

Prostate Cancer Recognition in MR-images with Keras Deep Learning

Research Project
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Introduction

- Motivation and Aim
- Image Database
- Keras and Tensorflow

Main Part

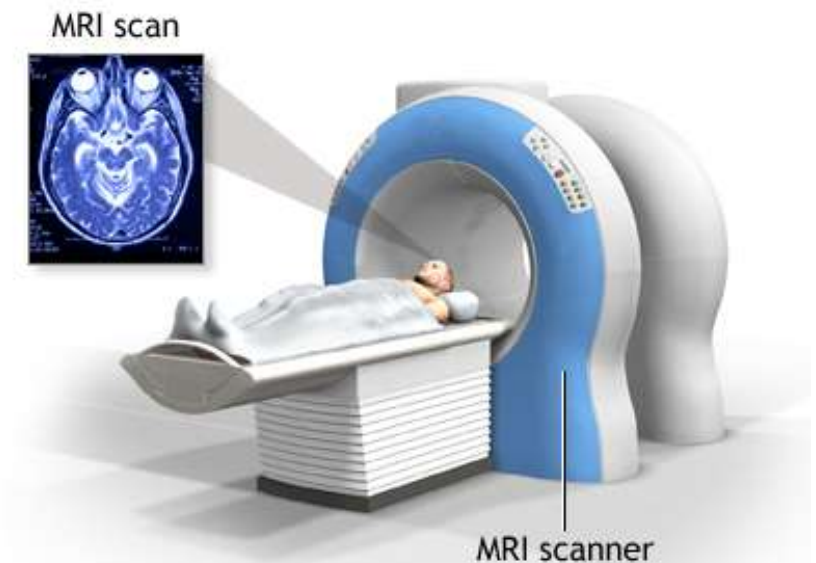
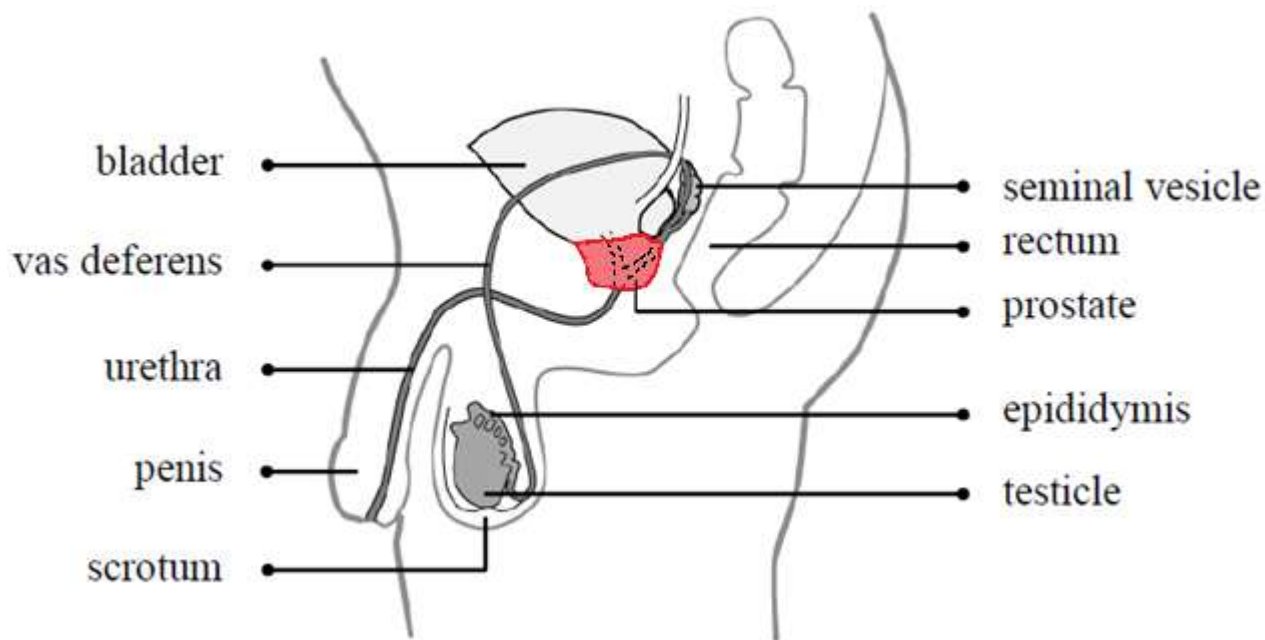
- Network Architecture
- Experiments

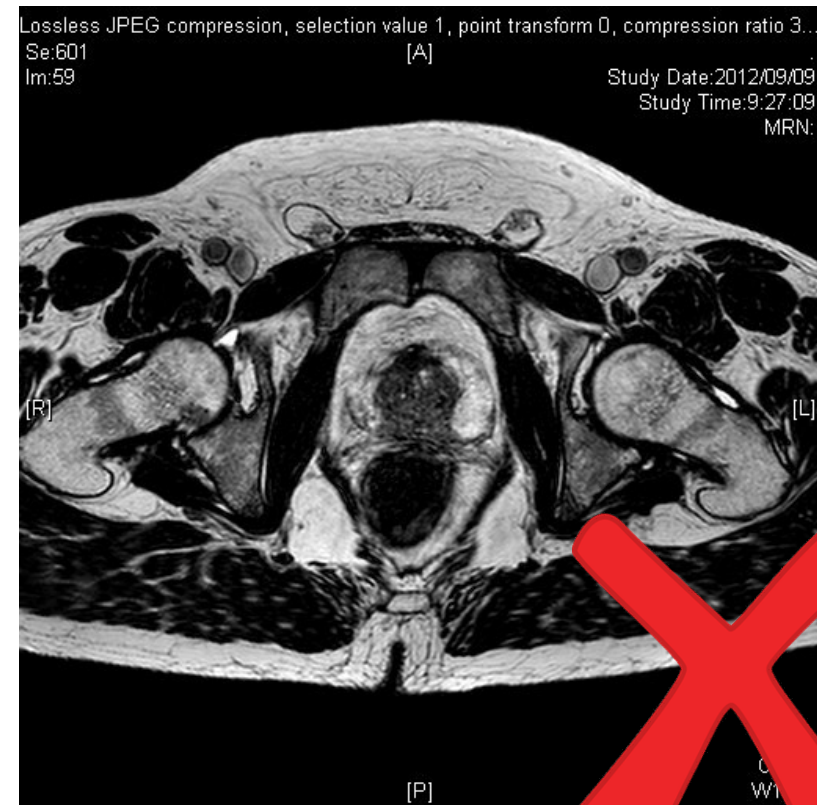
Summary

- Conclusion and Outlook

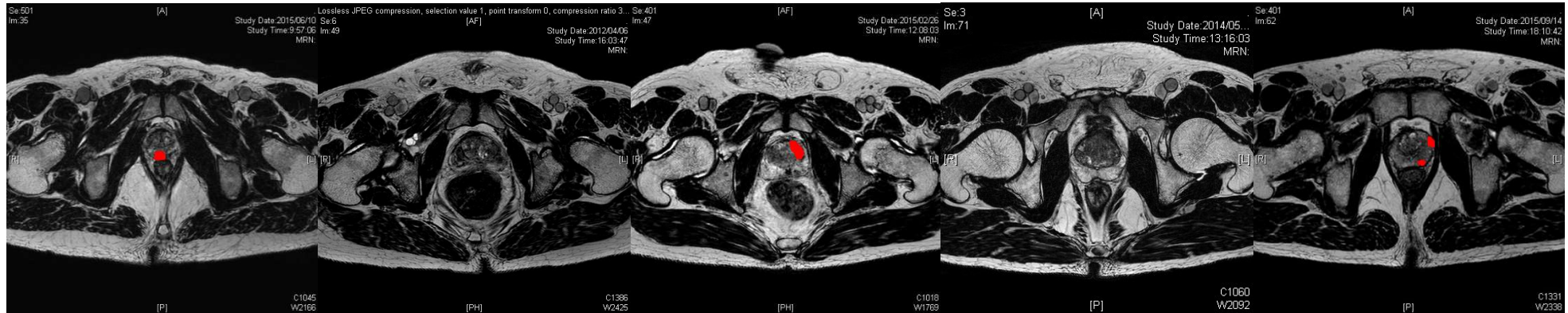
INTRODUCTION

- Prostate cancer is the second most common cancer in men
- Early detection is essential for successful treatment
- State-of-the-art imaging technique:
Multi Resonance Imaging (MRI)
- Requires expertise of experienced radiologists
- Computer aided diagnosis can help agreement among doctors





- Classification into cancer and healthy patient scans
- Usage of a simple neural network



- 1465 images of 218 cancer patients
- 3460 images of 128 healthy patients
- Cancer annotations included
- Training: 65% \triangleq 3216, validation: 15% \triangleq 741, testing: 20% \triangleq 968

Keras

- Open-source neural network library
- Written in Python
- Runs on top of Tensorflow, Theano or CNTK
- Enables fast experimentation with deep neural networks

Tensorflow

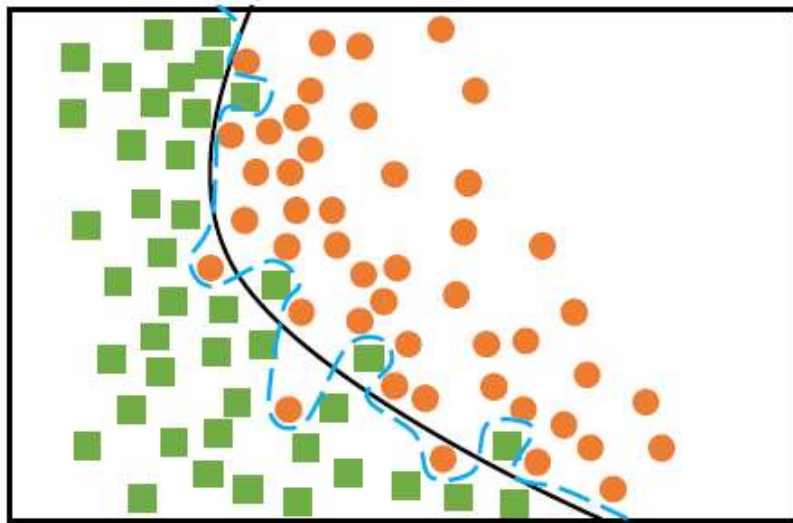
- Open-source software library developed by Google
- Can run on multiple CPUs and GPUs with optional CUDA extensions



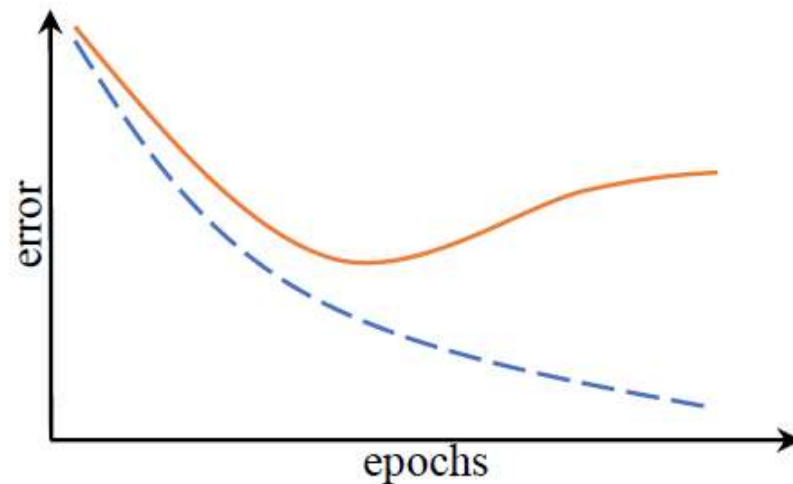
MAIN PART

Reasons for a small network

- Better understandable
- Easier to make adaptations
- Faster training times
- Less prone to overfitting ← main reason; what is overfitting?

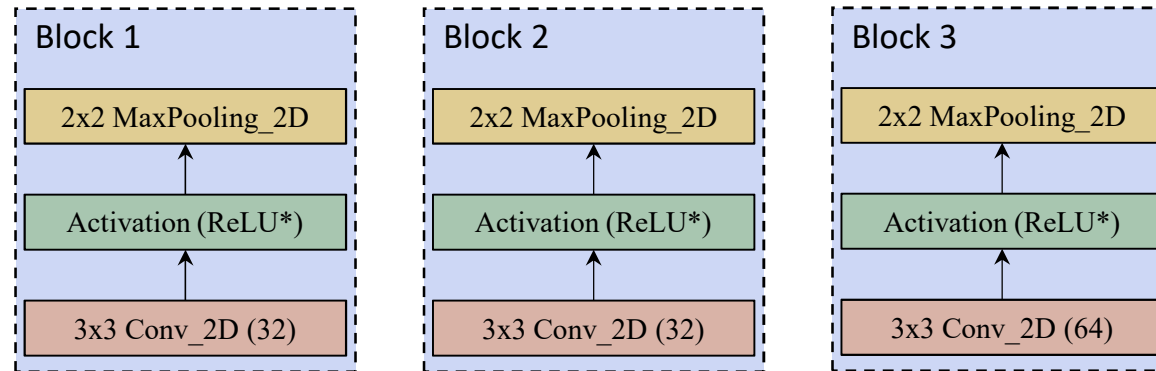


(a) Decision boundaries for an overfitted (dashed blue line) and an regularized model (black line)

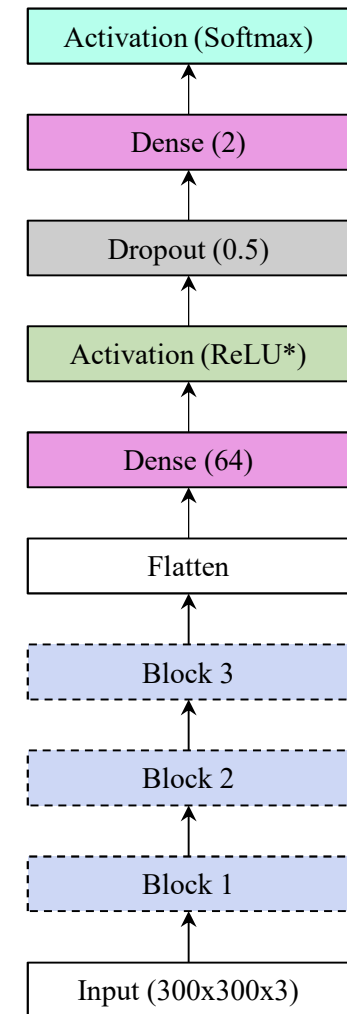


(b) Error rate for training (dashed blue) and test data (orange)

- Stack of three convolutional layers followed by two fully-connected layers
- Compiled with RMSprop** optimizer and a learning rate of 0.001



- Input: 300x300 pixel image in range [0, 1)
- Output: probability vector for the classes for each image (e.g. [0.34 0.66])



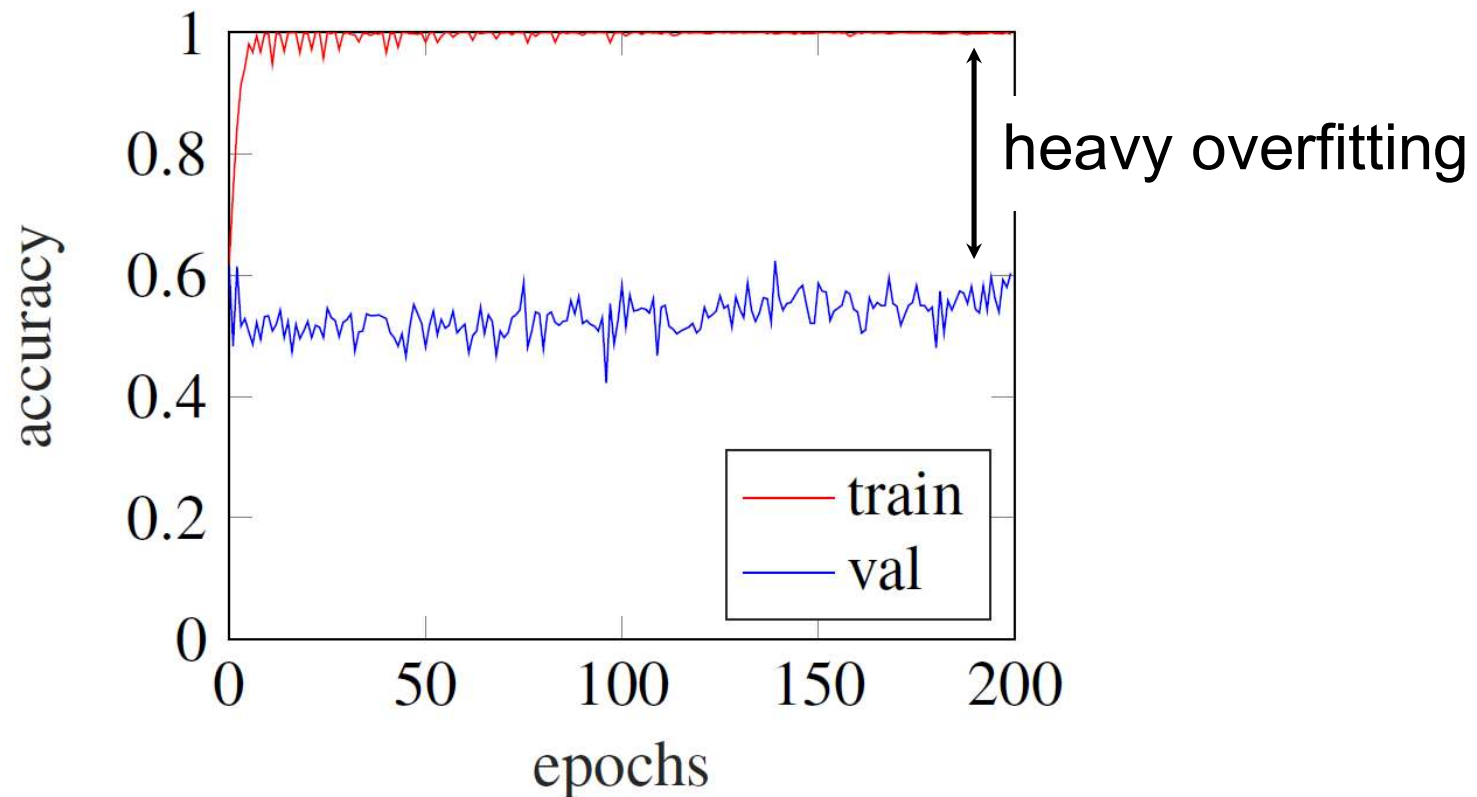
*Rectifier Linear Unit

**Root Mean Square Propagation

Experiment Setup and Baseline



- default parameter settings
- input images: 300x300 pixel sized crops
- 200 epochs with a batch size of 128
- RMSprop optimizer with a constant learning rate of 0.001

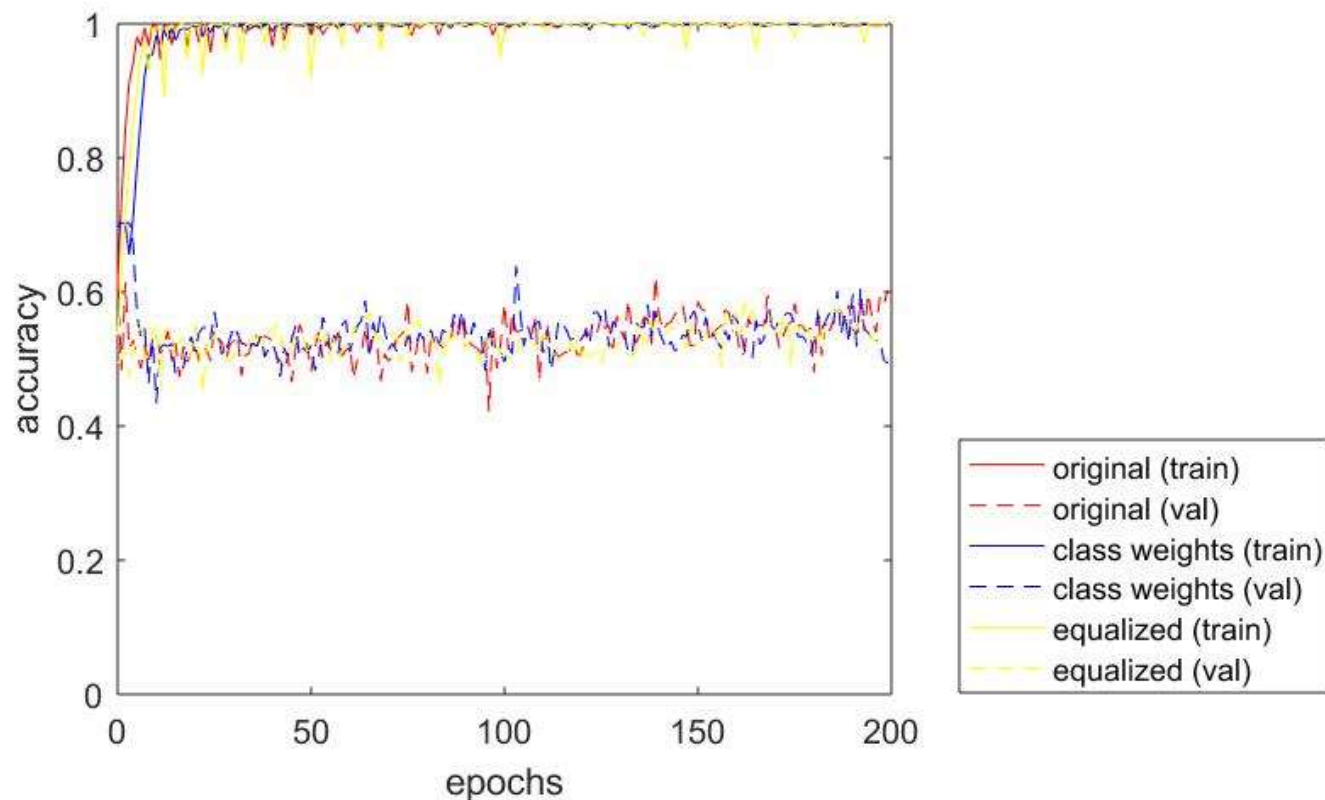


Experiment: Class weights



Dataset is not equally split into cancer and healthy cases, there exist three options to deal with it:

- Use original dataset
- Use original dataset, but set class weight dictionary for training
- Reduce dataset to equal size by discarding healthy images



Experiment: Optimizer Type & Learning Rate



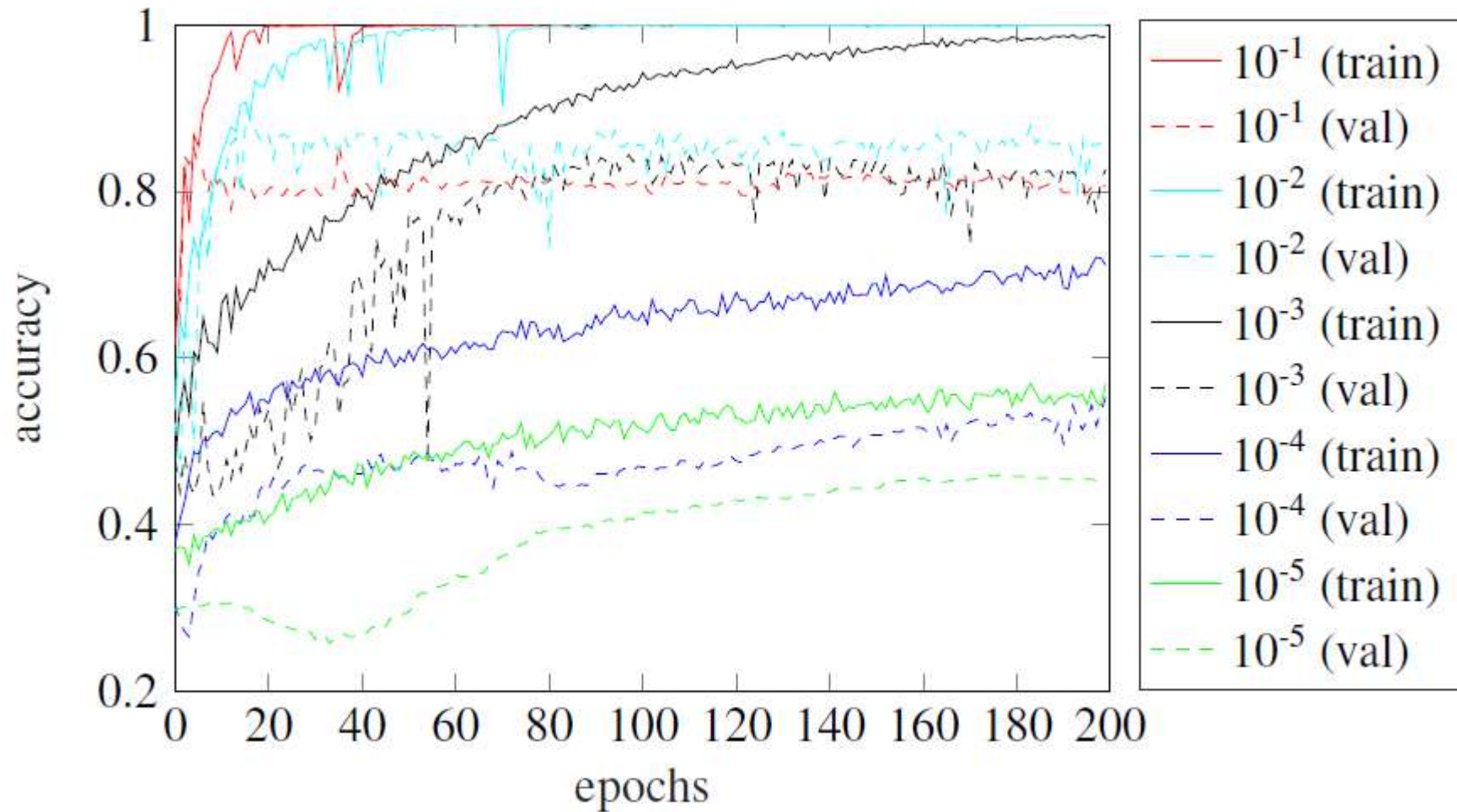
Three different optimizer types tested:

- Root Mean Squared Propagation (RMSprop)
- Stochastic Gradient Descent (SGD)
- Adaptive Moment Estimation (Adam)

Experiment: Optimizer Type & Learning Rate



Accuracy curves for training with SGD* optimizer with different learning rates (default: 0.01)



*Stochastic Gradient Descent

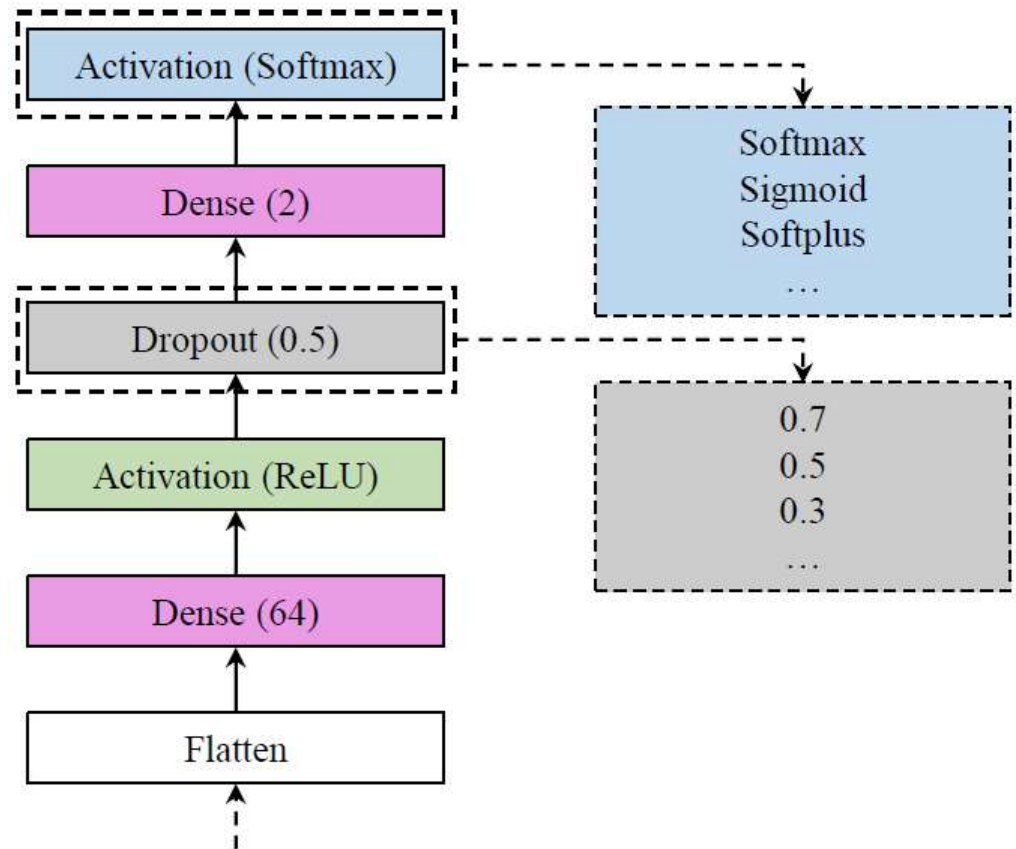
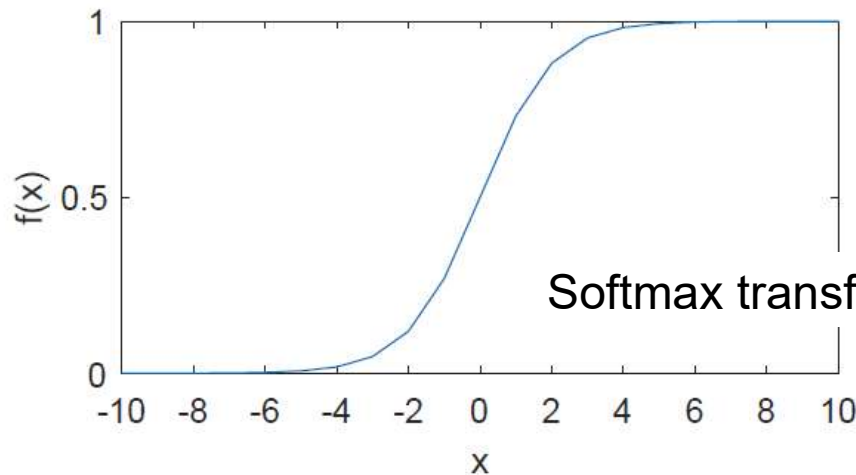
Experiment: Activation Function & Dropout



Which activation function do we change and where is the dropout applied?

Testing activations:

- Softmax
- Sigmoid
- Softplus

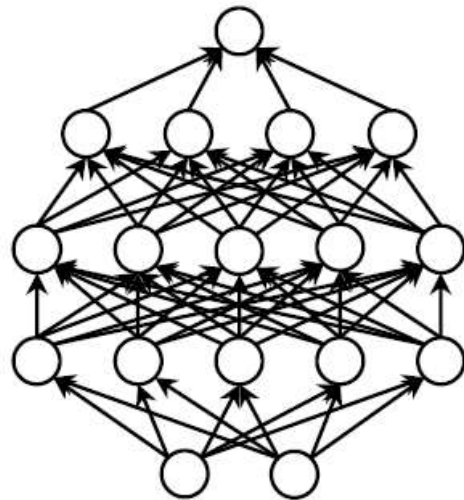


Experiment: Activation Function & Dropout

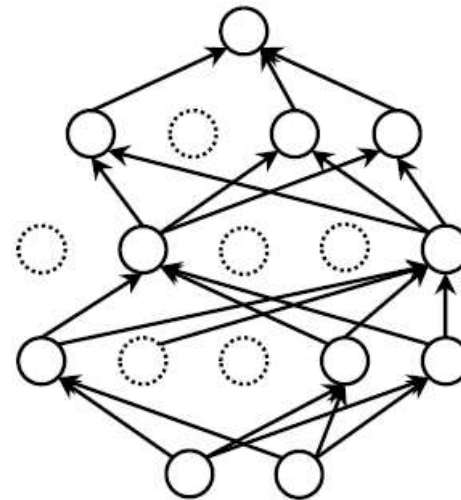


What is dropout?

- Technique to reduce overfitting
- At each training stage individual nodes are „dropped out“ (temporarily removed)
- Prevents co-adaption
- Only reduced network is trained, afterwards removed nodes are reinserted with their original weights



(a) Standard neural network

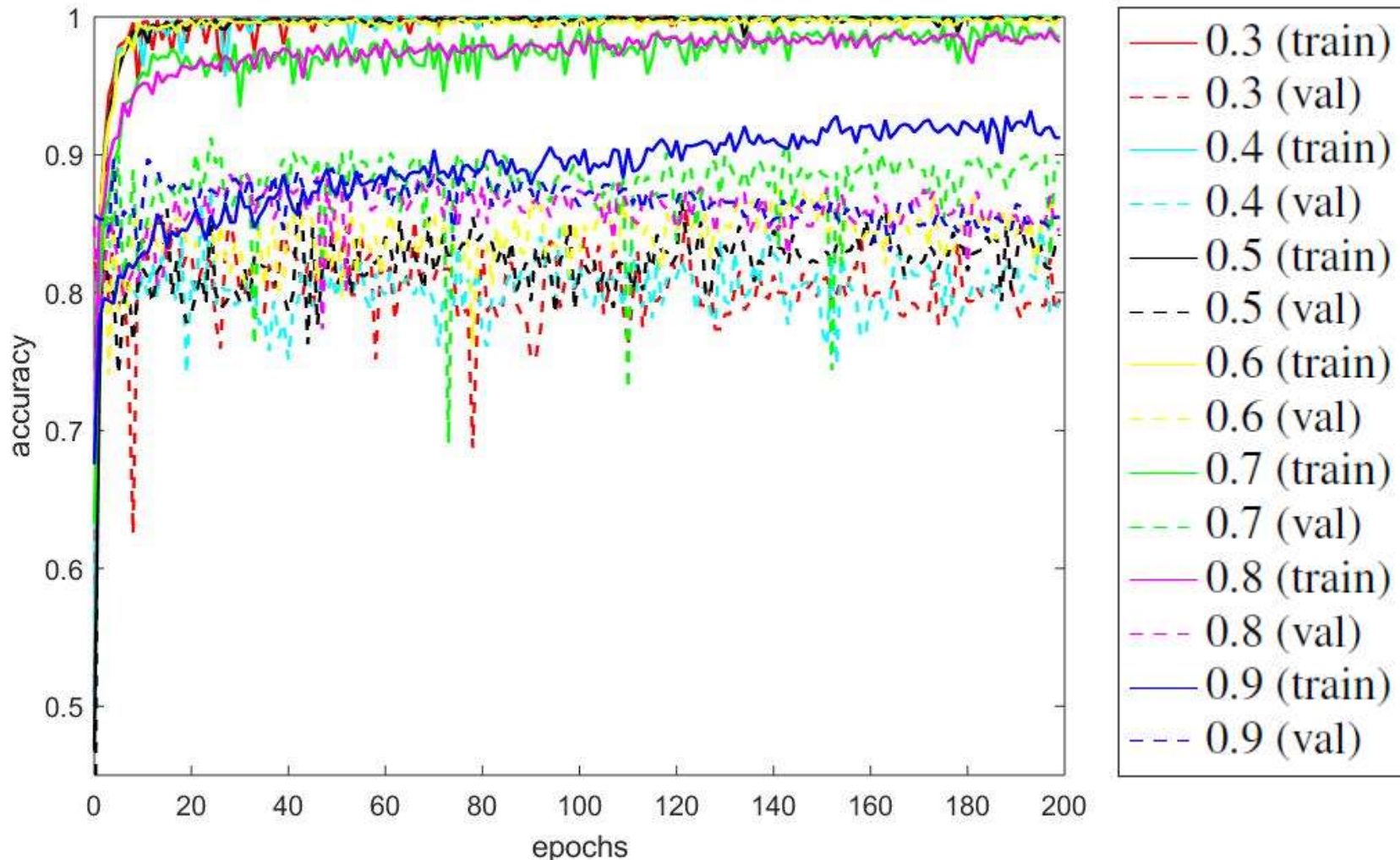


(b) Network after applying dropout

Experiment: Activation Function & Dropout



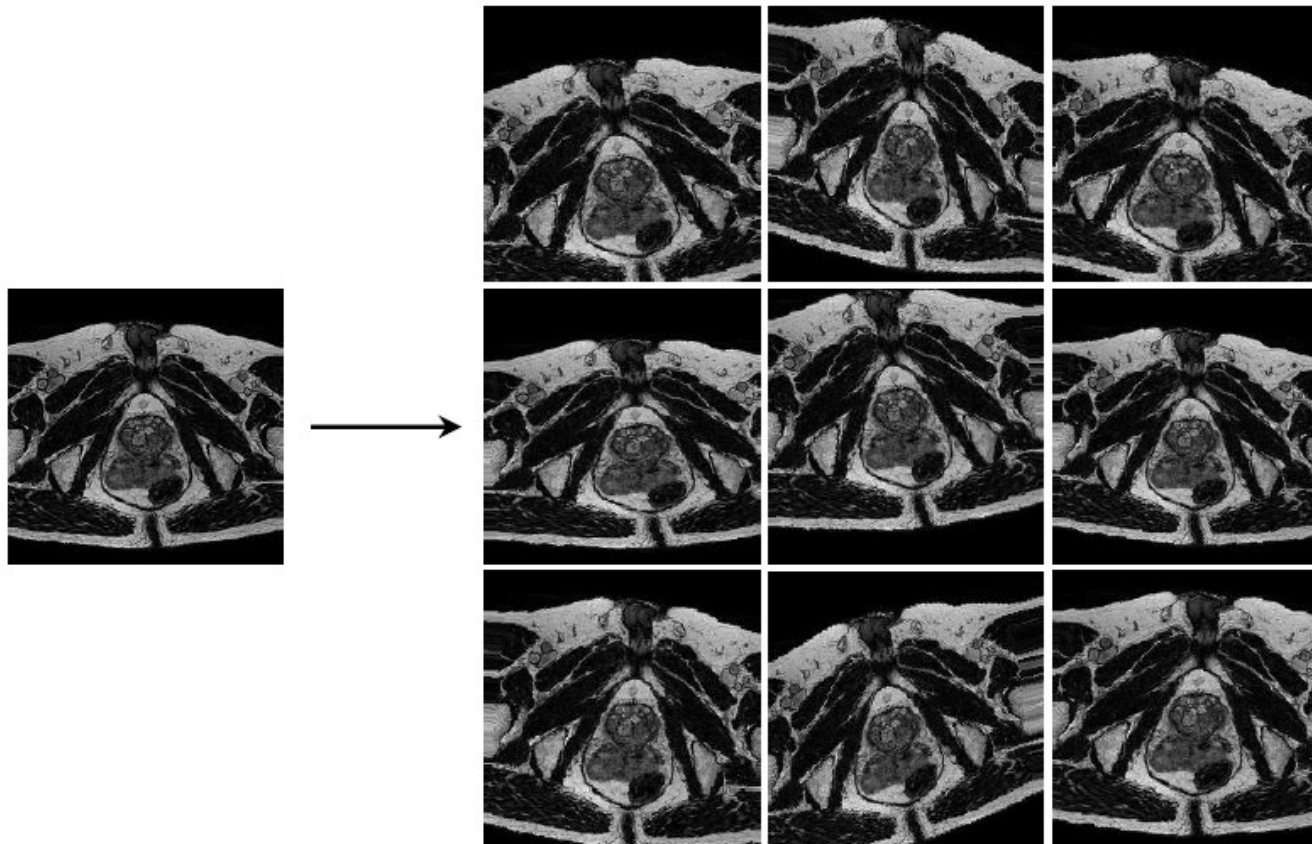
Training curve with sigmoid activation at the output layer and different dropout values



Experiment: Data Augmentation



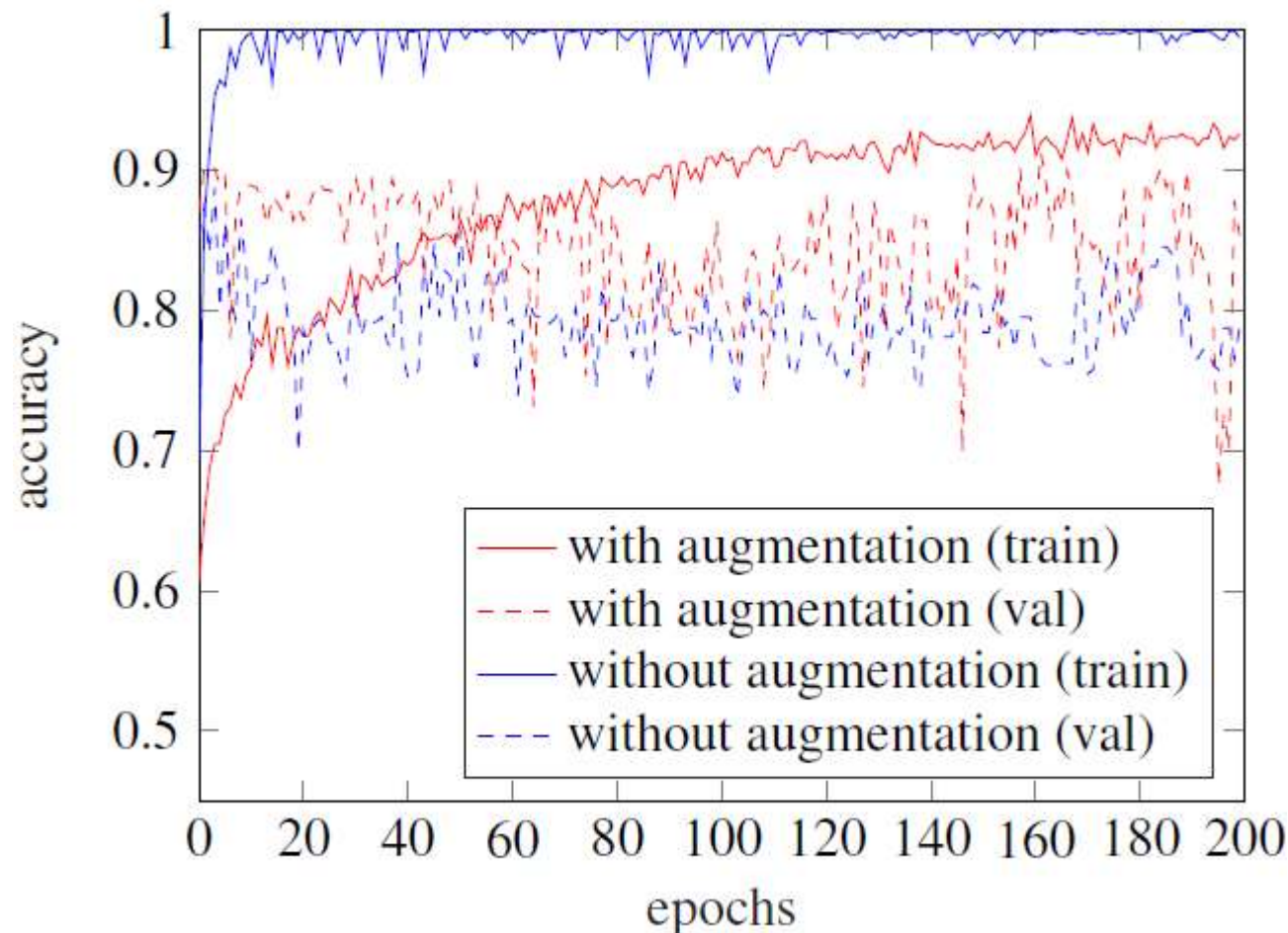
- Useful tool in many image classification tasks
- Artificial enlargement of the dataset → chance for a better classification result with reduced overfitting
- Difficult for medical applications, as input data is usually more homogeneous than real world images



Experiment: Data Augmentation



- Slower increase of training accuracy with data augmentation
- Higher fluctuations from epoch to epoch with data augmentation
- Higher validation accuracy and reduced overfitting



SUMMARY

The initial network architecture obtained the best classification results with an accuracy of 97%

Even though using a small network with a high dropout, overfitting is quite strong, probably due to the relatively small dataset

Data augmentation can reduce overfitting, but reduces training accuracies

- Larger dataset
- Use prostate segmentation as preprocessing step
- Include localization information of the cancer → extend network for cancer localization/segmentation

Thank you!

Questions?
