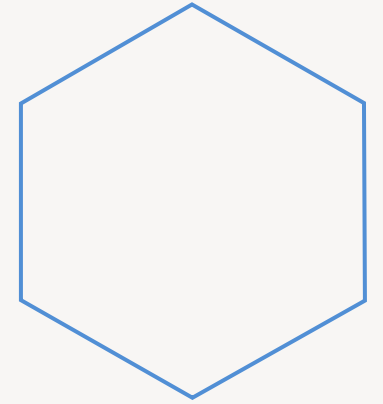


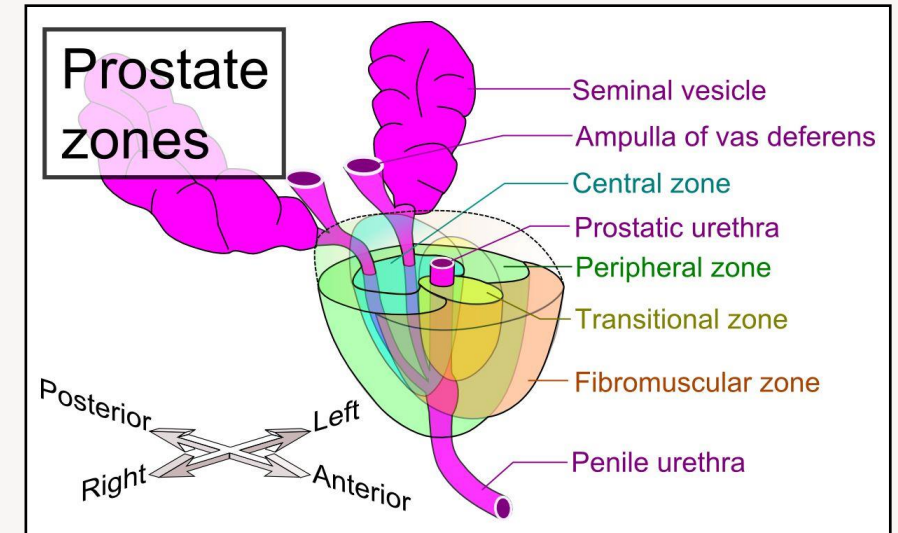
Segmentation of the prostate using U-net

Noam Sprei
Mor Tzadok



The Prostate Gland

- ❖ Small, walnut-sized gland.
- ❖ Part of the male reproductive system (produce a component of the semen fluid).
- ❖ Located below the bladder and surrounds the urethra.
- ❖ Health conditions that affect the prostate gland, such as:
 1. **Prostatitis**: Inflammation or infection of the prostate gland.
 2. **BPH**: Benign Prostatic Hyperplasia: Non-cancerous enlargement of the prostate gland.
 3. **Prostate Cancer**: The most common cancer in men, characterized by the abnormal growth of cells in the prostate gland.



Jan (Janko) Ondras



Computer Science Ph.D. Student



Cornell University



Ithaca, NY

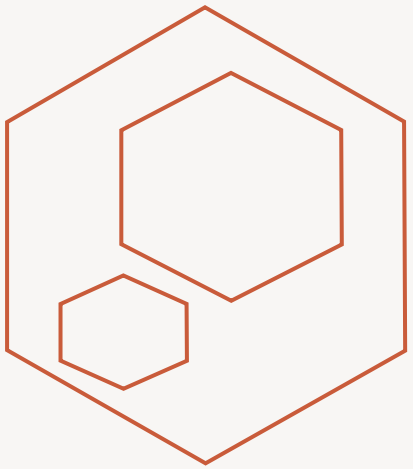


<https://janondras.wordpress.com/>



[@JankOndras](https://twitter.com/JankOndras)



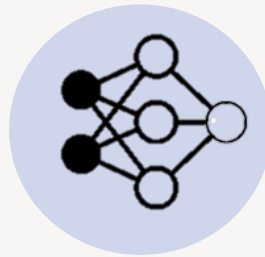


Prostate Segmentation



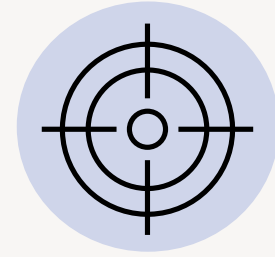
INPUT

3D MRI scan



METHOD

3D U-net




AIM

automatically annotate the
peripheral and central zones


The slide features a decorative graphic on the left side consisting of three hexagons: a large orange one at the top, a light blue one at the bottom, and a white one with a thin orange border in the middle. Overlapping the white hexagon is a small image of a desk with several papers, each containing different types of data visualizations like bar charts, line graphs, and tables.

NCI-ISBI 2013 Challenge

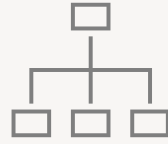
Automated Segmentation of Prostate Structures

A simple grey curved arrow pointing downwards from the 'NCI-ISBI 2013 Challenge' header to the descriptive text.

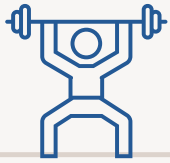
The National Cancer Institute's
(**NCI's**) Cancer Imaging Program in
collaboration with the International
Society for Biomedical Imaging
(**ISBI**) launched a challenge
involving prostate gland MRI

A simple grey curved arrow pointing downwards from the 'Automated Segmentation of Prostate Structures' header to the descriptive text.

The challenge aims to
improve automatically
annotate the peripheral and
central zones



The dataset's structure



Training set

Size: 60-case prostate MRI 3D series

files: Digital Imaging and Communications in Medicine (**DICOM**) images with matched **NRRD** markups that define the tissue CG and PZ are available from the NCI via The Cancer Imaging Archive (TCIA).



Leaderboard set

Size: 10 case prostate MRI 3D series

files: Digital Imaging and Communications in Medicine (**DICOM**).

Then, no NRRD segmentations were available for download for these leaderboard cases until after the conclusion of the challenge.



Test set

Size: 10 case prostate MRI 3D series

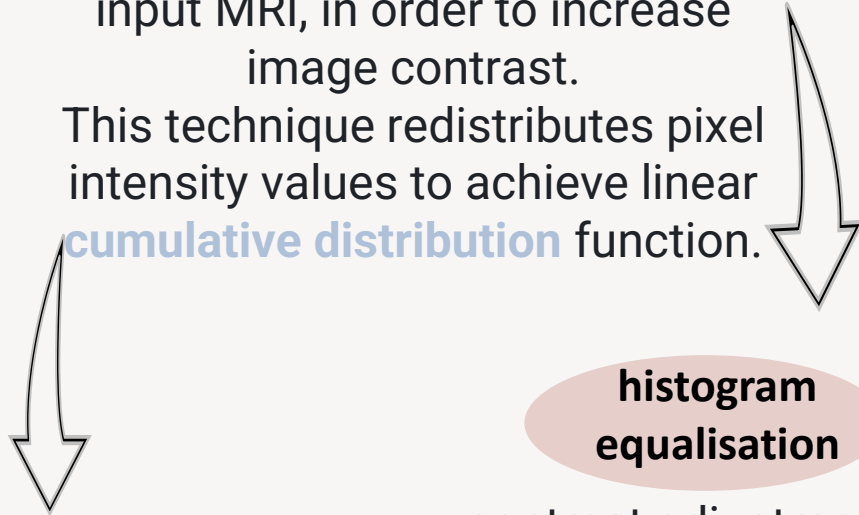
files: This data (DICOMS & NNRD) was available only after the challenge.

Each case consist of 15+ (4 mm thick slices at 3T) or 28+ (3 mm thick at 1.5T) adjacent axial cross-section cuts. They were acquired as T2-weighted MR axial pulse sequences.

Preprocessing

Used **histogram equalisation** on the input MRI, in order to increase image contrast.

This technique redistributes pixel intensity values to achieve linear **cumulative distribution** function.



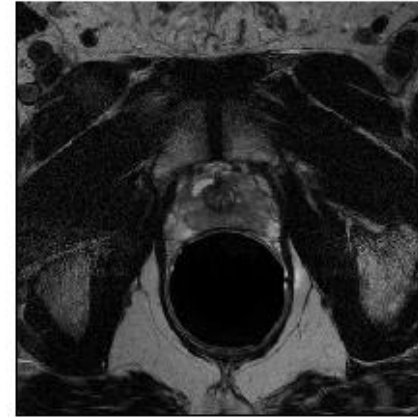
cumulative distribution

histogram equalisation

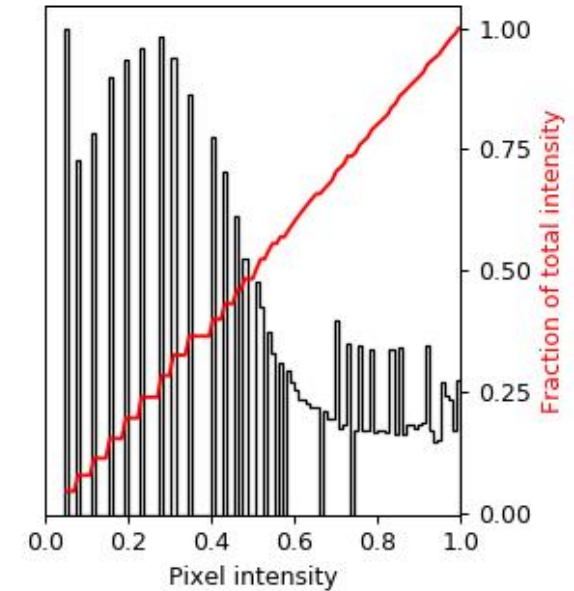
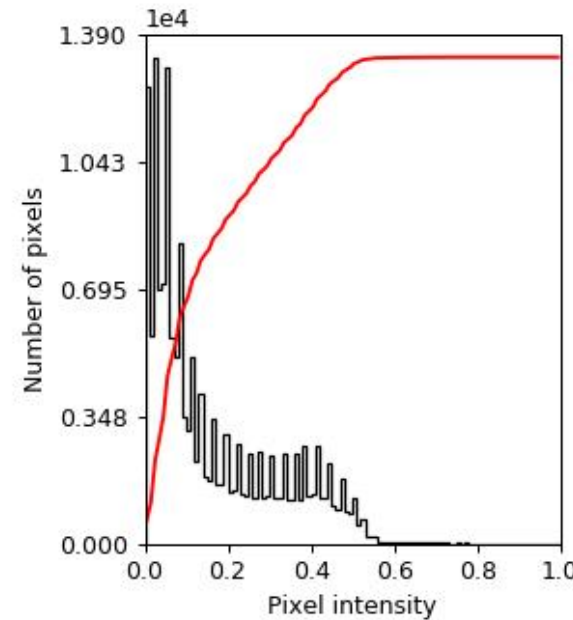
contrast adjustment using the image's histogram.

Histogram in which each bin equals to the count of the its value plus all bins for smaller values

Original low contrast image



Histogram equalised image





Data augmentation

Using the 3D Unet paper augmentations:



scalings



random rotations



gray value variations



elastic distortion

(same for voxel tile and its annotation)

Original scan



Elastic deformation



Original annotation



Elastic deformation



3D U-Net Architecture

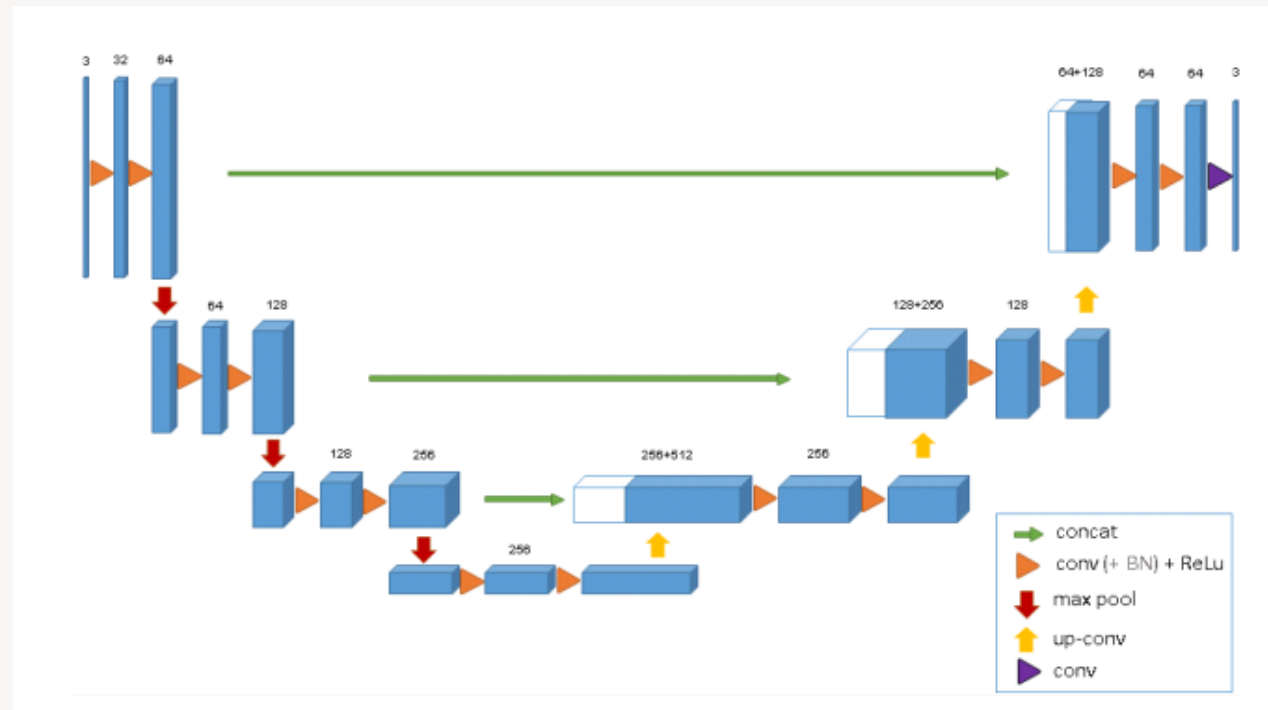


Segmentation to three classes:

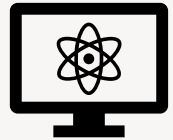
background, cg, pz



The structure:



Used the Unet from - [1606.06650.pdf \(arxiv.org\)](#)



Hyper parameters

Lost function: weighted cross entropy

Dropout: 0.3

Batch normalization: stride of one and kernel of three

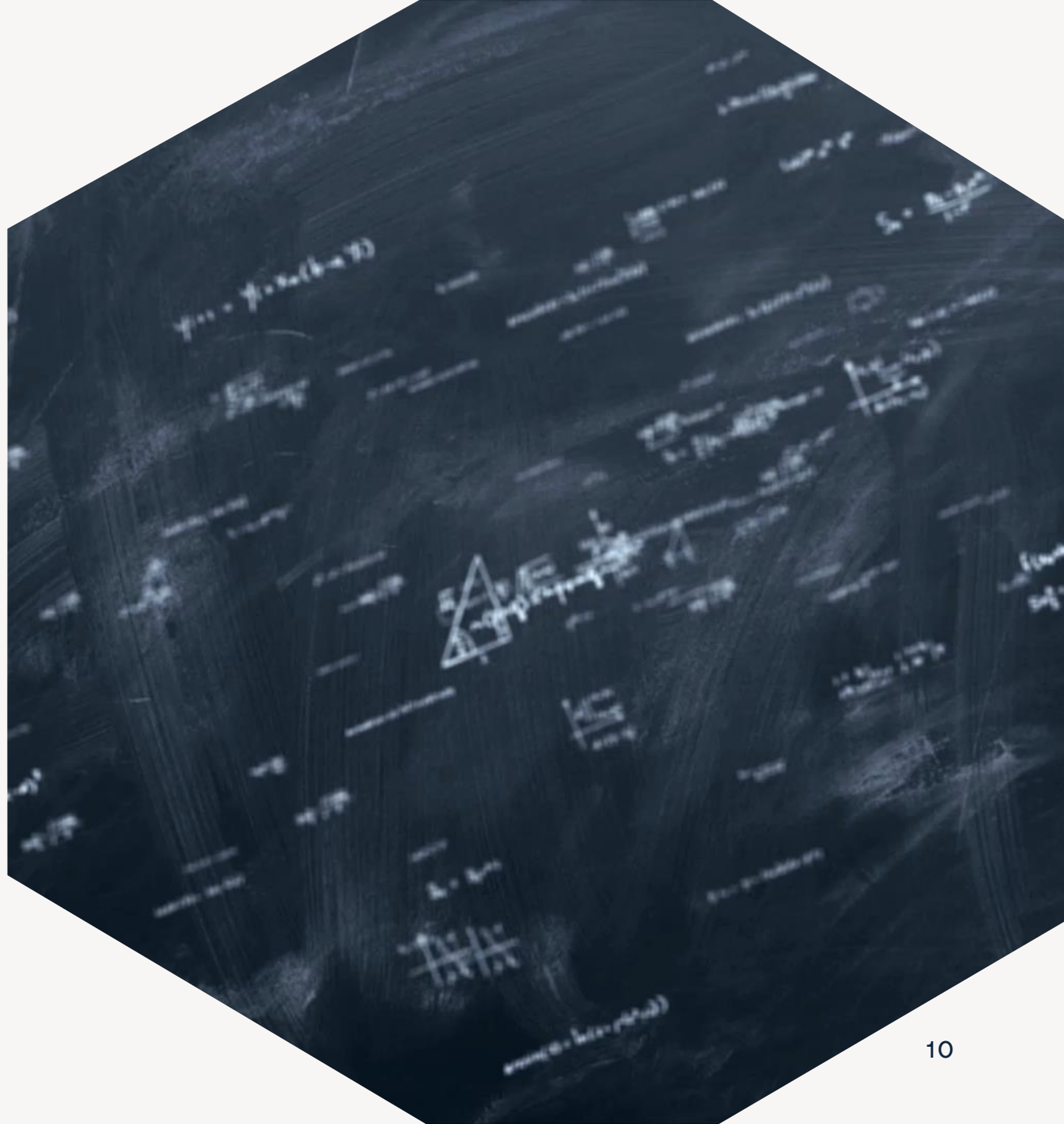
Padding: “same”- padding chosen s.t the output the same size as the input

Max pooling: [1,2,2], [1,2,2]

Optimizer: Adam, learning rate $\eta = 0.001$

Batch size: 32

Number of epochs: 322

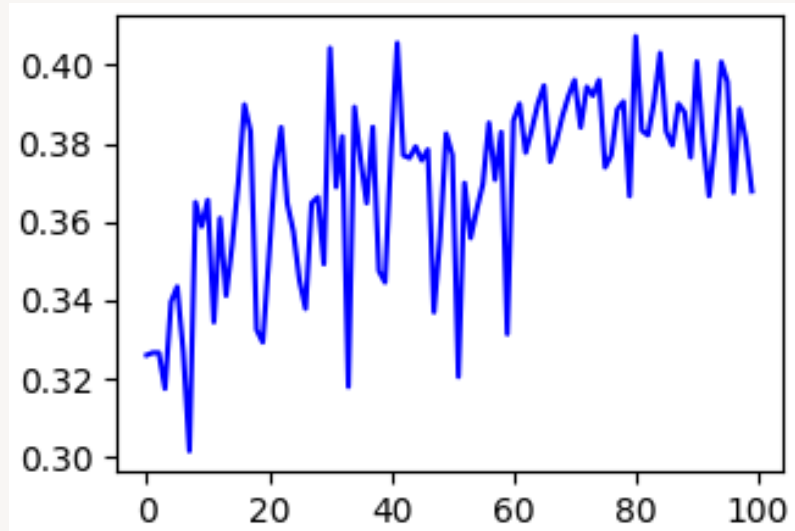


Evaluation & Results



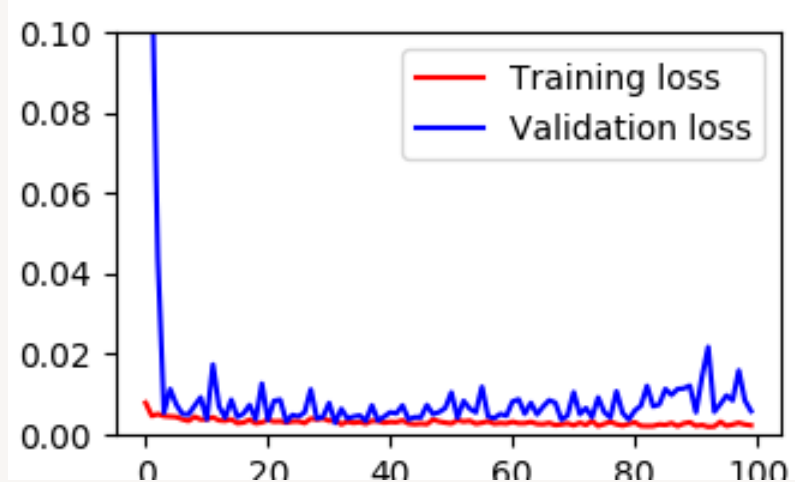
↻ 100 epochs

Validation mean IOU as a function of epochs



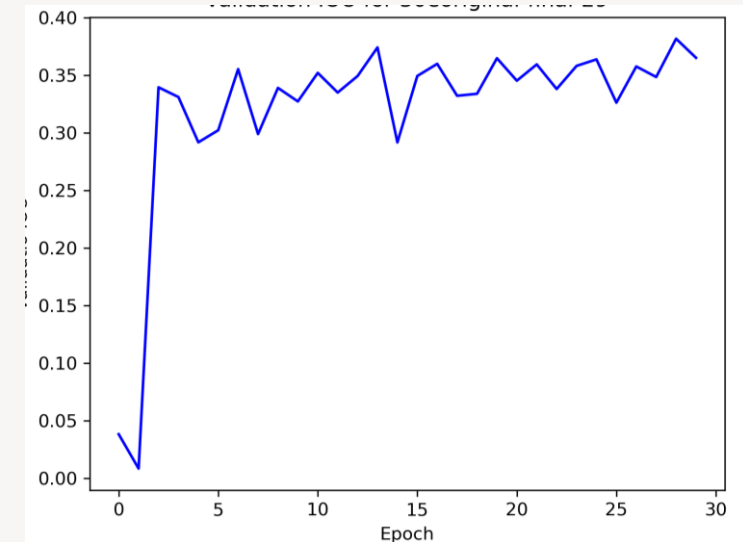
Mean test
IOU 0.41029

Train and validation loss as a function of epochs



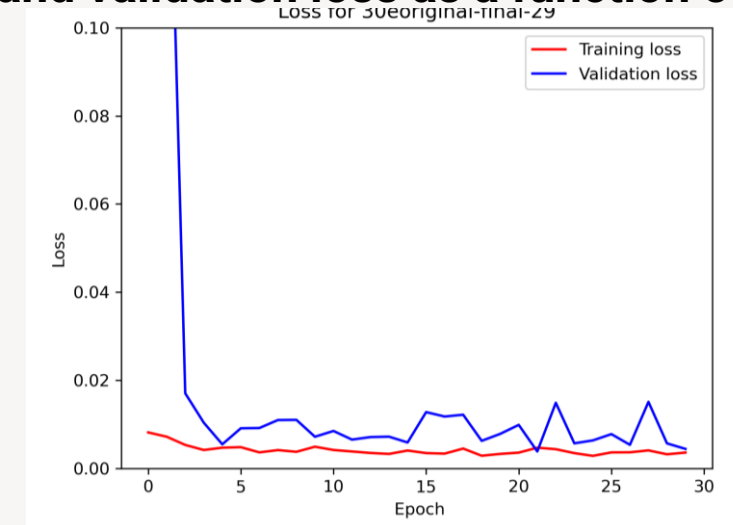
↻ 30 epochs

Validation mean IOU as a function of epochs



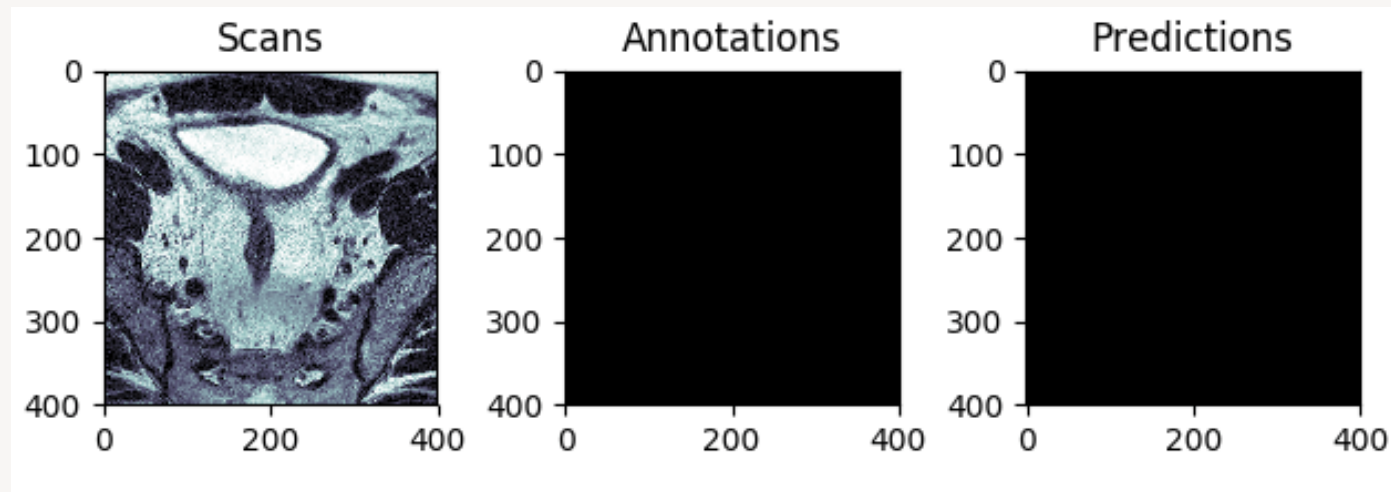
Mean test
IOU 0.37005

Train and validation loss as a function of epochs

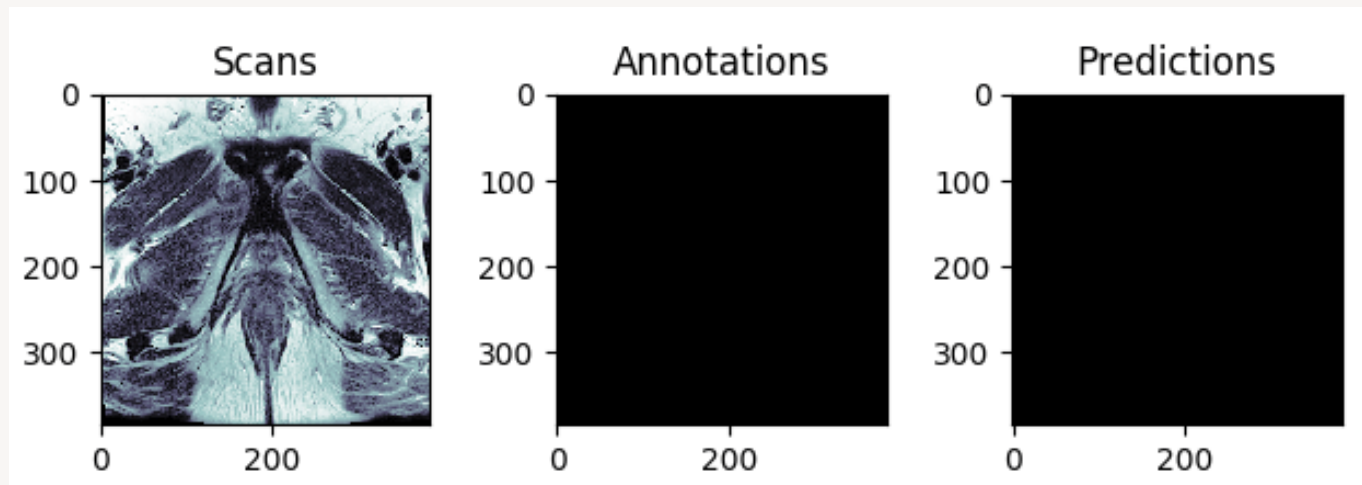


Best Result

↻ 100 epochs



v.s



Worst Result



Best Result

 30 epochs

v.s

Worst Result

14

Our mini-trials

We had some mini-trials of 4 epochs with one of the following changes:

- Weighted Dice loss instead of weighted cross entropy
- Adding augmentation – intensity shift
- Smaller learning rate
- More augmented images and larger scale of their hyperparameters





Results of our mini-trials

**Weighted Dice
loss instead of
weighted cross
entropy**

The dice loss decreased the
IOU score of the validation to
0.159

**Adding
augmentation :
intensity shift**

Adding the intensity shift
resulted in 0.34 IOU score on
the validation set

**Smaller learning
rate**

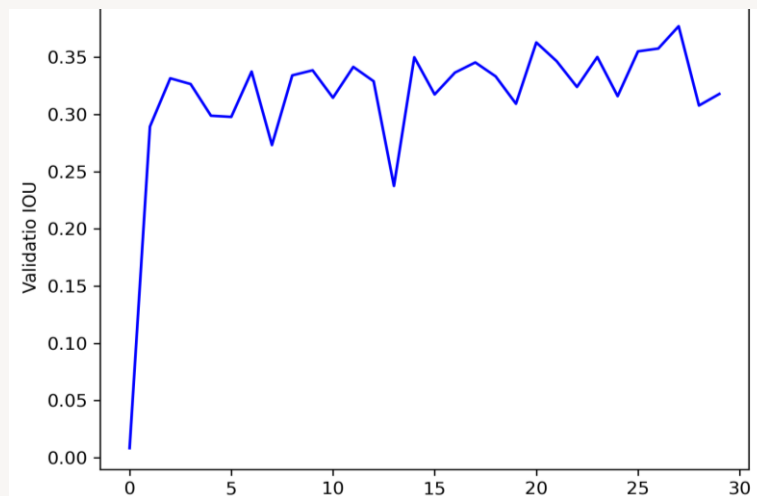
Decreasing the lr by half to
0.0005 resulted in 0.34 IOU
score on the validation set

**More augmented
images and larger
scale of their
hyperparameters**

Those modifications caused a
decrease in the IOU score of
the validation to 0.28

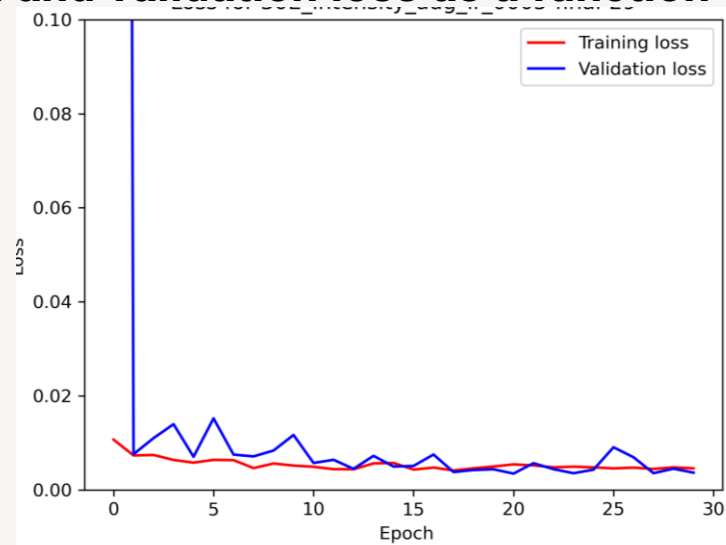
Our 30 epochs

Validation mean IOU as a function of epochs



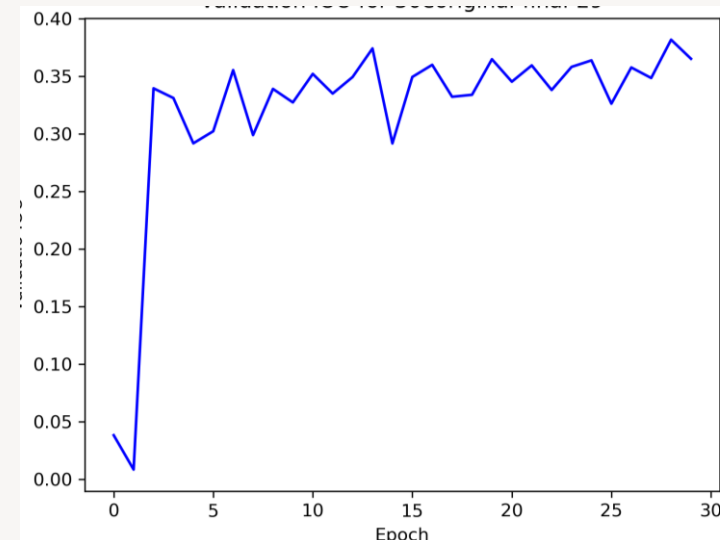
Mean test
IOU 0.32179

Train and validation loss as a function of epochs



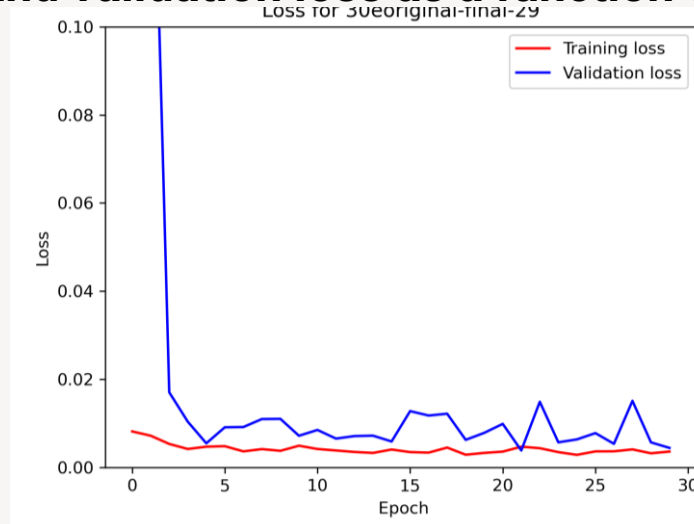
Original 30 epochs

Validation mean IOU as a function of epochs



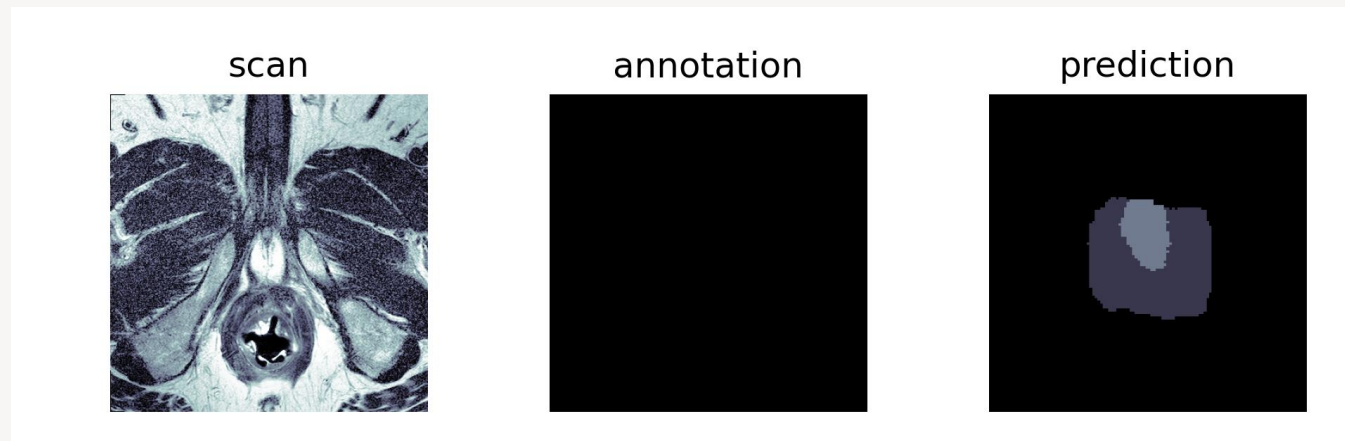
Mean test
IOU 0.37005

Train and validation loss as a function of epochs

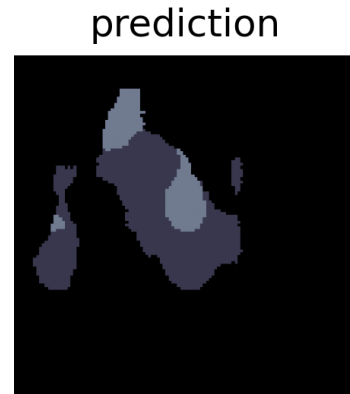
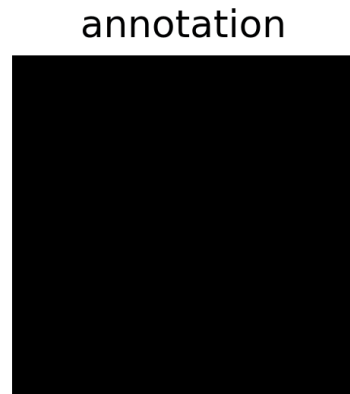
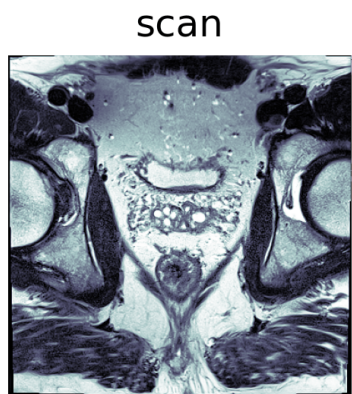


Best Result

↻ Our 30 epochs



v.s



Worst Result



Conclusions



The original project can be reconstructed

- We were able to get similar results to the reported results.



Lower number of epochs yields sufficient results

- The IOU on the test set after 30 epochs is not significantly lower than after 100 epochs.
- In addition, we can see signs of overfitting in the loss plot of the 100 epochs in the last epochs.



Further experiments with augmentations are needed

- We did see promising initial results when adding augmentations
- The augmentation we added didn't improve the results, but there are a lot of directions to explore with this augmentation and others.



Thank you

Noam Sprei & Mor Tzadok



References

The Data:

[NCI-ISBI 2013 Challenge - Automated Segmentation of Prostate Structures - The Cancer Imaging Archive \(TCIA\) Public Access - Cancer Imaging Archive Wiki](#)

The Original Code (Jancio's code):

[GitHub - jancio/3D-U-Net-Prostate-Segmentation: Implementation of the 3D U-Net model for segmentation of prostate structures](#)

Our Code:

https://github.com/MorTzadok/MRI_seminar/tree/main