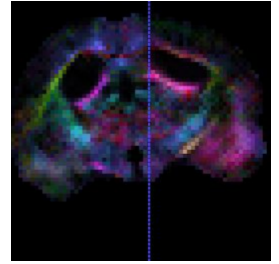


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Creating Diffusion Tensor Images (DTI)

DOI

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Protocol status: In development

We are still developing and optimizing this protocol

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Abstract

This procedure describes the steps required to create diffusion tensor images (DTI) from data from the MRI scanner.

The following programs are required to go through the protocol:

- MRtrix3
- TORTOISE
- ITKsnap

Note: If you are working from a Windows computer, you will need to follow X2Go Client Set-Up (dx.doi.org/10.17504/protocols.io.6tvhen6) before you can begin this protocol.

Troubleshooting

Set Up Directory with Required Files

- 1 Open a new terminal.
- 2 Confirm what directory you are currently in.

Command

```
pwd (0-58-generic #64-Ubuntu SMP)
```

```
pwd
```

```
pwd
```

- 3 Create a directory to place data. *(If not already created.)*

Command

```
Create directory. (0-58-generic #64-Ubuntu SMP)
```

```
mkdir
```

```
mkdir ~/Data/Projects/
```

- 4 Go to the directory that you plan to use.

Command

Go to a specified directory. (0-58-generic #64-Ubuntu SMP)

```
cd
```

```
cd
```

- 5 Copy directory where your raw data files that will be acquired, reside in. You will need the following file types:
 - Bvals
 - Bvecs
 - data.img
 - data.hdr* - (*may not need this in all cases - used for FOV origin*)
- 5.1 Find where the files you plan to copy are located. *Make sure to use the original root i.e /home/... as you will get an error if you start from another location.*

Command

To list files in a specific directory. (0-58-generic #64-Ubuntu SMP)

```
ls
```

```
ls
```

- 5.2 Copy the specific files that you need. Alternatively, you can use . to specify the current directory you are in.

Command

Copy file or entire directory to a specified location. (0-58-generic #64-Ubuntu SMP)

cp

cp

- 6 Perform conversion between different file types.

Command

mrconvert (MRtrix3)

mrconvert

mrconvert

- 7 Import data from nifti format to TORTOISE list file format. A new directory will be created and the folder will be labeled **____proc**.
Example:



Command

ImportNIFTI (TORTOISE)

ImportNIFTI

ImportNIFTI

- 8 Go to the new directory that was created from the previous step.
- 9 Correct motion. Use **DIFFPREP** command. (*This is a command from TORTOISE*). **Note:** ***This step is time & core consuming. Also, this step removes Gibbs ringing and noise***

Command

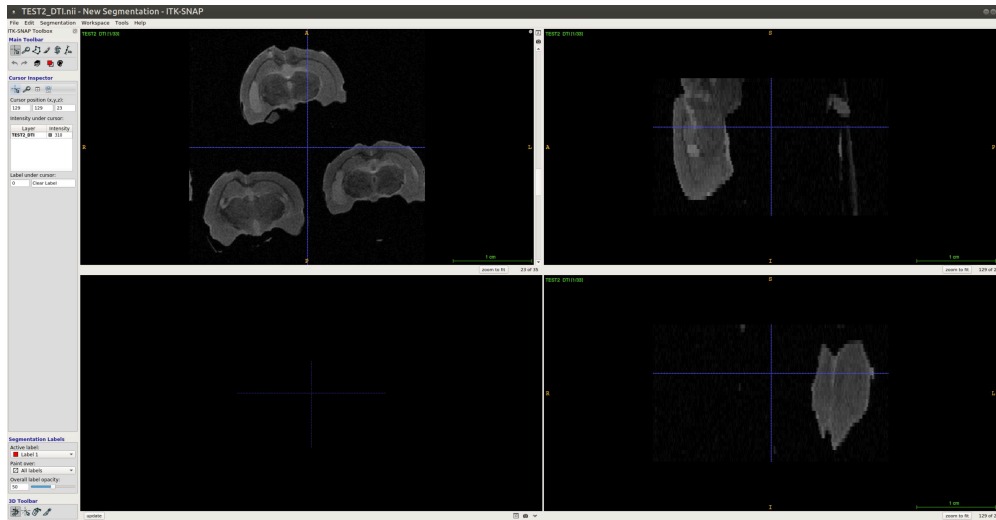
Corrects motion. (TORTOISE)

DIFFPREP

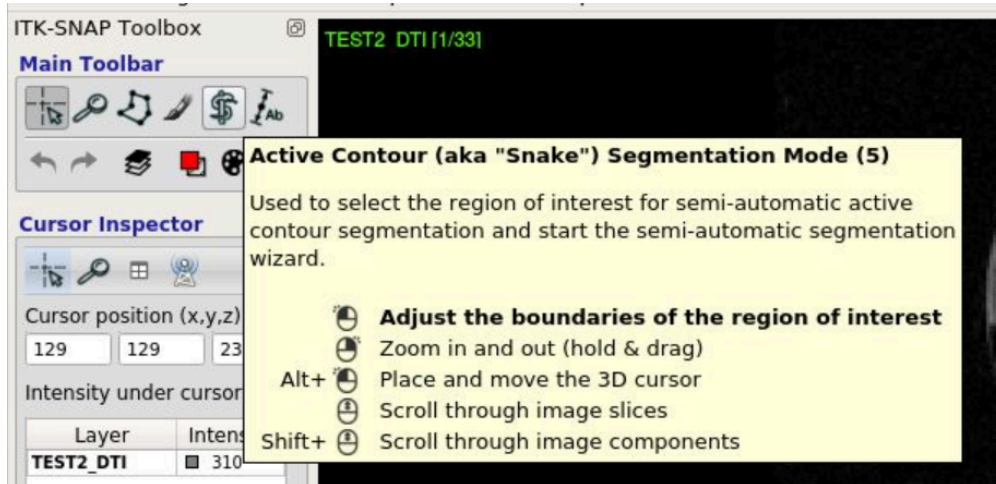
DIFFPREP

Create Mask and Segmentation

- 10 Open one of the populated files using ITKsnap.



- 11 Within ITKsnap, select **Active Contour (aka Snake) Segmentation Mode** from the toolbar.



- 12 A new side window will appear. Select **Segment 3D**.

Snake Inspector



ROI for auto-segmentation:

Position (x,y,z):

Size (x,y,z):

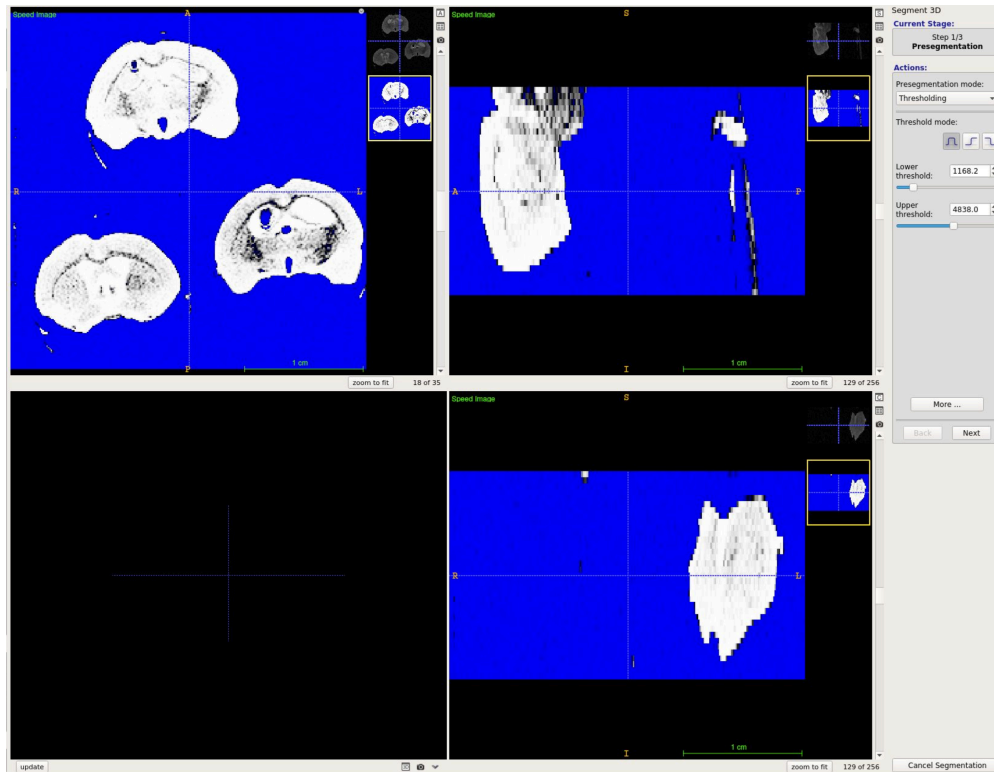
Reset ROI

☐ Resample ROI

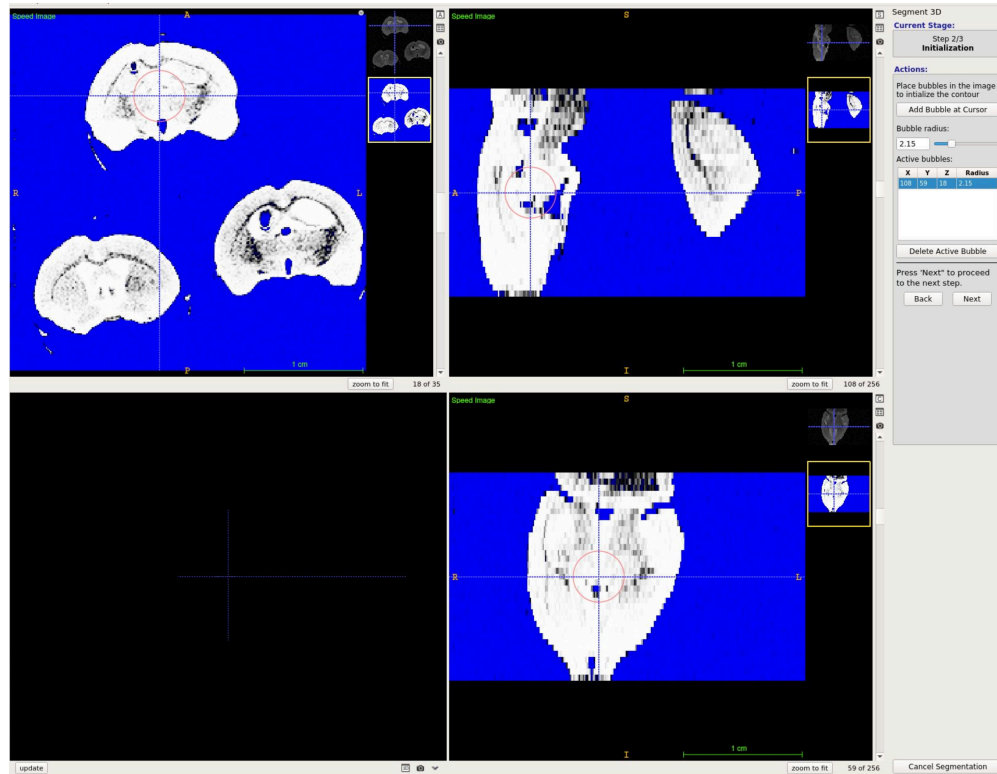
☐ Initialize with current segmentation

Segment 3D

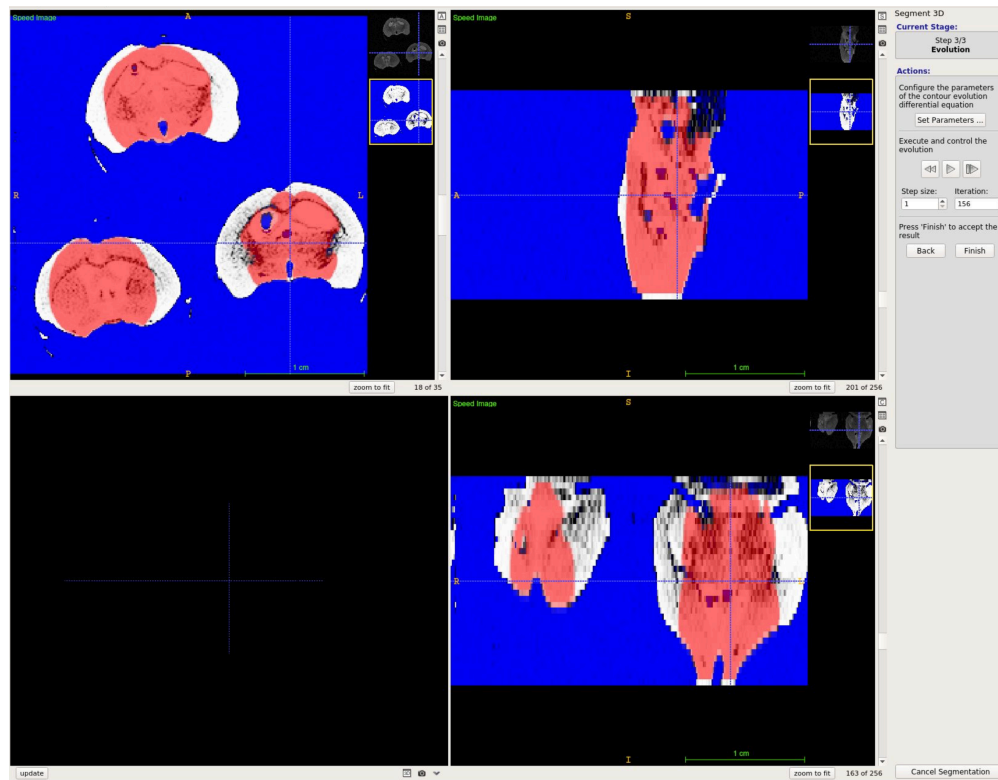
- 13 Toggle the thresholds (lower & upper) so that the blue mask covers the region you want repelled. (*Regions that are empty space or not the brain of interest should have this mask*). **Note: Try not to change the defaults.**



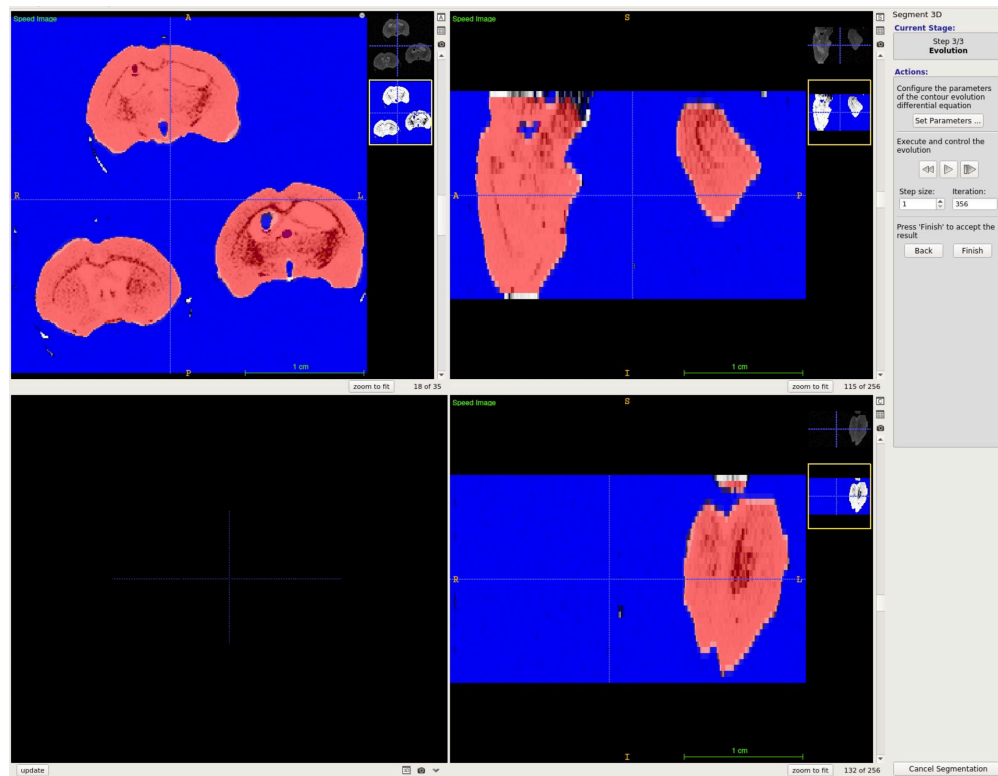
- 14 When the thresholds have been finalized, select **Next**.
- 15 Set the cursor in the relative center of your object of interest. Select **Add bubble at cursor**. A red circle(seed/bubble) will be placed at the location.
Note: You may need to repeat this if you have multiple objects of interest.



- 16 When the bubbles have been set and you have no more to add, select **Next**.
- 17 Press the play button to have the seed grow within the mask that was set-up in a previous step. **Note: Red should propagate and fill the entire region(s) of interest. This may require viewing the images from the different planes to confirm.**

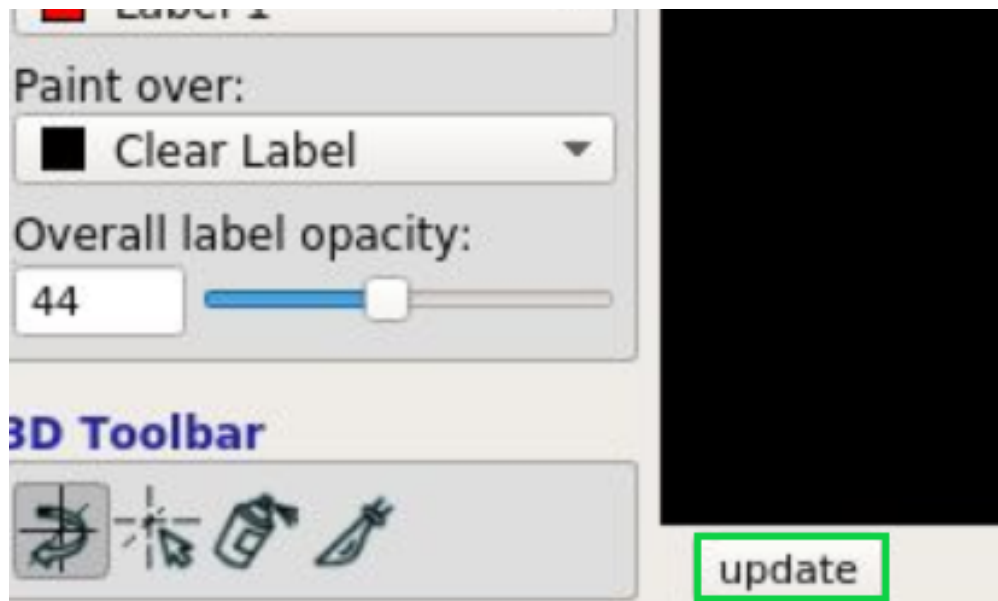


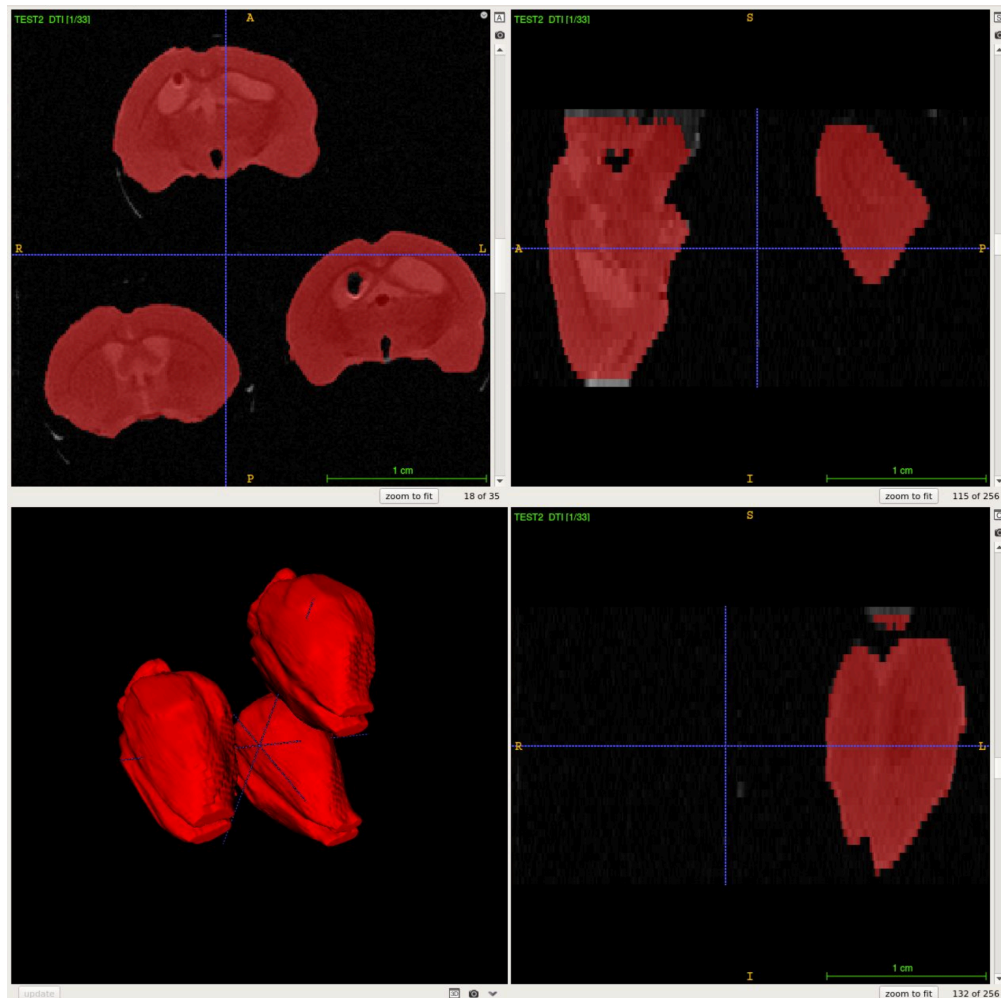
18



Once the region(s) of interest are completely filled, select **Finish**.

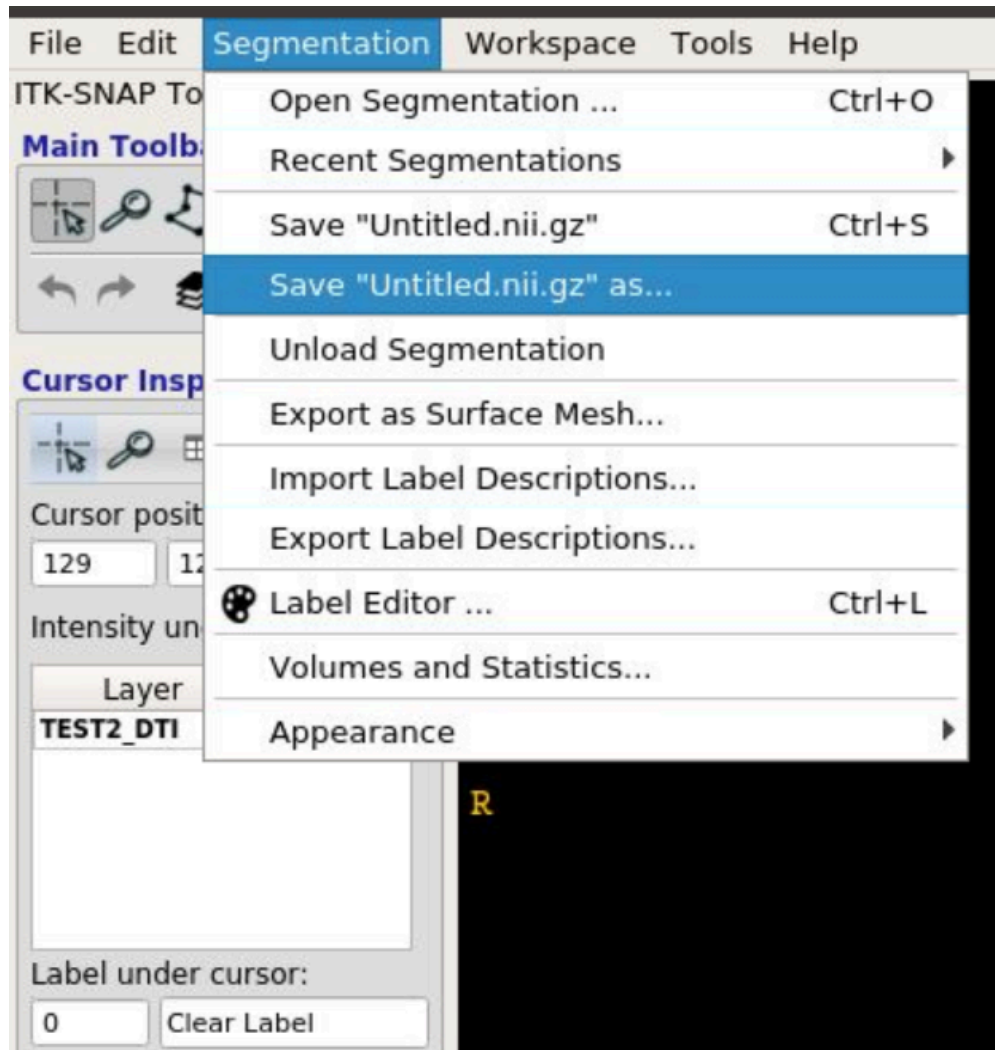
- 19 Press **Update** in the lower left corner of the ITKsnap interface to view the new 3D segmentation image.





- 20 Select **Segmentation** and **Save As** from the drop-down menu. You will be prompted to save your file as a **__.nii.gz** file type.

It is recommended you name your file __mask to clarify segmentation has already been performed.



- 21 Close out of ITKsnap.
- 22 In an open terminal, unzip the **____.nii.gz** file that was segmented in the previous step.

Command

Unzip file(s) of interest.

gunzip

gunzip __.nii.gz

- 23 Estimate diffusion tensor and Bo amplitude.

Command

Generates diffusion tensor and B0 amplitude. (TORTOISE)

EstimateTensorNLLS

EstimateTensorNLLS -i __DTI_DMC.list -m __mask.nii

This command will generate 2 files:

- **__N1_DT.nii** (diffusion tensor)
- **__DT_AM.nii** (B0 amplitude)

- 24 Compute all tensor maps on the **__N1_DT.nii** output file from the previous step.

Command

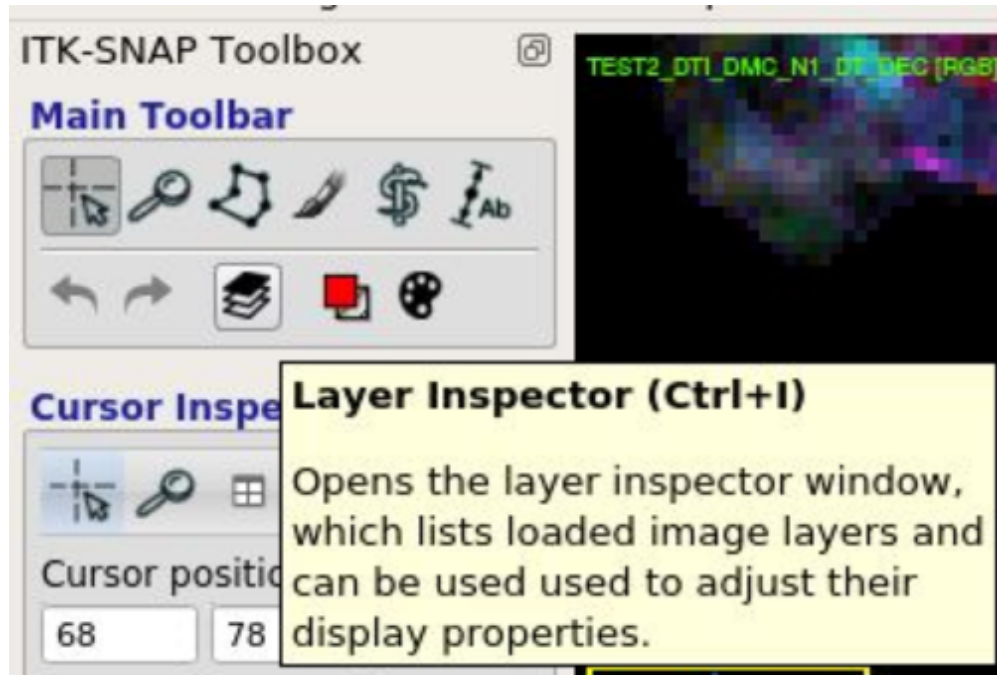
Computer all tensor maps. (TORTOISE)

ComputeAllTensorMaps.bash

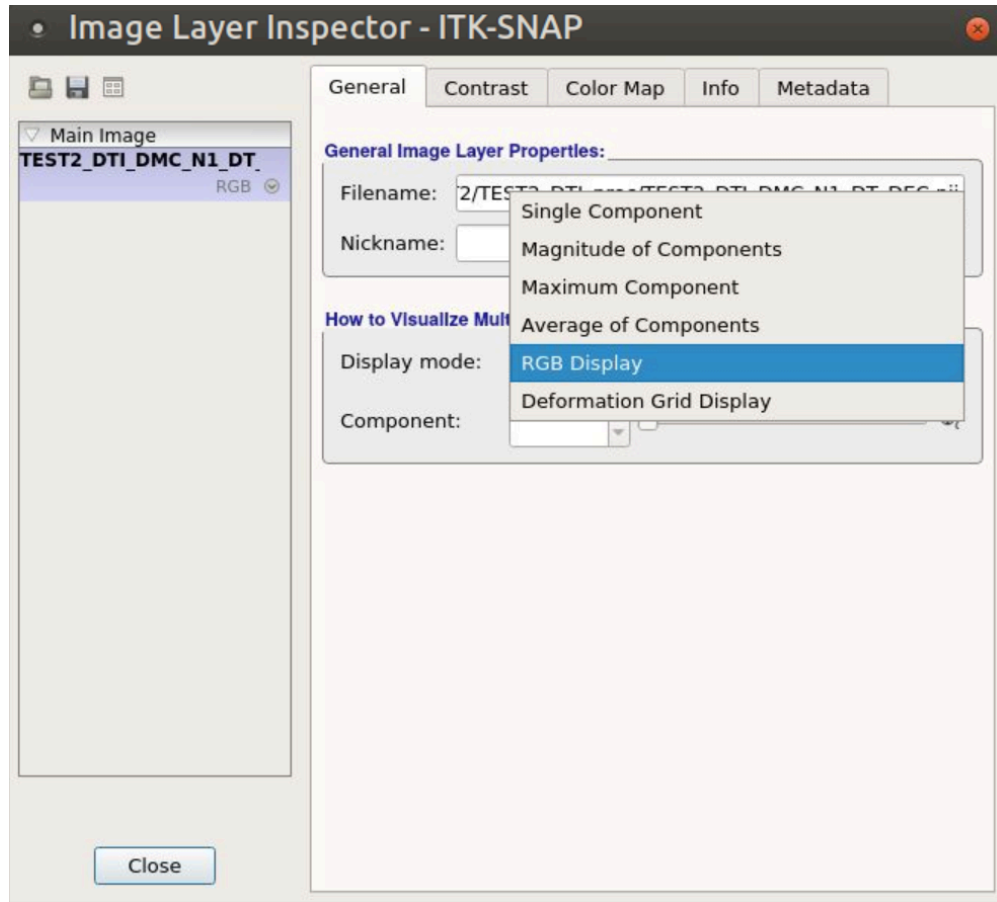
ComputeAllTensorMaps.bash __N1_DT.nii

25 Open the **__DEC.nii** output file in ITKsnap to view the primary eigenvector map.

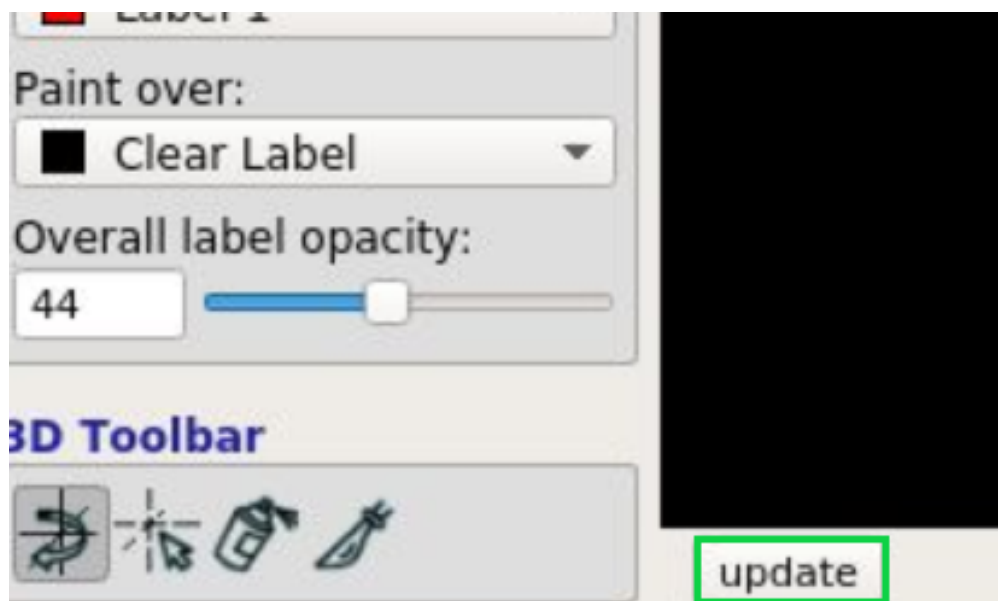
26 Within ITKsnap, select **Layer Inspector**.



27 From the **Layer Inspector** window, select the **Display mode** drop-down menu and select **RGB Display**.



- 28 Press **Update** in the lower left corner of the ITKsnap interface to view the image in the new display mode.



- 29 The final result will show a colorized image of your region of interest that allows a visual guide that estimates location, orientation, and anisotropy of the brain's white matter tracts based on the diffusion of water molecules.

