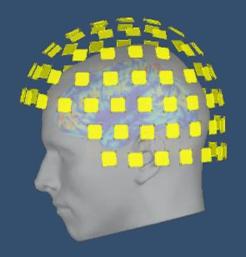
MEG and EEG analysis using Brainstorm http://neuroimage.usc.edu/brainstorm









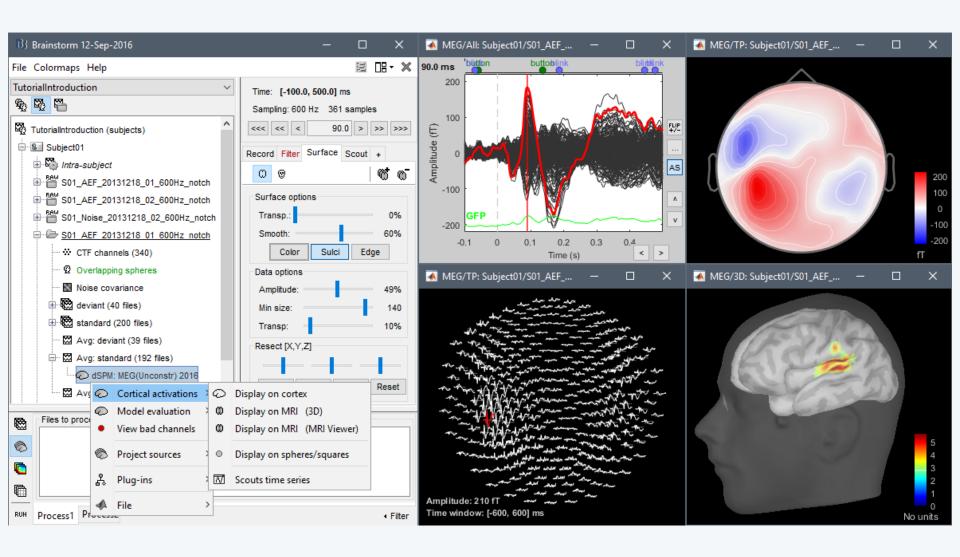


Beijing - April 2017

Francoi: Tadel

Sylvain Baillet

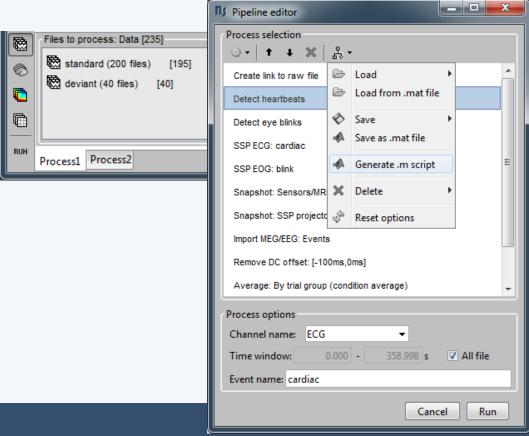
Graphic interface





Scripting environment

- Rapid selection of files and processes to apply
- Automatic generation of Matlab scripts
- Plug-in structure: easy to add custom processes



```
% Script generated by Brainstorm v3.1 (17-Dec-2010).
FileNamesA = {'SubjectO1\Left\data average 101213 1558.mat', ...
              'Subject01\Right\data average 101213 1559.mat');
FileNamesB = [];
% Process: Detect bad trials: Peak-to-peak MEGGRAD(0-2000)
sFiles = bst process(...
    'CallProcess', 'process detectbad', ...
   FileNamesA, FileNamesB, ...
    'timewindow', [-0.0998, 0.3000], ...
    'meggrad', {[0, 2000], 'fT/cm (x 0.04)', 1e-015), ...
    'rejectmode', 2);
% Process: Remove baseline: [-100ms,-1ms]
sFiles = bst process(...
    'CallProcess', 'process baseline', ...
    sFiles, [], ...
    'baseline', [-0.09983, -0.00056], ...
    'overwrite', 1);
% Process: Band-pass filter: 1Hz - 80Hz
sFiles = bst process(...
    'CallProcess', 'process_bandpass', ...
    sFiles, [], ...
    'f1' 1. ...
    'f2', 80, ...
    'overwrite', 1);
% Process: Average by condition
sFiles = bst process(...
    'CallProcess', 'process average', ...
    sFiles, [], ...
    'avgtype', 3, ...
    'isstd', 0);
```



Brain/torm

- Free and open-source application
- Matlab & Java: Platform-independent
- Designed for Matlab
- Stand-alone version available
- Interface-based: click, drag, drop
- No programming experience required
- Daily updates of the software
- Supports most common file formats











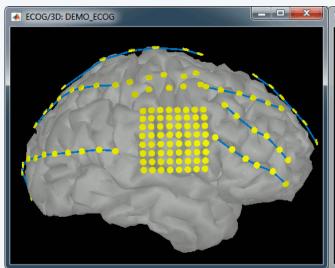


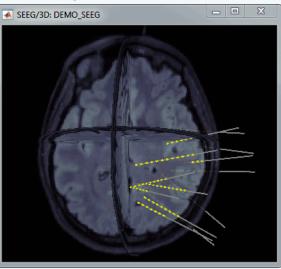


Multi-modal imaging

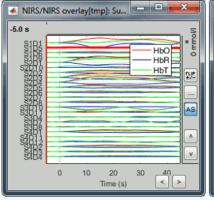
ECoG

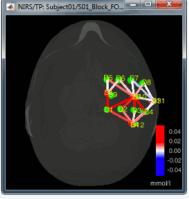
Depth electrodes

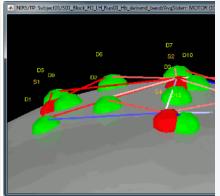




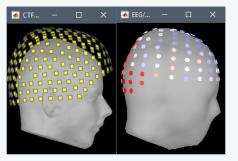
fNIRS







MEG/EEG



Electrophysiology

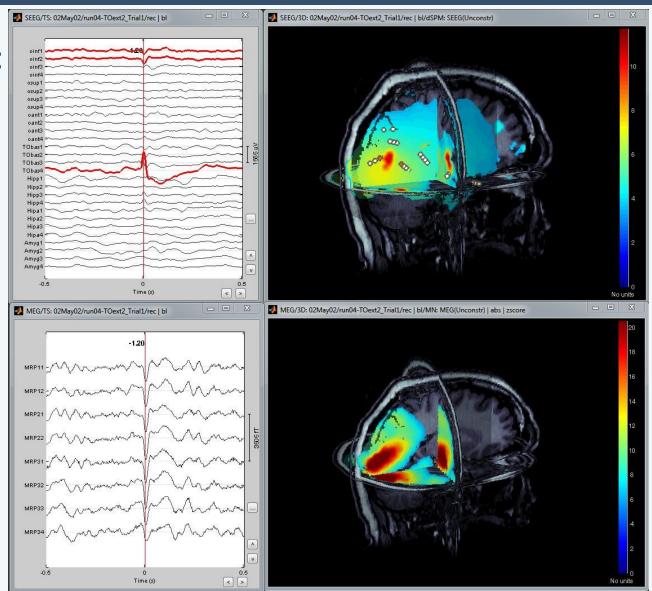




Multi-modal imaging

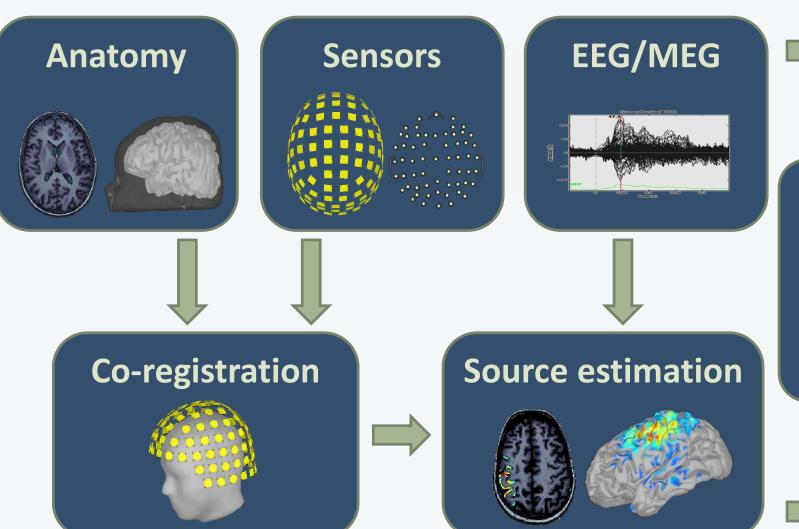
Easy integration of:

- MEG
- EEG
- ECoG
- SEEG
- Animal LFP
- NIRS





Workflow





Analysis

Averages
Contrasts
Group analysis
Time-frequency
Connectivity



Anatomy
Link recordings
MRI registration

Importing

PSD

Filters

Bad channels

Artifacts

Correction

Bad segments

Pre-processing

Markers

Epoching

Averaging

Sources

Time-frequency

Analysis of the experimental data

Loop:
all acquisition runs
all subjects



Anatomy

Link recordings MRI registration

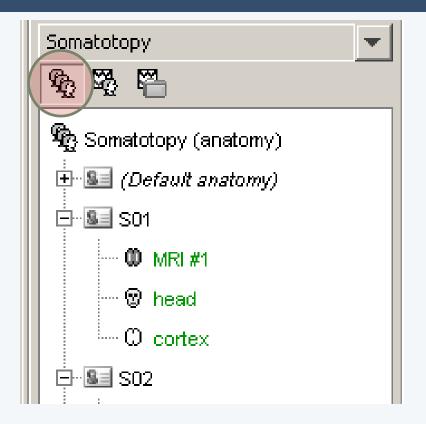
PSD Filters Bad channels Artifacts Correction Bad segments

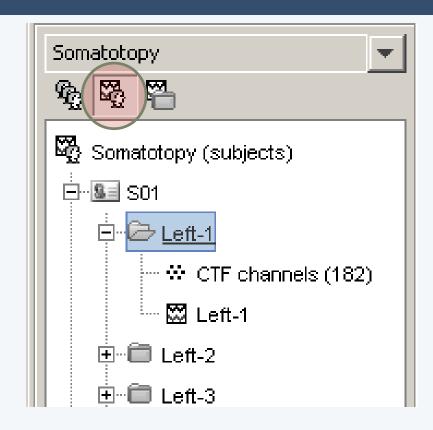
Markers Epoching Averaging Sources Time-frequency

- One-click import of the T1 segmentation: FreeSurfer, BrainSuite, BrainVISA, CIVET
- Import and place fiducials in the MRI (N,L,R)



Database





- Three levels:
 - Protocol
 - Subject
 - Condition

- Popup menus
- All files saved in Matlab .mat
- Same architecture on the disk

Anatomy

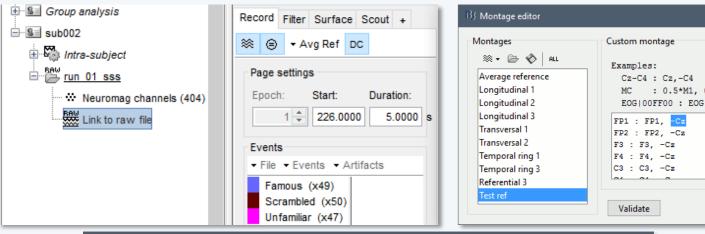
Link recordings

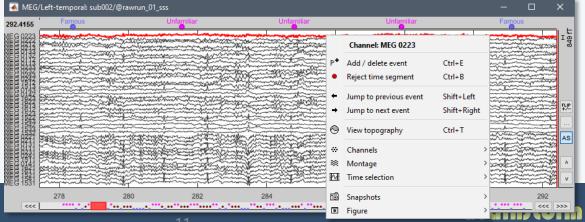
MRI registration

PSD Filters Bad channels **Artifacts** Correction Bad segments

Markers Epoching Averaging Sources Time-frequency

- Original files linked to the database (no copy)
- Rich data viewer with flexible montage editor
- Optimized reading functions







: 0.5*M1, 0.5*M2

Co-registration MEG / MRI (I)

Anatomy
Link recordings
MRI registration

PSD
Filters
Bad channels
Artifacts
Correction
Bad segments

Markers
Epoching
Averaging
Sources
Time-frequency

- Basic estimation based on three points:
 Nasion (NAS), Left ear (LPA), Right ear (RPA)
- MRI: Marked in the volume with the MRI Viewer
- MEG: Obtained with a tracking system (Polhemus)





Co-registration MEG / MRI (2)

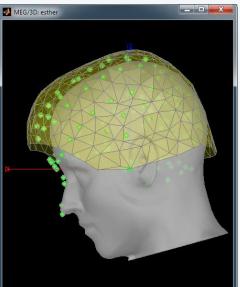
Anatomy
Link recordings
MRI registration

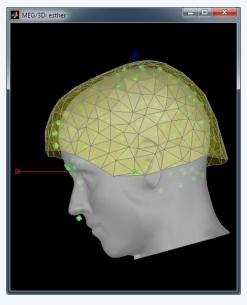
PSD
Filters
Bad channels
Artifacts
Correction
Bad segments

Markers
Epoching
Averaging
Sources
Time-frequency

- Automatic adjustment based on head shape:
 Fitting Polhemus points on the MRI head surface
- Final registration must be checked manually
- Polhemus driver included in Brainstorm







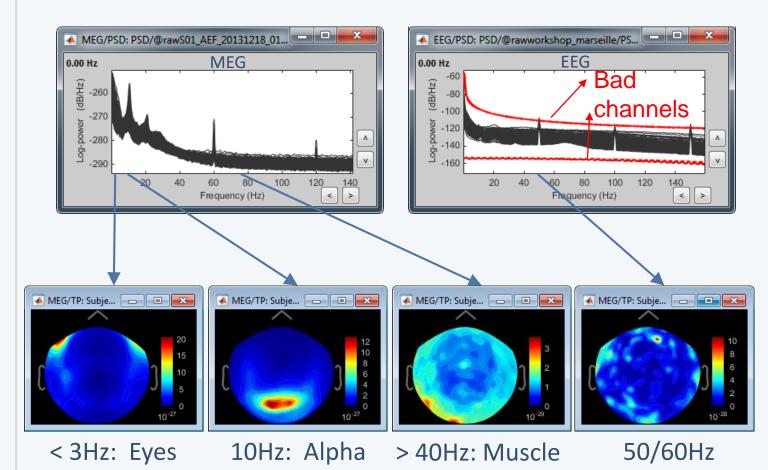
Anatomy
Link recordings
MRI registration

PSD

Filters
Bad channels
Artifacts
Correction
Bad segments

Markers
Epoching
Averaging
Sources
Time-frequency

Power spectrum density for quality control



Anatomy Link recordings MRI registration

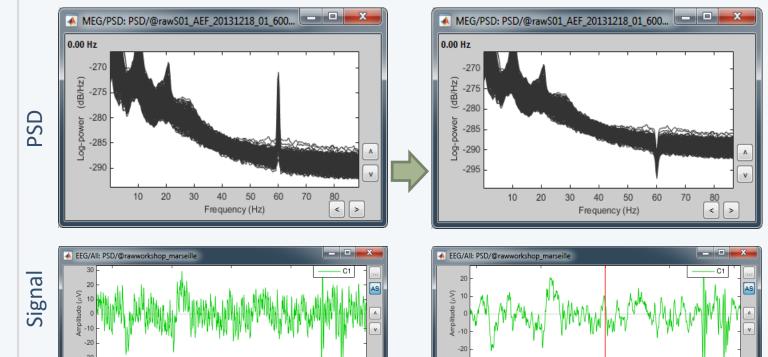
 Notch filter: Removes 50Hz/60Hz power line noise (and harmonics)

PSD

Filters

Bad channels
Artifacts
Correction
Bad segments

Markers
Epoching
Averaging
Sources
Time-frequency



Anatomy
Link recordings
MRI registration

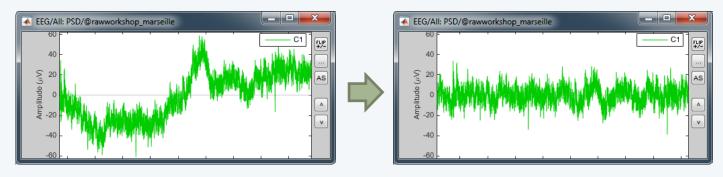
PSD

Filters

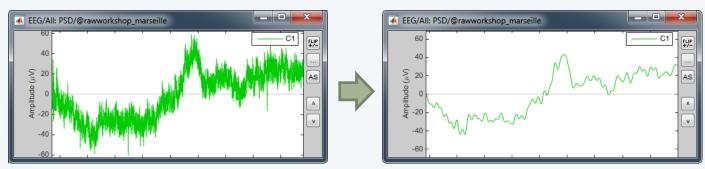
Bad channels
Artifacts
Correction
Bad segments

Markers
Epoching
Averaging
Sources
Time-frequency

 High-pass filter: Removes slow components (eye movements, breathing, sensor drifts...)



Low-pass filter: Remove high-frequencies



Anatomy
Link recordings
MRI registration

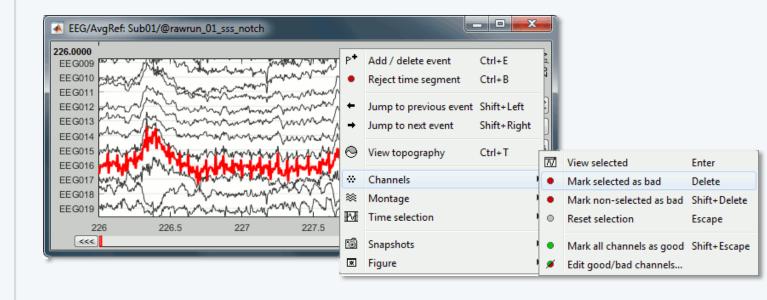
PSD Filters

Bad channels

Artifacts
Correction
Bad segments

Markers
Epoching
Averaging
Sources
Time-frequency

- Manual inspection of the recordings
- Interactive selection of bad channels
- Re-reference the EEG if necessary (Average ref)



Anatomy
Link recordings
MRI registration

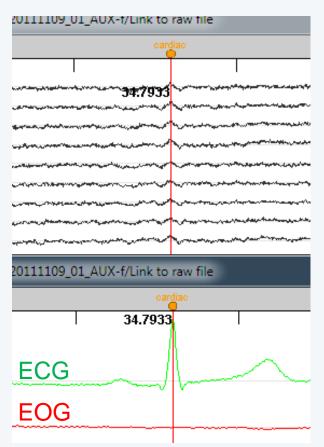
PSD Filters Bad channels

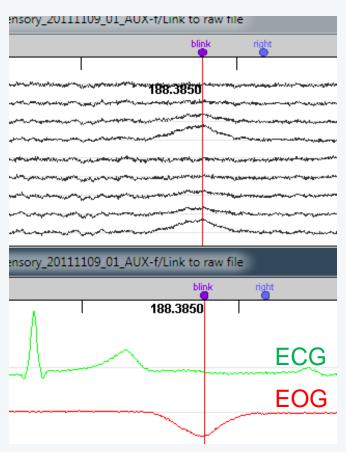
Artifacts

Correction
Bad segments

Markers
Epoching
Averaging
Sources
Time-frequency

 Automatic detection of blinks and heartbeats (peak detection, or explicit amplitude threshold)







Artifact correction

- Two categories of artifacts:
 - Well defined, reproducible, short, frequent:
 - Heartbeats, eye blinks, eye movements, some stimulators
 - Unavoidable and frequent: we cannot just ignore them
 - Can be modeled and removed from the signal efficiently
 - All the other events that can alter the recordings:
 - Movements, building vibrations, metro nearby...
 - Too complex or not repeated enough to be modeled
 - Safer to mark them as bad segments, and ignore them



Anatomy
Link recordings
MRI registration

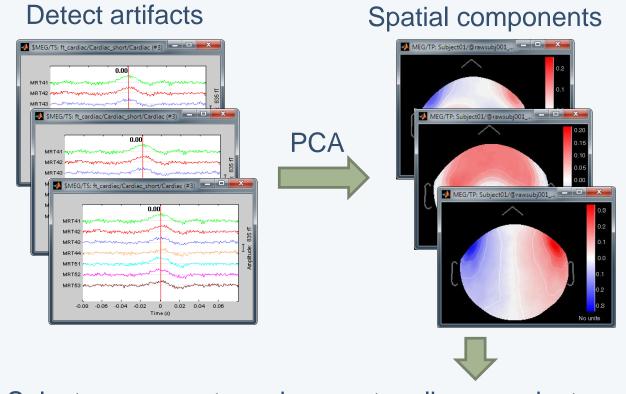
PSD Filters Bad channels Artifacts

Correction

Bad segments

Markers
Epoching
Averaging
Sources
Time-frequency

Correction with Signal Space Projections (SSP)



Select components and compute a linear projector to remove their contribution from the recordings

Anatomy
Link recordings
MRI registration

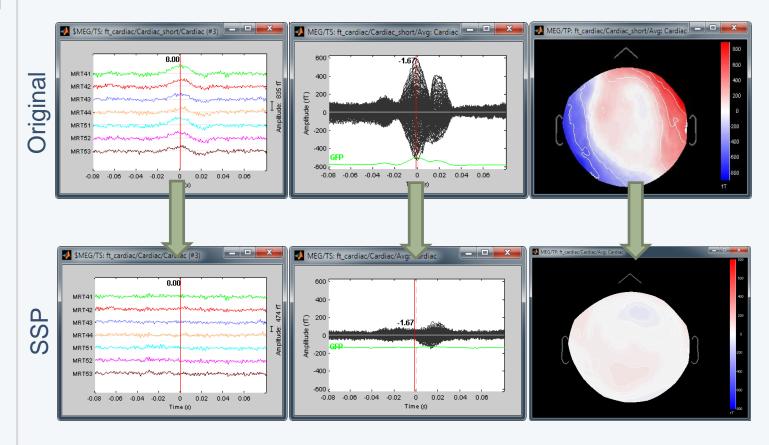
PSD Filters Bad channels Artifacts

Correction

Bad segments

Markers
Epoching
Averaging
Sources
Time-frequency

Example: Cardiac artifact



Anatomy
Link recordings
MRI registration

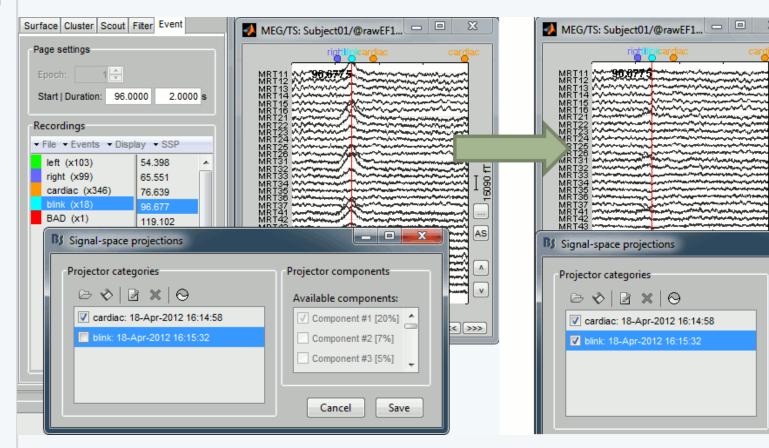
PSD Filters Bad channels Artifacts

Correction

Bad segments

Markers
Epoching
Averaging
Sources
Time-frequency

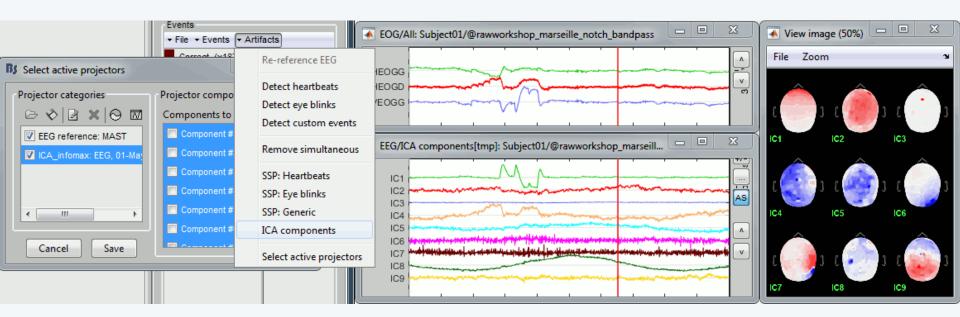
Example: Blink





Artifact correction

- Independent component analysis (ICA):
 - Popular in the EEG literature
 - Alternative to SSP for low number of sensors
 - Already implemented: Infomax and JADE (EEGLAB)





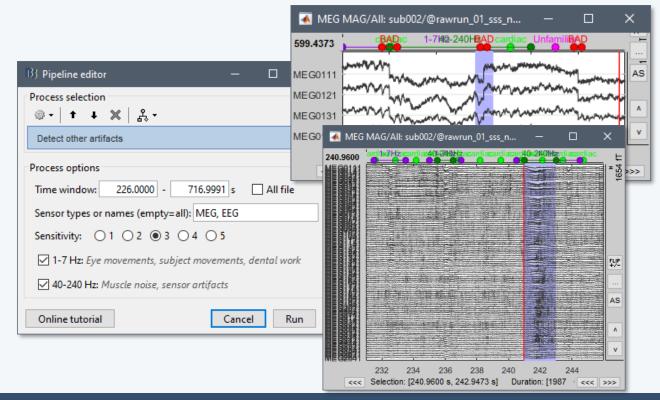
Anatomy
Link recordings
MRI registration

PSD
Filters
Bad channels
Artifacts
Correction

Bad segments

Markers
Epoching
Averaging
Sources
Time-frequency

- Automatic detection of artifacts (RMS-based)
- Manual screening of all the recordings is advised (scroll all the sensors by pages of 10-20s)
- Exclude: Blinks, movements, SQUID jumps

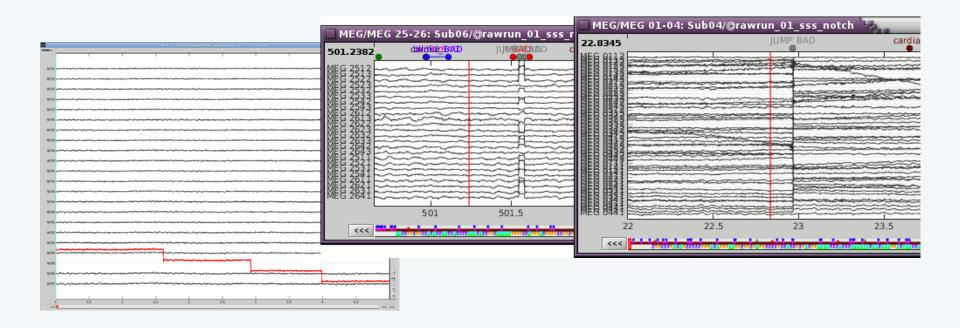




Elekta-Nevromag

SQUID jumps

- Sharps steps followed by a change of baseline value
- Mark the channels as bad before running MaxFilter
- Or mark the segments as bad in Brainstorm



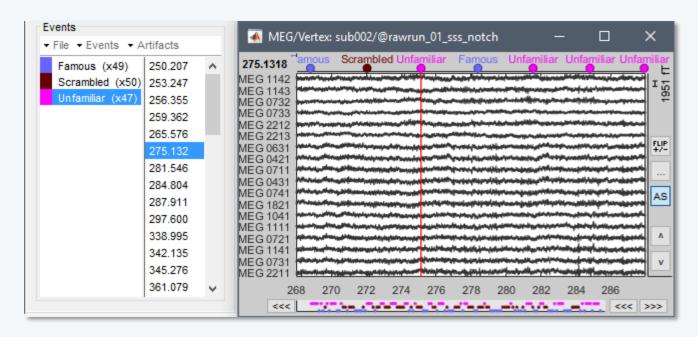
Anatomy
Link recordings
MRI registration

PSD
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Correction
Bad segments

Markers

Presentation Sensor Manual

- Two types of experiments:
 - Steady-state or resting-state (ongoing activity)
 - Event-based (stimulus, response, spike...)
- How to get event markers in the recordings?



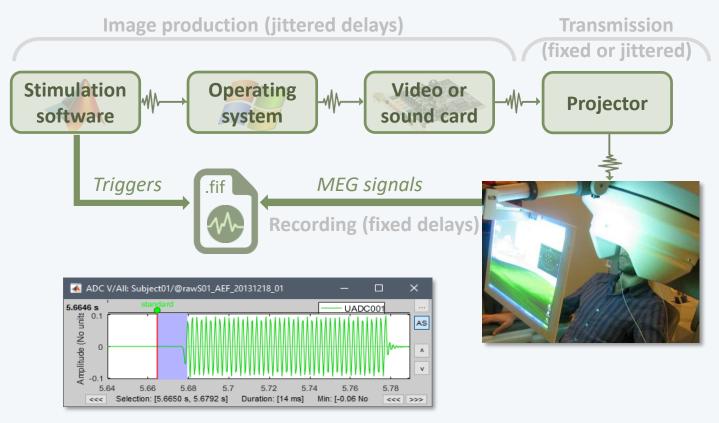
Anatomy
Link recordings
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PSD
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Correction
Bad segments

Markers

Presentation

Sensor Manual Reading the triggers saved by the presentation software (includes jittered OS delays)



File triggers are never aligned with the real stim



Anatomy
Link recordings
MRI registration

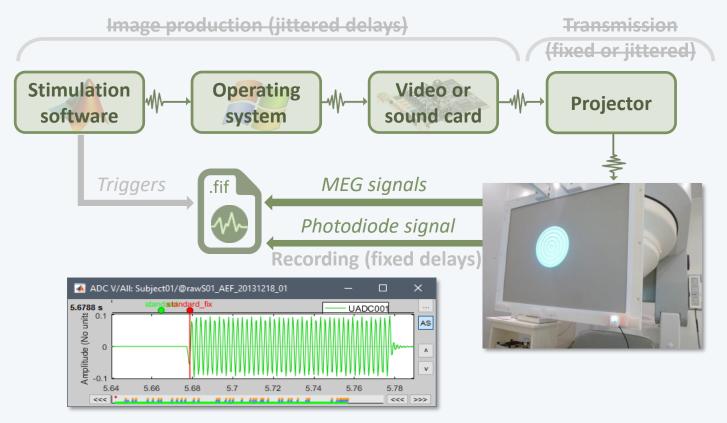
PSD
Filters
Bad channels
Artifacts
Correction
Bad segments

Markers
Presentation

Sensor

Manual

 Reading information recorded on the subject side (photodiode, microphone, response box...)



Avoids most uncontrollable jittered delays



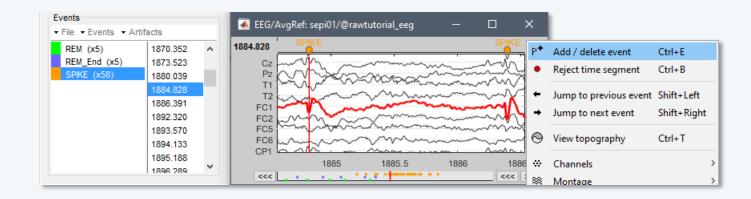
Anatomy
Link recordings
MRI registration

PSD
Filters
Bad channels
Artifacts
Correction
Bad segments

Markers
Presentation
Sensor

Manual

- Manual or automatic marking of biological or behavioral events, post-acquisition (epileptic spikes, sleep spindles, rat position in a box...)
- Optimized workflow for clinicians (keyboard and mouse shortcuts, workspace...)



Anatomy
Link recordings
MRI registration

PSD
Filters
Bad channels
Artifacts
Correction
Bad segments

Markers Epoching

Combine

Extract Length Process • In some experiments, the event of interest is a combination of presentation markers.

Example: Priming experiments



- In Brainstorm, the events of interest are saved as new marker types in the continuous files.
- In other environments, the events are sometimes defined only at the epoching time and not saved in the continuous file.

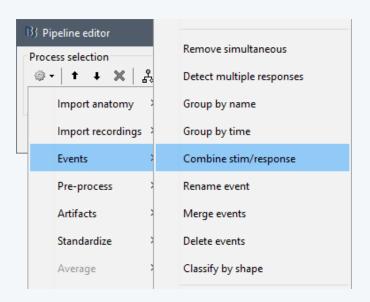
Anatomy
Link recordings
MRI registration

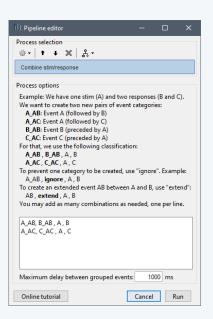
PSD
Filters
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Artifacts
Correction
Bad segments

Markers Epoching

Combine

Extract Length Process In Brainstorm, the events of interest are saved as new marker types in the continuous files.





• In other environments, the events are sometimes defined only at the epoching time and not saved in the continuous file.

Anatomy
Link recordings
MRI registration

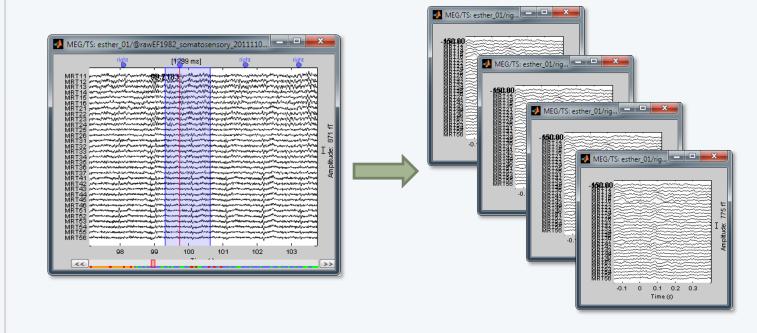
PSD
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Bad channels
Artifacts
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Bad segments

Markers
Epoching
Combine

Extract

Length Process

- Epochs = Trials = Short blocks of recordings around an event of interest.
- Epoching = Extracting epochs from the continuous recordings and saving them.





Anatomy
Link recordings
MRI registration

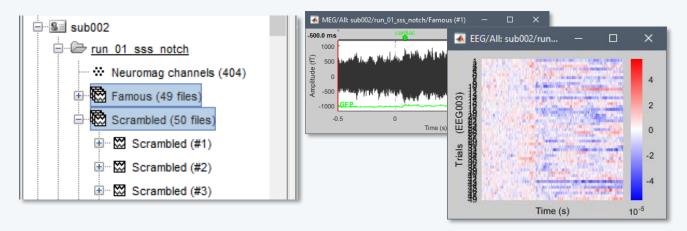
PSD
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Bad segments

Markers
Epoching
Combine

Extract

Length Process

- In Brainstorm, each imported epoch is an independent file in the database.
- Accessible by event type or individually.



 In other programs, all the epochs from one run are saved in one single file (one file per event type, or one file with all the events).



Anatomy
Link recordings
MRI registration

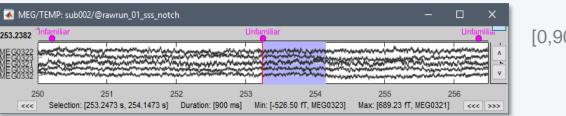
PSD
Filters
Bad channels
Artifacts
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Bad segments

Markers
Epoching
Combine
Extract

Length

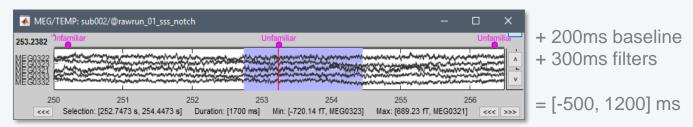
Process

- How to define the optimal epoch length?
- Experimental design:
 Expected effect duration, inter-stimulus interval



[0,900] ms

 Analysis: Frequency filters and amplitude normalizations may require longer epochs



Computational limitations: Size and time



Anatomy
Link recordings
MRI registration

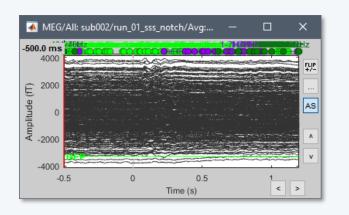
PSD
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Correction
Bad segments

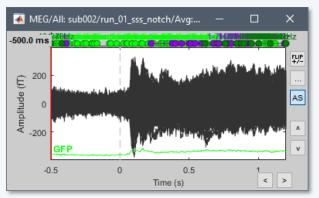
Markers
Epoching
Combine
Extract
Length

Process

Processing steps that can be applied on epochs:

- DC offset correction: Subtract the average estimated over a baseline period
- Detrending: Subtract a linear trend estimated over a reference period
- Resampling: Decrease the sampling rate
- This dataset: DC correction, baseline=[-500,0]ms







Anatomy
Link recordings
MRI registration

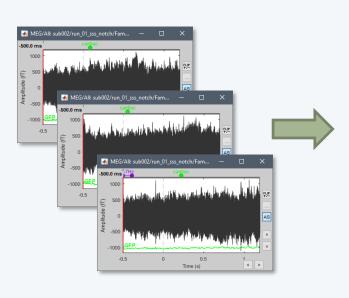
PSD
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Bad segments

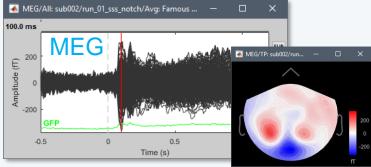
Markers Epoching

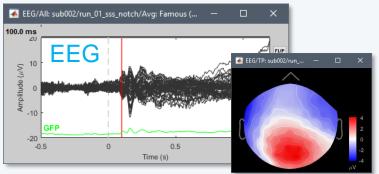
Averaging

Sources
Time-frequency

- Averaging the trials: Reveals the features of the signals that are locked in time to a given event
 - = Event-related field / potential
 - = Evoked response
 - = ERF/ERP









Anatomy
Link recordings
MRI registration

PSD
Filters
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Correction
Bad segments

Markers
Epoching
Averaging

Sources Time-frequency

- **EEG**: Averaging data across runs and subjects OK.
- MEG: Averaging across runs is not always accurate
 - Head shapes differ between subjects.
 - Head positions different between runs.
 - One sensor does not record the same thing in two different runs.
 - Coregistration of runs with Elekta MaxFilter helps but modifies a lot the recordings.
 Never use this to average across subjects.
 - Recommended: Estimate the sources for each run separately, then average in source space.



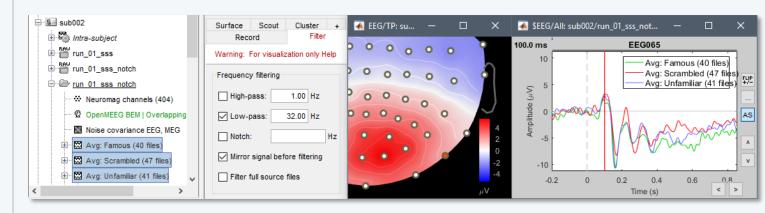
Anatomy
Link recordings
MRI registration

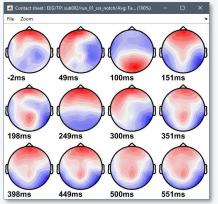
PSD
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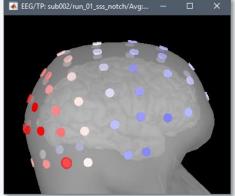
Markers Epoching

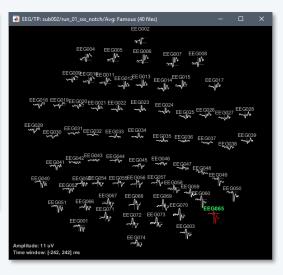
Averaging

Sources Time-frequency EEG ERP: Famous faces











Anatomy
Link recordings
MRI registration

PSD
Filters
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Correction
Bad segments

Markers Epoching Averaging

Sources

Time-frequency

Source space:

Cortex or full head volume

Forward model:

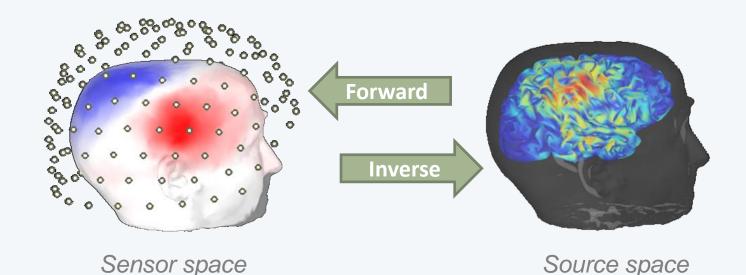
Overlapping spheres (MEG)
OpenMEEG BEM (EEG)

Inverse model:

Minimum norm estimates

Beamformers

Separately for MEG and EEG



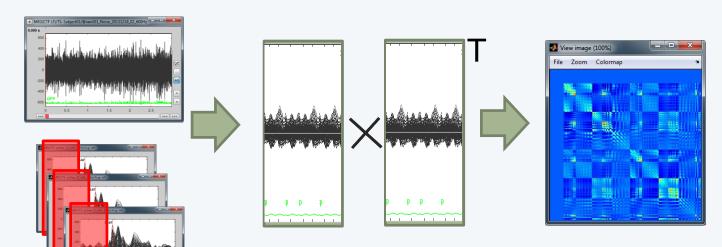
Anatomy
Link recordings
MRI registration

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Markers Epoching Averaging

Sources

- The MNE model requires an estimation of the level of noise of the sensors
- Noise covariance matrix = covariance of segments that do not contain any "meaningful" data
- Empty room, pre-stim baseline, resting



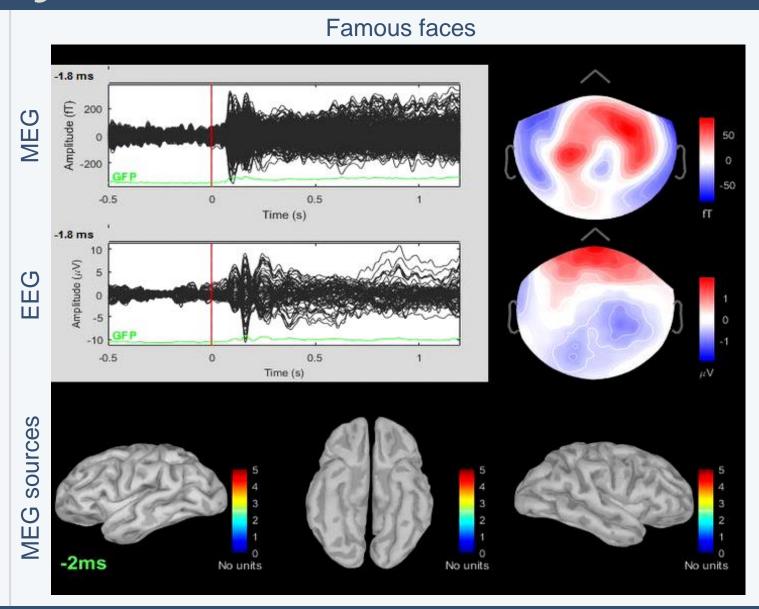


Anatomy
Link recordings
MRI registration

PSD
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Markers Epoching Averaging

Sources



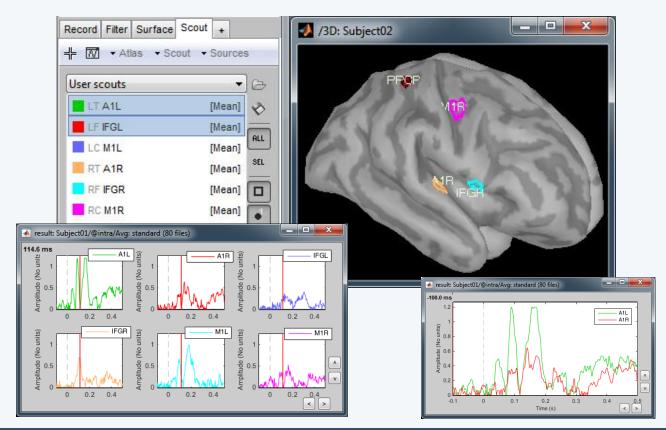
Anatomy
Link recordings
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Correction
Bad segments

Markers Epoching Averaging

Sources

- Regions of interest at cortical level (scouts)
 - = Subset of a few dipoles in the brain
 - = Group of vertices of the cortex surface





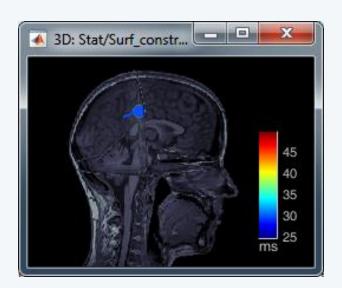
Anatomy
Link recordings
MRI registration

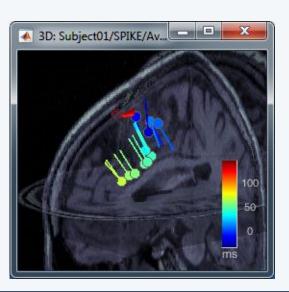
PSD
Filters
Bad channels
Artifacts
Correction
Bad segments

Markers Epoching Averaging

Sources

- Dipole scanning
 Compute a distributed source model, then find the most significant dipole at each time sample.
- Dipole fitting (FieldTrip)
 Non-linear search of the dipoles that minimizes the residuals (data explained - recordings)







Anatomy
Link recordings
MRI registration

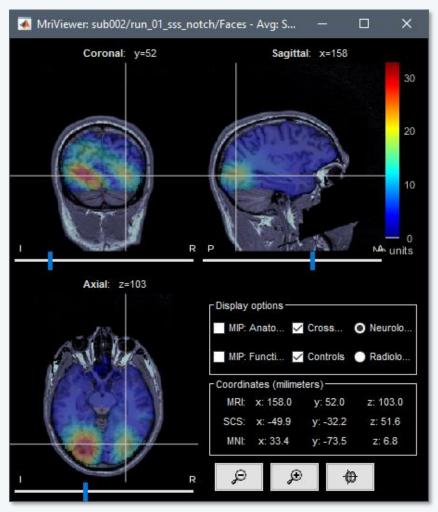
PSD
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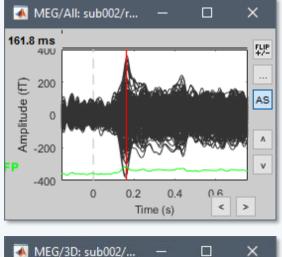
Markers Epoching Averaging

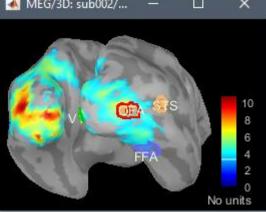
Sources

Time-frequency

Faces – Scrambled, 160ms



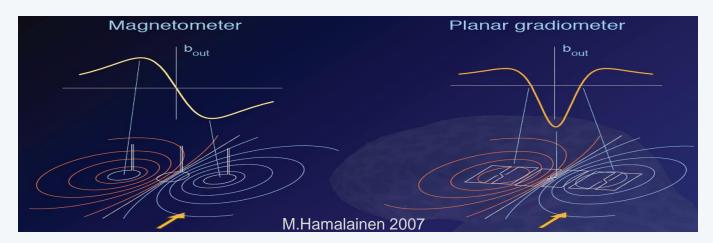






Source estimation: MEG

- Recommended in MEG analysis:
 - The subject head can move in the helmet
 - One sensor is not corresponding to one brain region
 - Different types of sensors (magneto / gradiometers)
 - Difficult to read, reproduce or compare
- Converting to source space helps solving those issues





Source estimation: EEG

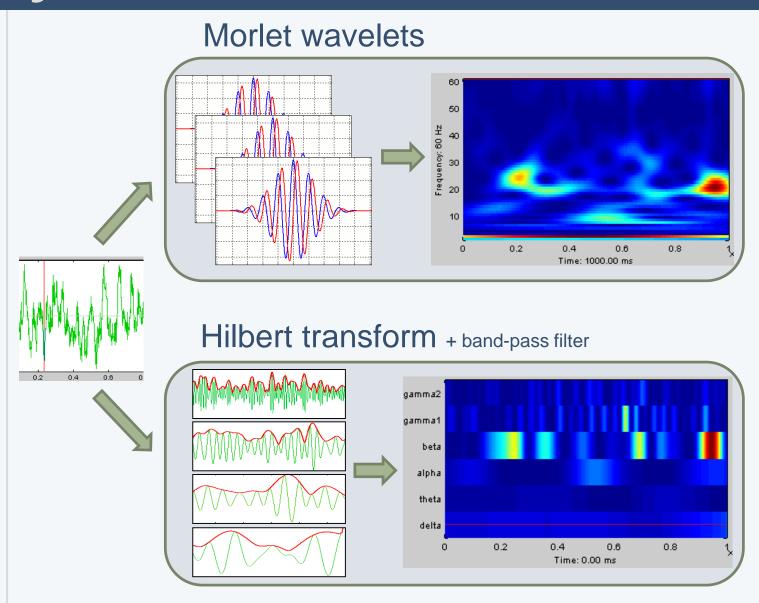
- In EEG, those problems don't exist:
 - Electrodes positions are fixed and known
 - More reproducible signal shapes and topographies
 - Clinicians are trained to work at the sensor level
- But the source reconstruction is still interesting:
 - Localize the signal generators in the brain (epilepsy and pre-surgical functional mapping)
 - Spatial separation of simultaneous sources



Anatomy
Link recordings
MRI registration

PSD
Filters
Bad channels
Artifacts
Correction
Bad segments

Markers
Epoching
Averaging
Sources

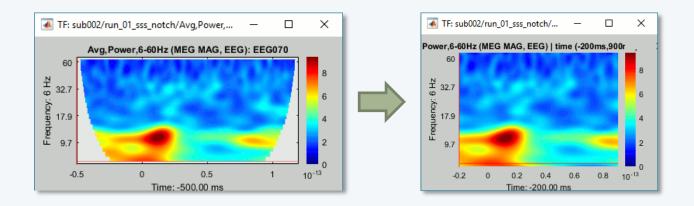


Anatomy
Link recordings
MRI registration

PSD
Filters
Bad channels
Artifacts
Correction
Bad segments

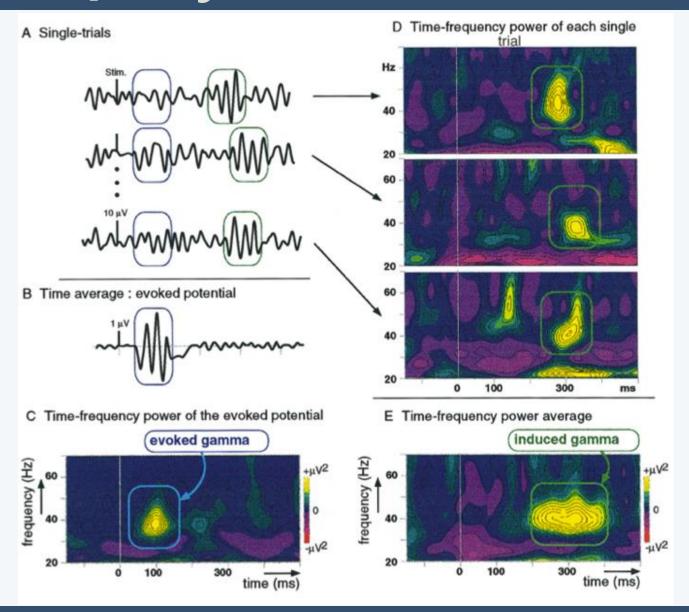
Markers
Epoching
Averaging
Sources

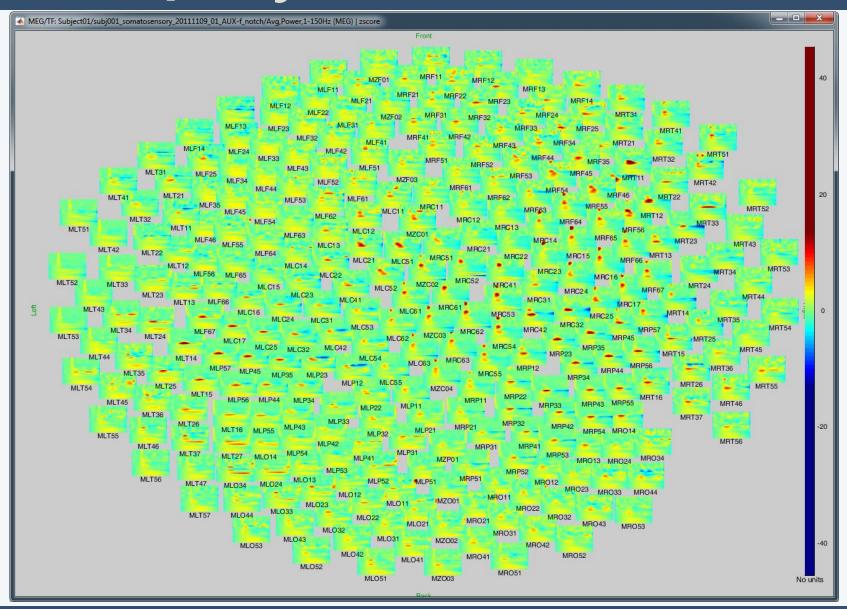
- Make sure you know the transient periods for all the frequency bins, and do not include them in your analysis.
- An easy solution is to cut them out immediately after the computation.



Time-frequency

Average





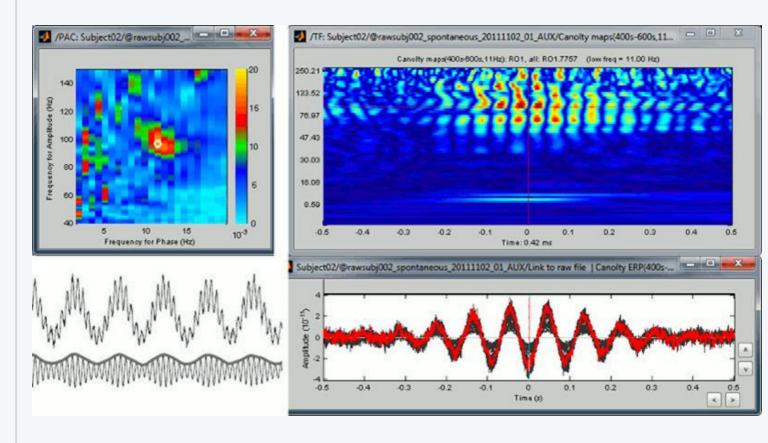
Anatomy
Link recordings
MRI registration

PSD
Filters
Bad channels
Artifacts
Correction
Bad segments

Markers
Epoching
Averaging
Sources
Time-frequency

Other measures

Phase-amplitude coupling



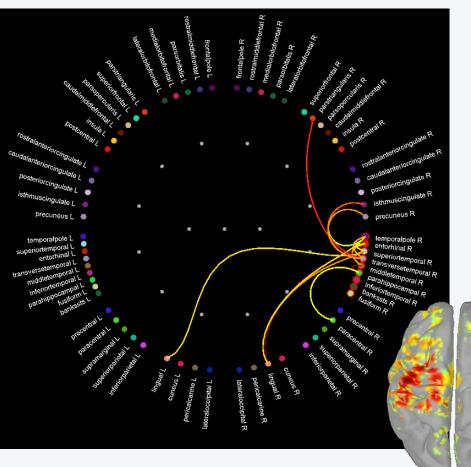
Anatomy
Link recordings
MRI registration

PSD
Filters
Bad channels
Artifacts
Correction
Bad segments

Markers
Epoching
Averaging
Sources
Time-frequency

Other measures

Connectivity measures



- Correlation
- Coherence
- Phase locking value
- Granger causality





Anatomy
Link recordings
MRI registration

PSD
Filters
Bad channels
Artifacts
Correction
Bad segments

Markers
Epoching
Averaging

Sources
Time-frequency

Project to template (sources only)



Normalize (TF and sources)



Low-pass 32Hz



Subject averages



Weighted avg

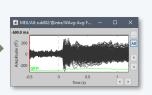
Run averages



Group averages

Statistical tests





N subjects x M runs

N averages



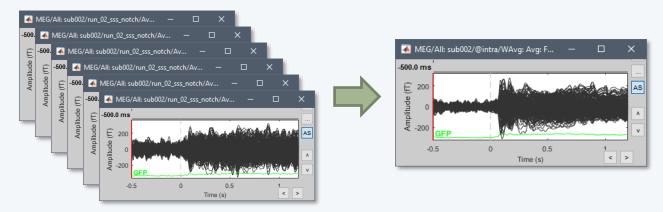
Subject averages

Low-pass Normalize Project

Group averages
Group statistics

Quality control Workflow

 Weighted average of 6 runs per subject (recordings, sources MEG/EEG, time-frequency)



- Sources: Compute within-subject differences
 - (Faces Scrambled) and (Famous Unfamiliar)
 - The sign of the MNE source amplitude is ambiguous, we will apply an absolute value before comparing between subjects



Subject averages

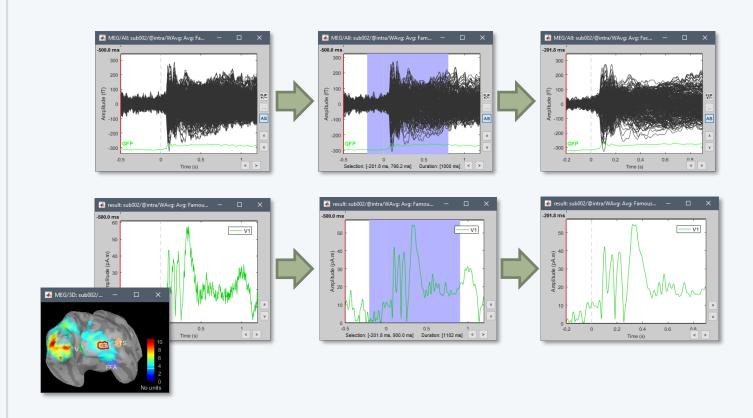
Low-pass

Normalize Project

Group averages
Group statistics

Quality control Workflow

- Low-pass filter the subject averages: 32Hz
- Remove 300ms on each side (edge effects)



Subject averages Low-pass

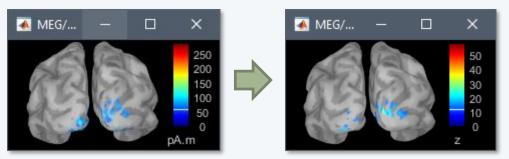
Normalize

Project

Group averages
Group statistics

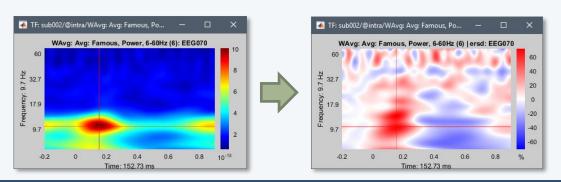
Quality control Workflow

- Amplitude normalization before group analysis
 Baseline = [-200,0]ms
- Sources: Z-score normalization wrt baseline S = (S - mean(baseline)) / std(baseline)



Time-frequency: Event-related sync/desync

TF = (TF - mean(baseline)) / mean(baseline) * 100





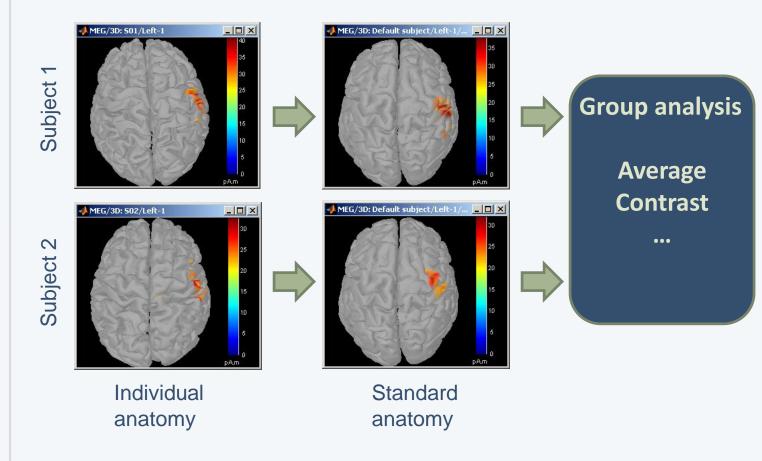
Subject averages
Low-pass
Normalize

Project

Group averages
Group statistics

Quality control Workflow

 Registration of individual sources on a template (ICBM152, Colin27, DNI, infants...)



Subject averages Low-pass Normalize

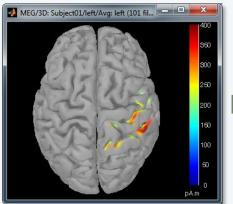
Project

Group averages
Group statistics

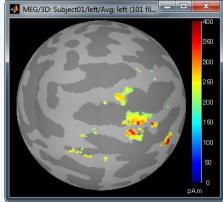
Quality control Workflow

Using FreeSurfer registration

Subject

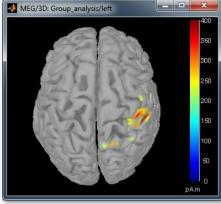


FreeSurfer registration



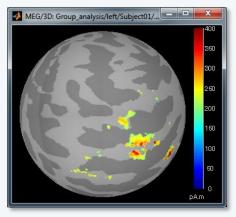
3D interpolation

Template





FreeSurfer registration



Subject averages
Low-pass
Normalize

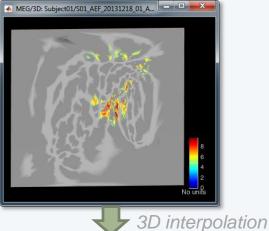
Project

Group averages
Group statistics

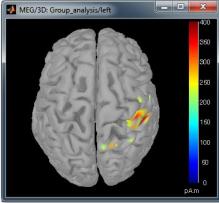
Quality control Workflow

Using BrainSuite registration



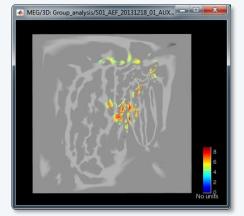


Template





BrainSuite registration



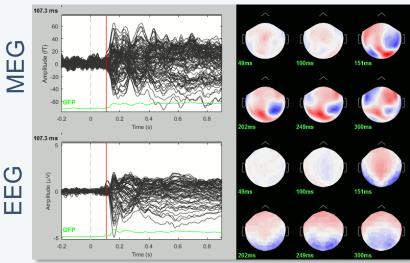
Subject averages
Low-pass
Normalize
Project

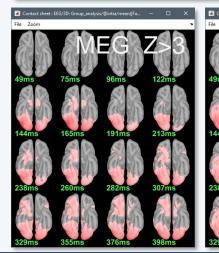
Group averages

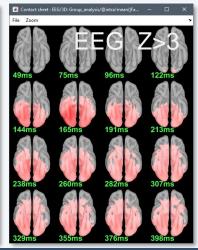
Group statistics

Quality control Workflow

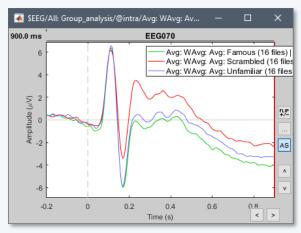
Faces - Scrambled

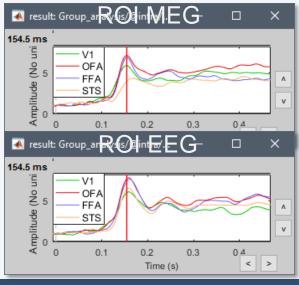






EEG 070







EEG

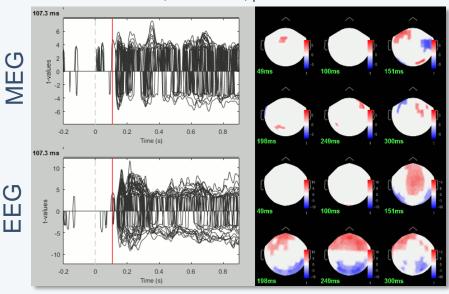
Subject averages
Low-pass
Normalize
Project

Group averages
Group statistics

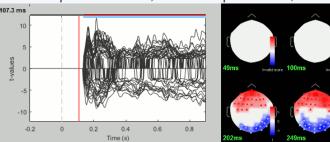
Quality control Workflow

Faces - Scrambled

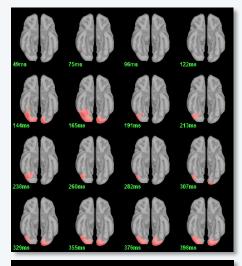
Permutation t-test, 1000 rand, p<0.05 FDR-corrected

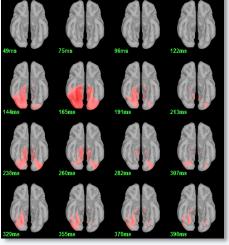


FieldTrip cluster test, cluster alpha=0.05, 1000 rand



Parametric Chi2-test log(|Faces-Scrambled|)=0 p<0.05 FDR-corrected





EG

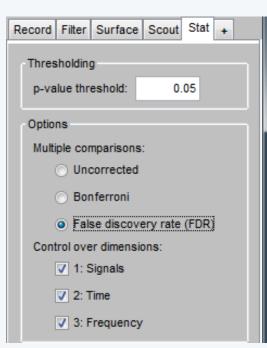


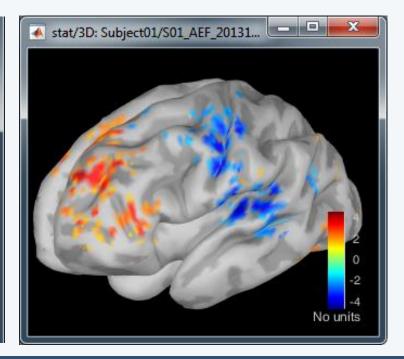
Subject averages
Low-pass
Normalize
Project

Group averages Group statistics

Quality control Workflow

- Contrasts between subjects or conditions
- Parametric t-test
- Cluster-based non-parametric tests
- Export to: **SPM**, R, Excel, SPSS, Matlab...





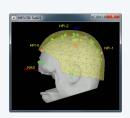


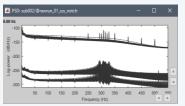
Subject averages Low-pass Normalize Project

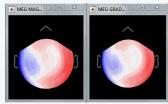
Group averages
Group statistics

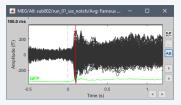
Quality control
Workflow

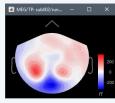
- When scripting the analysis, we recommend you always check visually the following items for each run separately:
 - MRI/sensors registration
 - PSD before and after filters
 - SSP and ICA component topographies
 - ERP/ERF: Sensors time series
 - ERP/ERF: Sensors topo of primary response
 - ERP/ERF: Sources of primary response
 - Any other metric of interest

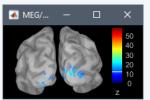


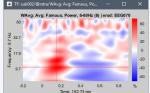












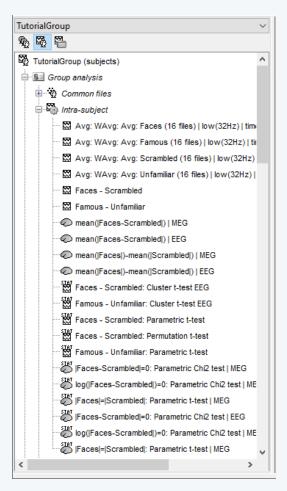


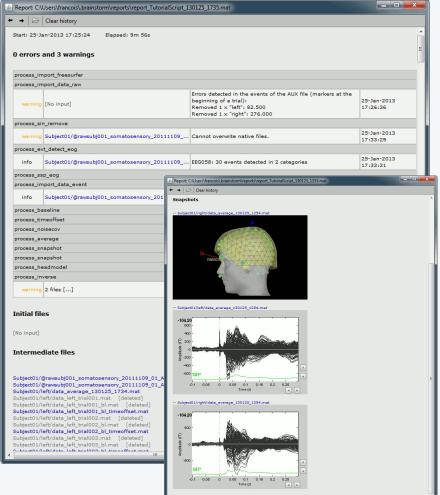
Subject averages
Low-pass
Normalize
Project

Group averages
Group statistics

Quality control
Workflow

Execution reports with snapshots saved in HTML







Subject averages
Low-pass
Normalize
Project

Group averages
Group statistics

Quality control **Workflow**

- Prototype the analysis interactively with one subject.
 Write down all the parameters to reproduce with a script.
- Set the anatomical fiducials for all the subjects.
- Script #1: Import the anatomy for all the subjects
- Script #2: Pre-processing (loop on subjects)
 - Link the raw files, register with MRI, compute PSD
 - Detect or import event markers
 - Pre-processing: Filtering, detect artifacts, SSP, ICA

Manual inspection:

- Check the execution reports for script 2
- Fix the SSP/ICA (re-compute, manual selection)
- Mark bad channels and bad segments



Subject averages
Low-pass
Normalize
Project

Group averages
Group statistics

Quality control **Workflow**

- Script #3: Subject-level analysis (loop on subjects)
 - Importing and averaging
 - Source estimation
 - Time-frequency
- Manual inspection:
 - Check the execution reports for script #3
 - Define regions of interest
- **Script #4**: Group analysis
 - Subject-level averages
 - Group-level averages
 - Statistics
 - Anything that involves regions of interest

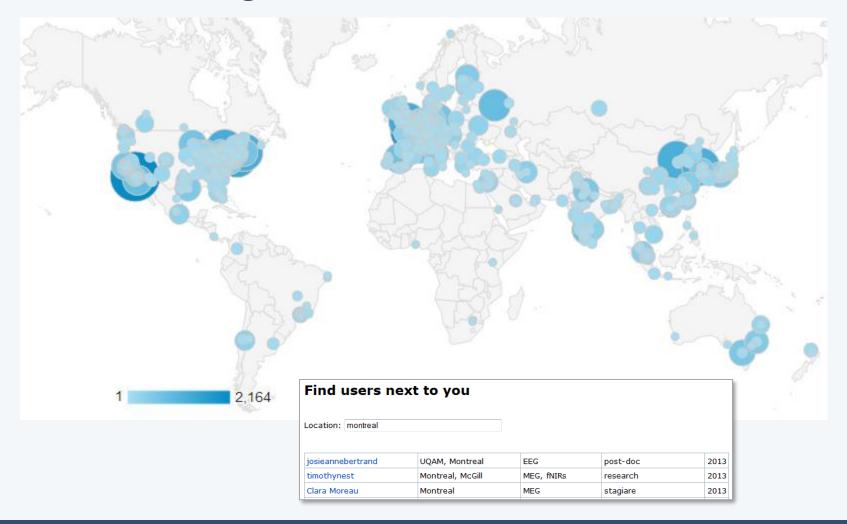


Add your code to Brainstorm

- Direct manipulation of the files in Matlab
- Use the menu "Run Matlab command"
- Write a plugin:
 - Well documented API
 - Lots of example (170 functions written as plugins)
- Examples of recent external contributions:
 - MVPA decoding (Oliva, MIT)
 - Microstate segmentation (Cacioppo, UChicago)
 - Eyetracker/EEG synchronization (Uni Freiburg)

User community

• 15,500 users registered on the website





User support

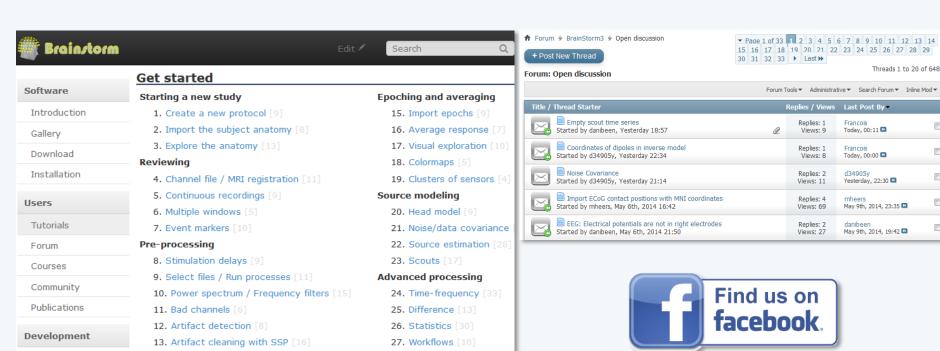
What's new

Additional bad segments [7

30-hour self-training program Online tutorials:

Active user forum: 150 posts/month

750 downloads/month Daily updates:





28. Scripting [31]

Contributors

Investigators

And...



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Richard Leahy USC



John Mosher Cleveland Clinic



François Tadel Inserm Grenoble



Martin Cousineau *MNI*

Matti Hamalainen Ghislaine Dehaene Antoine Ducorps Denis Schwartz

EG @ McGill



Elizabeth Bock MEG engineer



Guiomar Niso Post-doc



Soheila Samiee PhD student



Jeremy Moreau *PhD student*



Peter Donhauser PhD student



Dimitrios Pantazis *MIT*



collaborators

Alexandre Gramfort Telecom / Neurospin



Anand Joshi USC



Hossein Shahabi *USC*



Esther Florin
Univ Hosp Cologne



Anne-Sophie Dubarry Aix-Marseille Univ



Rey Ramirez *UW*



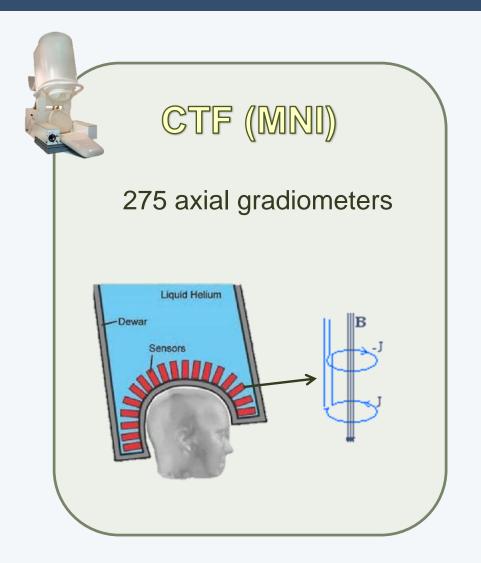
Sample data

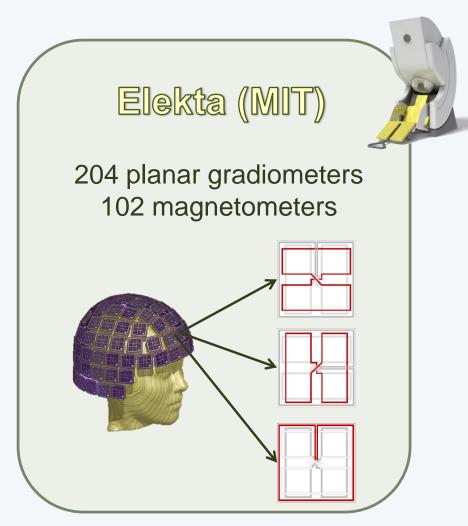
Oddball auditory task (Dec 2013, MNI, McGill)

- Binaural stimulation with intra-aural earphones
- 200 standard beeps (400Hz) + 40 deviant (554Hz)
- Inter-stimulus interval: Random in [0.7 1.7] s
- Subject taps the right index when a deviant is heard
- Acquisition at 2400 Hz Downsampled at 600Hz
- Recorded on CTF 275 MEG sensors
 - + 2 EEG (Pz,Cz) + 26 MEG reference sensors
 - + EOG + ECG + STIM + ... = 302 channels
- 6 minutes of recordings
- MRI processed with FreeSurfer 5.3

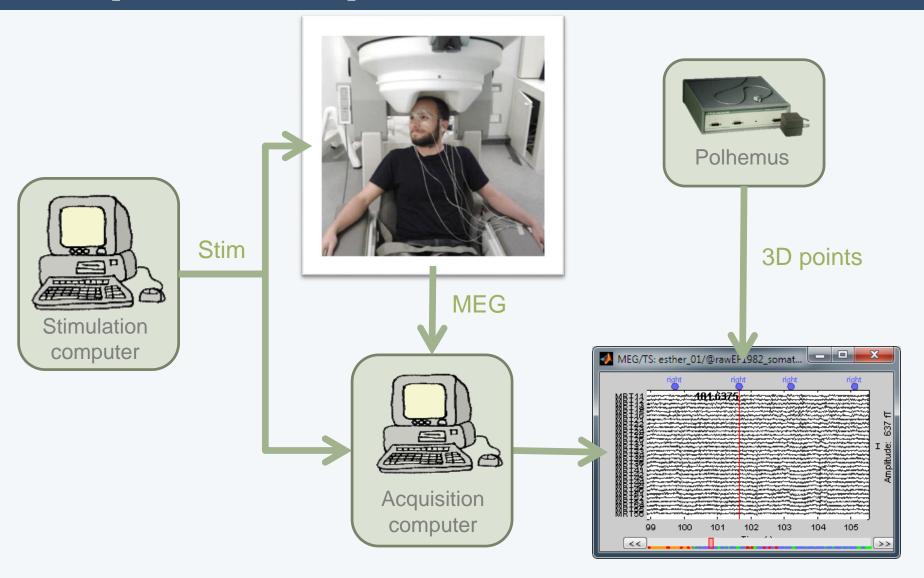


MEG sensors





Acquisition setup



Morning

- Creation of a new protocol, with one subject
- Preparation of the anatomy (MRI, surfaces)
- Anatomical atlases
- Co-registration MRI / MEG
- Reviewing the continuous file
- Correcting for artifacts with SSP
- Epoching and averaging

Afternoon

- Source estimation
- Regions of interest (scouts)
- Time-frequency decomposition
- Phase-amplitude coupling
- Connectivity
- Scripting interface
- Group analysis and statistics
- Registration on default anatomy

