

# **Complex Networks fMRI Preprocessing Pipeline**

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The preprocessing is in six parts

# The preprocessing is in six parts

## ***Signal preprocessing***

1. Preprocessing of anatomical images
2. Preprocessing of functional images
3. Anatomical standardization of functional images
4. Removal of noise signal

## ***Network construction***

5. Construction of nodes
6. Construction of links

We use the  
AFNI, FSL and WMTSA  
software packages

# We do most analyses with AFNI and FSL

- Two C software packages for processing MRI data.
- AFNI: Analysis of Functional NeuroImages.  
Made by the NIH.
- FSL: FMRI Software Library.  
Made by the FMRI in Oxford.

# We do most analyses with AFNI and FSL

1. Preprocessing of anatomical images
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# We construct links with WMTSA

- WMTSA: Wavelet Methods for Time-Series Analysis
- A Matlab and R program for computing frequency-band specific “wavelet” correlations.

# We construct links with WMTSA

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An overview of  
the main steps

# Preprocessing steps

- 1. Preprocessing of anatomical images**
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# Skull-stripping in anatomical image

```
## Get skull-strip mask of anatomical image
```

```
3dSkullStrip \
```

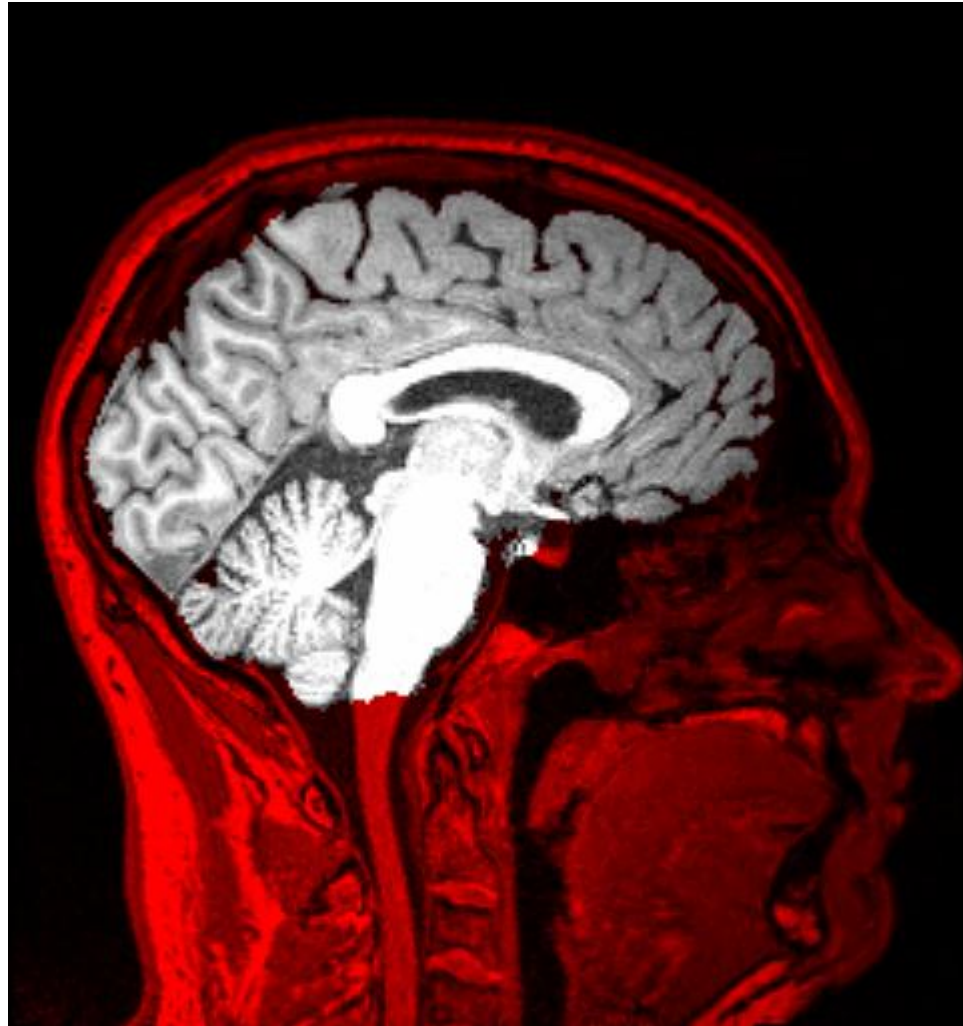
```
    -o_ply anatomical_skullstrip_mask.nii.gz \
    -input anatomical_image.nii.gz \
```

```
## Apply skull-strip mask to anatomical image
```

```
3dcalc \
```

```
    -prefix anatomical_skullstrip.nii.gz \
        -expr 'a*step(b)' \
        -b anatomical_skullstrip_mask.nii.gz \
    -a anatomical_image.nii.gz \
```

# Skull-stripping in anatomical image



# Preprocessing steps

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# Slice-timing correction of functional image

```
## Correct functional image for slice-timing
```

```
3dTshift \
```

```
    -prefix ${func}_st_F.nii.gz \
```

```
        -tpattern altpus \
```

```
${func}_ro_F.nii.gz \
```

# Motion correction (realignment) of functional image

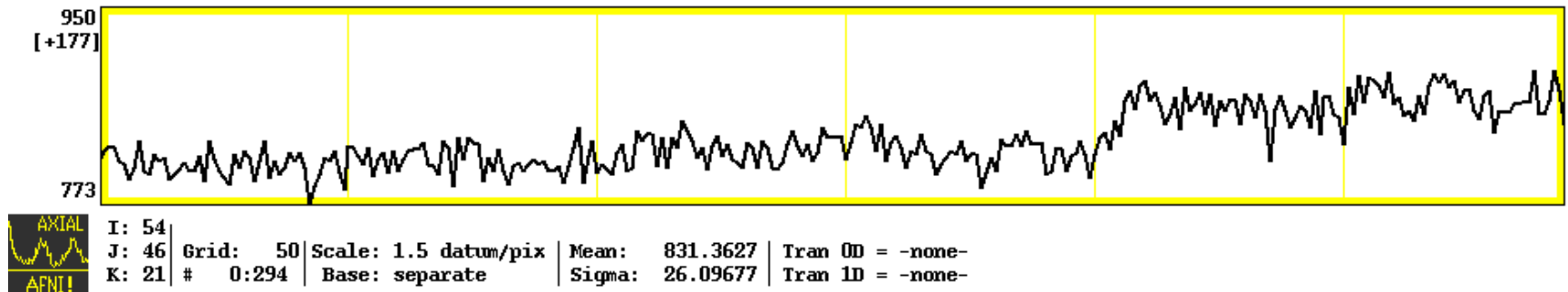
## Correct functional image for motion

**3dvolreg \**

```
-prefix ${func}_mc_F.nii.gz \  
  -1Dfile ${func}_motion.1D \  
  -Fourier -twopass -zpad 4 \  
  -base ${func}_st_F.nii.gz[$ind] \  
  ${func}_st_F.nii.gz \  

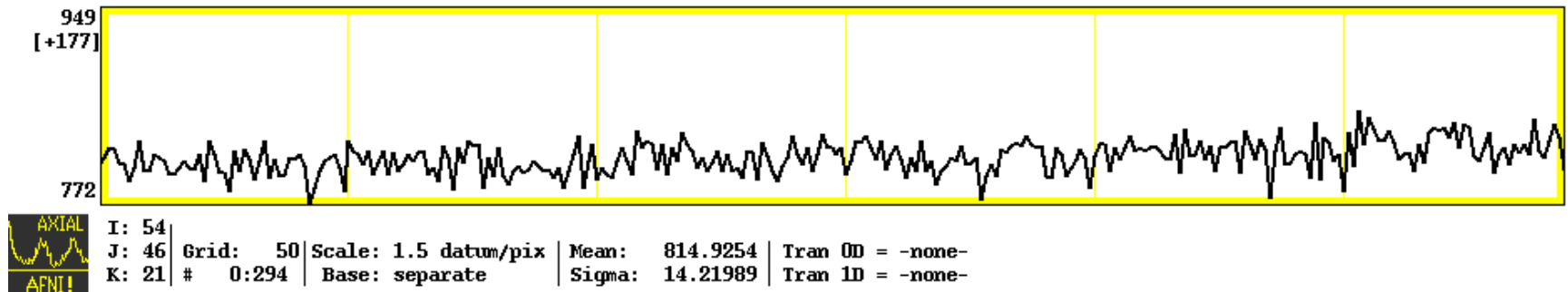
```

# Motion correction (realignment) of functional image





# Motion correction (realignment) of functional image



# Despiking of functional image

```
## Despike functional image
```

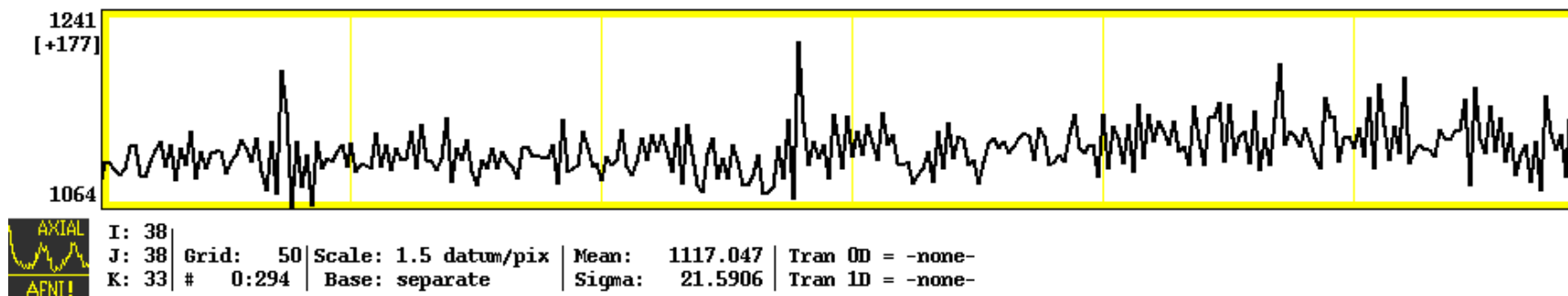
```
3dDespike \
```

```
    -prefix ${func}_ds_F.nii.gz \
```

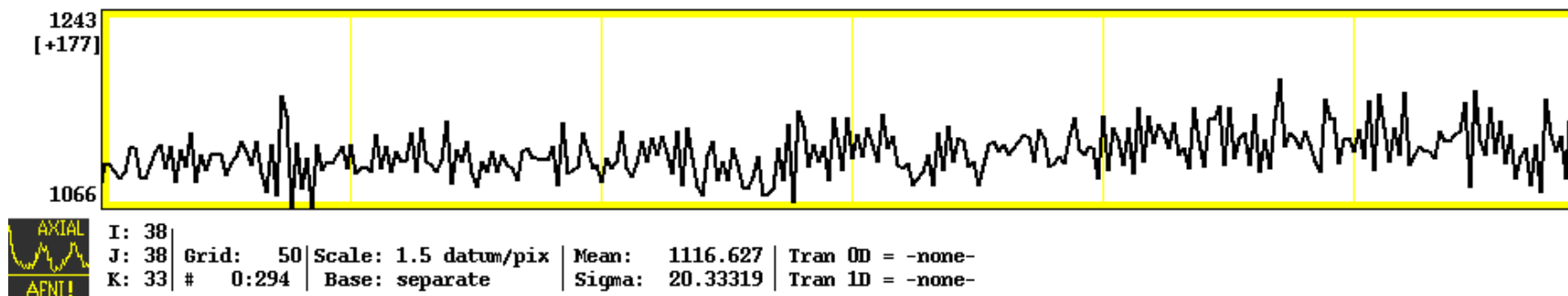
```
        -ssave ${func}_spikiness.nii.gz \
```

```
${func}_mc_F.nii.gz \
```

# Despiking of functional image



# Despiking of functional image



# Preprocessing steps

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# Registration of functional image to high-resolution anatomical image

```
## Get functional-to-anatomical image registration
```

```
flirt \
```

```
    -omat func2anat.mat \
```

```
        -cost corratio -dof 12 -interp trilinear \
```

```
        -ref ${anat}_pp_A.nii.gz \
```

```
    -in ${func}_pp_${ind}_F.nii.gz \
```

```
## Apply functional-to-anatomical image registration.
```

```
flirt \
```

```
    -out ${func}_pp_A.nii.gz \
```

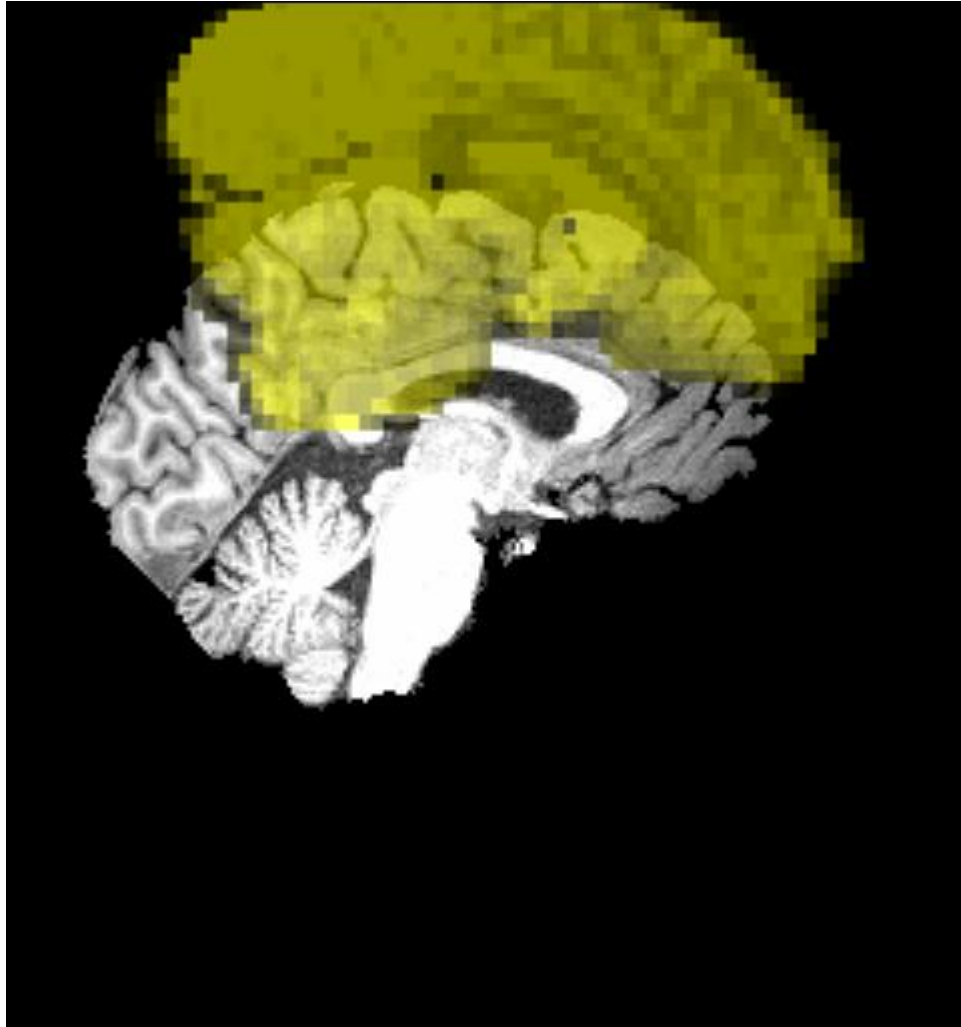
```
        -interp trilinear \
```

```
        -applyxfm -init func2anat.mat \
```

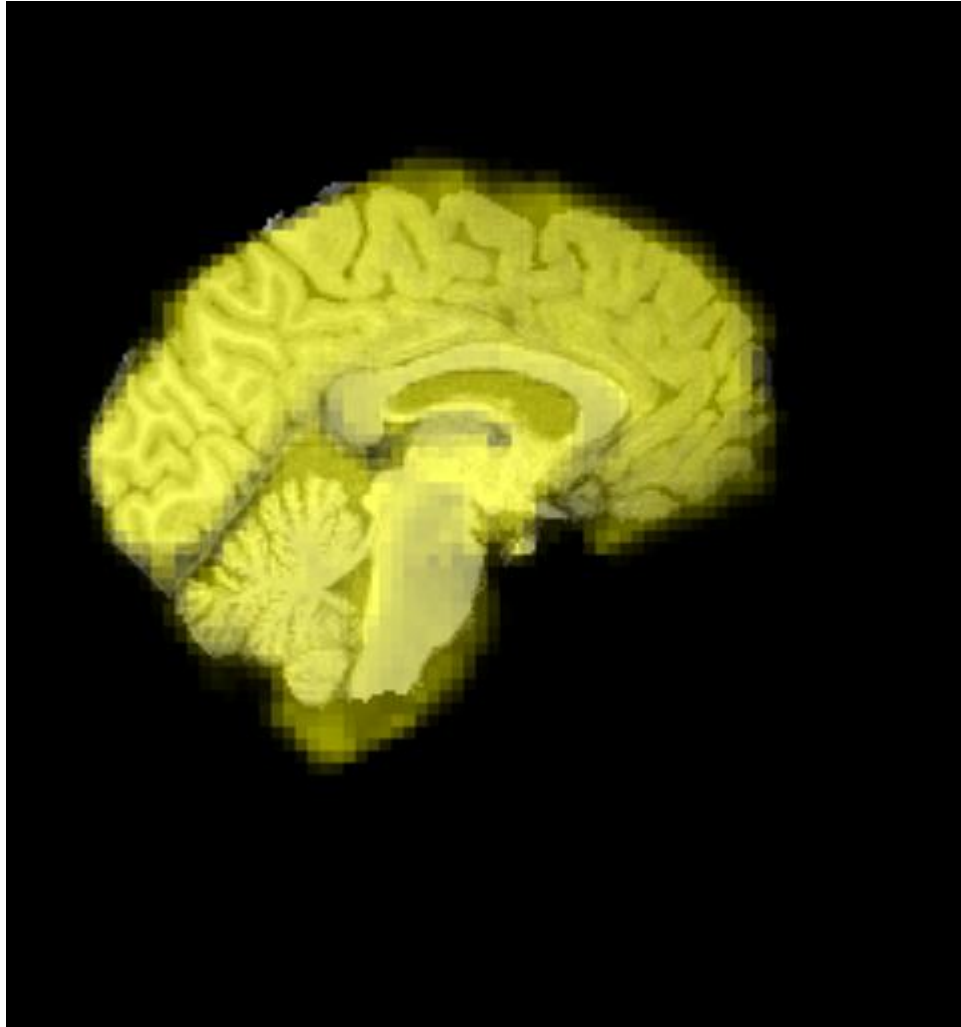
```
        -ref ${anat}_pp_rs_A.nii.gz \
```

```
    -in ${func}_pp_F.nii.gz \
```

# Registration of functional image to high-resolution anatomical image



# Registration of functional image to high-resolution anatomical image





# Standardization of functional image

```
## Get anatomical-to-standard image registration
```

```
flirt \
```

```
    -omat anat2stnd.mat \
```

```
        -cost corratio -dof 12 -interp trilinear \
```

```
        -ref stnd_pp_T.nii.gz \
```

```
    -in ${anat}_pp_A.nii.gz \
```

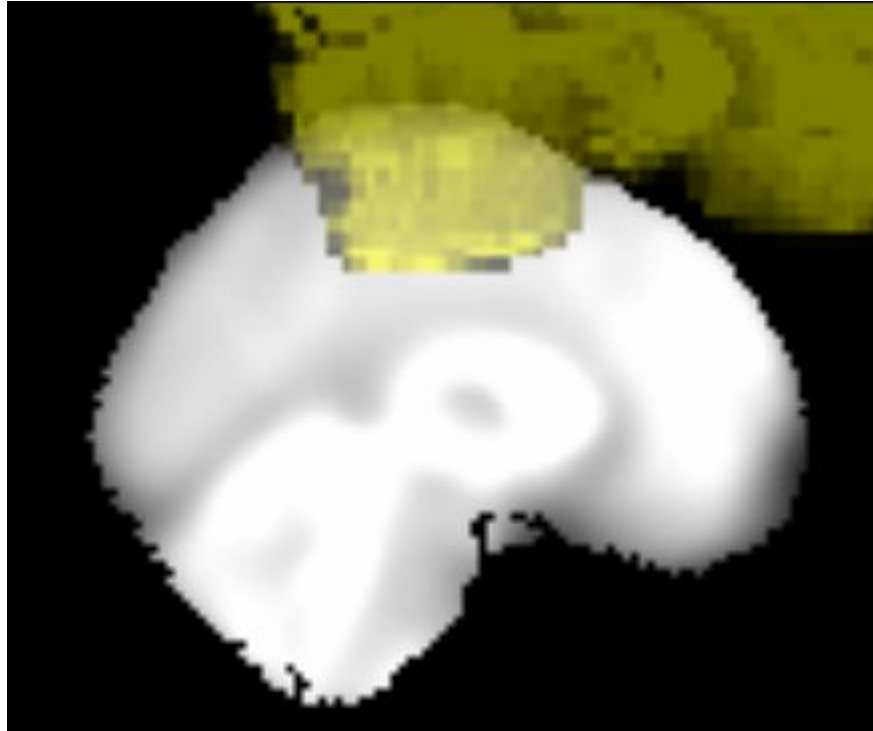
```
## Get functional-to-standard image transformation
```

```
convert_xfm \
```

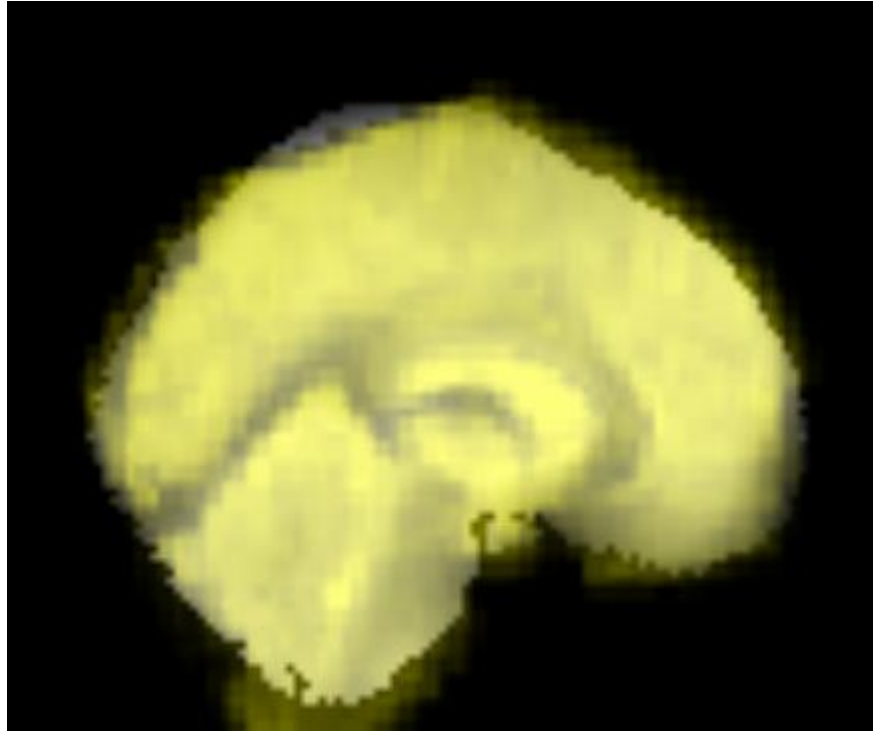
```
    -omat func2stnd.mat \
```

```
    -concat anat2stnd.mat func2anat.mat \
```

# Standardization of functional image



# Standardization of functional image



# Preprocessing steps

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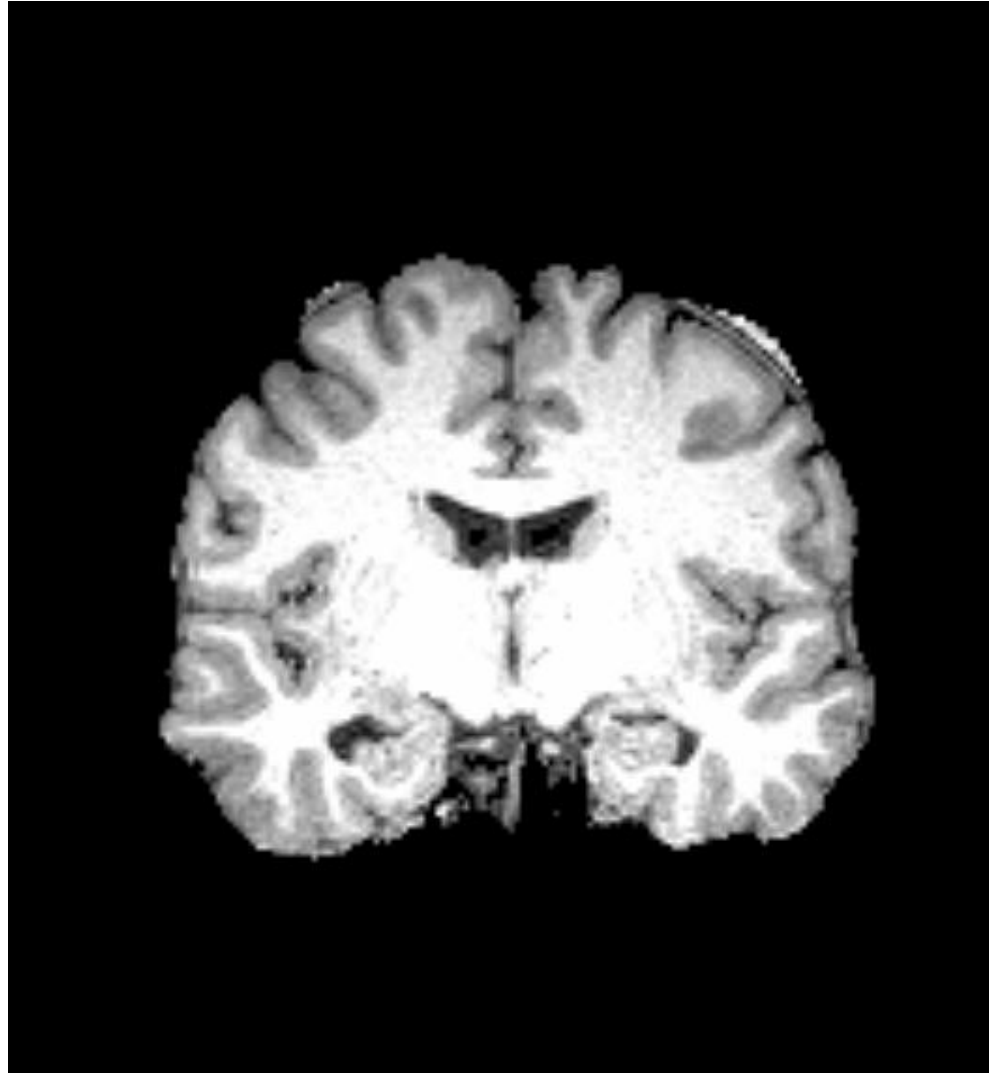
# Segmentation of anatomical image into grey matter, white matter and CSF

```
## Segment anatomical image
```

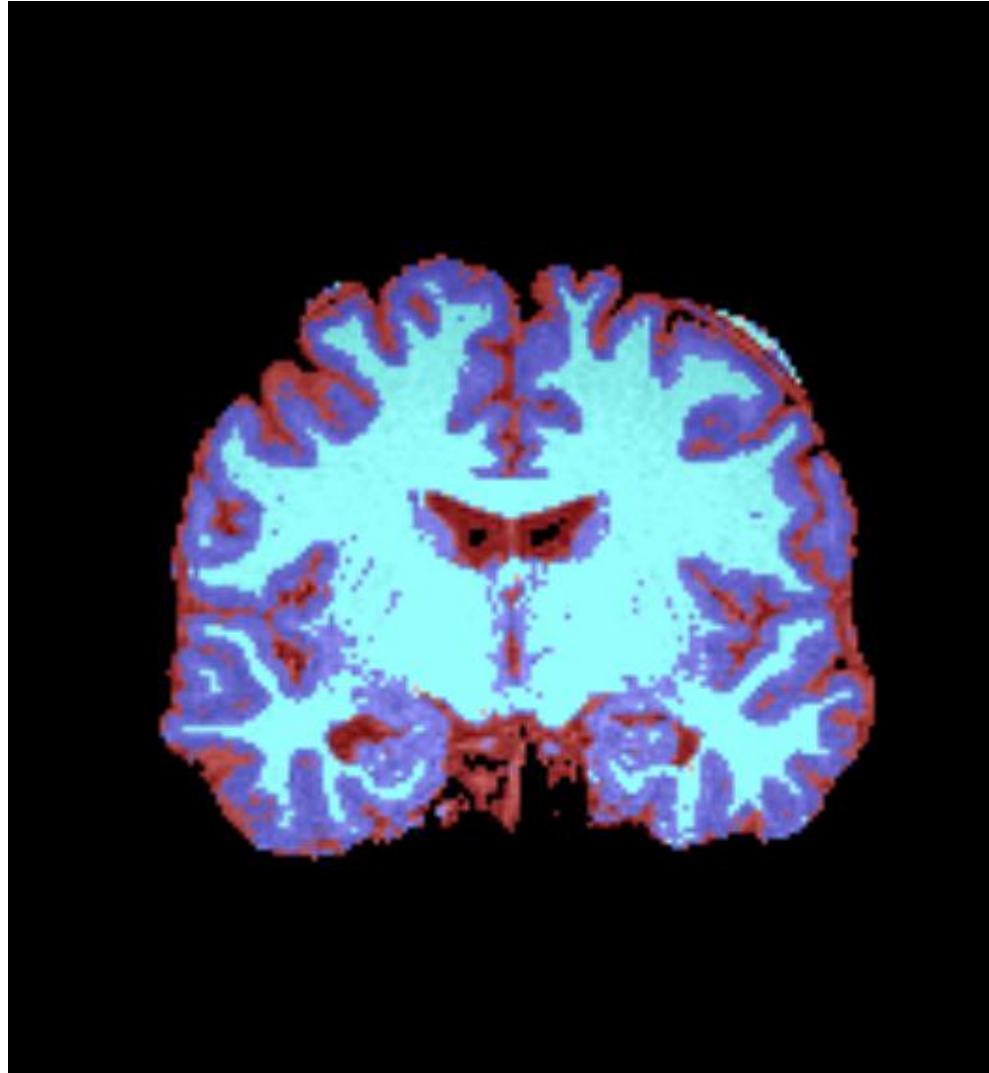
```
fast \  
    -o ${anat}_segm_A \  
        -t 1 -n 3 -g -p \  
    ${anat}_pp_A.nii.gz \  

```

# Segmentation of anatomical image into grey matter, white matter and CSF



# Segmentation of anatomical image into grey matter, white matter and CSF



# Mean CSF signal

```
## Get mean signal of CSF segment
```

```
3dmaskave \
```

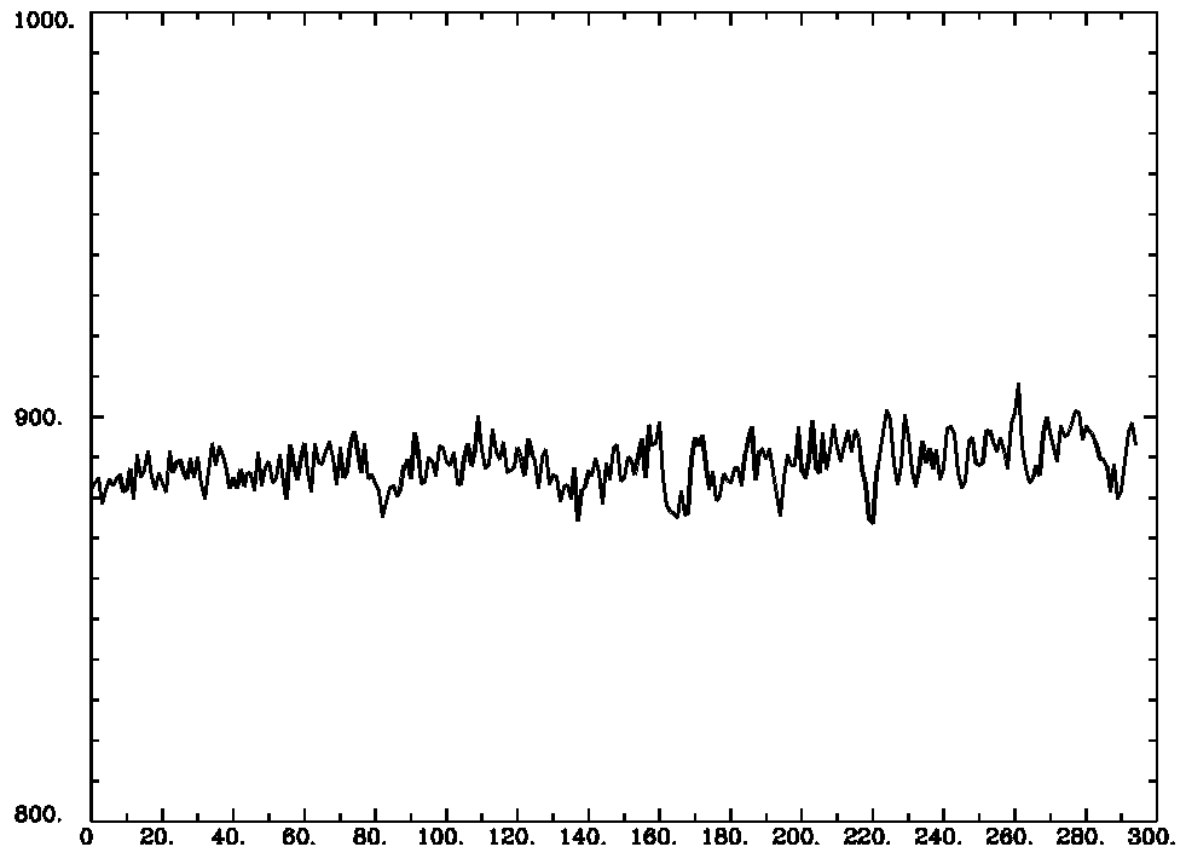
```
-quiet \
```

```
-mask ${func}_csf_A.nii.gz \
```

```
${func}_pp_A.nii.gz > ${func}_csf.1D \
```



# Mean CSF Signal



# Preprocessing steps

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# Motion parameters

## Correct functional image for motion

**3dvolreg \**

-prefix **\${func}\_mc\_F.nii.gz \**

-1Dfile **\${func}\_motion.1D \**

-Fourier -twopass -zpad 4 \

-base **\${func}\_st\_F.nii.gz[\$ind] \**

**\${func}\_st\_F.nii.gz \**

# Motion derivative

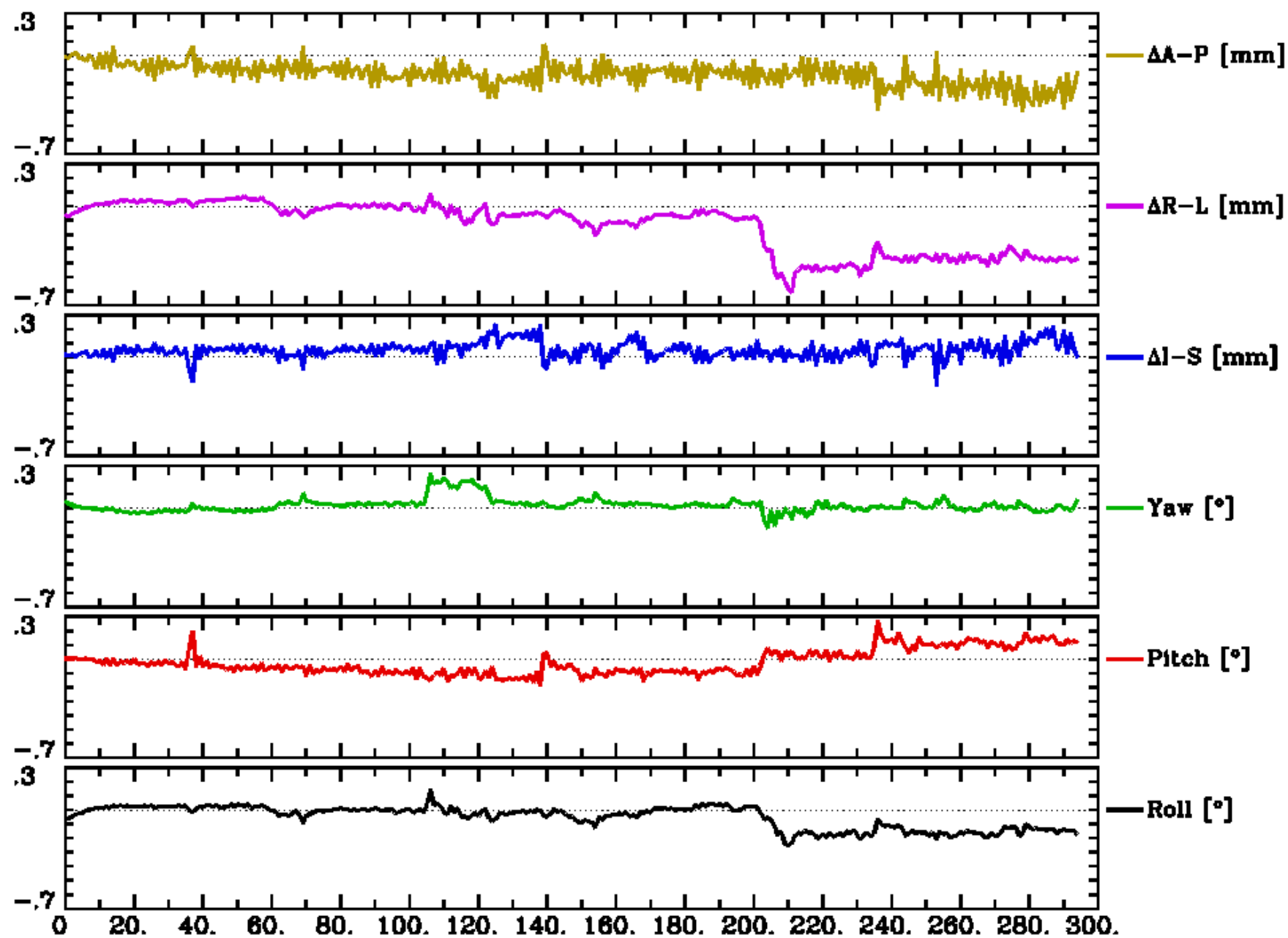
```
## Get motion derivative
```

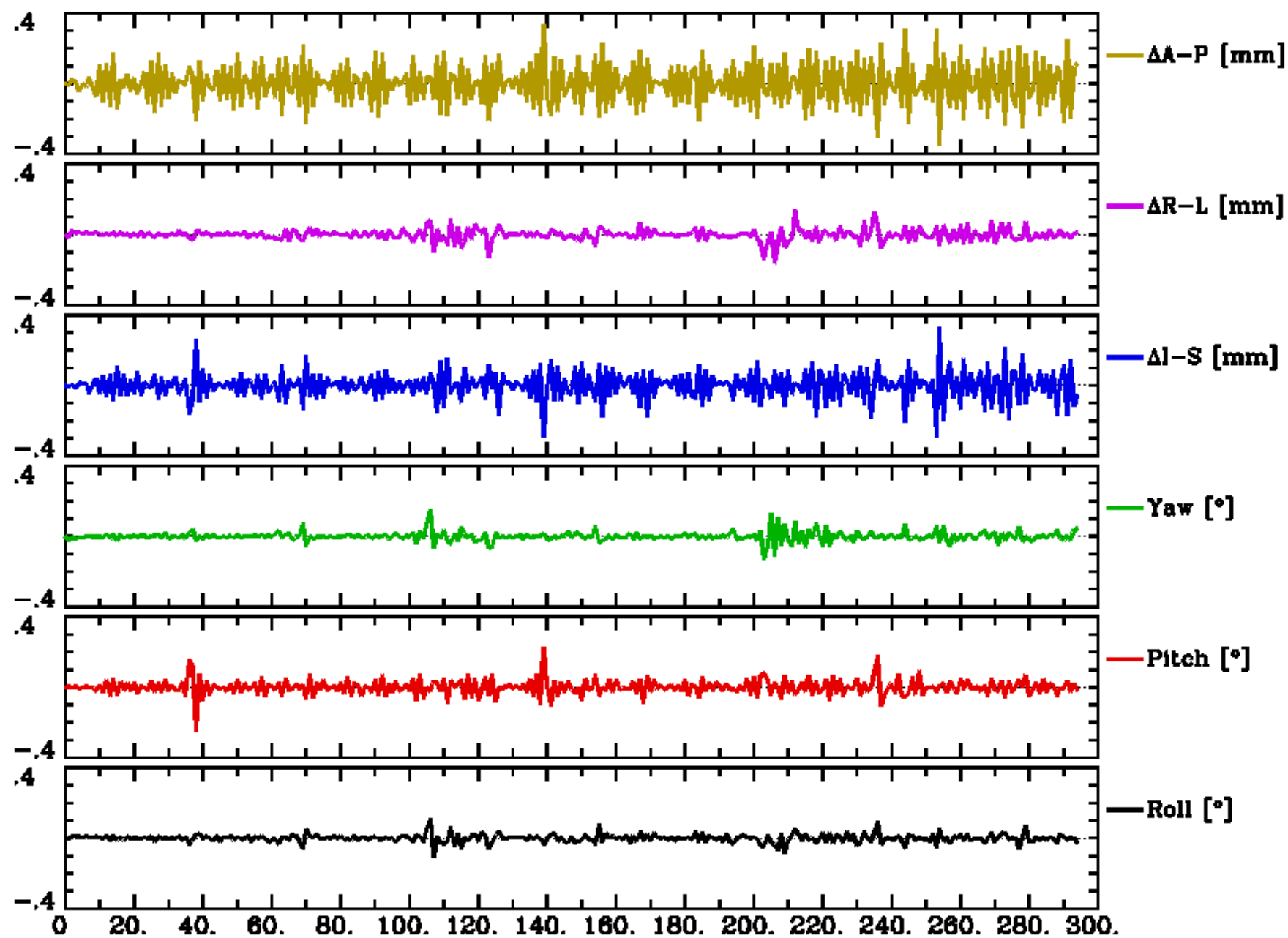
```
1d_tool.py \
```

```
    -write ${func}_motion_deriv.1D \
```

```
        -derivative \
```

```
    -infile ${func}_motion.1D \
```





# Removal of the noise signal

```
## Concatenate CSF signal, motion parameters, motion  
derivative into 'noise signal'
```

```
1dcat
```

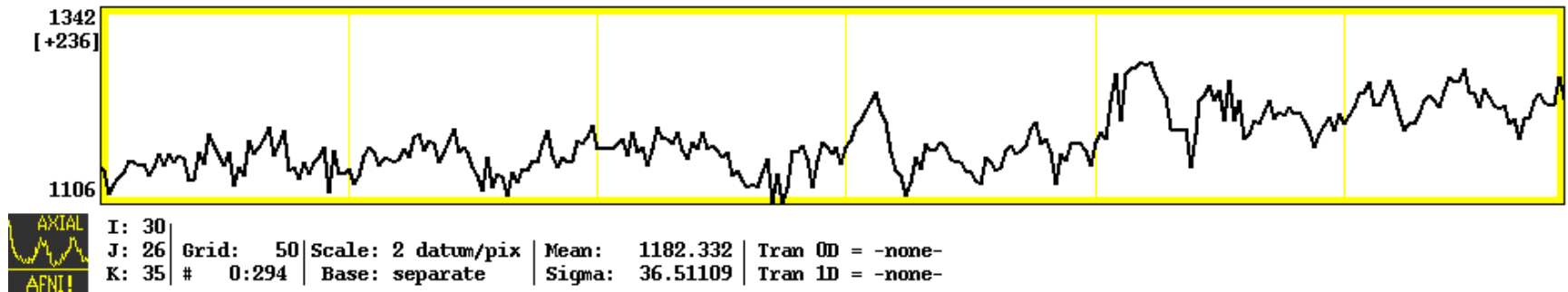
```
    ${func}_csf.1D ${func}_motion.1D  
    ${func}_motion_deriv.1D > ${func}_noise.1D
```

```
## Regress out the 'noise signal' from functional image
```

```
3dBandpass \
```

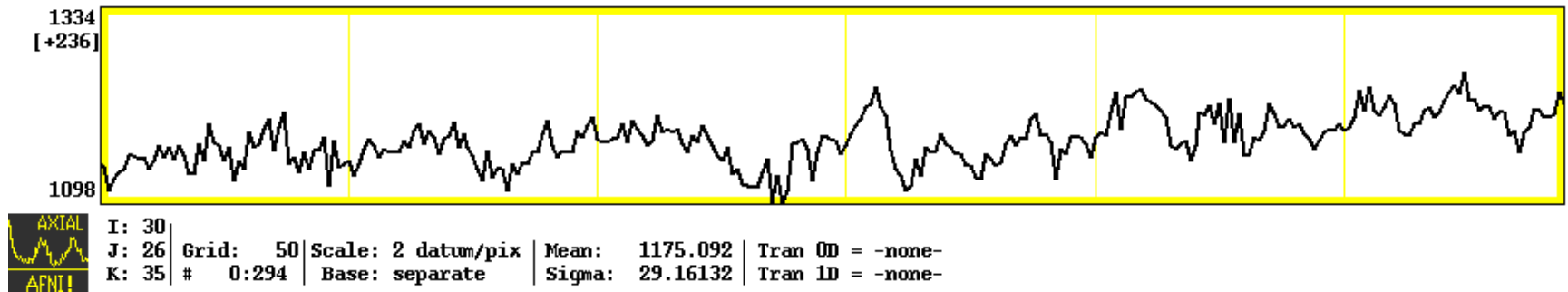
```
    -prefix ${func}_cl_F.nii.gz \  
        -mask ${func}_pp_F.nii.gz[$ind] \  
        -ort ${func}_noise.1D \  
        0.02 99999 \  
    ${func}_pp_F.nii.gz \  
    \
```

# Example uncorrected signal

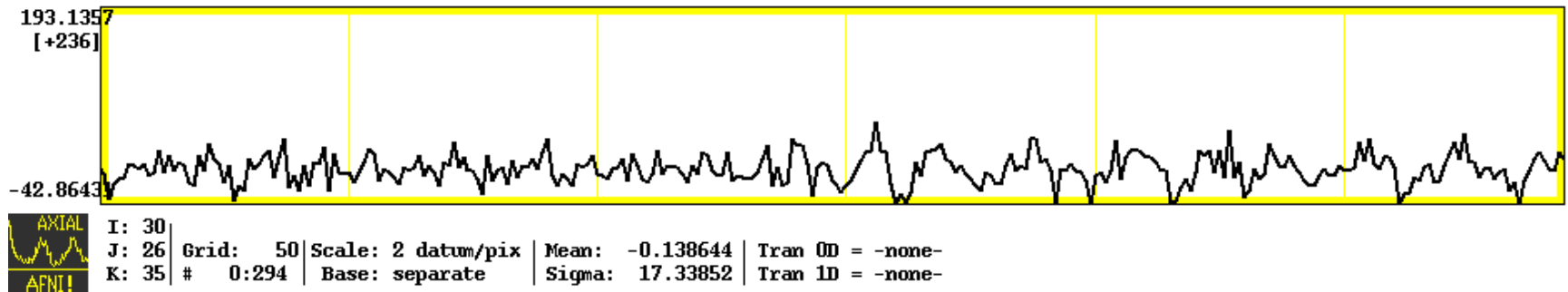




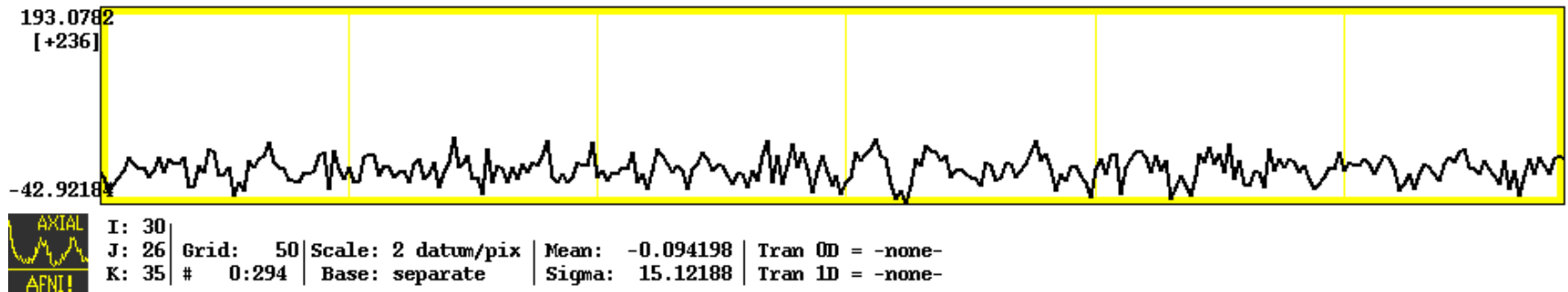
# Motion-corrected and despiked signal



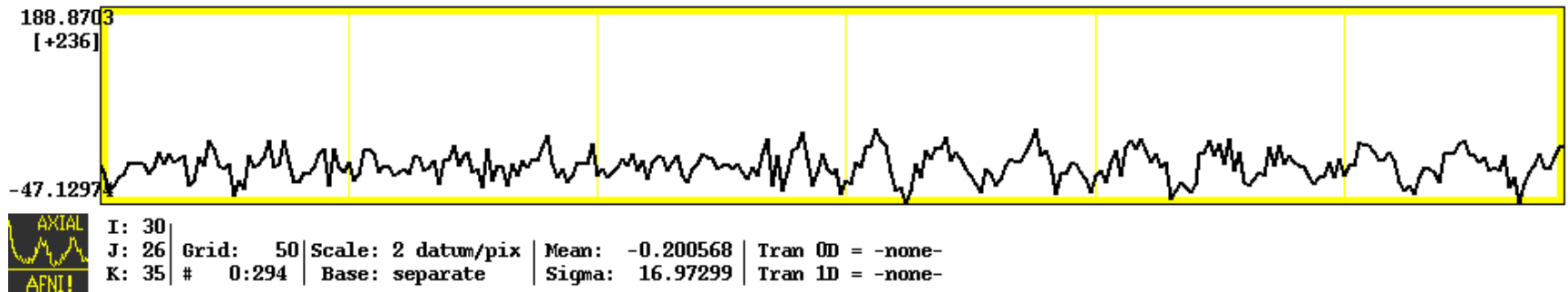
# CSF signal is regressed out



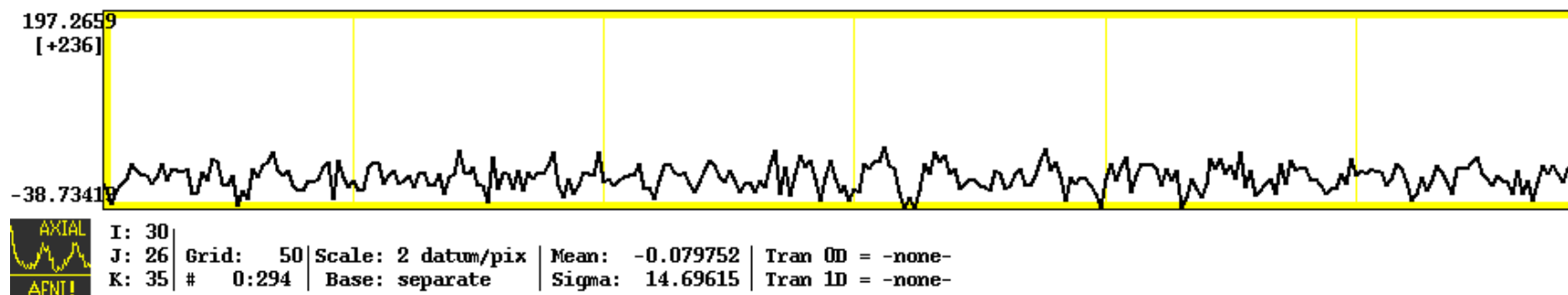
# Motion parameters are regressed out



# Motion derivative is regressed out



# CSF signal, motion parameters and motion derivative are regressed out



# Preprocessing steps

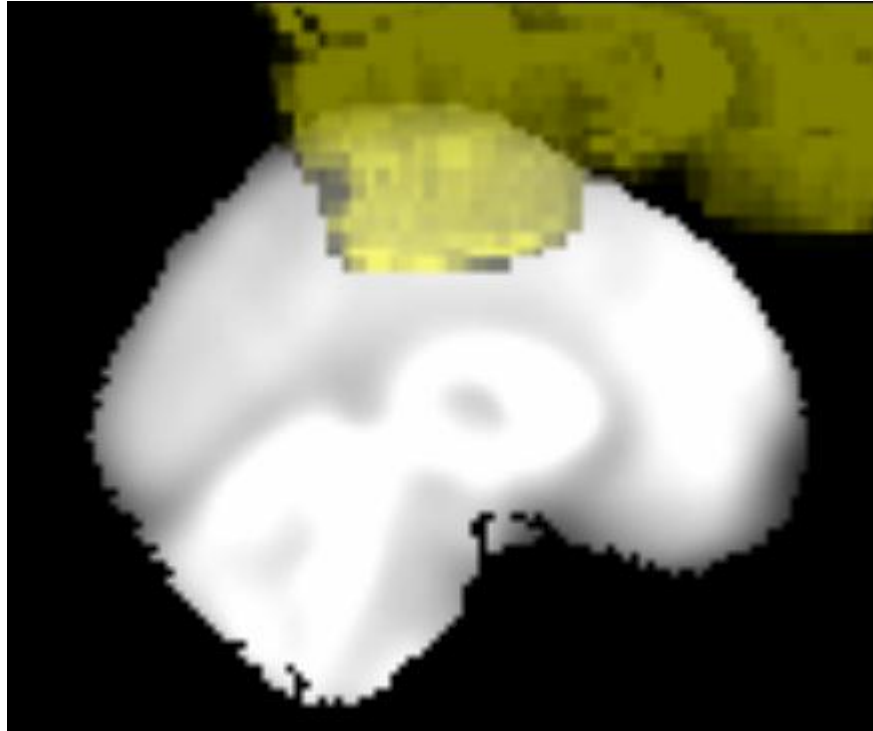
1. Preprocessing of anatomical images
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# Standardization of functional image

```
## Apply functional-to-standard image  
registration.
```

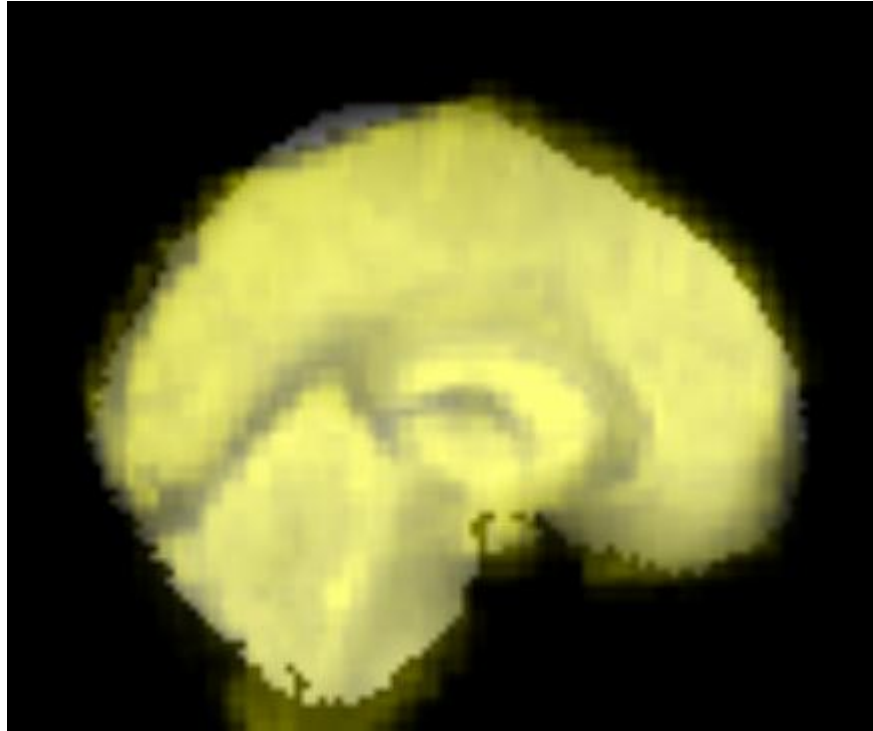
```
flirt \  
    -out ${func}_cl_norm_T.nii.gz \  
        -interp trilinear \  
        -applyxfm -init func2std.mat \  
        -ref std_pp_rs_T.nii.gz \  
    -in ${func}_cl_norm_F.nii.gz \  
    -out ${func}_cl_norm_T.nii.gz
```

# Standardization of functional image





# Standardization of functional image



# Parcellation of functional image



# Computation of mean signal in parcel $i$

```
## Get mean parcel signal
```

```
3dmaskave \
```

```
-quiet \
```

```
-mrange $i $i \
```

```
-mask atls_pp_T.nii.gz \
```

```
${func}_cl_norm_T.nii.gz > _t${i}.1D
```

# Preprocessing steps

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# Computation of wavelet correlations

```
wf = 'd4';                    % wavelet filter
s = 3;                        % wavelet scale
[n t] = size(M);
ts = modwt_num_nonboundary_coef(wf,t,s);
Ms=zeros([n ts]);
for j = 1:n
    ms = modwt(M(j,:), wf, s, 'circular').';
    Ms(j,:) = ms(s,(t+1-ts):t);
end
M_net = corrcoef(Ms.');
```

# The final network

