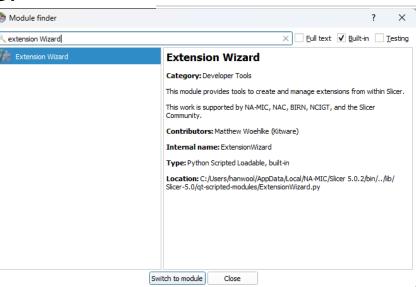
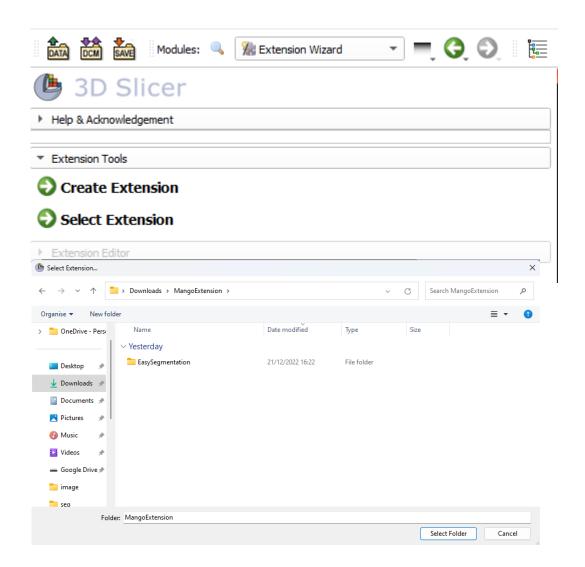
# EasySegmentation v1.6 Tutorial

- 1. Download MangoExtension folder
  - Method 1: git clone <a href="https://gitlab.com/mangomedical/extra/slicer\_gui.git">https://gitlab.com/mangomedical/extra/slicer\_gui.git</a>
  - Method 2: Download from <u>link</u>. (You need to unzip this file)
- 2. Open Slicer
- 3. Find Extension Wizard Module in Module finder



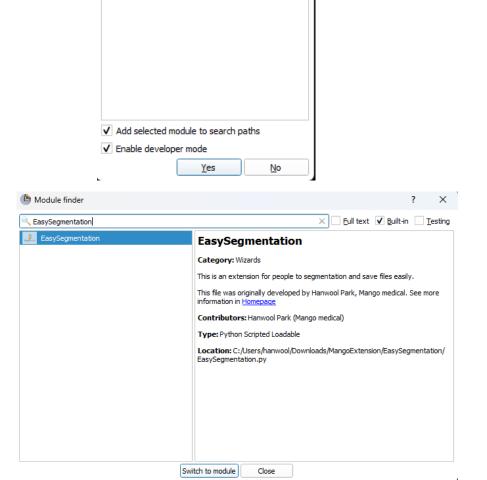
Modules:

Data Data



4. Click "Select Extension" button

5. Select "MangoExtension" folder. (Not EasySegmentation folder)



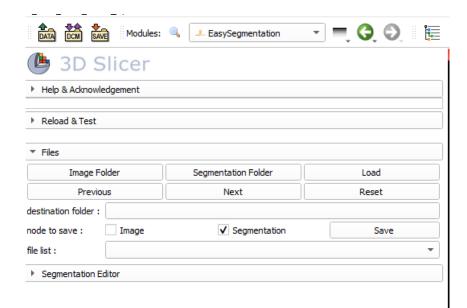
The following module can be loaded. Would

you like to load it now?

✓ EasySegmentation

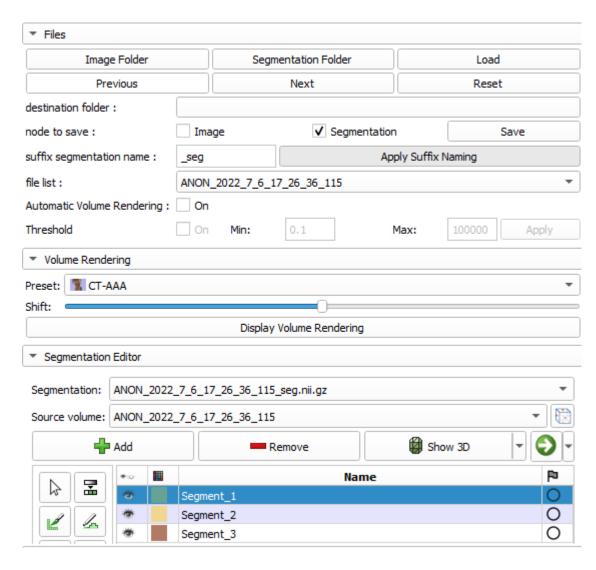
6. Click "Yes" button

7. Go to Module finder and search "EasySegmentation" module



8. If you can see this module, you succeed to import this module

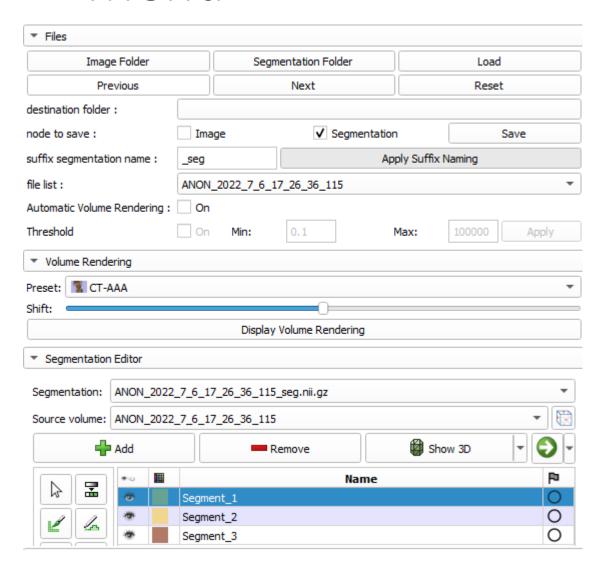
#### Menu



#### • Files:

- Image Folder:
  - choose the folder includes image volume files
- Segmentation Folder :
  - choose the folder includes segmentation files corresponding image volume files
  - need to include image file name. Ex) image: test1.nii.gz, segmentation: test1 seg.nii.gz
  - The file extensions can be .nii.gz, .nii or .nrrd
- Load :
  - after choose image and segmentation folder, we can load file in slicer
- Previous :
  - move previous file
- Next:
  - move next file
- Reset :
  - reset all setup this module
- Destination folder:
  - choose destination folder to save image or segmentation files
- Node to save :
  - you can choose to image or segmentation file to be saved
- Save:
  - save your file into destination folder
- Suffix segmentation name :
  - You can set the suffix name for segmentation file.
  - If you don't want to use suffix function, you can click "Apply Suffix Naming" button
- File list :
  - after you load files, you can see file list in combo box and you can click specific file.
- Automatic Volume Rendering :
  - You can check to load volume rendering view automatically
- Threshold :
  - When there is no segmentation data corresponding volume data, you can set up threshold values for new segmentation

## Menu

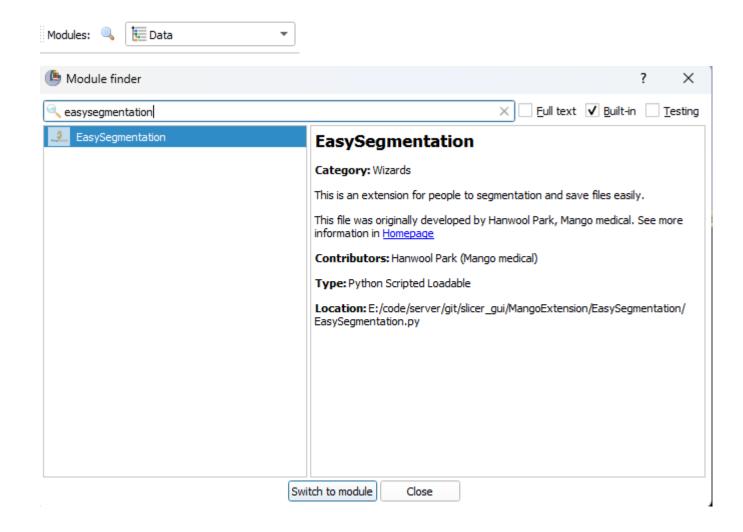


#### • Volume Rendering:

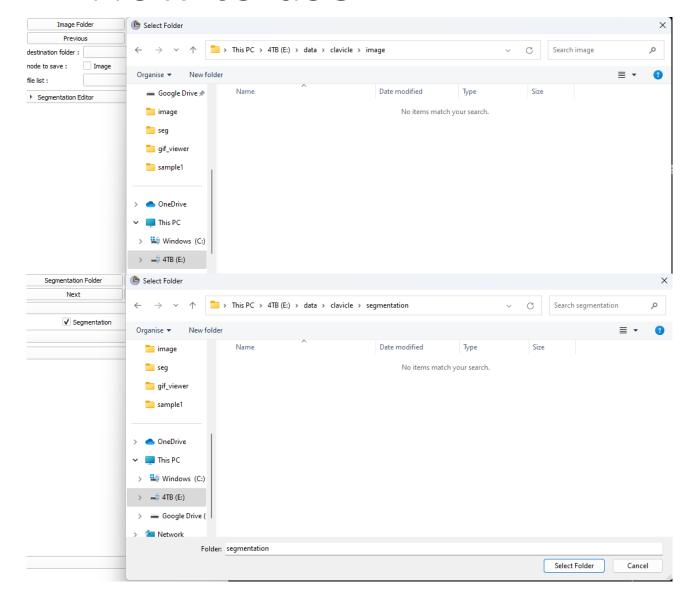
• This is volume Rendering module. You can check information here.

#### • Segmentation Editor:

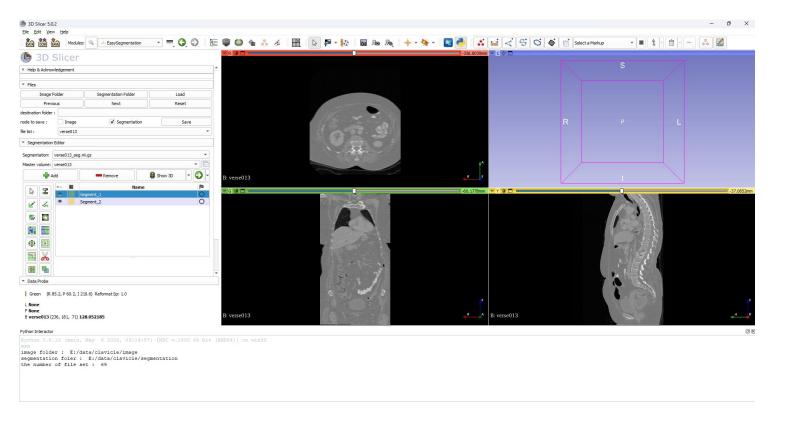
• This is Slicer Segmentation Editor module. You can check information <a href="here">here</a>.



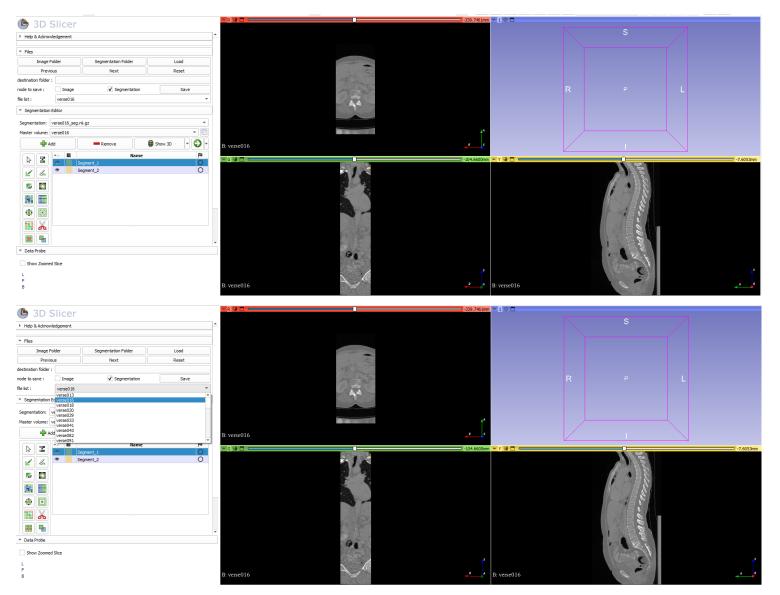
Click Module finder on Toolbar and search Easysegmentation



Choose image folder and segmentation folder

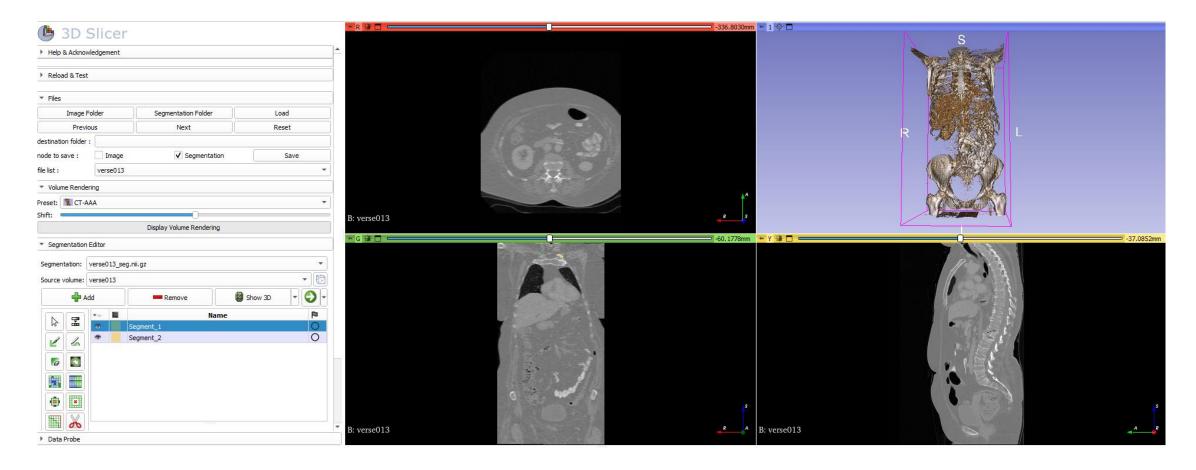


Click Load button. You can see how many files are found in Python Interactor.

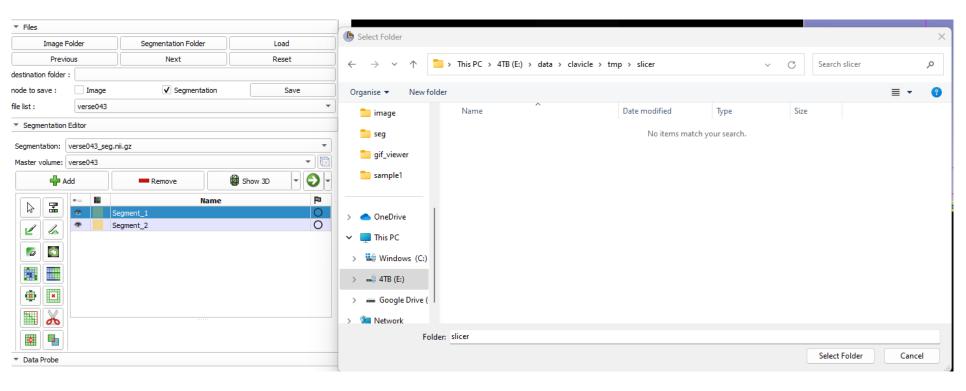


Click Next or Previous Button. You can move next or previous file.

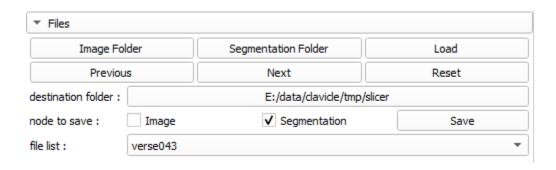
You can also click file list and choose specific file.



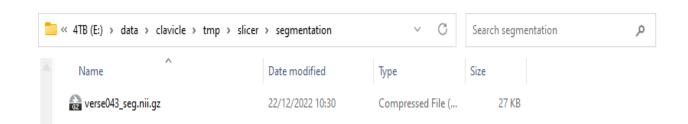
You can see volume rendering view to click "Display Volume Rendering" button You can also check Preset to see different types of views



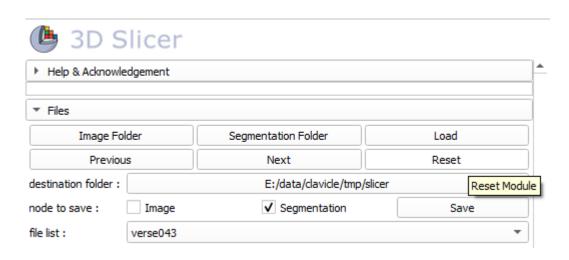
To save modified files, you need to choose destination folder.



You can choose Image or Segmentation to save.



You can check save result into destination folder





You can reset this module with Reset button.

Everything is reset. You can start module again

## **Version Notes**

#### Version 1.1

- fixed warning issues regarding deprecated functions
- Now, all image data can be loaded without corresponding segmentation file. You can create your own segmentation now.

#### Version 1.2

• add volume rendering feature

#### Version 1.3

• add file extensions (nii.gz, nii or nrrd)

#### Version 1.4

- add automatic rendering volume checkbox
- center the 3D view on the scene

#### Version 1.5

• Add threshold feature to create segmentation data when there is no segmentation data corresponding volume data

#### Version 1.6

• add suffix naming feature to save segmentation file with specific name

#### Notes

- Build Slicer version: 5.2.1
- OS: Windows, Linux (I don't know it works in MacOS)
- If you have any problem, please write email to hanwool@mangomedical.io