

# Medical Image processing

## Ex 1

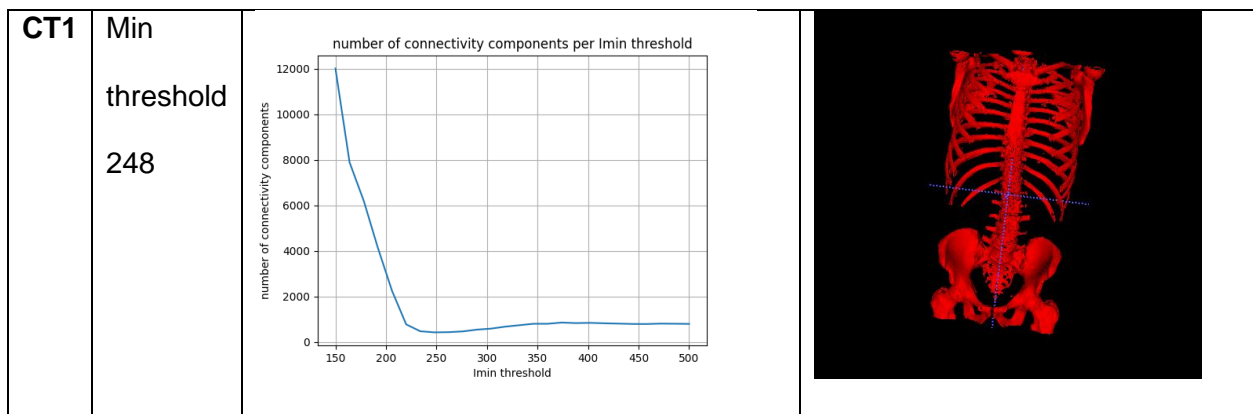
Amitai Ovadia 312244254

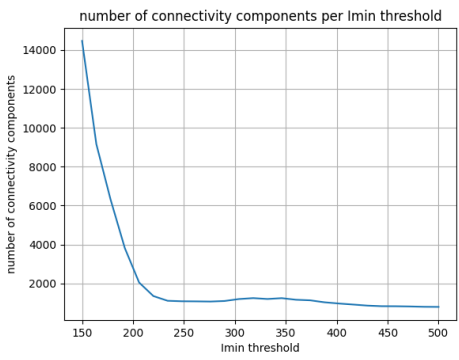
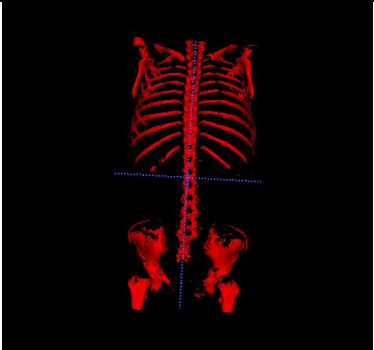
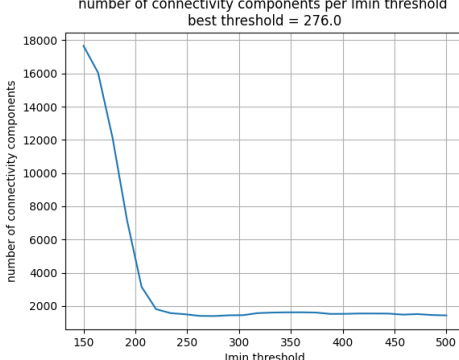
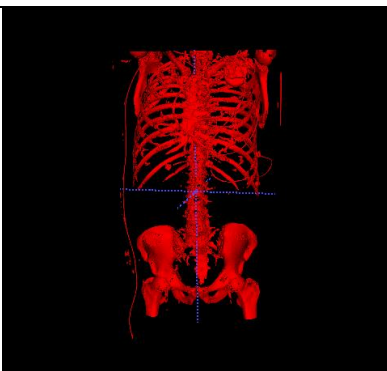
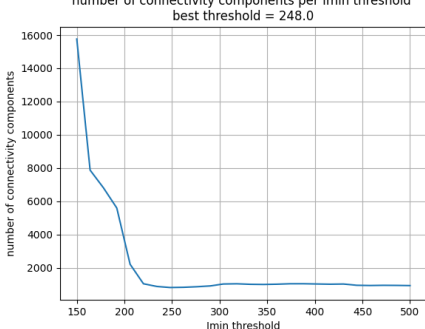
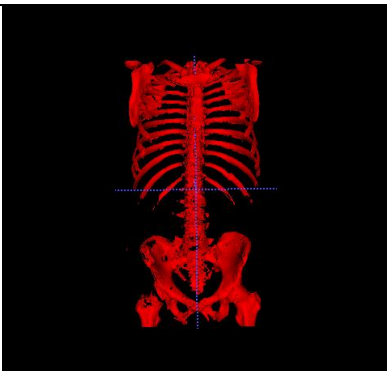
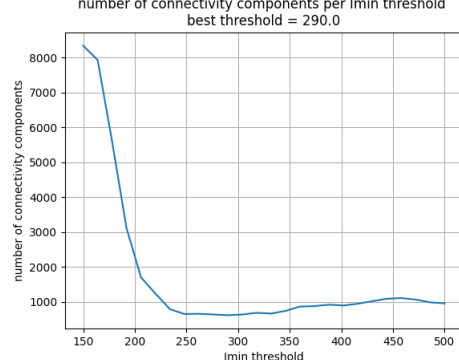
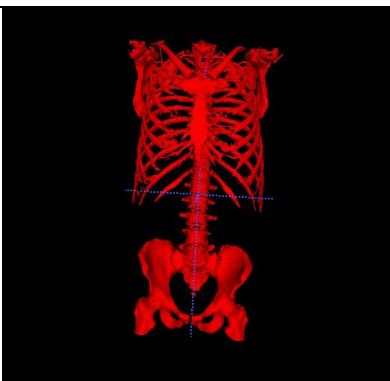
### Part 1:

#### Design:

- The solution thresholds all voxel values in the CT between minimum and maximum values using np.where function.
- It performs the thresholding for all values between [150, 500] in jumps of 14 and returns the result with the lowest number of connectivity components
- Then I performed morphological operations such as remove small objects, remove small holes and binary closing for several iterations, and stopped after a fixed amount of iterations so I won't erode all the image. Then I saved the resulting image.

#### Results:



CT2	Min threshold  500	<p>number of connectivity components per lmin threshold</p>  <table><tr><th>lmin threshold</th><th>number of connectivity components</th></tr><tr><td>150</td><td>14000</td></tr><tr><td>200</td><td>3000</td></tr><tr><td>250</td><td>2000</td></tr><tr><td>300</td><td>1800</td></tr><tr><td>350</td><td>1600</td></tr><tr><td>400</td><td>1500</td></tr><tr><td>450</td><td>1400</td></tr><tr><td>500</td><td>1300</td></tr></table>	lmin threshold	number of connectivity components	150	14000	200	3000	250	2000	300	1800	350	1600	400	1500	450	1400	500	1300	
lmin threshold	number of connectivity components																				
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200	3000																				
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350	1600																				
400	1500																				
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500	1300																				
CT3	Min threshold  276	<p>number of connectivity components per lmin threshold best threshold = 276.0</p>  <table><tr><th>lmin threshold</th><th>number of connectivity components</th></tr><tr><td>150</td><td>17000</td></tr><tr><td>200</td><td>4000</td></tr><tr><td>250</td><td>2000</td></tr><tr><td>300</td><td>1800</td></tr><tr><td>350</td><td>1600</td></tr><tr><td>400</td><td>1500</td></tr><tr><td>450</td><td>1400</td></tr><tr><td>500</td><td>1300</td></tr></table>	lmin threshold	number of connectivity components	150	17000	200	4000	250	2000	300	1800	350	1600	400	1500	450	1400	500	1300	
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CT4	Min threshold  248	<p>number of connectivity components per lmin threshold best threshold = 248.0</p>  <table><tr><th>lmin threshold</th><th>number of connectivity components</th></tr><tr><td>150</td><td>15000</td></tr><tr><td>200</td><td>3000</td></tr><tr><td>250</td><td>2000</td></tr><tr><td>300</td><td>1800</td></tr><tr><td>350</td><td>1600</td></tr><tr><td>400</td><td>1500</td></tr><tr><td>450</td><td>1400</td></tr><tr><td>500</td><td>1300</td></tr></table>	lmin threshold	number of connectivity components	150	15000	200	3000	250	2000	300	1800	350	1600	400	1500	450	1400	500	1300	
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CT5	Min threshold  290	<p>number of connectivity components per lmin threshold best threshold = 290.0</p>  <table><tr><th>lmin threshold</th><th>number of connectivity components</th></tr><tr><td>150</td><td>8000</td></tr><tr><td>200</td><td>2000</td></tr><tr><td>250</td><td>1000</td></tr><tr><td>300</td><td>800</td></tr><tr><td>350</td><td>700</td></tr><tr><td>400</td><td>600</td></tr><tr><td>450</td><td>500</td></tr><tr><td>500</td><td>400</td></tr></table>	lmin threshold	number of connectivity components	150	8000	200	2000	250	1000	300	800	350	700	400	600	450	500	500	400	
lmin threshold	number of connectivity components																				
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## libraries:

- Skimage :
- Numpy

## Methods:

class BonesSegmentation;

**Function:** SkeletonTHFinder

Takes the CT in the nifty\_file and segments the skeleton using threshold and morphological operations.

**Input:** self.nifty\_file\_path

**Output:** saves the new skeleton segmentation as a nifty file

**Function:** get\_best\_threshold

Searching through the threshold space and returning the threshold that gives the lowest number of connected components in the 3D CT image

**Input:** self.nifty\_file\_path

**Output:** best\_threshold, int representing the best minimum threshold

**Function:** SegmentationByTH

loads the CT image and thresholds it according to lmin and lmax values

**Input:**

lmin: minimal threshold

lmax: maximal threshold

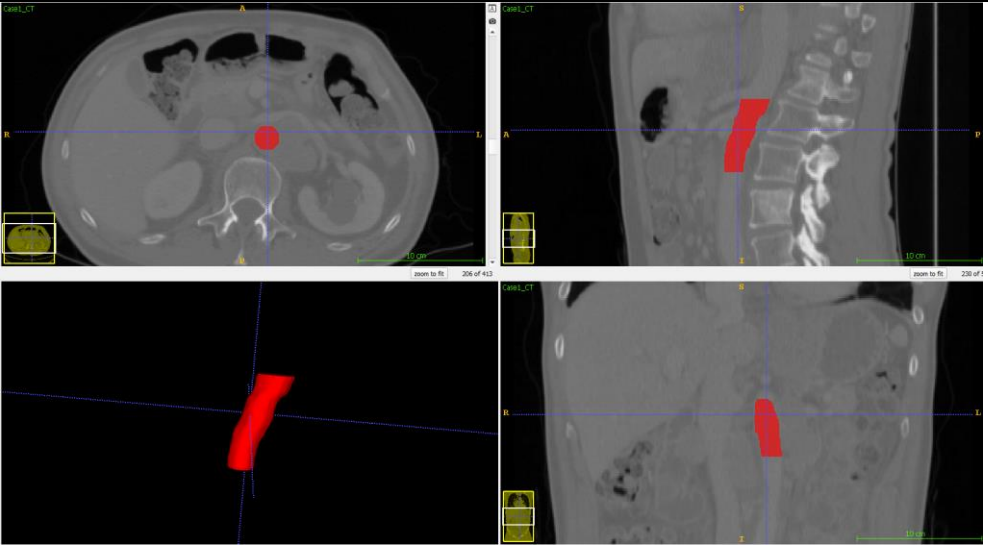
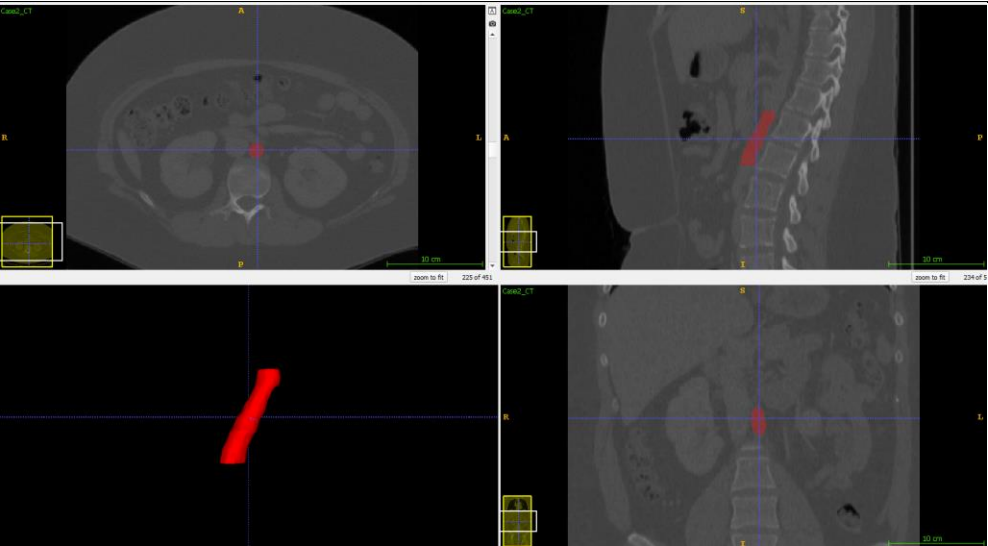
**Output:** save the segmented nifty file and returns 1 is succeeded 0 if failed

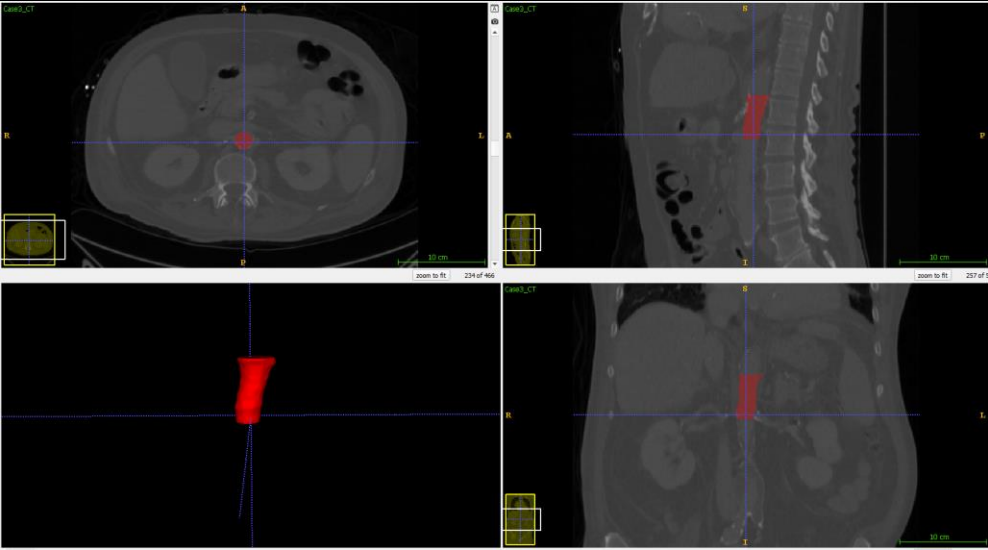
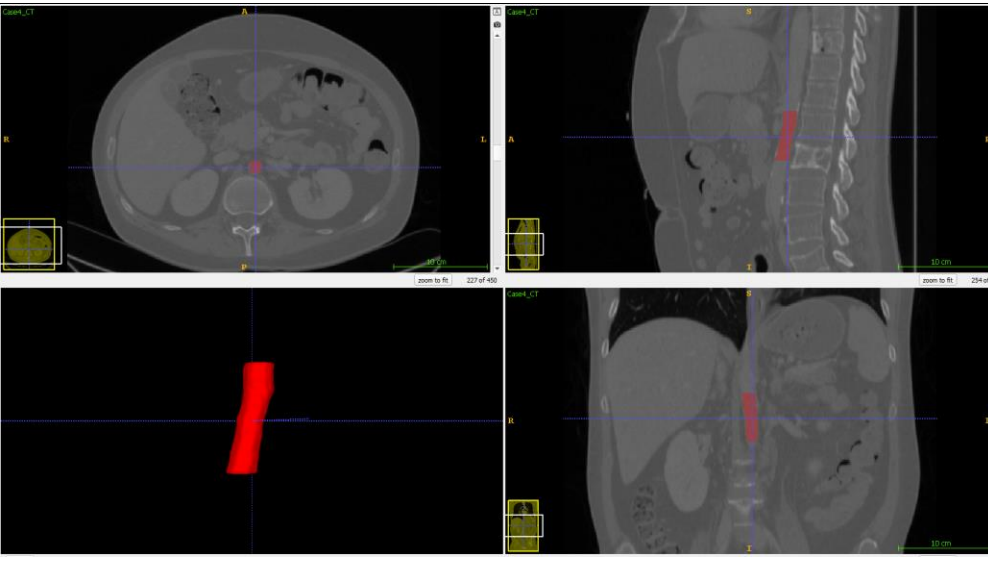
## **Part 2:**

### **Design:**

- CT scan is cut, and the remaining slices are the ones that include L1
- CT is cut again, according to assumed Aorta location in x, y: close the front of L1.
- Assuming that the Aorta is round, or close to round I performed the following trick:
- Each of this cut CT scans is then analyzed using the Circles Hough Transform. The supplied parameters are the search radii: between 7 and 20 pixels, and the number of desired circles is 20.  
  
overall I got 20 circle candidates for Aorta location per slice, ordered by their likelihood.
- Then in order to choose the most fitting circle for Aorta location in each slice I assumed that the Aorta is more or less in the same location in consecutive slices and used the following method:
- Fixing the first chosen circle in the lowest slice to be the most likely circle, I then chose the above circle, from the 20 different candidates, to be the one that is closest to the previous circle in terms of location and radius.
- I did this iteratively, choosing for every slice it's Aorta location circle according to it's previous neighbor.
- If no close circle was found, according to a specified error rate, a circle was then created in the location and radius of the previous circle (almost never happened).
- Then I created the Aorta segmentation by filling the circles per slice and stacking them on top of each other, and performed binary closing in order to smooth the result.

Results:

CT1	DICE = 0.93 VOD = 0.12	 <p>The image displays three panels for Case1_CT. The top-left panel is an axial CT scan showing a red segmented region in the center. The top-right panel is a sagittal CT scan showing a red segmented region along the spine. The bottom-left panel is a coronal CT scan showing a red segmented region. Each panel includes a small inset image in the bottom-left corner and a scale bar in the bottom-right corner.</p>
CT2	DICE = 0.82 VOD = 0.30	 <p>The image displays three panels for Case2_CT. The top-left panel is an axial CT scan showing a red segmented region in the center. The top-right panel is a sagittal CT scan showing a red segmented region along the spine. The bottom-left panel is a coronal CT scan showing a red segmented region. Each panel includes a small inset image in the bottom-left corner and a scale bar in the bottom-right corner.</p>

CT3	DICE = 0.87 VOD = 0.22	 <p>The image displays three panels for Case3_CT. The top-left panel is an axial view showing a cross-section of the abdomen with a red segmented region in the center. The top-right panel is a sagittal view showing the spine and abdominal organs with a red segmented region. The bottom-left panel is a coronal view showing the abdominal cavity with a red segmented region. Each panel includes a 20 cm scale bar and a small inset image in the bottom-left corner.</p>
CT4	DICE = 0.91 VOD = 0.16	 <p>The image displays three panels for Case4_CT. The top-left panel is an axial view showing a cross-section of the abdomen with a red segmented region in the center. The top-right panel is a sagittal view showing the spine and abdominal organs with a red segmented region. The bottom-left panel is a coronal view showing the abdominal cavity with a red segmented region. Each panel includes a 20 cm scale bar and a small inset image in the bottom-left corner.</p>

### Library:

- Morphology, transform, draw, feature, exposure and measure from skimage
- Numpy

## Methods:

class AortaSegmentation:

<b>Function:</b> get_L1_boundaries(self)
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find x,y,z boundaries of L1 to cut the CT accordingly
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<b>Input:</b> self.L1_img
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<b>Output:</b> min_x, max_x, min_y, max_y, min_z, max_z
---

<b>Function:</b> AortaSegmentation(self)
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Segment Aorta image and save segmentation, uses the function self.find_Aorta()
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<b>Input:</b> self.L1_img, self.CT_img
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<b>Output:</b> saves the Aorta segmentation
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<b>Function:</b> find_Aorta(self)
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finds all Aorta pixels in CT: first per slice and then for all 3D image
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uses the functions:
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self.find_possible_Aorta_circles()
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self.extract_best_Aorta_circles(circles)
--

self.construct_3d_Aorta_segmentation(final_circles)
---

<b>Input:</b> self
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<b>Output:</b> segmented_Aorta
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**Function:** find\_possible\_Aorta\_circles(self)

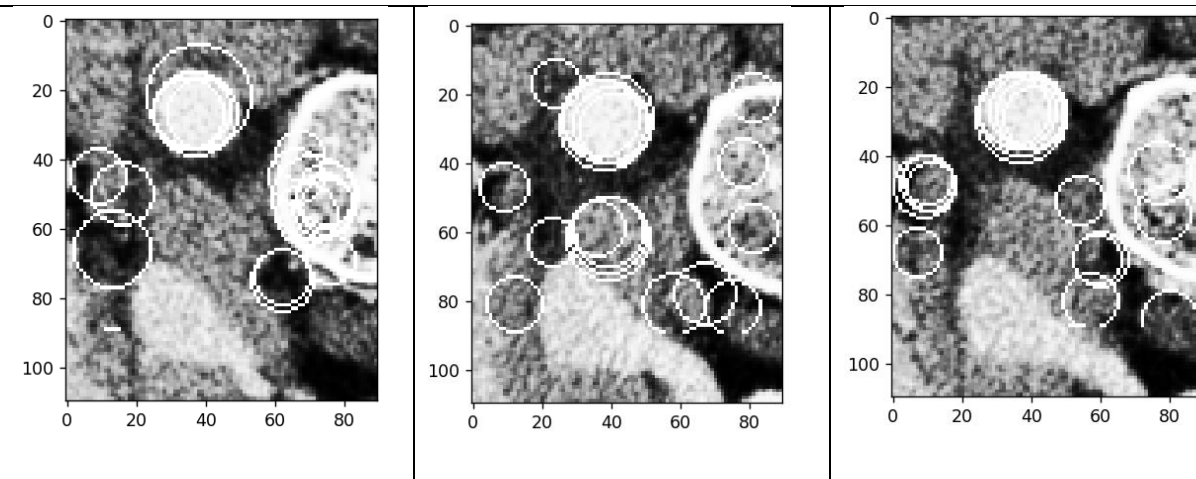
for each L1 slice finds 20 possible Aorta's circle locations  
using hough transform

**Input:** self.cut\_CT\_img

**Output:** circles: a list of circles information arrays per slice

In the images:

all the candidate circles for the Aorta locations in a specific slice are encircled in white.



**Function:** extract\_best\_Aorta\_circles(self, circles)

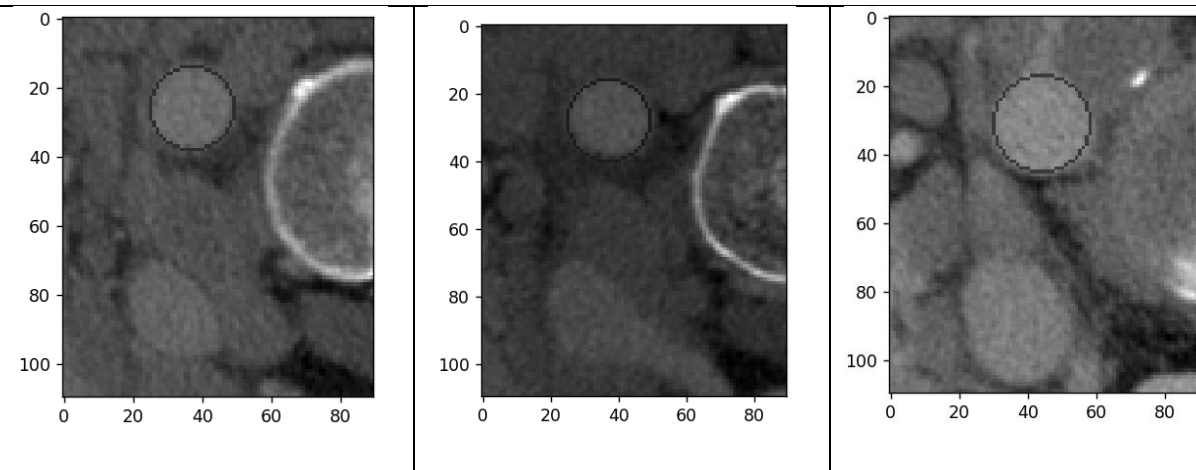
extract the most fitting Aorta circle for each slice

**Input:** circles: the output of find\_possible\_Aorta\_circles

**Output:** final\_circles: an array of the selected 1 circle per slice for Aorta location

In the images:

the estimated Aorta location is encircled in black.





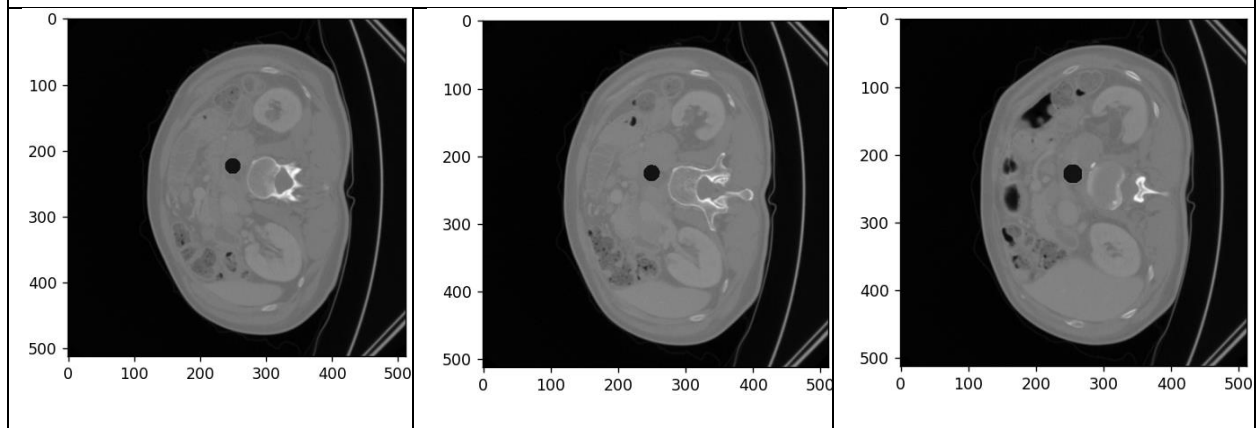
**Function:** `construct_3d_Aorta_segmentation(self, final_circles)`

construct 3D Aorta segmentation from Aorta circles information from each slice

**Input:** `final_circles`: Aorta circles information from each slice

**Output:** 3D Aorta segmentation

The images are of the Aorta estimated location in some CT slices



**Function:** `evaluateSegmentation(GT_seg, est_seg)`

evaluate performance of segmentation compared to the ground truth segmentation

**Input:** `GT_seg`: ground truth segmentation  
`est_seg`: computed segmentation

**Output:**  
DICE  $((2 * \text{intersection\_size}) / (\text{size\_A} + \text{size\_B}))$   
and  
VOD  $(1 - \text{intersection\_size} / \text{union\_size})$   
scores for evaluating the computed segmentation