	Access to Information
-	L01.737.150	Book Industry
-	L01.737.150.317	Bookbinding
-	L01.737.150.340	Bookselling
-	L01.737.150.340.250	Book Prices
-	L01.737.360	Copyright
-	L01.737.471	Duplicate Publication as Topic
-	L01.737.484	Editorial Policies
-	L01.737.498	Journalism
-	L01.737.498.500	Journalism, Dental
-	L01.737.498.550	Journalism, Medical
-	L01.737.569	Open Access Publishing
-	L01.737.640	Peer Review, Research
-	L01.737.684	Plagiarism

## MeSH Tree Changes for 2017

Type	Tree - heading
-	L01.737.787                      Printing
-	L01.737.813                      Publication Bias
-	L01.737.840                      Retraction of Publication as Topic
-	L01.906                              Systems Analysis
-	L01.906.575                      Operations Research
-	L01.906.575.422                  Monte Carlo Method
-	L01.906.575.622                  Probability Theory
-	L01.906.575.748                  Programming, Linear
-	L01.906.787                      Systems Integration
-	L01.906.893                      Workflow
-	M01                                  Persons
-	M01.050                              Abortion Applicants
-	M01.055                              Adult Children
-	M01.060                              Age Groups
-	M01.060.057                      Adolescent
-	M01.060.116                      Adult
-	M01.060.116.100                  Aged
-	M01.060.116.100.080              Aged, 80 and over
-	M01.060.116.100.540              Frail Elderly
-	M01.060.116.630                  Middle Aged
-	M01.060.116.815                  Young Adult
-	M01.060.406                      Child
-	M01.060.406.448                  Child, Preschool
-	M01.060.703                      Infant
-	M01.060.703.520                  Infant, Newborn
-	M01.060.703.520.460              Infant, Low Birth Weight
-	M01.060.703.520.460.560          Infant, Small for Gestational Age
-	M01.060.703.520.460.600          Infant, Very Low Birth Weight
-	M01.060.703.520.460.600.500      Infant, Extremely Low Birth Weight
-	M01.060.703.520.500              Infant, Postmature
-	M01.060.703.520.520              Infant, Premature
-	M01.060.703.520.520.500          Infant, Extremely Premature
-	M01.066                              Alcoholics
-	M01.072                              Athletes
-	M01.085                              Caregivers
-	M01.097                              Child, Abandoned

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	M01.102	Child, Exceptional
-	M01.102.350	Child, Gifted
-	M01.106	Child of Impaired Parents
-	M01.108	Child, Orphaned
-	M01.111	Child, Unwanted
-	M01.120	Consultants
-	M01.135	Crime Victims
-	M01.135.500	Adult Survivors of Child Abuse
-	M01.142	Criminals
-	M01.150	Disabled Persons
-	M01.150.100	Amputees
-	M01.150.200	Disabled Children
-	M01.150.600	Mentally Disabled Persons
-	M01.150.725	Mentally Ill Persons
-	M01.150.750	Persons With Hearing Impairments
-	M01.150.850	Visually Impaired Persons
-	M01.159	Disaster Victims
-	M01.169	Drug Users
-	M01.189	Emigrants and Immigrants
-	M01.189.500	Undocumented Immigrants
-	M01.228	Famous Persons
-	M01.252	Friends
-	M01.264	Grandparents
-	M01.276	Homebound Persons
-	M01.325	Homeless Persons
-	M01.325.400	Homeless Youth
-	M01.352	Jehovah's Witnesses
-	M01.380	Legal Guardians
-	M01.380.600	Proxy
-	M01.385	Medically Uninsured
-	M01.390	Men
-	M01.390.660	Nurses, Male
-	M01.395	Mentors
-	M01.416	Minors
-	M01.427	Missionaries
-	M01.438	Multiple Birth Offspring

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	M01.438.486	Quadruplets
-	M01.438.587	Quintuplets
-	M01.438.768	Triplets
-	M01.438.873	Twins
-	M01.438.873.920	Twins, Dizygotic
-	M01.438.873.940	Twins, Monozygotic
-	M01.526	Occupational Groups
-	M01.526.070	Administrative Personnel
New Heading	<b>M01.526.070.245</b>	<b>Case Managers</b>
-	M01.526.070.490	Health Facility Administrators
-	M01.526.070.490.490	Hospital Administrators
-	M01.526.070.490.490.100	Chief Executive Officers, Hospital
-	M01.526.070.670	Nurse Administrators
-	M01.526.070.700	Physician Executives
-	M01.526.070.940	Trustees
New Tree	<a href="#">M01.526.173</a>	<a href="#">Astronauts</a>
New Heading	<b>M01.526.225</b>	<b>Counselors</b>
-	M01.526.276	Ethicists
Old Tree	<b>M01.526.339</b>	<b>Faculty</b>
Old Tree	<b>M01.526.339.273</b>	<b>Faculty, Dental</b>
Old Tree	<b>M01.526.339.373</b>	<b>Faculty, Medical</b>
Old Tree	<b>M01.526.339.473</b>	<b>Faculty, Nursing</b>
-	M01.526.373	Emergency Responders
-	M01.526.373.250	Emergency Medical Technicians
-	M01.526.373.400	Firefighters
-	M01.526.373.750	Police
-	M01.526.390	Farmers
-	M01.526.407	Foreign Professional Personnel
-	M01.526.407.435	Foreign Medical Graduates
-	M01.526.407.717	Nurses, International
New Heading	<b>M01.526.446</b>	<b>Government Employees</b>
-	M01.526.485	Health Personnel
-	M01.526.485.067	Allied Health Personnel



## MeSH Tree Changes for 2017

Type	Tree - heading	
-	M01.526.485.067.040	Animal Technicians
-	M01.526.485.067.080	Community Health Workers
-	M01.526.485.067.105	Dental Auxiliaries
-	M01.526.485.067.105.250	Dental Assistants
-	M01.526.485.067.105.376	Dental Hygienists
-	M01.526.485.067.105.503	Dental Technicians
-	M01.526.485.067.105.751	Denturists
-	M01.526.485.067.150	Emergency Medical Technicians
-	M01.526.485.067.350	Home Health Aides
-	M01.526.485.067.450	Licensed Practical Nurses
-	M01.526.485.067.550	Medical Record Administrators
-	M01.526.485.067.600	Medical Secretaries
-	M01.526.485.067.600.630	Medical Receptionists
-	M01.526.485.067.652	Nurses' Aides
-	M01.526.485.067.652.628	Psychiatric Aides
Old Tree	<b>M01.526.485.067.676</b>	<b>Nutritionists</b>
-	M01.526.485.067.700	Operating Room Technicians
-	M01.526.485.067.735	Pharmacists' Aides
-	M01.526.485.067.735	Pharmacy Technicians
-	M01.526.485.067.738	Physical Therapist Assistants
Old Tree	<b>M01.526.485.067.739</b>	<b>Physical Therapists</b>
-	M01.526.485.067.740	Physician Assistants
-	M01.526.485.067.740.690	Ophthalmic Assistants
-	M01.526.485.067.740.750	Pediatric Assistants
-	M01.526.485.133	Anatomists
New Heading	<b>M01.526.485.140</b>	<b>Anesthetists</b>
New Heading	<b>M01.526.485.140.040</b>	<b>Anesthesiologists</b>
New Tree	<b>M01.526.485.140.650</b>	<b>Nurse Anesthetists</b>
New Heading	<b>M01.526.485.170</b>	<b>Audiologists</b>
-	M01.526.485.200	Caregivers
New Heading	<b>M01.526.485.215</b>	<b>Case Managers</b>
-	M01.526.485.230	Coroners and Medical Examiners

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	M01.526.485.290	Dental Staff
-	M01.526.485.290.490	Dental Staff, Hospital
-	M01.526.485.330	Dentists
-	M01.526.485.330.310	Dentists, Women
New Heading	<b>M01.526.485.330.397</b>	<b>Endodontists</b>
-	M01.526.485.330.483	Oral and Maxillofacial Surgeons
-	M01.526.485.330.655	Orthodontists
-	M01.526.485.345	Doulas
New Heading	<b>M01.526.485.349</b>	<b>Emergency Medical Dispatcher</b>
New Heading	<b>M01.526.485.353</b>	<b>Epidemiologists</b>
-	M01.526.485.360	Faculty, Dental
-	M01.526.485.375	Faculty, Medical
-	M01.526.485.390	Faculty, Nursing
-	M01.526.485.410	Health Educators
-	M01.526.485.430	Health Facility Administrators
-	M01.526.485.430.490	Hospital Administrators
-	M01.526.485.430.490.100	Chief Executive Officers, Hospital
-	M01.526.485.490	Infection Control Practitioners
-	M01.526.485.500	Medical Chaperones
-	M01.526.485.510	Medical Laboratory Personnel
-	M01.526.485.630	Medical Staff
-	M01.526.485.630.490	Medical Staff, Hospital
-	M01.526.485.630.490.400	Hospitalists
-	M01.526.485.650	Nurses
-	M01.526.485.650.580	Nurse Administrators
Old Tree	<b>M01.526.485.650.600</b>	<b>Nurse Anesthetists</b>
Old Tree	<b>M01.526.485.650.610</b>	<b>Nurse Clinicians</b>
Old Tree	<b>M01.526.485.650.620</b>	<b>Nurse Midwives</b>
-	M01.526.485.650.640	Nurse Practitioners
-	M01.526.485.650.640.249	Family Nurse Practitioners
-	M01.526.485.650.640.500	Pediatric Nurse Practitioners
New Heading	<b>M01.526.485.650.648</b>	<b>Nurse Specialists</b>
New	<b>M01.526.485.650.648.500</b>	<b>Nurse Anesthetists</b>

## MeSH Tree Changes for 2017

Type	Tree - heading	
Tree		
New Tree	<a href="#">M01.526.485.650.648.525</a>	<a href="#">Nurse Clinicians</a>
New Tree	<a href="#">M01.526.485.650.648.762</a>	<a href="#">Nurse Midwives</a>
New Heading	<b>M01.526.485.650.648.940</b>	<b>Nurses, Pediatric</b>
New Heading	<b>M01.526.485.650.648.940.500</b>	<b>Nurses, Neonatal</b>
-	M01.526.485.650.655	Nurses, Community Health
-	M01.526.485.650.662	Nurses, International
-	M01.526.485.650.670	Nurses, Male
-	M01.526.485.650.835	Nurses, Public Health
-	M01.526.485.680	Nursing Staff
-	M01.526.485.680.490	Nursing Staff, Hospital
New Tree	<a href="#">M01.526.485.695</a>	<a href="#">Nutritionists</a>
New Heading	<b>M01.526.485.710</b>	<b>Occupational Therapists</b>
New Heading	<b>M01.526.485.725</b>	<b>Optometrists</b>
-	M01.526.485.740	Personnel, Hospital
-	M01.526.485.740.322	Dental Staff, Hospital
-	M01.526.485.740.380	Hospital Administrators
-	M01.526.485.740.380.100	Chief Executive Officers, Hospital
-	M01.526.485.740.400	Hospital Volunteers
-	M01.526.485.740.422	Medical Staff, Hospital
-	M01.526.485.740.422.400	Hospitalists
-	M01.526.485.740.523	Nursing Staff, Hospital
-	M01.526.485.780	Pharmacists
New Tree	<a href="#">M01.526.485.790</a>	<a href="#">Physical Therapists</a>
-	M01.526.485.800	Physician Executives
-	M01.526.485.810	Physicians
New Heading	<b>M01.526.485.810.020</b>	<b>Allergists</b>
New Heading	<b>M01.526.485.810.040</b>	<b>Anesthesiologists</b>
New Heading	<b>M01.526.485.810.128</b>	<b>Cardiologists</b>

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Heading	<b>M01.526.485.810.215</b>	<b>Dermatologists</b>
New Heading	<b>M01.526.485.810.303</b>	<b>Endocrinologists</b>
-	M01.526.485.810.390	Foreign Medical Graduates
New Heading	<b>M01.526.485.810.438</b>	<b>Gastroenterologists</b>
-	M01.526.485.810.485	General Practitioners
New Heading	<b>M01.526.485.810.533</b>	<b>Geriatricians</b>
-	M01.526.485.810.580	Hospitalists
New Heading	<b>M01.526.485.810.628</b>	<b>Nephrologists</b>
New Heading	<b>M01.526.485.810.652</b>	<b>Neurologists</b>
-	M01.526.485.810.675	Occupational Health Physicians
New Heading	<b>M01.526.485.810.699</b>	<b>Oncologists</b>
New Heading	<b>M01.526.485.810.699.500</b>	<b>Radiation Oncologists</b>
New Heading	<b>M01.526.485.810.705</b>	<b>Ophthalmologists</b>
-	M01.526.485.810.722	Osteopathic Physicians
New Heading	<b>M01.526.485.810.734</b>	<b>Otolaryngologists</b>
New Heading	<b>M01.526.485.810.746</b>	<b>Pathologists</b>
New Heading	<b>M01.526.485.810.758</b>	<b>Pediatricians</b>
New Heading	<b>M01.526.485.810.758.500</b>	<b>Neonatologists</b>
New Heading	<b>M01.526.485.810.764</b>	<b>Physiatrists</b>
-	M01.526.485.810.770	Physicians, Family
-	M01.526.485.810.800	Physicians, Primary Care
-	M01.526.485.810.820	Physicians, Women
New Heading	<b>M01.526.485.810.865</b>	<b>Pulmonologists</b>
New Heading	<b>M01.526.485.810.877</b>	<b>Radiologists</b>
New Heading	<b>M01.526.485.810.877.500</b>	<b>Radiation Oncologists</b>

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Heading	<b>M01.526.485.810.888</b>	<b>Rheumatologists</b>
-	M01.526.485.810.910	Surgeons
-	M01.526.485.810.910.500	Barber Surgeons
-	M01.526.485.810.910.750	Neurosurgeons
New Heading	<b>M01.526.485.810.910.875</b>	<b>Orthopedic Surgeons</b>
New Heading	<b>M01.526.485.810.955</b>	<b>Urologists</b>
-	M01.526.485.905	Veterinarians
-	M01.526.493	Inventors
-	M01.526.502	Laboratory Personnel
-	M01.526.502.500	Medical Laboratory Personnel
-	M01.526.520	Lawyers
-	M01.526.555	Librarians
-	M01.526.625	Military Personnel
-	M01.526.693	Miners
New Heading	<b>M01.526.702</b>	<b>Educational Personnel</b>
New Tree	<a href="#">M01.526.702.250</a>	<a href="#">Faculty</a>
New Tree	<a href="#">M01.526.702.250.273</a>	<a href="#">Faculty, Dental</a>
New Tree	<a href="#">M01.526.702.250.373</a>	<a href="#">Faculty, Medical</a>
New Tree	<a href="#">M01.526.702.250.473</a>	<a href="#">Faculty, Nursing</a>
New Heading	<b>M01.526.702.250.736</b>	<b>Faculty, Pharmacy</b>
New Heading	<b>M01.526.702.500</b>	<b>School Teachers</b>
New Heading	<b>M01.526.727</b>	<b>Pilots</b>
-	M01.526.760	Police
-	M01.526.799	Religious Personnel
-	M01.526.799.500	Clergy
-	M01.526.799.750	Monks
-	M01.526.799.875	Nuns
-	M01.526.839	Research Personnel
Old Tree	<b>M01.526.839.500</b>	<b>Astronauts</b>

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	M01.526.919	Social Workers
-	M01.620	Parents
-	M01.620.390	Fathers
-	M01.620.630	Mothers
-	M01.620.785	Single Parent
-	M01.620.892	Surrogate Mothers
-	M01.643	Patients
-	M01.643.050	Adolescent, Hospitalized
-	M01.643.154	Adolescent, Institutionalized
-	M01.643.259	Child, Hospitalized
-	M01.643.364	Child, Institutionalized
-	M01.643.470	Inpatients
-	M01.643.630	Outpatients
-	M01.665	Pedestrians
-	M01.686	Population Groups
-	M01.686.508	Continental Population Groups
-	M01.686.508.100	African Continental Ancestry Group
-	M01.686.508.100.100	African Americans
-	M01.686.508.150	American Native Continental Ancestry Group
New Heading	<b>M01.686.508.150.288</b>	<b>Alaska Natives</b>
-	M01.686.508.150.575	Indians, Central American
-	M01.686.508.150.600	Indians, North American
-	M01.686.508.150.625	Indians, South American
-	M01.686.508.150.675	Inuits
-	M01.686.508.200	Asian Continental Ancestry Group
-	M01.686.508.200.100	Asian Americans
-	M01.686.508.400	European Continental Ancestry Group
-	M01.686.508.600	Oceanic Ancestry Group
-	M01.686.754	Ethnic Groups
-	M01.686.754.100	African Americans
-	M01.686.754.133	Amish
-	M01.686.754.167	Arabs
-	M01.686.754.225	Asian Americans
-	M01.686.754.283	Roma
-	M01.686.754.441	Hispanic Americans

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	M01.686.754.441.500	Mexican Americans
-	M01.686.754.520	Inuits
-	M01.686.754.600	Jews
-	M01.729	Prisoners
-	M01.729.500	Prisoners of War
-	M01.755	Refugees
-	M01.774	Research Subjects
-	M01.774.500	Healthy Volunteers
-	M01.776	Sex Workers
New Heading	<b>M01.777</b>	<b>Sexual Minorities</b>
New Tree	<b>M01.777.500</b>	<b>Transgender Persons</b>
-	M01.778	Sexual Partners
-	M01.781	Siblings
-	M01.785	Single Person
-	M01.800	Slaves
-	M01.816	Spouses
-	M01.848	Students
-	M01.848.602	Student Dropouts
-	M01.848.769	Students, Health Occupations
-	M01.848.769.519	Students, Dental
-	M01.848.769.602	Students, Medical
-	M01.848.769.685	Students, Nursing
-	M01.848.769.768	Students, Pharmacy
-	M01.848.769.851	Students, Premedical
-	M01.848.769.925	Students, Public Health
-	M01.860	Survivors
-	M01.860.300	Adult Survivors of Child Adverse Events
-	M01.860.300.500	Adult Survivors of Child Abuse
-	M01.860.400	HIV Long-Term Survivors
-	M01.873	Terminally Ill
-	M01.898	Tissue Donors
-	M01.898.313	Blood Donors
-	M01.898.656	Living Donors
-	M01.898.828	Unrelated Donors

## MeSH Tree Changes for 2017

Type	Tree - heading	
Old Tree	M01.909	Transgender Persons
-	M01.920	Transients and Migrants
-	M01.925	Transplant Recipients
-	M01.928	Vegetarians
-	M01.928.500	Vegans
-	M01.930	Veterans
-	M01.935	Visitors to Patients
-	M01.955	Volunteers
-	M01.955.236	Healthy Volunteers
-	M01.955.473	Hospital Volunteers
-	M01.965	Vulnerable Populations
-	M01.975	Women
-	M01.975.155	Battered Women
-	M01.975.310	Dentists, Women
-	M01.975.790	Physicians, Women
-	M01.975.807	Pregnant Women
-	M01.975.825	Women, Working
-	M01.987	Working Poor
-	N01	Population Characteristics
-	N01.224	Demography
-	N01.224.033	Age Distribution
-	N01.224.175	Censuses
-	N01.224.317	Ethnic Groups
-	N01.224.361	Family Characteristics
-	N01.224.361.150	Birth Intervals
-	N01.224.361.160	Birth Order
-	N01.224.361.500	Marital Status
-	N01.224.361.500.300	Divorce
-	N01.224.361.500.500	Marriage
-	N01.224.361.500.725	Single Person
-	N01.224.361.500.725.700	Single Parent
-	N01.224.361.500.862	Widowhood
-	N01.224.425	Health Status
-	N01.224.425.350	Geriatric Assessment
-	N01.224.425.437	Health Status Disparities
-	N01.224.425.525	Nutritional Status



## MeSH Tree Changes for 2017

Type	Tree - heading
-	N01.224.425.762 Social Determinants of Health
-	N01.224.600 Population Density
-	N01.224.625 Population Dynamics
-	N01.224.625.400 Health Transition
-	N01.224.625.525 Human Migration
-	N01.224.625.525.500 Emigration and Immigration
-	N01.224.625.650 Population Control
-	N01.224.625.660 Population Growth
-	N01.224.667 Population Forecast
-	N01.224.708 Population Groups
-	N01.224.791 Residence Characteristics
-	N01.224.791.200 Catchment Area (Health)
-	N01.224.791.400 Housing
-	N01.224.791.400.410 Housing for the Elderly
-	N01.224.791.400.650 Public Housing
-	N01.224.791.550 Independent Living
-	N01.224.803 Sex Distribution
-	N01.224.803.815 Sex Ratio
-	N01.224.935 Vital Statistics
-	N01.224.935.464 Life Expectancy
-	N01.224.935.530 Life Tables
-	N01.224.935.530.700 Quality-Adjusted Life Years
-	N01.224.935.597 Morbidity
-	N01.224.935.597.080 Basic Reproduction Number
-	N01.224.935.597.500 Incidence
-	N01.224.935.597.750 Prevalence
-	N01.224.935.698 Mortality
-	N01.224.935.698.100 Cause of Death
-	N01.224.935.698.150 Child Mortality
-	N01.224.935.698.201 Fatal Outcome
-	N01.224.935.698.300 Fetal Mortality
-	N01.224.935.698.400 Hospital Mortality
-	N01.224.935.698.489 Infant Mortality
New Tree	<a href="#">N01.224.935.698.489.500</a> Perinatal Mortality
-	N01.224.935.698.653 Maternal Mortality

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	N01.224.935.698.700	Mortality, Premature
Old Tree	<b>N01.224.935.698.739</b>	<b>Perinatal Mortality</b>
-	N01.224.935.698.826	Survival Rate
-	N01.224.935.849	Pregnancy Rate
-	N01.224.935.849.500	Birth Rate
-	N01.400	Health
-	N01.400.075	Adolescent Health
New Heading	<b>N01.400.150</b>	<b>Cardiorespiratory Fitness</b>
-	N01.400.225	Child Health
-	N01.400.300	Family Health
-	N01.400.337	Global Health
-	N01.400.350	Holistic Health
-	N01.400.388	Infant Health
-	N01.400.425	Men's Health
-	N01.400.500	Mental Health
-	N01.400.512	Minority Health
-	N01.400.525	Occupational Health
-	N01.400.535	Oral Health
-	N01.400.545	Physical Fitness
New Heading	<b>N01.400.545.500</b>	<b>Cardiorespiratory Fitness</b>
-	N01.400.550	Public Health
-	N01.400.625	Reproductive Health
-	N01.400.650	Rural Health
-	N01.400.675	Social Determinants of Health
-	N01.400.700	Suburban Health
-	N01.400.800	Urban Health
-	N01.400.850	Veterans Health
-	N01.400.900	Women's Health
-	N01.400.900.500	Maternal Health
-	N01.600	Population
-	N01.600.725	Rural Population
-	N01.600.775	Suburban Population
-	N01.600.900	Urban Population
-	N01.824	Socioeconomic Factors

## MeSH Tree Changes for 2017

Type	Tree - heading
-	N01.824.196 Educational Status
-	N01.824.196.500 Literacy
-	N01.824.245 Employment
-	N01.824.245.175 Career Mobility
-	N01.824.245.350 Employment, Supported
-	N01.824.245.600 Personnel Downsizing
-	N01.824.245.725 Return to Work
-	N01.824.245.850 Unemployment
-	N01.824.245.925 Workplace
-	N01.824.308 Family Characteristics
-	N01.824.308.500 Marital Status
-	N01.824.308.500.300 Divorce
-	N01.824.308.500.500 Marriage
-	N01.824.308.500.725 Single Person
-	N01.824.308.500.725.700 Single Parent
-	N01.824.308.500.862 Widowhood
-	N01.824.417 Income
-	N01.824.417.510 Pensions
-	N01.824.417.510.300 Employee Retirement Income Security Act
-	N01.824.417.605 Remuneration
-	N01.824.417.700 Salaries and Fringe Benefits
-	N01.824.417.700.162 Family Leave
-	N01.824.417.700.162.500 Parental Leave
-	N01.824.417.700.325 Health Benefit Plans, Employee
-	N01.824.417.700.325.300 Employee Retirement Income Security Act
-	N01.824.417.700.662 Sick Leave
-	N01.824.460 Medical Indigency
-	N01.824.547 Occupations
-	N01.824.547.330 Career Mobility
-	N01.824.600 Poverty
-	N01.824.600.550 Poverty Areas
-	N01.824.737 Social Change
-	N01.824.782 Social Class
-	N01.824.782.673 Social Mobility
-	N01.824.827 Social Conditions
-	N02 Health Care Facilities, Manpower, and Services

## MeSH Tree Changes for 2017

Type	Tree - heading
-	N02.138 Capacity Building
-	N02.208 Health Communication
-	N02.278 Health Facilities
-	N02.278.020 Academic Medical Centers
-	N02.278.020.300 Hospitals, Teaching
-	N02.278.020.300.310 Hospitals, University
-	N02.278.020.578 Schools, Medical
-	N02.278.035 Ambulatory Care Facilities
-	N02.278.035.128 Community Health Centers
-	N02.278.035.128.800 Substance Abuse Treatment Centers
-	N02.278.035.158 Community Mental Health Centers
-	N02.278.035.158.300 Child Guidance Clinics
-	N02.278.035.310 Maternal-Child Health Centers
-	N02.278.035.380 Outpatient Clinics, Hospital
-	N02.278.035.380.600 Pain Clinics
-	N02.278.035.652 Surgicenters
-	N02.278.050 Bed Occupancy
-	N02.278.065 Biological Specimen Banks
-	N02.278.065.200 Blood Banks
-	N02.278.065.600 Milk Banks
-	N02.278.065.650 Seed Bank
-	N02.278.065.700 Sperm Banks
-	N02.278.065.900 Tissue Banks
-	N02.278.065.900.205 Bone Banks
-	N02.278.065.900.400 Eye Banks
-	N02.278.080 Birthing Centers
-	N02.278.192 Dental Facilities
-	N02.278.192.250 Dental Clinics
-	N02.278.192.377 Dental Offices
-	N02.278.192.631 Laboratories, Dental
-	N02.278.200 Facility Design and Construction
-	N02.278.200.100 Building Codes
-	N02.278.200.252 Evidence-Based Facility Design
-	N02.278.200.403 Hospital Design and Construction
-	N02.278.207 Fitness Centers
-	N02.278.215 Health Facilities, Proprietary

## MeSH Tree Changes for 2017

Type	Tree - heading
-	N02.278.215.110                      Hospitals, Proprietary
-	N02.278.216                              Health Facility Administration
-	N02.278.216.500                        Hospital Administration
-	N02.278.216.500.500                    Ancillary Services, Hospital
-	N02.278.216.500.750                    Centralized Hospital Services
-	N02.278.216.500.875                    Financial Management, Hospital
-	N02.278.216.500.937                    Hospital Communication Systems
-	N02.278.216.500.968                    Hospital Departments
-	N02.278.216.500.968.040                Admitting Department, Hospital
-	N02.278.216.500.968.070                Anesthesia Department, Hospital
-	N02.278.216.500.968.215                Cardiology Service, Hospital
-	N02.278.216.500.968.260                Central Supply, Hospital
-	N02.278.216.500.968.270                Chaplaincy Service, Hospital
-	N02.278.216.500.968.298                Dental Service, Hospital
-	N02.278.216.500.968.310                Education Department, Hospital
-	N02.278.216.500.968.336                Emergency Service, Hospital
-	N02.278.216.500.968.336.500            Trauma Centers
-	N02.278.216.500.968.374                Food Service, Hospital
-	N02.278.216.500.968.412                Housekeeping, Hospital
-	N02.278.216.500.968.420                Laboratories, Hospital
-	N02.278.216.500.968.428                Laundry Service, Hospital
-	N02.278.216.500.968.450                Maintenance and Engineering, Hospital
-	N02.278.216.500.968.465                Medical Records Department, Hospital
New Heading	<b>N02.278.216.500.968.467                Morgue</b>
-	N02.278.216.500.968.472                Nuclear Medicine Department, Hospital
-	N02.278.216.500.968.489                Nursing Service, Hospital
-	N02.278.216.500.968.495                Obstetrics and Gynecology Department, Hospital
-	N02.278.216.500.968.500                Occupational Therapy Department, Hospital
-	N02.278.216.500.968.513                Oncology Service, Hospital
-	N02.278.216.500.968.527                Outpatient Clinics, Hospital
-	N02.278.216.500.968.527.600            Pain Clinics
-	N02.278.216.500.968.535                Pathology Department, Hospital
-	N02.278.216.500.968.565                Personnel Administration, Hospital
-	N02.278.216.500.968.603                Pharmacy Service, Hospital
-	N02.278.216.500.968.615                Physical Therapy Department, Hospital

## MeSH Tree Changes for 2017

Type	Tree - heading
-	N02.278.216.500.968.641      Psychiatric Department, Hospital
-	N02.278.216.500.968.679      Purchasing, Hospital
-	N02.278.216.500.968.679.375      Group Purchasing
-	N02.278.216.500.968.695      Radiology Department, Hospital
-	N02.278.216.500.968.712      Respiratory Therapy Department, Hospital
-	N02.278.216.500.968.730      Social Work Department, Hospital
-	N02.278.216.500.968.750      Surgery Department, Hospital
-	N02.278.216.500.968.790      Urology Department, Hospital
-	N02.278.216.500.984      Hospital Distribution Systems
-	N02.278.216.500.988      Hospital Restructuring
-	N02.278.216.500.988.380      Hospital-Physician Joint Ventures
-	N02.278.216.500.988.380.500      Physician Self-Referral
-	N02.278.216.500.990      Hospital Shared Services
-	N02.278.216.500.991      Hospital Shops
-	N02.278.216.500.992      Hospital-Patient Relations
-	N02.278.216.500.996      Hospital-Physician Relations
-	N02.278.216.500.996.500      Medical Staff Privileges
-	N02.278.216.500.998      Libraries, Hospital
-	N02.278.216.500.999      Materials Management, Hospital
-	N02.278.216.500.999.300      Inventories, Hospital
-	N02.278.218      Health Facility Closure
-	N02.278.220      Health Facility Environment
-	N02.278.220.640      Patients' Rooms
-	N02.278.235      Health Facility Merger
-	N02.278.255      Health Facility Moving
-	N02.278.306      Health Facility Size
-	N02.278.306.472      Hospital Bed Capacity
-	N02.278.306.472.080      Hospital Bed Capacity, under 100
-	N02.278.306.472.120      Hospital Bed Capacity, 100 to 299
-	N02.278.306.472.180      Hospital Bed Capacity, 300 to 499
-	N02.278.306.472.300      Hospital Bed Capacity, 500 and over
New Tree	<a href="#">N02.278.330</a> <a href="#">Health Resorts</a>
-	N02.278.354      Hospital Administration
-	N02.278.354.675      Medication Systems, Hospital
-	N02.278.354.700      Product Line Management

## MeSH Tree Changes for 2017

Type	Tree - heading
-	N02.278.388 Hospital Units
-	N02.278.388.150 Delivery Rooms
-	N02.278.388.200 Hemodialysis Units, Hospital
-	N02.278.388.493 Intensive Care Units
-	N02.278.388.493.160 Burn Units
-	N02.278.388.493.211 Coronary Care Units
-	N02.278.388.493.390 Intensive Care Units, Pediatric
-	N02.278.388.493.390.380 Intensive Care Units, Neonatal
-	N02.278.388.493.592 Recovery Room
-	N02.278.388.493.696 Respiratory Care Units
-	N02.278.388.540 Nurseries, Hospital
-	N02.278.388.620 Nursing Stations
-	N02.278.388.700 Operating Rooms
-	N02.278.388.740 Self-Care Units
-	N02.278.421 Hospitals
-	N02.278.421.280 Hospitals, Animal
-	N02.278.421.306 Hospitals, Community
-	N02.278.421.389 Hospitals, General
-	N02.278.421.395 Hospitals, Group Practice
-	N02.278.421.414 Hospitals, High-Volume
-	N02.278.421.434 Hospitals, Low-Volume
-	N02.278.421.481 Hospitals, Private
-	N02.278.421.481.500 Hospitals, Proprietary
-	N02.278.421.481.600 Hospitals, Religious
-	N02.278.421.481.800 Hospitals, Voluntary
-	N02.278.421.510 Hospitals, Public
-	N02.278.421.510.100 Hospitals, County
-	N02.278.421.510.140 Hospitals, District
-	N02.278.421.510.180 Hospitals, Federal
-	N02.278.421.510.180.250 Hospitals, Military
-	N02.278.421.510.180.400 Hospitals, Veterans
-	N02.278.421.510.210 Hospitals, Municipal
-	N02.278.421.510.300 Hospitals, State
-	N02.278.421.518 Hospitals, Rural
-	N02.278.421.530 Hospitals, Satellite
-	N02.278.421.556 Hospitals, Special

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	N02.278.421.556.070	Cancer Care Facilities
-	N02.278.421.556.110	Cardiac Care Facilities
-	N02.278.421.556.185	Hospices
-	N02.278.421.556.200	Hospitals, Chronic Disease
-	N02.278.421.556.294	Hospitals, Convalescent
-	N02.278.421.556.330	Hospitals, Isolation
-	N02.278.421.556.366	Hospitals, Maternity
-	N02.278.421.556.386	Hospitals, Osteopathic
-	N02.278.421.556.437	Hospitals, Pediatric
-	N02.278.421.556.508	Hospitals, Psychiatric
-	N02.278.421.556.868	Surgicenters
-	N02.278.421.639	Hospitals, Teaching
-	N02.278.421.639.725	Hospitals, University
-	N02.278.421.660	Hospitals, Urban
-	N02.278.421.660.400	Hospitals, Municipal
-	N02.278.421.702	Mobile Health Units
-	N02.278.421.745	Secondary Care Centers
-	N02.278.421.830	Tertiary Care Centers
-	N02.278.487	Laboratories
-	N02.278.487.410	Laboratories, Hospital
-	N02.278.524	Leper Colonies
-	N02.278.562	Medical Office Buildings
New Heading	<b>N02.278.590</b>	<b>Morgue</b>
-	N02.278.617	Nurseries
-	N02.278.617.473	Nurseries, Hospital
-	N02.278.678	Pharmacies
-	N02.278.692	Physicians' Offices
-	N02.278.726	Poison Control Centers
-	N02.278.808	Rehabilitation Centers
-	N02.278.808.860	Sheltered Workshops
-	N02.278.808.930	Substance Abuse Treatment Centers
-	N02.278.825	Residential Facilities
-	N02.278.825.187	Assisted Living Facilities
-	N02.278.825.375	Group Homes
Old Tree	<b>N02.278.825.408</b>	<b>Halfway Houses</b>



## MeSH Tree Changes for 2017

Type	Tree - heading	
-	N02.278.825.462	Homes for the Aged
-	N02.278.825.610	Nursing Homes
-	N02.278.825.610.480	Intermediate Care Facilities
-	N02.278.825.610.770	Skilled Nursing Facilities
Old Tree	N02.278.825.700	Orphanages
New Heading	N02.278.912	Student Run Clinic
-	N02.350	Health Manpower
-	N02.360	Health Personnel
-	N02.360.067	Allied Health Personnel
-	N02.360.067.040	Animal Technicians
-	N02.360.067.080	Community Health Workers
-	N02.360.067.105	Dental Auxiliaries
-	N02.360.067.105.250	Dental Assistants
-	N02.360.067.105.376	Dental Hygienists
-	N02.360.067.105.503	Dental Technicians
-	N02.360.067.105.751	Denturists
-	N02.360.067.150	Emergency Medical Technicians
-	N02.360.067.350	Home Health Aides
-	N02.360.067.450	Licensed Practical Nurses
-	N02.360.067.550	Medical Record Administrators
-	N02.360.067.600	Medical Secretaries
-	N02.360.067.600.630	Medical Receptionists
-	N02.360.067.652	Nurses' Aides
-	N02.360.067.652.628	Psychiatric Aides
Old Tree	N02.360.067.676	Nutritionists
-	N02.360.067.700	Operating Room Technicians
-	N02.360.067.730	Pharmacists' Aides
-	N02.360.067.730	Pharmacy Technicians
-	N02.360.067.732	Physical Therapist Assistants
Old Tree	N02.360.067.735	Physical Therapists
-	N02.360.067.740	Physician Assistants
-	N02.360.067.740.690	Ophthalmic Assistants
-	N02.360.067.740.750	Pediatric Assistants
-	N02.360.133	Anatomists
New	N02.360.140	Anesthetists

## MeSH Tree Changes for 2017

Type	Tree - heading	
Heading		
New Heading	<b>N02.360.140.040</b>	<b>Anesthesiologists</b>
New Tree	<a href="#">N02.360.140.650</a>	<a href="#">Nurse Anesthetists</a>
New Heading	<b>N02.360.170</b>	<b>Audiologists</b>
-	N02.360.200	Caregivers
New Heading	<b>N02.360.215</b>	<b>Case Managers</b>
-	N02.360.230	Coroners and Medical Examiners
-	N02.360.290	Dental Staff
-	N02.360.290.490	Dental Staff, Hospital
-	N02.360.330	Dentists
-	N02.360.330.310	Dentists, Women
New Heading	<b>N02.360.330.397</b>	<b>Endodontists</b>
-	N02.360.330.483	Oral and Maxillofacial Surgeons
-	N02.360.330.655	Orthodontists
New Tree	<a href="#">N02.360.338</a>	<a href="#">Doulas</a>
New Heading	<b>N02.360.345</b>	<b>Epidemiologists</b>
-	N02.360.360	Faculty, Dental
-	N02.360.375	Faculty, Medical
-	N02.360.390	Faculty, Nursing
-	N02.360.430	Health Facility Administrators
-	N02.360.430.490	Hospital Administrators
-	N02.360.430.490.100	Chief Executive Officers, Hospital
-	N02.360.490	Infection Control Practitioners
New Tree	<a href="#">N02.360.500</a>	<a href="#">Medical Chaperones</a>
-	N02.360.510	Medical Laboratory Personnel
-	N02.360.630	Medical Staff
-	N02.360.630.490	Medical Staff, Hospital
-	N02.360.630.490.400	Hospitalists
-	N02.360.650	Nurses
-	N02.360.650.580	Nurse Administrators
Old Tree	<b>N02.360.650.600</b>	<b>Nurse Anesthetists</b>

## MeSH Tree Changes for 2017

Type	Tree - heading	
Old Tree	<b>N02.360.650.610</b>	<b>Nurse Clinicians</b>
Old Tree	<b>N02.360.650.620</b>	<b>Nurse Midwives</b>
-	N02.360.650.640	Nurse Practitioners
-	N02.360.650.640.249	Family Nurse Practitioners
-	N02.360.650.640.500	Pediatric Nurse Practitioners
New Heading	<b>N02.360.650.648</b>	<b>Nurse Specialists</b>
New Tree	<b>N02.360.650.648.500</b>	<b>Nurse Anesthetists</b>
New Tree	<b>N02.360.650.648.525</b>	<b>Nurse Clinicians</b>
New Tree	<b>N02.360.650.648.762</b>	<b>Nurse Midwives</b>
New Heading	<b>N02.360.650.648.940</b>	<b>Nurses, Pediatric</b>
New Heading	<b>N02.360.650.648.940.500</b>	<b>Nurses, Neonatal</b>
-	N02.360.650.655	Nurses, Community Health
-	N02.360.650.662	Nurses, International
-	N02.360.650.670	Nurses, Male
-	N02.360.650.835	Nurses, Public Health
-	N02.360.680	Nursing Staff
-	N02.360.680.490	Nursing Staff, Hospital
New Tree	<b>N02.360.695</b>	<b>Nutritionists</b>
New Heading	<b>N02.360.710</b>	<b>Occupational Therapists</b>
New Heading	<b>N02.360.725</b>	<b>Optometrists</b>
-	N02.360.740	Personnel, Hospital
-	N02.360.740.322	Dental Staff, Hospital
-	N02.360.740.380	Hospital Administrators
-	N02.360.740.380.100	Chief Executive Officers, Hospital
-	N02.360.740.400	Hospital Volunteers
Old Tree	<b>N02.360.740.400.650</b>	<b>Patient Escort Service</b>
-	N02.360.740.422	Medical Staff, Hospital
-	N02.360.740.422.400	Hospitalists
-	N02.360.740.523	Nursing Staff, Hospital
-	N02.360.780	Pharmacists

## MeSH Tree Changes for 2017

Type	Tree - heading	
New Tree	<a href="#">N02.360.790</a>	<a href="#">Physical Therapists</a>
-	N02.360.800	Physician Executives
-	N02.360.810	Physicians
New Heading	<b>N02.360.810.020</b>	<b>Allergists</b>
New Heading	<b>N02.360.810.040</b>	<b>Anesthesiologists</b>
New Heading	<b>N02.360.810.128</b>	<b>Cardiologists</b>
New Heading	<b>N02.360.810.215</b>	<b>Dermatologists</b>
New Heading	<b>N02.360.810.303</b>	<b>Endocrinologists</b>
-	N02.360.810.390	Foreign Medical Graduates
New Heading	<b>N02.360.810.438</b>	<b>Gastroenterologists</b>
-	N02.360.810.485	General Practitioners
New Heading	<b>N02.360.810.533</b>	<b>Geriatricians</b>
-	N02.360.810.580	Hospitalists
New Heading	<b>N02.360.810.628</b>	<b>Nephrologists</b>
New Heading	<b>N02.360.810.652</b>	<b>Neurologists</b>
-	N02.360.810.675	Occupational Health Physicians
New Heading	<b>N02.360.810.699</b>	<b>Oncologists</b>
New Heading	<b>N02.360.810.699.500</b>	<b>Radiation Oncologists</b>
New Heading	<b>N02.360.810.705</b>	<b>Ophthalmologists</b>
-	N02.360.810.722	Osteopathic Physicians
New Heading	<b>N02.360.810.734</b>	<b>Otolaryngologists</b>
New Heading	<b>N02.360.810.746</b>	<b>Pathologists</b>
New Heading	<b>N02.360.810.758</b>	<b>Pediatricians</b>
New Heading	<b>N02.360.810.758.500</b>	<b>Neonatologists</b>
New	<b>N02.360.810.764</b>	<b>Physiatrists</b>

## MeSH Tree Changes for 2017

Type	Tree - heading	
Heading		
-	N02.360.810.770	Physicians, Family
-	N02.360.810.795	Physicians, Primary Care
-	N02.360.810.820	Physicians, Women
New Heading	<b>N02.360.810.865</b>	<b>Pulmonologists</b>
New Heading	<b>N02.360.810.877</b>	<b>Radiologists</b>
New Heading	<b>N02.360.810.877.500</b>	<b>Radiation Oncologists</b>
New Heading	<b>N02.360.810.888</b>	<b>Rheumatologists</b>
-	N02.360.810.910	Surgeons
-	N02.360.810.910.500	Barber Surgeons
-	N02.360.810.910.750	Neurosurgeons
New Heading	<b>N02.360.810.910.875</b>	<b>Orthopedic Surgeons</b>
New Heading	<b>N02.360.810.955</b>	<b>Urologists</b>
-	N02.360.905	Veterinarians
-	N02.370	Health Promotion
-	N02.370.300	Healthy People Programs
-	N02.370.650	Weight Reduction Programs
-	N02.421	Health Services
-	N02.421.044	Adolescent Health Services
-	N02.421.088	Child Care
-	N02.421.088.120	Infant Care
-	N02.421.088.120.059	Facilitated Tucking
-	N02.421.088.120.140	Kangaroo-Mother Care Method
-	N02.421.088.120.180	Perinatal Care
-	N02.421.088.120.240	Rooming-in Care
-	N02.421.143	Community Health Services
-	N02.421.143.065	Adult Day Care Centers
-	N02.421.143.098	Child Day Care Centers
-	N02.421.143.130	Child Health Services
-	N02.421.143.130.320	Early Intervention (Education)
-	N02.421.143.130.660	Maternal-Child Health Services
-	N02.421.143.150	Community Health Nursing

## MeSH Tree Changes for 2017

Type	Tree - heading
-	N02.421.143.150.500 Home Health Nursing
-	N02.421.143.150.750 Parish Nursing
-	N02.421.143.183 Community Mental Health Services
-	N02.421.143.202 Community Networks
-	N02.421.143.221 Community Pharmacy Services
-	N02.421.143.262 Community Participation
-	N02.421.143.262 Consumer Participation
-	N02.421.143.262.300 Patient Participation
-	N02.421.143.303 Counseling
-	N02.421.143.303.175 Distance Counseling
-	N02.421.143.303.700 Sex Counseling
-	N02.421.143.401 Family Planning Services
-	N02.421.143.452 Foster Home Care
-	N02.421.143.524 Home Care Services
-	N02.421.143.524.337 Hemodialysis, Home
-	N02.421.143.524.403 Home Care Services, Hospital-Based
-	N02.421.143.524.419 Home Health Nursing
-	N02.421.143.524.436 Home Infusion Therapy
-	N02.421.143.524.470 Home Nursing
-	N02.421.143.524.470.610 Respite Care
-	N02.421.143.524.595 Homemaker Services
-	N02.421.143.524.650 Parenteral Nutrition, Home
-	N02.421.143.524.650.565 Parenteral Nutrition, Home Total
-	N02.421.143.550 Hospices
-	N02.421.143.620 Maternal Health Services
-	N02.421.143.620.275 Maternal-Child Health Services
-	N02.421.143.620.550 Perinatal Care
-	N02.421.143.620.550.500 Postnatal Care
-	N02.421.143.620.620 Preconception Care
-	N02.421.143.620.704 Prenatal Care
-	N02.421.143.740 Occupational Health Services
-	N02.421.143.913 Senior Centers
Old Tree	<b>N02.421.192</b> Continuity of Patient Care
Old Tree	<b>N02.421.192.125</b> Patient Discharge
Old Tree	<b>N02.421.192.249</b> Patient Handoff
Old Tree	<b>N02.421.192.624</b> Patient Transfer

## MeSH Tree Changes for 2017

Type	Tree - heading	
Old Tree	N02.421.192.718	Transition to Adult Care
Old Tree	N02.421.192.812	Transitional Care
-	N02.421.240	Dental Health Services
-	N02.421.240.190	Dental Care
-	N02.421.240.190.210	Dental Care for Aged
-	N02.421.240.190.215	Dental Care for Children
-	N02.421.240.190.220	Dental Care for Chronically Ill
-	N02.421.240.190.230	Dental Care for Disabled
-	N02.421.240.200	Dental Service, Hospital
-	N02.421.242	Dietary Services
-	N02.421.242.472	Food Service, Hospital
-	N02.421.242.472.310	Menu Planning
-	N02.421.297	Emergency Medical Services
-	N02.421.297.028	Advanced Trauma Life Support Care
New Heading	N02.421.297.043	Emergency Medical Dispatch
-	N02.421.297.058	Emergency Medical Service Communication Systems
New Heading	N02.421.297.127	Emergency Police Dispatcher
-	N02.421.297.195	Emergency Service, Hospital
-	N02.421.297.195.480	Trauma Centers
-	N02.421.297.200	Emergency Services, Psychiatric
-	N02.421.297.695	Poison Control Centers
-	N02.421.297.879	Transportation of Patients
-	N02.421.297.879.049	Ambulance Diversion
-	N02.421.297.879.100	Ambulances
-	N02.421.297.879.100.100	Air Ambulances
-	N02.421.297.879.550	Stretchers
-	N02.421.297.900	Triage
-	N02.421.308	Genetic Services
-	N02.421.308.400	Genetic Counseling
-	N02.421.308.430	Genetic Testing
New Heading	N02.421.308.430.500	Pharmacogenomic Testing
-	N02.421.314	Health Services for Persons with Disabilities
-	N02.421.320	Health Services for the Aged
-	N02.421.325	Health Services for Transgender Persons

## MeSH Tree Changes for 2017

Type	Tree - heading
-	N02.421.330 Health Services, Indigenous
-	N02.421.380 Health Services Misuse
-	N02.421.380.450 Medical Overuse
-	N02.421.380.450.500 Unnecessary Procedures
-	N02.421.450 Medical Errors
-	N02.421.450.280 Diagnostic Errors
-	N02.421.450.500 Medication Errors
-	N02.421.450.500.249 Inappropriate Prescribing
-	N02.421.450.500.500 Medication Reconciliation
-	N02.421.450.500.750 Near Miss, Healthcare
-	N02.421.450.550 Near Miss, Healthcare
-	N02.421.450.600 Observer Variation
-	N02.421.450.800 Radiotherapy Setup Errors
-	N02.421.461 Mental Health Services
-	N02.421.461.232 Community Mental Health Services
-	N02.421.461.363 Counseling
-	N02.421.461.363.349 Directive Counseling
-	N02.421.461.363.349.500 Motivational Interviewing
-	N02.421.461.363.437 Distance Counseling
-	N02.421.461.363.700 Sex Counseling
-	N02.421.461.798 Social Work, Psychiatric
-	N02.421.533 Nursing Care
-	N02.421.533.099 Cardiovascular Nursing
-	N02.421.533.149 Critical Care Nursing
-	N02.421.533.200 Emergency Nursing
-	N02.421.533.245 Geriatric Nursing
-	N02.421.533.282 Holistic Nursing
-	N02.421.533.320 Home Nursing
-	N02.421.533.320.610 Respite Care
-	N02.421.533.390 Hospice and Palliative Care Nursing
-	N02.421.533.460 Maternal-Child Nursing
-	N02.421.533.460.600 Neonatal Nursing
-	N02.421.533.475 Medical-Surgical Nursing
-	N02.421.533.490 Nephrology Nursing
-	N02.421.533.505 Neuroscience Nursing
-	N02.421.533.520 Nursing, Practical



## MeSH Tree Changes for 2017

Type	Tree - heading	
-	N02.421.533.571	Obstetric Nursing
-	N02.421.533.585	Occupational Health Nursing
-	N02.421.533.600	Oncology Nursing
-	N02.421.533.635	Orthopedic Nursing
-	N02.421.533.691	Pediatric Nursing
-	N02.421.533.691.600	Neonatal Nursing
-	N02.421.533.710	Perioperative Nursing
-	N02.421.533.710.625	Operating Room Nursing
-	N02.421.533.710.650	Postanesthesia Nursing
-	N02.421.533.760	Primary Nursing
-	N02.421.533.778	Psychiatric Nursing
-	N02.421.533.889	Rehabilitation Nursing
-	N02.421.539	Nursing Services
-	N02.421.539.089	Home Care Services
-	N02.421.539.089.500	Home Health Nursing
-	N02.421.539.180	Nursing Service, Hospital
-	N02.421.585	Patient Care
Old Tree	N02.421.585.058	Aftercare
-	N02.421.585.106	Ambulatory Care
-	N02.421.585.106.500	Peritoneal Dialysis, Continuous Ambulatory
-	N02.421.585.148	Bloodless Medical and Surgical Procedures
New Tree	N02.421.585.169	Continuity of Patient Care
New Tree	N02.421.585.169.063	Aftercare
New Tree	N02.421.585.169.125	Patient Discharge
New Tree	N02.421.585.169.249	Patient Handoff
New Tree	N02.421.585.169.624	Patient Transfer
New Tree	N02.421.585.169.718	Transition to Adult Care
New Tree	N02.421.585.169.812	Transitional Care
-	N02.421.585.190	Critical Care
-	N02.421.585.190.500	Intensive Care, Neonatal
-	N02.421.585.215	Custodial Care

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	N02.421.585.246	Day Care, Medical
-	N02.421.585.300	Episode of Care
-	N02.421.585.352	Foster Home Care
-	N02.421.585.400	Hospitalization
-	N02.421.585.400.480	Length of Stay
-	N02.421.585.400.600	Patient Admission
-	N02.421.585.400.610	Patient Discharge
-	N02.421.585.400.615	Patient Handoff
-	N02.421.585.400.620	Patient Readmission
-	N02.421.585.400.630	Patient Transfer
-	N02.421.585.415	Institutionalization
-	N02.421.585.415.280	Deinstitutionalization
-	N02.421.585.440	Life Support Care
-	N02.421.585.440.028	Advanced Cardiac Life Support
-	N02.421.585.440.040	Advanced Trauma Life Support Care
-	N02.421.585.476	Long-Term Care
-	N02.421.585.525	Moving and Lifting Patients
-	N02.421.585.564	Night Care
-	N02.421.585.666	Palliative Care
New Heading	<b>N02.421.585.683</b>	<b>Patient Comfort</b>
-	N02.421.585.700	Patient Positioning
-	N02.421.585.700.500	Kangaroo-Mother Care Method
-	N02.421.585.703	Perinatal Care
-	N02.421.585.703.500	Postnatal Care
-	N02.421.585.722	Perioperative Care
-	N02.421.585.722.400	Intraoperative Care
-	N02.421.585.722.500	Perioperative Nursing
-	N02.421.585.722.700	Postoperative Care
-	N02.421.585.753	Perioperative Period
-	N02.421.585.753.374	Intraoperative Period
-	N02.421.585.753.374.500	Operative Time
-	N02.421.585.753.750	Postoperative Period
-	N02.421.585.753.750.055	Anesthesia Recovery Period
-	N02.421.585.753.937	Preoperative Period
-	N02.421.585.775	Preconception Care

## MeSH Tree Changes for 2017

Type	Tree - heading
-	N02.421.585.786 Prenatal Care
-	N02.421.585.795 Preoperative Care
-	N02.421.585.850 Subacute Care
-	N02.421.585.905 Terminal Care
-	N02.421.585.905.199 Euthanasia
-	N02.421.585.905.199.500 Euthanasia, Active
-	N02.421.585.905.199.500.200 Euthanasia, Active, Voluntary
-	N02.421.585.905.199.500.750 Euthanasia, Animal
-	N02.421.585.905.199.625 Euthanasia, Passive
-	N02.421.585.905.400 Hospice Care
-	N02.421.585.905.700 Resuscitation Orders
-	N02.421.585.905.850 Suicide, Assisted
-	N02.421.585.928 Time-to-Treatment
-	N02.421.585.940 Transitional Care
-	N02.421.585.952 Withholding Treatment
-	N02.421.585.952.500 Euthanasia, Passive
New Tree	<a href="#">N02.421.608</a> <a href="#">Patient Escort Service</a>
-	N02.421.631 Personal Health Services
-	N02.421.668 Pharmaceutical Services
-	N02.421.668.274 Community Pharmacy Services
-	N02.421.668.320 Drug Information Services
-	N02.421.668.320.120 Adverse Drug Reaction Reporting Systems
-	N02.421.668.320.200 Clinical Pharmacy Information Systems
-	N02.421.668.438 Medication Therapy Management
-	N02.421.668.475 Pharmaceutical Services, Online
-	N02.421.668.556 Pharmacy Service, Hospital
-	N02.421.668.778 Prescriptions
-	N02.421.668.778.500 Drug Prescriptions
-	N02.421.668.778.500.312 Drug Substitution
-	N02.421.668.778.750 Electronic Prescribing
-	N02.421.726 Preventive Health Services
-	N02.421.726.233 Diagnostic Services
-	N02.421.726.233.110 Clinical Laboratory Services
-	N02.421.726.233.166 Direct-To-Consumer Screening and Testing
-	N02.421.726.233.221 Genetic Testing

## MeSH Tree Changes for 2017

Type	Tree - heading
New Heading	<b>N02.421.726.233.221.500</b> <b>Pharmacogenomic Testing</b>
-	N02.421.726.233.443 Mass Screening
-	N02.421.726.233.443.221 Anonymous Testing
-	N02.421.726.233.443.443 Mass Chest X-Ray
-	N02.421.726.233.443.633 Multiphasic Screening
-	N02.421.726.233.443.816 Neonatal Screening
-	N02.421.726.233.628 Mobile Health Units
-	N02.421.726.320 Early Intervention (Education)
-	N02.421.726.363 Early Medical Intervention
-	N02.421.726.407 Health Education
-	N02.421.726.407.229 Consumer Health Information
-	N02.421.726.407.229.500 Health Literacy
-	N02.421.726.407.229.500.500 Patient Medication Knowledge
-	N02.421.726.407.457 Health Education, Dental
-	N02.421.726.407.478 Health Fairs
-	N02.421.726.407.680 Patient Education as Topic
-	N02.421.726.407.680.500 Prenatal Education
-	N02.421.726.507 Health Promotion
-	N02.421.726.507.300 Healthy People Programs
-	N02.421.726.507.650 Weight Reduction Programs
-	N02.421.726.608 Immunization Programs
-	N02.421.726.608.500 Mass Vaccination
-	N02.421.726.708 Needle-Exchange Programs
-	N02.421.726.755 Post-Exposure Prophylaxis
-	N02.421.726.758 Primary Prevention
-	N02.421.726.758.310 Immunization
-	N02.421.726.758.310.890 Vaccination
-	N02.421.726.758.310.890.500 Mass Vaccination
-	N02.421.726.758.655 Pre-Exposure Prophylaxis
-	N02.421.726.809 School Health Services
-	N02.421.726.809.576 School Dentistry
-	N02.421.726.809.742 School Nursing
-	N02.421.726.825 Secondary Prevention
-	N02.421.726.860 Tertiary Prevention
-	N02.421.784 Rehabilitation

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	N02.421.784.110	Activities of Daily Living
New Heading	<b>N02.421.784.244</b>	<b>Cardiac Rehabilitation</b>
-	N02.421.784.377	Correction of Hearing Impairment
-	N02.421.784.511	Neurological Rehabilitation
New Heading	<b>N02.421.784.511.500</b>	<b>Stroke Rehabilitation</b>
-	N02.421.784.578	Psychiatric Rehabilitation
-	N02.421.784.644	Rehabilitation, Vocational
-	N02.421.784.644.350	Employment, Supported
-	N02.421.784.680	Self Care
-	N02.421.784.840	Telerehabilitation
-	N02.421.800	Reproductive Health Services
-	N02.421.800.249	Family Planning Services
-	N02.421.800.500	Maternal Health Services
-	N02.421.816	Rural Health Services
-	N02.421.816.500	Rural Nursing
-	N02.421.849	Social Work
-	N02.421.849.673	Social Work, Psychiatric
-	N02.421.897	Student Health Services
-	N02.421.908	Suburban Health Services
-	N02.421.911	Tissue and Organ Procurement
-	N02.421.911.200	Directed Tissue Donation
-	N02.421.911.600	Donor Selection
-	N02.421.914	Urban Health Services
-	N02.421.920	Women's Health Services
-	N02.421.920.660	Preconception Care
-	N02.508	Housekeeping
-	N02.508.472	Housekeeping, Hospital
-	N02.628	Maintenance
-	N02.628.472	Maintenance and Engineering, Hospital
Old Tree	<b>N02.720</b>	<b>Religious Missions</b>
Old Tree	<b>N02.860</b>	<b>Senior Centers</b>
-	N03	Health Care Economics and Organizations
-	N03.109	Dual MEDICAID MEDICARE Eligibility
-	N03.219	Economics

## MeSH Tree Changes for 2017

Type	Tree - heading
-	N03.219.075 Compensation and Redress
-	N03.219.151 Costs and Cost Analysis
-	N03.219.151.080 Cost Allocation
-	N03.219.151.125 Cost-Benefit Analysis
-	N03.219.151.160 Cost Control
-	N03.219.151.160.200 Cost Savings
-	N03.219.151.165 Cost of Illness
-	N03.219.151.170 Cost Sharing
-	N03.219.151.170.300 Deductibles and Coinsurance
-	N03.219.151.170.650 Medical Savings Accounts
-	N03.219.151.400 Health Care Costs
-	N03.219.151.400.325 Direct Service Costs
-	N03.219.151.400.350 Drug Costs
-	N03.219.151.400.375 Employer Health Costs
-	N03.219.151.400.687 Hospital Costs
-	N03.219.151.450 Health Expenditures
-	N03.219.151.450.200 Capital Expenditures
-	N03.219.188 Economic Competition
-	N03.219.206 Economic Development
-	N03.219.215 Economics, Behavioral
-	N03.219.224 Economics, Dental
-	N03.219.224.426 Fees, Dental
-	N03.219.262 Economics, Hospital
-	N03.219.262.300 Hospital Charges
-	N03.219.262.500 Hospital Costs
-	N03.219.300 Economics, Medical
-	N03.219.300.426 Fees, Medical
-	N03.219.338 Economics, Nursing
-	N03.219.390 Economics, Pharmaceutical
-	N03.219.416 External Debt
-	N03.219.442 Fees and Charges
-	N03.219.442.090 Capitation Fee
-	N03.219.442.195 Fee-for-Service Plans
-	N03.219.442.301 Fees, Dental
-	N03.219.442.426 Fees, Medical
-	N03.219.442.551 Fees, Pharmaceutical

## MeSH Tree Changes for 2017

Type	Tree - heading
-	N03.219.442.551.621 Prescription Fees
-	N03.219.442.613 Hospital Charges
-	N03.219.442.800 Rate Setting and Review
-	N03.219.463 Financial Management
-	N03.219.463.030 Accounting
-	N03.219.463.030.080 Accounts Payable and Receivable
-	N03.219.463.030.100 Banking, Personal
-	N03.219.463.030.120 Depreciation
-	N03.219.463.030.140 Financial Statements
-	N03.219.463.030.160 Financial Audit
-	N03.219.463.030.320 Patient Credit and Collection
-	N03.219.463.045 Bankruptcy
-	N03.219.463.060 Budgets
-	N03.219.463.060.800 Rate Setting and Review
-	N03.219.463.085 Capital Financing
-	N03.219.463.100 Contract Services
-	N03.219.463.100.110 Competitive Bidding
-	N03.219.463.100.600 Outsourced Services
-	N03.219.463.280 Financial Management, Hospital
-	N03.219.463.300 Financing, Construction
-	N03.219.463.360 Fund Raising
-	N03.219.463.430 Group Purchasing
-	N03.219.463.500 Leasing, Property
-	N03.219.463.548 Marketing of Health Services
-	N03.219.463.800 Risk Management
-	N03.219.463.800.800 Risk Sharing, Financial
-	N03.219.483 Financial Support
-	N03.219.483.311 Foundations
-	N03.219.483.408 Health Planning Support
-	N03.219.483.500 Healthcare Financing
-	N03.219.483.645 Research Support as Topic
-	N03.219.483.838 Training Support
-	N03.219.483.838.276 Fellowships and Scholarships
-	N03.219.521 Financing, Organized
-	N03.219.521.346 Financing, Government
-	N03.219.521.346.506 Public Assistance

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	N03.219.521.346.506.281	Food Assistance
-	N03.219.521.346.506.564	Medical Assistance
-	N03.219.521.346.506.564.655	Medicaid
-	N03.219.521.346.506.564.663	Medicare
New Heading	<b>N03.219.521.346.506.564.663.250</b> <b>Reauthorization Act of 2015</b>	<b>Medicare Access and CHIP</b>
-	N03.219.521.346.506.564.663.500	Medicare Part A
-	N03.219.521.346.506.564.663.515	Medicare Part B
-	N03.219.521.346.506.564.663.757	Medicare Part C
-	N03.219.521.346.506.678	Old Age Assistance
-	N03.219.521.346.506.849	Social Security
-	N03.219.521.346.506.849.095	Aid to Families with Dependent Children
-	N03.219.521.346.733	Veterans Disability Claims
-	N03.219.521.346.866	Workers' Compensation
-	N03.219.521.576	Insurance
-	N03.219.521.576.090	Cost Sharing
-	N03.219.521.576.090.300	Deductibles and Coinsurance
-	N03.219.521.576.090.650	Medical Savings Accounts
-	N03.219.521.576.130	Insurance Benefits
-	N03.219.521.576.160	Insurance Carriers
-	N03.219.521.576.210	Insurance Claim Reporting
-	N03.219.521.576.215	Insurance Claim Review
-	N03.219.521.576.265	Insurance Coverage
-	N03.219.521.576.265.500	Universal Coverage
-	N03.219.521.576.300	Insurance, Disability
-	N03.219.521.576.300.900	Workers' Compensation
-	N03.219.521.576.343	Insurance, Health
-	N03.219.521.576.343.071	Children's Health Insurance Program
-	N03.219.521.576.343.144	For-Profit Insurance Plans
-	N03.219.521.576.343.290	Health Benefit Plans, Employee
-	N03.219.521.576.343.290.300	Employee Retirement Income Security Act
-	N03.219.521.576.343.349	Health Insurance Portability and Accountability Act
-	N03.219.521.576.343.409	Insurance, Accident
-	N03.219.521.576.343.450	Insurance, Dental
-	N03.219.521.576.343.480	Insurance, Health, Reimbursement
-	N03.219.521.576.343.492	Insurance, Hospitalization



## MeSH Tree Changes for 2017

Type	Tree - heading	
-	N03.219.521.576.343.508	Insurance, Long-Term Care
-	N03.219.521.576.343.533	Insurance, Major Medical
-	N03.219.521.576.343.539	Insurance, Medigap
-	N03.219.521.576.343.545	Insurance, Nursing Services
-	N03.219.521.576.343.575	Insurance, Pharmaceutical Services
-	N03.219.521.576.343.575.500	Medicare Part D
-	N03.219.521.576.343.575.500.500	Medication Therapy Management
-	N03.219.521.576.343.616	Insurance, Physician Services
-	N03.219.521.576.343.616.508	Medicare Assignment
-	N03.219.521.576.343.658	Insurance, Psychiatric
-	N03.219.521.576.343.700	Insurance, Surgical
-	N03.219.521.576.343.800	Managed Care Programs
-	N03.219.521.576.343.800.175	Competitive Medical Plans
-	N03.219.521.576.343.800.400	Health Maintenance Organizations
-	N03.219.521.576.343.800.450	Independent Practice Associations
-	N03.219.521.576.343.800.700	Patient Freedom of Choice Laws
-	N03.219.521.576.343.800.750	Preferred Provider Organizations
-	N03.219.521.576.343.800.875	Provider-Sponsored Organizations
-	N03.219.521.576.343.820	Managed Competition
-	N03.219.521.576.343.830	Medical Savings Accounts
-	N03.219.521.576.343.840	Medicare
New Heading	<b>N03.219.521.576.343.840.254 Act of 2015</b>	<b>Medicare Access and CHIP Reauthorization</b>
-	N03.219.521.576.343.840.508	Medicare Assignment
-	N03.219.521.576.343.840.725	Medicare Part A
-	N03.219.521.576.343.840.754	Medicare Part B
-	N03.219.521.576.343.840.877	Medicare Part C
-	N03.219.521.576.343.840.938	Medicare Part D
-	N03.219.521.576.343.840.938.500	Medication Therapy Management
-	N03.219.521.576.343.900	National Health Insurance, United States
-	N03.219.521.576.343.912	Not-For-Profit Insurance Plans
-	N03.219.521.576.343.918	Patient Protection and Affordable Care Act
-	N03.219.521.576.343.918.500	Health Insurance Exchanges
-	N03.219.521.576.343.925	Prepaid Health Plans
-	N03.219.521.576.343.925.400	Health Maintenance Organizations
-	N03.219.521.576.343.962	Single-Payer System

## MeSH Tree Changes for 2017

Type	Tree - heading
New Heading	<b>N03.219.521.576.343.972</b> <b>Value-Based Insurance</b>
-	N03.219.521.576.343.981 Value-Based Purchasing
-	N03.219.521.576.443 Insurance, Liability
-	N03.219.521.576.543 Insurance, Life
-	N03.219.521.576.570 Insurance Pools
-	N03.219.521.576.585 Insurance Selection Bias
-	N03.219.521.576.585.500 Preexisting Condition Coverage
-	N03.219.521.576.823 Social Security
-	N03.219.521.710 Insurance, Health, Reimbursement
-	N03.219.521.710.305 Reimbursement Mechanisms
-	N03.219.521.710.305.090 Fee-for-Service Plans
-	N03.219.521.710.305.090.125 Blue Cross Blue Shield Insurance Plans
-	N03.219.521.710.305.180 Physician Payment Review Commission
-	N03.219.521.710.305.200 Prospective Payment System
-	N03.219.521.710.305.200.080 Diagnosis-Related Groups
-	N03.219.521.710.305.200.080.550 Outliers, DRG
-	N03.219.521.710.305.200.500 Medicare Payment Advisory Commission
-	N03.219.521.710.305.200.605 Prospective Payment Assessment Commission
-	N03.219.521.710.305.290 Reimbursement, Disproportionate Share
-	N03.219.521.710.305.380 Reimbursement, Incentive
-	N03.219.521.710.305.500 Relative Value Scales
-	N03.219.521.855 Single-Payer System
-	N03.219.559 Financing, Personal
-	N03.219.559.500 Medical Savings Accounts
-	N03.219.650 Health Care Sector
-	N03.219.680 Inflation, Economic
-	N03.219.702 Investments
-	N03.219.780 Medical Indigency
-	N03.219.900 Taxes
-	N03.219.900.250 Income Tax
-	N03.219.900.880 Tax Equity and Fiscal Responsibility Act
-	N03.219.900.940 Tax Exemption
-	N03.349 Health Planning
-	N03.349.270 Health Care Rationing
-	N03.349.285 Health Care Reform

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	N03.349.300	Health Plan Implementation
-	N03.349.315	Health Planning Guidelines
-	N03.349.325	Health Planning Technical Assistance
-	N03.349.330	Health Priorities
-	N03.349.340	Health Resources
-	N03.349.380	Health Services Research
-	N03.349.380.210	Health Care Surveys
-	N03.349.380.210.500	Health Impact Assessment
New Heading	<b>N03.349.380.210.750</b>	<b>Patient Reported Outcome Measures</b>
-	N03.349.380.420	Health Services Needs and Demand
-	N03.349.380.565	Needs Assessment
-	N03.349.380.710	Organizational Case Studies
-	N03.349.550	National Health Programs
-	N03.349.550.610	National Health Insurance, United States
-	N03.349.550.805	Single-Payer System
-	N03.349.550.902	State Medicine
-	N03.349.650	Regional Health Planning
-	N03.349.650.095	Catchment Area (Health)
-	N03.349.650.120	Certificate of Need
-	N03.349.650.185	Community Health Planning
-	N03.349.650.250	Health Facility Planning
-	N03.349.650.250.080	Bed Conversion
-	N03.349.650.250.200	Hospital Planning
-	N03.349.650.270	Health Systems Plans
-	N03.349.650.340	Medically Underserved Area
-	N03.349.650.400	Regional Medical Programs
-	N03.349.650.480	State Health Plans
-	N03.349.650.480.500	Children's Health Insurance Program
-	N03.540	Organizations
-	N03.540.052	Academies and Institutes
-	N03.540.052.500	National Academy of Sciences (U.S.)
-	N03.540.052.500.500	Institute of Medicine (U.S.)
-	N03.540.052.500.500	National Academies of Science, Engineering, and Medicine (U.S.) Health and Medicine Division
-	N03.540.052.750	National Institutes of Health (U.S.)

## MeSH Tree Changes for 2017

Type	Tree - heading
-	N03.540.052.750.150 National Cancer Institute (U.S.)
-	N03.540.052.750.200 National Eye Institute (U.S.)
-	N03.540.052.750.300 National Heart, Lung, and Blood Institute (U.S.)
-	N03.540.052.750.325 National Human Genome Research Institute (U.S.)
-	N03.540.052.750.400 National Institute of Allergy and Infectious Diseases (U.S.)
-	N03.540.052.750.425 National Institute of Arthritis and Musculoskeletal and Skin Diseases (U.S.)
-	N03.540.052.750.435 National Institute of Biomedical Imaging and Bioengineering (U.S.)
-	N03.540.052.750.440 National Institute of Child Health and Human Development (U.S.)
-	N03.540.052.750.445 National Institute of Dental and Craniofacial Research (U.S.)
-	N03.540.052.750.450 National Institute of Diabetes and Digestive and Kidney Diseases (U.S.)
-	N03.540.052.750.455 National Institute of Environmental Health Sciences (U.S.)
-	N03.540.052.750.457 National Institute of General Medical Sciences (U.S.)
-	N03.540.052.750.460 National Institute of Mental Health (U.S.)
-	N03.540.052.750.461 National Institute of Neurological Disorders and Stroke
-	N03.540.052.750.462 National Institute of Nursing Research (U.S.)
-	N03.540.052.750.465 National Institute on Aging (U.S.)
-	N03.540.052.750.467 National Institute on Alcohol Abuse and Alcoholism (U.S.)
-	N03.540.052.750.470 National Institute on Deafness and Other Communication Disorders (U.S.)
-	N03.540.052.750.485 National Institute on Drug Abuse (U.S.)
-	N03.540.052.750.490 National Library of Medicine (U.S.)
-	N03.540.089 African Union
-	N03.540.125 Charities
-	N03.540.162 College Fraternities and Sororities
-	N03.540.199 Congresses as Topic
-	N03.540.199.205 Consensus Development Conferences as Topic
-	N03.540.199.205.500 Consensus Development Conferences, NIH as Topic
-	N03.540.245 Consumer Organizations
-	N03.540.245.360 Community Participation
-	N03.540.245.360 Consumer Participation
-	N03.540.245.360.300 Patient Participation

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	N03.540.348	Government
-	N03.540.348.500	Federal Government
-	N03.540.348.500.500	United States Government Agencies
-	N03.540.348.500.500.099	Peace Corps
-	N03.540.348.500.500.199	United States Agency for International Development
-	N03.540.348.500.500.400	United States Department of Agriculture
-	N03.540.348.500.500.500	United States Department of Defense
-	N03.540.348.500.500.600	United States Dept. of Health and Human Services
-	N03.540.348.500.500.600.310 (U.S.)	Centers for Medicare and Medicaid Services
-	N03.540.348.500.500.600.480	National Institutes of Health (U.S.)
-	N03.540.348.500.500.600.480.150	National Cancer Institute (U.S.)
New Heading	<b>N03.540.348.500.500.600.480.175 Translational Sciences (U.S.)</b>	<b>National Center for Advancing</b>
New Heading	<b>N03.540.348.500.500.600.480.188 Integrative Health (U.S.)</b>	<b>National Center for Complementary and</b>
-	N03.540.348.500.500.600.480.200	National Eye Institute (U.S.)
-	N03.540.348.500.500.600.480.300 (U.S.)	National Heart, Lung, and Blood Institute
-	N03.540.348.500.500.600.480.325 Institute (U.S.)	National Human Genome Research
-	N03.540.348.500.500.600.480.400 Diseases (U.S.)	National Institute of Allergy and Infectious
-	N03.540.348.500.500.600.480.425 Musculoskeletal and Skin Diseases (U.S.)	National Institute of Arthritis and
-	N03.540.348.500.500.600.480.435 and Bioengineering (U.S.)	National Institute of Biomedical Imaging
-	N03.540.348.500.500.600.480.440 Human Development (U.S.)	National Institute of Child Health and
-	N03.540.348.500.500.600.480.445 Research (U.S.)	National Institute of Dental and Craniofacial
-	N03.540.348.500.500.600.480.450 and Kidney Diseases (U.S.)	National Institute of Diabetes and Digestive
-	N03.540.348.500.500.600.480.455 Sciences (U.S.)	National Institute of Environmental Health
-	N03.540.348.500.500.600.480.457 Sciences (U.S.)	National Institute of General Medical
-	N03.540.348.500.500.600.480.460	National Institute of Mental Health (U.S.)
-	N03.540.348.500.500.600.480.461 and Stroke	National Institute of Neurological Disorders
-	N03.540.348.500.500.600.480.462	National Institute of Nursing Research

## MeSH Tree Changes for 2017

Type	Tree - heading	
	(U.S.)	
-	N03.540.348.500.500.600.480.465	National Institute on Aging (U.S.)
-	N03.540.348.500.500.600.480.467 Alcoholism (U.S.)	National Institute on Alcohol Abuse and
-	N03.540.348.500.500.600.480.470 Communication Disorders (U.S.)	National Institute on Deafness and Other
-	N03.540.348.500.500.600.480.485	National Institute on Drug Abuse (U.S.)
-	N03.540.348.500.500.600.480.490	National Library of Medicine (U.S.)
-	N03.540.348.500.500.600.650	United States Public Health Service
-	N03.540.348.500.500.600.650.225 (U.S.)	Centers for Disease Control and Prevention
New Tree	<a href="#">N03.540.348.500.500.600.650.225.260 (U.S.)</a>	<a href="#">National Center for Health Statistics</a>
-	N03.540.348.500.500.600.650.225.520 Safety and Health (U.S.)	National Institute for Occupational
-	N03.540.348.500.500.600.650.400 Technology (U.S.)	National Center for Health Care
Old Tree	<del>N03.540.348.500.500.600.650.425</del>	<del>National Center for Health Statistics (U.S.)</del>
-	N03.540.348.500.500.600.650.592 Research and Quality	United States Agency for Healthcare
-	N03.540.348.500.500.600.650.760 Administration	United States Food and Drug
-	N03.540.348.500.500.600.650.790 Services Administration	United States Health Resources and
-	N03.540.348.500.500.600.650.790.525 Center (U.S.)	National Health Planning Information
-	N03.540.348.500.500.600.650.825	United States Indian Health Service
-	N03.540.348.500.500.600.650.912	United States Office of Research Integrity
-	N03.540.348.500.500.600.650.920 Health Services Administration	United States Substance Abuse and Mental
-	N03.540.348.500.500.650	United States Department of Homeland Security
-	N03.540.348.500.500.700	United States Department of Veterans Affairs
-	N03.540.348.500.500.937	United States Environmental Protection Agency
-	N03.540.348.500.500.968	United States Federal Trade Commission
-	N03.540.348.500.500.984 Administration	United States National Aeronautics and Space
-	N03.540.348.500.500.992 Administration	United States Occupational Safety and Health
-	N03.540.348.500.500.996	United States Office of Economic Opportunity
-	N03.540.348.500.500.997	United States Office of National Drug Control Policy

## MeSH Tree Changes for 2017

Type	Tree - heading
-	N03.540.348.500.500.998 United States Office of Technology Assessment
-	N03.540.348.500.500.999 United States Social Security Administration
-	N03.540.348.750 Local Government
-	N03.540.348.875 State Government
-	N03.540.400 Government Agencies
-	N03.540.400.500 Child Protective Services
-	N03.540.400.750 Federal Government
-	N03.540.452 Health Planning Organizations
-	N03.540.452.188 Health Care Coalitions
-	N03.540.452.210 Health Planning Councils
-	N03.540.452.508 State Health Planning and Development Agencies
-	N03.540.452.508.440 Health Systems Agencies
-	N03.540.483 Home Care Agencies
-	N03.540.514 International Agencies
-	N03.540.514.320 International Planned Parenthood Federation
-	N03.540.514.481 Organisation for Economic Co-Operation and Development
-	N03.540.514.642 Red Cross
-	N03.540.514.718 United Nations
-	N03.540.514.718.800 World Health Organization
-	N03.540.514.718.800.628 Pan American Health Organization
-	N03.540.571 Labor Unions
-	N03.540.571.356 Collective Bargaining
-	N03.540.571.608 Strikes, Employee
-	N03.540.630 Organizations, Nonprofit
-	N03.540.630.180 Foundations
-	N03.540.630.480 Health Level Seven
-	N03.540.630.630 Hospital Auxiliaries
-	N03.540.630.780 Voluntary Health Agencies
-	N03.540.630.780.080 American Cancer Society
-	N03.540.630.780.110 American Heart Association
-	N03.540.630.780.305 International Planned Parenthood Federation
-	N03.540.630.780.500 Mental Health Associations
-	N03.540.630.780.608 Red Cross
-	N03.540.630.780.850 Tuberculosis Societies
-	N03.540.706 Public-Private Sector Partnerships
-	N03.540.782 Self-Help Groups

## MeSH Tree Changes for 2017

Type	Tree - heading
-	N03.540.782.270 Alcoholics Anonymous
-	N03.540.828 Societies
-	N03.540.828.052 American Public Health Association
-	N03.540.828.105 American Speech-Language-Hearing Association
-	N03.540.828.506 Societies, Dental
-	N03.540.828.506.274 American Dental Association
-	N03.540.828.540 Societies, Hospital
-	N03.540.828.540.080 American Hospital Association
-	N03.540.828.589 Societies, Medical
-	N03.540.828.589.274 American Medical Association
-	N03.540.828.672 Societies, Nursing
-	N03.540.828.672.274 American Nurses' Association
-	N03.540.828.672.463 International Council of Nurses
-	N03.540.828.755 Societies, Pharmaceutical
-	N03.540.828.838 Societies, Scientific
-	N03.623 Policy
-	N03.623.500 Social Control Policies
-	N03.623.500.550 Organizational Policy
-	N03.623.500.608 Public Policy
-	N03.623.500.608.180 Environmental Policy
-	N03.623.500.608.214 Family Planning Policy
-	N03.623.500.608.428 Health Policy
-	N03.623.500.608.428.285 Health Care Reform
-	N03.623.500.608.428.650 Nutrition Policy
-	N03.623.500.608.428.650.500 Recommended Dietary Allowances
-	N03.706 Social Control, Formal
-	N03.706.110 Credentialing
-	N03.706.110.070 Accreditation
-	N03.706.110.070.410 Joint Commission on Accreditation of Healthcare Organizations
-	N03.706.110.120 Certification
-	N03.706.110.120.580 Specialty Boards
-	N03.706.110.510 Licensure
-	N03.706.110.510.180 Licensure, Dental
-	N03.706.110.510.290 Licensure, Hospital
-	N03.706.110.510.410 Licensure, Medical



## MeSH Tree Changes for 2017

Type	Tree - heading	
-	N03.706.110.510.490	Licensure, Nursing
-	N03.706.110.510.560	Licensure, Pharmacy
-	N03.706.280	Facility Regulation and Control
-	N03.706.358	Government Regulation
-	N03.706.437	Human Rights
-	N03.706.437.300	Child Advocacy
-	N03.706.437.352	Civil Rights
-	N03.706.437.352.500	Privacy
-	N03.706.437.352.500.320	Genetic Privacy
-	N03.706.437.368	Consumer Advocacy
-	N03.706.437.380	Freedom
-	N03.706.437.380.500	Personal Autonomy
-	N03.706.437.650	Patient Rights
-	N03.706.437.650.124	Confidentiality
-	N03.706.437.650.124.160	Data Anonymization
-	N03.706.437.650.124.320	Genetic Privacy
-	N03.706.437.650.312	Informed Consent
-	N03.706.437.650.312.500	Informed Consent By Minors
-	N03.706.437.650.406	Patient Access to Records
-	N03.706.437.650.750	Right to Die
-	N03.706.437.650.875	Treatment Refusal
New Heading	<b>N03.706.437.650.875.500</b>	<b>Vaccination Refusal</b>
-	N03.706.437.675	Reproductive Rights
-	N03.706.437.700	Social Justice
-	N03.706.437.850	Women's Rights
-	N03.706.535	Jurisprudence
-	N03.706.535.020	Advance Directives
-	N03.706.535.020.500	Living Wills
-	N03.706.535.125	Compensation and Redress
-	N03.706.535.230	Confidentiality
-	N03.706.535.230.270	Duty to Warn
-	N03.706.535.230.320	Genetic Privacy
-	N03.706.535.241	Contracts
-	N03.706.535.247	Duty to Recontact
-	N03.706.535.253	Expert Testimony

## MeSH Tree Changes for 2017

Type	Tree - heading
-	N03.706.535.351 Forensic Psychiatry
-	N03.706.535.351.200 Commitment of Mentally Ill
-	N03.706.535.351.528 Insanity Defense
-	N03.706.535.489 Informed Consent
-	N03.706.535.489.134 Consent Forms
-	N03.706.535.489.384 Informed Consent By Minors
-	N03.706.535.489.635 Third-Party Consent
-	N03.706.535.489.635.500 Parental Consent
-	N03.706.535.518 Intellectual Property
-	N03.706.535.518.300 Copyright
-	N03.706.535.518.650 Patents as Topic
-	N03.706.535.533 International Law
-	N03.706.535.547 Liability, Legal
-	N03.706.535.606 Malpractice
-	N03.706.535.606.300 Defensive Medicine
-	N03.706.535.606.528 Professional Impairment
-	N03.706.535.606.528.500 Physician Impairment
-	N03.706.535.615 Mandatory Reporting
-	N03.706.535.625 Mental Competency
-	N03.706.535.664 Presumed Consent
-	N03.706.535.704 Resuscitation Orders
-	N03.706.535.803 Wrongful Life
-	N03.706.615 Legislation as Topic
-	N03.706.615.049 American Recovery and Reinvestment Act
-	N03.706.615.100 Antitrust Laws
-	N03.706.615.215 Employee Retirement Income Security Act
-	N03.706.615.273 Health Insurance Portability and Accountability Act
-	N03.706.615.302 Health Planning
-	N03.706.615.331 Legislation, Dental
-	N03.706.615.402 Legislation, Drug
-	N03.706.615.402.250 Drug and Narcotic Control
-	N03.706.615.412 Legislation, Food
-	N03.706.615.420 Legislation, Hospital
-	N03.706.615.420.249 Medicare Part C
-	N03.706.615.473 Legislation, Medical
-	N03.706.615.473.500 Medical Device Legislation

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	N03.706.615.544	Legislation, Nursing
-	N03.706.615.615	Legislation, Pharmacy
-	N03.706.615.686	Legislation, Veterinary
-	N03.706.615.693	Medicaid
-	N03.706.615.696	Medicare
New Heading	<b>N03.706.615.696.500 2015</b>	<b>Medicare Access and CHIP Reauthorization Act of</b>
-	N03.706.615.698	Medicare Part A
-	N03.706.615.699	Medicare Part B
-	N03.706.615.752	Medicare Part D
-	N03.706.615.805	Patient Freedom of Choice Laws
-	N03.706.615.806	Patient Protection and Affordable Care Act
-	N03.706.615.808	Patient Self-Determination Act
-	N03.706.615.862	Privacy
-	N03.706.615.889	Tax Equity and Fiscal Responsibility Act
-	N03.706.615.917	Threshold Limit Values
-	N03.706.657	Mandatory Programs
-	N03.706.657.249	Mandatory Reporting
-	N03.706.657.500	Mandatory Testing
-	N03.706.678	Patient Advocacy
-	N03.706.700	Peer Review
-	N03.706.700.690	Peer Review, Health Care
-	N03.706.700.700	Peer Review, Research
-	N03.706.742	Policy Making
-	N03.706.742.500	Advisory Committees
-	N03.787	State Dentistry
-	N03.858	State Medicine
-	N03.880	Technology Assessment, Biomedical
-	N03.880.502	Technology, High-Cost
-	N04	Health Services Administration
New Heading	<b>N04.226</b>	<b>Intersectoral Collaboration</b>
-	N04.452	Organization and Administration
-	N04.452.070	Annual Reports as Topic
-	N04.452.095	Appointments and Schedules
-	N04.452.095.738	Waiting Lists

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	N04.452.105	Capacity Building
-	N04.452.115	Clinical Governance
-	N04.452.122	Committee Membership
-	N04.452.150	Constitution and Bylaws
-	N04.452.190	Decision Making, Organizational
-	N04.452.227	Efficiency, Organizational
-	N04.452.264	Eligibility Determination
-	N04.452.313	Fee Schedules
-	N04.452.313.500	Relative Value Scales
-	N04.452.394	Governing Board
-	N04.452.394.600	Trustees
-	N04.452.418	Health Facility Administration
-	N04.452.442	Hospital Administration
-	N04.452.442.060	Ancillary Services, Hospital
-	N04.452.442.110	Centralized Hospital Services
-	N04.452.442.180	Financial Management, Hospital
-	N04.452.442.322	Hospital Communication Systems
-	N04.452.442.422	Hospital Departments
-	N04.452.442.422.040	Admitting Department, Hospital
-	N04.452.442.422.070	Anesthesia Department, Hospital
-	N04.452.442.422.215	Cardiology Service, Hospital
-	N04.452.442.422.260	Central Supply, Hospital
-	N04.452.442.422.270	Chaplaincy Service, Hospital
-	N04.452.442.422.298	Dental Service, Hospital
-	N04.452.442.422.310	Education Department, Hospital
-	N04.452.442.422.336	Emergency Service, Hospital
-	N04.452.442.422.336.400	Trauma Centers
-	N04.452.442.422.374	Food Service, Hospital
-	N04.452.442.422.412	Housekeeping, Hospital
-	N04.452.442.422.420	Laboratories, Hospital
-	N04.452.442.422.428	Laundry Service, Hospital
-	N04.452.442.422.450	Maintenance and Engineering, Hospital
-	N04.452.442.422.465	Medical Records Department, Hospital
New Heading	<b>N04.452.442.422.467</b>	<b>Morgue</b>
-	N04.452.442.422.472	Nuclear Medicine Department, Hospital

## MeSH Tree Changes for 2017

Type	Tree - heading
-	N04.452.442.422.489 Nursing Service, Hospital
-	N04.452.442.422.495 Obstetrics and Gynecology Department, Hospital
-	N04.452.442.422.500 Occupational Therapy Department, Hospital
-	N04.452.442.422.513 Oncology Service, Hospital
-	N04.452.442.422.527 Outpatient Clinics, Hospital
-	N04.452.442.422.527.600 Pain Clinics
-	N04.452.442.422.535 Pathology Department, Hospital
-	N04.452.442.422.565 Personnel Administration, Hospital
-	N04.452.442.422.603 Pharmacy Service, Hospital
-	N04.452.442.422.615 Physical Therapy Department, Hospital
-	N04.452.442.422.641 Psychiatric Department, Hospital
-	N04.452.442.422.679 Purchasing, Hospital
-	N04.452.442.422.679.375 Group Purchasing
-	N04.452.442.422.695 Radiology Department, Hospital
-	N04.452.442.422.712 Respiratory Therapy Department, Hospital
-	N04.452.442.422.730 Social Work Department, Hospital
-	N04.452.442.422.750 Surgery Department, Hospital
-	N04.452.442.422.790 Urology Department, Hospital
-	N04.452.442.450 Hospital Distribution Systems
-	N04.452.442.452 Hospital Information Systems
-	N04.452.442.452.050 Ambulatory Care Information Systems
-	N04.452.442.452.500 Medical Order Entry Systems
-	N04.452.442.452.600 Operating Room Information Systems
-	N04.452.442.452.680 Point-of-Care Systems
-	N04.452.442.510 Hospital-Patient Relations
-	N04.452.442.530 Hospital-Physician Relations
-	N04.452.442.530.500 Medical Staff Privileges
-	N04.452.442.550 Hospital Restructuring
-	N04.452.442.550.380 Hospital-Physician Joint Ventures
-	N04.452.442.550.380.500 Physician Self-Referral
-	N04.452.442.565 Hospital Shared Services
-	N04.452.442.585 Hospital Shops
-	N04.452.442.600 Libraries, Hospital
-	N04.452.442.650 Materials Management, Hospital
-	N04.452.442.650.300 Inventories, Hospital
-	N04.452.442.675 Medication Systems, Hospital

## MeSH Tree Changes for 2017

Type	Tree - heading
-	N04.452.442.700 Product Line Management
-	N04.452.460 Institutional Management Teams
-	N04.452.480 Knowledge Management
-	N04.452.500 Management Audit
-	N04.452.500.150 Benchmarking
-	N04.452.515 Management Information Systems
-	N04.452.515.080 Clinical Laboratory Information Systems
-	N04.452.515.095 Clinical Pharmacy Information Systems
-	N04.452.515.110 Database Management Systems
-	N04.452.515.135 Decision Support Systems, Management
-	N04.452.515.350 Healthcare Common Procedure Coding System
-	N04.452.515.360 Hospital Information Systems
-	N04.452.515.360.050 Ambulatory Care Information Systems
-	N04.452.515.360.500 Medical Order Entry Systems
-	N04.452.515.360.555 Operating Room Information Systems
-	N04.452.515.360.652 Point-of-Care Systems
-	N04.452.515.708 Office Automation
-	N04.452.515.708.950 Word Processing
-	N04.452.515.800 Personnel Staffing and Scheduling Information Systems
-	N04.452.515.825 Radiology Information Systems
-	N04.452.515.825.500 Teleradiology
-	N04.452.521 Mandatory Programs
-	N04.452.528 Medication Systems
-	N04.452.528.460 Medication Reconciliation
-	N04.452.528.473 Medication Systems, Hospital
-	N04.452.534 Models, Organizational
-	N04.452.540 Multi-Institutional Systems
-	N04.452.540.308 Hospital Shared Services
-	N04.452.602 Organizational Affiliation
-	N04.452.602.713 Transfer Agreement
-	N04.452.606 Organizational Culture
-	N04.452.610 Organizational Innovation
-	N04.452.610.250 Entrepreneurship
-	N04.452.615 Organizational Objectives
-	N04.452.633 Ownership
-	N04.452.633.390 Private Sector

## MeSH Tree Changes for 2017

Type	Tree - heading
-	N04.452.633.780 Privatization
-	N04.452.633.890 Public Sector
-	N04.452.653 Patient Identification Systems
-	N04.452.653.250 Health Smart Cards
-	N04.452.653.500 Radio Frequency Identification Device
-	N04.452.677 Personnel Management
-	N04.452.677.356 Collective Bargaining
-	N04.452.677.358 Employee Discipline
-	N04.452.677.362 Employee Grievances
-	N04.452.677.368 Employee Incentive Plans
-	N04.452.677.370 Employee Performance Appraisal
-	N04.452.677.400 Job Application
-	N04.452.677.410 Job Description
-	N04.452.677.420 Management Quality Circles
-	N04.452.677.430 Negotiating
-	N04.452.677.440 Personnel Administration, Hospital
-	N04.452.677.445 Personnel Delegation
-	N04.452.677.450 Personnel Downsizing
-	N04.452.677.460 Personnel Loyalty
-	N04.452.677.500 Personnel Selection
-	N04.452.677.650 Personnel Staffing and Scheduling
-	N04.452.677.680 Personnel Turnover
-	N04.452.677.740 Physician Incentive Plans
-	N04.452.677.800 Salaries and Fringe Benefits
-	N04.452.677.800.162 Family Leave
-	N04.452.677.800.162.500 Parental Leave
-	N04.452.677.800.325 Health Benefit Plans, Employee
-	N04.452.677.800.325.300 Employee Retirement Income Security Act
-	N04.452.677.800.662 Sick Leave
-	N04.452.677.822 Staff Development
-	N04.452.677.842 Strikes, Employee
-	N04.452.677.950 Workload
-	N04.452.677.975 Workplace
-	N04.452.706 Pharmacy Administration
-	N04.452.706.310 Drug and Narcotic Control
-	N04.452.706.310.500 Drug Recalls

## MeSH Tree Changes for 2017

Type	Tree - heading
-	N04.452.706.310.750 Safety-Based Drug Withdrawals
-	N04.452.706.477 Drug Utilization
-	N04.452.706.477.400 Drug Utilization Review
-	N04.452.718 Planning Techniques
New Heading	<b>N04.452.718.500 Strategic Planning</b>
-	N04.452.758 Professional Practice
-	N04.452.758.244 Group Practice
-	N04.452.758.244.369 Group Practice, Dental
-	N04.452.758.244.400 Group Practice, Prepaid
-	N04.452.758.244.425 Health Maintenance Organizations
-	N04.452.758.244.450 Independent Practice Associations
-	N04.452.758.275 Hospital-Physician Relations
-	N04.452.758.275.500 Medical Staff Privileges
-	N04.452.758.307 House Calls
-	N04.452.758.367 Institutional Practice
-	N04.452.758.372 Management Service Organizations
-	N04.452.758.377 Nursing
-	N04.452.758.377.500 Nursing, Private Duty
-	N04.452.758.377.750 Nursing, Supervisory
-	N04.452.758.377.875 Office Nursing
-	N04.452.758.377.937 Telenursing
-	N04.452.758.377.968 Travel Nursing
-	N04.452.758.388 Nursing Faculty Practice
-	N04.452.758.635 Office Visits
-	N04.452.758.635.500 No-Show Patients
-	N04.452.758.671 Partnership Practice
-	N04.452.758.671.589 Partnership Practice, Dental
-	N04.452.758.708 Practice Management
-	N04.452.758.708.200 Office Management
-	N04.452.758.708.200.400 Forms and Records Control
-	N04.452.758.708.200.400.500 Clinical Coding
-	N04.452.758.708.300 Practice Management, Dental
-	N04.452.758.708.450 Practice Management, Medical
-	N04.452.758.708.450.500 Concierge Medicine
-	N04.452.758.708.525 Practice Management, Veterinary



## MeSH Tree Changes for 2017

Type	Tree - heading
-	N04.452.758.708.600 Practice Valuation and Purchase
-	N04.452.758.745 Private Practice
-	N04.452.758.745.225 Fee-for-Service Plans
-	N04.452.758.745.450 Independent Practice Associations
-	N04.452.758.752 Professional Autonomy
-	N04.452.758.760 Professional Corporations
-	N04.452.758.772 Professional Practice Location
-	N04.452.758.788 Professional Staff Committees
-	N04.452.758.788.300 Ethics Committees
-	N04.452.758.788.300.500 Ethics Committees, Clinical
-	N04.452.758.788.300.750 Ethics Committees, Research
-	N04.452.758.788.682 Pharmacy and Therapeutics Committee
-	N04.452.758.849 Referral and Consultation
-	N04.452.758.849.174 Ethics Consultation
-	N04.452.758.849.350 Gatekeeping
-	N04.452.758.849.500 Physician Self-Referral
-	N04.452.758.849.550 Remote Consultation
-	N04.452.758.849.550.500 Distance Counseling
-	N04.452.758.849.775 Secondary Care
-	N04.452.758.849.887 Tertiary Healthcare
-	N04.452.760 Program Development
-	N04.452.794 Public Health Administration
-	N04.452.822 Public Relations
-	N04.452.822.070 Anniversaries and Special Events
-	N04.452.822.210 Community-Institutional Relations
-	N04.452.822.370 Hospital-Patient Relations
-	N04.452.822.379 Hospital-Physician Relations
-	N04.452.822.388 Interdepartmental Relations
-	N04.452.822.400 Interinstitutional Relations
-	N04.452.822.700 Patient Satisfaction
-	N04.452.822.700.500 Patient Preference
-	N04.452.859 Records as Topic
-	N04.452.859.132 Birth Certificates
-	N04.452.859.198 Consent Forms
-	N04.452.859.264 Death Certificates
-	N04.452.859.341 Dental Records

## MeSH Tree Changes for 2017

Type	Tree - heading
-	N04.452.859.360 Diet Records
-	N04.452.859.380 Hospital Records
-	N04.452.859.564 Medical Records
-	N04.452.859.564.137 Administrative Claims, Healthcare
-	N04.452.859.564.274 Clinical Coding
-	N04.452.859.564.550 Medical Record Linkage
-	N04.452.859.564.600 Medical Records, Problem-Oriented
-	N04.452.859.564.650 Medical Records Systems, Computerized
-	N04.452.859.564.650.250 Health Smart Cards
-	N04.452.859.564.650.500 Medical Order Entry Systems
-	N04.452.859.564.725 Patient Discharge Summaries
-	N04.452.859.564.800 Trauma Severity Indices
-	N04.452.859.564.800.125 Abbreviated Injury Scale
-	N04.452.859.564.800.250 Glasgow Coma Scale
-	N04.452.859.564.800.260 Glasgow Outcome Scale
-	N04.452.859.564.800.500 Injury Severity Score
-	N04.452.859.675 Nursing Records
-	N04.452.859.819 Registries
-	N04.452.859.819.725 SEER Program
-	N04.452.871 Risk Management
-	N04.452.871.715 Risk Assessment
-	N04.452.871.715.400 Healthcare Failure Mode and Effect Analysis
-	N04.452.871.715.800 Risk Adjustment
-	N04.452.871.800 Risk Sharing, Financial
-	N04.452.871.900 Safety Management
-	N04.452.871.900.500 Material Safety Data Sheets
-	N04.452.910 Security Measures
-	N04.452.910.099 Biometric Identification
-	N04.452.910.200 Computer Security
-	N04.452.910.200.500 Data Anonymization
-	N04.452.932 Time Management
-	N04.452.955 Total Quality Management
-	N04.452.977 Voluntary Programs
-	N04.590 Patient Care Management
-	N04.590.233 Comprehensive Health Care
-	N04.590.233.250 Comprehensive Dental Care

## MeSH Tree Changes for 2017

Type	Tree - heading
-	N04.590.233.508                      Nursing Process
-	N04.590.233.508.480                  Nursing Assessment
-	N04.590.233.508.480.110              Nursing Diagnosis
-	N04.590.233.508.613                  Nursing Research
-	N04.590.233.508.613.234              Clinical Nursing Research
-	N04.590.233.508.613.385              Nursing Administration Research
-	N04.590.233.508.613.413              Nursing Education Research
-	N04.590.233.508.613.432              Nursing Evaluation Research
-	N04.590.233.508.613.634              Nursing Methodology Research
-	N04.590.233.624                      Patient Care Planning
-	N04.590.233.624.124                  Advance Care Planning
-	N04.590.233.624.124.050              Advance Directives
-	N04.590.233.624.124.050.500        Living Wills
-	N04.590.233.624.250                  Case Management
-	N04.590.233.624.625                  Critical Pathways
-	N04.590.233.727                      Primary Health Care
-	N04.590.233.727.210                  Continuity of Patient Care
New Tree	<a href="#">N04.590.233.727.210.063</a> <a href="#">Aftercare</a>
-	N04.590.233.727.210.125              Patient Discharge
-	N04.590.233.727.210.249              Patient Handoff
-	N04.590.233.727.210.624              Patient Transfer
-	N04.590.233.727.210.718              Transition to Adult Care
-	N04.590.233.727.210.812              Transitional Care
-	N04.590.233.727.407                  Patient-Centered Care
-	N04.590.233.727.407.500              Patient Navigation
-	N04.590.233.727.605                  Refusal to Treat
-	N04.590.233.799                      Progressive Patient Care
-	N04.590.254                          Crew Resource Management, Healthcare
-	N04.590.275                          Critical Pathways
-	N04.590.374                          Delivery of Health Care
-	N04.590.374.034                      After-Hours Care
-	N04.590.374.034.500                  Answering Services
-	N04.590.374.052                      Culturally Competent Care
-	N04.590.374.070                      Delegation, Professional
-	N04.590.374.142                      Delivery of Health Care, Integrated

## MeSH Tree Changes for 2017

Type	Tree - heading
-	N04.590.374.142.600      Provider-Sponsored Organizations
-	N04.590.374.285      Health Care Reform
-	N04.590.374.350      Health Services Accessibility
-	N04.590.374.350.500      Health Equity
-	N04.590.374.380      Healthcare Disparities
-	N04.590.374.410      Managed Care Programs
-	N04.590.374.410.175      Competitive Medical Plans
-	N04.590.374.410.400      Health Maintenance Organizations
-	N04.590.374.410.450      Independent Practice Associations
-	N04.590.374.410.700      Patient Freedom of Choice Laws
-	N04.590.374.410.750      Preferred Provider Organizations
-	N04.590.374.410.875      Provider-Sponsored Organizations
-	N04.590.374.505      Practice Patterns, Dentists'
-	N04.590.374.553      Practice Patterns, Nurses'
-	N04.590.374.577      Practice Patterns, Physicians'
-	N04.590.374.600      Product Line Management
New Heading	<b>N04.590.374.650      Professional Practice Gaps</b>
-	N04.590.374.700      Safety-net Providers
-	N04.590.374.800      Telemedicine
-	N04.590.374.800.550      Remote Consultation
-	N04.590.374.800.600      Telepathology
-	N04.590.374.800.700      Teleradiology
-	N04.590.374.800.850      Telerehabilitation
-	N04.590.374.900      Uncompensated Care
-	N04.590.607      Disease Management
-	N04.590.607.500      Pain Management
-	N04.590.656      Medication Reconciliation
-	N04.590.661      Medication Therapy Management
-	N04.590.715      Patient Care Team
-	N04.590.715.550      Hospital Rapid Response Team
-	N04.590.715.571      Nursing, Team
-	N04.590.731      Patient Selection
-	N04.590.874      Point-of-Care Systems
-	N04.590.874.500      Point-of-Care Testing
-	N04.761      Quality of Health Care

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	N04.761.040	Advance Directive Adherence
-	N04.761.210	Clinical Competence
-	N04.761.337	Guideline Adherence
-	N04.761.559	Outcome and Process Assessment (Health Care)
-	N04.761.559.590	Outcome Assessment (Health Care)
-	N04.761.559.590.200	Failure to Rescue, Health Care
-	N04.761.559.590.399	Patient Outcome Assessment
-	N04.761.559.590.399.250	Critical Care Outcomes
-	N04.761.559.590.399.500	Lysholm Knee Score
New Heading	<b>N04.761.559.590.399.750</b>	<b>Minimal Clinically Important Difference</b>
New Heading	<b>N04.761.559.590.399.875</b>	<b>Patient Reported Outcome Measures</b>
-	N04.761.559.590.800	Treatment Outcome
-	N04.761.559.590.800.379	Early Termination of Clinical Trials
-	N04.761.559.590.800.569	Response Evaluation Criteria in Solid Tumors
New Heading	<b>N04.761.559.590.800.665</b>	<b>Sustained Virologic Response</b>
-	N04.761.559.590.800.760	Treatment Failure
-	N04.761.559.590.800.760.500	Failure to Rescue, Health Care
-	N04.761.559.590.900	Watchful Waiting
-	N04.761.559.650	Process Assessment (Health Care)
-	N04.761.616	Peer Review, Health Care
-	N04.761.673	Professional Review Organizations
-	N04.761.685	Program Evaluation
-	N04.761.685.150	Benchmarking
-	N04.761.700	Quality Assurance, Health Care
New Heading	<b>N04.761.700.075</b>	<b>Alert Fatigue, Health Personnel</b>
-	N04.761.700.150	Benchmarking
-	N04.761.700.250	Clinical Audit
-	N04.761.700.250.295	Dental Audit
-	N04.761.700.250.500	Medical Audit
-	N04.761.700.250.500.100	Commission on Professional and Hospital Activities
-	N04.761.700.250.520	Nursing Audit
-	N04.761.700.350	Guidelines as Topic
-	N04.761.700.350.324	Codes of Ethics

## MeSH Tree Changes for 2017

Type	Tree - heading
-	N04.761.700.350.650 Practice Guidelines as Topic
-	N04.761.700.512 Laboratory Proficiency Testing
-	N04.761.700.594 Near Miss, Healthcare
-	N04.761.700.615 Potentially Inappropriate Medication List
-	N04.761.700.635 Time Out, Healthcare
-	N04.761.700.675 Total Quality Management
-	N04.761.744 Quality Improvement
-	N04.761.744.500 Meaningful Use
New Heading	<b>N04.761.744.750 Value-Based Insurance</b>
-	N04.761.789 Quality Indicators, Health Care
-	N04.761.789.800 Risk Adjustment
-	N04.761.789.900 Standard of Care
-	N04.761.879 Utilization Review
-	N04.761.879.200 Concurrent Review
-	N04.761.879.300 Drug Utilization Review
-	N05 Health Care Quality, Access, and Evaluation
-	N05.300 Delivery of Health Care
-	N05.300.049 After-Hours Care
-	N05.300.049.500 Answering Services
-	N05.300.100 Attitude of Health Personnel
New Heading	<b>N05.300.100.169 Alert Fatigue, Health Personnel</b>
-	N05.300.100.337 Nurse's Role
-	N05.300.100.675 Refusal to Treat
-	N05.300.125 Attitude to Death
-	N05.300.150 Attitude to Health
-	N05.300.150.395 Health Services Misuse
-	N05.300.150.395.450 Medical Overuse
-	N05.300.150.395.450.500 Unnecessary Procedures
-	N05.300.150.410 Health Knowledge, Attitudes, Practice
-	N05.300.150.600 Patient Acceptance of Health Care
-	N05.300.150.600.600 Patient Compliance
-	N05.300.150.600.600.500 Medication Adherence
-	N05.300.150.600.600.750 No-Show Patients
-	N05.300.150.600.610 Patient Dropouts

## MeSH Tree Changes for 2017

Type	Tree - heading
-	N05.300.150.600.620 Patient Participation
-	N05.300.150.600.630 Patient Satisfaction
-	N05.300.150.600.630.500 Patient Preference
-	N05.300.150.600.800 Treatment Refusal
New Heading	<b>N05.300.150.600.800.500 Vaccination Refusal</b>
-	N05.300.206 Culturally Competent Care
-	N05.300.262 Delivery of Health Care, Integrated
-	N05.300.262.600 Provider-Sponsored Organizations
-	N05.300.375 Health Care Costs
-	N05.300.375.250 Direct Service Costs
-	N05.300.375.300 Drug Costs
-	N05.300.375.350 Employer Health Costs
-	N05.300.375.500 Hospital Costs
-	N05.300.380 Health Care Reform
-	N05.300.380.500 Accountable Care Organizations
-	N05.300.385 Health Expenditures
-	N05.300.400 Health Priorities
-	N05.300.420 Health Resources
-	N05.300.420.400 Health Manpower
-	N05.300.420.700 Strategic Stockpile
-	N05.300.430 Health Services Accessibility
-	N05.300.430.375 Health Care Rationing
-	N05.300.430.383 Health Equity
-	N05.300.430.390 Health Facility Closure
-	N05.300.430.400 Health Facility Environment
-	N05.300.430.410 Health Facility Size
-	N05.300.430.500 Marketing of Health Services
-	N05.300.430.500.500 Social Marketing
-	N05.300.450 Health Services Needs and Demand
-	N05.300.450.520 Medically Underserved Area
-	N05.300.450.760 Surge Capacity
-	N05.300.493 Healthcare Disparities
-	N05.300.515 Medical Tourism
-	N05.300.537 Needs Assessment
-	N05.300.580 Practice Patterns, Dentists'

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	N05.300.581	Practice Patterns, Nurses'
-	N05.300.625	Practice Patterns, Physicians'
New Heading	<b>N05.300.643</b>	<b>Professional Practice Gaps</b>
-	N05.300.660	Professional-Patient Relations
-	N05.300.660.325	Dentist-Patient Relations
-	N05.300.660.560	Nurse-Patient Relations
-	N05.300.660.625	Physician-Patient Relations
-	N05.300.745	Secondary Care
-	N05.300.787	Tertiary Healthcare
-	N05.300.830	Uncompensated Care
-	N05.350	Ethics
-	N05.350.100	Bioethical Issues
-	N05.350.200	Bioethics
-	N05.350.200.500	Ethics, Clinical
-	N05.350.213	Codes of Ethics
-	N05.350.213.249	Helsinki Declaration
-	N05.350.213.400	Hippocratic Oath
-	N05.350.219	Complicity
-	N05.350.225	Conflict of Interest
-	N05.350.225.500	Physician Self-Referral
-	N05.350.238	Double Effect Principle
-	N05.350.244	Ethical Analysis
-	N05.350.244.150	Casuistry
-	N05.350.244.750	Retrospective Moral Judgment
-	N05.350.244.875	Wedge Argument
-	N05.350.247	Ethical Relativism
-	N05.350.250	Ethical Review
-	N05.350.250.200	Ethics Consultation
-	N05.350.256	Ethical Theory
-	N05.350.262	Ethicists
-	N05.350.268	Ethics Committees
-	N05.350.268.500	Ethics Committees, Clinical
-	N05.350.268.750	Ethics Committees, Research
-	N05.350.297	Ethics, Business
-	N05.350.325	Ethics, Institutional



## MeSH Tree Changes for 2017

Type	Tree - heading	
-	N05.350.340	Ethics, Professional
-	N05.350.340.080	Codes of Ethics
-	N05.350.340.080.249	Helsinki Declaration
-	N05.350.340.162	Ethics, Clinical
-	N05.350.340.162.249	Ethics, Dental
-	N05.350.340.162.500	Ethics, Medical
-	N05.350.340.162.500.500	Hippocratic Oath
-	N05.350.340.162.500.750	Therapeutic Equipoise
-	N05.350.340.162.750	Ethics, Nursing
-	N05.350.340.162.875	Ethics, Pharmacy
-	N05.350.340.581	Professionalism
-	N05.350.670	Ethics, Research
-	N05.350.670.250	Censorship, Research
-	N05.350.670.500	Helsinki Declaration
-	N05.350.670.750	Therapeutic Equipoise
-	N05.350.670.875	Therapeutic Misconception
-	N05.350.835	Humanism
Old Tree	N05.350.835.400	Feminism
-	N05.350.917	Personhood
-	N05.350.958	Principle-Based Ethics
-	N05.350.958.500	Beneficence
-	N05.350.958.650	Personal Autonomy
-	N05.350.958.750	Social Justice
-	N05.350.979	Professional Misconduct
-	N05.350.979.500	Scientific Misconduct
-	N05.425	Health Services Research
-	N05.425.104	Community-Based Participatory Research
-	N05.425.157	Comparative Effectiveness Research
New Heading	N05.425.184	Global Burden of Disease
-	N05.425.210	Health Care Surveys
New Heading	N05.425.210.500	Patient Reported Outcome Measures
New Heading	N05.632	Public Health Systems Research
-	N05.700	Quality Assurance, Health Care
New	N05.700.075	Alert Fatigue, Health Personnel

## MeSH Tree Changes for 2017

Type	Tree - heading
Heading	
-	N05.700.150                      Benchmarking
-	N05.700.175                      Clinical Audit
-	N05.700.175.250                  Dental Audit
-	N05.700.175.500                  Medical Audit
-	N05.700.175.500.200              Commission on Professional and Hospital Activities
-	N05.700.175.520                  Nursing Audit
-	N05.700.200                      Credentialing
-	N05.700.200.100                  Accreditation
-	N05.700.200.100.420              Joint Commission on Accreditation of Healthcare Organizations
-	N05.700.200.190                  Certification
-	N05.700.200.190.750              Specialty Boards
-	N05.700.200.450                  Licensure
-	N05.700.200.450.300              Licensure, Dental
-	N05.700.200.450.380              Licensure, Hospital
-	N05.700.200.450.500              Licensure, Medical
-	N05.700.200.450.550              Licensure, Nursing
-	N05.700.200.450.625              Licensure, Pharmacy
-	N05.700.300                      Facility Regulation and Control
-	N05.700.350                      Guidelines as Topic
-	N05.700.350.324                  Codes of Ethics
-	N05.700.350.650                  Practice Guidelines as Topic
-	N05.700.513                      Near Miss, Healthcare
-	N05.700.594                      Potentially Inappropriate Medication List
-	N05.700.675                      Professional Review Organizations
-	N05.700.685                      Professional Staff Committees
-	N05.700.685.149                  Clinical Trials Data Monitoring Committees
-	N05.700.685.300                  Ethics Committees
-	N05.700.685.300.500              Ethics Committees, Clinical
-	N05.700.685.300.750              Ethics Committees, Research
-	N05.700.685.625                  Pharmacy and Therapeutics Committee
-	N05.700.739                      Time Out, Healthcare
-	N05.700.792                      Total Quality Management
-	N05.700.900                      Utilization Review
-	N05.700.900.200                  Concurrent Review

## MeSH Tree Changes for 2017

Type	Tree - heading
-	N05.700.900.300 Drug Utilization Review
-	N05.715 Quality of Health Care
-	N05.715.175 Clinical Competence
-	N05.715.350 Epidemiologic Factors
-	N05.715.350.075 Age Factors
-	N05.715.350.075.100 Age of Onset
-	N05.715.350.075.550 Maternal Age
-	N05.715.350.150 Bias (Epidemiology)
-	N05.715.350.150.675 Observer Variation
-	N05.715.350.150.775 Selection Bias
-	N05.715.350.200 Causality
-	N05.715.350.200.650 Precipitating Factors
-	N05.715.350.200.675 Protective Factors
-	N05.715.350.200.700 Risk Factors
-	N05.715.350.225 Comorbidity
-	N05.715.350.240 Confounding Factors (Epidemiology)
-	N05.715.350.350 Effect Modifier, Epidemiologic
-	N05.715.350.350.225 Cohort Effect
-	N05.715.350.350.375 Healthy Worker Effect
-	N05.715.350.350.625 Placebo Effect
-	N05.715.350.350.625.500 Nocebo Effect
-	N05.715.350.575 Reproductive History
-	N05.715.350.675 Sex Factors
-	N05.715.360 Health Care Evaluation Mechanisms
-	N05.715.360.300 Data Collection
-	N05.715.360.300.179 Checklist
-	N05.715.360.300.202 Data Accuracy
-	N05.715.360.300.224 Datasets as Topic
-	N05.715.360.300.269 Focus Groups
-	N05.715.360.300.360 Geriatric Assessment
-	N05.715.360.300.400 Interviews as Topic
-	N05.715.360.300.480 Narration
-	N05.715.360.300.560 Nutrition Assessment
-	N05.715.360.300.715 Records as Topic
-	N05.715.360.300.715.175 Birth Certificates
-	N05.715.360.300.715.245 Consent Forms

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	N05.715.360.300.715.315	Death Certificates
-	N05.715.360.300.715.330	Dental Records
-	N05.715.360.300.715.380	Hospital Records
-	N05.715.360.300.715.500	Medical Records
-	N05.715.360.300.715.500.249	Clinical Coding
-	N05.715.360.300.715.500.500	Medical Record Linkage
-	N05.715.360.300.715.500.520	Medical Records, Problem-Oriented
-	N05.715.360.300.715.500.530	Medical Records Systems, Computerized
-	N05.715.360.300.715.500.530.500	Health Smart Cards
-	N05.715.360.300.715.500.665	Patient Discharge Summaries
-	N05.715.360.300.715.500.800	Trauma Severity Indices
-	N05.715.360.300.715.500.800.100	Abbreviated Injury Scale
-	N05.715.360.300.715.500.800.325	Glasgow Coma Scale
-	N05.715.360.300.715.500.800.335	Glasgow Outcome Scale
-	N05.715.360.300.715.500.800.400	Injury Severity Score
-	N05.715.360.300.715.550	Nursing Records
-	N05.715.360.300.715.700	Registries
-	N05.715.360.300.715.700.725	SEER Program
-	N05.715.360.300.800	Surveys and Questionnaires
-	N05.715.360.300.800.313	Contraceptive Prevalence Surveys
-	N05.715.360.300.800.344	Health Care Surveys
New Heading	<b>N05.715.360.300.800.344.500</b>	<b>Patient Reported Outcome Measures</b>
-	N05.715.360.300.800.438	Health Surveys
-	N05.715.360.300.800.438.149	Behavioral Risk Factor Surveillance System
-	N05.715.360.300.800.438.300	Dental Health Surveys
-	N05.715.360.300.800.438.300.325	Dental Plaque Index
-	N05.715.360.300.800.438.300.340	DMF Index
-	N05.715.360.300.800.438.300.505	Index of Orthodontic Treatment Need
-	N05.715.360.300.800.438.300.670	Oral Hygiene Index
-	N05.715.360.300.800.438.300.690	Periodontal Index
-	N05.715.360.300.800.438.375	Health Status Indicators
Old Tree	<b>N05.715.360.300.800.438.375.181</b>	<b>Organ Dysfunction Scores</b>
-	N05.715.360.300.800.438.375.364	Patient Acuity
-	N05.715.360.300.800.438.375.364.500	Severity of Illness Index
New	<b>N05.715.360.300.800.438.375.364.500.250</b>	<b>APACHE</b>

## MeSH Tree Changes for 2017

Type	Tree - heading	
Tree		
-	N05.715.360.300.800.438.375.364.500.500	Karnofsky Performance Status
New Tree	<a href="#">N05.715.360.300.800.438.375.364.500.625</a>	<a href="#">Organ Dysfunction Scores</a>
New Heading	<b>N05.715.360.300.800.438.375.364.500.750 Score</b>	<b>Simplified Acute Physiology</b>
-	N05.715.360.300.800.438.375.730	Sickness Impact Profile
-	N05.715.360.300.800.438.500	Mass Screening
-	N05.715.360.300.800.438.500.174	Anonymous Testing
-	N05.715.360.300.800.438.500.500	Mass Chest X-Ray
-	N05.715.360.300.800.438.500.540	Multiphasic Screening
-	N05.715.360.300.800.438.500.575	Neonatal Screening
-	N05.715.360.300.800.438.500.825	Vision Screening
-	N05.715.360.300.800.438.625	Population Surveillance
-	N05.715.360.300.800.438.625.324	Public Health Surveillance
-	N05.715.360.300.800.438.625.650	Sentinel Surveillance
-	N05.715.360.300.800.469	Nutrition Surveys
-	N05.715.360.300.800.469.300	Diet Surveys
-	N05.715.360.300.800.500	Self Report
-	N05.715.360.317	Advance Directive Adherence
-	N05.715.360.335	Evaluation Studies as Topic
-	N05.715.360.335.500	Validation Studies as Topic
-	N05.715.360.395	Guideline Adherence
New Heading	<b>N05.715.360.425</b>	<b>Independent Medical Evaluation</b>
-	N05.715.360.455	Organizational Case Studies
-	N05.715.360.575	Outcome and Process Assessment (Health Care)
-	N05.715.360.575.575	Outcome Assessment (Health Care)
-	N05.715.360.575.575.200	Critical Care Outcomes
-	N05.715.360.575.575.300	Failure to Rescue, Health Care
-	N05.715.360.575.575.399	Patient Outcome Assessment
New Tree	<a href="#">N05.715.360.575.575.399.250</a>	<a href="#">Critical Care Outcomes</a>
-	N05.715.360.575.575.399.500	Lysholm Knee Score
New Heading	<b>N05.715.360.575.575.399.750</b>	<b>Minimal Clinically Important Difference</b>
New Heading	<b>N05.715.360.575.575.399.875</b>	<b>Patient Reported Outcome Measures</b>

## MeSH Tree Changes for 2017

Type	Tree - heading
-	N05.715.360.575.575.800 Treatment Outcome
-	N05.715.360.575.575.800.379 Early Termination of Clinical Trials
-	N05.715.360.575.575.800.569 Response Evaluation Criteria in Solid Tumors
New Heading	<b>N05.715.360.575.575.800.665 Sustained Virologic Response</b>
-	N05.715.360.575.575.800.760 Treatment Failure
-	N05.715.360.575.575.800.760.500 Failure to Rescue, Health Care
-	N05.715.360.575.625 Process Assessment (Health Care)
-	N05.715.360.600 Patient Satisfaction
-	N05.715.360.650 Program Evaluation
-	N05.715.360.650.150 Benchmarking
-	N05.715.360.700 Root Cause Analysis
-	N05.715.360.750 Statistics as Topic
-	N05.715.360.750.100 Actuarial Analysis
-	N05.715.360.750.125 Analysis of Variance
-	N05.715.360.750.125.500 Multivariate Analysis
-	N05.715.360.750.200 Cluster Analysis
-	N05.715.360.750.200.750 Small-Area Analysis
-	N05.715.360.750.200.775 Space-Time Clustering
-	N05.715.360.750.220 Confidence Intervals
-	N05.715.360.750.300 Data Interpretation, Statistical
-	N05.715.360.750.325 Discriminant Analysis
-	N05.715.360.750.350 Factor Analysis, Statistical
-	N05.715.360.750.500 Matched-Pair Analysis
-	N05.715.360.750.530 Models, Statistical
-	N05.715.360.750.530.450 Likelihood Functions
-	N05.715.360.750.530.460 Linear Models
-	N05.715.360.750.530.480 Logistic Models
-	N05.715.360.750.530.500 Models, Economic
-	N05.715.360.750.530.500.500 Models, Econometric
-	N05.715.360.750.530.530 Nomograms
-	N05.715.360.750.530.650 Proportional Hazards Models
-	N05.715.360.750.540 Monte Carlo Method
-	N05.715.360.750.625 Probability
-	N05.715.360.750.625.150 Bayes Theorem
-	N05.715.360.750.625.450 Likelihood Functions

## MeSH Tree Changes for 2017

Type	Tree - heading
-	N05.715.360.750.625.500 Markov Chains
-	N05.715.360.750.625.590 Odds Ratio
-	N05.715.360.750.625.620 Propensity Score
-	N05.715.360.750.625.650 Proportional Hazards Models
-	N05.715.360.750.625.700 Risk
-	N05.715.360.750.625.700.450 Logistic Models
-	N05.715.360.750.625.700.570 Protective Factors
-	N05.715.360.750.625.700.690 Risk Assessment
-	N05.715.360.750.625.700.690.400 Healthcare Failure Mode and Effect Analysis
-	N05.715.360.750.625.700.690.800 Risk Adjustment
-	N05.715.360.750.625.700.700 Risk Factors
-	N05.715.360.750.625.850 Uncertainty
-	N05.715.360.750.695 Regression Analysis
-	N05.715.360.750.695.440 Least-Squares Analysis
-	N05.715.360.750.695.460 Linear Models
-	N05.715.360.750.695.470 Logistic Models
-	N05.715.360.750.695.650 Proportional Hazards Models
-	N05.715.360.750.695.825 Spatial Regression
-	N05.715.360.750.725 Sensitivity and Specificity
-	N05.715.360.750.725.500 Limit of Detection
-	N05.715.360.750.725.750 Signal-To-Noise Ratio
-	N05.715.360.750.746 Spatial Analysis
-	N05.715.360.750.746.249 Geographic Mapping
-	N05.715.360.750.746.375 Spatial Regression
-	N05.715.360.750.746.500 Spatio-Temporal Analysis
-	N05.715.360.750.750 Statistical Distributions
-	N05.715.360.750.750.150 Binomial Distribution
-	N05.715.360.750.750.200 Chi-Square Distribution
-	N05.715.360.750.750.565 Normal Distribution
-	N05.715.360.750.750.620 Poisson Distribution
-	N05.715.360.750.760 Statistics, Nonparametric
-	N05.715.360.750.770 Stochastic Processes
-	N05.715.360.750.770.500 Markov Chains
-	N05.715.360.750.795 Survival Analysis
-	N05.715.360.750.795.300 Disease-Free Survival

## MeSH Tree Changes for 2017

Type	Tree - heading
-	N05.715.360.750.795.650                      Kaplan-Meier Estimate
-	N05.715.360.750.795.825                      Proportional Hazards Models
-	N05.715.360.775                                  Epidemiologic Study Characteristics as Topic
-	N05.715.360.775.088                              Clinical Studies as Topic
-	N05.715.360.775.088.500                          Clinical Trials as Topic
-	N05.715.360.775.088.500.200                      Clinical Trials, Phase I as Topic
-	N05.715.360.775.088.500.210                      Clinical Trials, Phase II as Topic
-	N05.715.360.775.088.500.220                      Clinical Trials, Phase III as Topic
-	N05.715.360.775.088.500.230                      Clinical Trials, Phase IV as Topic
-	N05.715.360.775.088.500.387                      Controlled Clinical Trials as Topic
-	N05.715.360.775.088.500.387.250                      Non-Randomized Controlled Trials as Topic
-	N05.715.360.775.088.500.387.500                      Randomized Controlled Trials as Topic
-	N05.715.360.775.088.500.387.500.500                      Intention to Treat Analysis
-	N05.715.360.775.088.500.387.500.750                      Pragmatic Clinical Trials as Topic
-	N05.715.360.775.088.750                              Observational Studies as Topic
-	N05.715.360.775.175                                  Epidemiologic Studies
-	N05.715.360.775.175.200                              Case-Control Studies
-	N05.715.360.775.175.200.710                          Retrospective Studies
-	N05.715.360.775.175.250                              Cohort Studies
-	N05.715.360.775.175.250.350                          Follow-Up Studies
-	N05.715.360.775.175.250.450                          Longitudinal Studies
-	N05.715.360.775.175.250.450.500                      National Longitudinal Study of Adolescent Health
-	N05.715.360.775.175.250.630                          Prospective Studies
-	N05.715.360.775.175.250.815                          Retrospective Studies
-	N05.715.360.775.175.262                              Controlled Before-After Studies
-	N05.715.360.775.175.275                              Cross-Sectional Studies
-	N05.715.360.775.175.537                              Interrupted Time Series Analysis
-	N05.715.360.775.175.800                              Seroepidemiologic Studies
-	N05.715.360.775.175.800.375                          HIV Seroprevalence
-	N05.715.360.775.225                                  Clinical Protocols
-	N05.715.360.775.225.500                              Antineoplastic Protocols
New Heading	<b>N05.715.360.775.225.750                              Standing Orders</b>
-	N05.715.360.775.350                                  Feasibility Studies
-	N05.715.360.775.508                                  Multicenter Studies as Topic



## MeSH Tree Changes for 2017

Type	Tree - heading	
-	N05.715.360.775.625	Pilot Projects
-	N05.715.360.775.700	Sampling Studies
-	N05.715.360.775.700.500	Lot Quality Assurance Sampling
-	N05.715.360.775.750	Twin Studies as Topic
-	N05.715.360.780	Epidemiologic Research Design
-	N05.715.360.780.150	Cross-Over Studies
-	N05.715.360.780.320	Double-Blind Method
-	N05.715.360.780.500	Matched-Pair Analysis
-	N05.715.360.780.515	Meta-Analysis as Topic
New Heading	<b>N05.715.360.780.515.500</b>	<b>Network Meta-Analysis</b>
-	N05.715.360.780.675	Random Allocation
-	N05.715.360.780.685	Reproducibility of Results
-	N05.715.360.780.685.250	Data Accuracy
-	N05.715.360.780.685.500	Dimensional Measurement Accuracy
-	N05.715.360.780.692	Sample Size
-	N05.715.360.780.700	Sensitivity and Specificity
-	N05.715.360.780.700.640	Predictive Value of Tests
-	N05.715.360.780.700.680	ROC Curve
-	N05.715.360.780.700.840	Signal-To-Noise Ratio
-	N05.715.360.780.730	Single-Blind Method
-	N05.715.360.825	Technology Assessment, Biomedical
-	N05.715.680	Peer Review, Health Care
-	N05.715.840	Standard of Care
-	N05.715.840.500	Patient Handoff
-	N06	Environment and Public Health
-	N06.230	Environment
-	N06.230.058	Altitude
-	N06.230.063	Carbon Footprint
-	N06.230.066	Caves
-	N06.230.069	Cities
-	N06.230.074	Confined Spaces
-	N06.230.080	Conservation of Natural Resources
-	N06.230.080.078	Conservation of Energy Resources
-	N06.230.080.200	Endangered Species
-	N06.230.080.600	Environmental Restoration and Remediation

## MeSH Tree Changes for 2017

Type	Tree - heading
-	N06.230.080.600.500 Biodegradation, Environmental
-	N06.230.100 Disasters
-	N06.230.100.035 Disaster Planning
-	N06.230.100.035.500 Strategic Stockpile
-	N06.230.100.083 Emergencies
-	N06.230.100.121 Emergency Shelter
-	N06.230.100.160 Mass Casualty Incidents
-	N06.230.100.300 Relief Work
-	N06.230.100.350 Rescue Work
-	N06.230.124 Ecosystem
-	N06.230.124.049 Biodiversity
-	N06.230.124.049.100 Biota
-	N06.230.124.049.100.500 Microbiota
-	N06.230.124.049.100.500.250 Gastrointestinal Microbiome
-	N06.230.124.049.100.500.500 Microbial Consortia
New Heading	<b>N06.230.124.049.100.500.750 Mycobiome</b>
-	N06.230.124.049.230 Ecotype
-	N06.230.124.049.250 Endangered Species
-	N06.230.124.049.400 Introduced Species
-	N06.230.124.100 Biomass
-	N06.230.124.130 Coral Reefs
-	N06.230.124.240 Ecological Systems, Closed
-	N06.230.124.250 Food Chain
-	N06.230.124.343 Forests
-	N06.230.124.343.500 Rainforest
-	N06.230.124.343.750 Taiga
-	N06.230.124.390 Grassland
-	N06.230.124.437 Rhizosphere
-	N06.230.124.531 Tundra
-	N06.230.124.531.500 Permafrost
-	N06.230.124.625 Wetlands
-	N06.230.132 Energy-Generating Resources
-	N06.230.132.258 Fossil Fuels
-	N06.230.132.258.108 Coal
-	N06.230.132.258.108.110 Coke

## MeSH Tree Changes for 2017

Type	Tree - heading
-	N06.230.132.258.369 Natural Gas
-	N06.230.132.258.630 Petroleum
-	N06.230.132.258.630.500 Fuel Oils
-	N06.230.132.258.630.540 Gasoline
-	N06.230.132.258.630.600 Kerosene
-	N06.230.132.580 Nuclear Energy
-	N06.230.132.580.500 Nuclear Fission
-	N06.230.132.580.520 Nuclear Fusion
-	N06.230.132.644 Renewable Energy
-	N06.230.132.644.124 Biofuels
-	N06.230.132.644.186 Geothermal Energy
-	N06.230.132.644.500 Solar Energy
-	N06.230.132.644.750 Water Movements
-	N06.230.132.644.750.500 Tidal Waves
-	N06.230.132.644.875 Wind
-	N06.230.150 Environment, Controlled
-	N06.230.150.050 Air Conditioning
-	N06.230.150.150 Diving
-	N06.230.150.225 Ecological Systems, Closed
-	N06.230.150.300 Heating
-	N06.230.150.360 Housing
-	N06.230.150.360.250 Housing, Animal
-	N06.230.150.360.250.200 Hospitals, Animal
-	N06.230.150.360.260 Housing for the Elderly
-	N06.230.150.360.650 Public Housing
-	N06.230.150.372 Humidity
-	N06.230.150.391 Life Support Systems
-	N06.230.150.410 Lighting
-	N06.230.150.440 Space Simulation
-	N06.230.150.440.950 Weightlessness Simulation
-	N06.230.150.440.950.400 Hindlimb Suspension
-	N06.230.150.450 Temperature
-	N06.230.150.520 Ventilation
-	N06.230.200 Environment Design
-	N06.230.204 Environmental Policy
-	N06.230.208 Explosions

## MeSH Tree Changes for 2017

Type	Tree - heading
-	N06.230.216 Fires
-	N06.230.216.500 Fire Extinguishing Systems
-	N06.230.216.750 Spontaneous Combustion
-	N06.230.232 Fresh Water
-	N06.230.232.500 Lakes
-	N06.230.232.625 Ponds
-	N06.230.232.650 Rivers
-	N06.230.265 Greenhouse Effect
-	N06.230.291 Ice
-	N06.230.291.500 Ice Cover
-	N06.230.295 Islands
-	N06.230.300 Meteorological Concepts
-	N06.230.300.100 Atmosphere
-	N06.230.300.100.150 Air
-	N06.230.300.100.150.100 Air Ionization
-	N06.230.300.100.150.185 Air Movements
-	N06.230.300.100.150.185.200 Wind
-	N06.230.300.100.150.592 Compressed Air
-	N06.230.300.100.185 Atmospheric Pressure
-	N06.230.300.100.185.100 Air Pressure
-	N06.230.300.100.185.902 Vacuum
-	N06.230.300.100.250 Climate
-	N06.230.300.100.250.275 Cold Climate
-	N06.230.300.100.250.325 Desert Climate
-	N06.230.300.100.250.450 Microclimate
-	N06.230.300.100.250.525 Seasons
-	N06.230.300.100.250.600 Tropical Climate
-	N06.230.300.100.300 Cosmic Radiation
-	N06.230.300.100.700 Stratospheric Ozone
-	N06.230.300.100.700.500 Ozone Depletion
-	N06.230.300.100.725 Weather
-	N06.230.300.100.725.154 Cold Temperature
-	N06.230.300.100.725.154.500 Extreme Cold
-	N06.230.300.100.725.232 Hot Temperature
-	N06.230.300.100.725.232.500 Extreme Heat
-	N06.230.300.100.725.310 Humidity

## MeSH Tree Changes for 2017

Type	Tree - heading
-	N06.230.300.100.725.375 Lightning
-	N06.230.300.100.725.450 Rain
-	N06.230.300.100.725.450.050 Acid Rain
-	N06.230.300.100.725.480 Snow
-	N06.230.300.100.725.525 Sunlight
-	N06.230.300.100.725.525.400 Infrared Rays
-	N06.230.300.100.725.525.600 Ultraviolet Rays
-	N06.230.300.100.725.710 Temperature
-	N06.230.300.100.725.710.300 Cold Temperature
-	N06.230.300.100.725.710.300.500 Extreme Cold
-	N06.230.300.100.725.710.380 Hot Temperature
-	N06.230.300.100.725.710.380.500 Extreme Heat
-	N06.230.300.100.725.780 Wind
-	N06.230.350 Natural Resources
-	N06.230.350.500 Water Resources
-	N06.230.400 Noise
-	N06.230.400.500 Noise, Occupational
-	N06.230.400.550 Noise, Transportation
-	N06.230.480 Odorants
-	N06.230.480 Odors
-	N06.230.520 Rain
-	N06.230.520.050 Acid Rain
-	N06.230.600 Soil
-	N06.230.600.500 Permafrost
-	N06.230.650 Steam
-	N06.230.750 Structure Collapse
-	N06.230.850 Water Movements
-	N06.850 Public Health
-	N06.850.135 Accidents
-	N06.850.135.060 Accident Prevention
-	N06.850.135.060.075 Safety
-	N06.850.135.060.075.199 Chemical Safety
-	N06.850.135.060.075.299 Hazard Analysis and Critical Control Points
-	N06.850.135.060.075.349 Patient Harm
-	N06.850.135.060.075.399 Patient Safety
-	N06.850.135.060.075.800 Safety Management



## MeSH Tree Changes for 2017

Type	Tree - heading	
-	N06.850.310.350.100.500	Insect Vectors
New Heading	<b>N06.850.310.350.100.500.500</b>	<b>Mosquito Vectors</b>
-	N06.850.310.405	Infectious Disease Incubation Period
-	N06.850.310.410	Infectious Disease Transmission, Patient-to-Professional
-	N06.850.310.415	Infectious Disease Transmission, Professional-to-Patient
-	N06.850.310.425	Infectious Disease Transmission, Vertical
-	N06.850.360	Drug Contamination
-	N06.850.376	Emergencies
-	N06.850.392	Endemic Diseases
-	N06.850.420	Environmental Medicine
-	N06.850.425	Environmental Microbiology
-	N06.850.425.110	Air Microbiology
-	N06.850.425.200	Food Microbiology
-	N06.850.425.300	Soil Microbiology
-	N06.850.425.450	Water Microbiology
-	N06.850.460	Environmental Pollution
-	N06.850.460.100	Air Pollution
-	N06.850.460.100.080	Air Pollution, Indoor
-	N06.850.460.100.110	Air Pollution, Radioactive
-	N06.850.460.100.555	Tobacco Smoke Pollution
-	N06.850.460.150	Biofouling
-	N06.850.460.200	Body Burden
-	N06.850.460.200.250	Drug Residues
-	N06.850.460.200.700	Pesticide Residues
-	N06.850.460.350	Environmental Exposure
-	N06.850.460.350.080	Environmental Monitoring
-	N06.850.460.350.080.500	Biological Oxygen Demand Analysis
-	N06.850.460.350.080.750	Water Quality
-	N06.850.460.350.112	Inhalation Exposure
-	N06.850.460.350.145	Maternal Exposure
-	N06.850.460.350.210	Maximum Allowable Concentration
-	N06.850.460.350.600	Occupational Exposure
-	N06.850.460.350.600.615	Maximum Allowable Concentration
-	N06.850.460.350.600.807	Threshold Limit Values
-	N06.850.460.350.600.903	War Exposure

## MeSH Tree Changes for 2017

Type	Tree - heading
-	N06.850.460.350.600.903.500 War-Related Injuries
-	N06.850.460.350.700 Paternal Exposure
-	N06.850.460.350.850 Radiation Exposure
-	N06.850.460.350.850.500 Radiation Injuries
-	N06.850.460.350.850.500.031 Abnormalities, Radiation-Induced
-	N06.850.460.350.850.500.158 Acute Radiation Syndrome
-	N06.850.460.350.850.500.285 Radiation Injuries, Experimental
-	N06.850.460.375 Environmental Restoration and Remediation
-	N06.850.460.375.500 Biodegradation, Environmental
-	N06.850.460.400 Food Contamination
-	N06.850.460.400.250 Food Contamination, Radioactive
-	N06.850.460.400.300 Food Microbiology
-	N06.850.460.400.650 Food Parasitology
-	N06.850.460.610 Noise
-	N06.850.460.610.526 Noise, Occupational
-	N06.850.460.610.680 Noise, Transportation
-	N06.850.460.660 Petroleum Pollution
-	N06.850.460.710 Waste Products
-	N06.850.460.710.189 Electronic Waste
-	N06.850.460.710.380 Hazardous Waste
-	N06.850.460.710.380.638 Radioactive Waste
-	N06.850.460.710.420 Industrial Waste
-	N06.850.460.710.460 Medical Waste
-	N06.850.460.710.460.150 Dental Waste
-	N06.850.460.710.460.300 Medical Waste Disposal
-	N06.850.460.710.730 Solid Waste
-	N06.850.460.710.865 Waste Water
-	N06.850.460.790 Water Pollution
-	N06.850.460.790.410 Water Pollution, Chemical
-	N06.850.460.790.410.050 Acid Rain
-	N06.850.460.790.460 Water Pollution, Radioactive
-	N06.850.460.790.730 Water Quality
-	N06.850.490 Epidemiologic Factors
-	N06.850.490.250 Age Factors
-	N06.850.490.250.100 Age of Onset
-	N06.850.490.250.550 Maternal Age



## MeSH Tree Changes for 2017

Type	Tree - heading
-	N06.850.490.500 Bias (Epidemiology)
-	N06.850.490.500.250 Observer Variation
-	N06.850.490.500.500 Selection Bias
-	N06.850.490.625 Causality
-	N06.850.490.625.500 Precipitating Factors
-	N06.850.490.625.625 Protective Factors
-	N06.850.490.625.750 Risk Factors
-	N06.850.490.687 Comorbidity
-	N06.850.490.718 Confounding Factors (Epidemiology)
-	N06.850.490.734 Effect Modifier, Epidemiologic
-	N06.850.490.734.500 Cohort Effect
-	N06.850.490.734.750 Healthy Worker Effect
-	N06.850.490.734.875 Placebo Effect
-	N06.850.490.734.875.500 Nocebo Effect
-	N06.850.490.812 Reproductive History
-	N06.850.490.812.250 Gravidity
-	N06.850.490.812.600 Parity
-	N06.850.490.875 Sex Factors
-	N06.850.505 Epidemiologic Measurements
-	N06.850.505.200 Biometry
-	N06.850.505.200.100 Anthropometry
-	N06.850.505.200.100.175 Body Mass Index
-	N06.850.505.200.100.300 Cephalometry
-	N06.850.505.200.100.400 Crown-Rump Length
New Heading	<b>N06.850.505.200.100.700 Kinanthropometry</b>
-	N06.850.505.400 Demography
-	N06.850.505.400.050 Age Distribution
-	N06.850.505.400.225 Censuses
-	N06.850.505.400.400 Family Characteristics
-	N06.850.505.400.400.160 Birth Order
-	N06.850.505.400.425 Health Status
-	N06.850.505.400.425.350 Geriatric Assessment
-	N06.850.505.400.425.675 Health Status Disparities
New Tree	<b>N06.850.505.400.425.837 Quality of Life</b>

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	N06.850.505.400.600	Population Density
-	N06.850.505.400.700	Population Dynamics
-	N06.850.505.400.700.400	Health Transition
-	N06.850.505.400.700.525	Human Migration
-	N06.850.505.400.700.525.500	Emigration and Immigration
-	N06.850.505.400.700.650	Population Control
-	N06.850.505.400.700.660	Population Growth
-	N06.850.505.400.800	Residence Characteristics
-	N06.850.505.400.800.200	Catchment Area (Health)
-	N06.850.505.400.800.400	Housing
-	N06.850.505.400.800.550	Independent Living
-	N06.850.505.400.850	Sex Distribution
-	N06.850.505.400.850.815	Sex Ratio
-	N06.850.505.400.975	Vital Statistics
-	N06.850.505.400.975.450	Life Expectancy
-	N06.850.505.400.975.475	Life Tables
-	N06.850.505.400.975.525	Morbidity
-	N06.850.505.400.975.525.080	Basic Reproduction Number
-	N06.850.505.400.975.525.375	Incidence
-	N06.850.505.400.975.525.750	Prevalence
-	N06.850.505.400.975.550	Mortality
-	N06.850.505.400.975.550.250	Cause of Death
-	N06.850.505.400.975.550.287	Child Mortality
-	N06.850.505.400.975.550.325	Fatal Outcome
-	N06.850.505.400.975.550.362	Fetal Mortality
-	N06.850.505.400.975.550.400	Hospital Mortality
-	N06.850.505.400.975.550.475	Infant Mortality
New Tree	<a href="#">N06.850.505.400.975.550.475.500</a>	<a href="#">Perinatal Mortality</a>
-	N06.850.505.400.975.550.500	Maternal Mortality
-	N06.850.505.400.975.550.550	Mortality, Premature
Old Tree	<del>N06.850.505.400.975.550.700</del>	<del>Perinatal Mortality</del>
-	N06.850.505.400.975.550.900	Survival Rate
-	N06.850.505.400.975.775	Pregnancy Rate
-	N06.850.505.400.975.775.500	Birth Rate
-	N06.850.505.557	Nutrition Assessment

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	N06.850.505.616	Nutrition Surveys
-	N06.850.505.616.300	Diet Surveys
-	N06.850.505.636	Pharmacovigilance
-	N06.850.505.715	Risk Assessment
-	N06.850.505.715.500	Healthcare Failure Mode and Effect Analysis
-	N06.850.520	Epidemiologic Methods
-	N06.850.520.270	Contact Tracing
-	N06.850.520.308	Data Collection
-	N06.850.520.308.056	Datasets as Topic
-	N06.850.520.308.112	Focus Groups
-	N06.850.520.308.225	Geriatric Assessment
-	N06.850.520.308.420	Interviews as Topic
-	N06.850.520.308.461	Lot Quality Assurance Sampling
-	N06.850.520.308.502	Narration
-	N06.850.520.308.585	Nutrition Assessment
-	N06.850.520.308.940	Records as Topic
-	N06.850.520.308.940.250	Birth Certificates
-	N06.850.520.308.940.350	Death Certificates
-	N06.850.520.308.940.375	Dental Records
-	N06.850.520.308.940.425	Hospital Records
-	N06.850.520.308.940.968	Medical Records
-	N06.850.520.308.940.968.249	Clinical Coding
-	N06.850.520.308.940.968.500	Medical Record Linkage
-	N06.850.520.308.940.968.550	Medical Records, Problem-Oriented
-	N06.850.520.308.940.968.625	Medical Records Systems, Computerized
-	N06.850.520.308.940.968.625.500	Health Smart Cards
-	N06.850.520.308.940.968.750	Patient Discharge Summaries
-	N06.850.520.308.940.968.875	Trauma Severity Indices
-	N06.850.520.308.940.984	Nursing Records
-	N06.850.520.308.970	Registries
-	N06.850.520.308.970.725	SEER Program
-	N06.850.520.308.980	Surveys and Questionnaires
-	N06.850.520.308.980.313	Contraceptive Prevalence Surveys
-	N06.850.520.308.980.344	Health Care Surveys
New Heading	<b>N06.850.520.308.980.344.500</b>	<b>Patient Reported Outcome Measures</b>

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	N06.850.520.308.980.438	Health Surveys
-	N06.850.520.308.980.438.149	Behavioral Risk Factor Surveillance System
-	N06.850.520.308.980.438.300	Dental Health Surveys
-	N06.850.520.308.980.438.300.300	Dental Plaque Index
-	N06.850.520.308.980.438.300.350	DMF Index
-	N06.850.520.308.980.438.300.512	Index of Orthodontic Treatment Need
-	N06.850.520.308.980.438.300.675	Oral Hygiene Index
-	N06.850.520.308.980.438.300.725	Periodontal Index
-	N06.850.520.308.980.438.475	Health Status Indicators
New Heading	<b>N06.850.520.308.980.438.475.091</b>	<b>Global Burden of Disease</b>
Old Tree	<b>N06.850.520.308.980.438.475.181</b>	<b>Organ Dysfunction Scores</b>
-	N06.850.520.308.980.438.475.364	Patient Acuity
-	N06.850.520.308.980.438.475.364.500	Severity of Illness Index
New Tree	<b>N06.850.520.308.980.438.475.364.500.250</b>	<b>APACHE</b>
-	N06.850.520.308.980.438.475.364.500.500	Karnofsky Performance Status
New Tree	<b>N06.850.520.308.980.438.475.364.500.625</b>	<b>Organ Dysfunction Scores</b>
New Heading	<b>N06.850.520.308.980.438.475.364.500.750 Score</b>	<b>Simplified Acute Physiology</b>
-	N06.850.520.308.980.438.475.730	Sickness Impact Profile
-	N06.850.520.308.980.438.580	Mass Screening
-	N06.850.520.308.980.438.580.174	Anonymous Testing
-	N06.850.520.308.980.438.580.510	Mass Chest X-Ray
-	N06.850.520.308.980.438.580.560	Multiphasic Screening
-	N06.850.520.308.980.438.580.580	Neonatal Screening
-	N06.850.520.308.980.438.580.925	Vision Screening
-	N06.850.520.308.980.438.700	Population Surveillance
-	N06.850.520.308.980.438.700.324	Public Health Surveillance
-	N06.850.520.308.980.438.700.650	Sentinel Surveillance
-	N06.850.520.308.980.469	Nutrition Surveys
-	N06.850.520.308.980.469.350	Diet Surveys
-	N06.850.520.308.980.500	Self Report
-	N06.850.520.308.985	Vital Statistics
-	N06.850.520.308.985.450	Life Expectancy
-	N06.850.520.308.985.475	Life Tables

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	N06.850.520.308.985.525	Morbidity
-	N06.850.520.308.985.525.080	Basic Reproduction Number
-	N06.850.520.308.985.525.375	Incidence
-	N06.850.520.308.985.525.750	Prevalence
-	N06.850.520.308.985.550	Mortality
-	N06.850.520.308.985.550.250	Cause of Death
-	N06.850.520.308.985.550.287	Child Mortality
-	N06.850.520.308.985.550.325	Fatal Outcome
-	N06.850.520.308.985.550.362	Fetal Mortality
-	N06.850.520.308.985.550.400	Hospital Mortality
-	N06.850.520.308.985.550.475	Infant Mortality
New Tree	N06.850.520.308.985.550.475.500	Perinatal Mortality
-	N06.850.520.308.985.550.500	Maternal Mortality
-	N06.850.520.308.985.550.550	Mortality, Premature
Old Tree	N06.850.520.308.985.550.700	Perinatal Mortality
-	N06.850.520.308.985.550.900	Survival Rate
-	N06.850.520.308.985.775	Pregnancy Rate
-	N06.850.520.308.985.775.500	Birth Rate
-	N06.850.520.373	Disease Notification
-	N06.850.520.445	Epidemiologic Research Design
-	N06.850.520.445.150	Cross-Over Studies
-	N06.850.520.445.300	Double-Blind Method
-	N06.850.520.445.392	Genome-Wide Association Study
-	N06.850.520.445.485	Matched-Pair Analysis
-	N06.850.520.445.500	Meta-Analysis as Topic
New Heading	N06.850.520.445.500.500	Network Meta-Analysis
-	N06.850.520.445.700	Random Allocation
-	N06.850.520.445.725	Reproducibility of Results
-	N06.850.520.445.725.500	Dimensional Measurement Accuracy
-	N06.850.520.445.762	Sample Size
-	N06.850.520.445.800	Sensitivity and Specificity
-	N06.850.520.445.800.650	Predictive Value of Tests
-	N06.850.520.445.800.750	ROC Curve
-	N06.850.520.445.800.875	Signal-To-Noise Ratio

## MeSH Tree Changes for 2017

Type	Tree - heading
-	N06.850.520.445.850                      Single-Blind Method
-	N06.850.520.450                              Epidemiologic Study Characteristics as Topic
-	N06.850.520.450.250                        Clinical Studies as Topic
-	N06.850.520.450.250.250                    Clinical Trials as Topic
-	N06.850.520.450.250.250.200              Clinical Trials, Phase I as Topic
-	N06.850.520.450.250.250.210              Clinical Trials, Phase II as Topic
-	N06.850.520.450.250.250.220              Clinical Trials, Phase III as Topic
-	N06.850.520.450.250.250.230              Clinical Trials, Phase IV as Topic
-	N06.850.520.450.250.250.365              Controlled Clinical Trials as Topic
-	N06.850.520.450.250.250.365.250            Non-Randomized Controlled Trials as Topic
-	N06.850.520.450.250.250.365.500            Randomized Controlled Trials as Topic
-	N06.850.520.450.250.250.365.500.500            Intention to Treat Analysis
-	N06.850.520.450.250.250.365.500.750            Pragmatic Clinical Trials as Topic
-	N06.850.520.450.250.500                    Observational Studies as Topic
-	N06.850.520.450.500                        Epidemiologic Studies
-	N06.850.520.450.500.500                    Case-Control Studies
-	N06.850.520.450.500.500.500                  Retrospective Studies
-	N06.850.520.450.500.750                    Cohort Studies
-	N06.850.520.450.500.750.350                  Follow-Up Studies
-	N06.850.520.450.500.750.500                  Longitudinal Studies
-	N06.850.520.450.500.750.500.500            National Longitudinal Study of Adolescent Health
-	N06.850.520.450.500.750.650                  Prospective Studies
-	N06.850.520.450.500.750.825                  Retrospective Studies
-	N06.850.520.450.500.812                    Controlled Before-After Studies
-	N06.850.520.450.500.875                    Cross-Sectional Studies
-	N06.850.520.450.500.912                    Interrupted Time Series Analysis
-	N06.850.520.450.500.950                    Seroepidemiologic Studies
-	N06.850.520.450.500.950.375                  HIV Seroprevalence
-	N06.850.520.450.550                        Feasibility Studies
-	N06.850.520.450.643                        Multicenter Studies as Topic
-	N06.850.520.450.720                        Pilot Projects
-	N06.850.520.450.875                        Sampling Studies
-	N06.850.520.450.900                        Twin Studies as Topic
-	N06.850.520.460                              Epidemiological Monitoring
-	N06.850.520.470                              Molecular Epidemiology

## MeSH Tree Changes for 2017

Type	Tree - heading
-	N06.850.520.470.500                      Genome-Wide Association Study
-	N06.850.520.699                              Sentinel Surveillance
-	N06.850.520.830                              Statistics as Topic
-	N06.850.520.830.100                          Actuarial Analysis
-	N06.850.520.830.150                          Analysis of Variance
-	N06.850.520.830.150.500                      Multivariate Analysis
-	N06.850.520.830.200                          Area Under Curve
-	N06.850.520.830.250                          Cluster Analysis
-	N06.850.520.830.250.675                      Small-Area Analysis
-	N06.850.520.830.250.700                      Space-Time Clustering
-	N06.850.520.830.275                          Confidence Intervals
-	N06.850.520.830.300                          Data Interpretation, Statistical
-	N06.850.520.830.350                          Discriminant Analysis
-	N06.850.520.830.400                          Factor Analysis, Statistical
-	N06.850.520.830.475                          Matched-Pair Analysis
-	N06.850.520.830.500                          Models, Statistical
-	N06.850.520.830.500.475                      Likelihood Functions
-	N06.850.520.830.500.500                      Linear Models
-	N06.850.520.830.500.525                      Logistic Models
-	N06.850.520.830.500.600                      Models, Economic
-	N06.850.520.830.500.600.500                      Models, Econometric
-	N06.850.520.830.500.625                      Nomograms
-	N06.850.520.830.500.700                      Proportional Hazards Models
-	N06.850.520.830.525                          Monte Carlo Method
-	N06.850.520.830.562                          Multilevel Analysis
-	N06.850.520.830.600                          Probability
-	N06.850.520.830.600.200                      Bayes Theorem
-	N06.850.520.830.600.400                      Likelihood Functions
-	N06.850.520.830.600.500                      Markov Chains
-	N06.850.520.830.600.600                      Odds Ratio
-	N06.850.520.830.600.650                      Propensity Score
-	N06.850.520.830.600.700                      Proportional Hazards Models
-	N06.850.520.830.600.800                      Risk
-	N06.850.520.830.600.800.450                      Logistic Models
-	N06.850.520.830.600.800.582                      Protective Factors
-	N06.850.520.830.600.800.715                      Risk Assessment

## MeSH Tree Changes for 2017

Type	Tree - heading
-	N06.850.520.830.600.800.715.500 Healthcare Failure Mode and Effect Analysis
-	N06.850.520.830.600.800.725 Risk Factors
-	N06.850.520.830.600.900 Uncertainty
-	N06.850.520.830.750 Regression Analysis
-	N06.850.520.830.750.400 Least-Squares Analysis
-	N06.850.520.830.750.425 Linear Models
-	N06.850.520.830.750.450 Logistic Models
-	N06.850.520.830.750.725 Proportional Hazards Models
-	N06.850.520.830.750.862 Spatial Regression
-	N06.850.520.830.872 Sensitivity and Specificity
-	N06.850.520.830.872.500 Limit of Detection
-	N06.850.520.830.872.750 Signal-To-Noise Ratio
-	N06.850.520.830.933 Spatial Analysis
-	N06.850.520.830.933.249 Geographic Mapping
-	N06.850.520.830.933.375 Spatial Regression
-	N06.850.520.830.933.500 Spatio-Temporal Analysis
-	N06.850.520.830.994 Statistical Distributions
-	N06.850.520.830.994.250 Binomial Distribution
-	N06.850.520.830.994.300 Chi-Square Distribution
-	N06.850.520.830.994.500 Normal Distribution
-	N06.850.520.830.994.750 Poisson Distribution
-	N06.850.520.830.995 Statistics, Nonparametric
-	N06.850.520.830.996 Stochastic Processes
-	N06.850.520.830.996.500 Markov Chains
-	N06.850.520.830.998 Survival Analysis
-	N06.850.520.830.998.300 Disease-Free Survival
-	N06.850.520.830.998.650 Kaplan-Meier Estimate
-	N06.850.540 Equipment Contamination
-	N06.850.540.500 Biofouling
-	N06.850.585 Equipment Reuse
-	N06.850.601 Food Quality
-	N06.850.601.500 Food Safety
-	N06.850.601.500.249 Food Contamination
-	N06.850.601.500.249.250 Food Contamination, Radioactive
-	N06.850.601.500.249.300 Food Microbiology



## MeSH Tree Changes for 2017

Type	Tree - heading
-	N06.850.601.500.249.650 Food Parasitology
-	N06.850.601.500.500 Food Inspection
-	N06.850.601.500.750 Hazard Analysis and Critical Control Points
-	N06.850.601.750 Nutritive Value
-	N06.850.601.750.500 Glycemic Index
-	N06.850.601.750.750 Glycemic Load
-	N06.850.650 Health Transition
-	N06.850.670 Hygiene
New Tree	<a href="#">N06.850.670.150</a> Hand Hygiene
New Tree	<a href="#">N06.850.670.150.500</a> Hand Disinfection
-	N06.850.670.300 Military Hygiene
-	N06.850.725 Patient Harm
-	N06.850.780 Public Health Practice
-	N06.850.780.200 Communicable Disease Control
-	N06.850.780.200.112 Animal Culling
-	N06.850.780.200.225 Contact Tracing
-	N06.850.780.200.262 Disease Notification
-	N06.850.780.200.300 Fumigation
Old Tree	<del>N06.850.780.200.412</del> Hand Hygiene
Old Tree	<del>N06.850.780.200.412.500</del> Hand Disinfection
-	N06.850.780.200.425 Immunization
-	N06.850.780.200.425.900 Vaccination
-	N06.850.780.200.425.900.500 Mass Vaccination
-	N06.850.780.200.450 Infection Control
-	N06.850.780.200.450.150 Antisepsis
-	N06.850.780.200.450.150.160 Asepsis
-	N06.850.780.200.450.325 Blood Safety
-	N06.850.780.200.450.500 Infection Control, Dental
-	N06.850.780.200.450.650 Patient Isolation
-	N06.850.780.200.450.700 Quarantine
-	N06.850.780.200.450.850 Sterilization
-	N06.850.780.200.450.850.375 Disinfection
-	N06.850.780.200.550 Mandatory Testing
-	N06.850.780.200.650 Pest Control

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	N06.850.780.200.650.425	Insect Control
-	N06.850.780.200.650.425.500	Mosquito Control
-	N06.850.780.200.650.650	Pest Control, Biological
-	N06.850.780.200.650.700	Rodent Control
-	N06.850.780.200.650.850	Tick Control
-	N06.850.780.200.650.925	Weed Control
-	N06.850.780.200.800	Sanitation
-	N06.850.780.200.800.325	Food Inspection
-	N06.850.780.200.800.800	Sanitary Engineering
-	N06.850.780.200.800.800.350	Drainage, Sanitary
-	N06.850.780.200.800.800.525	Recycling
-	N06.850.780.200.800.800.700	Refuse Disposal
New Tree	N06.850.780.200.800.800.795	Toilet Facilities
-	N06.850.780.200.800.800.890	Waste Disposal, Fluid
-	N06.850.780.200.800.800.900	Waste Management
-	N06.850.780.200.800.800.900.449	Waste Disposal Facilities
-	N06.850.780.200.800.800.900.449.500	Hazardous Waste Sites
-	N06.850.780.200.800.800.900.900	Water Purification
-	N06.850.780.200.925	Universal Precautions
-	N06.850.780.325	Decontamination
-	N06.850.780.375	Environmental Monitoring
-	N06.850.780.375.349	Biological Oxygen Demand Analysis
-	N06.850.780.375.700	Radiation Monitoring
-	N06.850.780.500	Mass Screening
-	N06.850.780.500.162	Anonymous Testing
-	N06.850.780.500.412	Mandatory Testing
-	N06.850.780.500.500	Mass Chest X-Ray
-	N06.850.780.500.560	Multiphasic Screening
-	N06.850.780.500.580	Neonatal Screening
-	N06.850.780.500.765	Substance Abuse Detection
-	N06.850.780.500.950	Vision Screening
-	N06.850.780.675	Population Surveillance
-	N06.850.780.675.324	Biosurveillance
-	N06.850.780.675.487	Public Health Surveillance
-	N06.850.780.675.650	Sentinel Surveillance

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	N06.850.780.680	Primary Prevention
-	N06.850.780.680.310	Immunization
-	N06.850.780.680.310.890	Vaccination
-	N06.850.780.680.310.890.500	Mass Vaccination
-	N06.850.780.680.655	Pre-Exposure Prophylaxis
-	N06.850.780.750	Secondary Prevention
-	N06.850.780.800	Tertiary Prevention
-	N06.850.810	Radiologic Health
-	N06.850.810.080	Air Pollution, Radioactive
-	N06.850.810.125	Food Contamination, Radioactive
-	N06.850.810.250	Radiation Dosage
-	N06.850.810.250.180	Dose-Response Relationship, Radiation
-	N06.850.810.250.275	Relative Biological Effectiveness
-	N06.850.810.300	Radiation Effects
-	N06.850.810.300.360	Radiation Injuries
-	N06.850.810.300.360.031	Abnormalities, Radiation-Induced
-	N06.850.810.300.360.158	Acute Radiation Syndrome
-	N06.850.810.300.360.285	Radiation Injuries, Experimental
-	N06.850.810.335	Radiation Genetics
-	N06.850.810.370	Radiation Monitoring
-	N06.850.810.370.310	Film Dosimetry
-	N06.850.810.370.365	Optically Stimulated Luminescence Dosimetry
-	N06.850.810.370.420	Thermoluminescent Dosimetry
-	N06.850.810.425	Radiation Protection
-	N06.850.810.460	Radioactive Fallout
-	N06.850.810.485	Radioactive Waste
-	N06.850.810.530	Water Pollution, Radioactive
-	N06.850.860	Sanitation
Old Tree	<b>N06.850.860.410</b>	<b>Public Facilities</b>
Old Tree	<b>N06.850.860.410.054</b>	<b>Airports</b>
Old Tree	<b>N06.850.860.410.110</b>	<b>Bathing Beaches</b>
Old Tree	<b>N06.850.860.410.250</b>	<b>Health Resorts</b>
Old Tree	<b>N06.850.860.410.375</b>	<b>Parks, Recreational</b>
Old Tree	<b>N06.850.860.410.500</b>	<b>Swimming Pools</b>
Old Tree	<b>N06.850.860.410.560</b>	<b>Toilet Facilities</b>
-	N06.850.860.510	Sanitary Engineering

## MeSH Tree Changes for 2017

Type	Tree - heading
-	N06.850.860.510.244                      Recycling
-	N06.850.860.510.490                      Toilet Facilities
-	N06.850.860.510.900                      Waste Management
-	N06.850.860.510.900.600                      Refuse Disposal
-	N06.850.860.510.900.600.400                      Garbage
-	N06.850.860.510.900.600.500                      Incineration
-	N06.850.860.510.900.600.600                      Medical Waste Disposal
-	N06.850.860.510.900.600.900                      Waste Disposal, Fluid
-	N06.850.860.510.900.700                      Waste Disposal Facilities
-	N06.850.860.510.900.700.500                      Hazardous Waste Sites
-	N06.850.860.510.900.900                      Water Purification
-	N06.850.860.510.910                      Water Softening
-	N06.890                      Public Health Dentistry
-	N06.890.160                      Dental Health Surveys
-	N06.890.160.090                      Dental Plaque Index
-	N06.890.160.100                      DMF Index
-	N06.890.160.134                      Index of Orthodontic Treatment Need
-	N06.890.160.168                      Oral Hygiene Index
-	N06.890.160.215                      Periodontal Index
-	N06.890.235                      Fluoridation
-	N06.890.410                      Health Education, Dental
-	V01                      Publication Components
-	V01.070                      Abstracts
-	V01.100                      Advertisements
-	V01.110                      Animation
-	V01.140                      Architectural Drawings
-	V01.165                      Bibliography
-	V01.165.150                      Biobibliography
-	V01.175                      Biography
-	V01.175.125                      Blogs
-	V01.175.250                      Interview
-	V01.175.500                      Personal Narratives
-	V01.175.500.100                      Autobiography
-	V01.185                      Book Illustrations
-	V01.185.500                      Caricatures
-	V01.185.625                      Drawings

## MeSH Tree Changes for 2017

Type	Tree - heading
-	V01.185.687 Maps
-	V01.185.843 Portraits
-	V01.195 Bookplates
-	V01.200 Charts
-	V01.205 Comment
-	V01.250 Editorial
-	V01.255 Electronic Supplementary Materials
-	V01.255.500 Video-Audio Media
-	V01.255.500.500 Instructional Films and Videos
-	V01.255.500.625 Interactive Tutorial
-	V01.255.500.750 Webcasts
-	V01.260 English Abstract
-	V01.550 Letter
-	V01.630 News
-	V01.665 Patient Education Handout
-	V01.700 Published Erratum
-	V01.775 Retraction of Publication
-	V02 Publication Formats
-	V02.025 Abbreviations
-	V02.035 Abstracts
-	V02.050 Academic Dissertations
-	V02.060 Account Books
-	V02.070 Addresses
-	V02.070.500 Lectures
-	V02.070.750 Sermons
-	V02.110 Almanacs
-	V02.115 Anecdotes
-	V02.120 Animation
-	V02.130 Annual Reports
-	V02.135 Aphorisms and Proverbs
-	V02.140 Architectural Drawings
-	V02.150 Atlases
-	V02.165 Bibliography
-	V02.165.500 Biobibliography
-	V02.170 Biography
-	V02.170.250 Blogs

## MeSH Tree Changes for 2017

Type	Tree - heading
-	V02.170.500 Interview
-	V02.170.750 Personal Narratives
-	V02.170.750.100 Autobiography
-	V02.180 Book Reviews
-	V02.200 Broadsides
New Heading	<b>V02.220 Calendars</b>
-	V02.240 Catalogs
-	V02.240.500 Price Lists
-	V02.245 Charts
-	V02.250 Chronology
-	V02.255 Classical Article
-	V02.260 Collected Works
-	V02.260.260 Collected Correspondence
-	V02.260.400 Festschrift
-	V02.265 Collections
-	V02.267 Comment
-	V02.270 Congresses
New Tree	<a href="#">V02.270.250 Consensus Development Conference</a>
New Tree	<a href="#">V02.270.250.500 Consensus Development Conference, NIH</a>
-	V02.270.500 Overall
-	V02.272 Cookbooks
-	V02.275 Corrected and Republished Article
-	V02.300 Database
-	V02.302 Dataset
-	V02.305 Diaries
-	V02.310 Dictionary
-	V02.310.500 Phrases
-	V02.310.750 Terminology
-	V02.312 Directory
-	V02.315 Documentaries and Factual Films
-	V02.320 Duplicate Publication
-	V02.335 Editorial
-	V02.350 Encyclopedias
-	V02.355 Ephemera

## MeSH Tree Changes for 2017

Type	Tree - heading
-	V02.355.249 Advertisements
-	V02.355.374 Letter
-	V02.355.500 Posters
-	V02.355.750 Programs
-	V02.355.775 Prospectuses
-	V02.365 Essays
-	V02.375 Eulogies
-	V02.375.400 Funeral Sermons
-	V02.395 Examination Questions
-	V02.415 Exhibitions
-	V02.450 Fictional Works
-	V02.460 Forms
-	V02.480 Formularies
-	V02.500 Government Publications
-	V02.503 Graphic Novels
-	V02.510 Guidebooks
-	V02.515 Guideline
-	V02.515.500 Practice Guideline
-	V02.520 Handbooks
-	V02.530 Historical Article
-	V02.530.150 Biography
-	V02.530.150.250 Blogs
-	V02.530.150.500 Interview
-	V02.530.150.750 Personal Narratives
-	V02.530.150.750.100 Autobiography
-	V02.530.250 Classical Article
-	V02.530.400 Festschrift
-	V02.540 Humor
-	V02.540.200 Cartoons
-	V02.545 Incunabula
-	V02.550 Indexes
-	V02.575 Instructional Films and Videos
-	V02.600 Journal Article
-	V02.600.249 Introductory Journal Article
-	V02.600.500 Review
-	V02.600.500.500 Consensus Development Conference

## MeSH Tree Changes for 2017

Type	Tree - heading
-	V02.600.500.500.500 Consensus Development Conference, NIH
-	V02.605 Laboratory Manuals
-	V02.607 Lecture Notes
-	V02.610 Legal Cases
-	V02.620 Legislation
-	V02.625 Letter
-	V02.630 Manuscripts
-	V02.650 Meeting Abstracts
-	V02.660 Monograph
-	V02.660.500 Textbooks
-	V02.665 News
-	V02.667 Newspaper Article
-	V02.668 Nurses' Instruction
-	V02.670 Outlines
-	V02.672 Overall
-	V02.672.500 Festschrift
-	V02.675 Patents
-	V02.682 Periodical Index
-	V02.690 Periodicals
-	V02.695 Pharmacopoeias
-	V02.695.500 Herbals
-	V02.697 Photographs
-	V02.700 Pictorial Works
-	V02.700.074 Caricatures
-	V02.700.149 Cartoons
-	V02.700.300 Drawings
-	V02.700.375 Graphic Novels
-	V02.700.450 Maps
-	V02.700.600 Portraits
-	V02.712 Poetry
-	V02.725 Popular Works
-	V02.725.500 Juvenile Literature
-	V02.725.750 Patient Education Handout
-	V02.731 Postcards
-	V02.734 Problems and Exercises
-	V02.735 Programmed Instruction



## MeSH Tree Changes for 2017

Type	Tree - heading	
-	V02.736	Public Service Announcements
-	V02.737	Published Erratum
-	V02.750	Resource Guides
-	V02.800	Retracted Publication
-	V02.825	Retraction of Publication
-	V02.912	Review
-	V02.912.500	Consensus Development Conference
-	V02.912.500.500	Consensus Development Conference, NIH
-	V02.925	Statistics
New Heading	<b>V02.928</b>	<b>Study Guide</b>
-	V02.930	Tables
-	V02.935	Technical Report
-	V02.940	Unedited Footage
-	V02.945	Union Lists
-	V02.950	Unpublished Works
-	V02.970	Video-Audio Media
-	V02.990	Webcasts
-	V03	Study Characteristics
-	V03.100	Case Reports
-	V03.150	Clinical Conference
-	V03.175	Clinical Study
-	V03.175.250	Clinical Trial
-	V03.175.250.100	Clinical Trial, Phase I
-	V03.175.250.200	Clinical Trial, Phase II
-	V03.175.250.300	Clinical Trial, Phase III
-	V03.175.250.400	Clinical Trial, Phase IV
-	V03.175.250.500	Controlled Clinical Trial
-	V03.175.250.500.500	Randomized Controlled Trial
-	V03.175.250.500.500.500	Pragmatic Clinical Trial
-	V03.175.500	Observational Study
-	V03.250	Comparative Study
-	V03.300	Consensus Development Conference
-	V03.300.300	Consensus Development Conference, NIH
-	V03.400	Evaluation Studies
-	V03.600	Meta-Analysis

## MeSH Tree Changes for 2017

Type	Tree - heading
-	V03.650 Multicenter Study
-	V03.800 Scientific Integrity Review
-	V03.900 Twin Study
-	V03.950 Validation Studies
-	V04 Support of Research
-	V04.124 Research Support, Non-U.S. Gov't
-	V04.750 Research Support, U.S. Government
-	V04.750.124 Research Support, American Recovery and Reinvestment Act
-	V04.750.249 Research Support, U.S. Gov't, Non-P.H.S.
-	V04.750.500 Research Support, U.S. Gov't, P.H.S.
-	V04.750.500.500 Research Support, N.I.H., Extramural
-	V04.750.500.750 Research Support, N.I.H., Intramural
-	Z01 Geographic Locations
-	Z01.058 Africa
-	Z01.058.266 Africa, Northern
-	Z01.058.266.104 Algeria
-	Z01.058.266.317 Egypt
-	Z01.058.266.513 Libya
-	Z01.058.266.629 Morocco
-	Z01.058.266.887 Tunisia
-	Z01.058.290 Africa South of the Sahara
-	Z01.058.290.100 Africa, Central
-	Z01.058.290.100.110 Cameroon
-	Z01.058.290.100.120 Central African Republic
-	Z01.058.290.100.130 Chad
-	Z01.058.290.100.140 Congo
-	Z01.058.290.100.220 Democratic Republic of the Congo
-	Z01.058.290.100.300 Equatorial Guinea
-	Z01.058.290.100.380 Gabon
-	Z01.058.290.120 Africa, Eastern
-	Z01.058.290.120.180 Burundi
-	Z01.058.290.120.275 Djibouti
-	Z01.058.290.120.292 Eritrea
-	Z01.058.290.120.310 Ethiopia
-	Z01.058.290.120.400 Kenya
-	Z01.058.290.120.680 Rwanda

## MeSH Tree Changes for 2017

Type	Tree - heading
-	Z01.058.290.120.730 Somalia
-	Z01.058.290.120.745 South Sudan
-	Z01.058.290.120.760 Sudan
-	Z01.058.290.120.840 Tanzania
-	Z01.058.290.120.880 Uganda
-	Z01.058.290.175 Africa, Southern
-	Z01.058.290.175.095 Angola
-	Z01.058.290.175.230 Botswana
-	Z01.058.290.175.400 Lesotho
-	Z01.058.290.175.500 Malawi
-	Z01.058.290.175.545 Mozambique
-	Z01.058.290.175.580 Namibia
-	Z01.058.290.175.735 South Africa
-	Z01.058.290.175.795 Swaziland
-	Z01.058.290.175.920 Zambia
-	Z01.058.290.175.960 Zimbabwe
-	Z01.058.290.190 Africa, Western
-	Z01.058.290.190.200 Benin
-	Z01.058.290.190.245 Burkina Faso
-	Z01.058.290.190.258 Cape Verde
-	Z01.058.290.190.272 Cote d'Ivoire
-	Z01.058.290.190.300 Gambia
-	Z01.058.290.190.320 Ghana
-	Z01.058.290.190.375 Guinea
-	Z01.058.290.190.380 Guinea-Bissau
-	Z01.058.290.190.425 Liberia
-	Z01.058.290.190.500 Mali
-	Z01.058.290.190.520 Mauritania
-	Z01.058.290.190.560 Niger
-	Z01.058.290.190.565 Nigeria
-	Z01.058.290.190.710 Senegal
-	Z01.058.290.190.725 Sierra Leone
-	Z01.058.290.190.800 Togo
-	Z01.107 Americas
-	Z01.107.084 Caribbean Region
-	Z01.107.084.900 West Indies

## MeSH Tree Changes for 2017

Type	Tree - heading
-	Z01.107.084.900.050 Antigua and Barbuda
-	Z01.107.084.900.100 Bahamas
-	Z01.107.084.900.140 Barbados
-	Z01.107.084.900.182 British Virgin Islands
-	Z01.107.084.900.225 Cuba
-	Z01.107.084.900.262 Dominica
-	Z01.107.084.900.300 Dominican Republic
-	Z01.107.084.900.362 Grenada
-	Z01.107.084.900.393 Guadeloupe
-	Z01.107.084.900.425 Haiti
-	Z01.107.084.900.525 Jamaica
-	Z01.107.084.900.600 Martinique
-	Z01.107.084.900.645 Netherlands Antilles
-	Z01.107.084.900.750 Puerto Rico
-	Z01.107.084.900.787 Saint Kitts and Nevis
-	Z01.107.084.900.825 Saint Lucia
-	Z01.107.084.900.862 Saint Vincent and the Grenadines
-	Z01.107.084.900.900 Trinidad and Tobago
-	Z01.107.084.900.970 United States Virgin Islands
-	Z01.107.169 Central America
-	Z01.107.169.133 Belize
-	Z01.107.169.238 Costa Rica
-	Z01.107.169.339 El Salvador
-	Z01.107.169.454 Guatemala
-	Z01.107.169.534 Honduras
-	Z01.107.169.690 Nicaragua
-	Z01.107.169.772 Panama
-	Z01.107.169.772.584 Panama Canal Zone
-	Z01.107.296 Gulf of Mexico
-	Z01.107.424 Latin America
-	Z01.107.567 North America
-	Z01.107.567.176 Canada
-	Z01.107.567.176.064 Alberta
-	Z01.107.567.176.160 British Columbia
-	Z01.107.567.176.410 Manitoba
-	Z01.107.567.176.494 New Brunswick

## MeSH Tree Changes for 2017

Type	Tree - heading
-	Z01.107.567.176.540 Newfoundland and Labrador
-	Z01.107.567.176.564 Northwest Territories
-	Z01.107.567.176.589 Nova Scotia
-	Z01.107.567.176.614 Nunavut
-	Z01.107.567.176.639 Ontario
-	Z01.107.567.176.732 Prince Edward Island
-	Z01.107.567.176.791 Quebec
-	Z01.107.567.176.858 Saskatchewan
-	Z01.107.567.176.929 Yukon Territory
-	Z01.107.567.403 Greenland
-	Z01.107.567.589 Mexico
-	Z01.107.567.875 United States
-	Z01.107.567.875.075 Appalachian Region
-	Z01.107.567.875.075.100 Alabama
-	Z01.107.567.875.075.250 Georgia
-	Z01.107.567.875.075.400 Kentucky
-	Z01.107.567.875.075.418 Maryland
-	Z01.107.567.875.075.437 New York
-	Z01.107.567.875.075.475 North Carolina
-	Z01.107.567.875.075.512 Ohio
-	Z01.107.567.875.075.550 Pennsylvania
-	Z01.107.567.875.075.662 South Carolina
-	Z01.107.567.875.075.775 Tennessee
-	Z01.107.567.875.075.837 Virginia
-	Z01.107.567.875.075.900 West Virginia
-	Z01.107.567.875.350 Great Lakes Region
-	Z01.107.567.875.350.350 Illinois
-	Z01.107.567.875.350.350.200 Chicago
-	Z01.107.567.875.350.360 Indiana
-	Z01.107.567.875.350.500 Michigan
-	Z01.107.567.875.350.510 Minnesota
-	Z01.107.567.875.350.530 New York
-	Z01.107.567.875.350.530.530 New York City
-	Z01.107.567.875.350.540 Ohio
-	Z01.107.567.875.350.550 Pennsylvania
-	Z01.107.567.875.350.900 Wisconsin

## MeSH Tree Changes for 2017

Type	Tree - heading
-	Z01.107.567.875.500 Mid-Atlantic Region
-	Z01.107.567.875.500.200 Delaware
-	Z01.107.567.875.500.210 District of Columbia
-	Z01.107.567.875.500.500 Maryland
-	Z01.107.567.875.500.500.100 Baltimore
-	Z01.107.567.875.500.525 New Jersey
-	Z01.107.567.875.500.530 New York
-	Z01.107.567.875.500.530.530 New York City
-	Z01.107.567.875.500.550 Pennsylvania
-	Z01.107.567.875.500.550.525 Philadelphia
-	Z01.107.567.875.510 Midwestern United States
-	Z01.107.567.875.510.350 Illinois
-	Z01.107.567.875.510.350.200 Chicago
-	Z01.107.567.875.510.360 Indiana
-	Z01.107.567.875.510.370 Iowa
-	Z01.107.567.875.510.390 Kansas
-	Z01.107.567.875.510.400 Kentucky
-	Z01.107.567.875.510.500 Michigan
-	Z01.107.567.875.510.510 Minnesota
-	Z01.107.567.875.510.515 Missouri
-	Z01.107.567.875.510.520 Nebraska
-	Z01.107.567.875.510.525 North Dakota
-	Z01.107.567.875.510.540 Ohio
-	Z01.107.567.875.510.550 Oklahoma
-	Z01.107.567.875.510.700 South Dakota
-	Z01.107.567.875.510.900 Wisconsin
-	Z01.107.567.875.550 New England
-	Z01.107.567.875.550.200 Connecticut
-	Z01.107.567.875.550.500 Maine
-	Z01.107.567.875.550.510 Massachusetts
-	Z01.107.567.875.550.510.210 Boston
-	Z01.107.567.875.550.580 New Hampshire
-	Z01.107.567.875.550.680 Rhode Island
-	Z01.107.567.875.550.880 Vermont
-	Z01.107.567.875.560 Northwestern United States
-	Z01.107.567.875.560.380 Idaho

## MeSH Tree Changes for 2017

Type	Tree - heading
-	Z01.107.567.875.560.500 Montana
-	Z01.107.567.875.560.550 Oregon
-	Z01.107.567.875.560.900 Washington
-	Z01.107.567.875.560.925 Wyoming
-	Z01.107.567.875.580 Pacific States
-	Z01.107.567.875.580.100 Alaska
-	Z01.107.567.875.580.200 California
-	Z01.107.567.875.580.200.450 Los Angeles
-	Z01.107.567.875.580.200.700 San Francisco
-	Z01.107.567.875.580.375 Hawaii
-	Z01.107.567.875.580.550 Oregon
-	Z01.107.567.875.580.900 Washington
-	Z01.107.567.875.750 Southeastern United States
-	Z01.107.567.875.750.100 Alabama
-	Z01.107.567.875.750.110 Arkansas
-	Z01.107.567.875.750.350 Florida
-	Z01.107.567.875.750.370 Georgia
-	Z01.107.567.875.750.480 Louisiana
-	Z01.107.567.875.750.480.500 New Orleans
-	Z01.107.567.875.750.500 Mississippi
-	Z01.107.567.875.750.530 North Carolina
-	Z01.107.567.875.750.700 South Carolina
-	Z01.107.567.875.750.870 Virginia
-	Z01.107.567.875.750.900 West Virginia
-	Z01.107.567.875.760 Southwestern United States
-	Z01.107.567.875.760.100 Arizona
-	Z01.107.567.875.760.200 California
-	Z01.107.567.875.760.200.450 Los Angeles
-	Z01.107.567.875.760.200.700 San Francisco
-	Z01.107.567.875.760.210 Colorado
-	Z01.107.567.875.760.550 Nevada
-	Z01.107.567.875.760.560 New Mexico
-	Z01.107.567.875.760.750 Texas
-	Z01.107.567.875.760.800 Utah
-	Z01.107.757 South America
-	Z01.107.757.077 Argentina

## MeSH Tree Changes for 2017

Type	Tree - heading
-	Z01.107.757.136 Bolivia
-	Z01.107.757.176 Brazil
-	Z01.107.757.235 Chile
-	Z01.107.757.284 Colombia
-	Z01.107.757.362 Ecuador
-	Z01.107.757.435 French Guiana
-	Z01.107.757.488 Guyana
-	Z01.107.757.656 Paraguay
-	Z01.107.757.702 Peru
-	Z01.107.757.833 Suriname
-	Z01.107.757.904 Uruguay
-	Z01.107.757.943 Venezuela
-	Z01.158 Antarctic Regions
-	Z01.208 Arctic Regions
-	Z01.252 Asia
-	Z01.252.100 Asia, Central
-	Z01.252.100.420 Kazakhstan
-	Z01.252.100.450 Kyrgyzstan
-	Z01.252.100.900 Tajikistan
-	Z01.252.100.940 Turkmenistan
-	Z01.252.100.975 Uzbekistan
-	Z01.252.122 Asia, Northern
-	Z01.252.122.500 Russia
-	Z01.252.122.500.500 Siberia
-	Z01.252.145 Asia, Southeastern
-	Z01.252.145.103 Borneo
-	Z01.252.145.130 Brunei
-	Z01.252.145.182 Cambodia
-	Z01.252.145.281 Timor-Leste
-	Z01.252.145.380 Indonesia
-	Z01.252.145.435 Laos
-	Z01.252.145.487 Malaysia
-	Z01.252.145.530 Mekong Valley
-	Z01.252.145.570 Myanmar
-	Z01.252.145.671 Philippines
-	Z01.252.145.774 Singapore



## MeSH Tree Changes for 2017

Type	Tree - heading
-	Z01.252.145.841 Thailand
-	Z01.252.145.945 Vietnam
-	Z01.252.245 Asia, Western
-	Z01.252.245.131 Bangladesh
-	Z01.252.245.162 Bhutan
-	Z01.252.245.393 India
-	Z01.252.245.393.750 Sikkim
-	Z01.252.245.500 Middle East
-	Z01.252.245.500.125 Afghanistan
-	Z01.252.245.500.175 Bahrain
-	Z01.252.245.500.350 Iran
-	Z01.252.245.500.360 Iraq
-	Z01.252.245.500.375 Israel
-	Z01.252.245.500.400 Jordan
-	Z01.252.245.500.425 Kuwait
-	Z01.252.245.500.450 Lebanon
-	Z01.252.245.500.600 Oman
-	Z01.252.245.500.690 Qatar
-	Z01.252.245.500.750 Saudi Arabia
-	Z01.252.245.500.795 Syria
-	Z01.252.245.500.850 Turkey
-	Z01.252.245.500.900 United Arab Emirates
-	Z01.252.245.500.950 Yemen
-	Z01.252.245.674 Nepal
-	Z01.252.245.723 Pakistan
-	Z01.252.245.840 Sri Lanka
-	Z01.252.474 Far East
-	Z01.252.474.164 China
-	Z01.252.474.164.225 Beijing
-	Z01.252.474.164.450 Hong Kong
-	Z01.252.474.164.675 Macau
-	Z01.252.474.164.900 Tibet
-	Z01.252.474.463 Japan
-	Z01.252.474.463.709 Tokyo
-	Z01.252.474.557 Korea
-	Z01.252.474.557.500 Democratic People's Republic of Korea

## MeSH Tree Changes for 2017

Type	Tree - heading
-	Z01.252.474.557.750                      Republic of Korea
-	Z01.252.474.557.750.500                      Seoul
-	Z01.252.474.651                      Mongolia
-	Z01.252.474.872                      Taiwan
-	Z01.433                      Cities
-	Z01.433.100                      Baltimore
-	Z01.433.114                      Beijing
-	Z01.433.128                      Berlin
-	Z01.433.210                      Boston
-	Z01.433.305                      Chicago
-	Z01.433.429                      District of Columbia
-	Z01.433.553                      London
-	Z01.433.565                      Los Angeles
-	Z01.433.653                      Moscow
-	Z01.433.697                      New Orleans
-	Z01.433.741                      New York City
-	Z01.433.800                      Paris
-	Z01.433.820                      Philadelphia
-	Z01.433.850                      Rome
-	Z01.433.875                      San Francisco
-	Z01.433.887                      Seoul
-	Z01.433.900                      Tokyo
-	Z01.542                      Europe
-	Z01.542.049                      Andorra
-	Z01.542.088                      Austria
-	Z01.542.101                      Balkan Peninsula
-	Z01.542.115                      Belgium
-	Z01.542.248                      Europe, Eastern
-	Z01.542.248.020                      Albania
-	Z01.542.248.136                      Baltic States
-	Z01.542.248.136.360                      Estonia
-	Z01.542.248.136.560                      Latvia
-	Z01.542.248.136.580                      Lithuania
-	Z01.542.248.160                      Bosnia and Herzegovina
-	Z01.542.248.180                      Bulgaria
-	Z01.542.248.295                      Croatia

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	Z01.542.248.395	Czech Republic
-	Z01.542.248.495	Hungary
-	Z01.542.248.506	Kosovo
-	Z01.542.248.517	Macedonia (Republic)
-	Z01.542.248.540	Moldova
-	Z01.542.248.609	Montenegro
-	Z01.542.248.679	Poland
-	Z01.542.248.700	Republic of Belarus
-	Z01.542.248.764	Romania
-	Z01.542.248.775	Russia
-	Z01.542.248.775.150	Bashkiria
-	Z01.542.248.775.310	Dagestan
-	Z01.542.248.775.510	Moscow
-	Z01.542.248.775.900	Tatarstan
-	Z01.542.248.786	Serbia
-	Z01.542.248.797	Slovakia
-	Z01.542.248.820	Slovenia
-	Z01.542.248.960	Ukraine
-	Z01.542.286	France
-	Z01.542.286.588	Paris
-	Z01.542.315	Germany
-	Z01.542.315.182	Berlin
-	Z01.542.315.315	Germany, East
-	Z01.542.315.440	Germany, West
-	Z01.542.335	Gibraltar
-	Z01.542.363	Great Britain
-	Z01.542.363	United Kingdom
-	Z01.542.363.161	Channel Islands
-	Z01.542.363.161.500	Guernsey
-	Z01.542.363.300	England
-	Z01.542.363.300.553	London
-	Z01.542.363.602	Northern Ireland
-	Z01.542.363.766	Scotland
-	Z01.542.363.766.500	Hebrides
-	Z01.542.363.914	Wales
-	Z01.542.383	Greece

## MeSH Tree Changes for 2017

Type	Tree - heading
-	Z01.542.467 Ireland
-	Z01.542.489 Italy
-	Z01.542.489.569 Rome
-	Z01.542.489.751 Sicily
-	Z01.542.524 Liechtenstein
-	Z01.542.550 Luxembourg
-	Z01.542.580 Mediterranean Region
-	Z01.542.580.500 Mediterranean Islands
-	Z01.542.580.500.300 Cyprus
-	Z01.542.580.500.650 Malta
-	Z01.542.580.500.751 Sicily
-	Z01.542.616 Monaco
-	Z01.542.651 Netherlands
-	Z01.542.727 Portugal
-	Z01.542.786 San Marino
-	Z01.542.816 Scandinavian and Nordic Countries
-	Z01.542.816.124 Denmark
-	Z01.542.816.124.500 Greenland
-	Z01.542.816.186 Finland
-	Z01.542.816.249 Iceland
-	Z01.542.816.374 Norway
-	Z01.542.816.374.500 Svalbard
-	Z01.542.816.500 Sweden
-	Z01.542.846 Spain
-	Z01.542.883 Switzerland
-	Z01.542.900 Transcaucasia
-	Z01.542.900.099 Armenia
-	Z01.542.900.120 Azerbaijan
-	Z01.542.900.420 Georgia (Republic)
-	Z01.542.961 Vatican City
-	Z01.586 Historical Geographic Locations
-	Z01.586.035 Ancient Lands
-	Z01.586.035.100 Arabia
-	Z01.586.035.150 Armenia
-	Z01.586.035.175 Byzantium
-	Z01.586.035.200 Egypt

## MeSH Tree Changes for 2017

Type	Tree - heading	
-	Z01.586.035.450	Greece
-	Z01.586.035.587	Mesopotamia
-	Z01.586.035.725	Persia
-	Z01.586.117	Austria-Hungary
-	Z01.586.200	Commonwealth of Independent States
-	Z01.586.200.099	Armenia
-	Z01.586.200.120	Azerbaijan
-	Z01.586.200.420	Georgia (Republic)
-	Z01.586.200.440	Kazakhstan
-	Z01.586.200.450	Kyrgyzstan
-	Z01.586.200.540	Moldova
-	Z01.586.200.650	Republic of Belarus
-	Z01.586.200.775	Russia
-	Z01.586.200.775.150	Bashkiria
-	Z01.586.200.775.310	Dagestan
-	Z01.586.200.775.510	Moscow
Old Tree	<b>Z01.586.200.838</b>	<b>Siberia</b>
-	Z01.586.200.900	Tajikistan
-	Z01.586.200.940	Turkmenistan
-	Z01.586.200.960	Ukraine
-	Z01.586.200.975	Uzbekistan
-	Z01.586.225	Confederate States of America
-	Z01.586.250	Czechoslovakia
-	Z01.586.315	Germany
-	Z01.586.315.315	Germany, East
-	Z01.586.315.440	Germany, West
-	Z01.586.315.720	Prussia
-	Z01.586.361	Holy Roman Empire
-	Z01.586.407	Korea
-	Z01.586.500	Middle East
-	Z01.586.575	Netherlands Antilles
-	Z01.586.650	New Guinea
-	Z01.586.687	Ottoman Empire
New Tree	<b>Z01.586.744</b>	<b>Panama Canal Zone</b>
-	Z01.586.800	Russia (Pre-1917)



## MeSH Tree Changes for 2017

Type	Tree - heading	
-	Z01.639.160	Borneo
Old Tree	Z01.639.280	Great Britain
Old Tree	Z01.639.280	United Kingdom
Old Tree	Z01.639.280.161	Channel Islands
Old Tree	Z01.639.280.161.500	Guernsey
Old Tree	Z01.639.280.300	England
Old Tree	Z01.639.280.602	Northern Ireland
Old Tree	Z01.639.280.766	Scotland
Old Tree	Z01.639.280.766.500	Hebrides
Old Tree	Z01.639.280.914	Wales
-	Z01.639.400	Greenland
-	Z01.639.490	Iceland
-	Z01.639.520	Indian Ocean Islands
-	Z01.639.520.200	Comoros
-	Z01.639.520.500	Madagascar
-	Z01.639.520.520	Mauritius
-	Z01.639.520.700	Reunion
-	Z01.639.520.750	Seychelles
-	Z01.639.520.875	Sri Lanka
-	Z01.639.580	Indonesia
-	Z01.639.587	Ireland
-	Z01.639.595	Japan
-	Z01.639.610	Macau
-	Z01.639.640	Mediterranean Islands
-	Z01.639.640.300	Cyprus
-	Z01.639.640.650	Malta
-	Z01.639.640.751	Sicily
-	Z01.639.760	Pacific Islands
-	Z01.639.760.590	Melanesia
-	Z01.639.760.590.373	Fiji
-	Z01.639.760.590.620	New Caledonia
-	Z01.639.760.590.715	Papua New Guinea
-	Z01.639.760.590.736	Vanuatu
-	Z01.639.760.680	Micronesia
-	Z01.639.760.680.435	Guam
-	Z01.639.760.680.717	Palau

## MeSH Tree Changes for 2017

Type	Tree - heading
-	Z01.639.760.747 New Zealand
-	Z01.639.760.815 Polynesia
-	Z01.639.760.815.482 Hawaii
-	Z01.639.760.815.700 Pitcairn Island
-	Z01.639.760.815.800 Samoa
-	Z01.639.760.815.800.100 American Samoa
-	Z01.639.760.815.800.400 Independent State of Samoa
-	Z01.639.760.815.890 Tonga
-	Z01.639.790 Philippines
-	Z01.639.820 Prince Edward Island
-	Z01.639.835 Svalbard
-	Z01.639.850 Taiwan
-	Z01.639.880 West Indies
-	Z01.639.880.050 Antigua and Barbuda
-	Z01.639.880.100 Bahamas
-	Z01.639.880.140 Barbados
-	Z01.639.880.225 Cuba
-	Z01.639.880.262 Dominica
-	Z01.639.880.300 Dominican Republic
-	Z01.639.880.362 Grenada
-	Z01.639.880.393 Guadeloupe
-	Z01.639.880.425 Haiti
-	Z01.639.880.525 Jamaica
-	Z01.639.880.600 Martinique
-	Z01.639.880.645 Netherlands Antilles
-	Z01.639.880.750 Puerto Rico
-	Z01.639.880.787 Saint Kitts and Nevis
-	Z01.639.880.825 Saint Lucia
-	Z01.639.880.862 Saint Vincent and the Grenadines
-	Z01.639.880.900 Trinidad and Tobago
-	Z01.639.880.970 United States Virgin Islands
-	Z01.678 Oceania
-	Z01.678.100 Australasia
-	Z01.678.100.373 Australia
-	Z01.678.100.373.500 Australian Capital Territory
-	Z01.678.100.373.750 New South Wales



## MeSH Tree Changes for 2017

Type	Tree - heading
-	Z01.678.100.373.875 Northern Territory
-	Z01.678.100.373.937 Queensland
-	Z01.678.100.373.968 South Australia
-	Z01.678.100.373.984 Tasmania
-	Z01.678.100.373.992 Victoria
-	Z01.678.100.373.996 Western Australia
-	Z01.678.100.747 New Zealand
-	Z01.678.100.873 Pacific Islands
-	Z01.756 Oceans and Seas
-	Z01.756.092 Atlantic Ocean
-	Z01.756.092.650 North Sea
-	Z01.756.217 Black Sea
-	Z01.756.342 Indian Ocean
-	Z01.756.592 Mediterranean Sea
-	Z01.756.700 Pacific Ocean