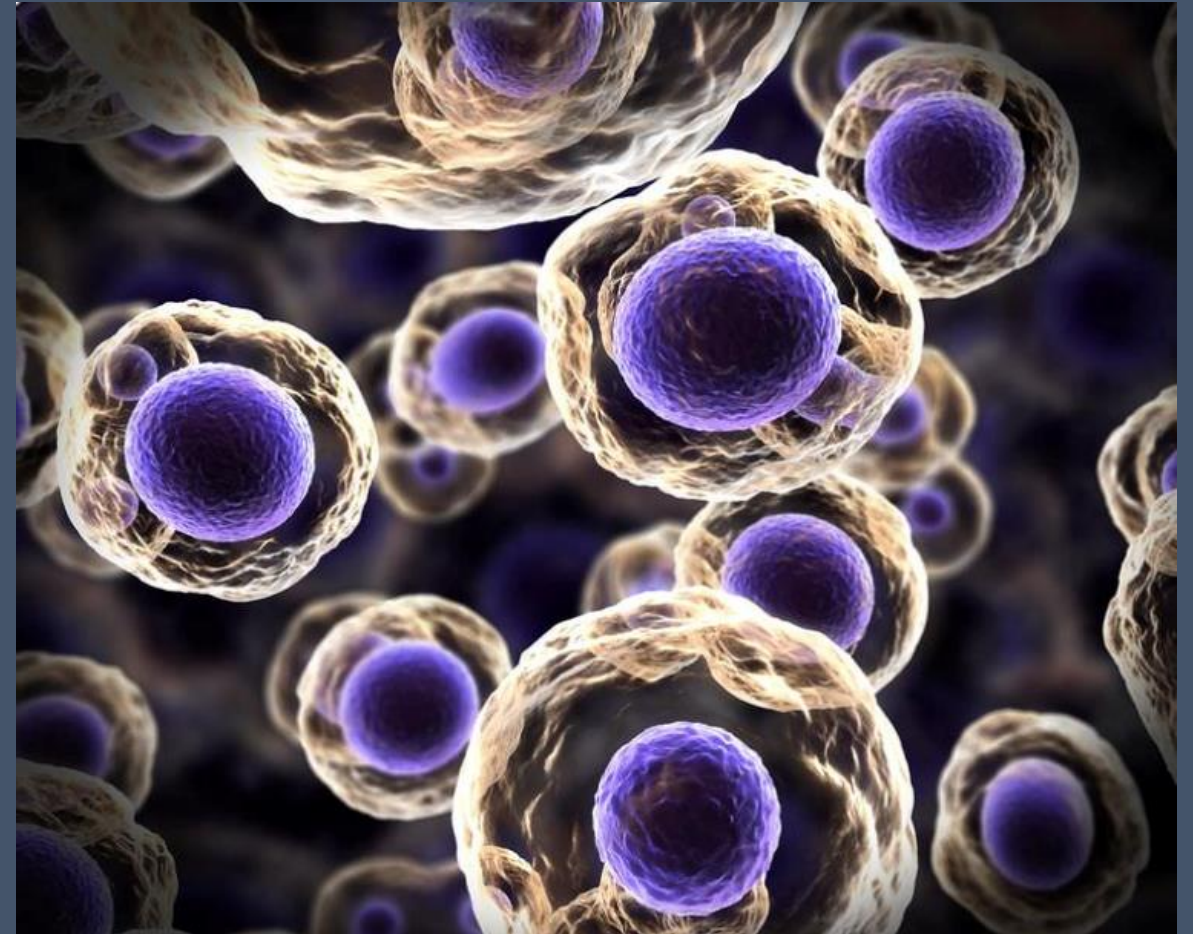


Implementation and evaluation of Otsu's thresholding

Final presentation

Elizaveta Chernova, Veronika Schuler,
Laura Wächter, Hannah L. Winter

21.07.2021



Cell nuclei segmentation

Outline

01

Otsu's
Thresholding



03

Preprocessing



05

Cell counting



02

Evaluation methods

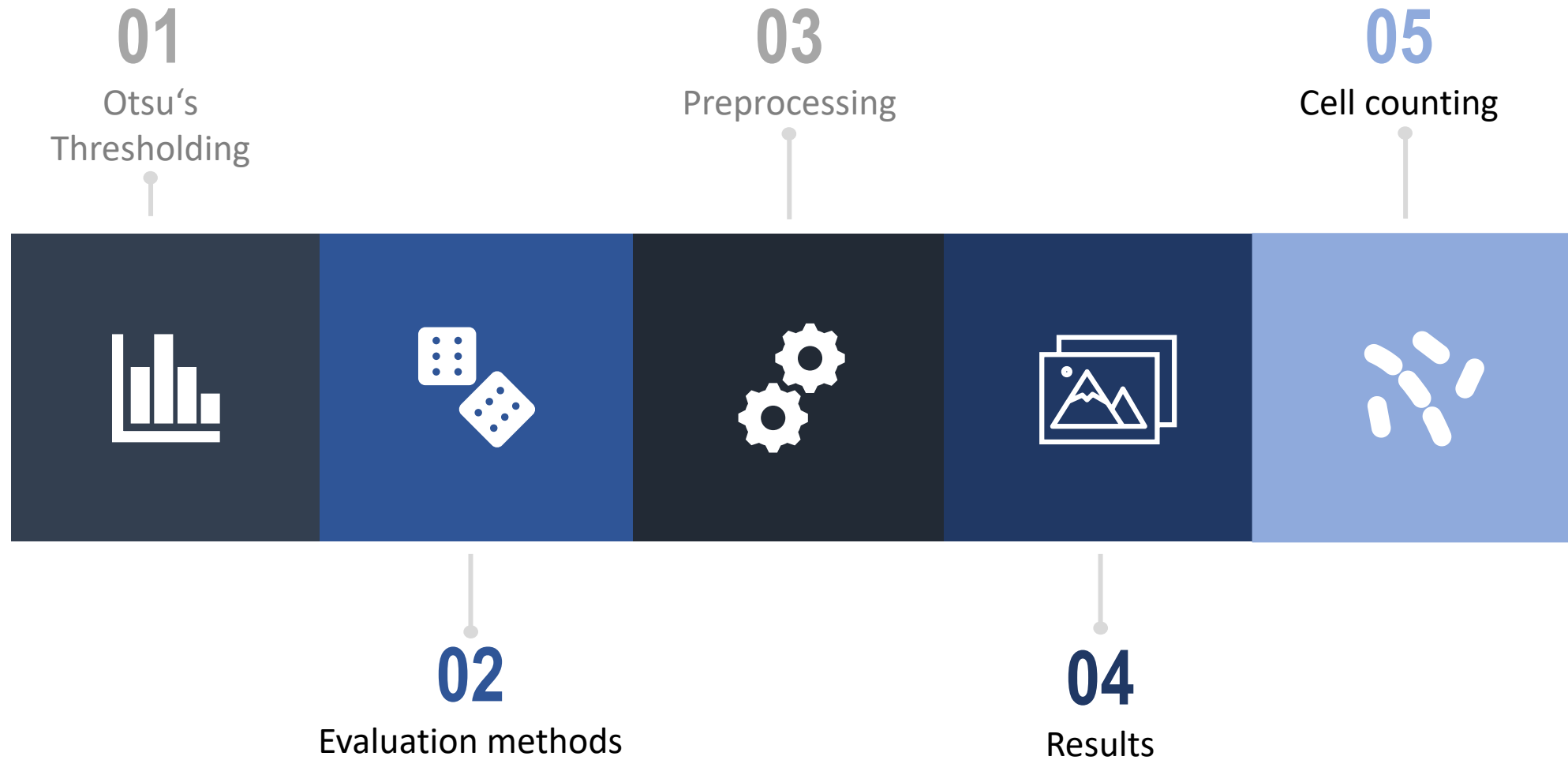


04

Results

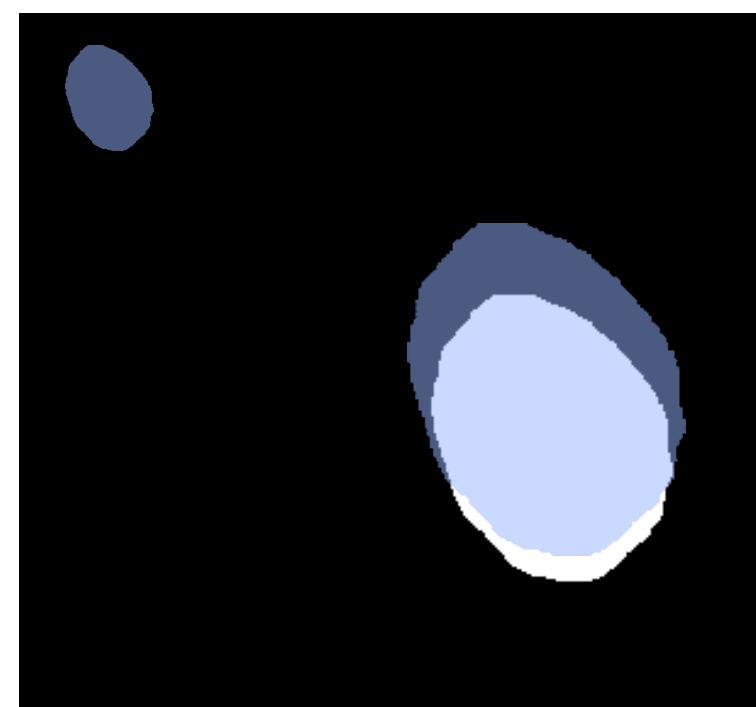
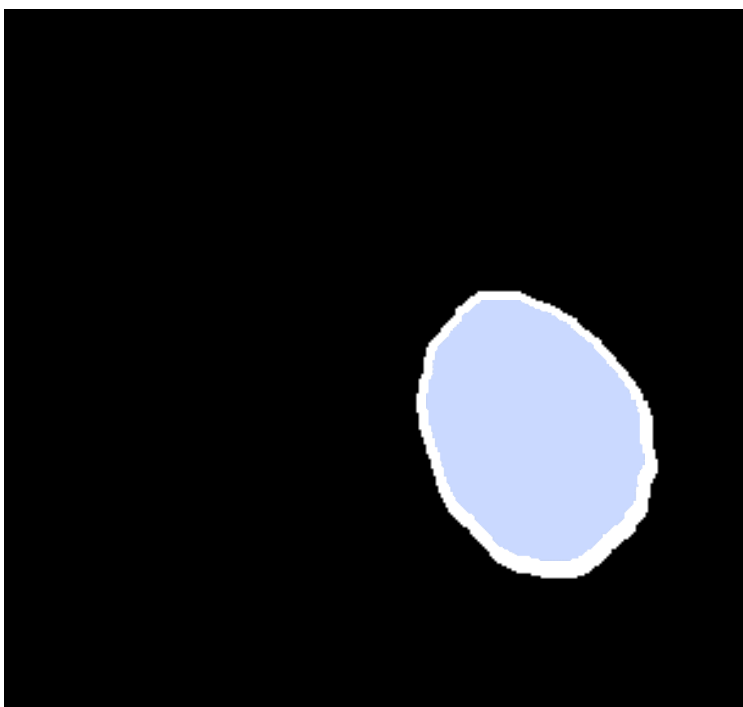
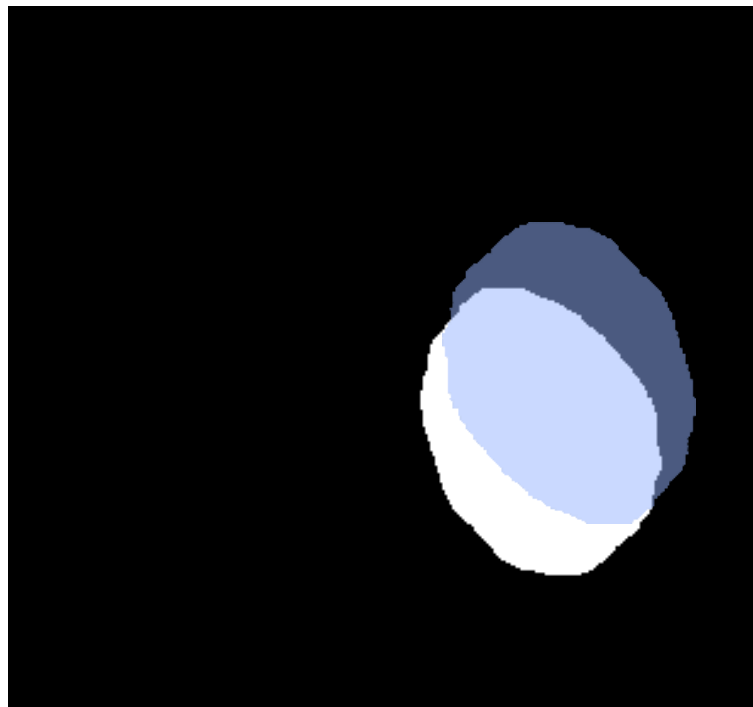


Outline





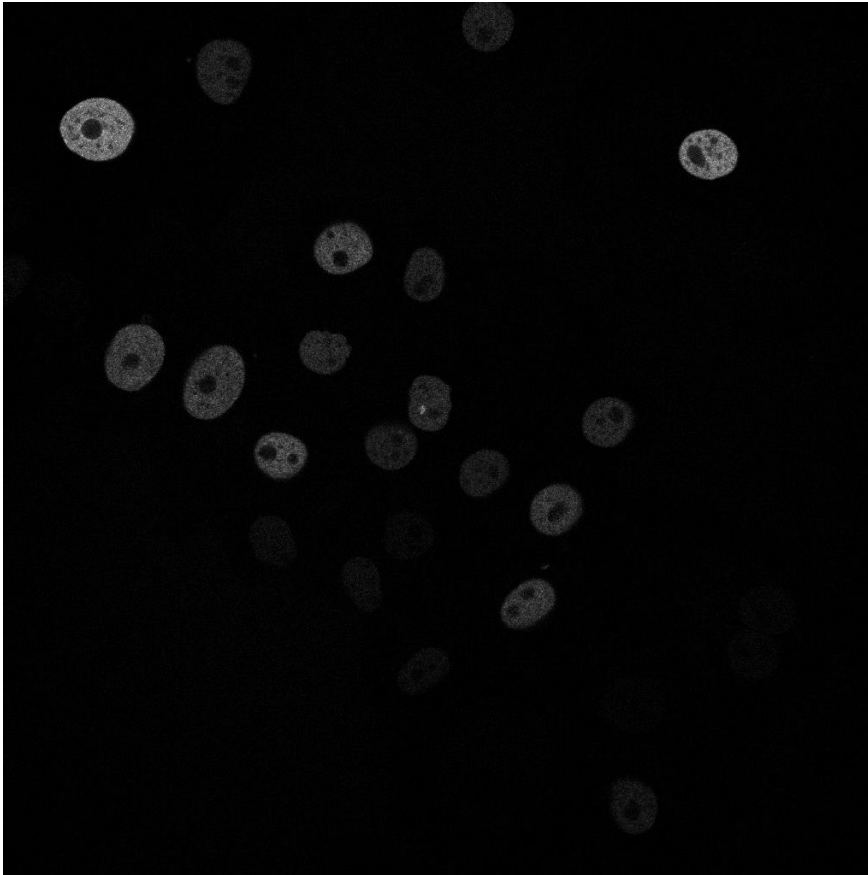
DSC vs. MSD vs. HD



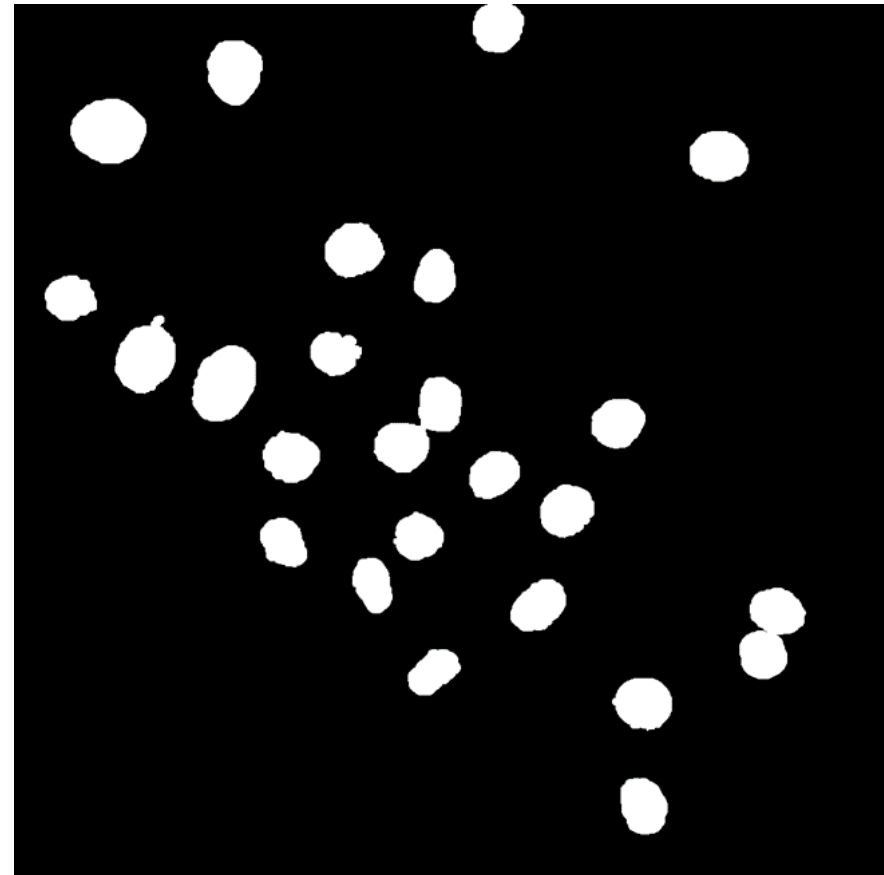


N2DH-GOWT1

Original Image

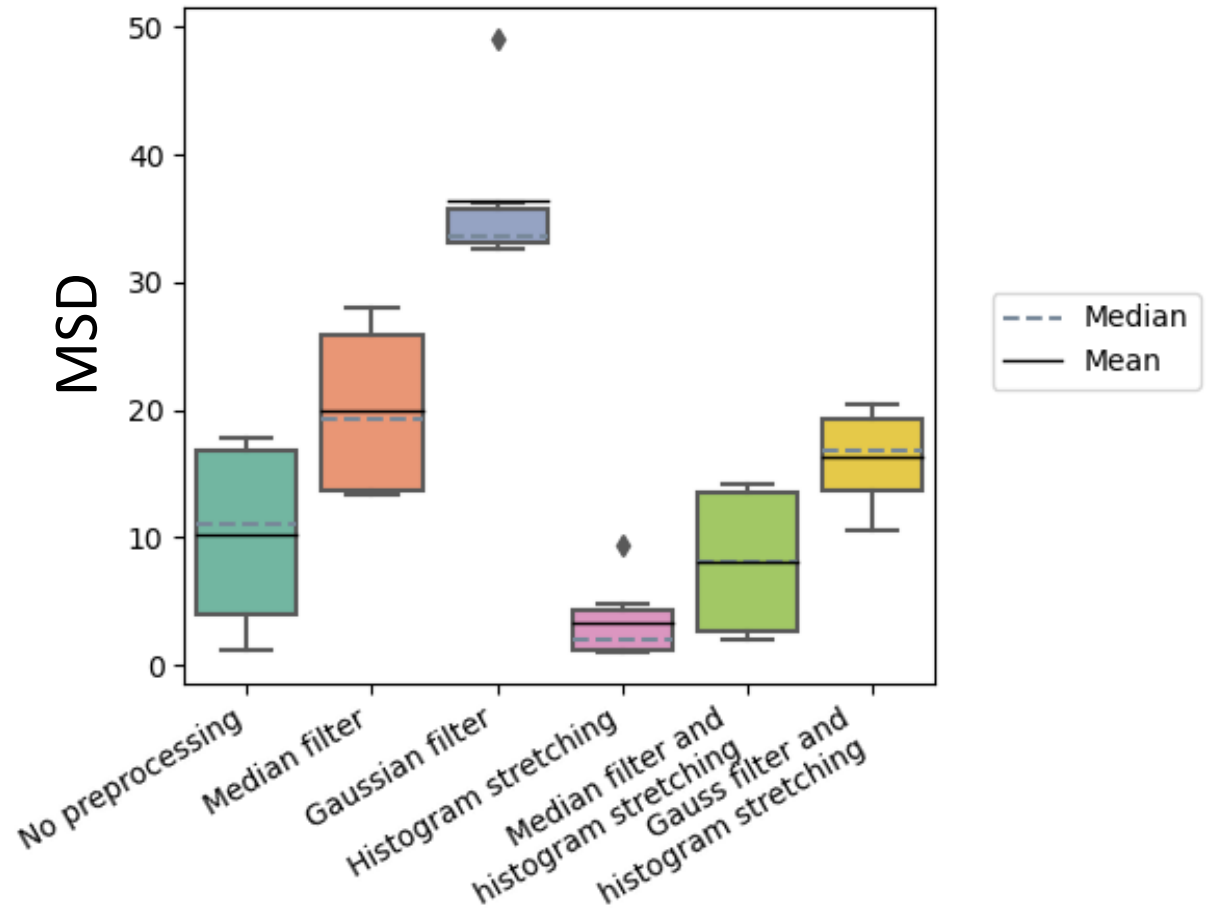
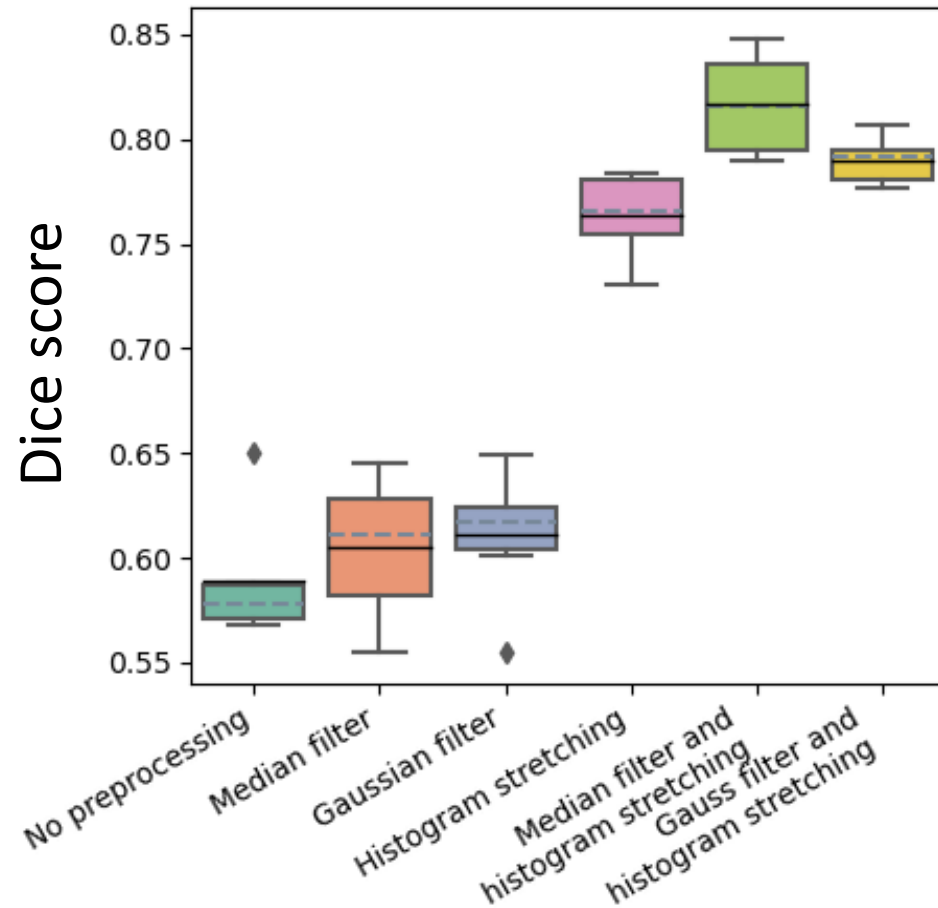


Ground truth





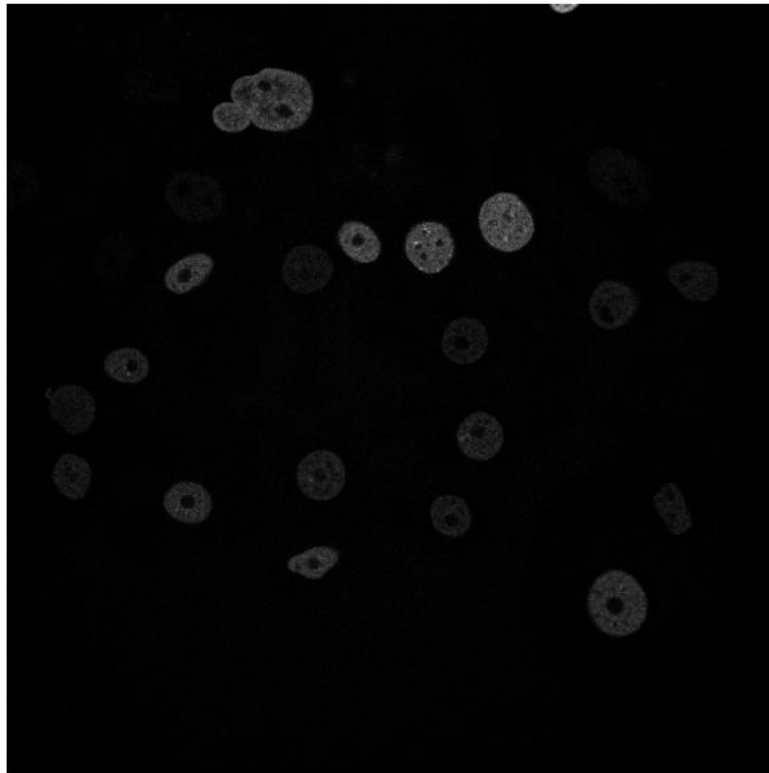
N2DH-GOWT1



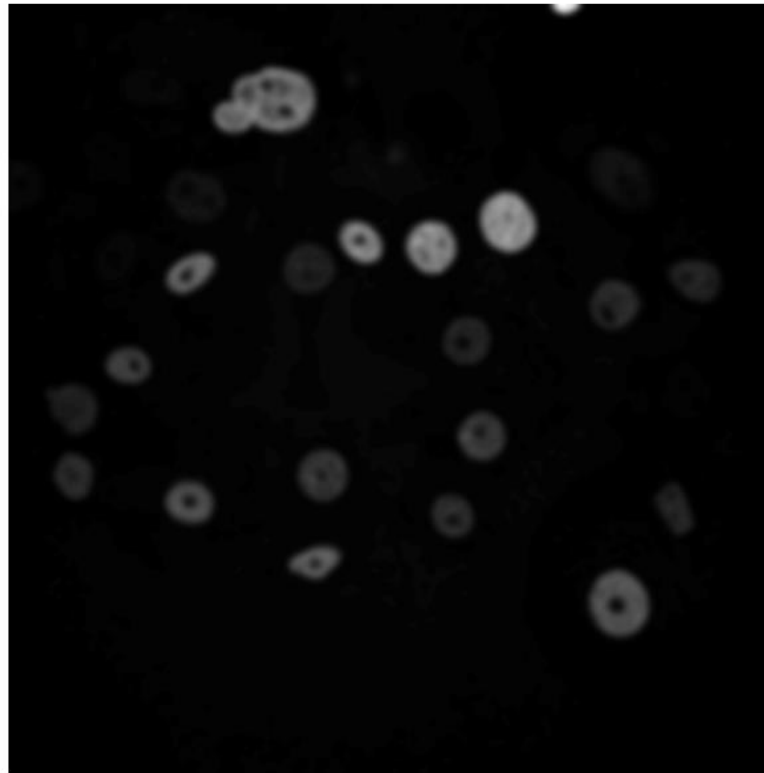


N2DH-GOWT1

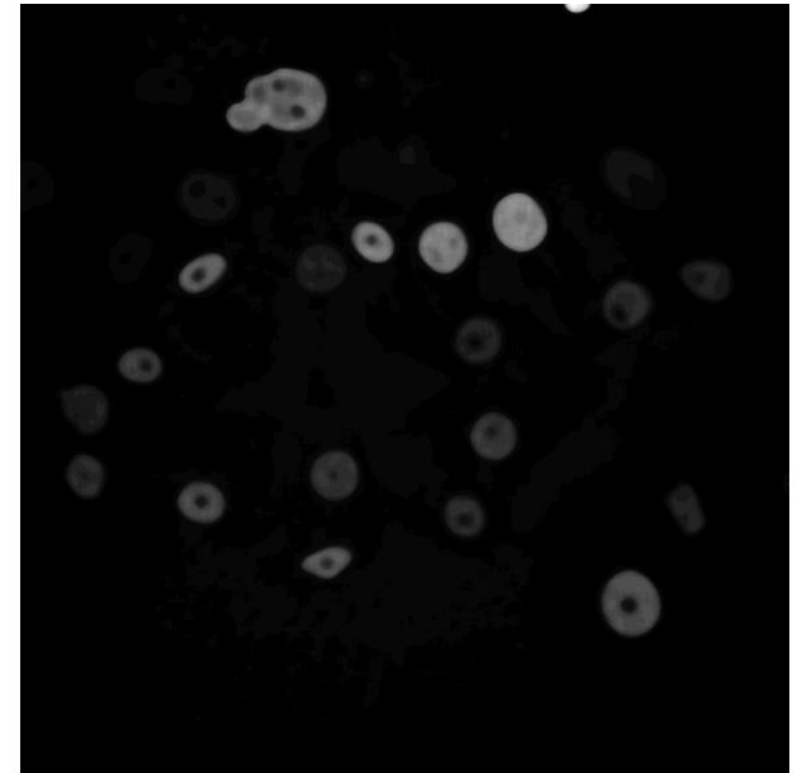
Original Image



Gaussian filter



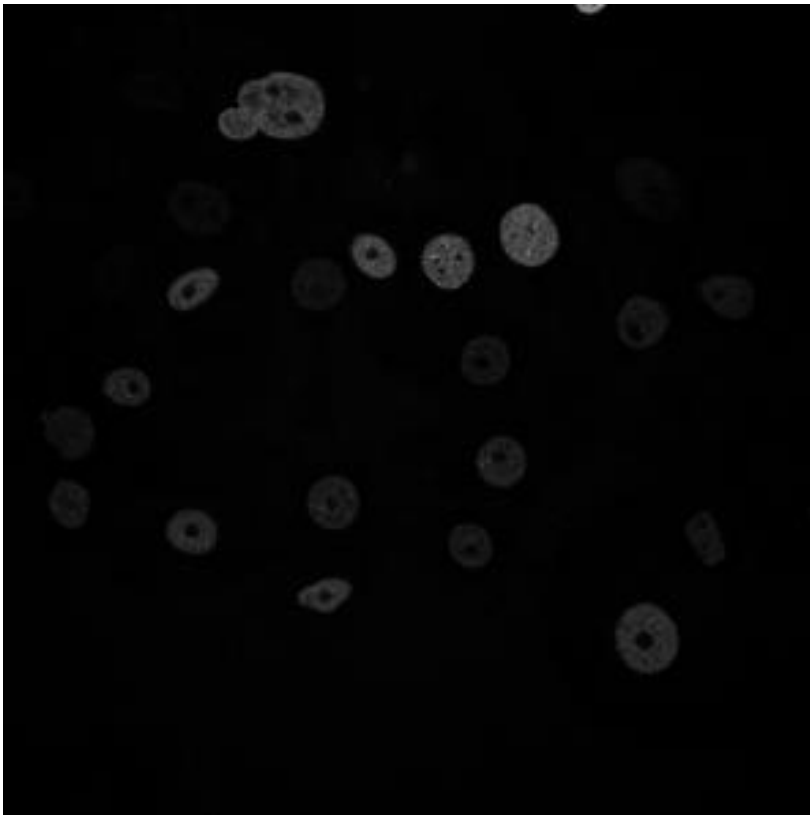
Median filter



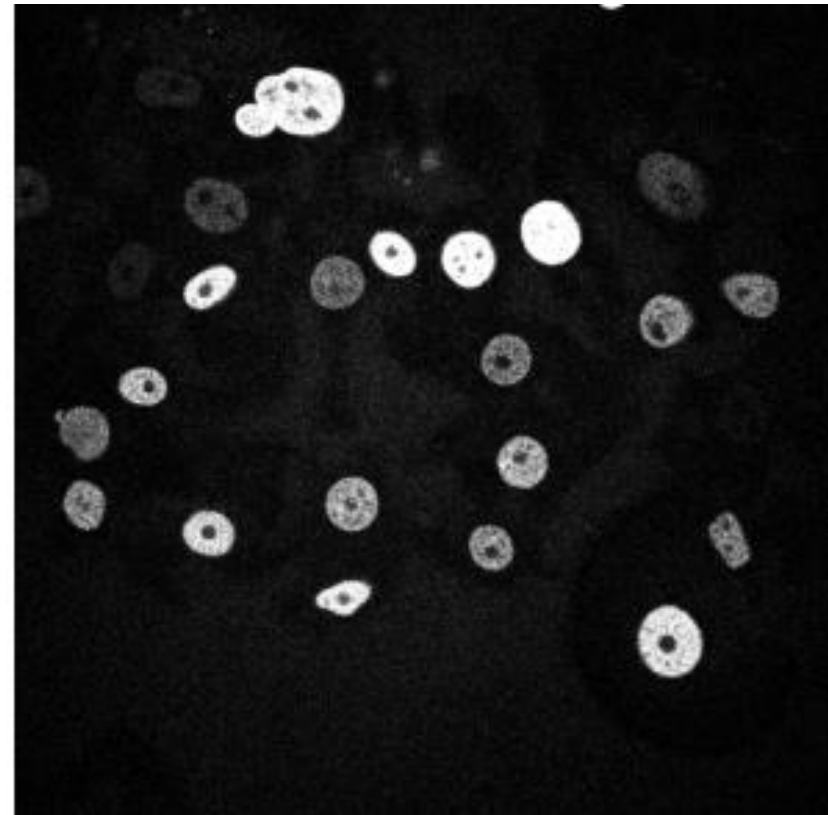


N2DH-GOWT1

Original Image



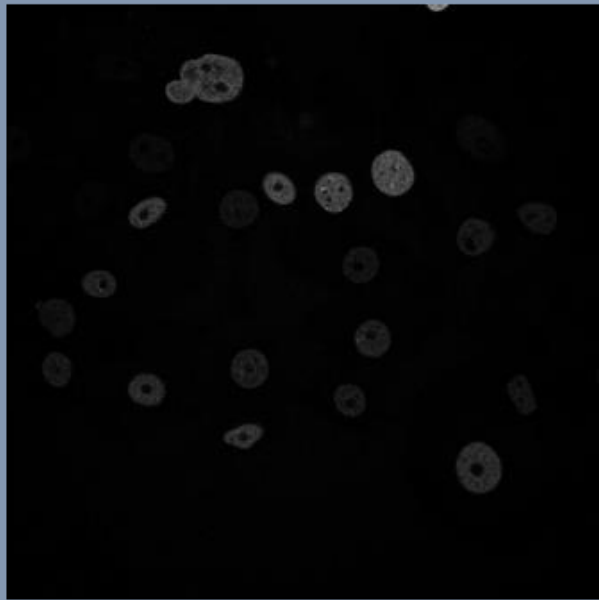
Histogram stretching



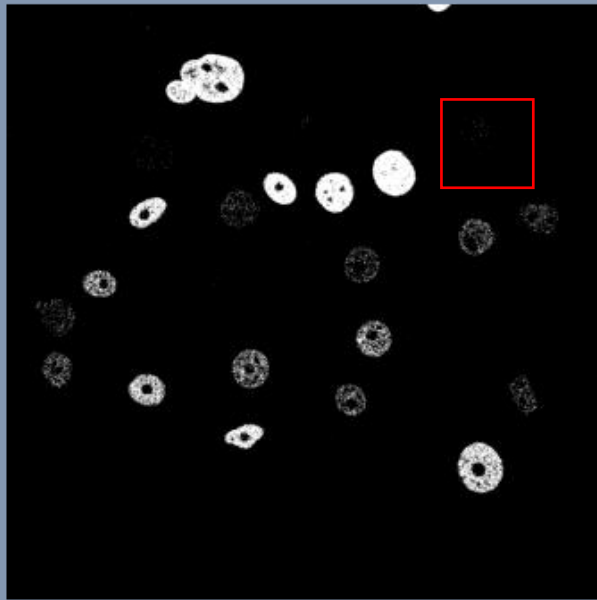


N2DH-GOWT1

Original image

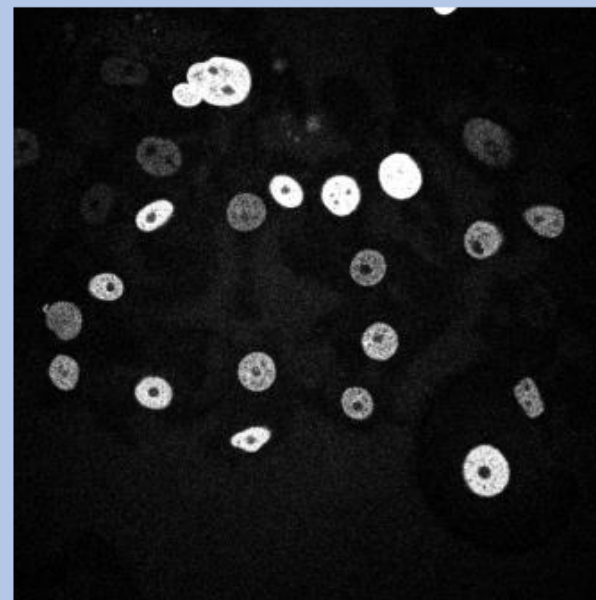


Segmentation

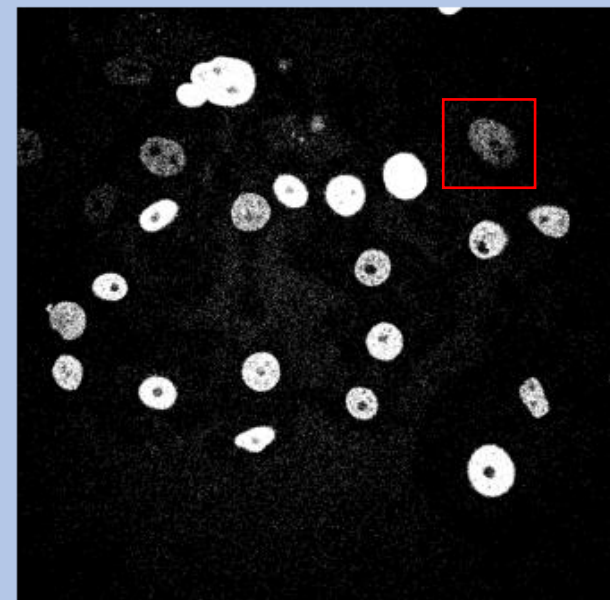


Dice Score
MSD

Histogram stretching



Segmentation

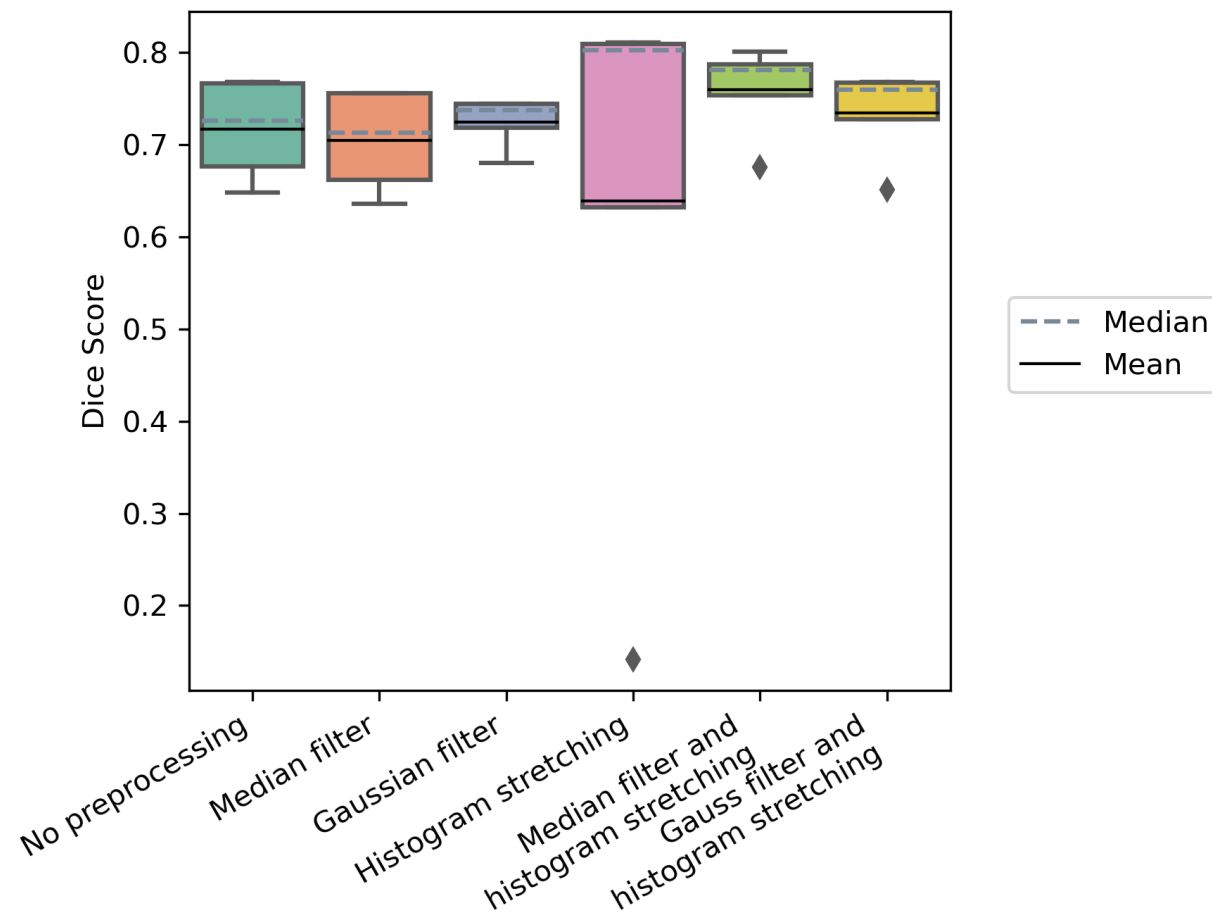


Dice Score
MSD



N2DL-HeLa

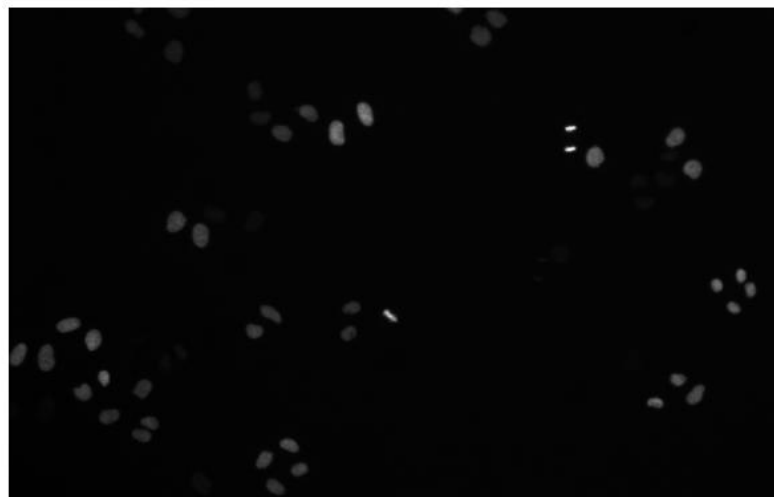
Best strategy:
Median filter and
histogram stretching
(0.76)





N2DL-HeLa

Original image



Gaussian filter

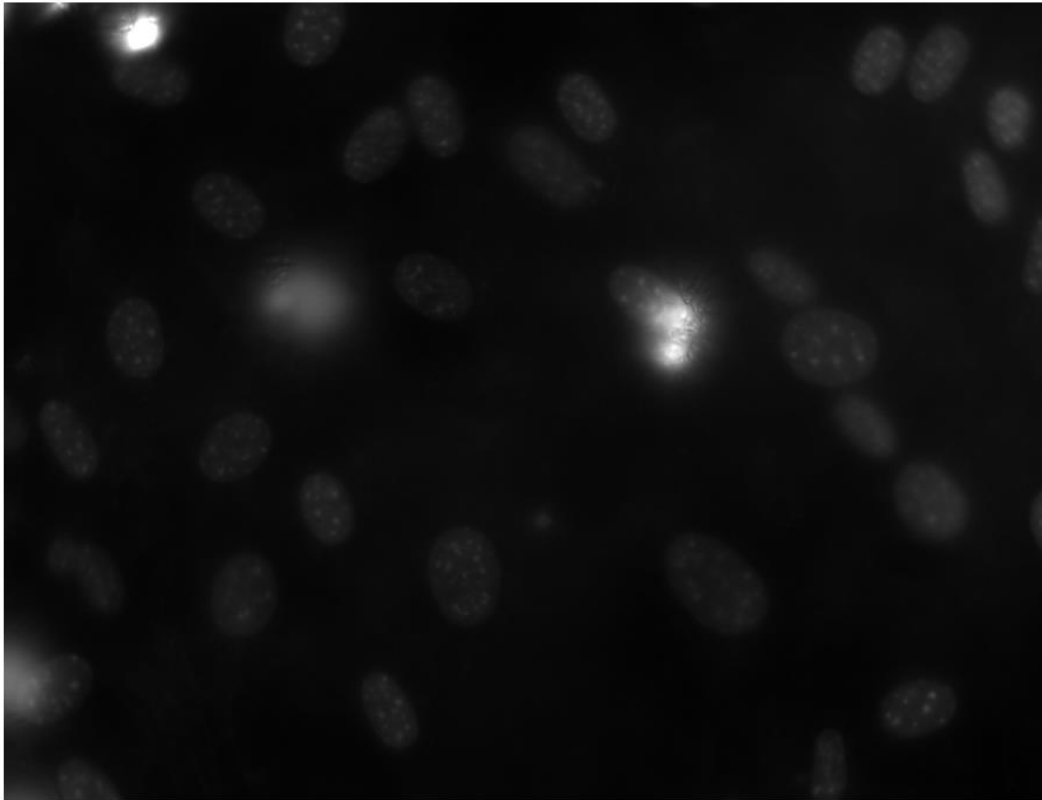


Segmentation

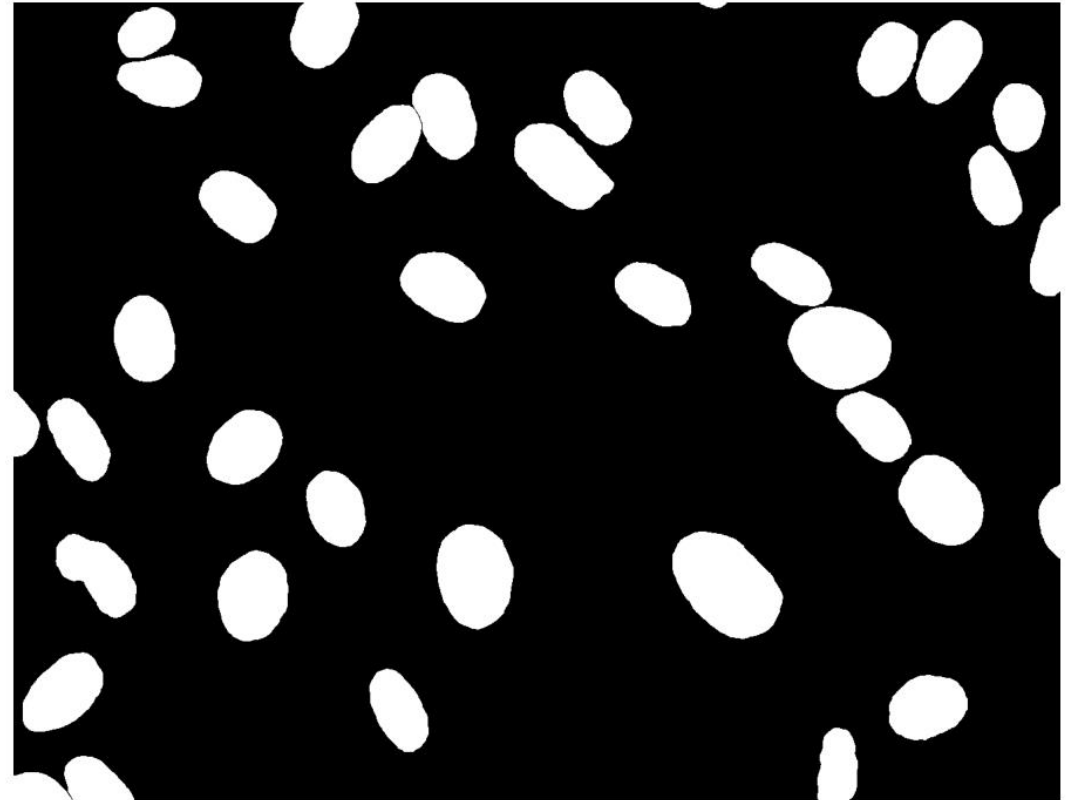


NIH3T3

Original image



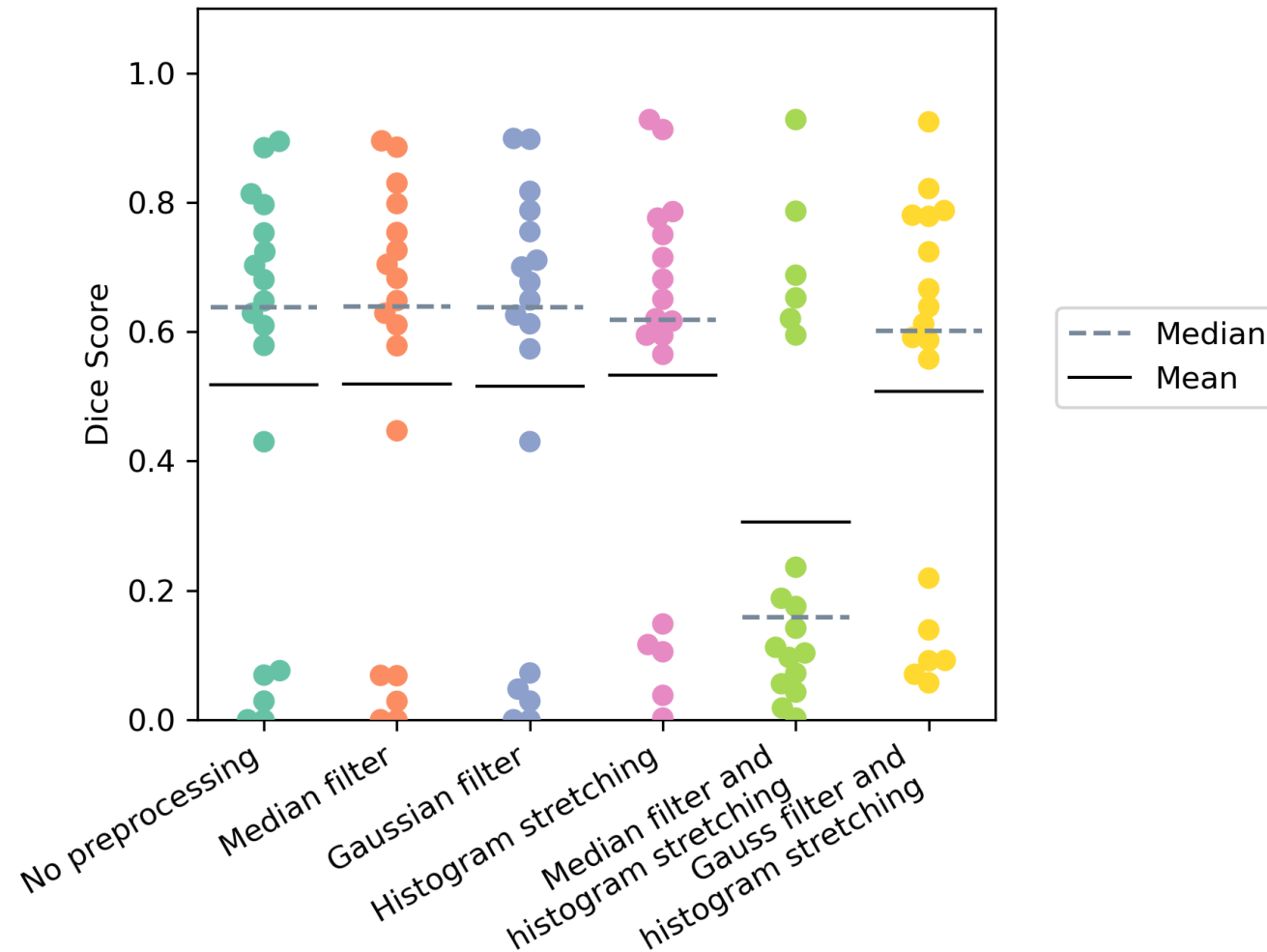
Ground truth





NIH3T3: One-level Otsu

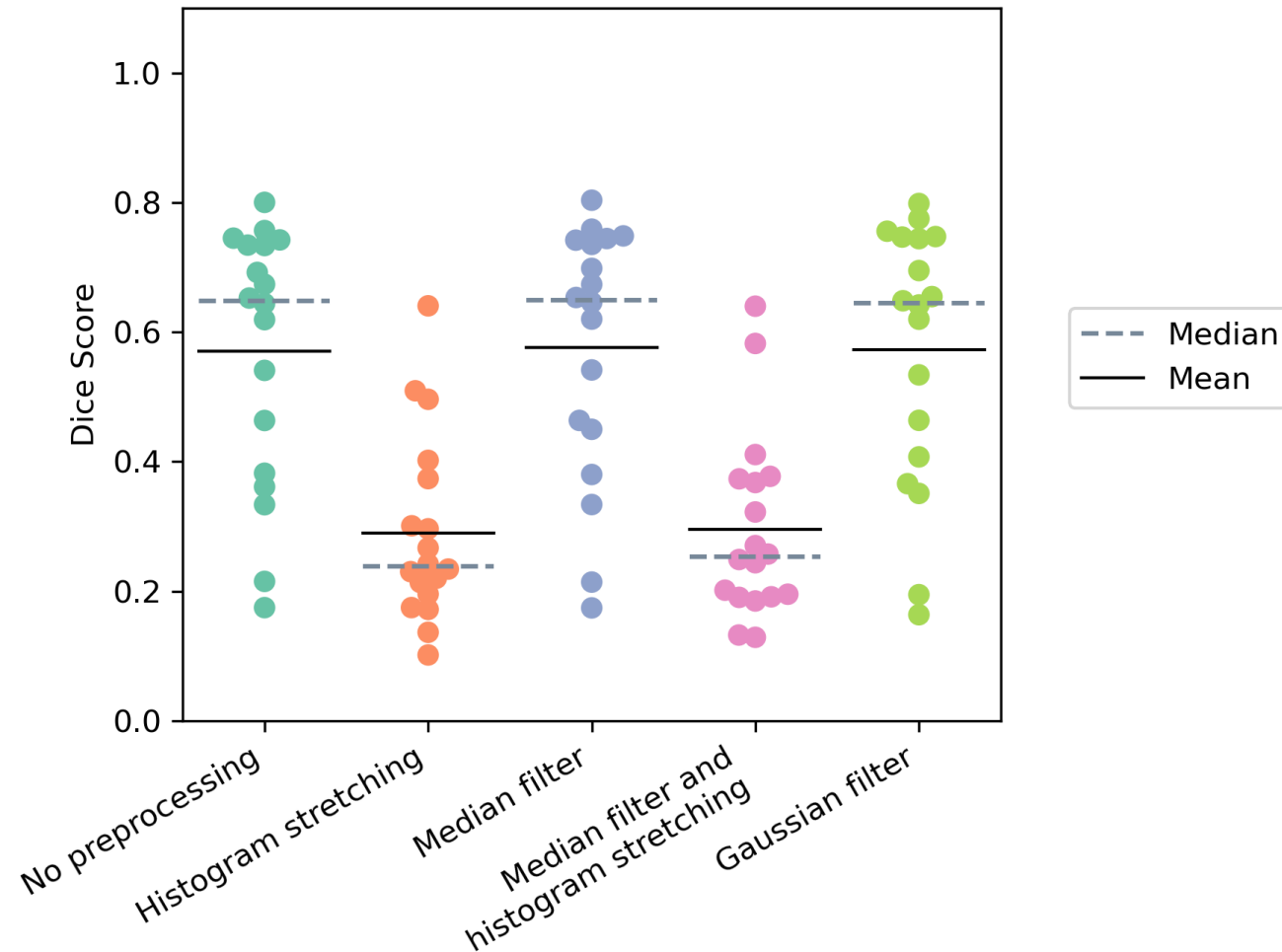
Best strategy:
Histogram stretching
(0.533)





NIH3T3: Two-level Otsu

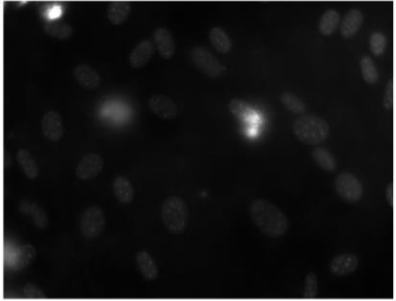
Best strategy:
Median filter
(0.577)



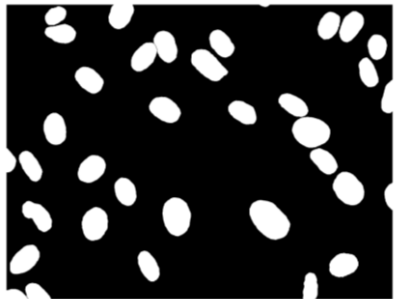


NIH3T3: Two-level Otsu

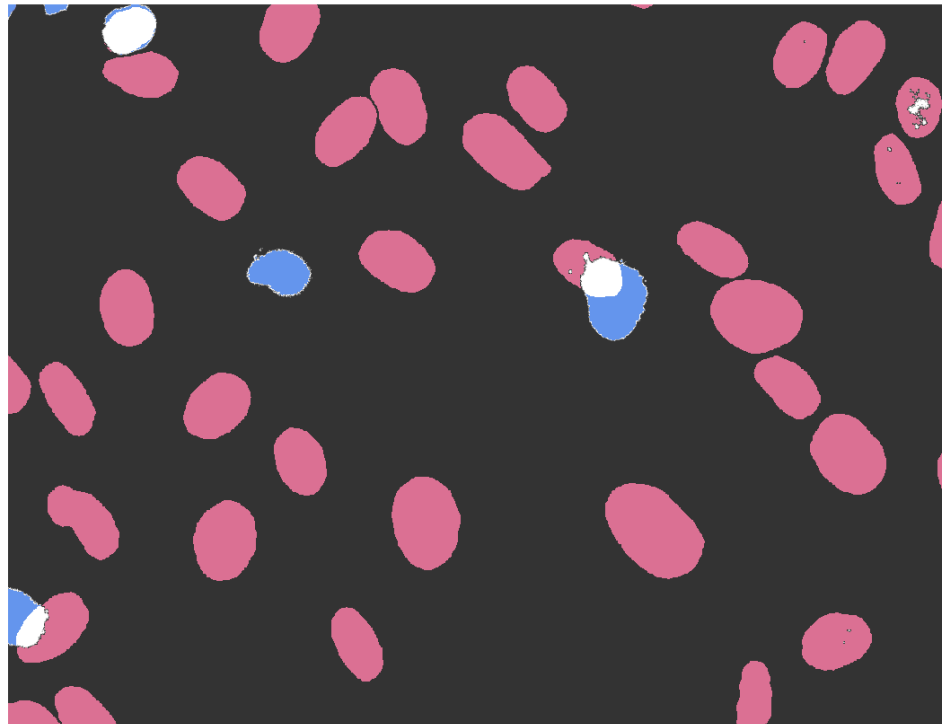
Original image



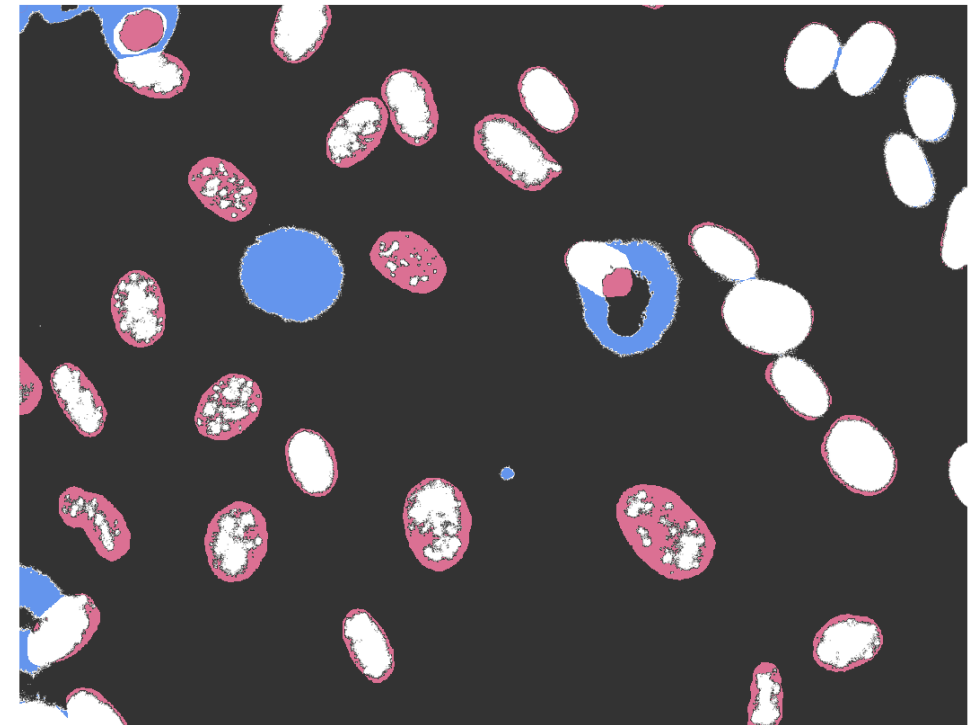
Ground truth



One-level Segmentation



Two-level Segmentation



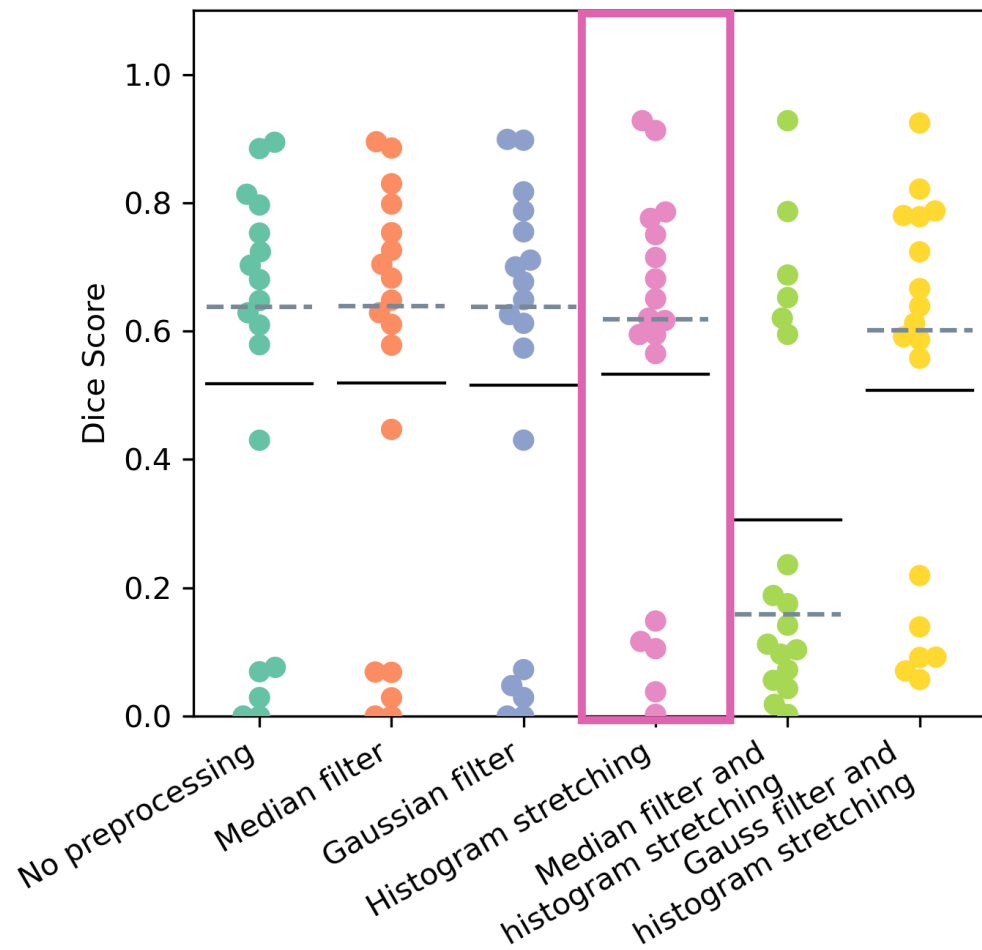
False negatives

False positives

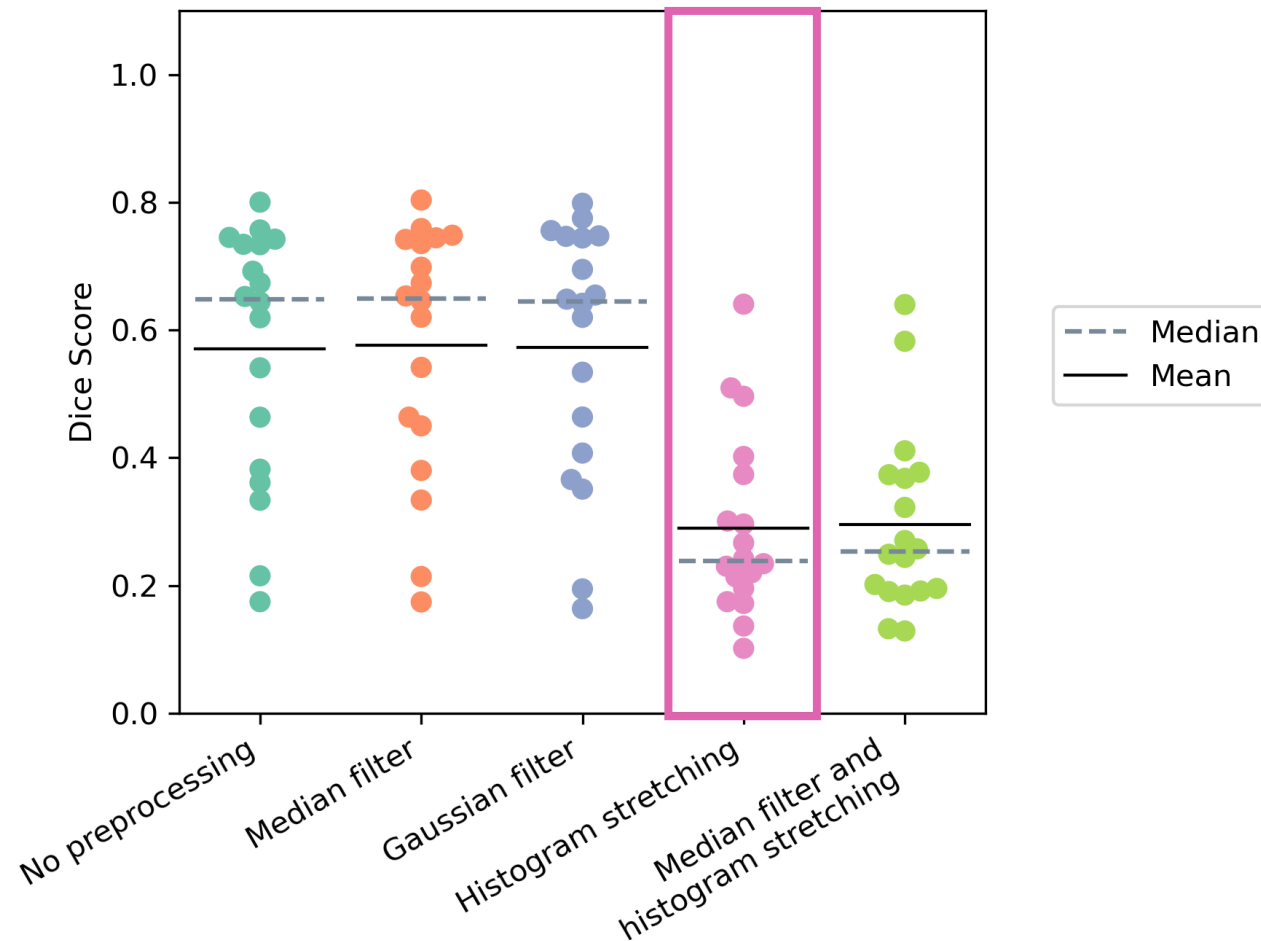


NIH3T3: Two-level Otsu

One-level Otsu



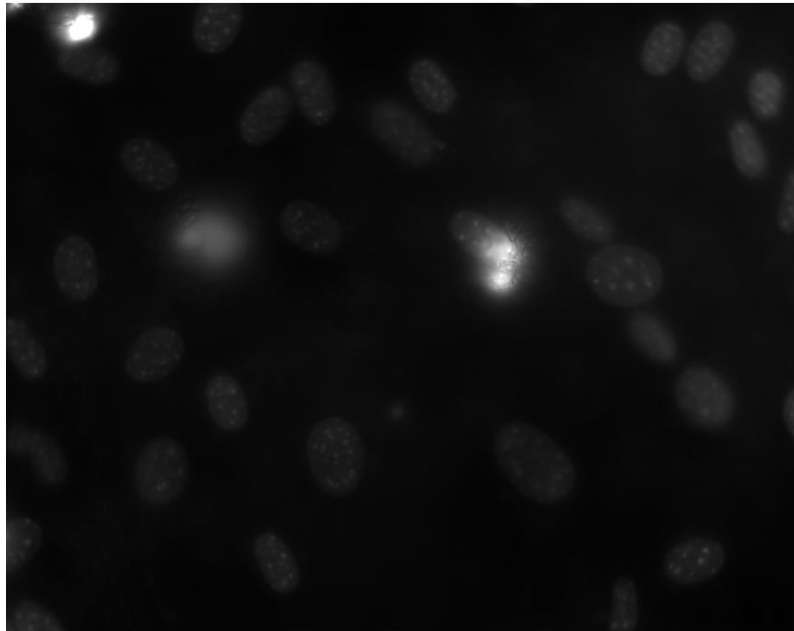
Two-level Otsu



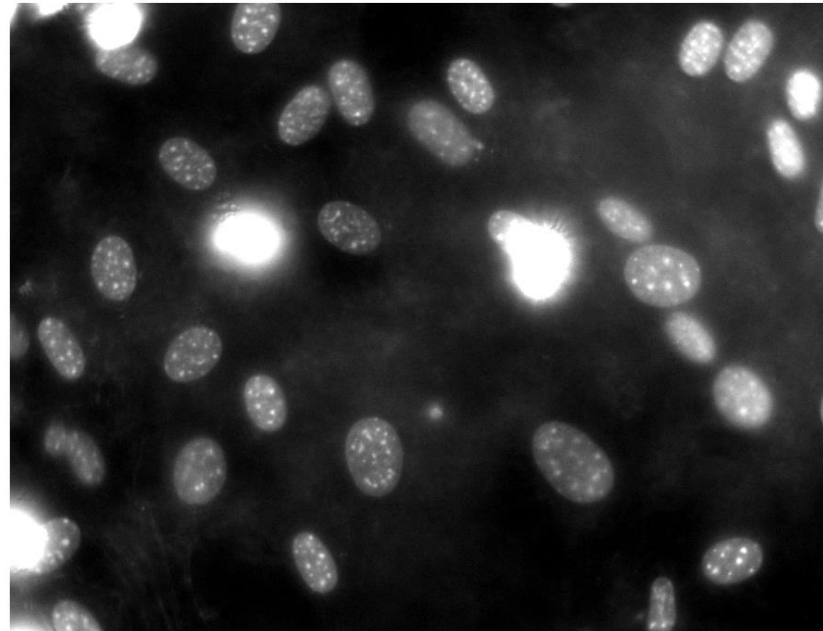


NIH3T3: Two-level Otsu

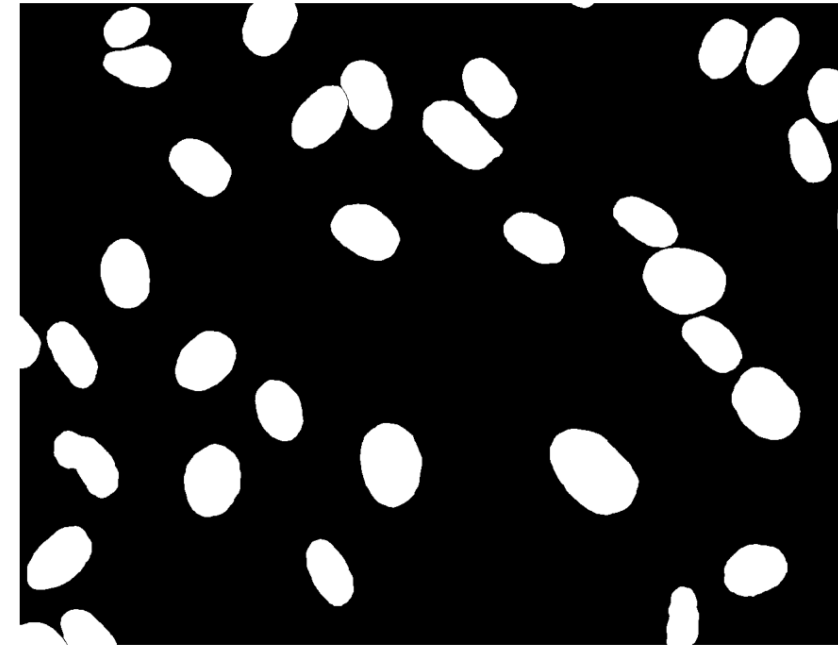
Original image



Histogram stretching

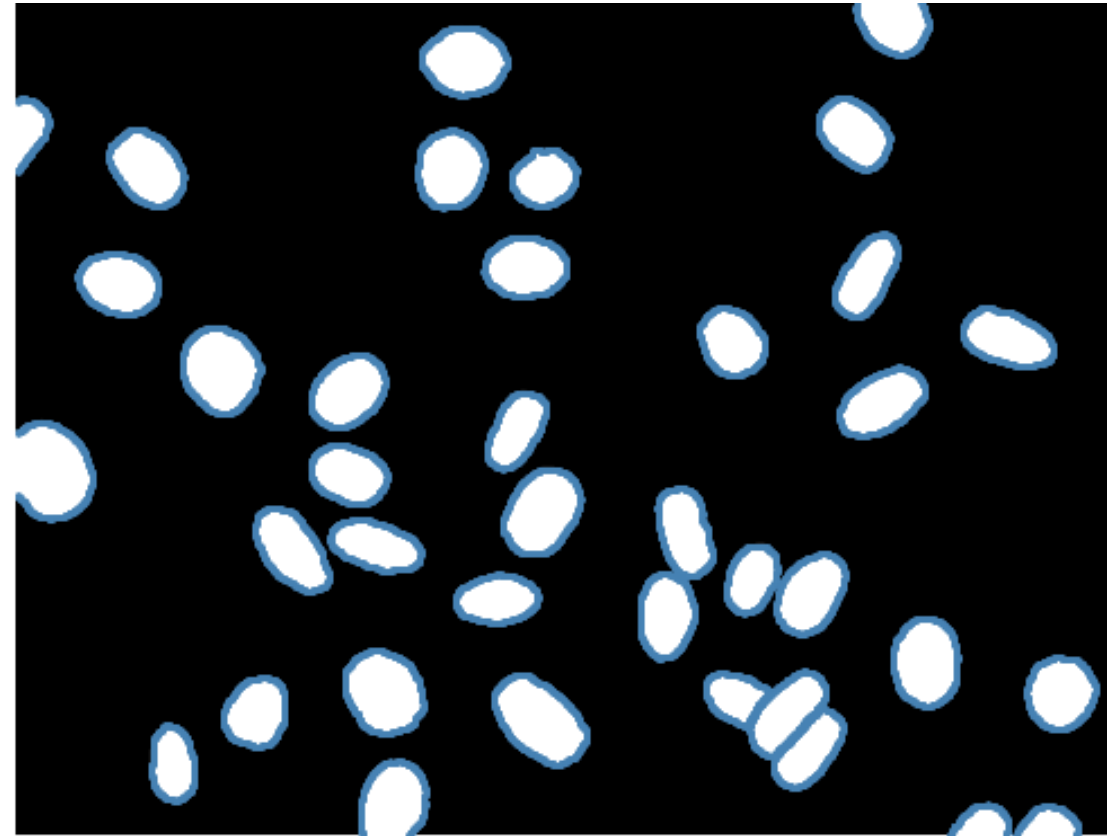
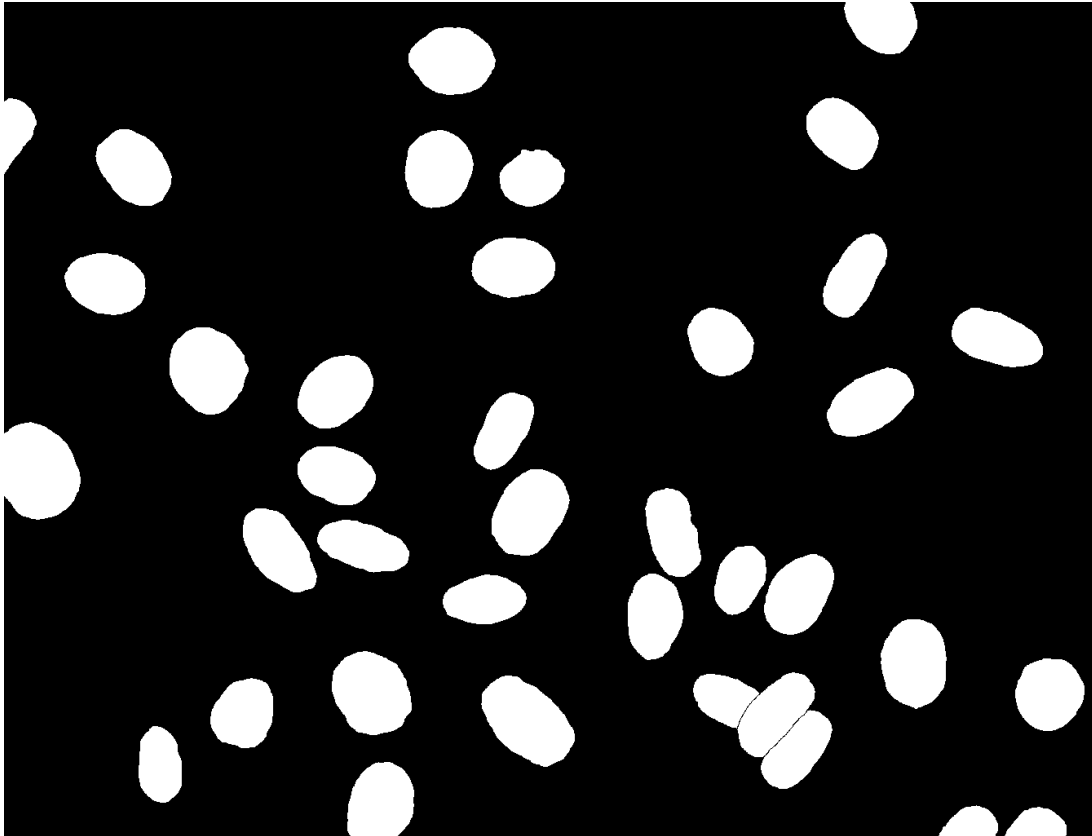


Ground truth



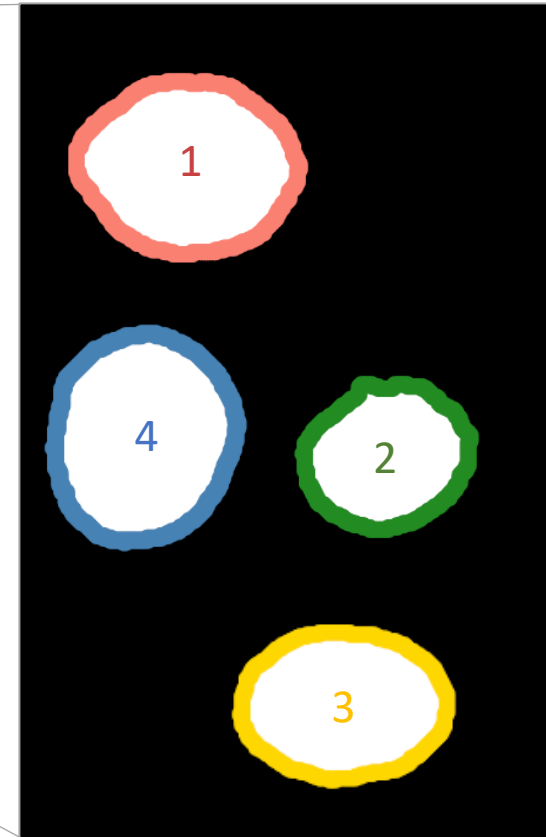
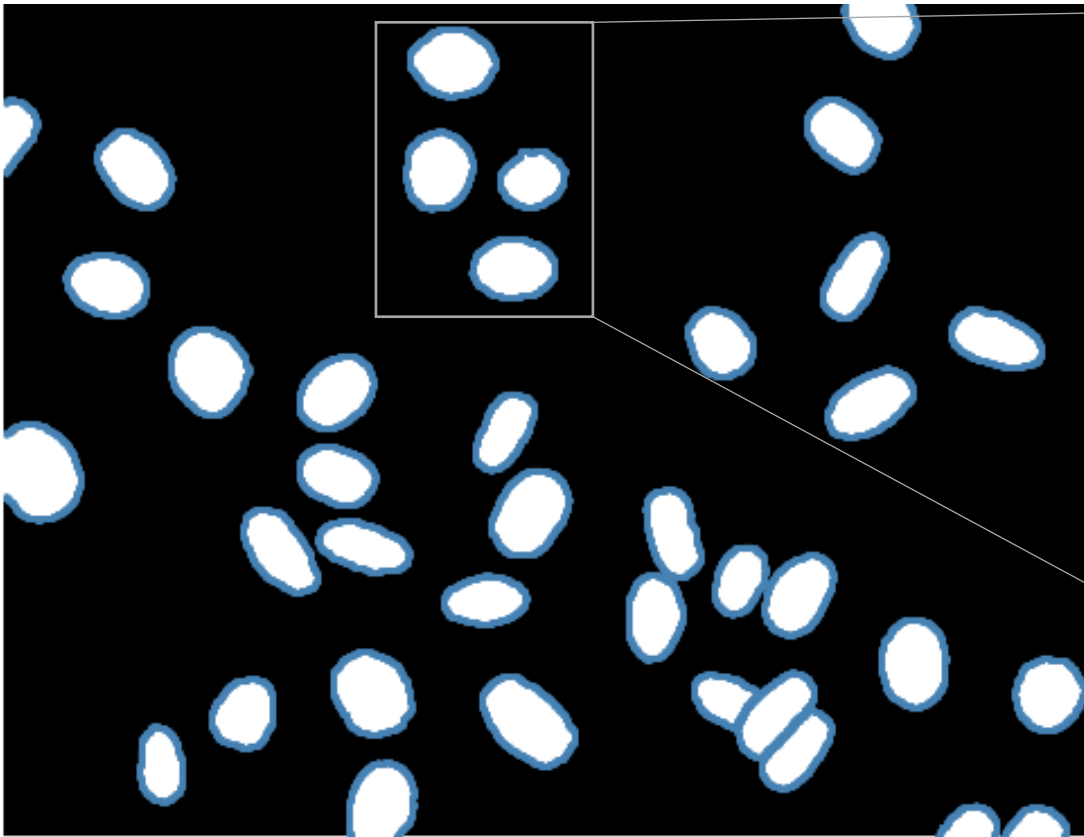


Cell nuclei counting



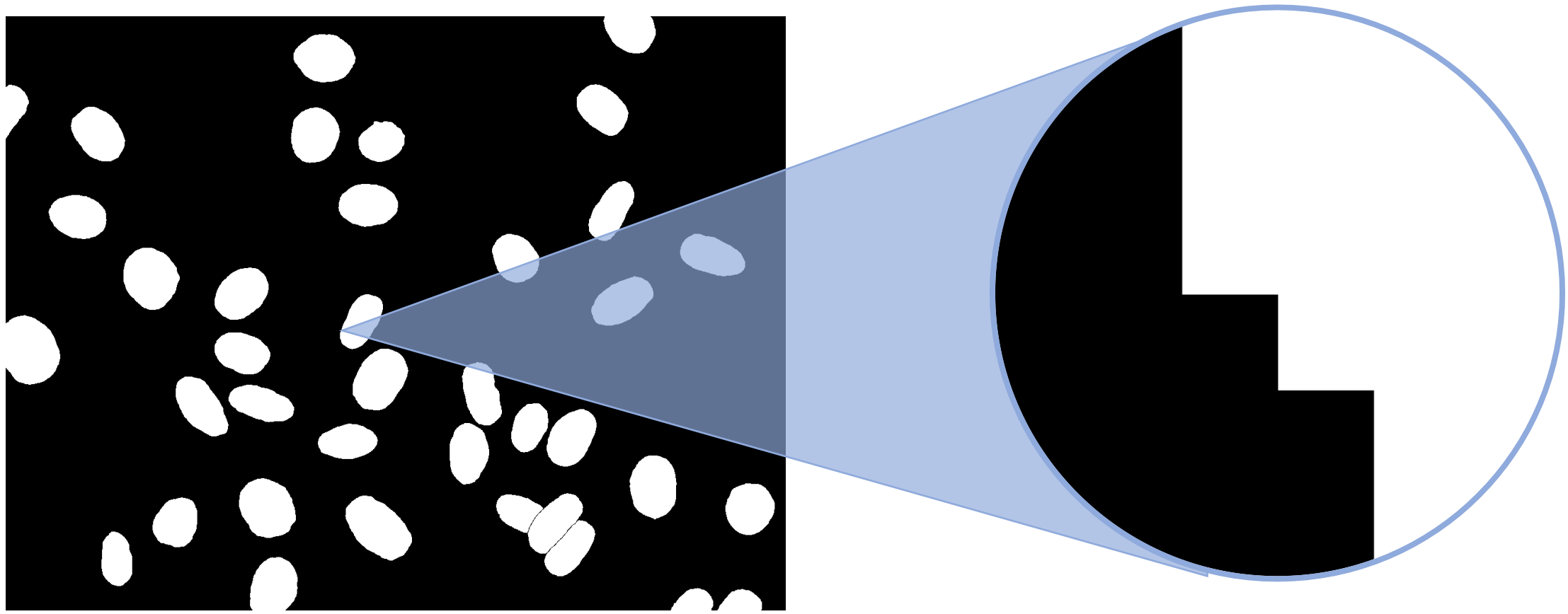


Cell nuclei counting



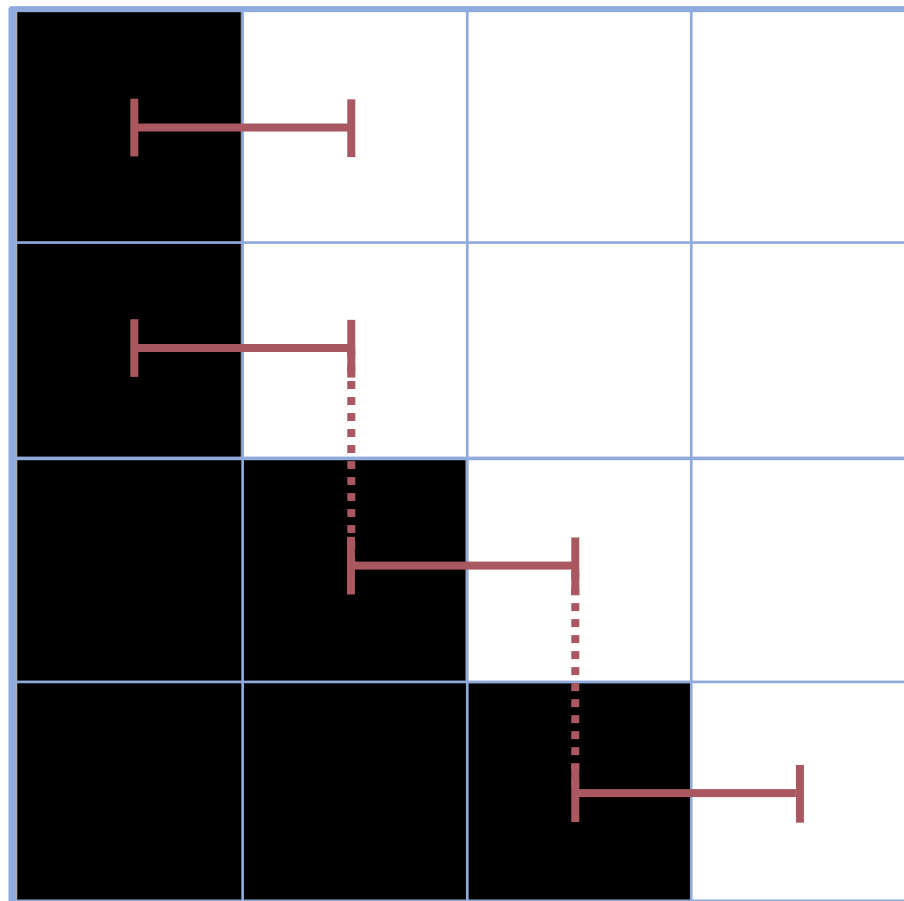


Cell nuclei counting





Cell nuclei counting



$$d = 1$$



Cell nuclei counting

```
edge_pixels = []

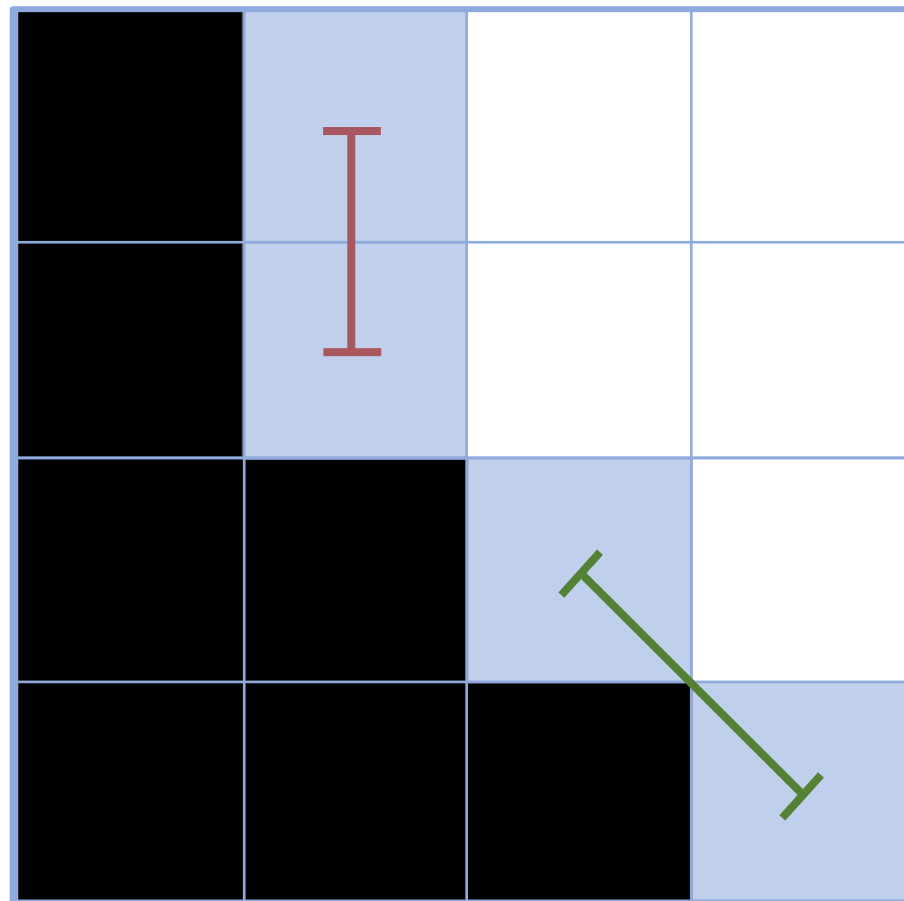
for index in np.ndindex(img.shape):
    if workimg[index[0]][index[1]] == 1:
        if 0 in workimg[(index[0] - 1):(index[0] + 2), (index[1] - 1):(index[1] + 2)]:
            edge_pixels.append(index)

return edge_pixels
```

$$\overbrace{\quad}^{d = 1}$$



Cell nuclei counting

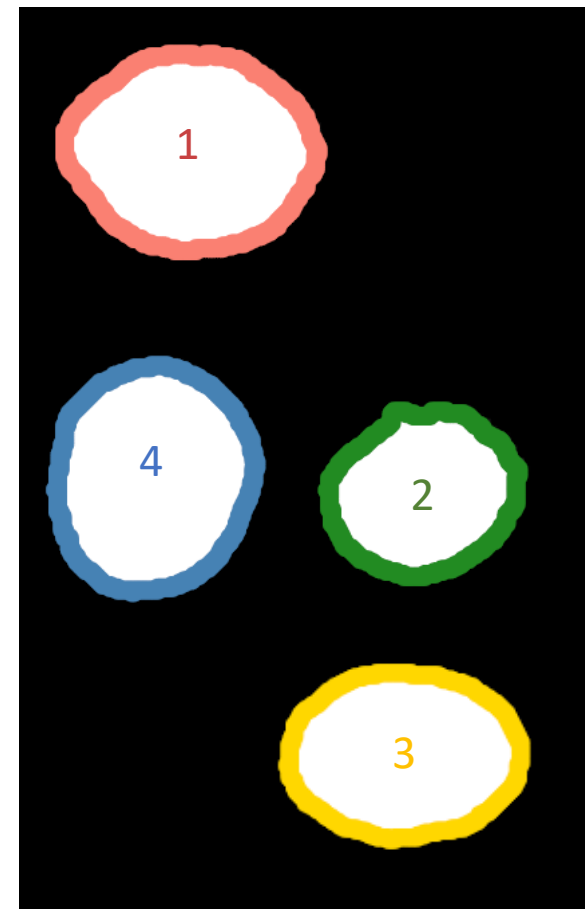


$$d = 1$$
$$d = \sqrt{2}$$



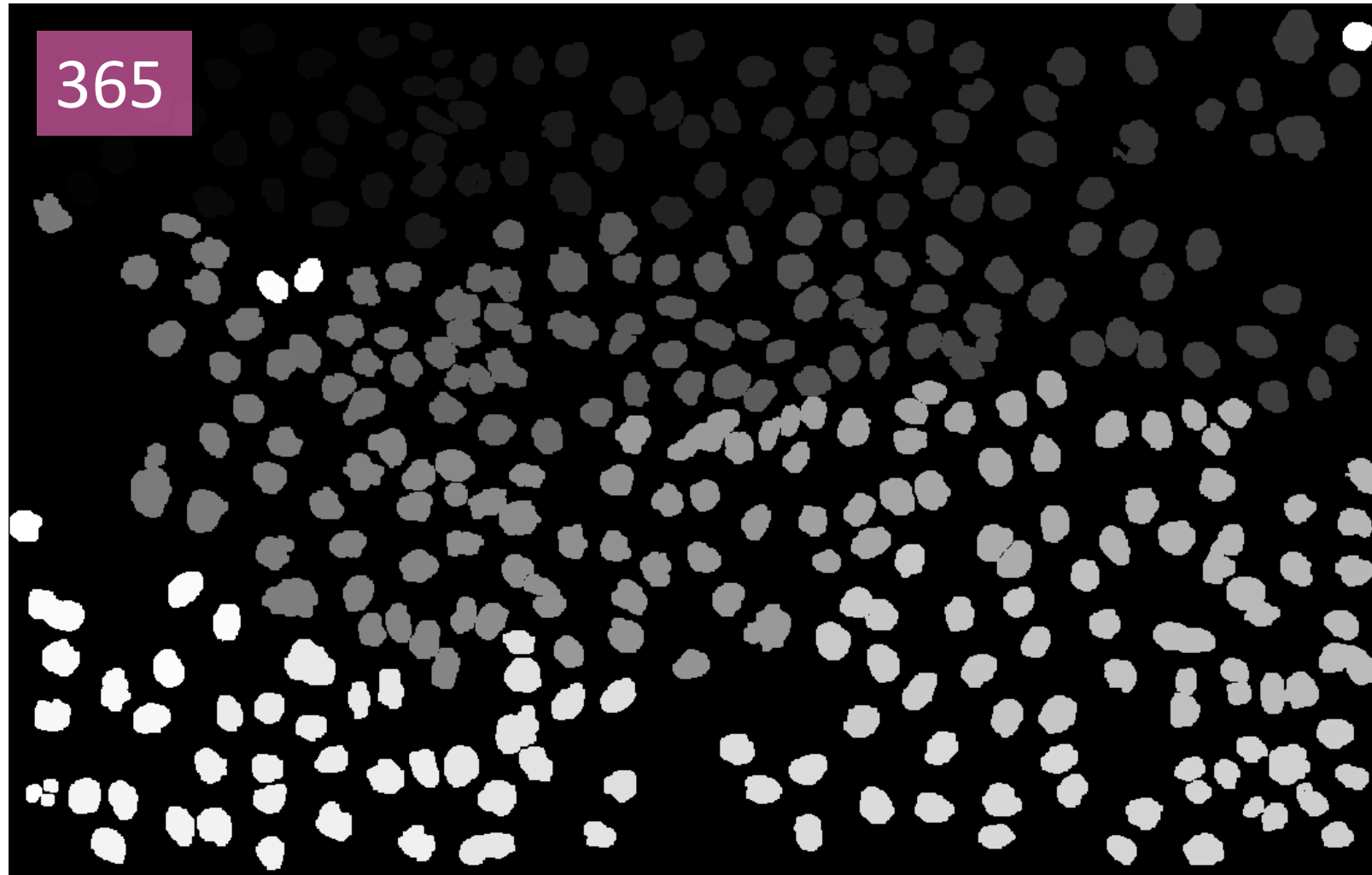
Cell nuclei counting

```
for start_pixel in border_pixels:
    old_group = [start_pixel]
    new_group = []
    first_run = True
    while old_group != new_group:
        if not first_run:
            old_group = new_group
        first_run = False
        for pixel in old_group:
            for other_pixel in border_pixels:
                if math.dist(pixel, other_pixel) < 2:
                    new_group.append(other_pixel)
                    border_pixels.remove(other_pixel)
            if pixel in border_pixels:
                border_pixels.remove(pixel)
    all_groups.append(new_group)
```





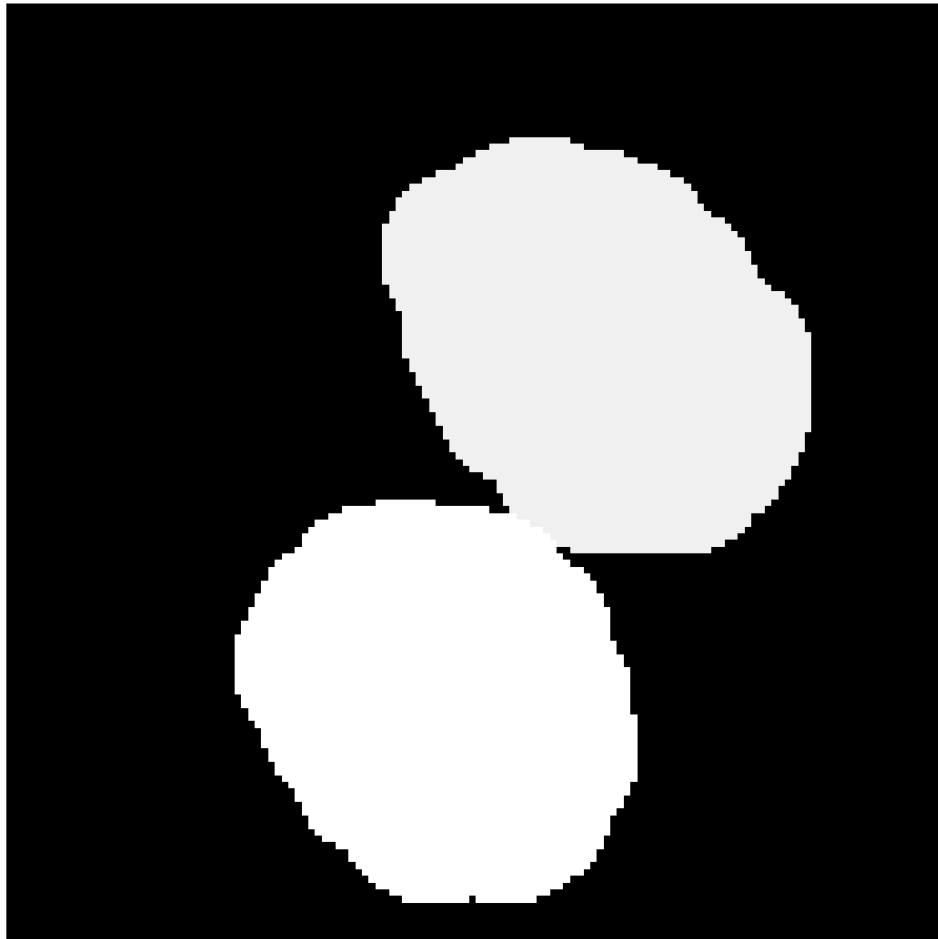
Evaluation cell nuclei counting



349



Evaluation cell nuclei counting





Conclusion

Successful
implementation

Evaluation

Future
improvements
possible



Thank you
for your
attention!

Laura Wächter, Veronika Schuler, Elizaveta Chernova, Hannah L. Winter

Additional slide – MSD code

```
# calculate minimum distances for each point in seg to the sets of points in gt
tree_seg_gt = spatial.cKDTree(gt_array)
mindist_seg_gt, minid_seg_gt = tree_seg_gt.query(seg_array)

# calculate sum and length of arrays with minimal distances
sum_seg_gt = np.sum(mindist_seg_gt)
size_seg_gt = len(mindist_seg_gt)

mean_surface_distance = (1/(size_gt_seg+size_seg_gt))*(sum_gt_seg + sum_seg_gt)

return mean_surface_distance
```

Additional slide – HD code

```
# calculate minimum distances for each point in seg to the sets of points in gt
tree_seg_gt = spatial.cKDTree(gt_array)
mindist_seg_gt, minid_seg_gt = tree_seg_gt.query(seg_array)

# calculate sum and length of arrays with minimal distances
sum_seg_gt = np.sum(mindist_seg_gt)
size_seg_gt = len(mindist_seg_gt)

hausdorff_distance = max(max_gt_seg, max_seg_gt)

return hausdorff_distance
```


Additional slide - cell counting dataset 1

Table 1: Results of the cell counting on the N2DH-GOWT1 dataset.

	Calculated number	Ground truth number	Absolute difference	Relative difference
man_seg01.tif	24	23	1	0.043478
man_seg21.tif	23	24	-1	-0.041667
man_seg31.tif	24	22	2	0.090909
man_seg39.tif	23	25	-2	-0.080000
man_seg52.tif	30	30	0	0.000000
man_seg72.tif	28	28	0	0.000000

Additional slide – cell counting dataset 2

	Calculated number	Ground truth number	Absolute difference	Relative difference
man_seg13.tif	58	59	-1	-0.016949
man_seg52.tif	107	109	-2	-0.018349
man_seg75.tif	365	349	16	0.045845
man_seg79.tif	329	342	-13	-0.038012



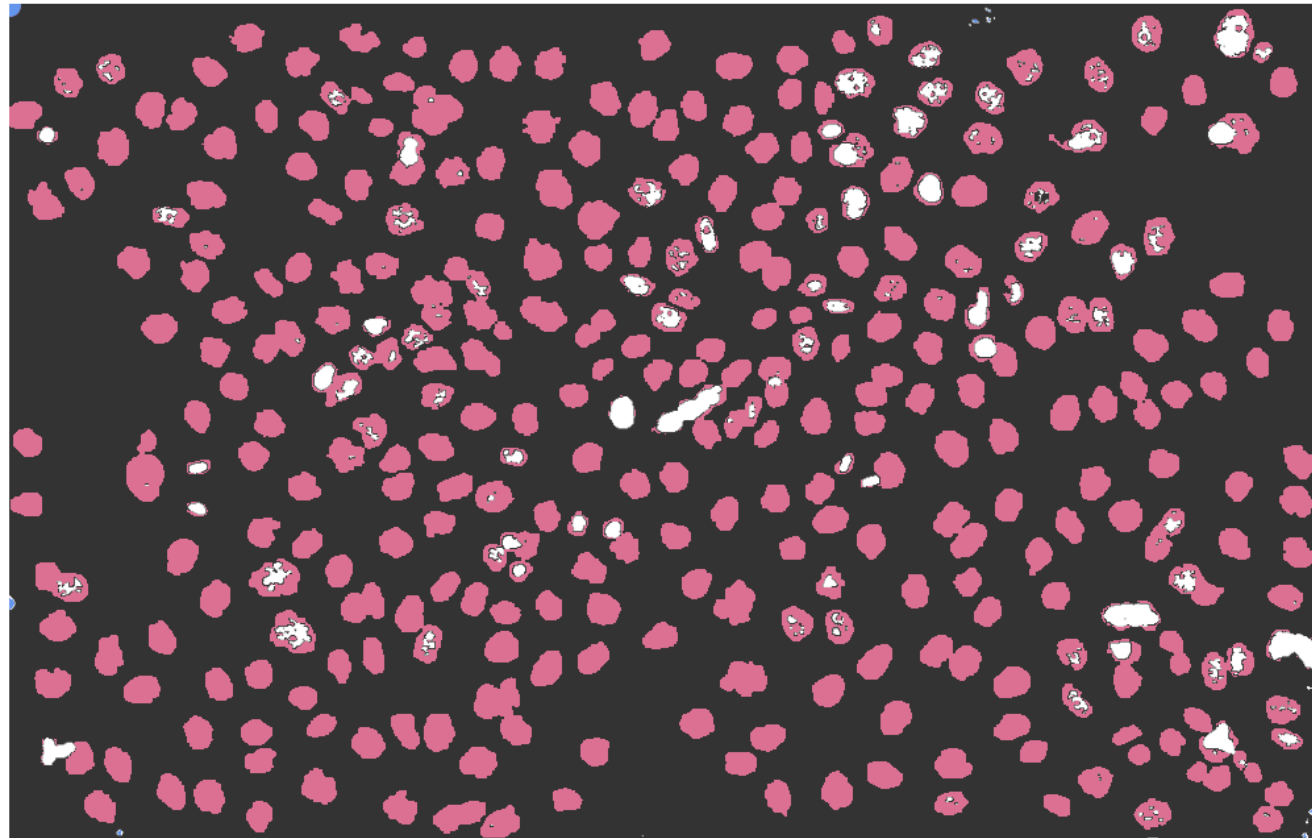
Cell Counting

```
for start_pixel in border_pixels:
    old_group = [start_pixel]
    new_group = []
    first_run = True
    while old_group != new_group:
        if not first_run:
            old_group = new_group
        first_run = False
        for pixel in old_group:
            for other_pixel in border_pixels:
                if math.dist(pixel, other_pixel) < 2:
                    new_group.append(other_pixel)
                    border_pixels.remove(other_pixel)
            if pixel in border_pixels:
                border_pixels.remove(pixel)
    all_groups.append(new_group)
```



N2DL-HeLa: Outlier

Overlay of ground truth and test image



- False negatives
- False positives

Additional slide – Histogram stretching

