

Image Masking and Cell Segmentation

BY JOSHUA PAIK

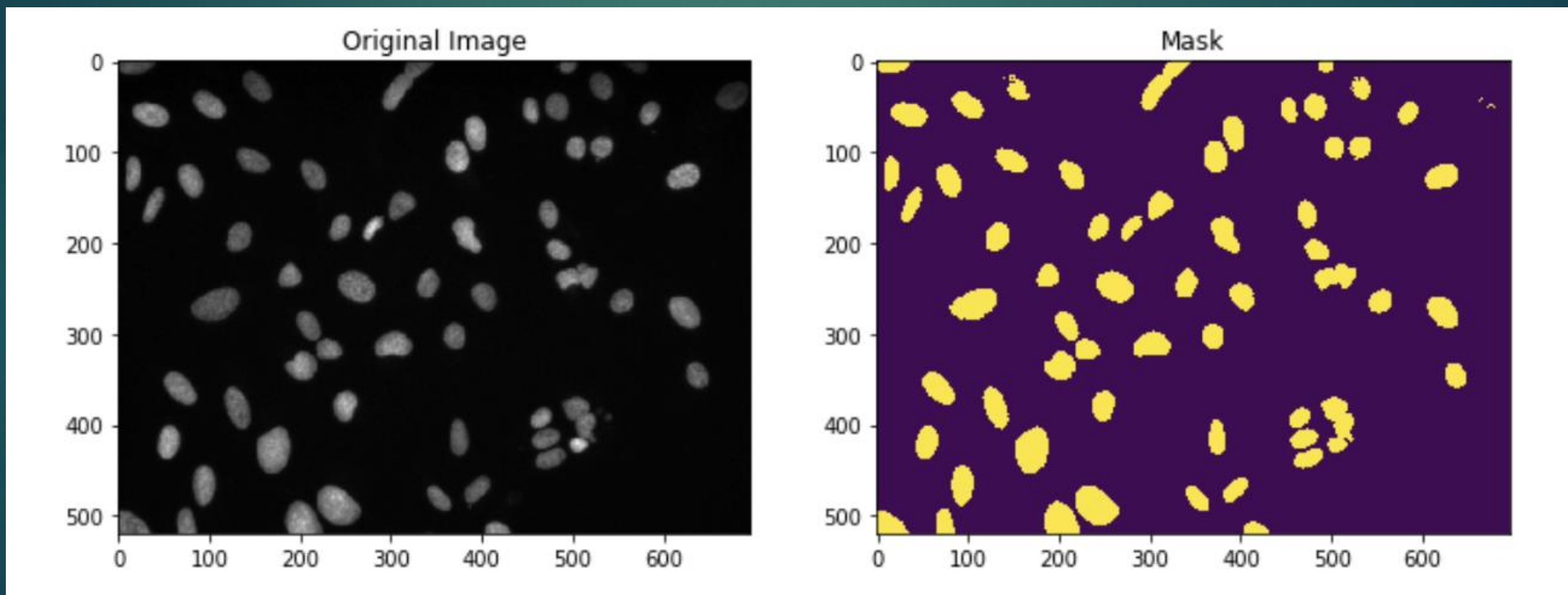
Outline of Today's Talk

- ▶ Introduction to the Kaggle Data Science Bowl 2018
- ▶ Exploratory Data Analysis
- ▶ Explanation of UNET
- ▶ Data Augmentation
- ▶ My Solution
- ▶ Winning Solutions

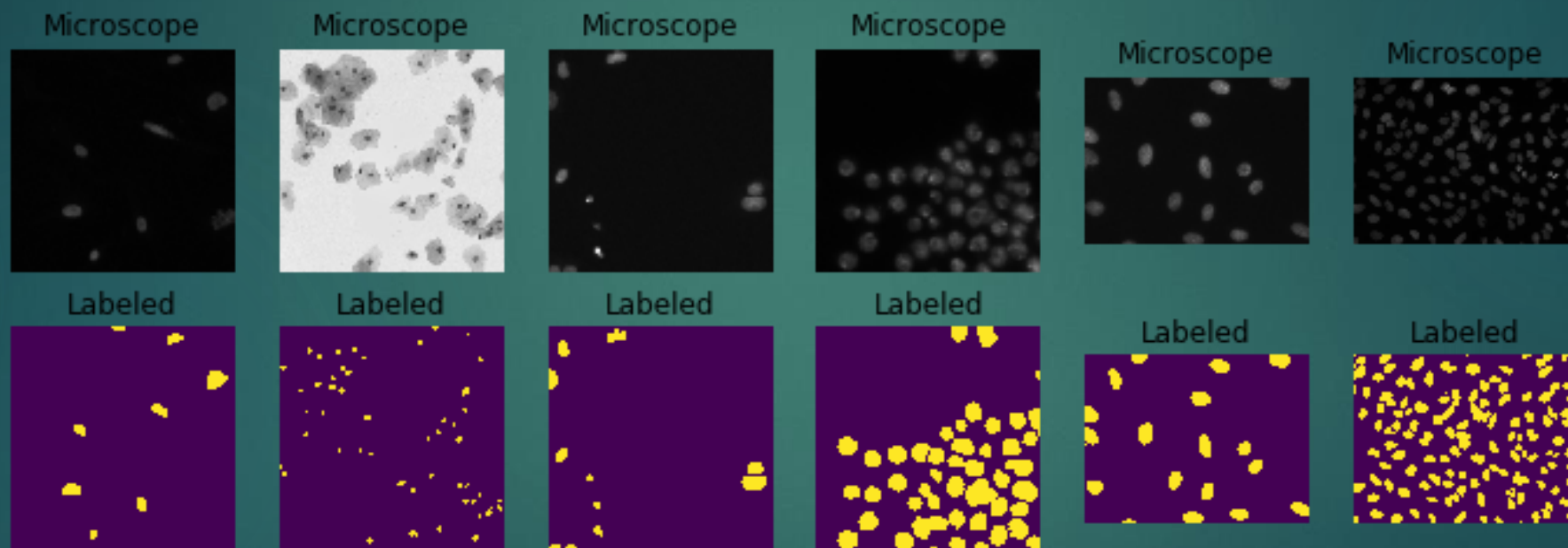
Introduction to the Kaggle Data Science Bowl 2018

- ▶ GOAL? Identify cell nuclei from microscopy data.
- ▶ WHY? To advance research. As the nuclei contains DNA, it is important to locate it in genetic and biological research.
- ▶ HOW? Use existing neural network architecture platforms and image processing techniques

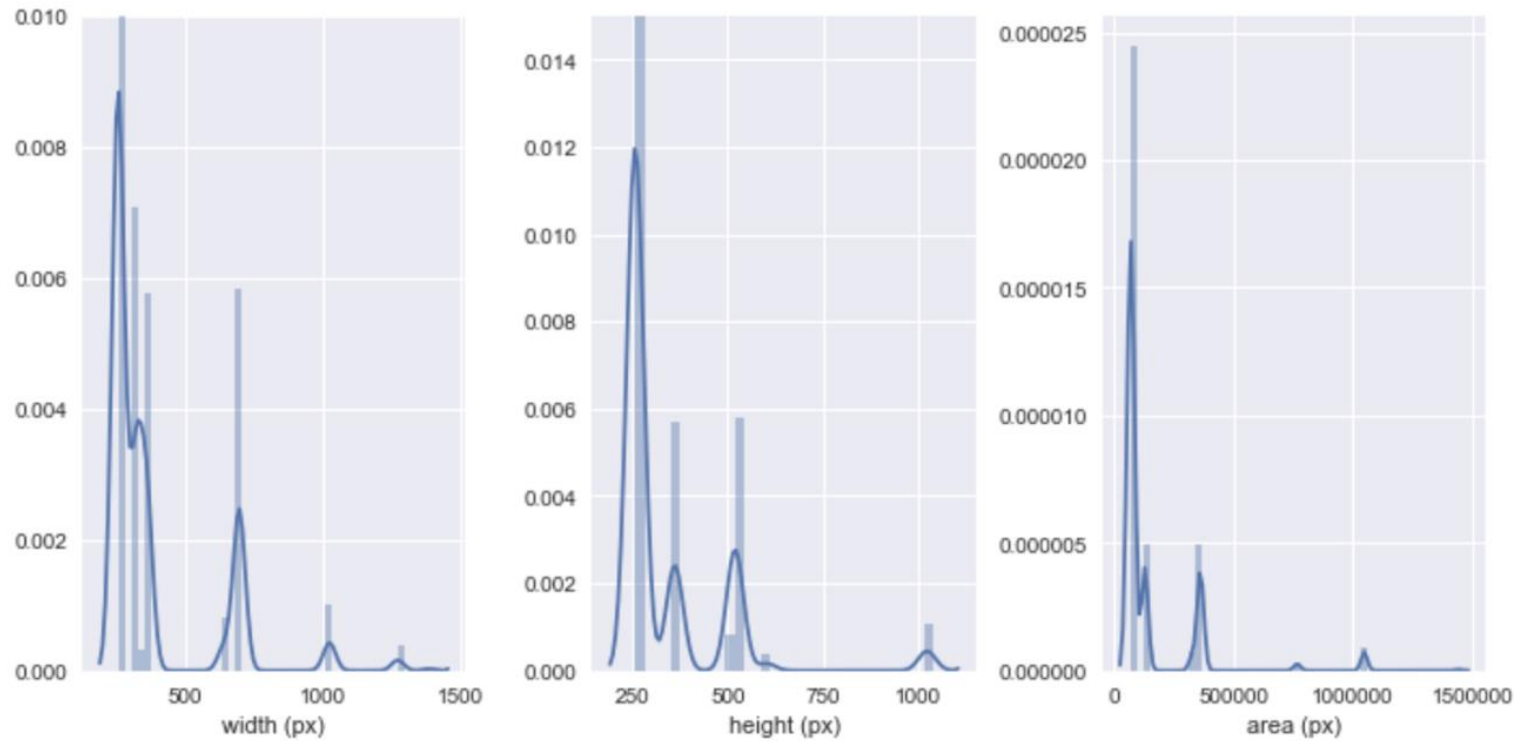
Goal: Create image masks for each cells



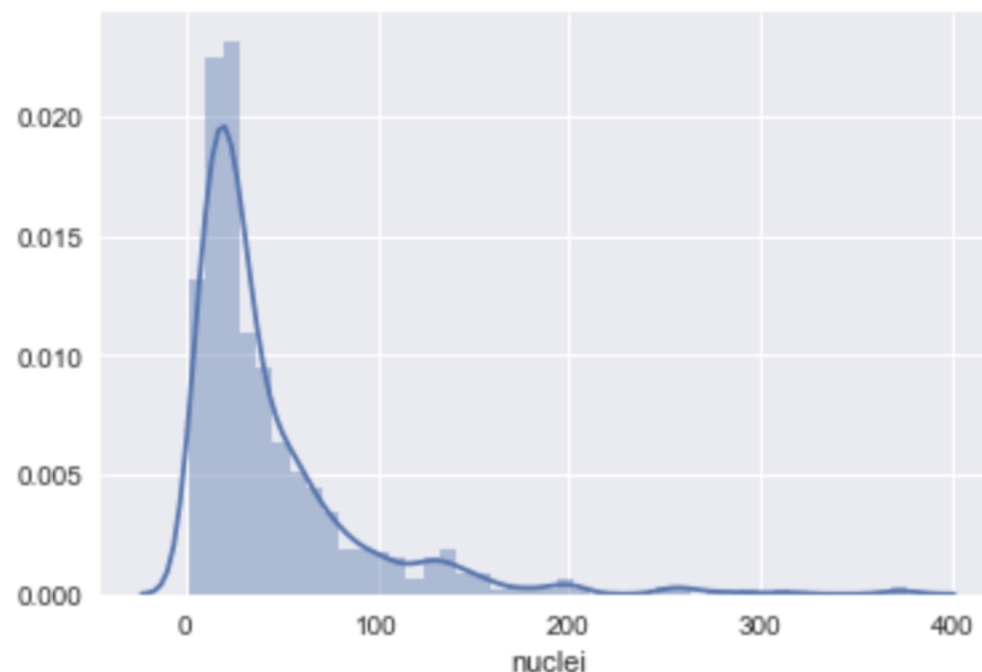
More Sample Data



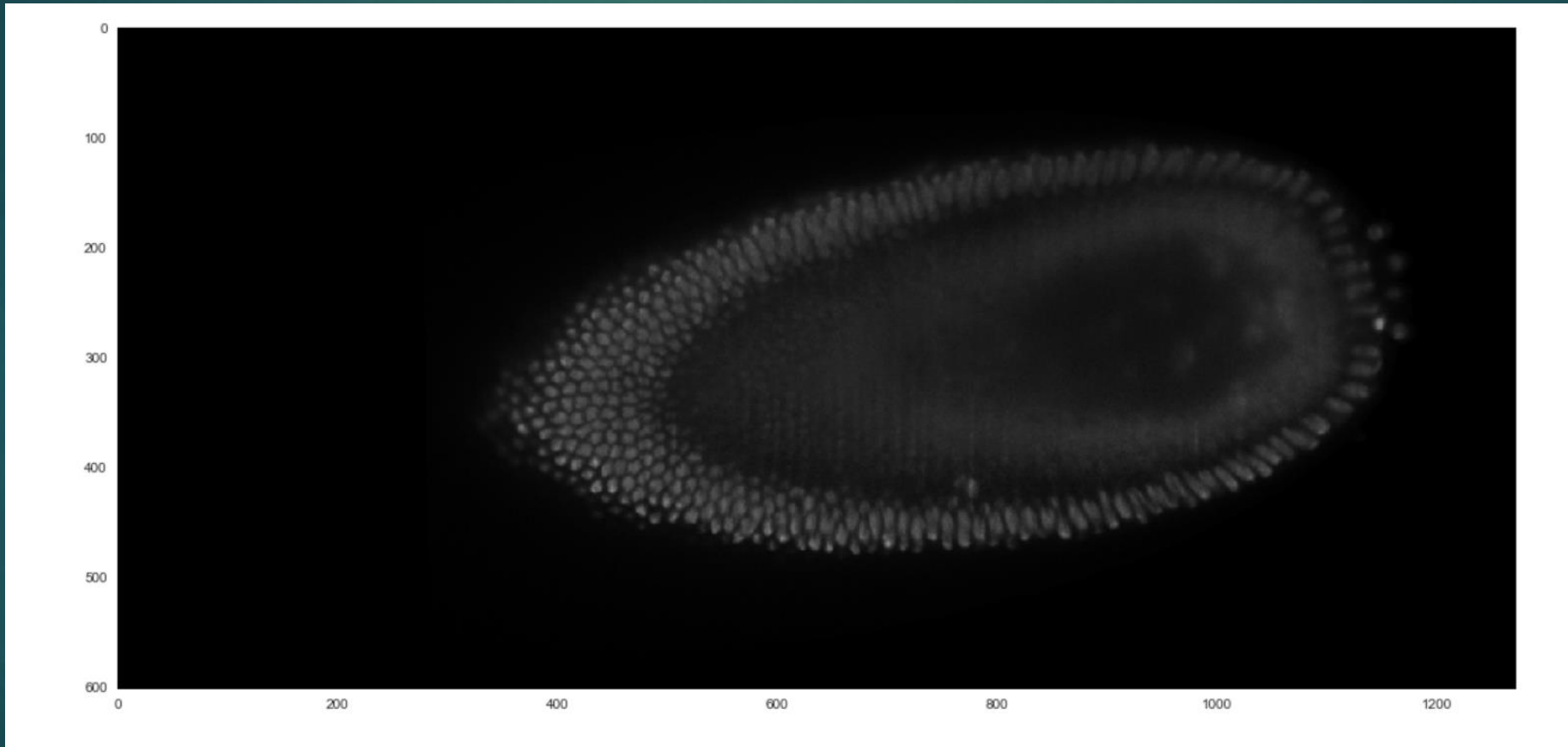
EDA: Distribution of image masks width, height, area



EDA: Distribution of number of nuclei in each images



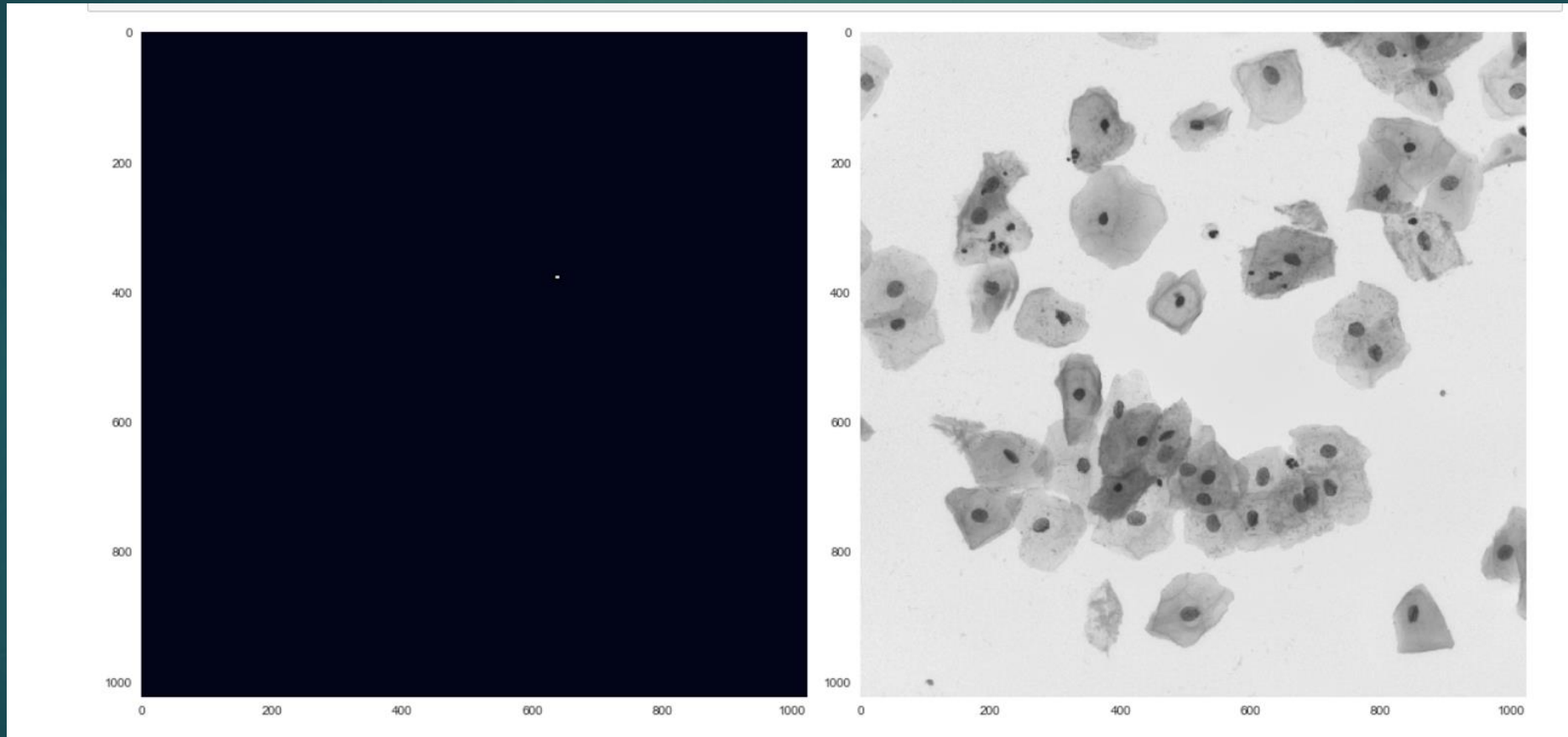
EDA: Image with most number of nuclei



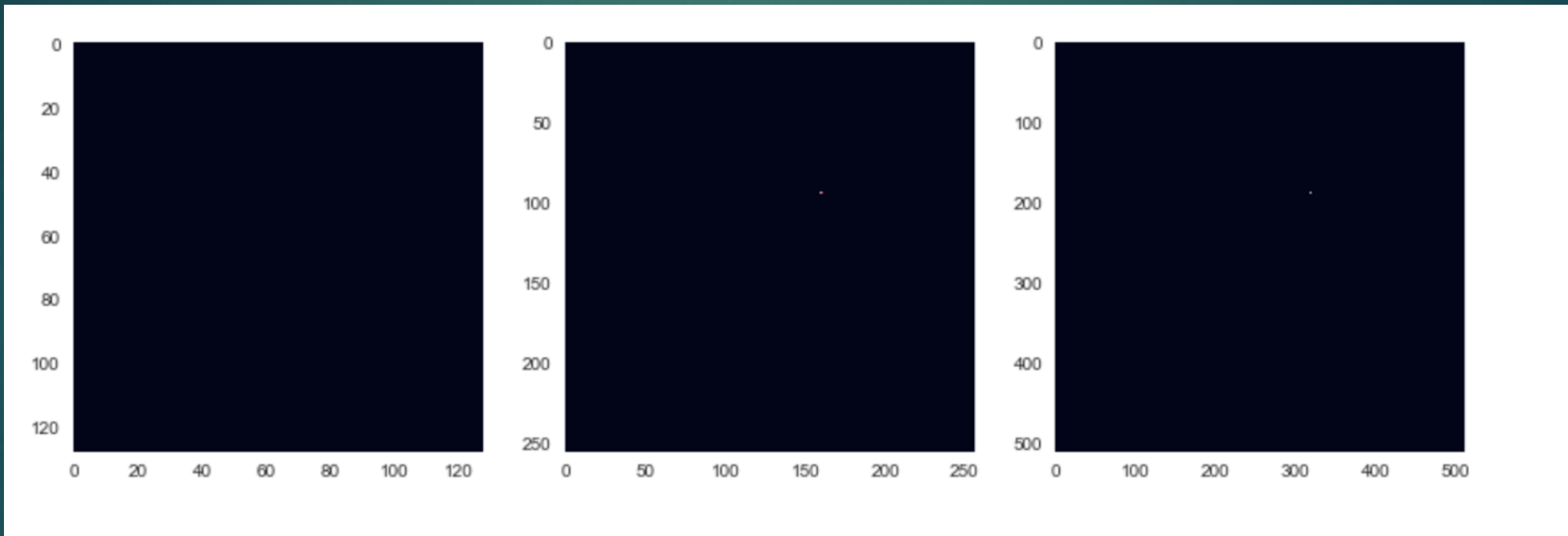
EDA: Image with Least number of nuclei



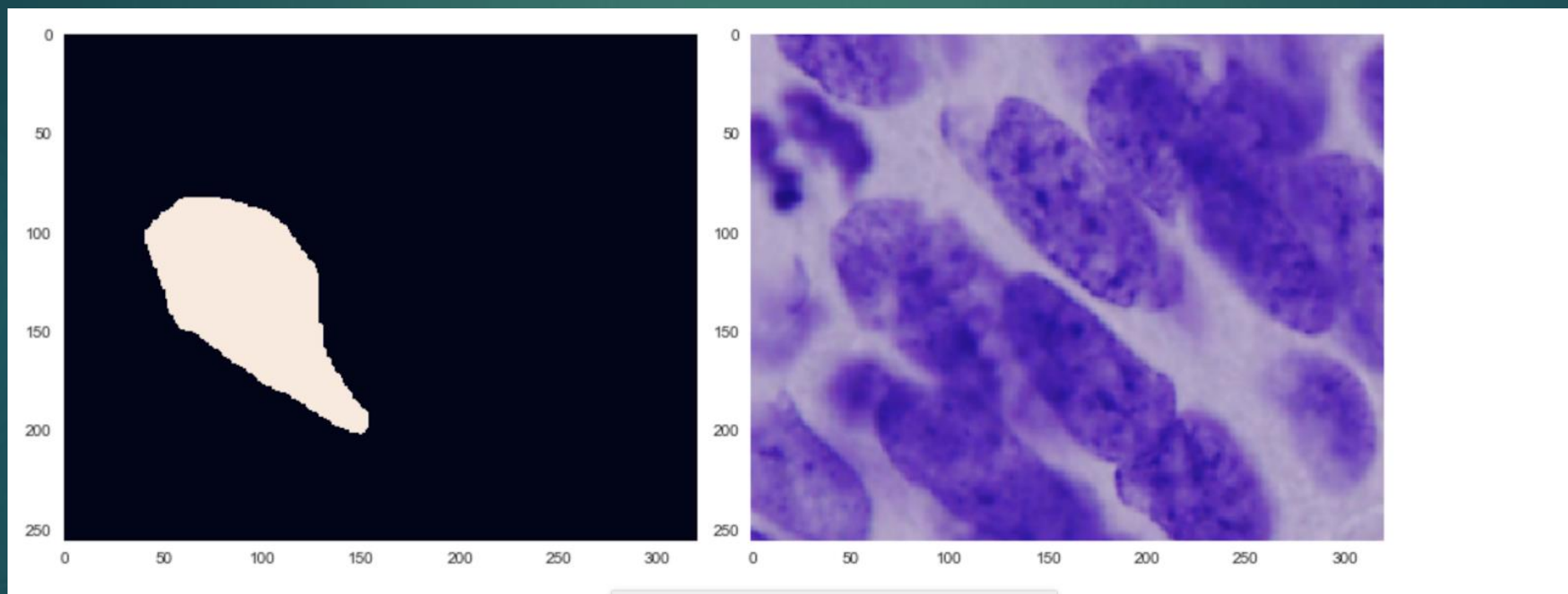
EDA: Smallest nuclei problem



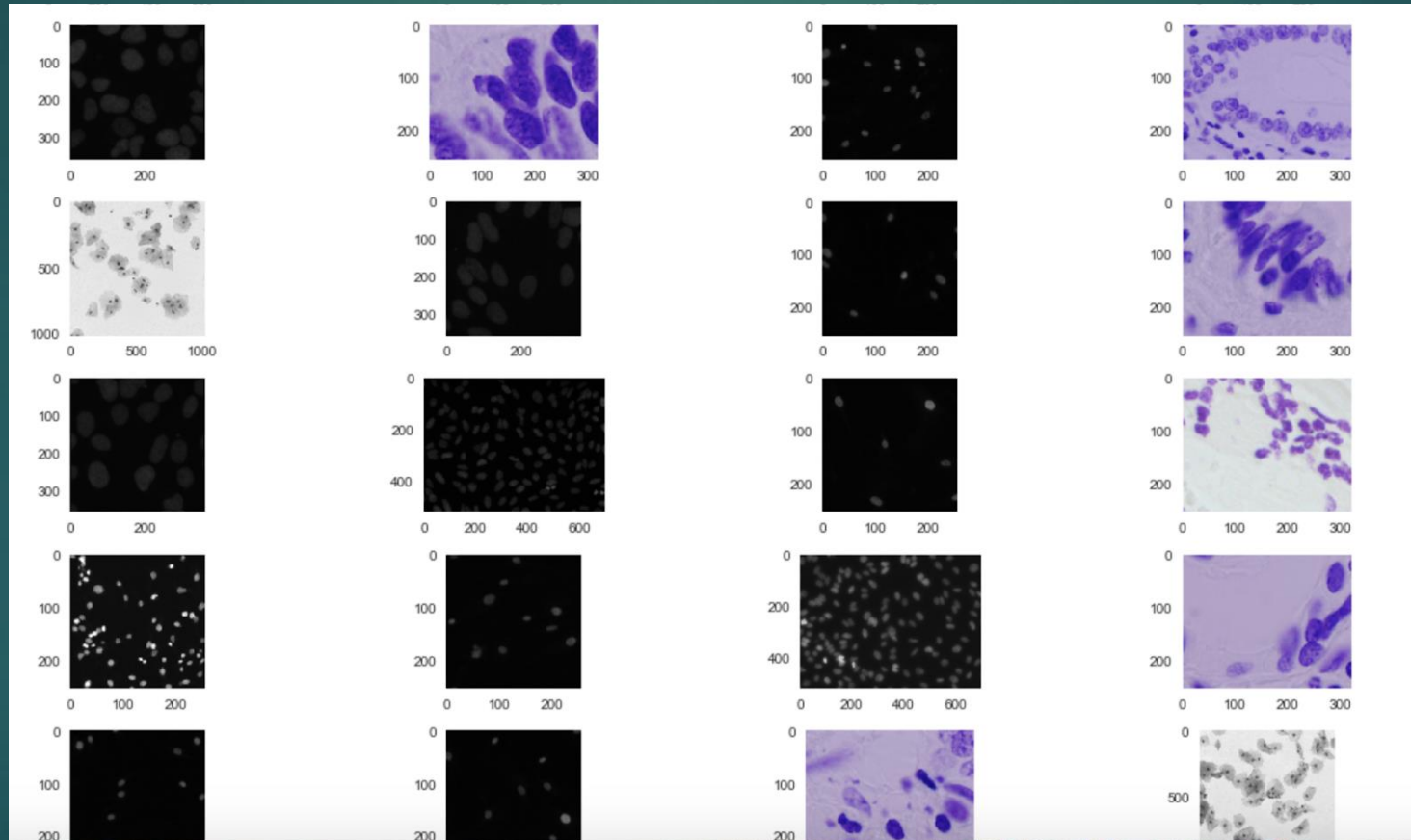
EDA: Smallest Nuclei Problem: Dependent on Scaling



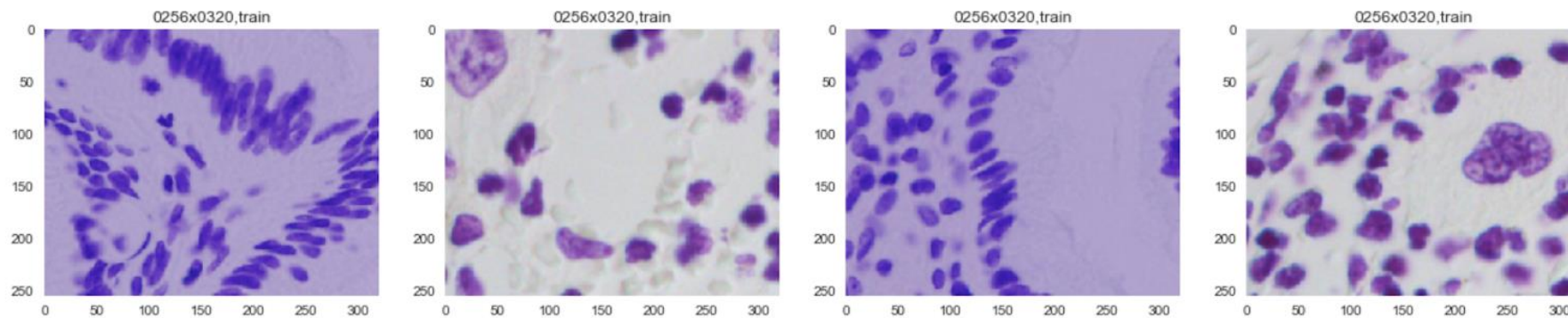
EDA: Overlapping cell problem



There are a lot of overlapping cells

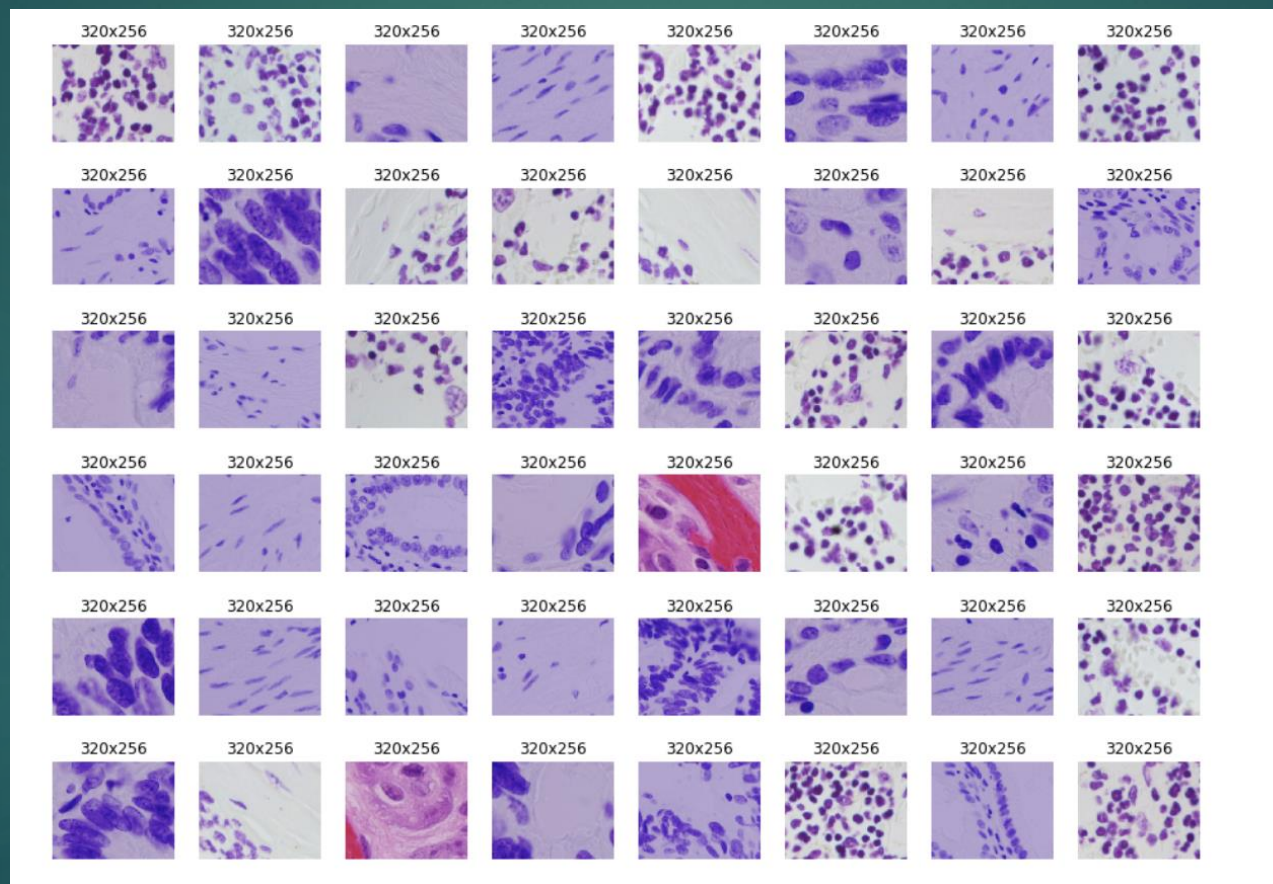


More Overlapping Cells



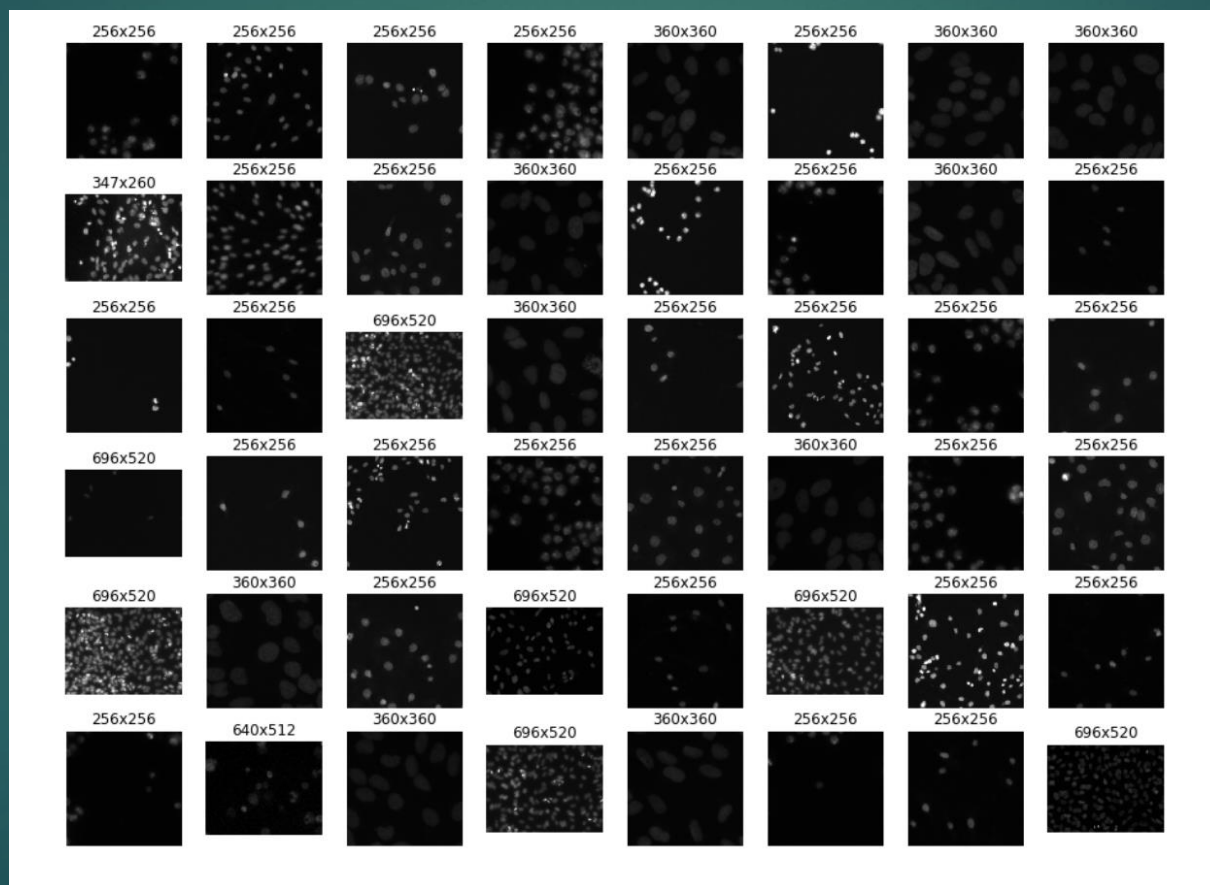
EDA: K Means Classification

Fluorescent: 81.5%



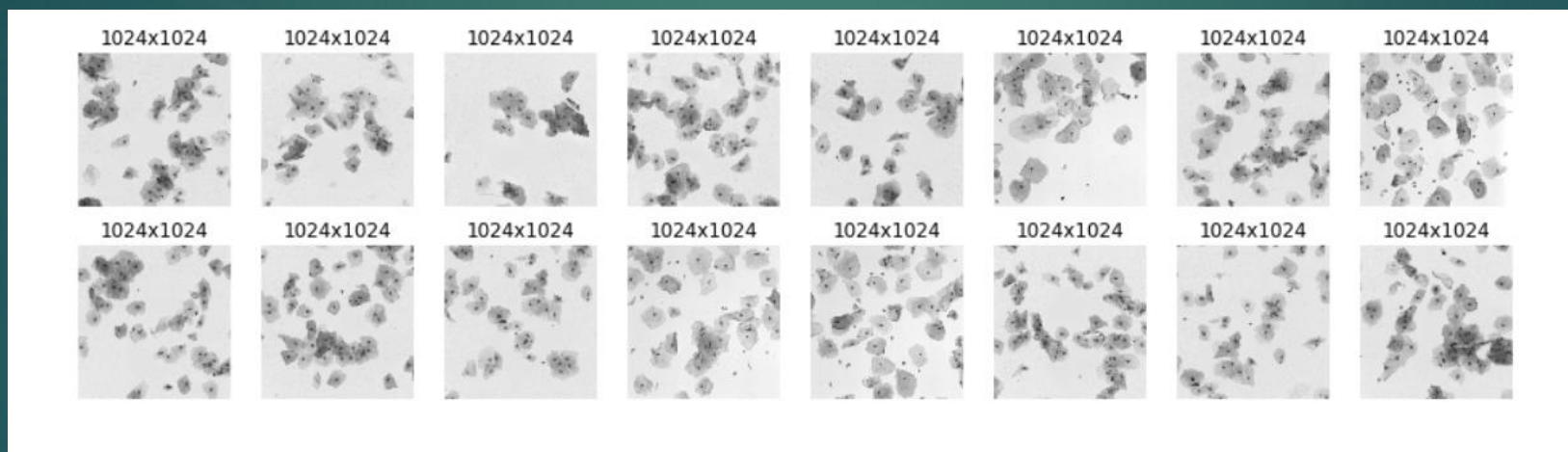
EDA: K Means Classification

Histological: 16.1%



EDA: K Means Classification

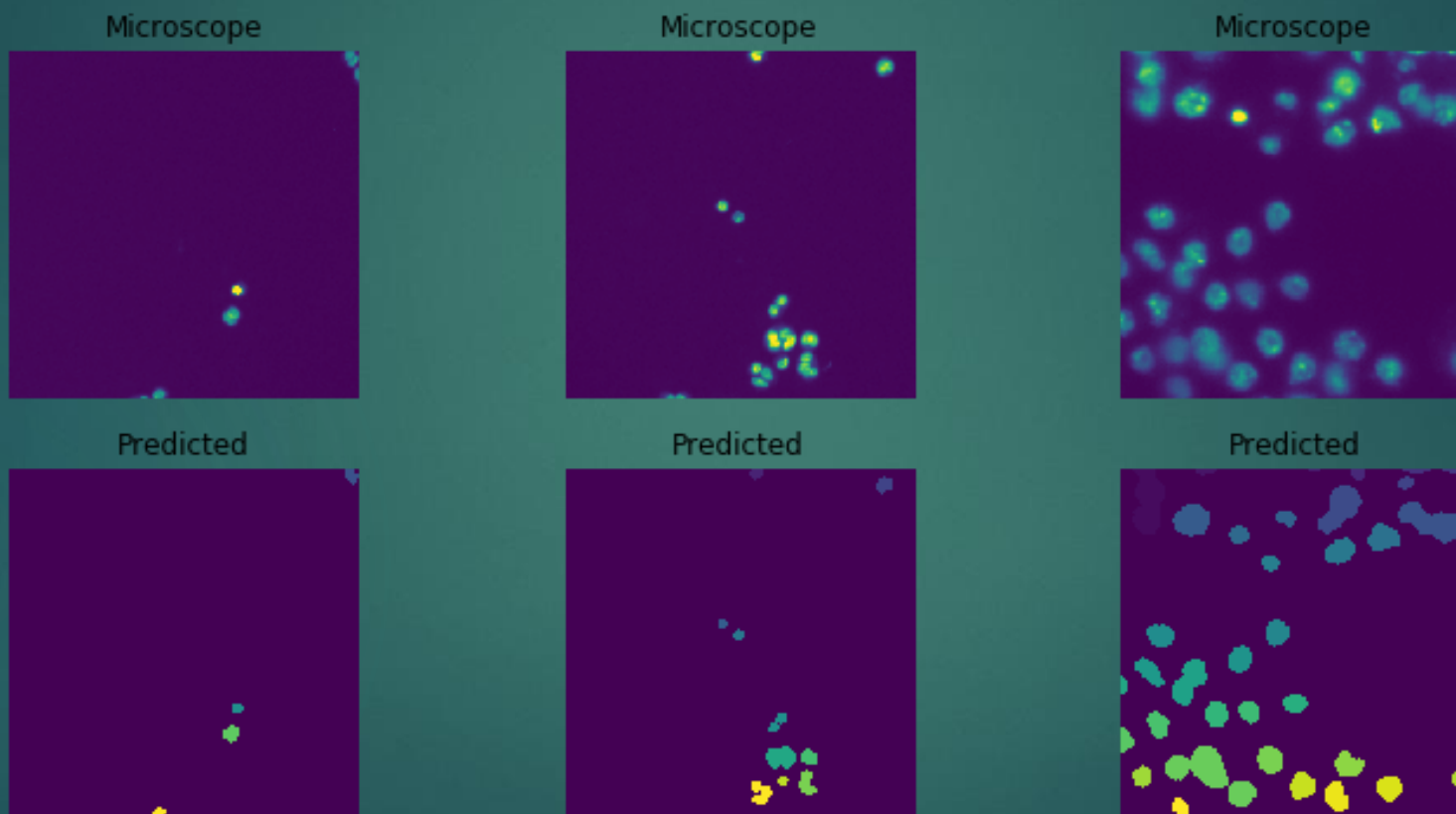
Bright-field: 2.4%



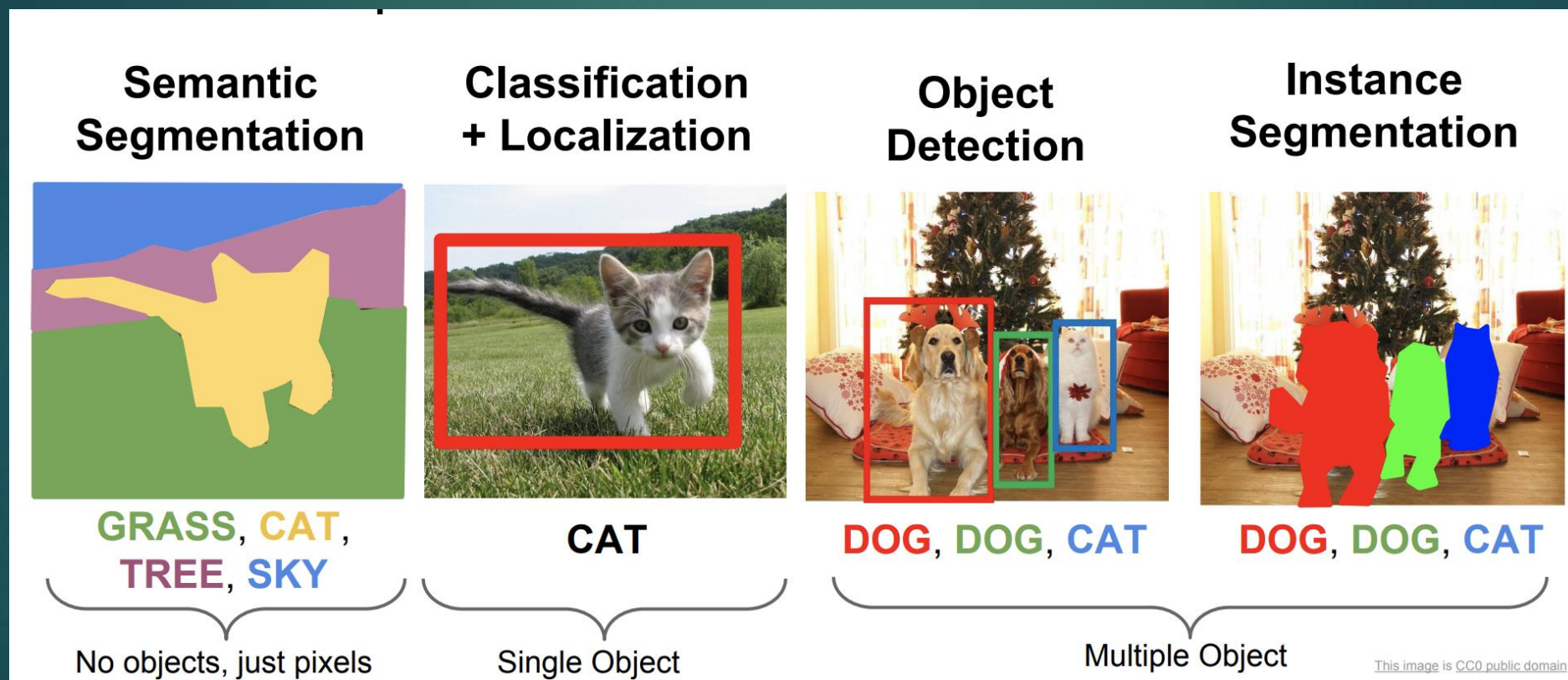
```
def parametric_pipeline(img_green,
                        invert_thresh_pd = 10,
                        circle_size_x = 7,
                        circle_size_y = 7,
                        ):
    circle_size_x = np.clip(int(circle_size_x), 1, 30)
    circle_size_y = np.clip(int(circle_size_y), 1, 30)

    #green channel happens to produce slightly better results
    #than the grayscale image and other channels
    #morphological opening (size tuned on training data)
    circle7=cv2.getStructuringElement(cv2.MORPH_ELLIPSE,(circle_size_x, circle_size_y))
    img_open=cv2.morphologyEx(img_green, cv2.MORPH_OPEN, circle7)
    #Otsu thresholding
    img_th=cv2.threshold(img_open,0,255,cv2.THRESH_OTSU)[1]
    #Invert the image in case the objects of interest are in the dark side
    if(np.sum(img_th==255)>((invert_thresh_pd/10.0)*np.sum(img_th==0))):
        img_th=cv2.bitwise_not(img_th)
    #second morphological opening (on binary image this time)
    bin_open=cv2.morphologyEx(img_th, cv2.MORPH_OPEN, circle7)
    #connected components
    cc=cv2.connectedComponents(bin_open)[1]
    #cc=segment_on_dt(bin_open,20)
    return cc
```

Non Machine Learning Approach

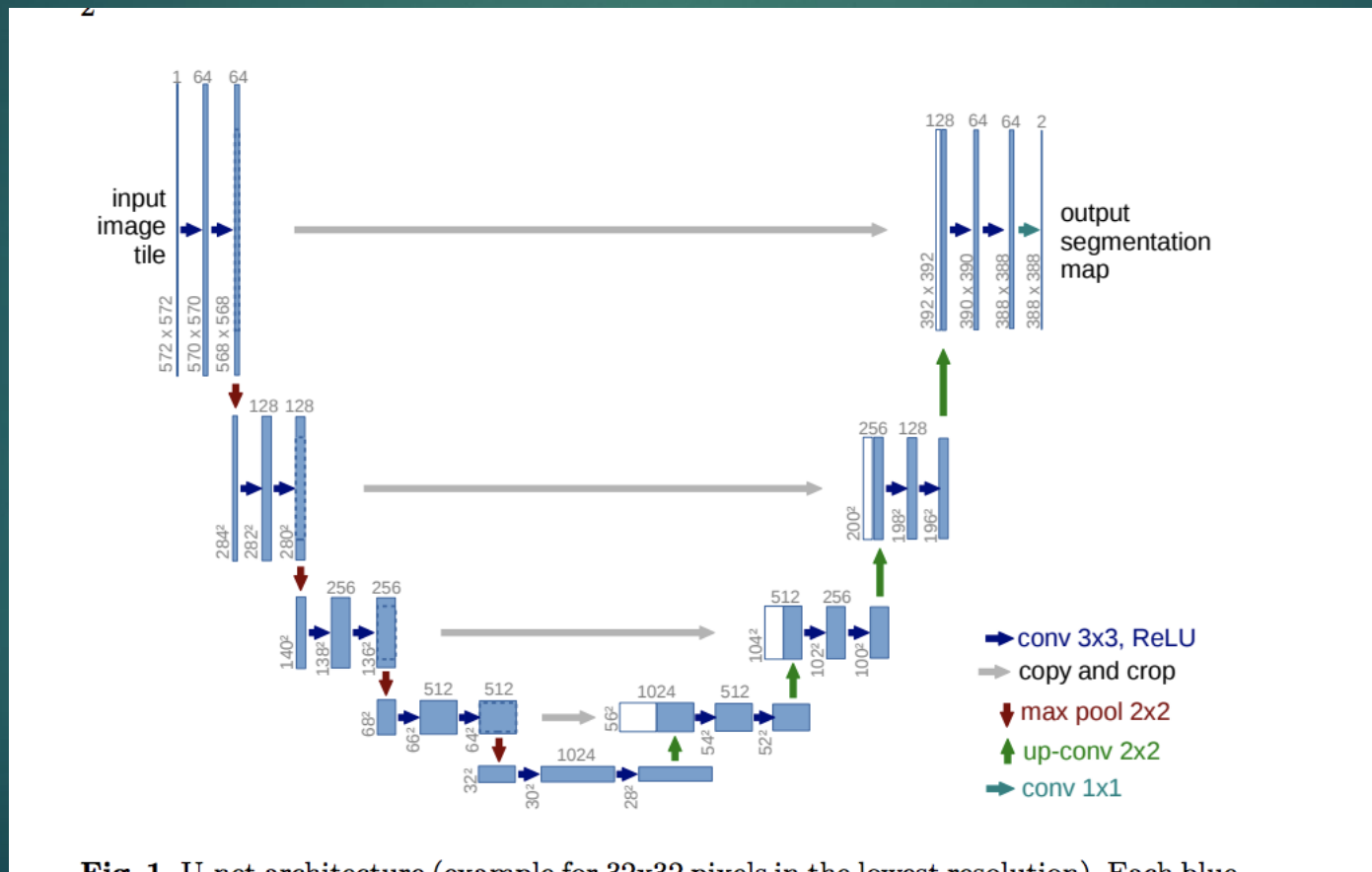


What type of problem is the Kaggle DSBowl

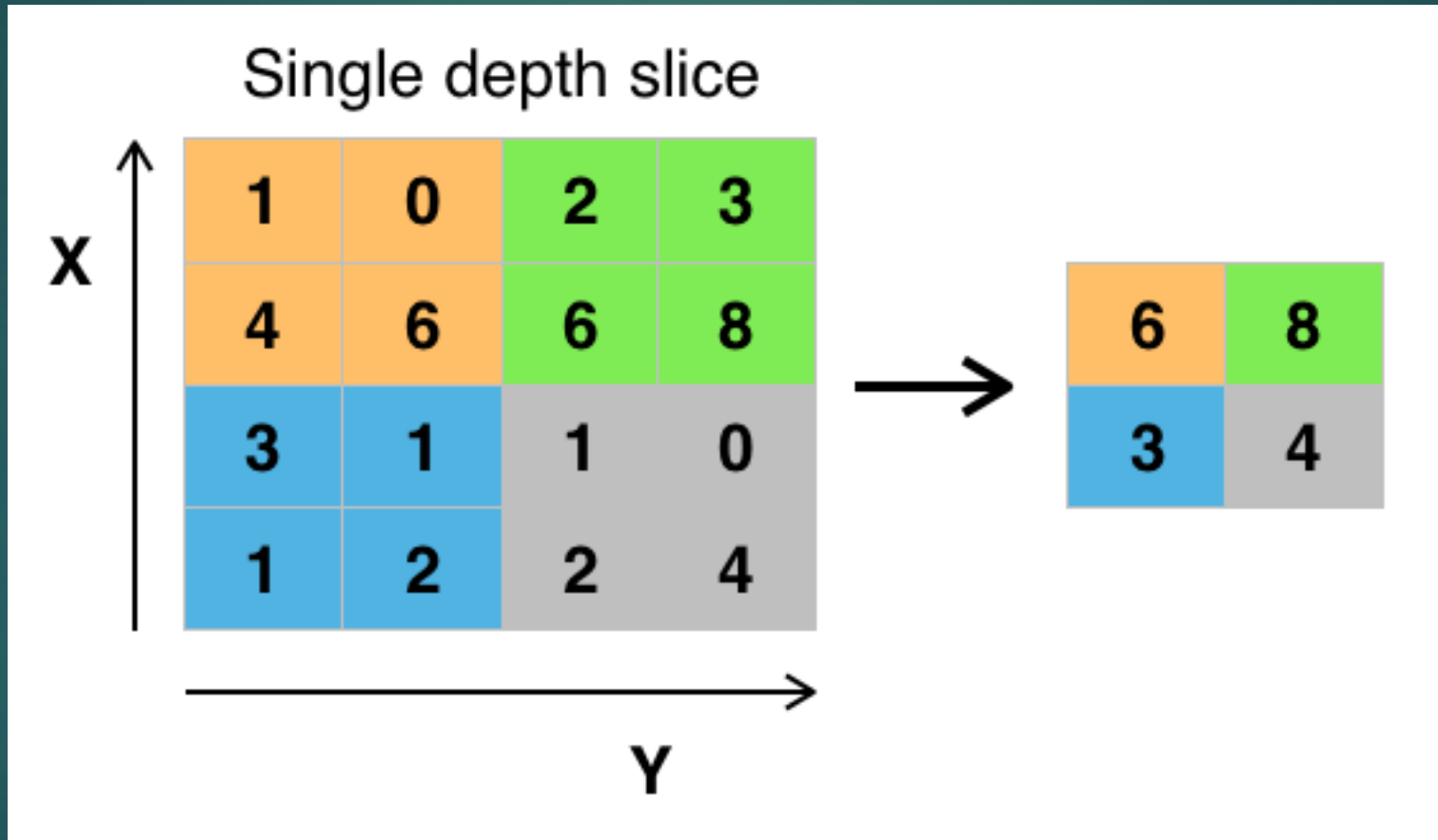


Explanation of UNET

<https://arxiv.org/pdf/1505.04597.pdf>

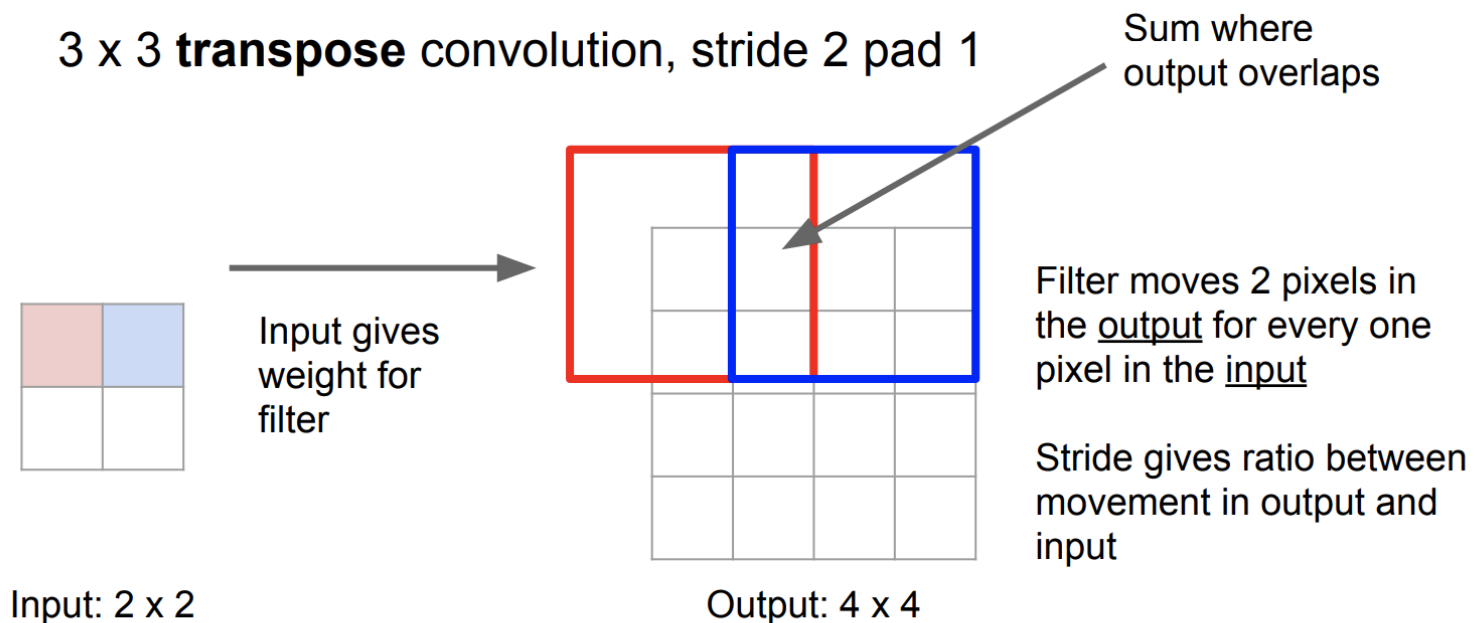


Max Pool

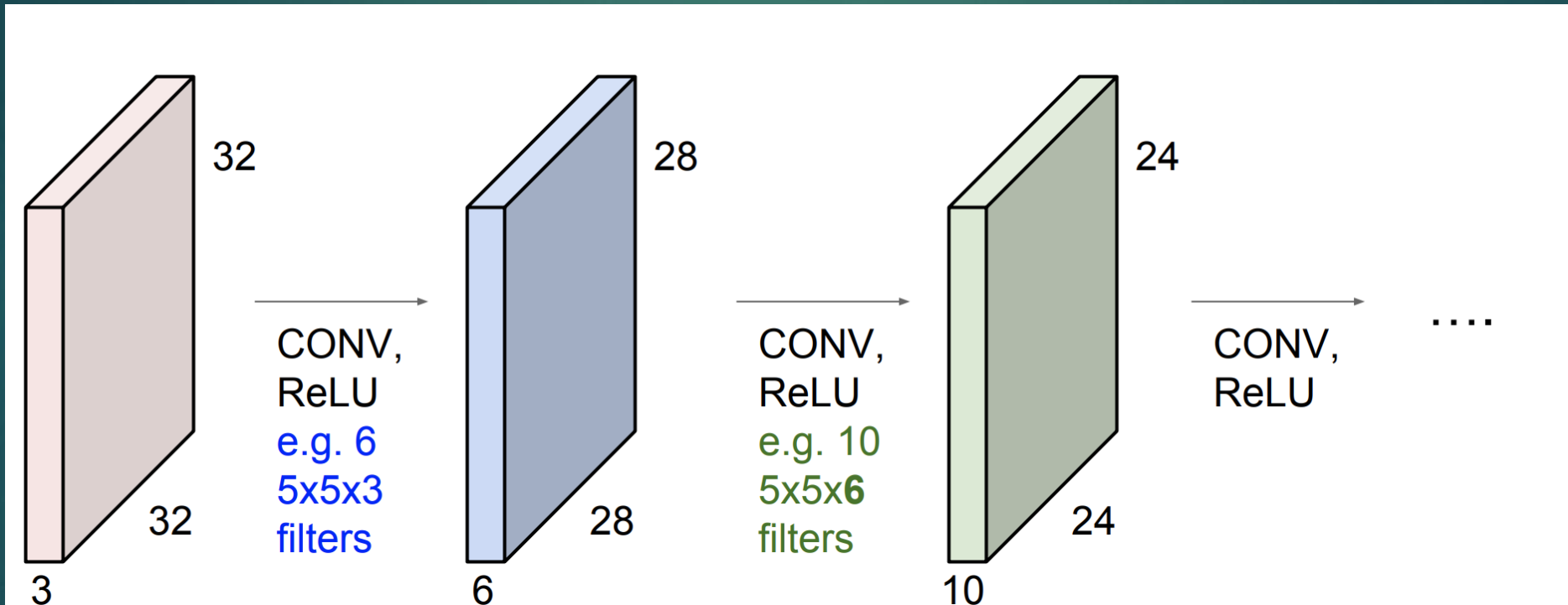


Up Convolution or Transpose Convolution

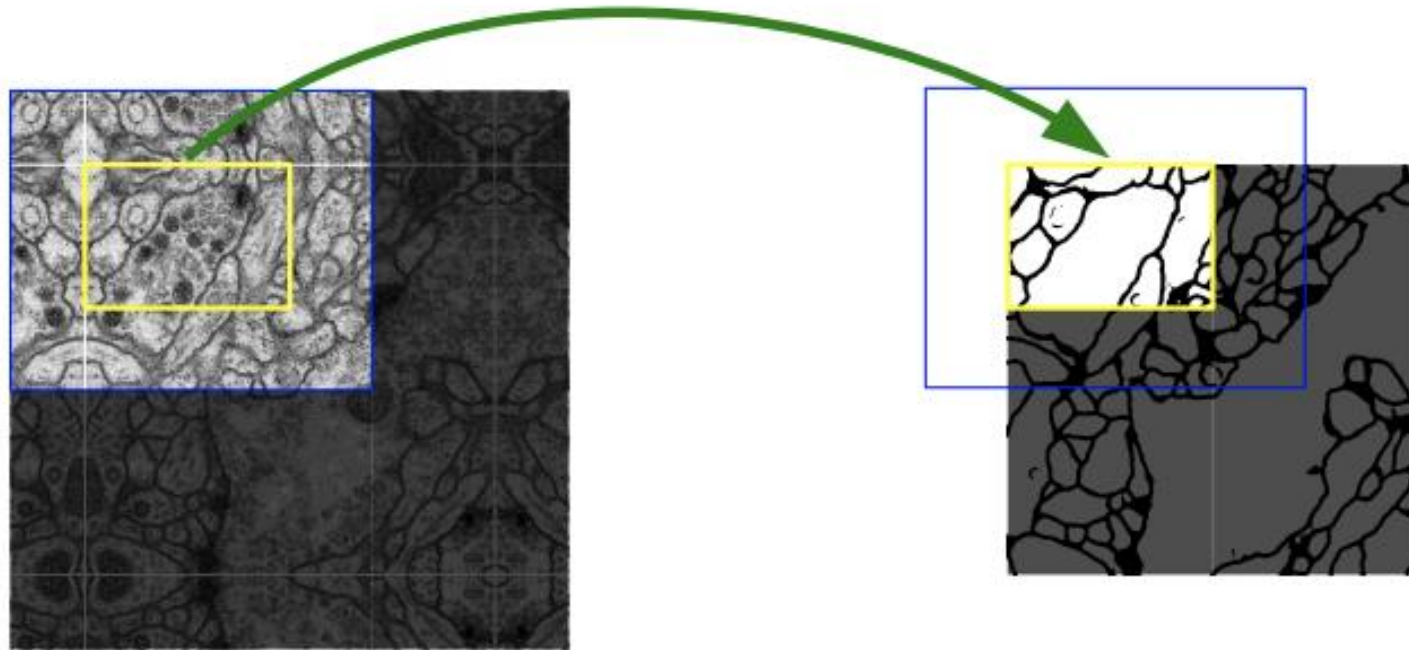
Learnable Upsampling: Transpose Convolution



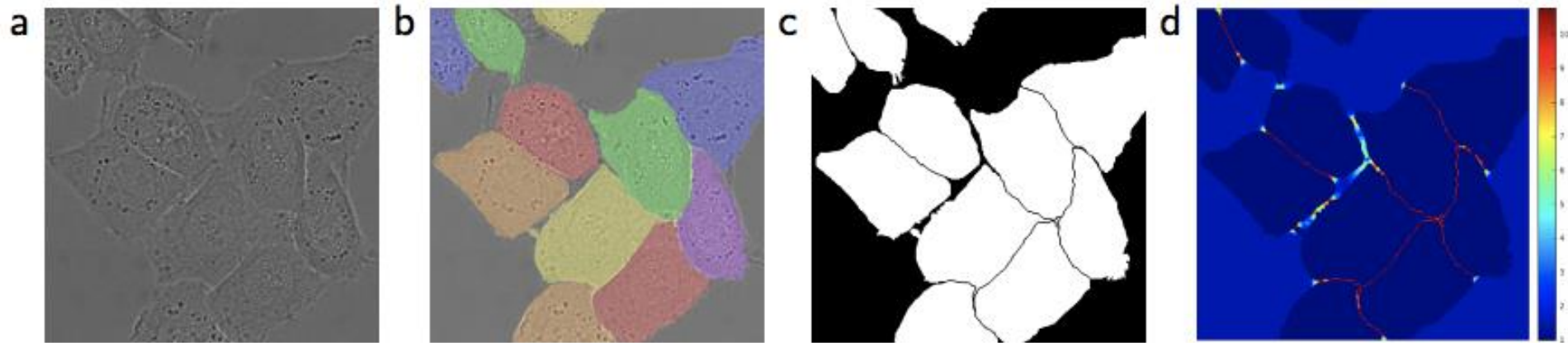
Conv 3x3 ReLU



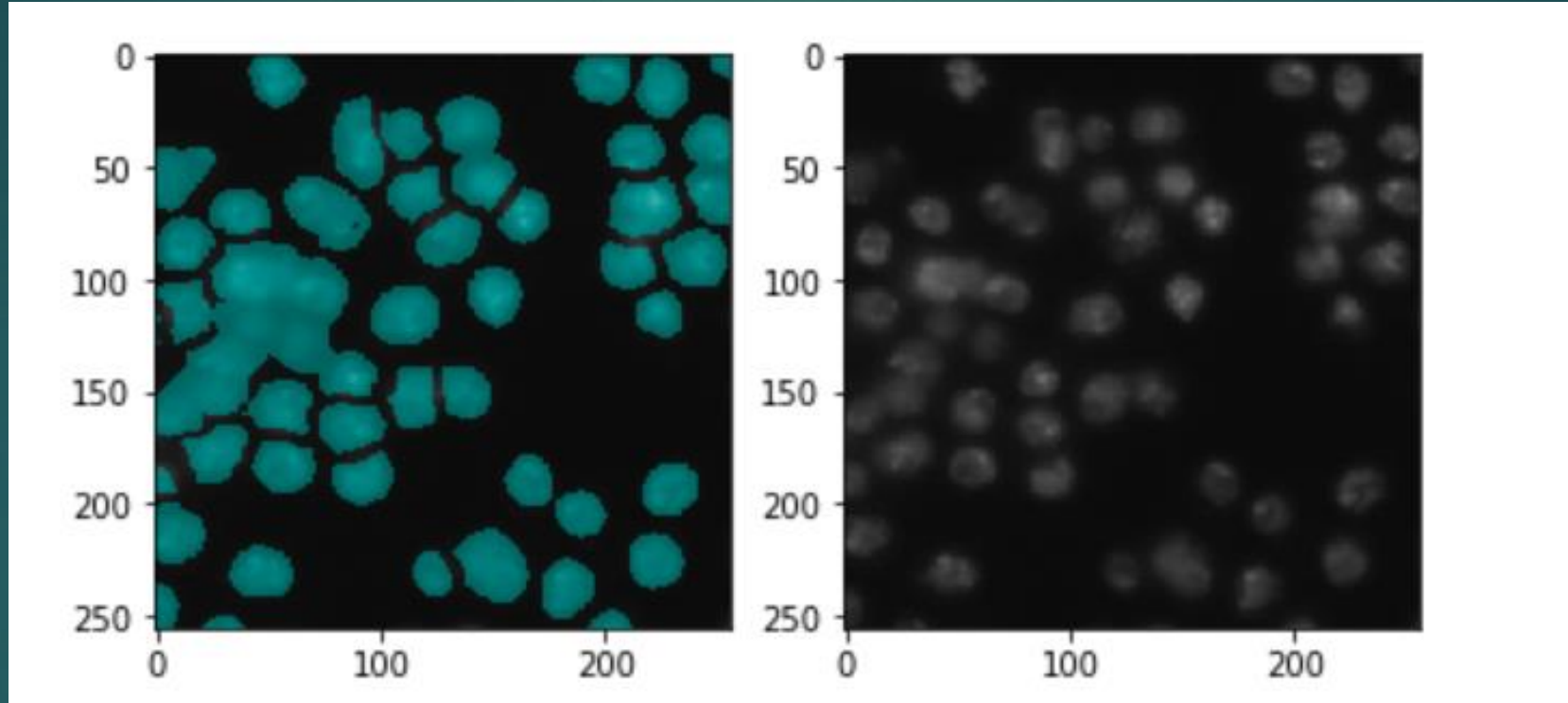
UNET - Overlapping Tile Strategy



UNET – Accuracy and Method

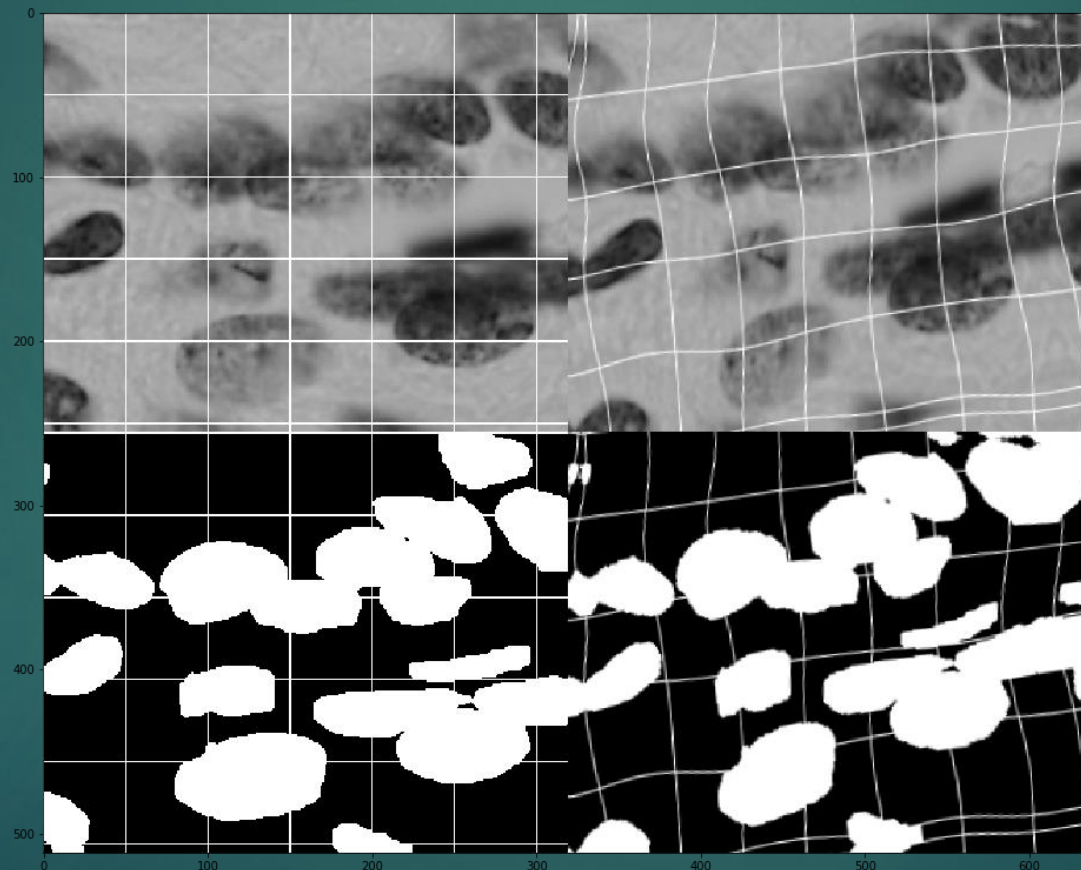


First Implementation



Pre Process: Data Augmentation

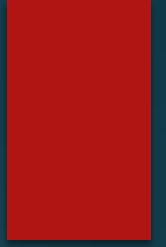
Elastic Deformations



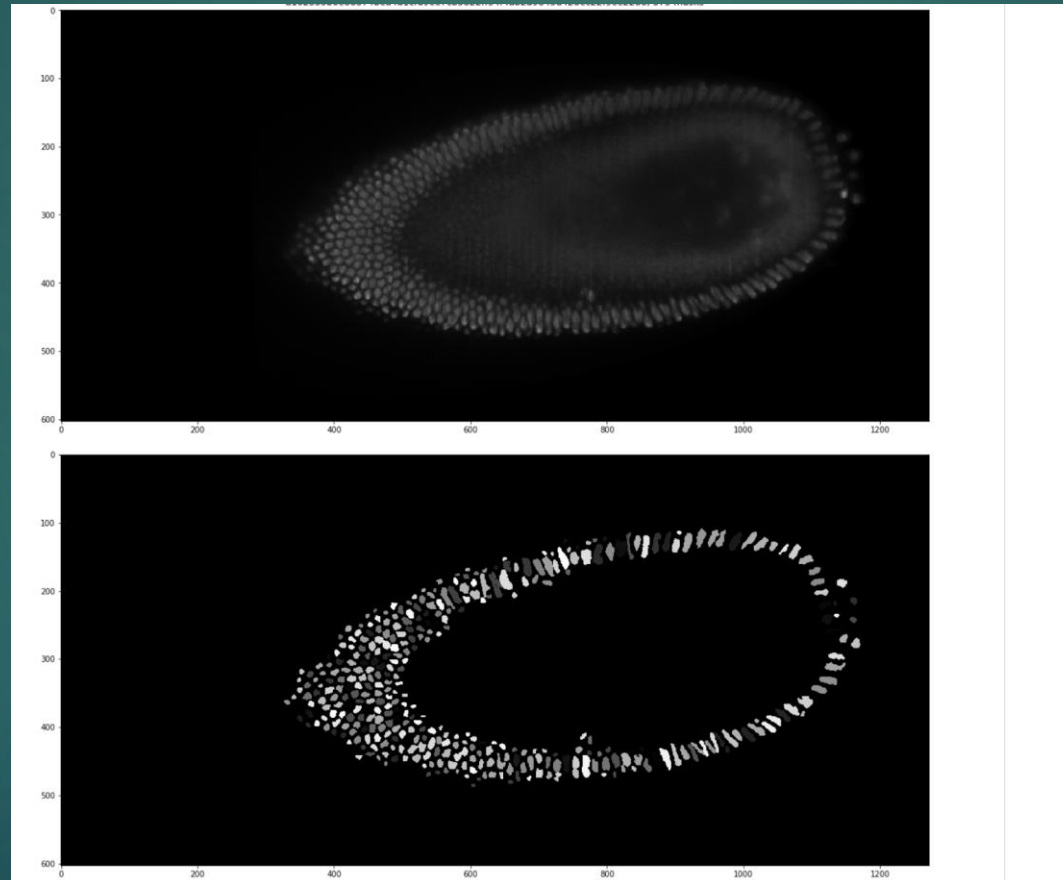
What did the winners do?

- ▶ Clache, Sharpen, Emboss
- ▶ Gaussian Noise
- ▶ Color to Gray
- ▶ Inverting
- ▶ Remapping grayscale images to random color images
- ▶ Blur, Median Blur, Motion Blur
- ▶ contrast and brightness
- ▶ random scale, rotates and flips
- ▶ Heavy geometric transformations: Elastic Transform, Perspective Transform, Piecewise Affine transforms, pincushion distortion
- ▶ Random HSV
- ▶ Channel shuffle - I guess this one was very important due to the nature of the data
- ▶ Nucleus copying on images. That created a lot of overlapping nuclei. It seemed to help networks to learn better borders for overlapping nuclei.
- ▶

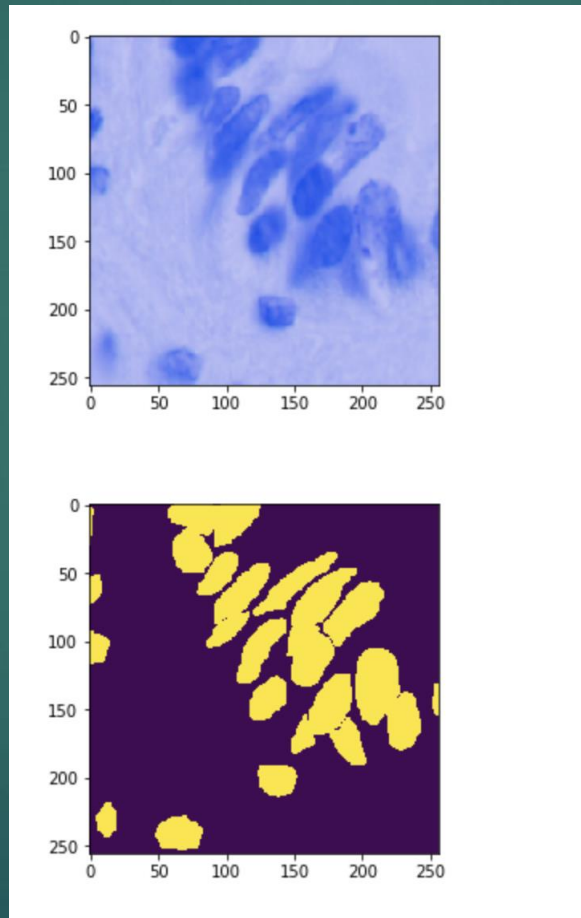
Post Processing: Watershed Algorithm



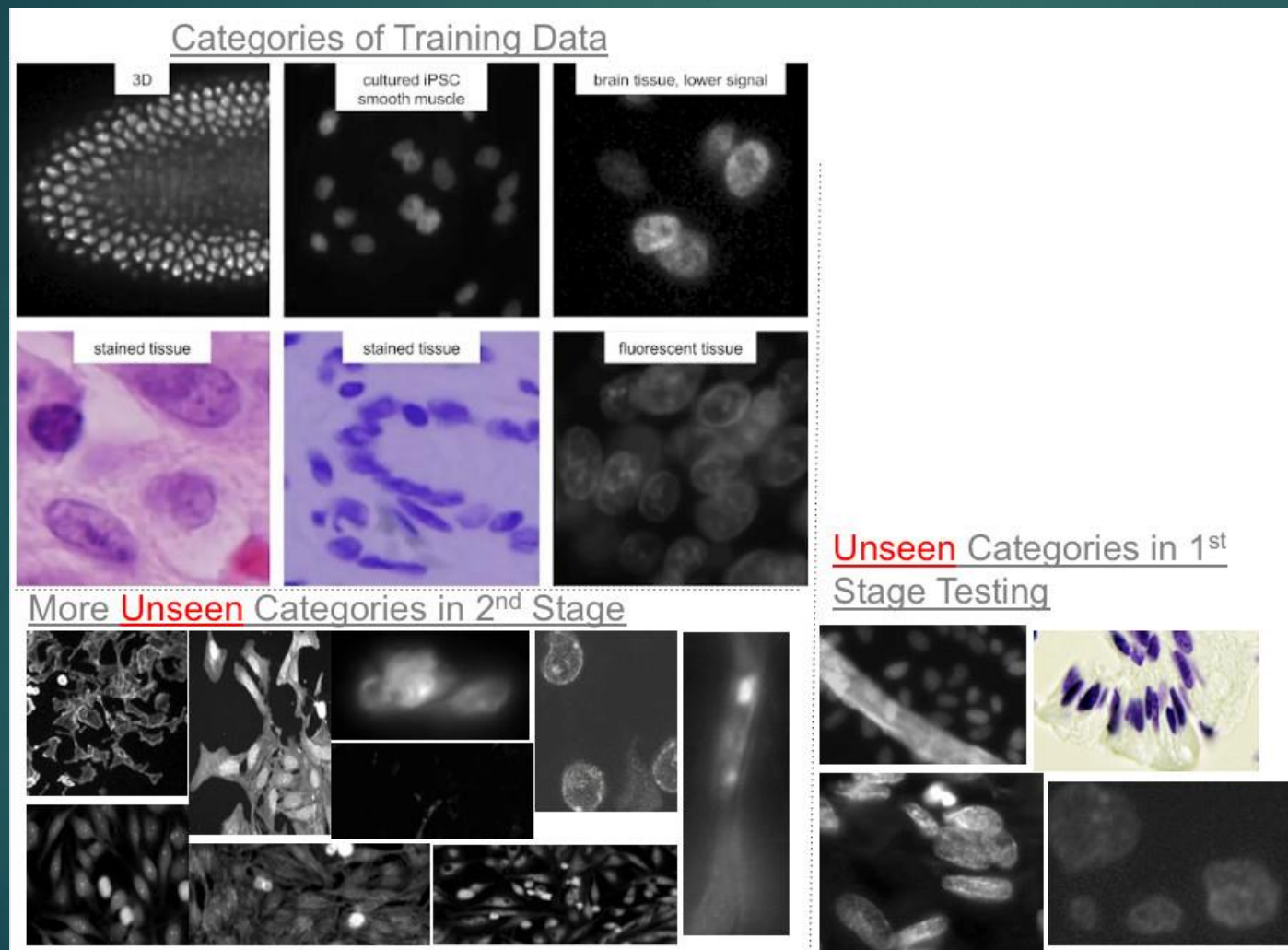
Analysis of Results



Analysis of Results



Data Variation in Stage 2!!!



Video of predictions

- ▶ <https://drive.google.com/open?id=1ZdNzQOTR83vrqNPeynUYueIWF-RxJ-35i>