



DPABISurf: A Surface-Based Resting-State fMRI Data Analysis Toolbox

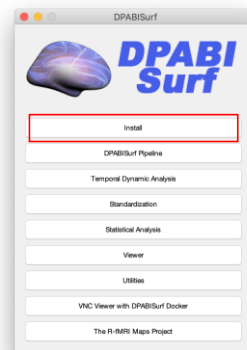
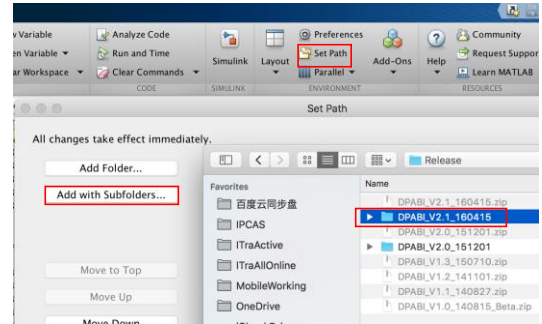
Chao-Gan YAN, Ph.D.

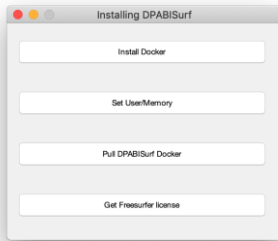
严超轶

ycg.yan@gmail.com

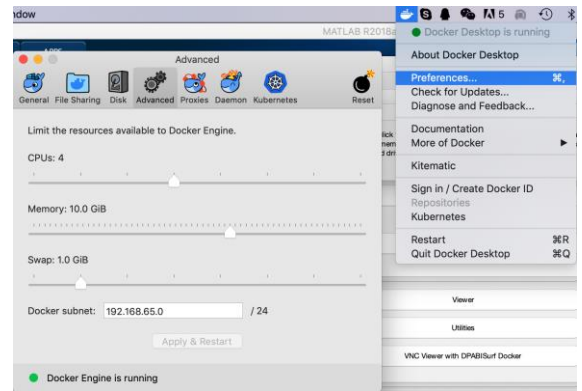
<http://rfmri.org>

Institute of Psychology, Chinese Academy of Sciences





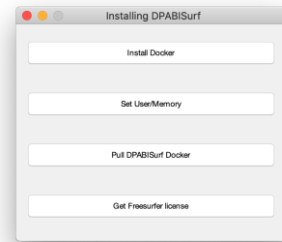
7



8

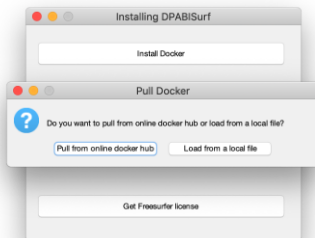


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Linux: `sudo groupadd docker`
`sudo usermod -aG docker $USER`

10

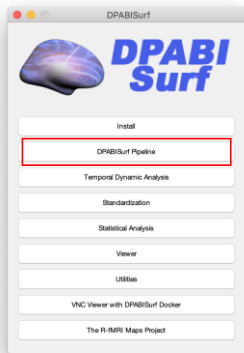


11

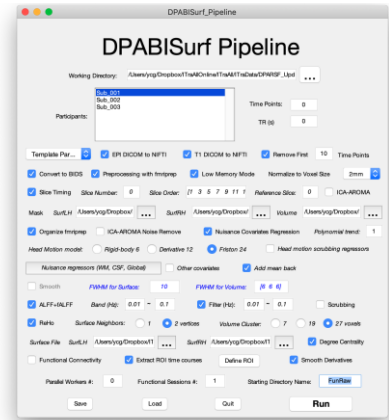


Please get Freesurfer license, and specify the license.txt you received in your email. Please visit:
<https://surfer.nmr.mgh.harvard.edu/registration>

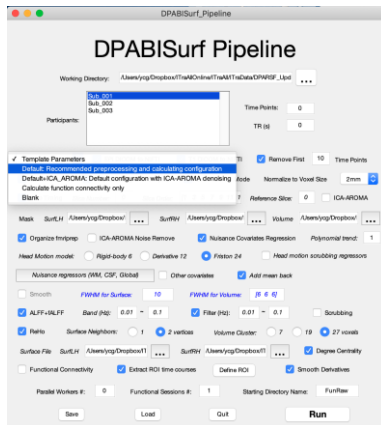
12



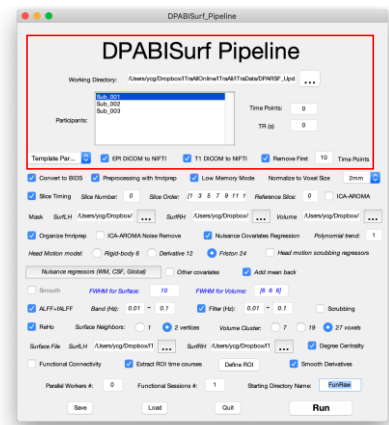
13



14



15

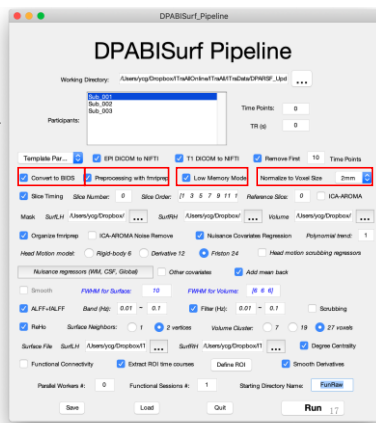


16

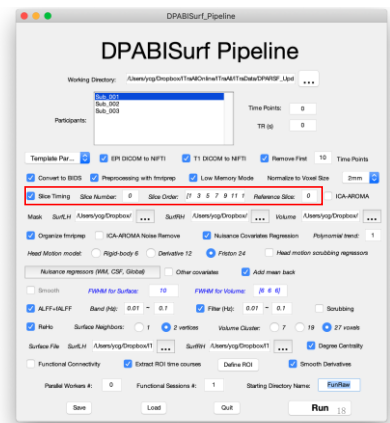
BIDS structure:
<http://bids.neuroimaging.io>
 fMRIPrep
 Esteban et al., 2018.
 Nat Methods.

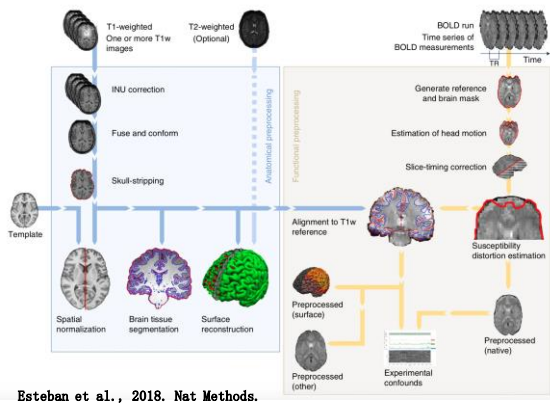
Low memory mode
 (fMRIPrep)

Voxel size
 (fMRIPrep)

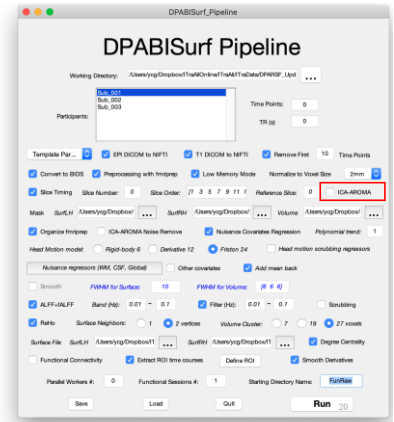


Slice timing
 (fMRIPrep)





ICA-AROMA
(fMRIPrep)



ICA-AROMA

ICA-AROMA

ICA
Register IC spatial maps
to MNI152 2mm

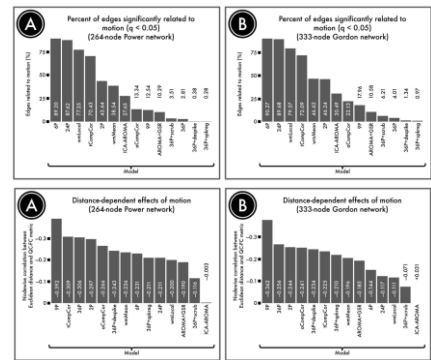
Motion Component Classification
Based on four features:
maximum *RP* correlation
Edge fraction
CSF fraction
High-frequency content

fMRI data denoising
Removal of classified ICs from
the fMRI data (fsl_regfilt)

Pruim et al., 2015. Neuroimage

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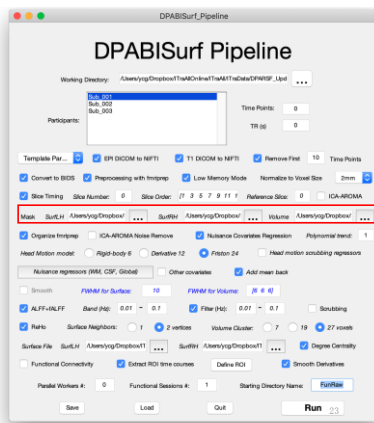
ICA-AROMA



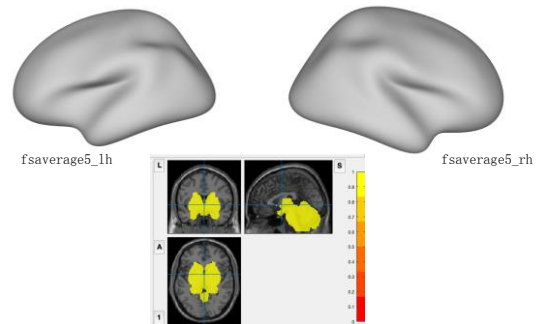
Ciric et al., 2017. Neuroimage

Mask

E. g.
(DPABIPath)/DPABISu
rf/SurfTemplates/fs
average5_lh_cortex.
label.gii



Three Spaces

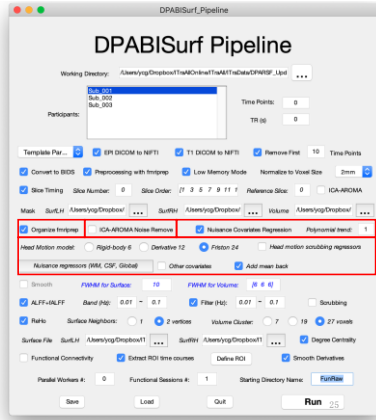


Volume space
can be restricted within freesurfer_subcortical_mask

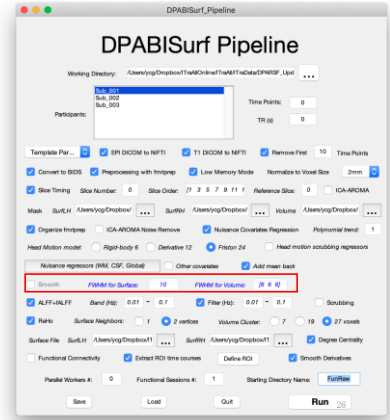
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Organize the output files from fMRIPrep
Non-aggressively regressed out the covariates of ICA-AROMA noises.

Nuisance Regression
Same as DPARSF



Smooth



Spatial Smoothing

Why should you smooth?

- Might Improve CNR/SNR
- Improve intersubject registration

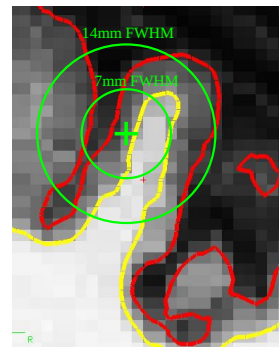
How much smoothing?

- Blob-size
- Typically 5-20 mm FWHM
- Surface smoothing more forgiving than volume-based

Based on Freesurfer Course

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Volume-based Smoothing

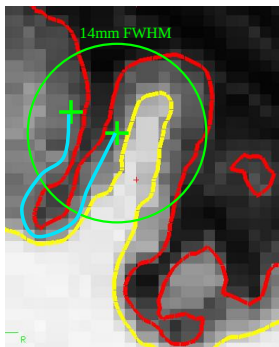


- Smoothing is averaging of “nearby” voxels

Based on Freesurfer Course

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Volume-based Smoothing



Based on Freesurfer Course

- 5 mm apart in 3D
- 25 mm apart on surface!
- Kernel much larger
- Averaging with other tissue types (WM, CSF)
- Averaging with other functional areas

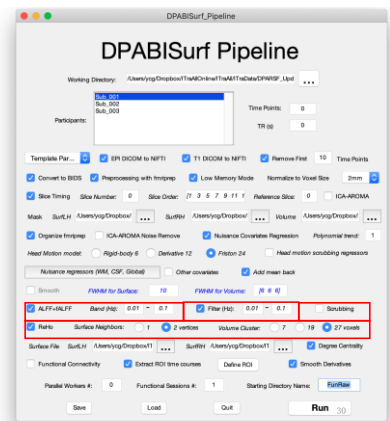
29

ALFF

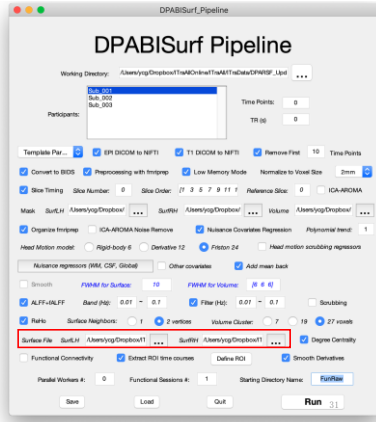
Filter

Scrubbing

ReHo



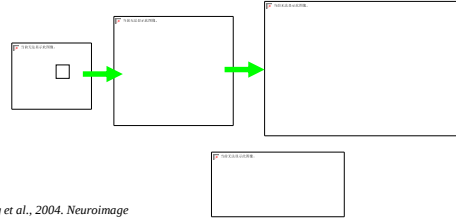
Provide surface file
for 2d ReHo
Calculation



3D ReHo

Regional Homogeneity (ReHo)

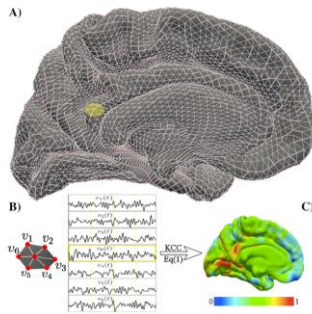
Similarity or coherence of the time courses
within a functional cluster



Zang et al., 2004. Neuroimage

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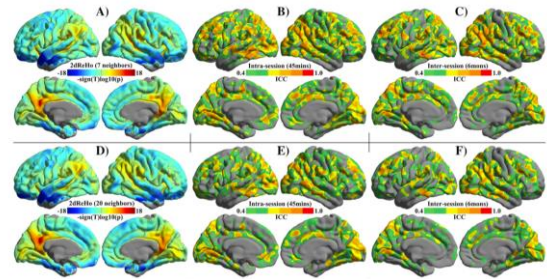
2D ReHo



Zuo et al., 2013. Neuroimage

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2D ReHo



Zuo et al., 2013. Neuroimage

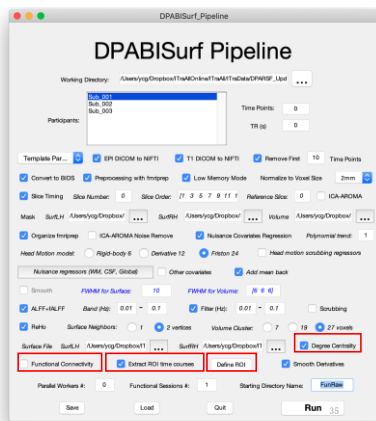
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Degree Centrality

Functional
Connectivity

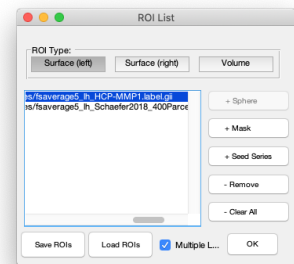
Extract ROI Time
Courses

Define ROI



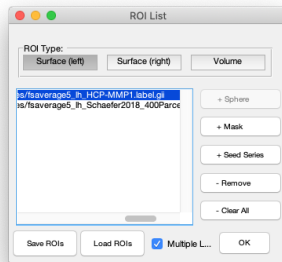
Define ROI

Left Surface
Right Surface
Volume



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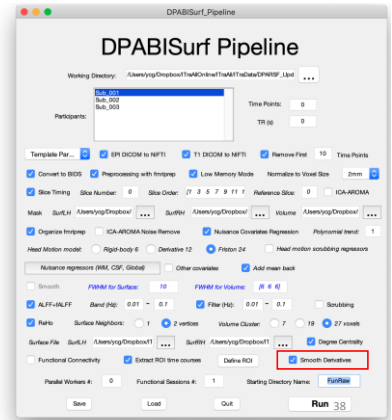
Define ROI



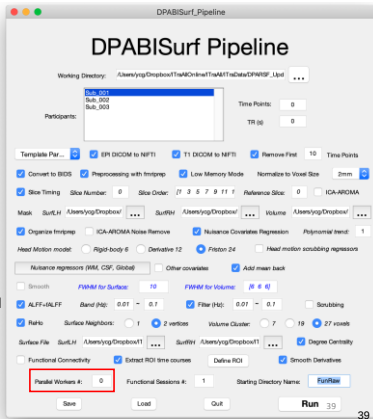
Left Surface
Right Surface
Volume

ROI order:
ROISignals_SurfLHSurfRHVolu_FunSurfWCF

Smooth Derivatives



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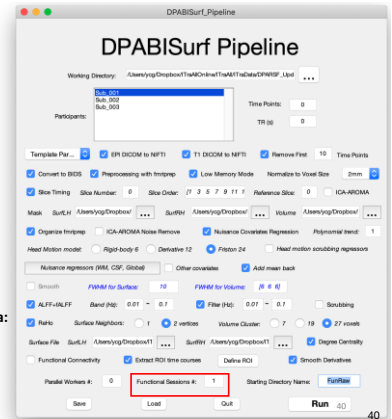
**Parallel Workers (if
parallel computing toolbox
is installed)**

Each subject is distributed into a different worker.
Do not be too much because of fMRIPrep processing.

Multiple functional sessions

1st session: FunRaw
2nd session: S2_FunRaw
3rd session: S3_FunRaw
...

If also have multiple sessions of T1 image data:
 1st session: T1Raw
 2nd session: S2_T1Raw
 3rd session: S3_T1Raw
 ...



**Starting
Directory Name**

If you do not start with raw DICOM images, you need to specify the **Starting Directory Name**.

E.g. "FunSurfWCF" means
normalized, covariates
regressed and filtered

W - Normalize

I - ICA-AROMA Noise non-aggressively regressed

C - Covariates Removed

S - Smooth

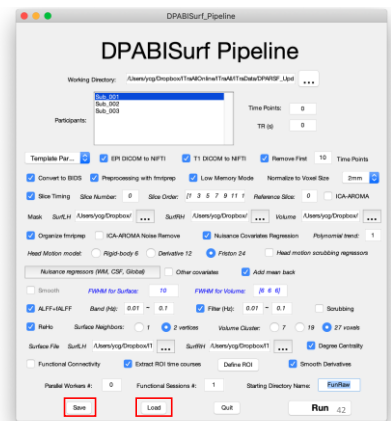
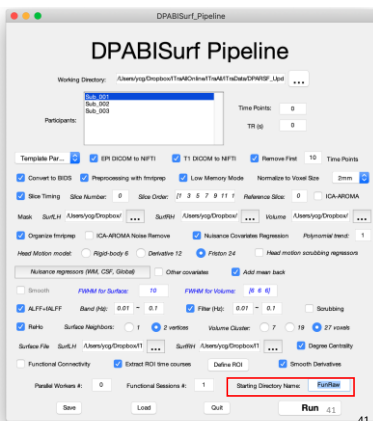
F - Filter

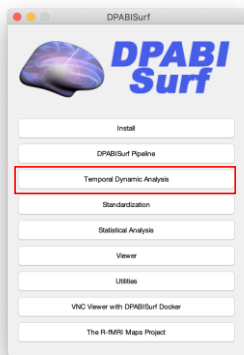
B - ScrubBBing

Save and Load Parameters

Save parameters to
*.mat

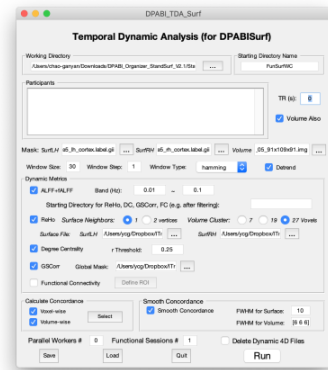
Load parameters
from *.mat





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DPABI_TDA_Surf



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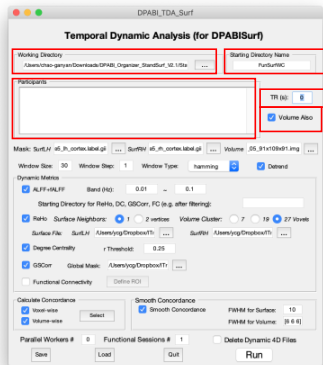
Based on DPABISurf
Preprocessed Data

Starting Directory
Name

Participants

TR

In addition to the
surface-based data,
also process the data
in volume space
processed by
DPABISurf



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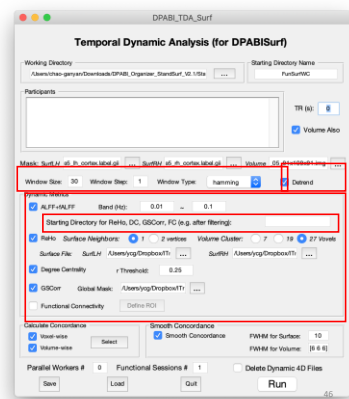
Based on DPARSF
Preprocessed Data

Window Setup

Detrend

Dynamic Indices

If need filtering

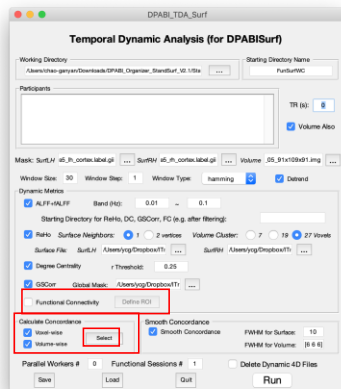
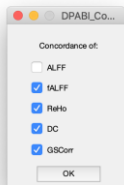


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Based on DPARSF
Preprocessed Data

Functional
Connectivity

Concordance
Settings

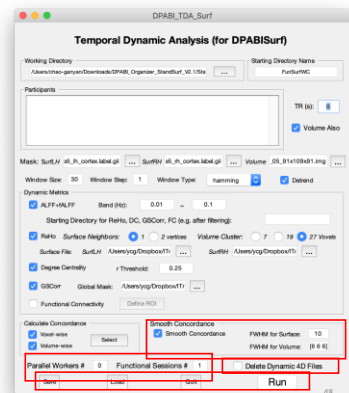


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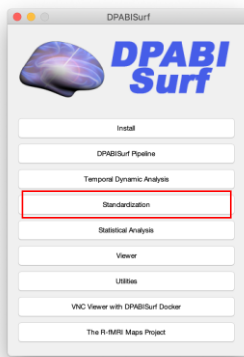
Smooth concordance
maps

Parallel and Sessions
Delete Dynamic 4D
Files to save space

Save/Load
parameters and Run



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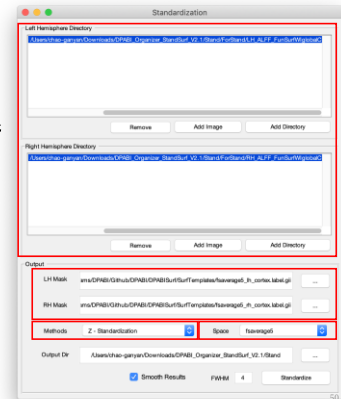


Standardization across both hemispheres

Set data for both left hemisphere and right hemisphere
Masks for both hemispheres

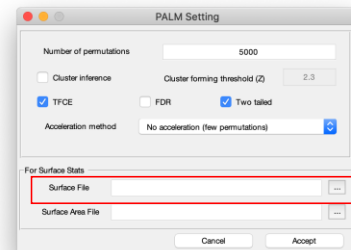
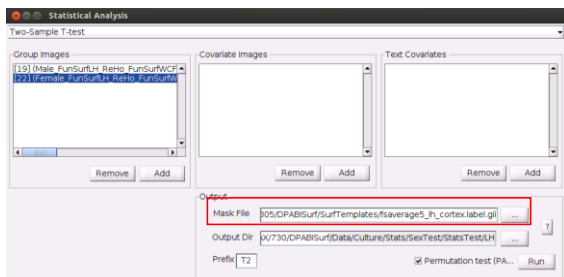
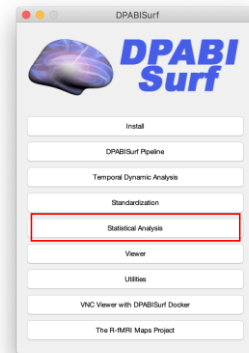
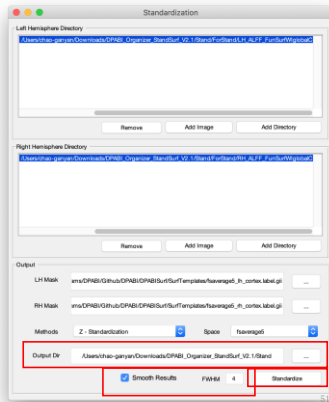
Standardization methods

Space: fsaverage (anatomical) or fsaverage5 (functional)

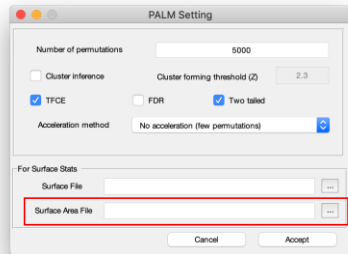


Output directory: left hemisphere and right hemisphere will be separated in different directories automatically
Smooth results: surface-based smoothing

Run

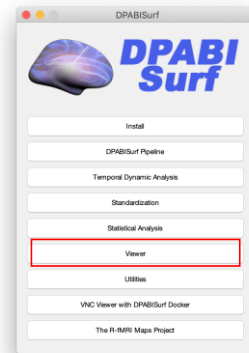


/DPABI/DPABISurf/SurfTemplates/fsaverage5_lh_white.surf.gii



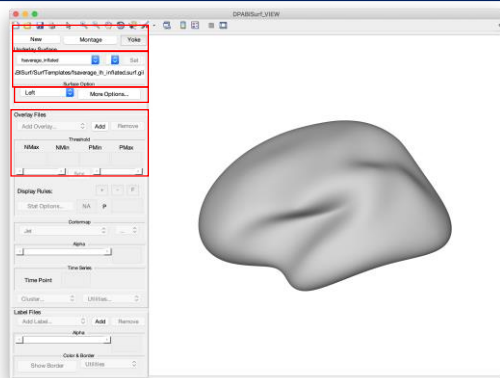
Optional: [DPABI]/DPABISurf/SurfTemplates/fsaverage5_lh_white_avg.area.gii

55



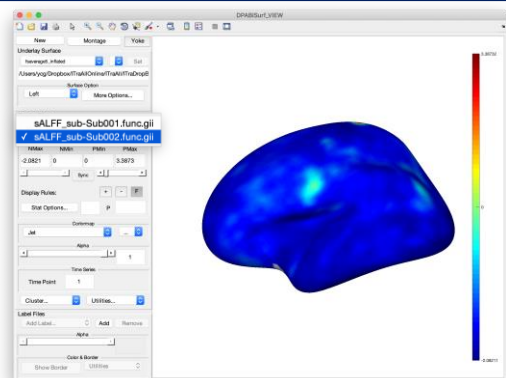
56

DPABISurf_VIEW



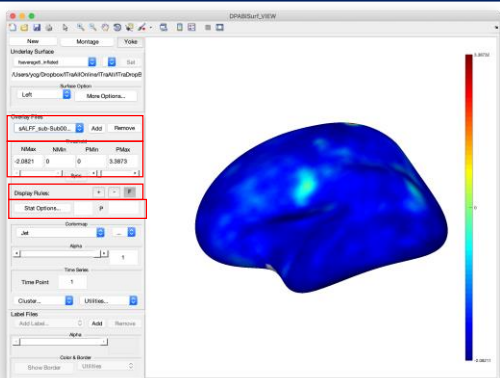
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DPABISurf_VIEW



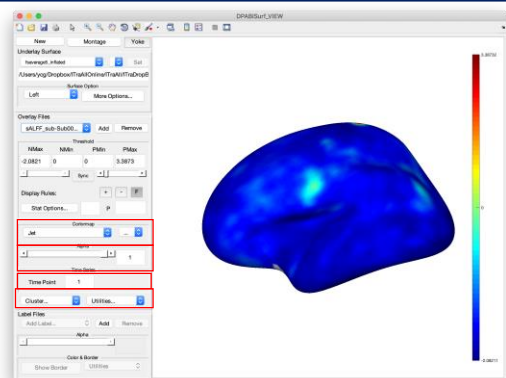
58

DPABISurf_VIEW



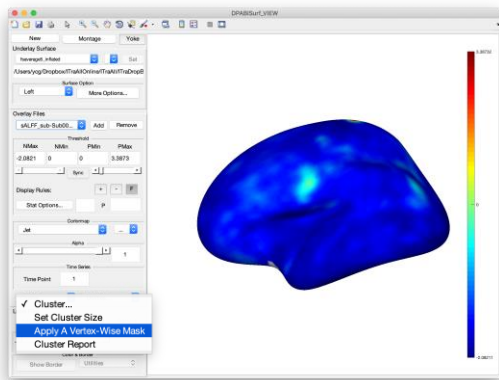
59

DPABISurf_VIEW



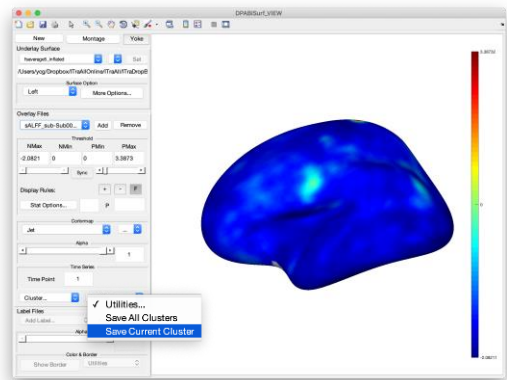
60

DPABISurf_VIEW



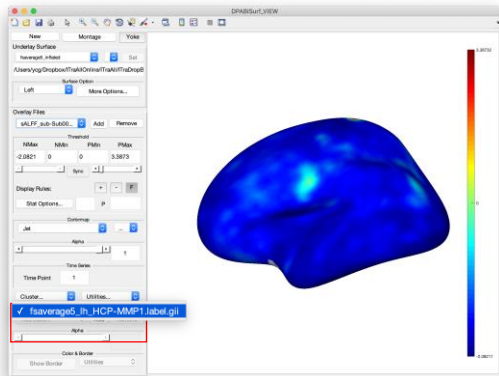
61

DPABISurf_VIEW



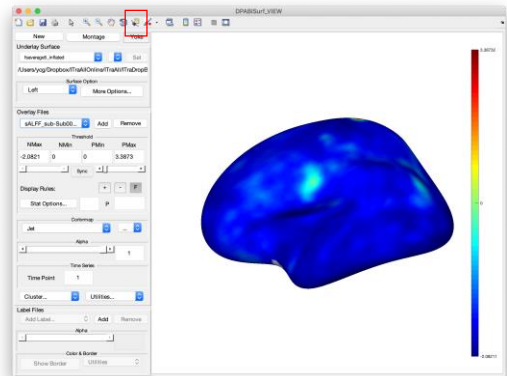
62

DPABISurf_VIEW



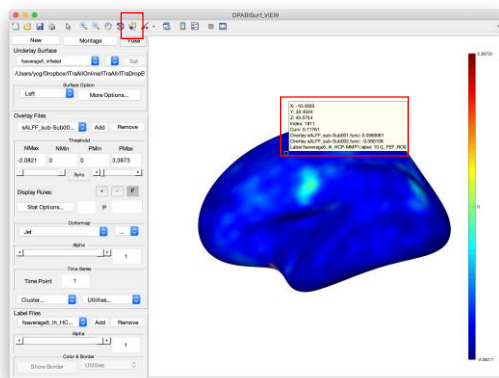
63

DPABISurf_VIEW



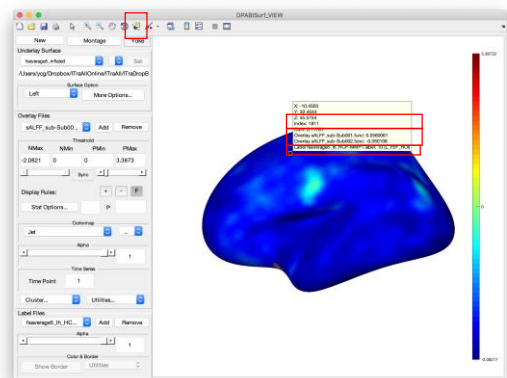
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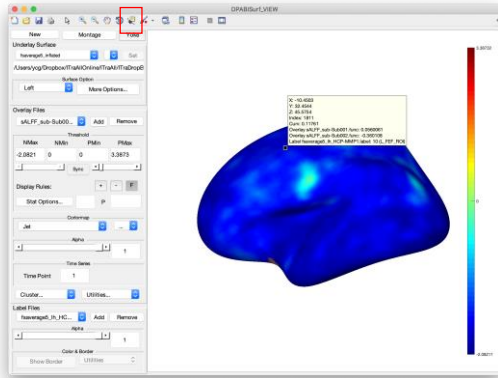
65

DPABISurf_VIEW



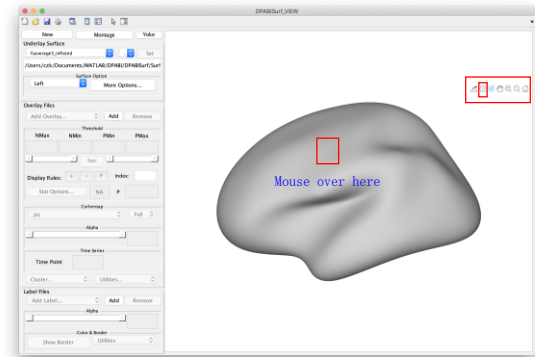
66

DPABISurf_VIEW



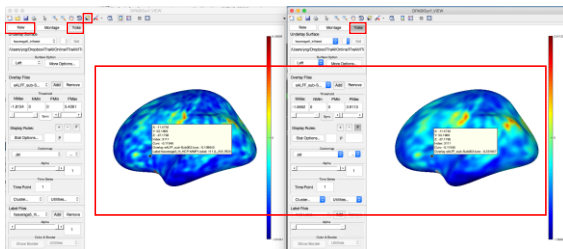
67

DPABISurf_VIEW

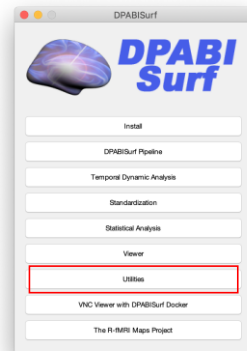


68

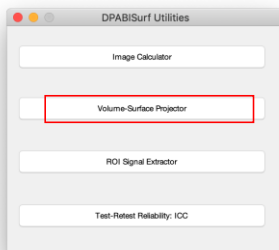
DPABISurf_VIEW



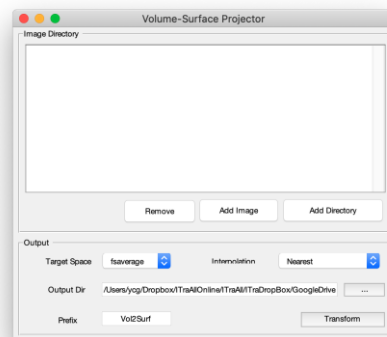
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Use DPABISurf Docker

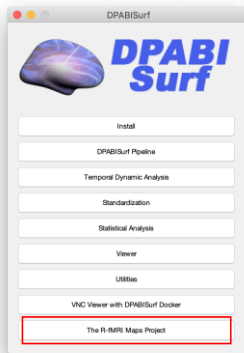
```

mri_convert bin subcort.mask.1mm.mgz freesurfer_subcortical_mask.1mm.nii
sid: mri_convert.c.v.1.226 2016/02/26 16:15:24 mreuter Exp $
reading from subcort.mask.1mm.mgz...
TR=0.00, TE=0.00, TI=0.00, flip angle=0.00
s_ras = (-1, 0, 0)
s_ras = (0, 0, -1)
s_ras = (0, 1, 0)
writing to freesurfer_subcortical_mask.1mm.nii...
root@506ad169e3c:/opt/freesurfer/subjects/fsaverage/mri# ls
T1.mgz          orig.mgz
aparc+aseg.mgz  parc.a2005+aseg.mgz
aparc.a2005+aseg.mgz  p.aseg.mgz
aparc.a2009+aseg.mgz  rh.ribbon.mgz
aseg.mgz           ribbon.mgz
brain.mgz          subcort.mask.1mm.README
brainmask.mgz      subcort.mask.1mm.mgz
freesurfer_subcortical_mask.1mm.nii  subcort.prob.log
freesurfer_subcortical_prob.1mm.nii  subcort.prob.mgz
rh.ribbon.mgz      transforms
un335.cor.mgz
root@506ad169e3c:/opt/freesurfer/subjects/fsaverage/mri# ls *.nii
freesurfer_subcortical_mask.1mm.nii  freesurfer_subcortical_prob.1mm.nii
root@506ad169e3c:/opt/freesurfer/subjects/fsaverage/mri# mv *.nii /data
root@506ad169e3c:/opt/freesurfer/subjects/fsaverage/mri#

```

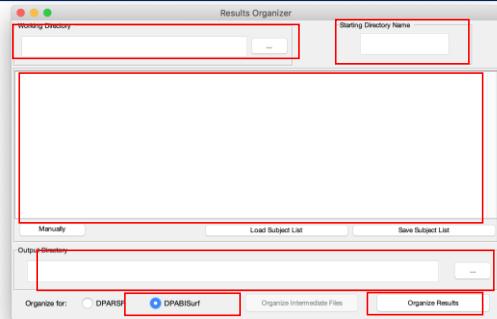
Call DPABI StandAlone:
/opt/DPABI/DPABI_StandAlone/run_DPABI_StandAlone.sh \$(MCRPath)

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The R-fMRI Maps Project



Reading and Writing functions

Reading:

```

[Data, VoxelSize, FileList, Header] = y_ReadAll('XX.func.gii');
Data - 10242*230 single
Header - GIFTI Structure

```

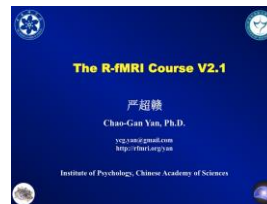
Processing:

```
MeanData = mean(Data, 2);
```

Writing:

```
y_Write(MeanData, Header, 'MeanData.gii');
```

Further Help



<http://rfmri.org/Course>



<http://wiki.rfmri.org>



The R-fMRI Journal Club



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Acknowledgments

Chinese Academy of Sciences **NYU Child Study Center**
Xi-Nian Zuo F. Xavier Castellanos

Hangzhou Normal University **Child Mind Institute**
Yu-Feng Zang Michael P. Milham

Beijing Normal University
Yong He
Xin-Di Wang

Peking University
Jia-Hong Gao
Wei-Wei Men

- Funding**
- National Natural Science Foundation of China
 - Chinese Academy of Sciences

Thanks for your attention!