Scripting 1st Level Analysis

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Sec: 2 BN:25 ID:9203396

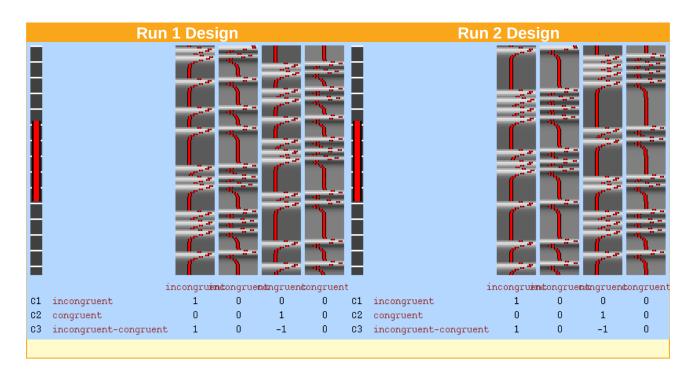
Make_FSL_Timings.sh:

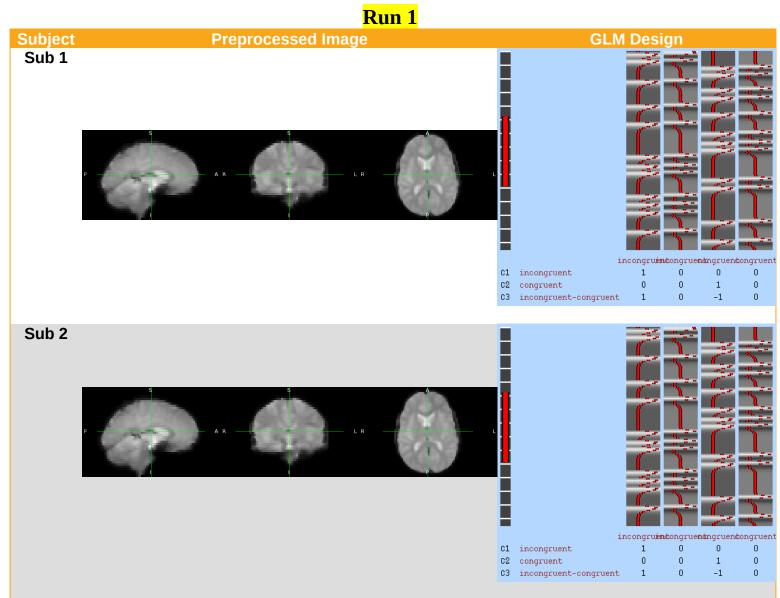
```
#!/bin/bash
#Check whether the file subjList.txt exists; if not, create it
if [!-f subjList.txt]; then
    ls -d sub-?? > subjList.txt
#Loop over all subjects and format timing files into FSL format
for subj in `cat subjList.txt`; do
    cd $subj/func #Navigate to the subject's func directory, which contains the timing
files
     #Extract the onset times for the incongruent and congruent trials for each run. NOTE:
This script only extracts the trials in which the subject made a correct response. Accuracy
is nearly 100% for all subjects, but as an exercise the student can modify this to extract the
incorrect trials as well.
     cat ${subj}_task-flanker_run-1_events.tsv | awk '{if ($3=="incongruent_correct")
{print $1, $2, "1"}}' > incongruent_run1.txt
     cat ${subj}_task-flanker_run-1_events.tsv | awk '{if ($3=="congruent_correct")
{print $1, $2, "1"}}' > congruent_run1.txt
     cat ${subj}_task-flanker_run-2_events.tsv | awk '{if ($3=="incongruent_correct")
{print $1, $2, "1"}}' > incongruent_run2.txt
     cat ${subj}_task-flanker_run-2_events.tsv | awk '{if ($3=="congruent_correct")
{print $1, $2, "1"}}' > congruent_run2.txt
    cd ../..
done
```

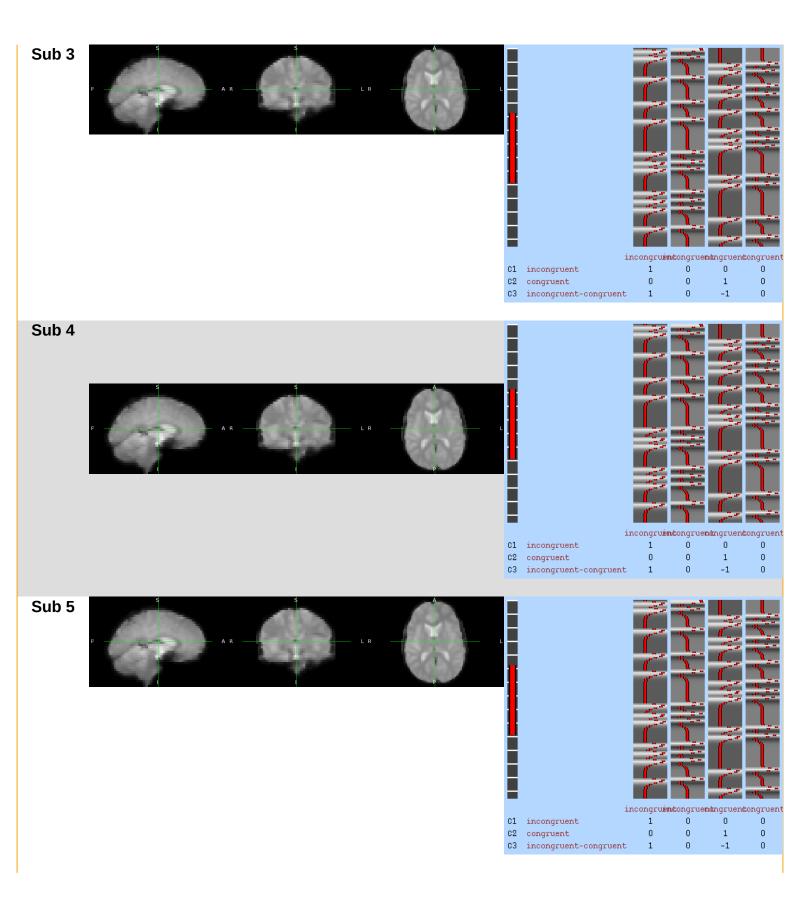
run_1stLevel_Analysis.sh:

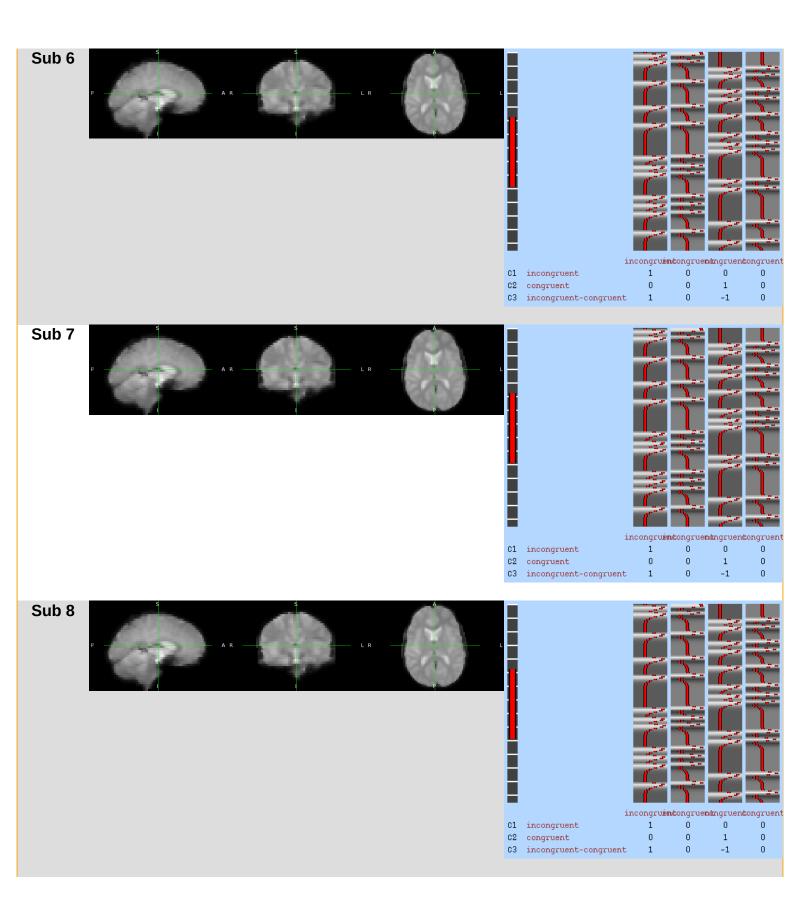
#!/bin/bash

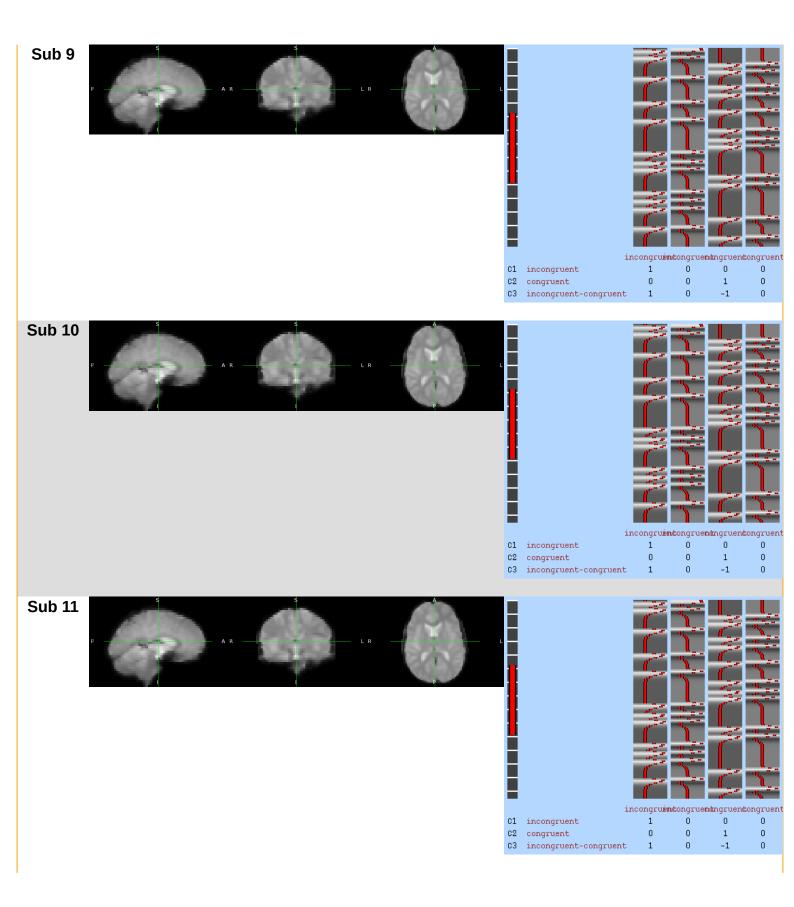
```
# Generate the subject list to make modifying this script
# to run just a subset of subjects easier.
for id in `seq -w 1 26`; do
  subj="sub-$id"
  echo "===> Starting processing of $subi"
  echo
  cd $subj
     # If the brain mask doesn't exist, create it
     if [!-f anat/${subj}_T1w_brain_f02.nii.gz]; then
       echo "Skull-stripped brain not found, using bet with a fractional intensity threshold
of 0.2"
       # Note: This fractional intensity appears to work well for most of the subjects in
the
       # Flanker dataset. You may want to change it if you modify this script for your own
study.
       bet2 anat/${subi} T1w.nii.gz \
          anat/${subj}_T1w_brain_f02.nii.gz -f 0.2
    fi
     # Copy the design files into the subject directory, and then
    # change "sub-08" to the current subject number
    cp ../design_run1.fsf .
    cp ../design_run2.fsf .
    # Note that we are using the | character to delimit the patterns
     # instead of the usual / character because there are / characters
    # in the pattern.
    sed -i " "s|sub-08|${subj}|g" \
       design_run1.fsf
    sed -i " "s|sub-08|${subj}|g" \
       design_run2.fsf
     # Now everything is set up to run feat
     echo "===> Starting feat for run 1"
     feat design run1.fsf
    echo "===> Starting feat for run 2"
    feat design_run2.fsf
          echo
  # Go back to the directory containing all of the subjects, and repeat the loop
  cd ..
done
echo
```

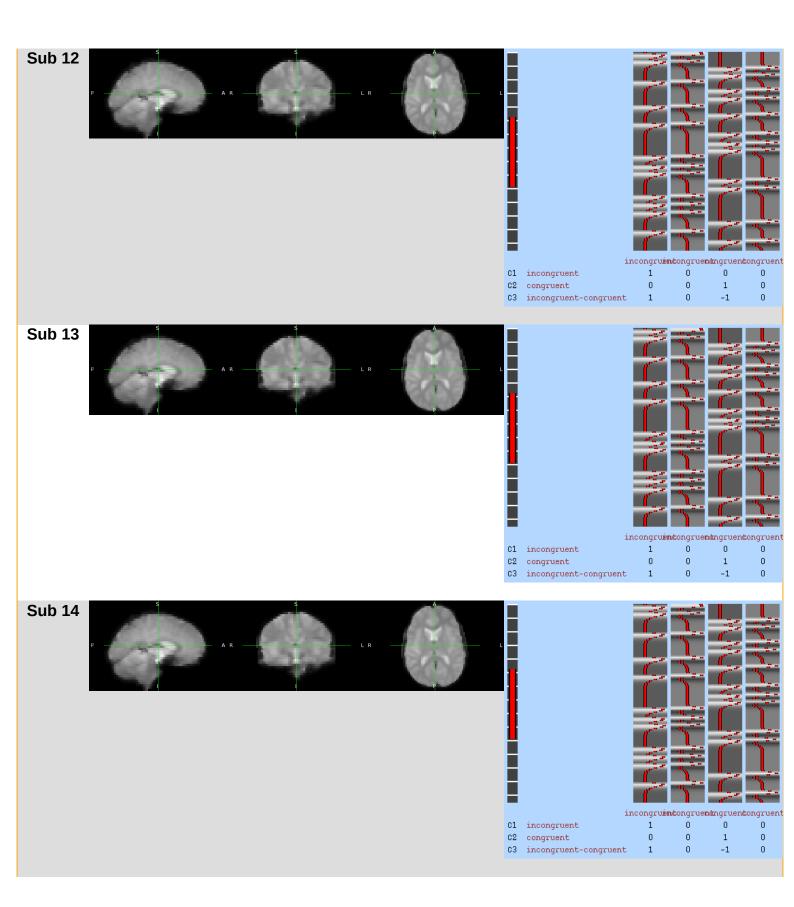


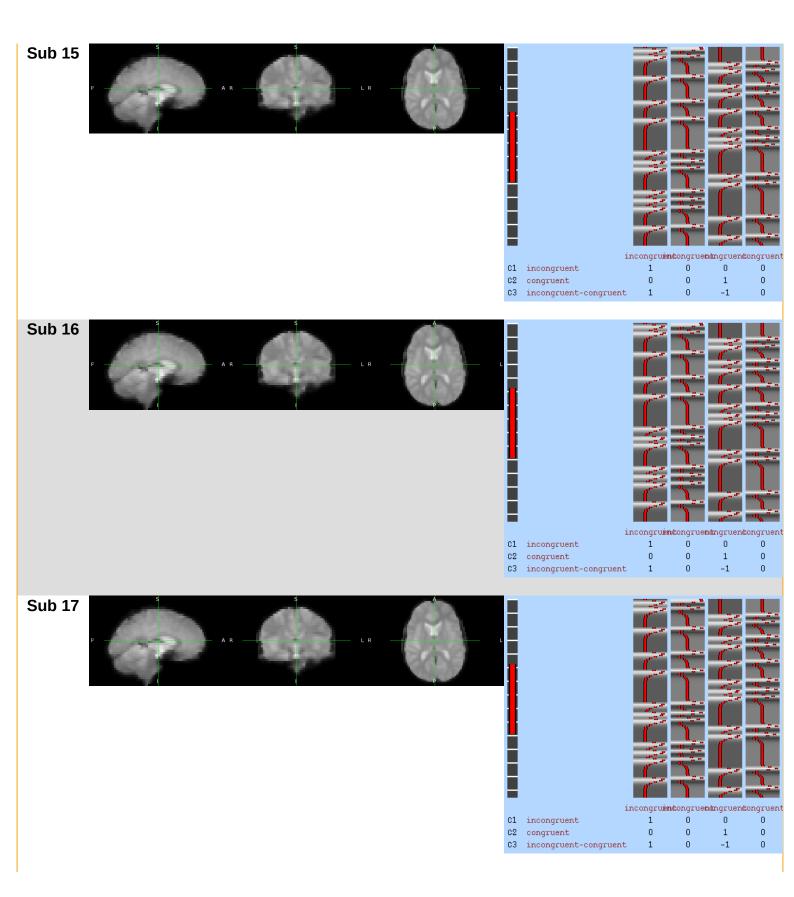


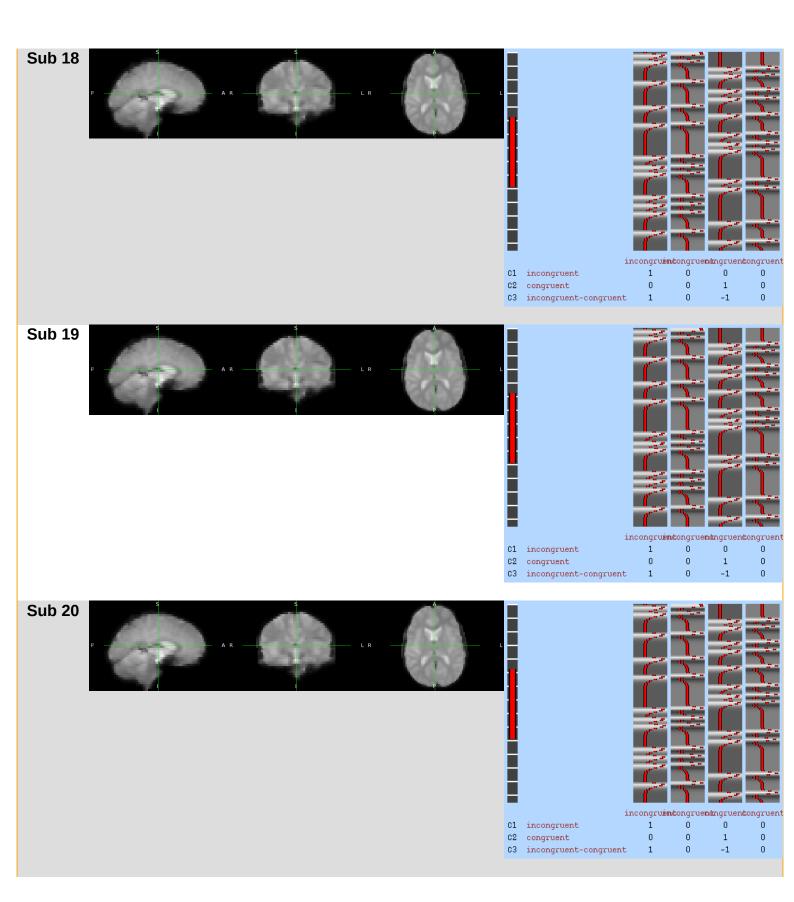


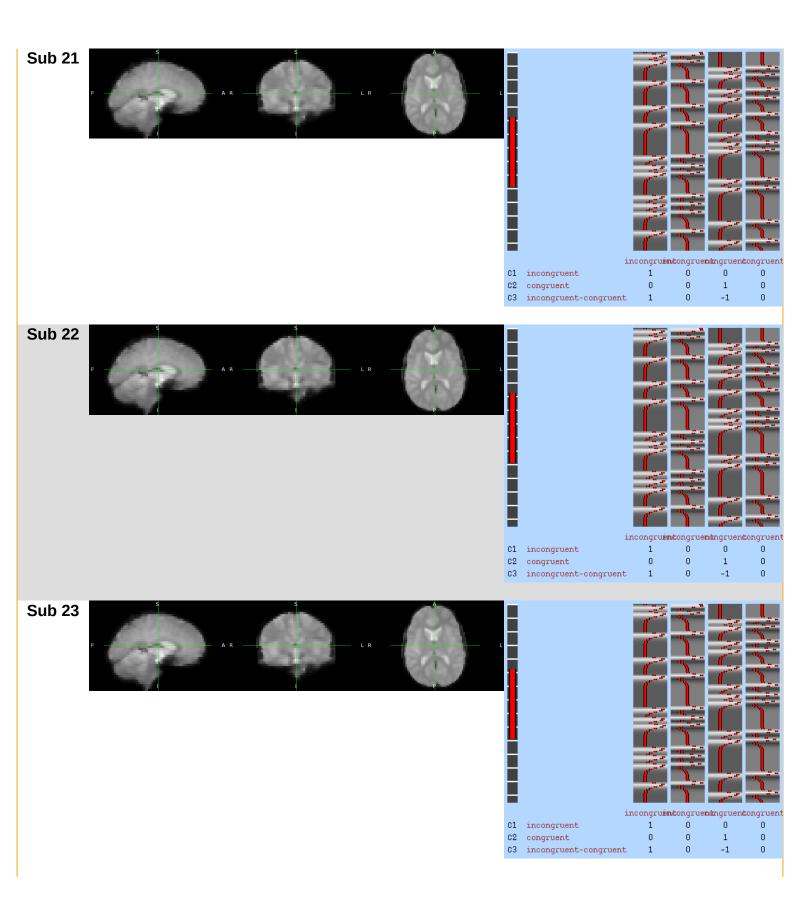


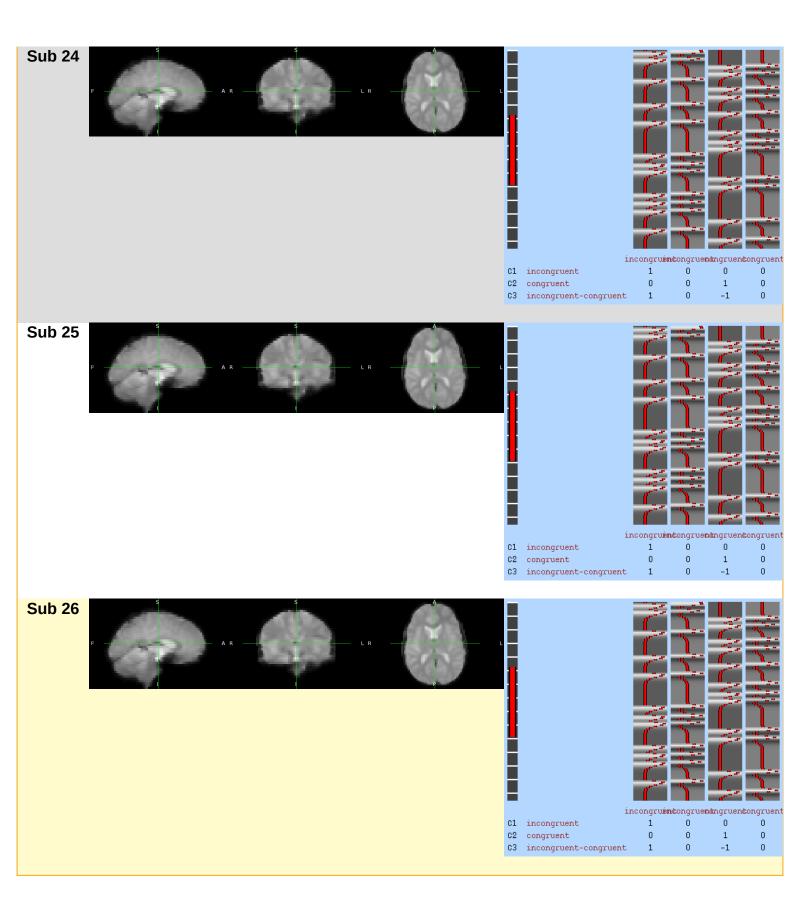












<mark>Run 2</mark>

