



Automatic segmentation of fibula bone by using deep learning

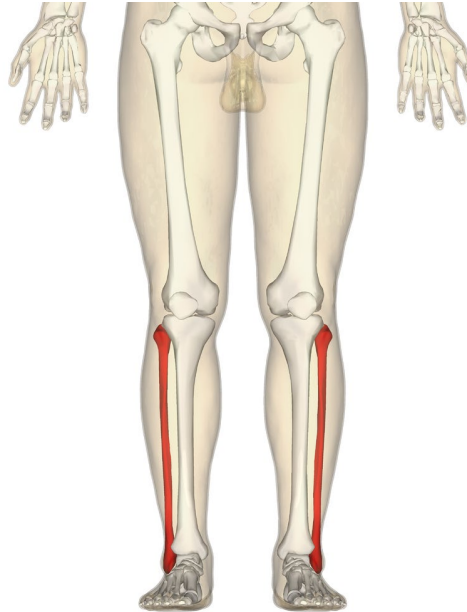
Promotor: Prof. Bart Vanrumste

Co-supervisor: Dr. Yi Sun

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Zechun Wang	r0736595

Subject of master thesis

- Proposes an **automatic fibula segmentation** approach in CT scans





Outline

1. Motivation
2. Dataset & Data preprocessing
3. Methodology & Experiments
4. Conclusion

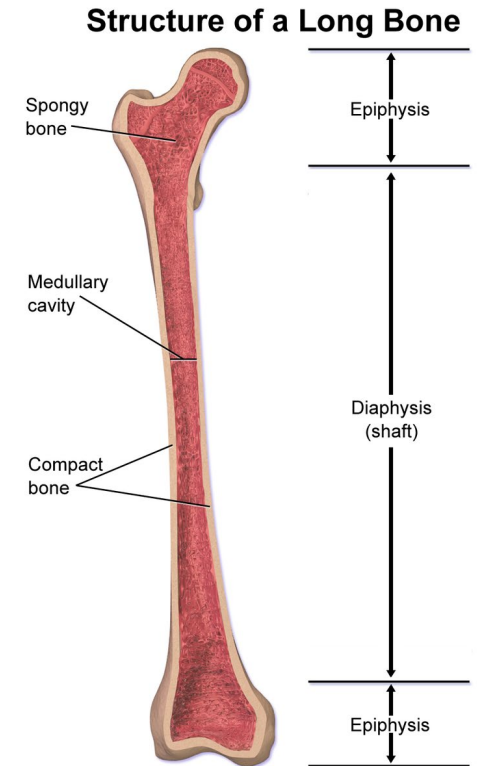
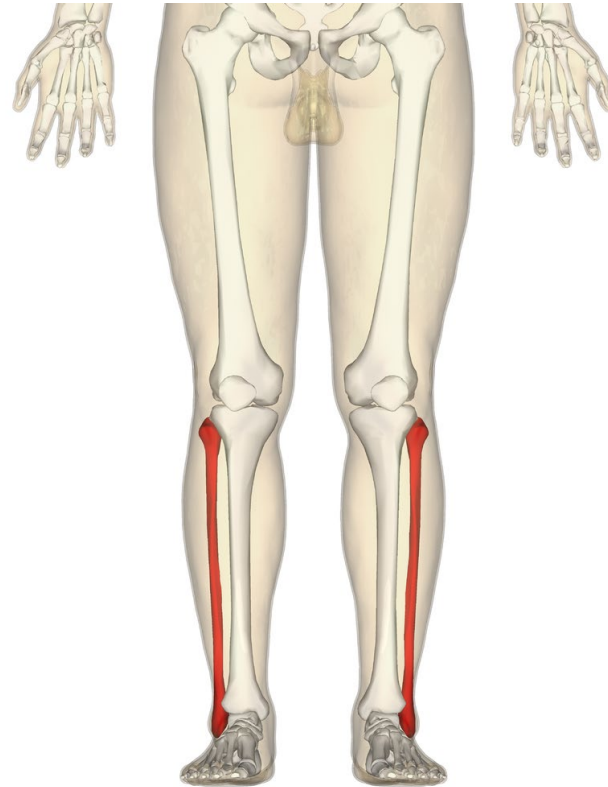
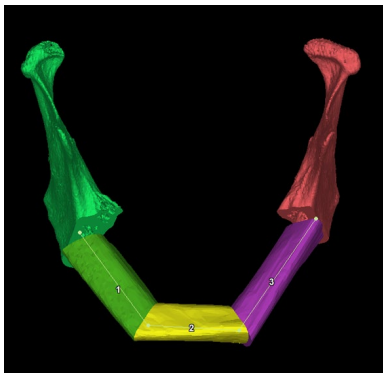


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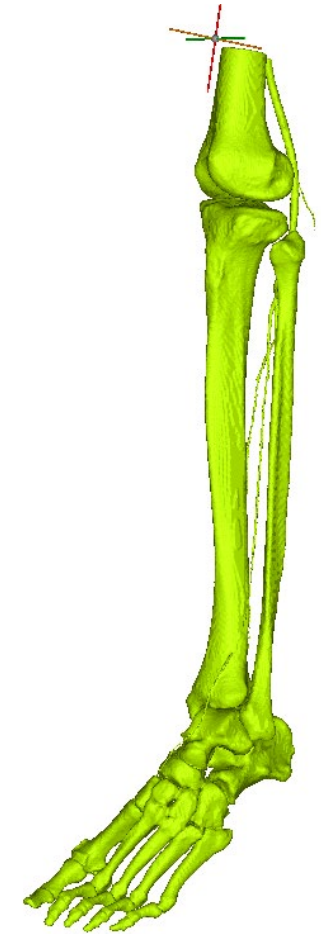
Concept of Fibula

- The fibula is a **leg bone** on the lateral side of the tibia.
- Segmentation of fibula is commonly used in the **mandibular reconstruction**.



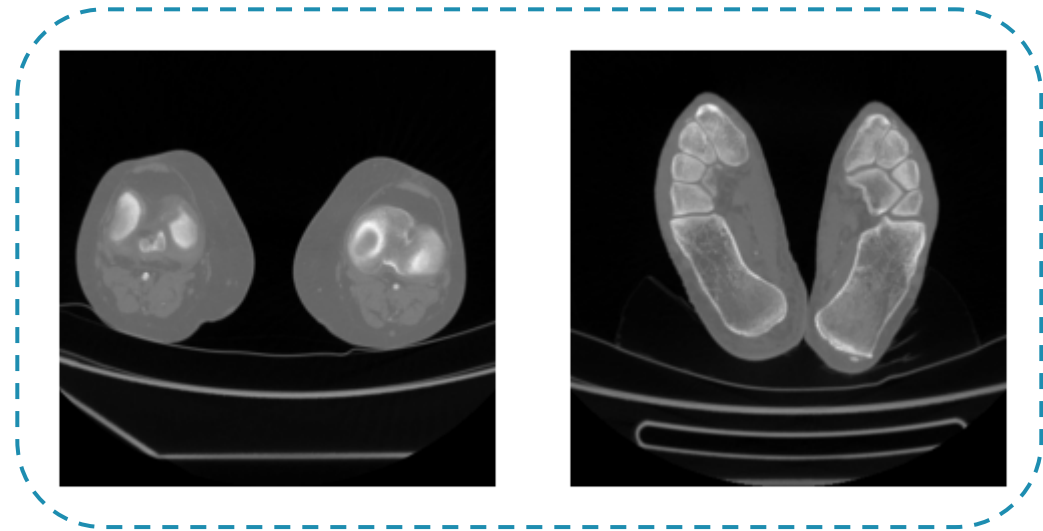
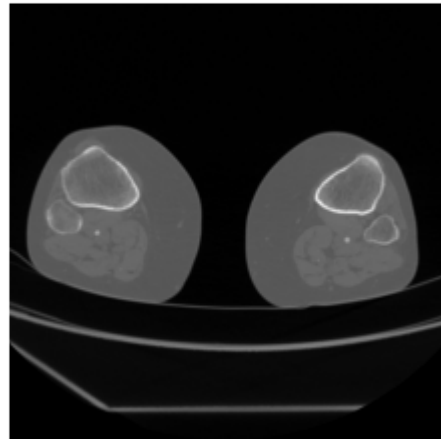
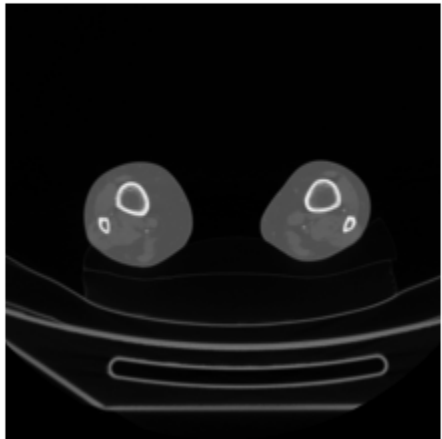
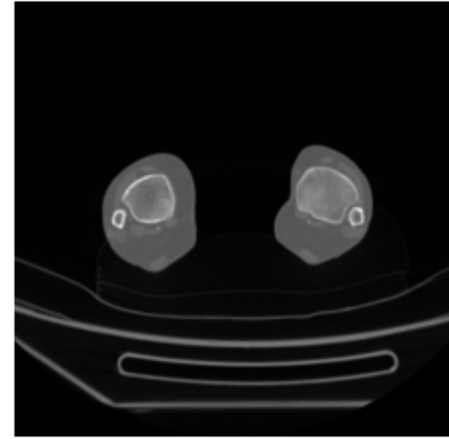
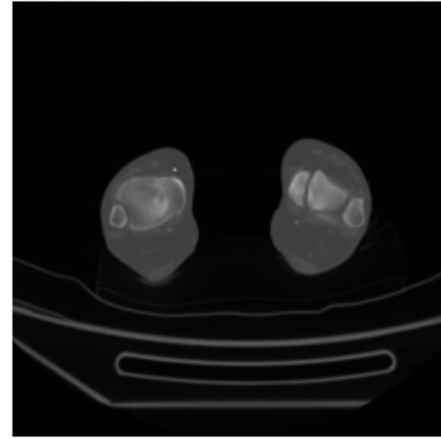
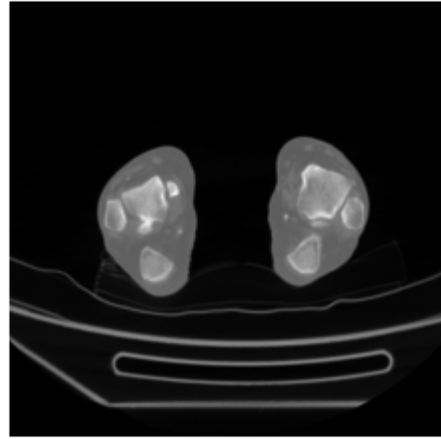
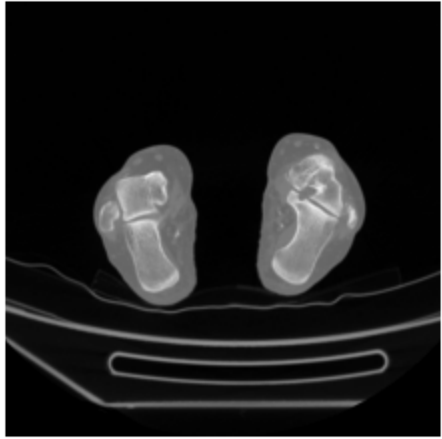
The traditional fibula segmentation method

- Traditional medical segmentation adopts **thresholding algorithm** method with the assistant of manual annotations.
- The segmentation results of thresholding algorithm **cannot be directly adopted** in clinical applications, experts are still required to spend lots of time to **further manually improve** on the results' quality. Zhou et al. (2019)



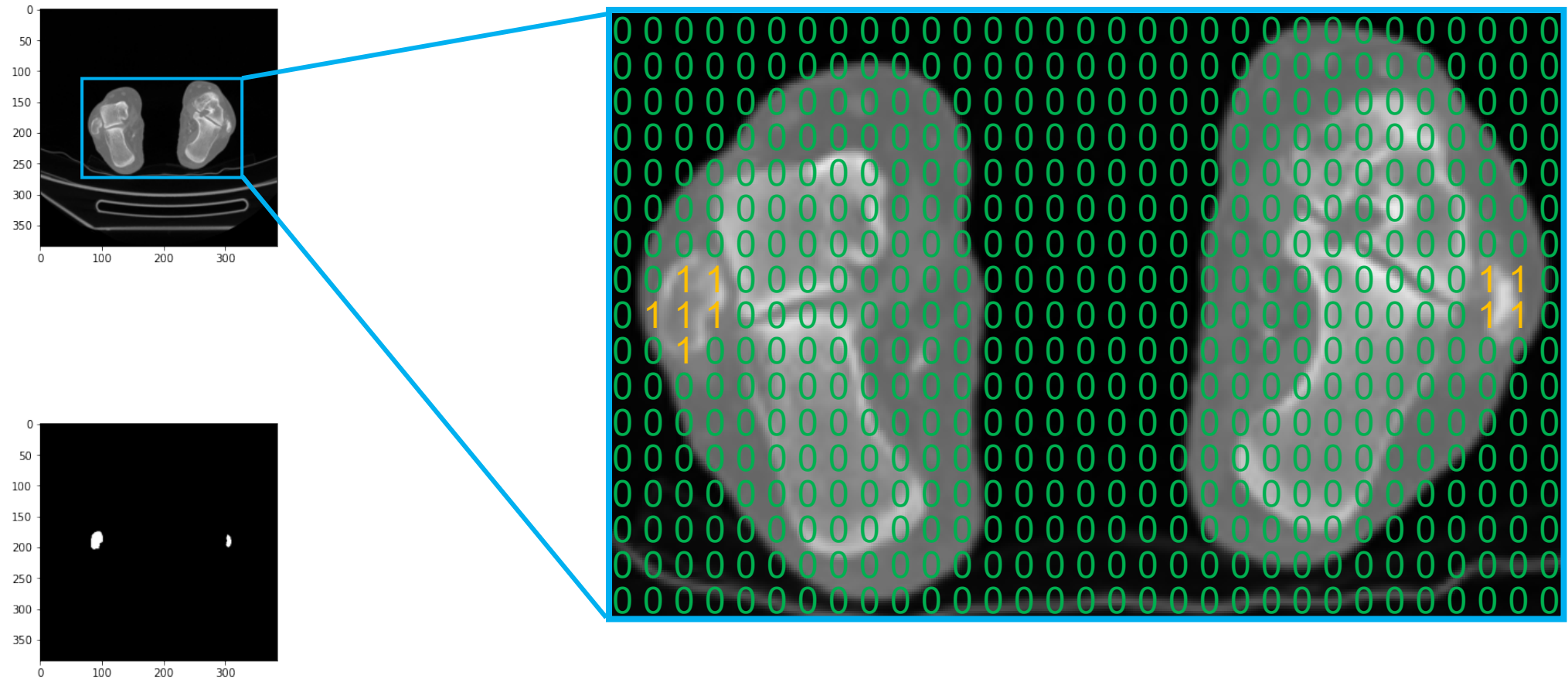


Challenge





What is segmentation





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Dataset

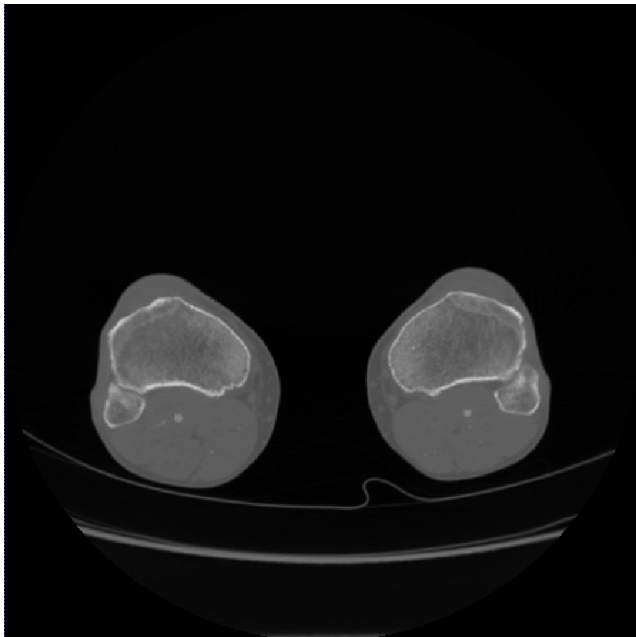
- Source: Sint-Rafael hospital
- Image Dimension: 512 x 512
- Data distribution
- Dataset description
 - 19 CT scans
 - 409 to 1543 CT slices
 - 15830 images
 - 20.4 GB

	Train set	Validation set	Test set
Number of CT scans	16	1	2
Number of images	13636	790	1404

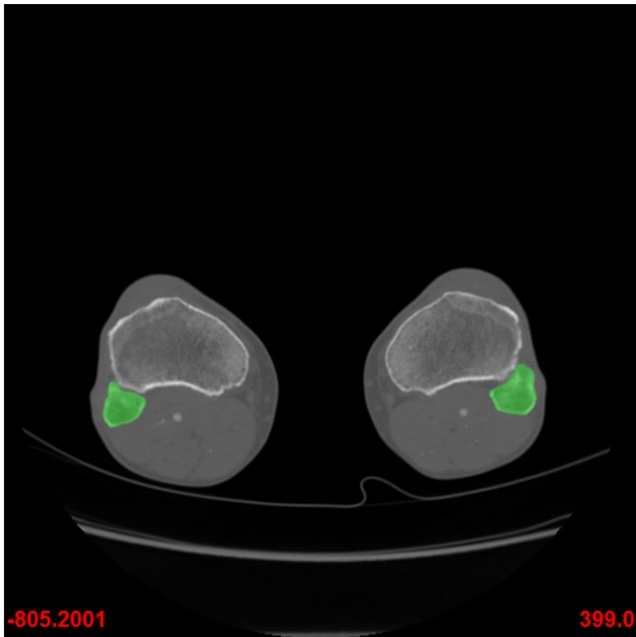
- Limitation: Only one validation set



Dataset



Original CT slice



Experts' manual annotations

Examples in the data set

Serial number	Number of CT slices in one scan	Distance between each slice (mm)
1	734	0.6
2	670	0.6
3	790	0.6
4	786	0.6
5	794	0.6
6	799	0.6
7	810	0.6
8	951	0.6
9	803	0.7
10	767	0.8
11	761	0.8
12	999	0.625
13	1543	0.4
14	901	0.7
15	759	0.7
16	797	0.7
17	774	0.8
18	409	1.5
19	983	0.6

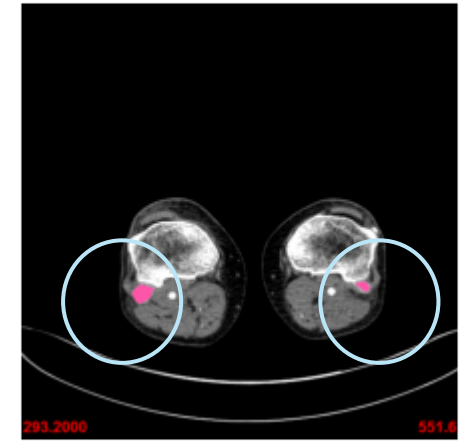
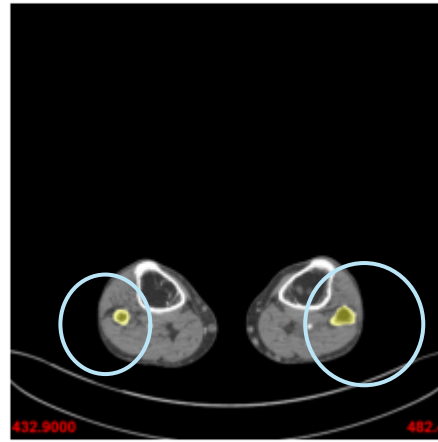
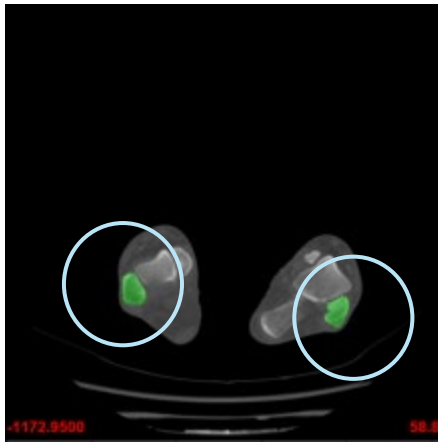


Outline

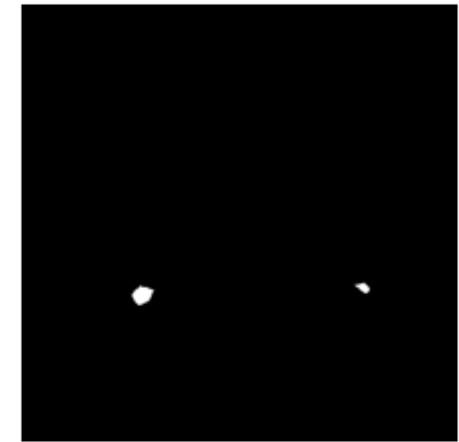
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Data preprocessing

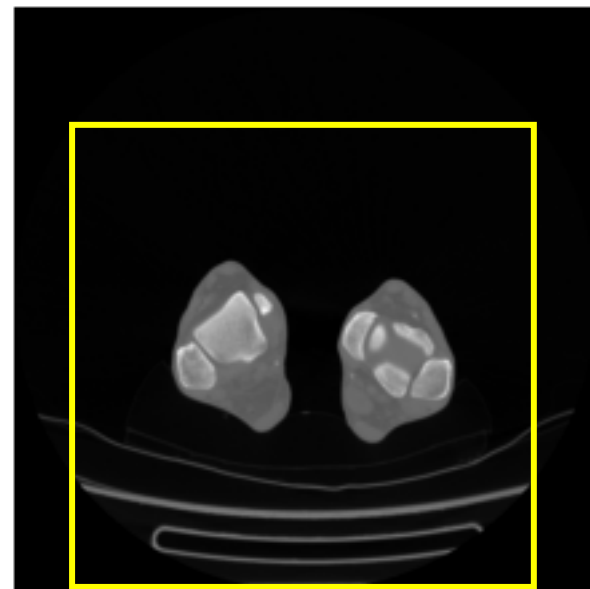


The original data of the manual annotations

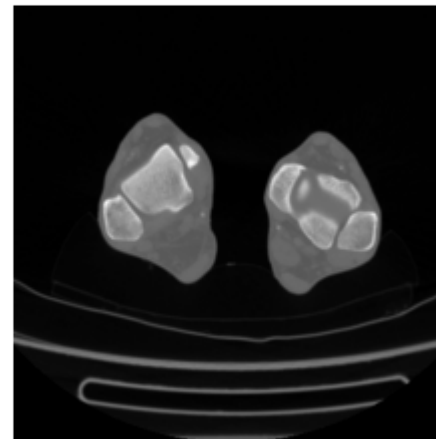
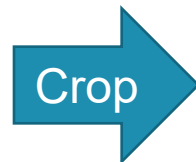


the ground truth after HSV processing

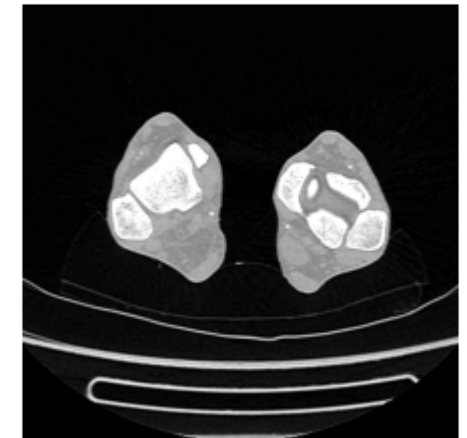
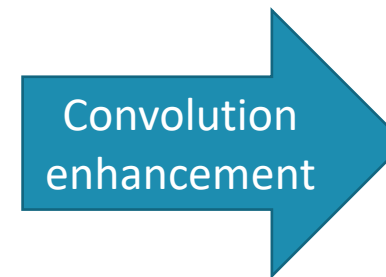
Data preprocessing



512 x 512



384 x 384



384 x 384

For boundary enhancement

Convolution kernel

$$\begin{bmatrix} -1 & -1 & -1 \\ -1 & 10 & -1 \\ -1 & -1 & -1 \end{bmatrix}$$



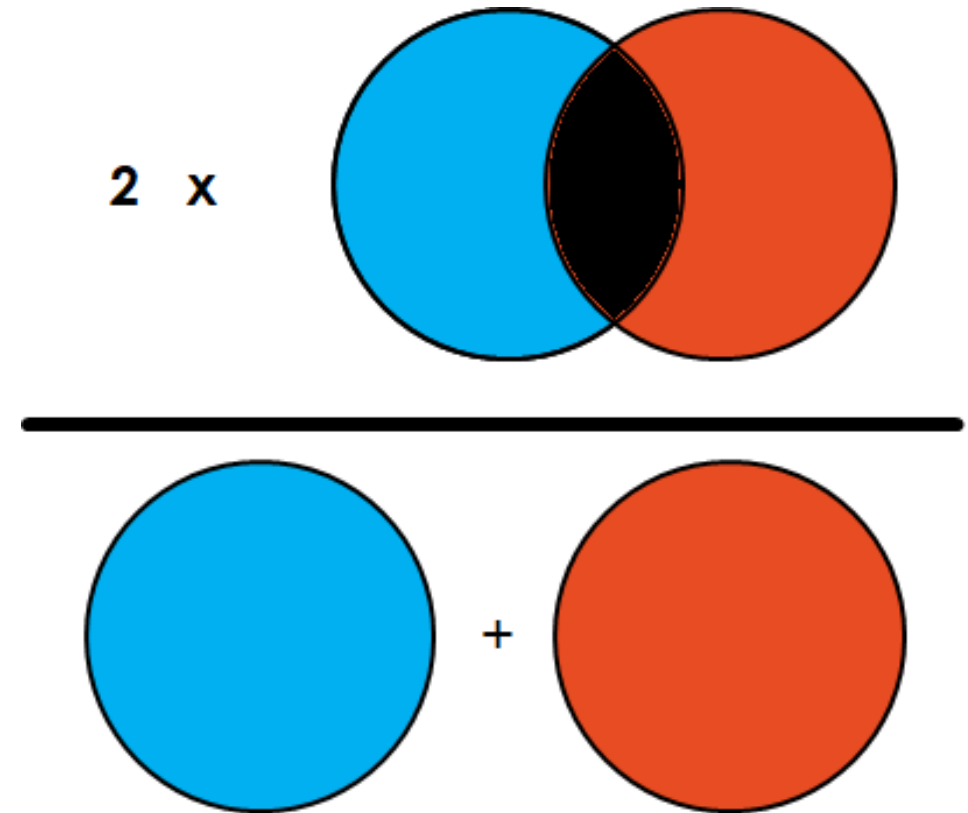
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Dice Score

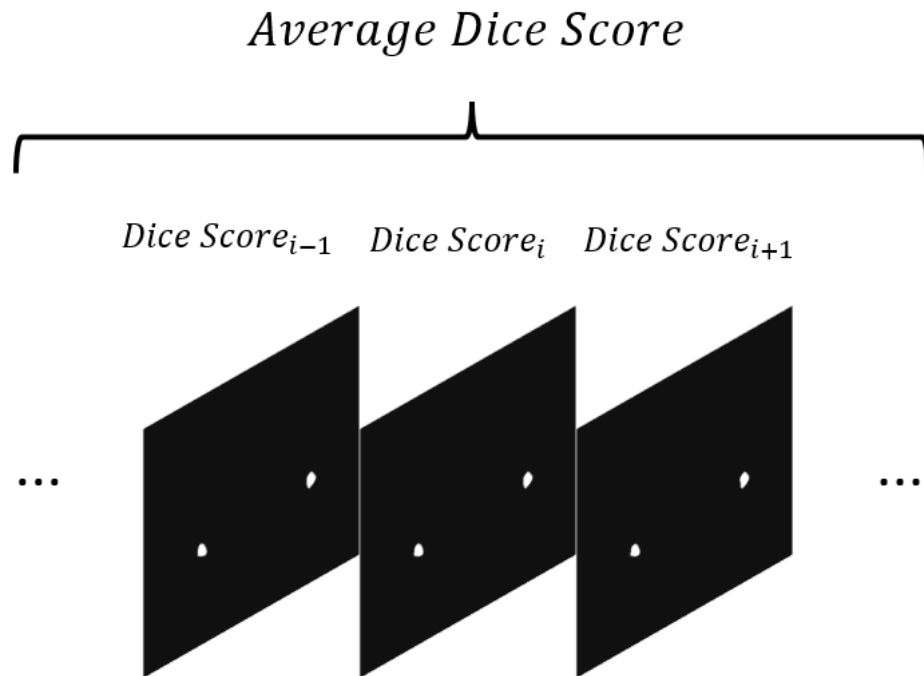
- $D(X, Y) = \frac{2 * |X \cap Y|}{|X| + |Y|}$
- By using this method, a score between 0 and 1 can be computed.



Two method of Dice Score

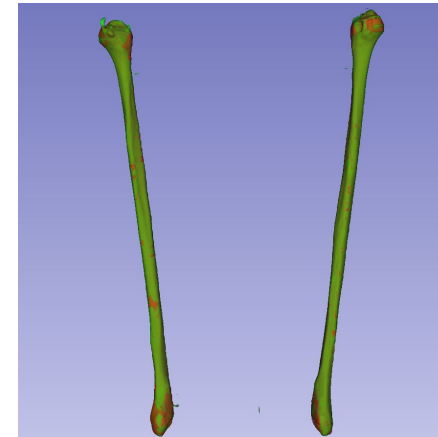
Average Dice score

- Average of each dice score in one CT scan.
- Focus more on evaluating the whole output



Volumetric Dice score

- Overlap voxel area
- Pay more attention to the evaluation of fibula.



$$\text{Volumetric Dice Score} = \frac{2 * |X \cap Y|}{|X| + |Y|}$$

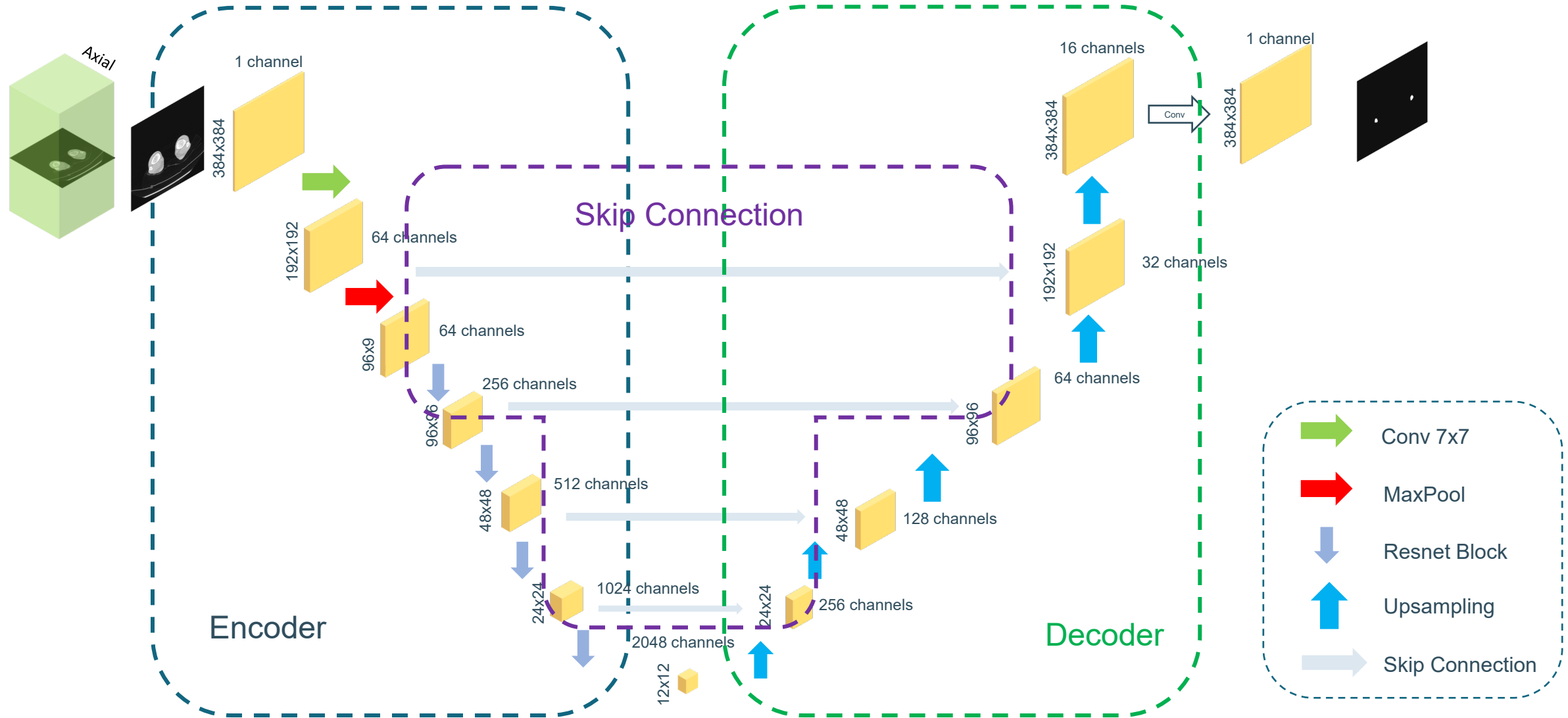


Outline

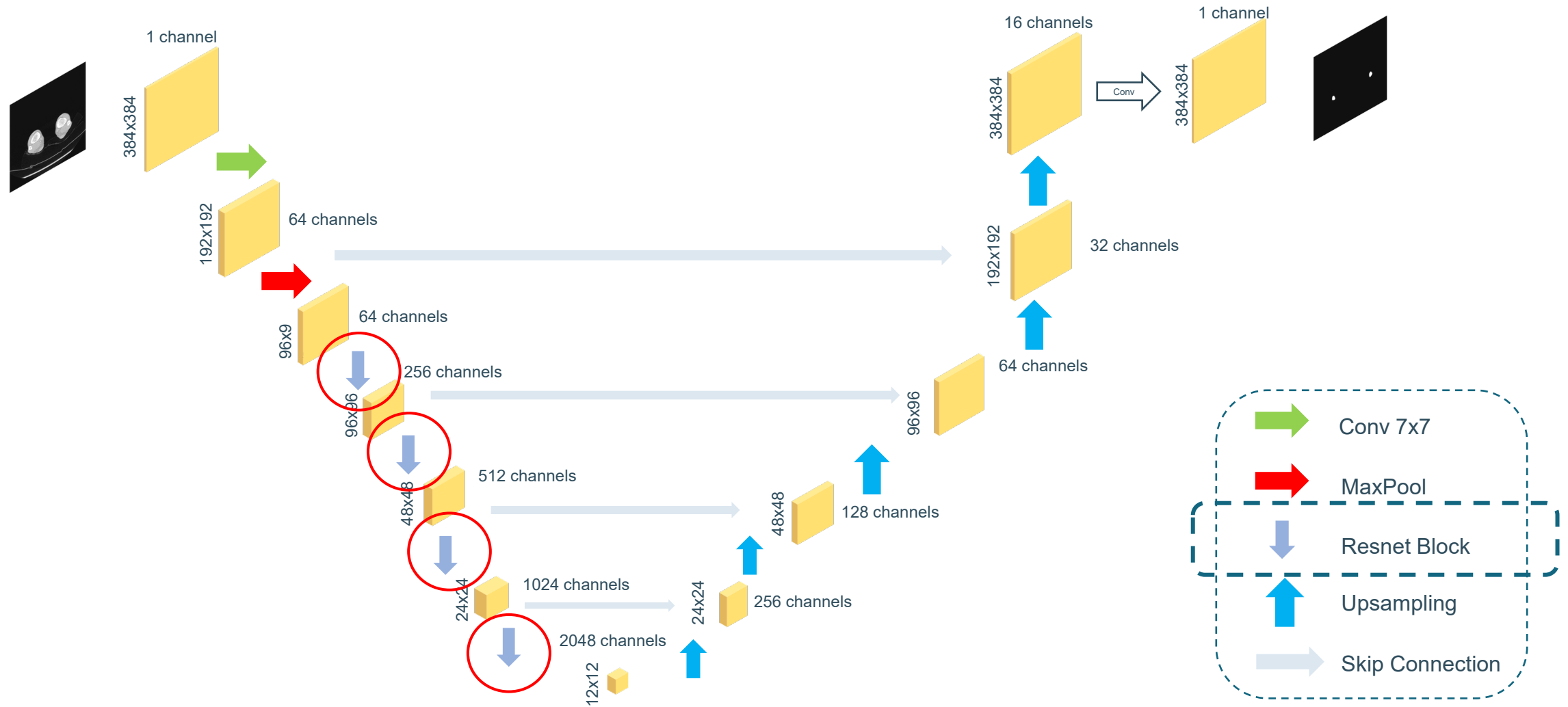
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Single-planar segmentation model

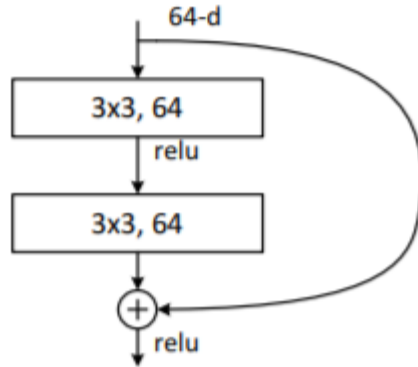


Single-planar segmentation model: Res-Net block

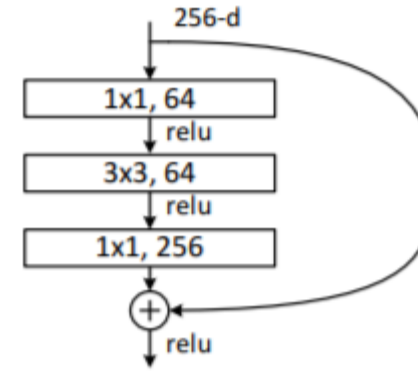


Res-Net block

- Traditional Res-Net consists of building blocks or bottlenecks



The structure of the building block for ResNet-18 and ResNet-34



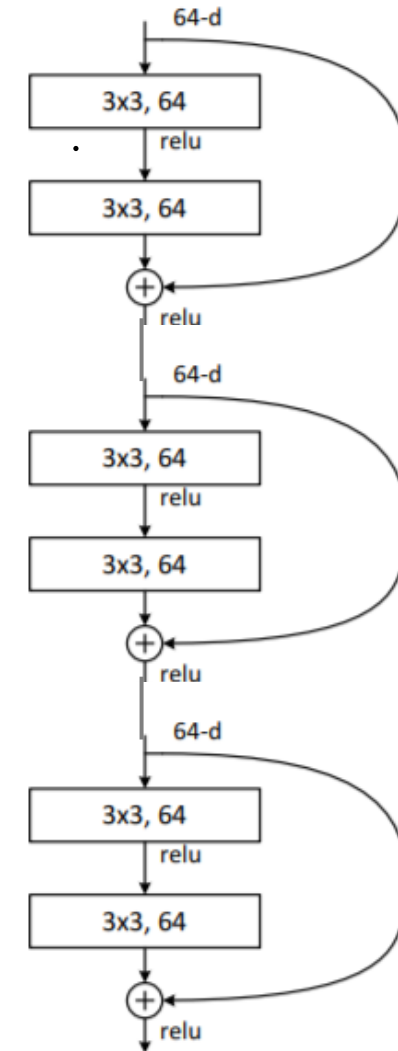
The structure of the bottleneck block for ResNet-50 and higher Res-Net



Res-Net architecture

- The Architectures for 34 layers and 50 layers Res-Net

Layer Name	34-layer	50-layer
Conv1	7 × 7, 64, stride 2	
Conv2-x	3 × 3 max pool, stride 2	
	$\begin{bmatrix} 3 \times 3, & 64 \\ 3 \times 3, & 64 \end{bmatrix} \times 3$	$\begin{bmatrix} 1 \times 1, & 64 \\ 3 \times 3, & 64 \\ 1 \times 1 & 256 \end{bmatrix} \times 3$
Conv3-x	$\begin{bmatrix} 3 \times 3, & 128 \\ 3 \times 3, & 128 \end{bmatrix} \times 4$	$\begin{bmatrix} 1 \times 1, & 128 \\ 3 \times 3, & 128 \\ 1 \times 1 & 512 \end{bmatrix} \times 4$
Conv4-x	$\begin{bmatrix} 3 \times 3, & 256 \\ 3 \times 3, & 256 \end{bmatrix} \times 6$	$\begin{bmatrix} 1 \times 1, & 256 \\ 3 \times 3, & 256 \\ 1 \times 1 & 1024 \end{bmatrix} \times 6$
Conv5-x	$\begin{bmatrix} 3 \times 3, & 512 \\ 3 \times 3, & 512 \end{bmatrix} \times 3$	$\begin{bmatrix} 1 \times 1, & 512 \\ 3 \times 3, & 512 \\ 1 \times 1 & 2048 \end{bmatrix} \times 3$





Upsampling

10	4	22
2	18	7
9	14	25

3 x 3

10		4		22	
2		18		7	
9		14		25	

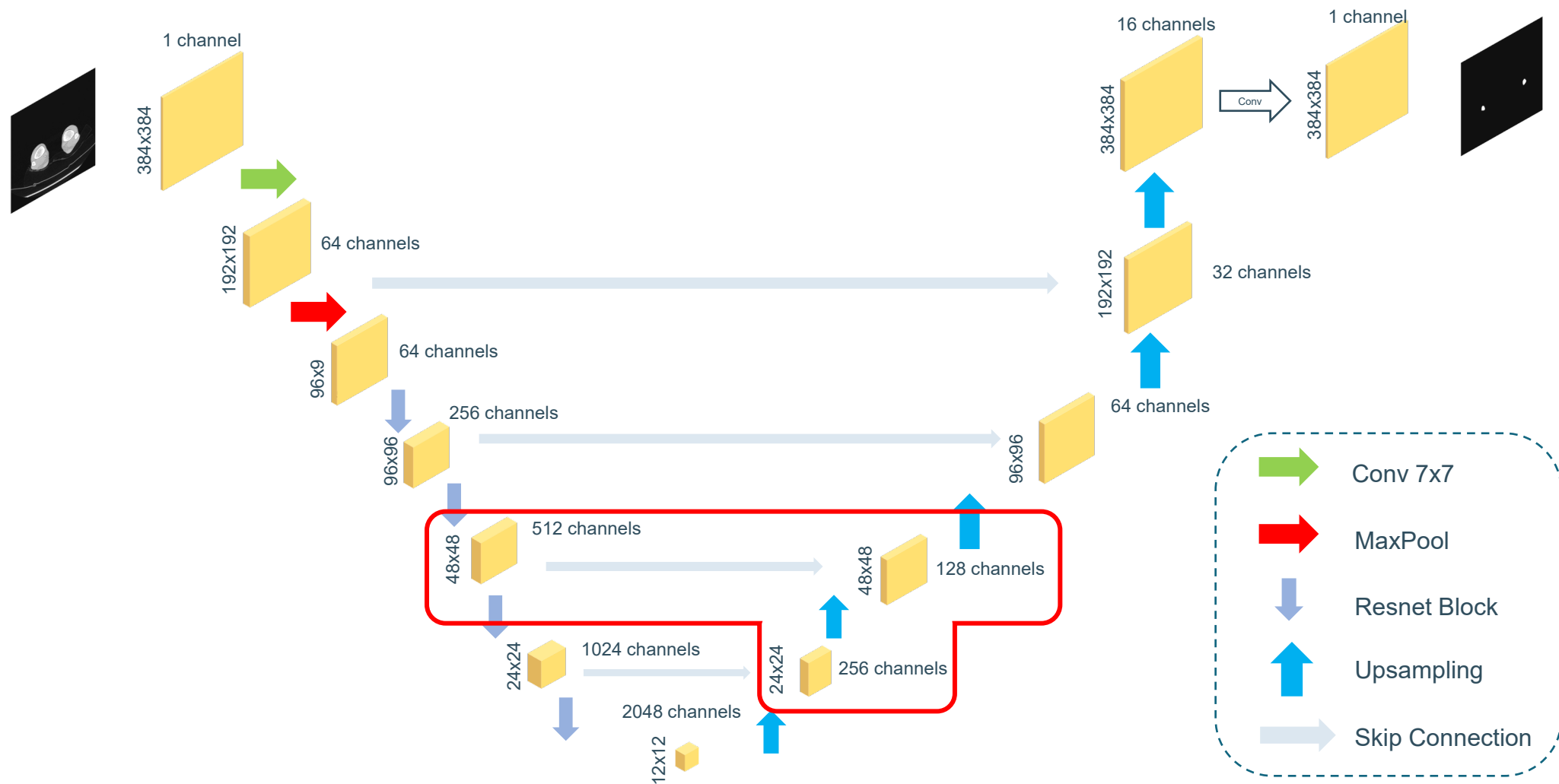
6 x 6

10	10	4	4	22	22
10	10	4	4	22	22
2	2	18	18	7	7
2	2	18	18	7	7
9	9	14	14	25	25
9	9	14	14	25	25

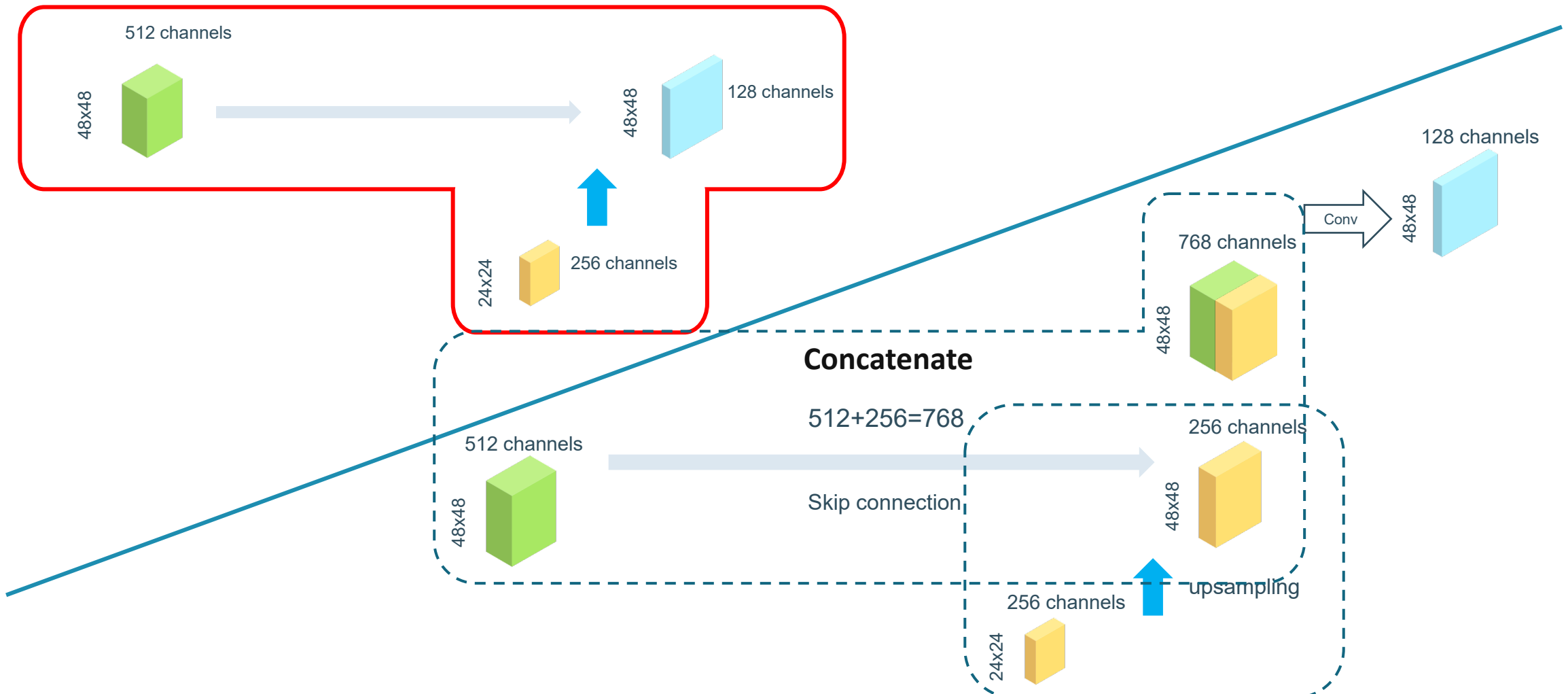
6 x 6

Nearest Neighbor Interpolation

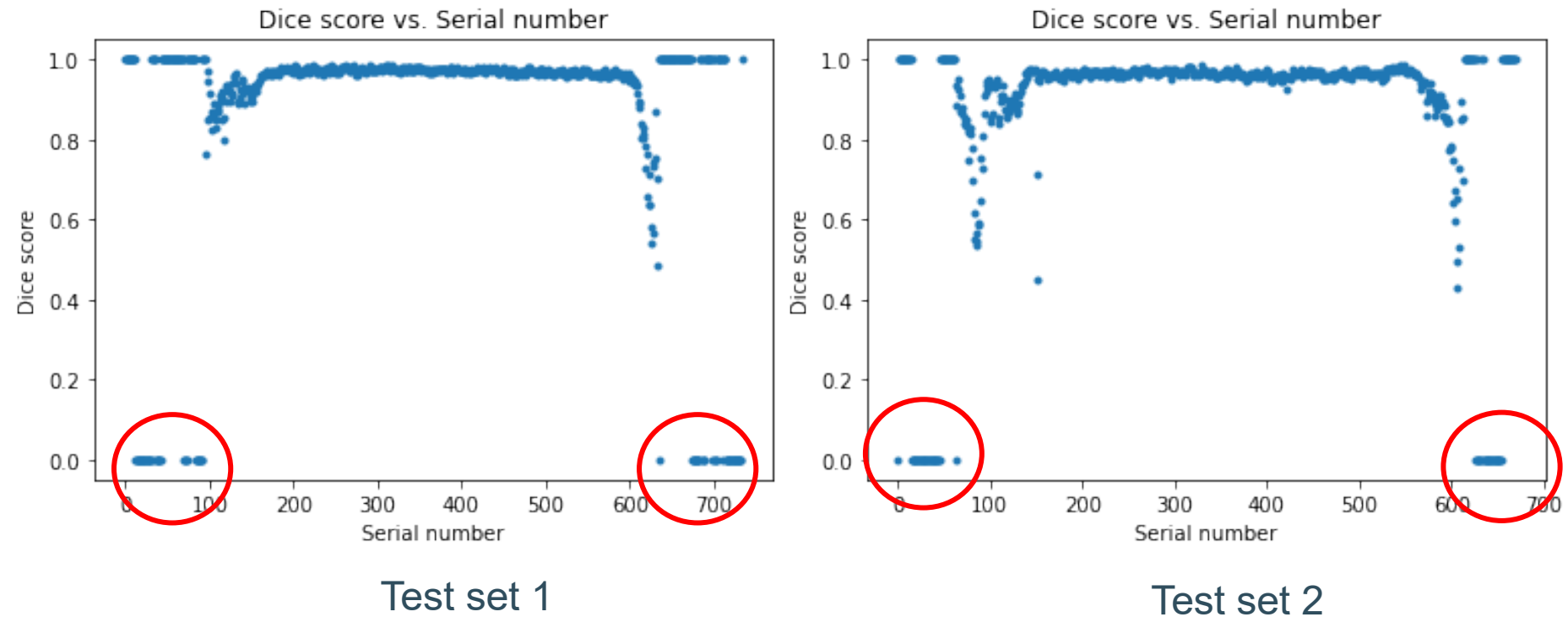
Skip connection



Skip connection

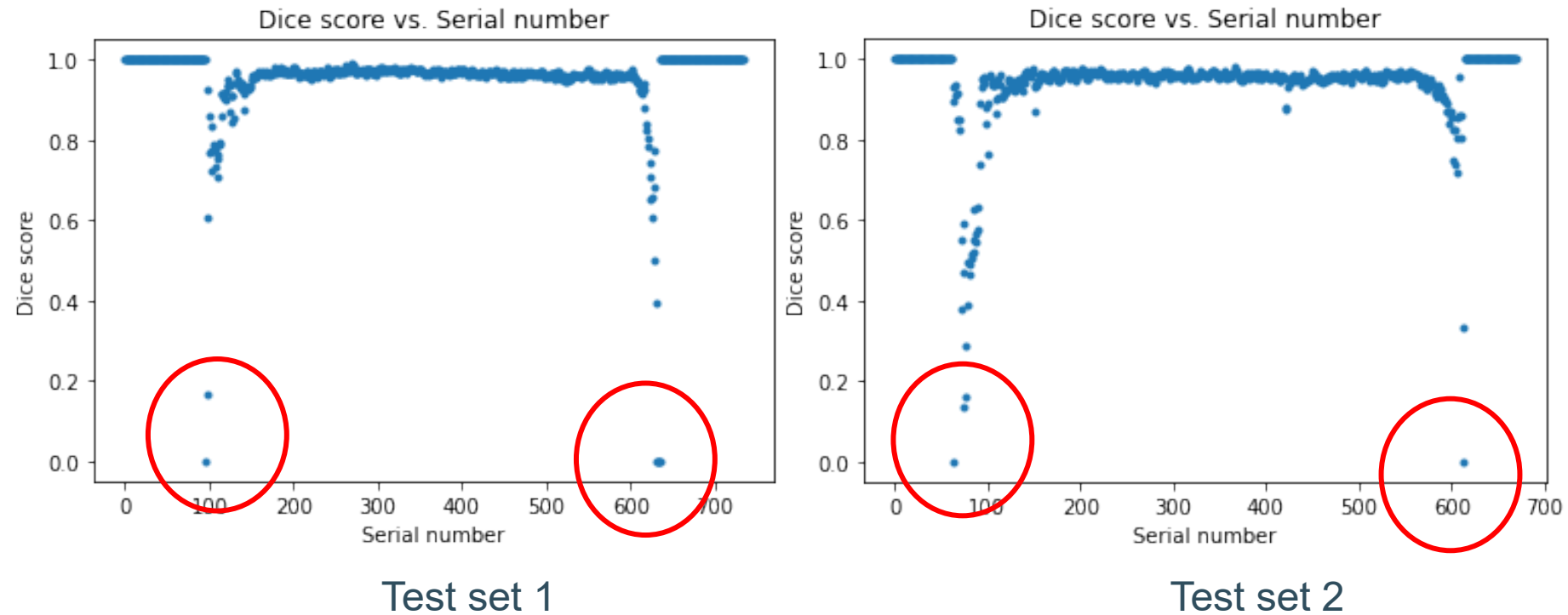


Experiment 1: 34-layer Res-Net in single-planar model



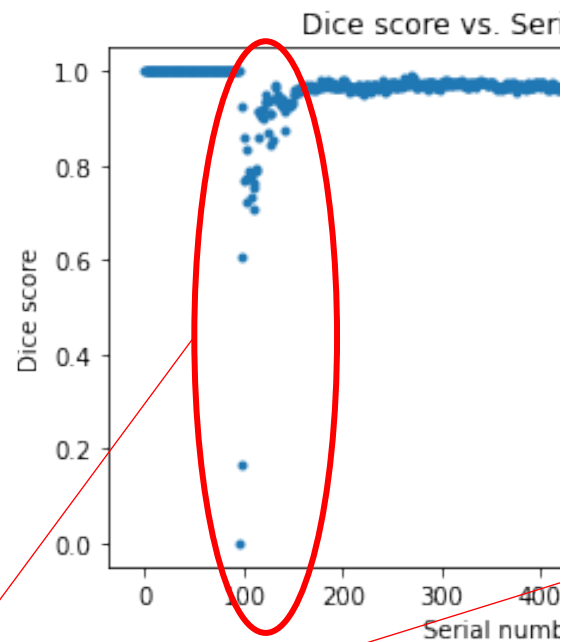
	Average Dice Score	Volumetric Dice Score
Test set 1	0.863	0.920
Test set 2	0.859	0.917

Experiment 2: 50-layer Res-Net in single-planar model



	Average Dice Score	Volumetric Dice Score
Test set 1	0.952	0.946
Test set 2	0.884	0.906

Experiment 2: 50-lay

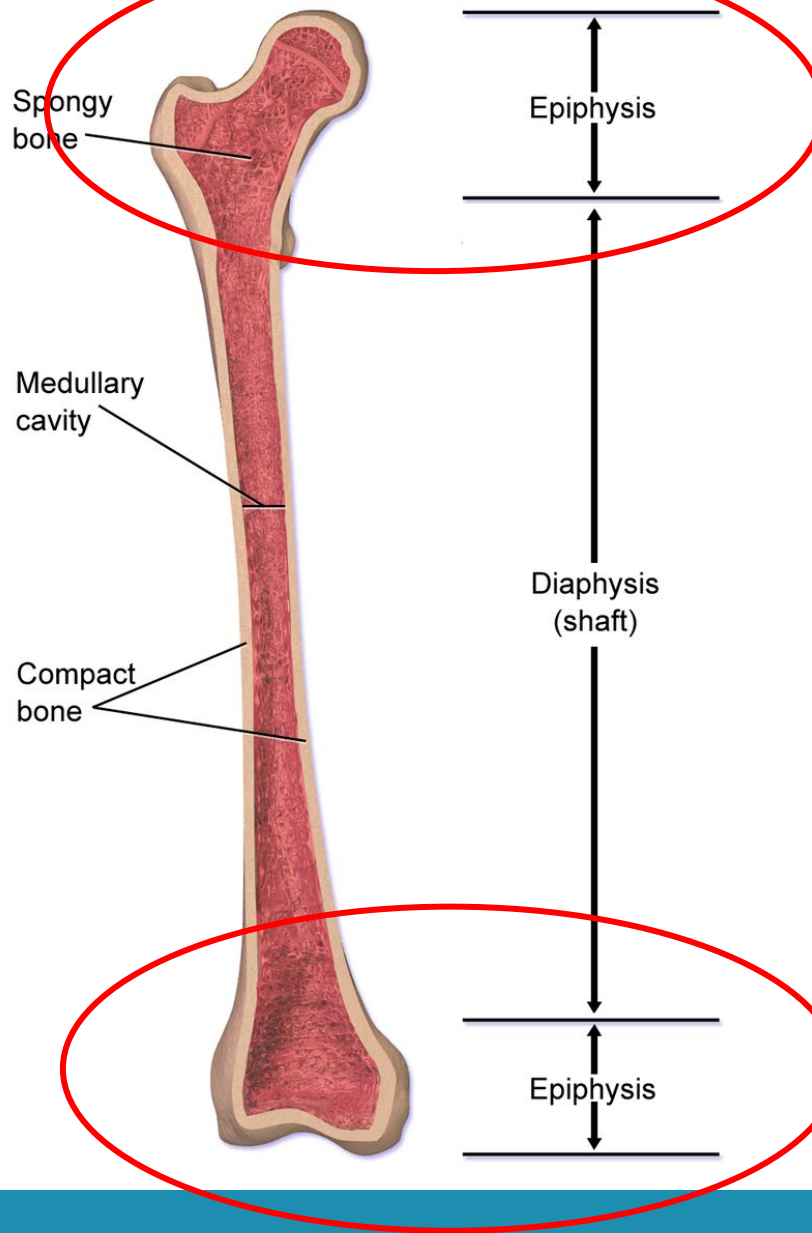


Epiphysis part

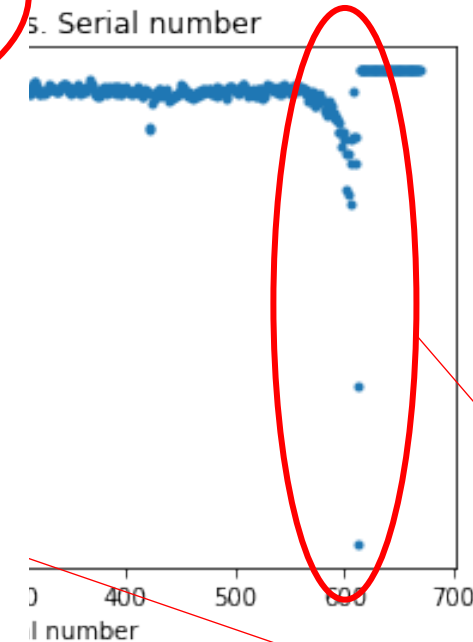
Test set 1

Test set 1
Test set 2

Structure of a Long Bone



planar model

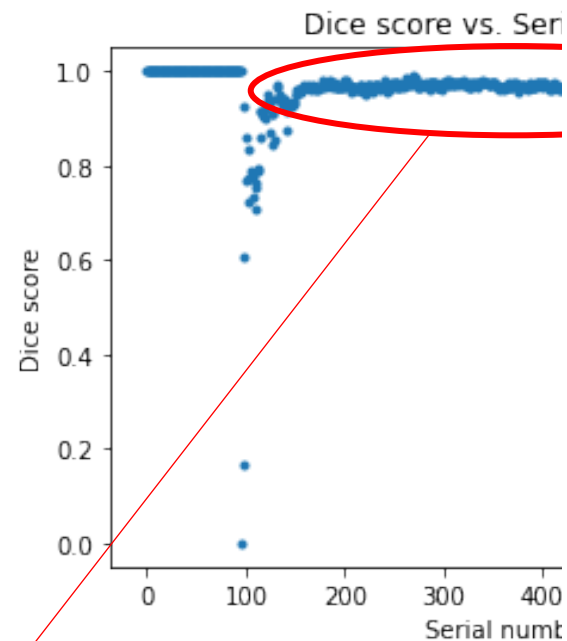


Epiphysis part

Test set 2

metric Dice Score
0.946
0.906

Experiment 2: 50-lay



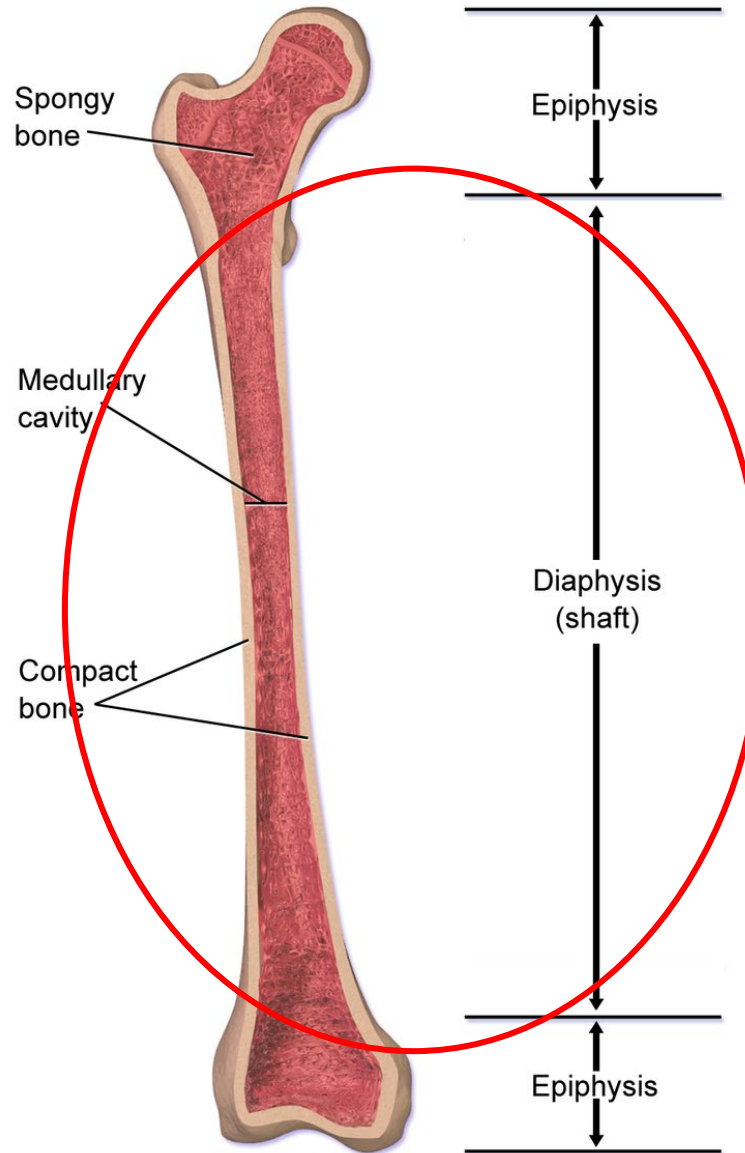
Diaphysis part

Test set 1

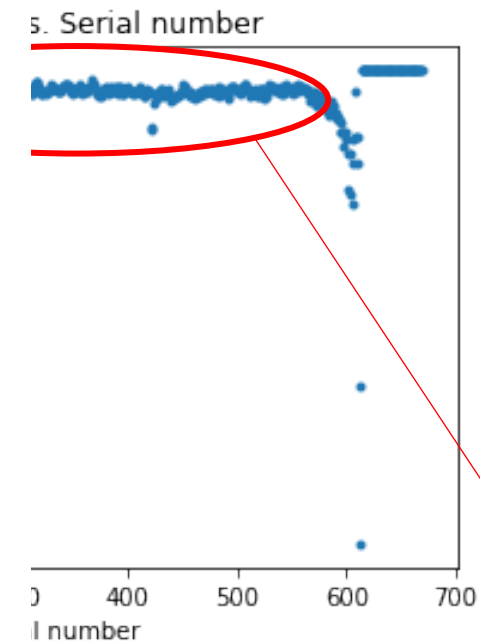
Test set 1

Test set 2

Structure of a Long Bone



planar model



Diaphysis part

Test set 2

metric Dice Score

0.946

0.906



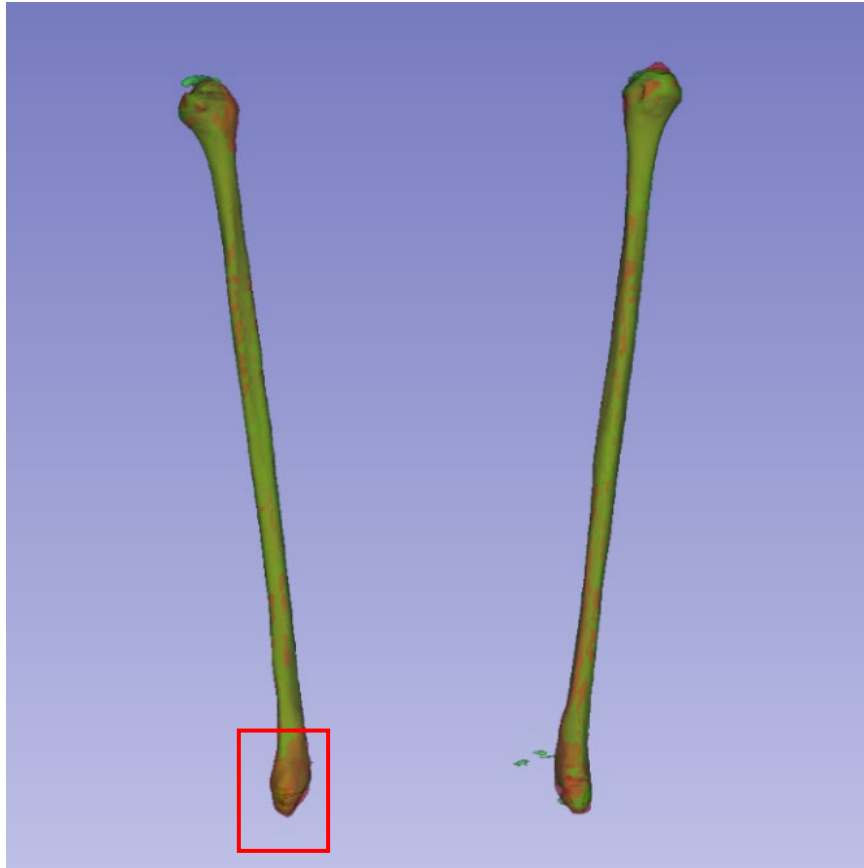
Comparison of Experiment 1 and 2

	Test set 1 average Dice score	Test set 2 average Dice score	Test set 1 volume Dice Score	Test set 2 volume Dice Score
34-layer Res-Net	0.863	0.859	0.920	0.917
50-layer Res-Net	0.952 ↑	0.884 ↑	0.946 ↑	0.906 ↓

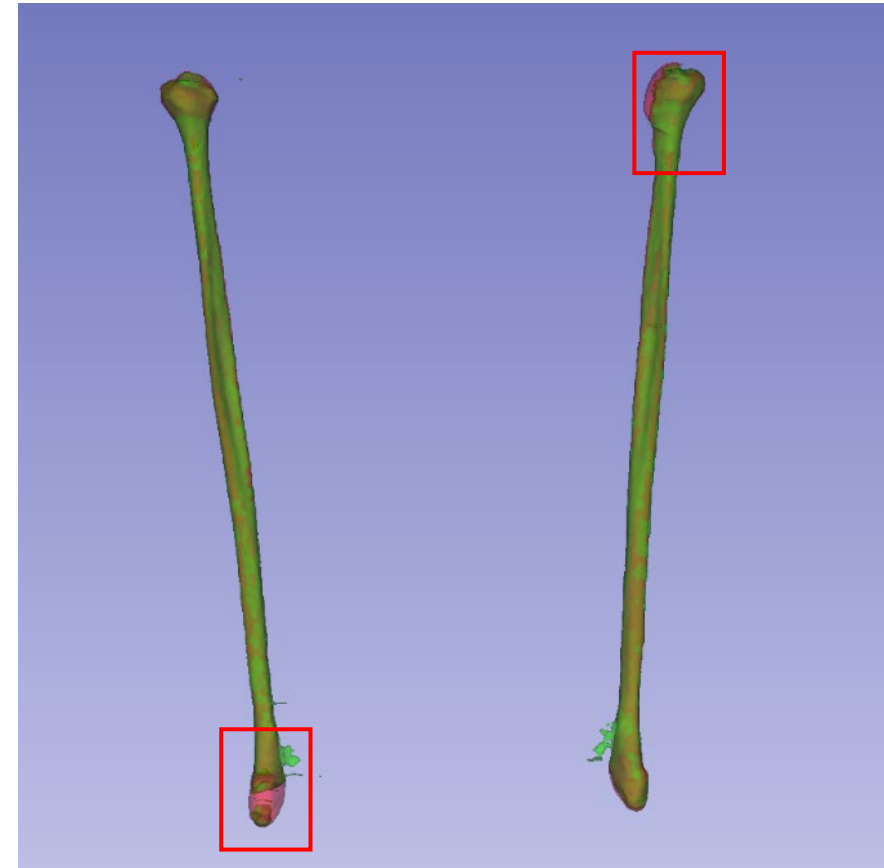
→ Adopt **50-layer Res-Net** structure for single –planar segmentation model



Discussion for Experiment 2: 50-layer Res-Net in single-planar model



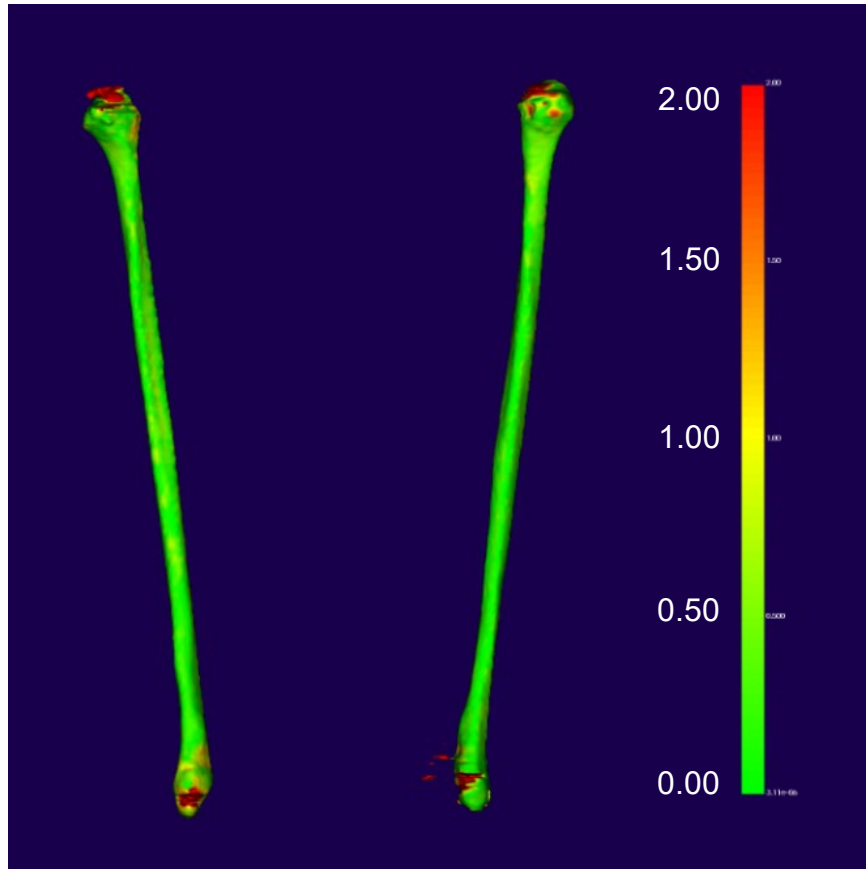
Test set 1



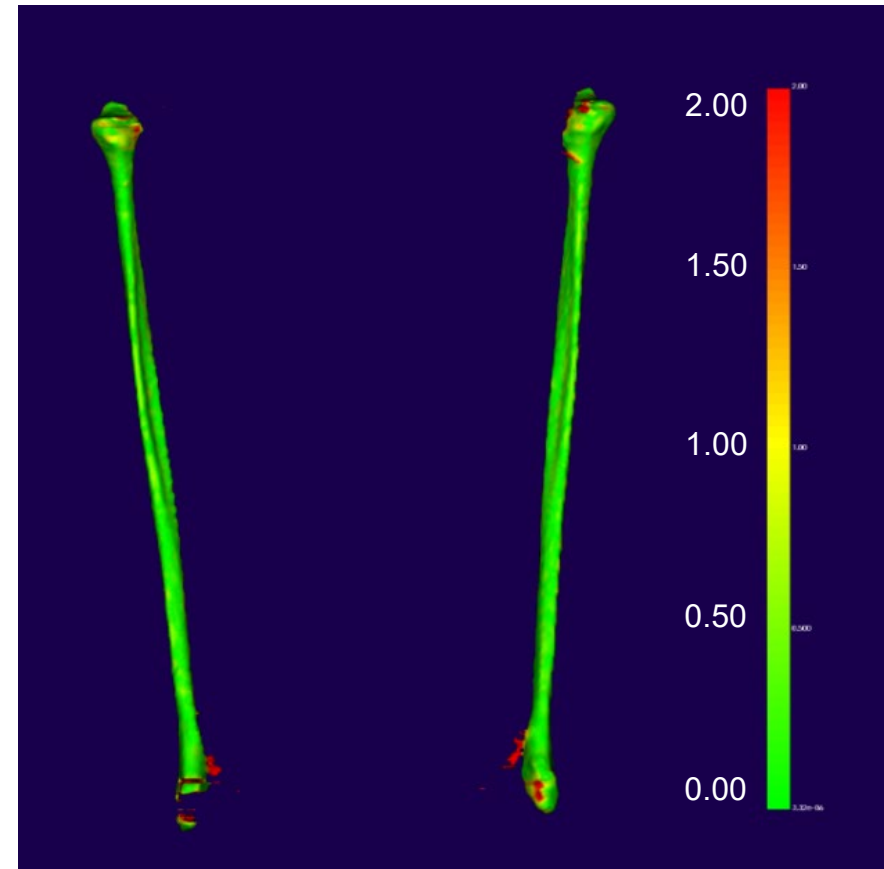
Test set 2



Discussion for Experiment 2



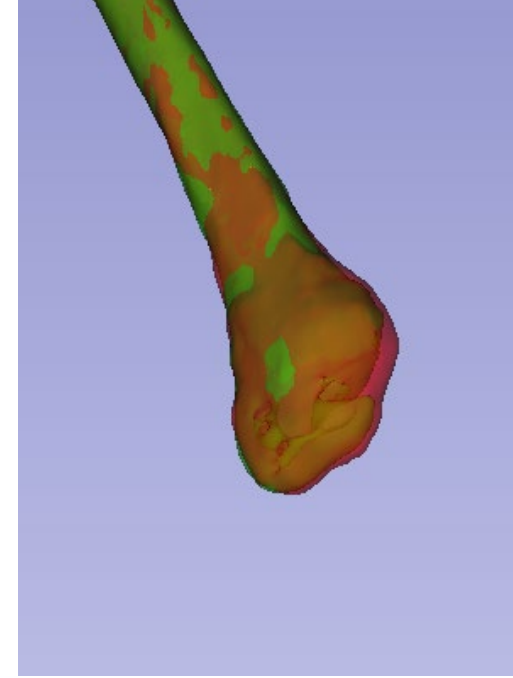
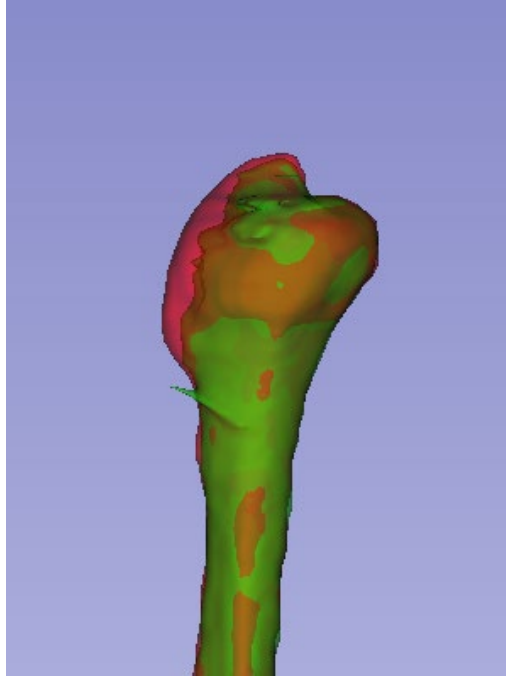
Test set 1



Test set 2



Two method to improve epiphysis segmentation



- Information combination
- Data augmentation



3D Network to 2.5D Network

3D network (3D Unet)

(Regardless of computing and VRAM)

- Combine the information between image slices
- Ensure the continuity
- Better than 2D network

Drawbacks for fibula segmentation

- **limitation of the VRAM** (800x384x384)
- Cannot take the entire 3D metric as input
- Crop to **a series of 3D patches**
- Network is difficult to learning the **overall structure**

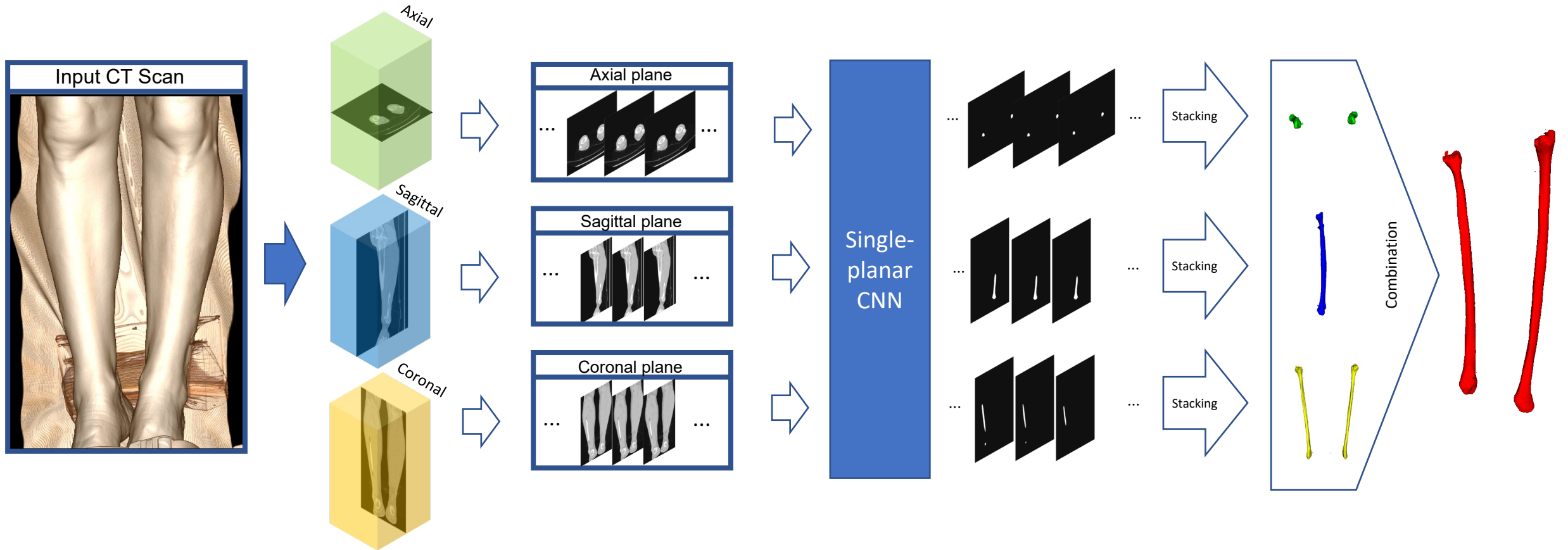
Multi-planar combination (2.5D Network)



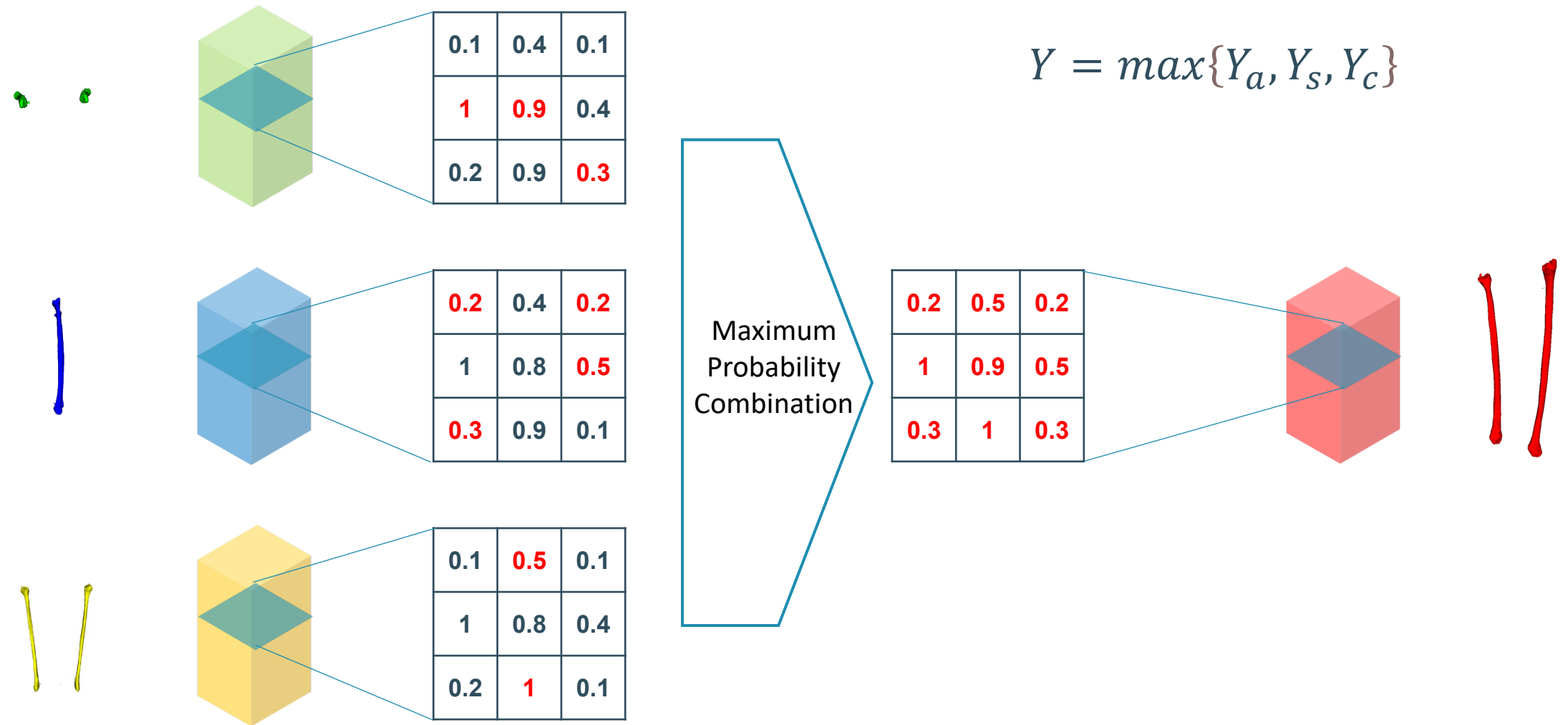
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Multi-planar segmentation model



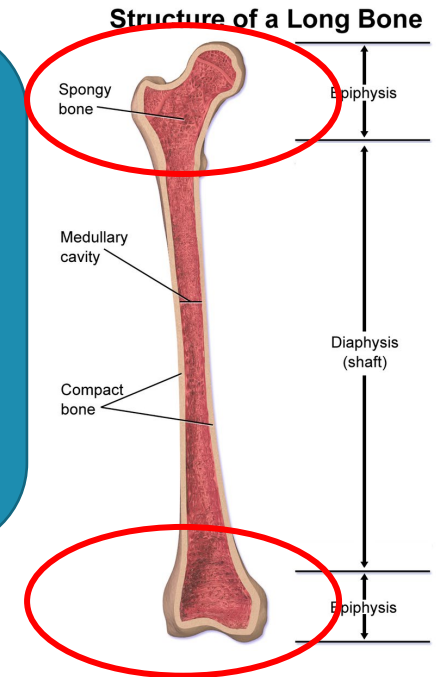
Maximum Probability Combination



Cropping the region of interest (CROI) method

- The input image for single-planar segmentation model is 384x384
- Different resolution (384x734, 384x670...)

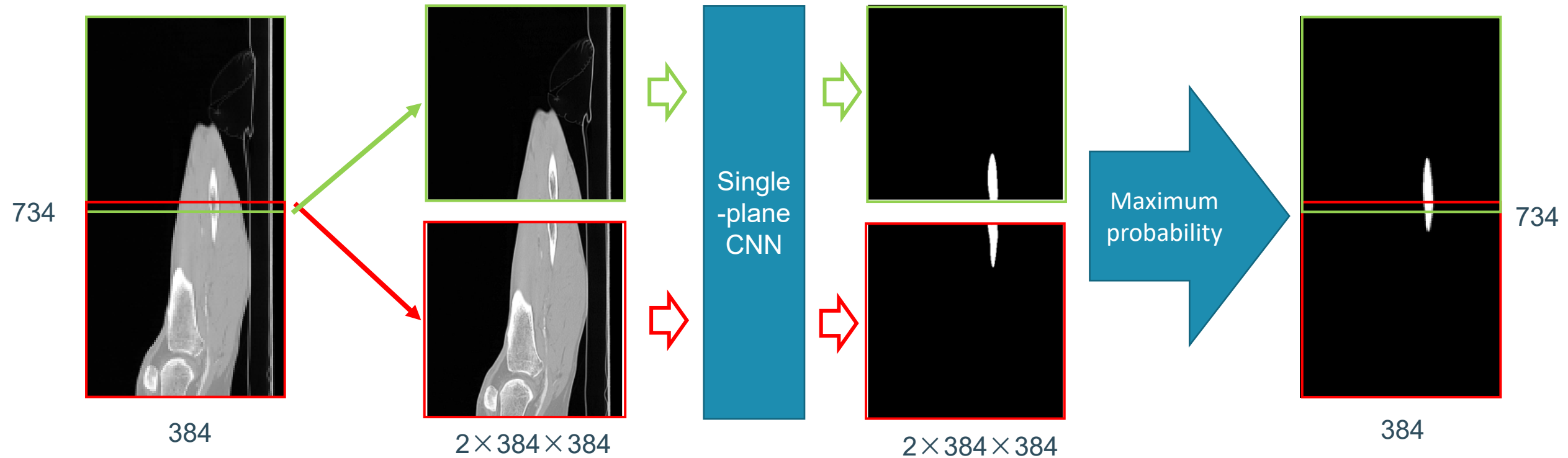
- Predicted performance of **epiphysis part** is worse than the diaphysis part
- Focus on data at **epiphysis part**



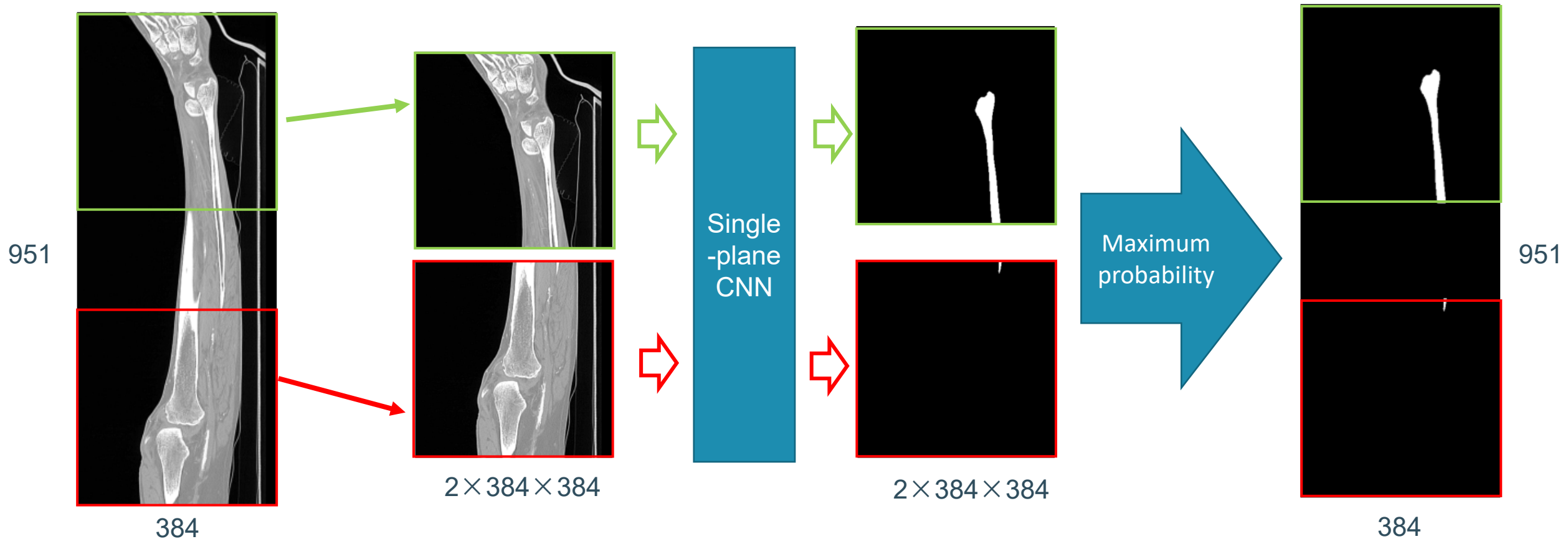
Cropping the region of interest (CROI)



Cropping the region of interest (CROI) method (For sagittal and coronal)



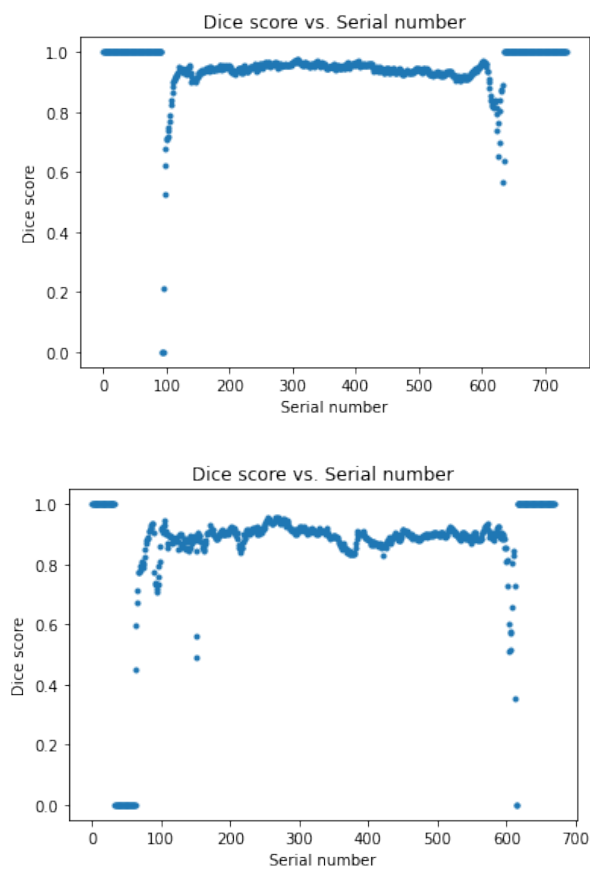
Cropping the region of interest (CROI) method (For sagittal and coronal)





Experiment 3: Multi-planar segmentation model

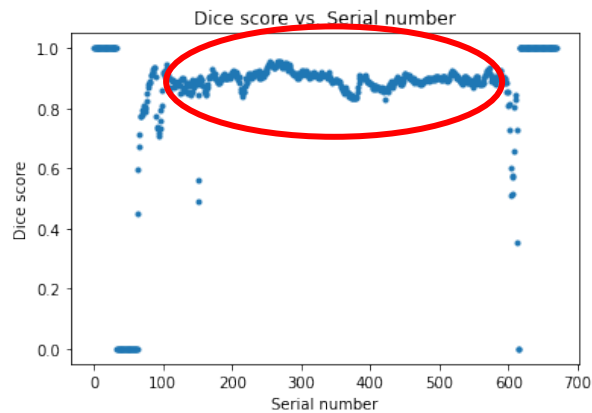
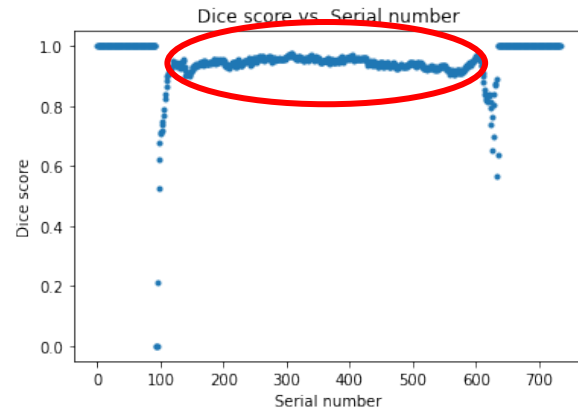
Combination result



	Test set 1 average Dice score	Test set 2 average Dice score	Test set 1 volume Dice Score	Test set 2 volume Dice Score
Axial plane	0.956	0.938	0.947	0.923
Sagittal plane	0.942	0.938	0.908	0.909
Coronal plane	0.946	0.945	0.918	0.896
Combination result	0.945 ▼	0.856 ▼	0.932 ▼	0.883 ▼

Experiment 3: Multi-planar segmentation model

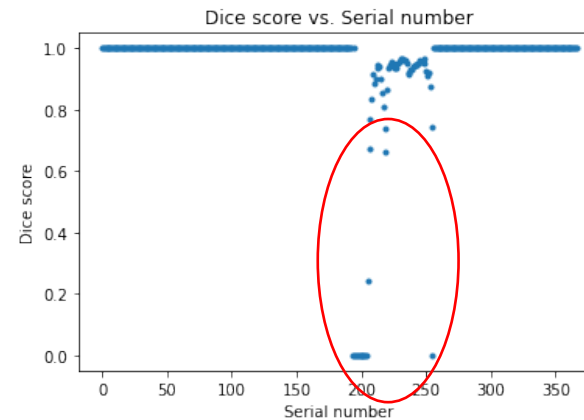
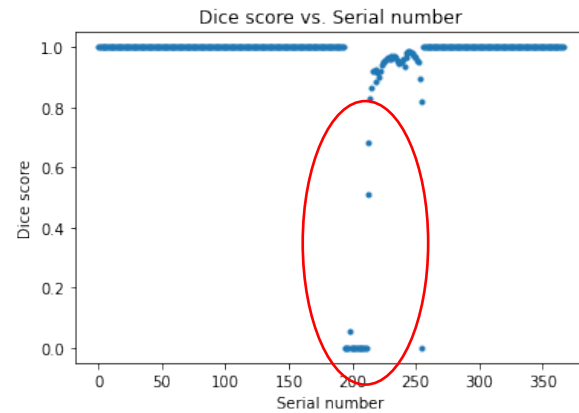
Combination result



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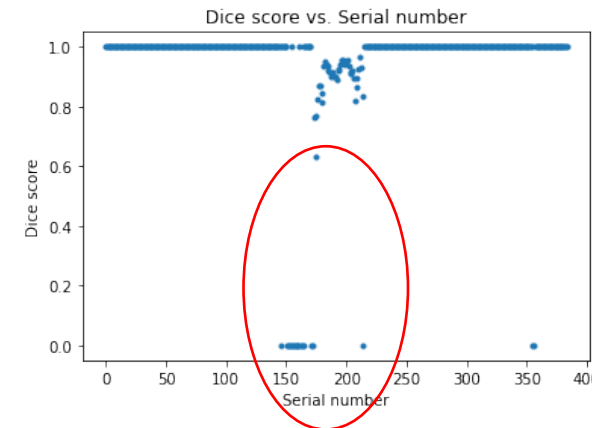
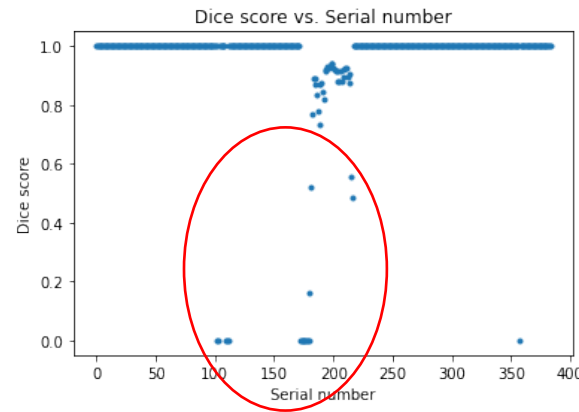
Experiment 3: Multi-planar segmentation model

Coronal plane subnetwork



Test set 1

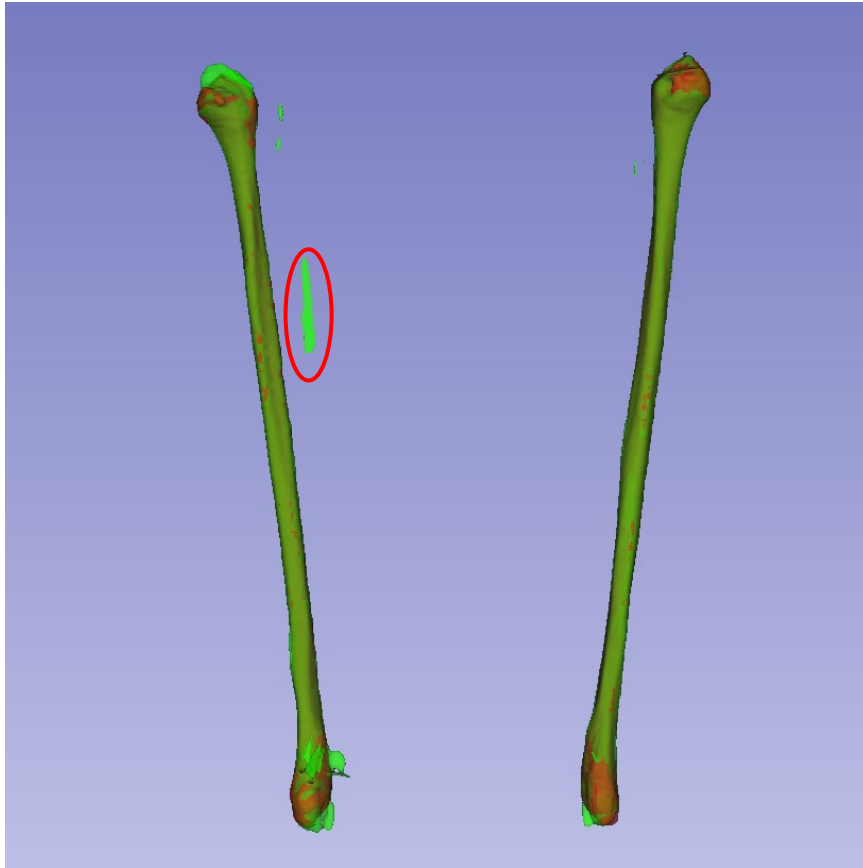
- Too few useful parts
- Result is poor



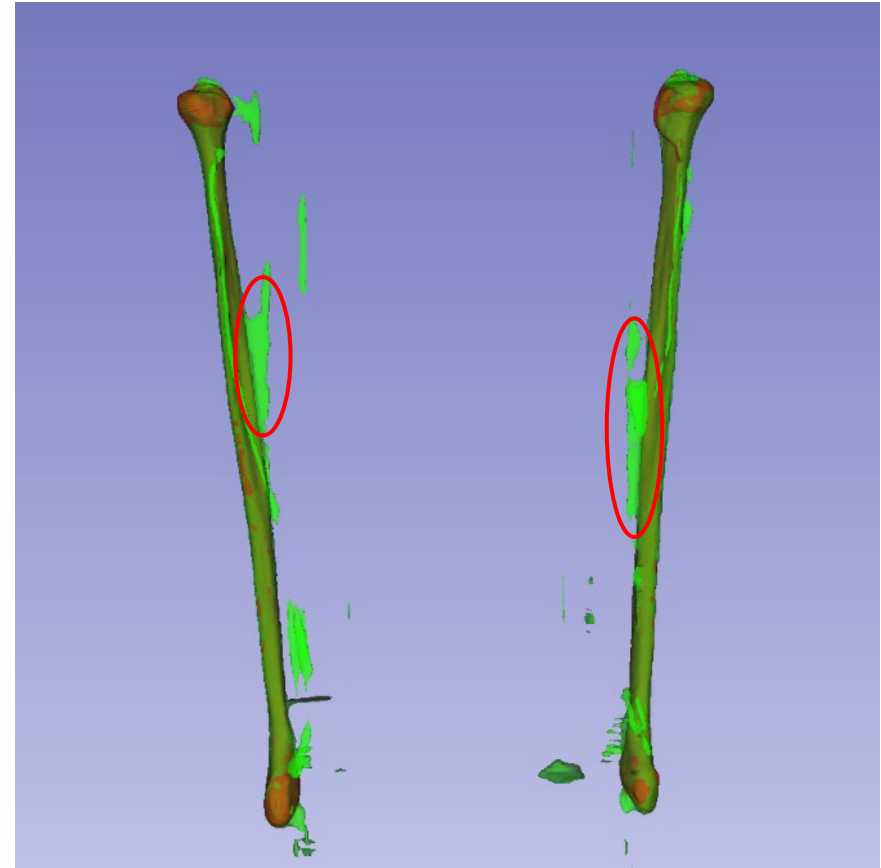
Test set 2



Discussion for Experiment 3: Multi-planar segmentation model



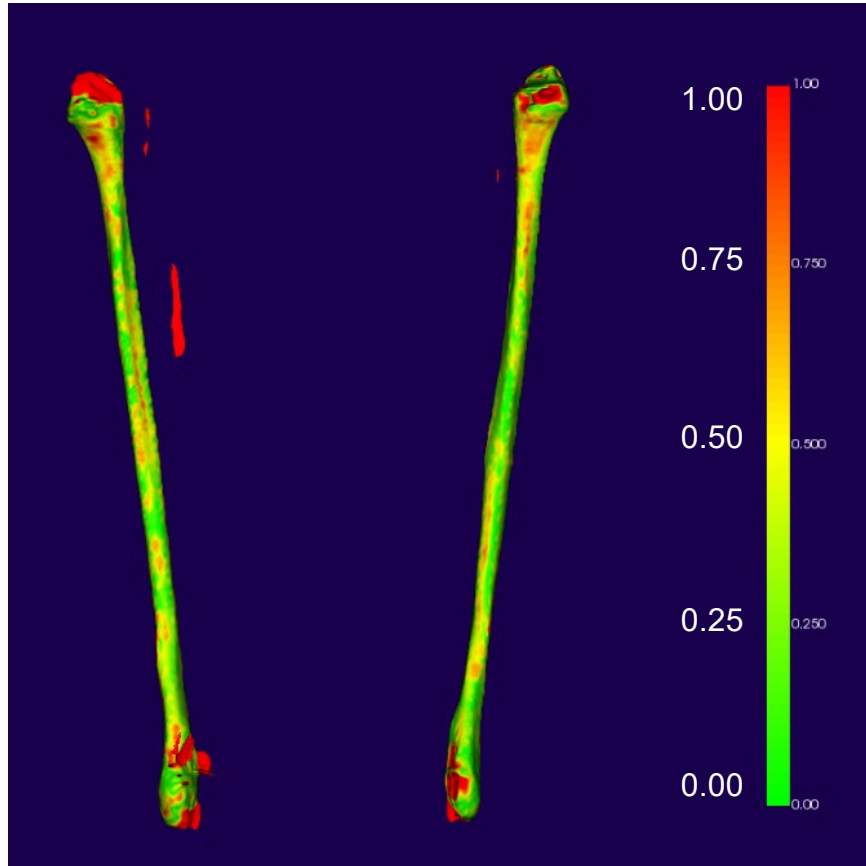
Test set 1



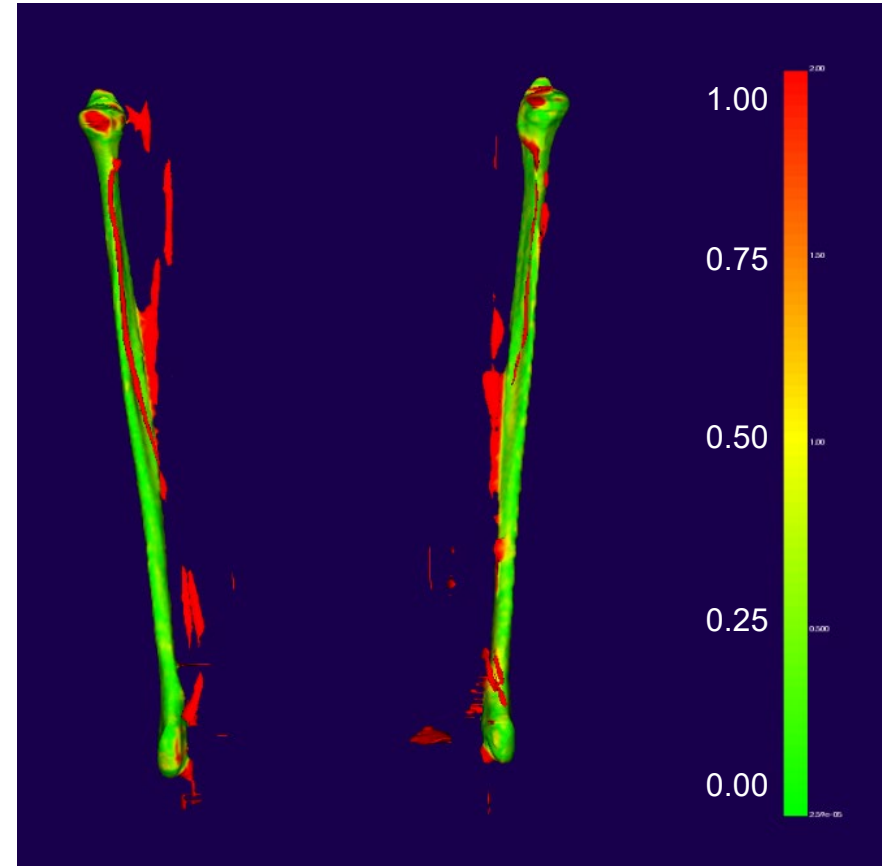
Test set 2



Discussion for Experiment 3



Test set 1



Test set 2



Outline

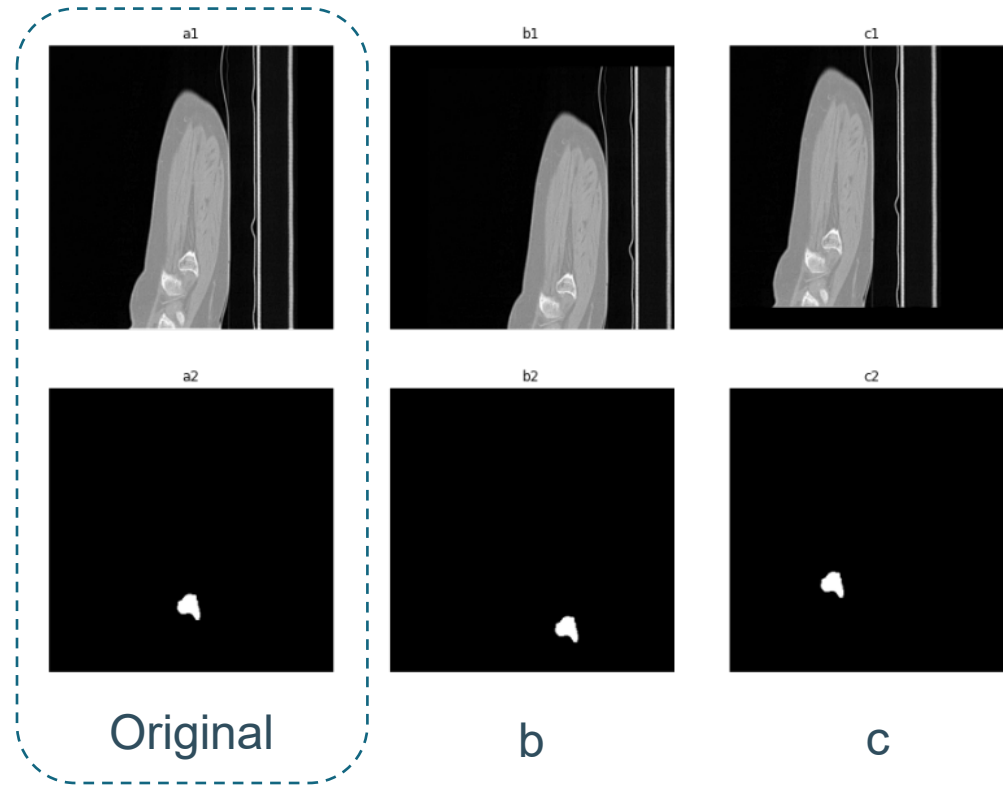
1. Motivation
2. Dataset & Data preprocessing
3. Methodology & Experiments
 - a. Evaluation criterion
 - b. Single-planar segmentation model
 - c. Multi-planar segmentation model
 - d. Multi-planar segmentation model with data augmentation
4. Conclusion



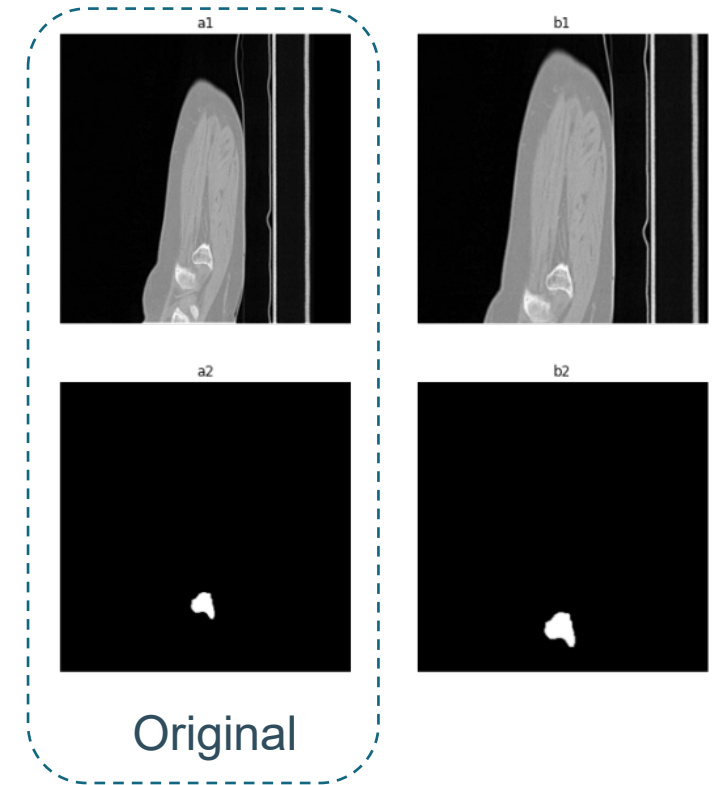
Data Augmentation

Images

Ground Truth



- **Translation:** b. 50 to the right and 30 downwards
c. 50 to the left and 30 upwards

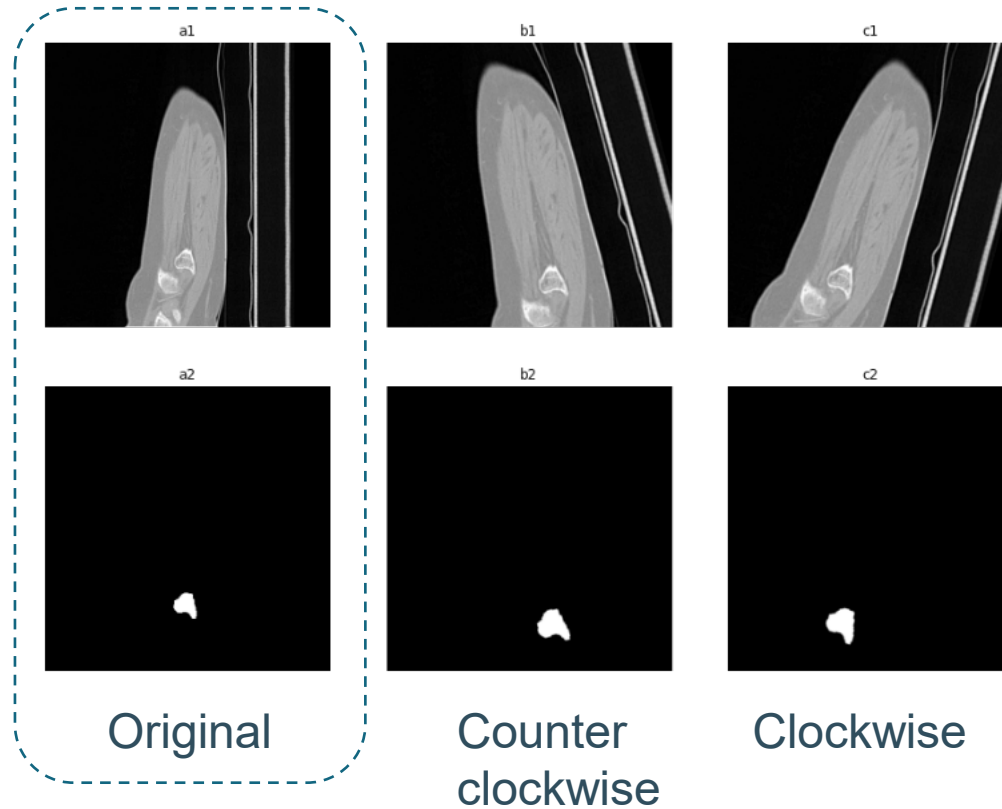


- **Scale:** 1.3 times.

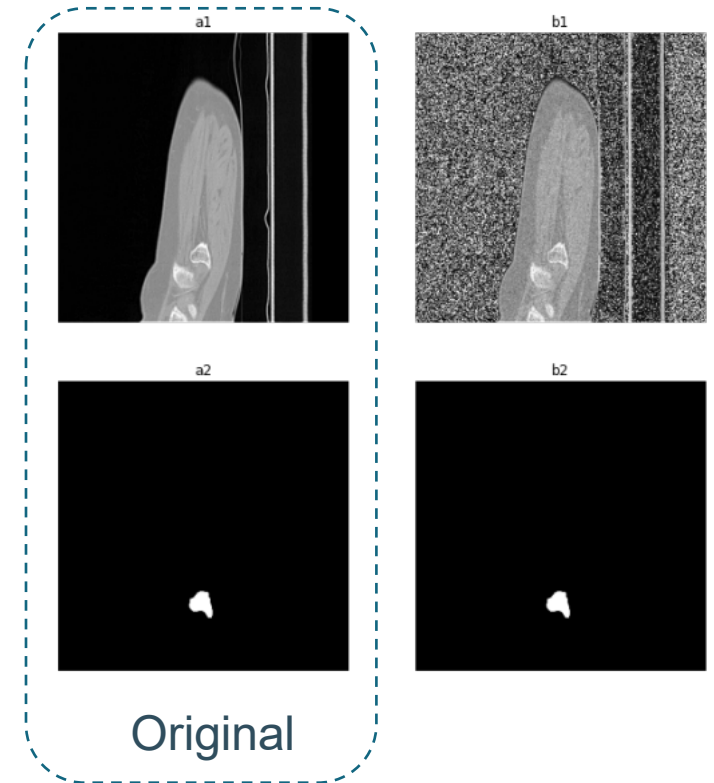
Data Augmentation

Images

Ground Truth

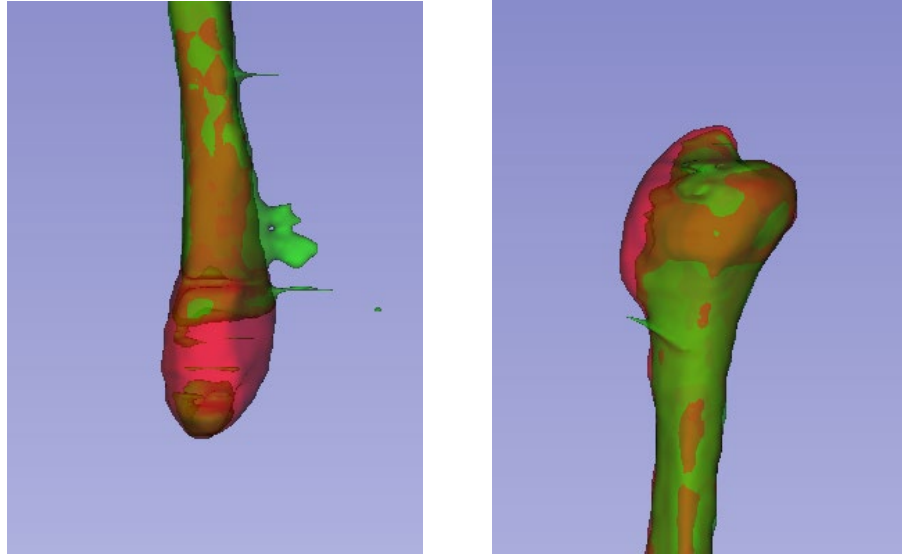


- **Rotate:** 15 degrees rotate, and 1.3 times enlarged



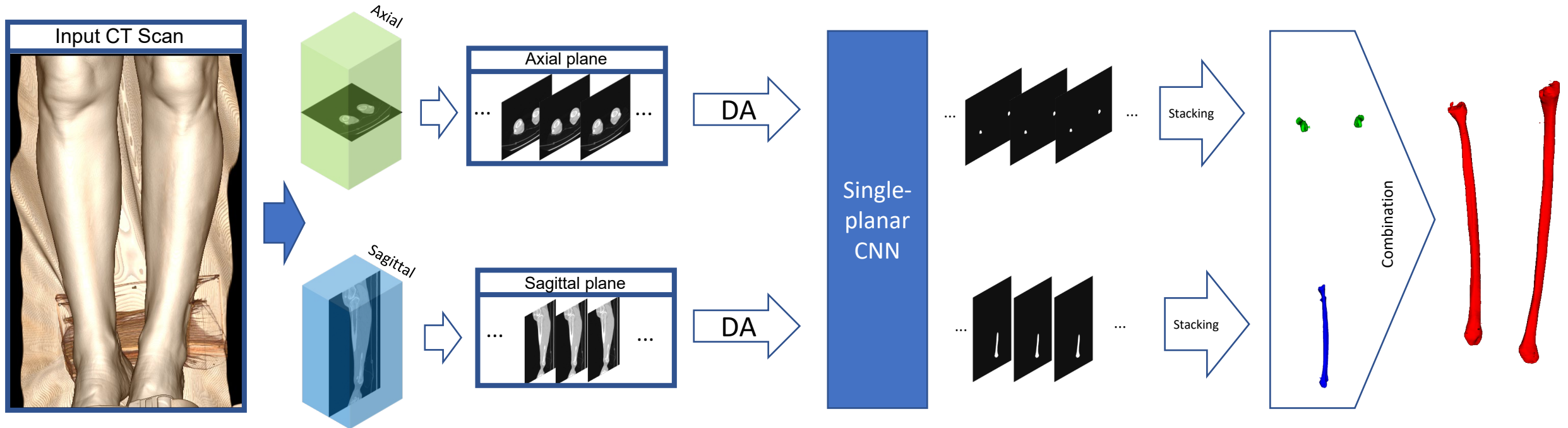
- **Gaussian noise:** an average of zero and a standard deviation of 1

Data Augmentation



- Axial:
 - Select **epiphysis part manually** from train set
 - 2160 to 10,800
- Sagittal:
 - Select slices included the **fibula bone**
 - 2830 to 14,150
- Add to original train set

Final segmentation model

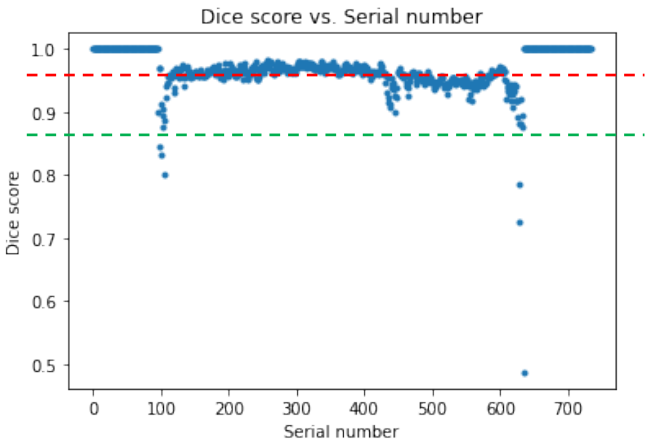


- Data augmentation
- Multi-planar segmentation only with axial plane and sagittal plane.

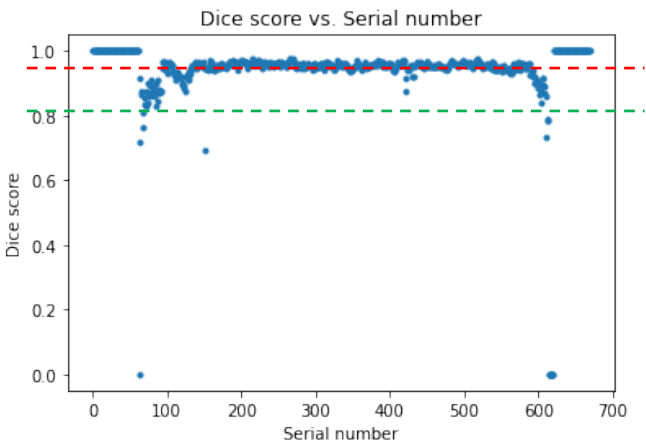


Experiment 4

Axial segmentation result on axial slices after data augmentation



Test set 1



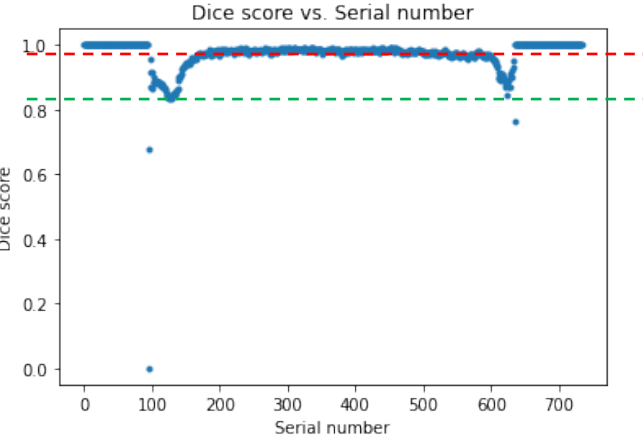
Test set 2

	Test set 1 average Dice score	Test set 2 average Dice score	Test set 1 volume Dice Score	Test set 2 volume Dice Score
Axial plane	0.956	0.938	0.947	0.923
Axial plane After Data Augmentation	0.967	0.943	0.957	0.944
Sagittal plane	0.942	0.938	0.908	0.909
Sagittal plane After Data Augmentation	0.961	0.972	0.955	0.964
Data Augmentation Combination	0.967	0.944	0.957	0.946

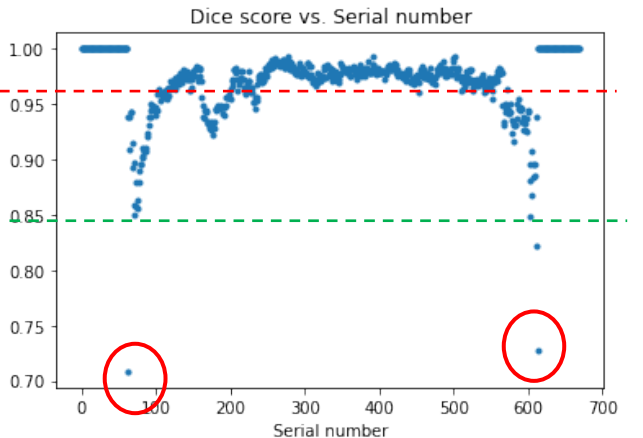


Experiment 4

Sagittal segmentation result on axial slices after data augmentation



Test set 1



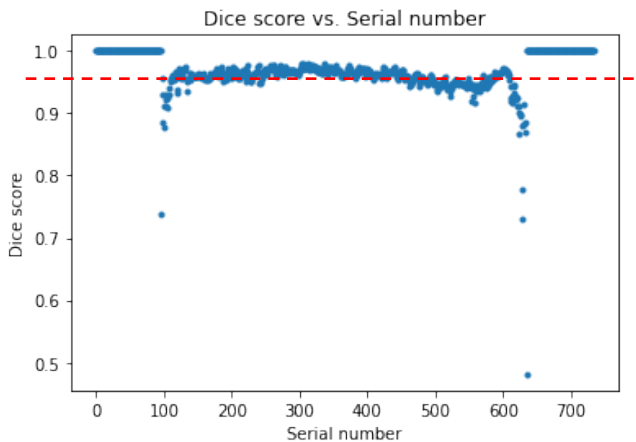
Test set 2

	Test set 1 average Dice score	Test set 2 average Dice score	Test set 1 volume Dice Score	Test set 2 volume Dice Score
Axial plane	0.956	0.938	0.947	0.923
Axial plane After Data Augmentation	0.967	0.943	0.957	0.944
Sagittal plane	0.942	0.938	0.908	0.909
Sagittal plane After Data Augmentation	0.961	0.972	0.955	0.964
Data Augmentation Combination	0.967	0.944	0.957	0.946

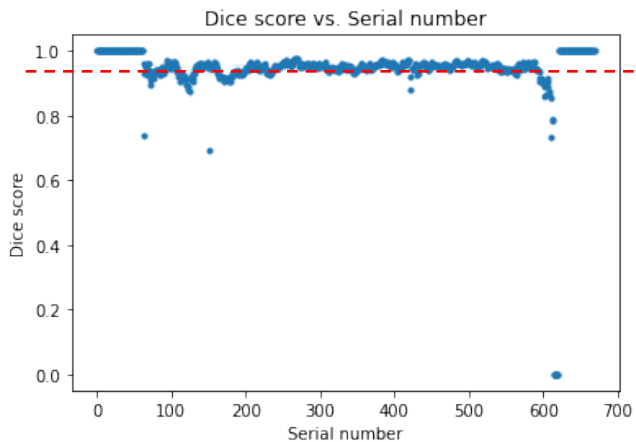


Experiment 4

Combination segmentation result on axial slices after data augmentation



Test set 1

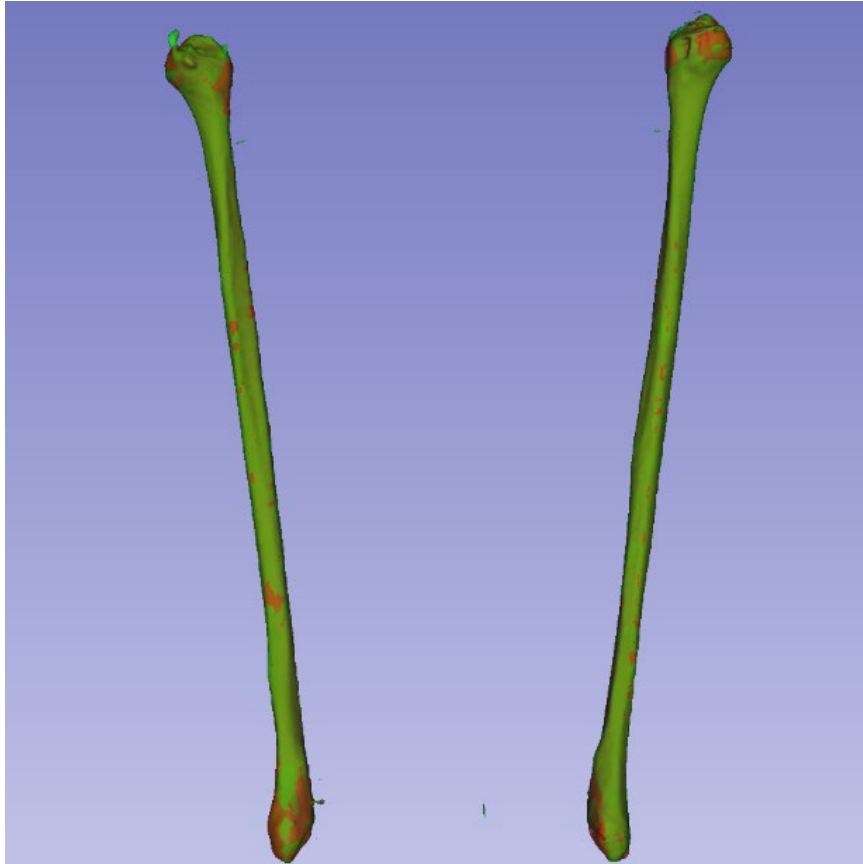


Test set 2

	Test set 1 average Dice score	Test set 2 average Dice score	Test set 1 volume Dice Score	Test set 2 volume Dice Score
Axial plane	0.956	0.938	0.947	0.923
Axial plane After Data Augmentation	0.967	0.943	0.957	0.944
Sagittal plane	0.942	0.938	0.908	0.909
Sagittal plane After Data Augmentation	0.961	0.972	0.955	0.964
Data Augmentation Combination	0.967	0.944	0.957	0.946

Discussion for Experiment 4:

Multi-planar segmentation model with data augmentation



Test set 1

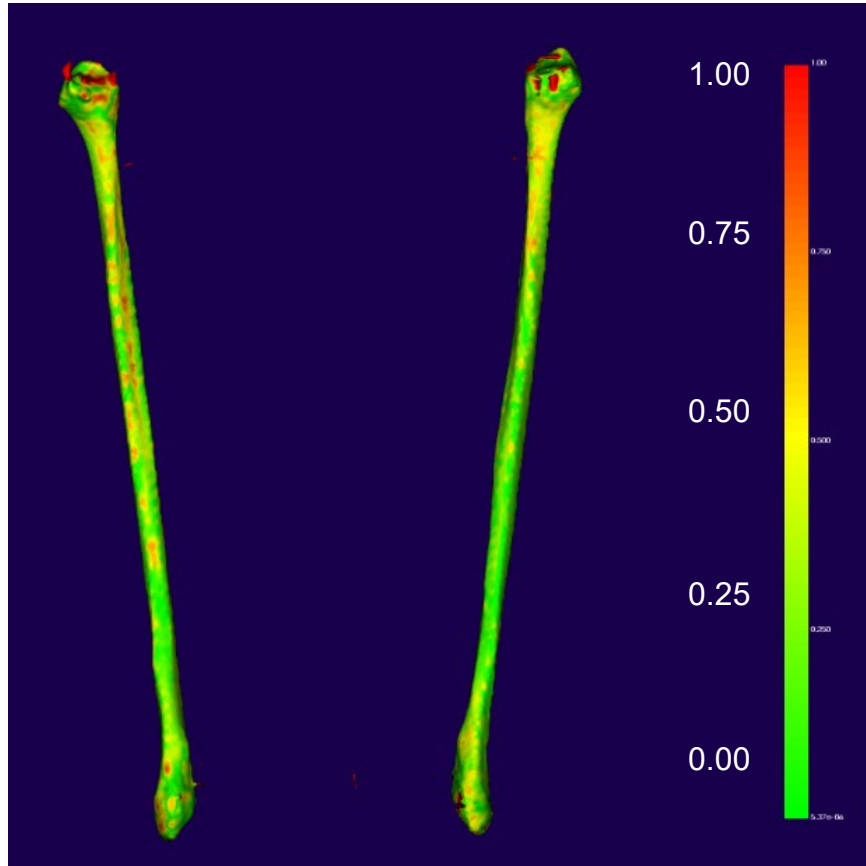


Test set 2

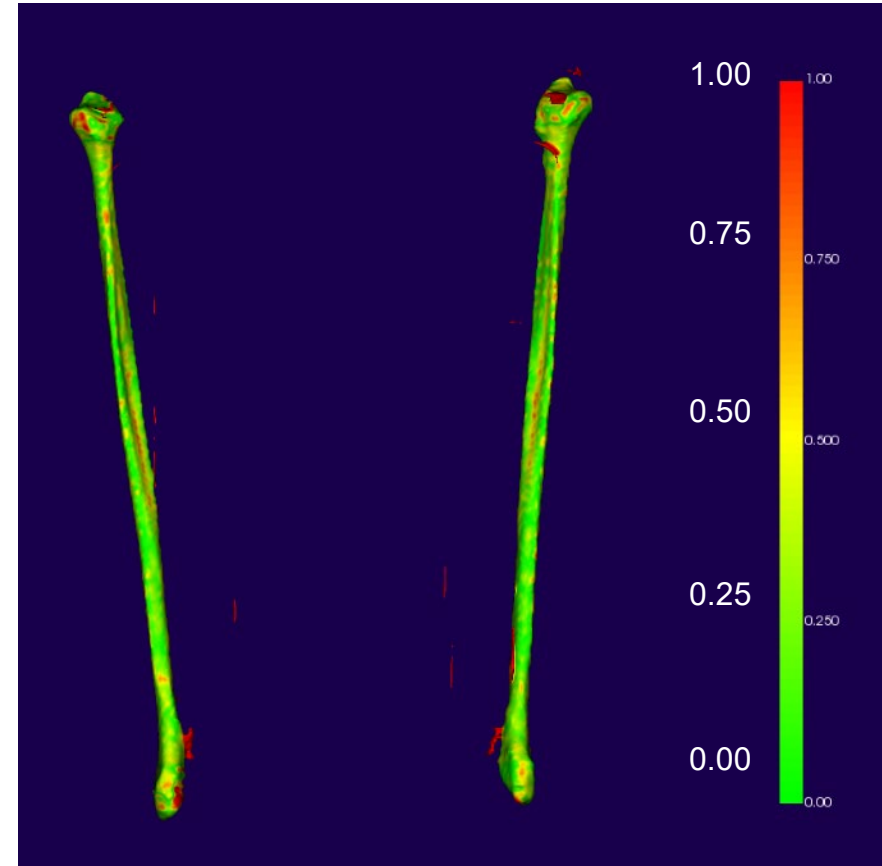


Discussion for Experiment 4:

Multi-planar segmentation model with data augmentation



Test set 1



Test set 2

Outline

1. Motivation
2. Dataset & Data preprocessing
3. Methodology & Experiments
4. Conclusion



Conclusion

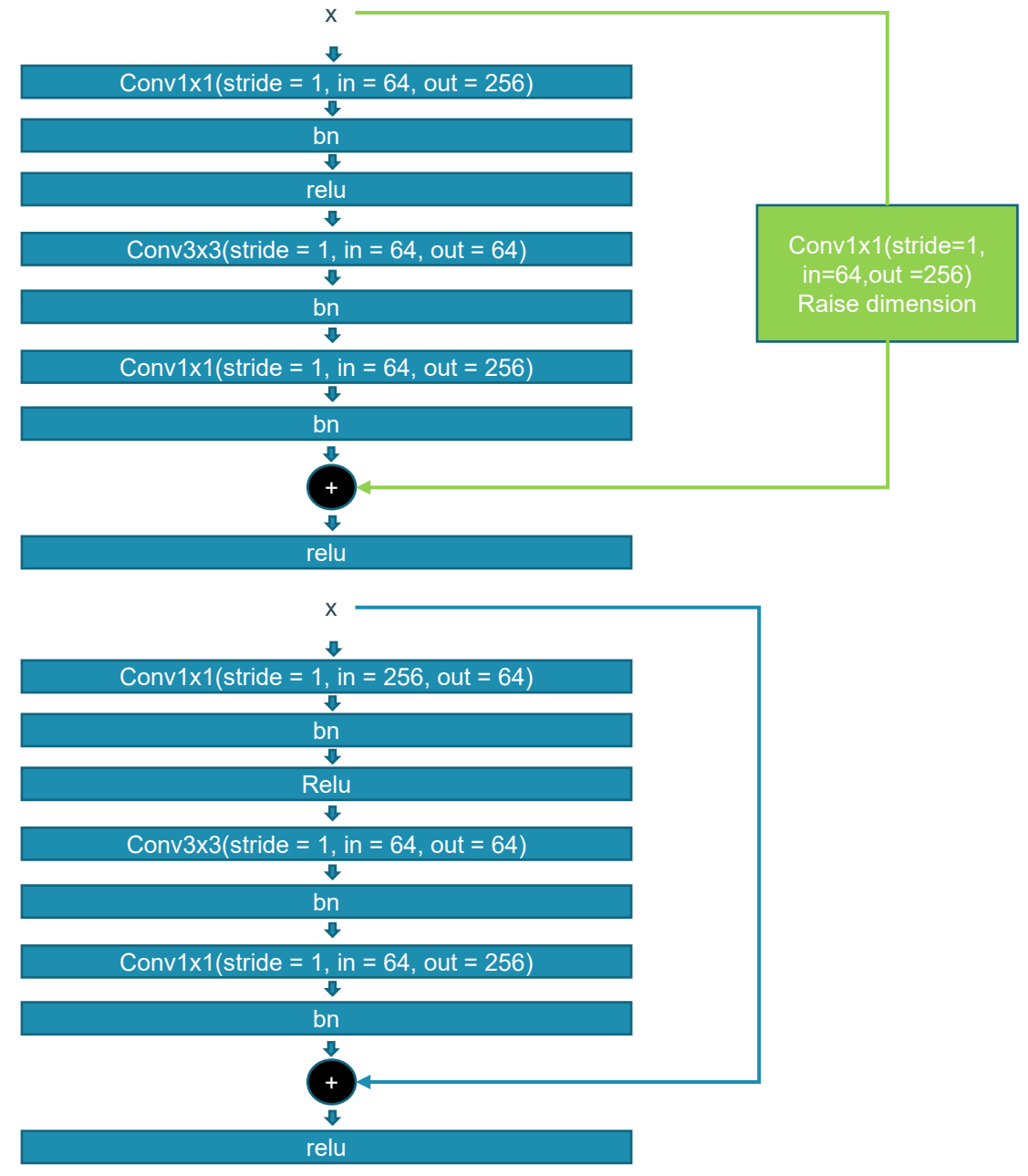
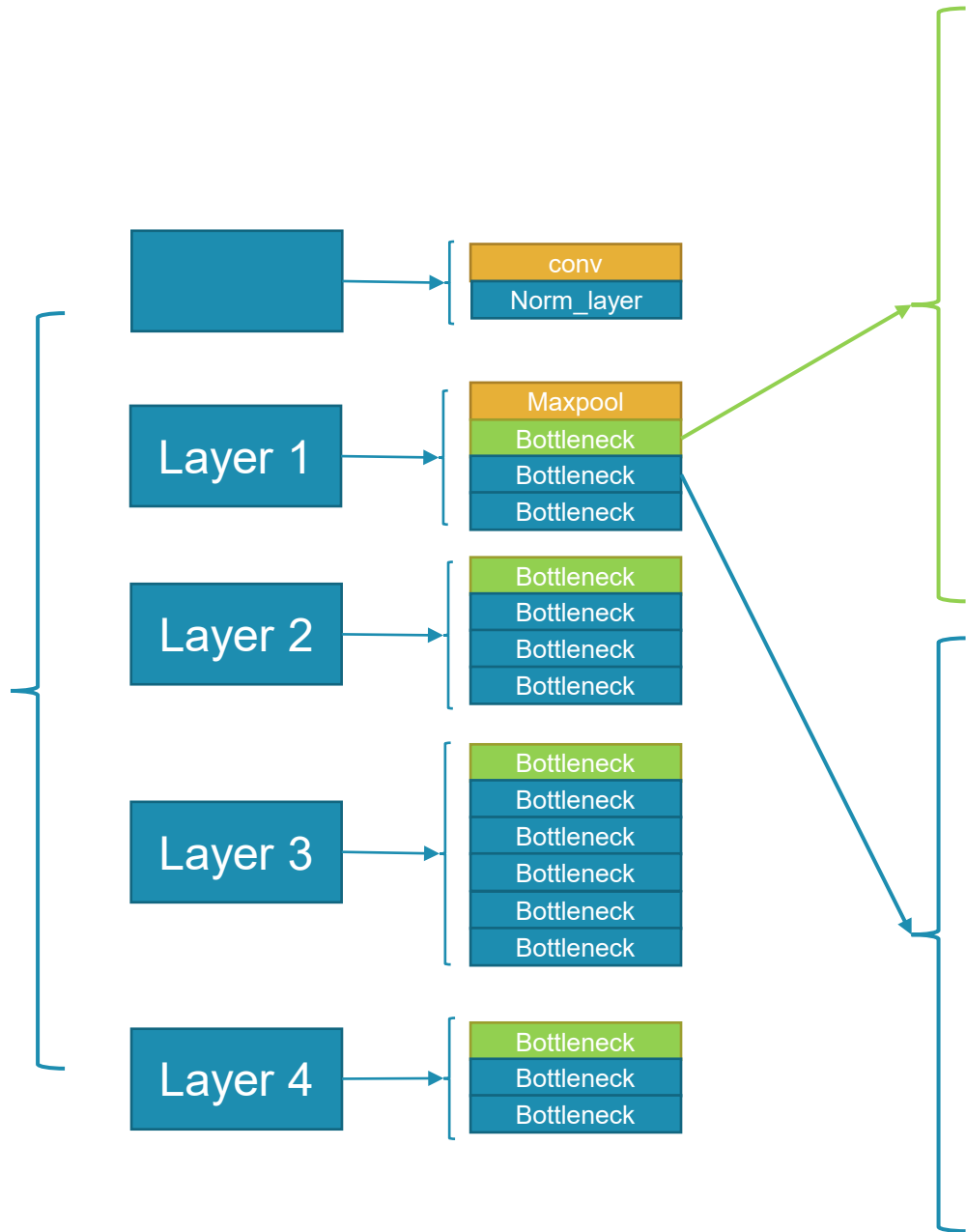
- An **automatic** fibula segmentation neural network
→ proven to be **accurate and efficient**
- The **multi-planar segmentation model**
→ increased the segmentation accuracy of epiphysis
→ decreased the segmentation accuracy of diaphysis
- The **data augmentation**
→ increased segmentation accuracy for single-planar segmentation and multi-planar segmentation
- Applicable for clinical medical fibula segmentation
→ Dice score higher than **0.95**



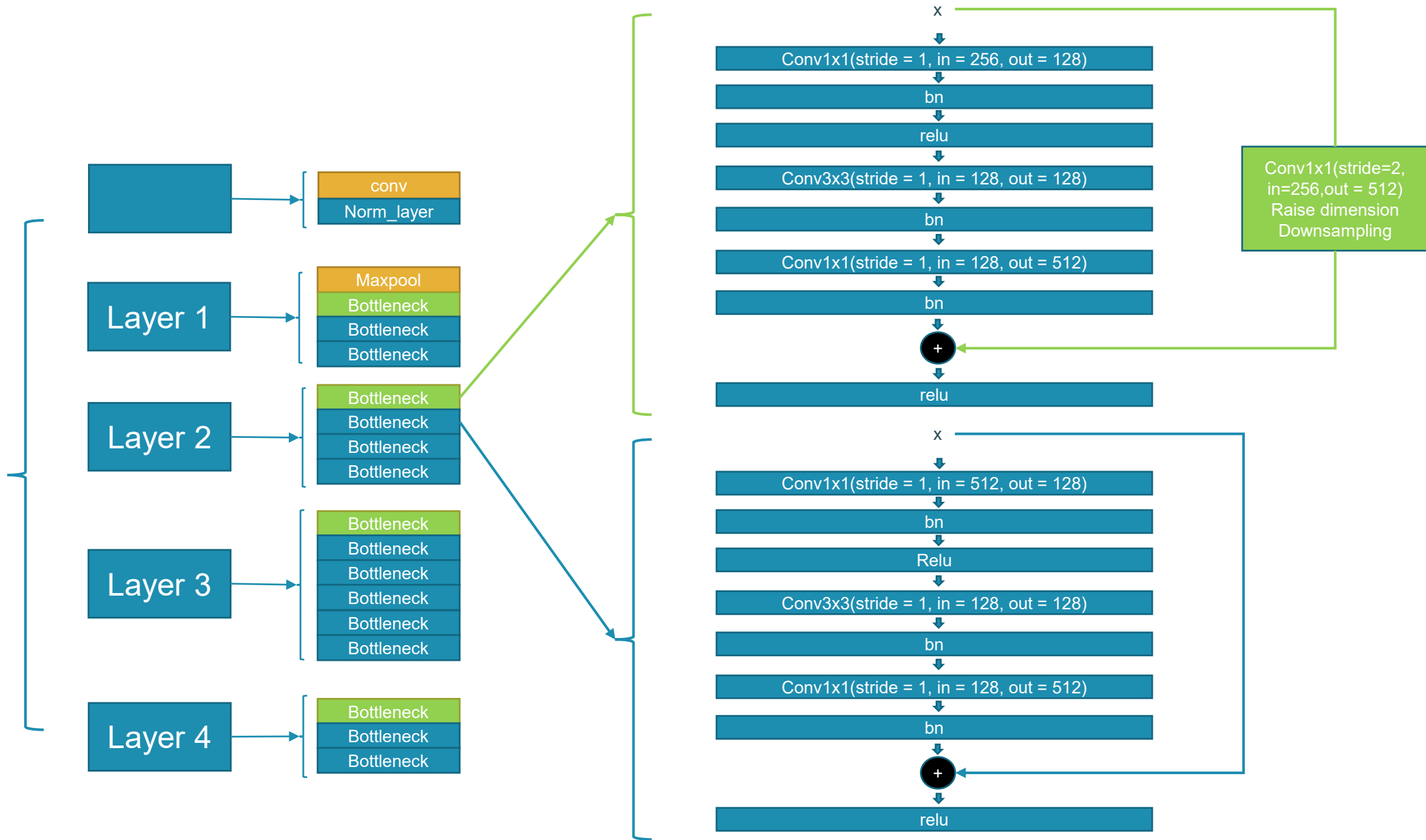
 Thank You
For Your Attention



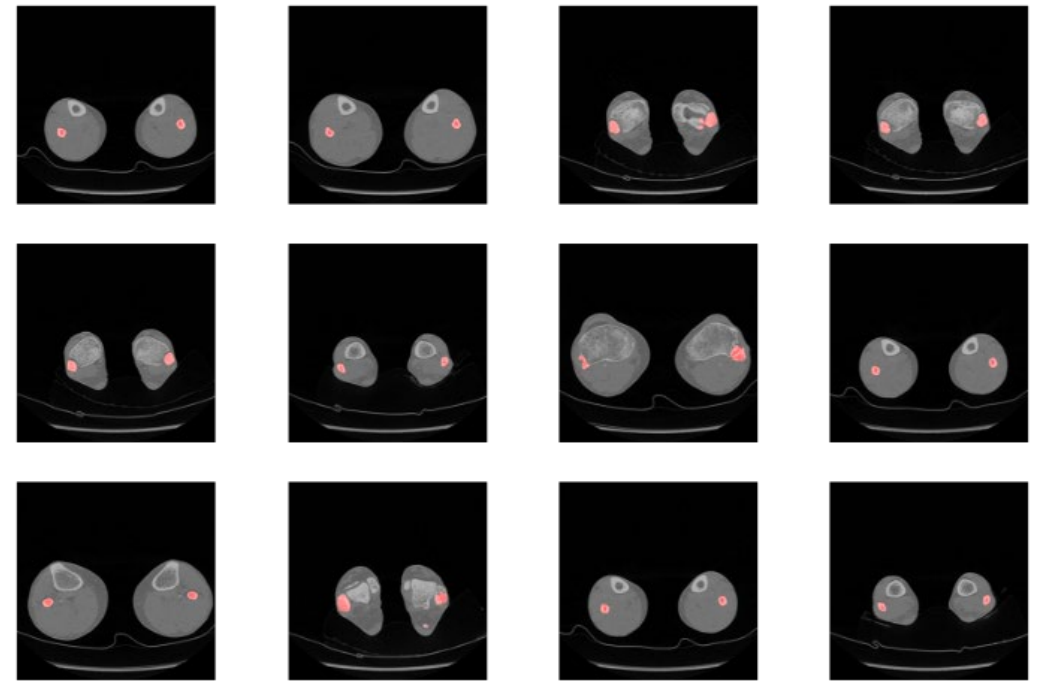
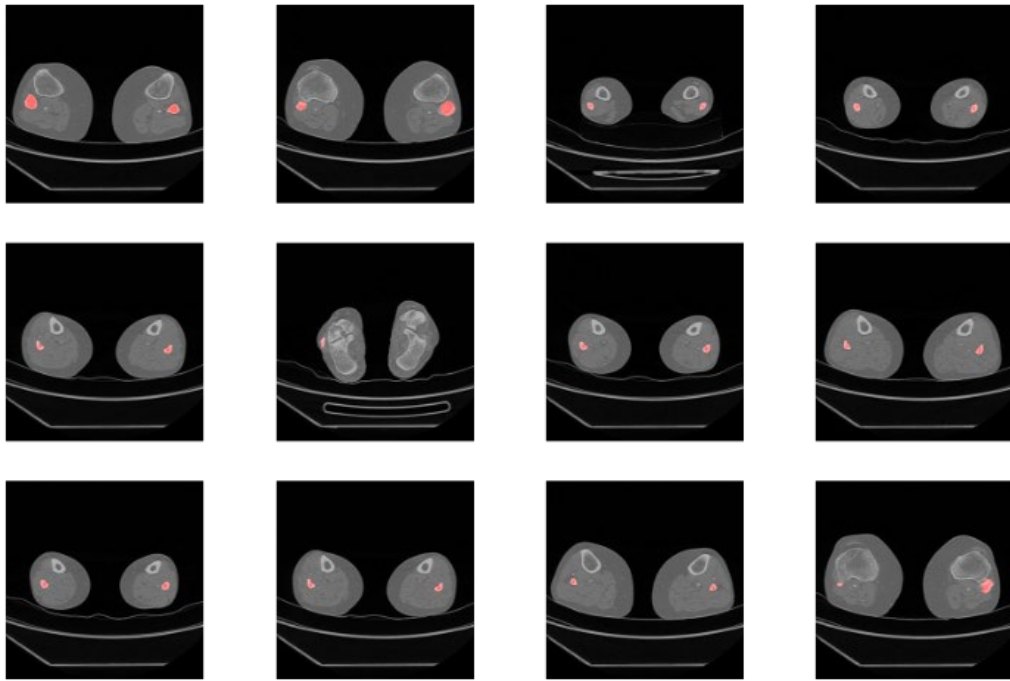
Resnet50

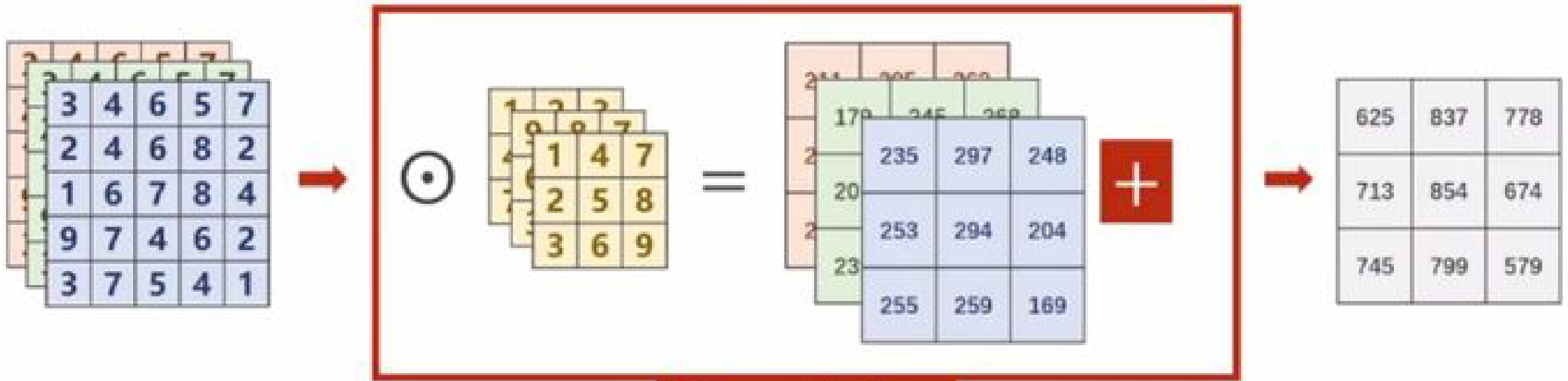


Resnet50

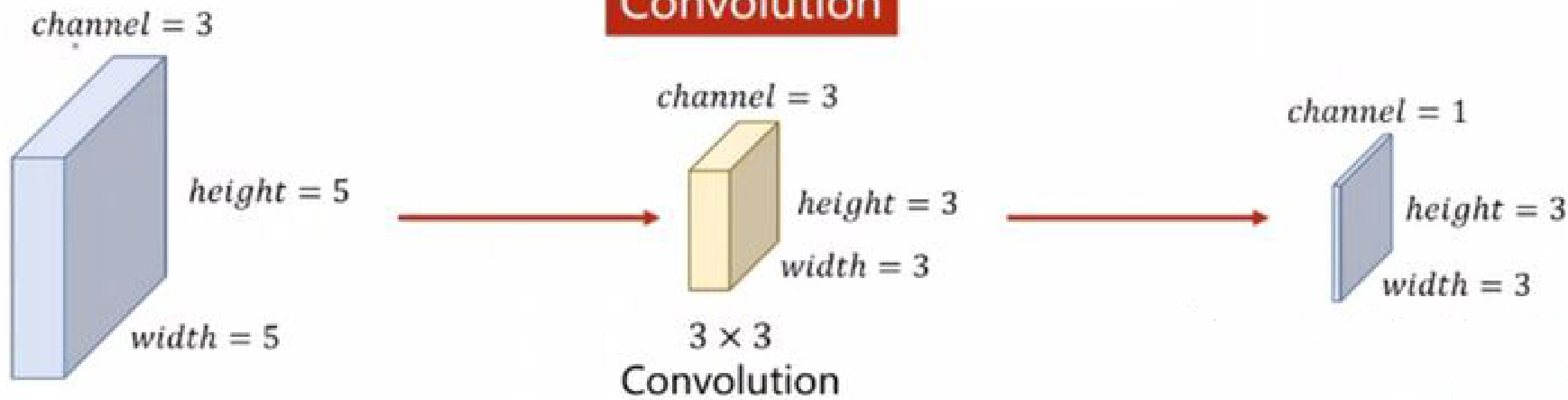


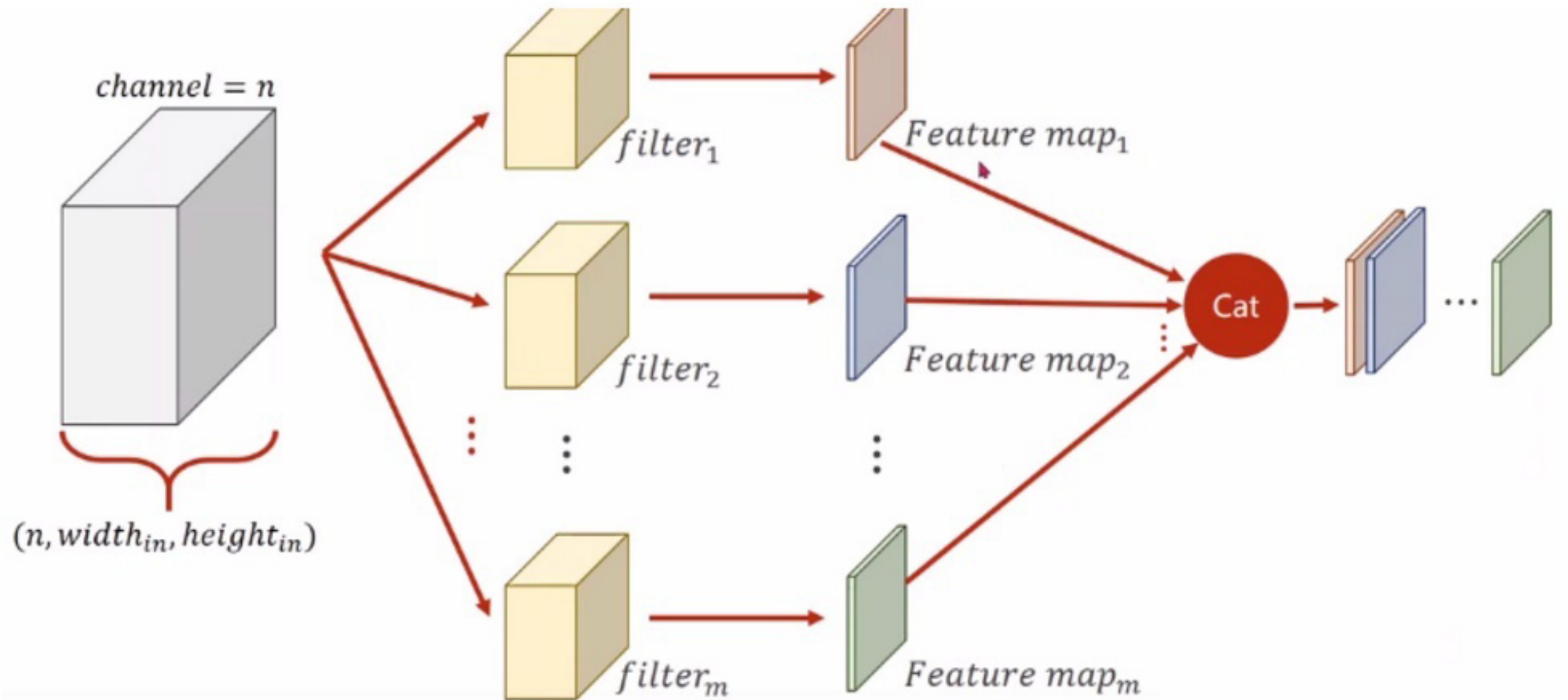
Results of Experiment 2





Convolution

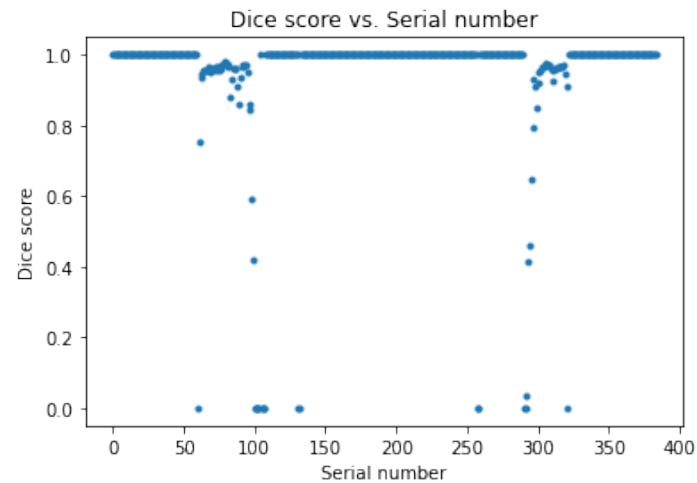
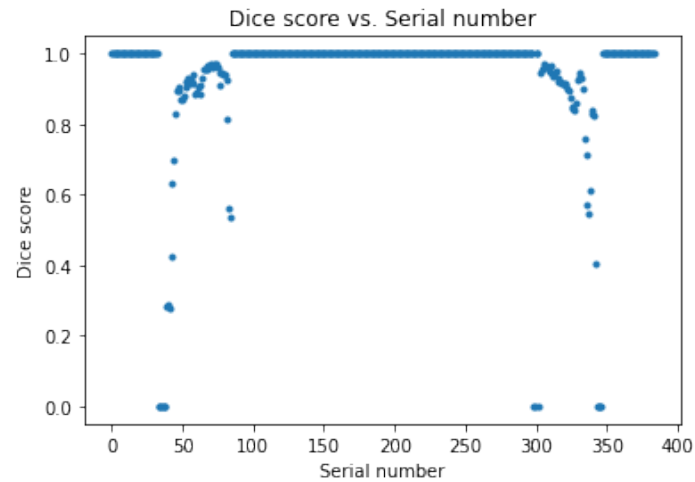
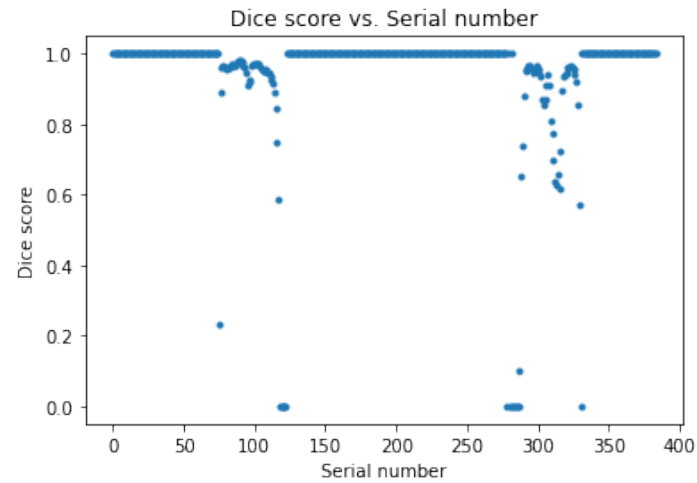
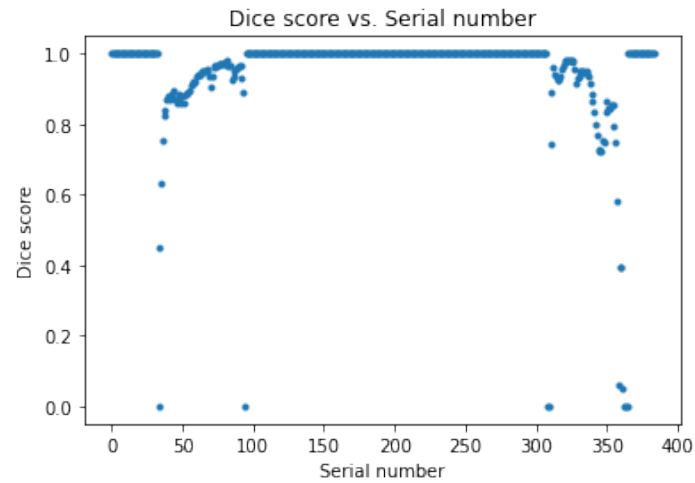






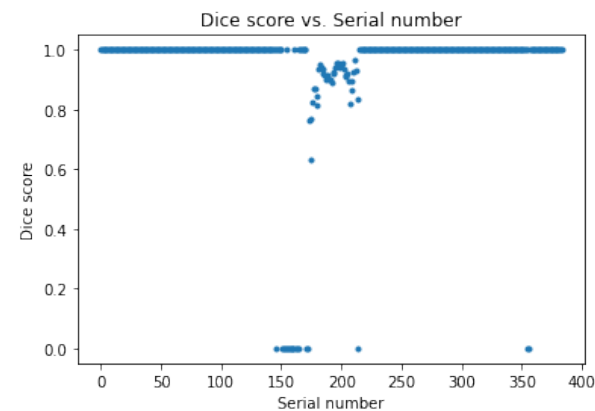
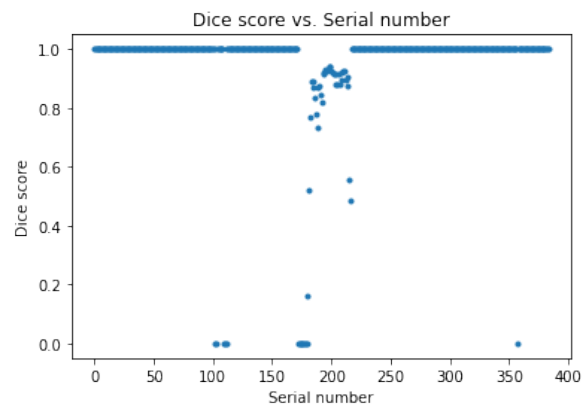
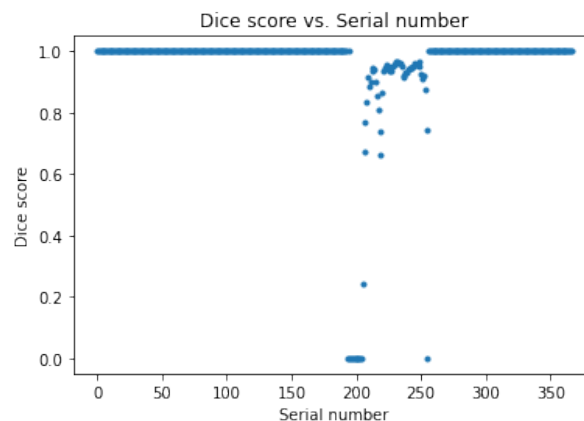
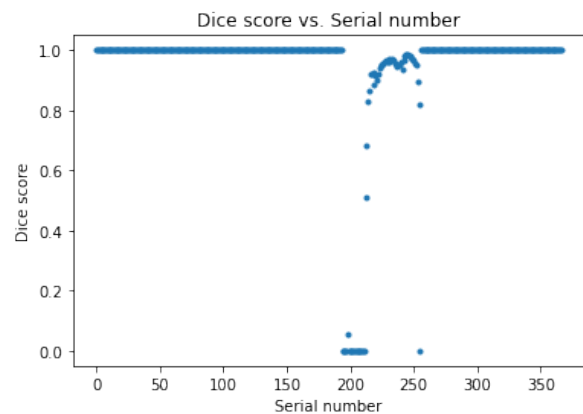
Experiment 3

Sagittal plane subnetwork



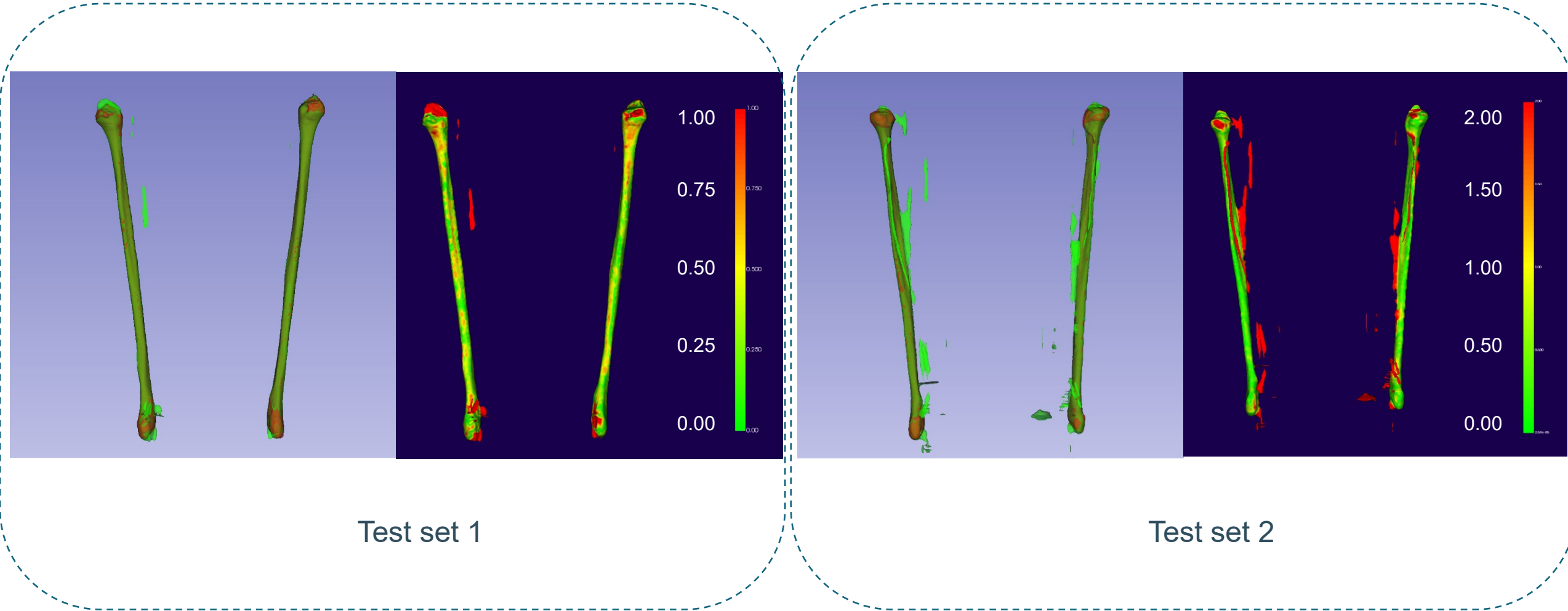
Experiment 3

Coronal plane subnetwork



1. Too few useful parts
2. Result is poor

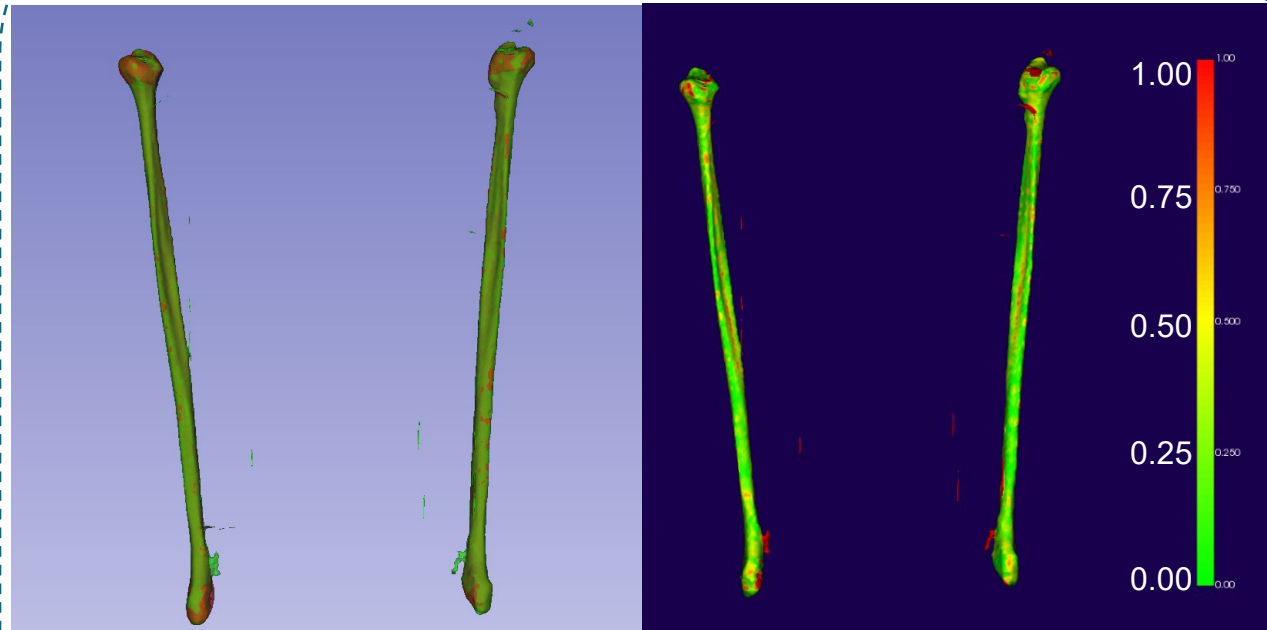
Discussion for Experiment 3



Discussion for Experiment 4



Test set 1



Test set 2

