

- First analysis: Simple regression with fixed hemodynamic model

- ◇ Will use 3 regressors to correspond to the three stimulus classes
- ◇ Step 1: Create 0-1 time series files to correspond to absence-presence of annular, anti-annular, and motor stimuli:

Files: ann_stim_07.1D antiann_stim_07.1D righthand_stim_07.1D

↪ Created with a text editor (in this case, vi)

- ◇ Step 2: Create hemodynamic response to stimuli time series, via program waver:

```
waver -GAM -dt 2.0 -input ann_stim_07.1D > ann_response_07.1D
```

```
waver -GAM -dt 2.0 -input antiann_stim_07.1D > antiann_response_07.1D
```

```
waver -GAM -dt 2.0 -input righthand_stim_07.1D > righthand_response_07.1D
```

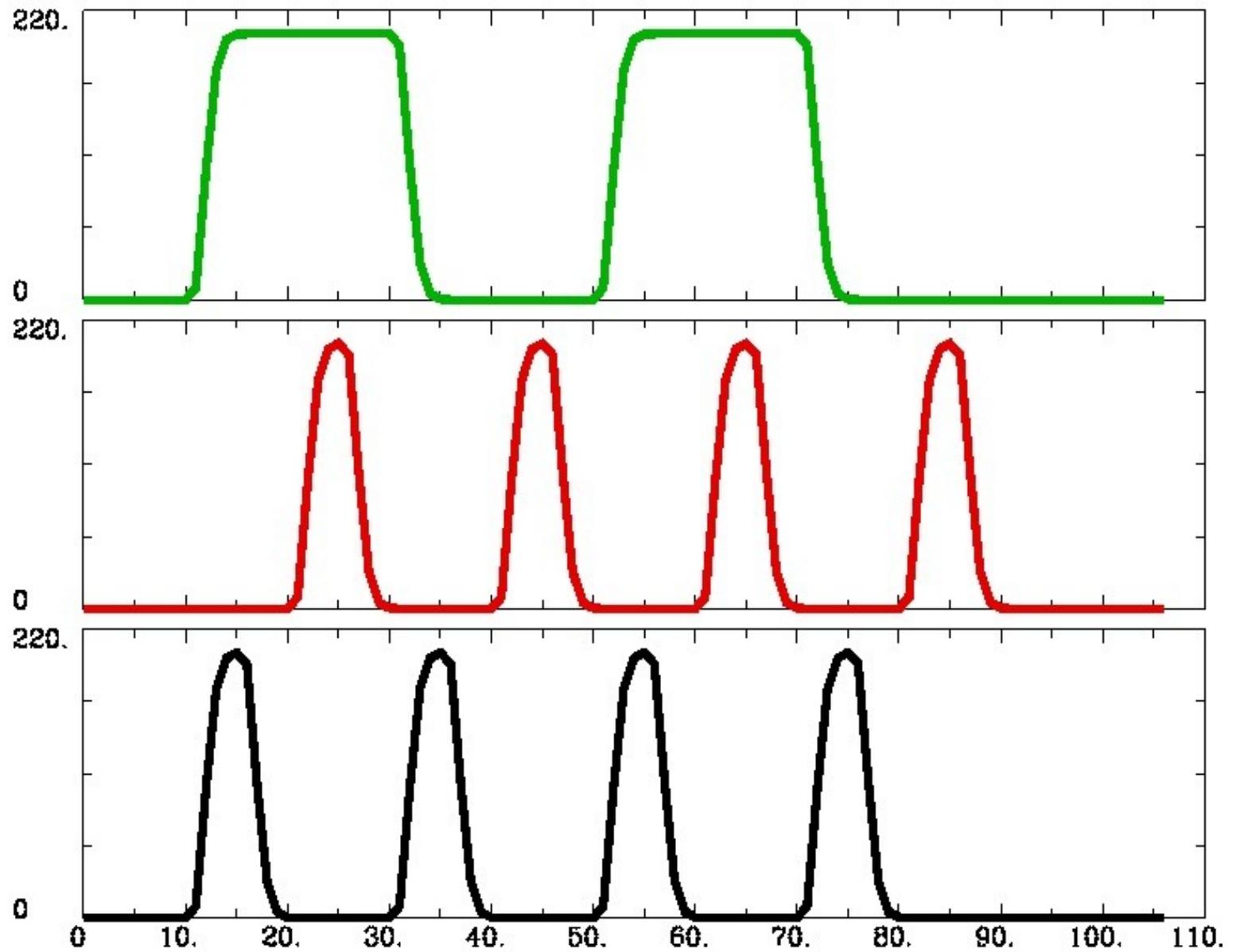
(cf. Script File waver_07a)

↪ Put all 3 responses together into 1 file with 3 columns:

```
1dcat ann_response_07.1D antiann_response_07.1D \  
righthand_response_07.1D > all3_response_07.1D
```

↪ Plot them (to make sure we haven't made a stupid mistake yet):

```
1dplot all3_response_07.1D
```



◇ Carry out regression analysis (cf. Script File decon_07a):

```
3dDeconvolve -input epi07+orig.HEAD \
             -num_stimts 3 \
             -stim_file 1 ann_response_07.1D \
             -stim_file 2 antiann_response_07.1D \
             -stim_file 3 righthand_response_07.1D \
             -stim_label 1 annulus \
             -stim_label 2 antiann \
             -stim_label 3 motor \
             -stim_minlag 1 0 -stim_maxlag 1 0 \
             -stim_minlag 2 0 -stim_maxlag 2 0 \
             -stim_minlag 3 0 -stim_maxlag 3 0 \
             -fitts epi07a_fitts \
             -fout -bucket epi07a_stats
```

↪ Each stimulus has minimum and maximum lags set to 0

⇒ Signal model is just the sum of the response time series we input

↪ We keep the stimulus labels short since the AFNI menus don't like long labels

↪ -fitts option ⇒ We get the fitted model as an output dataset

↪ -fout -bucket options ⇒ We get the F -statistics in a bucket dataset

↪ We skip the individual regressor t -statistics (no -tout): these are redundant with the partial-model F -statistics when we have only 1 lag

▷ Each t tests one coefficient; each partial- F tests all coefficients in that sub-model; 1 degree of freedom in a sub-model ⇒ $t^2 = F$

↪ We don't need -ignore option, since this dataset doesn't have the T1-equilibration spike at the beginning (must have been pre-saturated)

◇ Look at results with interactive AFNI program:

↪ Since data slices were axial, open Axial image and graph for dataset epi07

↪ Let's look at output of -fitts graph overlaid on data time series:

▷ In AFNI controller: Define Datamode→Plugins→Dataset#2



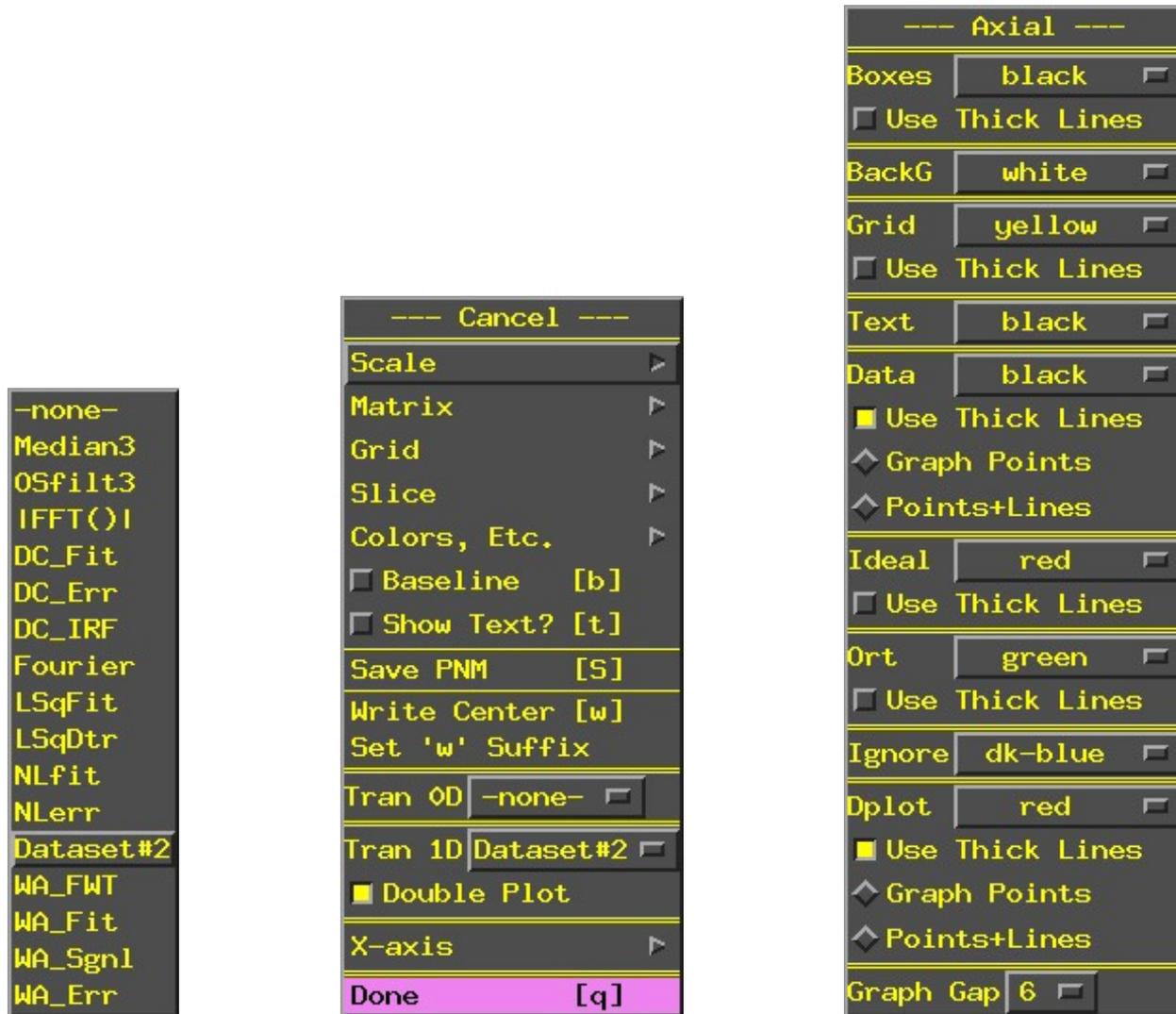
Dataset#2 Plugin Control window



Dataset Chooser window

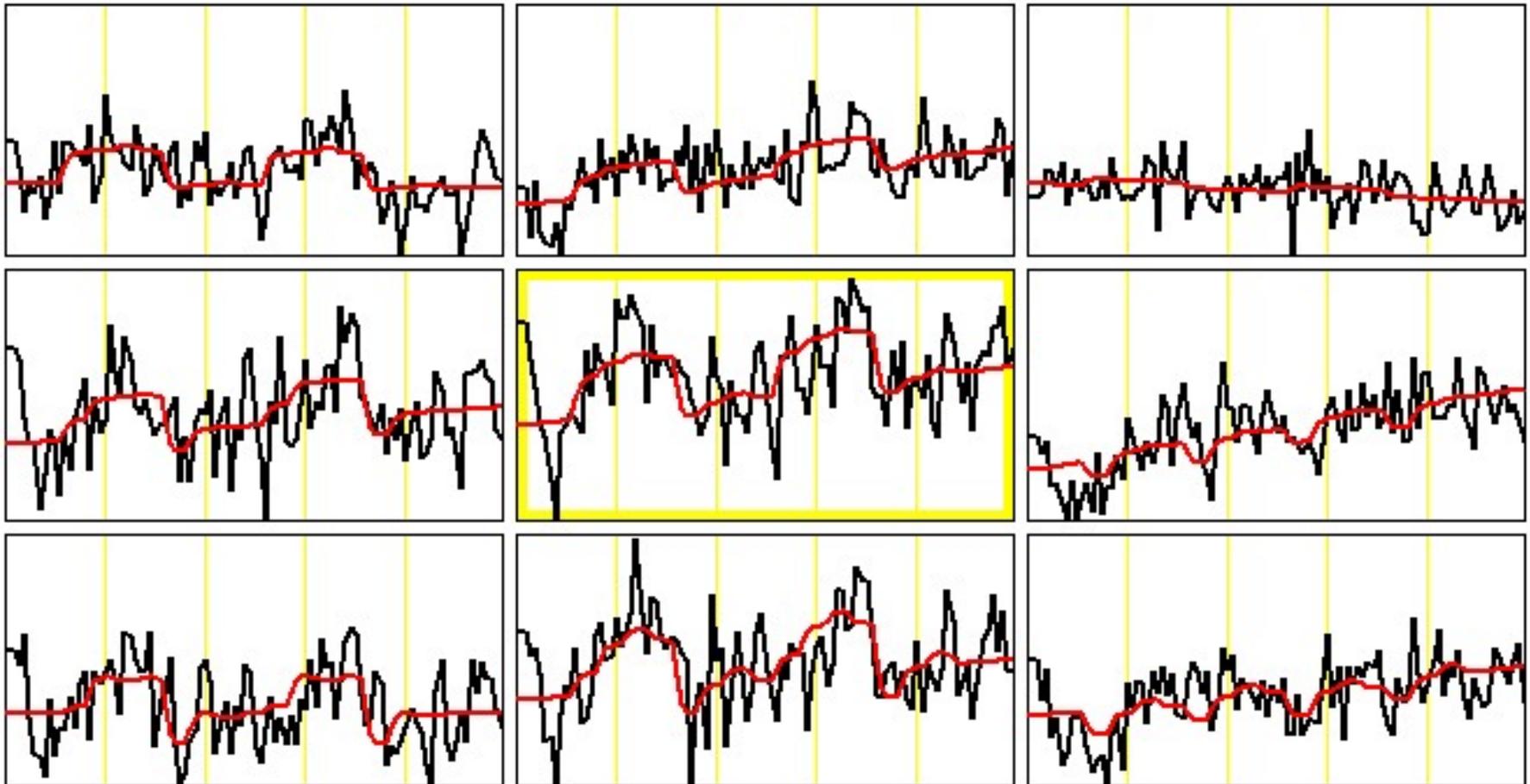
- ▷ Choose the fit dataset by clicking -- Choose Dataset --
Then click on the epi07a_fitts+orig choice in the chooser
Then Set in the chooser (will close chooser window)
Then Run+Close in the plugin's window (tells AFNI about your choice)

- ▷ From the graph window's Opt menu:
- select Tran 1D→Dataset#2
- press in Double Plot
- press in Opt→Use Thick Lines (under the Dplot item)



The menus you will encounter when setting up for Double Plot

▷ Sample Dataset#2 Double Plot (x,y,z=-48 [R], 14 [P], -26 [I]):



- ▷ Can get identical graphs using the Deconvolution plugin and Double Plotting with the Tran 1D→DC_Fit transformation function
- Regression analysis done “on-the-fly” for voxels shown in graph window
 - Button3 graph popup shows fit parameters/statistics for sub-graph fit

↳ Let's look at the color overlay from the functional dataset

▷ Open Define Function control panel

The image shows a software control panel for defining a function. It features a central color map with a vertical slider and a table of data. The panel is annotated with several labels:

- Color map**: Points to the vertical color bar on the left.
- Hidden popup menu here**: Points to the top of the color map.
- Choose which dataset makes the underlay image**: Points to the 'Options' section.
- Choose which sub-brick of Functional dataset makes the color**: Points to the 'Func' dropdown menu.
- Choose which sub-brick of Functional dataset is the threshold**: Points to the 'Thr' dropdown menu.
- Shows ranges of data in Anatomical and Functional dataset**: Points to the table of values.
- Shows automatic range for color scaling**: Points to the 'autoRange' field.
- Rotates color map**: Points to the 'Rota' field.
- Lets you choose range for color scaling**: Points to the 'Rota' field.
- Shows voxel values at focus**: Points to the 'Anat = 1077' value.
- Number of panes in color map**: Points to the '# 9' field.
- Positive-only or both signs of function?**: Points to the '** 0' checkbox.
- Choose range of threshold slider, in powers of 10**: Points to the '[N/A]' field.
- p-value of current threshold**: Points to the '.5000' value on the slider.

Base t^0 Inten

1.00
0.75
0.50
0.25
0.05
-0.05
-0.25
-0.50
-0.75
-1.00

Options

- Anat underlay
- Func underlay
- Func @Thr underlay

Func #0 Base t^0 Coef

Thr #0 Base t^0 Coef

Anat	0:	2111
Func	0.378448:	2066.769
Thr	0.378448:	2066.769

autoRange: 2066.769

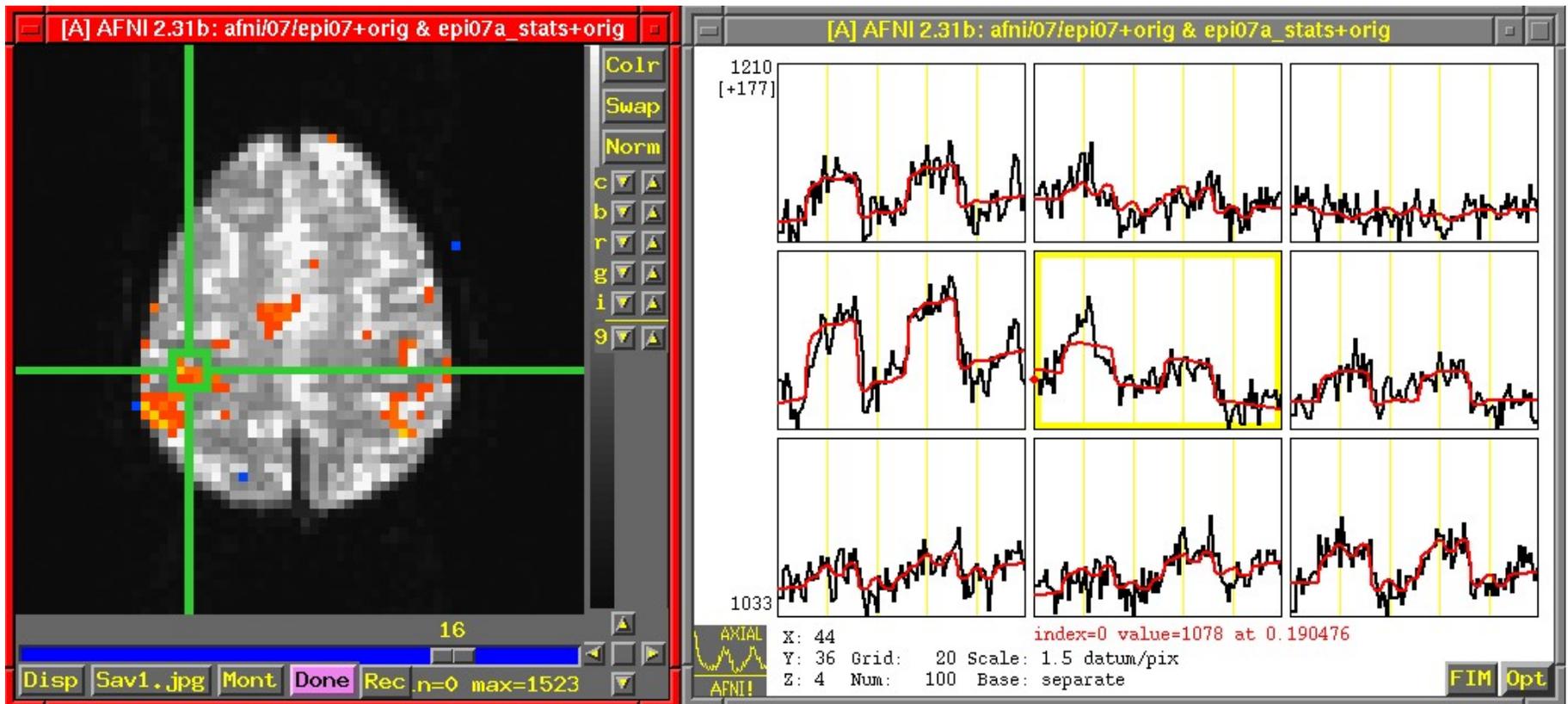
[N/A] # 9 Rota

** 0 Pos?

Anat = 1077
Func = ?
Thr = ?

See TT Atlas Regions

- ▷ Set the Func control to #6 motor [0] Coef
- Set the Thr control to #8 Full F-stat
- Set threshold slider range to ** 1
- Set threshold slider to about 9.8 (p -value about 1.0×10^{-5})
- Click on See Function
- Move Axial image viewer to slice 16 (where we'll see some motor activity)



- ▷ Exercises:
 - Find motor voxels that are modulated by visual stimuli
 - Find visual activation areas

- Second analysis: Deconvolution

- ◇ Since each stimulus interval for each stimulus class is the same in this experiment (4×10 s for visual, 2×40 s for motor), could try deconvolve whole response for each class

- ↳ Would treat experiment as having 4 visual events of each class (with about 10 lags), and 2 motor events (with about 25 lags)

- ↳ But wouldn't work very well, since we'd be estimating $2+10+10+25=57$ parameters from 100 time points \Rightarrow garbage results (over-fitting the data)

- ▷ Exercises:

- Explain this calculation

- Carry out this analysis anyway, and see what happens

- ◇ Instead, will treat each TR with a stimulus as a discrete event, and try to deconvolve hemodynamic response function

- ↳ Not an ideal experiment for this analysis, since only have 2 “turn on” and 2 “turn off” transitions for motor response

- ↳ Will use lags from 0 to 6 (0 s to 12 s), since expect hemodynamics to be about that long

- ↳ Will have $2+7+7+7=23$ unknown parameters (vs. $2+1+1+1=5$ in the simple regression analysis)

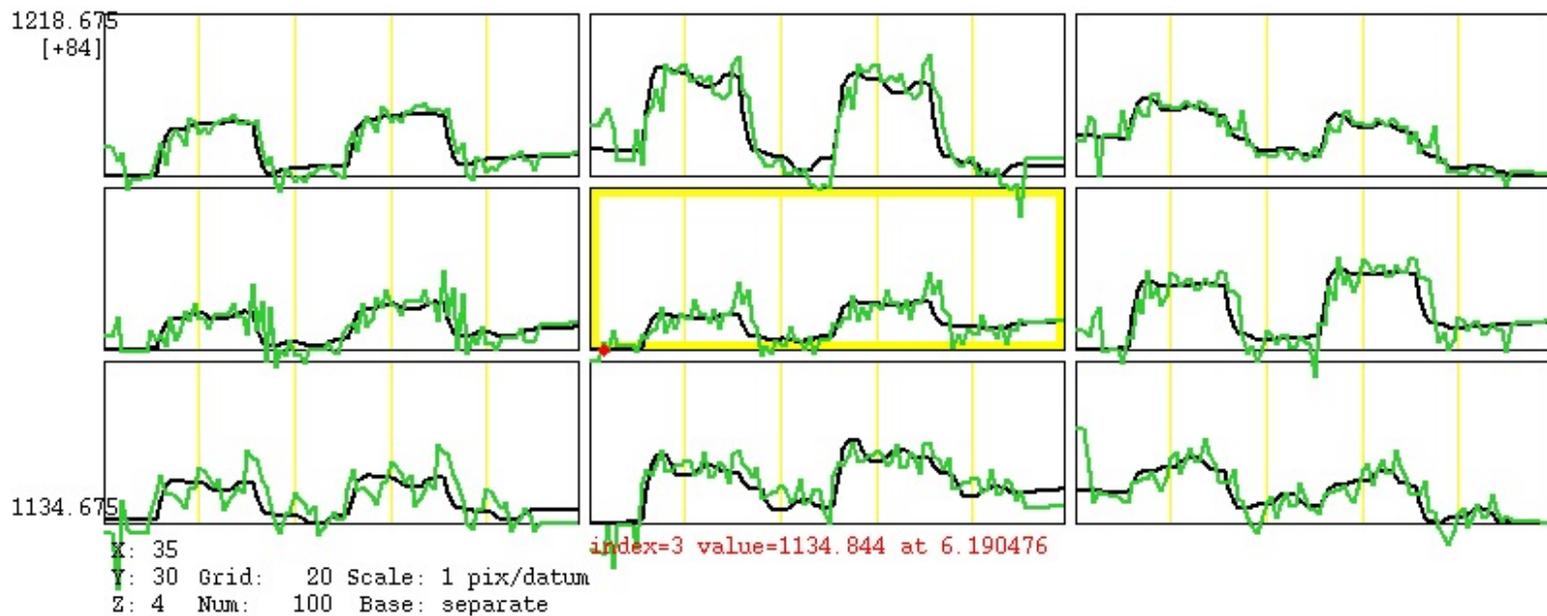
- ↳ Input functions will be the 0-1 stimulus time series files created earlier

- Carry out deconvolution analysis (cf. Script File decon_07b):

```
3dDeconvolve -input epi07+orig.HEAD \
             -num_stimts 3 \
             -stim_file 1 ann_stim_07.1D \
             -stim_file 2 antiann_stim_07.1D \
             -stim_file 3 righthand_stim_07.1D \
             -stim_label 1 annulus \
             -stim_label 2 antiann \
             -stim_label 3 motor \
             -stim_minlag 1 0 -stim_maxlag 1 6 \
             -stim_minlag 2 0 -stim_maxlag 2 6 \
             -stim_minlag 3 0 -stim_maxlag 3 6 \
             -fitts epi07b_fitts \
             -iresp 1 epi07b_iresp_ann \
             -iresp 2 epi07b_iresp_antiann \
             -iresp 3 epi07b_iresp_motor \
             -sresp 1 epi07b_sresp_ann \
             -sresp 2 epi07b_sresp_antiann \
             -sresp 3 epi07b_sresp_motor \
             -fout -tout -bucket epi07b_stats
```

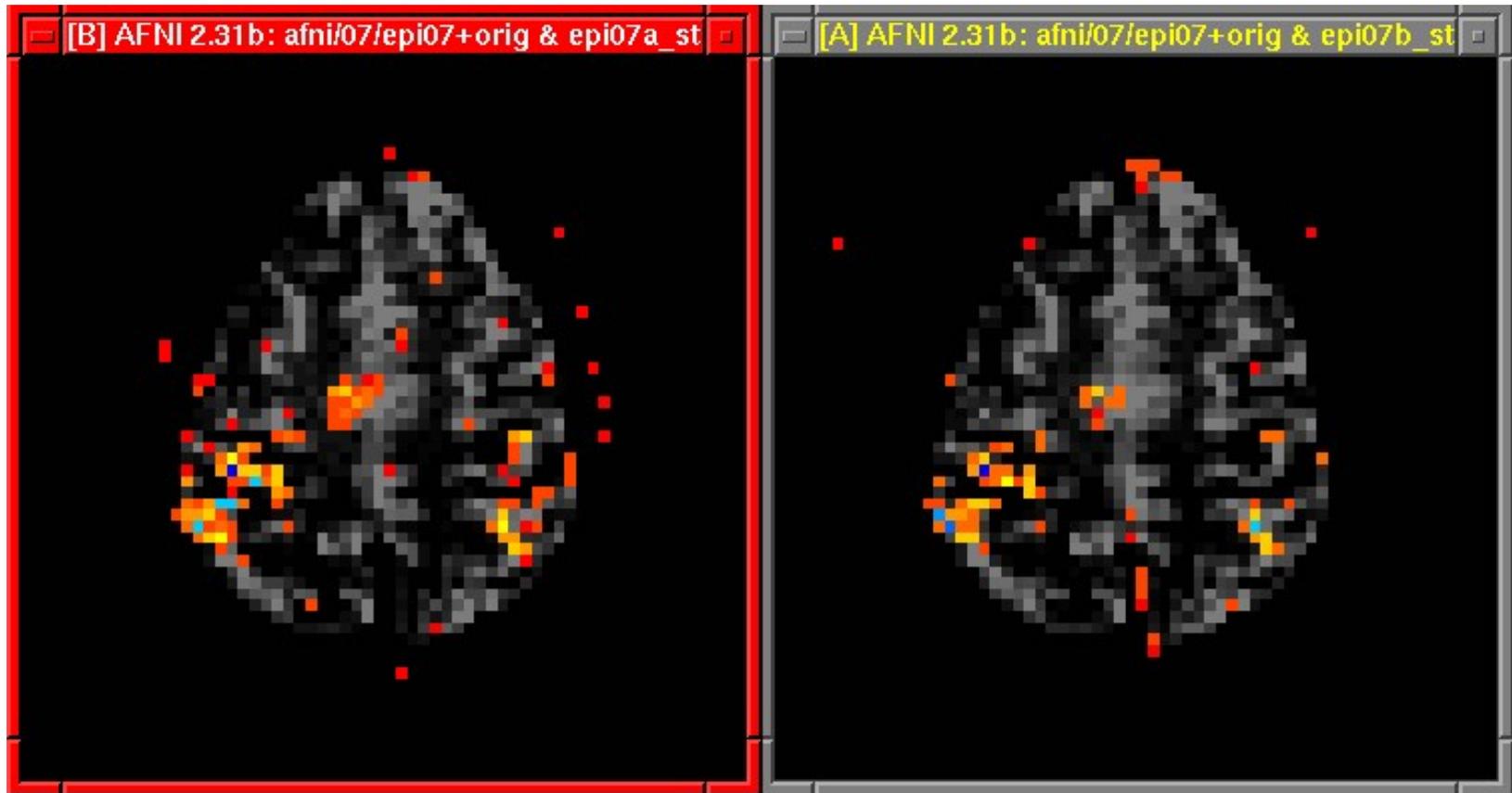
- ◇ Saving impulse response and its standard error for graphing (-iresp, -sresp)
- ◇ Saving individual lag coefficient t -statistics this time (-tout)

- Different ways to look at the results with AFNI:
 - ◇ Graph `epi07b_iresp_*` time series with `epi07b_sresp_*` Double Plot-ed to see estimate hemodynamic response and it standard error together
 - ◇ Set the threshold sub-brick to the full F-statistic to see where any activation was detected
 - ◇ Set the threshold sub-brick to the partial F-statistics to see where individual stimuli correlated activity was detected
 - ◇ Compare the simple regression and deconvolution time series fits



↪ Looks like deconvolution is “over-fitting” the data somewhat

- ◇ Compare the simple regression and deconvolution activation maps



Motor: Simple Regression

Motor: Deconvolution

- ↪ Displaying Axial slice 16 (again)
- ↪ Thresholding on Full F-stat, set to $p=1.0e-4=10^{-4}$ (per voxel)
- ↪ Color overlay is from motor F-stat (using Pos, autoRange, 11 colors)
- ↪ Exercise: Why is there somewhat less activation in the Deconvolution map?