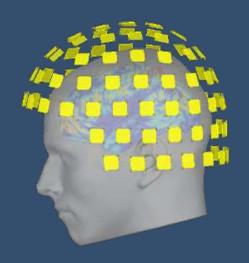
THI1

Username : x-hbpvisitor

Password: nitfes96

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







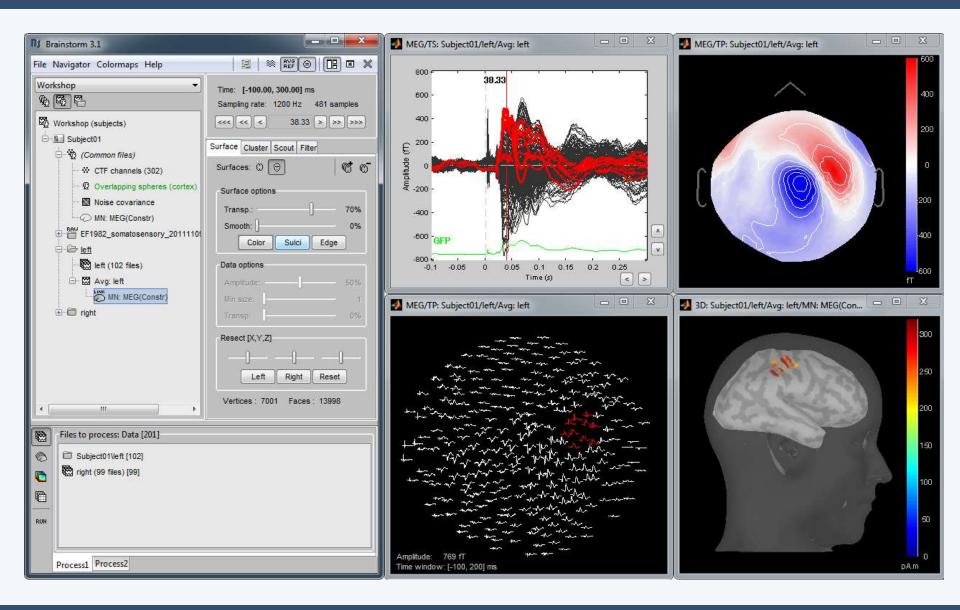




Geneva, December 2015

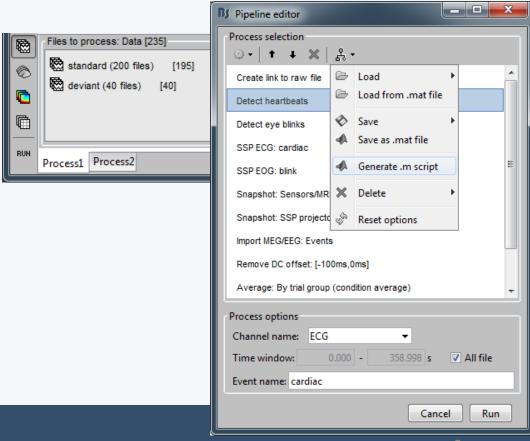
Francois Tadel Sylvain Baillet Anne-Sophie Dubarry

#### 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', ...
              'SubjectO1\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);
```



#### Brainstorm is...

A free and open-source application (GPL)







Designed for Matlab environment



Stand-alone version also available



Interface-based: click, drag, drop

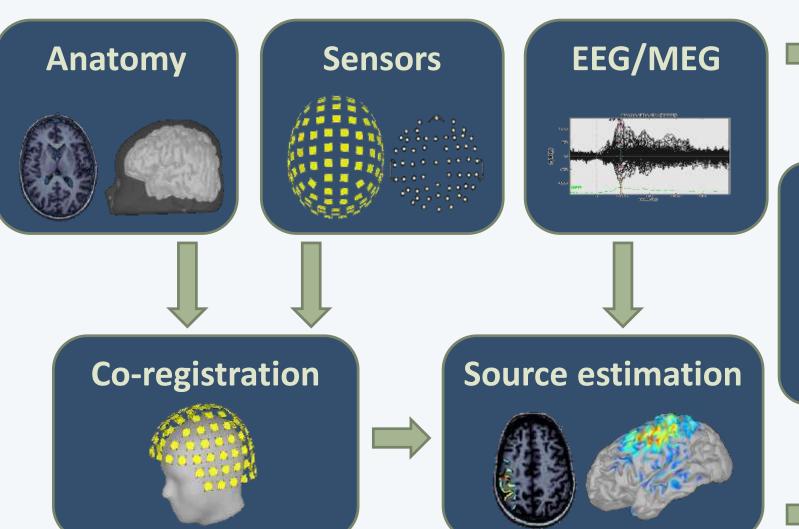


No Matlab experience required



- Daily updates of the software
- Supports most common file formats

#### Workflow





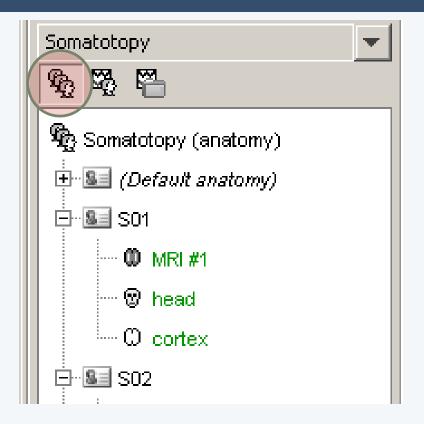
#### Analysis

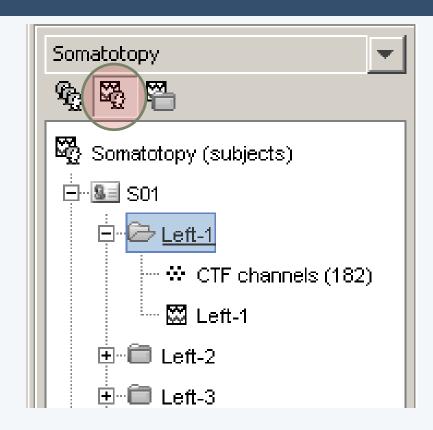
Averages
Contrasts
Group analysis
Time-frequency
Connectivity





#### Database





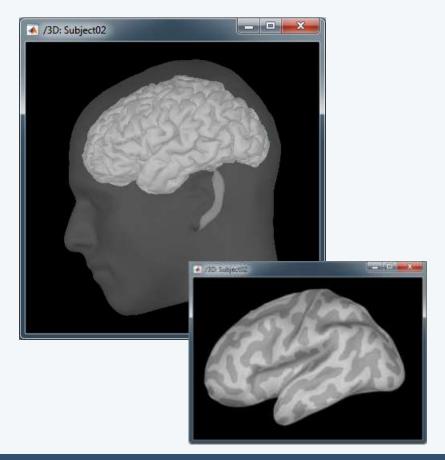
- Three levels:
  - Protocol
  - Subject
  - Condition

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

#### Anatomy

 One-click import of the T1 segmentation: FreeSurfer, BrainSuite, BrainVISA, CIVET

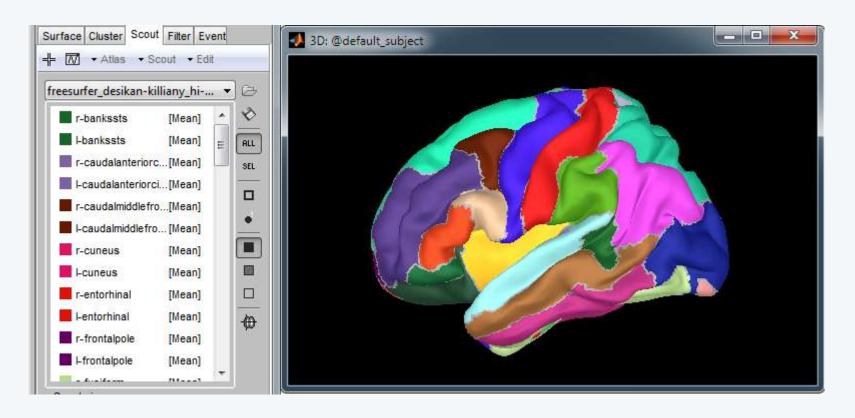






#### Atlases

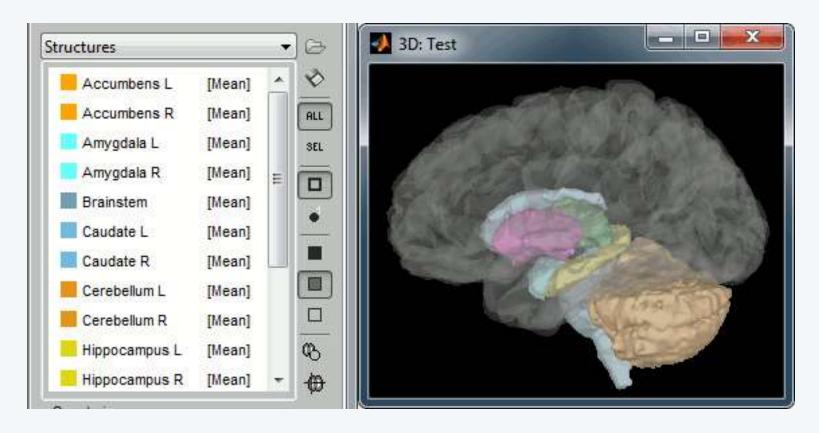
 Support for the surface-based atlases generated automatically by FreeSurfer and BrainSuite





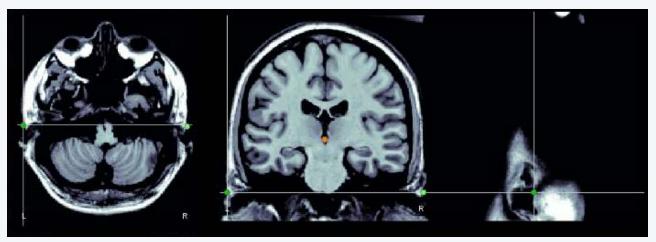
#### Atlases

 Support for the subcortical atlases generated automatically by FreeSurfer



#### Co-registration MEG / MRI (I)

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

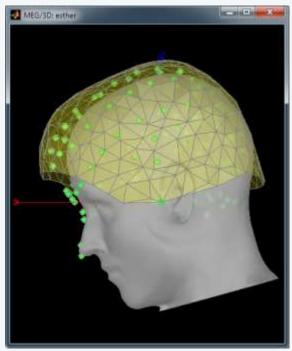


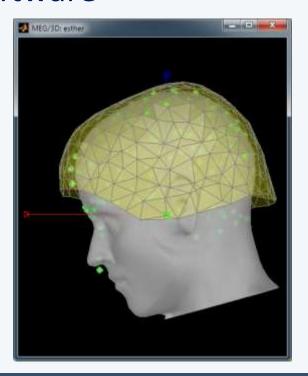


#### Co-registration MEG / MRI (2)

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

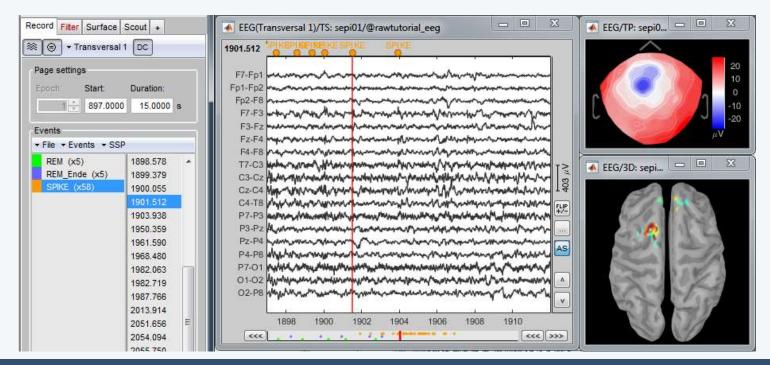






#### Continuous recordings

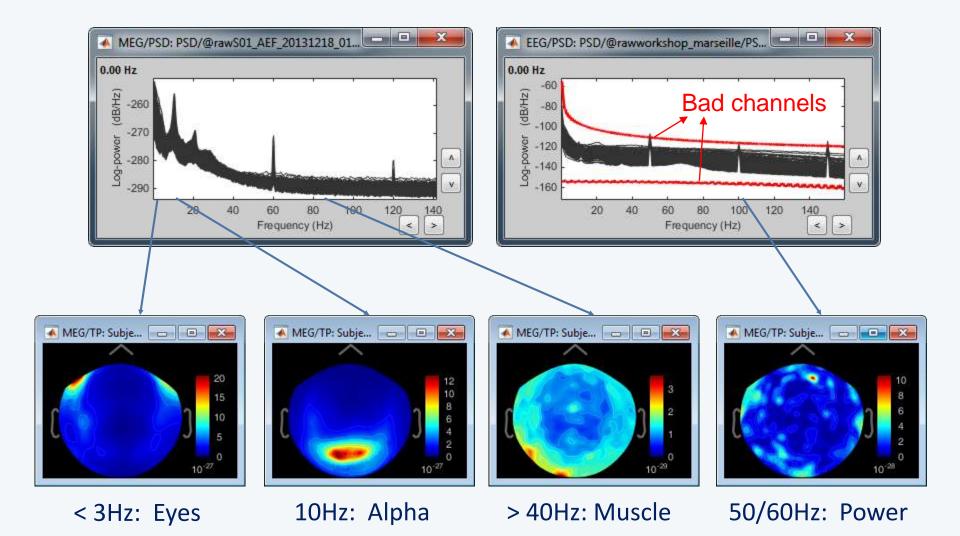
- Manual inspection of the recordings
- Identify noise sources, mark bad segments
- Check stimulus markers, add custom events
- Optimized workflow for clinicians (shortcuts, workspace...)



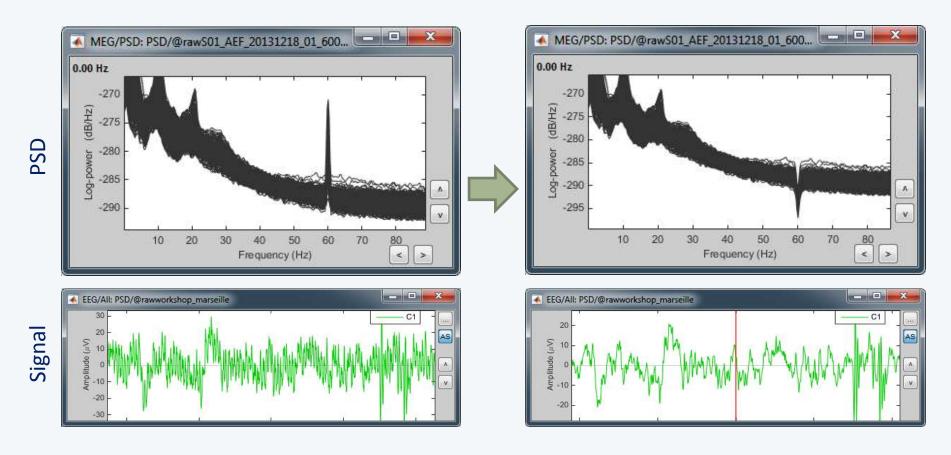


#### Quality control

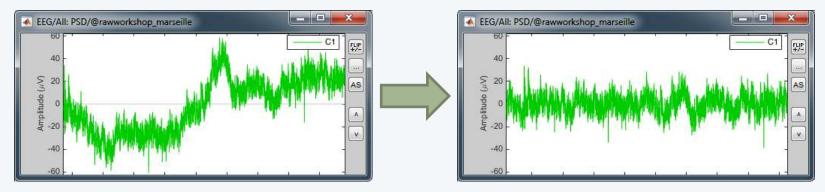
### Spectral density



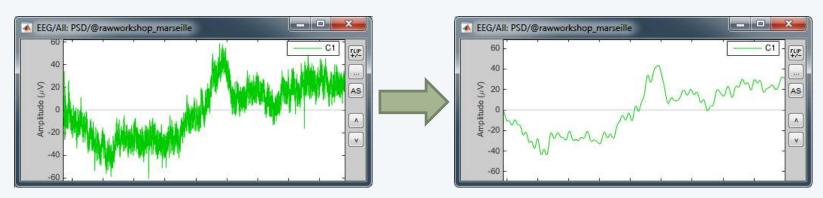
Notch filter
 Removes 50Hz/60Hz power line noise (and harmonics)



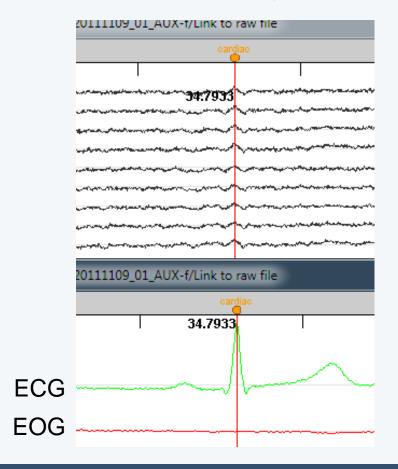
 High-pass filter: Removes slow components (eye movements, breathing, sensor drifts...)

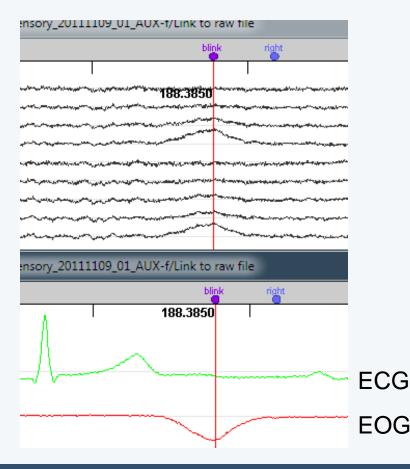


Low-pass filter: Remove high-frequencies



- Artifact detection and removal:
  - heartbeats, eye blinks, movements, ...



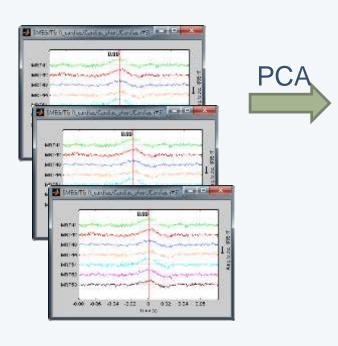




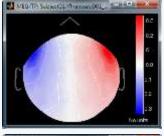
#### Signal-Space Projection (SSP)

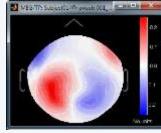
Detect artifacts

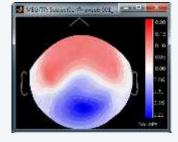
Concatenate epochs



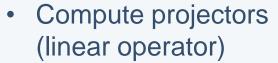
Spatial components







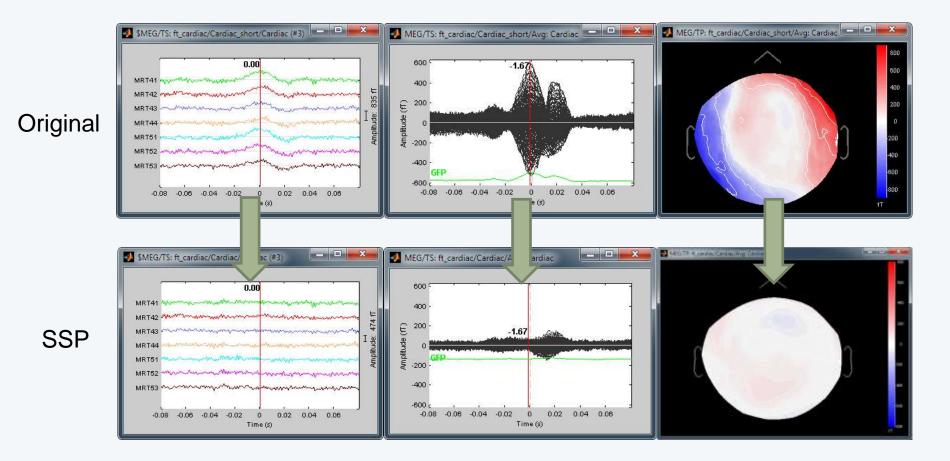
Select components

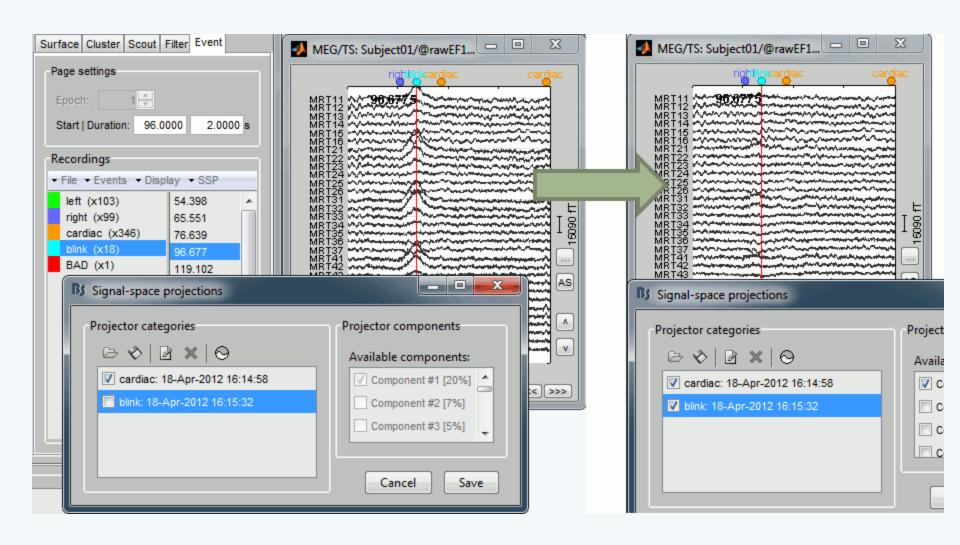


Apply to EEG/MEG

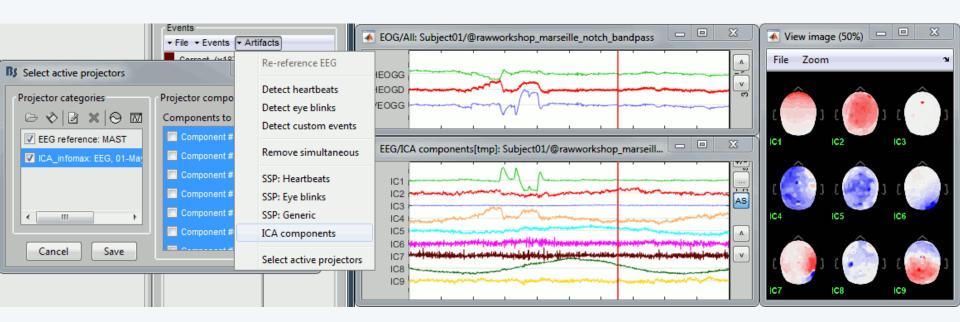


Example: Cardiac artifact





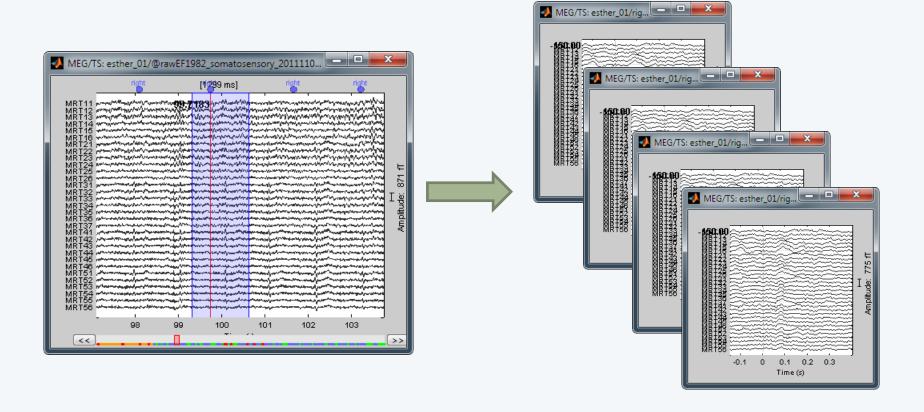
- Independent component analysis (ICA):
  - Popular in the EEG literature
  - Alternative to SSP for low number of sensors
  - Already implemented: Infomax and JADE (EEGLAB)





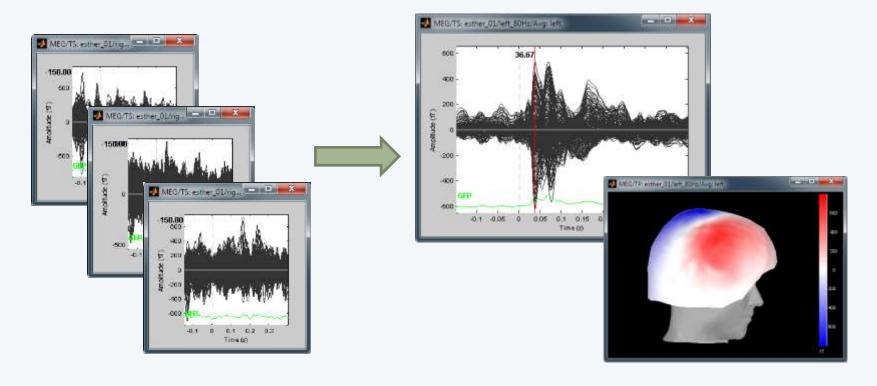
#### Epoching

 Epoching: extraction of small blocks of recordings around an event of interest (stimulus, spike...)



#### Averaging

- Averaging all the trials: Reveals the features of the signals that are locked in time to a given event
  - => Event-related field (or potential) = Evoked response

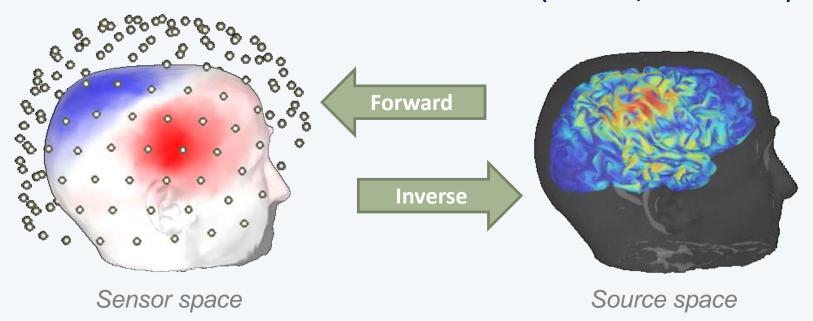


#### Source estimation

Source space: Cortex surface or full head volume

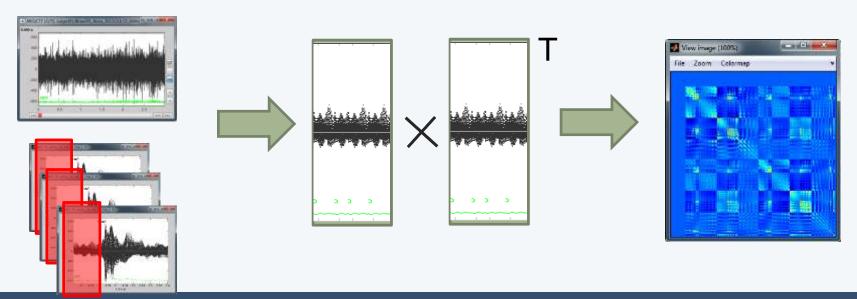
Forward model: Overlapping spheres (MEG)
 OpenMEEG BEM (EEG)

Inverse model: Minimum norm estimates
 + normalizations (dSPM, sLORETA)

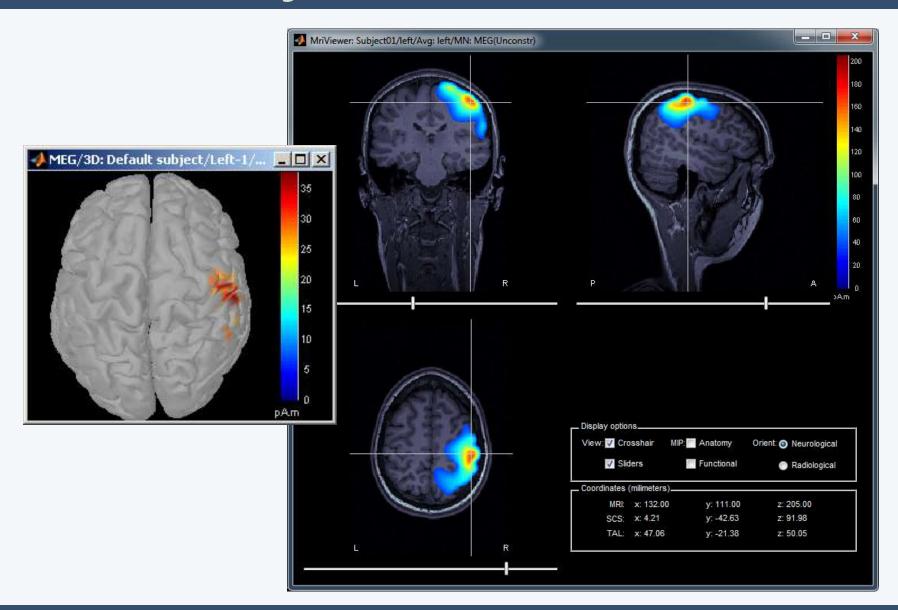


#### Noise covariance matrix

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



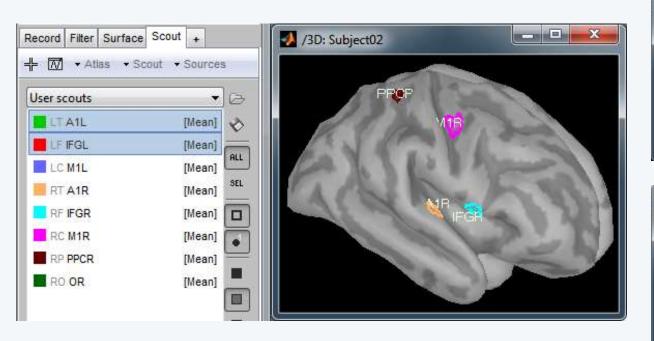
#### Source activity

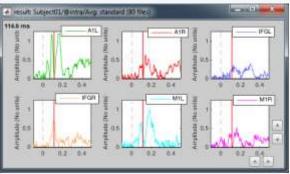


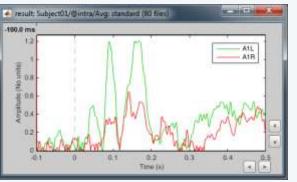


### Regions of interest

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



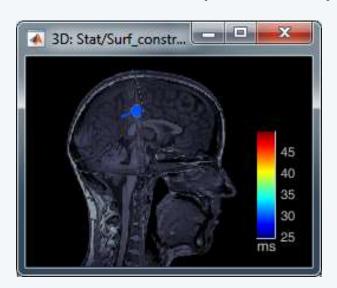


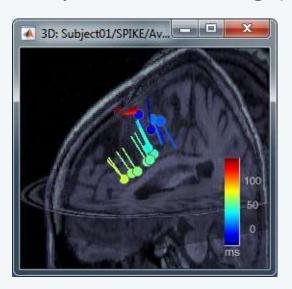




#### Single dipoles

- 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 (difference data explained by the dipole - recordings)

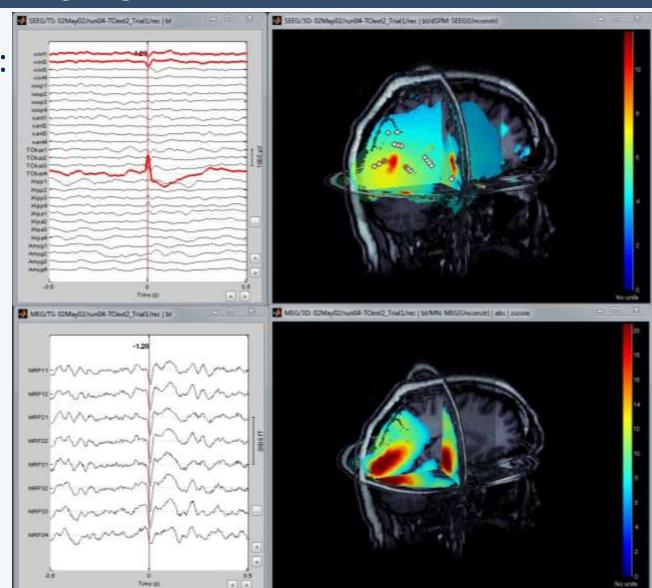




### Multi-modal imaging

#### Easy integration of:

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

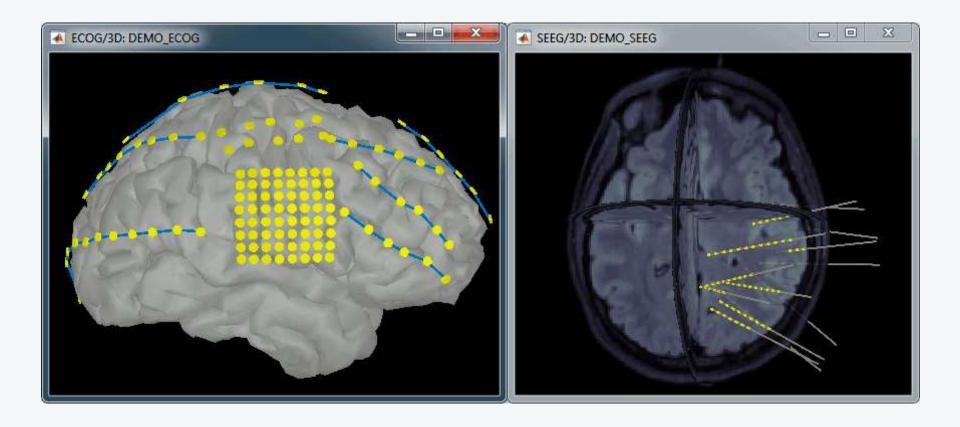




### Invasive recordings

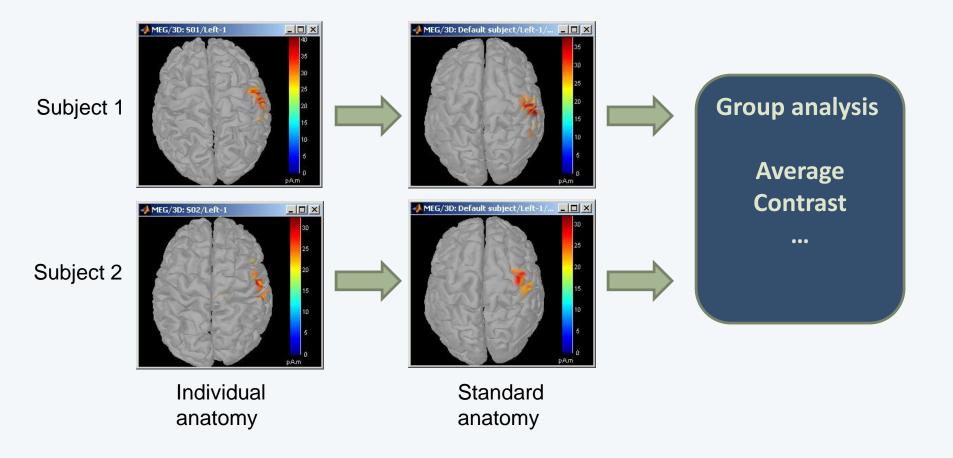
#### **ECoG**

#### Depth electrodes



#### Group analysis

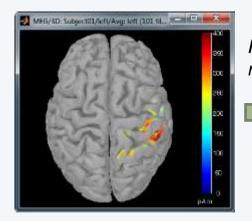
Registration of individual brains on a template



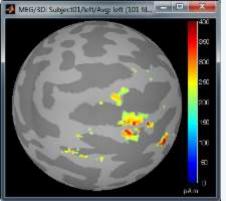
#### Group analysis

#### freeSurfer

Subject



FreeSurfer registration

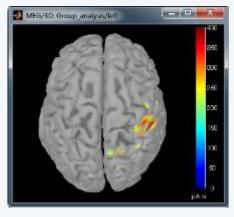


Subject anatomy Right hemisphere

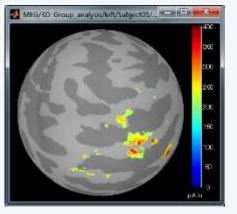


Shepard interpolation

Default anatomy



FreeSurfer registration

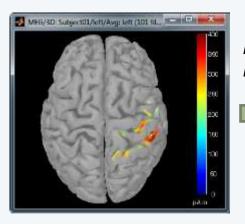


Template Right hemisphere

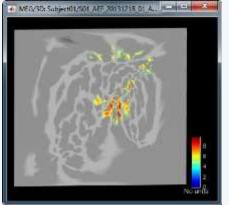
#### Group analysis

#### BrainSuite

Subject



BrainSuite registration

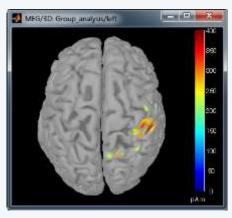


Subject anatomy Right hemisphere



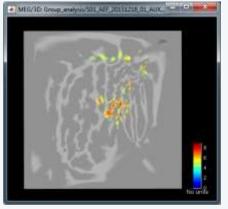
Shepard interpolation

Default anatomy





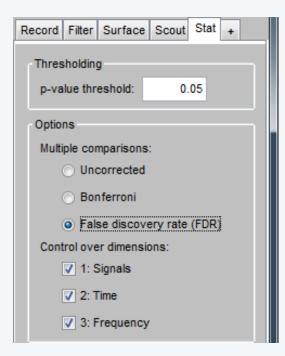
BrainSuite registration

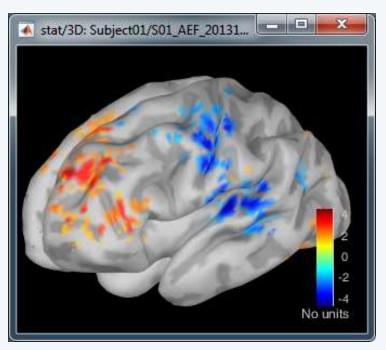


Template Right hemisphere

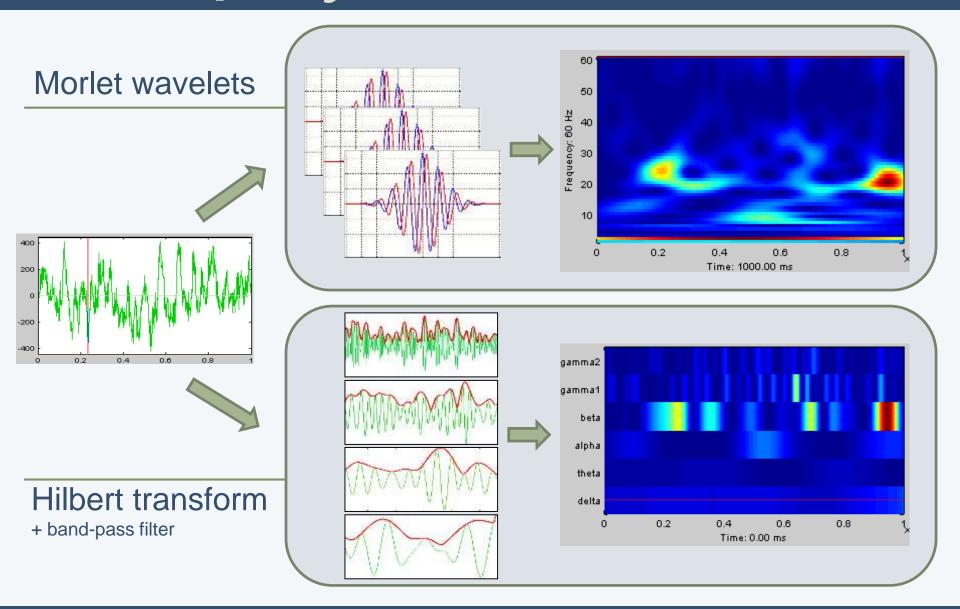
#### Statistics

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



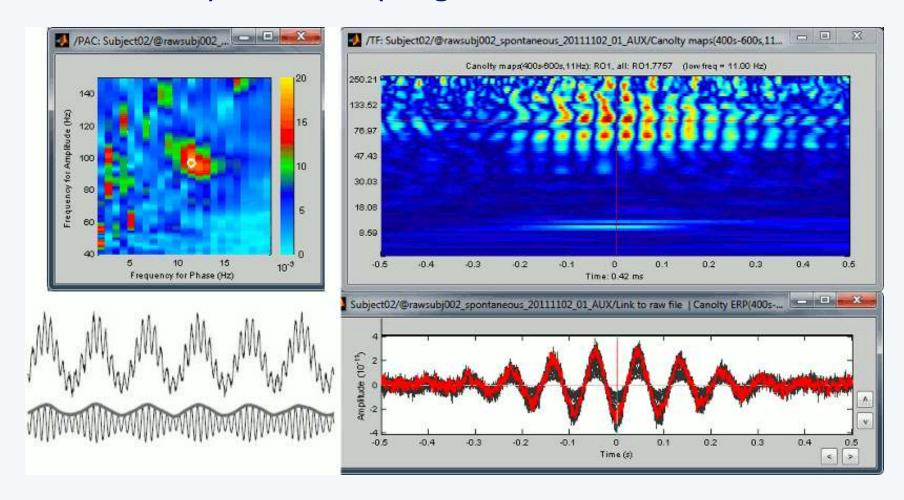


### Time-frequency



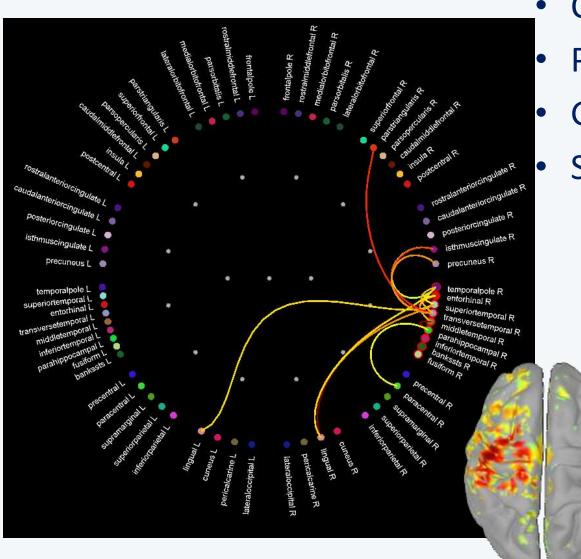
#### Cross - frequency coupling

Phase-amplitude coupling estimation





#### Connectivity



- Correlation / Coherence
- Phase locking value
- Granger causality
- Sensor or source levels



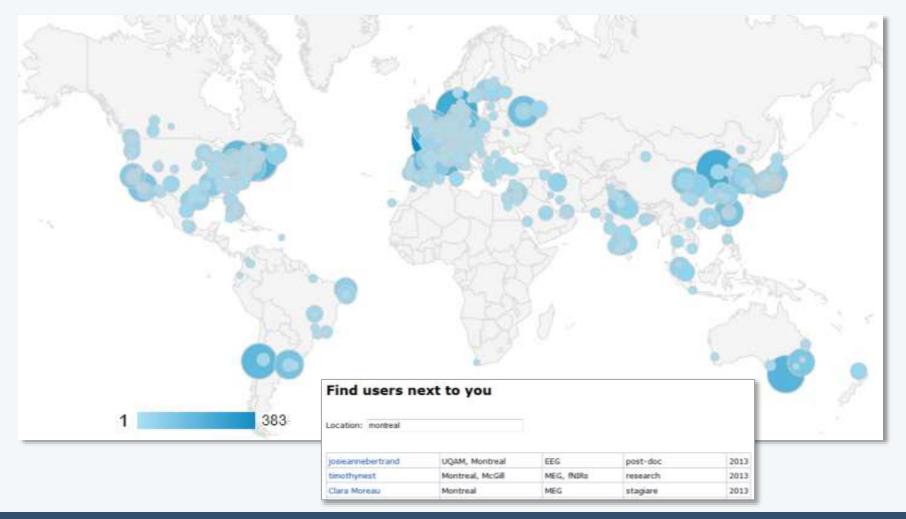
#### Add your code to Brainstorm

- 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

• 12,000 users from 70 countries registered on the website



#### User support

Online tutorials: 30-hour self-teaching program

Active user forum: 200 posts/month

Daily updates: 700 downloads/month



Contact us for specific questions and requests:
 We will help you adding the features you need



#### Contributors

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**Rey Ramirez** UW



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Sergül Aydore Columbia



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Anand Joshi USC





#### Sample data

#### **Epilepsy recordings: Localization of frontal spikes**

- Patient recorded at the Epilepsy Centre of the University Hospital in Freiburg, Germany
- Focal epilepsy with focal sensory, dyscognitive and secondarily generalized seizures since the age of eight years
- Histological analysis revealed a focal cortical dysplasia
- One hour of sleep recordings
- Neurofile NT EEG system, sampled at 256Hz
- 58 epileptic spikes marked by the epileptologists in Freiburg
- MRI processed with FreeSurfer 5.3

