

# Image Analysis and Artificial Intelligence

Pathology Support, Comparative & Molecular Pathogenesis Branch

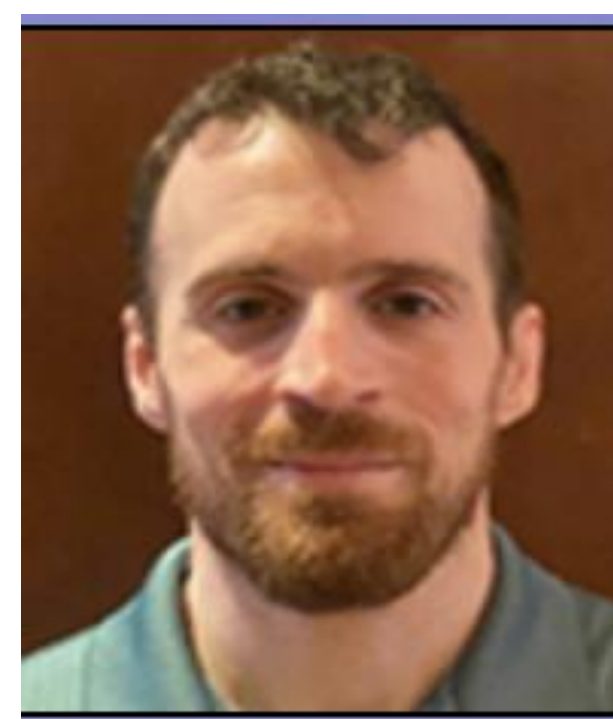
## Mission Statement

Provide automated image analysis and histopathological evaluation using artificial intelligence supported deep learning tools to enhance data rigor and reproducibility for DIR and DTT investigators and collaborating scientists

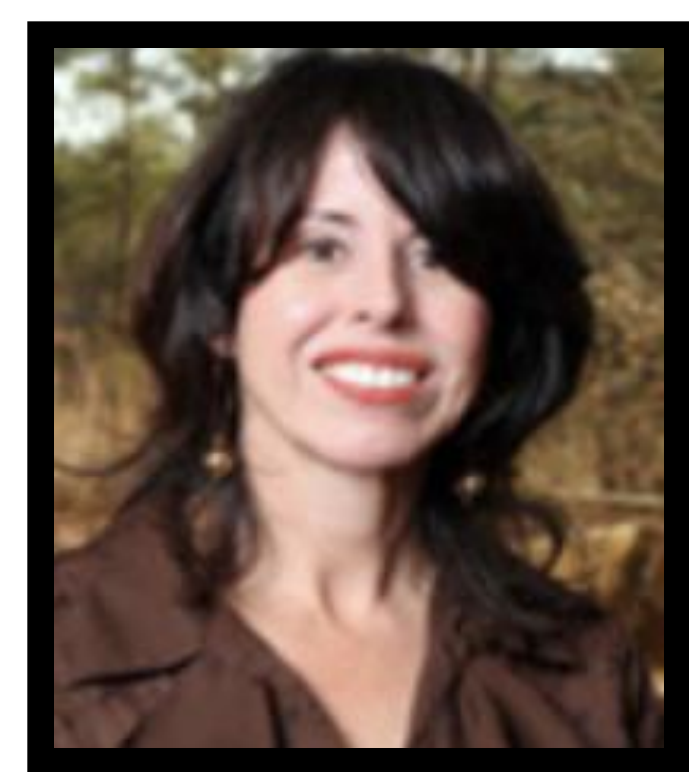
## Staff



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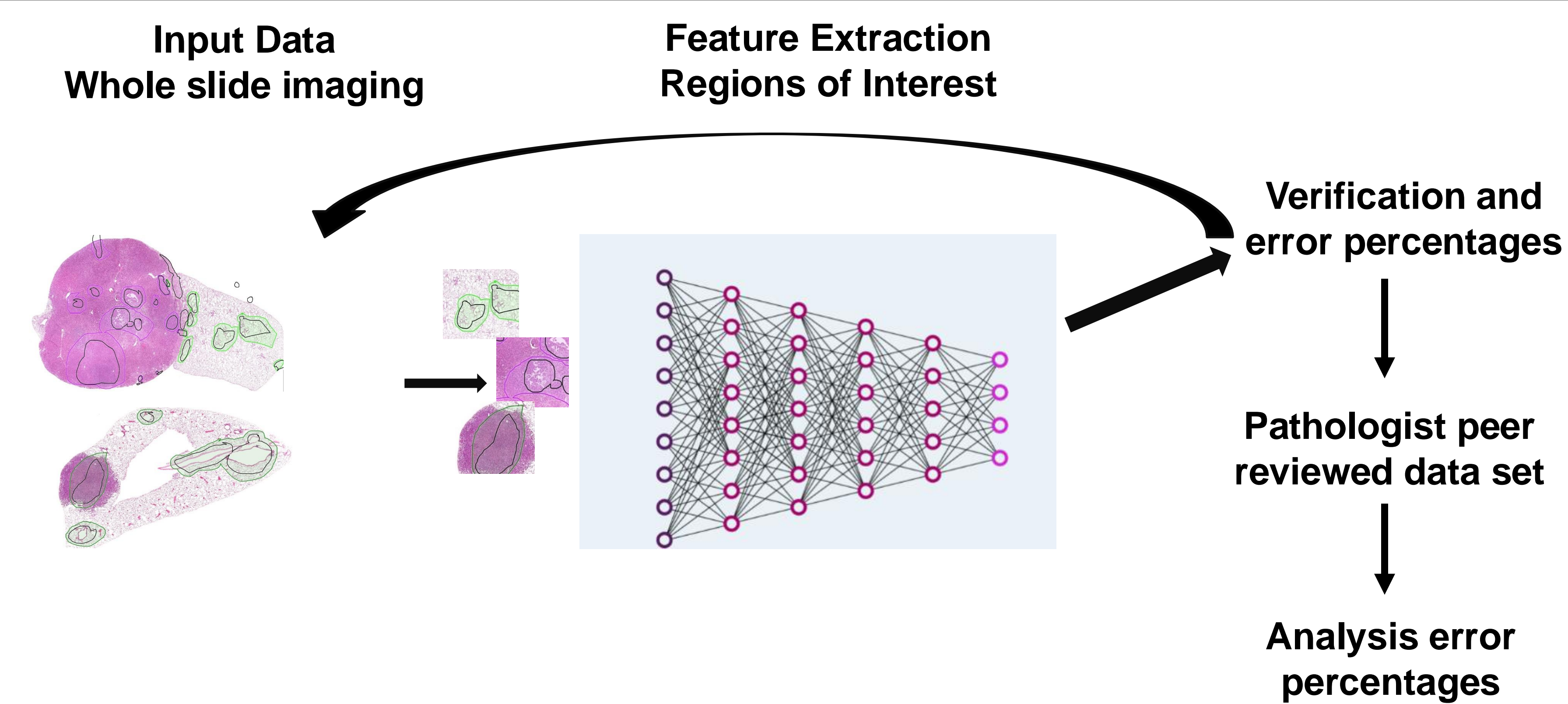


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## Methodology



## Capabilities

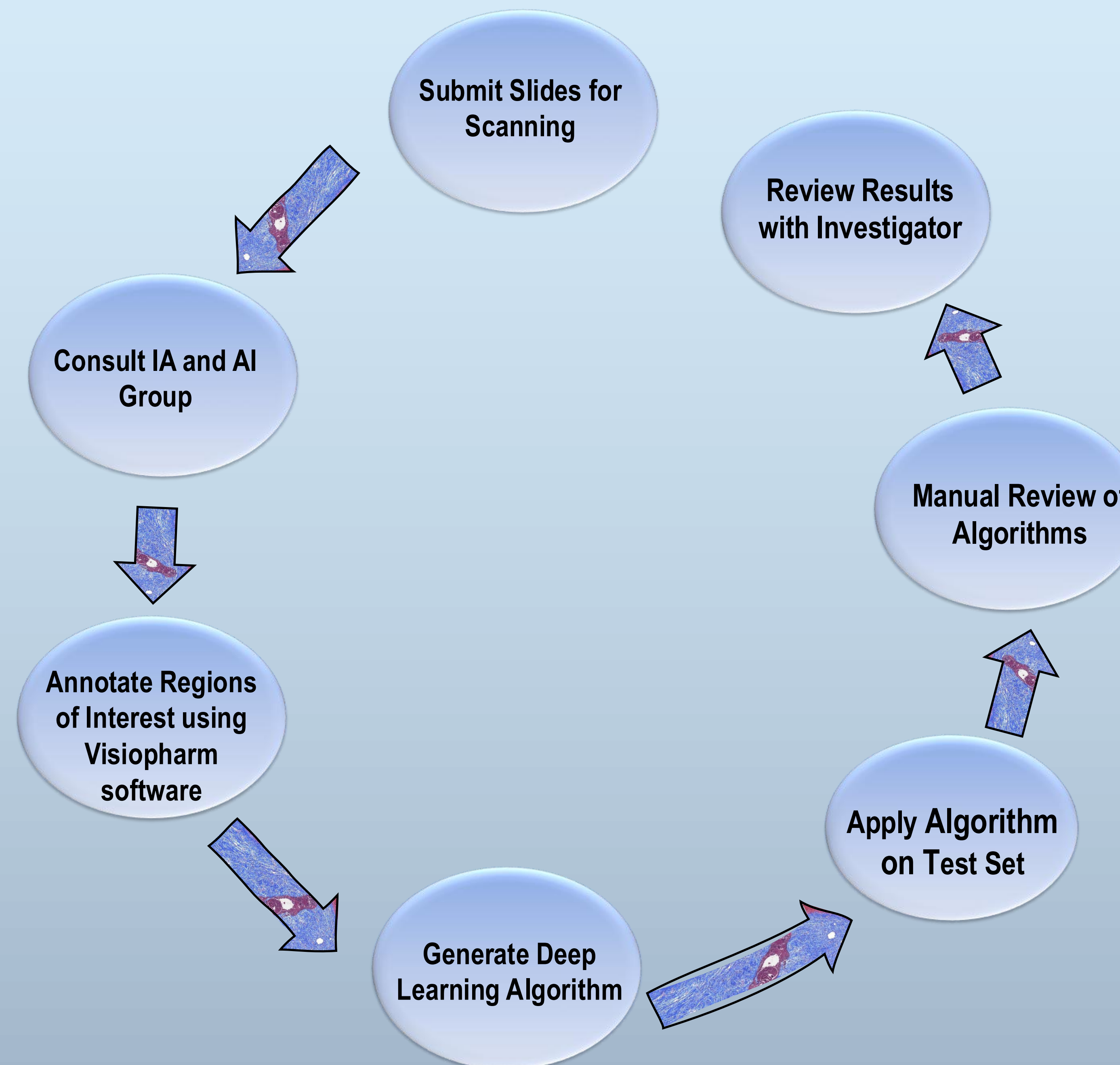
Supervised training of a convolutional neural network (CNN) using Visiopharm<sup>®</sup> Digital Pathology & AI platform

We develop deep learning algorithms for automated image analysis, which include identification and quantification of immunohistochemical and special stains

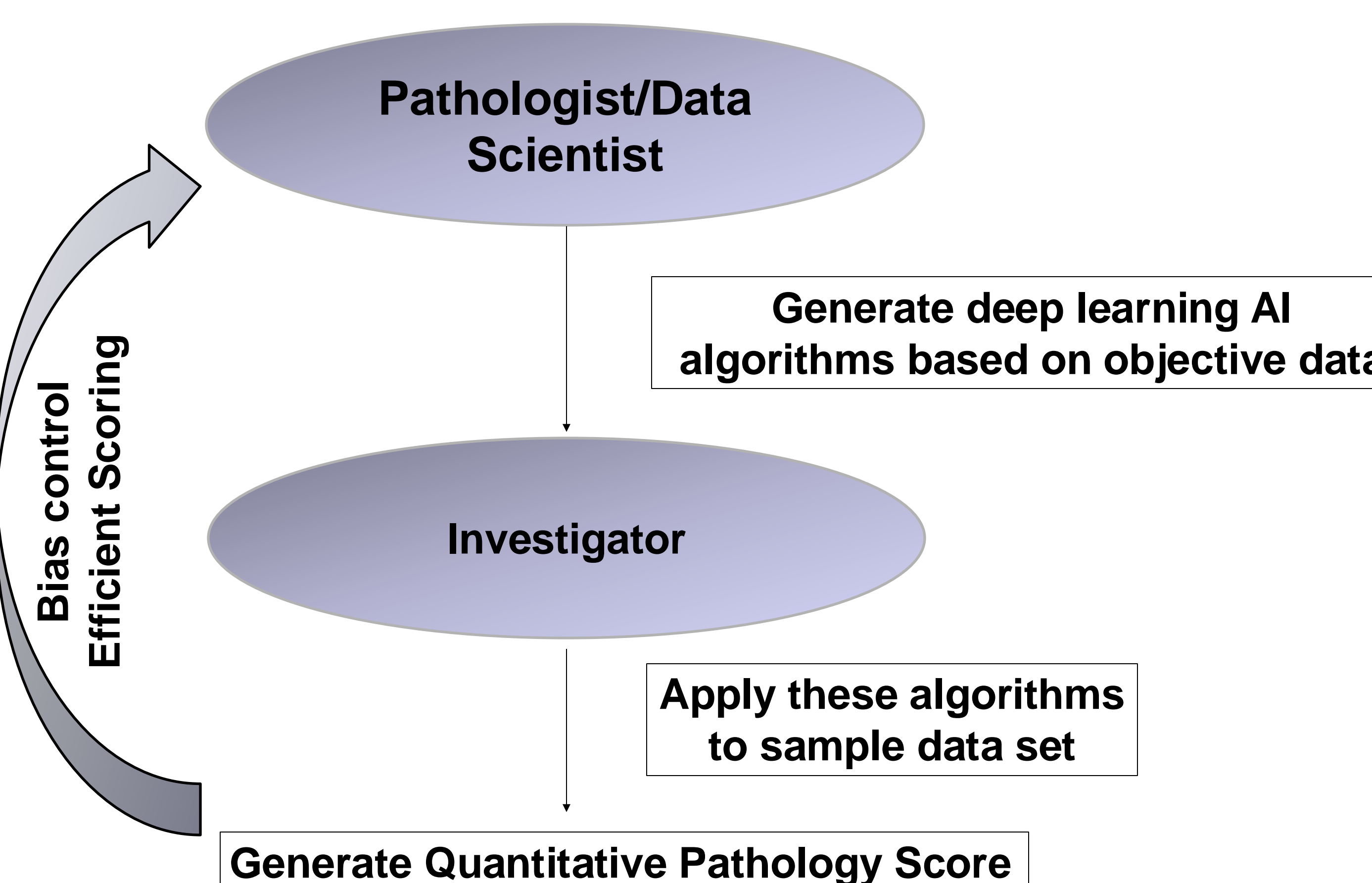
Develop deep learning algorithms for automated H&E whole slides image analysis for lesion identification and quantification

Consultation with investigators

## Work-Flow

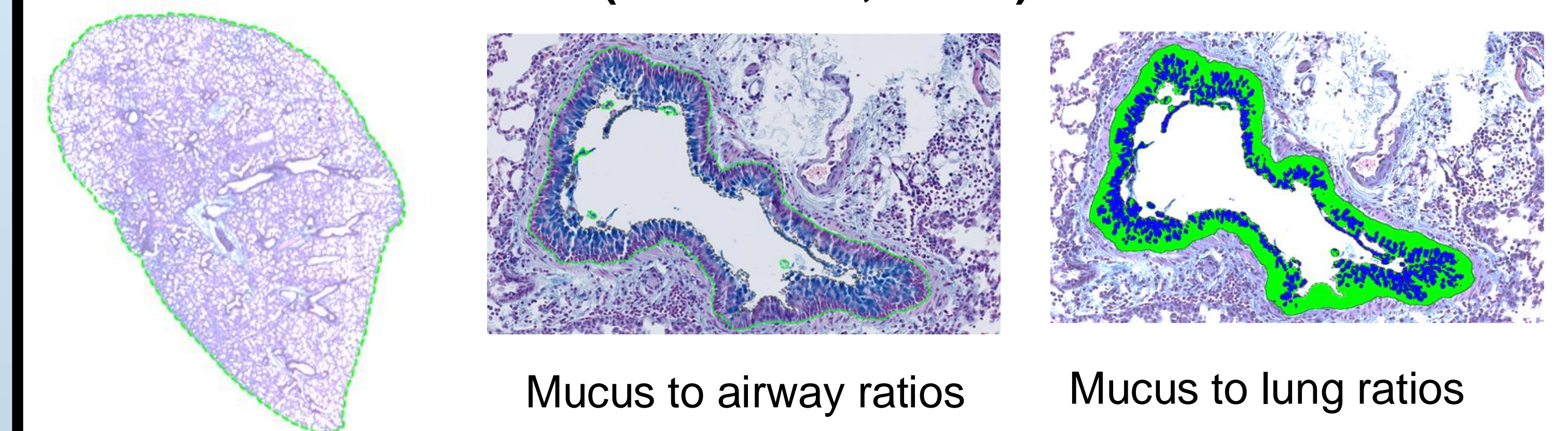


## Summary



## Projects Completed

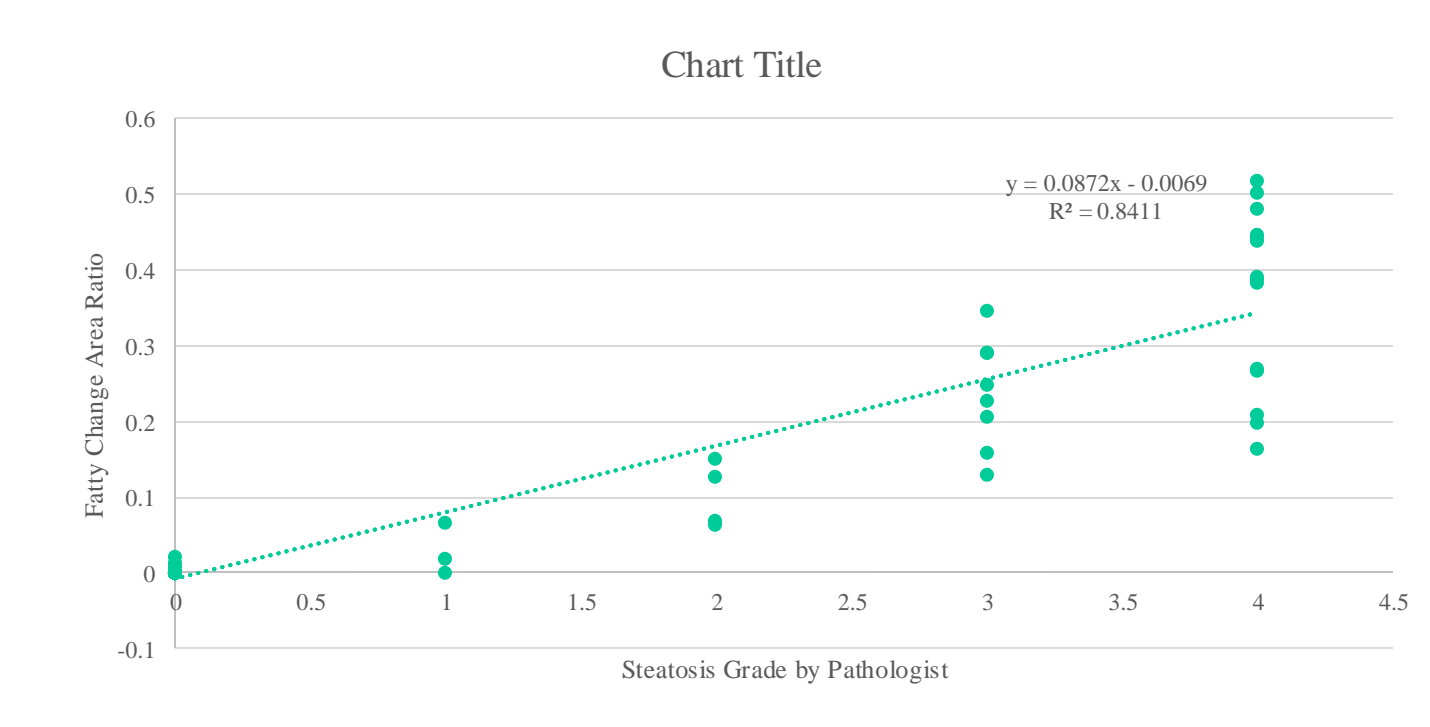
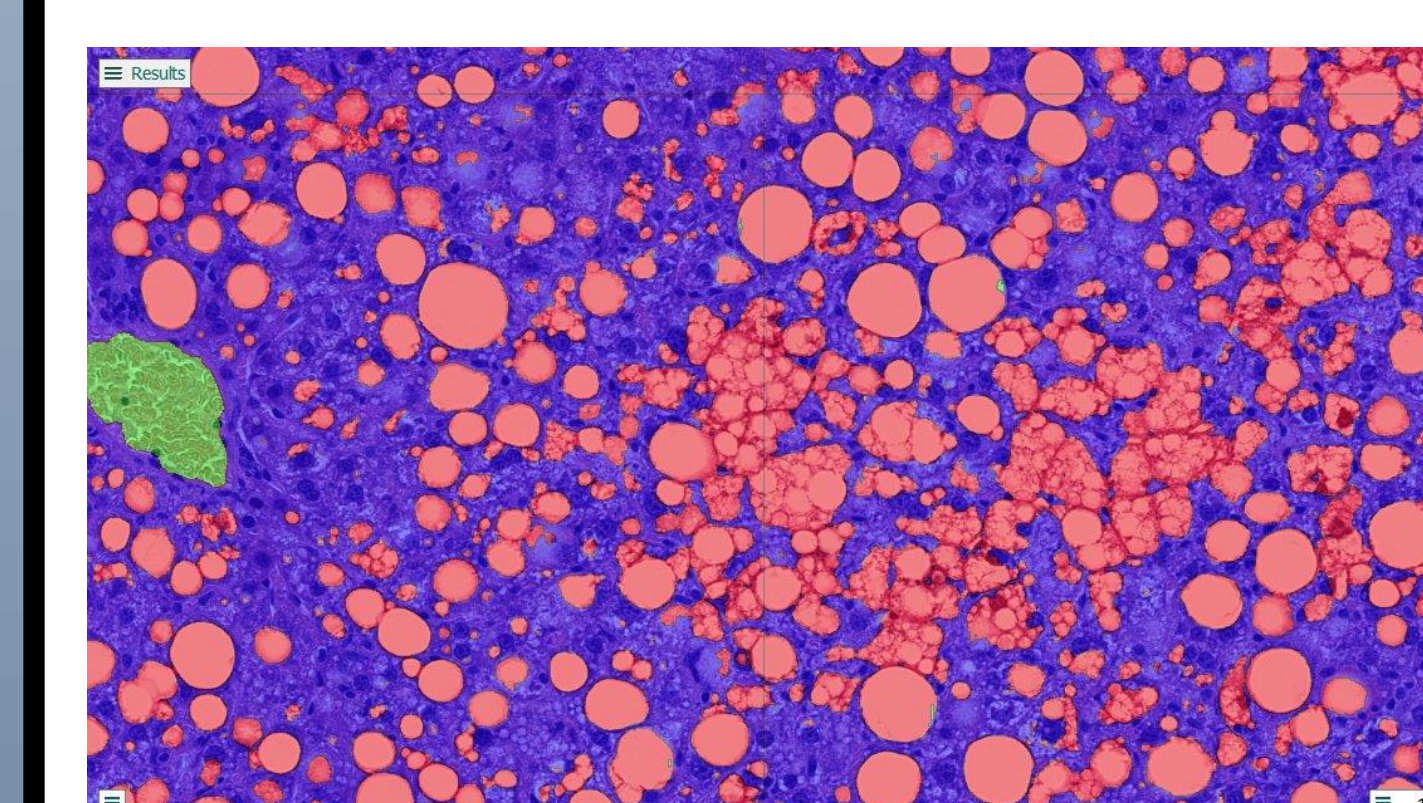
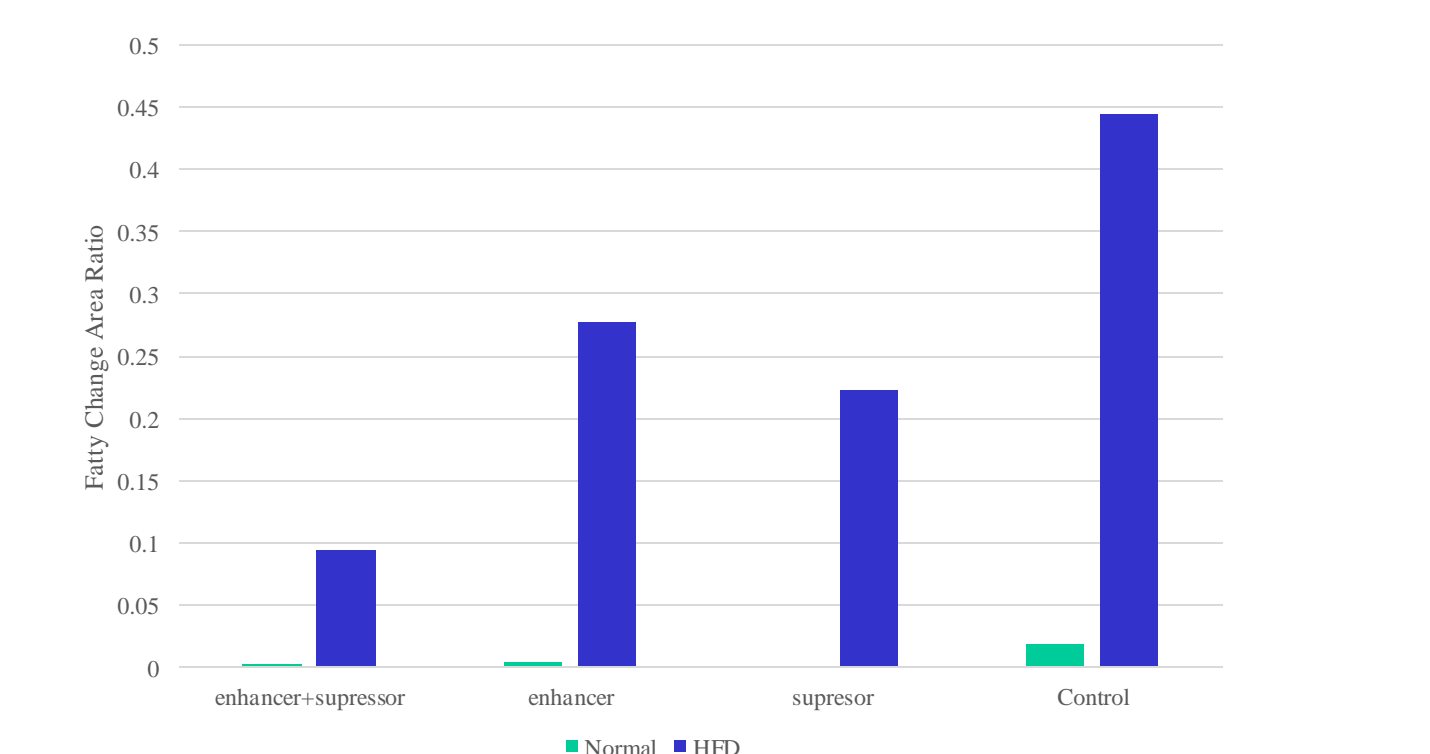
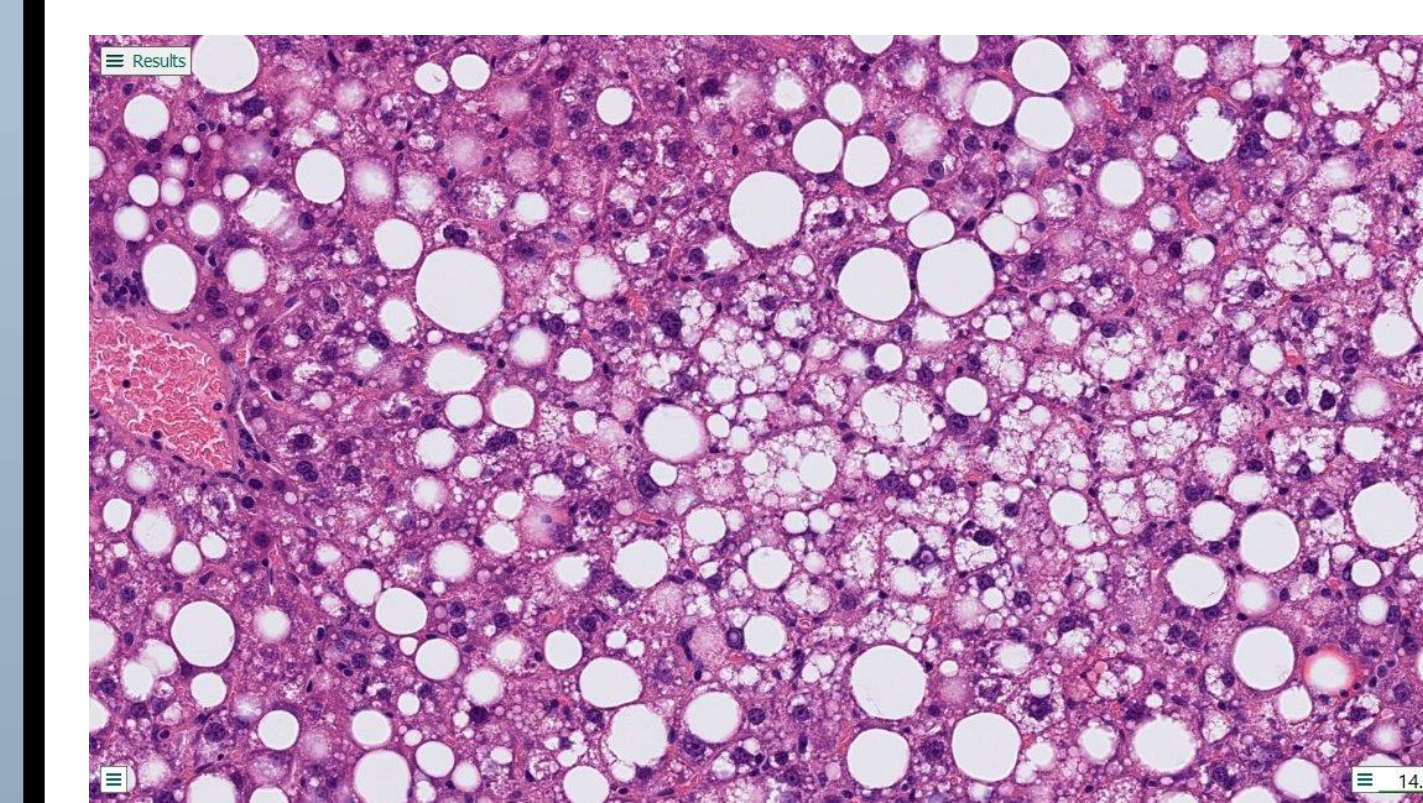
**AI-based image analysis in lung Alcian blue stained sections**  
Chemokine CCL19 promotes type 2 T cell.  
(Nakano H., et. al.)



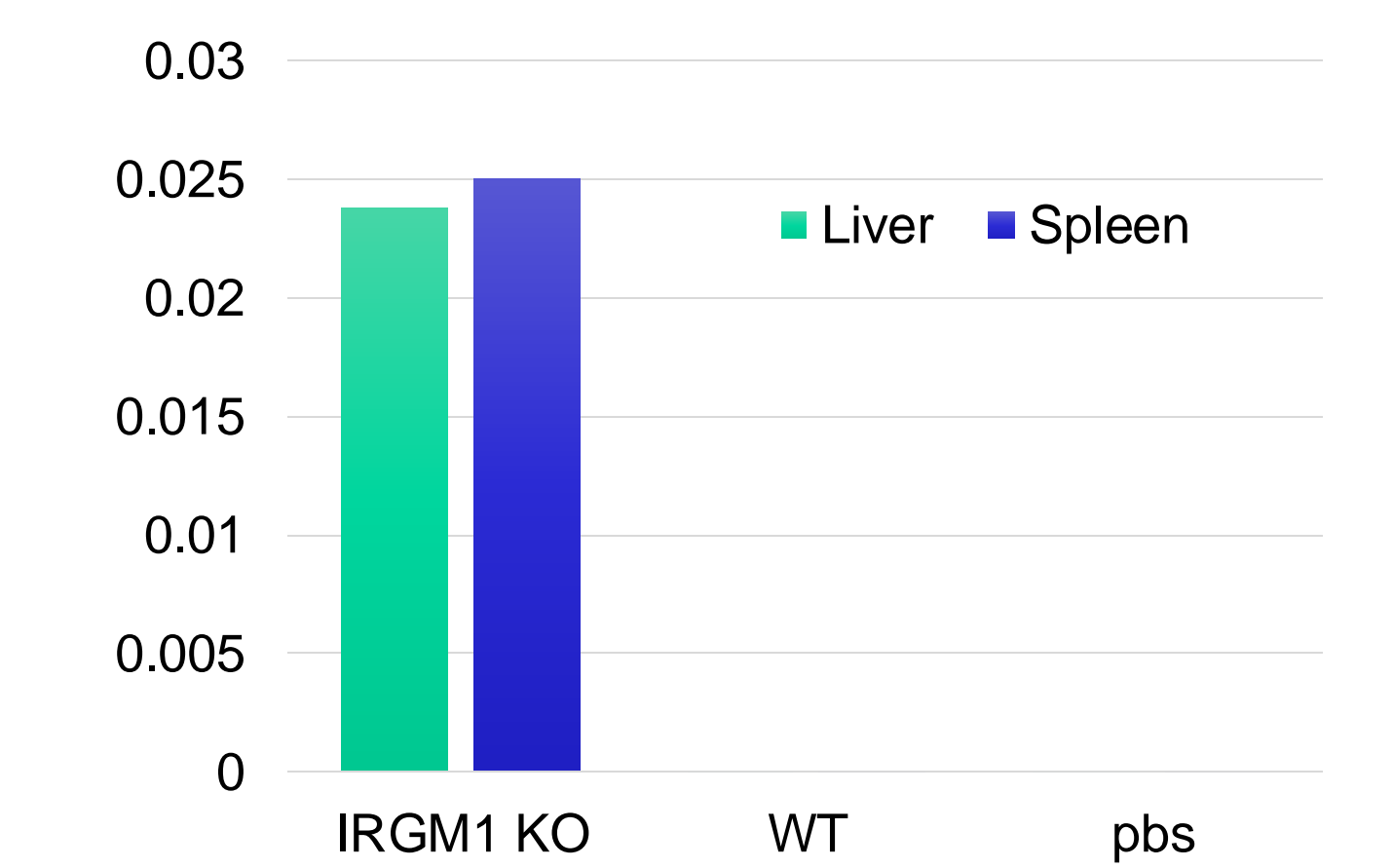
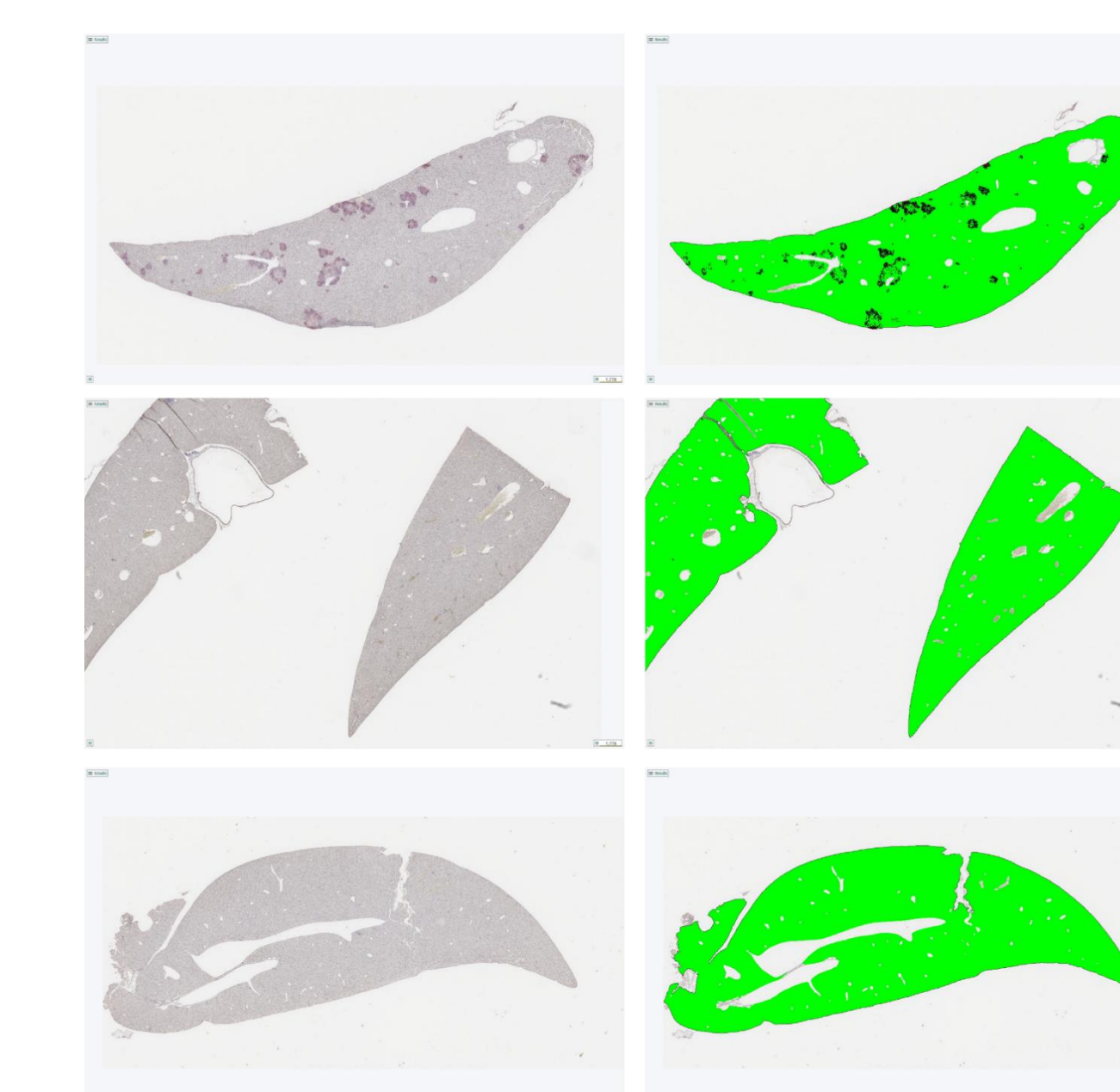
Group	2	3	5	6
Mean	0.00077	0.00087	0.07402	0.08027
SD	0.00176	0.00082	0.06871	0.06855

Group	2	3	5	6
Mean	0.000016	0.000014	0.002249	0.002120
SD	0.000037	0.000013	0.002751	0.002018

**Phenotypic characterization of genetically modified mice fed with normal and high fat diet.**  
(Drs. Blackshear and Arao et. al.)



**Quantification of Listeria IHC staining**



	IRGM1 KO	WT	pbs
Liver	0.0238468	0.0000313	0.0000038
Spleen	0.0250052	0.0000177	0.0000029