Open-source EEG/MEG analysis software: FieldTrip toolbox in detail

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FieldTrip

MNE

MEG + EEG ANALYSIS & VISUALIZATION

Brainstorm

http://neuroimage.usc.edu/brainstorm

sccn/eeglab

FreeSurfer

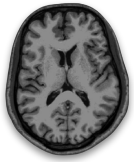
MEGIN

Greeting from MEGIN, Finland

- Providing MEG device and surgical mapping solutions for neuroscience research and clinical applications for more than 35 years.
- More than 120 MEGIN MEG systems located across globe, contributing to research and medical diagnosis.
- Learn more about MEGIN at megin.com

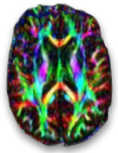
Human brain and neuroimaging

Structural imaging techniques



MRI
(Magnetic resonance imaging)

CT
(Computed tomography)



DTI
(Diffusion tensor imaging)

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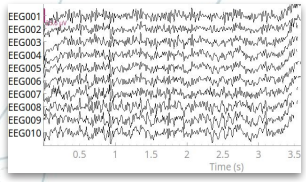
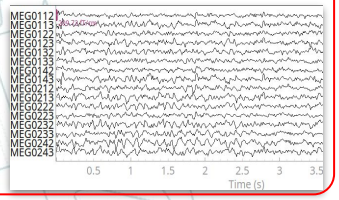
Functional imaging techniques

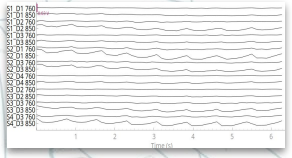
EEG
(Electroencephalography)



MEG
(Magnetoencephalography)



fNIRS
(Functional
near-infrared spectroscopy)



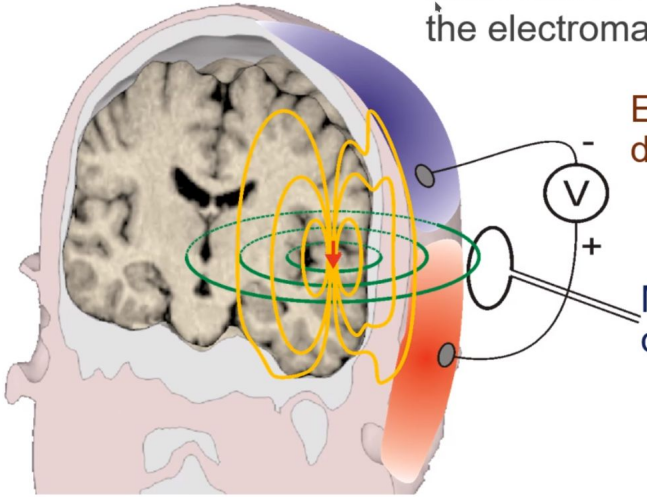
fMRI
(Functional
magnetic resonance imaging)

…

MEGIN

MEG/EEG signals measurement

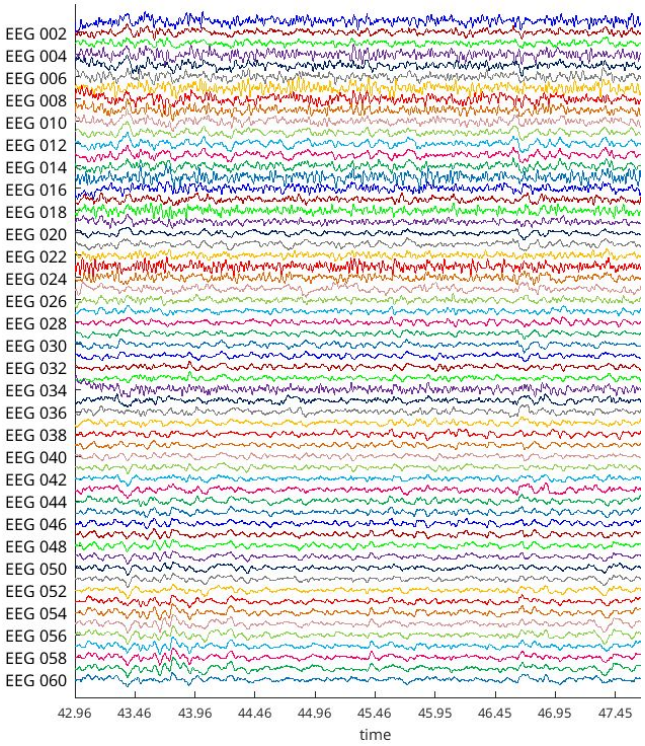
MEG and EEG track electric brain activity by measuring the electromagnetic fields generated by neurons



EEG = measuring the potential differences on the scalp

MEG = measuring neuromagnetic fields outside of the head

EEG data, analysis methods and software



- Evoked responses
- Induced responses
- Oscillatory responses
- Time-frequency/ spectral methods
- Cluster-based permutation test
- Source reconstruction
- Functional connectivity
- Microstates
- Signal complexities
- …

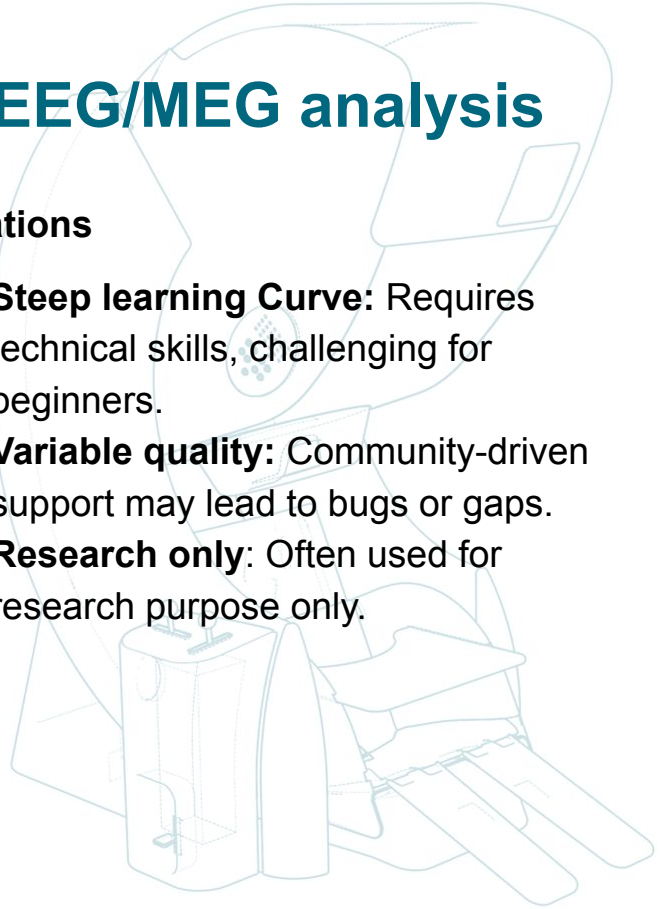
Why to use open-source software for EEG/MEG analysis

Advantages

- **Unlimited access:** Available to all, removing financial barriers to researchers.
- **Customizable:** Source code can be modified to suit specific research needs.
- **Collaborative:** Encourages global contributions, fostering innovation.
- **Up-to-date:** Quickly adapts to new methods and analysis techniques.
- **Transparent:** Open code ensures reproducibility and algorithm validation.
- **Well-documented:** Often includes tutorials, forums, and version control.

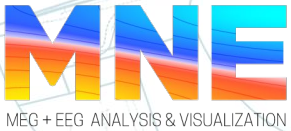
Limitations

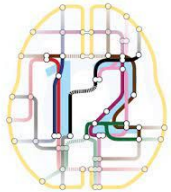
- **Steep learning Curve:** Requires technical skills, challenging for beginners.
- **Variable quality:** Community-driven support may lead to bugs or gaps.
- **Research only**: Often used for research purpose only.



Major open-source EEG/MEG analysis software

- **FieldTrip** https://www.fieldtriptoolbox.org/
  - MATLAB-based scripting.
  - Easy-to-learn but require some coding skills.
- **MNE-Python** https://mne.tools/stable/index.html
  - Python-based; requires Python scripting knowledge.
- **EEGLAB** https://sccn.ucsd.edu/eeglab/
  - Mainly GUI-based; scripting possible.
- **Brainstorm** https://neuroimage.usc.edu/brainstorm/
  - GUI-based; easy-to-use.
  - Scripting possible.
- **SPM12** https://www.fil.ion.ucl.ac.uk/spm/software/spm12/
  - Structural-functional analysis possible; scripting as well as GUI-based.

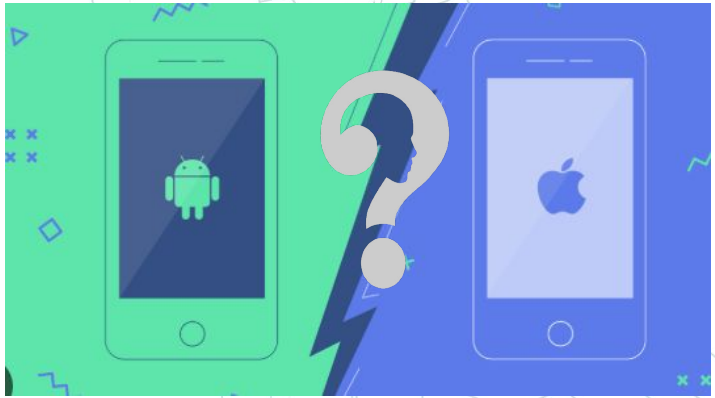




Differences and similarities

- Jaiswal, A., Nenonen, J., Stenroos, M., Oostenveld, R., ... & Parkkonen, L. (2020). Comparison of beamformer implementations for MEG source localization. *NeuroImage*, *216*, 116797.

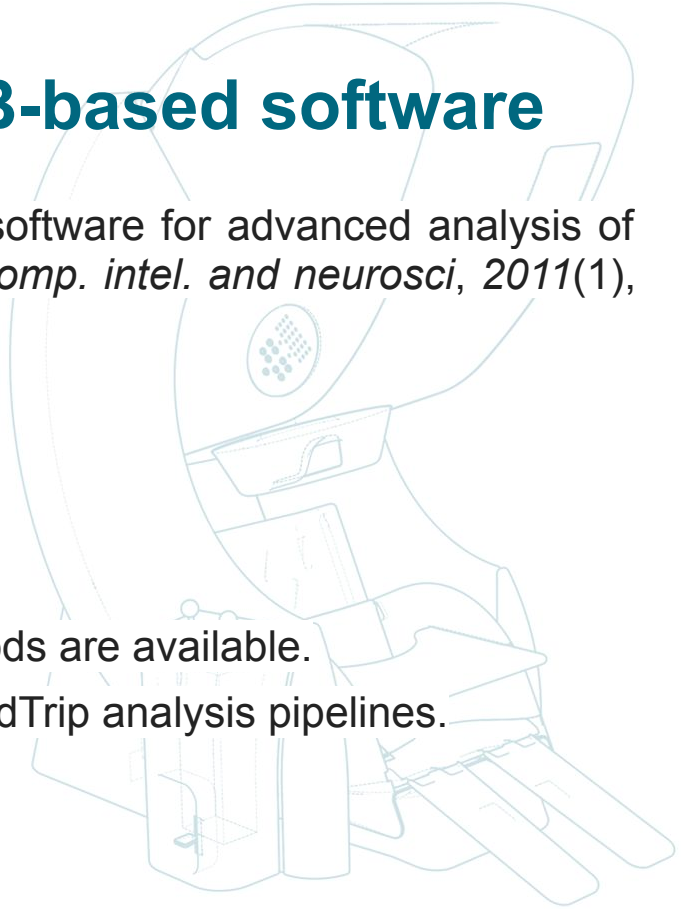
- **Platform:** MATLAB or Python or C/C++
- **Interface:** Coding or Graphical-UI.
- **Usability:** Easy or tedious -to-use.
- **Integration with other s/w**: Simple or complex.
- **Reproducibility:** Similar or different.



FieldTrip: an open-source MATLAB-based software

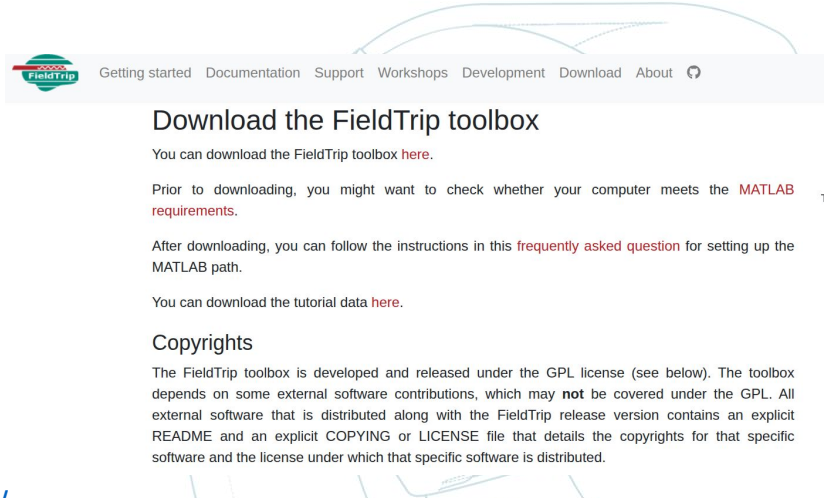
- Oostenveld, R. et al. (2011). FieldTrip: open source software for advanced analysis of MEG, EEG, and invasive electrophysiological data. *Comp. intel. and neurosci*, *2011*(1), 156869.

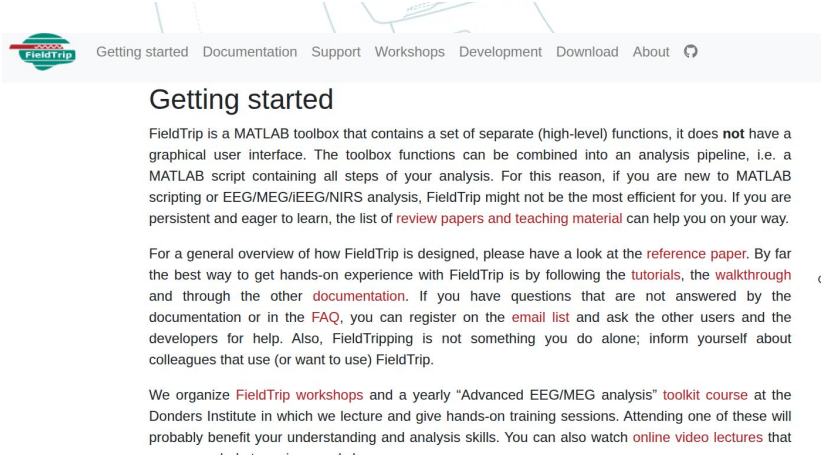
- **Why FieldTrip:**
  - Easy to learn. Simple and well structured code.
  - MATLAB is almost free for academia.
  - Many conventional and advanced analysis methods are available.
  - Several plugins provide easy integration with FieldTrip analysis pipelines.
  - Active developers community.



Starting with FieldTrip

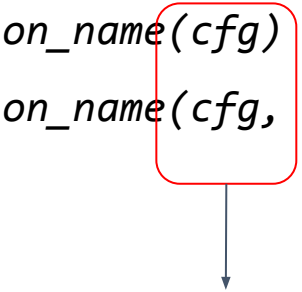
- Where to get it: https://github.com/fieldtrip/fieldtrip
- How to install:
  - *addpath('fieldtrip directory on your drive')*
  - *ft_defaults*
- Where to start it:
  - https://www.fieldtriptoolbox.org/documentation/
  - https://www.fieldtriptoolbox.org/getting_started/
- What can you do with it?
  - Time-domain analysis, spectral analysis, clustering, source reconstruction, connectivity analysis, and many more.





FieldTrip code structure

- *output = function_name(cfg)*
- *output = function_name(cfg, input)*




```
Example:
cfg             = [];
cfg.continuous  = 'yes';
cfg.channel     = 'all';
cfg.dataset     = filename;
raw = ft_preprocessing(cfg);
```

Data structures

Continuous data

disp(raw)

```
      hdr: [1×1 struct]
    label: {70×1 cell}
     time: {1x1 cell}
    trial: {[70×166800 double]}
  fsample: [600.6150]
sampleinfo: [1 166800]
     elec: [1×1 struct]
      cfg: [1×1 struct]
```

Segmented data

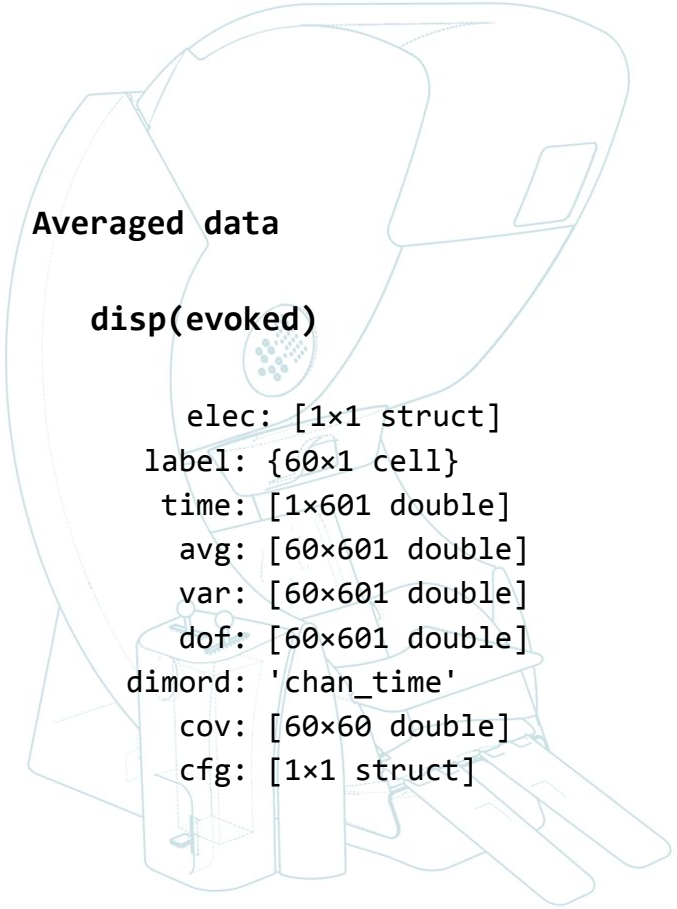
disp(epochs)

```
     elec: [1×1 struct]
  fsample: 600.6150
    label: {60×1 cell}
      cfg: [1×1 struct]
     hdr: [1×1 struct]
    trial: {1×71 cell}
     time: {1×71 cell}
 trialinfo: [71×1 double]
sampleinfo: [71×2 double]
```

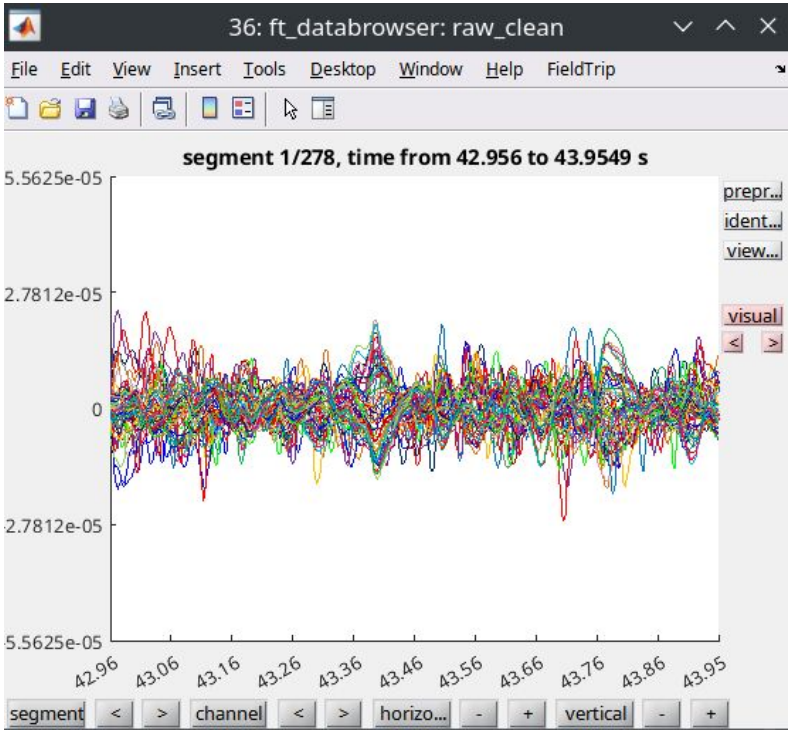
Averaged data

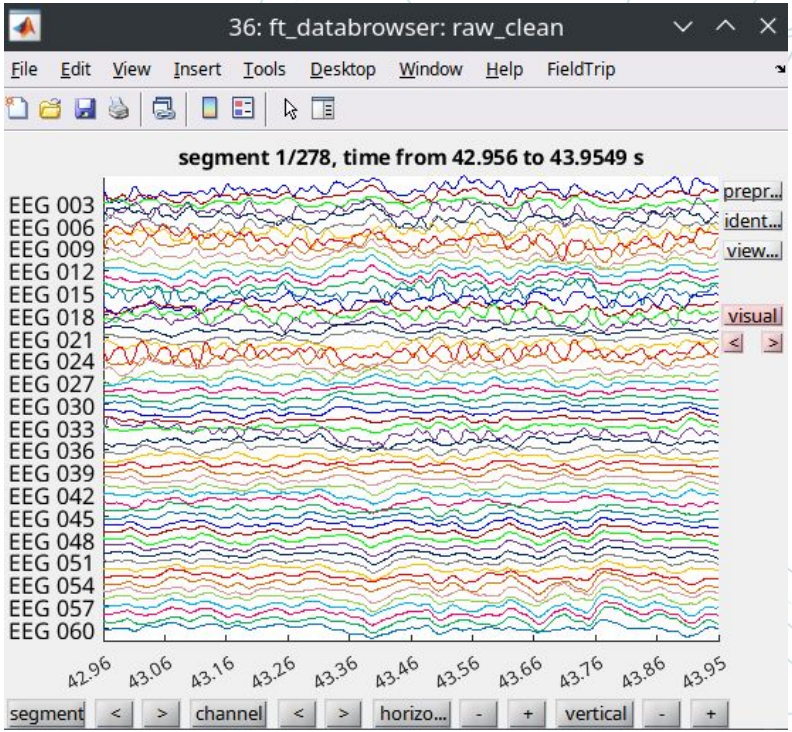
disp(evoked)

```
     elec: [1×1 struct]
    label: {60×1 cell}
     time: [1×601 double]
      avg: [60×601 double]
      var: [60×601 double]
      dof: [60×601 double]
  dimord: 'chan_time'
      cov: [60×60 double]
      cfg: [1×1 struct]
```

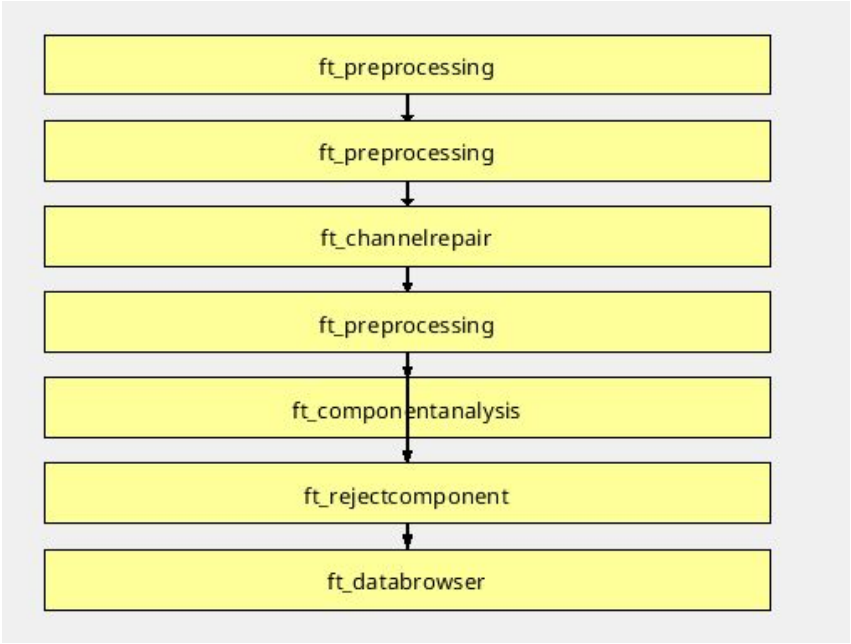


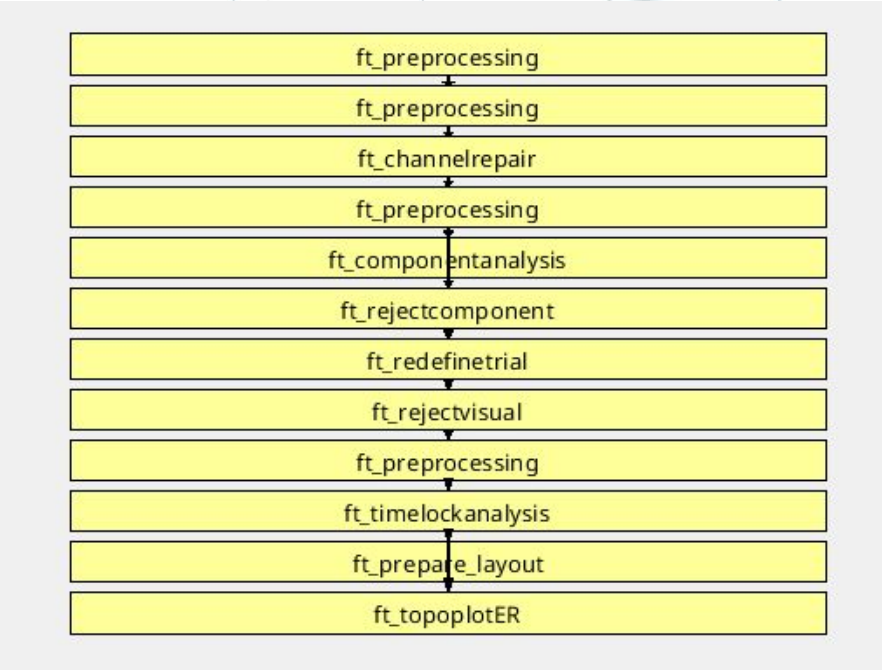
Data browser or reviewing tool





Processing pipelines





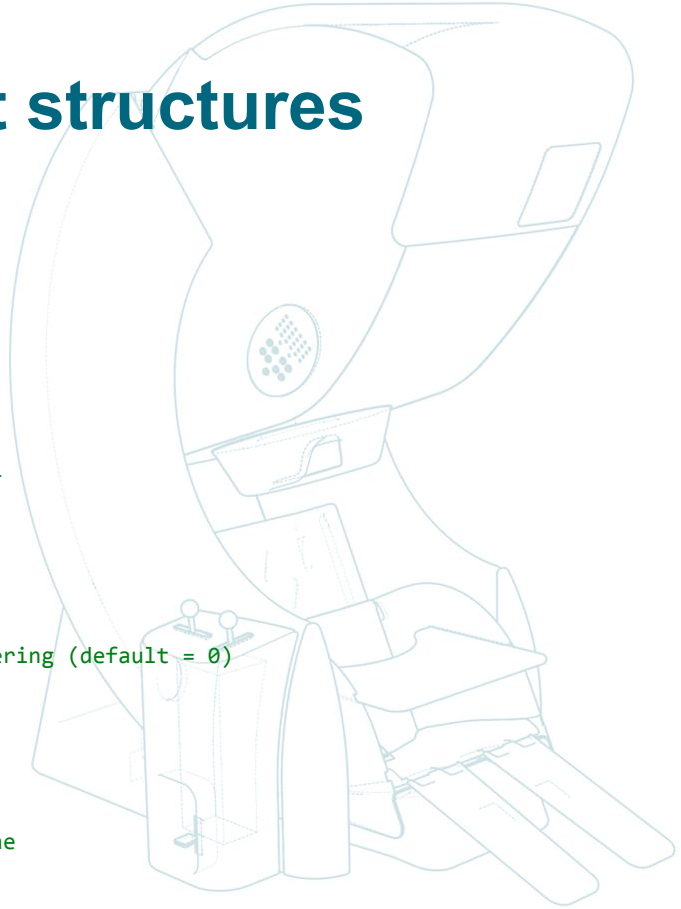
Sample scripts

```
addpath('..//..//TPTools//fieldtrip//)
ft_defaults

%% Set data directory and other parameters
data_dir = '..//..//..//Workshop_IITMandi/';
filename = [data_dir,
'sample_audvis_raw_eeg.fif'];
par             = [];
par.bpfreq      = [1 45];

%% Read the continuous data
cfg             = [];
cfg.continuous = 'yes';
cfg.channel     = 'all';
cfg.dataset     = filename;
raw       = ft_preprocessing(cfg);
disp(raw)

%% Review
cfg = [];
cfg.preproc.demean   = 'yes';
artf = ft_databrowser(cfg, raw);

%% Prepare and plot EEG layout/montage
cfg = [];
cfg.elec     = raw.elec;
cfg.channel = {'EEG*'};
lay2D = ft_prepare_layout(cfg, raw);
```

```
%% Preprocess continuous raw data
cfg = [];
cfg.bpfilter   = 'yes';
cfg.bpfiltord  = 2;
cfg.bpfilttype = 'but';
cfg.bpfreq     = par.bpfreq;
cfg.channel    = {'EEG*'};
rawfilt_eeg  = ft_preprocessing(cfg, raw);

%% Review
cfg_browse = [];
ft_databrowser(cfg_browse, rawfilt_eeg);

%% Reconstruct bad channels
cfg = [];
cfg.senstype   = 'eeg';
cfg.method     = 'spline';
cfg.badchannel = par.badch;
cfg.neighbours = neighbours;
rawfilt_eeg = ft_channelrepair(cfg, rawfilt_eeg);

%% Review
ft_databrowser(cfg_browse, rawfilt_eeg);
```

Function-specific details and input structures

```
function [data] = ft_preprocessing(cfg, data)
% FT_PREPROCESSING reads MEG and/or EEG data according to user-specified trials
% and applies several user-specified preprocessing steps to the signals.
%
% Use as
%   [data] = ft_preprocessing(cfg)
% or
%   [data] = ft_preprocessing(cfg, data)
%
% The first input argument "cfg" is the configuration structure, which contains all
% details for the dataset filename, trials and the preprocessing options.
%
% If you are calling FT_PREPROCESSING with only the configuration as first input
% argument and the data still has to be read from file, you should specify
%   cfg.dataset      = string with the filename
%   cfg.trl          = Nx3 matrix with the trial definition, see FT_DEFINETRIAL
%   cfg.padding      = length (in seconds) to which the trials are padded for filtering (default = 0)
%   cfg.padtype      = string, type of padding (default: 'data' padding or
%                      'mirror', depending on feasibility)
%   cfg.continuous   = 'yes' or 'no' whether the file contains continuous data
%                      (default is determined automatic)
%
% Instead of specifying the dataset in the configuration, you can also explicitly
% specify the name of the file containing the header information and the name of the
% file containing the data, using
%   cfg.datafile     = string with the filename
```



Thanks for your attention!

MEGIN