

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

- Run ft_defaults.m to add the right subfolders within FieldTrip.
- Create directories for project
- Initialize JSON config file

megne2setup.m

paths (struct)

out

- ft_defaults
- path_check
- path_generation
- save_to_json

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 fcp_1_output JSON):
 - All trials
 - Filtered for head motion (HM)
 - Filtered for HM, muscle, jump artifacts
- List of bad channels (saved to its own JSON)

- check_csv_has_empty
- detectBadChannels
- ds_pid_match
- ft_artifact_jump
- ft_artifact_muscle
- ft_definetrial
- ft_preproessing
- ft_rejectartifact
- ft_read_event
- ft_read_header
- HeadMotionTool
- load_config
- load_participants
- plotTriggers
- save_to_json
- write_match_if_not_empty

1,2,3. Processing

fcp_2_PreprocessingICA.m

1. Setup

- As in fcp_1
- 2. Noise reduction**
 - Load gradiometer info and compute 3rd order gradients from CTF to account for noise
- 3. ICA**
 - Remove bad channel signals altogether from analysis (they are dependent on neighbor sensor signals, thus redundant to include in ICA)
 - Run ICA

Outputs

- ICA components for each participant

fcp_2_5_checkpoint.m

- Human to identify ICA components that are artifacts or too noisy.
- Back-project to clean ICA

Outputs

- Bad ICA components to JSON file
- ICA-denoised data

- ds_pid_match
- ft_channelrepair
- ft_channelselection
- ft_componentanalysis
- ft_denoise_synthetic
- ft_prepare_neighbours
- ft_preprocessing
- ft_rejectcomponent
- ft_resampleddata
- ft_selectdata
- ft_topoplotIC
- inputBADchannels
- load_config
- load_participants
- saveSensorsToFile
- save_to_json
- write_match_if_not_empty

4. Beamforming

fcp_4_beamforming.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; note that participants may be removed after preprocessing due to insufficient number of trials leftover (e.g., too much noise)

2. Head model preparation

- Load and segment T1 template brain
- Construct head model and do necessary unit conversions
- Construct dipole grid in template brain
- 3. Check alignment**
 - Load and segment participant MRIs, load preprocessed 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
 - Load desired atlas and create binary masks to define valid voxels within head model
 - Interpolate functional data onto anatomical data
 - Further interpolation onto AAL-defined regions
 - Return these results

Outputs (for each participant)

- catmatrix = individual source timeseries in *.mat files: (timepoints in 1 trial) x (num. trials) x (num. AAL nodes)
- srate = sampling rate
- coords = coordinate of source points w/in
- Individual and template head model alignment, *.png
- Source and head model alignment, *.png

- ds_pid_match
- ft_convert_units
- ft_freqanalysis
- ft_plot_mesh
- ft_plot_sens
- ft_plot_vol
- ft_prepare_headmodel
- ft_prepare_leadfield
- ft_prepare_sourcemodel
- ft_read_atlas
- ft_read_mri
- ft_resampleddata
- ft_sourceanalysis
- ft_sourcedescriptives
- ft_sourceinterpolate
- ft_sourceplot
- ft_timelockanalysis
- ft_volumelookup
- ft_volumesegment
- load_config
- load_participants
- write_match_if_not_empty

5. Frequency and Connectivity Analysis

fcp_5_freqanalysis.m

1. Setup

- As in fcp_1 and load in virtual sensor data from fcp_4
- 2. Power spectrum**
 - Perform time window analysis
 - Baseline correct the data to control for general/random spikes in power

Outputs

- 4-D matrix containing power spectrum data. Matrix dimensions are: [participants] x [regions] x [frequency] x [time]
- Users can plot frequency by time for a specific region of a given participant's data by applying a plotting function on a slice of the matrix

fcp_5_taskconnectivity.m

1. Setup

- As in fcp_1
- Create filter coefficients

2. Analysis

- Load participant source timeseries matrices (catmatrix)
- Apply Hilbert filter to isolate frequency band
- Return PLI or PLV connectivity between pairs of AAL nodes
- Repeat for each trial and frequency band
- Save individual adjacency matrix
- Then, repeat for all participants
- Take average across trials for each participant and assemble all-participant adjacency matrix

Outputs

- Individual adjacency matrix (nodes x nodes x trial x freq. band)
- All participant adjacency matrix (nodes x nodes x participants x band)

- ds_pid_match
- ft_connectivityanalysis
- load_config
- load_participants
- write_match_if_not_empty