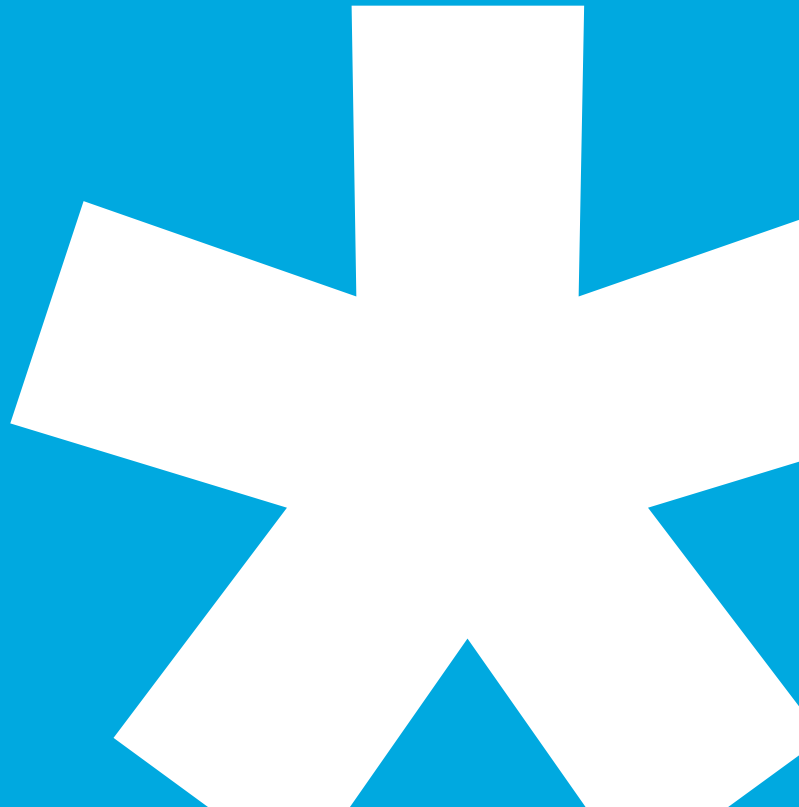


varian

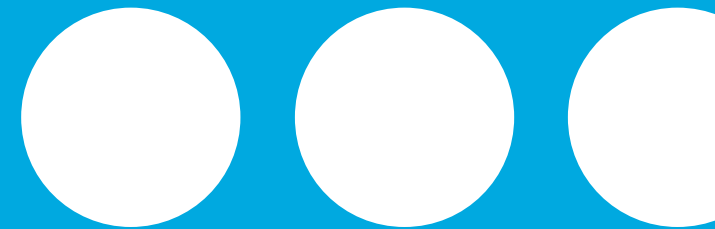
# Nuclei Detection

Sangyu Shen

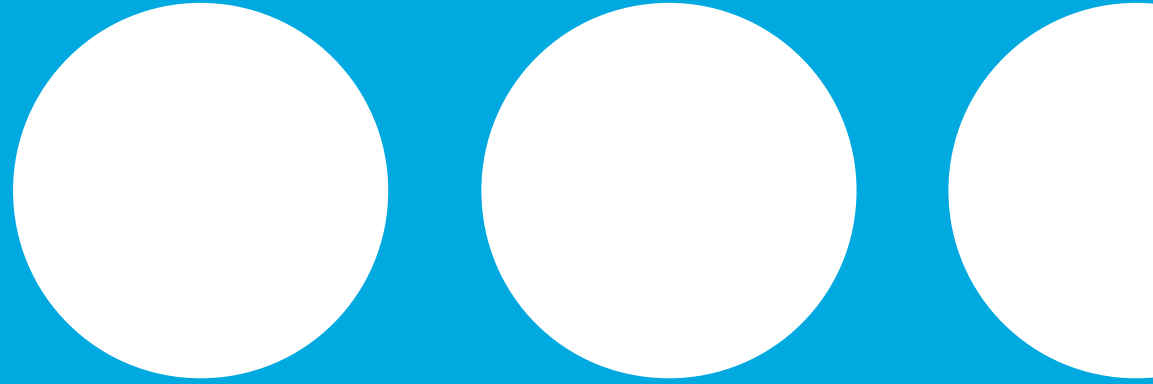


# Agenda

1. Goals of the project
2. Segmentation – Semantic vs Instance
3. Unet Model
4. Mask RCNN Model
5. Model Result
6. Conclusion



# Goals



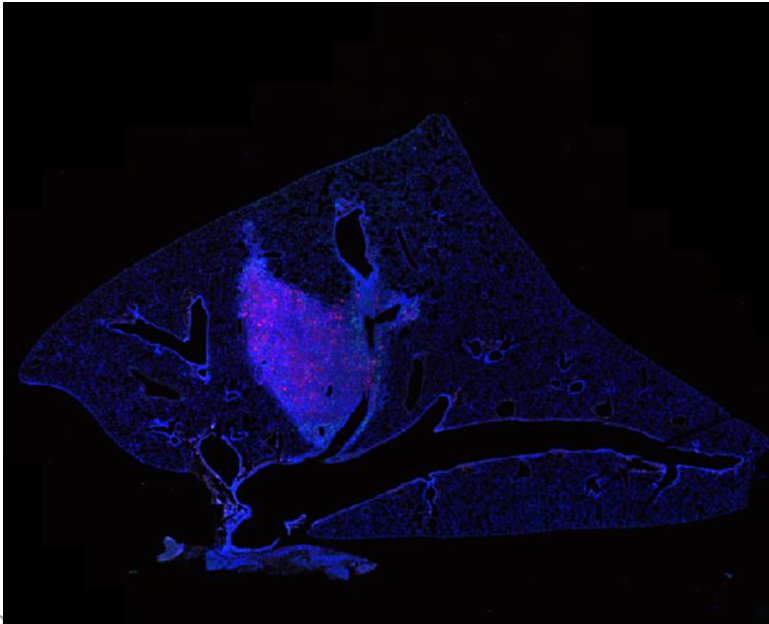
varian

# Nuclei Segmentation

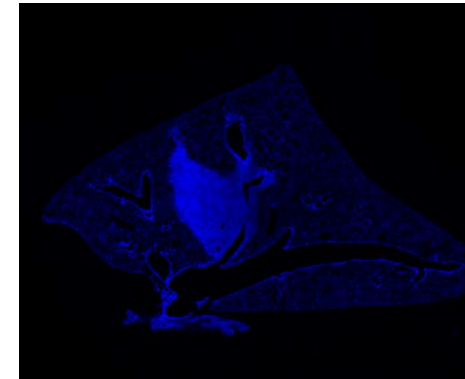
- **Project Goal:** Count and segment nuclei from microscope images

## Counting cells

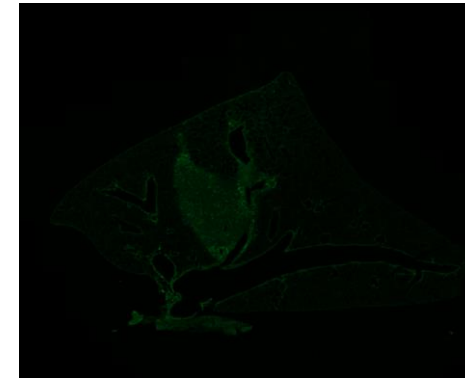
- The first step for many clinical purposes
- Enable extraction of high quality features for analysis in pathology



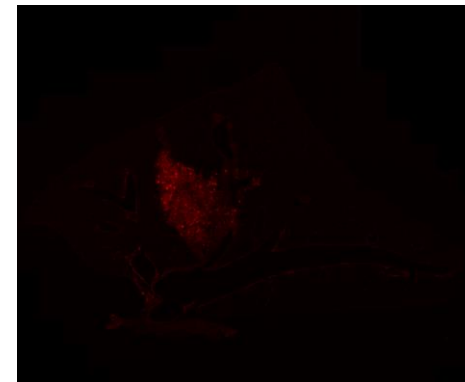
Fluorescently  
stained with  
different  
markers



DAPI:  
Nuclei



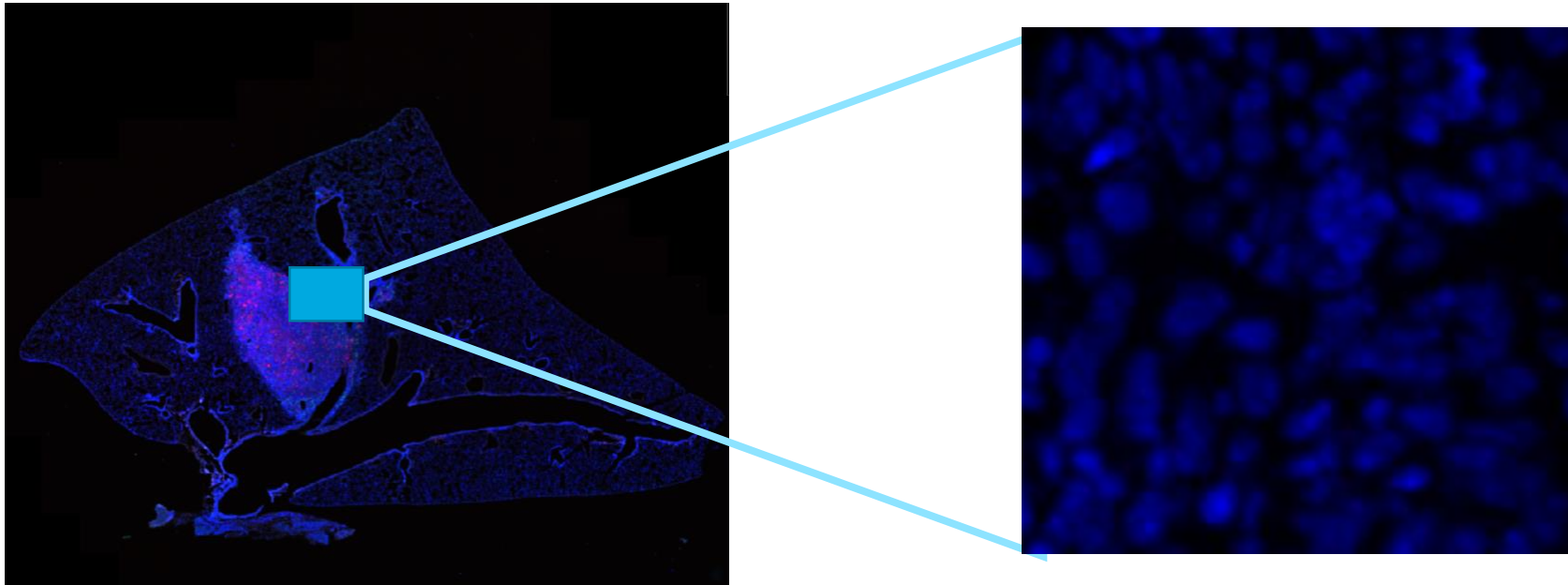
FITC:  
T cell



Cy3:  
Tumor

# Nuclei Segmentation

- **Project Goal:** Count and segment nuclei from microscope images



# Previous Approach

- QuPath
  - Open source software for quantitative pathology (scriptable)
  - Segment cells with filters and thresholds (traditional image processing)
  - Tunable parameters
  - Further analysis pipeline after segmentation

Watershed cell detection

**Setup parameters**

Choose detection channel

**Nucleus parameters**

Background radius  px

Median filter radius  px

Sigma  px

Minimum area  px<sup>2</sup>

Maximum area  px<sup>2</sup>

**Intensity parameters**

Threshold

☒ Split by shape

**Cell parameters**

Cell expansion  px

☒ Include cell nucleus

**General parameters**

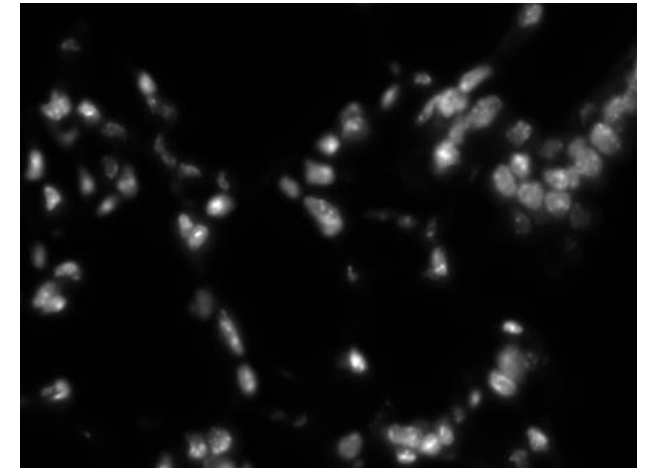
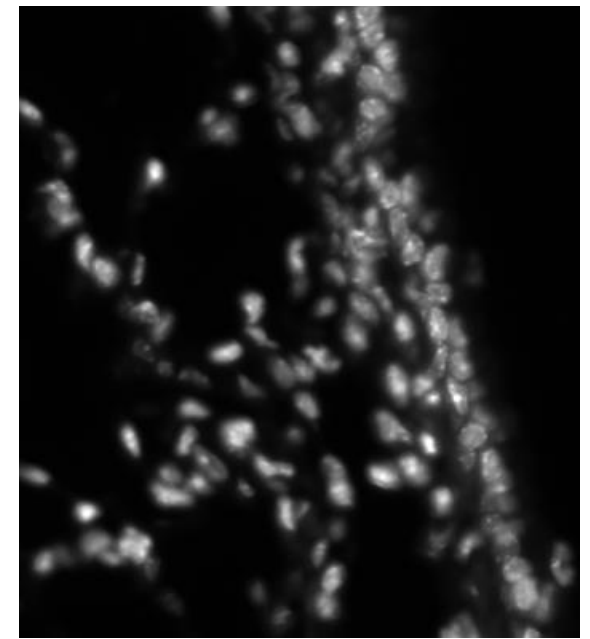
☒ Smooth boundaries

☒ Make measurements

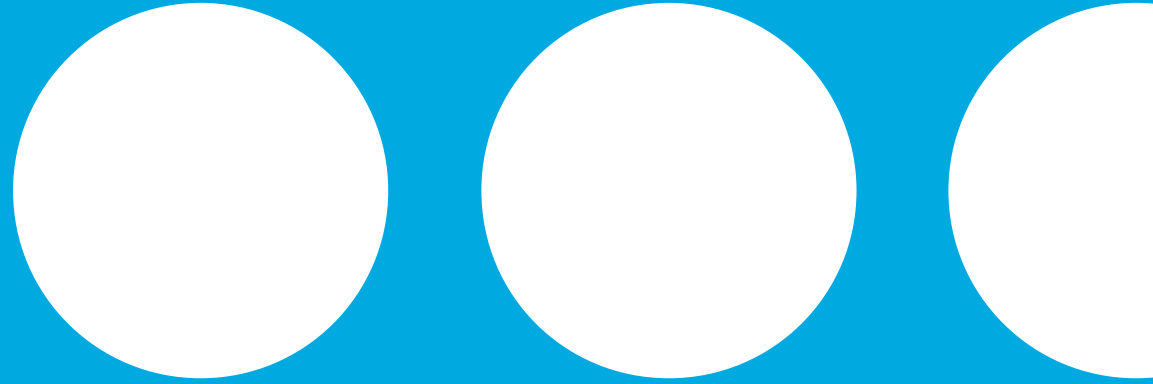
# Challenges

- Manual counting
  - Time consuming
  - Labor intensive
  - Impossible to count a whole tissue slice
  - Difficult quality control – sampling and random errors
- Thresholding techniques
  - Various staining conditions, cell types
  - Difficult to count overlapping cells
  - No standard requirements
  - Threshold parameters: not one size fits all

Need new method – fast, robust, general



Background

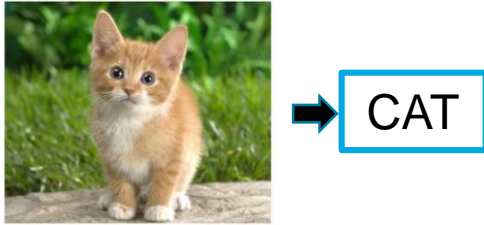


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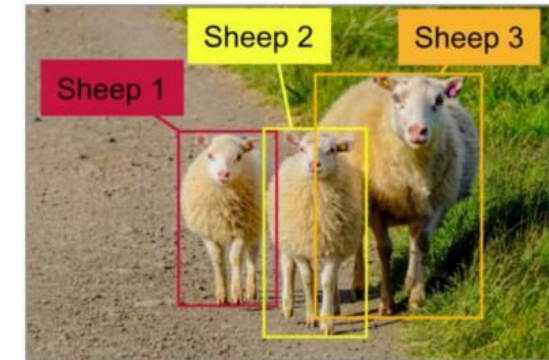


# Image Processing Problem

- Classification



- Object Detection



- Semantic Segmentation

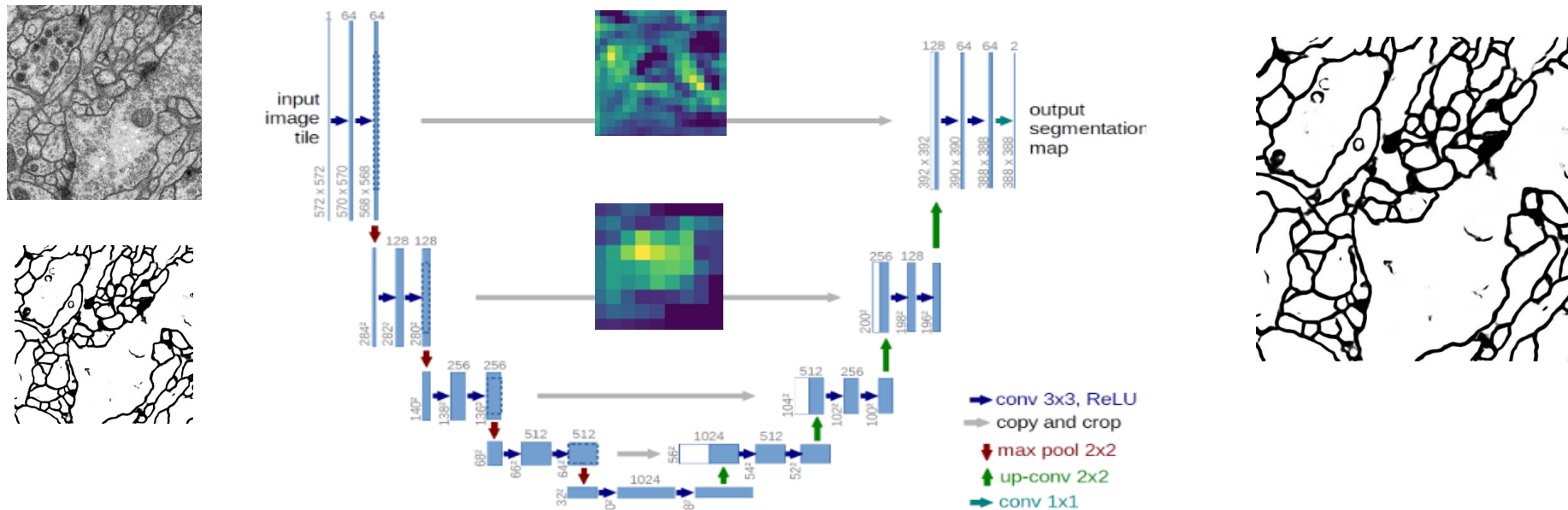


- Instance Segmentation



# Segmentation Algorithm - Unet

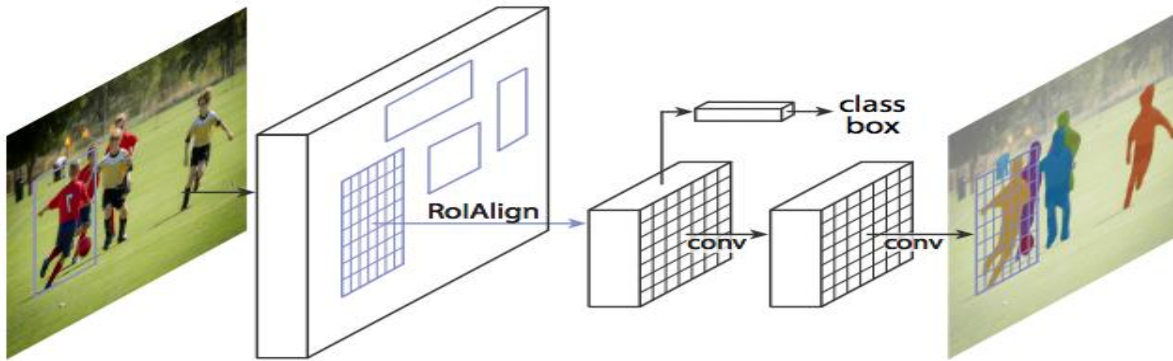
- Unet algorithm – semantic segmentation
  - Convolutional Networks for Biomedical Image Segmentation
  - Created by Olaf Ronneberger, Philipp Fischer, Thomas Brox in 2015



- Encoder: captures context from image
- Decoder: enables precise localization

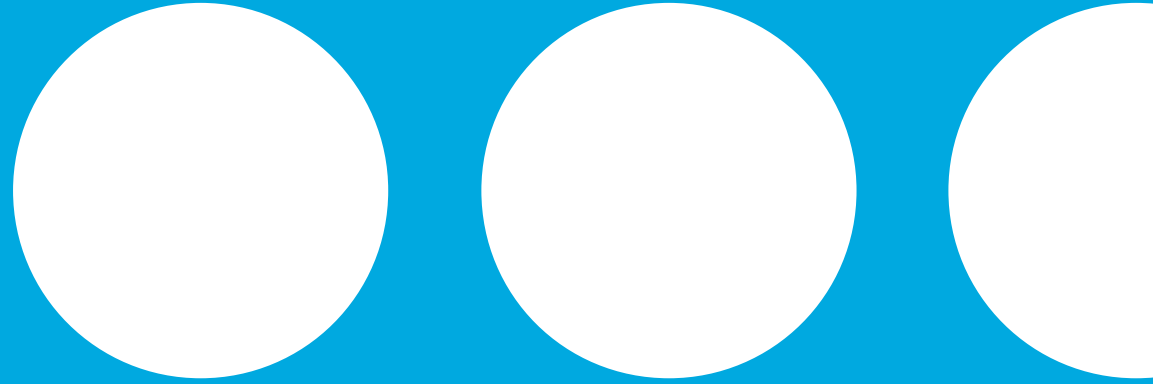
# Segmentation Algorithm – Mask RCNN

- Mask R CNN algorithm – instance segmentation
  - Created by Facebook AI research, 2017
  - Generates bounding boxes and segmentation masks for each instance of an object in the image



- Region Proposal Network: proposes candidate object bounding boxes
- Feature Extraction and Prediction: extracts from each candidate box; classify and re-define bounding-box
- Mask: also outputs a binary mask for each box

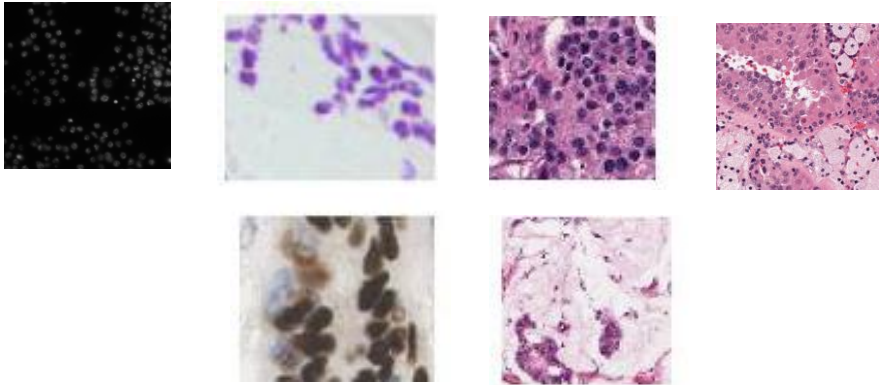
Methodology



varian

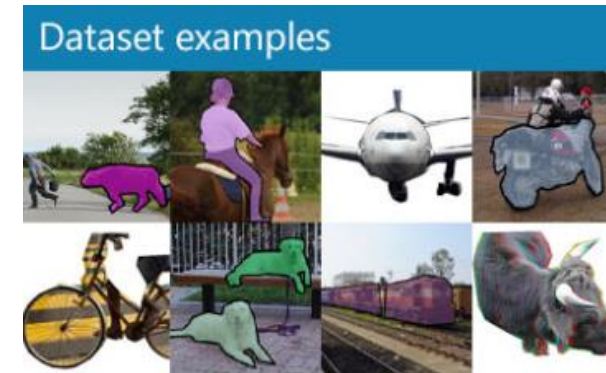
# Nuclei Segmentation Training

- Dataset:
  - Booz Allen Hamilton Nuclei Segmentation Dataset
    - ~900 images with masks
    - a variety of conditions, e.g, cell type, magnification, and imaging modality (brightfield vs. fluorescence)



- Augmentation:
  - horizontal or vertical flips
  - Random rotation: 90 or (-10, 10) degrees
  - Random resize image

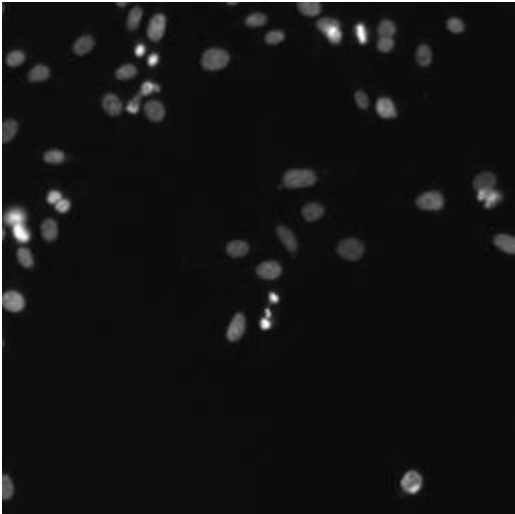
- Pre-train Weights:
  - Computed based on COCO dataset
  - Provided a better starting point



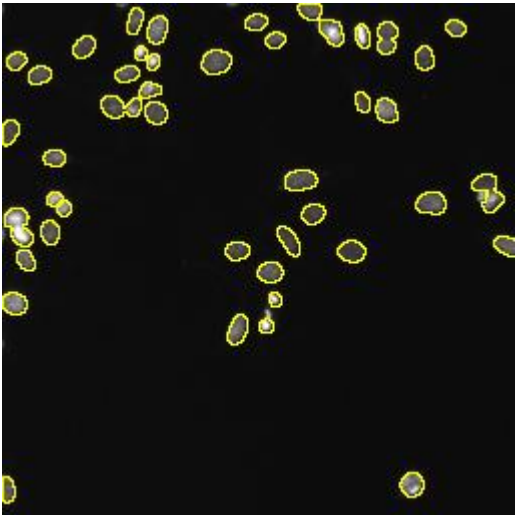
- a large-scale object detection, segmentation, and captioning dataset.
- 330K images (>200K labeled)
- 1.5 million object instances
- 80 object categories

# Nuclei Segmentation Training

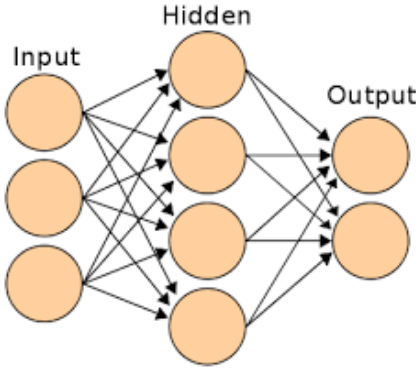
Image



Ground Truth



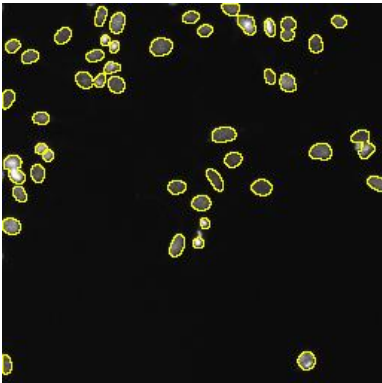
AL USE ONLY



Prediction	
Image ID	Pixels
Image-2019-01	59776 8 60126 12
	60469 22 60820 24
	61171 26 61523 27
	61875 27 62227
Image-2019-01	5233 3 5583 7 5934 9
	6286 9 6637 11 6989
...	...

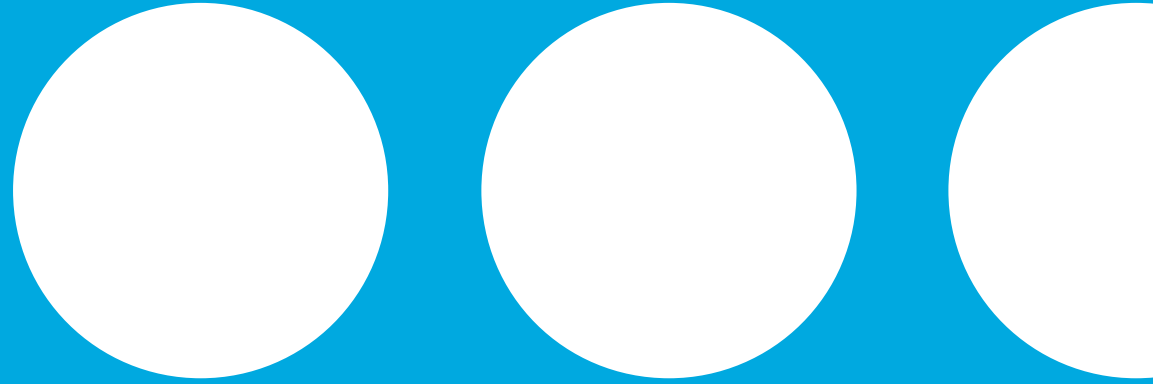
# rows = total cells

Mask Visualization



varian

Result



varian

# Nuclei Segmentation Training

Model	Training Time	Prediction Time	Complete Prediction	# total cells
Unet	< 24 hours	~ 40s	~ 24 hours *	-
Mask RCNN	~ 2 days	<1s	~ 2.5 hours	~ 150,000 *

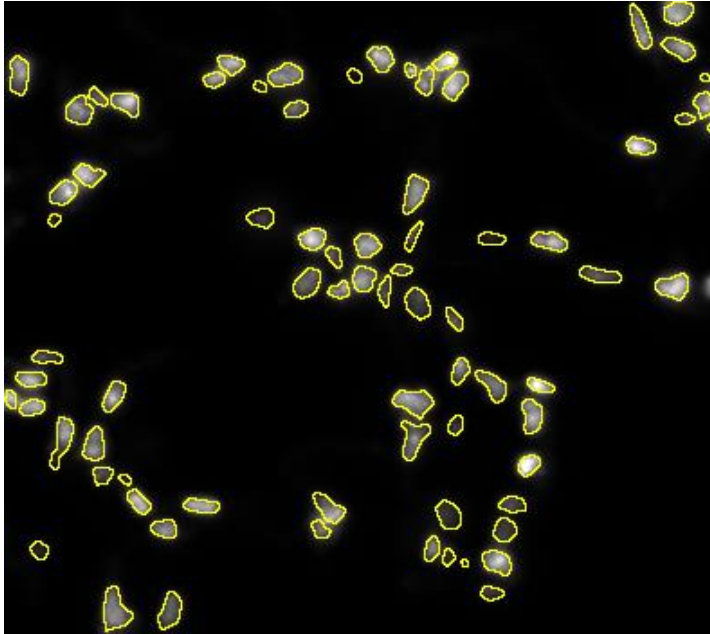
\* Estimated – 6 batches in 3 hours ( total 49 batches)

Execution: NVIDIA GK210 GPU 12GB

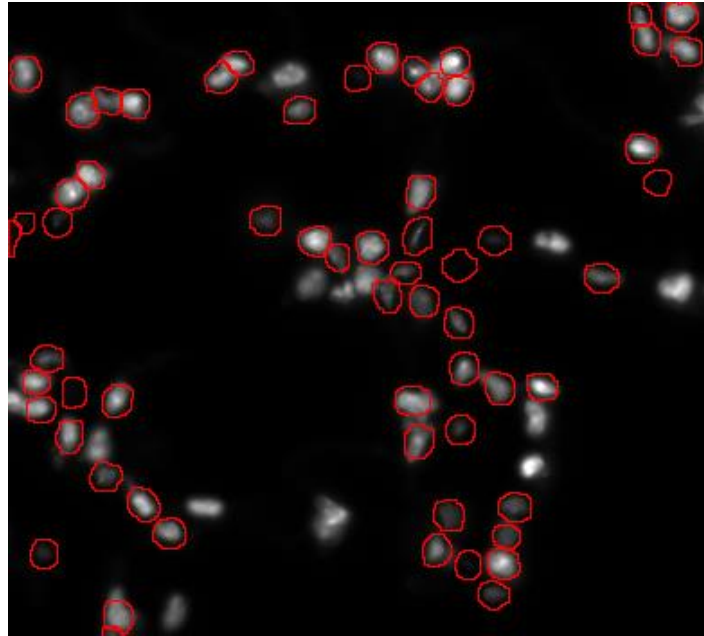


# Testing – Outcome between Unet and Mask RCNN

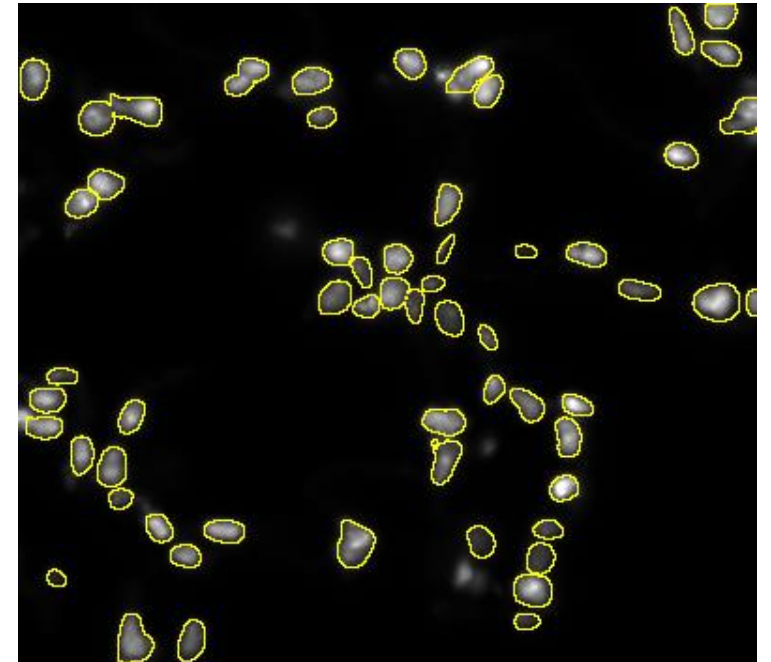
QuPath



Unet

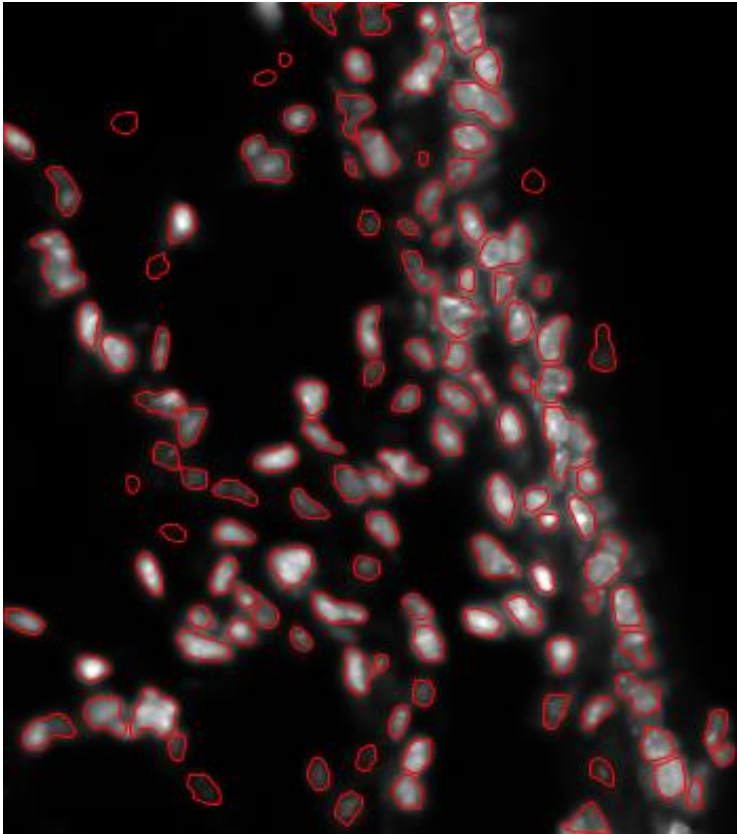


Mask  
RCNN

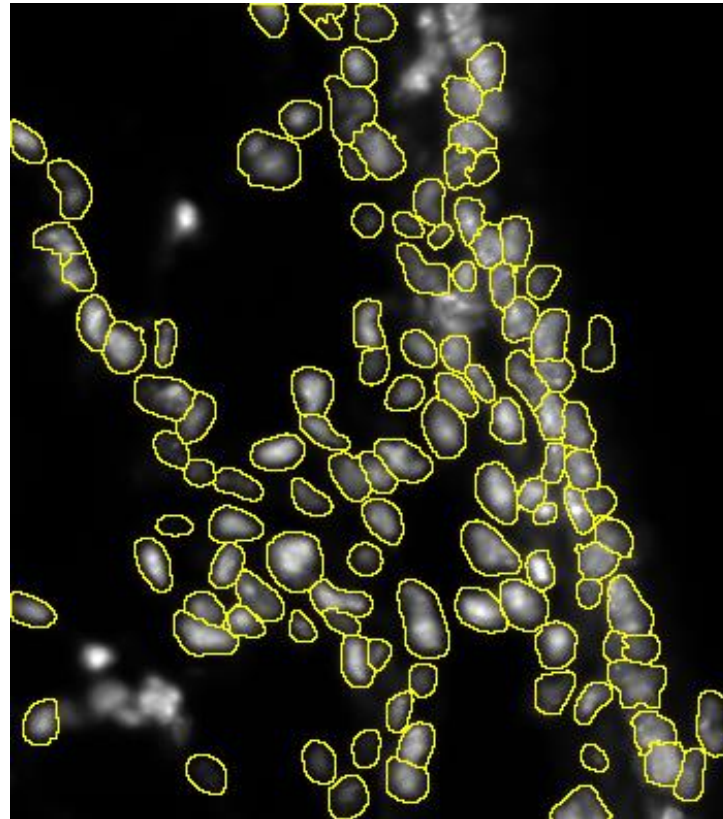


# Testing – Outcome between Unet and Mask RCNN

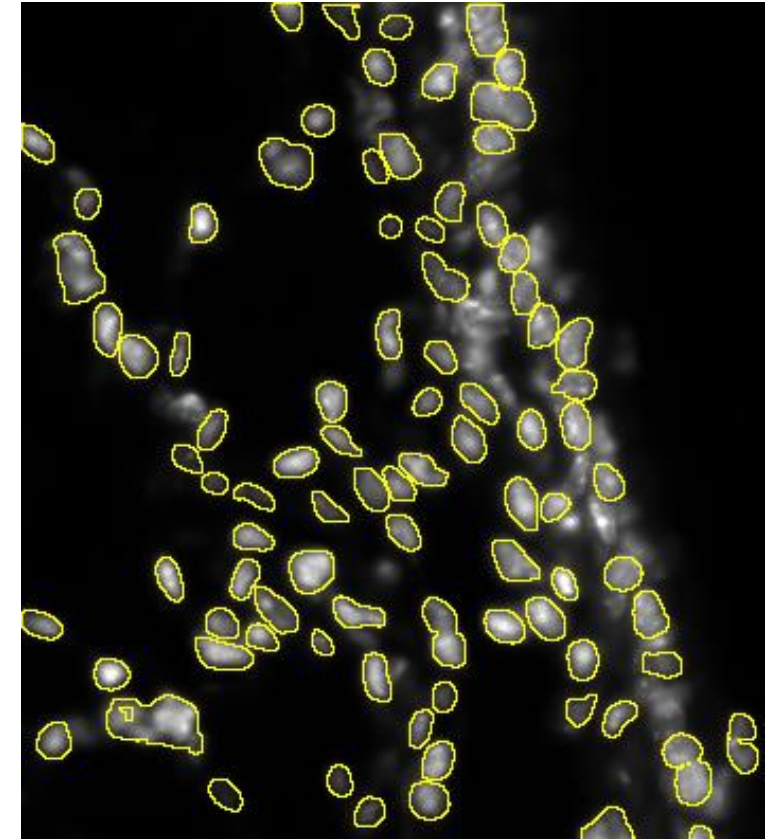
QuPath



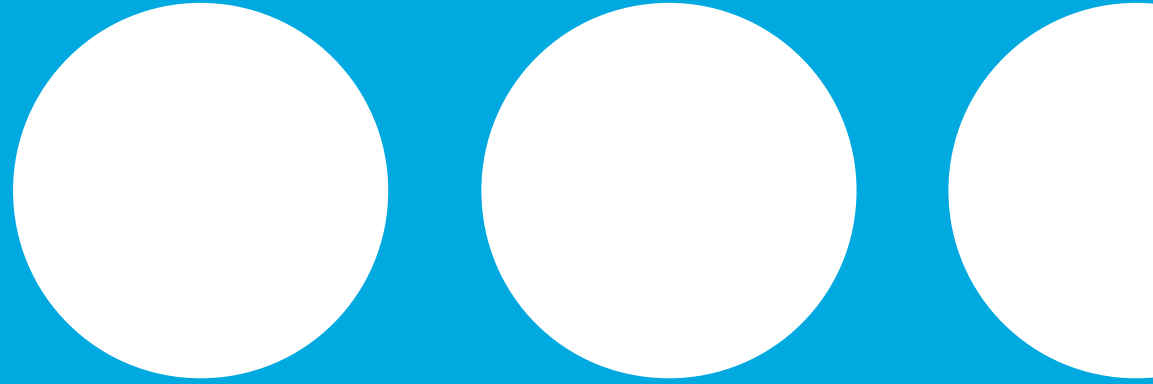
Unet



Mask RCNN



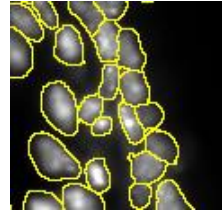
Conclusion



# Conclusion

## CNN Method:

- Better solution to overlapping cells
- Could perform for a whole image at once
  - Where threshold method fails
- Could be embedded into QuPath framework
- Fast prediction
- Stable and general outcomes
  - User independent
  - Train on various image types
- Framework for future process



# Possible Improvement

- More Data
  - Larger training data
  - Tweak by more specific data related to our project
- Validation and hyperparameters tuning – Generalized Ability
  - Stratified validation set
  - Other optimizer
  - More post-processing

( Thank you )

