## Region of Interest Drawing and Usage in AFNI

- Goal: manually select Regions of Interest (ROIs) based on anatomical structures, then analyze functional datasets within these regions
  - Alternative procedure for selection of ROIs: geometrically connected clusters of 'activity' (supra-threshold voxels in some functional statistical map)
  - $\diamond$  Even with this method, you might want to manually adjust the ROIs
- ROIs are stored as regular AFNI datasets; it is only the user (you) that decides whether a particular dataset is a ROI mask
  - $\diamond$  Nonzero voxels are "in" the mask; zero voxels are "outside"
  - ◊ It is best to store a ROI mask as a functional dataset, so you can overlay it in color on an anatomical dataset
- Outline of procedure:
  - ♦ On the main AFNI control panel, set the anatomical underlay dataset (with <u>Switch Anatomy</u>) to be what you want to draw on—usually a SPGR or MP-RAGE type of dataset
  - ♦ Start the Draw Dataset plugin (also called the Editor)
  - $\hookrightarrow \texttt{Define Datamode}{\rightarrow}\texttt{Plugins}{\rightarrow}\texttt{Draw Dataset}$

- $\hookrightarrow$  Creating an all zero dataset can be done with the Editor plugin, or with the Copy Dataset plugin
- $\hookrightarrow$  You could instead edit a ROI dataset that you had already created before
- $\hookrightarrow$  It is required that the dataset being edited and the dataset that is the anatomical underlay have the same geometry (voxel size, etc.), since drawing/editing is done on a voxel-by-voxel basis
- Set the functional overlay dataset (with <u>Switch Function</u>) to be the dataset you are editing, and turn the functional overlay on (with <u>See Function</u>)
- $\diamond$  Draw the ROIs into the functional dataset, eventually Save-ing the results
- ♦ Convert the anatomical-resolution ROI dataset into a dataset at the resolution of the functional datasets you want to analyze with the ROI
- → The ROI and functional datasets may already be at the same resolution, if you are operating in +tlrc coordinates
- $\hookrightarrow {\sf Resolution\ conversion\ of\ masks\ is\ done\ with\ program\ {\tt 3dfractionize}}$
- ♦ Use programs <u>3dmaskave</u>, <u>3dmaskdump</u>, and <u>3dROIstats</u> to extract ROIbased information about functional datasets
- $\hookrightarrow$  Also can use <u>ROI</u> <u>Average</u> plugin to extract interactively the average of a dataset over a ROI



- Critical things to remember:
  - Solution Solution Solution of the same function of the same function.
  - $\hookrightarrow$  Otherwise, you won't see anything when you edit!
  - ♦ When drawing, you are putting numbers into a dataset brick
    - $\hookrightarrow$  These numbers are written to disk only when you do <u>Save</u>, <u>SaveAs</u>, or <u>Done</u>; before then, you can <u>Quit</u> (or exit AFNI) to get the unedited dataset back

- Step 1: Load a dataset to be edited
  - $\diamond$  Choose Dataset button gives you a list of datasets that
    - (a) Actually have brick data with only one sub-brick; and
    - (b) Are at the same voxel dimension, grid size, etc., as the current anatomical underlay dataset
  - $\diamond$  When you are starting, you probably don't want to edit an existing dataset
  - ♦ To get an all zero copy of the anatomical underlay, click the <u>Copy</u> button on, and set the controls to its right to <u>Zero</u>, <u>Func</u>, <u>As</u> <u>Is</u>
    - 1. Data would make a copy of the dataset with the actual data values
    - 2. Func makes the copy be a functional dataset, no matter what the original is; Anat would make the copy be an anatomical dataset; As Is would leave the copy in whatever mode the original was
    - 3. As Is means to keep the voxel values in the copy as the same type as in the original; you can also change the voxels values to be stored as
      - $\triangleright$  Bytes (integer values: 0..255) 1 byte each
      - $\triangleright$  Shorts (integer values: -32767..32767) 2 bytes each
      - ▷ Floats (fractional values) 4 bytes each
      - Bytes and Shorts make the most sense for ROI masks, where you are essentially attaching labels to voxels
  - ♦ Then Choose Dataset, pick the dataset you want a copy of (e.g., the anatomical underlay), and then Set

- Step 2: Drawing the ROI (or ROIs)
  - $\diamondsuit$  Choose the value to draw into the dataset
    - $\hookrightarrow$  If you are drawing only 1 ROI, then the default value of 1 is good
    - $\hookrightarrow$  If you are drawing multiple ROIs, then you should choose a different numerical value for each so that they can be distinguished later
      - ▷ And write down which number corresponds to which ROI!
  - $\diamond$  Choose the drawing color
  - $\hookrightarrow$  This is the color that is shown while you are drawing
  - $\hookrightarrow$  Color choice is only important to give a good contrast with underlay
  - → After you finish a drawing motion, the voxels you drew will be filled with the drawing value, the image will be redisplayed, and the colors will be determined by the Define Function control panel
  - $\diamond$  Choose the drawing mode:
    - $\hookrightarrow$  Open Curve  $\Rightarrow$

Drawing action produces a continuous open-ended curve

 $\hookrightarrow \texttt{Closed Curve} \Rightarrow$ 

Drawing action produces a continuous closed-ended curve (not filled inside)

 $\hookrightarrow$  Points  $\Rightarrow$ 

Only points actually drawn over are filled (used to "touch up" an ROI)

 $\hookrightarrow$  Flood->Value  $\Rightarrow$ 

Flood fill outwards from drawn point, stopping when the flood hits the current drawing value (used to fill a closed curve)

 $\hookrightarrow \texttt{Flood}\texttt{-}\texttt{Nonzero} \Rightarrow$ 

Flood fill outwards from drawn point, stopping when the flood hits any nonzero voxel (used to fill between regions)

 $\hookrightarrow$  Zero->Value  $\Rightarrow$ 

Flood fill with zero—instead of the drawing value—stopping when the flood hits the current drawing value (I forget why this is useful)

 $\hookrightarrow$  Filled Curve  $\Rightarrow$ 

Drawing action produces a continuous closed-ended curve (filled inside)

- $\diamond$  Actually draw something
- $\hookrightarrow$  Drawing is done with mouse Button 2 ("middle" button) in a 2D slice image
- $\hookrightarrow$  Hold the button down in the image window during a single drawing action
- $\hookrightarrow$  While the drawing action is happening, the chosen drawing color will trace the screen pixels you draw over
- → When you release the button, these pixels are converted to voxels, and the dataset is actually changed, using the drawing value and drawing mode you selected
- $\hookrightarrow$  At this point, the color of the drawn region will change to reflect the drawing value and the setup of the Define Function control panel

♦ Undo button will let you take back the last drawing action

- $\hookrightarrow$  Only one level of undo is available: <u>Undo</u> then <u>Undo</u> will take you back to where you just finished drawing
- $\diamond$  You can only draw on one 2D slice image at a time
- $\hookrightarrow$  If you draw on a montage display, only screen pixels overlaying the first image you Button 2 click in will count
- $\hookrightarrow$  While drawing, if you cross over between sub-images in the montage, unexpected effects will result

▷ But there is always Undo to the rescue!

- Step 3: Save your results

  - $\hookrightarrow$  You could also then choose another dataset to edit
  - SaveAs will let you write the current dataset to disk under a new name, creating a new dataset, then continue editing the new dataset
  - $\diamond$  Quit exits editing and closes the plugin window, without saving to disk any changes since the last Save
  - $\hookrightarrow$  Exiting AFNI has the same effect
  - $\diamond$  Done is equivalent to Save then Quit

## • Optional Drawing Steps:

- ♦ Linear Fillin lets you draw a 3D ROI not in every slice, but in every third slice (say), and then go back and fill in the gaps
  - $\hookrightarrow$  For example, if you draw in coronal slices, then you want to fill in the <u>A-P</u> direction (the default)
  - $\hookrightarrow$  If you draw every  $n^{\text{th}}$  slice, then you want to set Gap to n-1
  - $\hookrightarrow$  Line segments of voxels in the fillin direction that have the current drawing value at each end, and have no more than <u>Gap</u> zero voxels in between, will get their gap voxels filled with the drawing value
  - $\hookrightarrow$  After you try this, you will probably have to touch up the dataset manually
  - $\hookrightarrow$  Fillin cannot be undone be careful!
  - ← This operation can also be done with program <u>3dRowFillin</u>, which creates a new dataset (and thus provides a way to undo: the <u>rm</u> command)
- ♦ TT Atlas Region to Load lets you load regions from the Talaraich Daemon database into the dataset voxels
- → Requires that you be drawing in +tlrc coordinates, or at least have a transformation from +orig→+tlrc computed in the current directory
- $\hookrightarrow$  Choose a region to draw into the dataset (e.g., Hippocampus)
- $\hookrightarrow$  Load: OverWrite will fill all voxels in the region with the drawing value
- $\hookrightarrow$  Load: InFill will fill only voxels in the region that are currently zero
- $\hookrightarrow$  You probably want to edit the results manually to fit the subject

- Drawing and Rendering at the Same Time (totally fun, and maybe useful):
  - $\diamond$  You cannot draw into the rendering plugin, but you can use it to see in 3D what you are drawing in 2D
  - $\hookrightarrow$  If you meet the dataset criteria for rendering (usually in +tlrc coordinates)
  - $\diamond$  How to set up the renderer:
  - ← Choose the underlay to be the current anatomical dataset (or a "scalped" version, from 3dIntracranial)
  - $\hookrightarrow$  Choose the overlay dataset to be the dataset you are editing
  - $\hookrightarrow \mathsf{Turn} \text{ on See Overlay}$
  - $\hookrightarrow$  Set Color Opacity to ShowThru (or ST+Dcue)
  - $\hookrightarrow \mathsf{Turn} \text{ on } \mathsf{DynaDraw}$
  - → Drawing in a 2D image window immediately triggers a redraw in the rendering window (if the 2D and 3D overlay datasets are the same)
  - $\hookrightarrow$  This is only useful if your computer is fast enough to render quickly (< 1 s per frame)



## Things to Do with ROI Datasets

- ROIs are used on a voxel-by-voxel basis to select parts of datasets
- If you draw at the anatomical resolution and want to use the ROI dataset at the functional resolution, you probably have to convert the high-resolution ROI dataset to a low-resolution dataset (unless you are working in +tlrc coordinates)
- Program <u>3dfractionize</u> does this resolution conversion: 3dfractionize -template func+orig -input ROI\_hires+orig -clip 0.5 -preserve -prefix ROI\_lores
  - $\diamond$  <u>-template</u> func+orig  $\Rightarrow$  New dataset is written at resolution of func+orig
  - $\diamond$  -input ROI\_hires+orig  $\Rightarrow$  Defines the input high-resolution dataset
  - $\diamond \frac{-\text{clip } 0.5}{50\%} \Rightarrow \text{Output voxels will only get a nonzero value if they are at least}$
- One large output voxel may be partially or fully covered by several nonzero smaller input voxels; <u>3dfractionize</u> computes the overlap fraction for each output voxel and uses that fraction to decide what value to put in the voxel

• <u>N.B.</u>: A ROI is defined by the values stored in the voxels of the 'mask' dataset

 $\diamond$  Contiguity of voxels has no meaning to the ROI software described below

 $\diamond$  Two voxels are in the same ROI if they have the same value in the mask dataset

• <u>3dmaskave</u>: program to compute the average of voxels from a dataset, with the voxels selected from a mask dataset (interactive version: ROI Average plugin)

 $\diamond$  Example:

will print out 1 line for each of the 3 selected sub-bricks in the input dataset func\_fred+tlrc:

44.287 [137 voxels] 27.021 [137 voxels] -33.22 [137 voxels]

Each line is the average of the sub-brick values over the selected voxels: all voxels whose value in the mask dataset are between 1 and 3 (inclusive), as specified by -mrange 1 3

♦ The <u>-q</u> option suppresses the voxel count printout; it is useful if you want to compute the average time series over a ROI and save it to a 1D file for later use in AFNI:

```
3dmaskave -mask ROI_fred+tlrc -mrange 1 1
    epi01_fred+tlrc > epi01_fred_ROI_1.1D
```

• <u>3dmaskdump</u>: program to dump out all voxels in a dataset (or datasets) that match some values given in mask dataset

 $\diamond$  Mask dataset is usually a ROI dataset

 $\diamond$  Example:

```
3dmaskdump -noijk -mask ROI_fred+tlrc
```

```
-mrange 1 1 'func_fred+tlrc[2,5,7]'
```

would produce output like so for each voxel whose value in ROI\_fred+tlrc was 1:

12 37 42

- 27 0 321
- 4 4 444

That is, each voxel that passes the -mask and -mrange tests (i.e., is nonzero in the mask and has its value in the range of numbers given by -mrange) gets one line printed with the values that are in sub-bricks #2, #5, and #7 from dataset func\_fred+tlrc at that voxel.

- $\diamondsuit$  More than one dataset can be given at the end of the command line
- Main application is to dump out the data or functional activation values that match a ROI so they can be processed in some other program (e.g., Excel)
- $\diamond$  If -noijk option is omitted, each output line starts with ijk-indexes of the voxel
- $\hookrightarrow {\sf Program} \ \underline{{\sf 3dUndump}} \ {\sf can} \ {\sf be} \ {\sf used} \ {\sf to} \ {\sf create} \ {\sf a} \ {\sf dataset} \ {\sf from} \ {\sf a} \ {\sf text} \ {\sf file} \ {\sf with} \ {\sf ijk-indexes} \ {\sf and} \ {\sf dataset} \ {\sf values}$

• <u>3dROIstats</u>: program to compute average of voxels from a dataset, using multiple regions selected by a single ROI dataset

♦ Averaging is done over each region defined by a distinct value in the ROI dataset

 $\diamond$  Example:

3dROIstats -mask ROI\_fred+tlrc 'func\_fred+tlrc'

produces output that looks like this:

File	Sub-brick	Mean_1	Mean_2	Mean_3
<pre>func_fred+tlrc</pre>	0	33.3	44.4	55.5
<pre>func_fred+tlrc</pre>	1	22.2	66.6	-21.2
<pre>func_fred+tlrc</pre>	2	2.3	9.7	666.666

The Mean\_1 column is the average over the ROI whose mask value is 1, Mean\_2 over ROI with mask value 2, etc.

- ◊ Very useful if you create ROI masks for a number of subjects, using the same number codes for the same anatomical regions (e.g., 1=hippocampus, 2=amygdala, 3=superior temporal gyrus ...).
- Solution of Sol

## Creating ROI Datasets from Activation Maps

• The program <u>3dmerge</u> can find contiguous supra-threshold voxel clusters in an activation map and then convert each cluster into a ROI with a separate data value

 $\diamondsuit$  These ROIs can then be used as starting points for some analysis

```
• Example:
```

```
3dmerge -1clust_order 1 500
-1tindex 1 -1thresh 0.4
-prefix ROI_func func+tlrc
```

Does the following:

- $\diamond$  Thresholds the dataset on sub-brick #1 at value 0.4
- $\diamond$  Clusters together all the surviving nonzero voxels using a contiguity test of 1 mm and keeping only clusters at least 500 mm<sup>3</sup> in volume
- ♦ All voxels in the largest cluster are assigned value 1, in the second largest assigned 2, etc., and the result is written to disk in dataset ROI\_func+tlrc
- You can now use this dataset as a mask, edit it with the drawing plugin, ...