

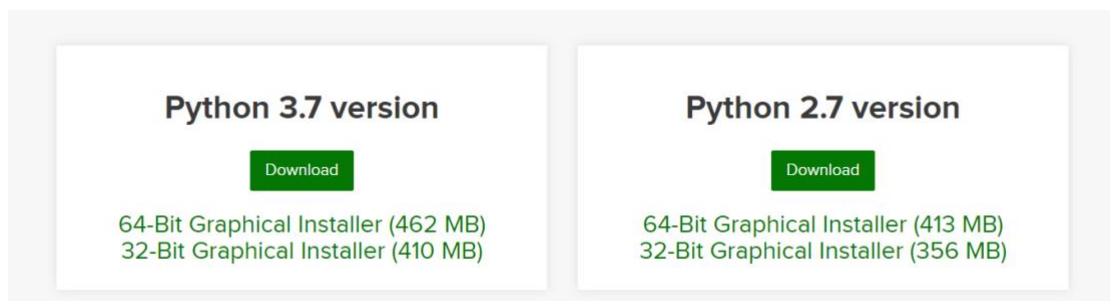
# Cardiovascular risk computed via Deep Learning (DL) on thoracic CT scans (Med3DResNet)- Evaluation tutorial in Windows

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## First of All

- Install Anaconda3 (python3.7) <https://www.anaconda.com/distribution/>.



- Create a new folder under a path of your choice and name it `Project`.  
(e.g. `E:\Project`).
- `cd` into the `Project` directory. Clone/Download our github repository <https://drive.google.com/file/d/1CmRXBzu2vuJOLgYvSWH-gTF79YiRhW-h/view?usp=sharing> to the Project folder.
- Extract it's contents inside the `Project` folder.

- You should now have a single folder named `ACVProject-master` under your `Project` folder, which contains code and another files as such:

	<code>data</code>	2019/12/10 1:43
	<code>documentation</code>	2019/12/16 16:35
	<code>images</code>	2019/12/10 1:43
	<code>videos</code>	2019/12/16 16:35
	<code>.gitignore</code>	2019/12/10 1:43
	<code>AnnotationWidget.py</code>	2019/12/10 1:43
	<code>CNN_batch_run.py</code>	2019/12/10 1:43
	<code>CNN_main.py</code>	2019/12/10 1:43
	<code>CNN_ops.py</code>	2019/12/10 1:43
	<code>CNN_ResNet.py</code>	2019/12/10 1:43

### Create a new Conda virtual environment (Optional)

- Open a new Anaconda/Command Prompt window

- `cd` into the `ACVProject-master` directory.
- Type the following command:

```
conda create -n Med3D pip python=3.6
```

- The above will create a new virtual environment with name `Med3D`
- Now lets activate the newly created virtual environment by running the following in the Anaconda Prompt window:

Activate Med3D

- Once you have activated your virtual environment, the name of the environment should be displayed within brackets at the beginning of your cmd path specifier, e.g:

```
(Med3D) E:\Project\ACVProject-master>
```

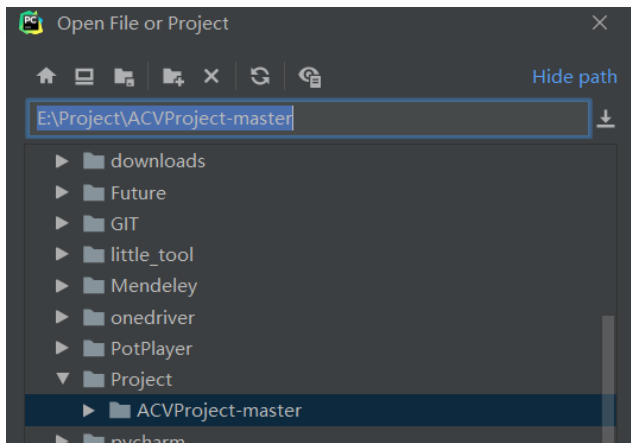
- Type the following command:

```
pip install -r requirements_win.txt
pip install keras
```

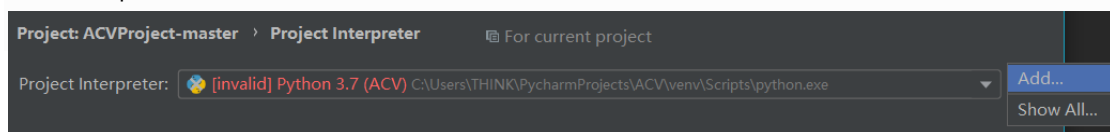
- Install Pycharm

<https://www.jetbrains.com/pycharm/download/#section=windows>

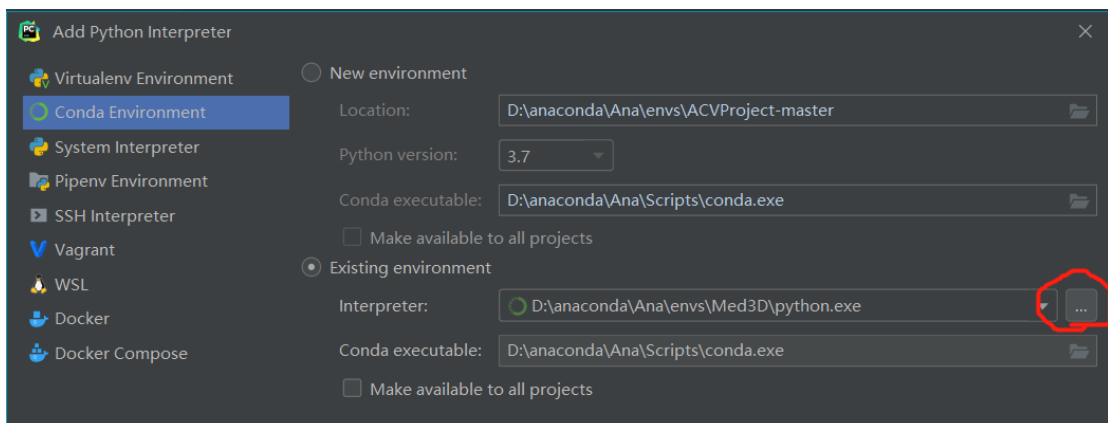
- Click File=>Open choose ACVProject-master directory.



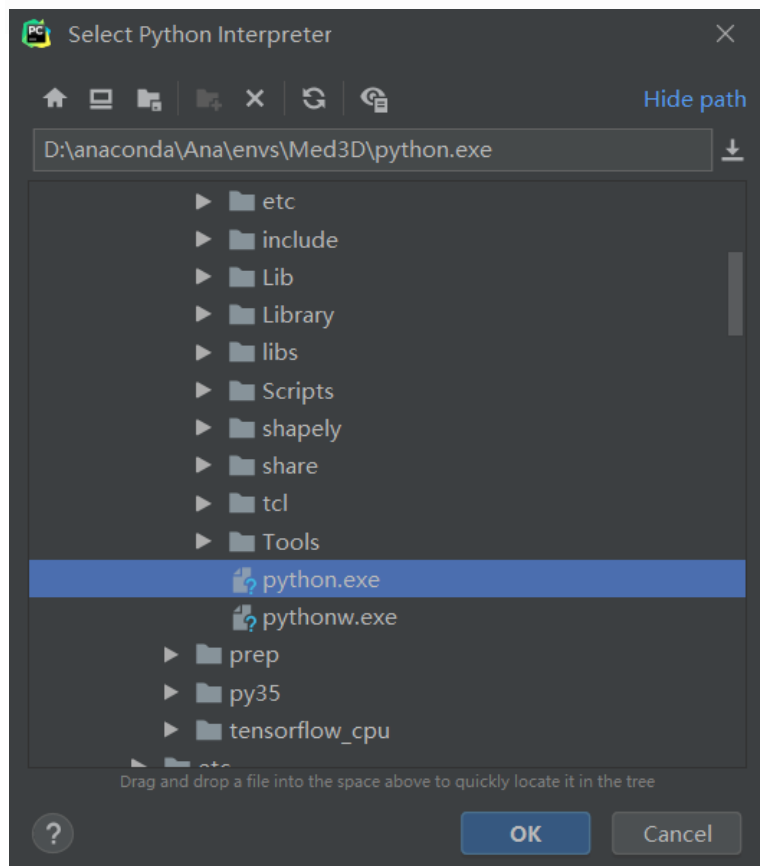
- Click File=>settings=>Project:ACVProject-master=>Project interpreter=>add



- Click Conda Environment=>Existing environment=>...

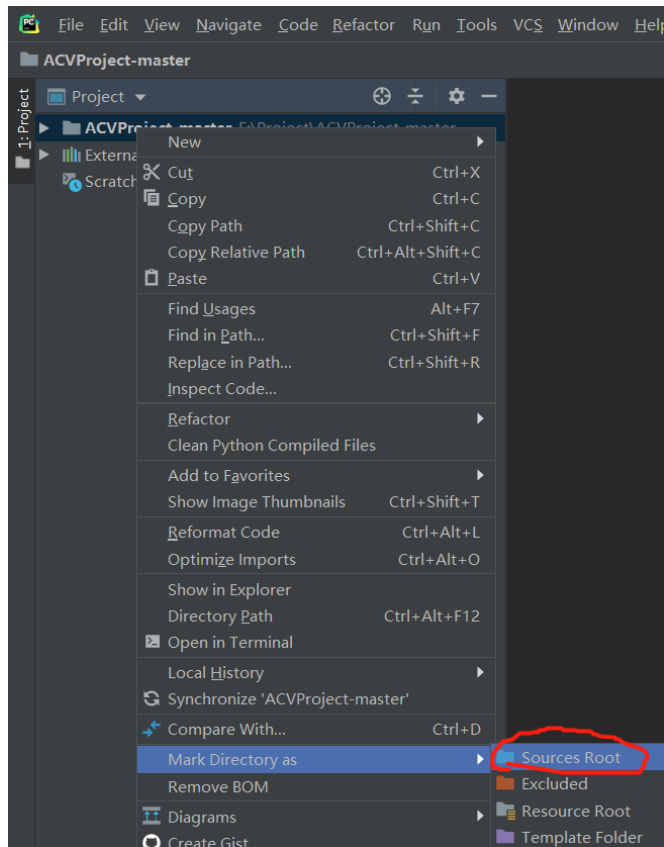


- Choose path to where you install anaconda\envs\Med3D\python.exe

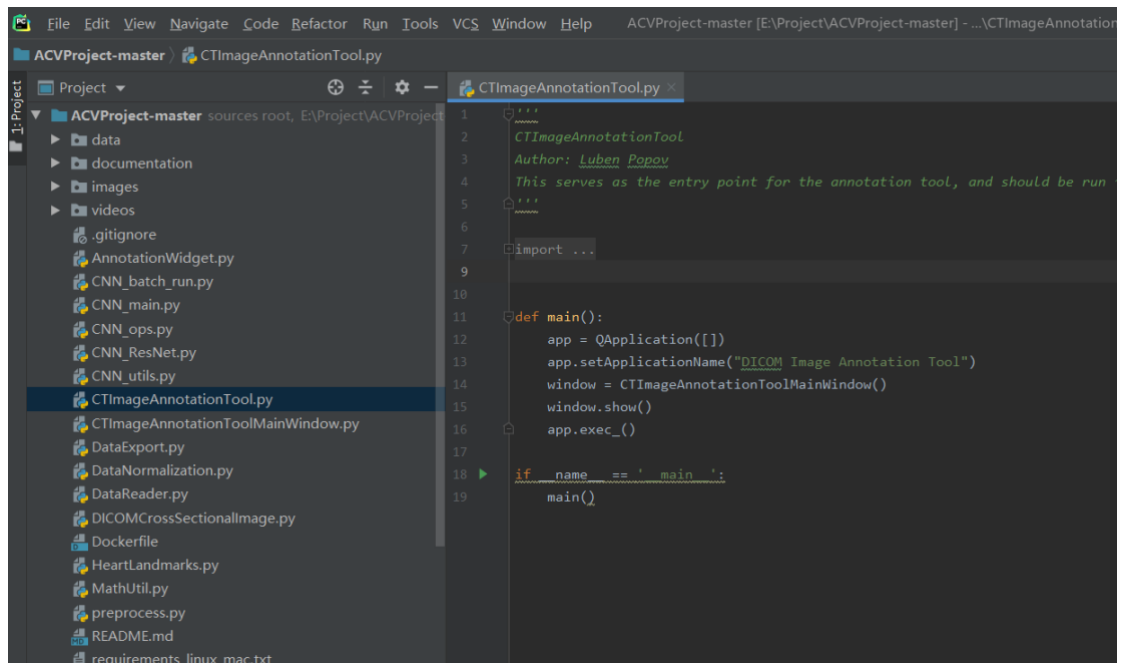


- Click OK=>OK

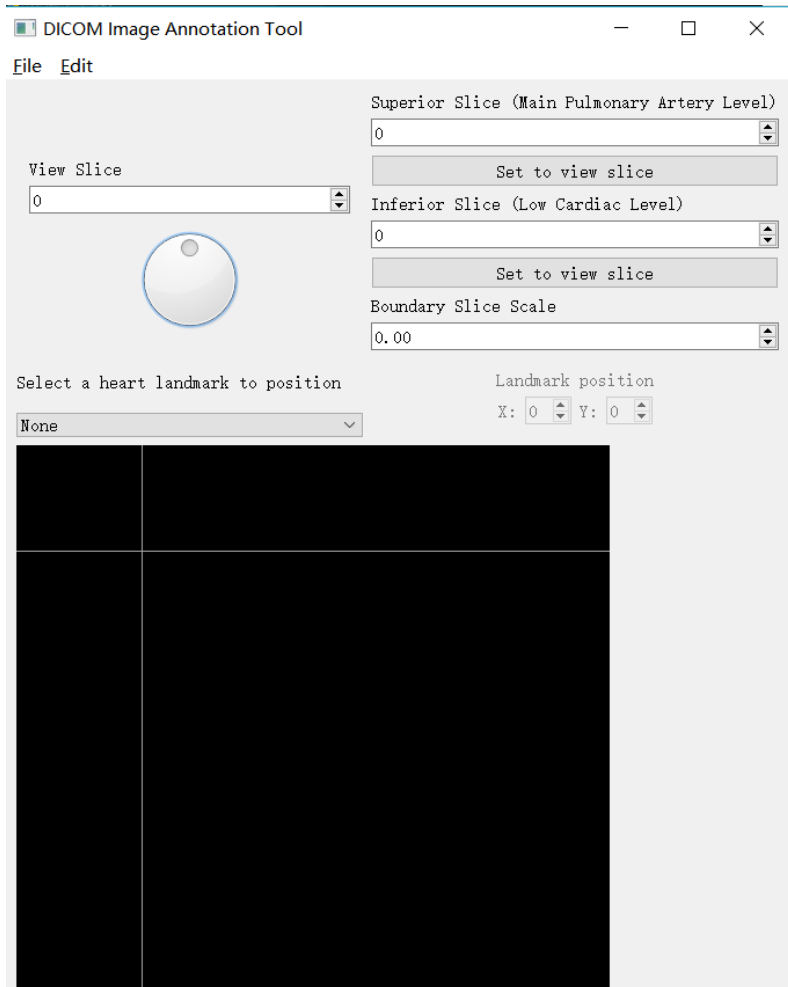
- Right click ACVProject folder and set it as source folder.



- Now Open `CTImageAnnotationTool.py` and run this script with Alt+Shift+F10



- Now you should open this annotation tool like below:



- For annotation tool using please check `E:\Project\ACVProject-master\documentation\CTImageAnnotationToolManual.pdf`
- Download dataset with this google drive link to the `E:\Project`: [https://uofh-my.sharepoint.com/:f:/g/personal/taburt\\_cougarnet\\_uh\\_edu/EiZNY6eYAINBkJFTpeljkzYBrS9p03lG64a\\_DcJW3GPZGw](https://uofh-my.sharepoint.com/:f:/g/personal/taburt_cougarnet_uh_edu/EiZNY6eYAINBkJFTpeljkzYBrS9p03lG64a_DcJW3GPZGw)
- Extract it's contents inside the `E:\Project` folder.
- You should now have folder as following:  
E:  
├─ ACVProject-master  
└─ ACV\_project\_Team1\_F19
- For dataset and CNN framework description please check `E:\Project\ACVProject-master\documentation\Med3DResNetManual.pdf`

- Open a new *Anaconda/Command Prompt* window
- `cd` into the `ACVProject_master` directory.
- Add two line code in the `CNN_utils.py`

```
Import csv
```

```
Import numpy as np
```

- Change the `CNN_utils.py` 108 line code from

```
MIN_HU, MAX_HU = update_hu_range(test_image_temp, MIN_HU, MAX_HU)
```

- To

```
MIN_HU, MAX_HU = Visualization.update_hu_range(test_image_temp, MIN_HU, MAX_HU)
```

- Type the following command:

```
python CNN_main.py --phase train --dataset ACV --res_n 18 --work_path  
E:/Project/ACV_project_Team1_F19/ --train_test_ratio 70_30 --batch_size 40 -  
-lr 0.1 --data_type projection --n_axial_channels=4 --epoch 25
```