

Cell nuclei segmentation

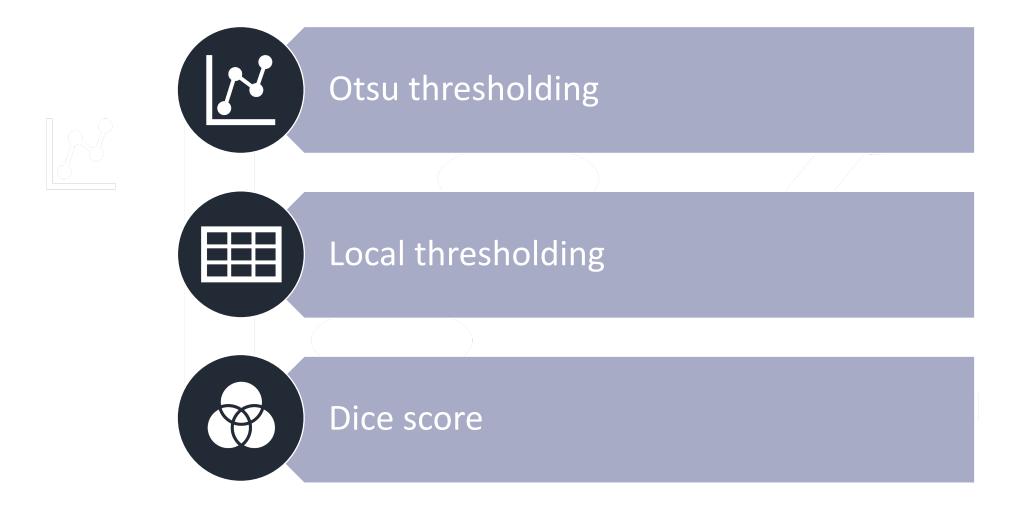
Implementation and evaluation of Otsu thresholding

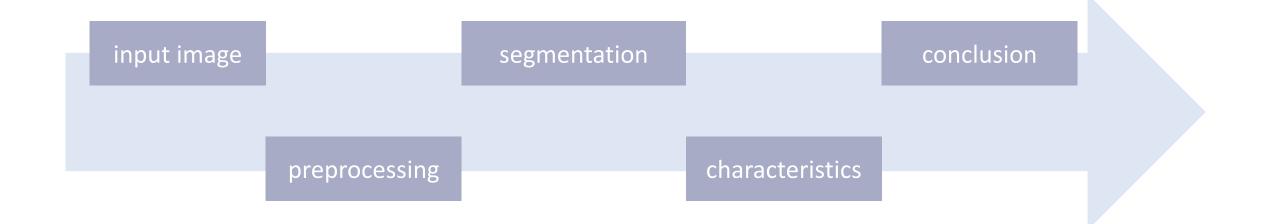
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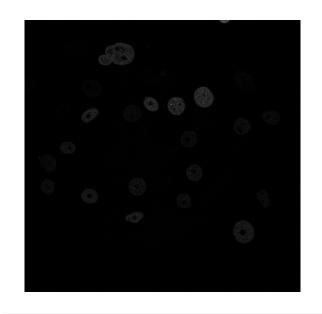
Tutor: Hannah Winte

Software requirements

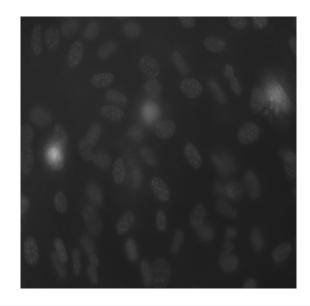




Datasets







	N2DH-GOWT1	N2DL-HeLa	NIH3T3
Organism	mouse (Mus musculus)	human (homo sapiens)	mouse (Mus musculus)
Cell type	embryonic stemm cells	epithelial cells of cervical cancer	embryonic fibroblast cells
Staining	Oct4-GFP	H2b-GFP	Hoechst
Microscope	timelapse confocal microscopy	Olympus lx81 microscope used for live imaging of fluorescently labeled chromosomes	fluorescence microscopy

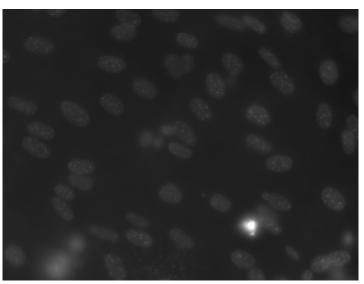
Input image Preprocessing

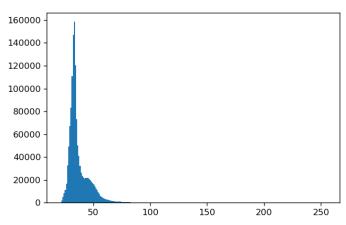
Segmentation

Characteristics

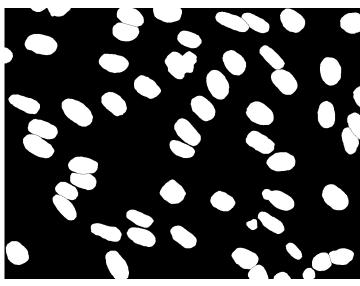
Image types

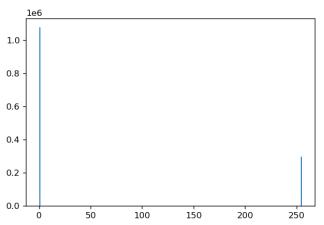
Image

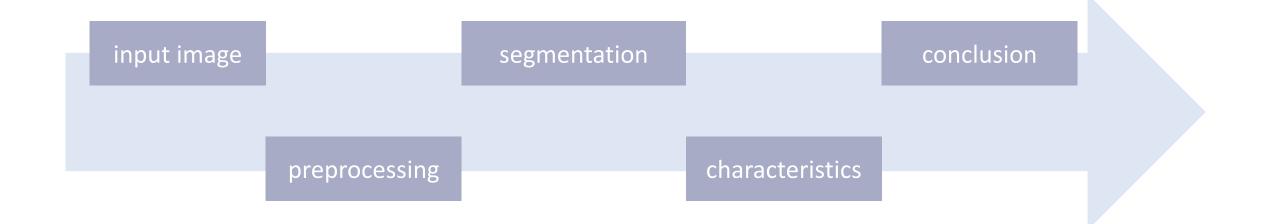




Ground Truth







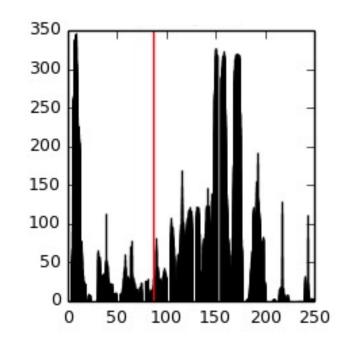
Otsu thresholding

what we need:

what we work with:

what we get:



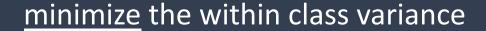




http://sharky93.github.io/docs/gallery/auto_examples/plot_otsu.html

Otsu thresholding





$$\sigma_w^2(t) = w_1(t)\sigma_1^2(t) + w_2(t)\sigma_2^2(t)$$

σ: variance of class

w: probabilities of the two classes

divided by a threshold

t: threshold

maximize the between class variance

$$\sigma_b^2(t) = w_1(t)w_2(t)[\mu_1(t) - \mu_2(t)]^2$$

σ: variance of class

w: probabilities of the two classes

divided by a threshold

t: threshold

μ: mean of class

Otsu thresholding



Process the input grayscale image



Generate a distribution of the pixel intensities (histogram)



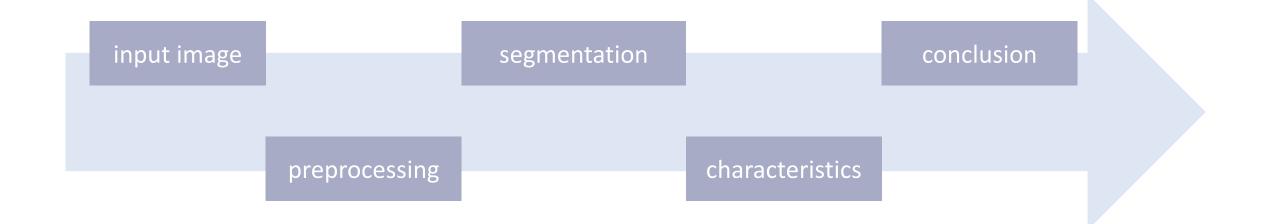
Calculate the threshold value t



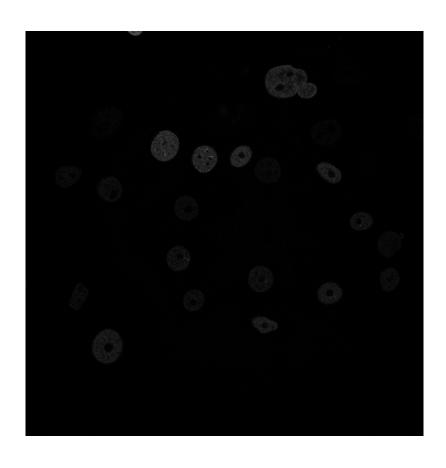
Replace the pixels values with white (>t) an black (<t)

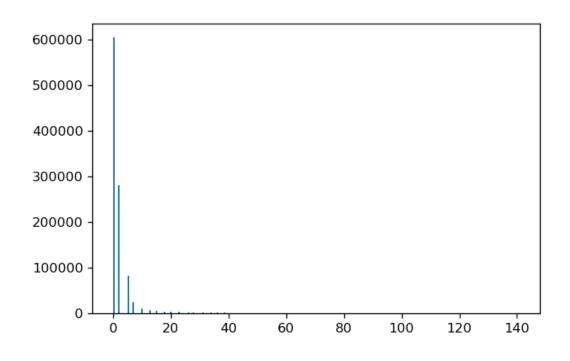
Conditions for optimal Otsu threasholding

- high contrast between objects and background
- bimodal distribution
- minimal image noise
- similar intensities of objects



Challenges: N2DH-GOWT1

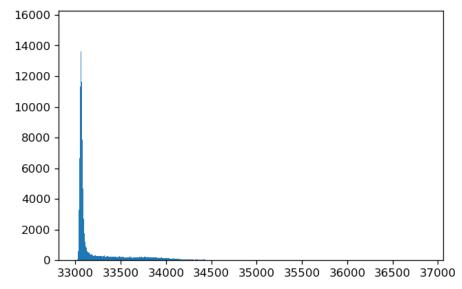




- noise
- low contrast

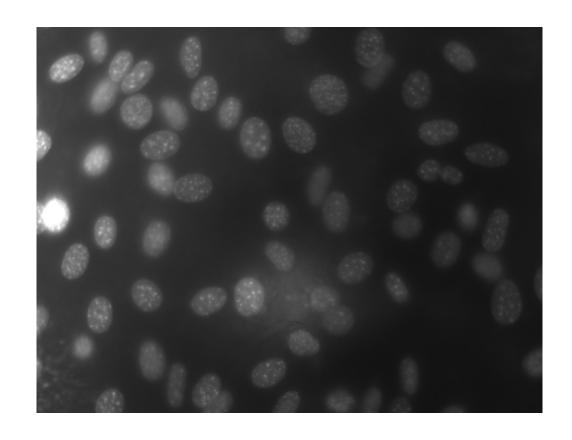
Challenges: N2DL-HeLa

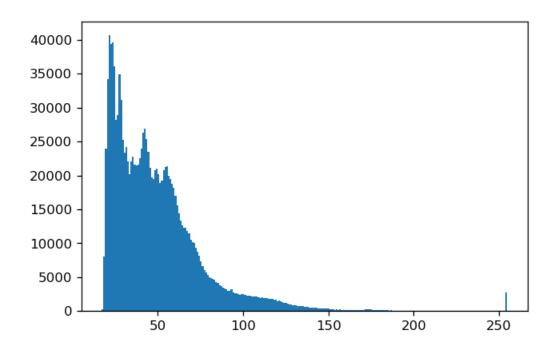




low contrast

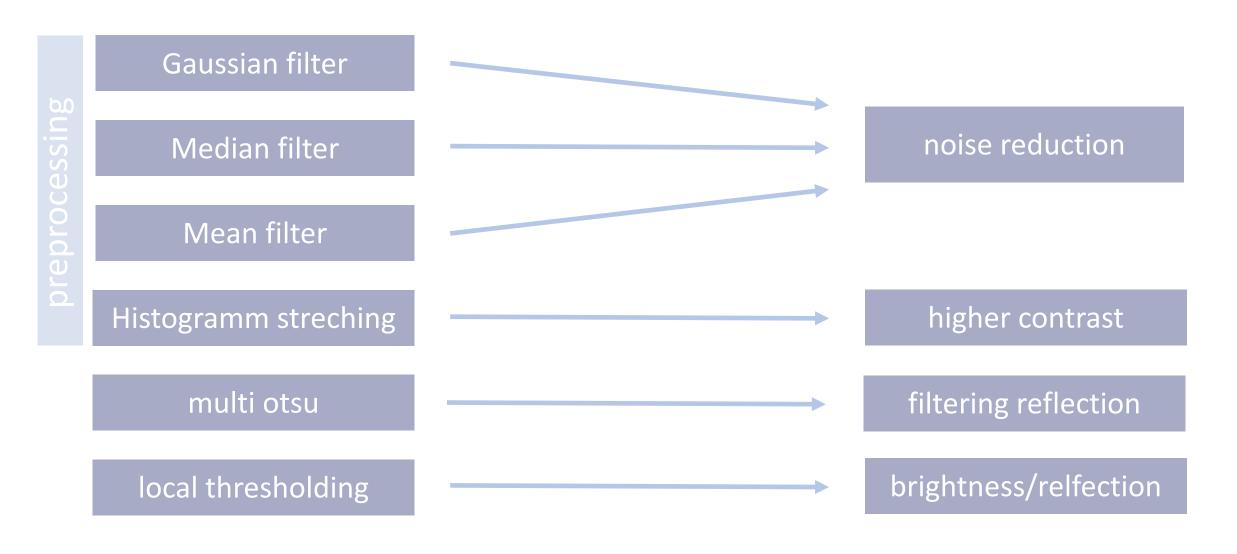
Challenges: NIH3T3





- varying brightness
- reflections

How to master these challenges?



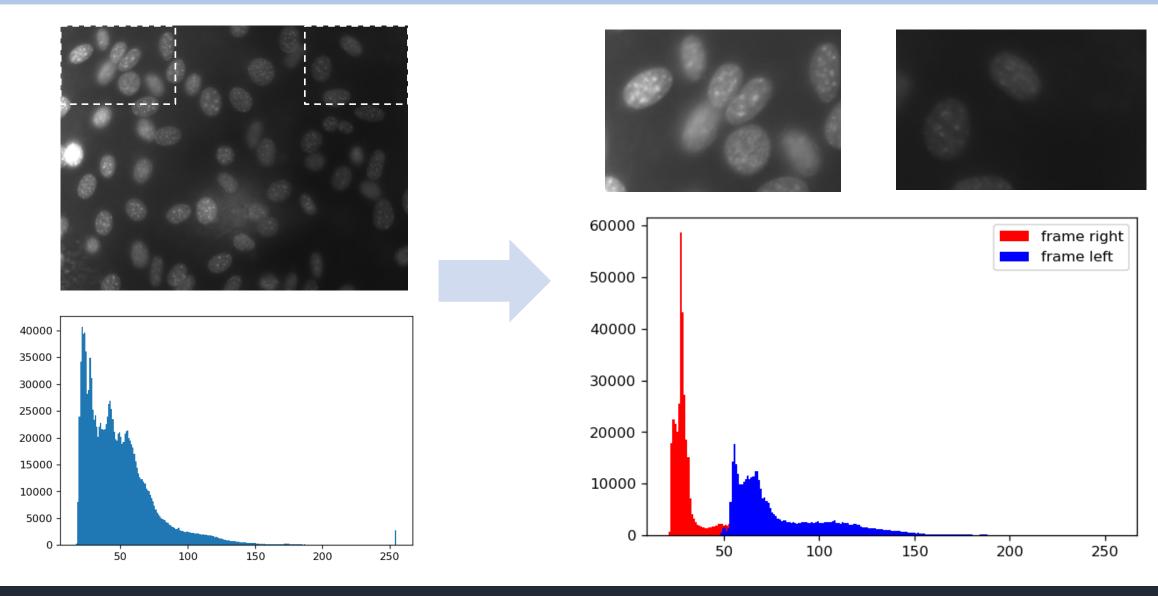
Input image

Preprocessing

Segmentation

Characteristics

Local thresholding



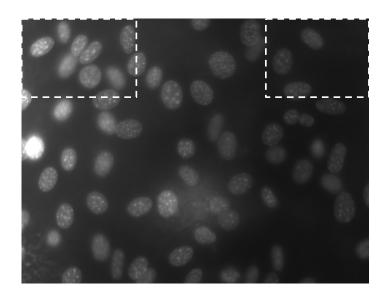
Input image

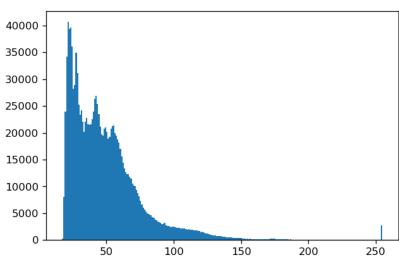
Preprocessing

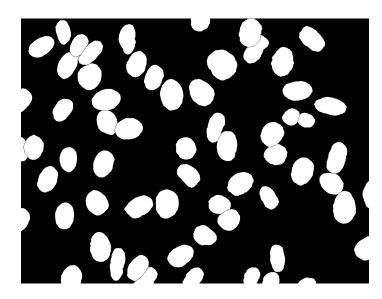
Segmentation

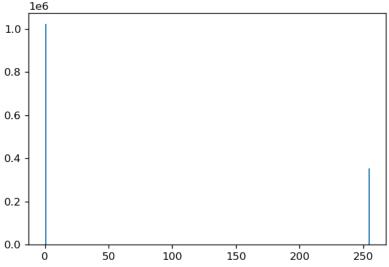
Characteristics

Local thresholding







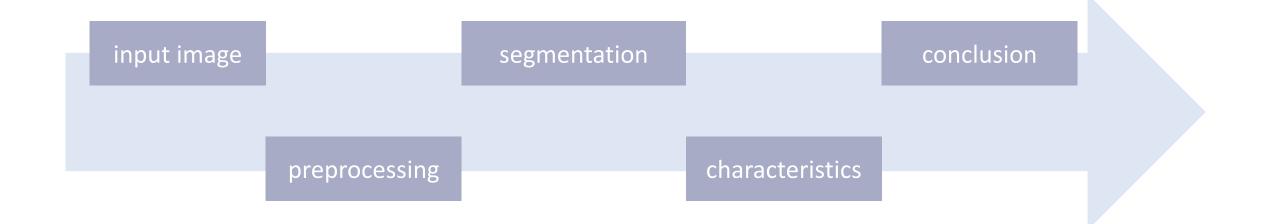


Input image

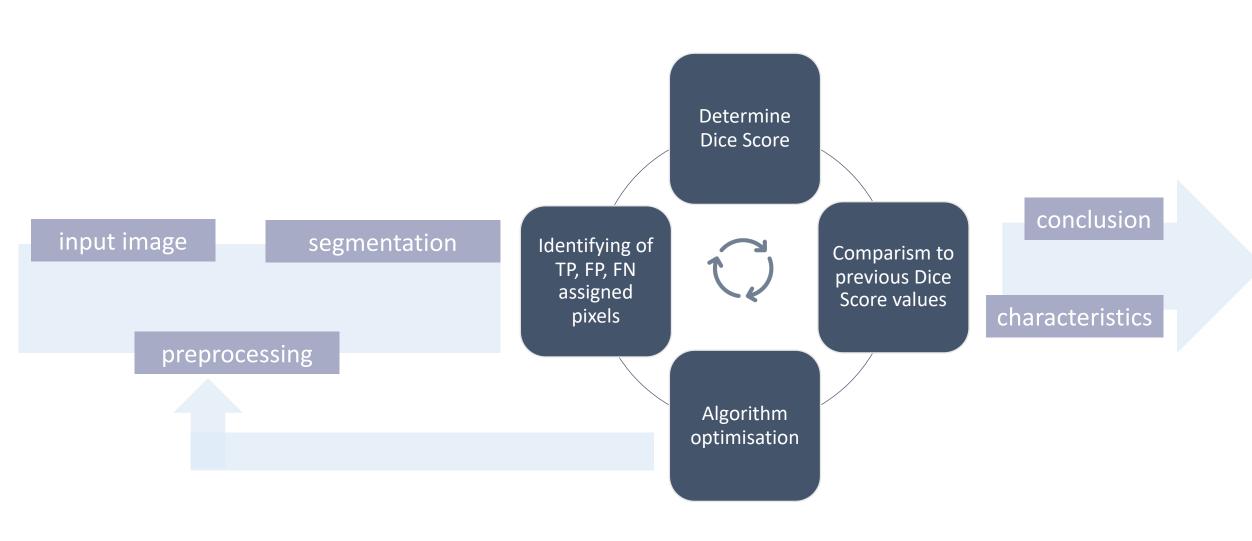
Preprocessing

Segmentation

Characteristics



Usage of Dice Score



Dice Score

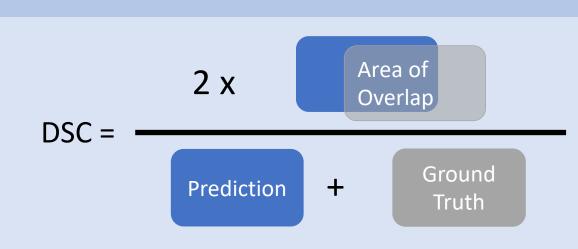
Measure of similarity between two sets of data



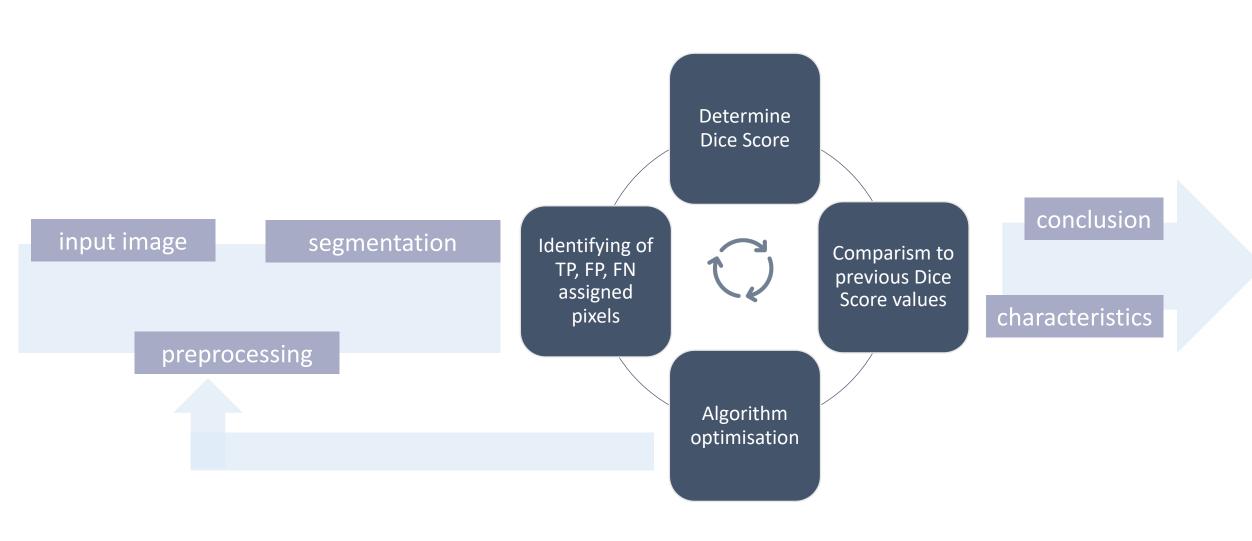
used to evaluate the similarity between a predicted segmentation mask and the ground truth segmentation mask

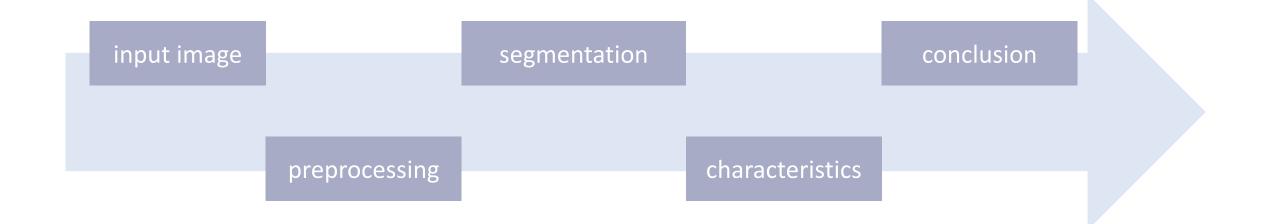
Boolean Data

Discrete Data



Usage of Dice Score





Additional ideas



Cell counting



Cell Size

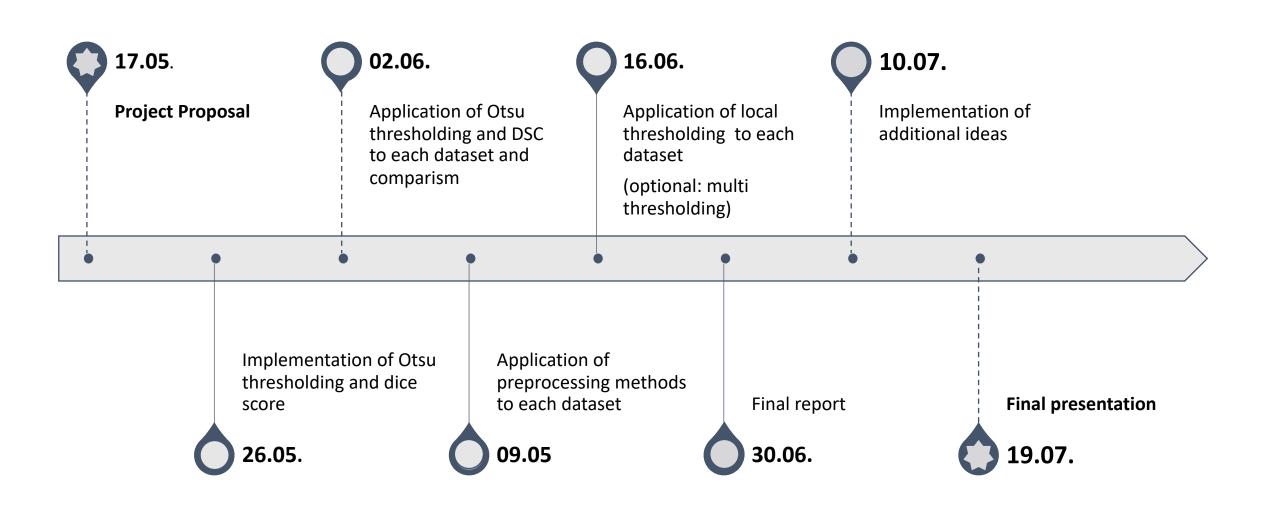


Cell arrangement



Other evaluation methods

Timeline



Python packages

- numpy (numerical python) → to work with numerical data
- skimage (scikit-image) → for image processing
- cv2 (OpenCV) → for solving computer vision problems
- matplotlib.pyplot → to visualize our data

→ and probably many more that we will discover while working on our project





