

afni_proc.py
is your *friend*

or it will be soon

Example scripts from

AFNI_data6/FT_analysis

Also see <https://arxiv.org/abs/1709.07471>

Appendix has processing scripts

What the hell is afni_proc.py?

- It is a Python program that
 - Takes as input a series of “options” describing processing steps to use to analyze datasets *from one subject*
 - Produces as output a Unix tcsh script file that runs all the **AFNI** programs to do the processing
- Reasons to use afni_proc.py
 - It is flexible and compact, to produce a long script
 - The output script not only does the data analysis, but also saves various diagnostic tools and files
 - All intermediate output datasets are saved to help diagnose things when results are confusing or just plain wrong
 - You can get help from us on the **AFNI** message board

Where do afni_proc.py command lines/scripts come from?

- **Method #1:**
 - take an existing script (from yourself or a friend), and modify it to meet your needs
- **Method #2:**
 - find an approximate fit to what you want in examples from afni_proc.py's help, and modify to meet your needs
- **Method # $\sqrt{-1}$:**
 - use GUI uber_subject.py
- **Method #666:**
 - beg for help on the **AFNI** message board

Starting Simple - 1

```
afni_proc.py -subj_id FT \
  -dsets FT/FT_epi_r?+orig.HEAD \
  -copy_anat FT/FT_anat+orig \
  -tcat_remove_first_trs 2 \
  -regress_stim_times FT/AV*.txt \
  -regress_stim_labels Vrel Arel \
  -regress_basis 'BLOCK(20,1)' \
  -regress_est_blur_errts \
  -regress_opts_3dD \
  -gltsym 'SYM: Vrel -Arel' \
  -glt_label 1 V-A
```

Script file
s01.ap.simple

Starting Simple - 2

`afni_proc.py` `-subj_id FT`

ID will label
output files

```
-dsets FT/FT_epi_r?+orig.HEAD \
-copy_anat FT/FT_anat+orig \
-tcat_remove_first_trs 2 \
-regress_stim_times FT/AV*.txt \
-regress_stim_labels Vrel Arel \
-regress_basis 'BLOCK(20,1)' \
-regress_est_blur_errts \
-regress_opts_3dD \
      -gltsym 'SYM: Vrel -Arel' \
      -glt_label 1 V-A
```

Script file
s01.ap.simple

Starting Simple - 3

```
afni_proc.py -subj_id FT
```

EPI time
series
datasets
to analyze

```
-dsets FT/FT_epi_r?+orig.HEAD
```

```
-copy_anat FT/FT_anat+orig
```

```
-tcat_remove_first_trs 2
```

```
-regress_stim_times FT/AV*.txt
```

```
-regress_stim_labels Vrel Arel
```

```
-regress_basis 'BLOCK(20,1)'
```

```
-regress_est_blur_errts
```

```
-regress_opts_3dD
```

```
    -gltsym 'SYM: Vrel -Arel'
```

```
    -glt_label 1 V-A
```

Script file
s01.ap.simple

Starting Simple - 4

```
afni_proc.py -subj_id FT \
  -dsets FT/FT_epi_r?+orig.HEAD \
  -copy_anat FT/FT_anat+orig \
  -tcat_remove_first_trs 2 \
  -regress_stim_times FT/AV*.txt \
  -regress_stim_labels Vrel Arel \
  -regress_basis 'BLOCK(20,1)' \
  -regress_est_blur_errts \
  -regress_opts_3dD \
  -gltsym 'SYM: Vrel -Arel' \
  -glt_label 1 V-A
```

T1-weighted
anatomical
dataset for
alignment to
EPI datasets

Script file
s01.ap.simple

Starting Simple - 5

```
afni_proc.py -subj_id FT \
  -dsets FT/FT_epi_r?+orig.HEAD \
  -copy_anat FT/FT_anat+orig \
  -tcat_remove_first_trs 2 \
  -regress_stim_times FT/AV*.txt \
  -regress_stim_labels Vrel Arel \
  -regress_basis 'BLOCK(20,1)' \
  -regress_est_blur_errts \
  -regress_opts_3dD \
  -gltsym 'SYM: Vrel -Arel' \
  -glt_label 1 V-A
```

Stimulus timing files, labels, and HRF model;
Note: timing files have *start* times for each task iteration

Script file
s01.ap.simple

Starting Simple - 6

```
afni_proc.py -subj_id FT \
  -dsets FT/FT_epi_r?+orig.HEAD \
  -copy_anat FT/FT_anat+orig \
  -tcat_remove_first_trs 2 \
  -regress_stim_times FT/AV*.txt \
  -regress_stim_labels Vrel Arel \
  -regress_basis 'BLOCK(20,1)' \
  -regress_est_blur_errts \
  -regress_opts_3dD \
  -gltsym 'SYM: Vrel -Arel' \
  -glt_label 1 V-A
```

Estimate
smoothness
of EPI noise
for group
analysis

Script file
s01.ap.simple

Starting Simple - 7

```
afni_proc.py -subj_id FT \
  -dsets FT/FT_epi_r?+orig.HEAD \
  -copy_anat FT/FT_anat+orig \
  -tcat_remove_first_trs 2 \
  -regress_stim_times FT/AV*.txt \
  -regress_stim_labels Vrel Arel \
  -regress_basis 'BLOCK(20,1)' \
  -regress_est_blur_errts \
  -regress_opts_3dD \
  -gltsym 'SYM: Vrel -Arel' \
  -glt_label 1 V-A \
```

Set up
General
Linear
Test
between 2
conditions

Script file
s01.ap.simple

A Real Case - 1

Script file
s05.ap.uber

```
#!/usr/bin/env tcsh
```

```
# creation date: Thu Sep 10 14:27:59 2015
```

```
# set data directories
```

```
set top_dir      = FT
```

```
# set subject and group identifiers
```

```
set subj         = FT
```

```
set group_id    = horses
```

Not actually
used here

Code subject level information
into shell variables:

Makes it easier to re-use this afni_proc.py command

A Real Case - 2

Script file
s05.ap.uber

```
afni_proc.py -subj_id $subj \
  -script proc.$subj -scr_overwrite \
  -blocks tshift align tlrc volreg blur mask scale regress \
  -copy_anat $top_dir/FT_anat+orig \
  -dsets \
    $top_dir/FT_epi_r1+orig.HEAD \
    $top_dir/FT_epi_r2+orig.HEAD \
    $top_dir/FT_epi_r3+orig.HEAD \
  -volreg_align_to MIN_OUTLIER \
  -volreg_align_e2a \
  -volreg_tlrc_warp \
  -blur_size 4.0 \
  -tcat_remove_first_trs 2 \
  -regress_stim_times \
    $top_dir/AV1_vis.txt \
    $top_dir/AV2_aud.txt \
  -regress_stim_labels \
    vis aud \
  -regress_basis 'BLOCK(20,1)' \
  -regress_censor_motion 0.3 \
  -regress_opts_3dD \
    -jobs 2 \
    -gltsym 'SYM: vis -aud' -glt_label 1 V-A \
    -gltsym 'SYM: 0.5*vis +0.5*aud' -glt_label 2 mean.VA \
  -regress_compute_fitts \
  -regress_make_ideal_sum sum_ideal.1D \
  -regress_est_blur_epits \
  -regress_est_blur_errts \
  -regress_run_clustsim yes \
  -execute
```

The entire afni_proc.py
command:
Font size will be bigger
on following slides!

A Real Case – 3b

Script file
s05.ap.uber

```
afni_proc.py -subj_id $subj \
  -script proc.$subj -scr_overwrite \
  -blocks tshift align tlrc volreg blur mask scale regress \
  -copy_anat $top_dir/FT_anat+orig \
  -dsets \
    $top_dir/FT_epi_r1+orig.HEAD \
    $top_dir/FT_epi_r2+orig.HEAD \
    $top_dir/FT_epi_r3+orig.HEAD \
  -volreg_align_to MIN_OUTLIER \
  -volreg_align_e2a \
  -volreg_tlrc_warp \
  -blur_size 4.0
```

Select input
datasets
(anat and EPI)

A Real Case – 3c

Script file
s05.ap.uber

```
afni_proc.py -subj_id $subj \
  -script proc.$subj -scr_overwrite \
  -blocks tshift align tlrc volreg blur mask scale regress \
  -copy_anat $top_dir/FT_anat+orig \
  -dsets \
    $top_dir/FT_epi_r1+orig.HEAD \
    $top_dir/FT_epi_r2+orig.HEAD \
    $top_dir/FT_epi_r3+orig.HEAD \
  -volreg_align_to MIN_OUTLIER \
  -volreg_align_e2a \
  -volreg_tlrc_warp \
  -blur_size 4.0
```

Specify how
“volreg” step
will operate

A Real Case – 3d

Script file
s05.ap.uber

```
afni_proc.py -subj_id $subj \
  -script proc.$subj -scr_overwrite \
  -blocks tshift align tlrc volreg blur mask scale regress \
  -copy_anat $top_dir/FT_anat+orig \
  -dsets \
    $top_dir/FT_epi_r1+orig.HEAD \
    $top_dir/FT_epi_r2+orig.HEAD \
    $top_dir/FT_epi_r3+orig.HEAD \
  -volreg_align_to MIN_OUTLIER \
  -volreg_align_e2a \
  -volreg_tlrc_warp \
  -blur_size 4.0
```

Specify how
much spatial
blurring will
be used
(FWHM mm)

A Real Case – 4b

Script file
s05.ap.uber

```
-tcat_remove_first_trs 2
-regress_stim_times
    $stop_dir/AV1_vis.txt
    $stop_dir/AV2_aud.txt
-regress_stim_labels
    vis aud
-regress_basis 'BLOCK(20,1)'
-regress_censor_motion 0.3
-regress_opts_3dD
    -jobs 2
    -gltsym 'SYM: vis -aud' -glt_label 1 V-A
    -gltsym 'SYM: 0.5*vis +0.5*aud' -glt_label 2 mean.VA
-regress_compute_fitts
-regress_make_ideal_sum sum_ideal.1D
-regress_est_blur_epits
-regress_est_blur_errts
-regress_run_clustsim yes
-execute
```

Maximum motion
(in mm) to accept
between
successive TRs

A Real Case – 4d

Script file
s05.ap.uber

```
-tcat_remove_first_trs 2
-regress_stim_times
    $stop_dir/AV1_vis.txt
    $stop_dir/AV2_aud.txt
-regress_stim_labels
    vis aud
-regress_basis 'BLOCK(20,1)'
-regress_censor_motion 0.3
-regress_opts_3dD
    -jobs 2
    -gltsym 'SYM: vis -aud' -glt_label 1 V-A
    -gltsym 'SYM: 0.5*vis +0.5*aud' -glt_label 2 mean.VA
-regress_compute_fitts
-regress_make_ideal_sum sum_ideal.1D
-regress_est_blur_epits
-regress_est_blur_errts
-regress_run_clustsim yes
-execute
```

Other regression options:
Compute fitted model
(best fit to data);
Create sum of task ideal
response time series
(for display purposes)

A Real Case – 4e

Script file
s05.ap.uber

```
-tcat_remove_first_trs 2
-regress_stim_times
    $stop_dir/AV1_vis.txt
    $stop_dir/AV2_aud.txt
-regress_stim_labels
    vis aud
-regress_basis 'BLOCK(20,1)'
-regress_censor_motion 0.3
-regress_opts_3dD
    -jobs 2
    -gltsym 'SYM: vis -aud' -glt_label 1 V-A
    -gltsym 'SYM: 0.5*vis +0.5*aud' -glt_label 2 mean.VA
-regress_compute_fitts
-regress_make_ideal_sum sum_ideal.1D
-regress_est_blur_epits
-regress_est_blur_errts
-regress_run_clustsim yes
-execute
```

Estimate smoothness
of *noise* in the data:
From the dataset itself,
From the *residuals*
(=data-model fit).

And estimate cluster-
size thresholds from
smoothness estimates

A Real Case – 4f

Script file
s05.ap.uber

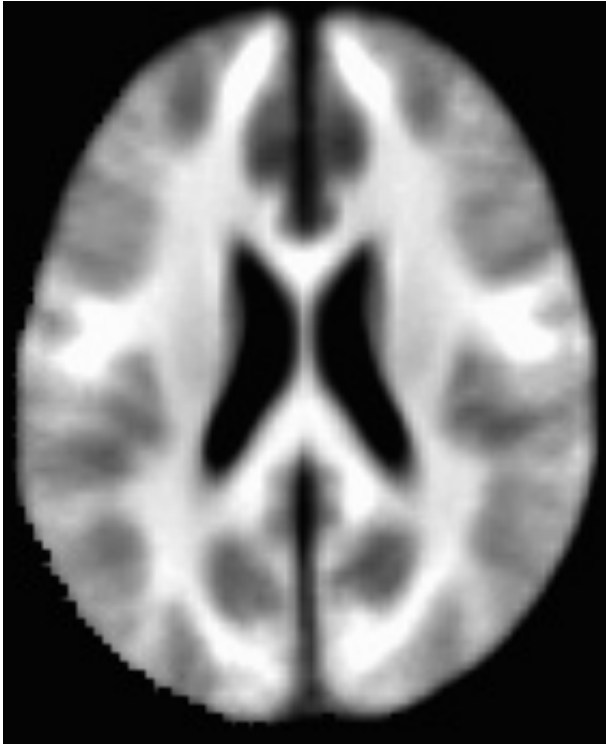
```
-tcat_remove_first_trs 2
-regress_stim_times
    $stop_dir/AV1_vis.txt
    $stop_dir/AV2_aud.txt
-regress_stim_labels
    vis aud
-regress_basis 'BLOCK(20,1)'
-regress_censor_motion 0.3
-regress_opts_3dD
    -jobs 2
    -gltsym 'SYM: vis -aud' -glt_label 1 V-A
    -gltsym 'SYM: 0.5*vis +0.5*aud' -glt_label 2 mean.VA
-regress_compute_fitts
-regress_make_ideal_sum sum_ideal.1D
-regress_est_blur_epits
-regress_est_blur_errts
-regress_run_clustsim yes
-execute
```

Run script after
it is created

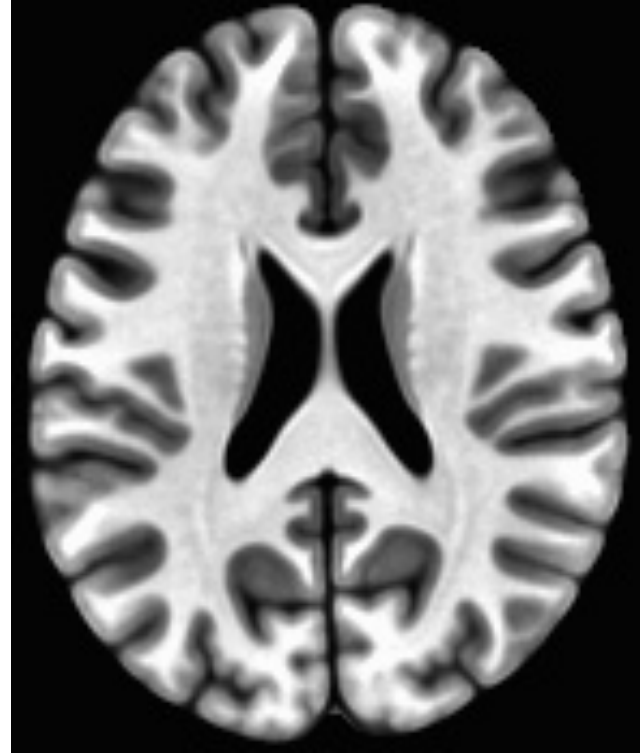
Nonlinear Warping to MNI Template

- afni_proc.py *can* do the nonlinear warping for you
 - But, nonlinear warping is slow (in fact, slowly slow)
 - If you need to re-rerun subject analysis, nonlinear warping will slow the re-run script down *a lot*
- Solution: do the nonlinear warping *before* using afni_proc.py, then supply the warping results so that afni_proc.py will skip doing the warping itself
- Mechanism: the **@SSwarper** script (tcsh)
 - Does Skull Stripping ("SS") and nonlinear warping
 - Base dataset is **MNI152_2009_template.nii.gz**
 - Nonlinearly warped, not too blurry

Two MNI Templates



MNI152_1mm_uni+tlrc
Affine alignments

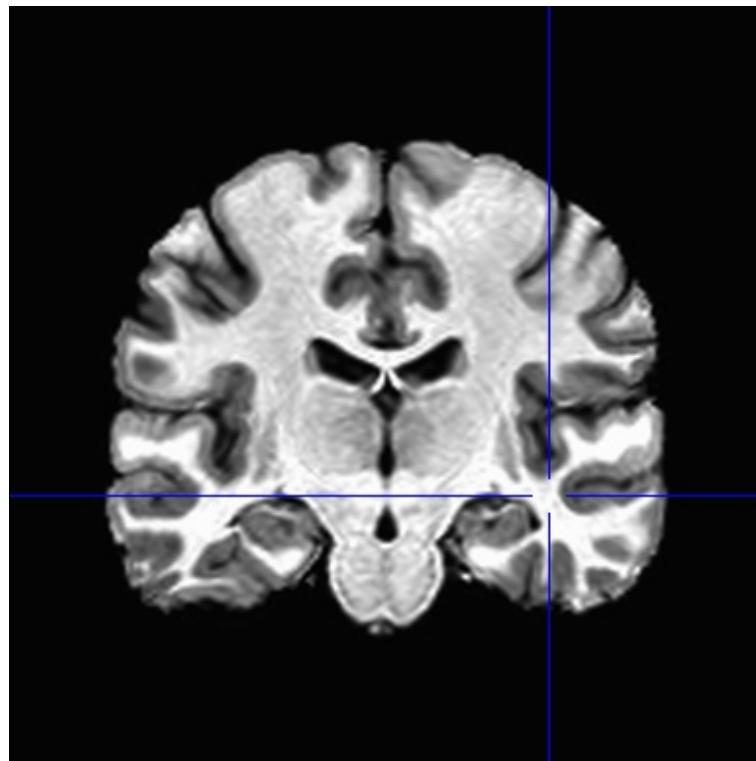
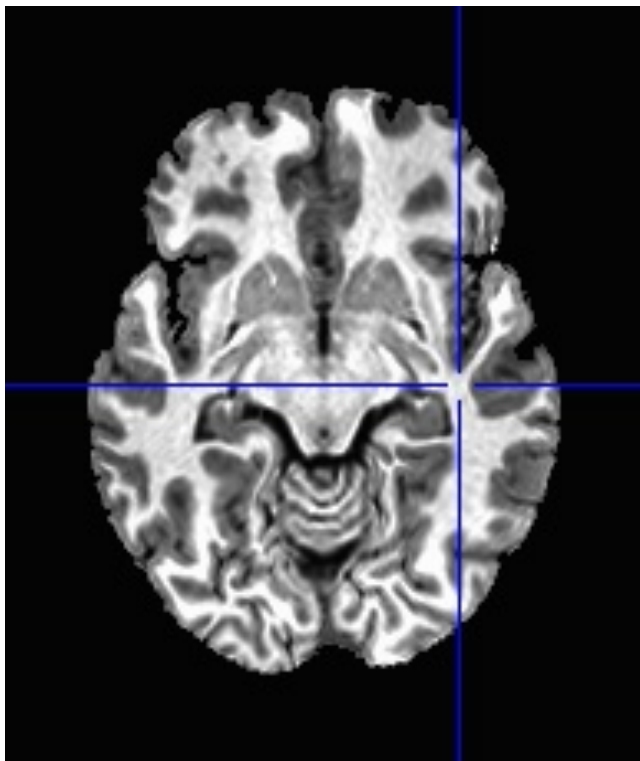


MNI152_2009_template.nii.gz
Nonlinear alignments

What @SSwarper Reads and Writes

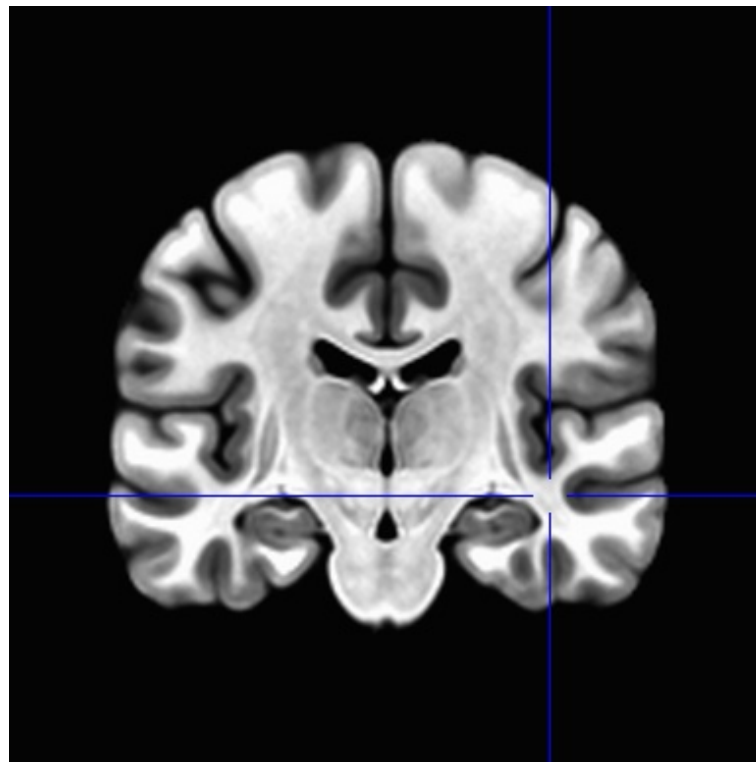
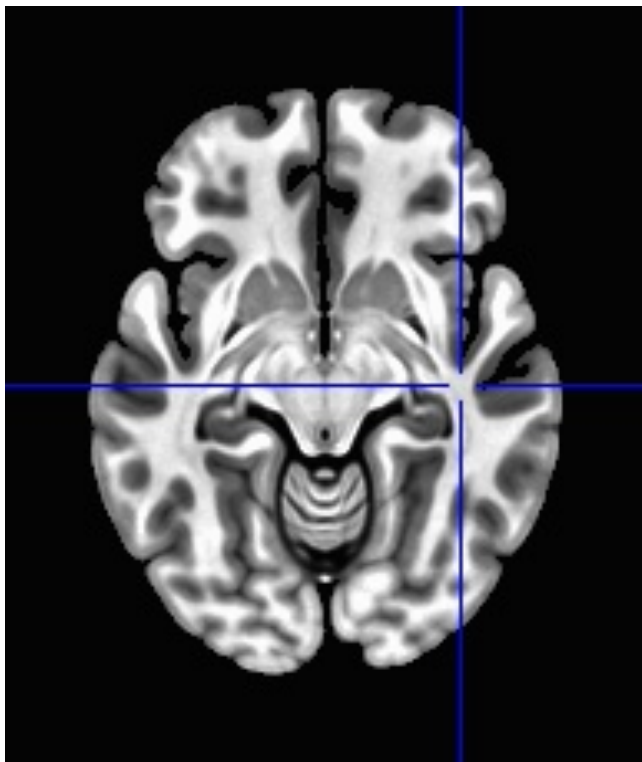
- Inputs:
 - T1-weighted anatomical image of subject (skull-on)
 - Subject ID code, for names of output files
- Outputs (subject ID = **sub007**):
 - **anatSS.sub007.nii**
 - skull-stripped dataset in original coordinates
 - **anatQQ.sub007.nii**
 - skull-stripped dataset, nonlinearly warped to MNI template
 - **anatQQ.sub007.aff12.1D**
 - affine matrix to transform original dataset to MNI template
 - **anatQQ.sub007_WARP.nii**
 - incremental warp from affine transformation to nonlinearly aligned dataset
- These files are needed for later use in `afni_proc.py`

@SSwarper Results



sub00440 from Beijing-Zang
in the FCON-1000 collection

MNI Template Slices



For comparison

Nonlinear Registration Script

- What follows is a script for doing nonlinear warping (registration) of *one* anatomical dataset to an MNI template
- In a real study, this script is run once for each subject
- Takes a long time, so the script should be submitted to a multi-node cluster

Nonlinear Registration - 1

```
#!/bin/tcsh
### This script nonlinear warps one anatomical dataset,
### taken from the anat_orig directory, to the MNI 2009
### nonlinear template (supplied with AFNI binaries), and
### puts the resulting files into anat_warped directory.
### The only command line argument is the subject ID
```

```
set subj = $argv[1]
```

```
set tempdir = .
```

```
# don't log AFNI programs in ~/.afni.log
```

```
# don't try any version checks
```

```
# don't auto-compress output files
```

```
setenv AFNI_DONT_LOGFILE YES
```

```
setenv AFNI_VERSION_CHECK NO
```

```
setenv AFNI_COMPRESSOR NONE
```

Nonlinear Registration - 2

```
### go to data directory
```

```
# topdir = directory above this Scripts directory
```

```
set topdir = `dirname $cwd`
```

```
cd $topdir/anat_orig
```

```
### create final output directories
```

```
mkdir -p $topdir/anat_warped
```

```
mkdir -p $topdir/anat_warped/snapshots
```

```
### create temp directory to hold work, and copy anat there
```

```
mkdir -p temp_$subj
```

```
cp anat_$subj.nii.gz temp_$subj
```

```
cd temp_$subj
```

Nonlinear Registration - 3

```
### process the anat dataset, using the AFNI script
### that does the warping and skull-stripping
@SSwarper anat_${subj}.nii.gz $subj
# compress the output datasets
gzip -lv *.nii
### move the results to where they belong
# skull-stripped original, Q-warped dataset, and the warps
\mv -f anatSS.${subj}.nii.gz      anatQQ.${subj}.nii.gz      \
    anatQQ.${subj}.aff12.1D anatQQ.${subj}_WARP.nii.gz \
    $topdir/anat_warped
# snapshots for visual inspection
\mv -f *.jpg $topdir/anat_warped/snapshots
# delete the temporary directory
cd ..
\rm -rf temp_${subj}
exit 0
```

Nonlinear Registration - 4

Add these lines above `afni_proc.py` command:

```
set basedset = MNI152_2009_template.nii.gz
set tpath = `@FindAfnIDsetPath $basedset`
if( "$tpath" == '' ) then
    echo "***** @SSwarper -- Failed to find $basedset :("
    exit 1
endif
set basedset = $tpath/$basedset
```

Add these options to `afni_proc.py` command:

```
-copy_anat anat_warped/anatSS.${subj}.nii          \  
-tlrc_base $basedset                                \  
-tlrc_NL_warp                                       \  
-tlrc_NL_warped_dsets                              \  
$warpdir/anatQQ.${subj}.nii.gz                     \  
$warpdir/anatQQ.${subj}.aff12.1D                   \  
$warpdir/anatQQ.${subj}_WARP.nii.gz                \  

```


Use 3dREMLfit (GLSQ) instead of 3dDeconvolve (OLSQ)

Add these options to `afni_proc.py` command:

```
-regress_3dD_stop \
-regress_reml_exec
```

Use @snapshot_volreg to check alignments

In the output directory

@snapshot_volreg

A television test pattern background with the text "AND NOW BACK TO OUR REGULARLY SCHEDULED PROGRAMMING" overlaid. The background consists of a grid of colored squares: a top black bar, a row of seven colored squares (grey, yellow, cyan, green, magenta, red, blue), a row of four colored squares (blue, black, black, grey), a row of seven colored squares (dark blue, white, purple, black, black, black, black), and a bottom black bar. The text is in a bold, white, sans-serif font with a black outline, centered horizontally and spanning across the colored squares.

AND NOW BACK TO
OUR REGULARLY
SCHEDULED
PROGRAMMING