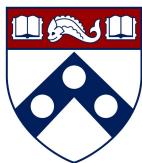




CHILD MIND®
INSTITUTE
big data analytics

C-PAC

 **TEXAS**
The University of Texas at Austin

 Penn
UNIVERSITY OF PENNSYLVANIA



COLUMBIA UNIVERSITY
IN THE CITY OF NEW YORK

<http://fcp-indi.github.io>

- What is C-PAC?
- Why is C-PAC?
- How does C-PAC work?
- How to run C-PAC in a container?
- Demo

Magnetic Resonance Imaging (MRI)

What's MRI?

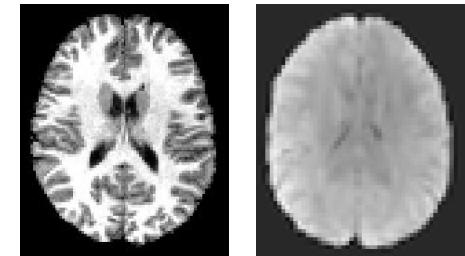
- Medical imaging technique used in radiology to form pictures of the anatomy and the physiological processes of the body

Why is MRI preprocessing important?

- Remove data collection artifacts
- Prepare for group-level analysis
- Improve computational reproducibility

How is MRI preprocessed?

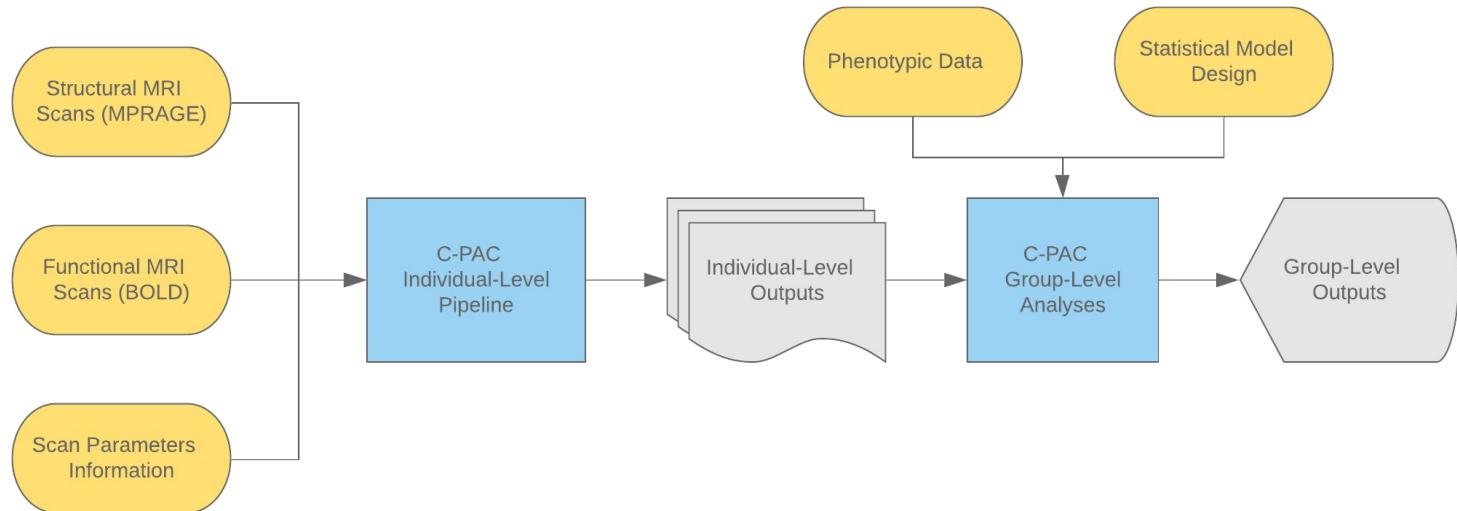
- Anatomical preprocessing: brain extraction, tissue segmentation, registration
- Functional preprocessing: slice timing correction, motion correction, coregistration, functional image to template warp, spatial smoothing, statistical analysis



- What is C-PAC?
- Why is C-PAC?
- How does C-PAC work?
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C-PAC Overview

- **C-PAC (Configurable Pipeline for the Analysis of Connectomes)**
 - An open-source **functional connectomics package**
 - Designed for ease-of-use (no programming required), but also deep configurability
 - The C-PAC pipeline employs many common neuroimaging tools (AFNI, FSL, ITK/ANTs, C3d etc.) and Python-based neuroimaging packages (PyBASC, PyPEER etc.)



World-wide Usage



Academic Usage

Contents lists available at ScienceDirect

 NeuroImage
journal homepage: www.elsevier.com/locate/neuroimage 

Deriving reproducible biomarkers from multi-site resting-state data: An Autism-based example

Alexandre Abraham^{a,b,*}, Michael P. Milham^{a,f}, Adriana Di Martino^e, R. Cameron Craddock^{a,f}, Dimitris Samaras^{a,d}, Bertrand Thirion^{a,b}, Gael Varoquaux^{a,b}

^a Parital Team, Saclay-INRIA In de France, Saclay, France
^b CEA, NeuroSpin bld 145, 91191 Gif-Sur-Yvette, France
^c Stony Brook University, NY 11794, USA
^d Ecole Centrale, 92290 Châtenay-Malabry, France
^e Center for the Developing Brain Child Mind Institute, New York, USA
^f Center for Biomedical Imaging and Neuromodulation, Nathan S. Kline Institute for Psychiatric Research, Orangeburg, NY, USA
^{* The Child Study Center at NYU Langone Medical Center, New York, NY, USA}

Contents lists available at ScienceDirect

 NeuroImage
journal homepage: www.elsevier.com/locate/neuroimage 

CrossMark 

Contents lists available at ScienceDirect

 Medical Image Analysis
journal homepage: www.elsevier.com/locate/media 

Disease prediction using graph convolutional networks: Application to Autism Spectrum Disorder and Alzheimer's disease

Sarah Parisot^{b,1,*}, Sofia Ira Ktena^{a,1}, Enzo Ferrante^c, Matthew Lee^c, Ricardo Guerrero^{a,3}, Ben Glocker^a, Daniel Rueckert^a

^a Biomedical Image Analysis Group, Imperial College London, UK
^b Alimbra Solutions Ltd, London, UK
^c Research Institute for Signals, Systems and Computational Intelligence, sinc{I}, FICH-UNL/CONICET, Santa Fe, Argentina
¹ StoryStream Ltd, London, UK

Contents lists available at ScienceDirect

 NeuroImage
journal homepage: www.elsevier.com/locate/neuroimage 

Check for updates 

Contents lists available at ScienceDirect

 NeuroImage
journal homepage: www.elsevier.com/locate/neuroimage 

Benchmarking functional connectome-based predictive models for resting-state fMRI

Kamalaker Dadi^{a,b,*}, Mehdi Rahim^{a,b}, Alexandre Abraham^{a,b}, Darya Chyzhyk^{a,b,c}, Michael Milham^a, Bertrand Thirion^{a,b}, Gael Varoquaux^{a,b}, for the Alzheimer's Disease Neuroimaging Initiative¹

^a Parital Project team, INRIA Saclay In de France, France
^b CEA/NeuroSpin bld 145, 91191 Gif-Sur-Yvette, France
^c Center for the Developing Brain Child Mind Institute, Center for Biomedical Imaging and Neuromodulation, Nathan S. Kline Institute for Psychiatric Research, USA

Contents lists available at ScienceDirect

 NeuroImage
journal homepage: www.elsevier.com/locate/neuroimage 

CrossMark 

Contents lists available at ScienceDirect

 Journal of Affective Disorders
journal homepage: www.elsevier.com/locate/jad 

Optimising network modelling methods for fMRI

Usama Pervaiz^{b,*}, Diego Vidaurre^{b,c}, Mark W. Woolrich^b, Stephen M. Smith^a

^a Oxford Centre for Functional MRI of the Brain (FMRIB), Wellcome Centre for Integrative Neuroimaging, Nuffield Department of Clinical Neurosciences, University of Oxford, Oxford, OX3 9QH, United Kingdom
^b Oxford Centre for Human Brain Activity (OHBA), Wellcome Centre for Integrative Neuroimaging, Department of Psychiatry, University of Oxford, OX3 7JX, United Kingdom
^c Department of Clinical Medicine, Aarhus University, Denmark

Contents lists available at ScienceDirect

 Journal of Affective Disorders
journal homepage: www.elsevier.com/locate/jad 

Check for updates 

The real-time fMRI neurofeedback based stratification of Default Network Regulation Neuroimaging data repository

Amalia R. McDonald^a, Jordan Muraskin^a, Nicholas T. Van Dam^b, Caroline Froehlich^a, Benjamin Puccio^a, John Pelman^b, Clemens C.C. Bauer^c, Alexis Akeyson^d, Melissa M. Breland^a, Vince D. Calhoun^{a,e}, Steven Carter^e, Tiffany P. Chang^e, Chelsea Gessner^e, Alyssa Giannone^e, Steven Giavasis^e, Jamie Glass^e, Steven Homann^e, Margaret King^e, Melissa Kramer^e, Drew Landis^e, Alexis Lieval^e, Jonathan Lisinski^e, Anna Mackay-Brandt^e, Brittny Miller^e, Laura Panek^e, Hayley Reed^e, Christine Santiago^e, Eszter Schoell^e, Richard Sinnig^e, Melissa Sital^e, Elise Taverna^e, Russell Tobe^e, Kristin Trautman^e, Betty Vargheese^e, Lauren Walden^e, Runtang Wang^e, Abigail B. Waters^e, Dylan C. Wood^d, F.Xavier Castellanos^{a,h}, Bennett Leventhal^a, Stanley J. Colcombe^a, Stephen LaConte^{f,i,k}, Michael P. Milham^{a,b}, R. Cameron Craddock^{a,b,*}

^a Nathan S. Kline Institute for Psychiatric Research, Orangeburg, NY, USA
^b Child Mind Institute, New York, NY, USA
^c Department of Brain and Cognitive Sciences, Massachusetts Institute of Technology, Boston, MA, USA
^d The Mind Research Network, Albuquerque, New Mexico, USA
^e Department of Electrical and Computer Engineering, The University of New Mexico, Albuquerque, New Mexico, USA
^f Virginia Tech Carilion Research Institute, Roanoke, VA, USA
^g Children's Hospital of Philadelphia, Philadelphia, PA, USA
^h The Child Study Center, NYU Langone Medical Center, New York, NY, USA
ⁱ Department of Psychiatry, University of California - San Francisco, San Francisco, CA, USA
^j School of Biomedical Engineering and Sciences, Virginia Polytechnic Institute and State University, Blacksburg, VA, USA
^k Departments of Emergency Medicine and Emergency Radiology, Virginia Tech Carilion School of Medicine, Roanoke, VA, USA

Contents lists available at ScienceDirect

 NeuroImage
journal homepage: www.elsevier.com/locate/neuroimage 

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ORIGINAL ARTICLE

Evaluating fMRI-Based Estimation of Eye Gaze During Naturalistic Viewing

Jake Son^{1,2}, Lei Ai¹, Ryan Lim³, Ting Xu¹, Stanley Colcombe³, Alexandre Rosa Franco^{1,3}, Jessica Cloud³, Stephen LaConte⁴, Jonathan Lisinski⁴, Arno Klein^{1,2}, R. Cameron Craddock^{1,3,5} and Michael Milham^{1,3,*}

¹Center for the Developing Brain, Child Mind Institute, New York, NY 10022, USA, ²MATTER Lab, Child Mind Institute, New York, NY 10022, USA, ³Center for Biomedical Imaging and Neuromodulation, Nathan S. Kline Institute for Psychiatric Research, New York, NY 10962, USA, ⁴Fralin Biomedical Research Institute, Virginia Tech Carilion Research Institute, Blacksburg, VA 24016, USA and ⁵Department of Diagnostic Medicine, Dell Medical School, Austin, TX 78701, USA

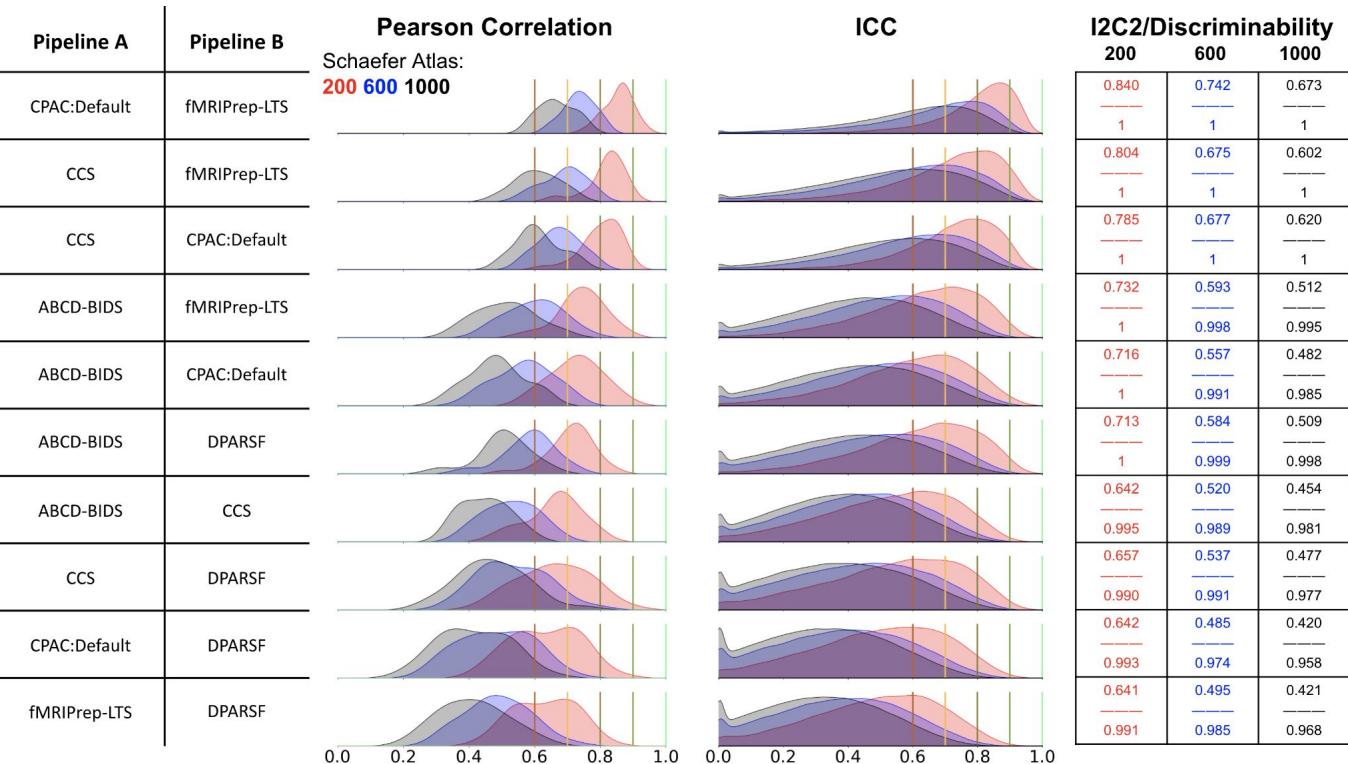
Contents lists available at ScienceDirect

 NeuroImage
journal homepage: www.elsevier.com/locate/neuroimage 

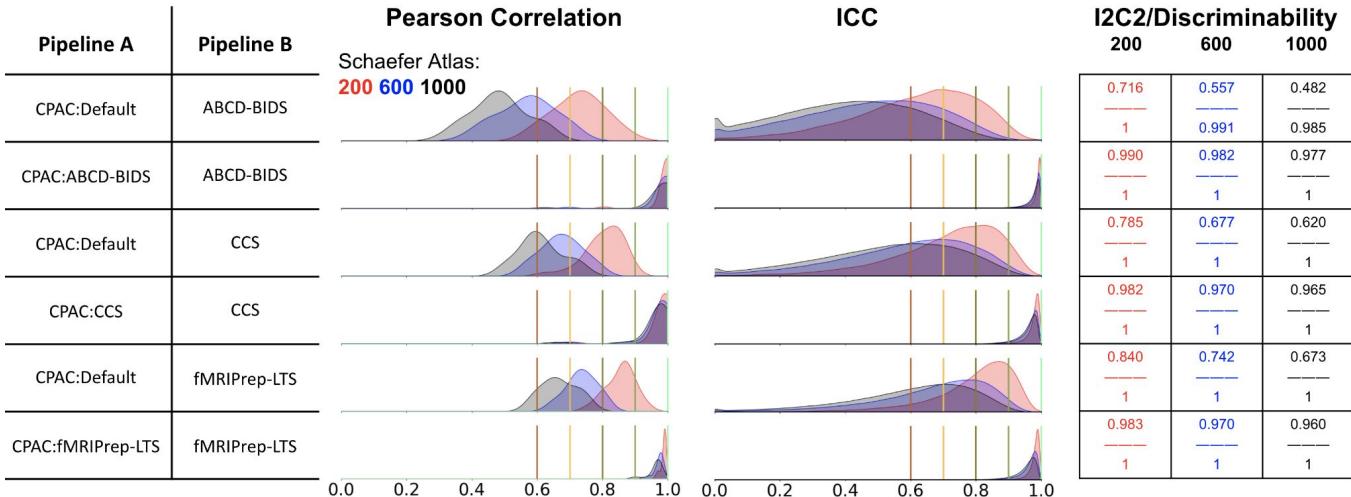
Check for updates 

- What is C-PAC?
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- Demo

Minimal Preprocessing



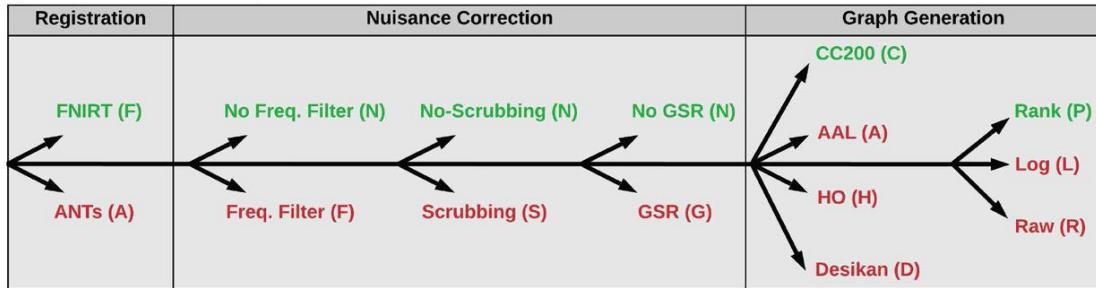
Harmonized Minimal Preprocessing



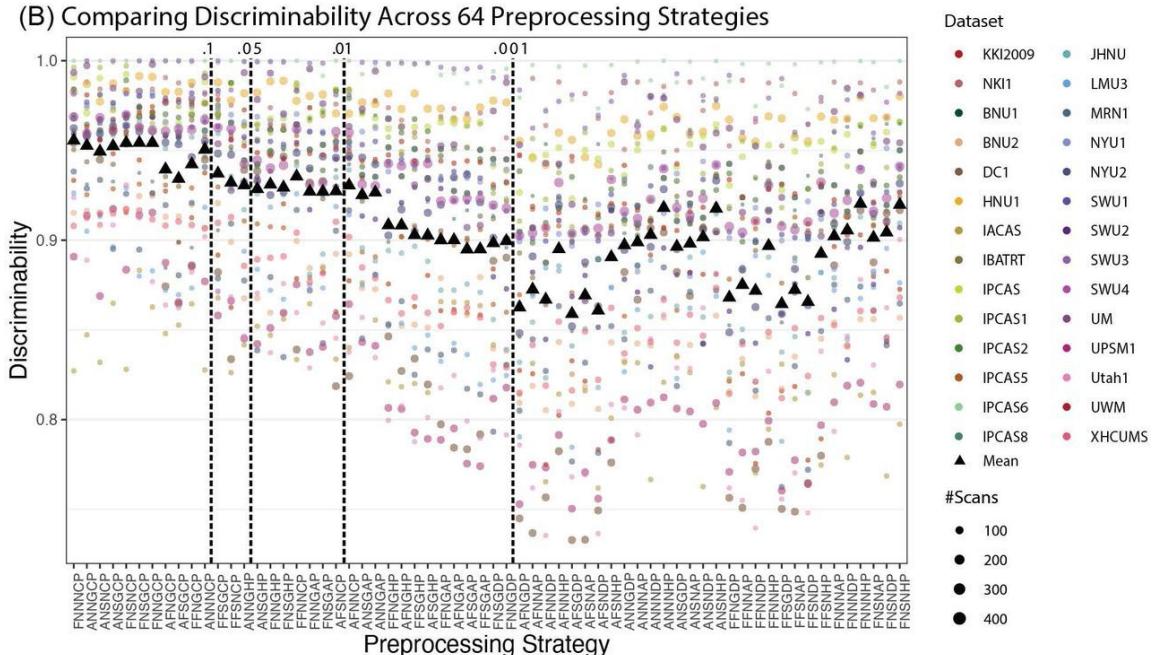
Configurable

Fully customize your pipeline

(A) Processing Strategies Evaluated



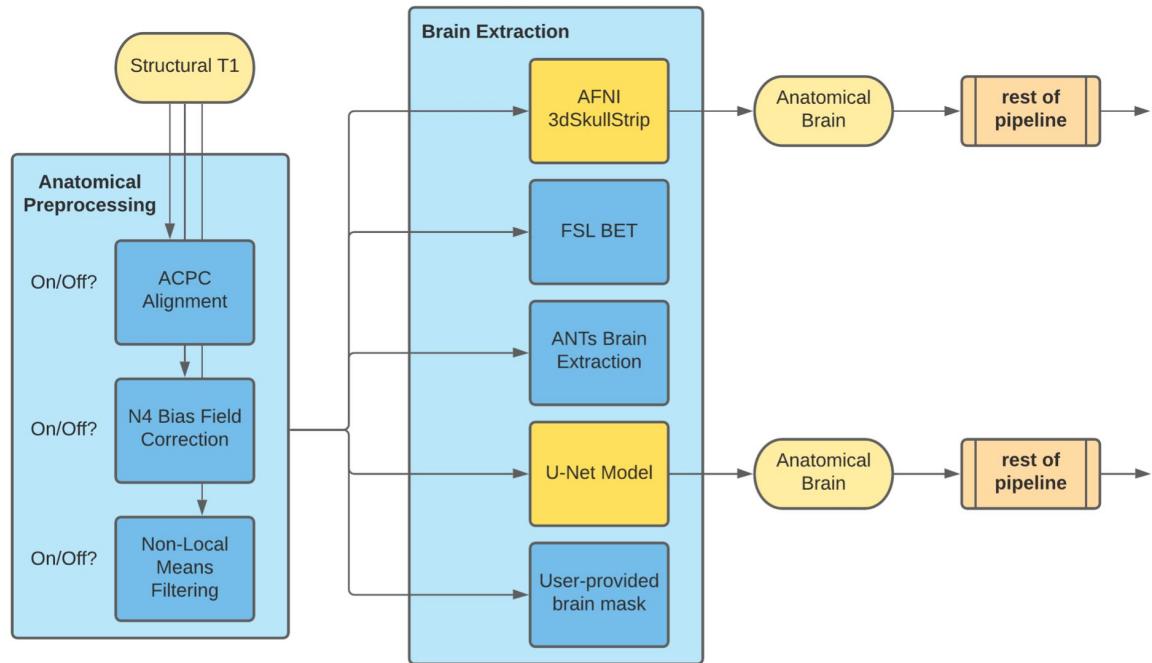
(B) Comparing Discriminability Across 64 Preprocessing Strategies



Forking - On, Off, and On/Off

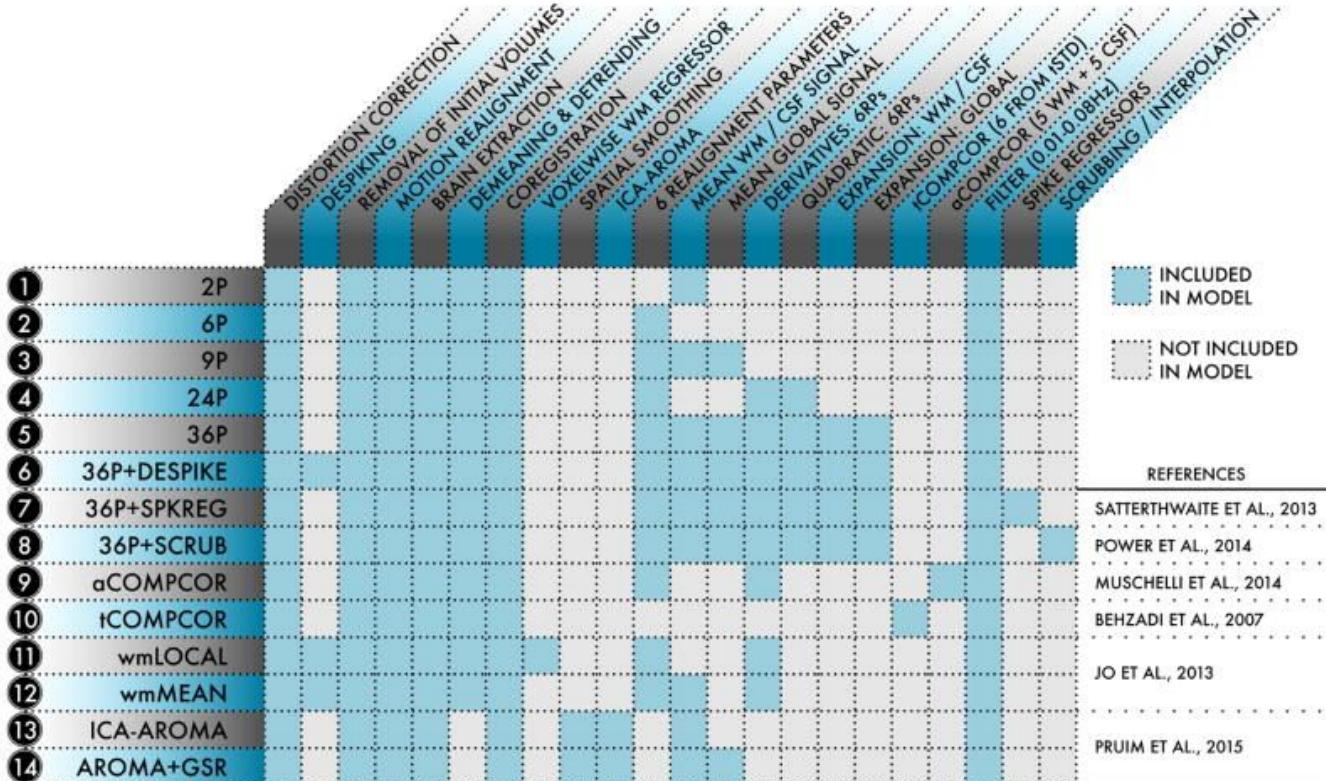
Forkable

Provide fork options,
parallelize forks with
multiple CPUs



Forkable

Provide fork options,
parallelize forks with
multiple CPUs



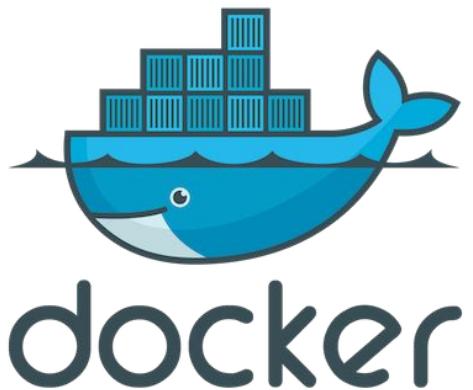
Easy-of-use

- GUI
- Text-based YAML file can be easily share with others.

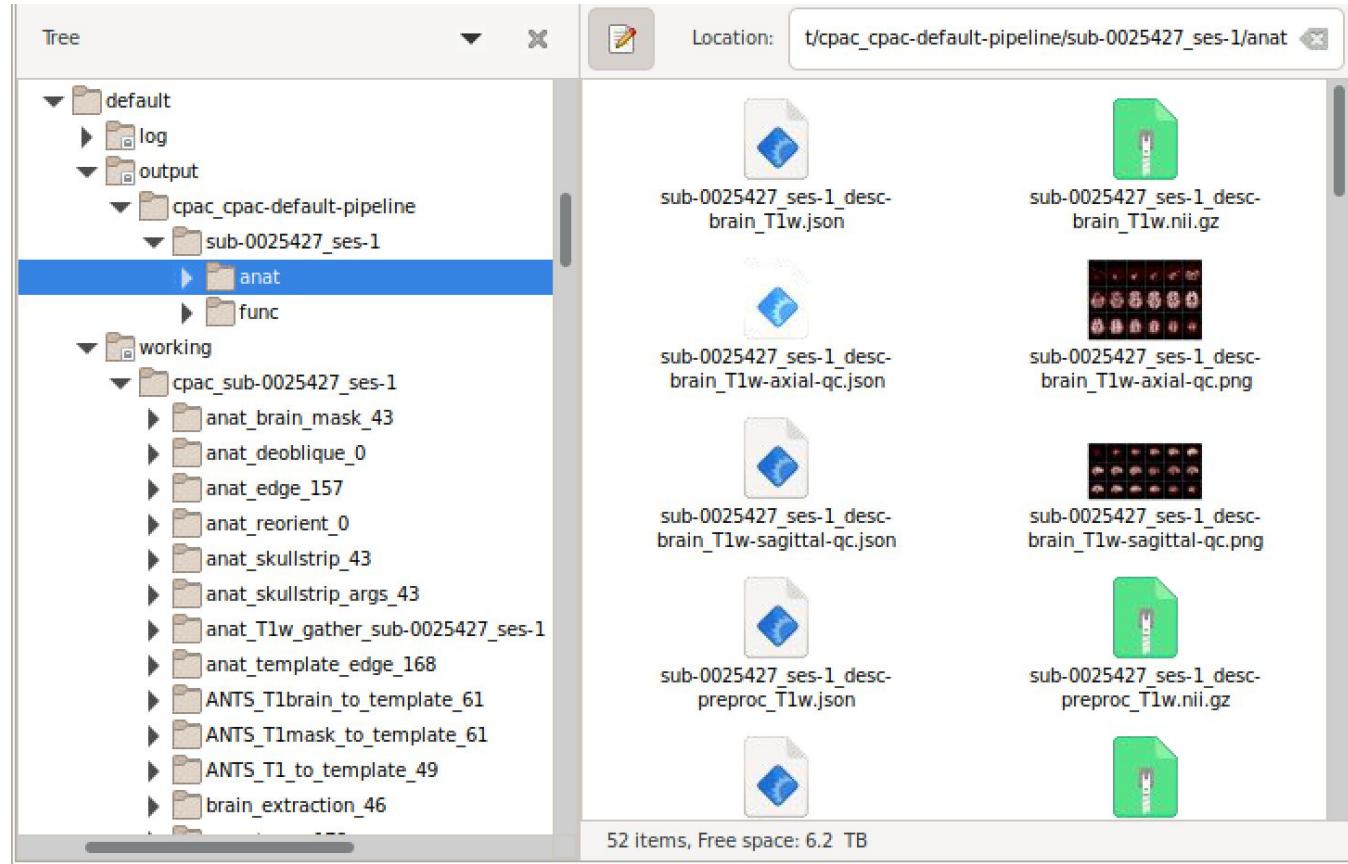
The screenshot shows the C-PAC web application interface. At the top, there's a header with the Child Mind Institute logo, the text "big data analytics", and the "C-PAC" logo. Below the header, a navigation bar has a "HOME" icon and link. The main area is divided into two sections: "Pipelines" on the left and "About C-PAC" on the right. The "Pipelines" section lists three items: "Default C-PAC 1.8.0", "Anatomical Preproc", and "Functional Preproc". Each item has a small icon next to it. To the right of the pipelines, the "About C-PAC" section contains a brief description of what C-PAC is and how it builds upon other software packages. Below these sections, a large code editor window displays a YAML configuration file for anatomical preprocessing:

```
anatomical_preproc:  
  run: On  
  brain_extraction:  
    run: On  
    using:  
      - 3dSkullStrip  
      - BET
```

Standardized
Environment



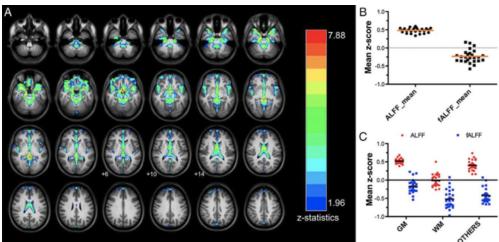
Warm Restart



End-to-end Preprocessing

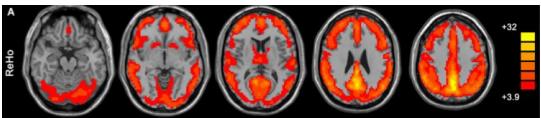
Amplitude of Low Frequency Fluctuations (ALFF) and fractional ALFF (f/ALFF)

Zang et al., 2007. Cited by 1982
Zou et al., 2008. Cited by 1242



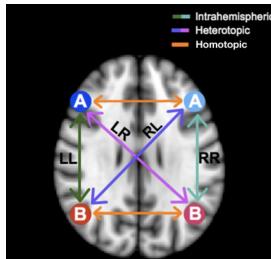
Regional Homogeneity (ReHo)

Zang et al., 2004. Cited by 1841



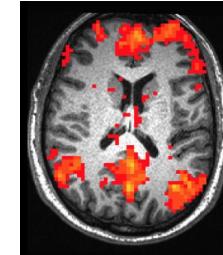
Voxel-Mirrored Homotopic Connectivity (VMHC)

Stark et al., 2008. Cited by 241
Zuo et al., 2010. Cited by 549



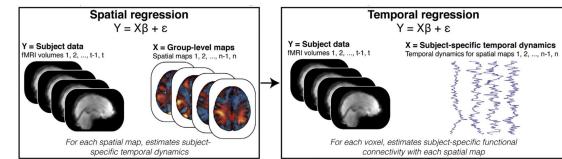
Seed-based Correlation Analysis (SCA)

Biswal et al., 1995. Cited by 8935



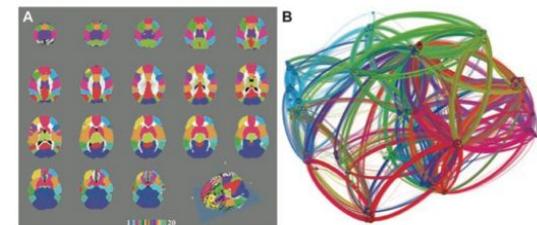
Dual Regression

Beckmann et al., 2009. Cited by 651
Filippini et al., 2009. Cited by 1635



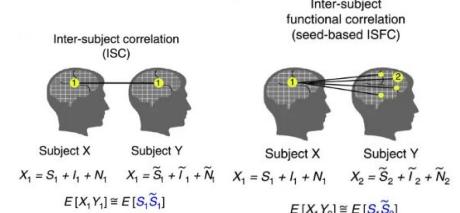
Network Centrality

Buckner et al., 2009. Cited by 2476
Bullmore and Sporns, 2009. Cited by 9619



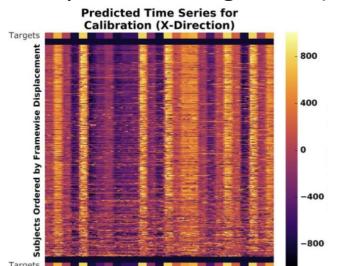
Innovative Analysis

Intersubject Correlation Analyses (ISC) and Inter-subject Functional Correlation (ISFC)



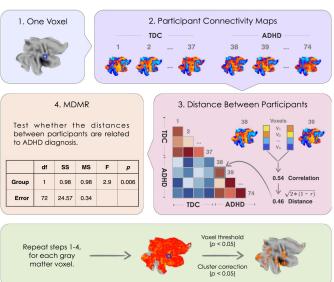
Simony et al., 2016.

Predictive Eye Estimate Regression (PEER)



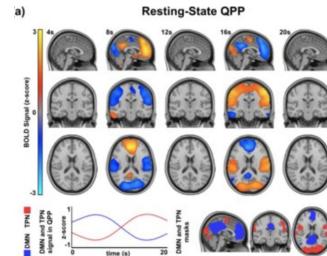
Son et al., 2020.

Multivariate Distance Matrix Regression (MDMR)



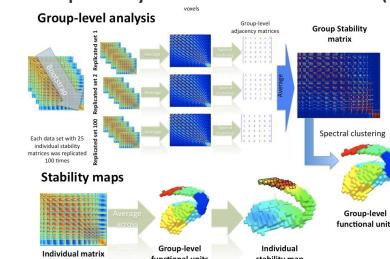
Shehzad et al., 2014

Quasi-Periodic Pattern Analysis (QPP)



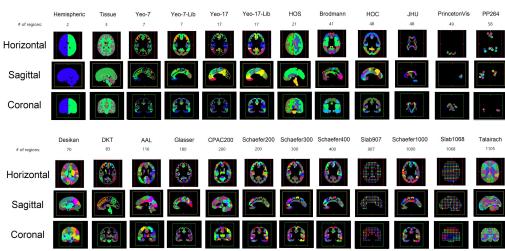
Abbas et al., 2019

Bootstrap Analysis of Stable Clusters (BASC)



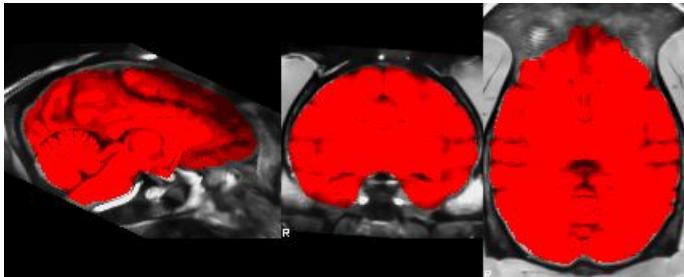
Bellec et al., 2010.

Automated Atlas Extraction (Neuroparc Library)



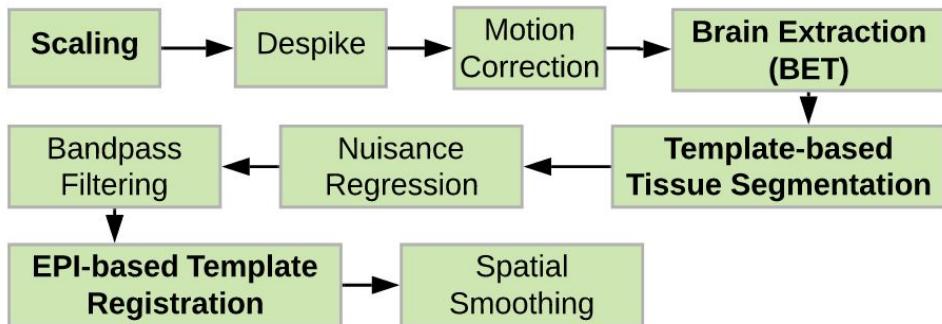
NeuroData, 2020

Non-human Preprocessing



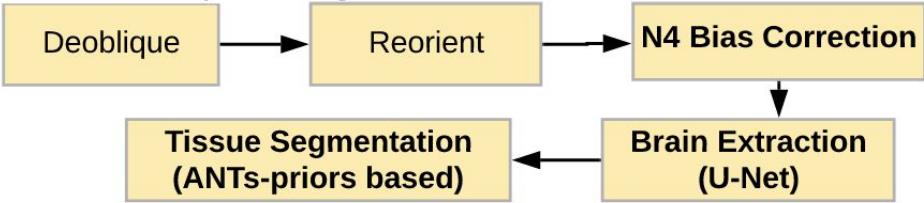
Wang et al., 2021

Rodent

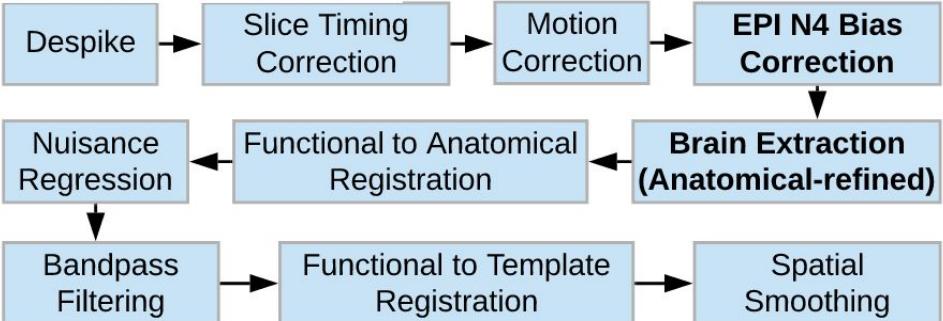


Macaque

Anatomical Preprocessing



Functional Preprocessing

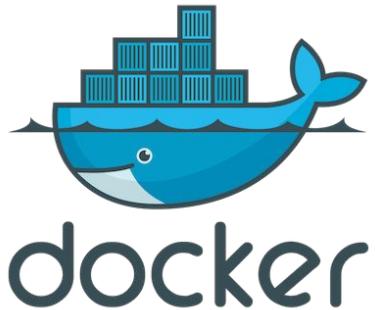


...then how do you run C-PAC?

Get Running Fast

Quickly pull and run C-PAC via Docker and Singularity containers.

- No need to install anything except Docker or Singularity.
- Self-contained replicated environments for reproducibility.
- *>> docker pull fcpindi/c-pac:latest*



Get Running Big

Spin up big analyses on the cloud using Amazon Web Services (AWS).

- No need to install anything - only needs an AWS account.
- Large variety of performance and space options available depending on budget.
- C-PAC AMI available publicly on the Amazon AMI Marketplace.



How to run C-PAC?

>> *docker run <dir mapping> <container image>*

```
docker pull fcpindi/c-pac:latest
```

```
docker run -i --rm \
    -v /Users/You/local_bids_data:/bids_dataset \
    -v /Users/You/some_folder:/outputs \
    -v /tmp:/scratch \
    fcpindi/c-pac:latest /bids_dataset /outputs participant
```



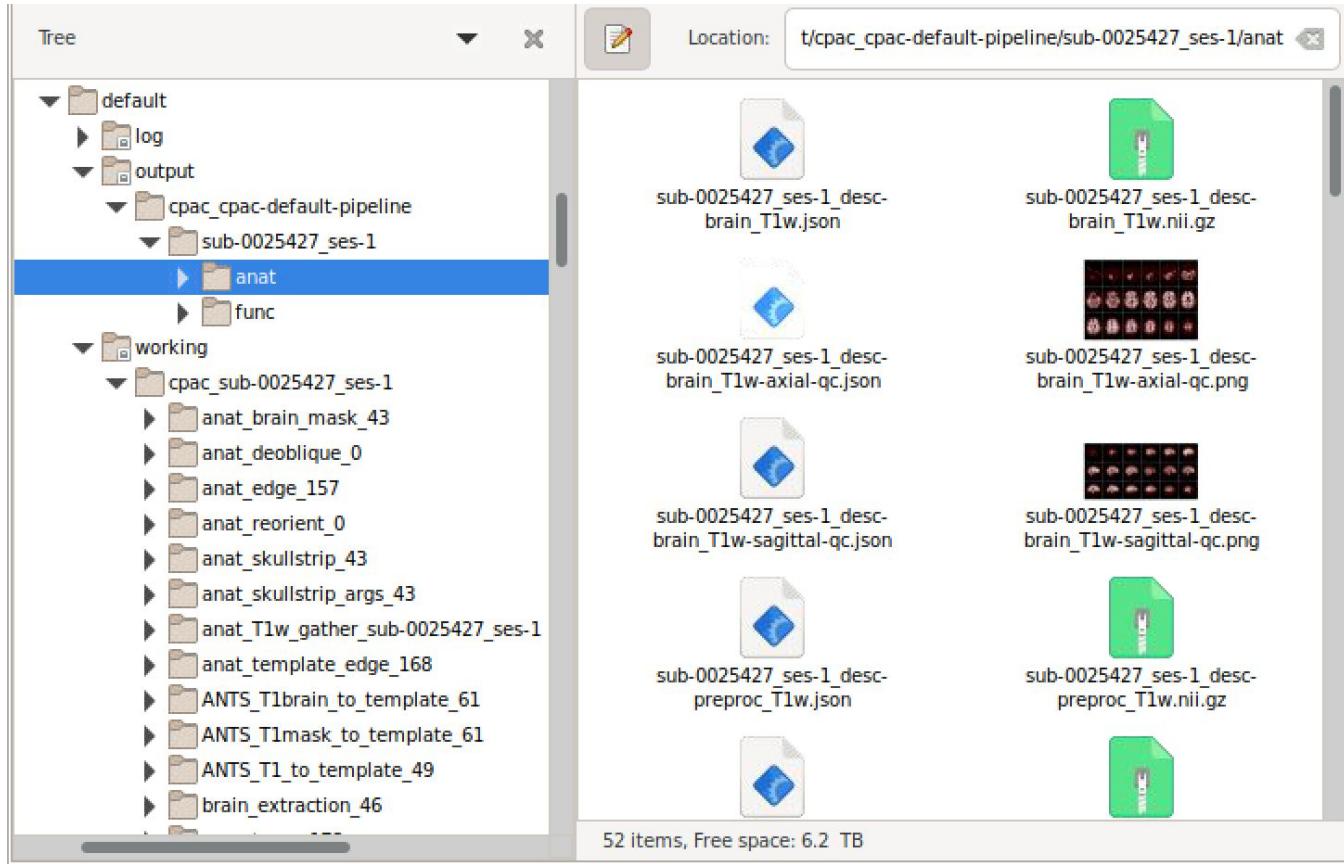
>> *singularity run <dir mapping> <container image>*

```
singularity pull shub://FCP-INDI/C-PAC
```

```
singularity run \
    -B /Users/You/local_bids_data:/bids_dataset \
    -B /Users/You/some_folder:/outputs \
    -B /tmp:/scratch \
    FCP-INDI-C-PAC-master-latest.simg /bids_dataset /outputs participant
```

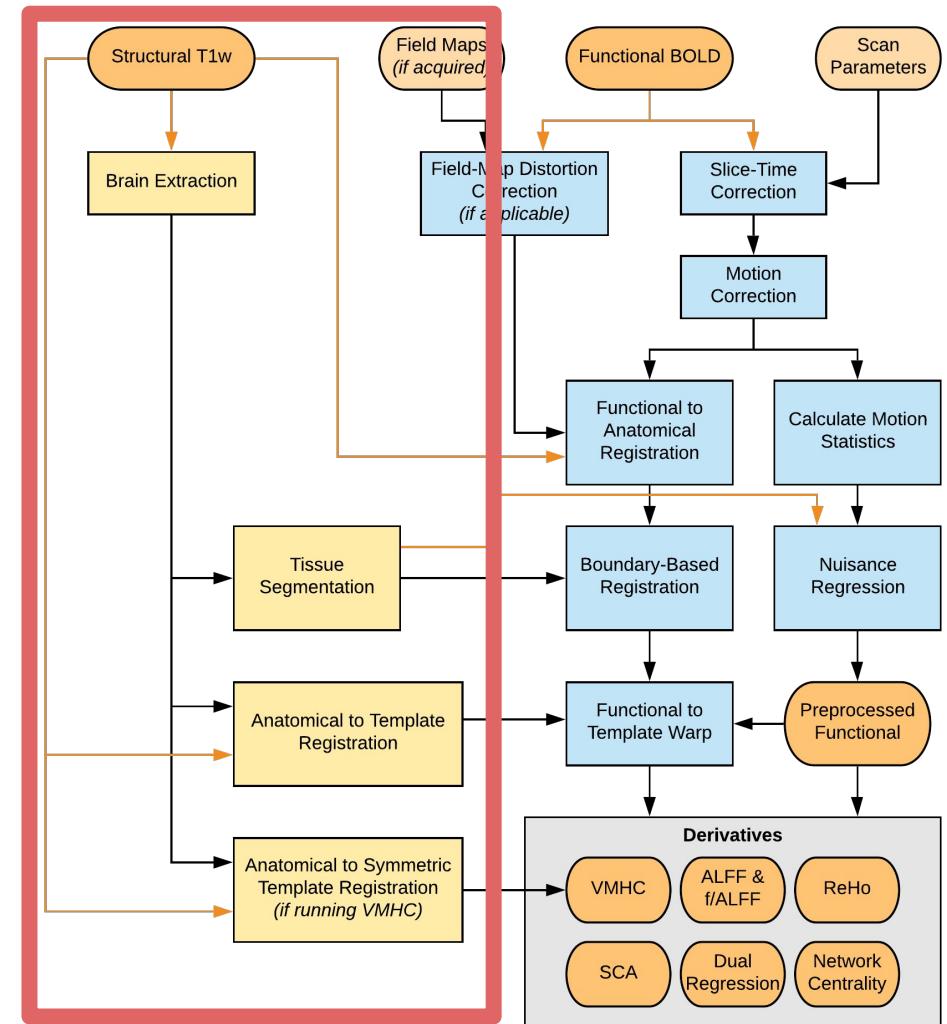


Get Output (BIDS derivatives)



- What is C-PAC?
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Anatomical Preprocessing



Anatomical Preprocessing: Brain Extraction

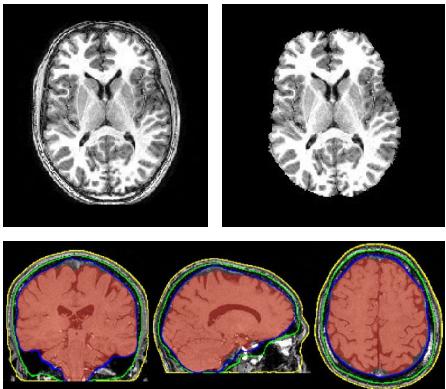
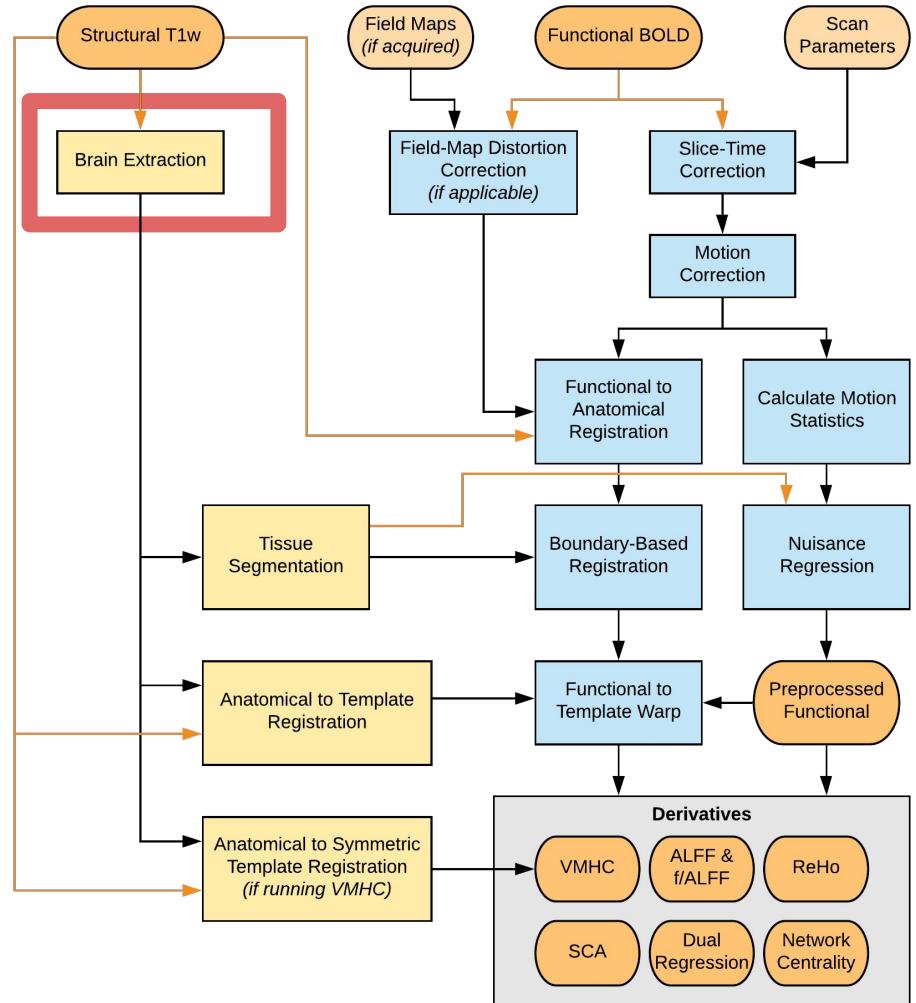
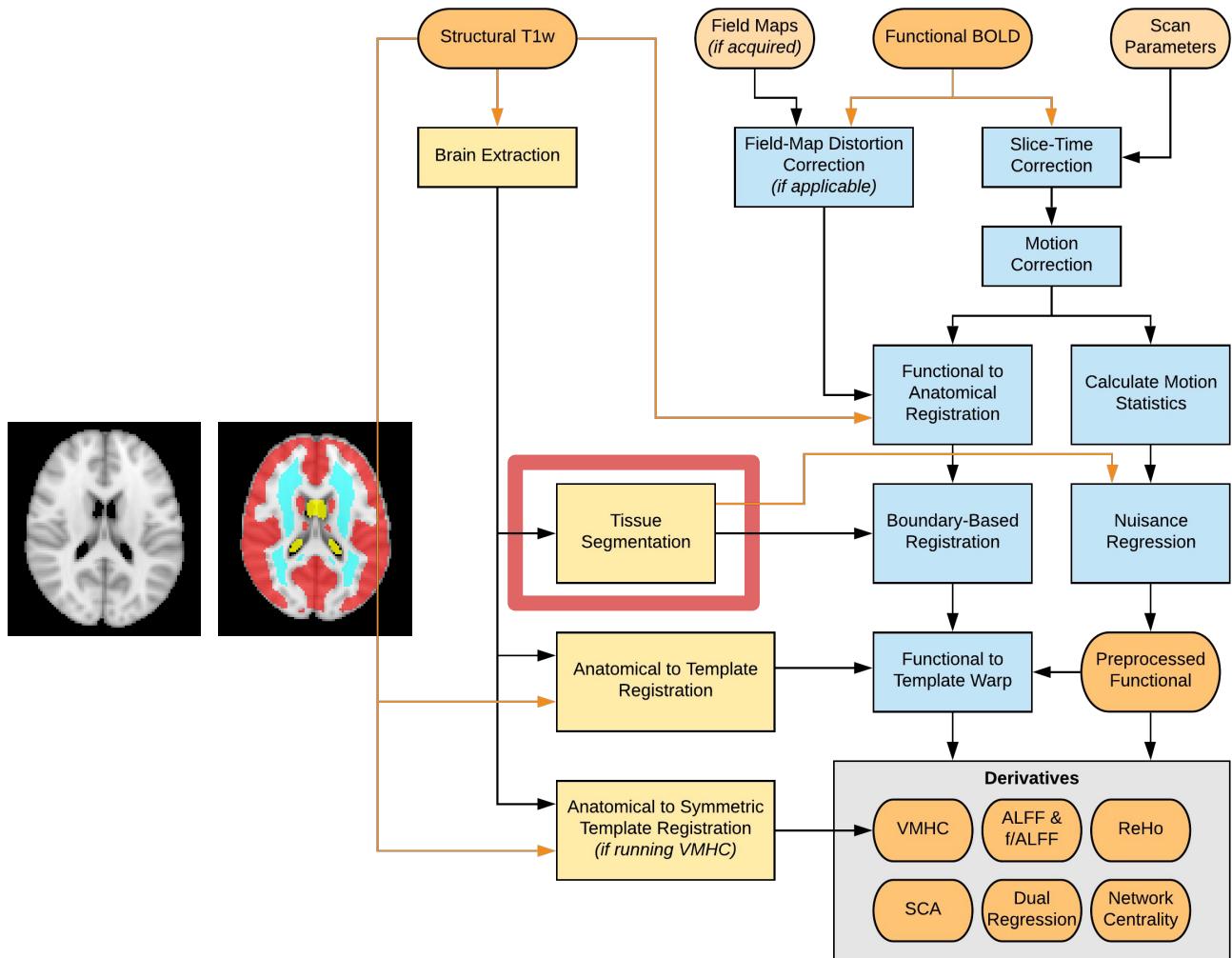
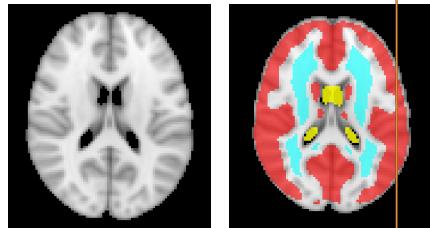


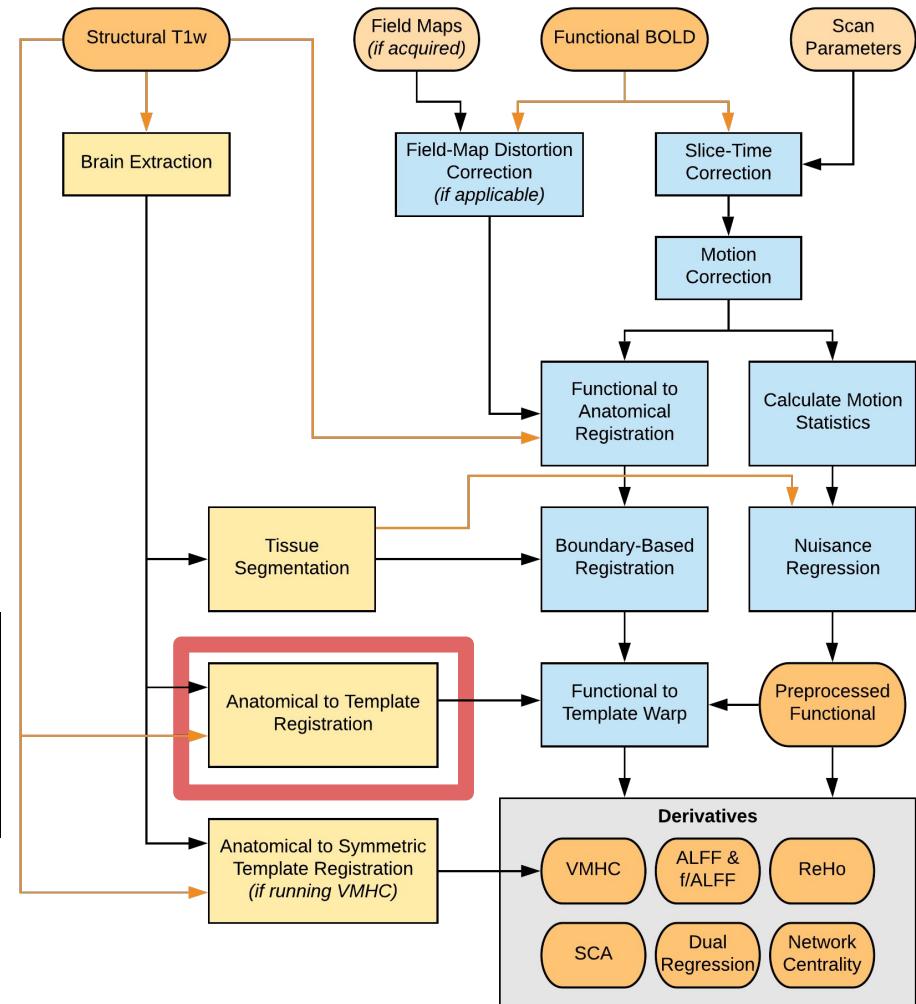
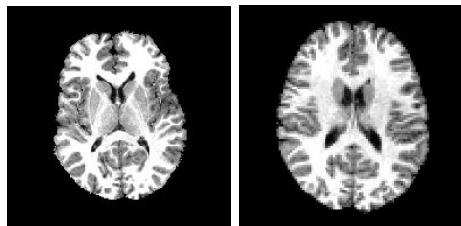
Image from <https://fsl.fmrib.ox.ac.uk/fsl/fslwiki/BET>



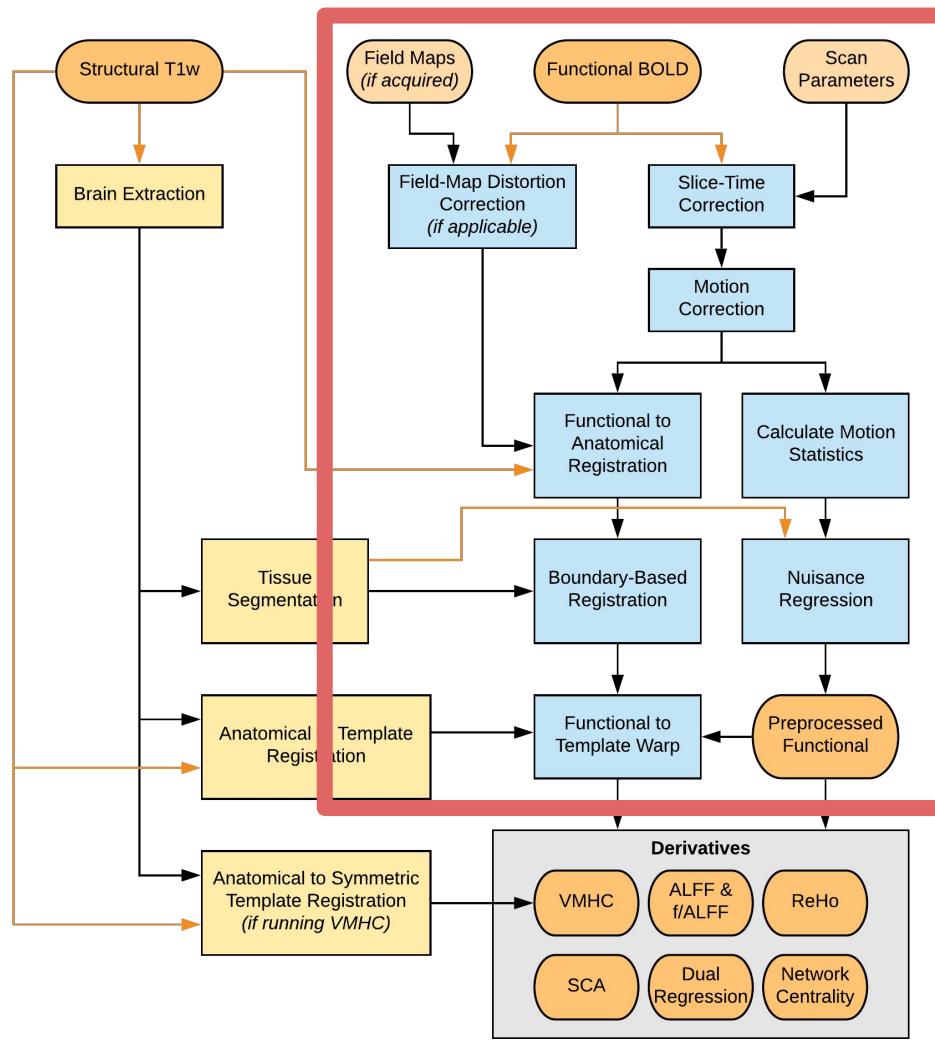
Anatomical Preprocessing: Tissue Segmentation



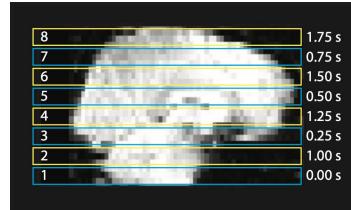
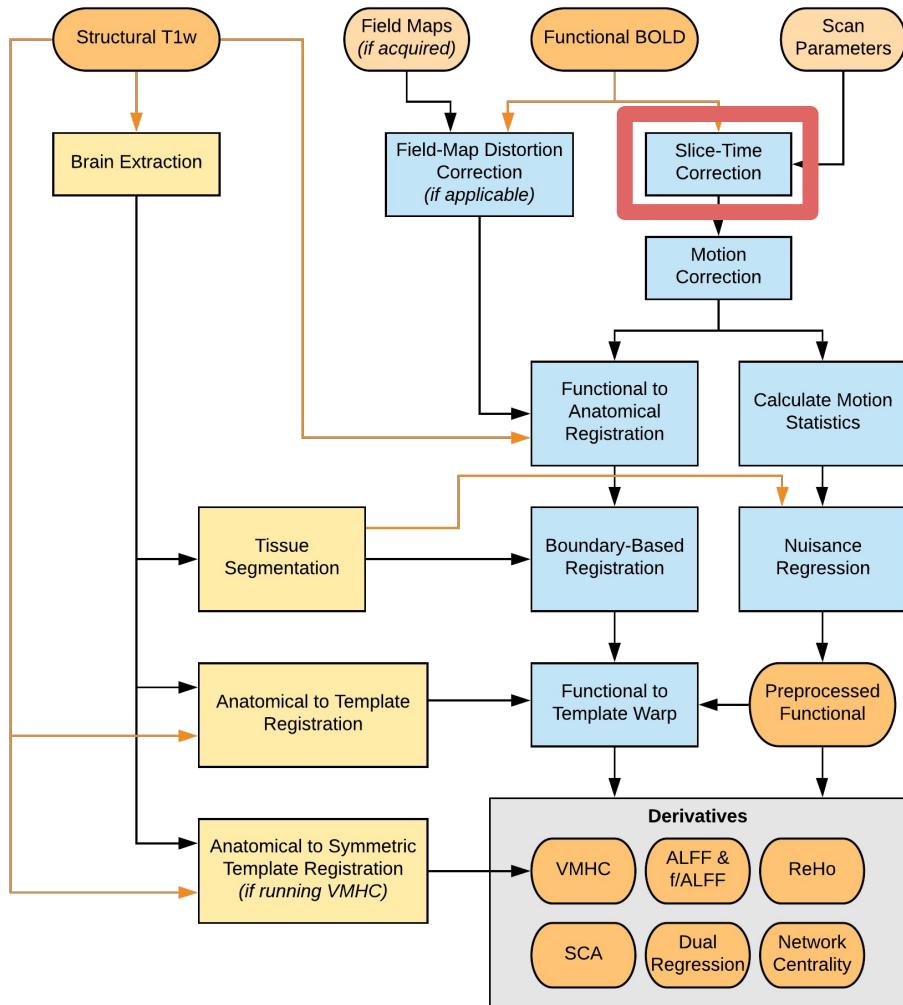
Anatomical Preprocessing: Anatomical Registration



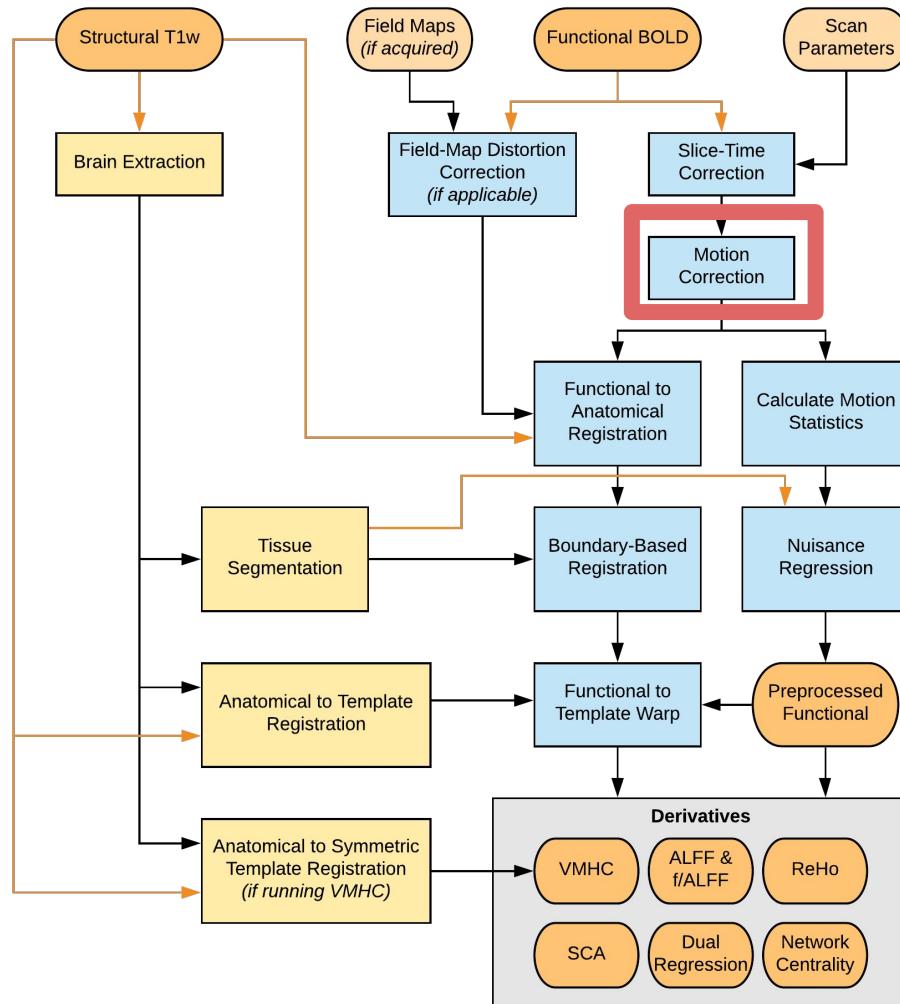
Functional Preprocessing



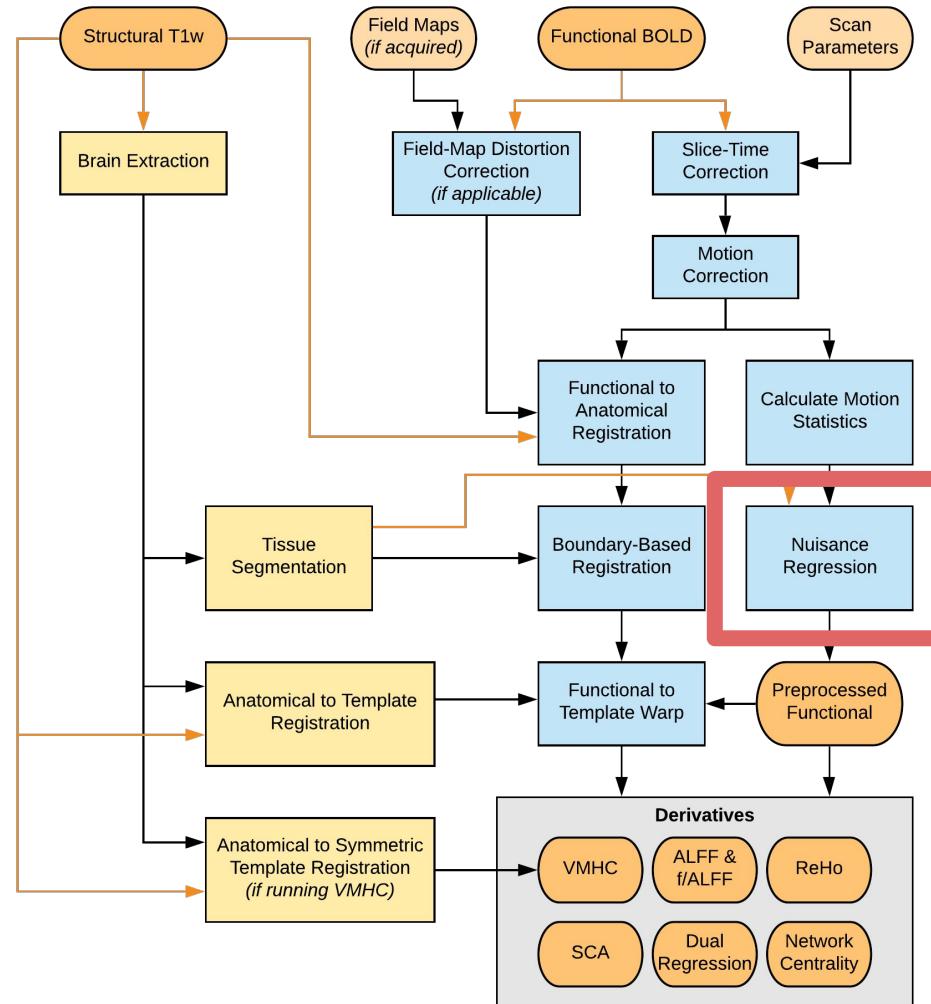
Functional Preprocessing: Slice-Time Correction



Functional Preprocessing: Motion Correction



Functional Preprocessing: Nuisance Regression

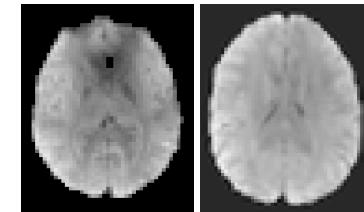
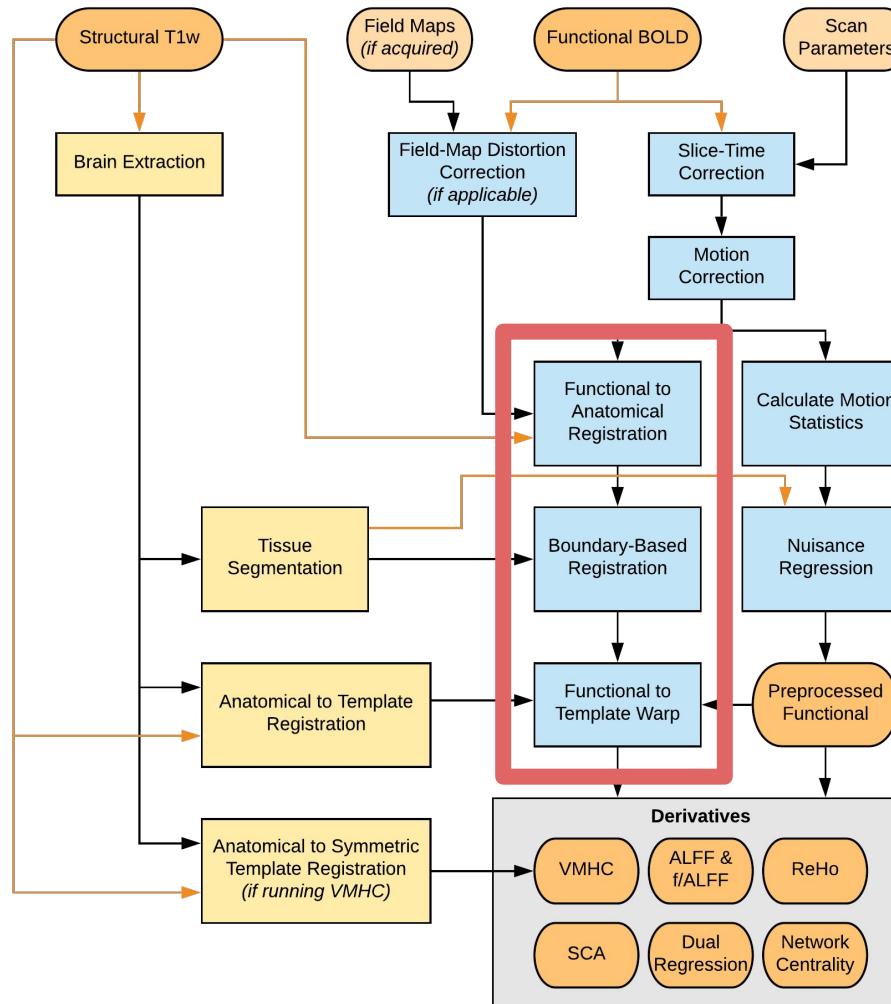


Functional Preprocessing: Nuisance Regression

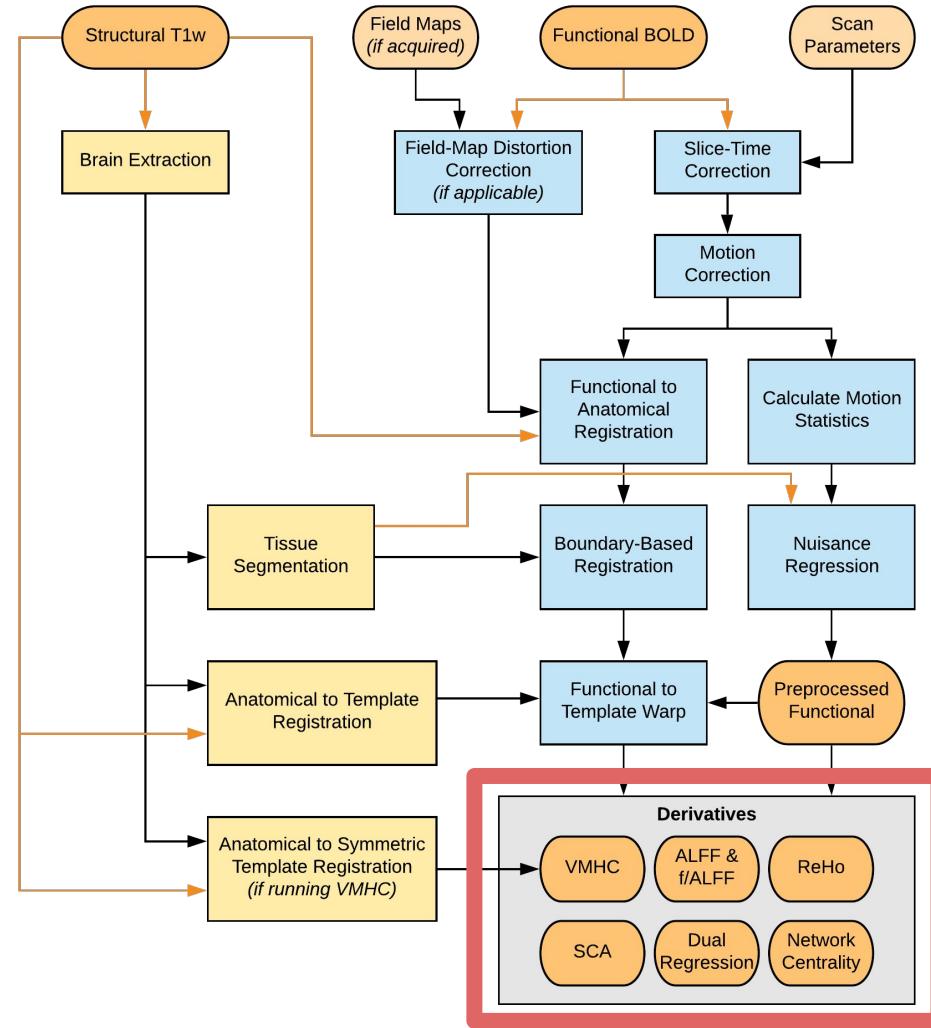
- Motion
 - 6-parameter model
 - 24-parameter model Friston et al., 1996. Cited by 2789
- Global Signal Regression (GSR)
 - Fox et al., 2005. Cited by 7779
- WM and CSF regression
- aCompCor and tCompCor
 - Noise ROI component-based noise correction
 - Behzadi et al., 2007. Cited by 2633
- Polynomial Detrending
 - Removes linear or quadratic trend
- Temporal Filtering
 - Bandpass low-frequency
- Censor
 - Threshold FD (Framewise Displacement) and DVARS
 - Power et al., 2012. Cited by 5382
- ICA-AROMA Denoising
 - ICA-based Automatic Removal of Motion Artifacts
 - Pruim et al., 2015. Cited by 840

Regressor Name
My Regressor
<input type="checkbox"/> Motion
<input type="checkbox"/> GlobalSignal
<input type="checkbox"/> Gray Matter
<input type="checkbox"/> White Matter
<input type="checkbox"/> Cerebrospinal Fluid
<input type="checkbox"/> aCompCor
<input type="checkbox"/> tCompCor
<input type="checkbox"/> PolyOrt
<input type="checkbox"/> BandPass
<input type="checkbox"/> Censor

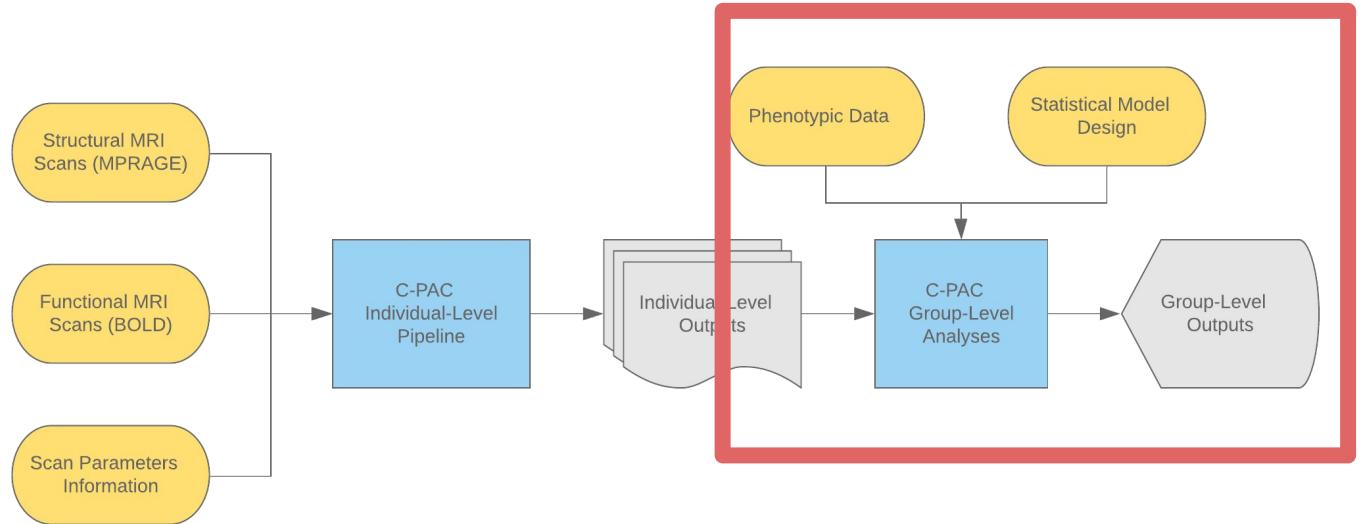
Functional Preprocessing: Registration



Functional Preprocessing: Derivatives



Group-Level Analyses



- FSL FEAT
- Non-Parametric Permutation Inference (FSL Randomise)

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    -v /Users/You/local_bids_data:/bids_dataset \
    -v /Users/You/some_folder:/outputs \
    -v /tmp:/scratch \
    fcpindi/c-pac:latest /bids_dataset /outputs participant
```



>> *singularity run <dir mapping> <container image>*

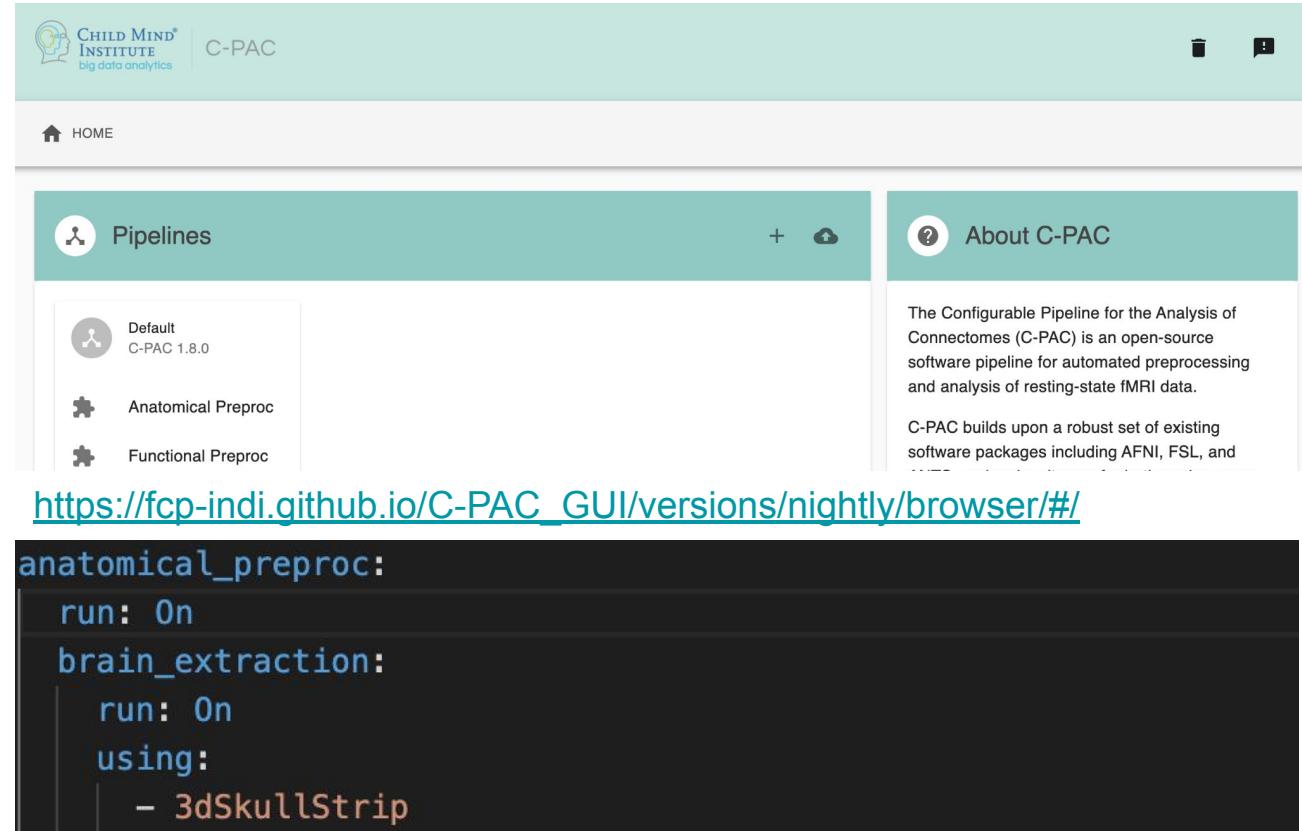
```
singularity pull shub://FCP-INDI/C-PAC
```

```
singularity run \
    -B /Users/You/local_bids_data:/bids_dataset \
    -B /Users/You/some_folder:/outputs \
    -B /tmp:/scratch \
    FCP-INDI-C-PAC-master-latest.simg /bids_dataset /outputs participant
```



Pipeline Config

- GUI Editor
- Text-based YAML file Editor



The screenshot shows the C-PAC GUI interface. At the top, there's a header with the Child Mind Institute logo and the text "C-PAC". Below the header, a navigation bar has a "HOME" icon and the word "HOME". The main content area is titled "Pipelines". It lists three pipeline configurations: "Default C-PAC 1.8.0", "Anatomical Preproc", and "Functional Preproc". To the right of the pipelines is a sidebar titled "About C-PAC" which contains a brief description of the software and its dependencies.

https://fcp-indi.github.io/C-PAC_GUI/versions/nightly/browser/#/

```
anatomical_preproc:
  run: On
  brain_extraction:
    run: On
    using:
      - 3dSkullStrip
```

```
docker run -i --rm \
-v /Users/You/local_bids_data:/bids_dataset \
-v /Users/You/some_folder:/outputs \
-v /tmp:/scratch \
-v /Users/You/Documents:/configs \
-v /Users/You/resources:/resources \
fcpindi/c-pac:latest /bids_dataset /outputs participant --pipeline_file /configs/pipeline_config.yml
```

Data Config



Gorgolewski, Krzysztof J., et al. "The brain imaging data structure, a format for organizing and describing outputs of neuroimaging experiments." *Scientific data* 3.1 (2016): 1-9.

```
my_dataset/
participants.tsv
sub-01/
anat/
    sub-01_T1w.nii.gz
func/
    sub-01_task-rest_bold.nii.gz
    sub-01_task-rest_bold.json
dwi/
    sub-01_dwi.nii.gz
    sub-01_dwi.json
    sub-01_dwi.bval
    sub-01_dwi.bvec
sub-02/
sub-03/
```

Data Config



- Data Structure



anat.nii.gz



func.nii.gz

- Non-BIDS
Data Config

```
- anat: /non_bids_dataset/sub-0025427/anat.nii.gz
  func:
    rest_run-1:
      | scan: /non_bids_dataset/sub-0025427/func.nii.gz
    site_id: site-none
    subject_id: sub-0025427
    unique_id: ses-1
```

```
docker run -i --rm \
  -v /Users/You/any_directory:/bids_dataset \
  -v /Users/You/some_folder:/outputs \
  -v /tmp:/scratch \
  -v /Users/You/Documents:/configs \
  fcpind/c-pac:latest /bids_dataset /outputs participant --data_config_file /configs/data_config.yml
```

Optional arguments in C-PAC container

--pipeline_file	--mem_mb /--mem_gb
--data_config_file	--save_working_dir
--n_cpus	--anat_only
	--participant_label

>> docker run fcpindi/c-pac:latest --help

```
docker run -i --rm \
-v /Users/You/local_bids_data:/bids_dataset \
-v /Users/You/some_folder:/outputs \
-v /tmp:/scratch \
-v /Users/You/Documents:/configs \
-v /Users/You/resources:/resources \
fcpindi/c-pac:latest /bids_dataset /outputs participant --pipeline_file /configs/pipeline_config.yml
```

- What is C-PAC?
- Why is C-PAC?
- How does C-PAC work?
- How to run C-PAC in a container?
- Demo

Demo Time!

<https://github.com/XinhuiLi/OpenTutorials>

For More Details

<https://fcp-indi.github.io>

C-PAC forum:

https://groups.google.com/forum/?utm_medium=email&utm_source=footer#!forum/cpax_forum

C-PAC GitHub issues:

<https://github.com/FCP-INDI/C-PAC/issues>