

## 0. Setup

### Initialization:

- FieldTrip top-level folder for ft\_defaults.m
- MEGneto top level for megne2setup.m
- SPM T1 template visible

- project\_path
- analysis\_name
- rawdata\_path
- mri\_path
- overwrite

MATLAB input

- Add proper paths for FieldTrip folder
- Create directories for project
- Initialize JSON config file

megne2setup.m

paths (struct)

out

- ft\_defaults
- path\_generation
- save\_to\_json
- path\_check

## 1,2. Processing

fcp\_1\_TaskEpoching.m

### 1. Setup

- Read in config settings
- Load and match PIDs from MEG and MRI folders
- Create list of subjects with full sets of data

### 2. Plot triggers

- Generate and save plot of event triggers

### 3. Epoch continuous data into trials

- Set up trial definitions for epoching
- Epoch events list to define time windows corresponding to each trial and overall number of trials; save to config

### 4. Head motion

- Identify trials with excessive head motion
- Reject those trials, save filtered version

### 5. Artifacts

- Check for muscle, jump artifacts
- Reject those trials with artifacts, save filtered version

### 6. Bad channels

- Identify and save list of bad channels
- Throw warning if there are many

### Outputs

- Plots:
  - Triggers for each event type
  - Head motion visualization
- Epochs (into JSON):
  - All trials
  - Filtered for head motion (HM)
  - Filtered for HM, muscle, jump artifacts
- List of bad channels (saved to its own JSON)

- load\_config
- load\_participants
- ds\_pid\_match
- write\_match\_if\_not\_empty
- ft\_definetrial
- plotTriggers
- HeadMotionTool
- ft\_artifact\_muscle
- ft\_artifact\_jump
- ft\_rejectartifact
- save\_to\_json
- detectBadChannels

fcp\_2\_PreprocessingICA.m

### 1. Setup

- As in fcp\_1

### 2. Noise reduction

- Load gradiometer info, 3<sup>rd</sup> order gradients from CTF to account for noise

### 3. Bad channel removal

- Remove bad channels; replace signal with average of neighbours

### 4. ICA

- Remove bad channel signal altogether from consideration
- Run ICA
- Interactive decision on whether to keep or reject component
- Reject all selected components at once
- Save

### Outputs

- Final preprocessed data output
- Bad ICA components to JSON file

- load\_config
- load\_participants
- ds\_pid\_match
- write\_match\_if\_not\_empty
- save\_to\_json

## 3. Beamforming

fcp\_3\_beamforming\_sourcegrid.m

### 1. Setup

- Read in config settings
- Load and match PIDs from MEG and MRI folders
- Create list of subjects with full sets of data

### 2. Head model preparation

- Load and segment T1 template brain
- Construct head model and do necessary unit conversions
- Construct dipole grid in template brain
- Load desired atlas and create binary masks to define valid voxels within head model

### 3. Check alignment

- Load and segment participant MRIs, load preproc. MEG data
- Construct subject-specific head, source models
- Check alignment between subject and template head model
- Check alignment source model and head model
- Save images

### 4. Source reconstruction

- Compute lead field matrix
- Run source analysis, reduce to dominant orientation
- Interpolate functional data onto anatomical data
- Return these results

### Outputs

- Individual and template head model alignment, \*.png
- Source and head model alignment, \*.png

- Source analysis output
- Source descriptives output
- Source interpolation onto MRI data

out

## 4. Connectivity Analysis

fcp\_4\_connectivity.m

### 1. Setup

- As in fcp\_1

### 2. Analysis

- Compute connectivity w/ FT code

### Outputs

- Connectivity statistics