

fMRI Preprocessing  
“RCW”  
**spm8Batch**  
methodology

## What you need:

- spm8Batch system
- spm8 with at least revision 4290
- FSL 4.1.7 or FSL 4.1.8

# **spm8Batch processing tools:**

Standard Directory Structure

bash commands from unix shell

launches to background

email/txt msg notification

# Expected directory structure

```
/diskdrive/  
  [experiment]/  
    Subjects/  
      [subject]/  
        anatomy/  
          htloverlay.nii  
          htspgr.nii  
        func/  
          run_XX/  
            run_XX.nii  
        connect/func/  
          func/  
            run_XX/  
              run_XX.nii
```

## Auxiliary Commands

- getfMRI
- UNNIFTI
- tarBET

## Main Processing Commands

- [next page]

Login

Launch command

(automatically builds components goes into background)

Logout (if you wish)

Check email, check the log for any errors.

```
/diskdrive/  
  [experiment]/  
    matlabScripts/  
      spm8batch/  
        [command]/  
          YYYY_MM/  
  
warpfMRI_XXXXX_...sh  
warpfMRI_XXXXX_...log  
warpfMRI_XXXXX_...m
```

processing order



*[physioCorr]*

*sliceTime*

*realignfMRI*

bestBET

fslCheck

mvBestBET

tarBET

coregOverlay

coregHiRes

warpHiRes - *DARTEL option*

warpfMRI

smoothfMRI

**all commands have built-in help**

# Command Options

-A	all runs present
-a [directory]	anatomy directory e.g. anatomy/BET
-b	also put best BET one picked
-D	super debug flag
-d	debug flag
-F [TR]	fMRI TR
-f [directory]	functional directory e.g. connect/func
-g [gradient]	gradient value to pass to 'bet'
-h [name]	high resolution image
-i [run number]	include this run number
-M [directory]	master subject directory
-m ["options"]	mcflirt options
-n [name]	name prepend
-O [name]	other object to drag into process
-o [name]	overlay image name
-R	reslice flag set to 2
-r	reslice flag set to 1
-S [#]	standard volume number for mcflirt
-s [directory]	sub-directory name to search for images
-T [name]	template image name
-t	test flag
-U [unique]	user email name/txt msg address
-u [#]	BET threshold override value
-v [name]	volume to use for coregistration
-w [directory]	coregistration output directory
-z [#]	voxel reslice size
-# [#-#]	inclusive run list

Expected data:

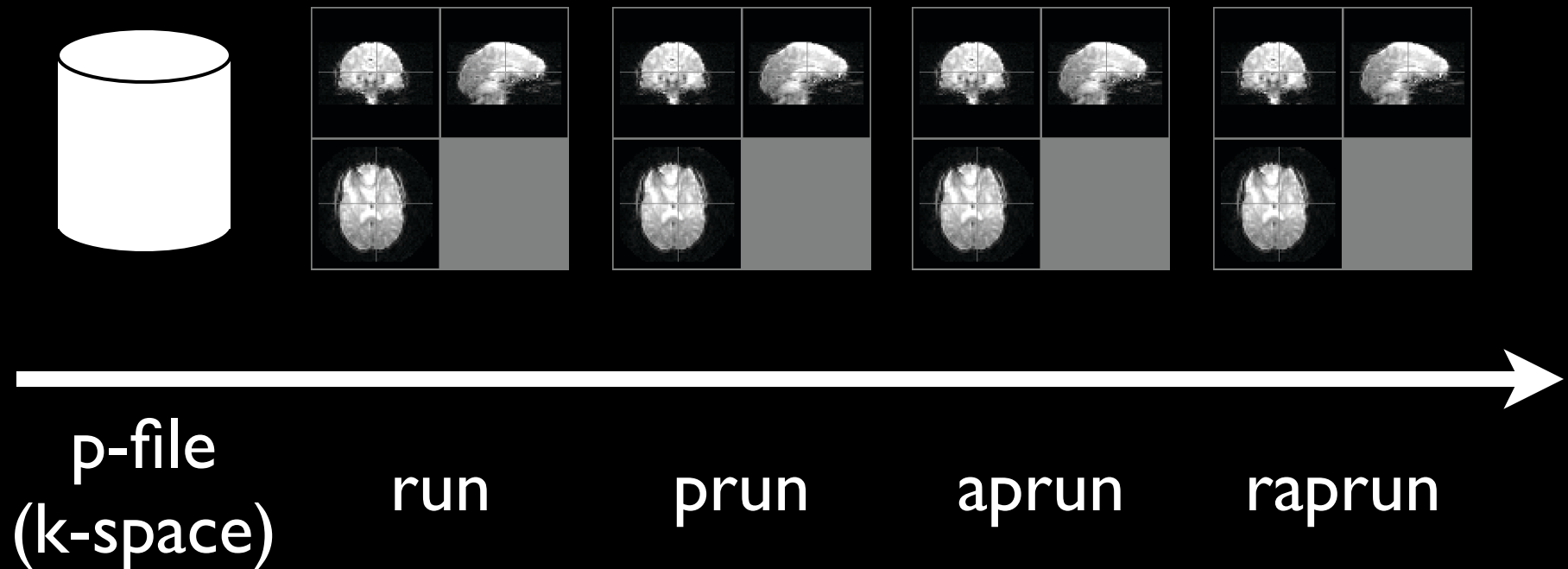
overlay image in the same rough space as  
the functional data

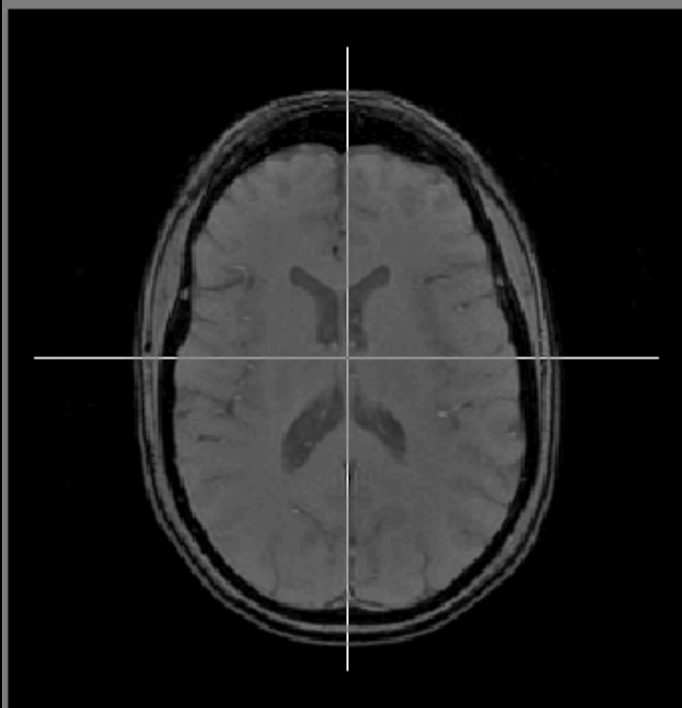
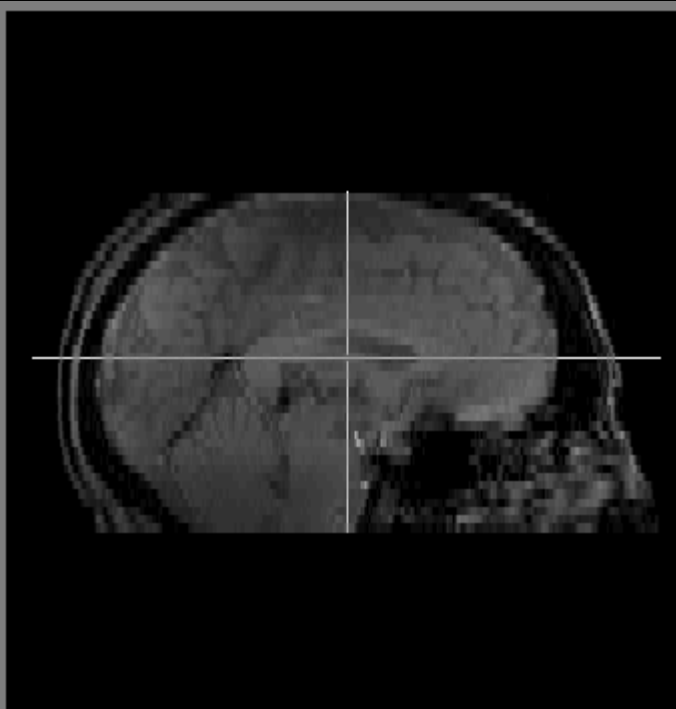
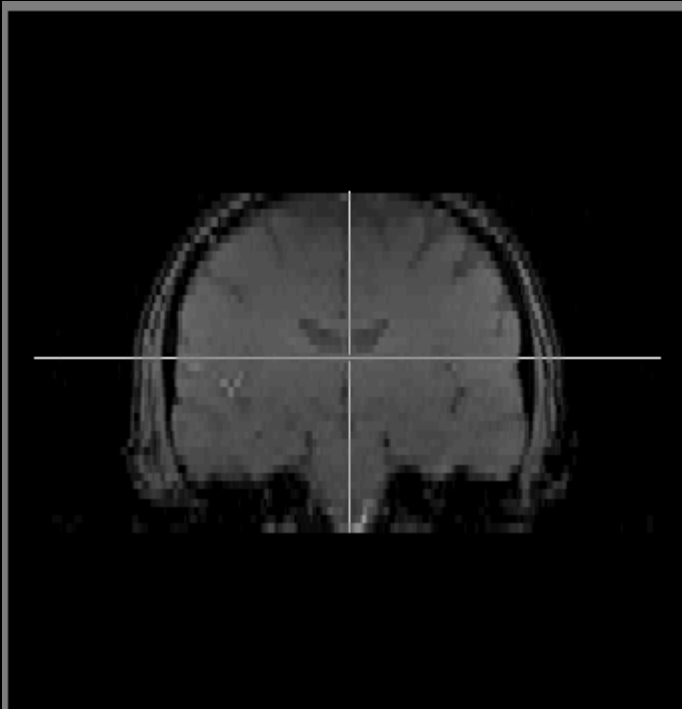
hi resolution image needed for warping

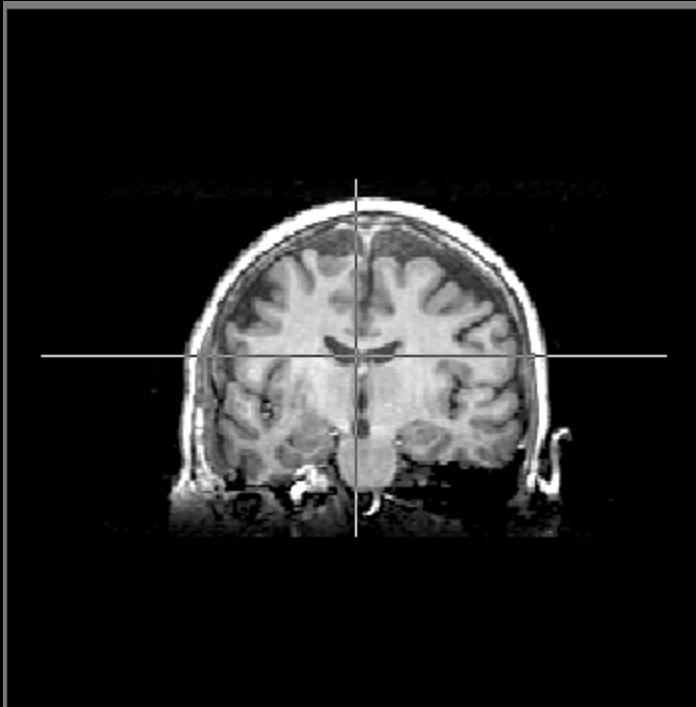
time-series data



# Pre-processing done at fMRI Lab

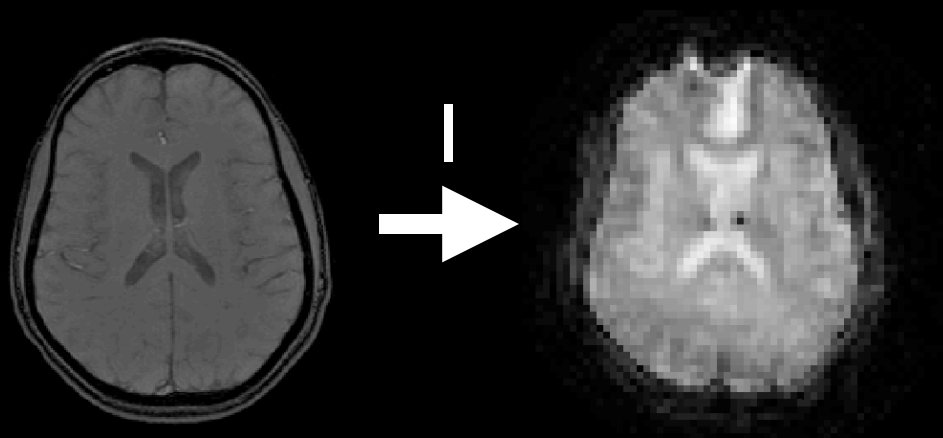






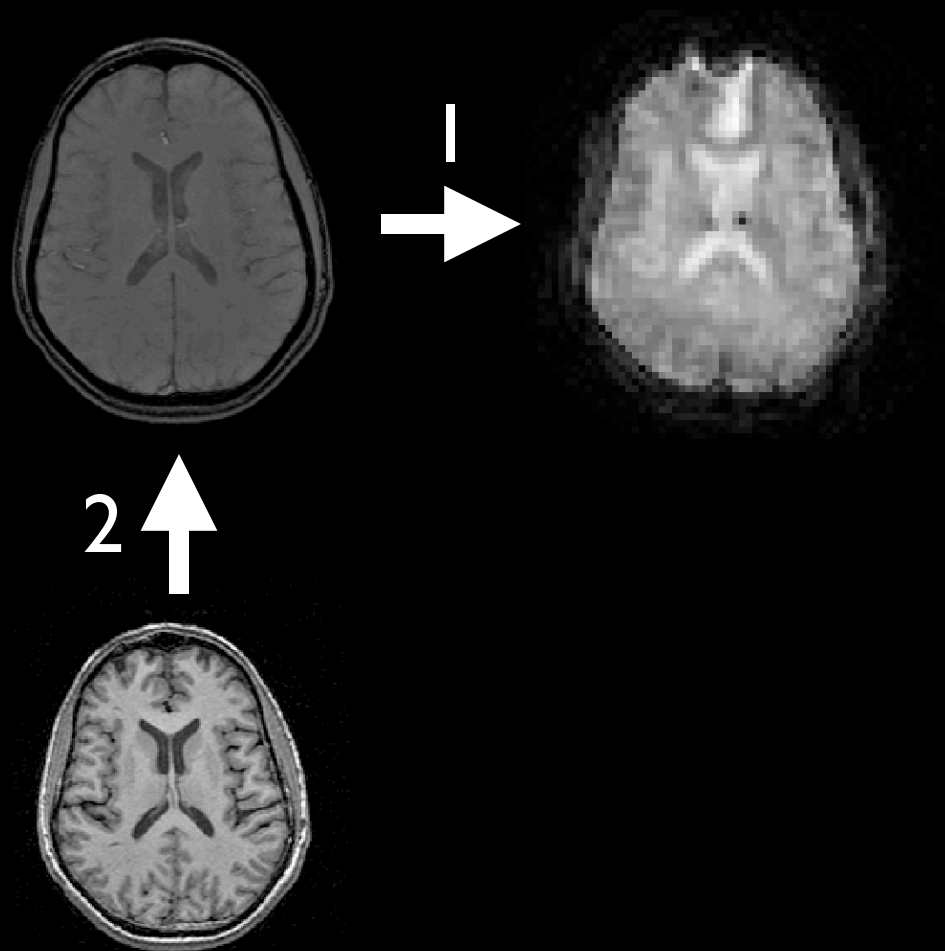
Hi Resolution

# Strategy for normalization of functionals



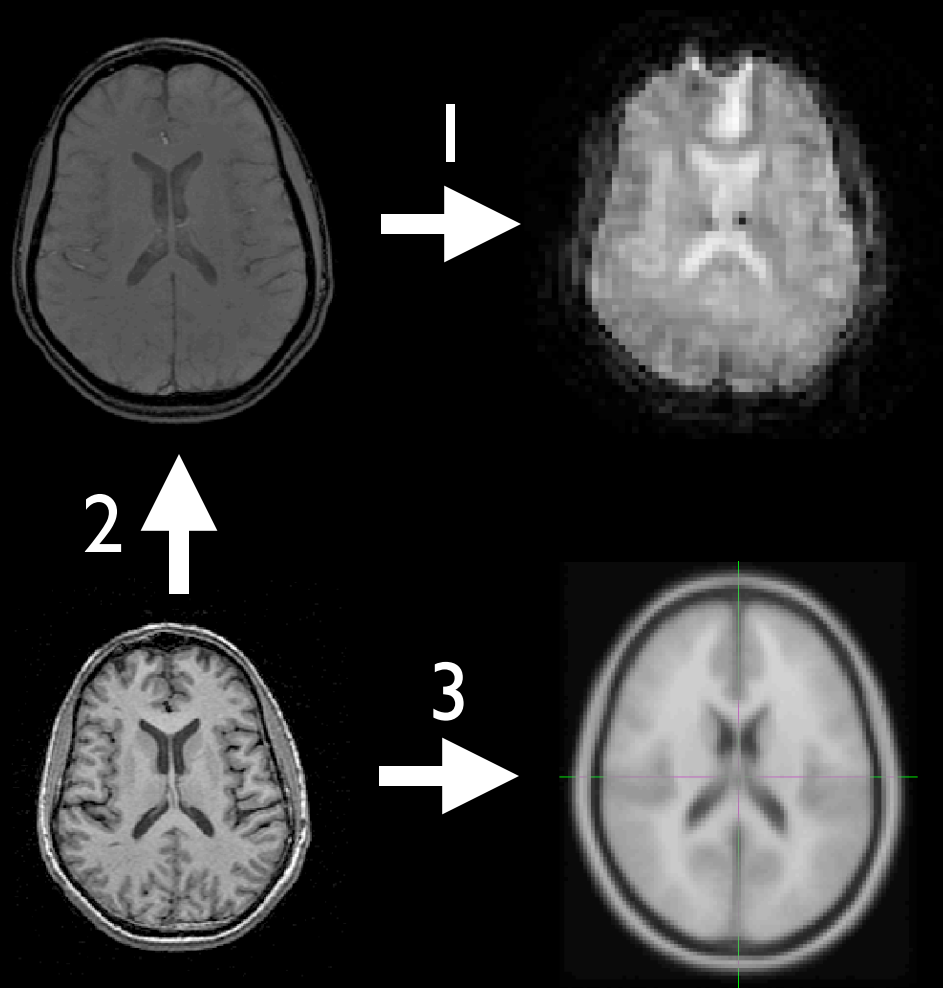
- 1 - coregOverlay
- 2 - coregHiRes
- 3 - warpHiRes
- 4 - warpfMRI

# Strategy for normalization of functionals



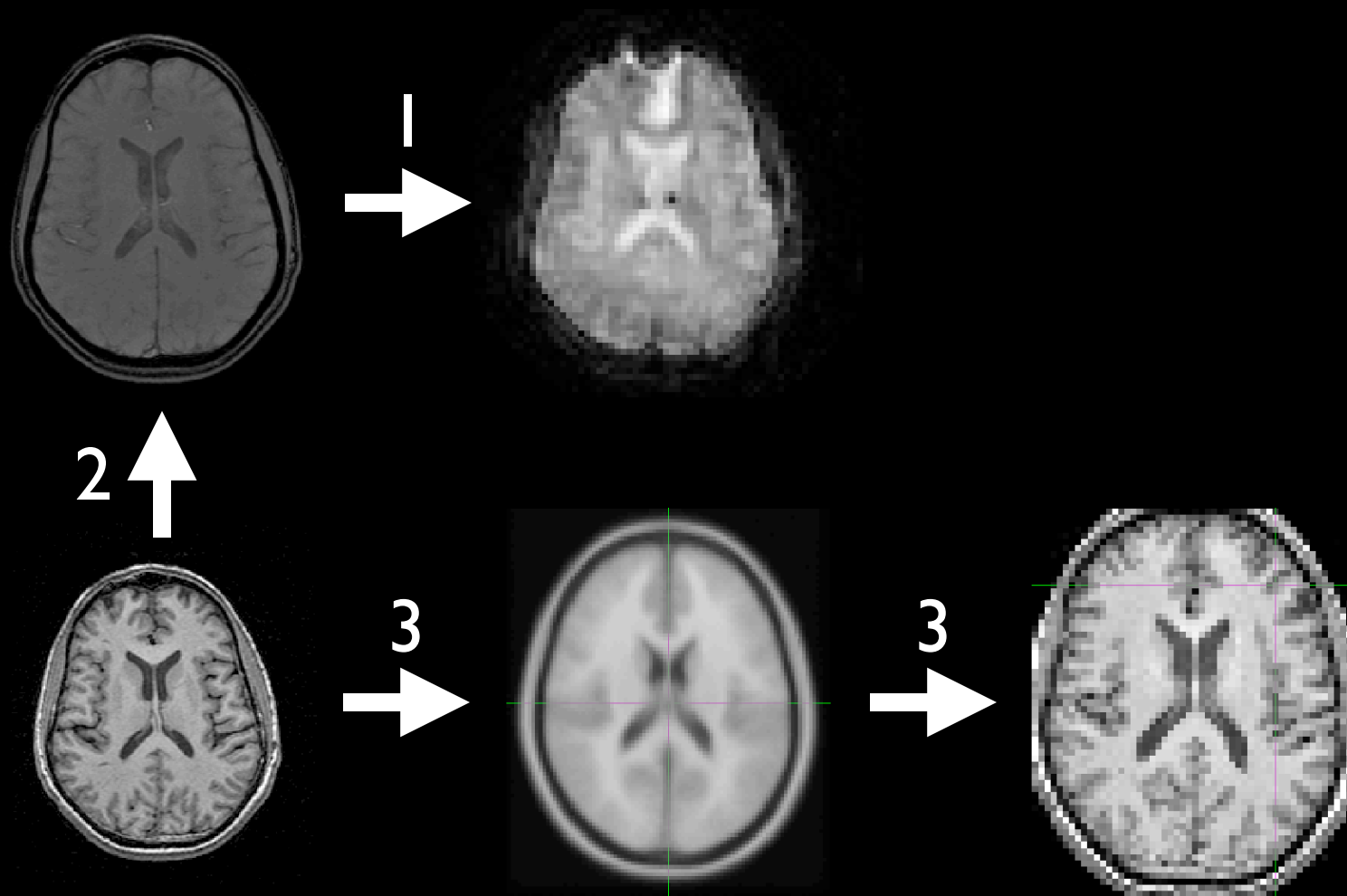
- 1 - coregOverlay
- 2 - coregHiRes
- 3 - warpHiRes
- 4 - warpfMRI

# Strategy for normalization of functionals



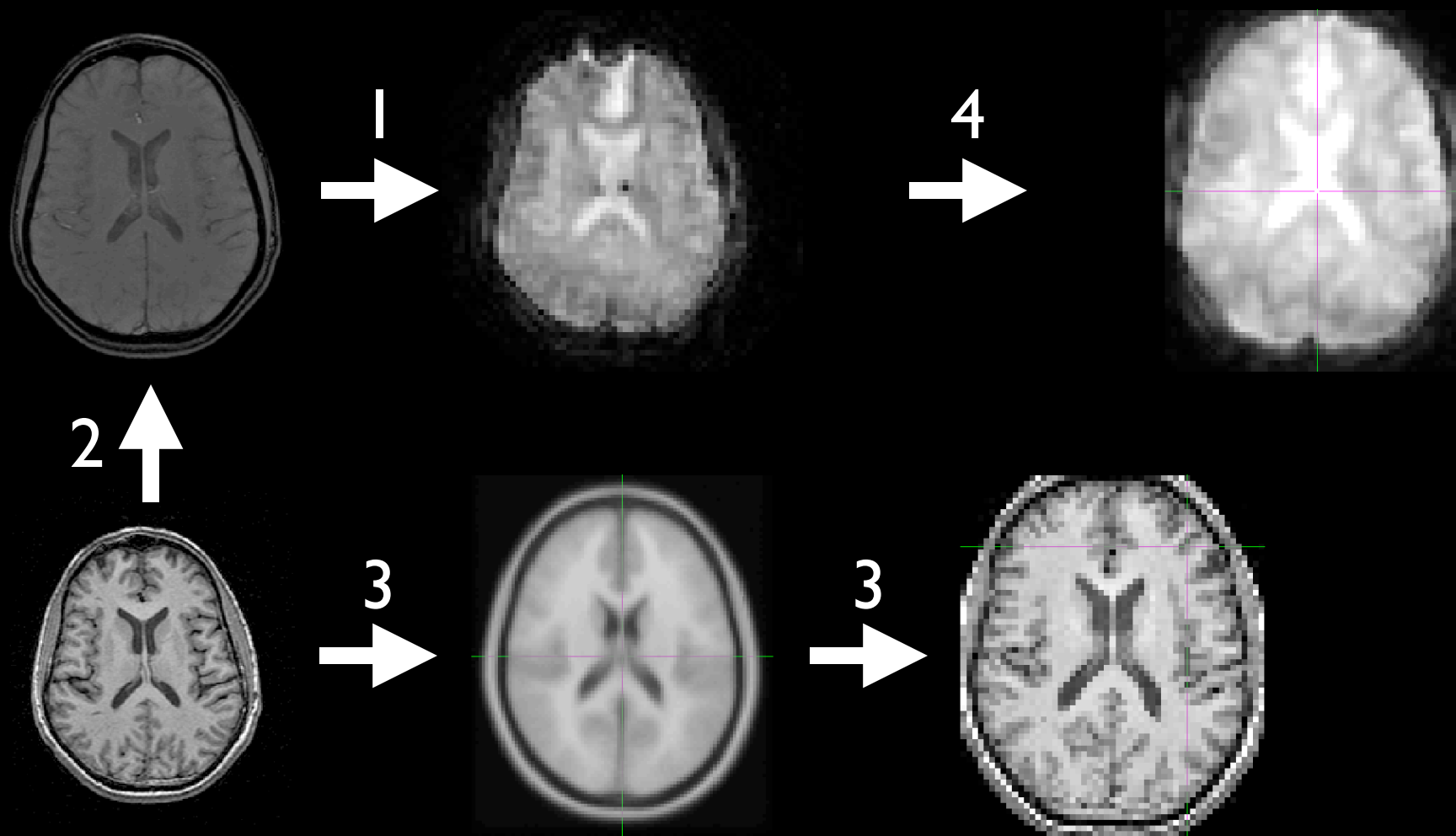
- 1 - coregOverlay
- 2 - coregHiRes
- 3 - warpHiRes
- 4 - warpfMRI

# Strategy for normalization of functionals



- 1 - coregOverlay
- 2 - coregHiRes
- 3 - warpHiRes
- 4 - warpfMRI

# Strategy for normalization of functionals



- 1 - coregOverlay
- 2 - coregHiRes
- 3 - warpHiRes
- 4 - warpfMRI



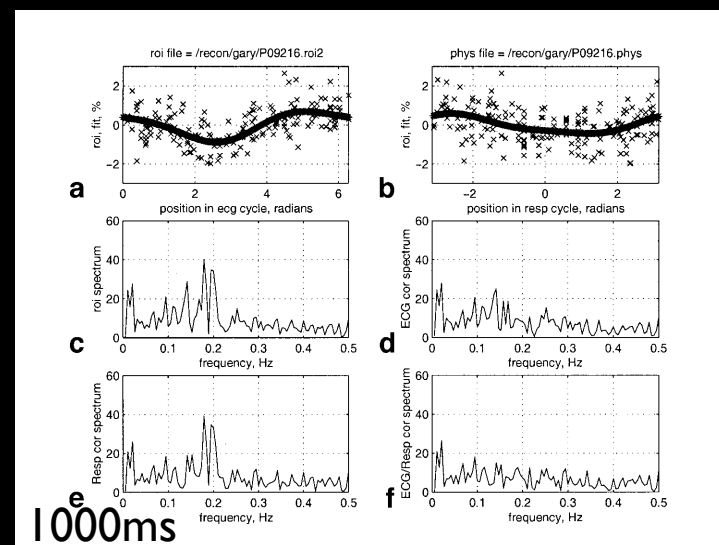
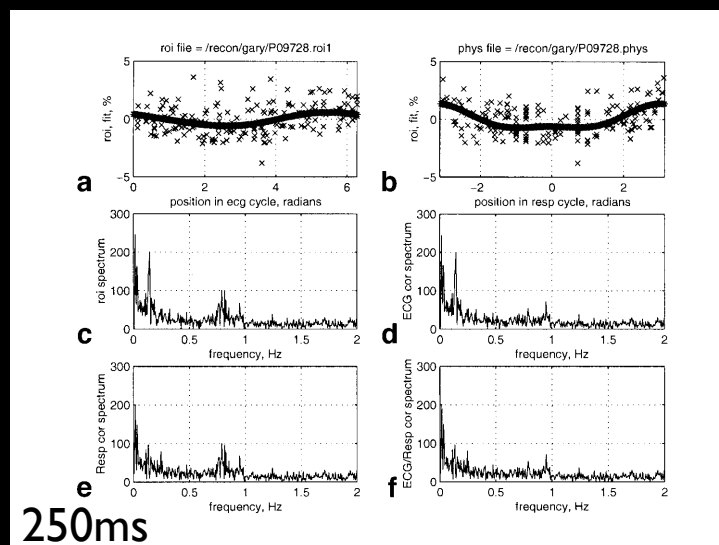
# *physioCorr [subject]*

## physiological correction:

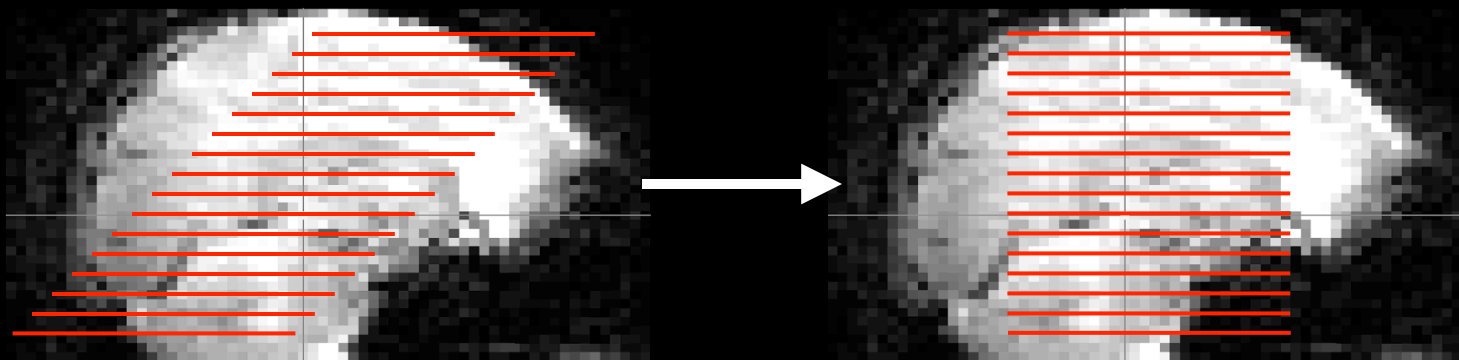
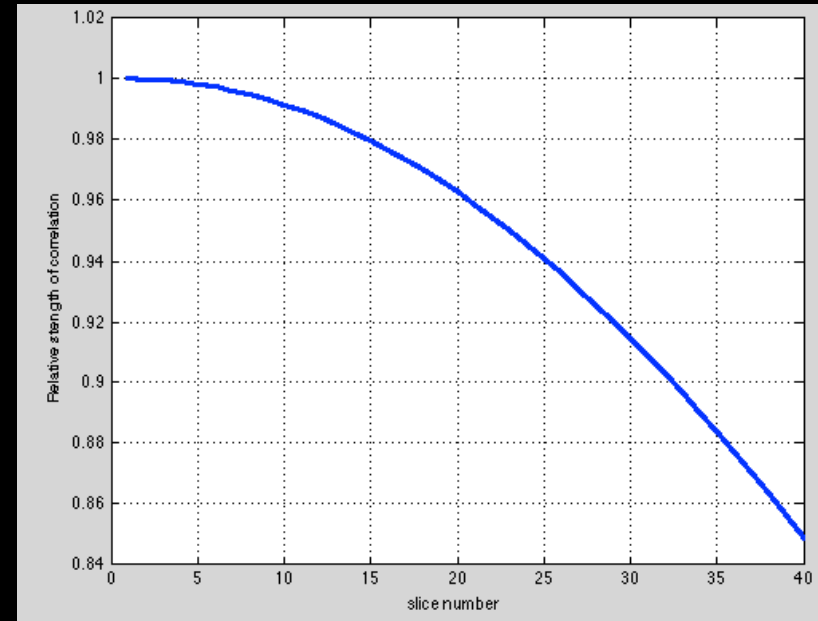
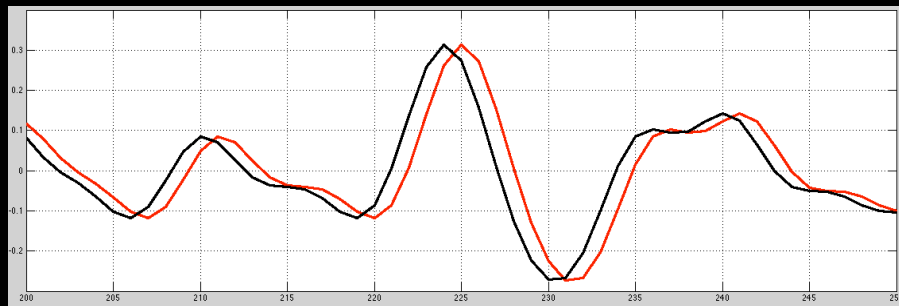
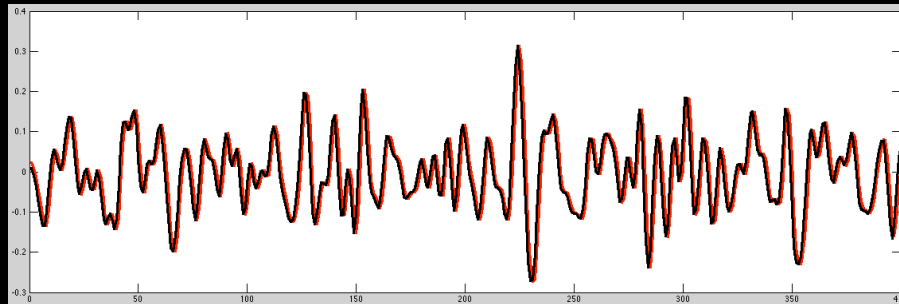
In the current paper, we have shown that the NVR model composed of a comprehensive set of nuisance regressors substantially reduces the structured noise in fMRI residuals. The NVR model is based on a number of effects which are known to contribute to the non-white noise in fMRI (hardware drift, residual movement artefacts, respiration and cardiac pulsation). In fact, the proposed NVR model is only new in the sense that we for the first time have used a combination of several already published models in the same analysis.

It was furthermore found that our approach, in general, was superior to the covariance estimation currently implemented in SPM2. In particular, we found the global AR(1) model of SPM to be inadequate near larger arteries which is not surprising given the inability of a first-order AR model to account for oscillatory noise. -- Lund, Neuroimage 29, 2006

## RETROICOR - implemented at fMRI BIRB

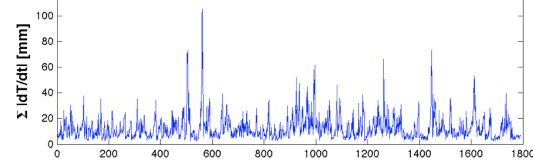


# *sliceTime [subject]*

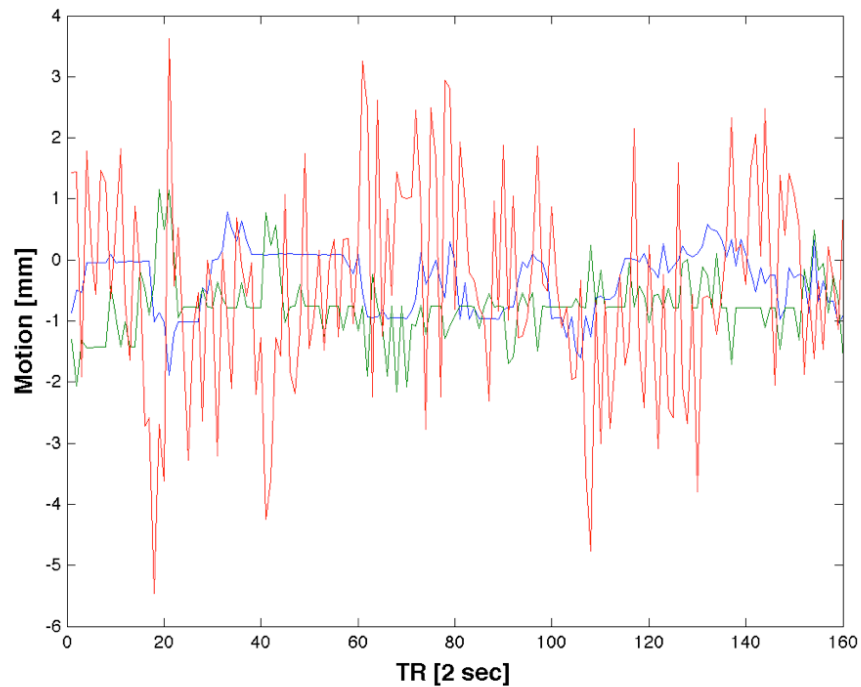
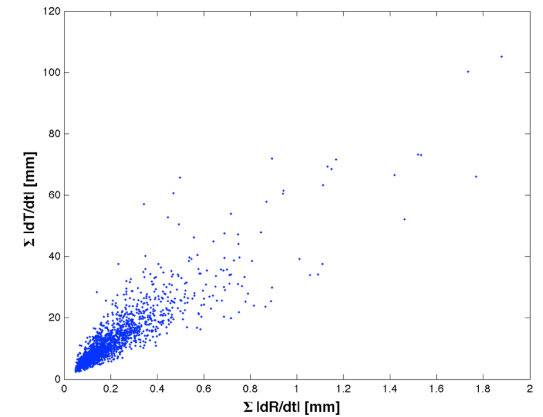
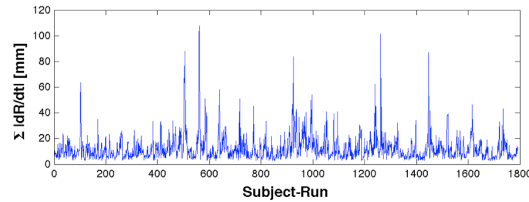
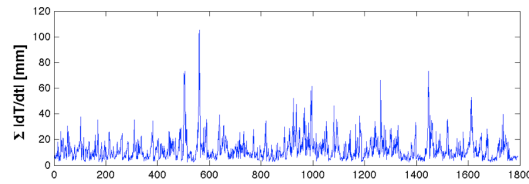
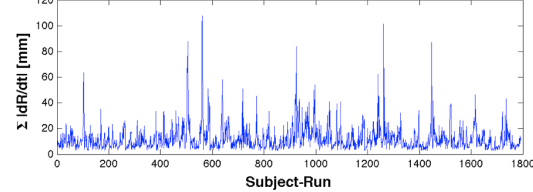


# *realignfMRI [subject]*

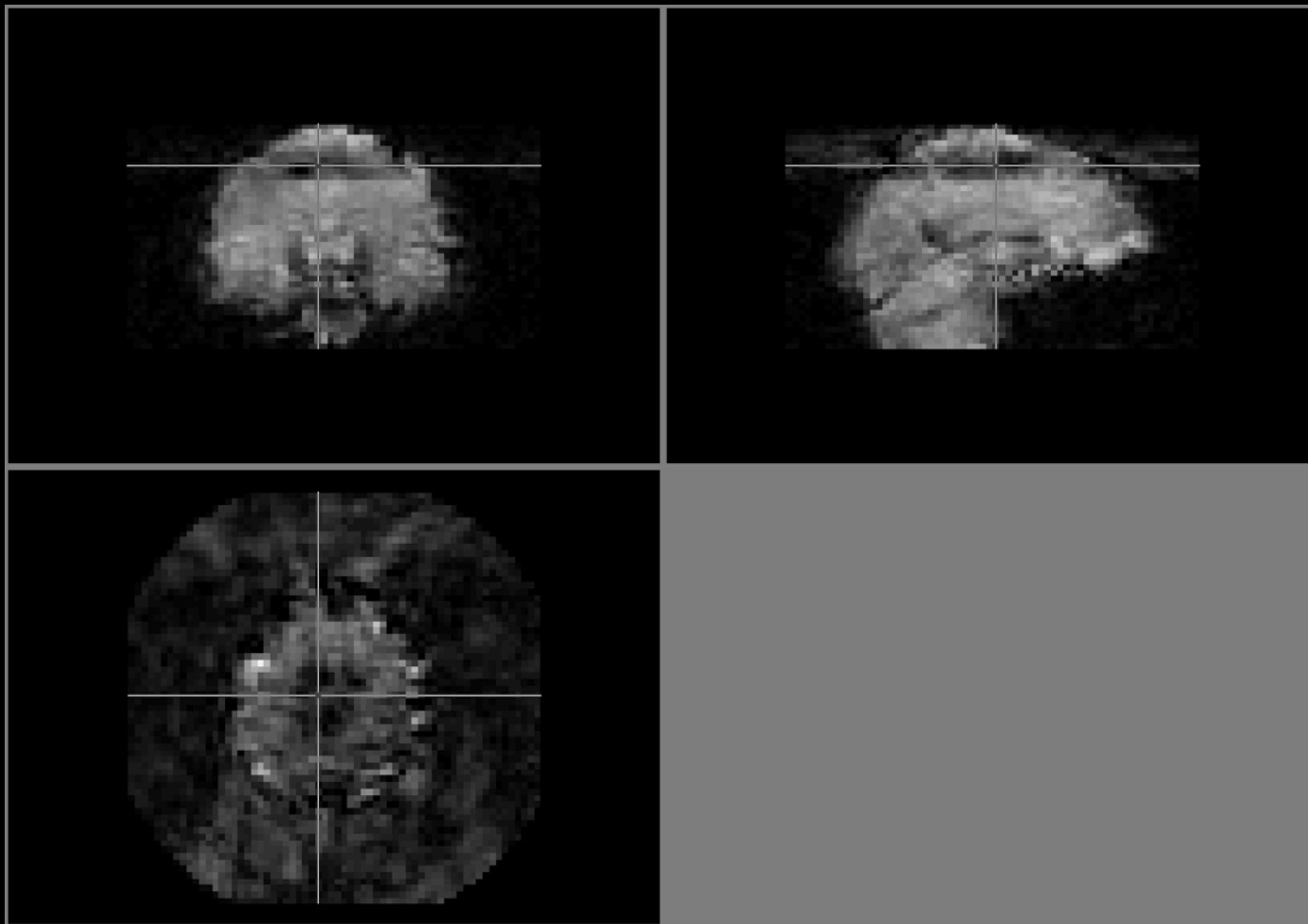
$\langle \text{rotParm} \rangle = 12.17^\circ \pm 10.00^\circ$



$\langle \text{trnParm} \rangle = 12.61 \text{ mm} \pm 9.63 \text{ mm}$



Always check for bad data. Arduous task  
but critical.



*physioCorr [subject]*

***run\_XX.nii → prun\_XX.nii***

*sliceTime [subject]*

***prun\_XX.nii → aprun\_XX.nii***

*realignfMRI [subject]*

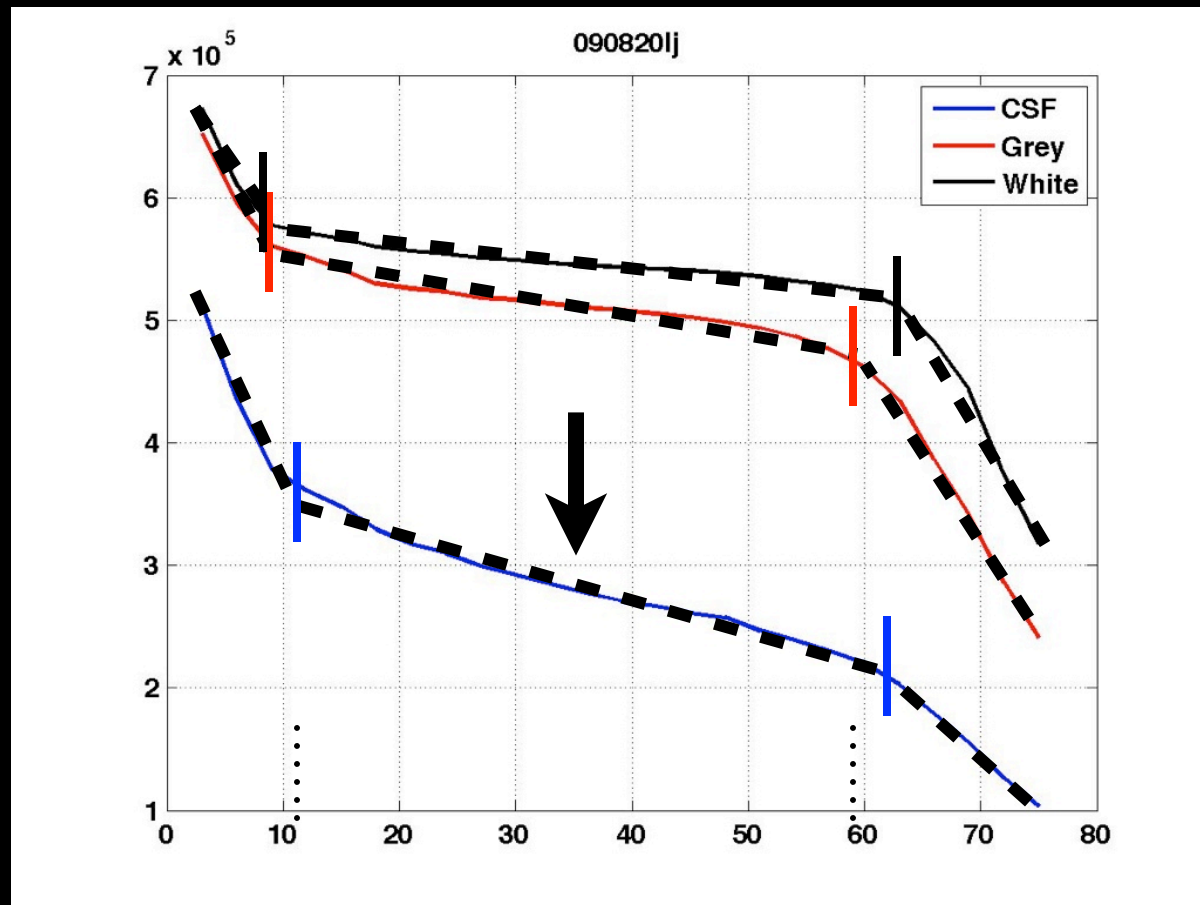
***aprun\_XX.nii → raprun\_XX.nii***

# Getting the best brain extraction

`bestBET, fs1Check, mvBestBET`

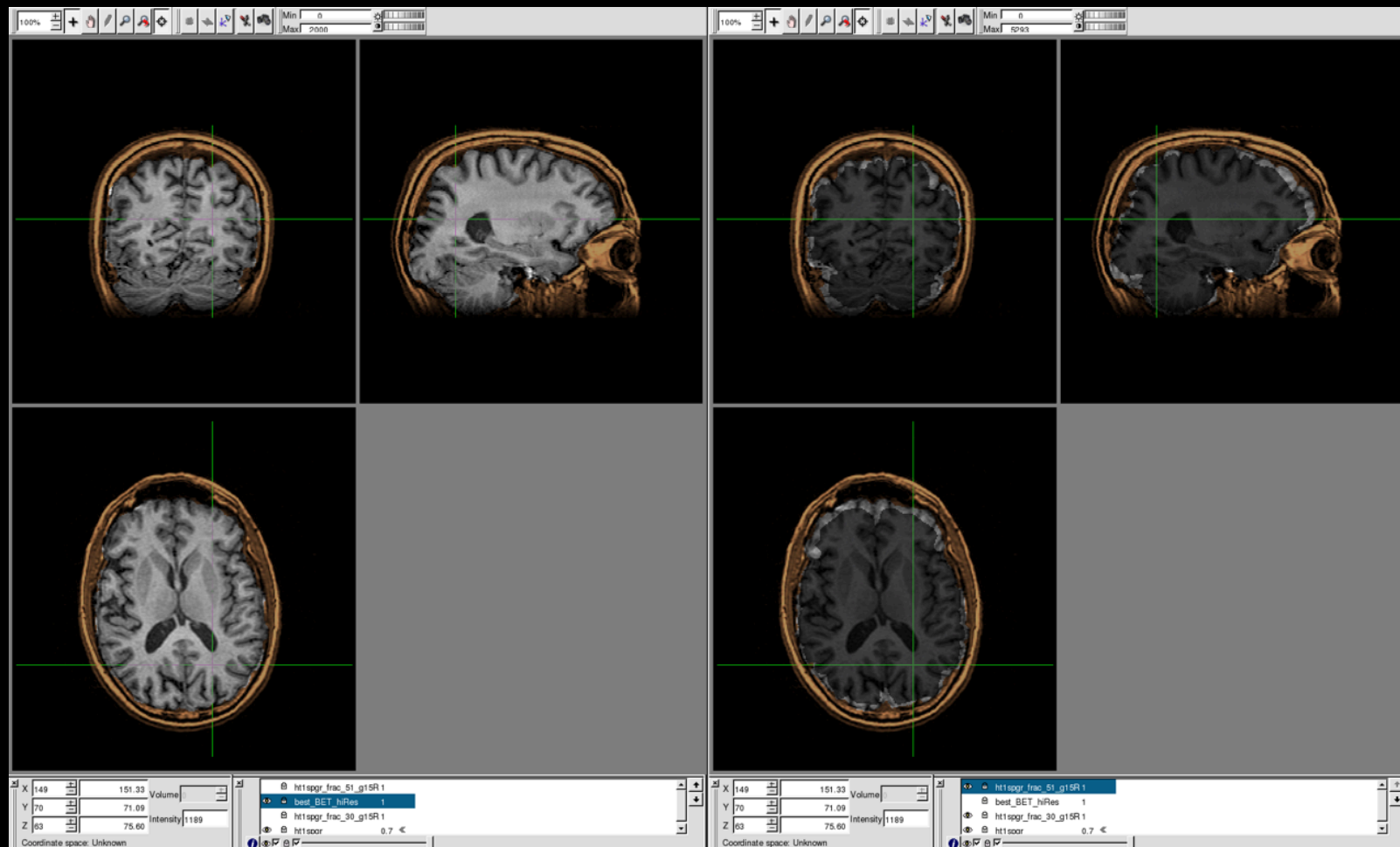
bestBET [subject]

voxel count



extraction fraction

```
fs1Check -h ht1spgr [subject][-b]
```

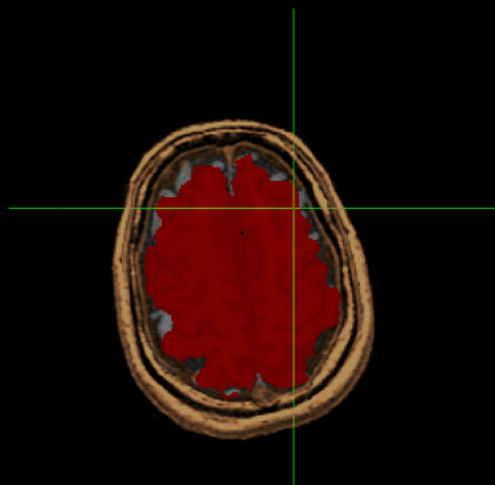
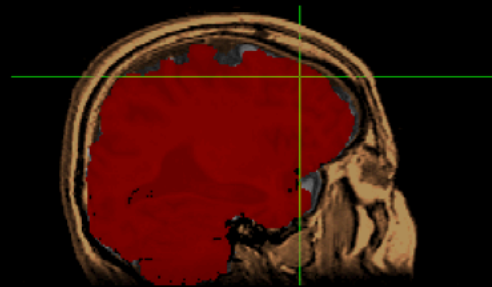
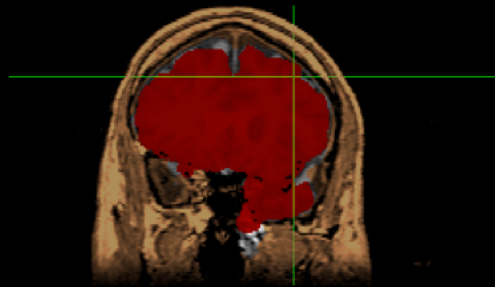


```
mvBestBET -h ht1spgr [subject] [-u value]
```

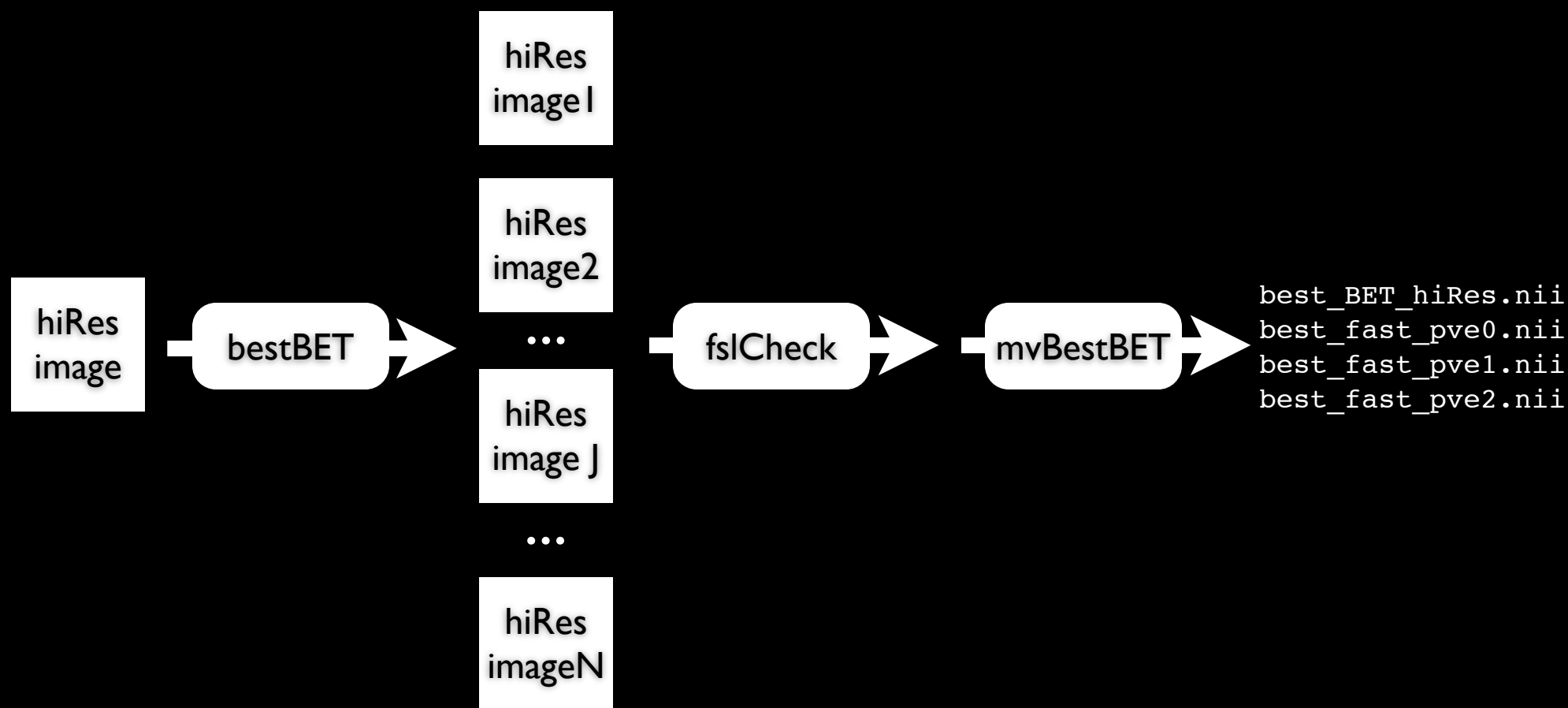


# Example of different thresholds.

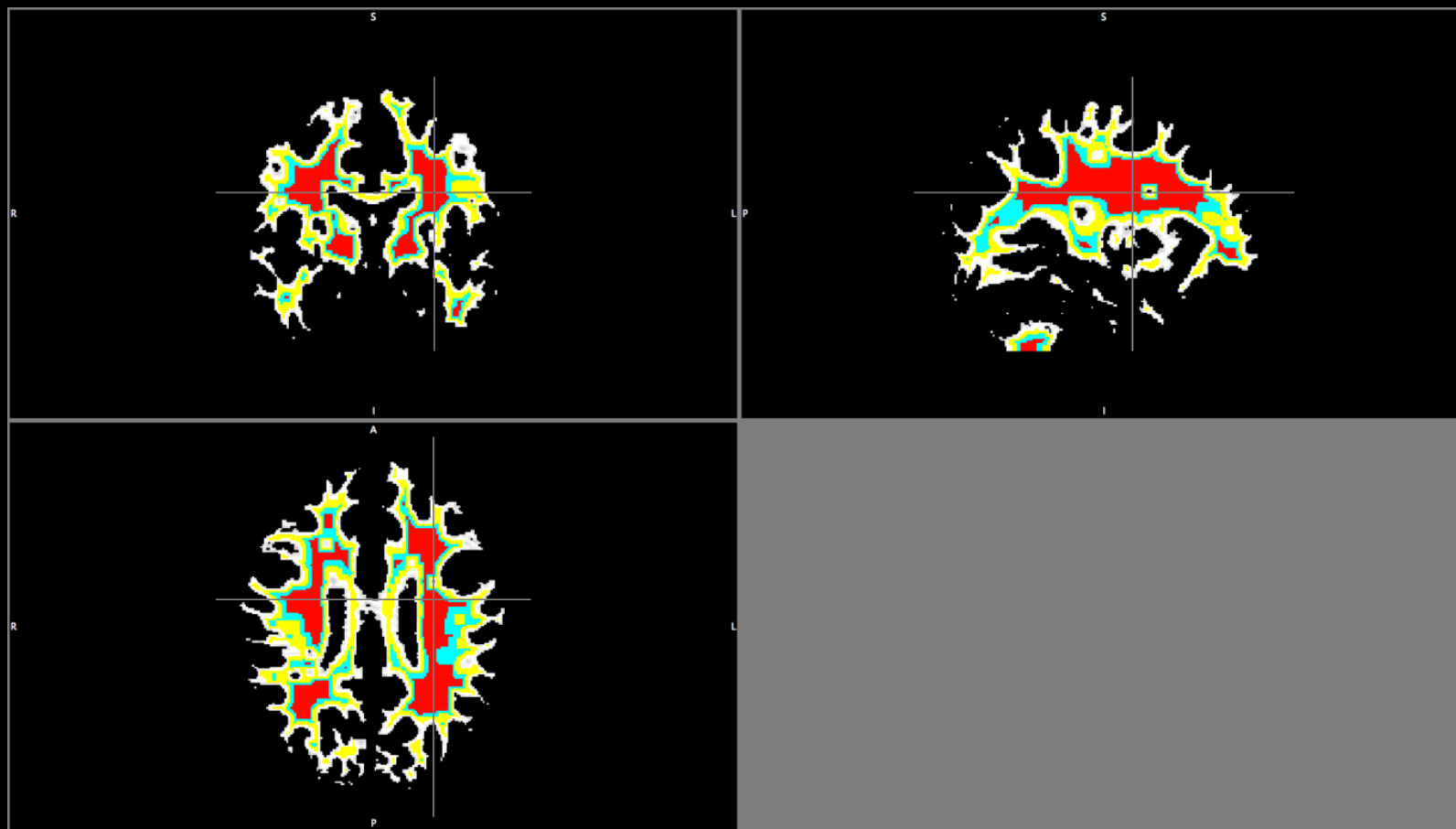
Red is not great.



As part of bestBET, different tissue segments are created. The mvBestBET command will automatically label the ones you choose for further processing.

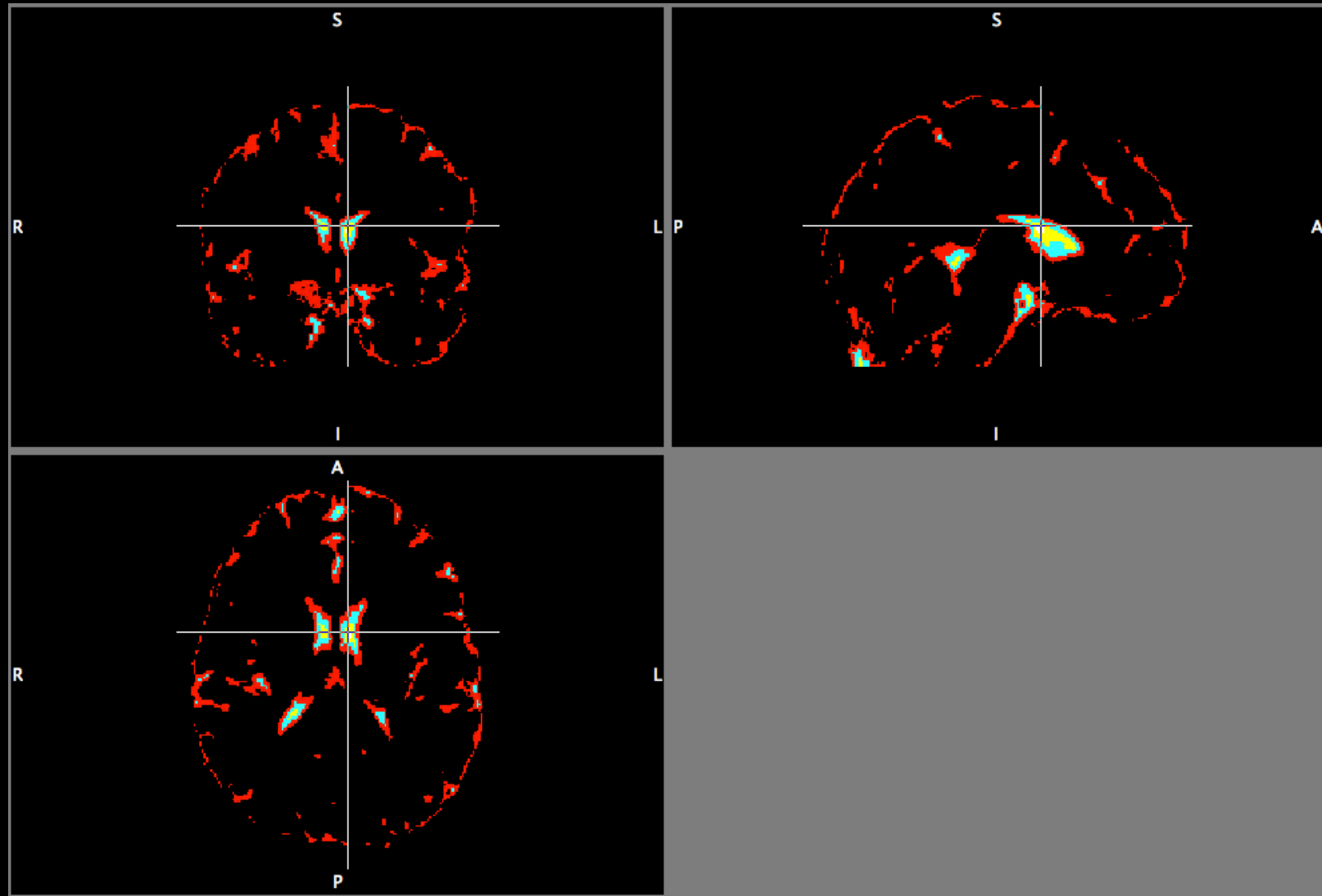


# White matter segment with various erosions.



anatomy/BET/best\_BET\_hiRes.nii  
anatomy/BET/best\_fast\_pve0.nii  
anatomy/BET/best\_fast\_pve1.nii  
anatomy/BET/best\_fast\_pve2.nii

# CSF segment with various erosions.



# coregOverlay [subject]

Prior

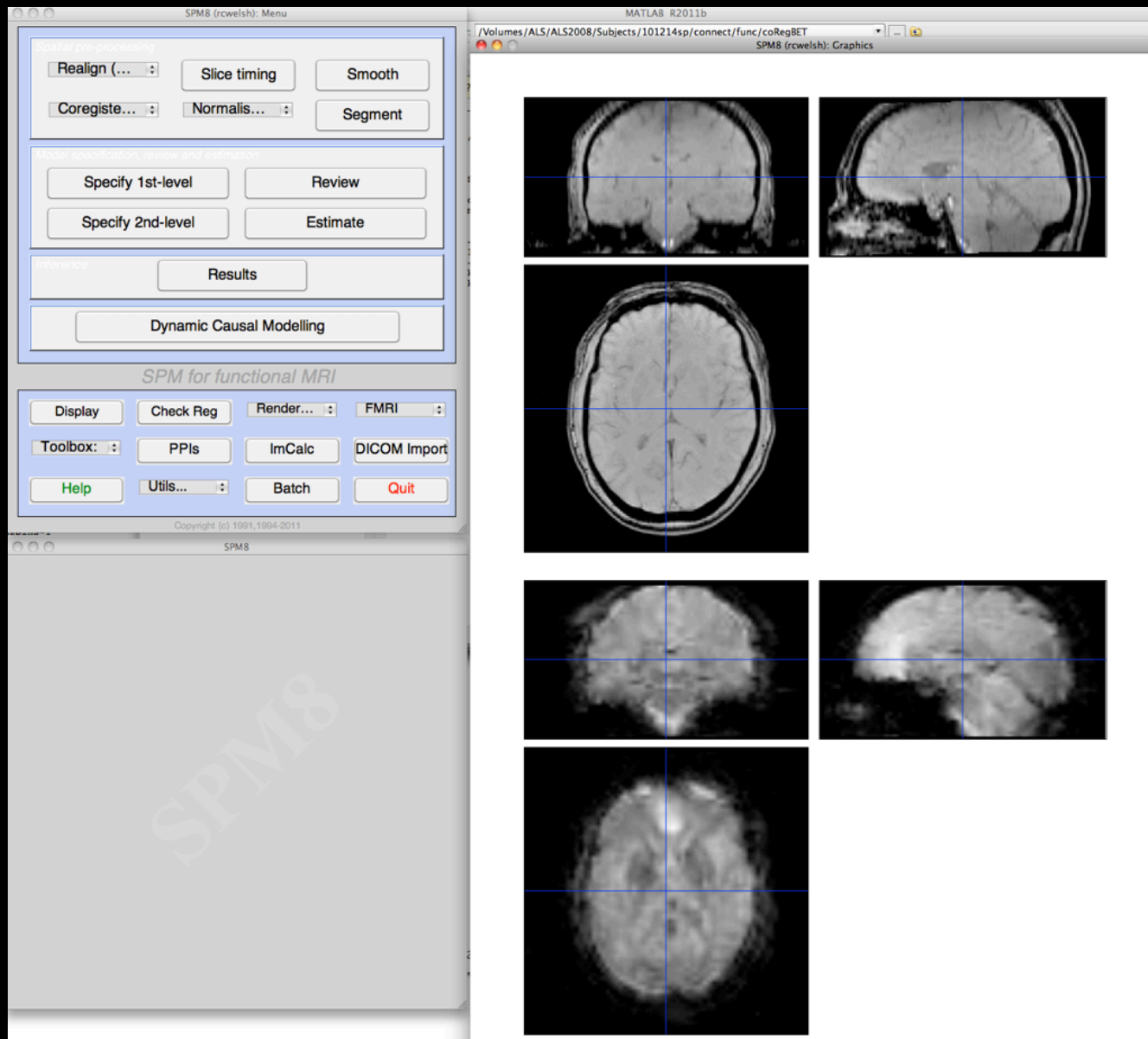


After

```
[subject]/  
  anatomy/  
    htlooverlay.nii  
    htspgr.nii  
func/  
  run_XX/  
    run_XX.nii  
    raprun_XX.nii
```

```
[subject]/  
  anatomy/  
    htlooverlay.nii  
    htspgr.nii  
func/  
  run_XX/  
    run_XX.nii  
    raprun_XX.nii  
coReg/  
  htlooverlay.nii
```

# Check registration



# coregHiRes [subject]

Prior



After

```
[subject]/  
  anatomy/  
    htlooverlay.nii  
    htspgr.nii  
    BET/  
      best_BET_hiRes.nii  
func/  
  run_XX/  
    run_XX.nii  
    raprun_XX.nii
```

```
[subject]/  
  anatomy/  
    htlooverlay.nii  
    htspgr.nii  
func/  
  run_XX/  
    run_XX.nii  
    raprun_XX.nii  
coReg/  
  htlooverlay.nii  
  best_BET_hiRes.nii  
  best_fast_pve0.nii  
  best_fast_pve1.nii  
  best_fast_pve2.nii
```

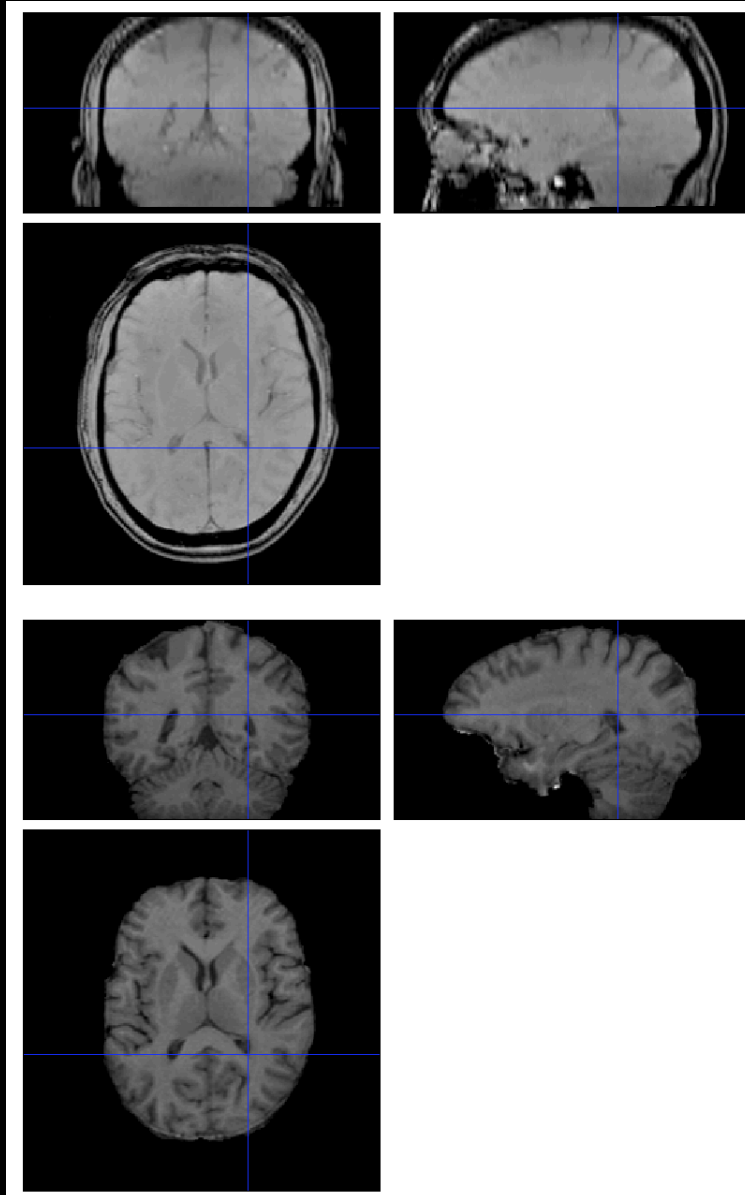
warpHiRes [subject]

After

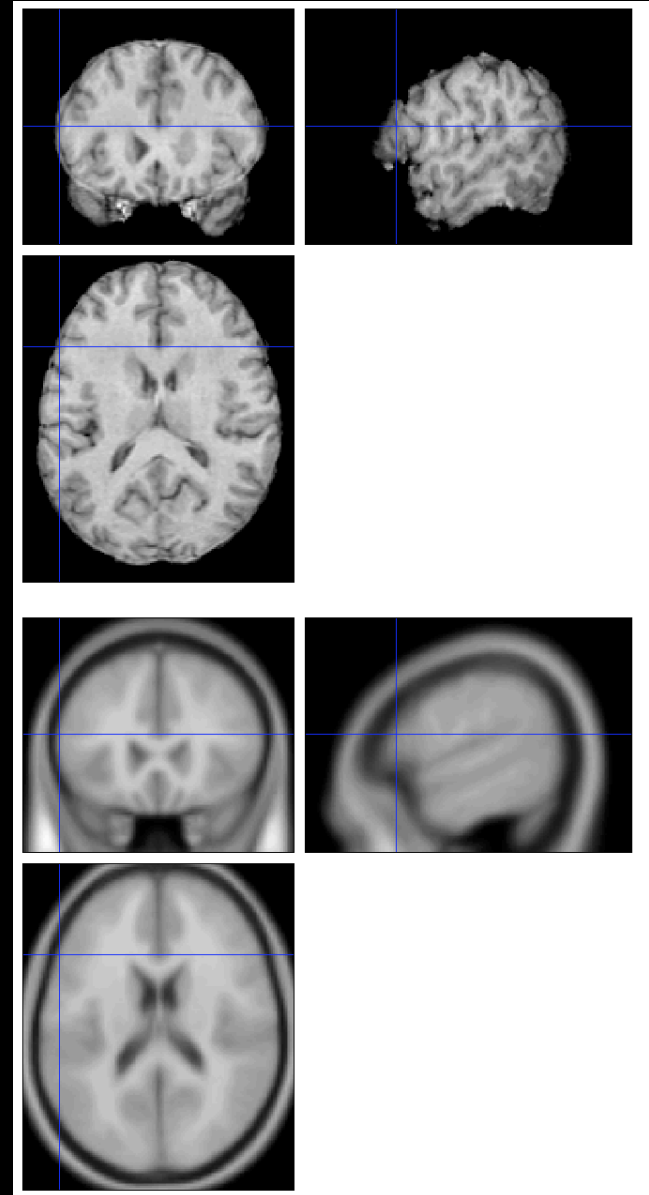
```
[subject]/  
  anatomy/  
    ...  
  func/  
    run_XX/  
      ...  
    coReg/  
      w1mm_best_BET_hiRes.nii  
      w1mm_best_fast_pve0.nii  
      w1mm_best_fast_pve1.nii  
      w1mm_best_fast_pve2.nii  
      best_BET_hiRes_sn.mat
```



## Overlay to hires



## Warped HiRes



# warpfMRI [subject]

Prior



After

```
[subject]/  
  anatomy/  
    htloverlay.nii  
    htspgr.nii  
func/  
  run_XX/  
    run_XX.nii  
    raprun_XX.nii
```

```
[subject]/  
  anatomy/  
    htloverlay.nii  
    htspgr.nii  
func/  
  run_XX/  
    run_XX.nii  
    raprun_XX.nii  
    w3mm_raprun_XX.nii
```

# smoothfMRI [subject]

Prior



After

```
[subject]/  
  anatomy/  
    htloverlay.nii  
    htspgr.nii  
  func/  
    run_XX/  
      run_XX.nii  
      raprun_XX.nii
```

```
[subject]/  
  anatomy/  
    htloverlay.nii  
    htspgr.nii  
  func/  
    run_XX/  
      run_XX.nii  
      raprun_XX.nii  
      w3mm_raprun_XX.nii  
      s5mm_w3mm_raprun_XX.nii
```

# Warped functionals    Smoothed functionals

