

UNIVERSITY OF TWENTE.

CARDIAC CINE MR SEGMENTATION ISMRM MACHINE LEARNING TUTORIAL

JELMER WOLTERINK, PHD
DEPARTMENT OF APPLIED MATHEMATICS & TECHNICAL MEDICINE CENTER

15-20 May 2021

Declaration of Financial Interests or Relationships

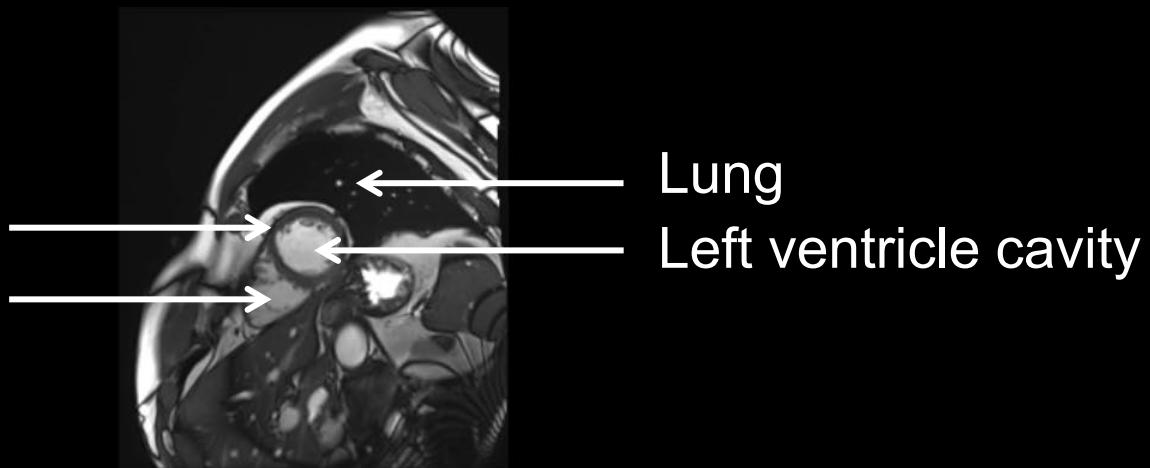
Jelmer Wolterink:

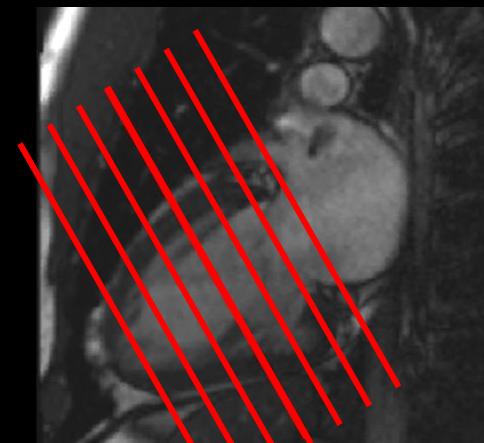
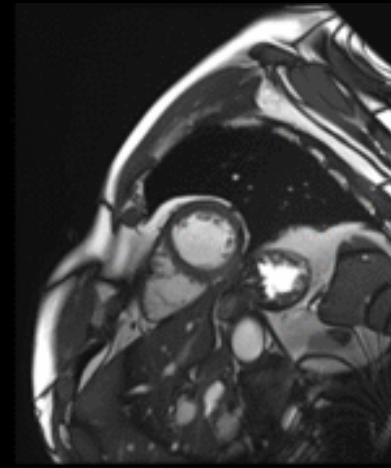
I have no financial interests or relationships to disclose with regard to the subject matter of this presentation.

CONTENT

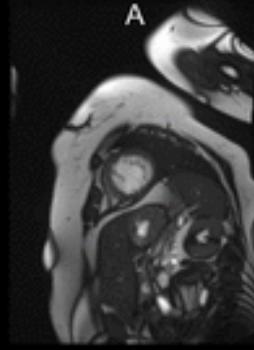
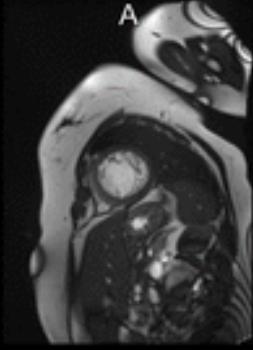
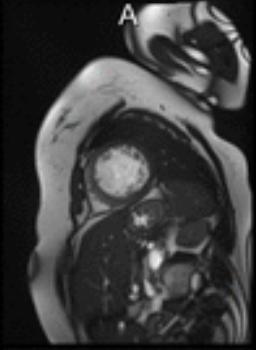
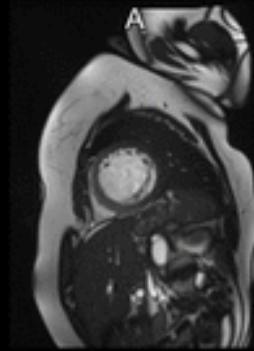
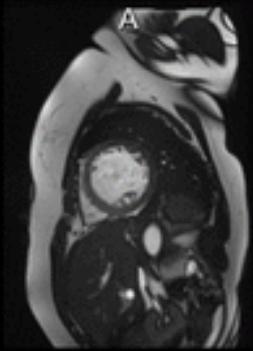
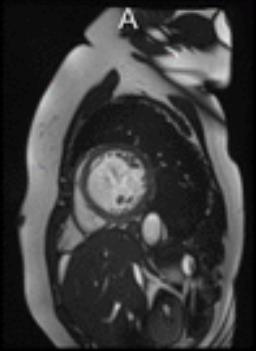
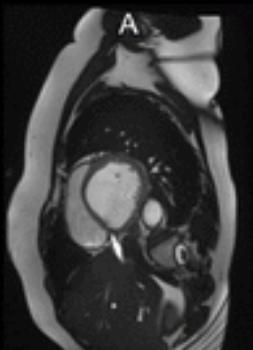
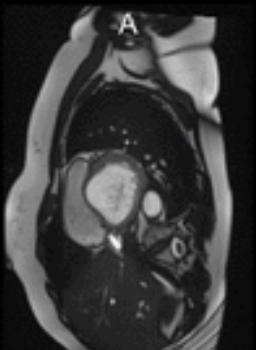
- Hands-on tutorial cardiac cine MR segmentation
- In this presentation
 - 1. Data and task
 - 2. Crash course deep learning
 - 3. Getting started with Python + PyTorch + MONAI

Left ventricle myocardium
Right ventricle





Ejection fraction: 67.2%
Normal

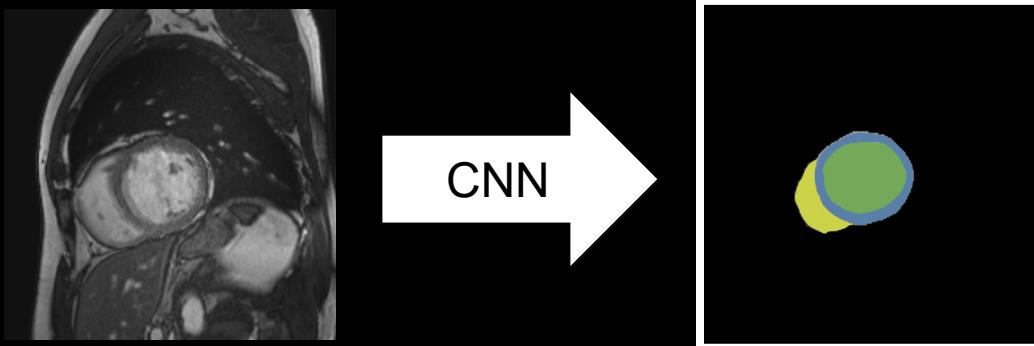


Ejection fraction: 24.4%
Heart failure

TASK IN THIS TUTORIAL

Train a convolutional neural network (CNN) for automatic segmentation in cardiac cine MRI of

- left ventricle cavity
- left ventricle myocardium
- right ventricle



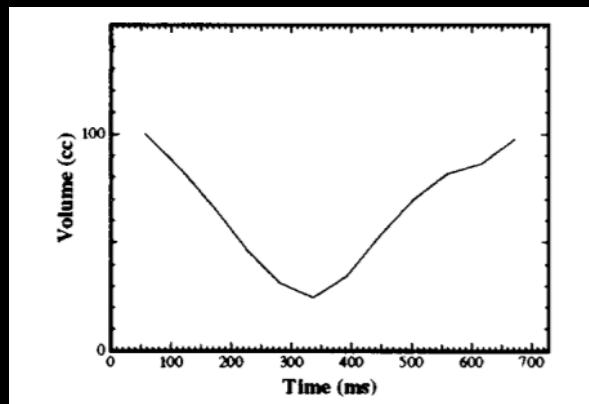
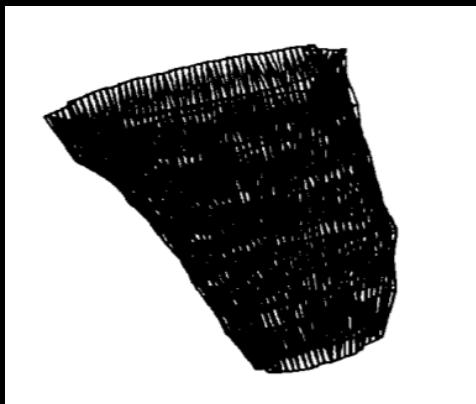
AUTOMATIC SEGMENTATION: 1993

Cardiac MR Image Segmentation Using Deformable Models

A Gupta, L von Kurowski, A Singh, D Geiger, C-C Liang, M-Y Chiu*
LP Adler, M Haacke, and DL Wilson†

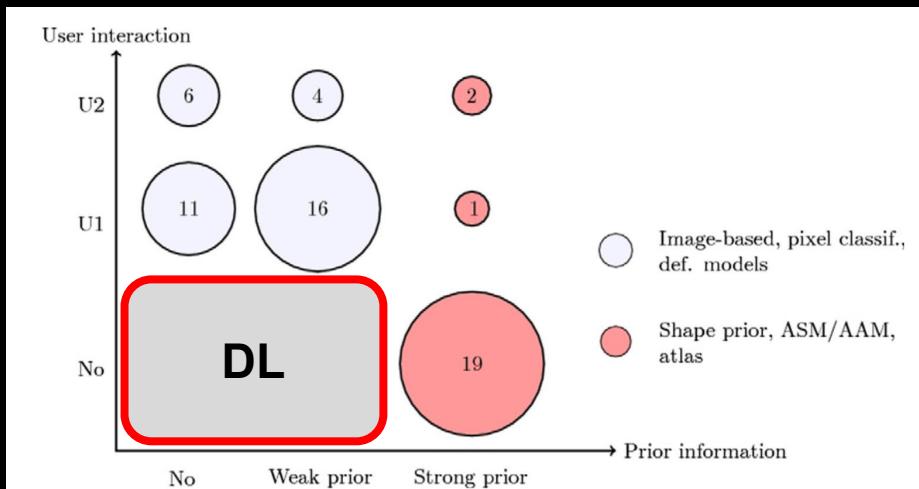
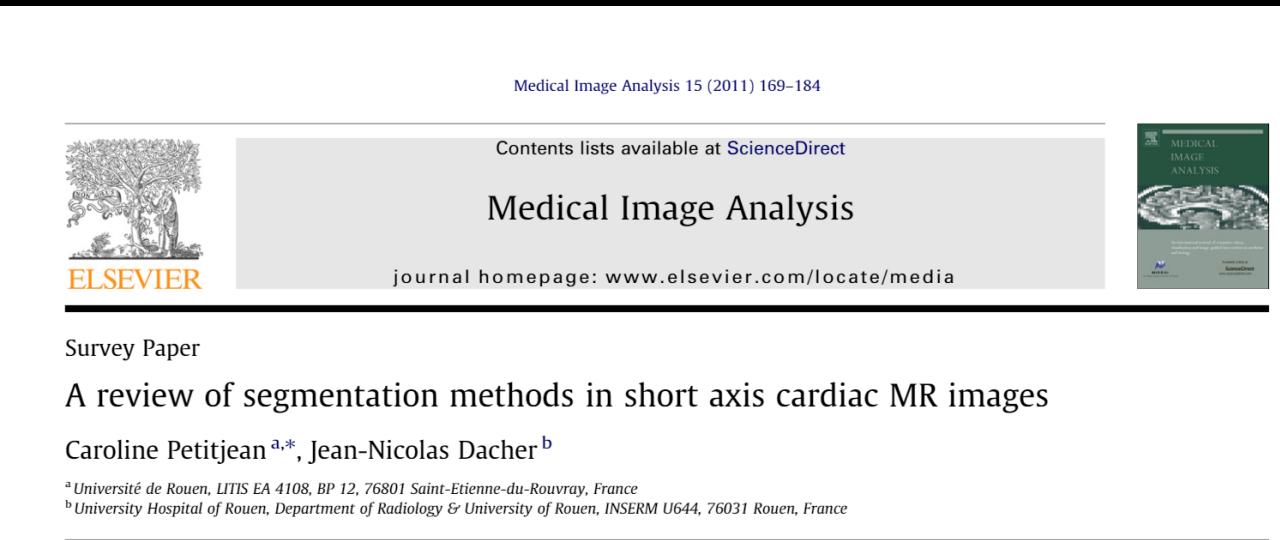
* Siemens Corp. Research, Princeton, USA

† Case Western Reserve Univ., Cleveland, USA



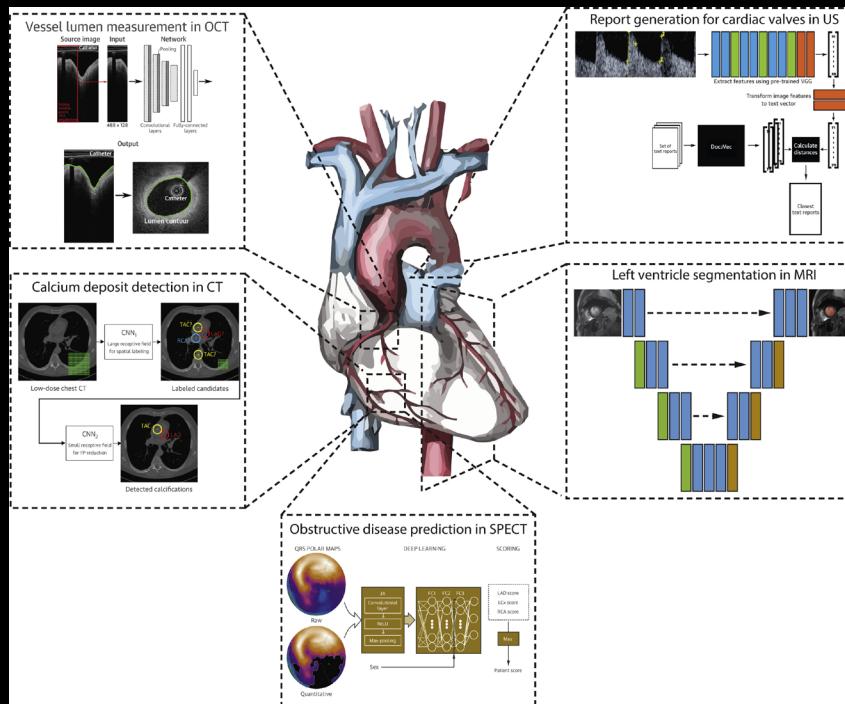
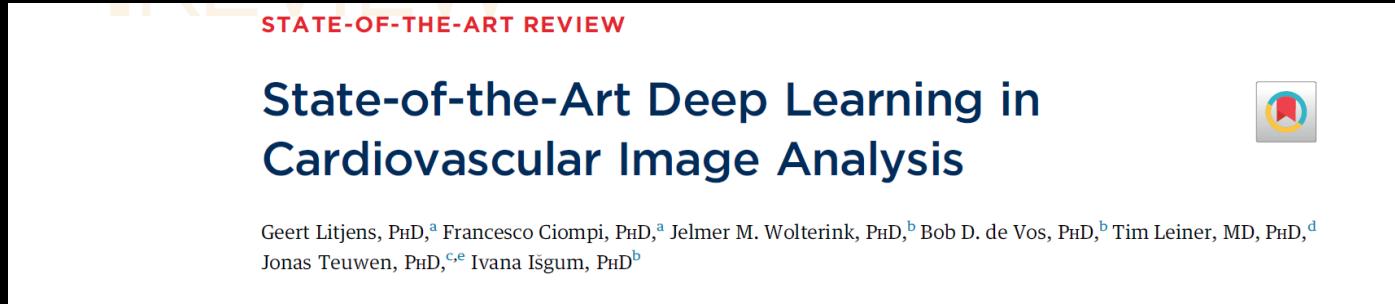
UNIVERSITY
OF TWENTE.

REVIEW: 2011



UNIVERSITY
OF TWENTE.

DEEP LEARNING FOR CVD



UNIVERSITY
OF TWENTE.

PROBLEM SOLVED? 2018

2514

IEEE TRANSACTIONS ON MEDICAL IMAGING, VOL. 37, NO. 11, NOVEMBER 2018



Deep Learning Techniques for Automatic MRI Cardiac Multi-Structures Segmentation and Diagnosis: Is the Problem Solved?

Olivier Bernard¹, Alain Lalande, Clement Zotti¹, Frederick Cervenansky, Xin Yang, Pheng-Ann Heng, Irem Cetin, Karim Lekadir, Oscar Camara, Miguel Angel Gonzalez Ballester, Gerard Sanroma, Sandy Napel, Steffen Petersen, Georgios Tziritas, Elias Grinias, Mahendra Khened, Varghese Alex Kollerathu, Ganapathy Krishnamurthi, Marc-Michel Rohé, Xavier Pennec, Maxime Sermesant¹, Fabian Isensee, Paul Jäger, Klaus H. Maier-Hein, Peter M. Full, Ivo Wolf, Sandy Engelhardt, Christian F. Baumgartner¹, Lisa M. Koch, Jelmer M. Wolterink¹, Ivana Išgum, Yeonggul Jang, Yoonmi Hong, Jay Patravali, Shubham Jain, Olivier Humbert, and Pierre-Marc Jodoin

UNIVERSITY
OF TWENTE.

PARTIALLY

Methods *	ED						ES					
	LV		RV		MYO		LV		RV		MYO	
	D	d_H										
	val.	mm										
O_{1a} vs O_2 (inter-obs)	0.956	5.6	0.930	12.6	0.870	6.7	0.898	8.1	0.866	14.0	0.891	7.6
O_2 vs O_{1b} (inter-obs)	0.950	6.2	0.931	12.1	0.868	7.2	0.895	8.5	0.861	14.1	0.886	8.0
O_{1a} vs O_{1b} (intra-obs)	0.967	4.0	0.957	7.6	0.900	5.1	0.941	5.4	0.930	9.1	0.917	6.0
Average DL methods vs GT	0.965	7.6	0.947	13.2	0.906	10.1	0.927	9.2	0.886	15.2	0.898	10.9
Isensee <i>et al.</i> [44] vs GT	0.968	7.4	0.946	10.1	0.902	8.7	0.931	6.9	0.906	12.1	0.919	8.7
<hr/>												
O_{1a} vs O_2 (inter-obs)	0.956	4.4	0.938	7.7	0.867	5.0	0.913	5.5	0.890	8.7	0.894	5.5
O_2 vs O_{1b} (inter-obs)	0.953	4.9	0.937	8.6	0.864	5.5	0.905	5.8	0.898	9.4	0.886	6.1
O_{1a} vs O_{1b} (intra-obs)	0.971	3.1	0.960	5.8	0.905	3.6	0.950	3.9	0.940	6.9	0.923	4.4
Average DL methods vs GT	0.972	3.7	0.951	8.1	0.896	5.2	0.929	4.2	0.899	9.9	0.915	6.1
Isensee <i>et al.</i> [44] vs GT	0.972	3.7	0.969	6.4	0.910	4.6	0.945	4.2	0.912	8.6	0.930	5.1

* ED: End diastole; ES: End systole; LV: Endocardial contour of the left ventricle; RV: Endocardial contour of the right Ventricle; Myo: Myocardium contours; D: Dice Index; d_H : Hausdorff distance; GT: Ground-truth.

NOT ENTIRELY

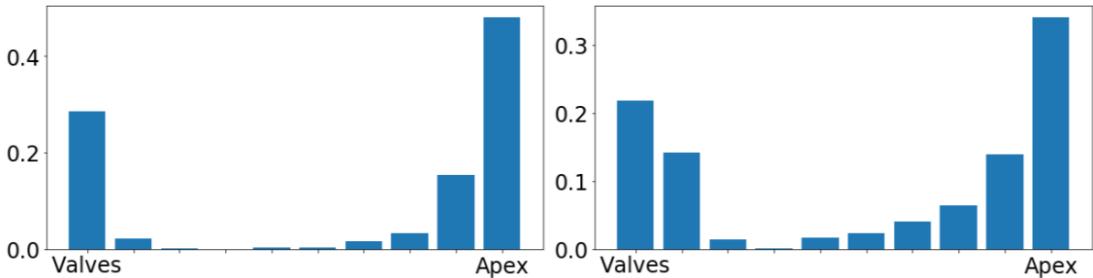


Fig. 3. Histogram of degenerated slices ED (left), and ES (right).

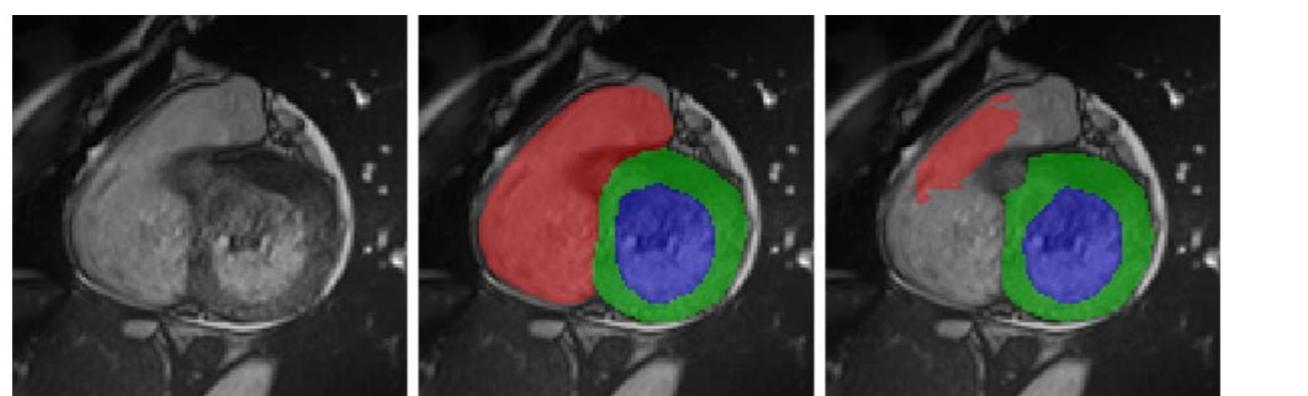


Fig. 4. Typical degenerated result at the base of the heart. [Left] input image; [Middle] ground truth; [Right] prediction.

UNIVERSITY
OF TWENTE.

NEURAL NETWORK TRAINING

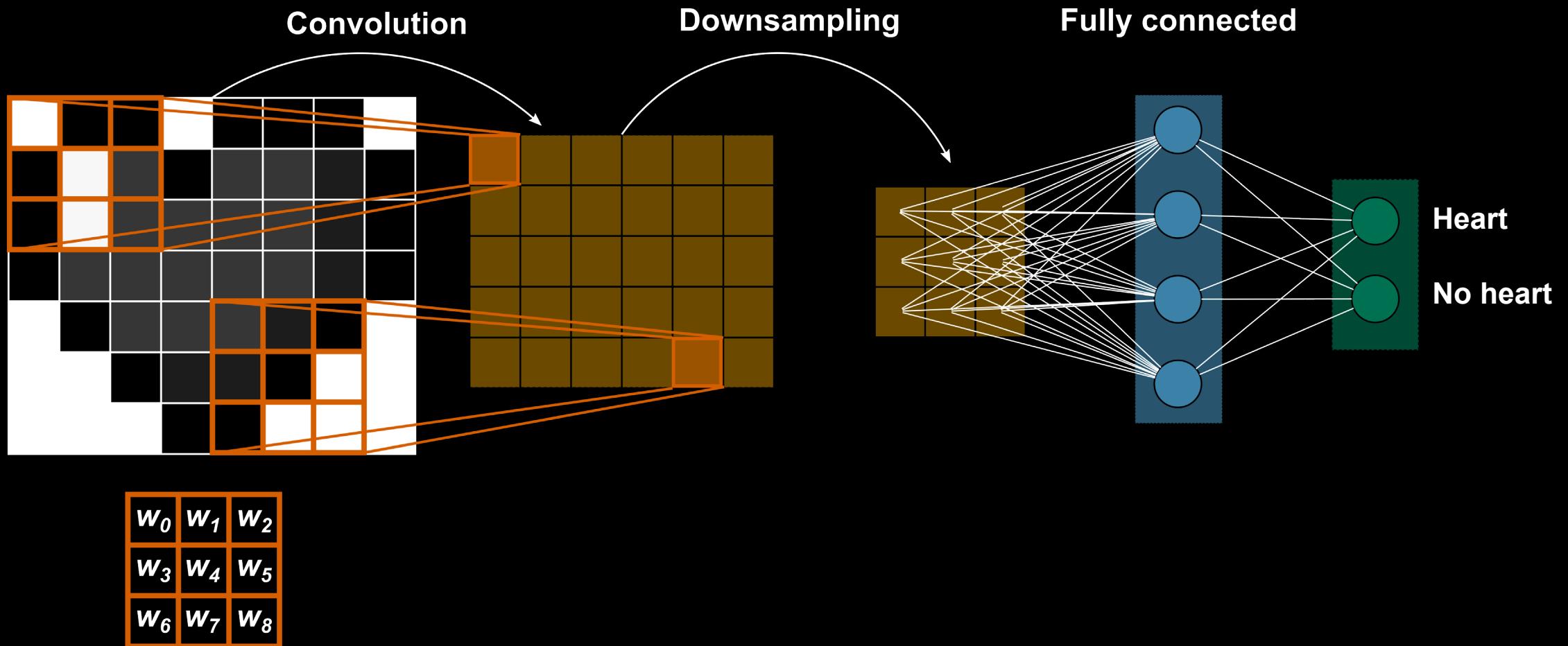
1. Training stage

- **Optimize** the parameters of the **network**
- Minimize **loss**, backpropagation of errors
- Required: **images** with reference **segmentations** (training set)

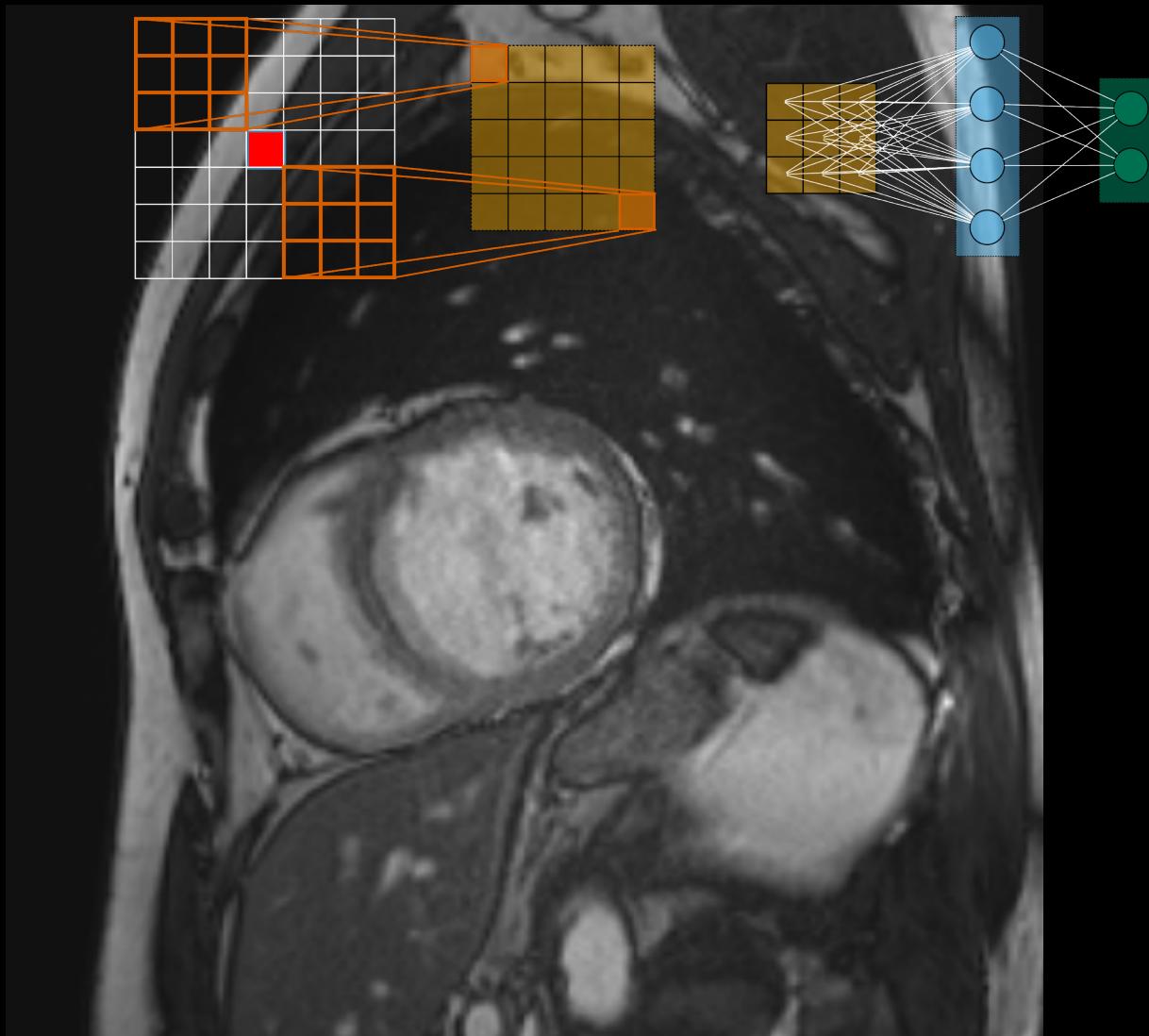
2. Test stage

- Apply the trained networks
- Required: images with reference segmentations (test set)

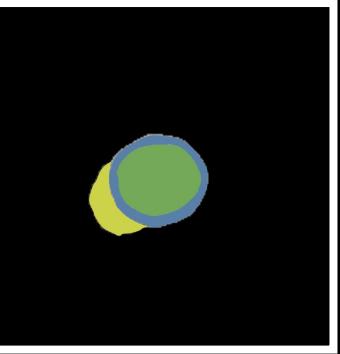
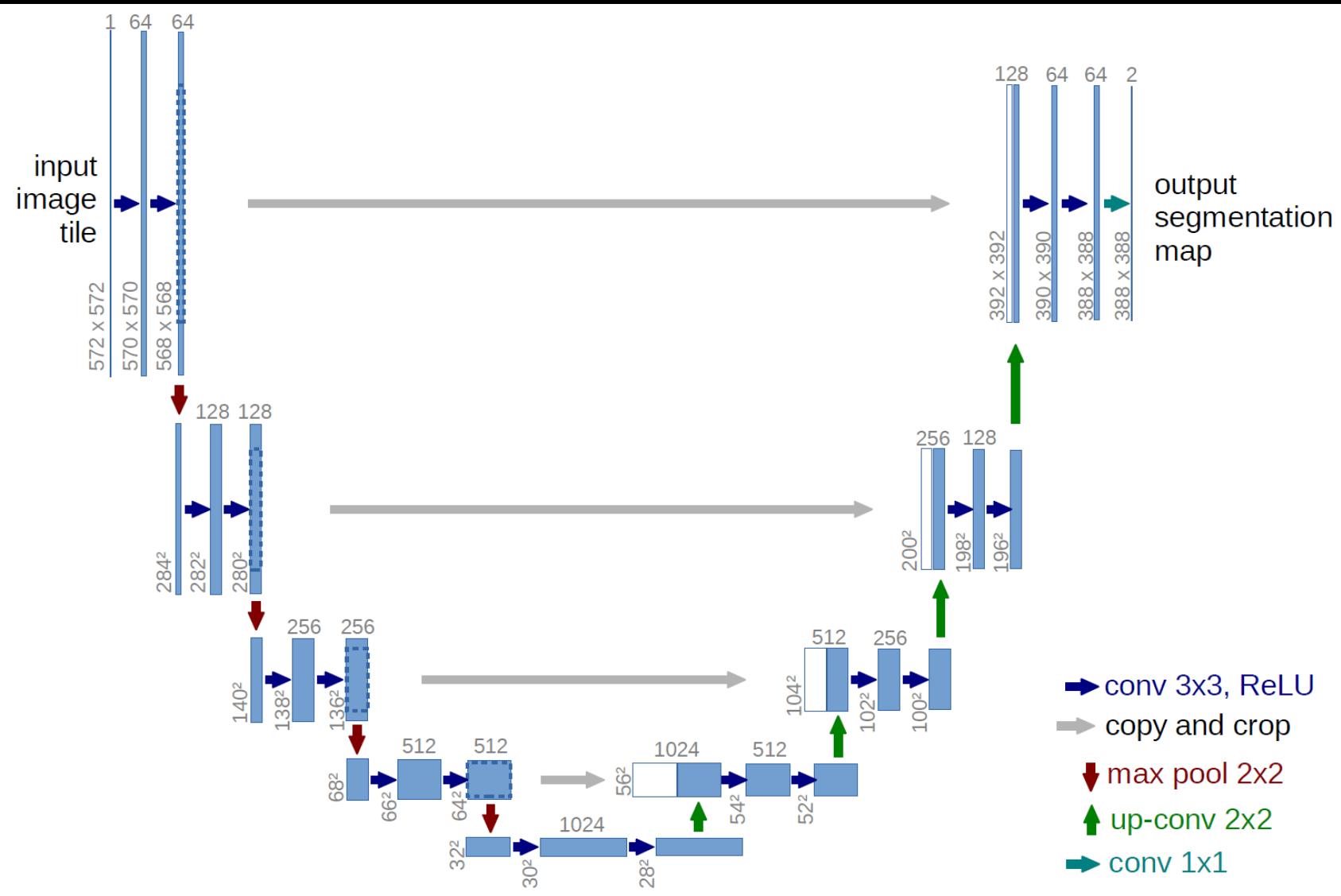
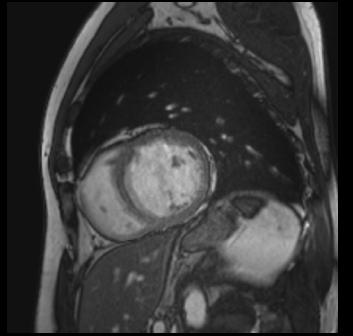
CONVOLUTIONAL NEURAL NETWORK (CNN)



SLIDING-WINDOW PIXEL CLASSIFICATION

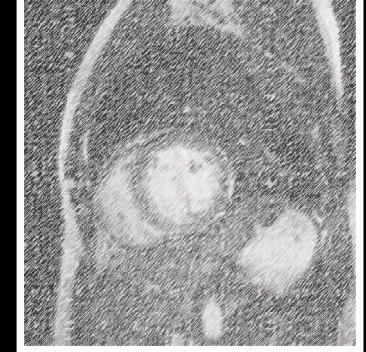
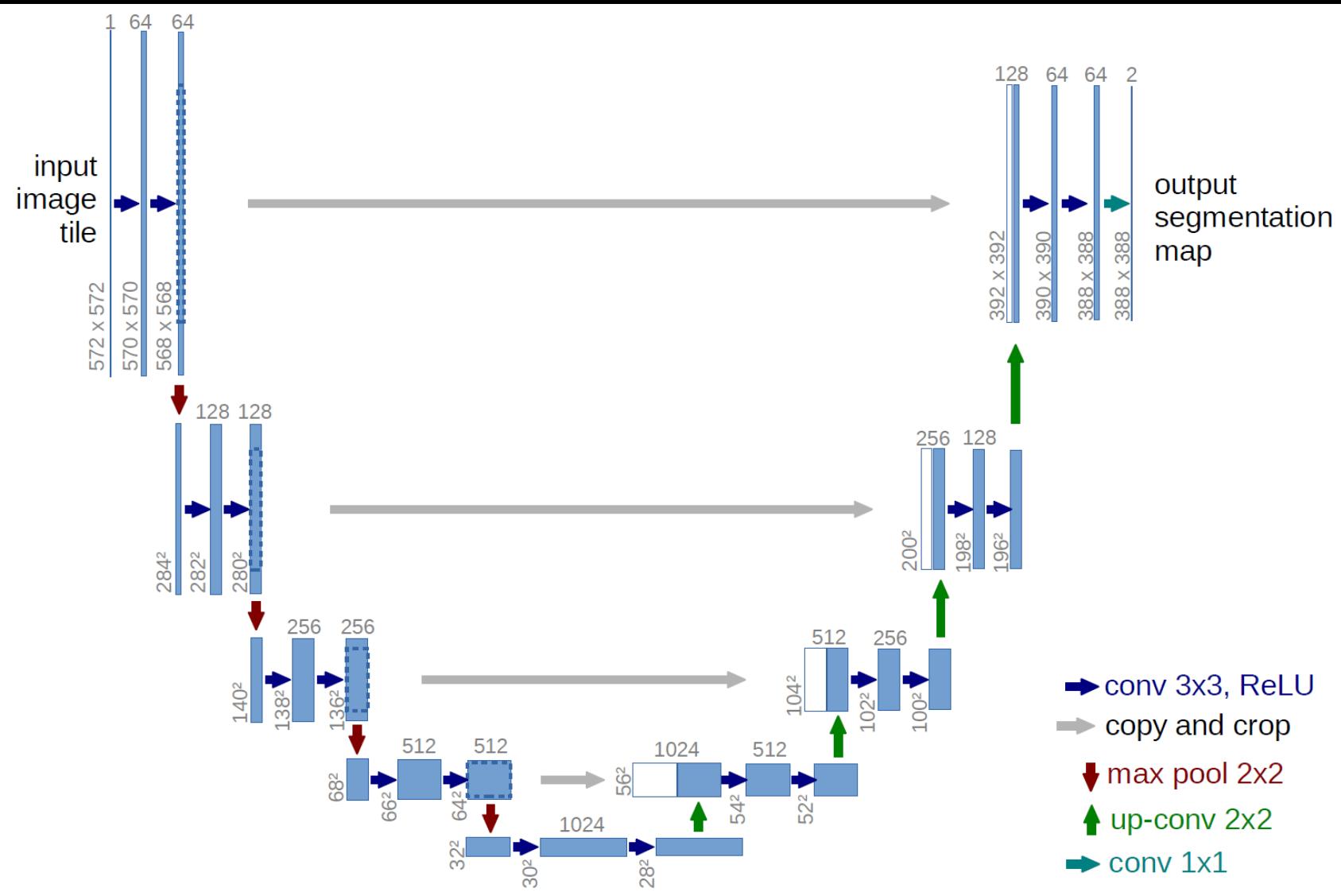
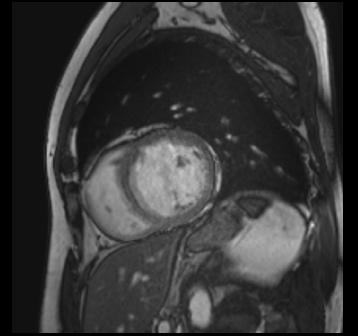


2D U-NET



UNIVERSITY
OF TWENTE.

2D U-NET



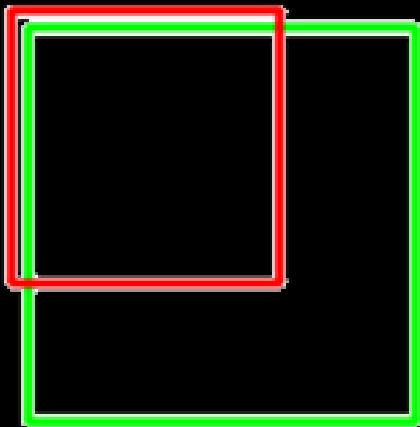
UNIVERSITY
OF TWENTE.

DICE LOSS

Based on Dice similarity coefficient (DSC)

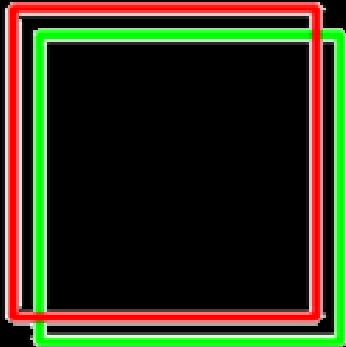
$$DSC = \frac{2|X \cap Y|}{|X| + |Y|}$$

DSC = 0.57



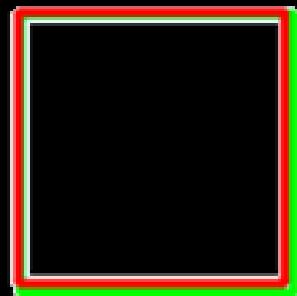
Poor

DSC = 0.83



Good

DSC = 0.96

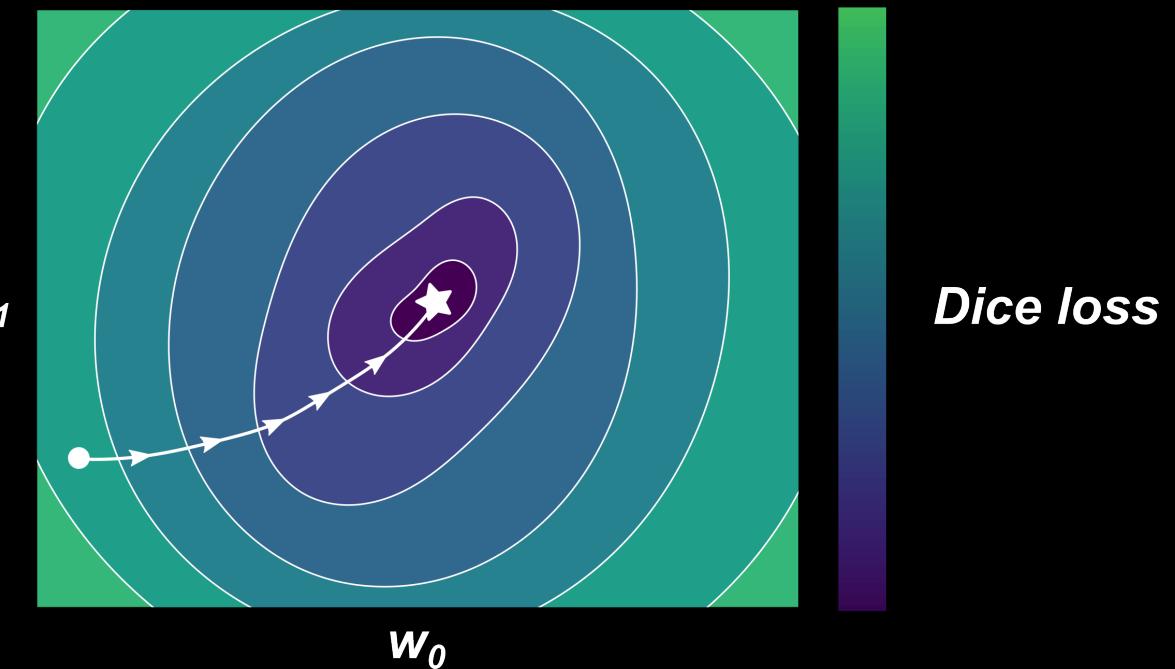


Excellent

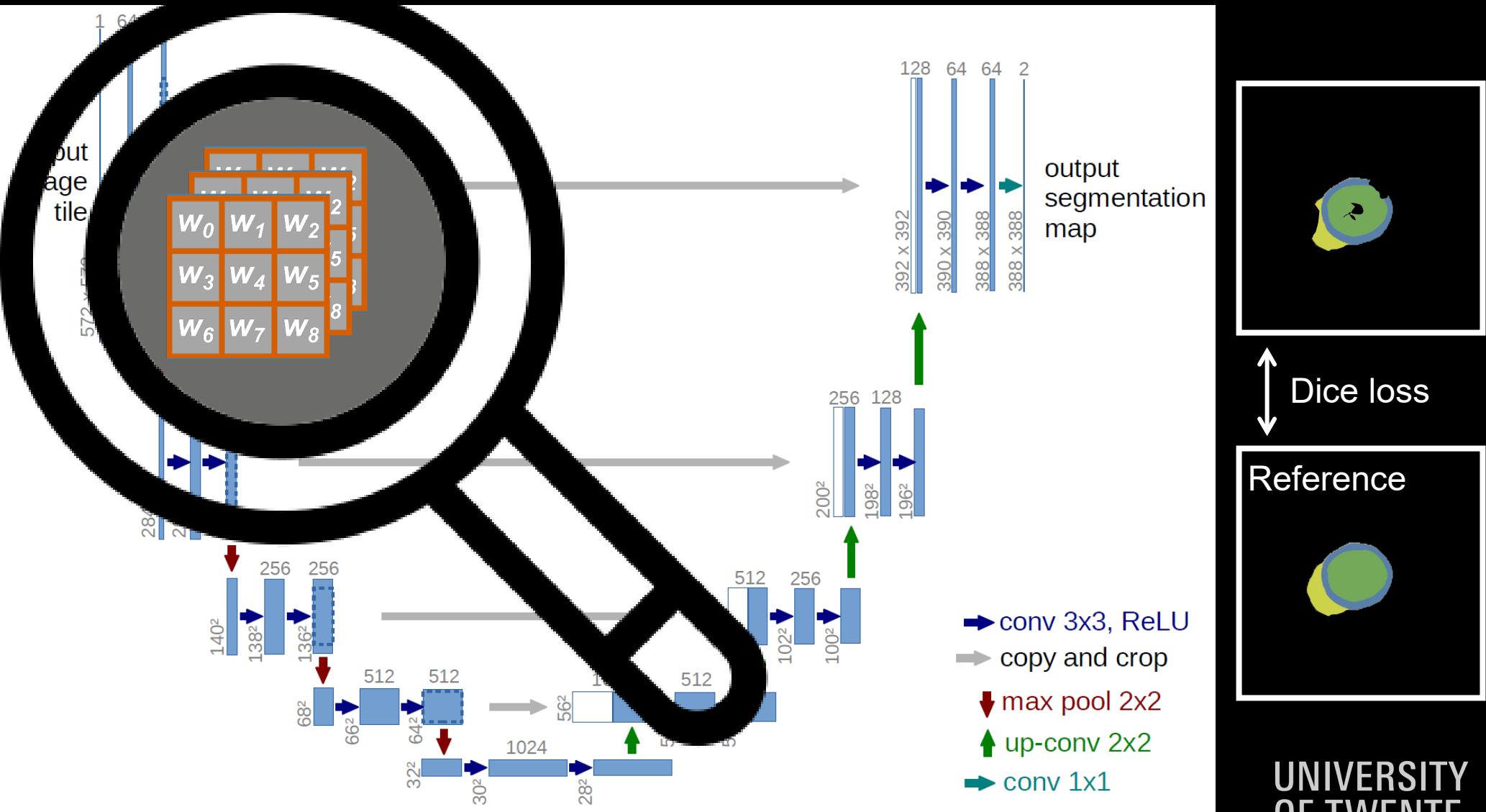
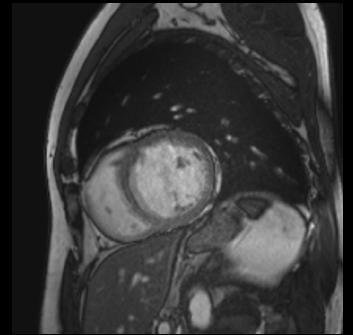
OPTIMIZATION

Iterative process, in each iteration

1. Take random ‘mini-batch’ of samples
2. Obtain prediction on samples and compute loss w.r.t. reference
3. Backpropagate loss
4. Update weights
5. Repeat



BACKPROPAGATION



UNIVERSITY
OF TWENTE.

SUMMARY

- What do we (at least) need
 1. A data set
 2. A neural network architecture
 3. A loss function
 4. An optimization algorithm

DATA SET

- Sunnybrook cardiac MR data set¹
 - 45 patients, 4 pathologies
- Multi-Centre, Multi-Vendor & Multi-Disease Cardiac Image Segmentation Challenge (M&Ms)²
 - 350 patients, 4 scanners, 8 pathologies
- Automatic Cardiac Diagnosis Challenge (ACDC)³
 - 150 patients, 4 pathologies

¹ <https://www.cardiacatlas.org/studies/sunnybrook-cardiac-data/>

² <https://www.ub.edu/mnms/>

³ <https://www.creatis.insa-lyon.fr/Challenge/acdc/>

DATA SET

- Sunnybrook cardiac MR data set¹
 - 45 patients, 4 pathologies
- Multi-Centre, Multi-Vendor & Multi-Disease Cardiac Image Segmentation Challenge (M&Ms)²
 - 350 patients, 4 scanners, 8 pathologies
- Automatic Cardiac Diagnosis Challenge (ACDC)³
 - 150 patients, 4 pathologies

¹ <https://www.cardiacatlas.org/studies/sunnybrook-cardiac-data/>

² <https://www.ub.edu/mnms/>

³ <https://www.creatis.insa-lyon.fr/Challenge/acdc/>

ACDC DATA SET

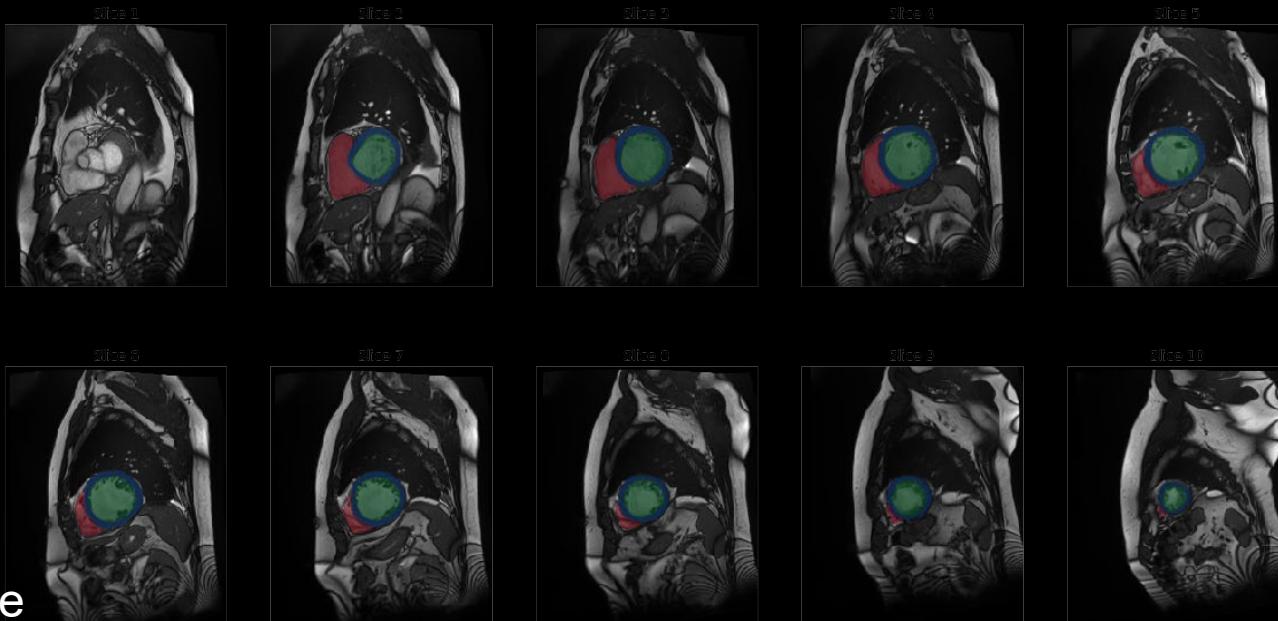
For each patient

- 20-30 frames
- Annotations in end-diastole/end-systole frames

Training set: 100 patients

Test set: 50 patients

Pt 1, end-diastole



- Left ventricle
- Myocardium
- Right ventricle

ACDC DATA SET

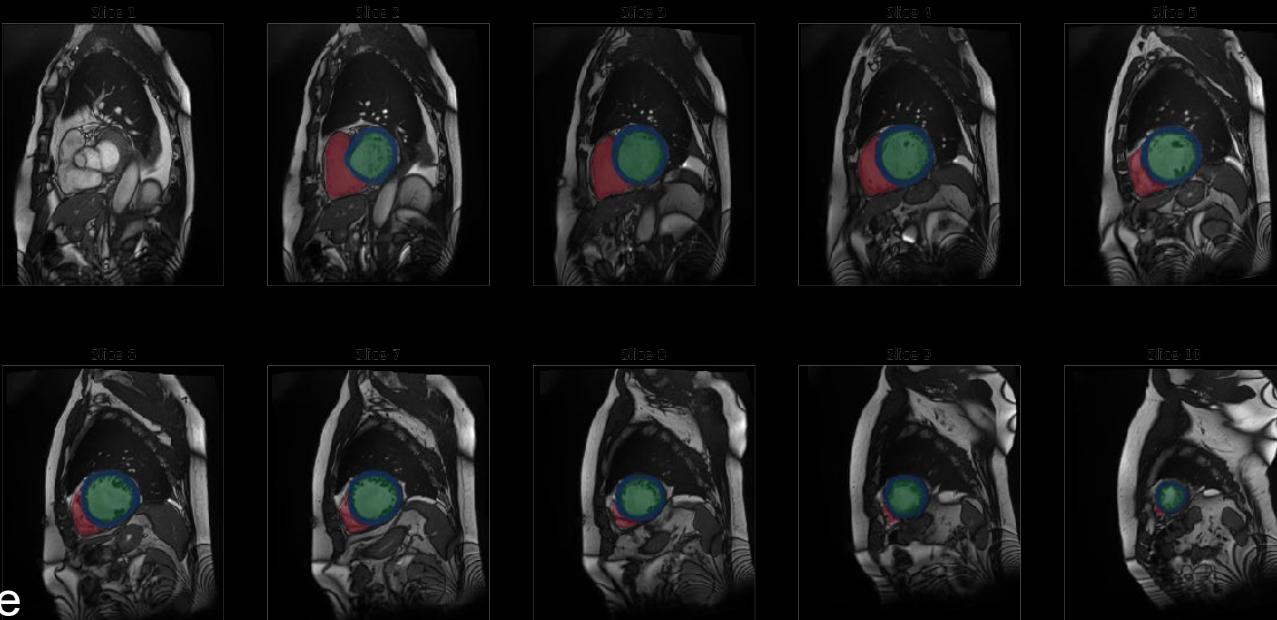
For each patient

- 20-30 frames
- Annotations in end-diastole/end-systole frames

Training set: 100 patients

Test set: 50 patients

Pt 1, end-diastole



- Left ventricle
- Myocardium
- Right ventricle

PYTHON GETTING STARTED

1. Install a Python distribution: Anaconda¹



2. Install a Python editor: JupyterLab²



3. Start Jupyter

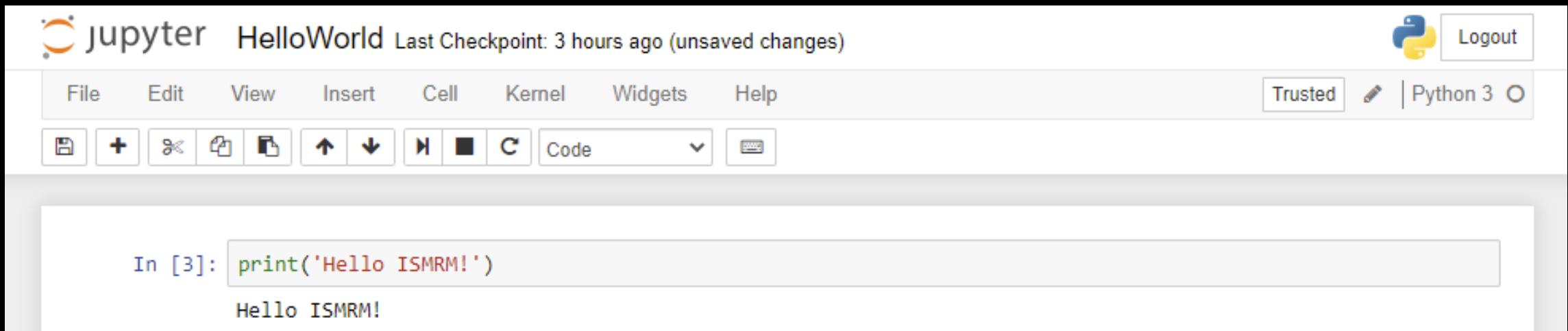
```
$ jupyter notebook
```

¹ <https://www.anaconda.com/products/individual>

² <https://jupyter.org/>, <https://jupyterlab.readthedocs.io/en/stable/>

PYTHON GETTING STARTED

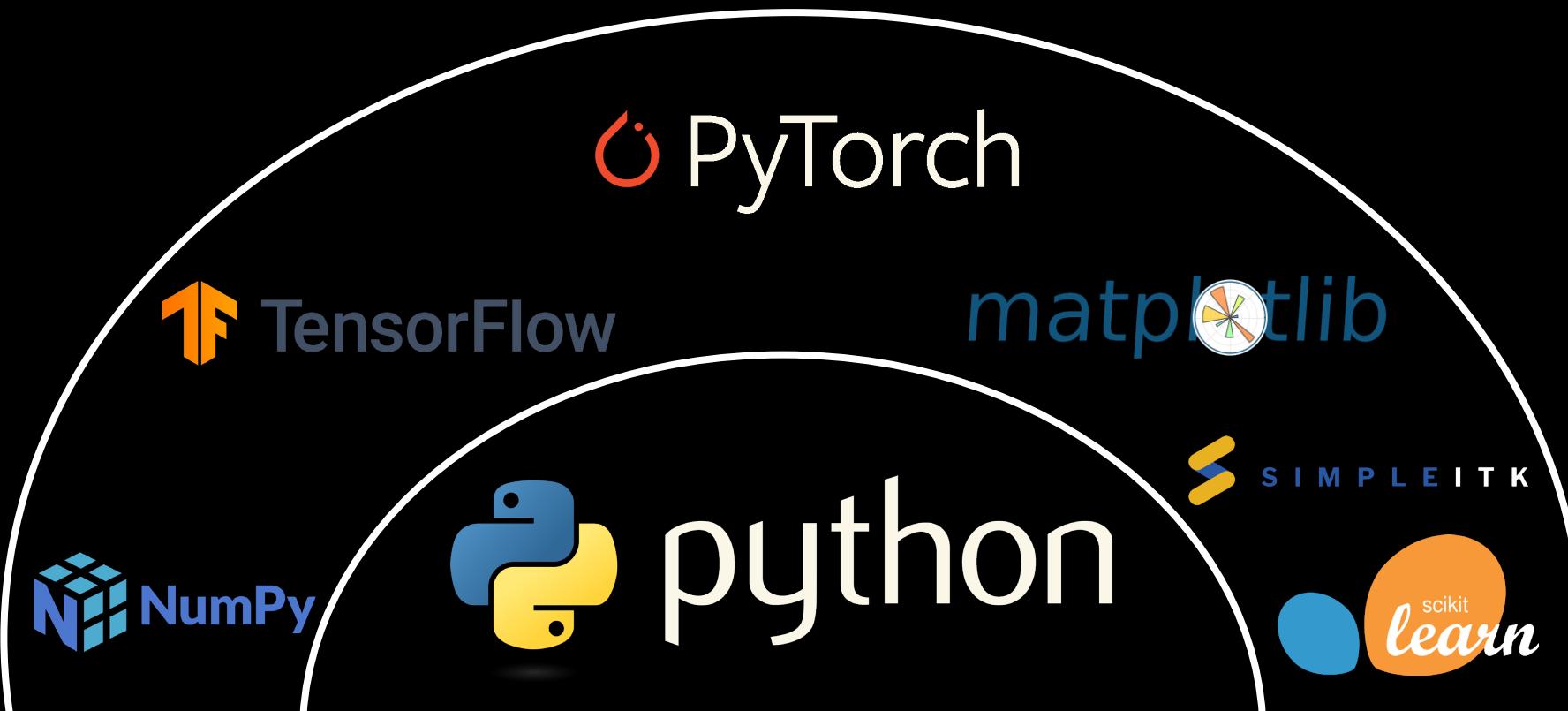
Jupyter allows you to edit and run Python in a browser



¹ <https://www.anaconda.com/products/individual>

² <https://jupyter.org/>

PYTHON PACKAGES



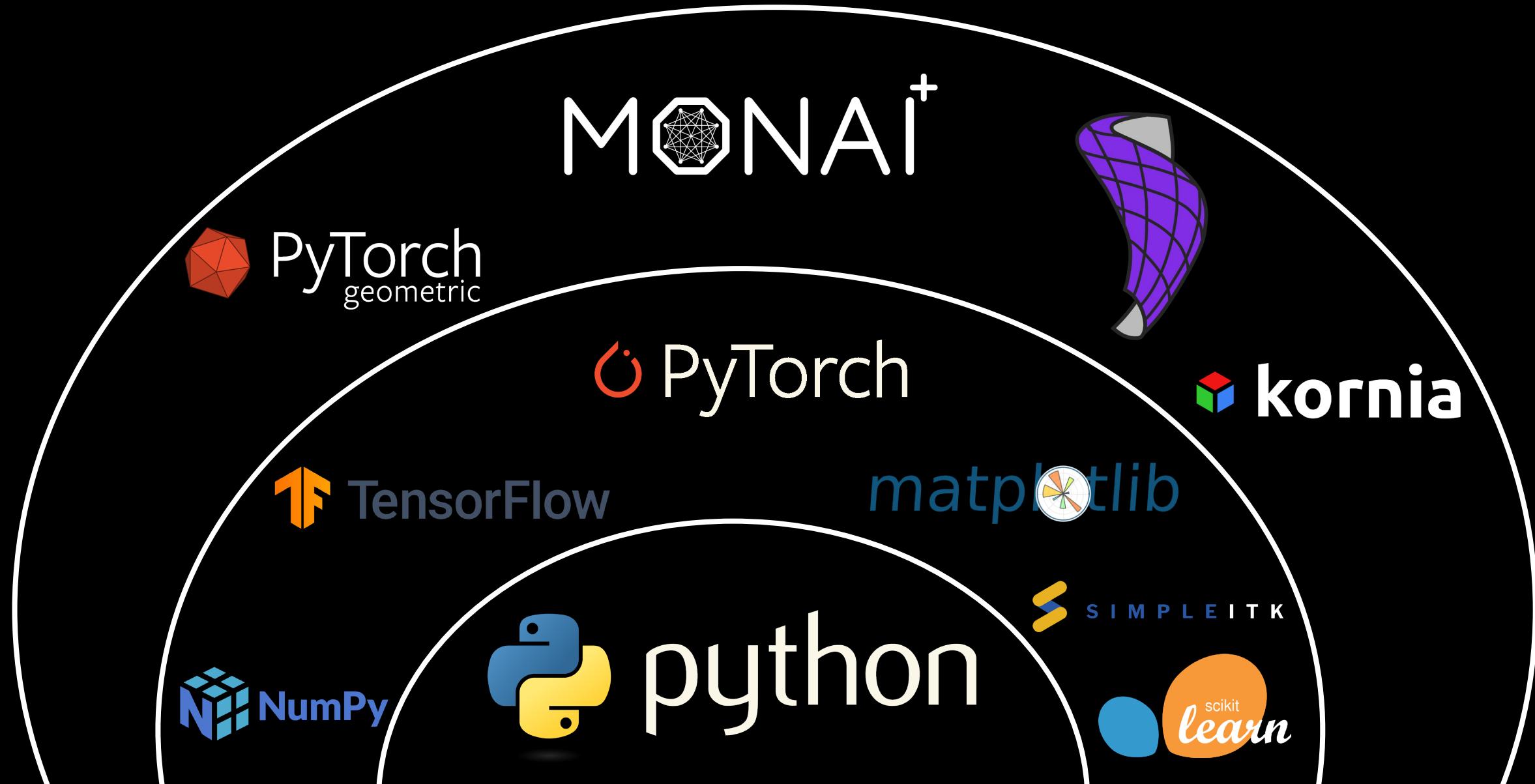
PYTORCH¹

- Provides important deep learning building blocks
 - Optimizers
 - Neural network layers
 - Running on different devices
 - Data augmentation
 - Data loaders
 - ...
- Very versatile, well-documented²

¹ <https://pytorch.org/>

² <https://pytorch.org/tutorials/beginner/basics/intro.html>

PYTORCH ECOSYSTEM



MONAI¹

- PyTorch for healthcare data
- Medical image loaders
 - NIfTI, DICOM, MHD, ...
- Medical imaging networks
 - U-Net, V-Net, ...
- Medical imaging data augmentation
 - Random deformations
 - Random bias fields (for MRI)

¹ <https://monai.io/>

LOADING AND VIEWING DATA

```
In [3]: transform = monai.transforms.LoadImageD(("image", "label"))

file_dict = {"image": "{}/patient001/patient001_frame01.nii.gz".format(datapath),
             "label": "{}/patient001/patient001_gt.nii.gz".format(datapath)}

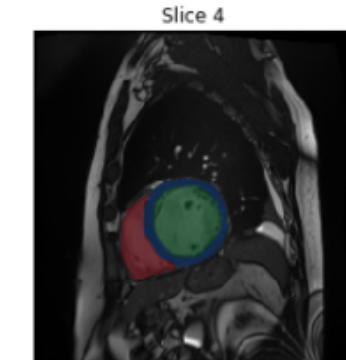
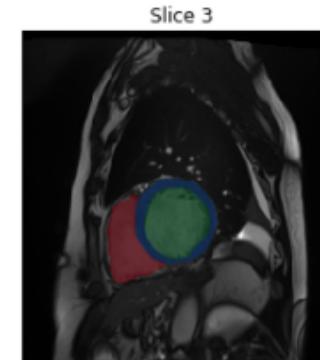
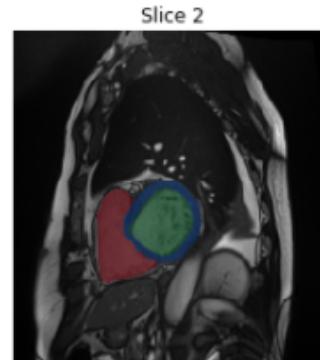
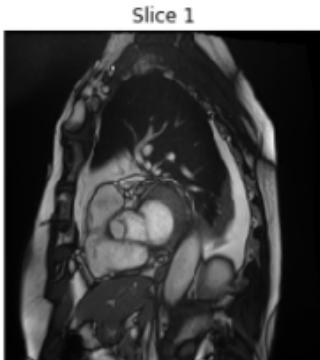
data_dict = transform(file_dict)
```

We then use Matplotlib to visualize the images in a rectangular grid. The for-loop iterates over z, the slice index. Each slice is shown, and the reference label is shown as colored overlay.

```
In [4]: def visualize_data(pt_dict):
    plt.figure(figsize=(20,20))

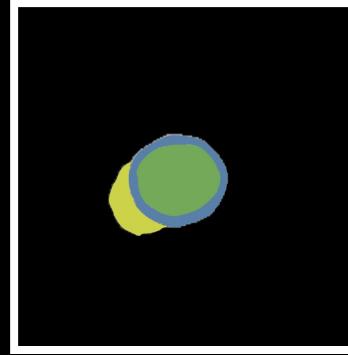
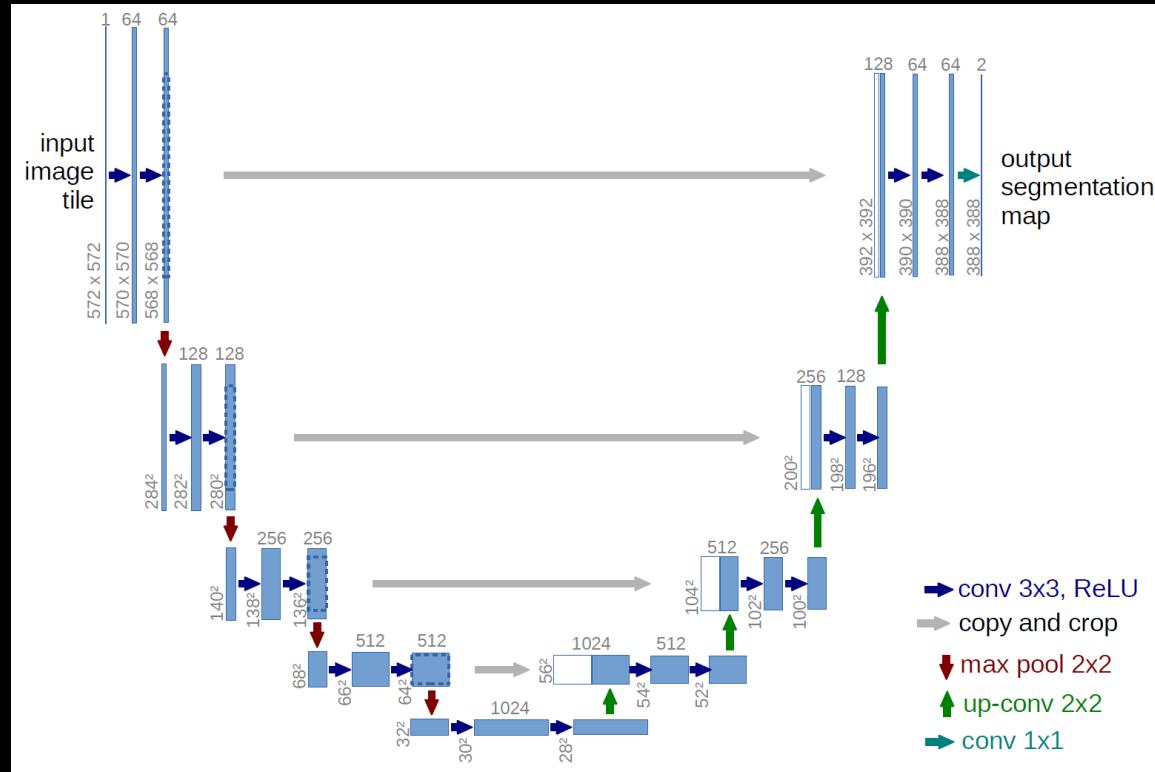
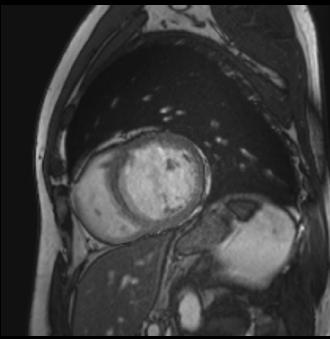
    for z in range(pt_dict["image"].shape[0]):
        plt.subplot(np.ceil(np.sqrt(pt_dict["image"].shape[0])), np.ceil(np.sqrt(pt_dict["image"].shape[0])), 1 + z)
        plt.imshow(pt_dict["image"][z, :, :], cmap='gray')
        plt.axis('off')
        plt.imshow(np.ma.masked_where(pt_dict["label"][[z, :, :]]!=2, pt_dict["label"][[z, :, :]]==2), alpha=0.6, cmap='Blues', clim=(0, 1))
        plt.imshow(np.ma.masked_where(pt_dict["label"][[z, :, :]]!=3, pt_dict["label"][[z, :, :]]==3), alpha=0.6, cmap='Greens', clim=(0, 1))
        plt.imshow(np.ma.masked_where(pt_dict["label"][[z, :, :]]!=1, pt_dict["label"][[z, :, :]]==1), alpha=0.6, cmap='Reds', clim=(0, 1))
        plt.title('Slice {}'.format(z + 1))
    plt.show()

visualize_data(data_dict)
```



CODE

NETWORK

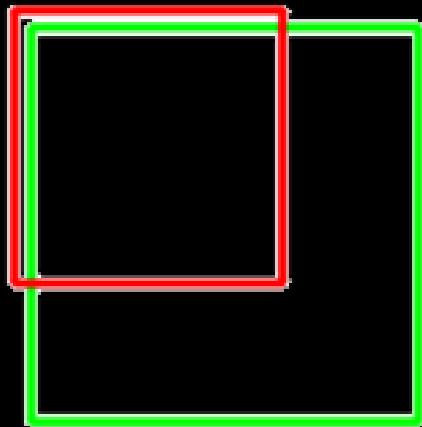


```
model = monai.networks.nets.UNet(dimensions=2,  
        in_channels=1, out_channels=4,  
        channels=(64, 128, 256, 512, 1024))
```

CODE

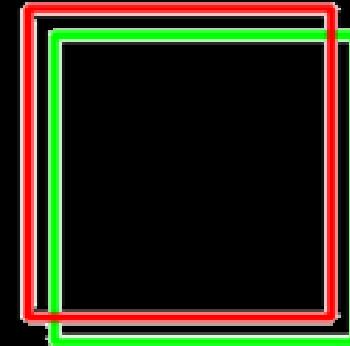
DICE LOSS

DSC = 0.57



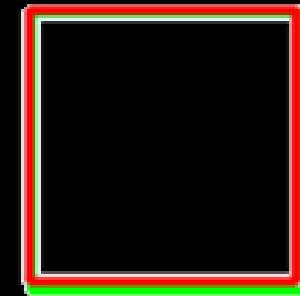
Poor

DSC = 0.83



Good

DSC = 0.96



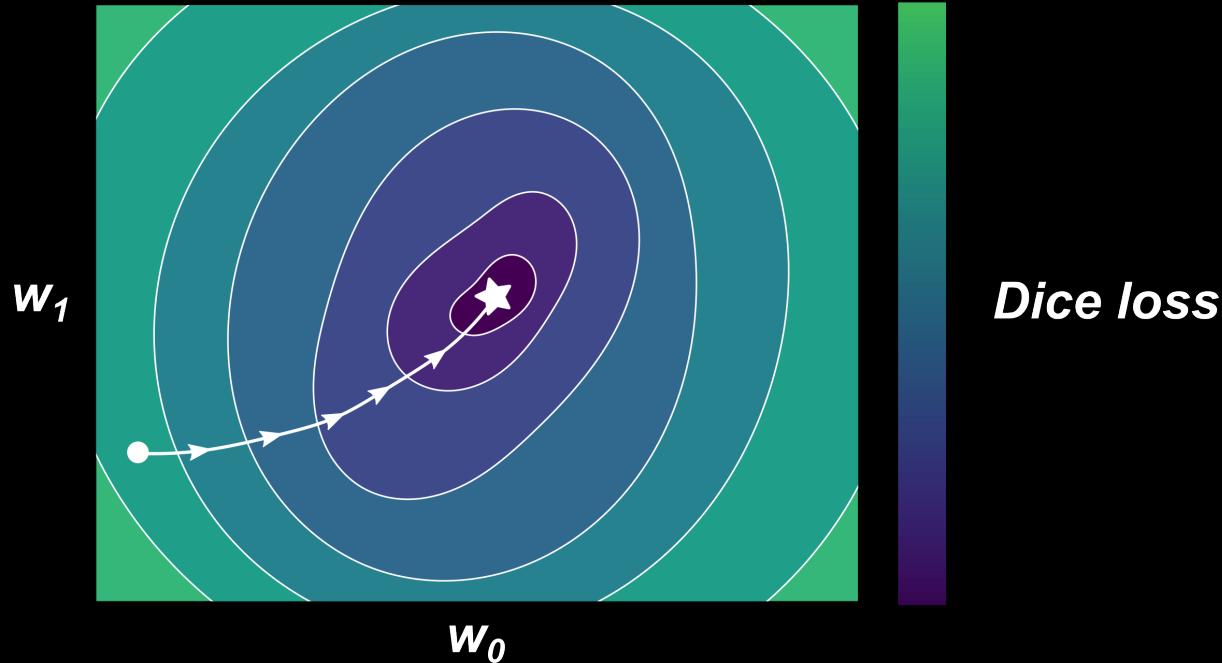
Excellent

```
loss_function = monai.losses.DiceLoss(softmax=True)
loss = loss_function(input, target)
```

CODE

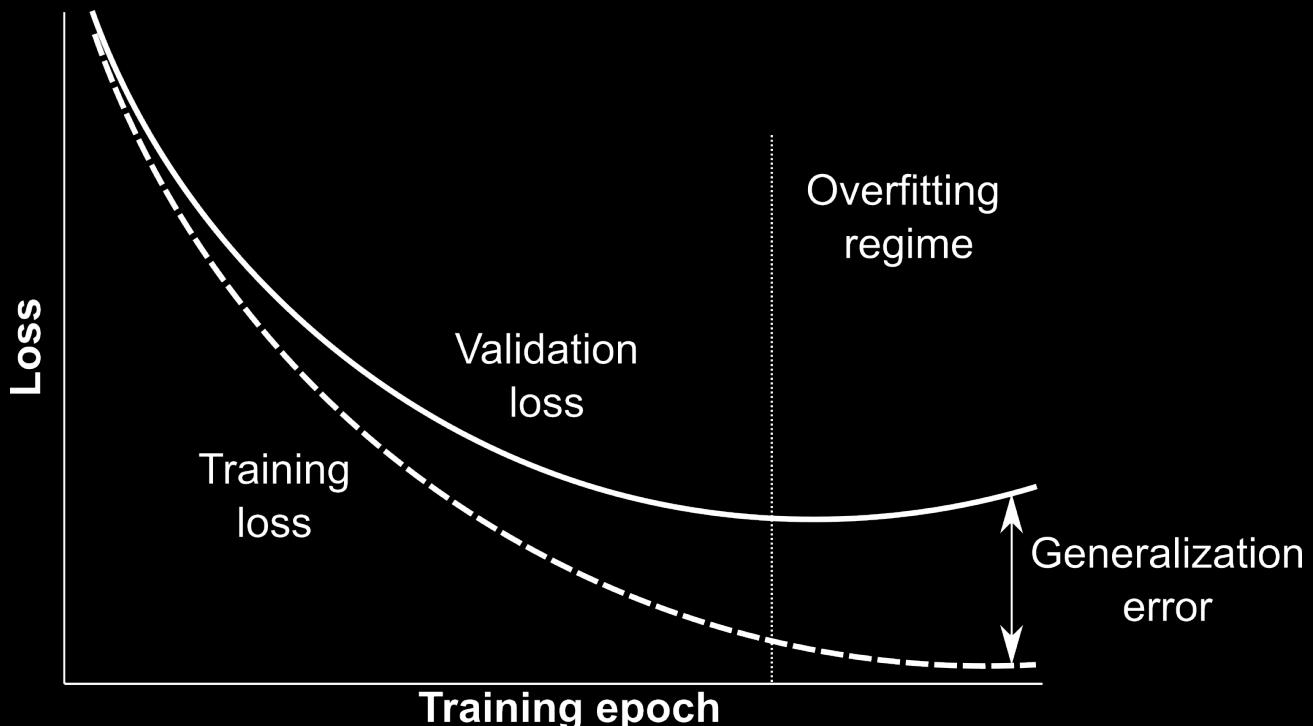
```
optimizer = torch.optim.Adam(model.parameters(), 1e-3)
loss.backward()
optimizer.step()
```

OPTIMIZER



MONITOR YOUR TRAINING PROCESS

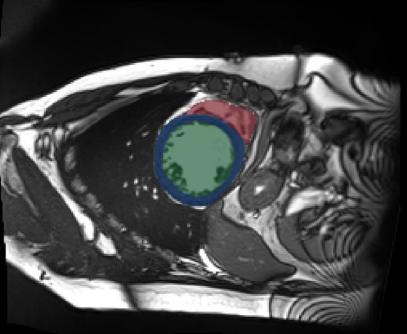
- Monitor loss curves and identify if the model has converged
- Spot overfitting/generalization error



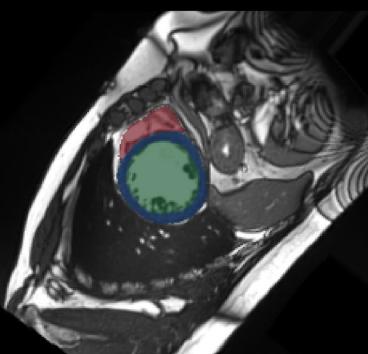
DATA AUGMENTATION

- Add artificial samples to the training data set
- Wide range of randomized image transforms possible

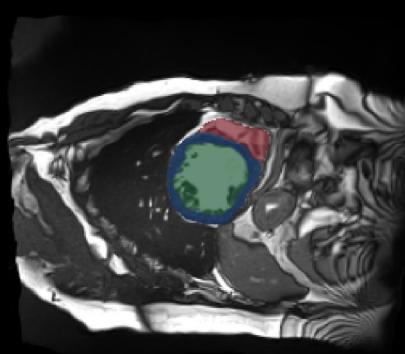
Original



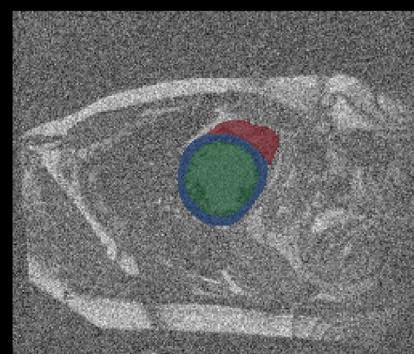
Rotate



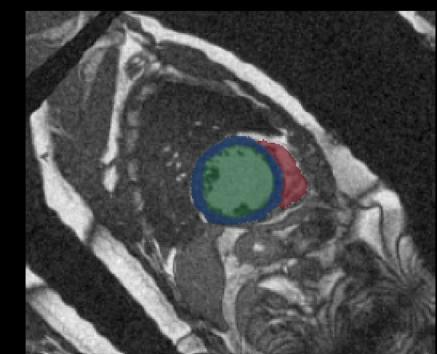
Elastic deformation



Gaussian noise



All three



VALIDATING YOUR RESULTS

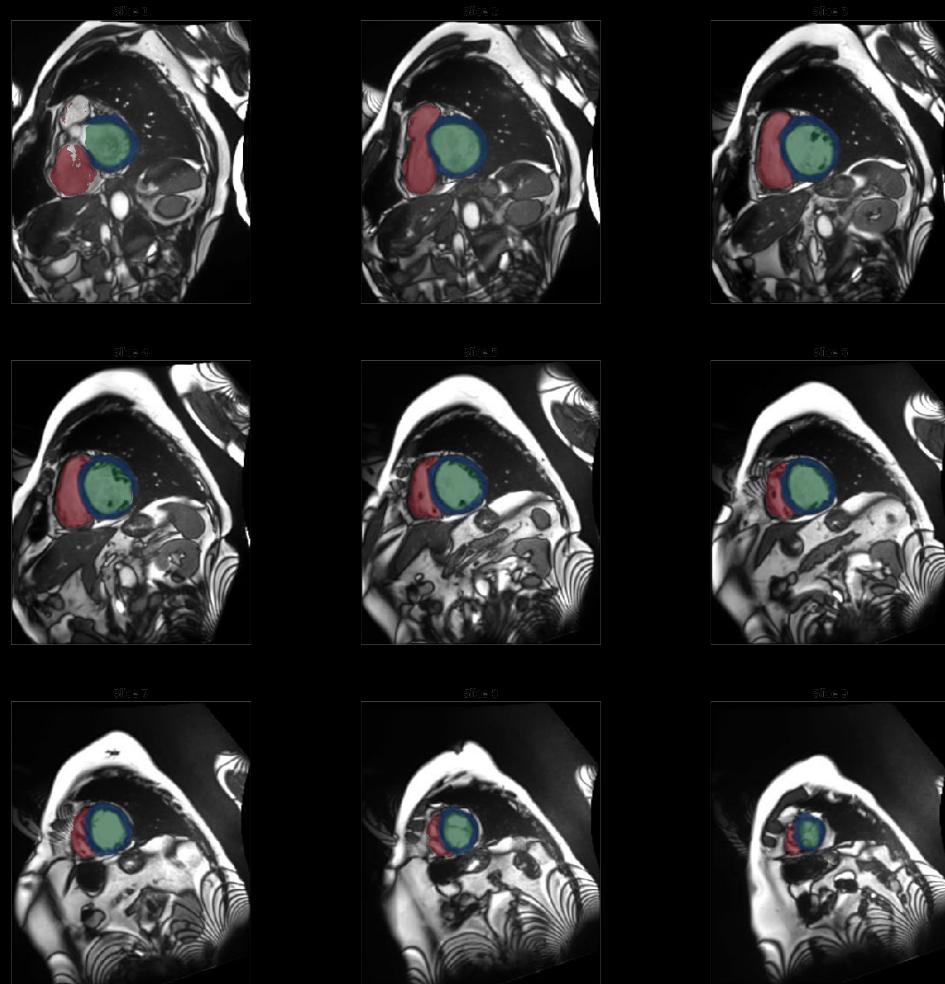
Compare automatic and reference segmentations

- Dice similarity coefficient

```
monai.metrics.DiceMetric()
```

- Hausdorff distance

```
monai.metrics.HausdorffDistanceMetric()
```



Average DSC 0.92, average Hausdorff distance 5.97 mm

CODE

Example code on



<https://github.com/jelmerwolterink/ISMRMTutorial>

- Loading and viewing images
- Training a U-Net in Python + PyTorch + MONAI
- Evaluating and visualizing your results

¹ <https://pytorch.org/>

² <https://pytorch.org/tutorials/beginner/basics/intro.html>

ALTERNATIVELY



- Google Colaboratory¹ provides Python notebooks in the cloud
- Free GPU utilization²
- Load notebook from GitHub



The screenshot shows the Google Colaboratory interface. At the top, there's a navigation bar with tabs for Examples, Recent, Google Drive, GitHub (which is selected), and Upload. Below the navigation bar, there's a search bar with the placeholder "Enter a GitHub URL or search by organization or user". A checkbox labeled "Include private repos" is available. The main area displays a GitHub URL: "https://github.com/jelmerwolterink/ISMRMTutorial". Underneath the URL, there are dropdown menus for "Repository:" set to "jelmerwolterink/ISMRMTutorial" and "Branch:" set to "main". A "Path" input field is below these. A list of notebooks is shown, with the first one being "CardiacMRSegmentation_ISMRM.ipynb". At the bottom right, there are "NEW NOTEBOOK" and "CANCEL" buttons.

¹ <https://colab.research.google.com/>

² <https://colab.research.google.com/notebooks/gpu.ipynb>

WHAT CAN YOU DO?

- Training set augmentation (using PyTorch¹, MONAI², TorchIO³)
- Modify U-Net architecture (3D vs. 2D)
- Other networks⁴, e.g. DynUNet, VNet
- Other optimizers⁵, e.g. SGD, AdaGrad, ..
- Add data from other sources, e.g. M&Ms
- Segment the full cardiac cycle in ACDC
- ...

¹ <https://pytorch.org/vision/stable/transforms.html>

² <https://docs.monai.io/en/latest/transforms.html>

³ <https://torchio.readthedocs.io/transforms/augmentation.html>

⁴ <https://docs.monai.io/en/latest/networks.html>

⁵ <https://pytorch.org/docs/stable/optim.html>

SUMMARY

- ACDC data set
- Python + PyTorch + MONAI
- Minimal code provided
- Q&A on Thursday, post questions as issues on GitHub

Code: <https://github.com/jelmerwolterink/ISMRMTutorial>

Thank you for your attention!