

# HW2 Report — Daehyun Cho

## [1] Preprocessing and GLM

### **(a) Preprocess and GLM**

(b) Review overall results with afni\_proc.py QC with a html format.

## [2] Statistical Analysis

(a) One-sample t-test

(b) Paired t-test for all pairs of tasks

(c) One-way repeated ANOVA

## [3] Discuss Overall results obtained from [1] and [2]

## [1] Preprocessing and GLM

### (a) Preprocess and GLM

Here is my `afni_proc.py` command and I've automated them into one tcsh file since it takes too long time...

```
#!/bin/tcsh

foreach subj (s*)
    foreach task (ad_evt lh_evt rh_evt vs_evt)

        set prefix = $subj$task
        afni_proc.py \
            -subj_id $prefix \
            -dsets ./$subj/$task+orig.HEAD \
            -copy_anat ./$subj/anat/anat_orig/anat_$subj.nii.gz \
            -blocks despike tshift align tlrc volreg blur mask scale

        regress \

            -tlrc_base /home/jj/abin/MNI152_T1_2009c+tlrc.BRIK.gz \
            -volreg_align_to MIN_OUTLIER \
            -volreg_align_e2a \
            -volreg_warp_dxyz 3 \
            -mask_epi_anat yes \
            -blur_size 8 \
            -regress_reml_exec \
            -regress_apply_mask \
            -html_review_style pythonic \
            -execute

    end
end
```

Required preprocessing steps and their corresponding commands are below

---

1. **despiking** `-blocks despike`
2. **slice timing correction** `-blocks tshift`
3. **motion correction** `-blocks align`
4. **spatial normalization to the MNI space** `-blocks tlrc volreg`  
 with **3mm isotropic voxel size** `-volreg_warp_dxyz 3`
  - so this needed an extra option `-tlrc_base`  
`/home/jj/abin/MNI152_T1_2009c+tlrc.BRIK.gz` because I thought the default was Talairach space. But later found out that the default space was MNI. What a waste of time...
  - `-volreg_align_to MIN_OUTLIER` It's to specify the base position for volume reg but normally we use third. Since I didn't know when MP data was taken, I chose `MIN_OUTLIER`
  - `-volreg_align_e2a` Aligned EPI to anatomy at volreg step.
5. **Spatial smoothing** `-blocks -blur`  
 using **8mm isotropic FWHM Gaussian kernel** `-blur_size 8`
6. **Scaling** of the voxel intensity to an average of 100 `-blocks scale`
  - default average was 100, no extra options needed.
  - `-blocks mask` and `-mask_epi_anat` options were added since default scaling processes were set to extended mask, which is not our desirable results.
7. **temporal detrending**
  - ss
8. **General Linear Model** `-blocks regress`
  -

**(b) Review overall results with afni\_proc.py QC with a html format.**

This was done with `-html_review_style pythonic`

Those results

## [2] Statistical Analysis

The first thing to do was to resample all the `stats.` files after GLM since they had number of voxels and stuffs. I referenced the link below. Beforehand, I've put all the `stats.` from the same task into same folder(i.e. to have auditory tasks GLM results were all in the `ad` folder)

Re: T-Test Error: "Does match first one in size"

I'm currently in the process of adding more data to my current sample. The new data was collected during a different session. I am having issues with the t-test (one and two sample) and keep getting this error (see below, "does match first one in size") for ONLY the newly added data.

<https://afni.nimh.nih.gov/afni/community/board/read.php?1,107517,148717#msg-148717>

With `tcsh` script below, I've gone through all the resampling process.

```
#!/bin/tcsh

foreach file (*.gz)

    echo $file
    3dresample -master ../ad/stats.s02ad_evt_REMLvar+orig.BRIK.gz \
              -prefix res$file \
              -input $file

end
```

### (a) One-sample t-test

Command — `3dtttest++ -setA STATS*` ← file that starts with stats, results from GLM

### (b) Paired t-test for all pairs of tasks

Command — `3dtttest++ -setA STATSforTask1 -setB STATSforTask2` ← two file

### (c) One-way repeated ANOVA

## [3] Discuss Overall results obtained from [1] and [2]

### ***Gap between theory and practice***

Even though I get to know what's going on in theory, I can't directly match them to practice right away. There are too many exceptions, extra options for

minor edge cases which is good, but documentations are not intuitive to newly-users, which is sad...

tcsh

I used bash more than 5 years ago and it was good to automate some works with tcsh script file.