

# Vascular Segmentation of Renal Cell Carcinoma



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Bioinformatics A.A. 2021/2022

# Overview

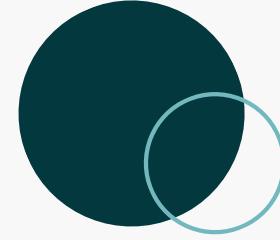
- Dataset analysis and preprocessing techniques

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- Evaluation metrics

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- Segmentation models
  - Autoencoder
  - U-Net
  - Pix2Pix GAN

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- Results and conclusions

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- Next works



# Dataset

Composed of 210 histological images (2000x2000) with associated segmentation mask, depicting two types of cancer:

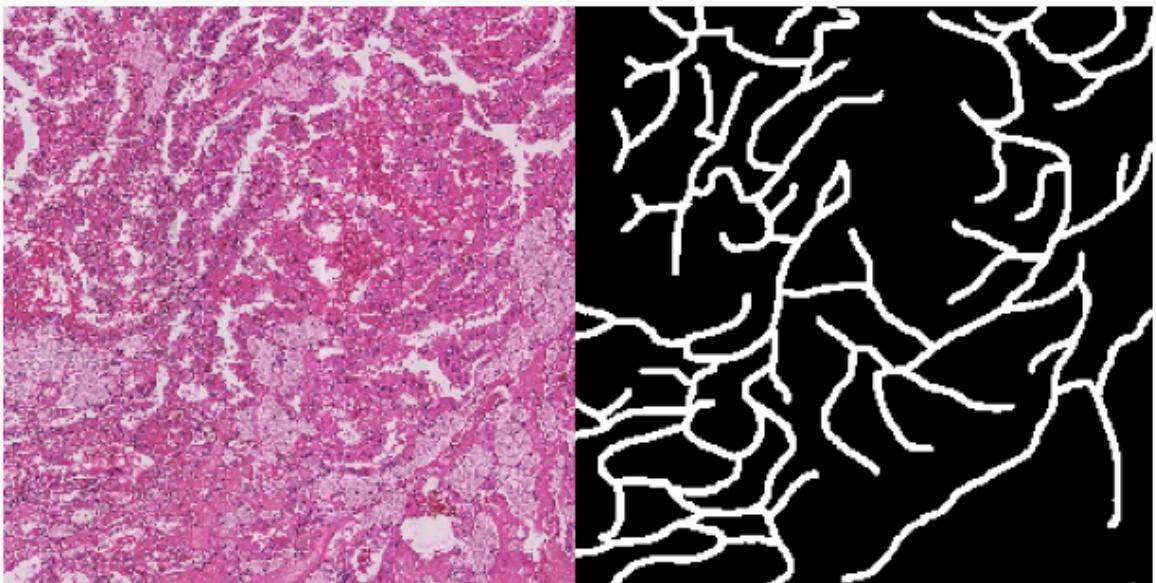
## Papillary Renal Cell Carcinoma (pRCC)

### TRAIN

100 images

### TEST

5 images



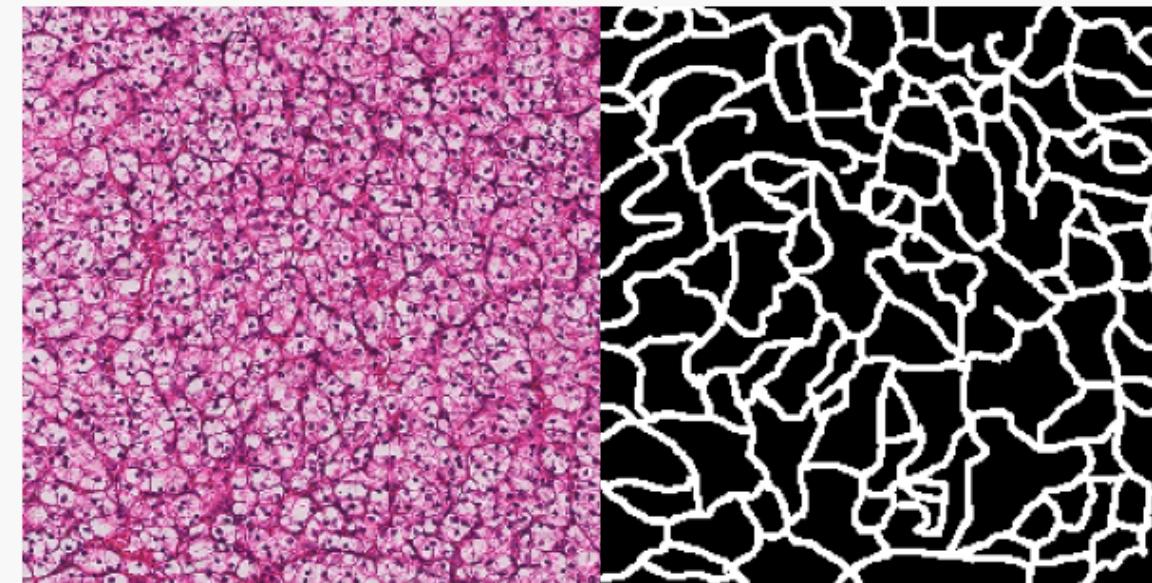
## Clear Cell Renal Cell Carcinoma (ccRCC)

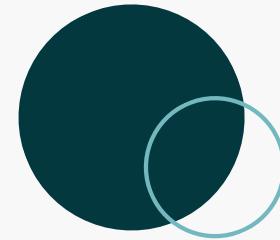
### TRAIN

100 images

### TEST

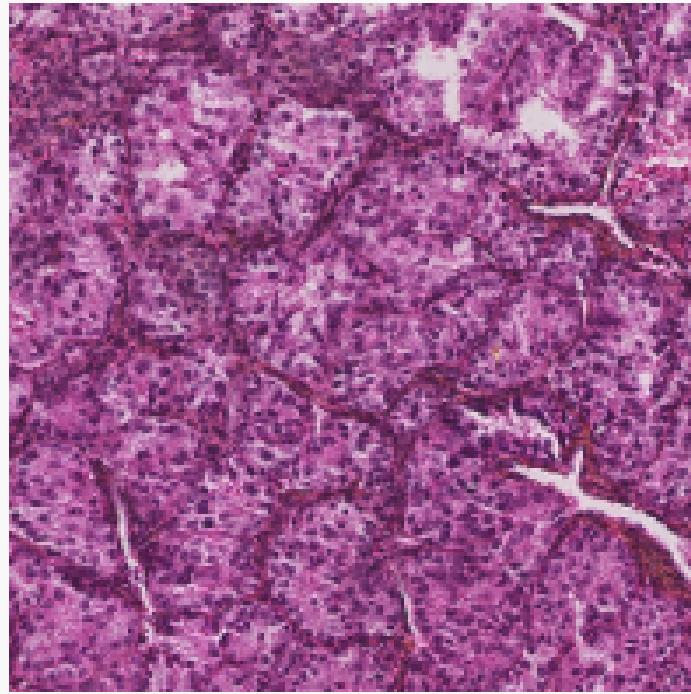
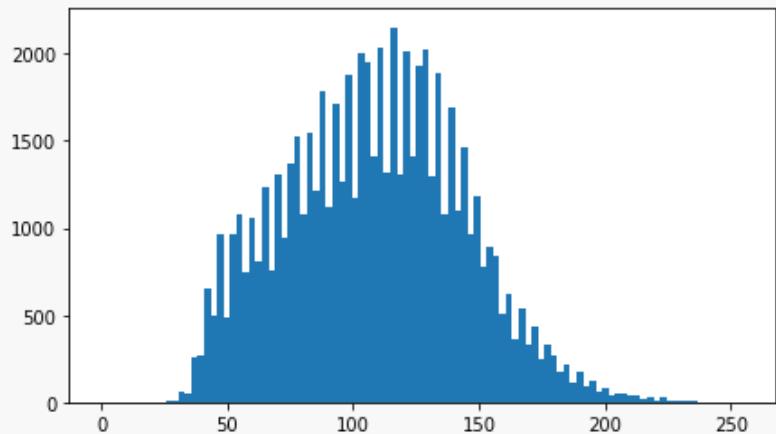
5 images





# Preprocessing - Contrast

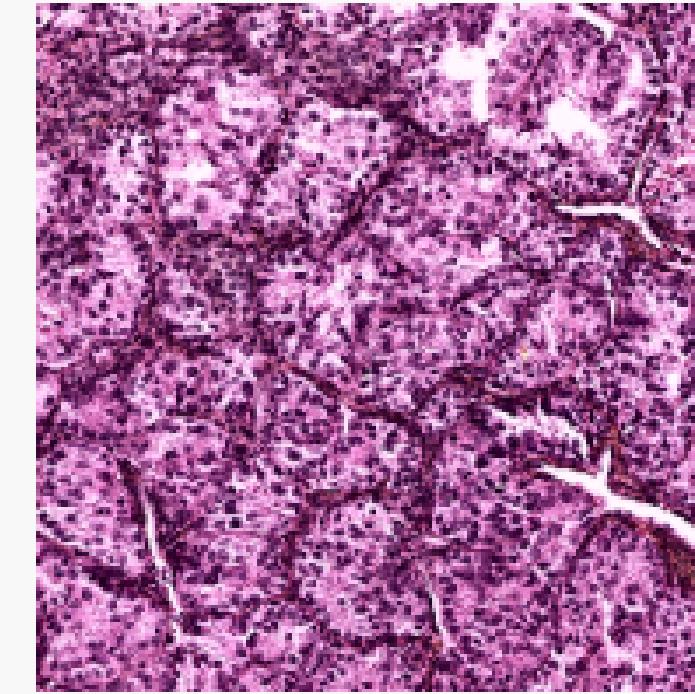
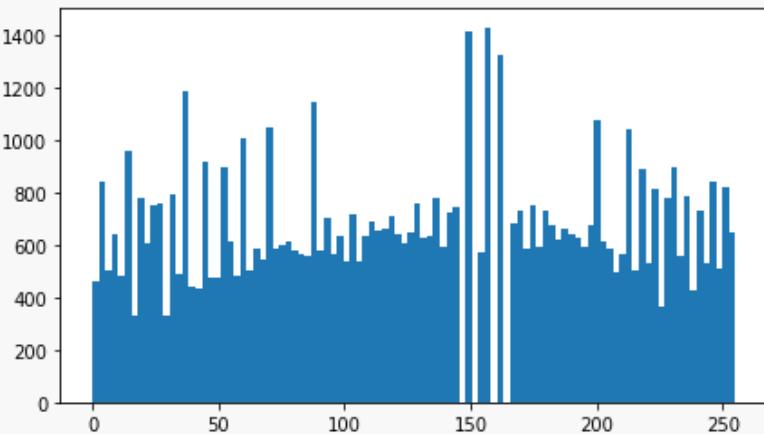
Original Image



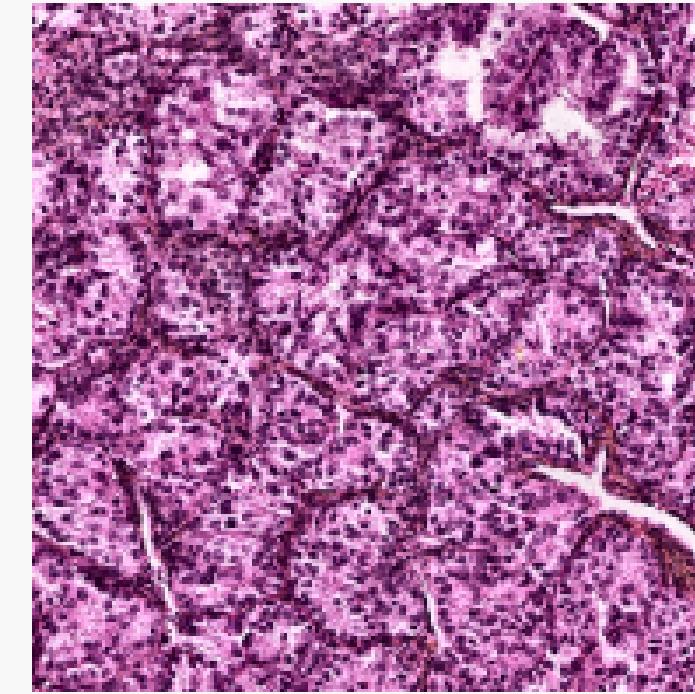
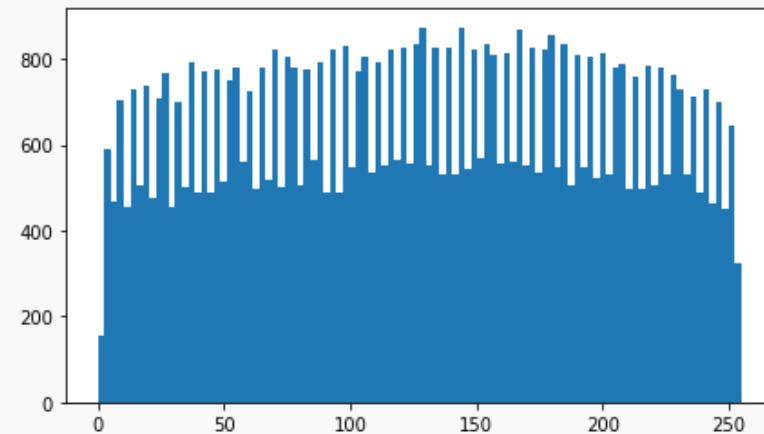
Improve  
contrast

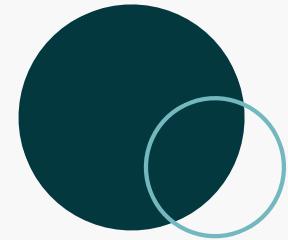


Histogram Equalization

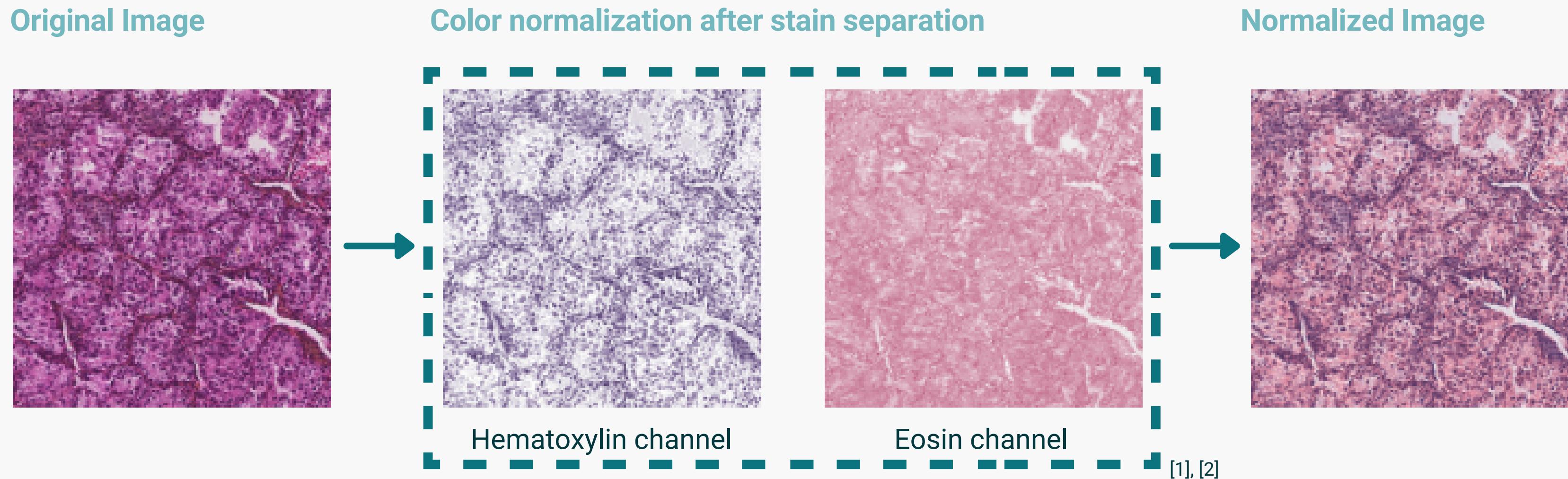


Contrast Limiting Adaptive  
Histogram Equalization (CLAHE)

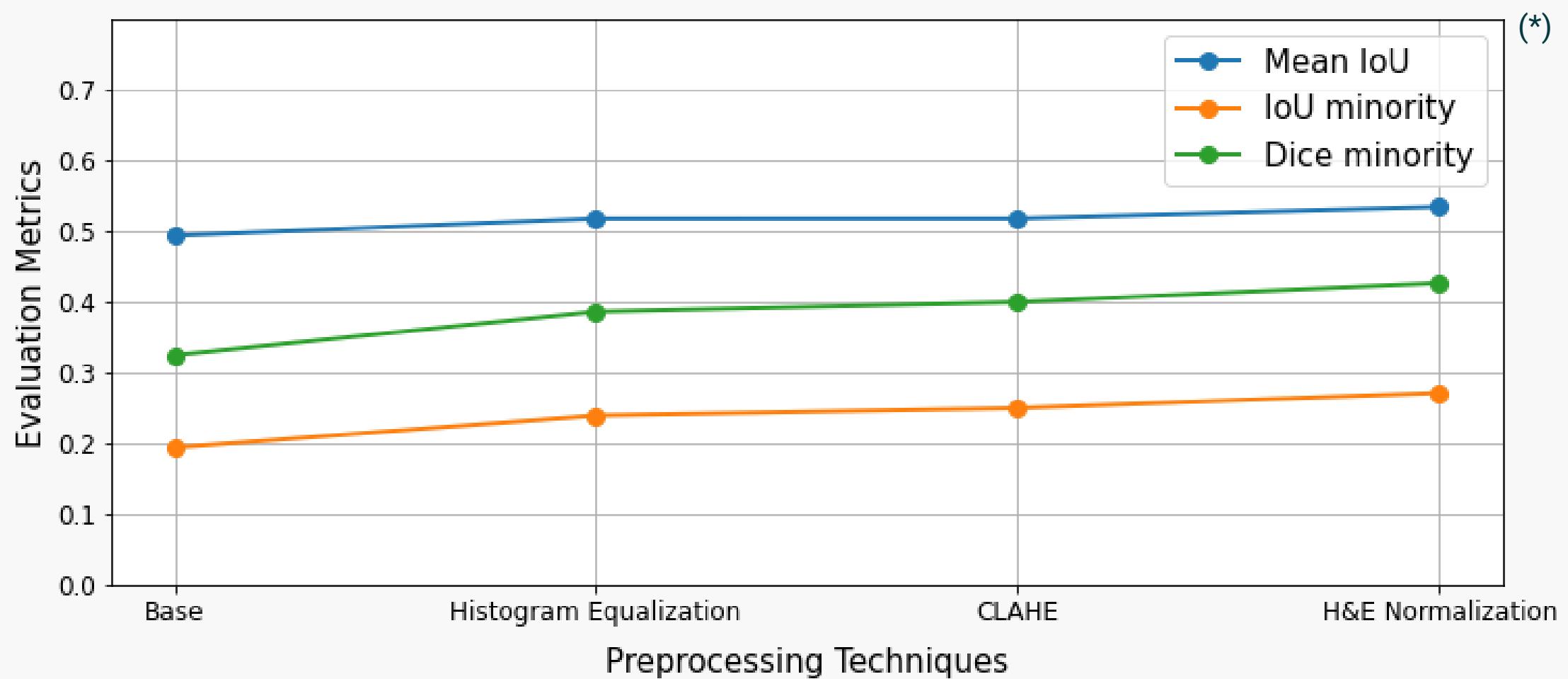




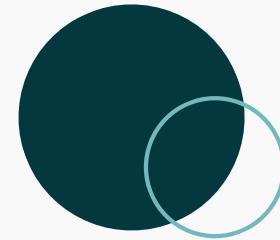
# Preprocessing - Stain



# Preprocessing - Results



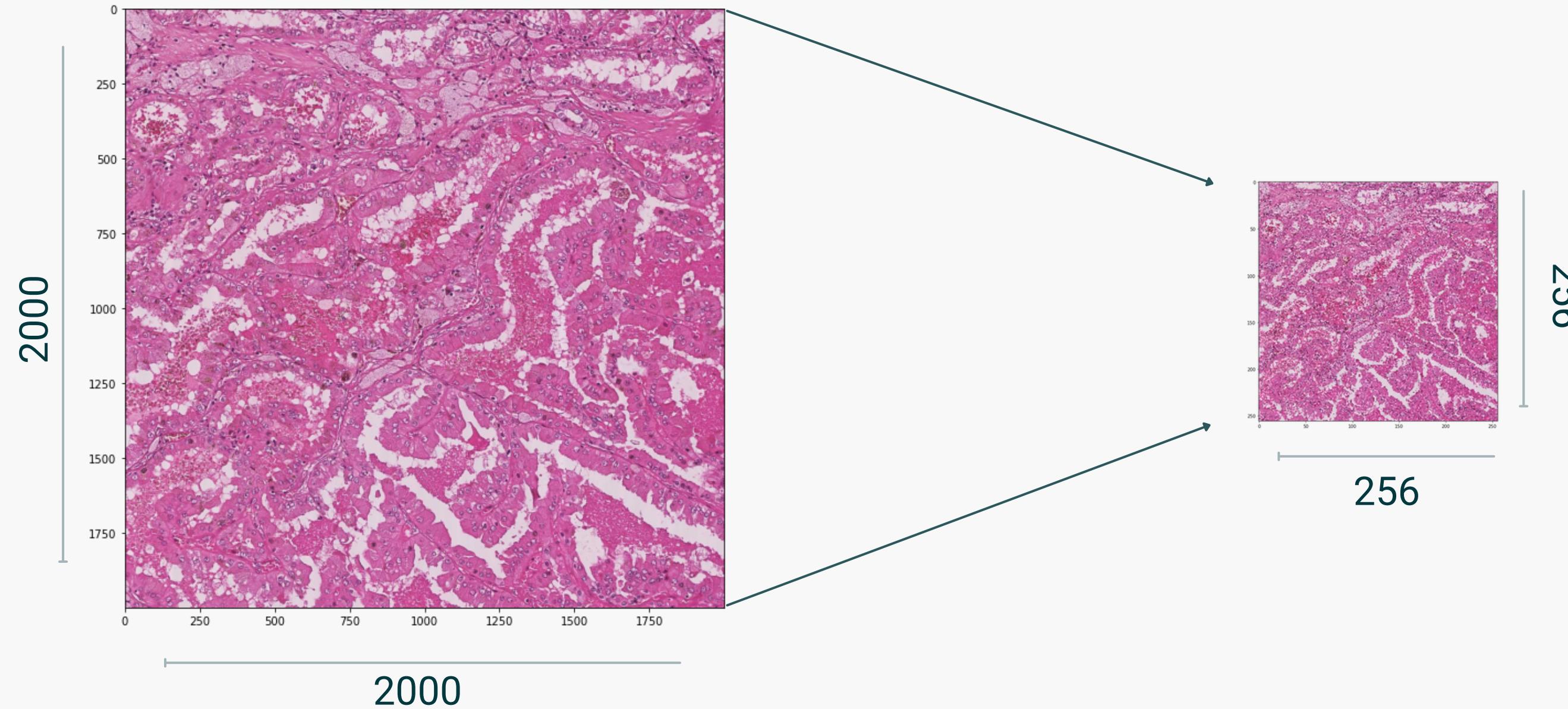
(\*) Results refer to pRCC dataset through Autoencoder

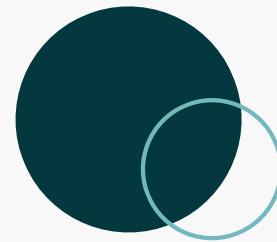


# Preprocessing - Resize

1

Resize 256x256

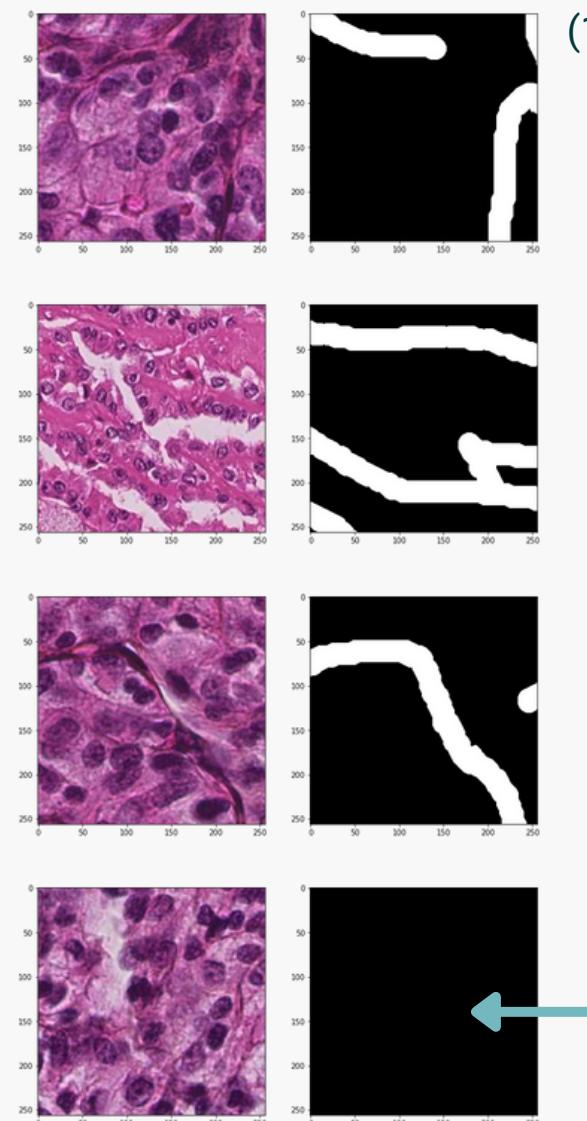




# Preprocessing - Patches

2

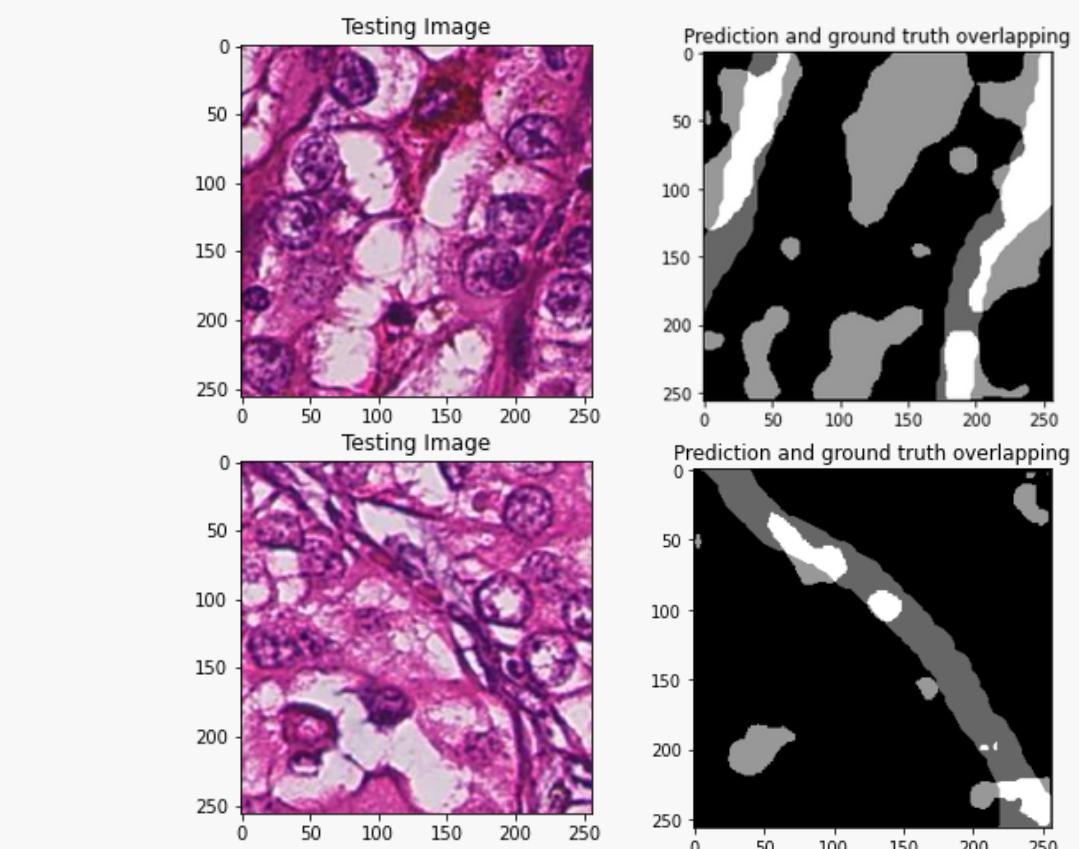
## Patchify



From 100 pRCC samples to **4900** pRCC patches

Predict

Totally black  
[2]



Dataset	Epochs	Batch size	Learning Rate	Mean IoU	IoU minority	Dice minority
Original pRCC	1000	8	0.001	0.53379	0.27096	0.42638
Patches pRCC	1000	64	0.0001	0.42382	0.15898	0.27434

(2\*)

(1\*) Patches from pRCC dataset

(2\*) Results obtained through Autoencoder

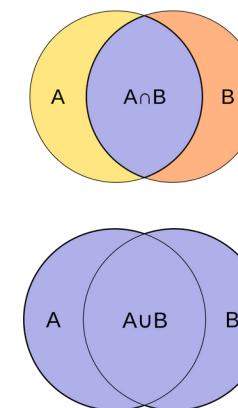
# Evaluation Metrics

## 1 IoU - Intersection over Union

Measures similarity between images. It is defined as the area of the intersection divided by the area of the union between predicted and ground truth segmentations.

### IoU

$$IoU = \frac{OverlapArea}{UnionArea}$$



### IoU - Jaccard coefficient

$$IoU = \frac{truePositives}{truePositives + falsePositives + falseNegatives}$$

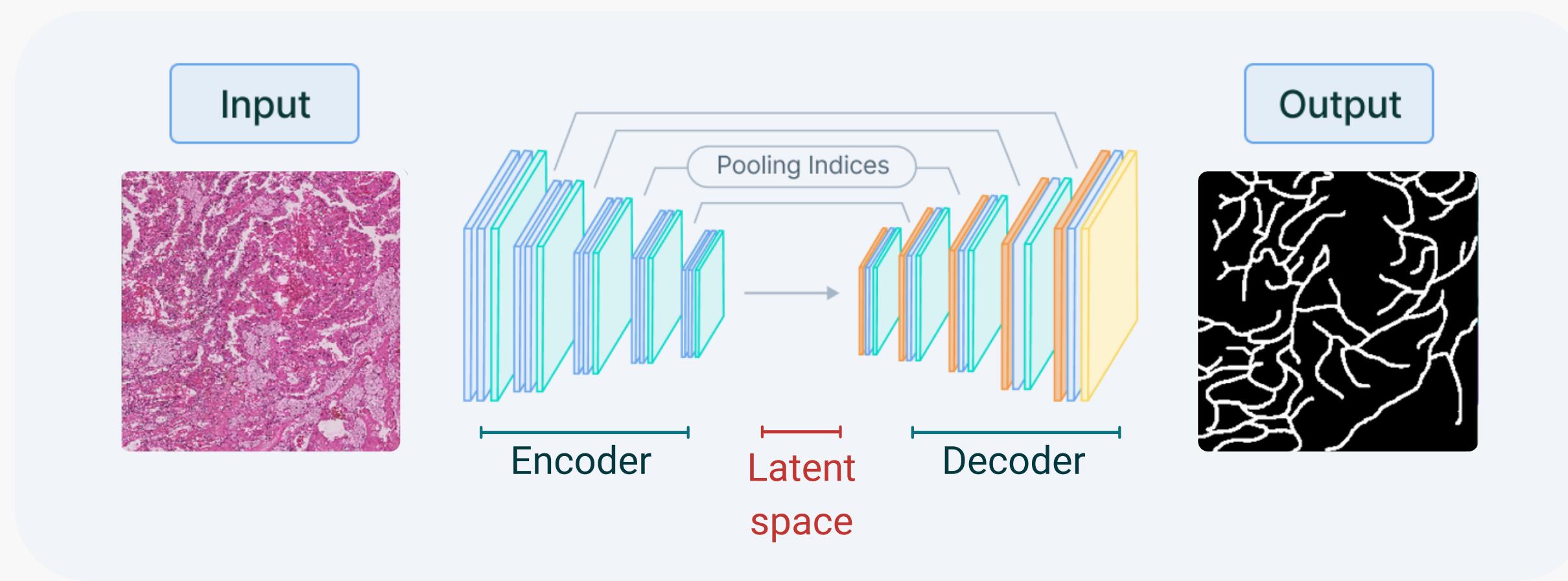
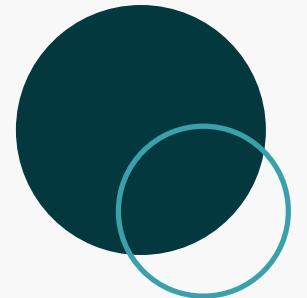
## 2 Dice

Similar to IoU but composed of the double intersection over the sum between intersection and union.

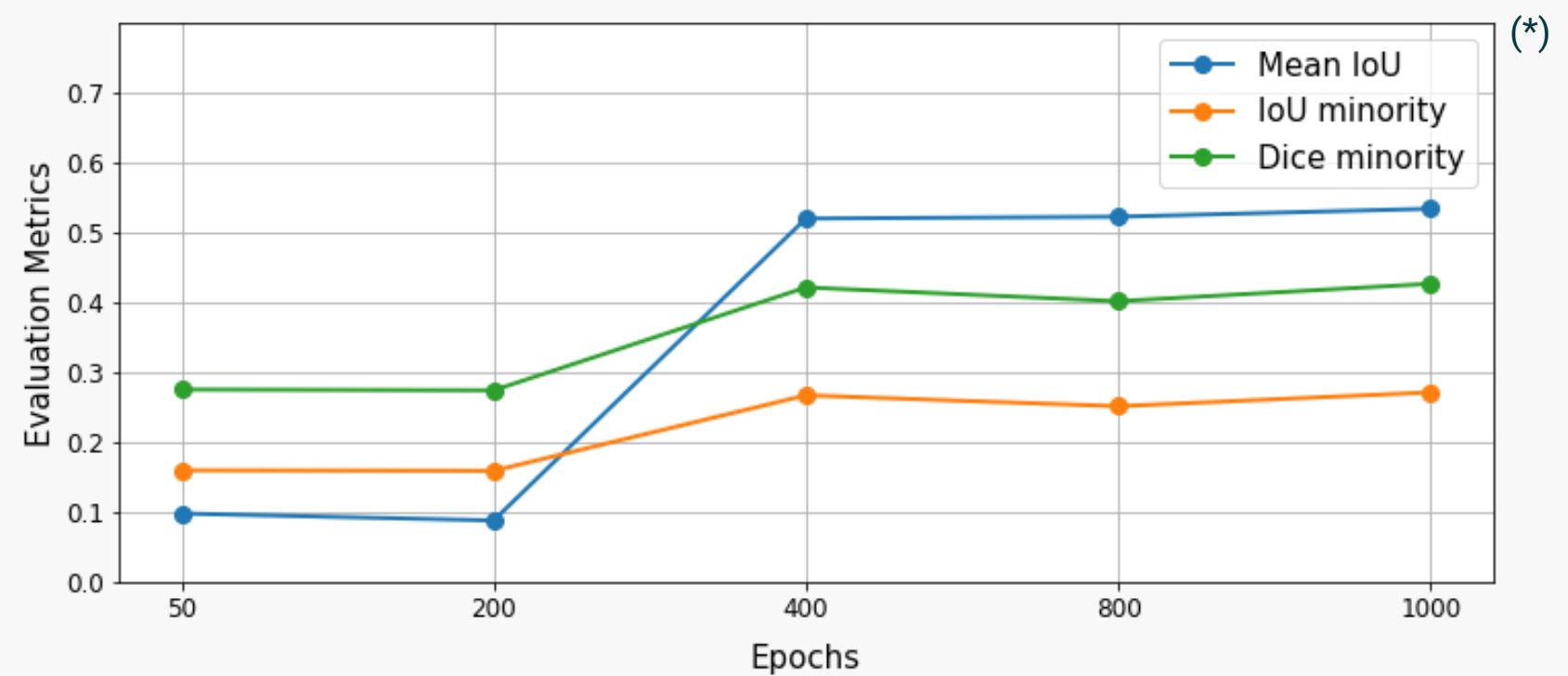
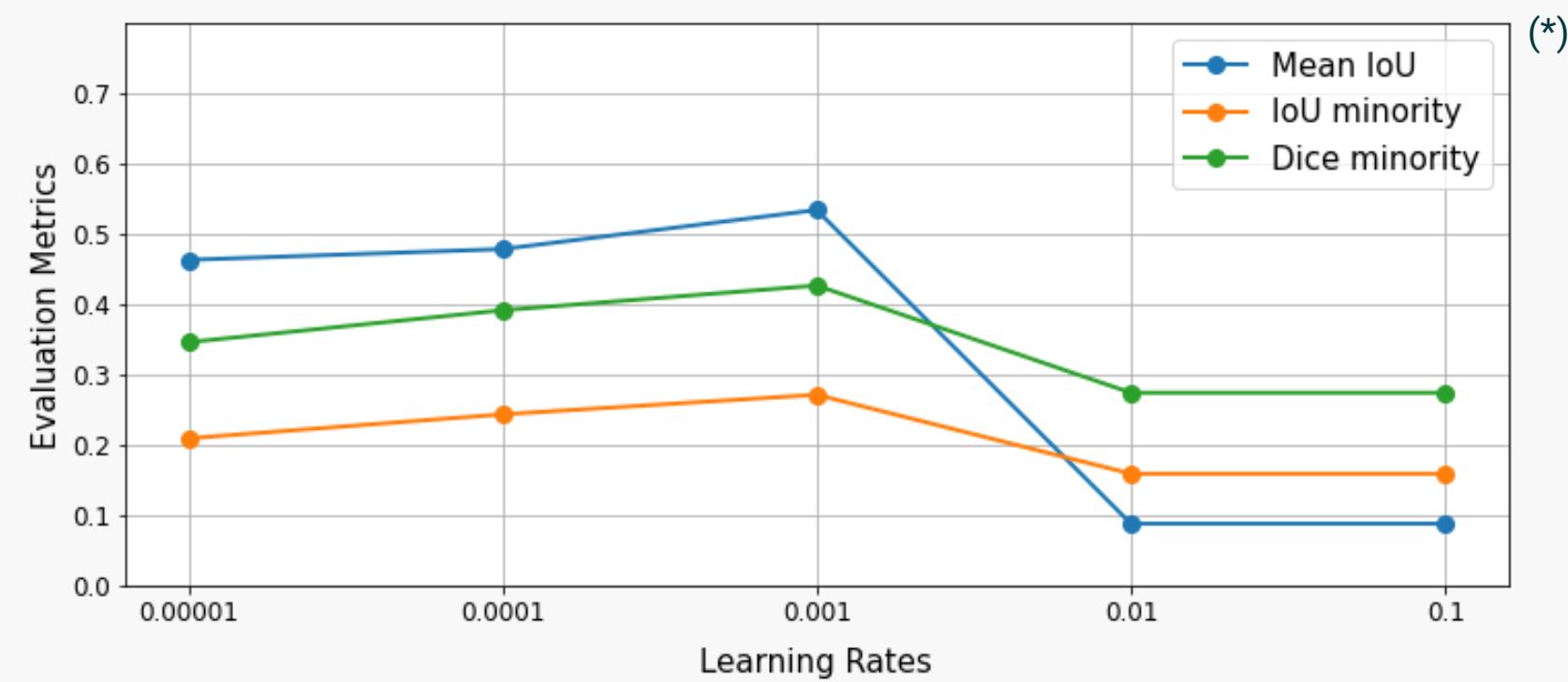
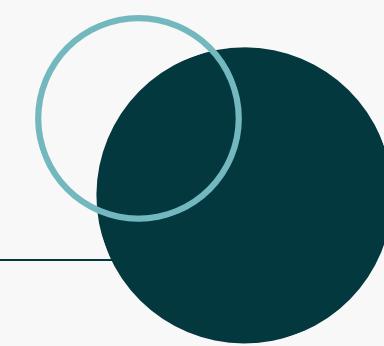
### Dice score

$$Dice = 2 \frac{truePositives}{(truePositives + falsePositives) + (truePositive + falseNegatives)}$$

# Autoencoder - Structure

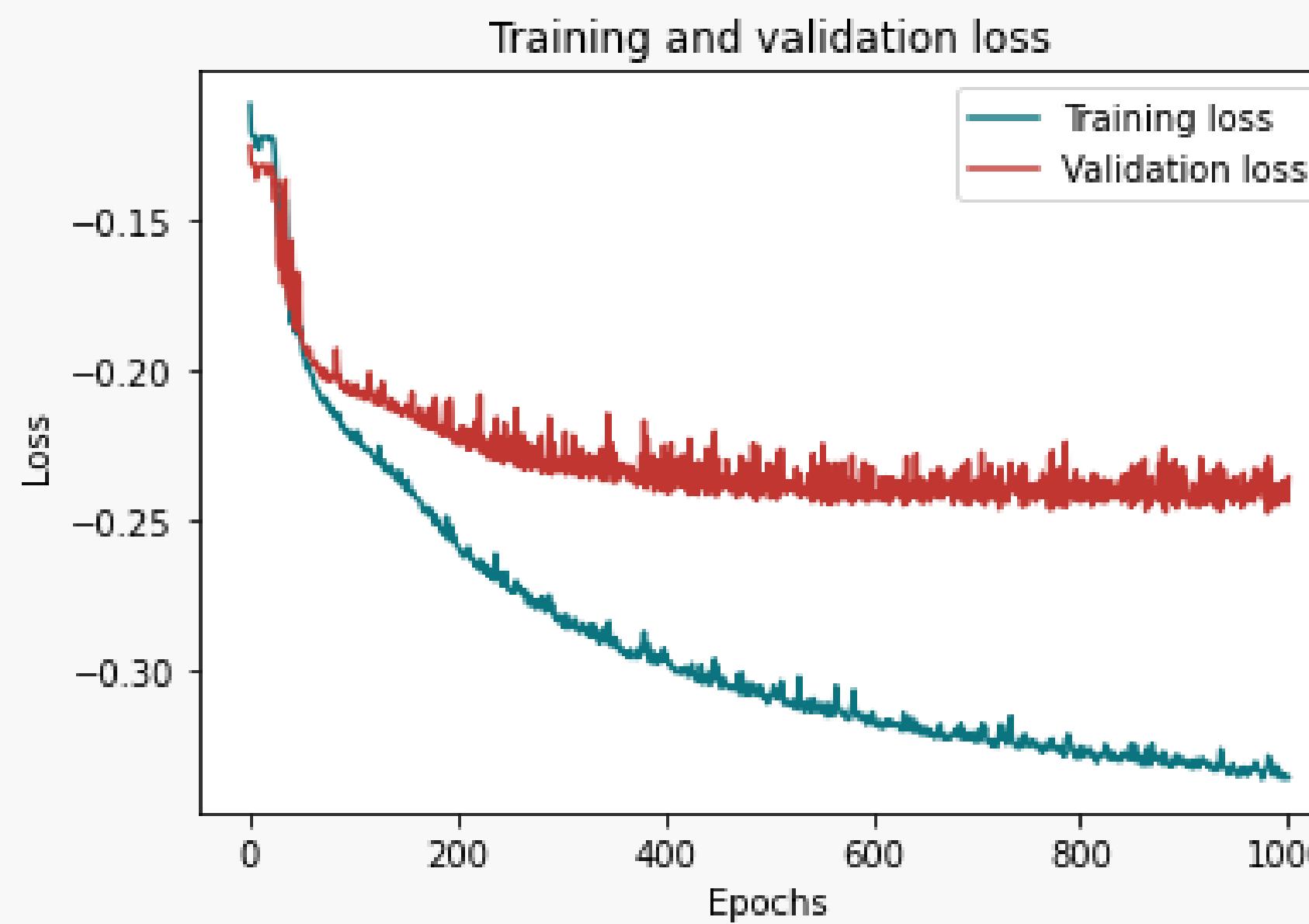
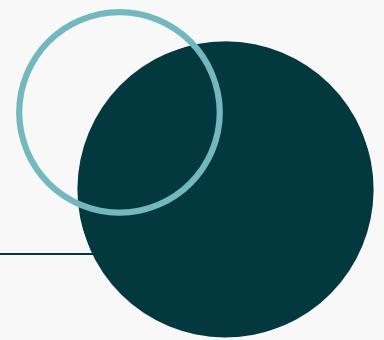


# Autoencoder - Tuning



(\*) Results refer to pRCC dataset

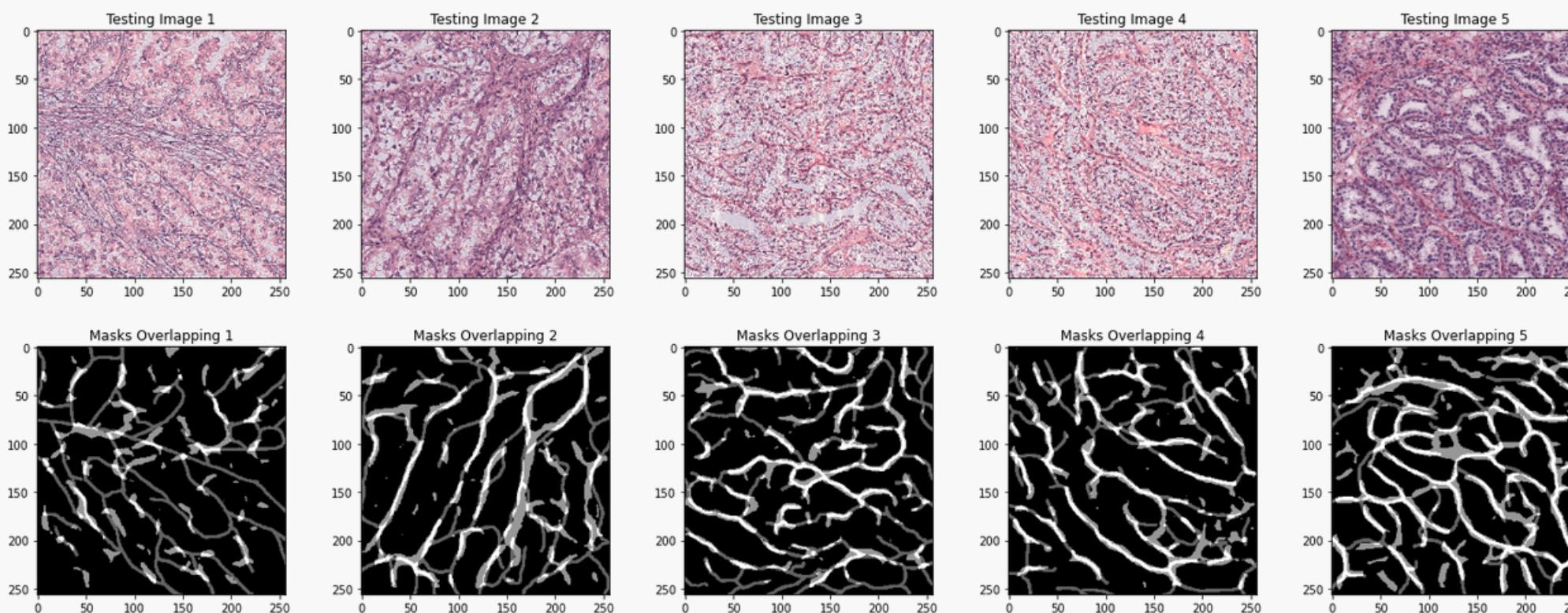
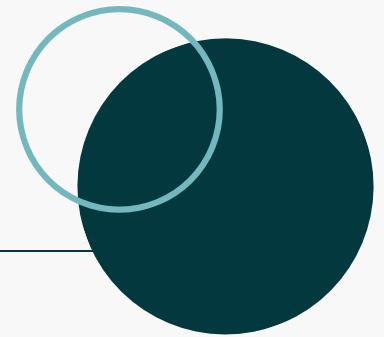
# Autoencoder - Loss



**DATASET** : pRCC - resize  
**PREPROCESSING**: H&E norm  
**LR**: 1e-3  
**BATCH**: 8  
**TEST MEAN IOU**: 0.5338

$$IoU_{Loss} = - \sum_{n=1}^N \frac{\hat{y}_n y_n}{\hat{y}_n + y_n - (\hat{y}_n y_n)}$$

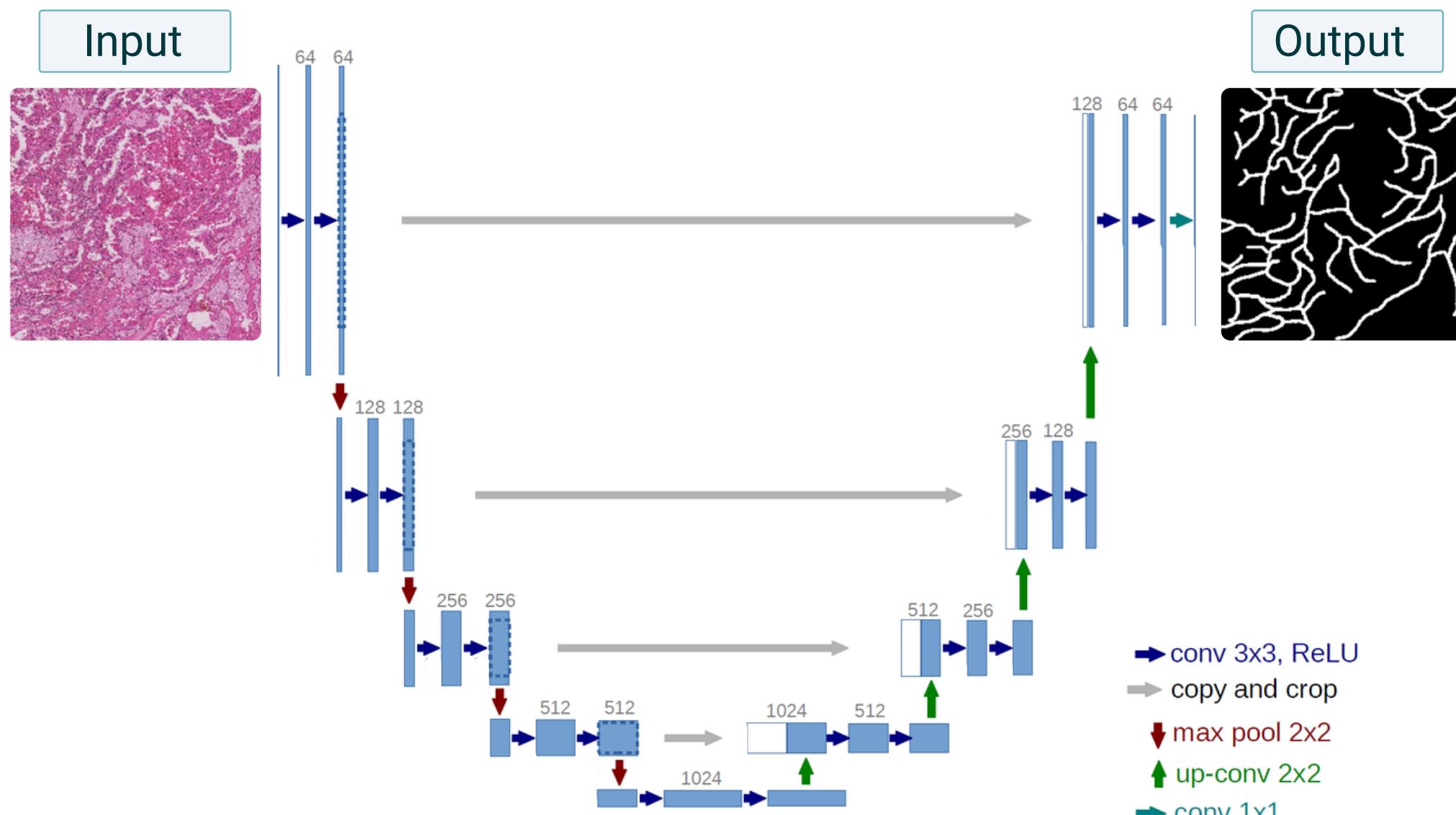
# Autoencoder - Best Results



**DATASET :** pRCC - resize  
**PREPROCESSING:** H&E norm  
**LR:** 1e-3  
**BATCH:** 8  
**LOSS:** Jaccard Loss

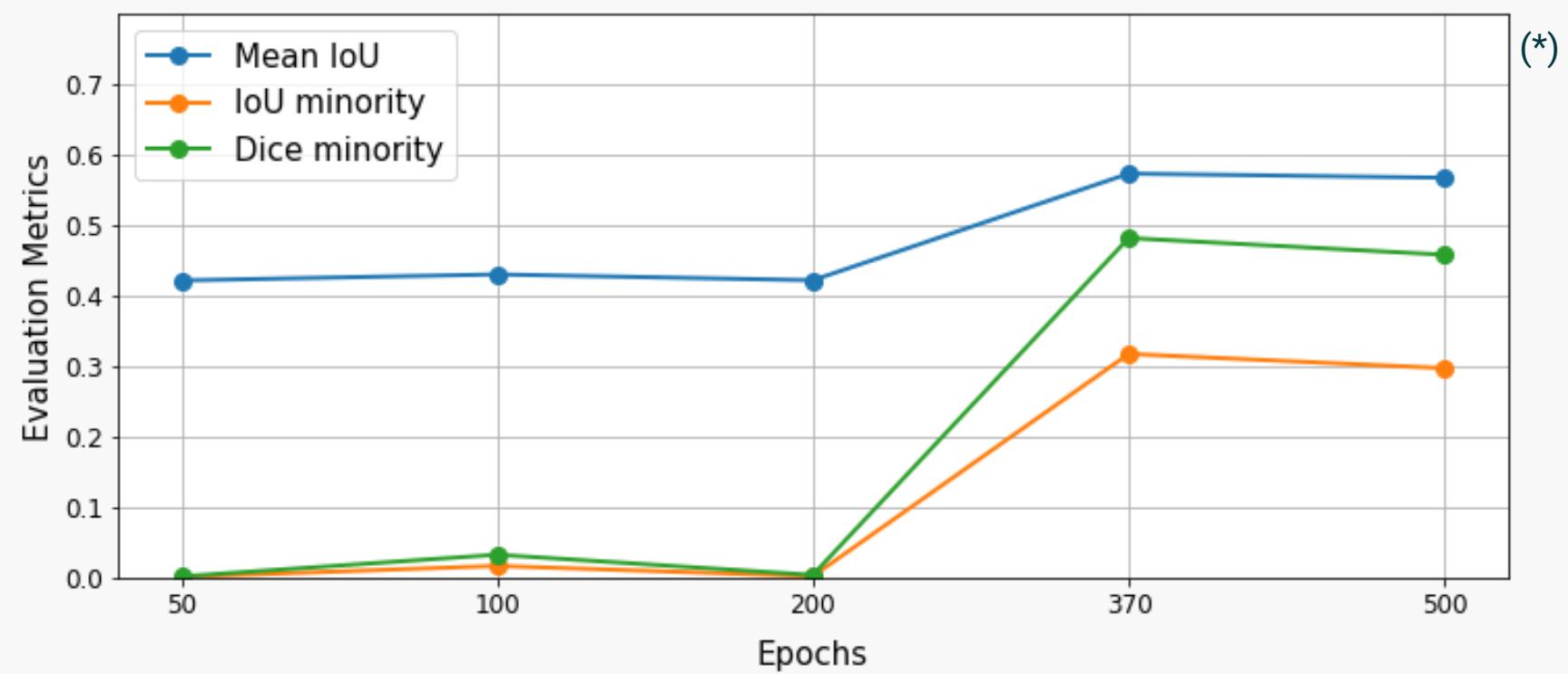
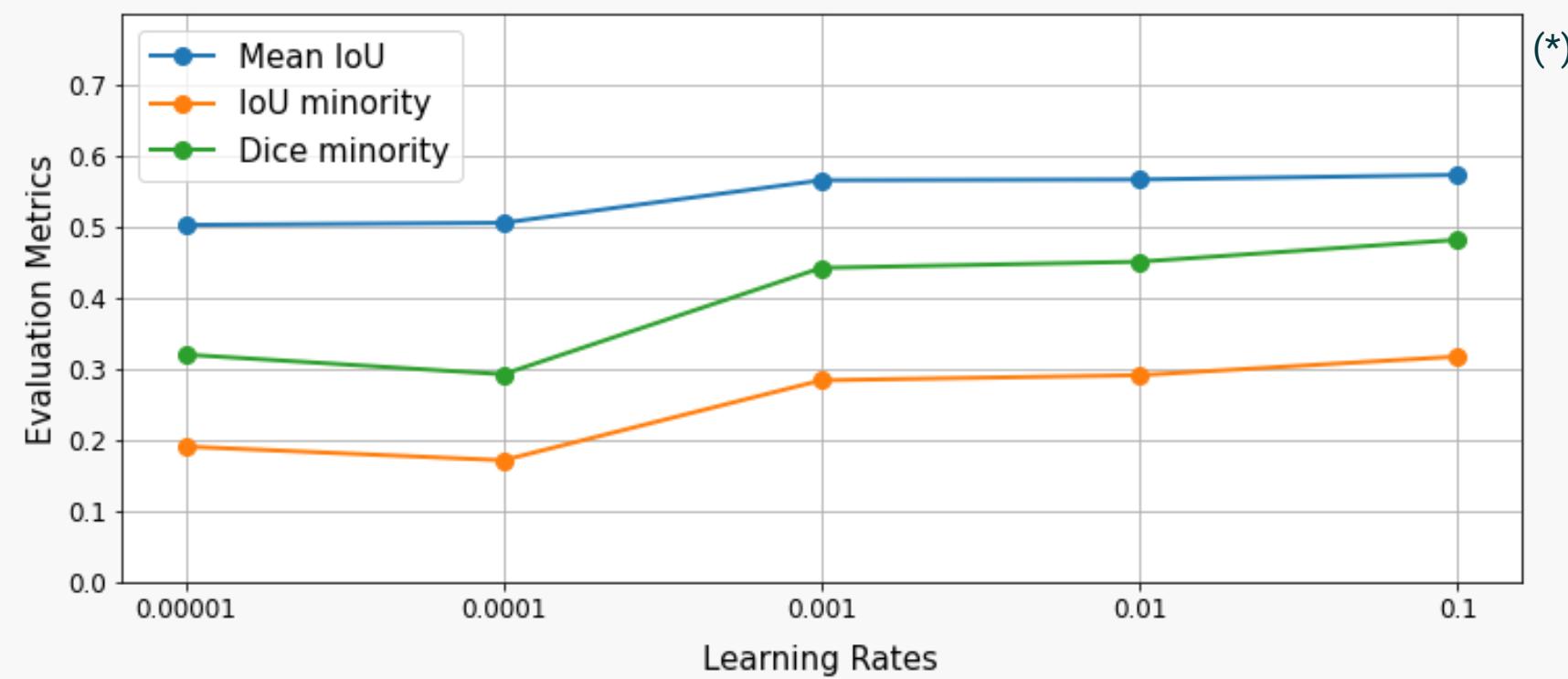
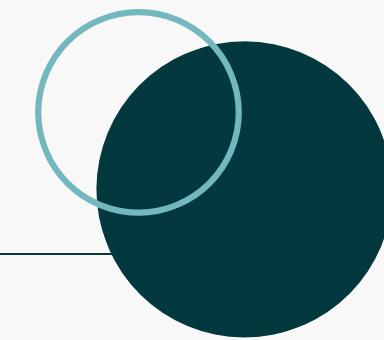
Dataset	Mean IoU	IoU Minority	Dice Minority
pRCC	0.53378814	0.2709558996564221	0.4263812768478744
ccRCC	0.5476335	0.37982286231997253	0.5505385838894643
pRCC+ccRCC	0.3365001	0.2234816844656442	0.36532085000234366

# U-Net - Structure



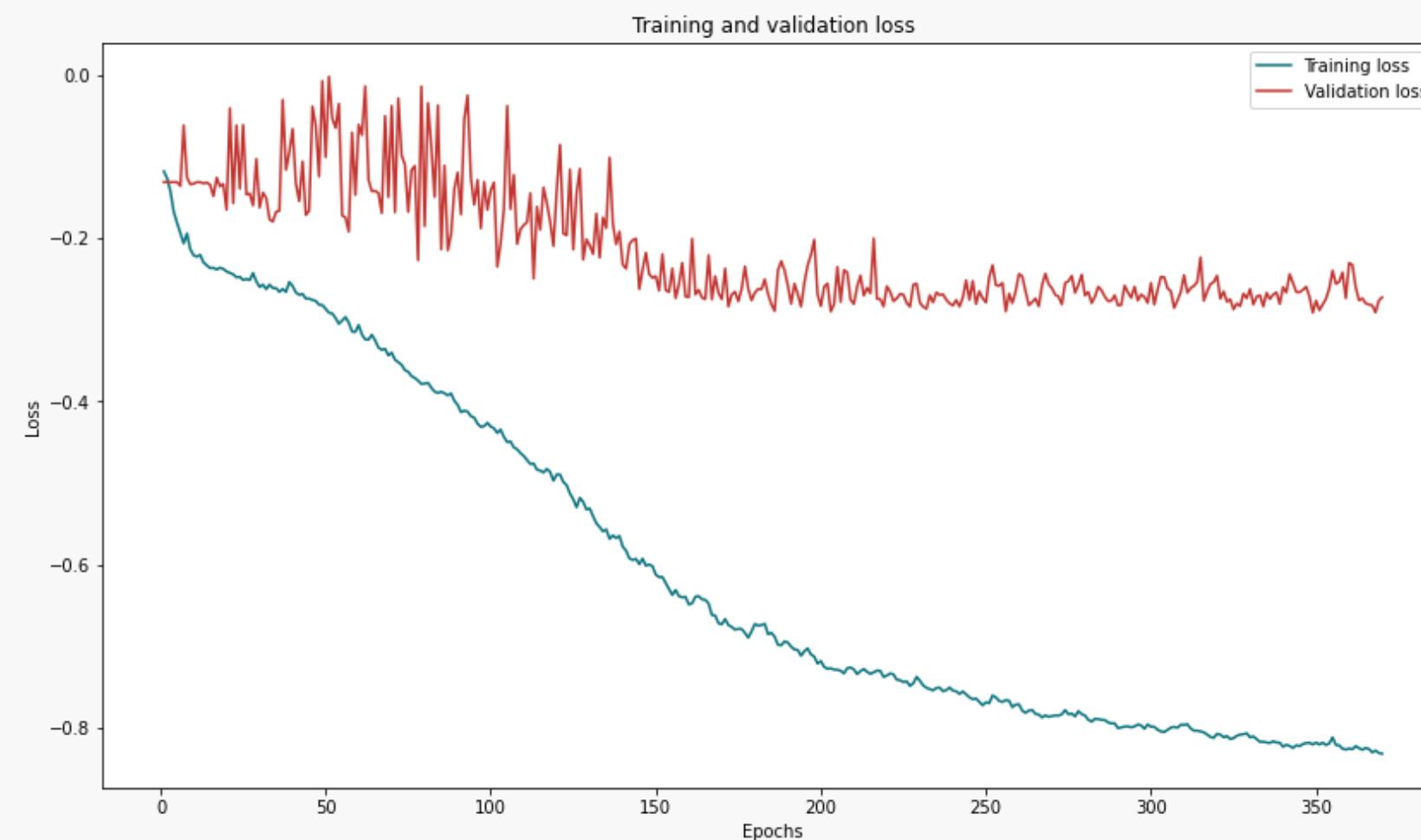
[3]

# U-Net - Tuning



(\*) Results refer to pRCC dataset

# U-Net - Loss

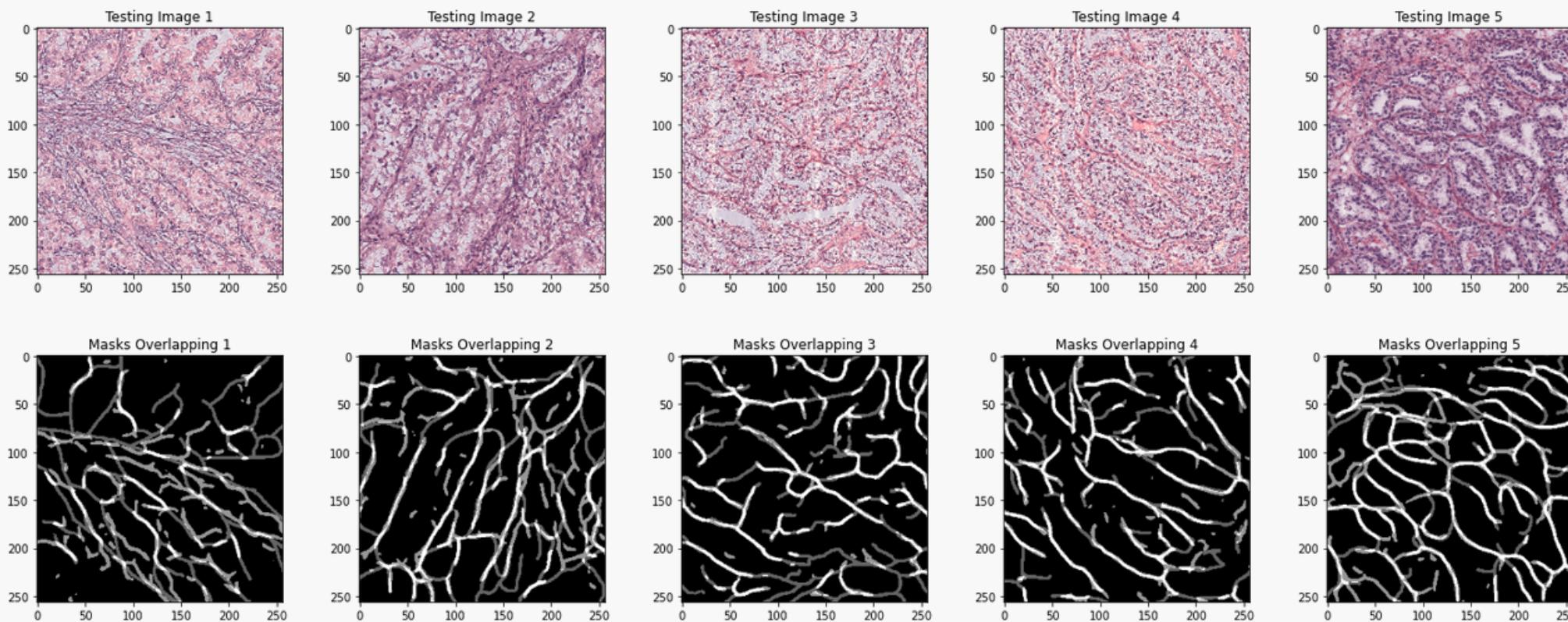
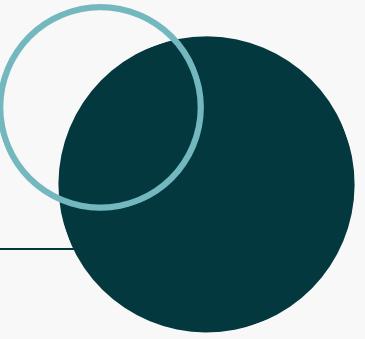


**DATASET** : pRCC - resize  
**PREPROCESSING**: H&E norm  
**LR**: 1e-1  
**BATCH**: 16  
**TEST MEAN IOU**: 0.5730

## Loss Score

$$IoU_{Loss} = - \sum_{n=1}^N \frac{\hat{y}_n y_n}{\hat{y}_n + y_n - (\hat{y}_n y_n)}$$

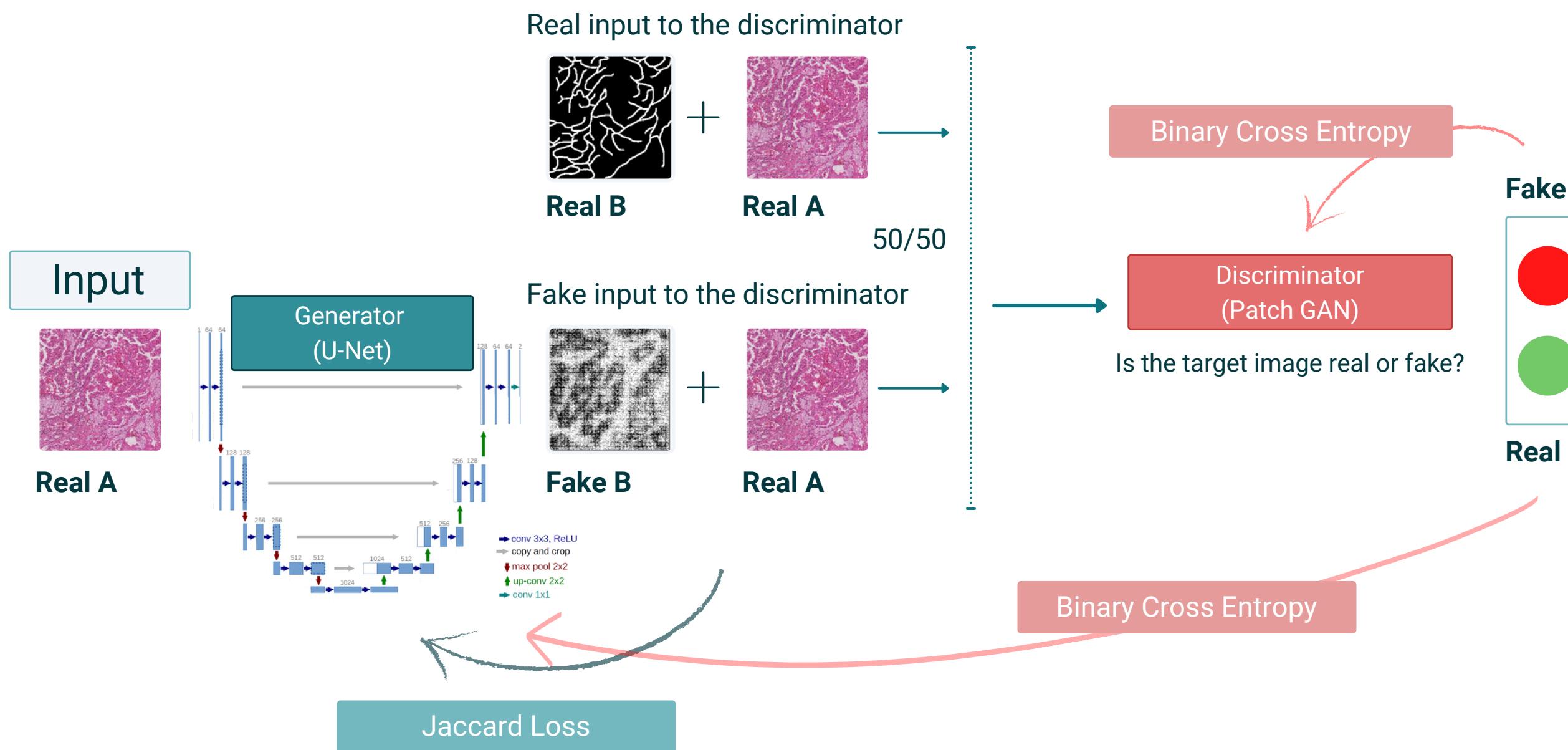
# U-Net - Best Results



**DATASET :** pRCC - resize  
**PREPROCESSING:** H&E norm  
**LR:** 1e-1  
**BATCH:** 16  
**LOSS:** Jaccard Loss

Dataset	Mean IoU	IoU Minority	Dice Minority
pRCC	0.5730026	0.3171680532008569	0.4815908682724352
ccRCC	0.5610693	0.3882488006888351	0.5593360505641207
pRCC+ccRCC	0.55349886	0.28494036	0.4435075454438602

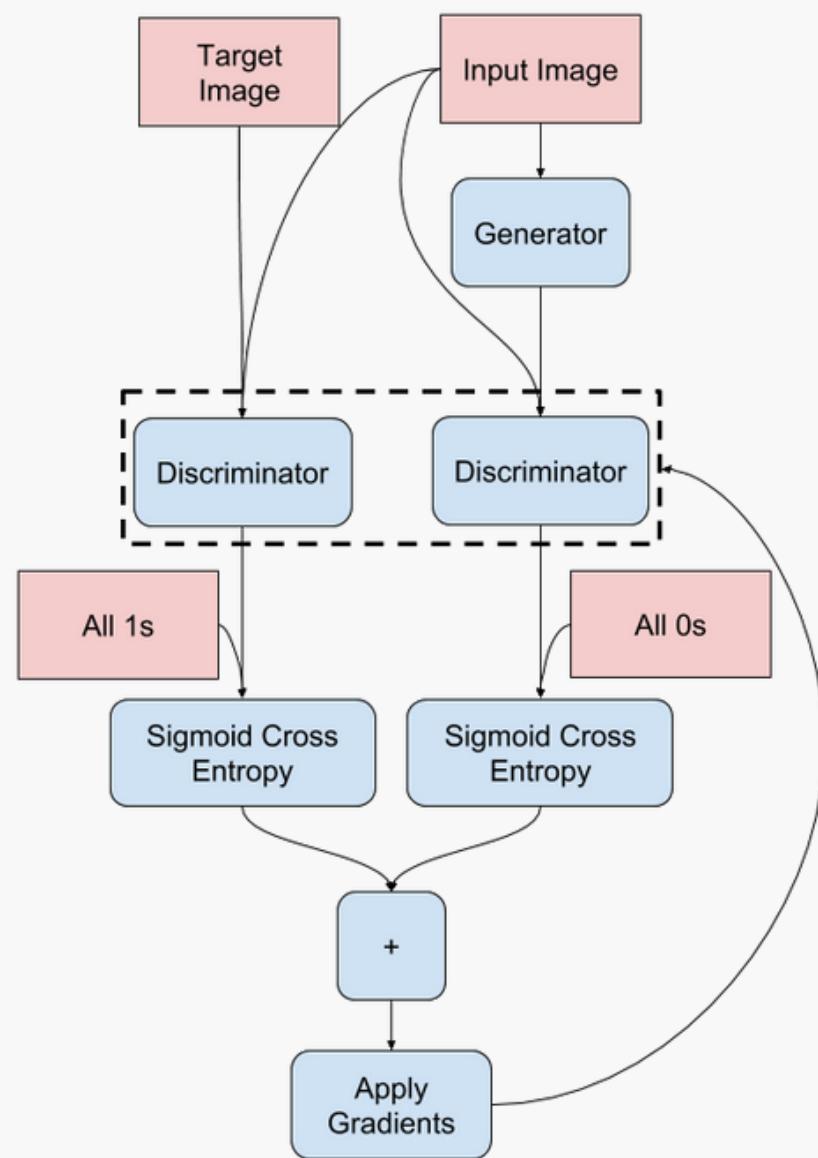
# Pix2Pix GAN - Structure



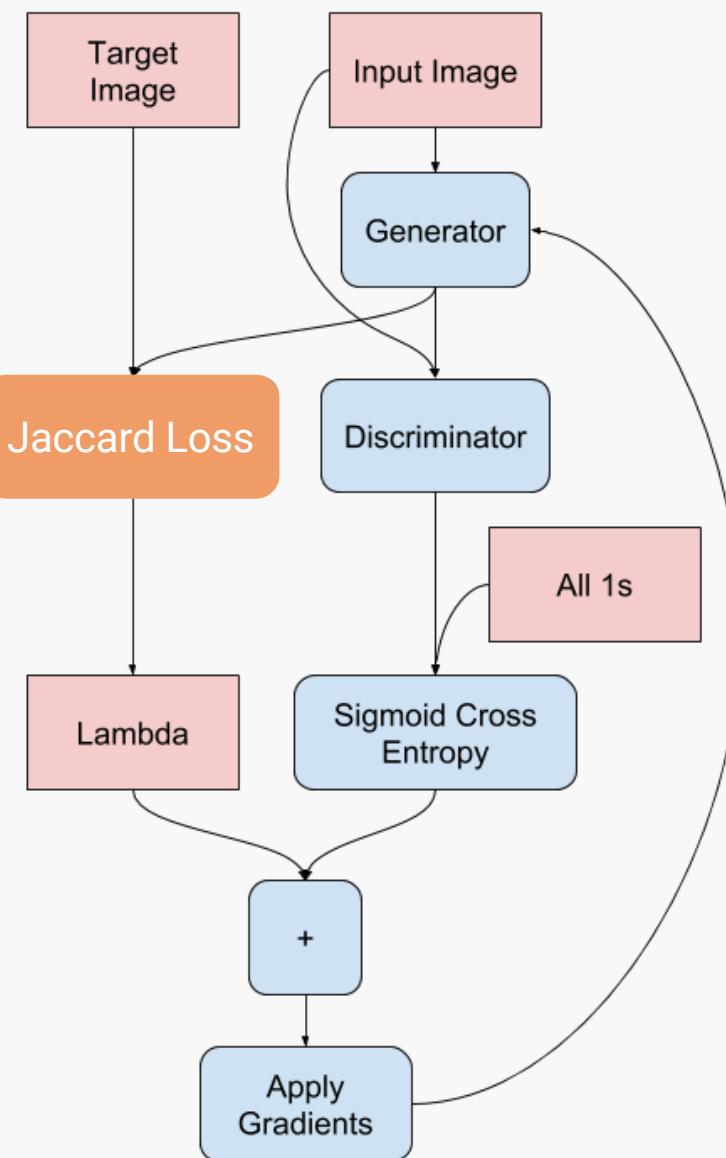
[4]

# Pix2Pix - Adversarial loss

## Discriminator Updating



## Generator Updating



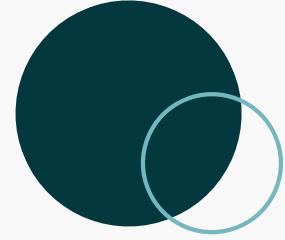
## Binary cross entropy

$$BCE_{Loss} = - \sum_{n=1}^N [y_n \log(\hat{y}_n) + (1 - y_n) \log(1 - \hat{y}_n)]$$

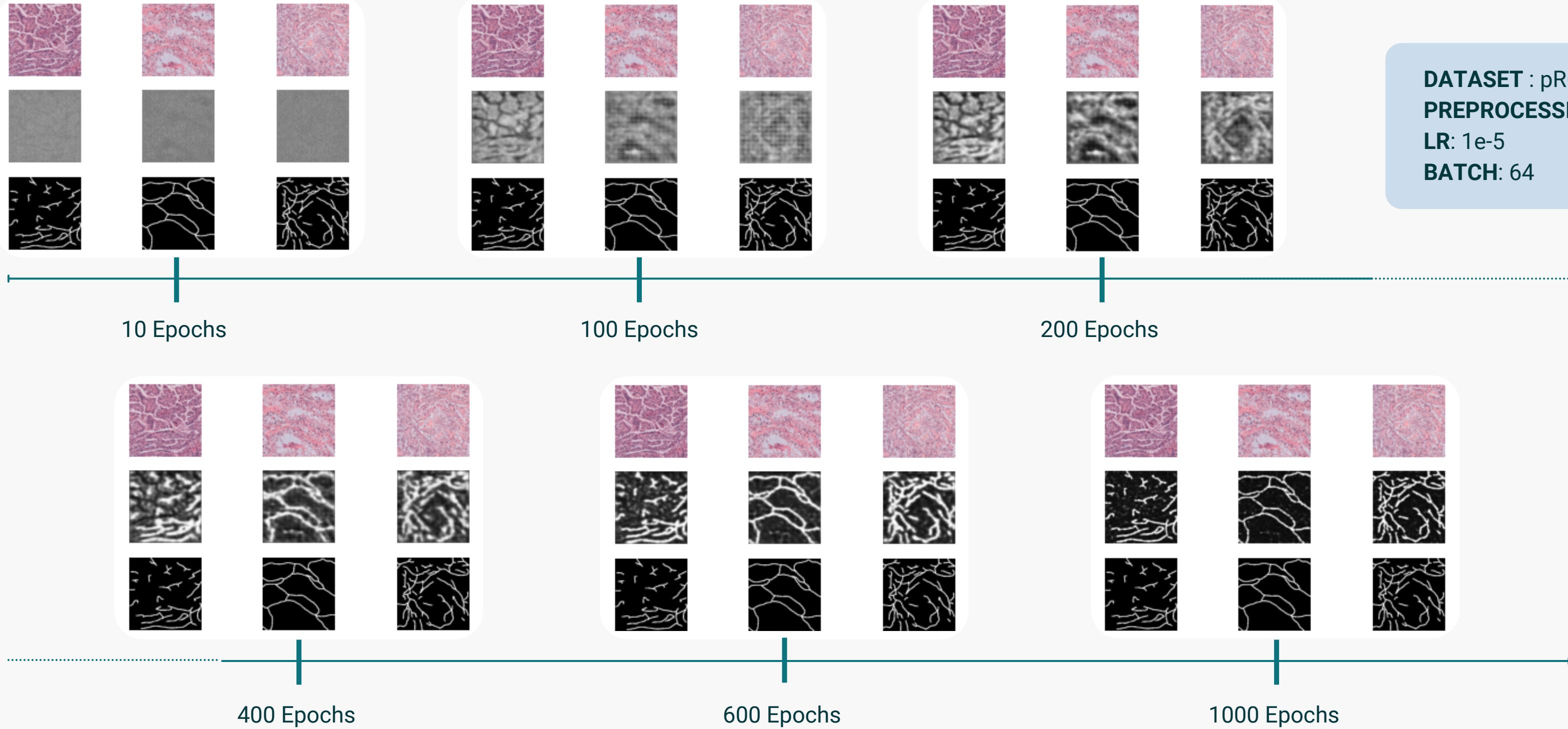
## Jaccard Loss

$$IoU_{Loss} = - \sum_{n=1}^N \frac{\hat{y}_n y_n}{\hat{y}_n + y_n - (\hat{y}_n y_n)}$$

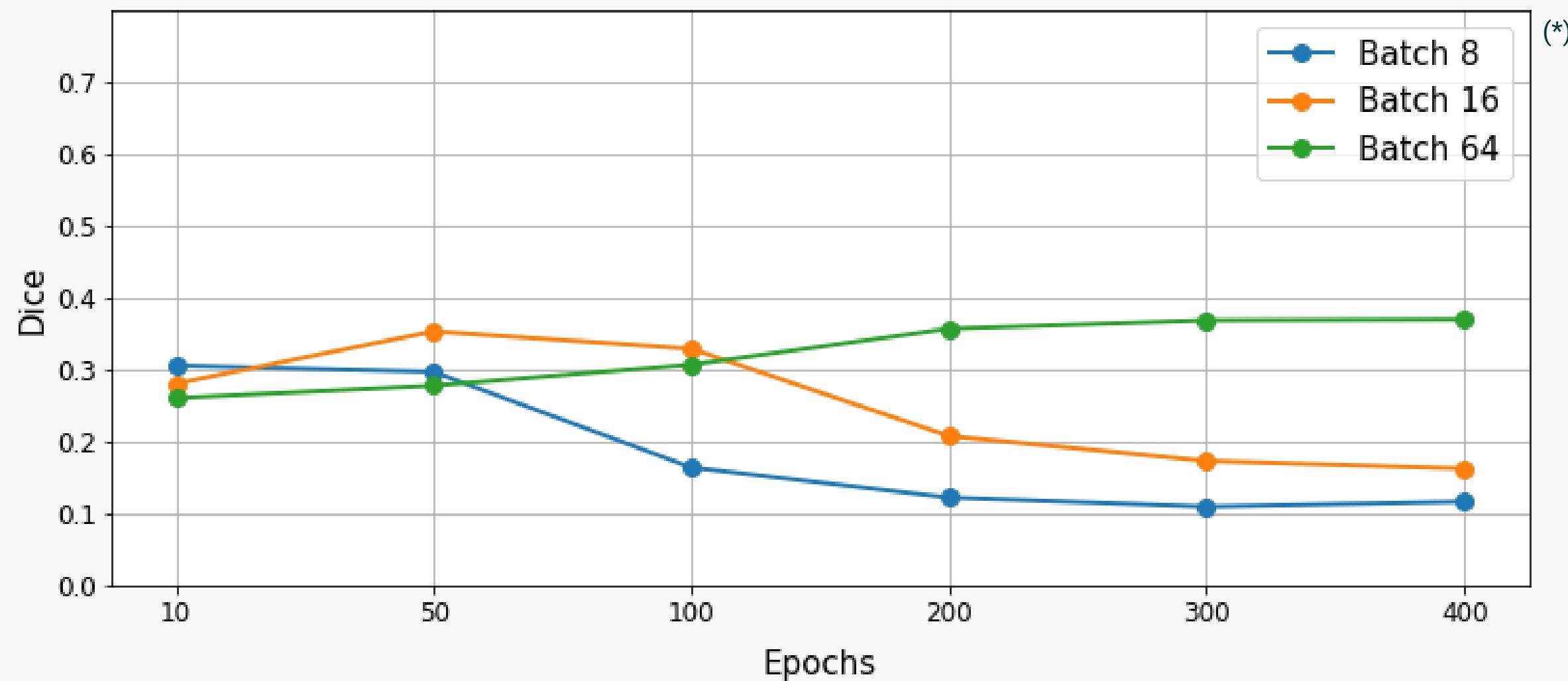
Originally L1 loss was used by Pix2Pix paper. After some experiments, it has been decided to use the Jaccard Loss and also to have a Sigmoid activation function instead of a Tanh activation function.



# Pix2Pix - Training

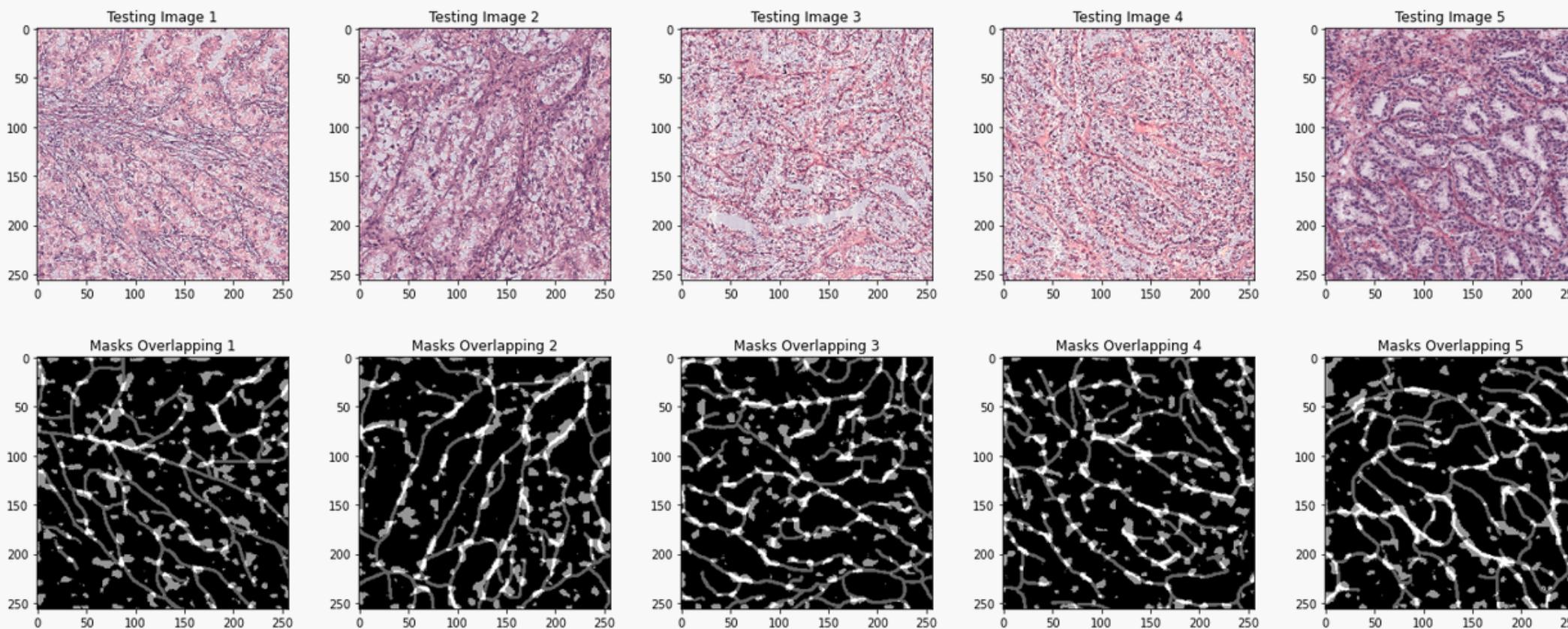
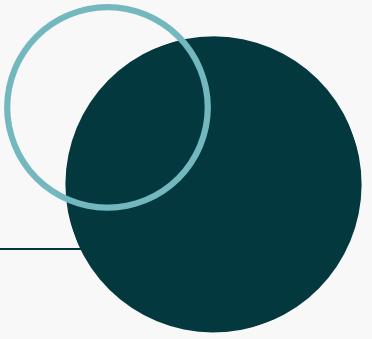


# Pix2Pix - Tuning batch size



(\*) Results refer to pRCC dataset with LR 1e-5

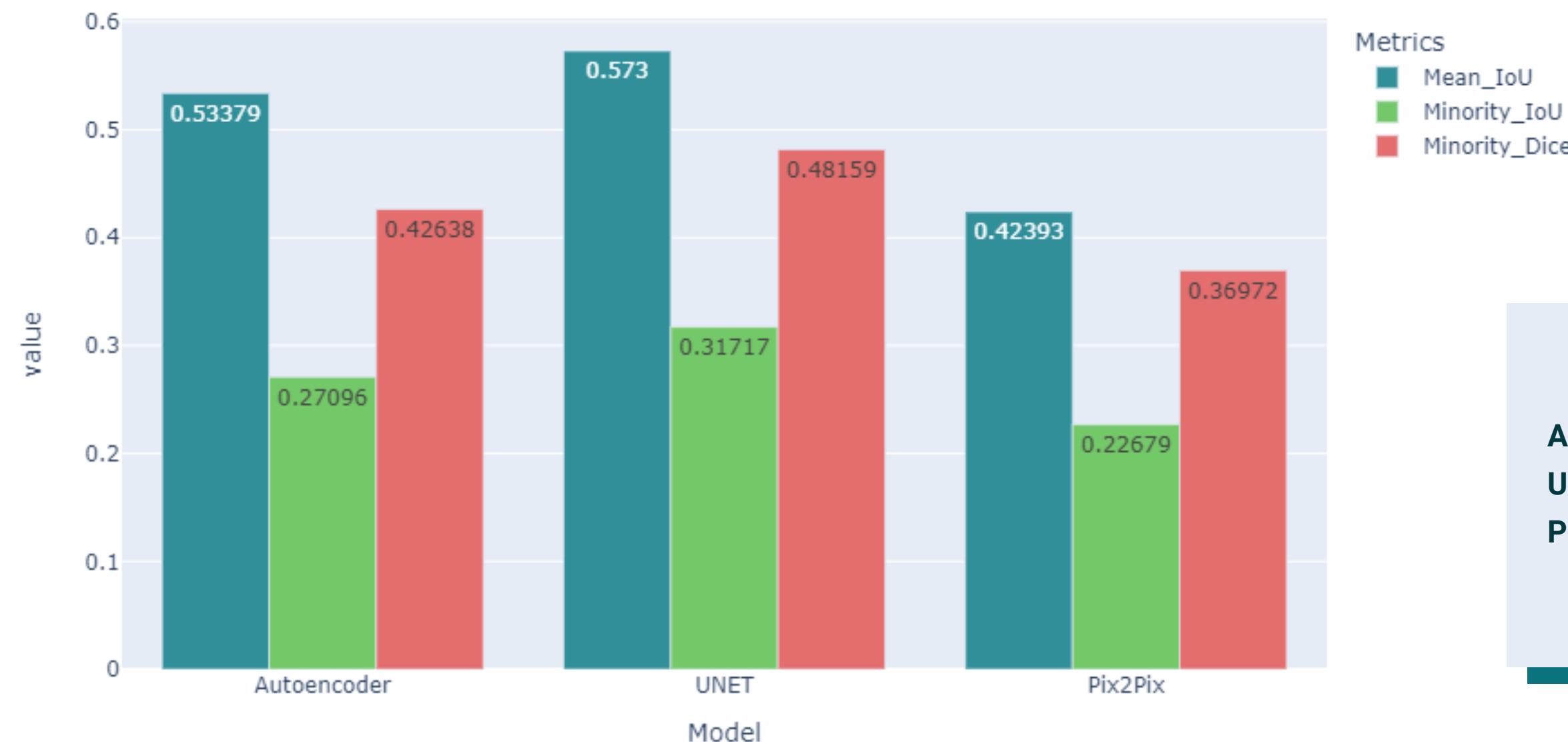
# Pix2Pix - Best Results



**DATASET :** pRCC - resize  
**PREPROCESSING:** H&E norm  
**LR:** 1e-5  
**BATCH:** 64

Dataset	Mean IoU	IoU Minority	Dice Minority	Epochs
pRCC	0.42393038	0.226787	0.3697257	400
ccRCC	0.37890804	0.297376	0.458426	400
pRCC+ccRCC	0.43870416	0.2272085	0.370285	300

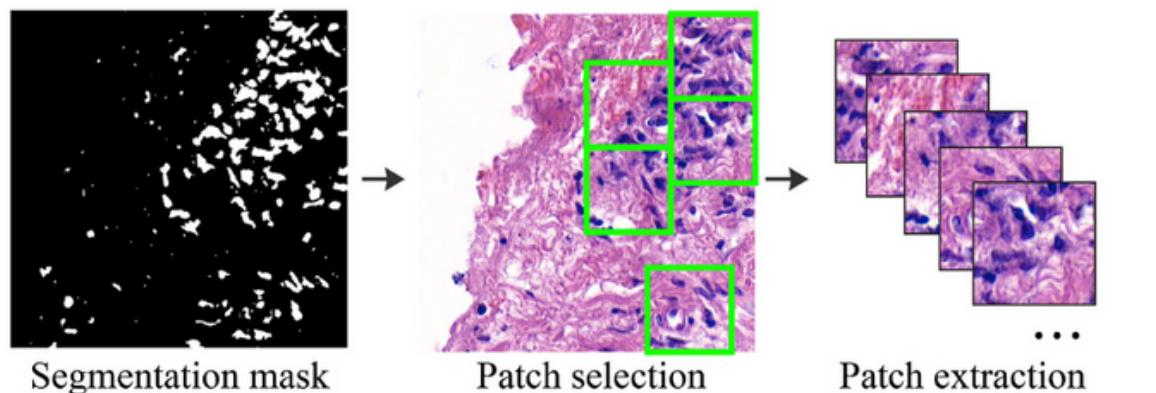
# Conclusion



**Autoencoder:** baseline  
**U-Net:** best performing model  
**Pix2Pix:** requires more data

(\*) Results refer to pRCC dataset

# Next Works

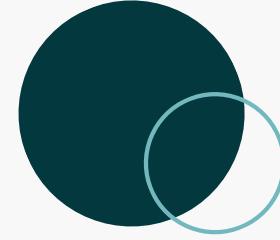


✓ Perform classification before segmentation

✓ Increase samples maintaining vascular network information

✓ Patch Smart Sampling

✓ Remove noise (e.g. through hematoxylin channel)



# References

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- [1] A method for normalizing histology slides for quantitative analysis. *Marc Macenko, Marc Niethammer, J. S. Marron, David Borland, John T. Woosley, Xiaojun Guan, Charles Schmitt, Nancy E. Thomas*
- [2] Impact of stain normalization and patch selection on the performance of convolutional neural networks in histological breast and prostate cancer classification. *Massimo Salvi, Filippo Molinari, U Rajendra Acharya, Luca Molinaro, Kristen M Meiburger*
- [3] U-Net: Convolutional Networks for Biomedical Image Segmentation. *Olaf Ronneberger, Philipp Fischer, and Thomas Brox*
- [4] Image-to-Image Translation with Conditional Adversarial Networks. *Phillip Isola, Jun-Yan Zhu, Tinghui Zhou, Alexei A. Efros*

# **Thank you for your attention!**

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