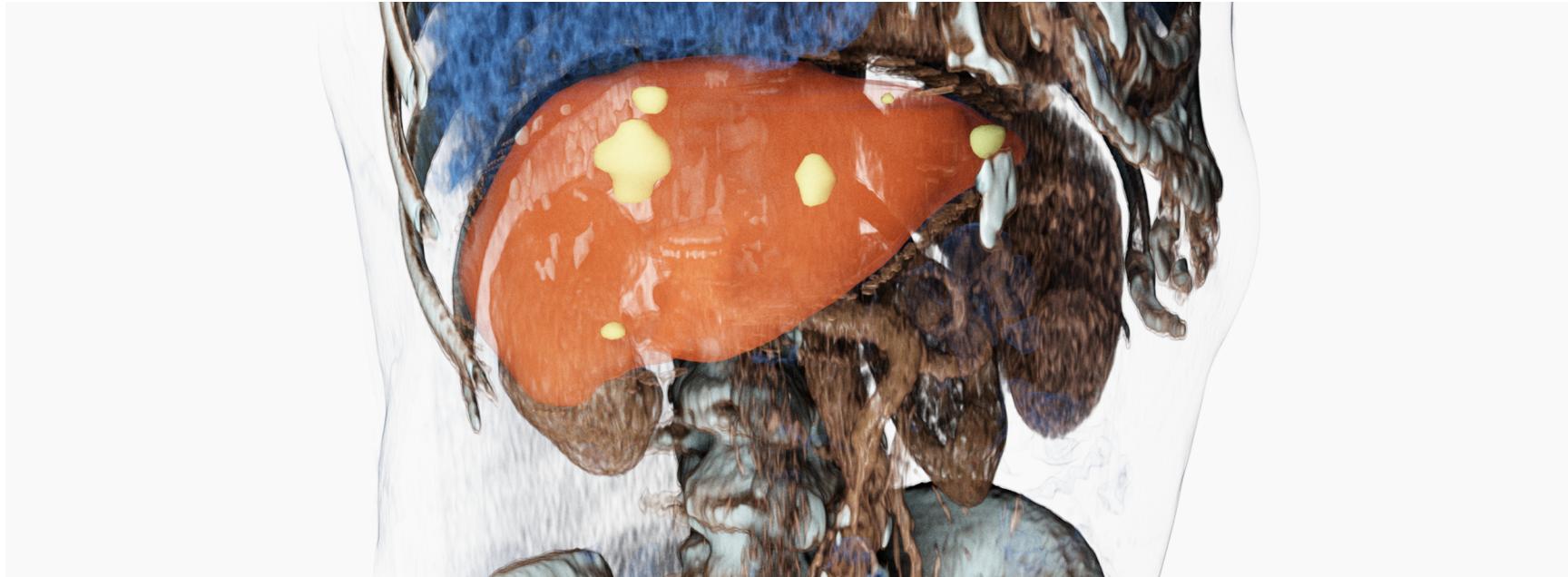
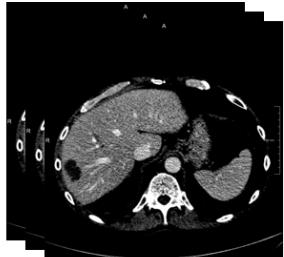


NEURAL NETWORK BASED AUTOMATIC LIVER AND LIVER TUMOR SEGMENTATION

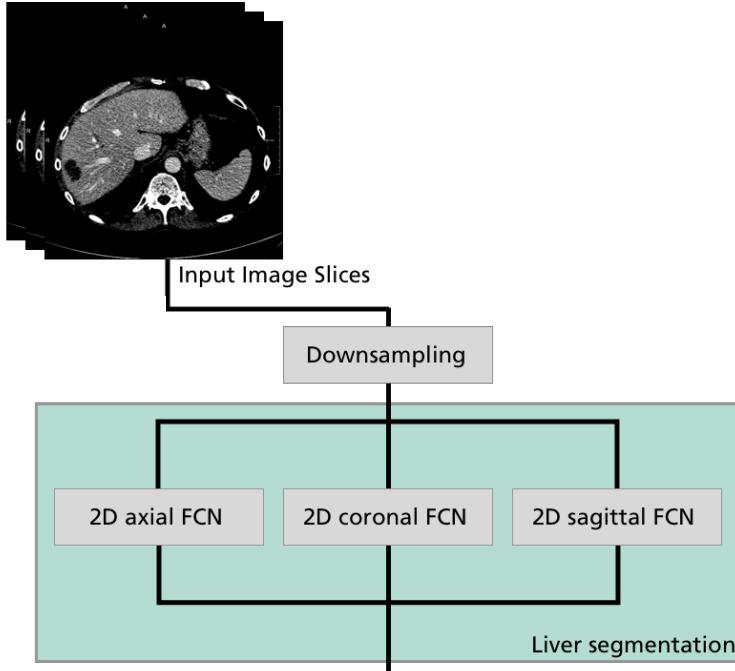
Grzegorz Chlebus, Hans Meine, Jan Hendrik Moltz, Andrea Schenk

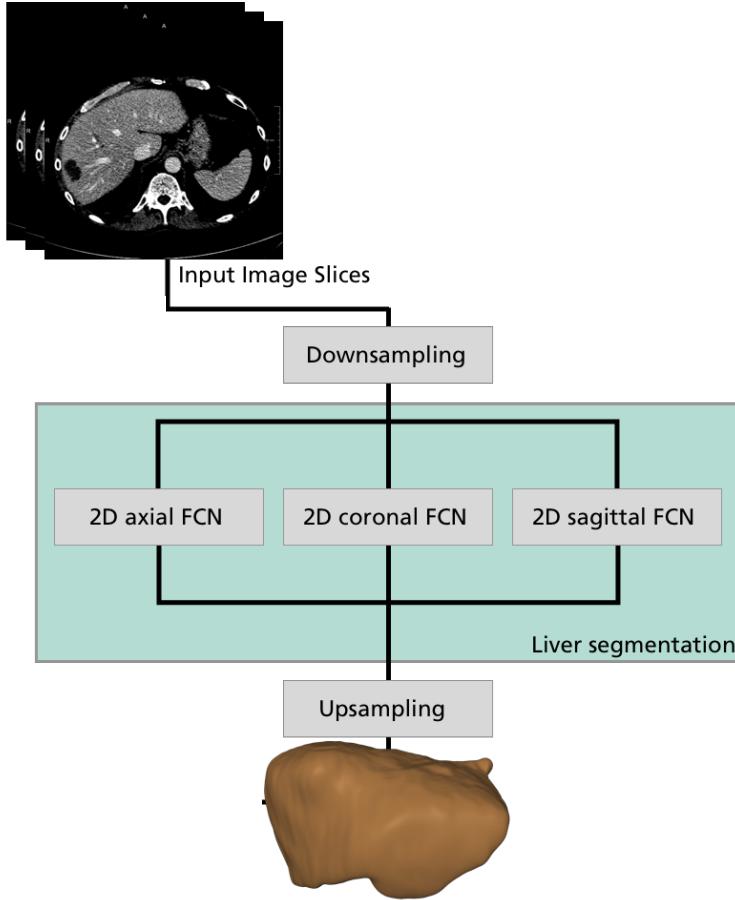


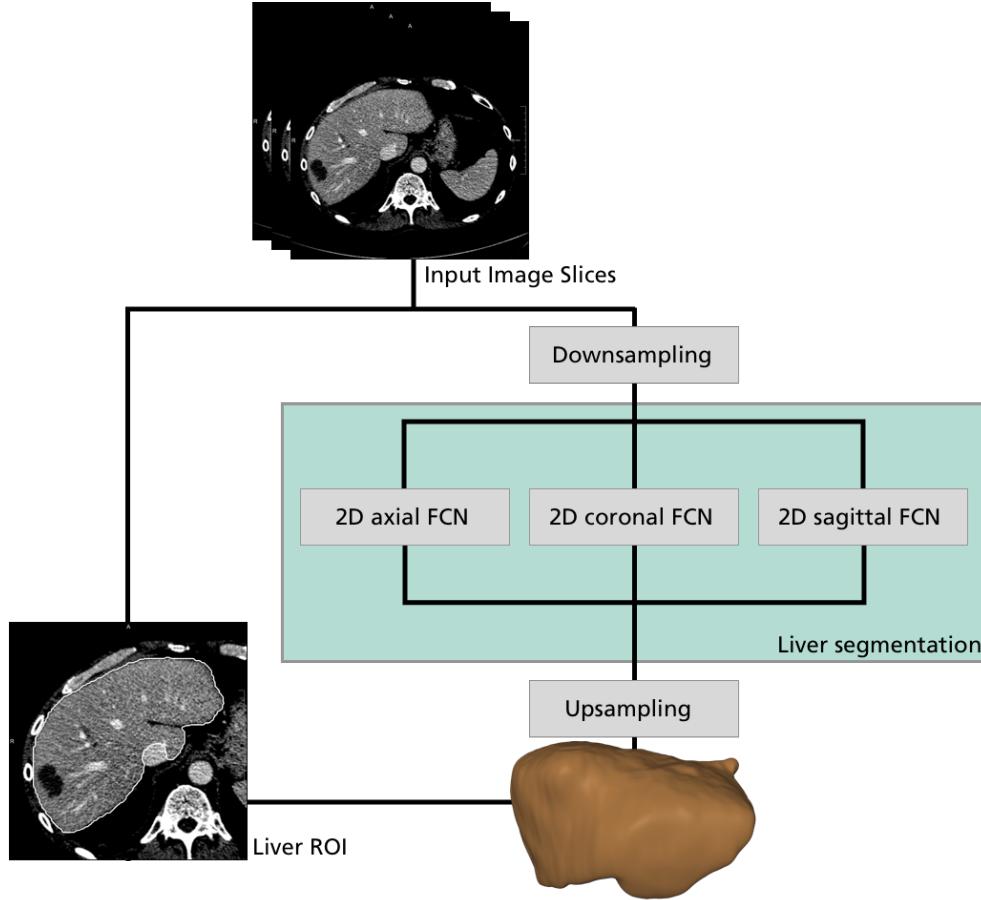
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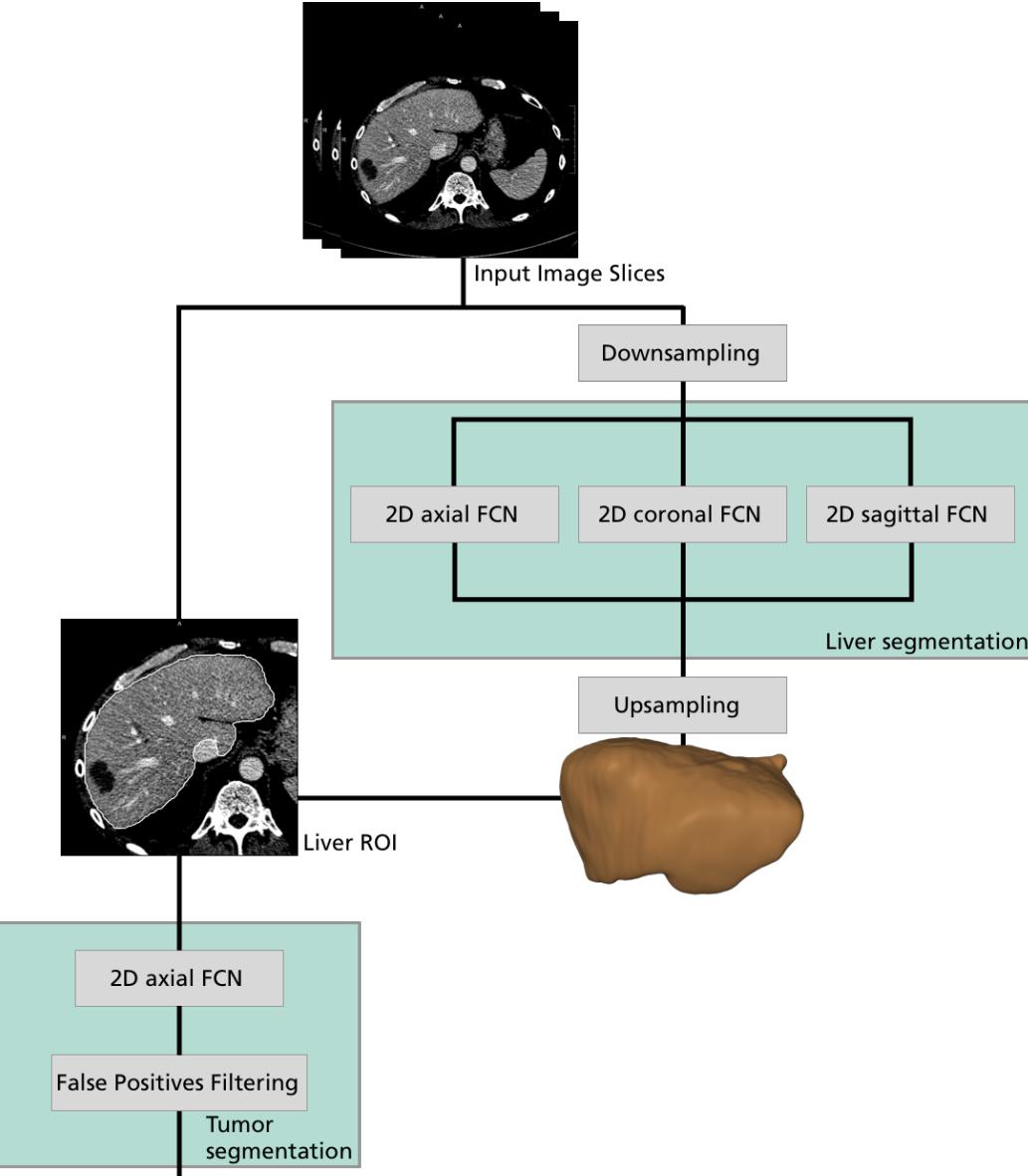


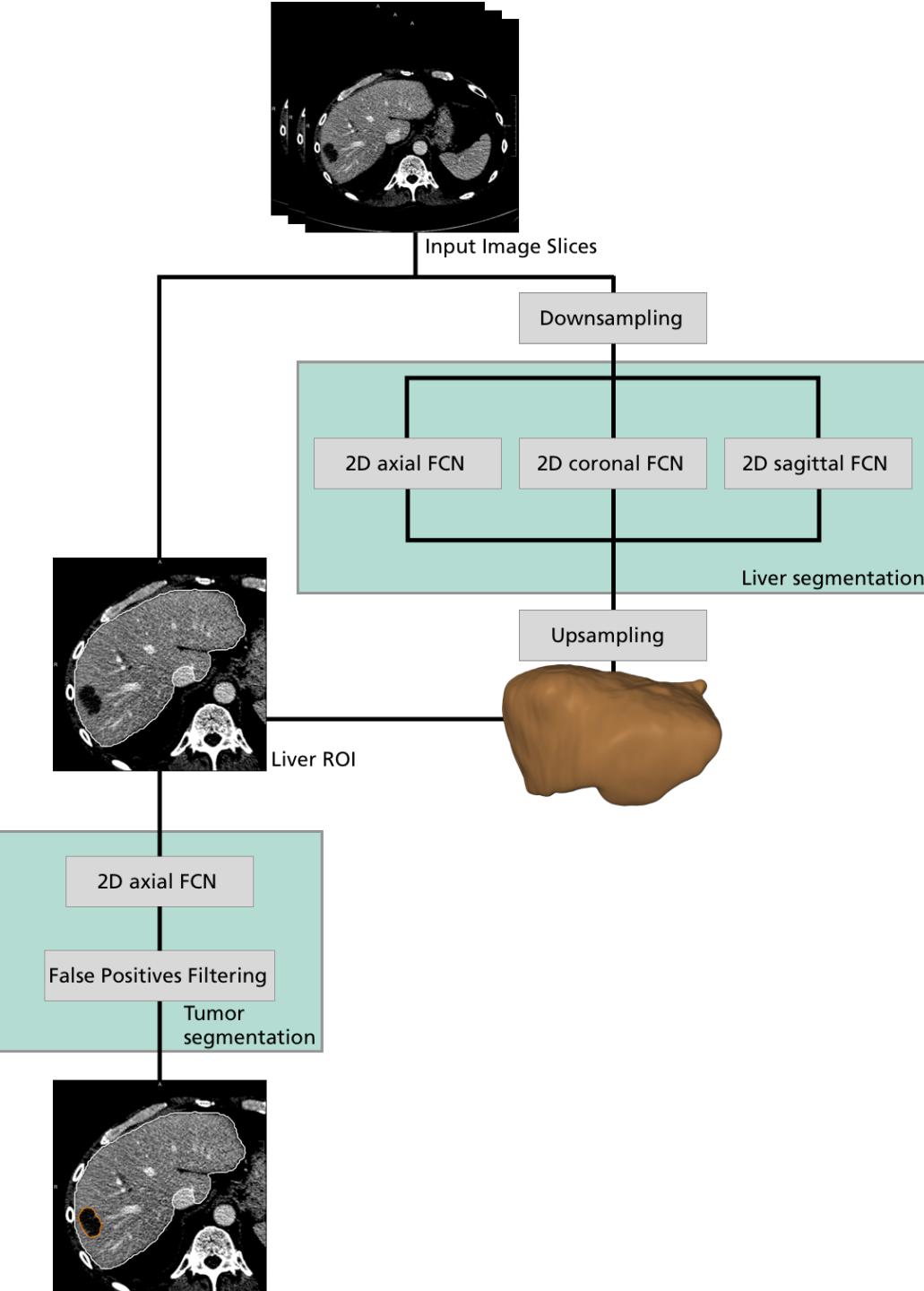
Input Image Slices











Datasets

- LiTS
 - 131 CTs:
 - 105 - training
 - 15 - validation
 - 11 - testing
 - ~0.8 mm in-plane resolution
 - ~1.5 mm slice thickness
- Liver surgery planning
 - 179 CTs all used for training
 - ~0.6 mm in plane-resolution
 - ~0.8 mm slice thickness
 - Livers segmented by radiological experts



Liver Segmentation Data and Preprocessing

- Two training datasets
 - LiTS
 - Liver surgery planning
- Preprocessing
 - Rescaling raw GV to HU
 - Resampling to 2 mm isotropic voxel size
 - Padding with -1000 HU

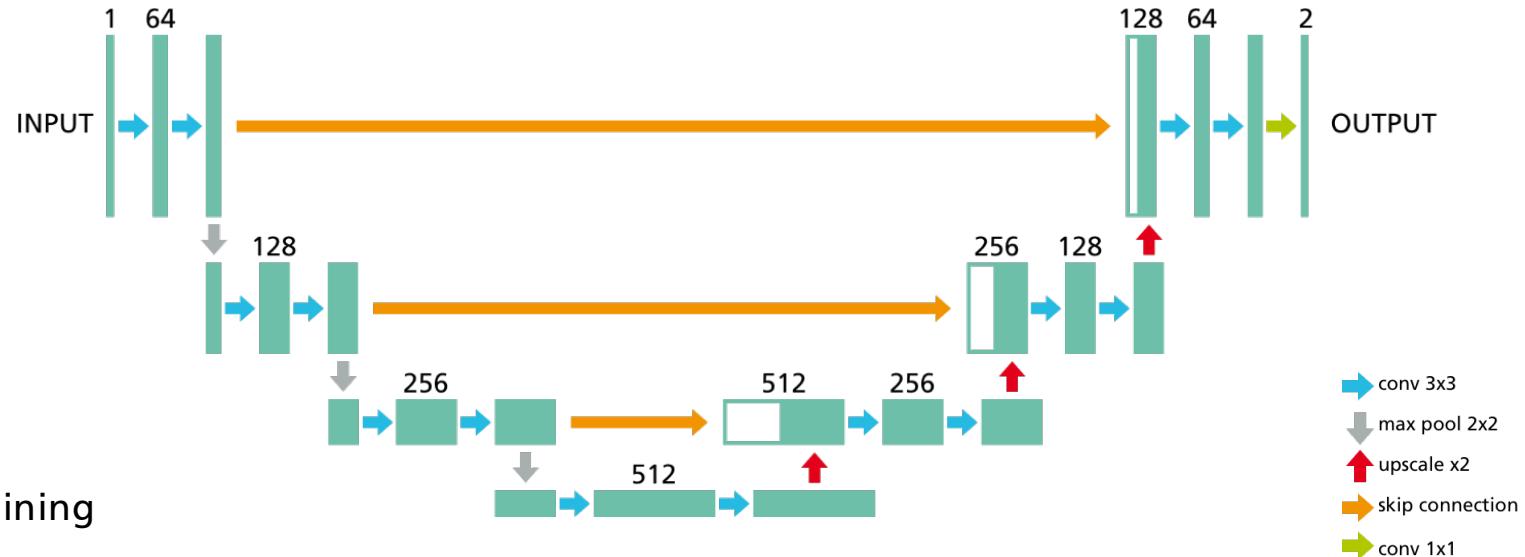
FCN General Info

- Convolution block
 - Dropout p=0.5 in the upscaling path
 - ReLU activation function
 - Batch normalization
- Softmax as the final layer
- Training
 - Dice loss function
 - Adam optimizer

Liver Segmentation

Network Architecture and Training

- 2D U-net [1] with 4 resolution levels



- Training
 - Patch size: 148x148 (axial) or 148x44
 - Batch size 15
 - 10^{-5} learning rate
 - ~30k iterations / ~43 epochs / ~19 h

[1] Ronneberger O. et al., "U-Net: Convolutional Networks for Biomedical Image Segmentation", 2015.

Medical Knowledge Through Research

Liver Segmentation Postprocessing

- Biggest connected components of the majority vote mask

Liver Segmentation Postprocessing

- Biggest connected components of the majority vote mask



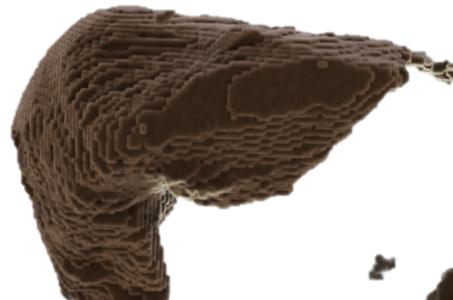
transversal

Liver Segmentation Postprocessing

- Biggest connected components of the majority vote mask



transversal



coronal

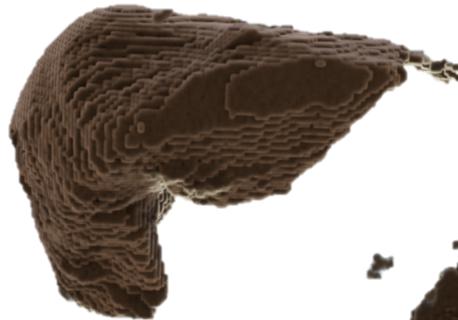


Liver Segmentation Postprocessing

- Biggest connected components of the majority vote mask



transversal



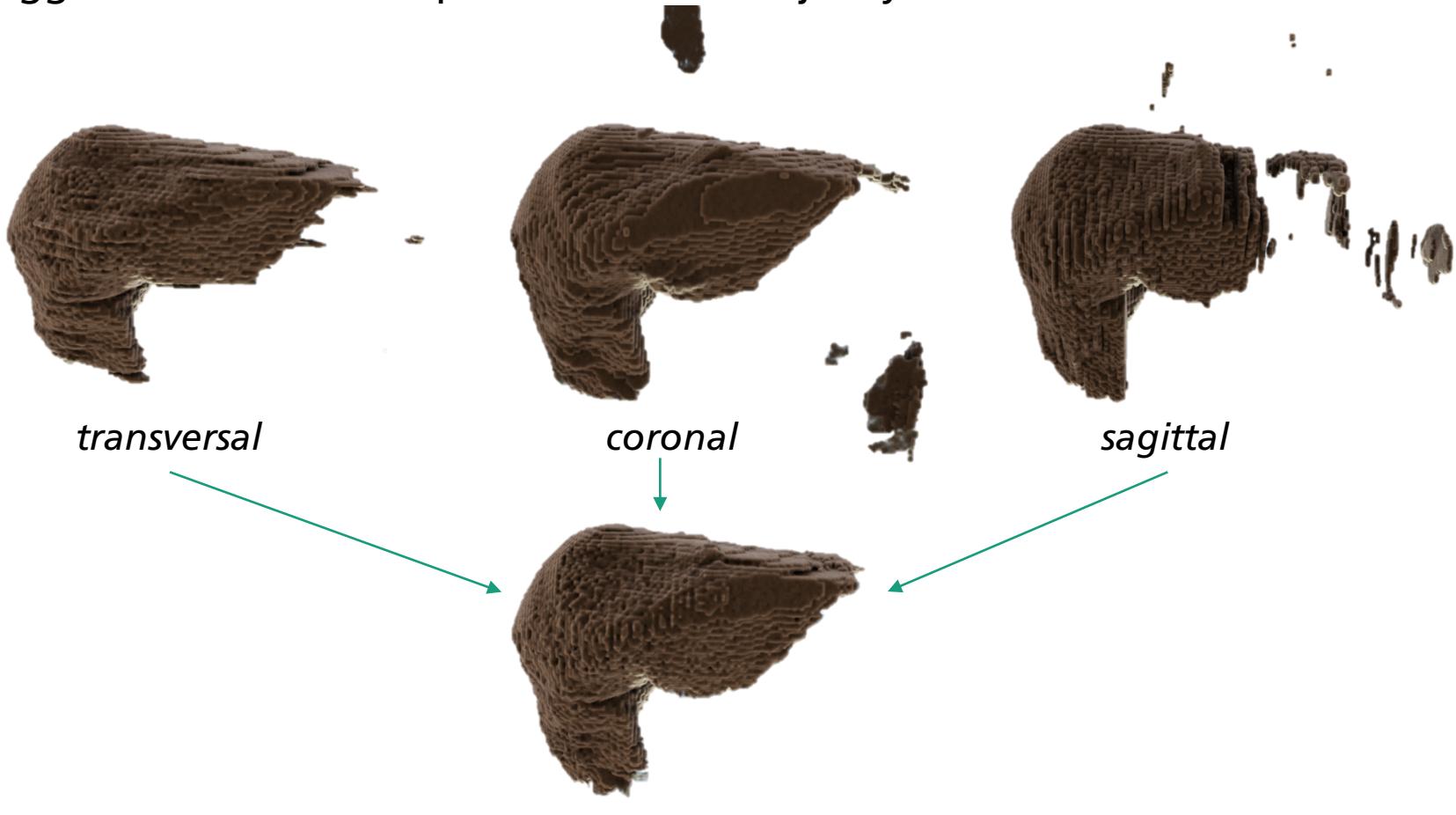
coronal



sagittal

Liver Segmentation Postprocessing

- Biggest connected components of the majority vote mask



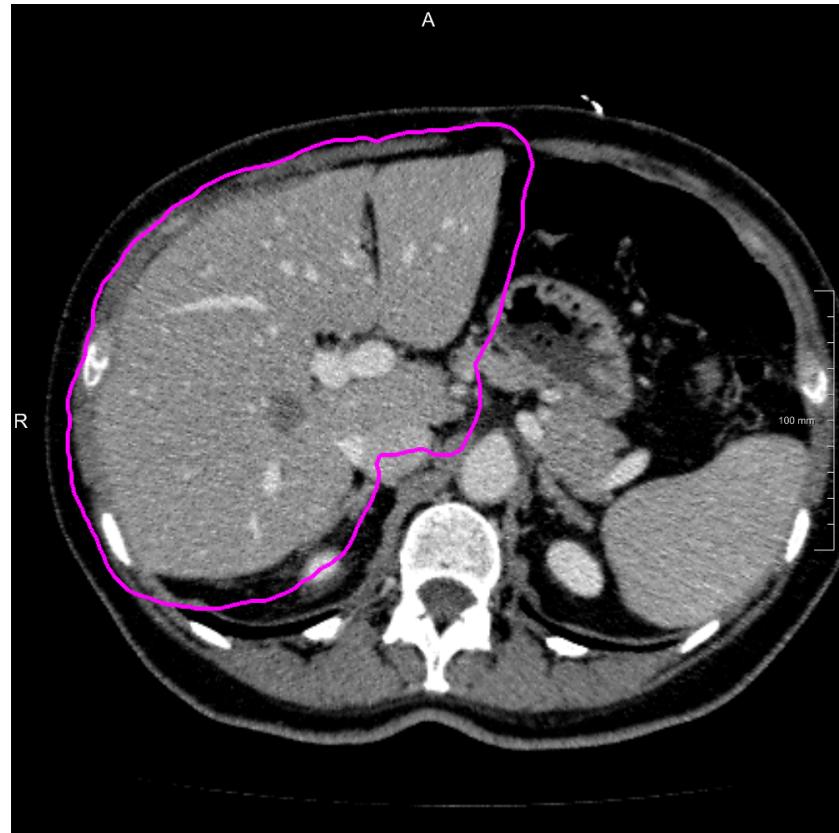
Tumor Segmentation Data and Preprocessing

- LiTS dataset
- Preprocessing
 - Padding with -1000 HU



Tumor Segmentation Data and Preprocessing

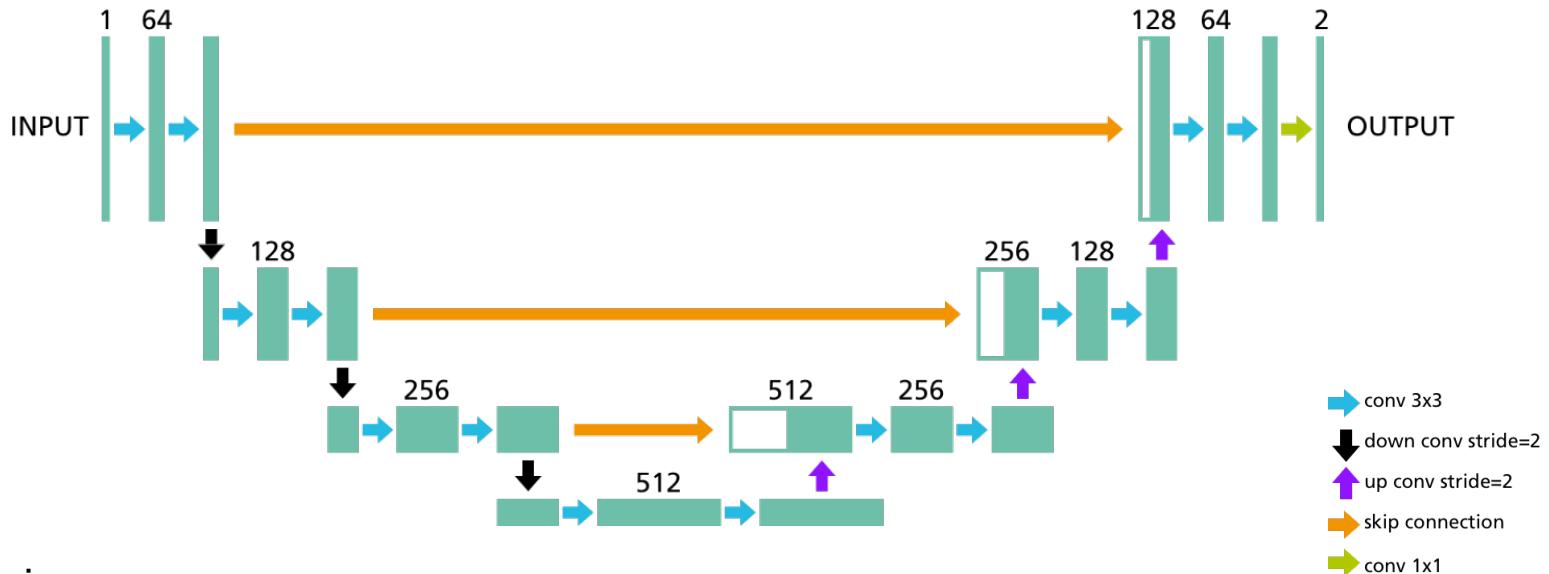
- LiTS dataset
- Preprocessing
 - Padding with -1000 HU
- Masked loss



Tumor Segmentation

Network Architecture and Training

- Modified 2D U-net with 4 resolution levels



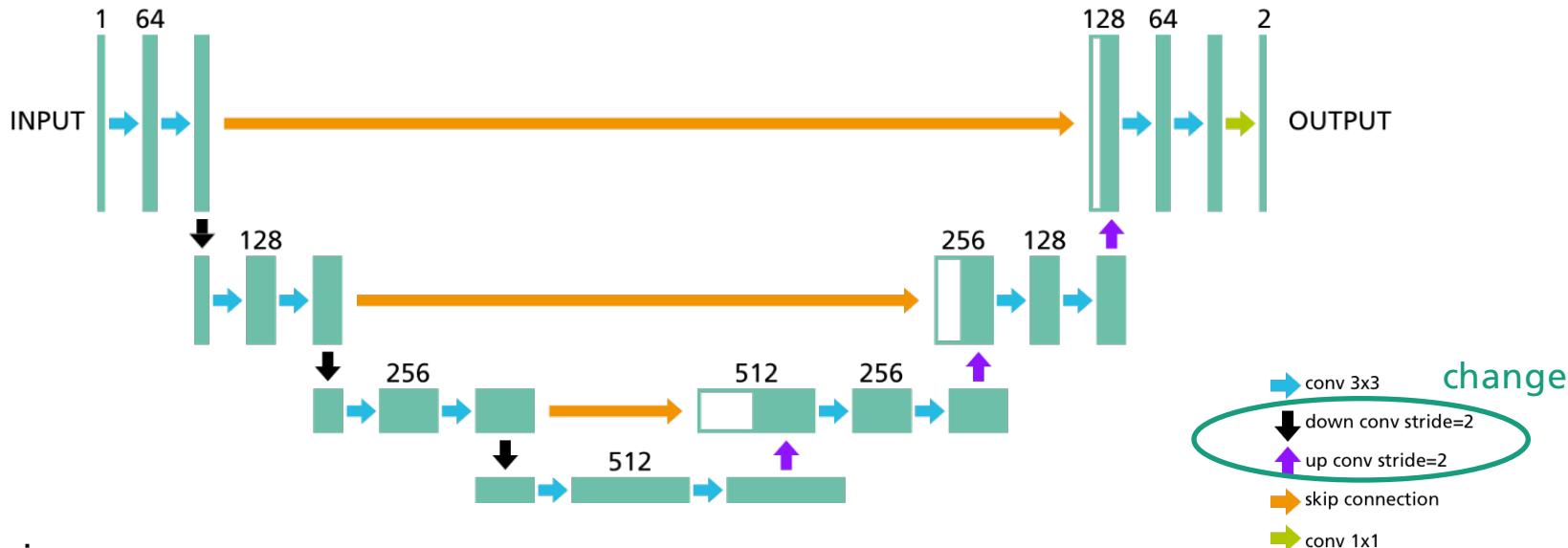
- Training

- Patch size: 252x252 (axial)
- Only tumor patches
- Batch size 6
- ~230k iterations / ~32 epochs / ~38 h
- 5^{-5} learning rate
- Random flipping

Tumor Segmentation

Network Architecture and Training

- Modified 2D U-net with 4 resolution levels



- Training

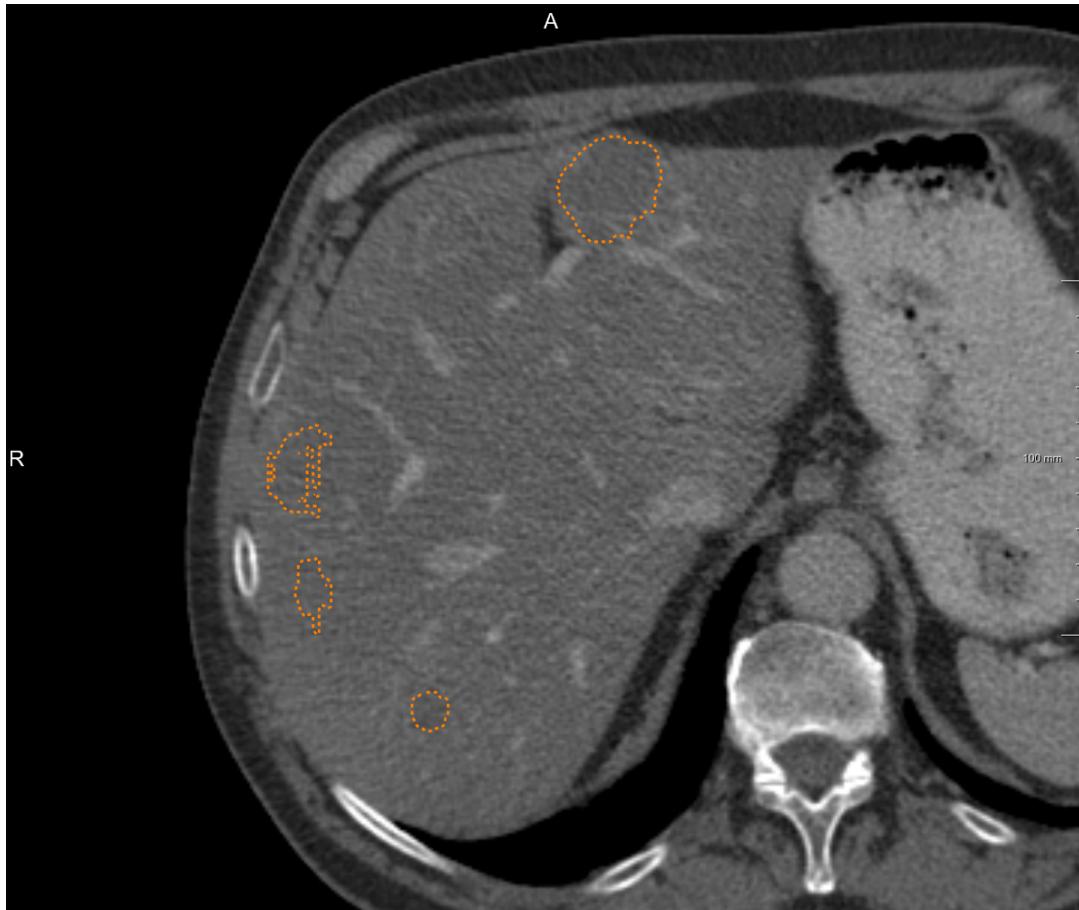
- Patch size: 252x252 (axial)
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- ~230k iterations / ~32 epochs / ~38 h
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Tumor Segmentation Output of the FCN



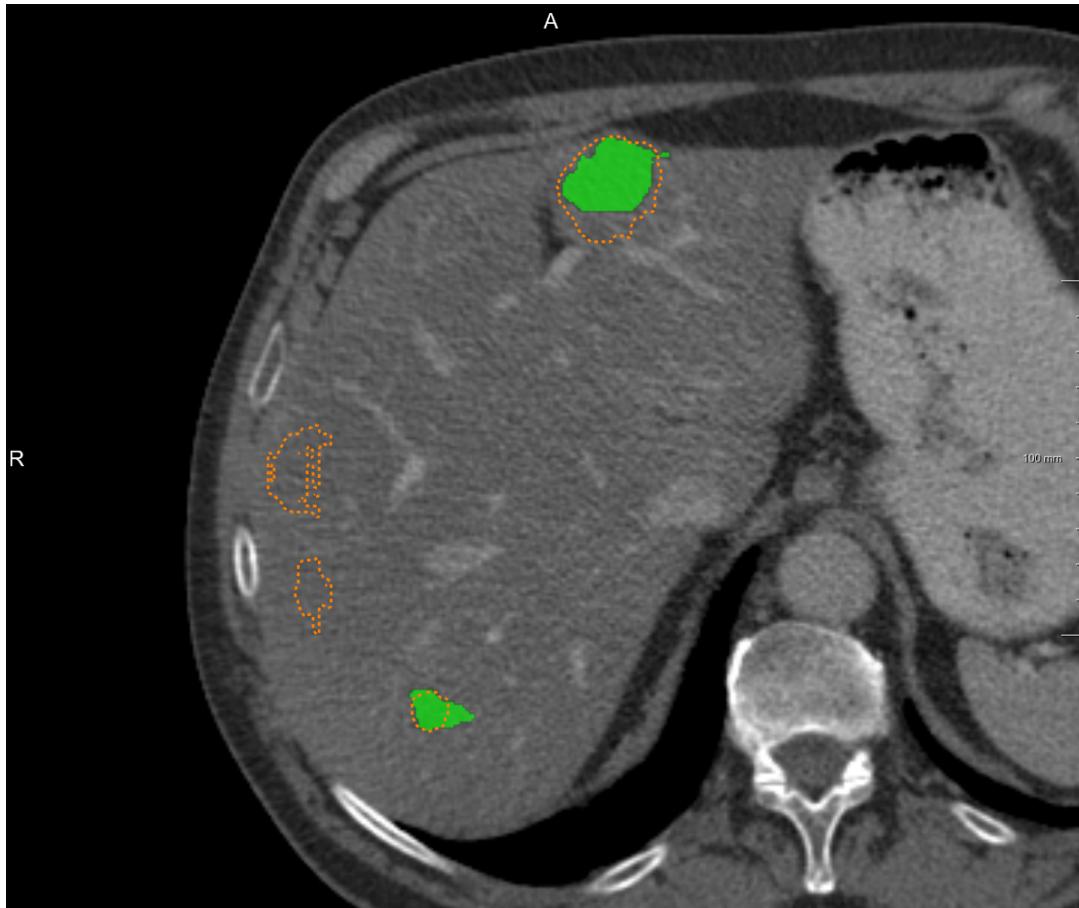
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Tumor Segmentation Output of the FCN



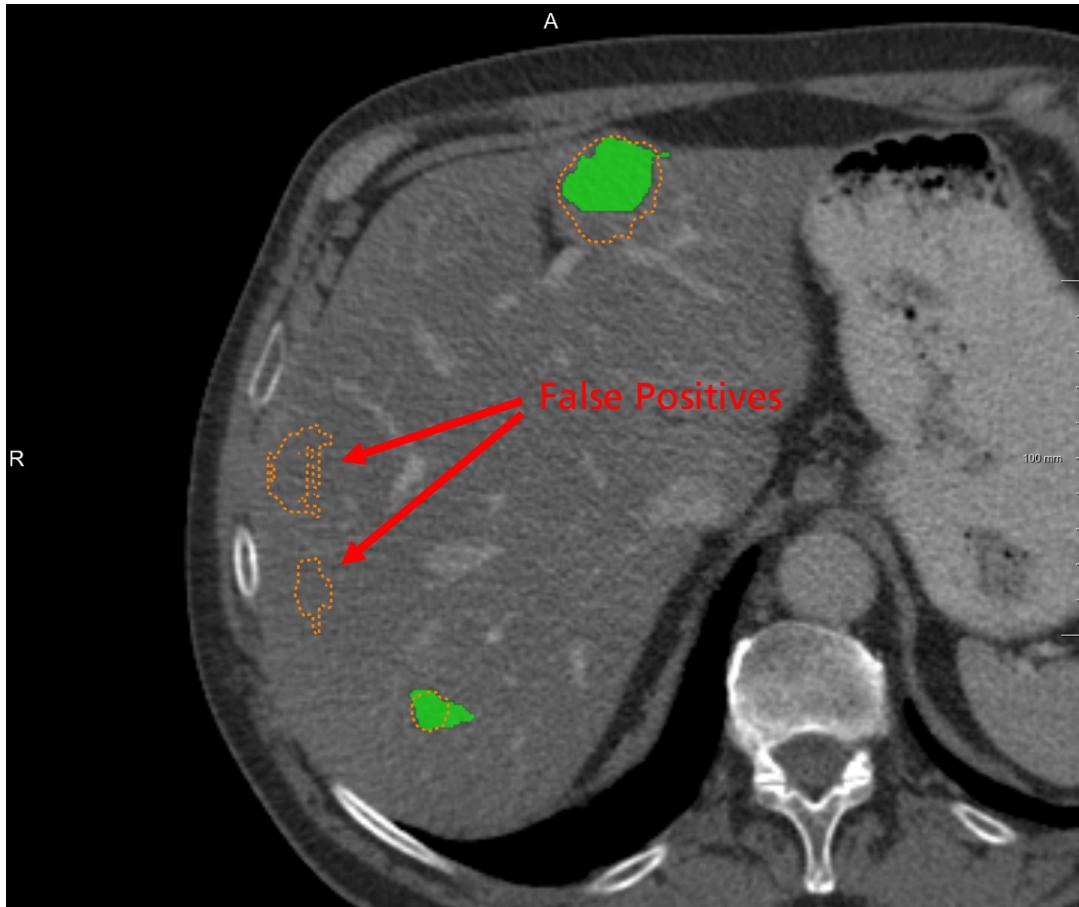
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Tumor Segmentation Output of the FCN



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Tumor Segmentation Output of the FCN



■ False Positives (FPs) Problem

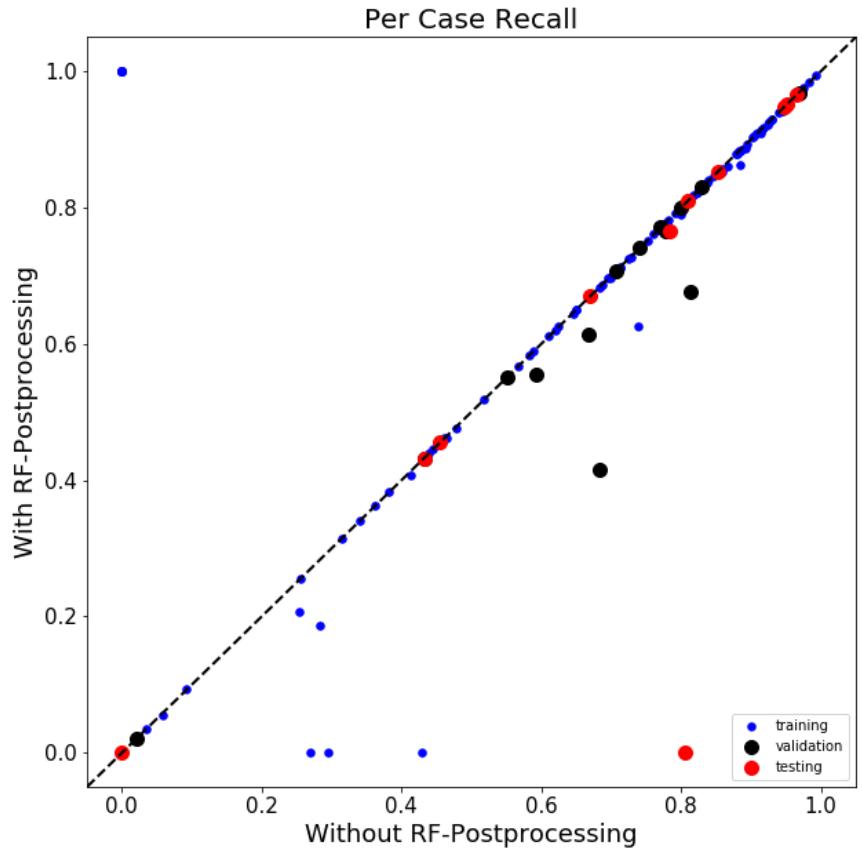
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Tumor Segmentation FPs Filtering

- Train another classifier to detect FPs
- 46 features based on:
 - CT intensity
 - Shape
 - DTF of the liver mask
- Random Forest (RF) accuracy ~90%

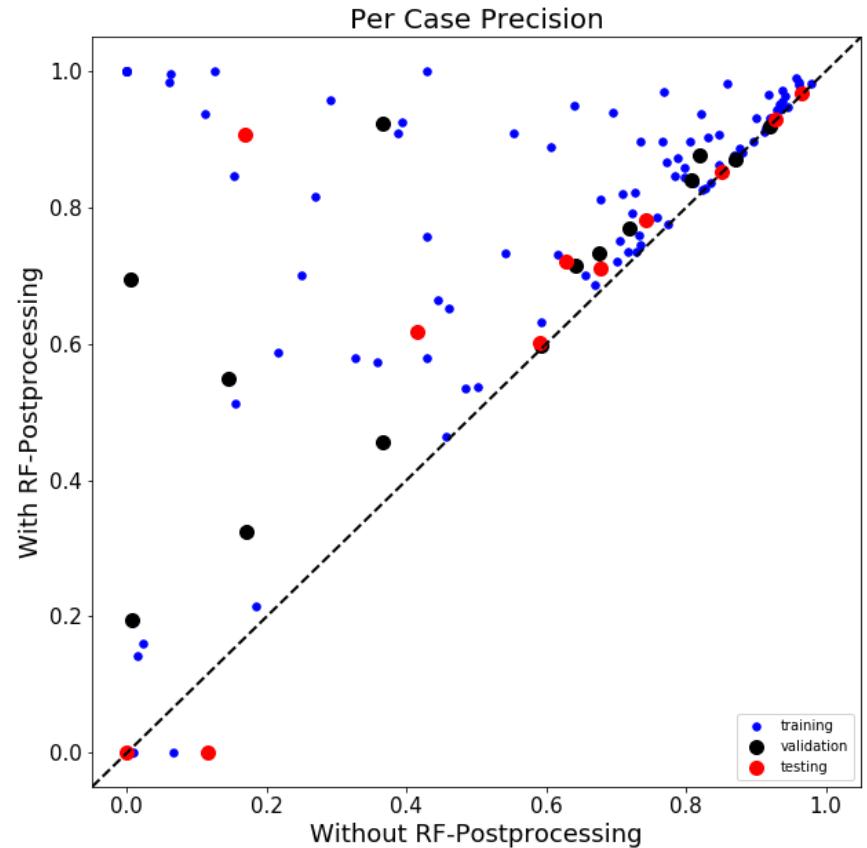
Tumor Segmentation FPs Filtering

- Train another classifier to detect FPs
- 46 features based on:
 - CT intensity
 - Shape
 - DTF of the liver mask
- Random Forest (RF) accuracy ~90%
- RF vs No-RF:
 - Recall



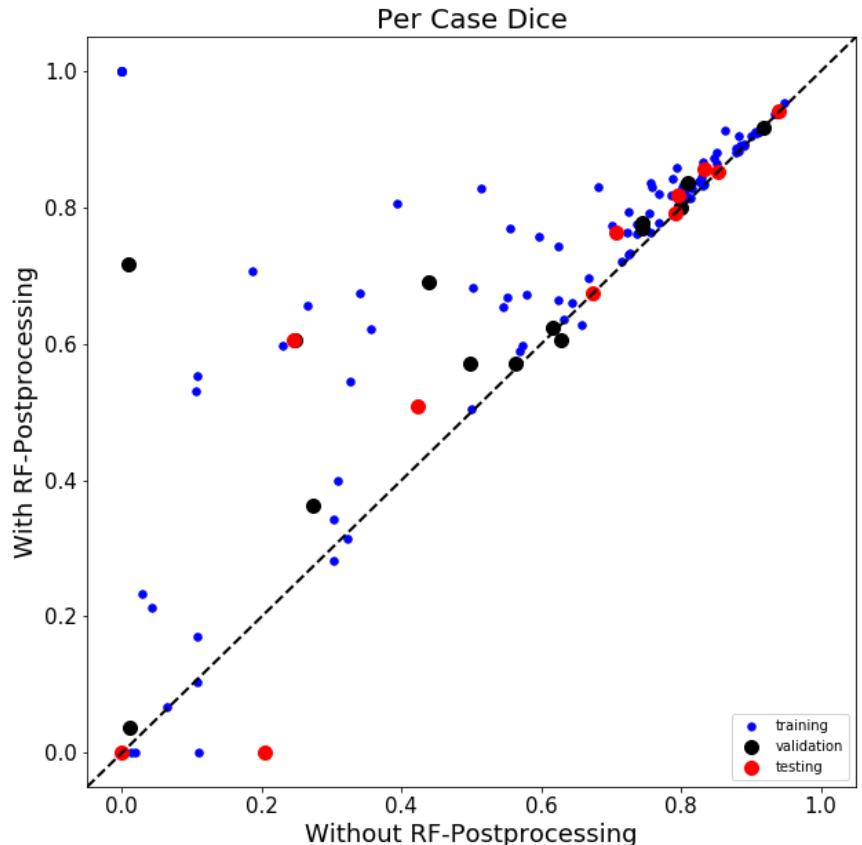
Tumor Segmentation FPs Filtering

- Train another classifier to detect FPs
- 46 features based on:
 - CT intensity
 - Shape
 - DTF of the liver mask
- Random Forest (RF) accuracy ~90%
- RF vs No-RF:
 - Recall
 - Precision

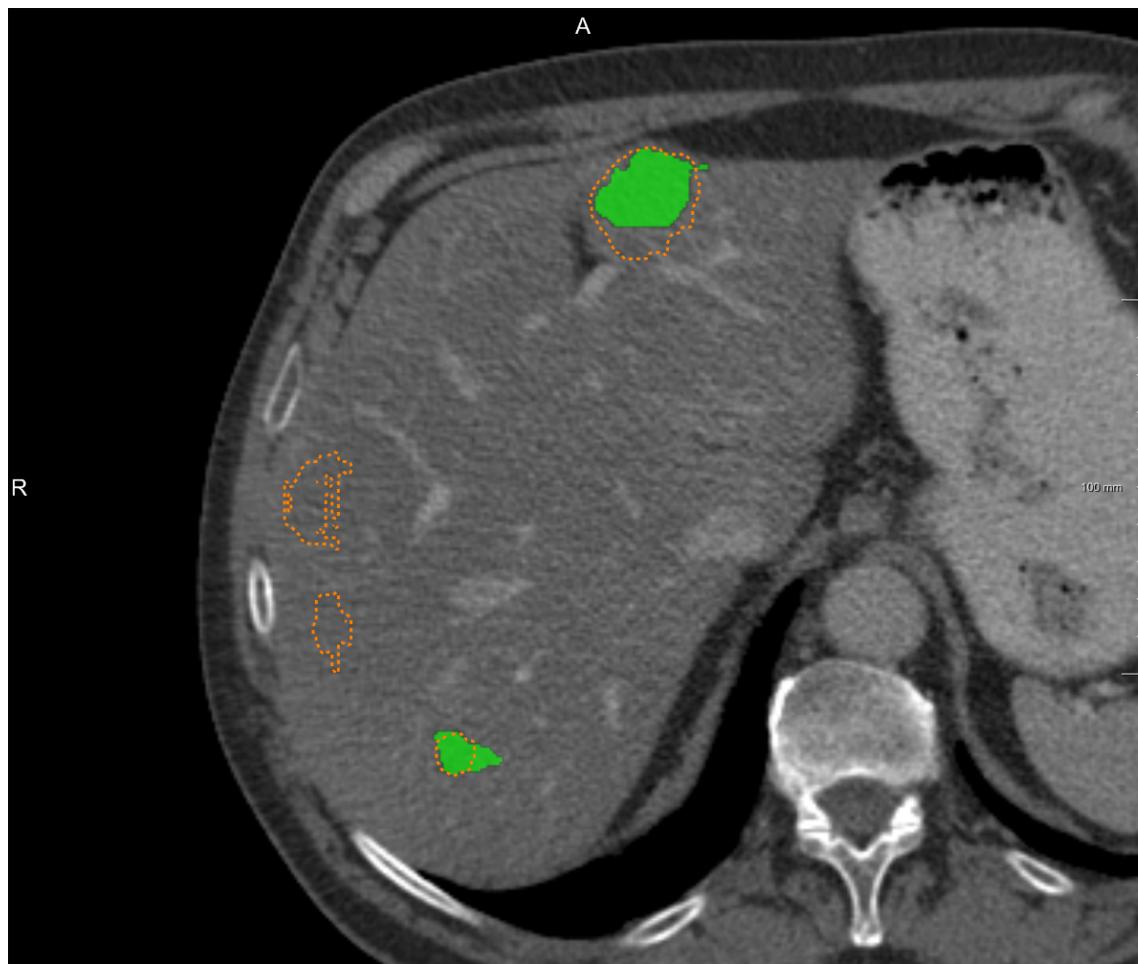


Tumor Segmentation FPs Filtering

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 - Precision 
 - Dice 



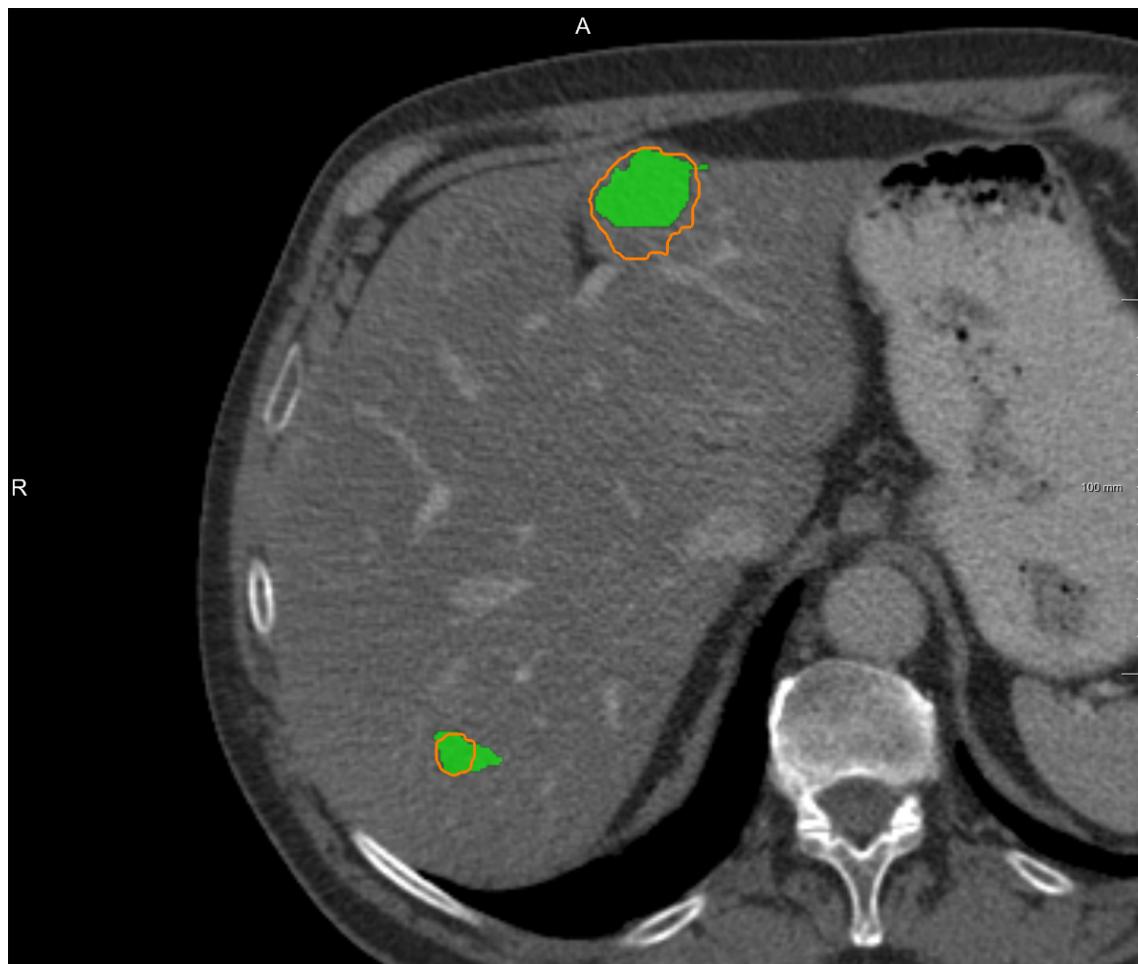
Tumor Segmentation FPs Filtering Examples



Without RF-Postprocessing

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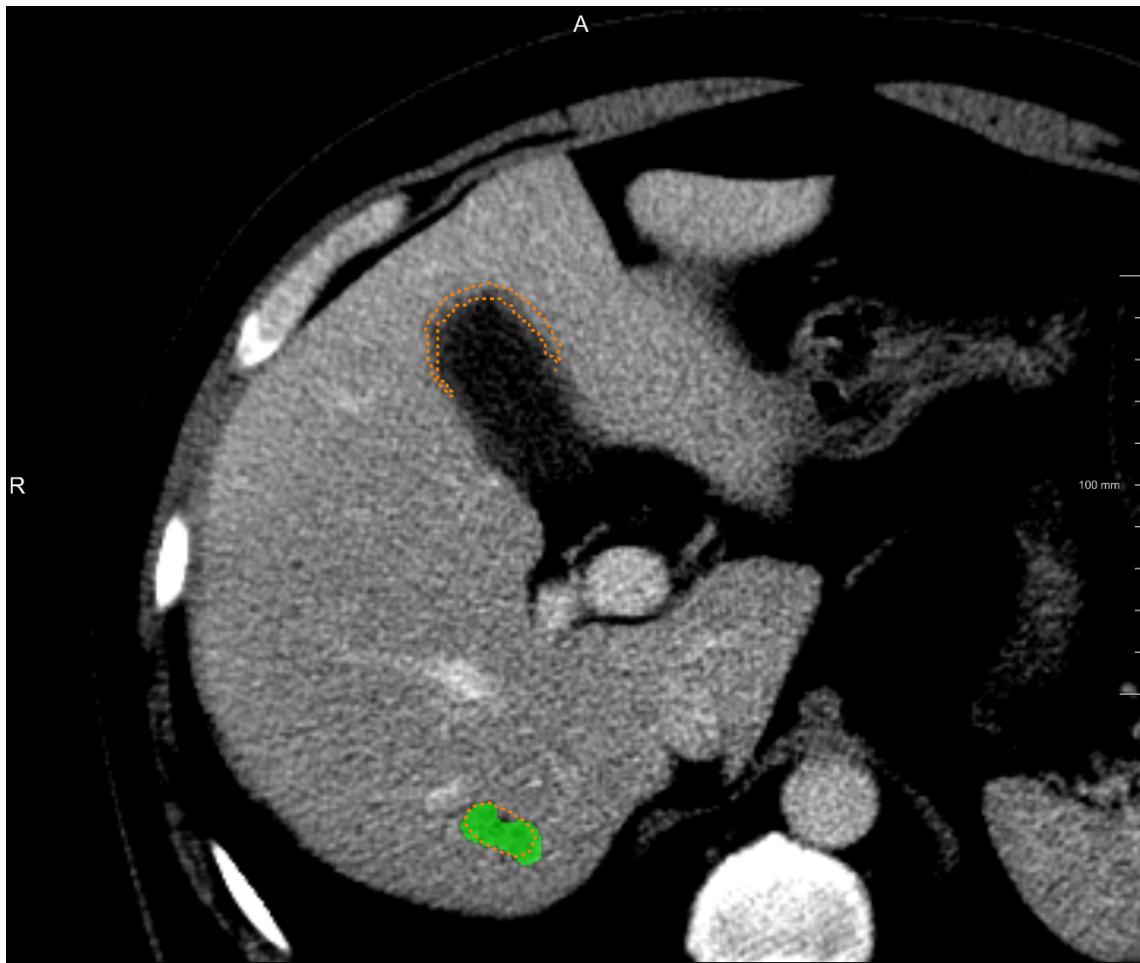
Tumor Segmentation FPs Filtering Examples



With RF-Postprocessing

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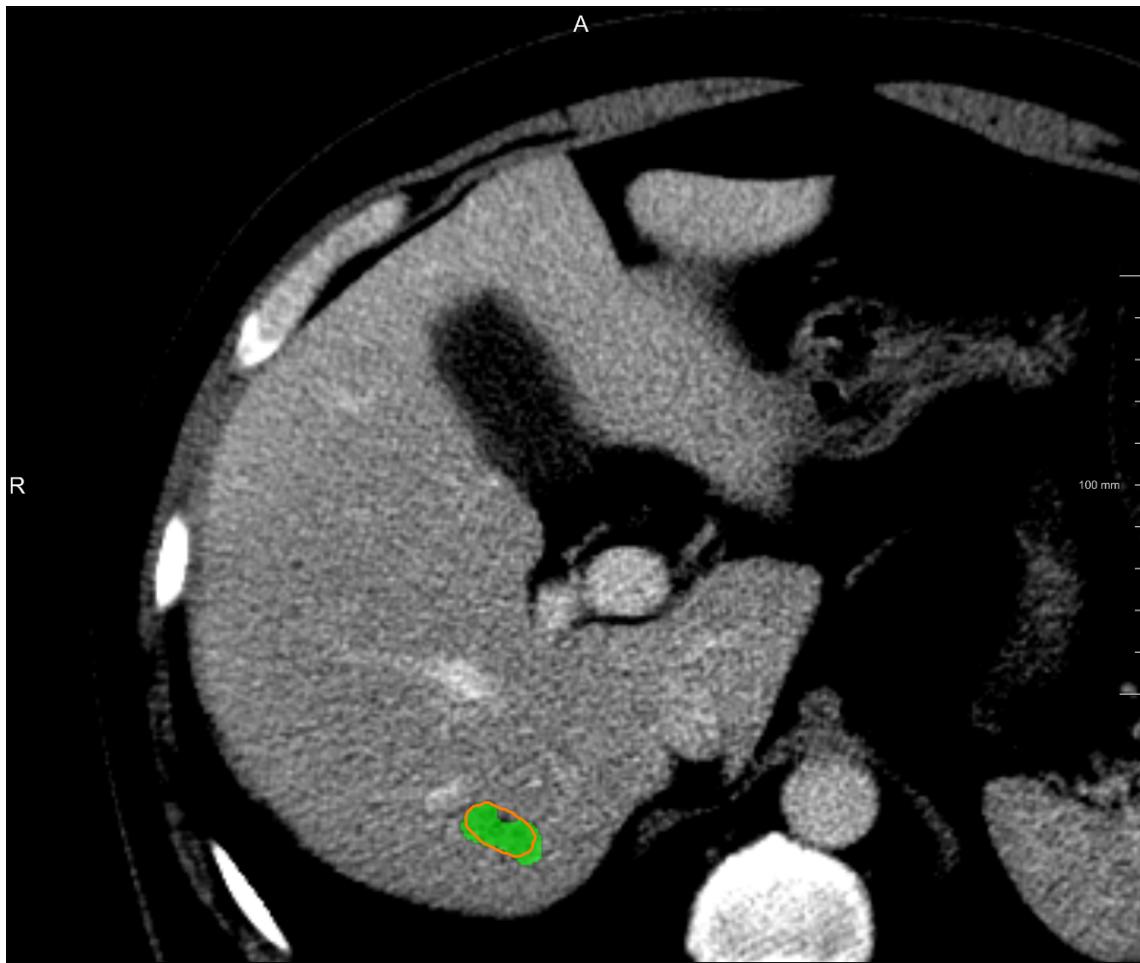
Tumor Segmentation FPs Filtering Examples



Without RF-Postprocessing

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Tumor Segmentation FPs Filtering Examples

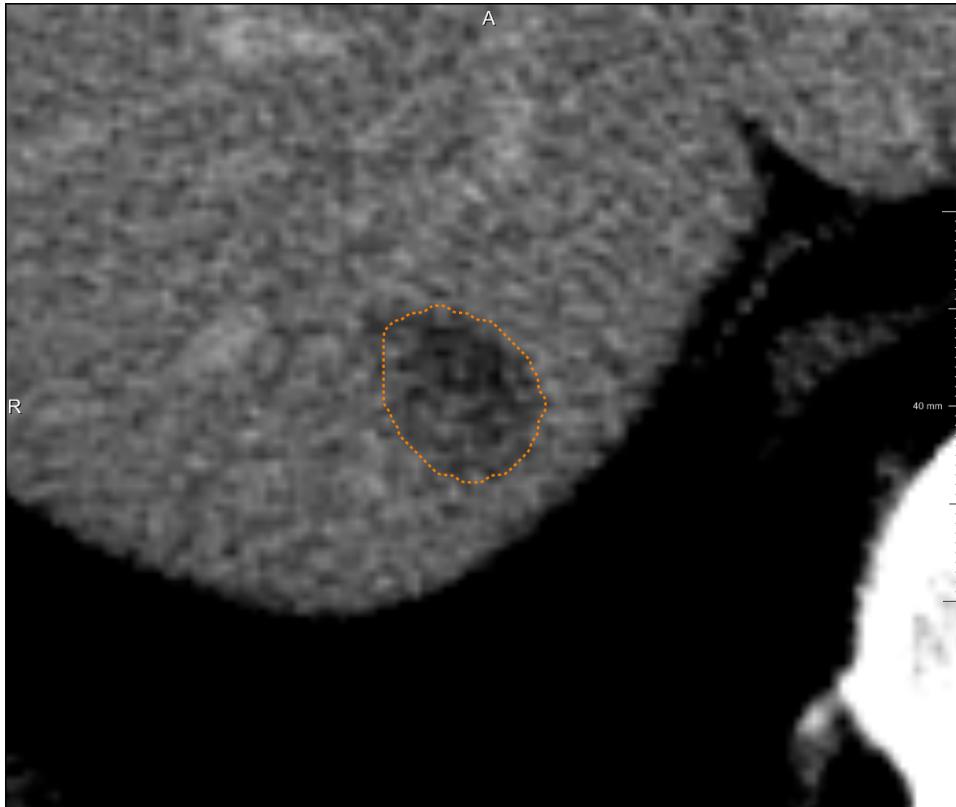


With RF-Postprocessing

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Tumor Segmentation FPs Filtering with Tumor Refinement

- Use tumor candidates to initialize stroke-based semi-automatic segmentation tool [2]

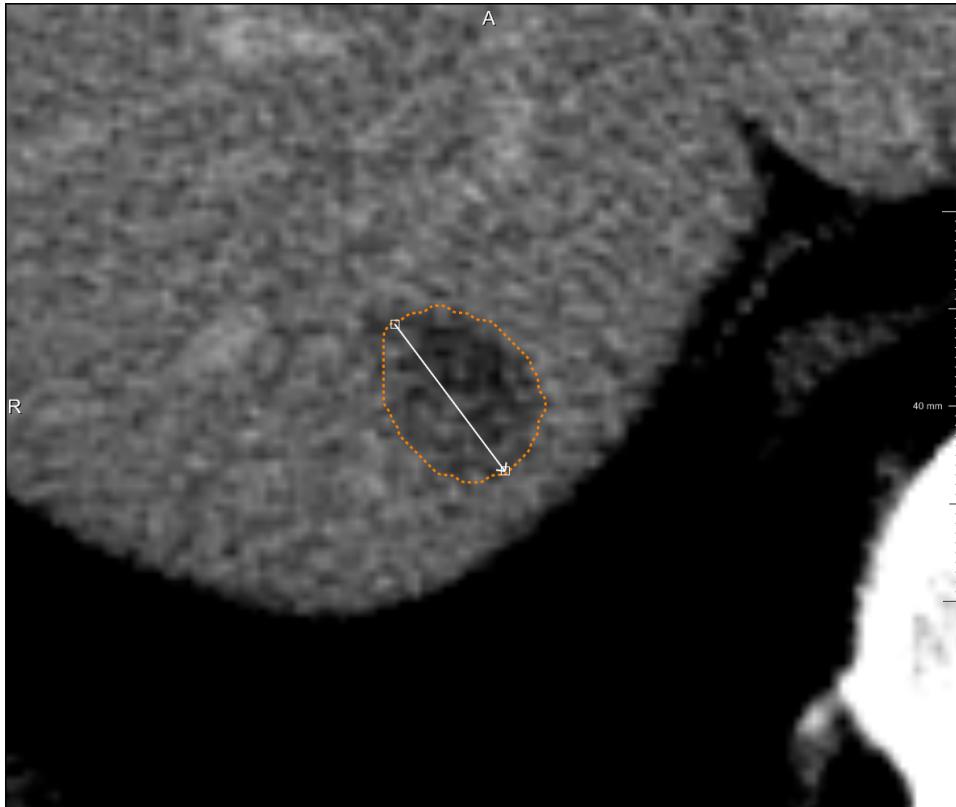


[2] Moltz J.H. et al., "Advanced segmentation techniques for lung nodules, liver metastases, and enlarged lymph nodes in CT scans", 2009.

Medical Knowledge Through Research

Tumor Segmentation FPs Filtering with Tumor Refinement

- Use tumor candidates to initialize stroke-based semi-automatic segmentation tool [2]

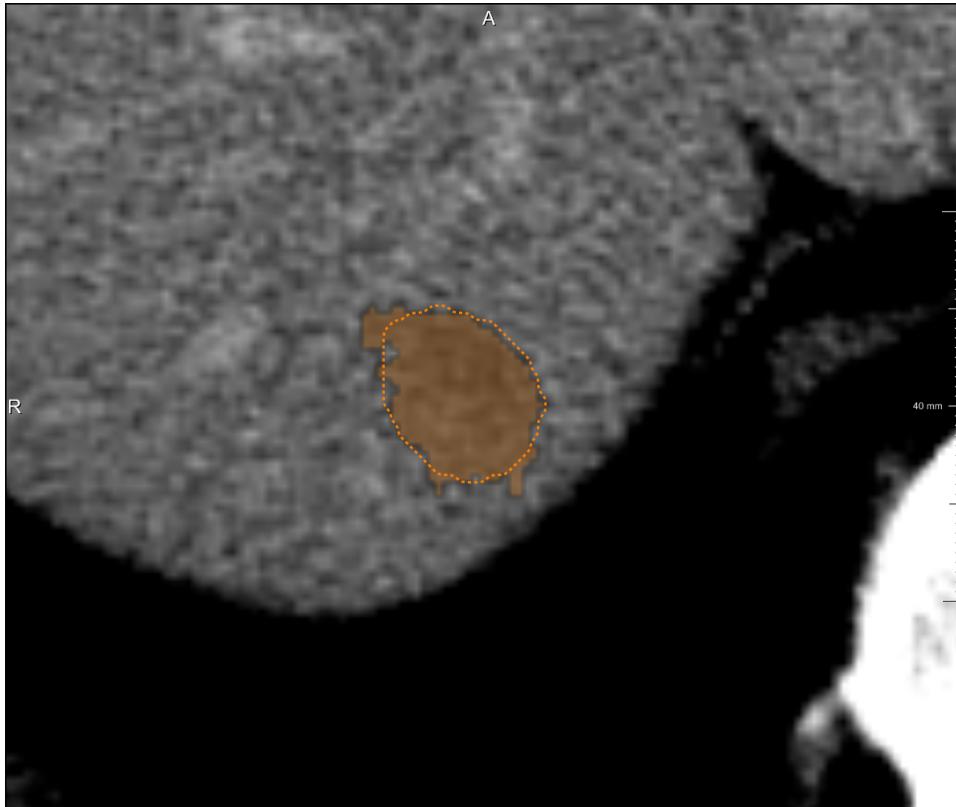


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Tumor Segmentation FPs Filtering with Tumor Refinement

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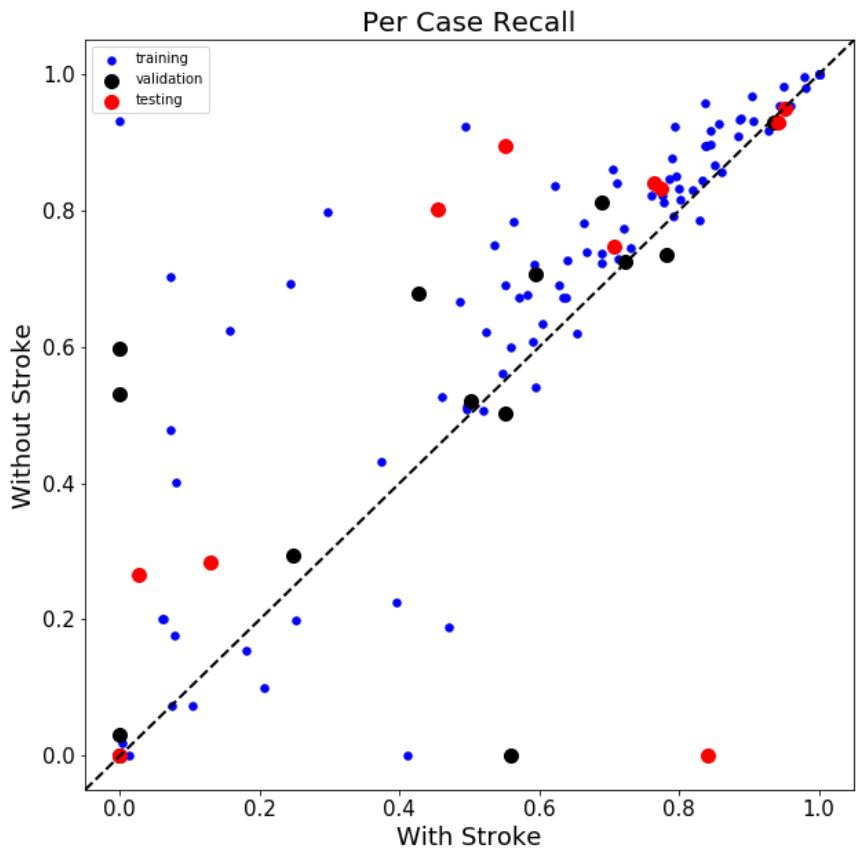
Stroke application

- Segmentation refinement
- Additional features for RF classifier of the refined tumors

Tumor Segmentation FPs Filtering with Tumor Refinement

Stroke application

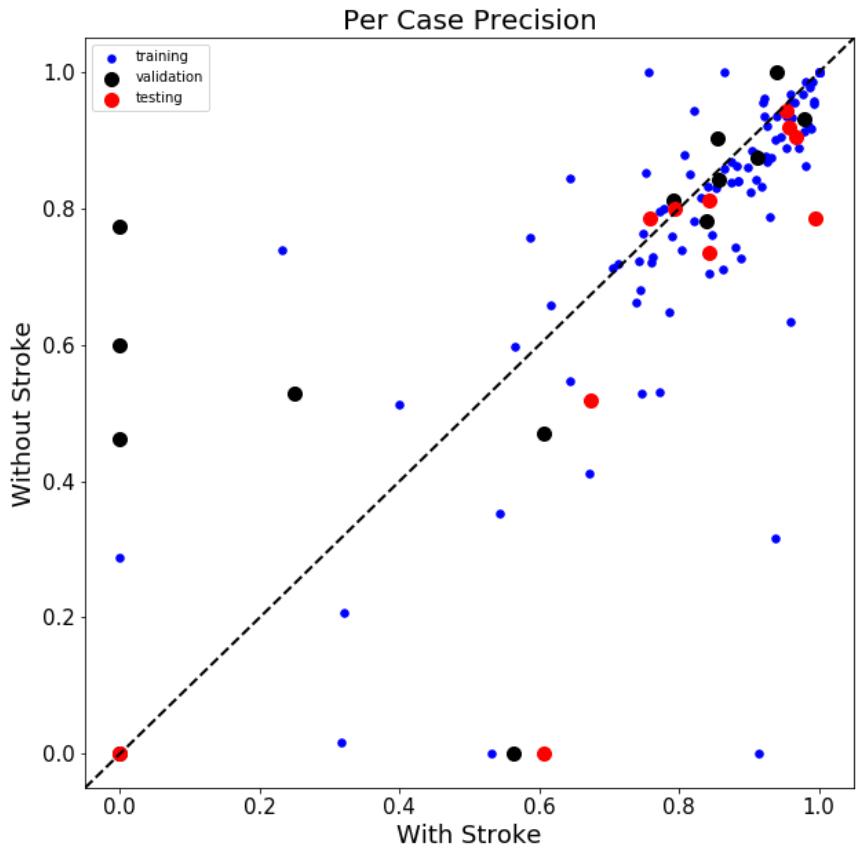
- Segmentation refinement
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- Stroke vs No-Stroke:
 - Recall



Tumor Segmentation FPs Filtering with Tumor Refinement

Stroke application

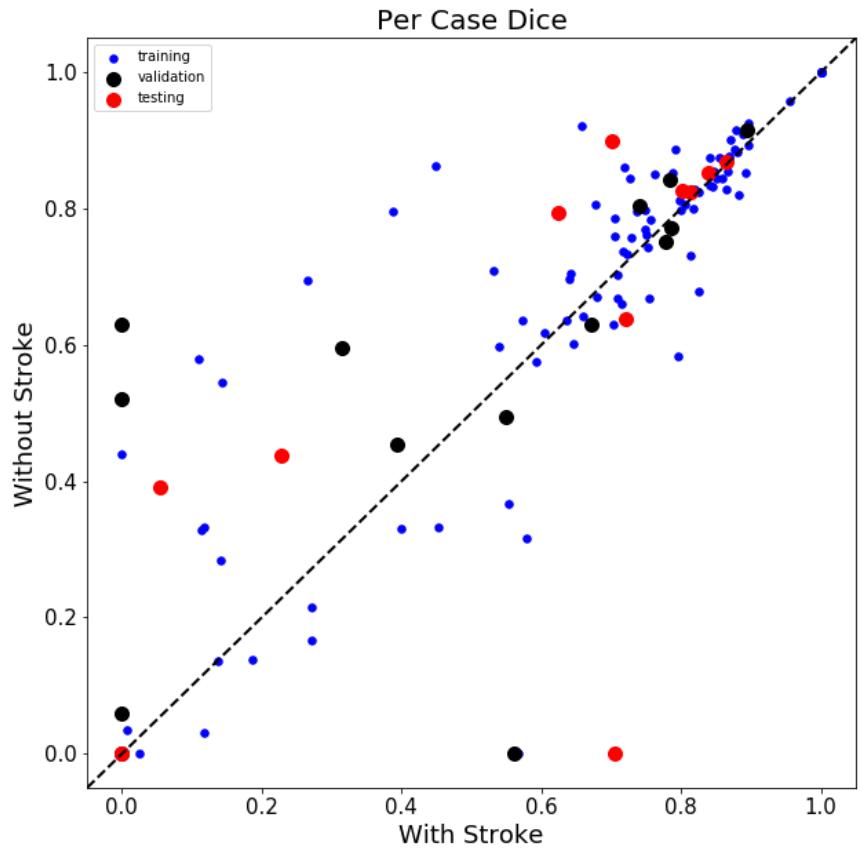
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- Additional features for RF classifier of the refined tumors
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 - Recall 
 - Precision 



Tumor Segmentation FPs Filtering with Tumor Refinement

Stroke application

- Segmentation refinement
- Additional features for RF classifier of the refined tumors
- Stroke vs No-Stroke:
 - Recall 
 - Precision 
 - Dice 



2.5D FCN Architecture

Idea:

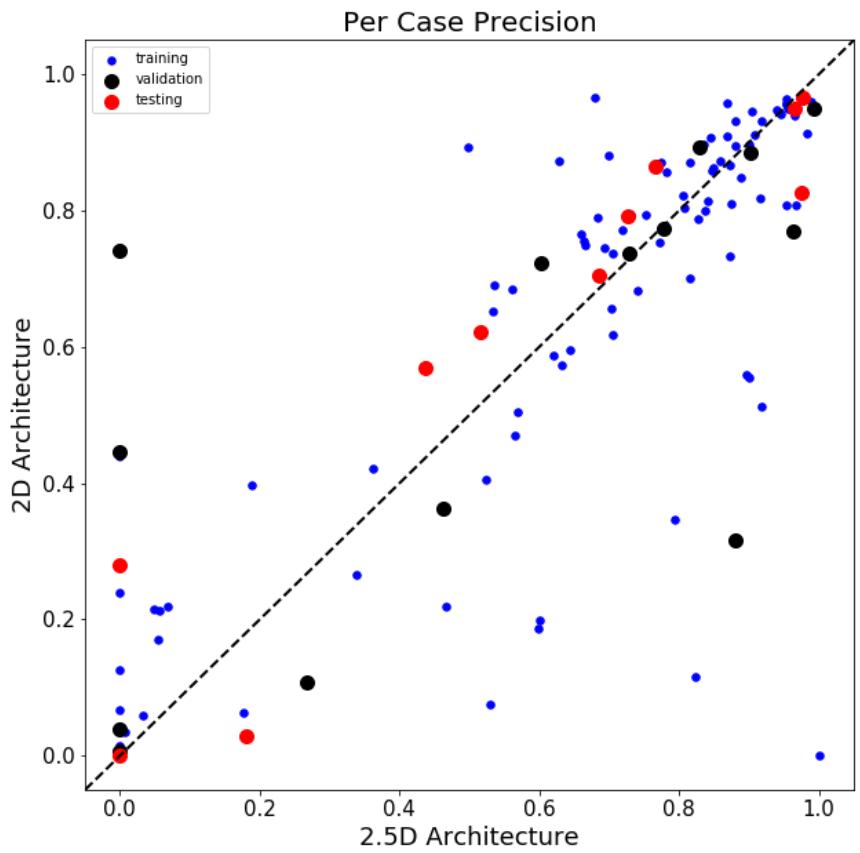
- Add more context information
- 1st LiTS round winner used 2.5D [3]

[3] Han X. "Automatic Liver Lesion Segmentation Using A Deep Convolutional Neural Network Method." 2017.
Medical Knowledge Through Research

2.5D FCN Architecture

Idea:

- Add more context information
- 1st LiTS round winner used 2.5D [3]
- 2.5D vs 2D:
 - Precision ↑



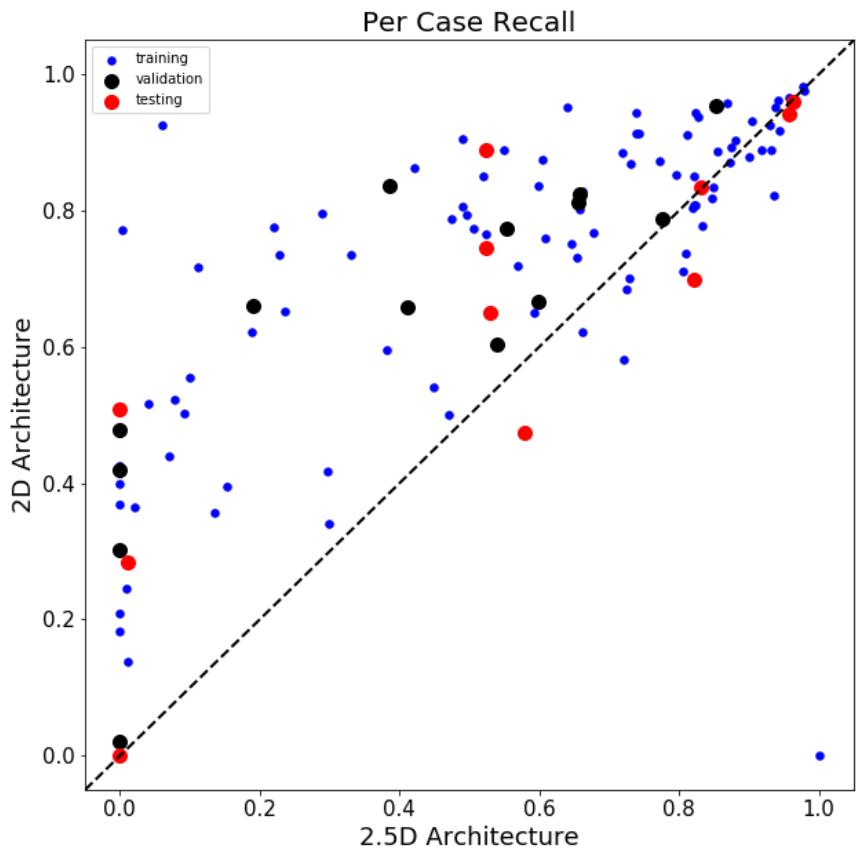
[3] Han X. "Automatic Liver Lesion Segmentation Using A Deep Convolutional Neural Network Method." 2017.

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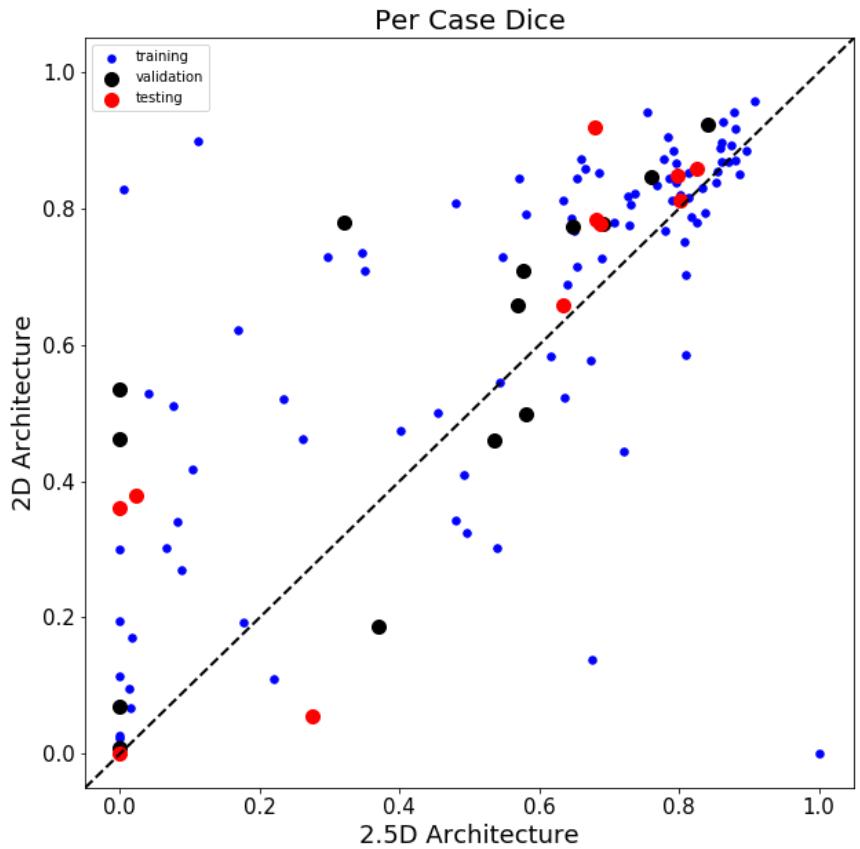
[3] Han X. "Automatic Liver Lesion Segmentation Using A Deep Convolutional Neural Network Method." 2017.

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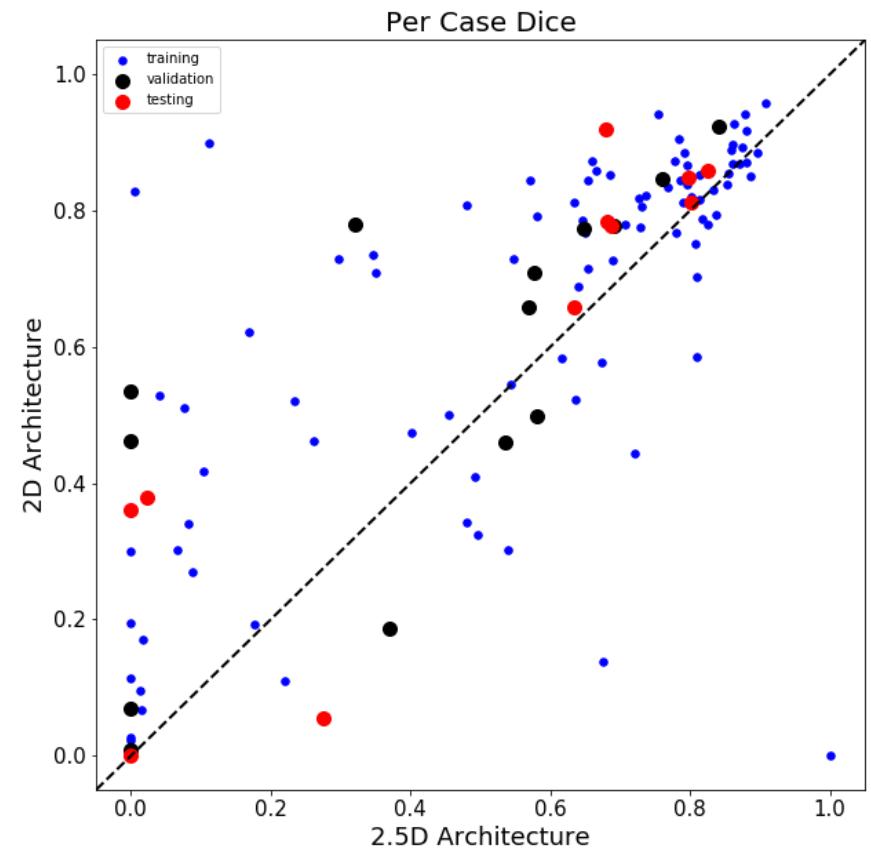
[3] Han X. "Automatic Liver Lesion Segmentation Using A Deep Convolutional Neural Network Method." 2017.

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2.5D FCN Architecture

Idea:

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- 1st LiTS round winner used 2.5D [3]
- 2.5D vs 2D:
 - Precision
 - Recall
 - Dice
- Same observations for liver!



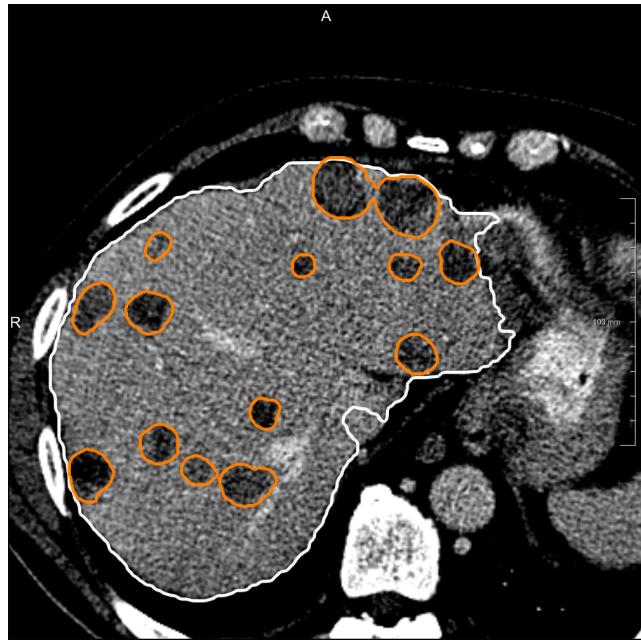
[3] Han X. "Automatic Liver Lesion Segmentation Using A Deep Convolutional Neural Network Method." 2017.

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Examples

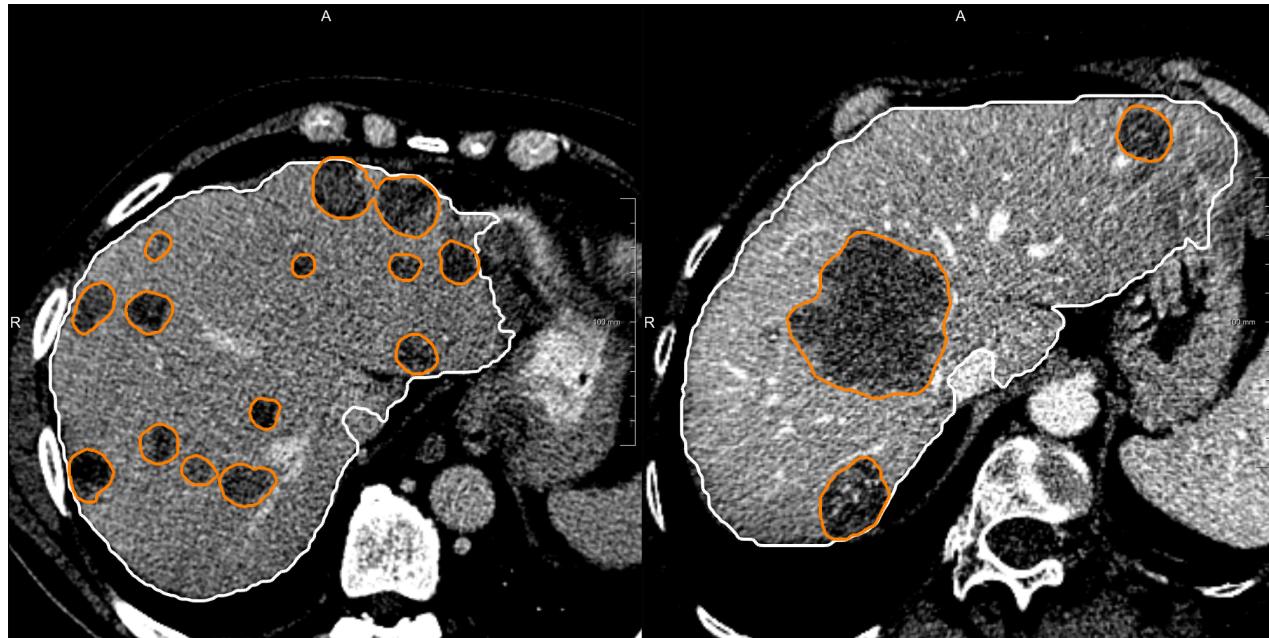
Medical Knowledge Through Research

Examples



Medical Knowledge Through Research

Examples



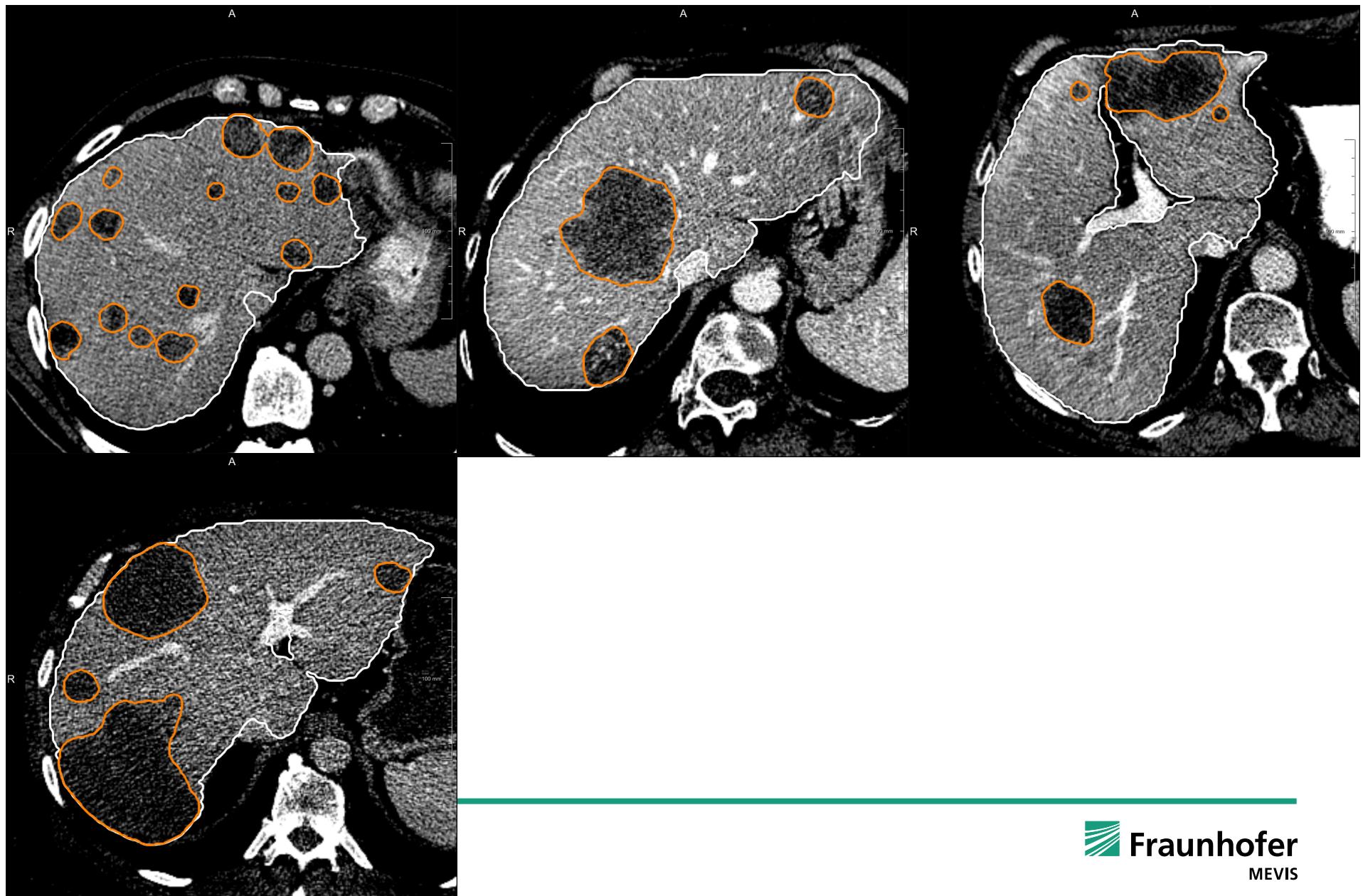
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Examples

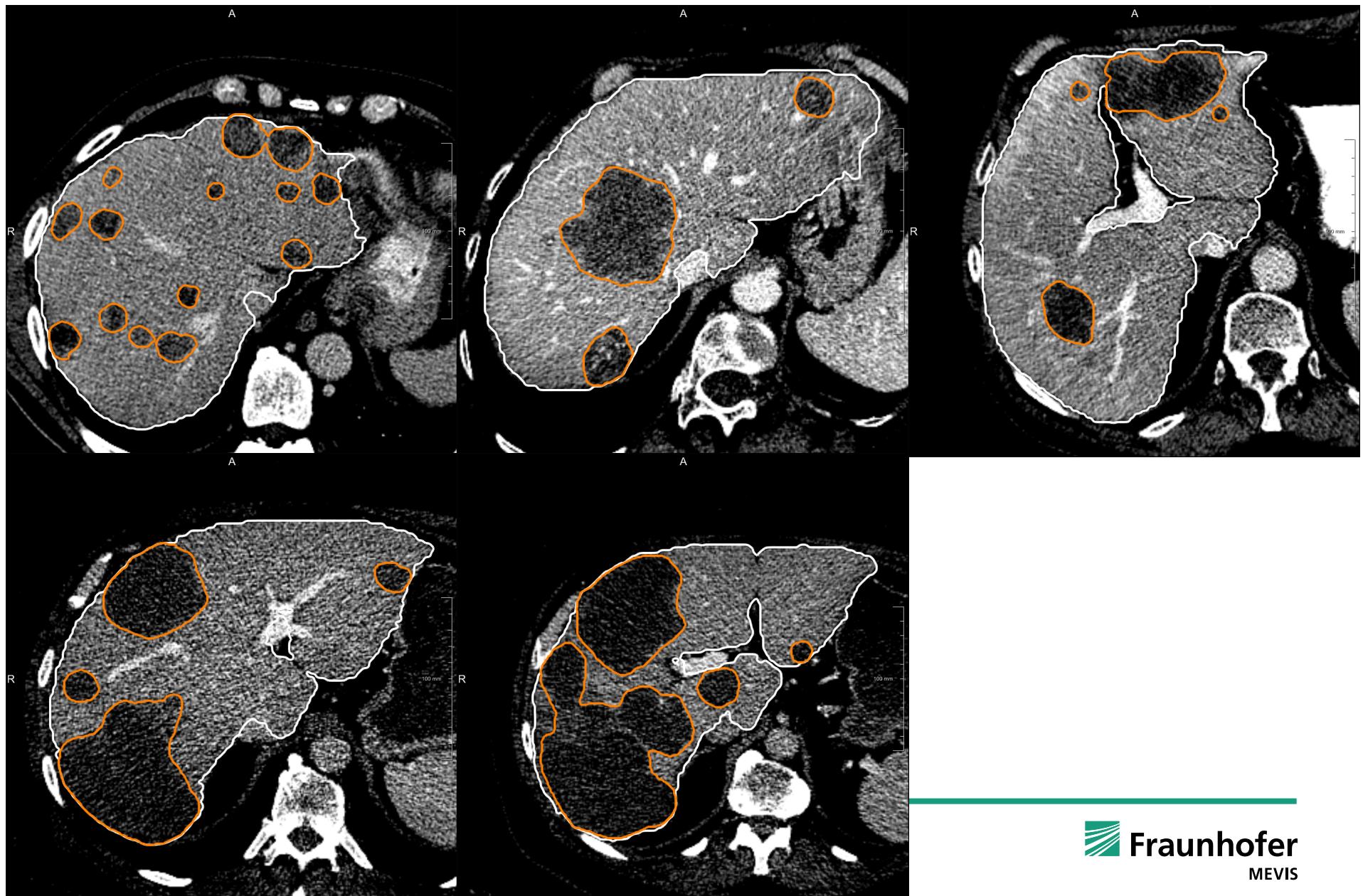


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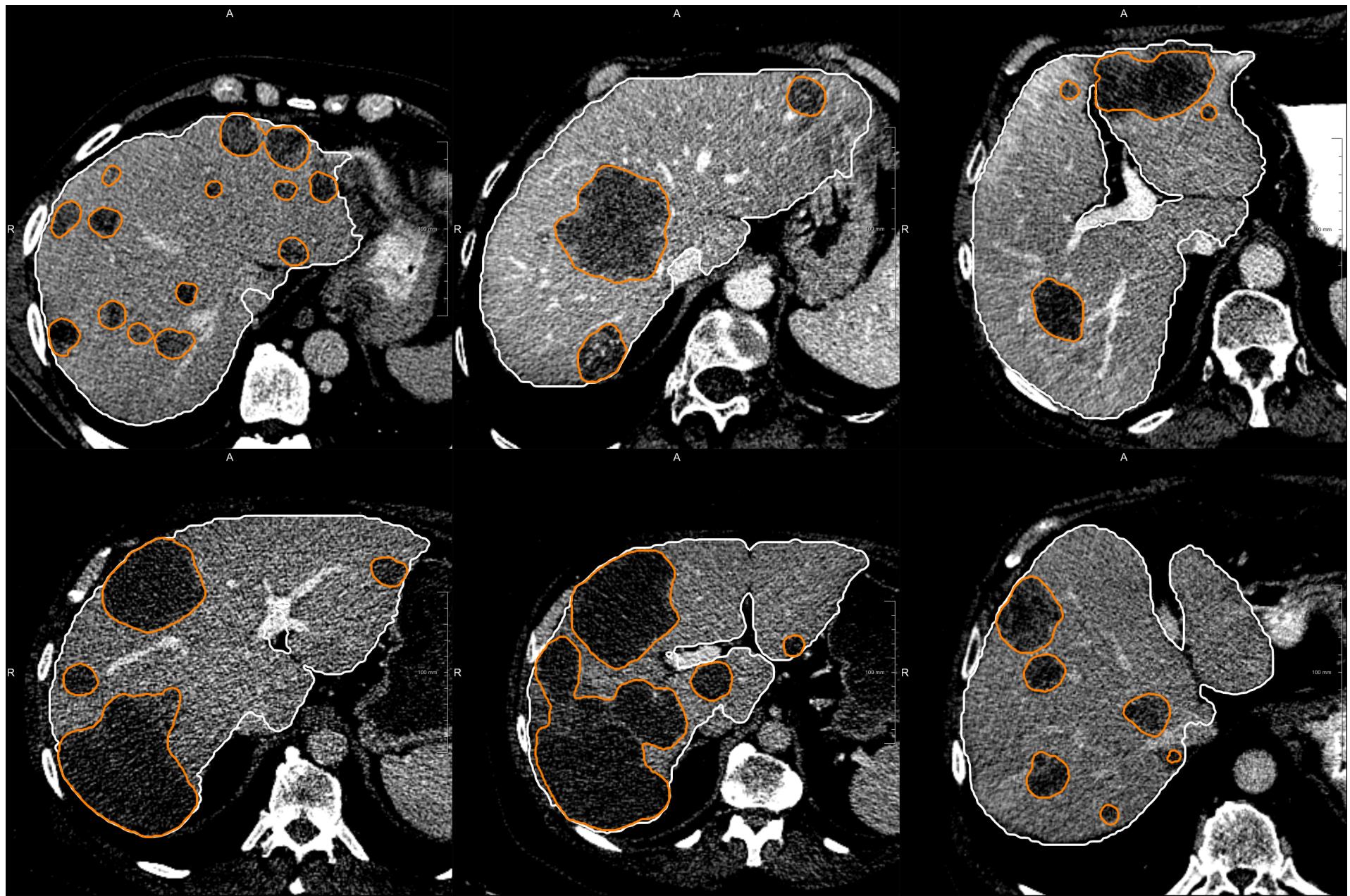
Examples



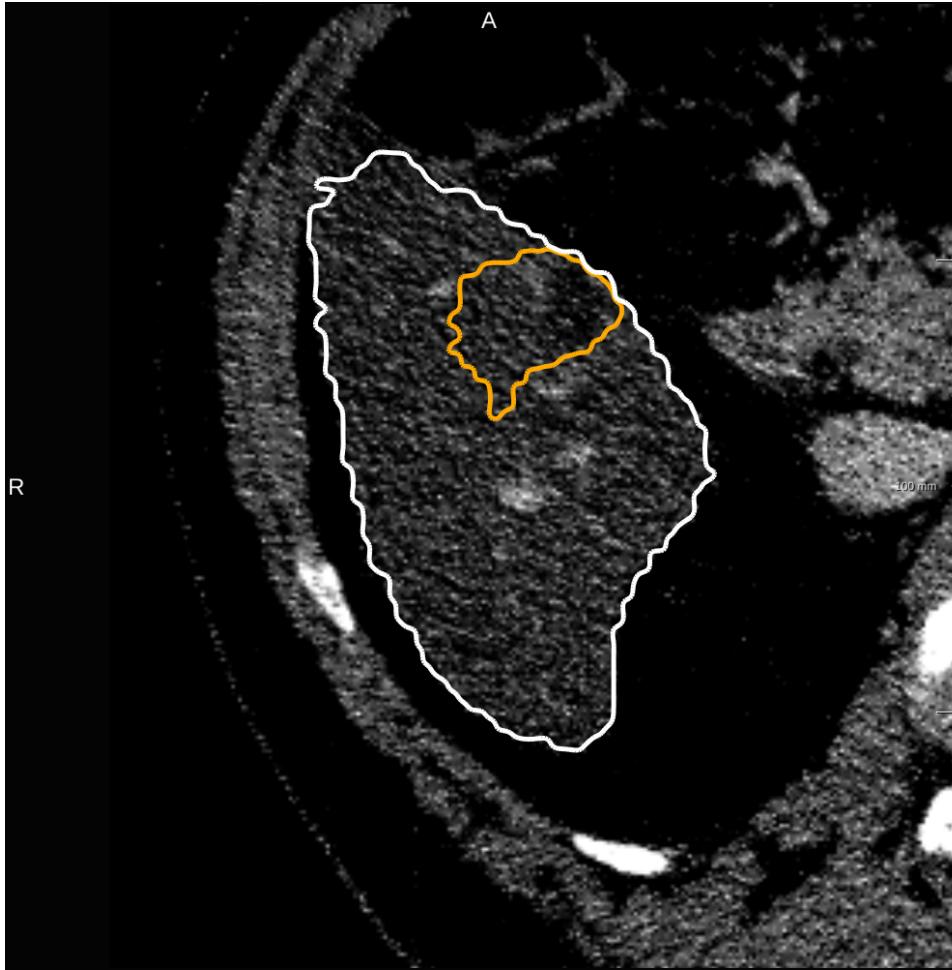
Examples



Examples



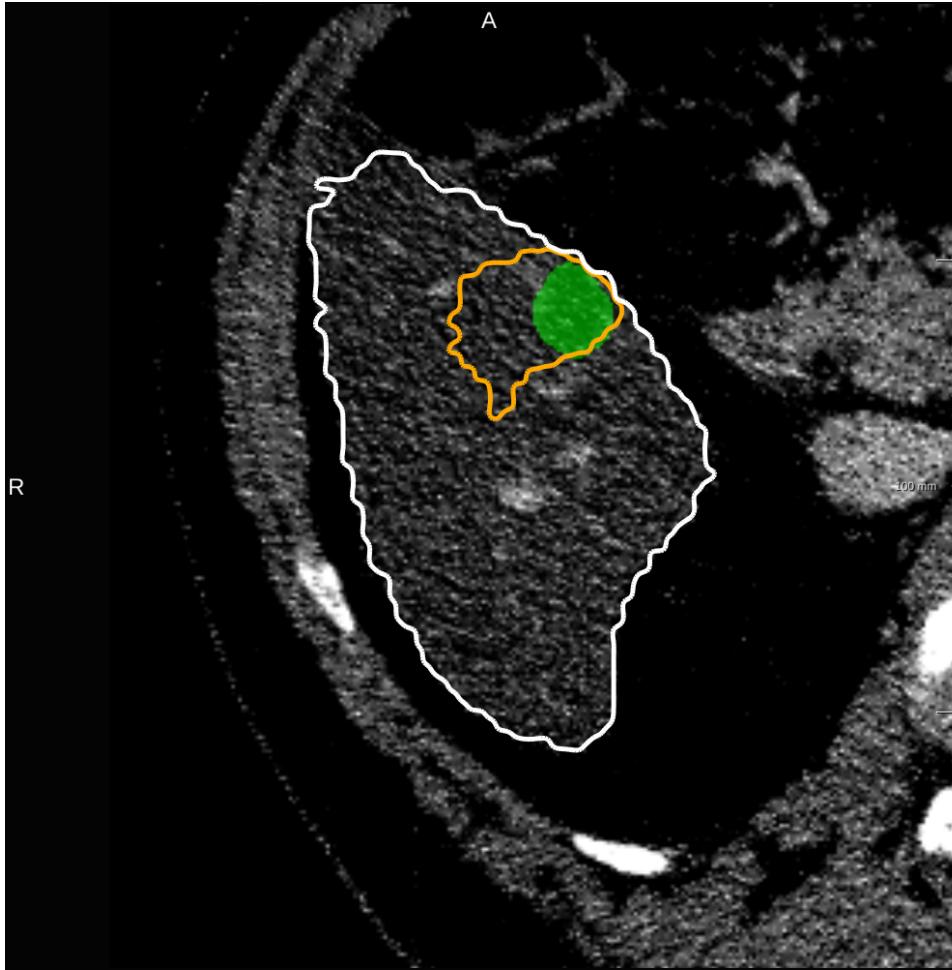
Problematic Cases



- Found tumor bigger than the reference

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Problematic Cases



- Found tumor bigger than the reference

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Problematic Cases



- Liver mask misses tumors located near organ's border

Problematic Cases



- Big tumors are not fully segmented

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Problematic Cases



- Obvious(?) tumors are completely missed

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LiTS Results

- Liver segmentation
 - Dice per case: 0.96
 - Relative volume difference: -0.4%

LiTS Results

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- Tumor segmentation
 - Dice per case: 0.68
 - Precision at > 0% overlap: 0.72
 - Recall at > 0% overlap: 0.57

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 - RMSE: 0.02
 - Max: 0.07

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 - Liver segmentation: ~43 s
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OK



Further work
required

Technical Setup

- Data loading and preprocessing

- MeVisLab



- Deep Learning Toolkits

- RedLeaf
 - Lasagne
 - Theano

Lasagne



theano

- Evaluation

- Challengr



Conclusions

- We proposed a fully automatic method for liver and liver lesion segmentation based on FCNs
- False positive tumors were filtered with a high accuracy using image intensity and shape based features
- Providing more context to the network (2.5D) decreased the segmentation quality
- Further work is required to make tumor segmentation clinically applicable

Outlook

- Different architectures
 - Adversarial networks
 - Recurrent networks
- Other training strategies
 - Curriculum learning



Outlook

- Different architectures
 - Adversarial networks
 - Recurrent networks
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Thank you for your attention 😊

Questions?

