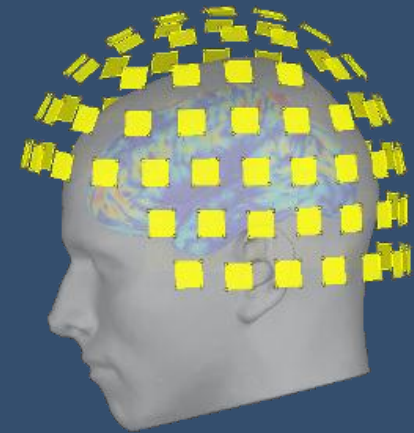


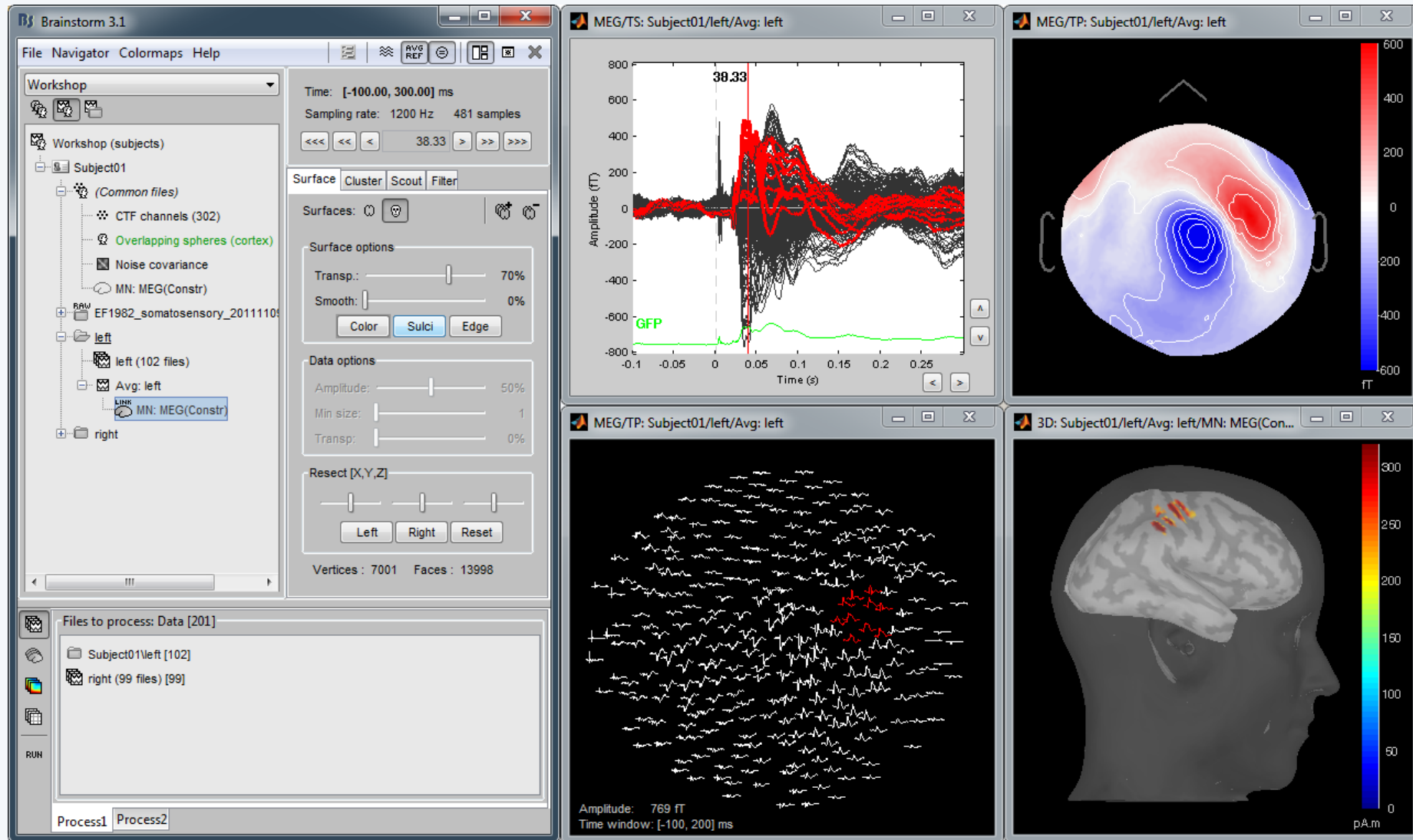
MEG and EEG analysis using

Brainstorm 3.1

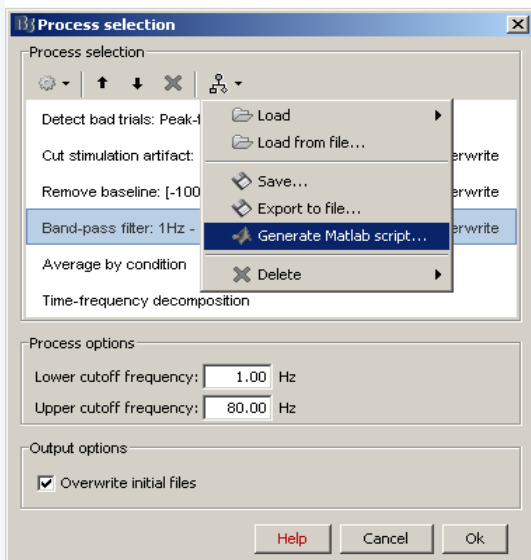
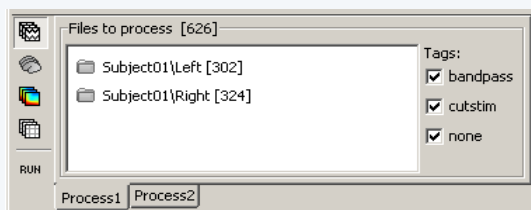
<http://neuroimage.usc.edu/brainstorm>



François Tadel
MIT – Boston – 04.28.2012



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



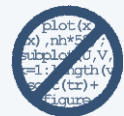
```

1 % Script generated by Brainstorm v3.1 (17-Dec-2010).
2 FileNamesA = ('Subject01\Left\data_average_101213_1558.mat', ...
3               'Subject01\Right\data_average_101213_1559.mat');
4 FileNamesB = [];
5
6 % Process: Detect bad trials: Peak-to-peak MEGGRAD(0-2000)
7 sFiles = bst_process(...
8     'CallProcess', 'process_detectbad', ...
9     FileNamesA, FileNamesB, ...
10    'timewindow', [-0.0998, 0.3000], ...
11    'meggrad', { [0, 2000], 'fT/cm (x 0.04)', 1e-015}, ...
12    'rejectmode', 2);
13
14 % Process: Remove baseline: [-100ms, -1ms]
15 sFiles = bst_process(...
16     'CallProcess', 'process_baseline', ...
17     sFiles, [], ...
18     'baseline', [-0.09983, -0.00056], ...
19     'overwrite', 1);
20
21 % Process: Band-pass filter: 1Hz - 80Hz
22 sFiles = bst_process(...
23     'CallProcess', 'process_bandpass', ...
24     sFiles, [], ...
25     'f1', 1, ...
26     'f2', 80, ...
27     'overwrite', 1);
28
29 % Process: Average by condition
30 sFiles = bst_process(...
31     'CallProcess', 'process_average', ...
32     sFiles, [], ...
33     'avgttype', 3, ...
34     'isstd', 0);

```

Brainstorm is...

- A free and open-source application (GPL)
- Matlab & Java: Platform-independent
- Designed for Matlab environment
- Stand-alone version also available
- Interface-based: click, drag, drop
- No Matlab experience required
- Daily updates of the software

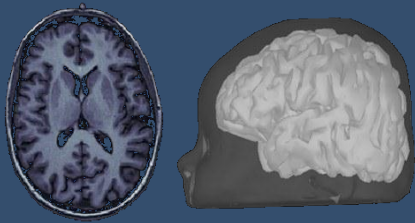


A bit of history

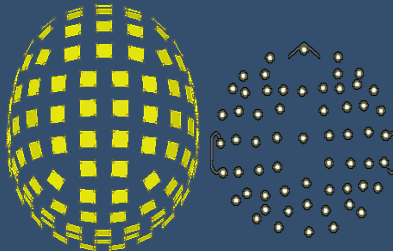
- 12 years of research and development
- Active collaboration between multiple groups:
 - University of Southern California, Los Angeles
 - La Salpetriere Hospital / CNRS, Paris
 - Neurospin / Inserm / CEA, Paris
 - Los Alamos National Lab, NM
 - Medical College of Wisconsin, Milwaukee
 - Cleveland Clinic, OH
 - Martinos Center / MGH, MA
 - McGovern Institute / MIT, MA
 - Montreal Neurological Institute / McGill, QC
- New interface released in 2009
- Over 7000 user accounts / 60 countries

Workflow

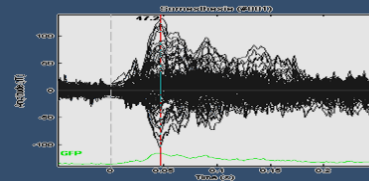
Anatomy



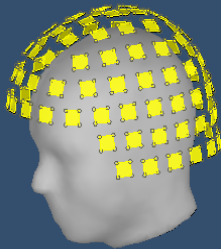
Sensors



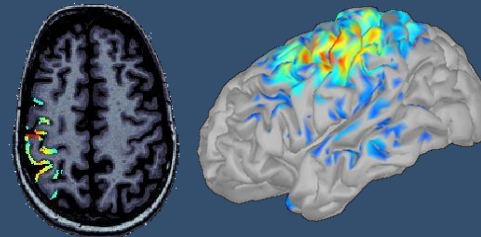
EEG/MEG



Co-registration

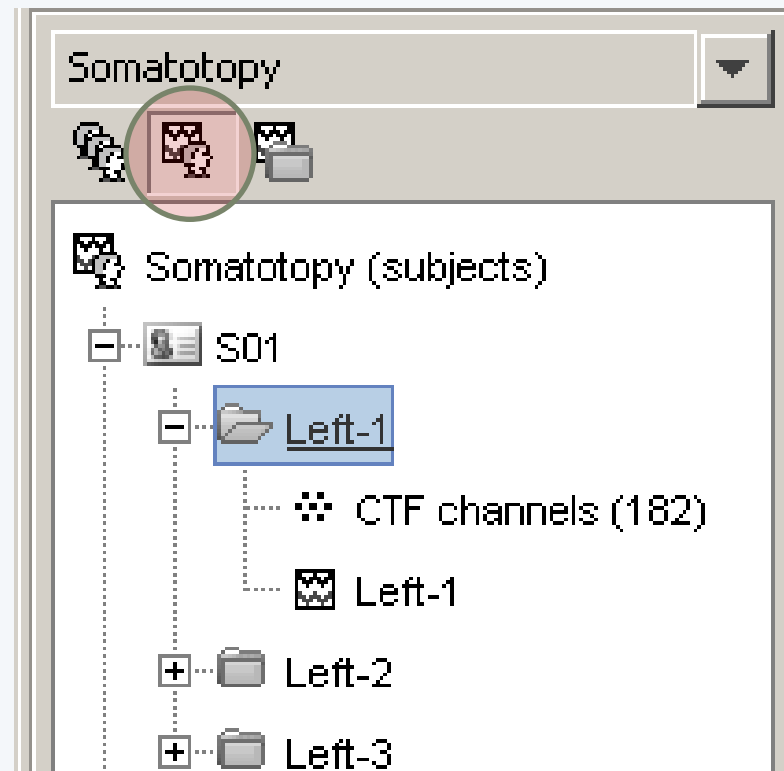
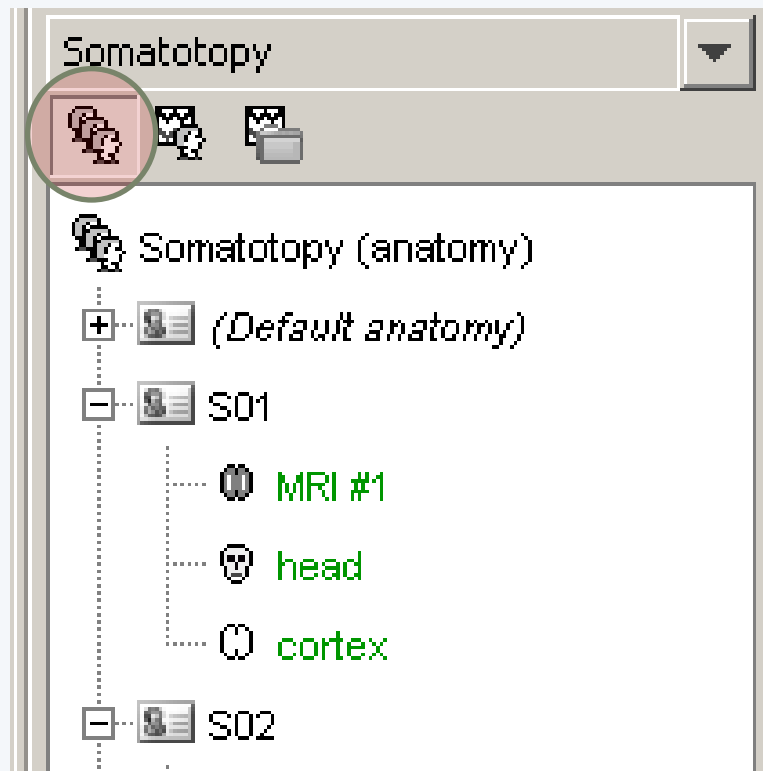


Source estimation



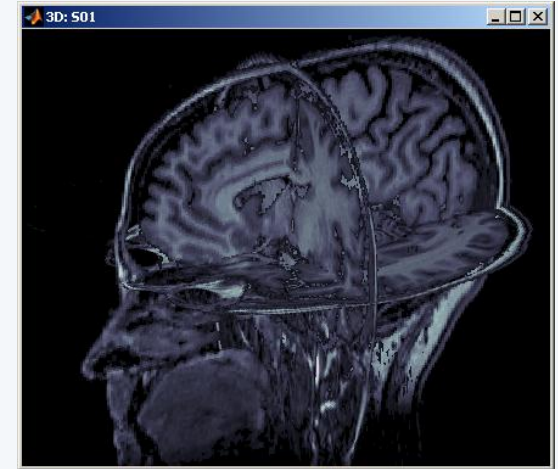
Analysis



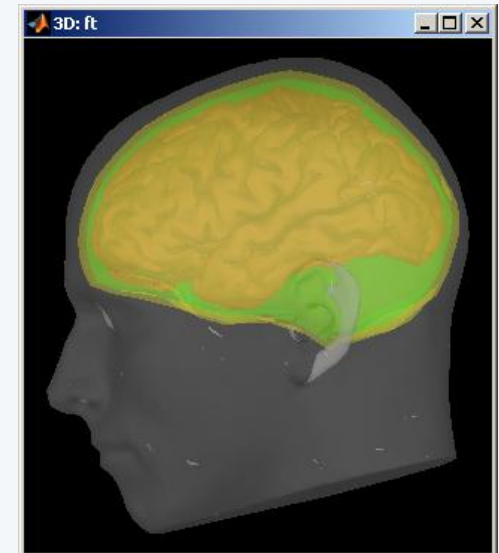
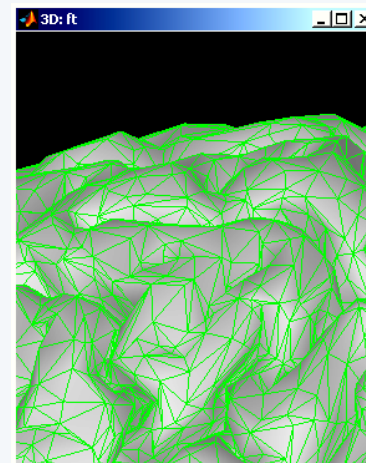
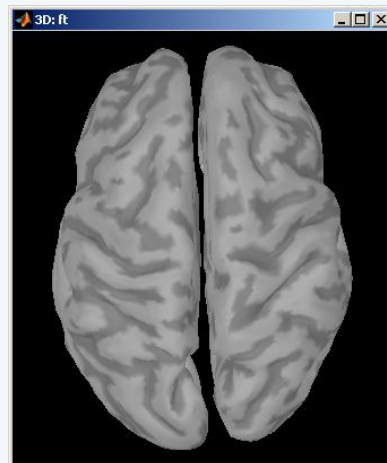


- Three levels:
 - Protocol
 - Subject
 - Condition
- Popup menus
- All files saved in Matlab .mat
- Same architecture on the disk

- T1-MRI volume

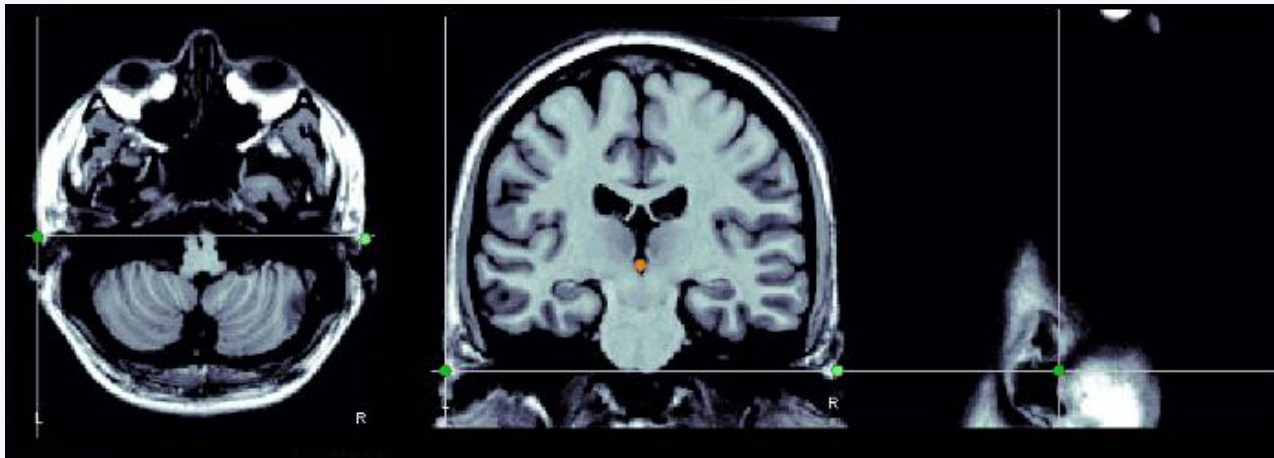


- Surfaces extracted with a dedicated software: BrainVISA, FreeSurfer, BrainSuite



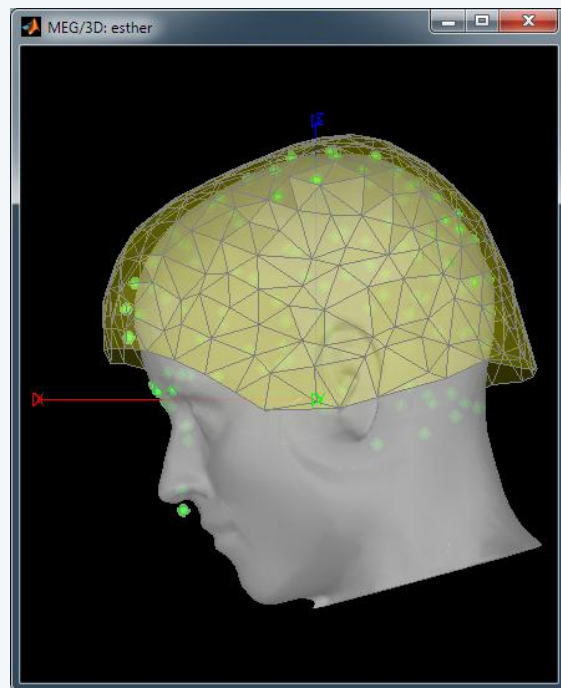
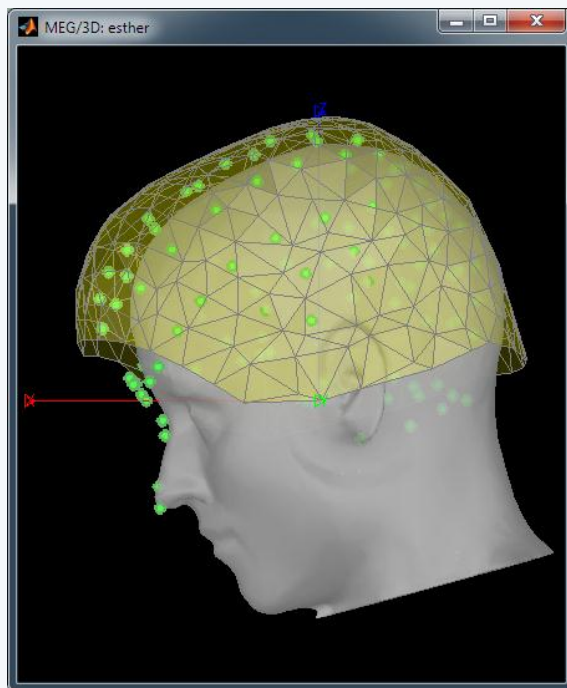
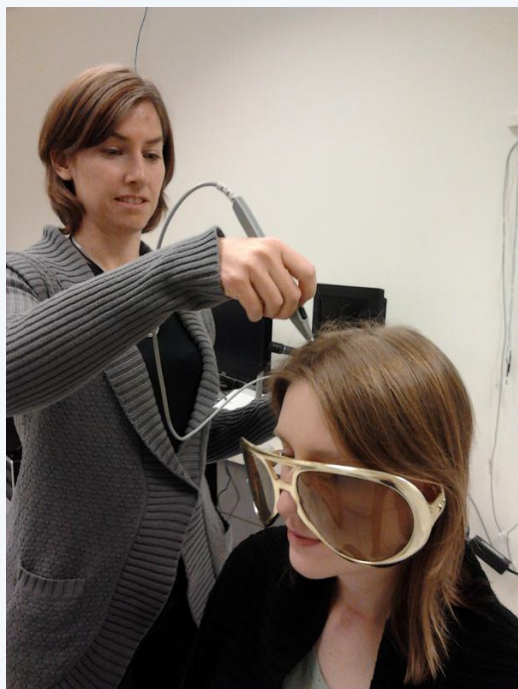
Co-registration MEG / MRI (I)

- 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)



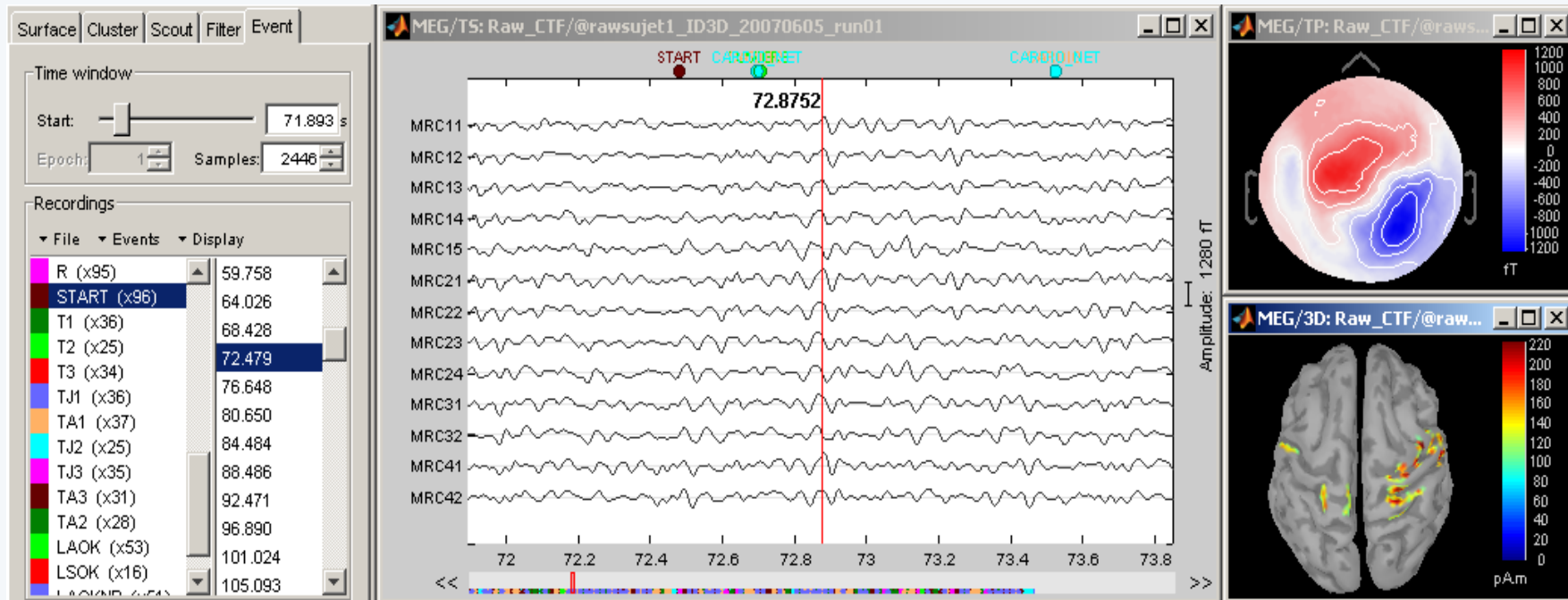
Co-registration MEG / MRI (2)

- Automatic adjustment based on head shape
 - Trying to fit the head points (digitized with the Polhemus) with the head surface (from the MRI)
- Final registration must be checked manually



Continuous recordings

- Review continuous file
- Supports most common EEG/MEG file formats
- Edit markers, display 2D projections and sources



- Frequency filtering

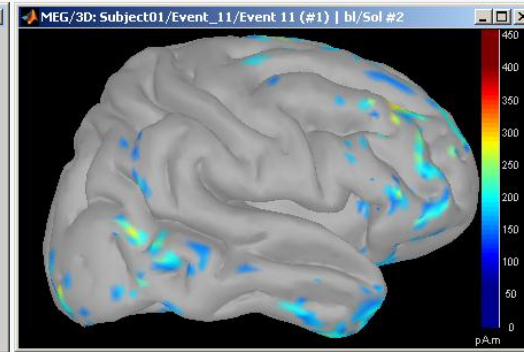
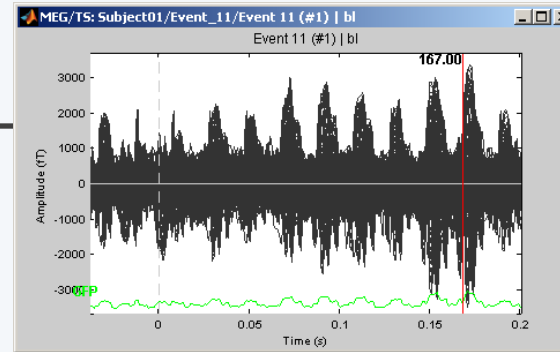
Frequency filtering

☐ High-pass: 1.00 Hz

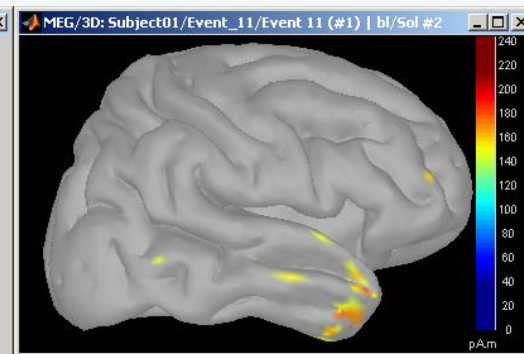
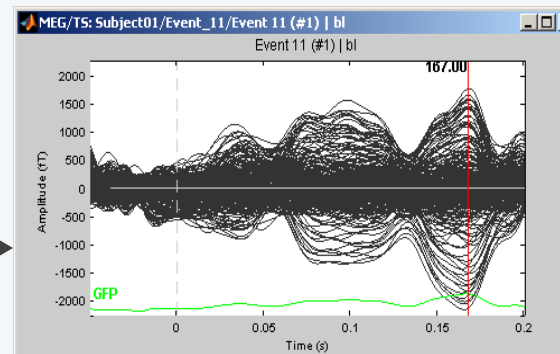
☒ Low-pass: 40.00 Hz

☐ Sin. removal: Hz

☒ Mirror signal before filtering

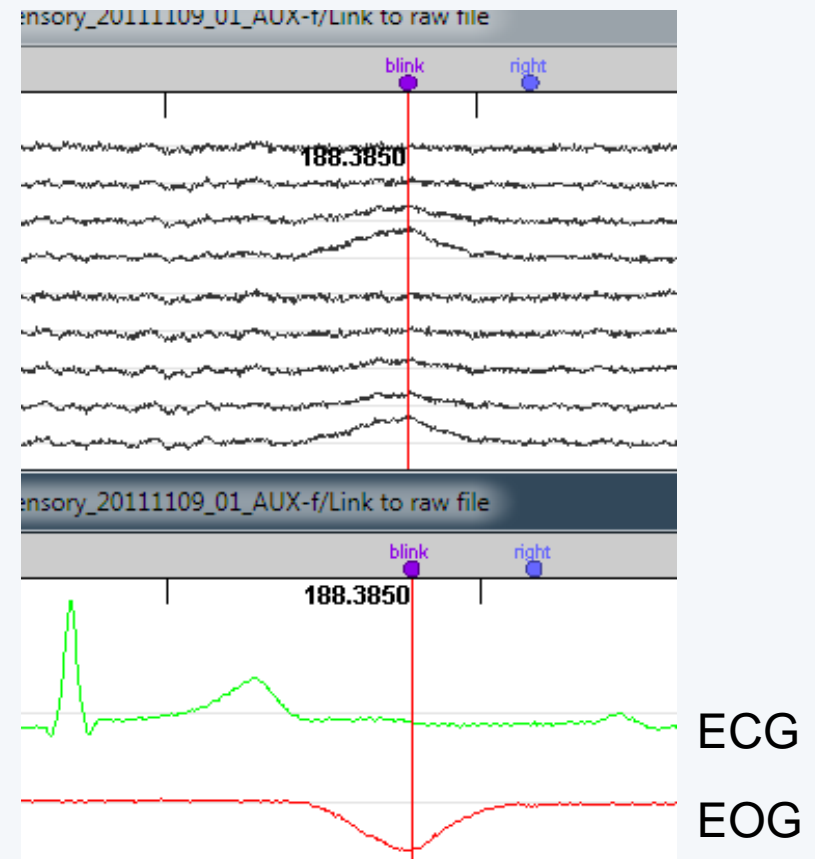
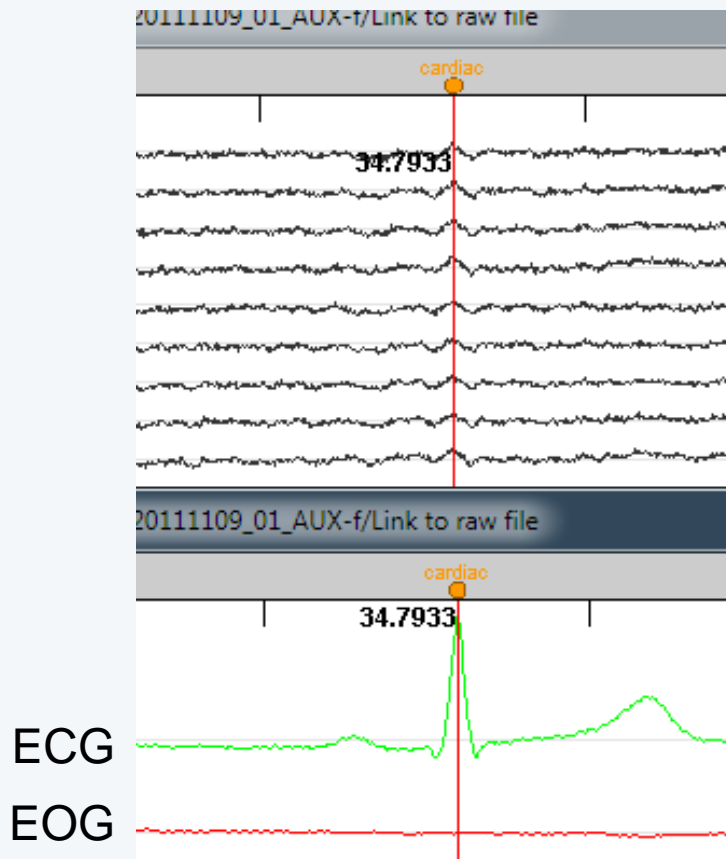


Not filtered



Low-pass
40Hz

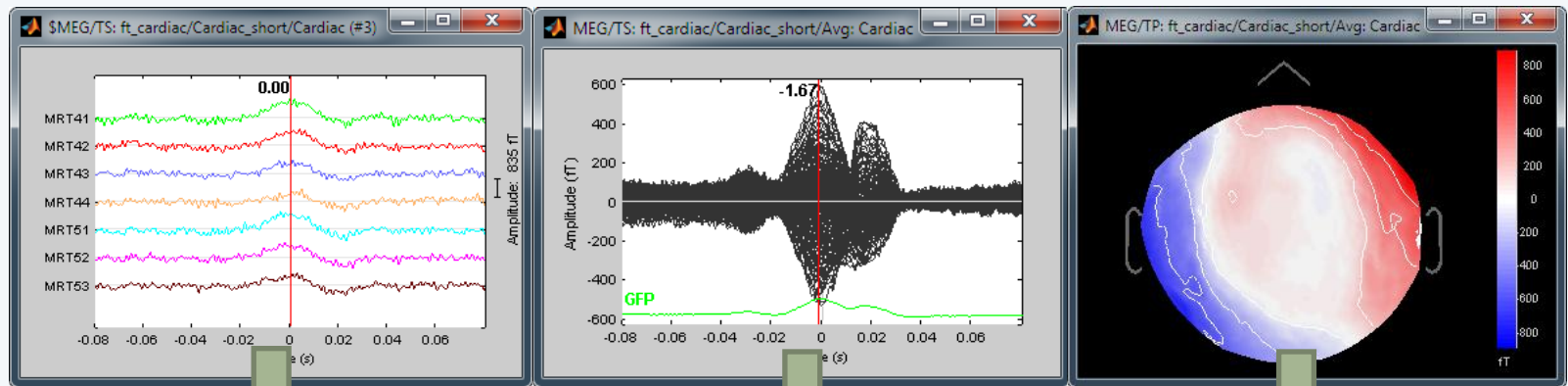
- Artifact detection and removal:
 - heartbeats, eye blinks, movements, ...



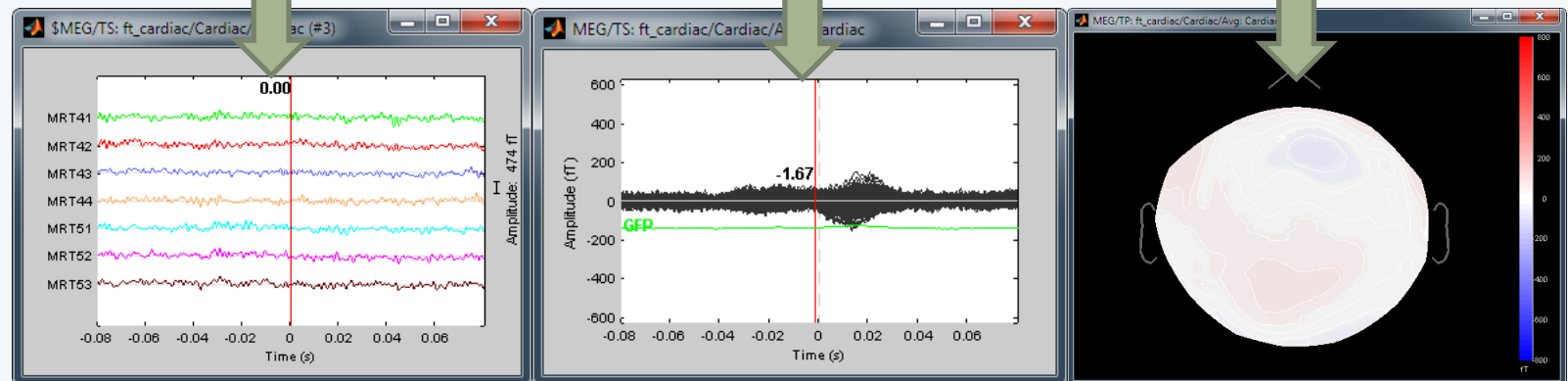
- Two categories of artifacts:
 - Well defined, reproducible, short, frequent:
 - Heartbeats, eye blinks, stimulator
 - Unavoidable and frequent: we cannot just ignore them
 - Can be modeled and removed from the signal efficiently
 - All the other events that can alter the recordings:
 - Movements, building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

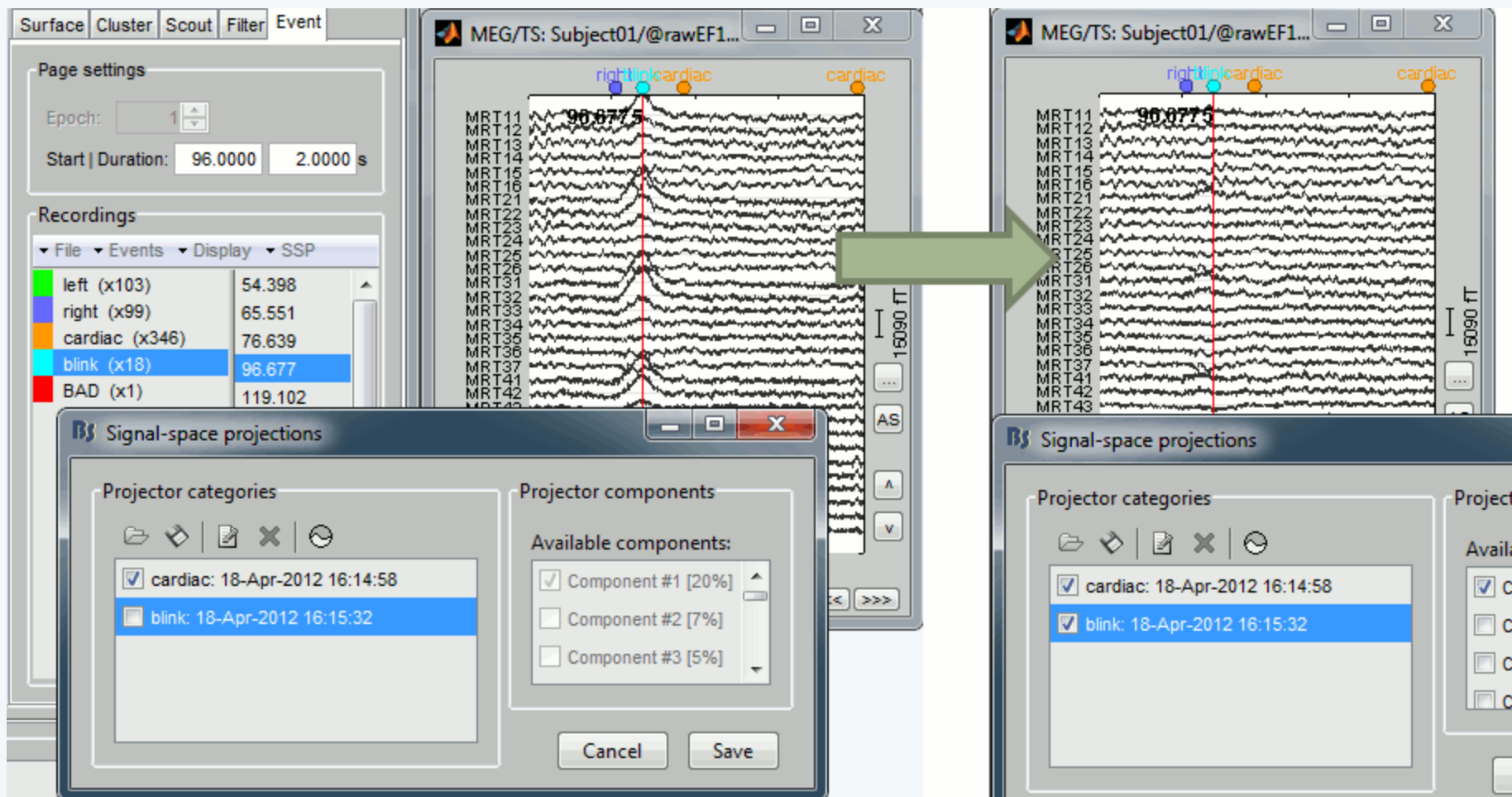
- Example of the cardiac artifact
=> Computation of a Signal-Space Projection (SSP)

Original

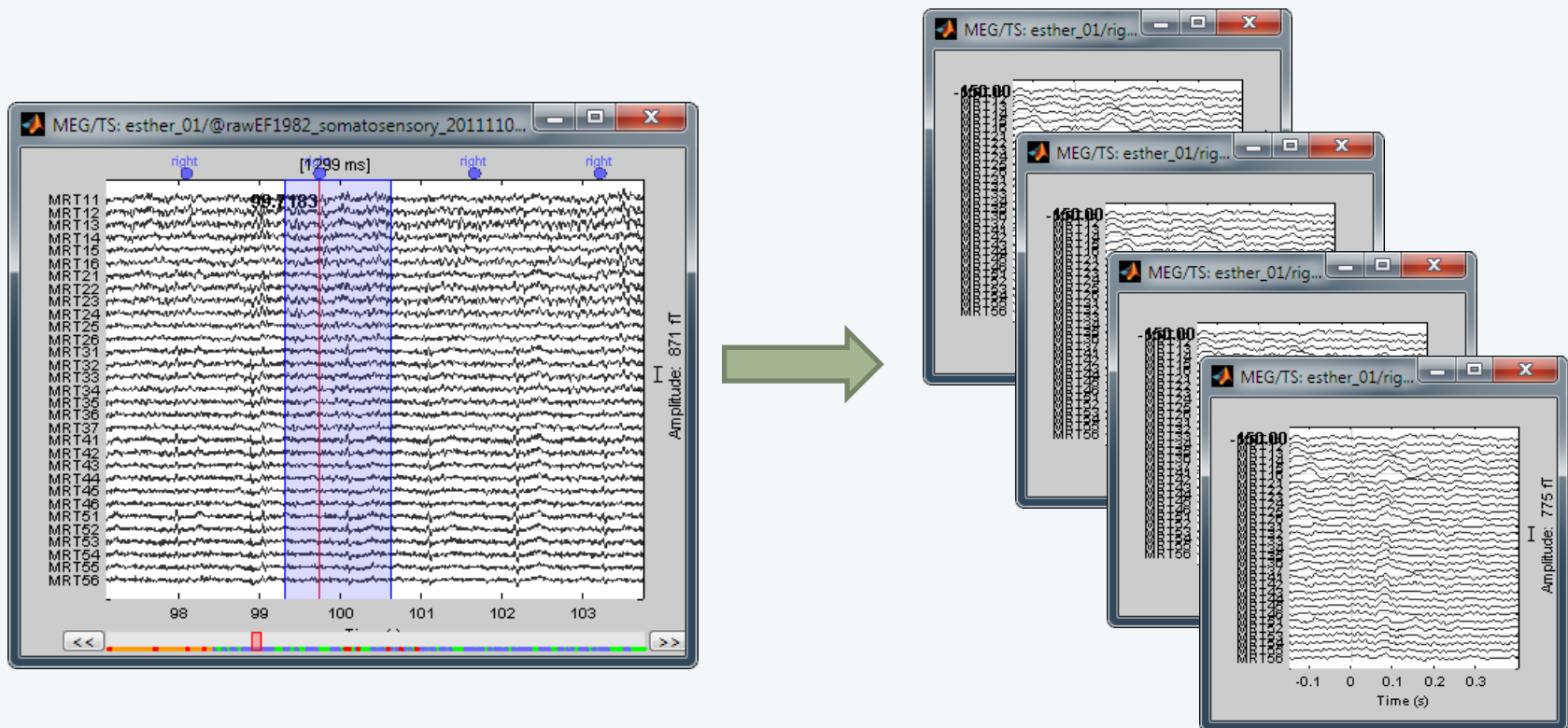


SSP

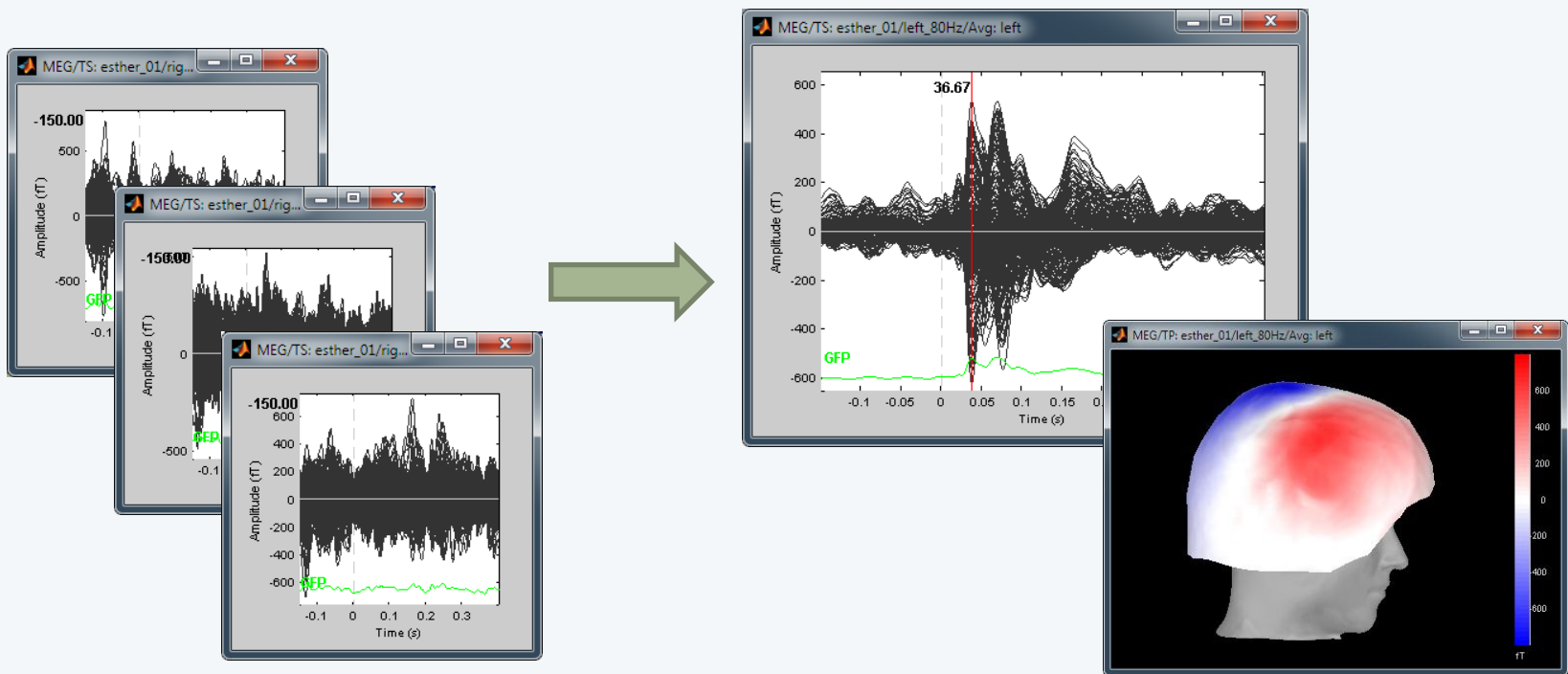




- Epoching: extraction of small blocks of recordings around an event of interest (stimulus, spike...)

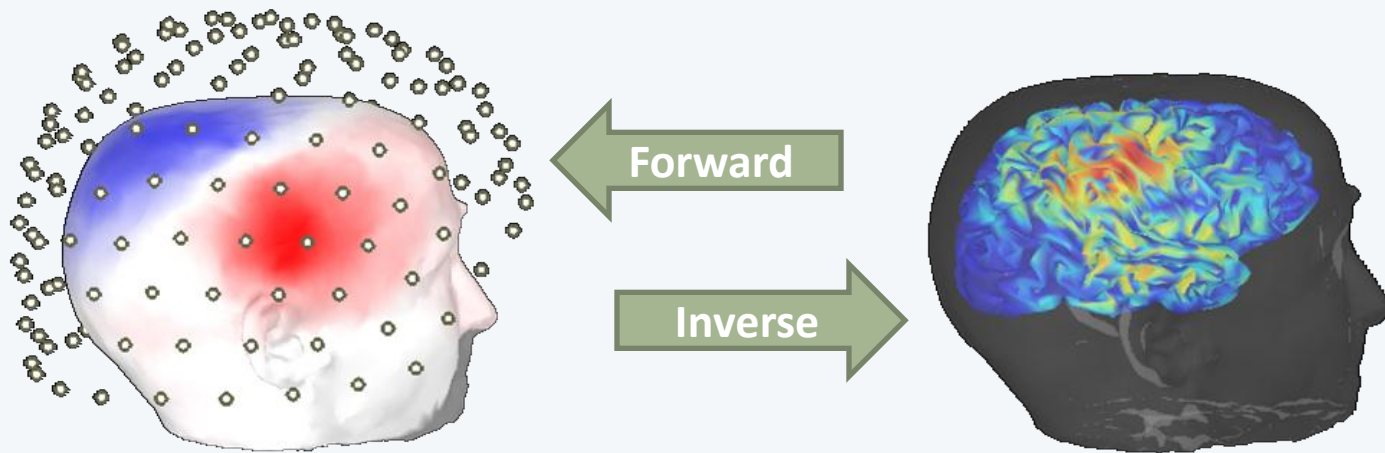


- Averaging all the trials: Reveals the features of the signals that are locked in time to a given event
=> Evoked-response field (or potential)

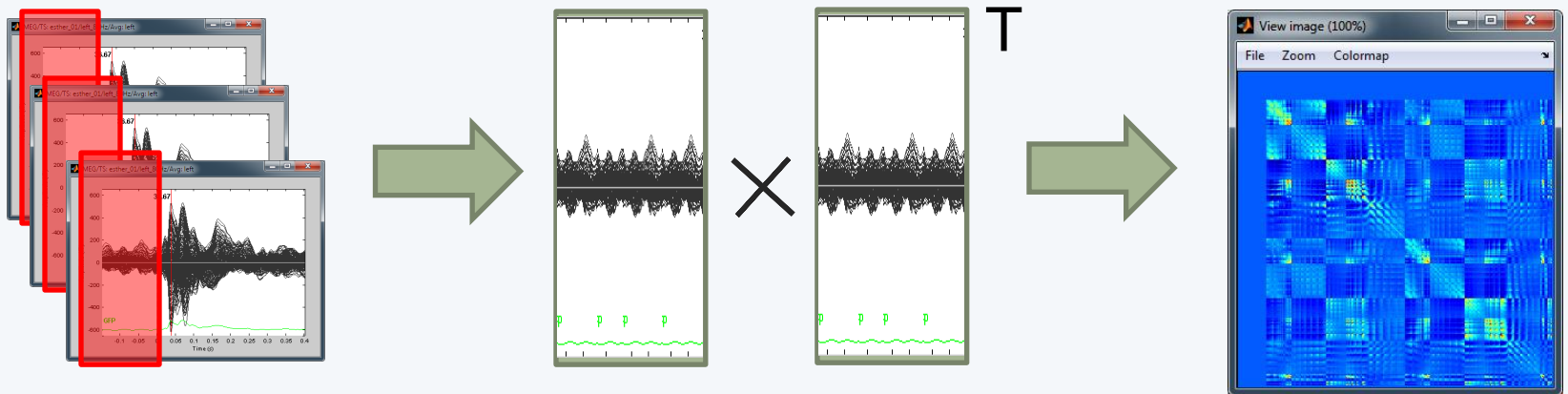


Source estimation

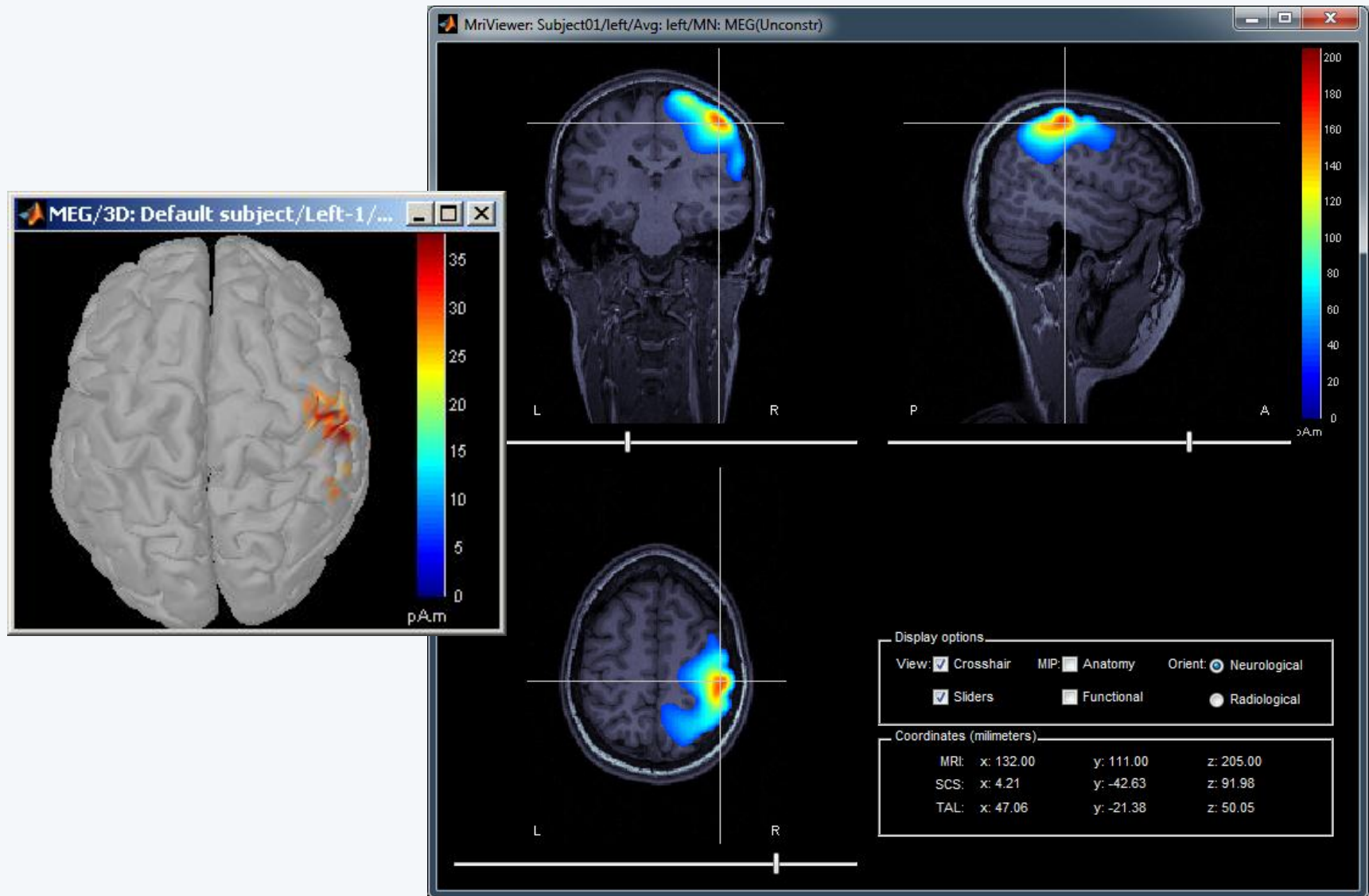
- Source space: cortex surface (or full head volume)
- Forward model = head model
Sources => Sensors
- Inverse model: Minimum norm estimates
Sensors => Sources



- Inverse model (minimum norm estimates) requires an estimation of the level of noise on the sensors
- Noise covariance matrix = covariance of the segments that do not contain any “meaningful” data
- Typically: empty room measures, or pre-stim baseline

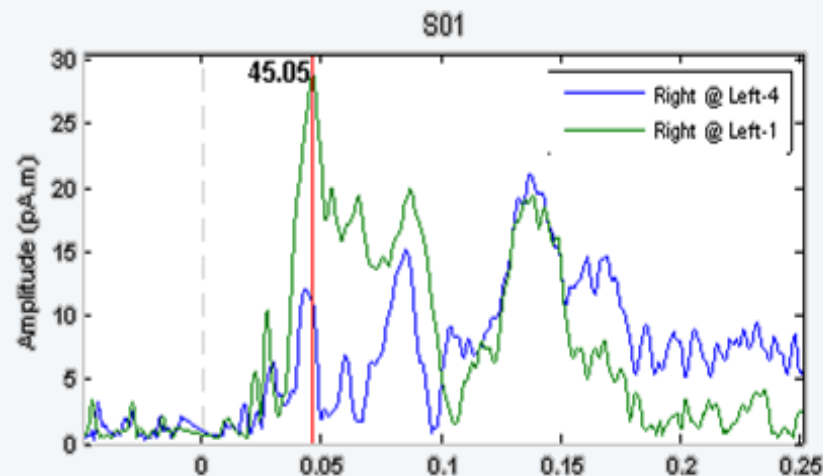
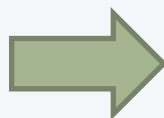
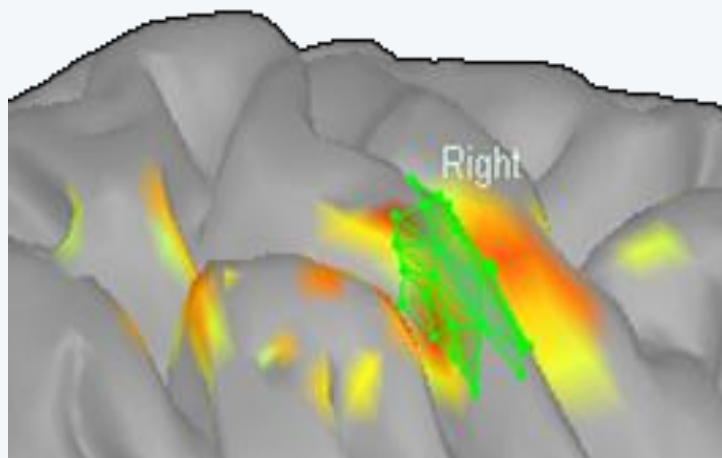


Sources activity



Regions of interest

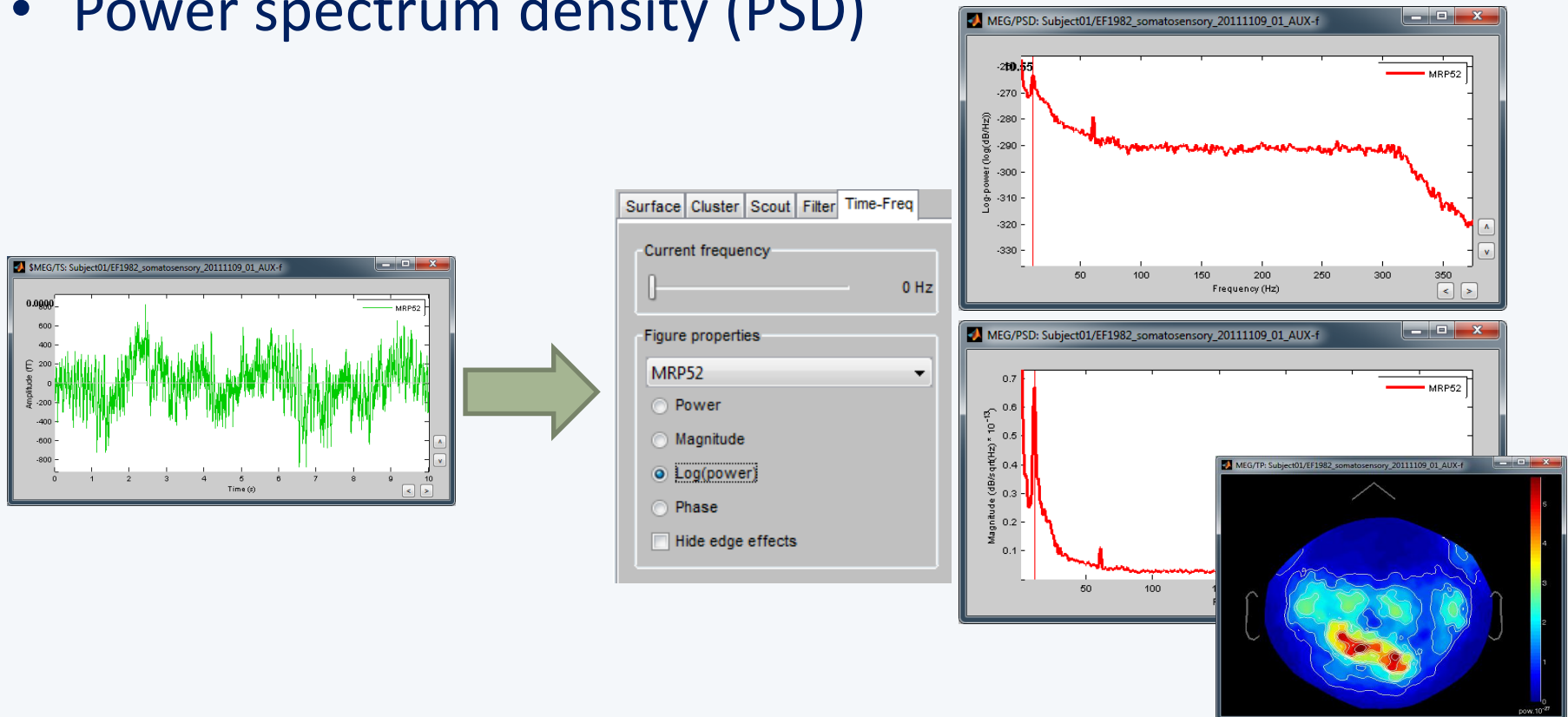
- Regions of interest at cortical level (scouts)
 - = Subset of a few dipoles in the brain
 - = Group of vertices of the cortex surface



- Noise normalization (z-score)
- Spectral and time-frequency analysis
- Group analysis:
 - Anatomical registration and normalization
 - Statistical inference
- Connectivity measures

Spectral analysis

- Fast Fourier transform (FFT)
- Power spectrum density (PSD)



Time-frequency / Hilbert transform

Time-frequency (Morlet wavelets)

Comment:

Time definition

☒ Same as input files
 [-500.00 : 0.42 : 1000.00] ms

☐ Group in time bands (ms)

t1: 0.00, 999.17
 t2: 1000.00, 1999.17
 t3: 2000.00, 2999.17
 t4: 3000.00, 3999.17
 t5: 4000.00, 4999.17
 t6: 5000.00, 5999.17

Generate

Frequency definition

☐ Linear (start:step:stop)
 1:1:80

☒ Group in frequency bands (Hz)

delta: 2, 4
 theta: 5, 7
 alpha: 8, 12
 beta: 15, 29
 gamma1: 30, 59
 gamma2: 60, 90

Reset

Morlet wavelet options

Central frequency: Hz (default=1)

Time resolution (FWHM): s (default=3)

Processing options

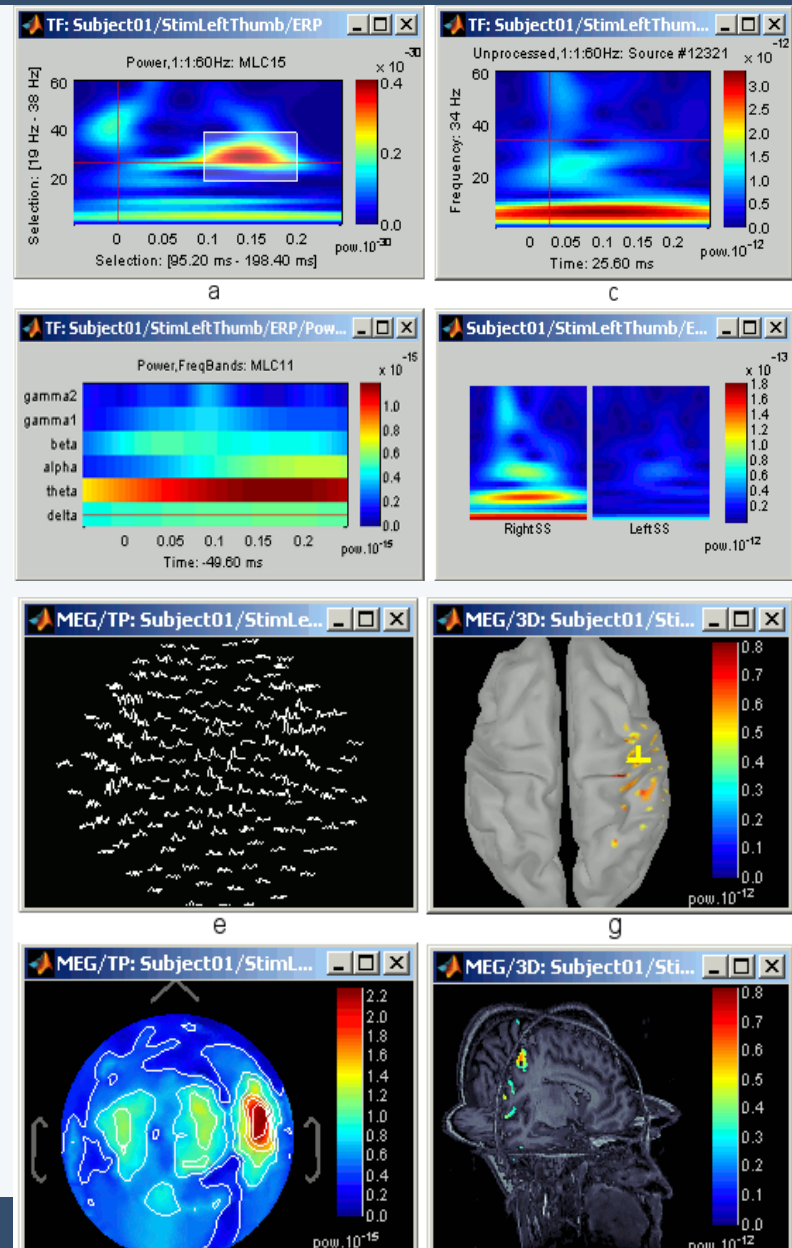
Compute the following measure:

☐ None (save complex values)

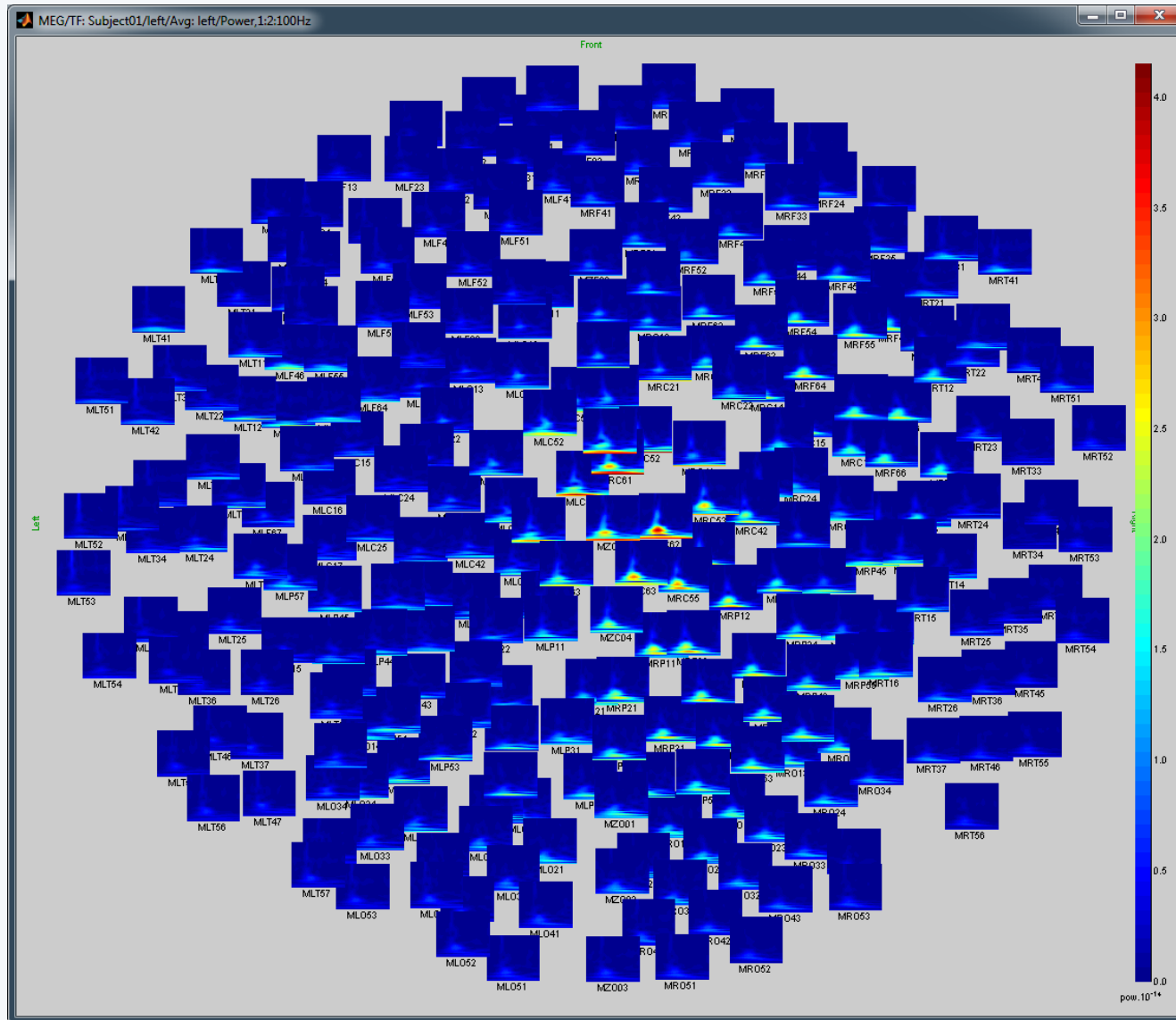
☒ Power

Estimated output file size: 45 Mb

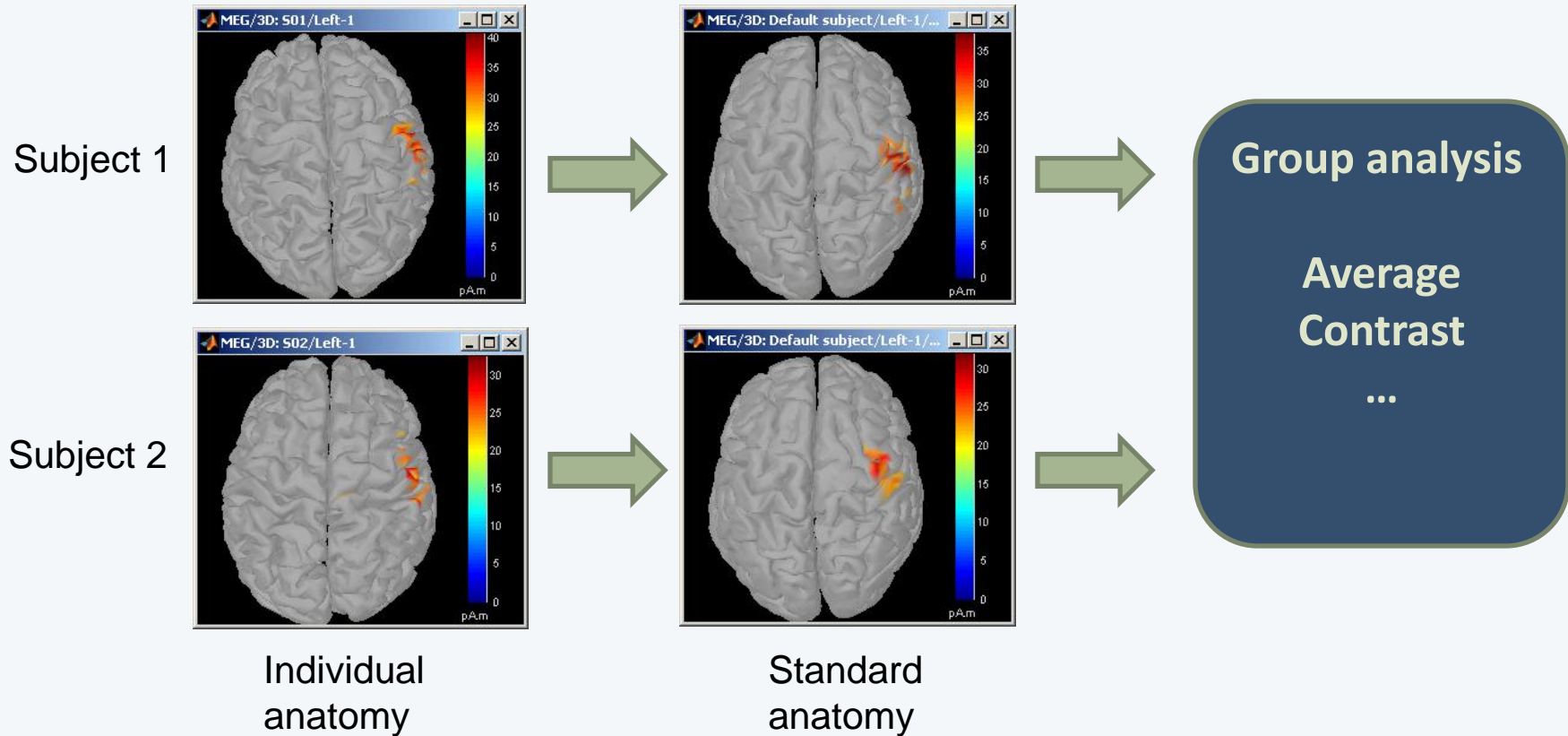
OK



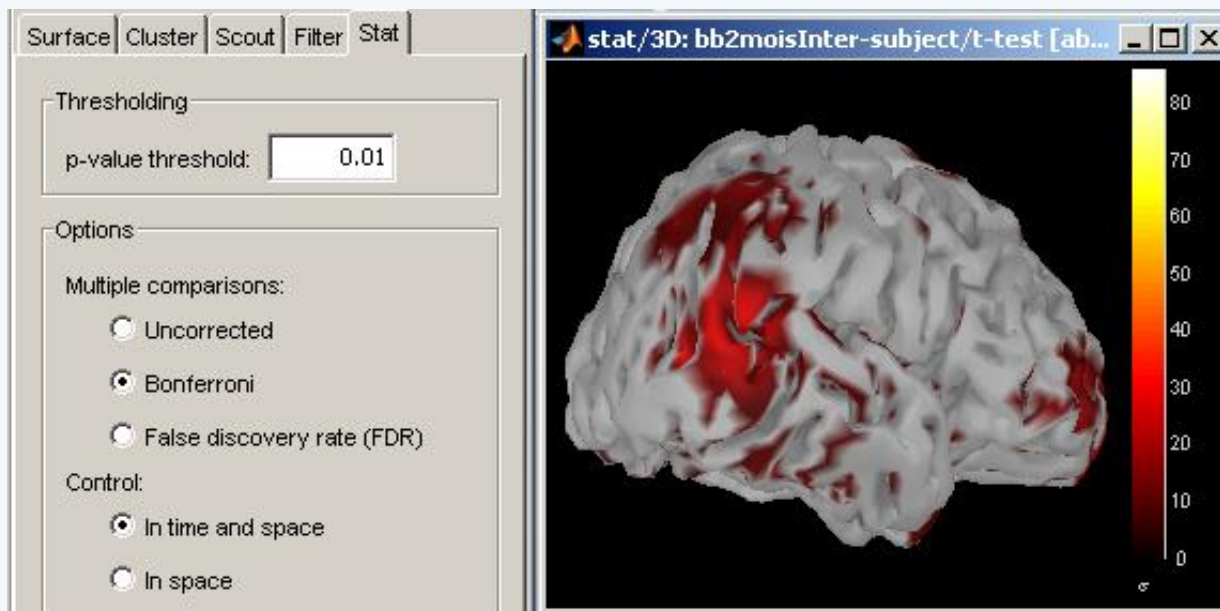
Time-frequency



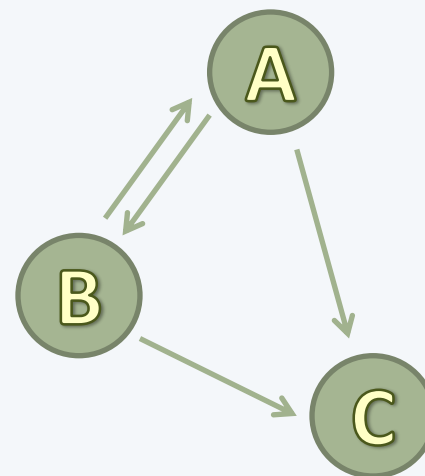
- Registration of individual brains on a template

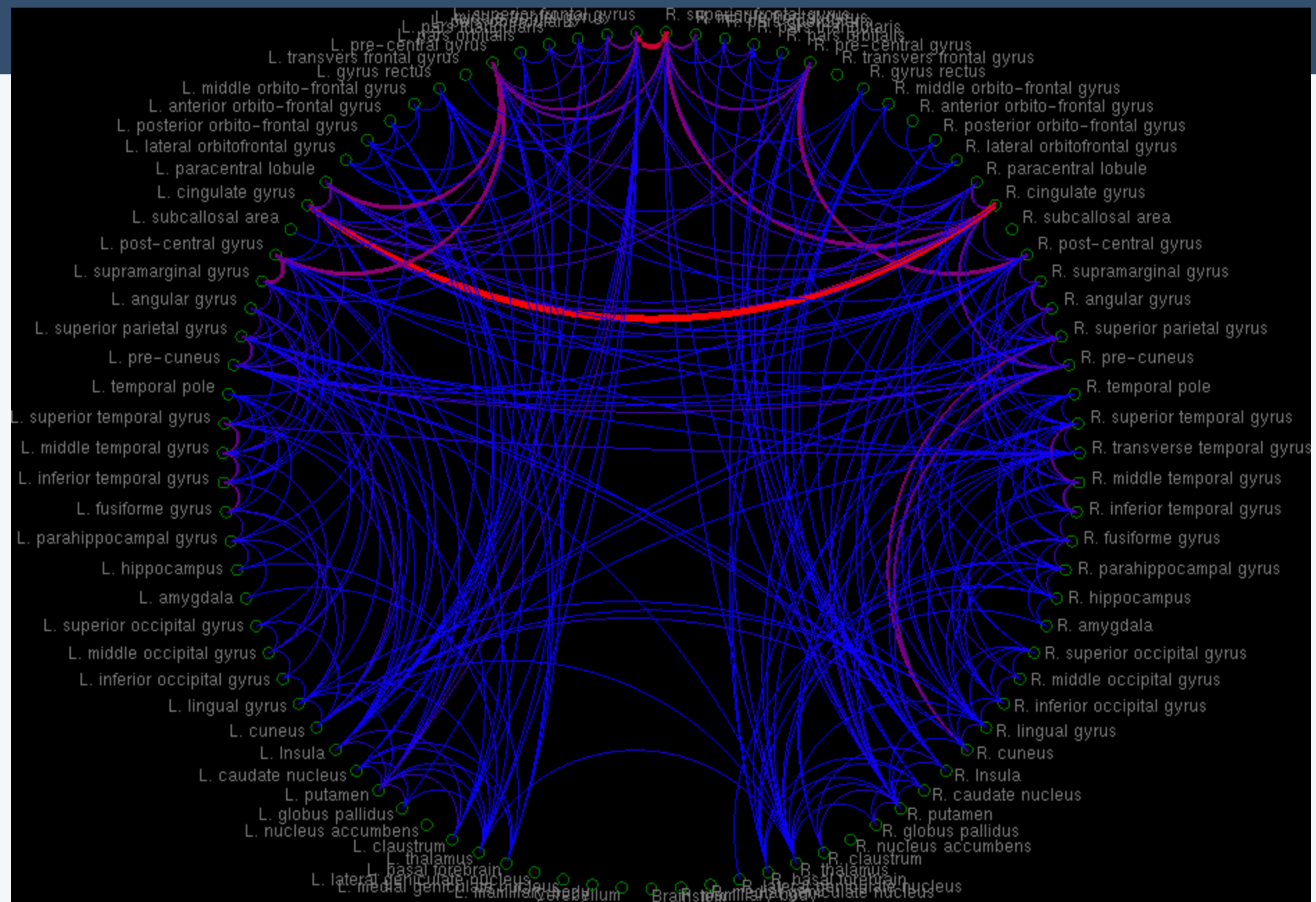


- Contrasts between subjects or conditions
- Statistical analysis: z-score, t-test
- Quick extraction of measures from complex paradigms
=> Export to: R, Excel, Statistica, SPSS, Matlab...

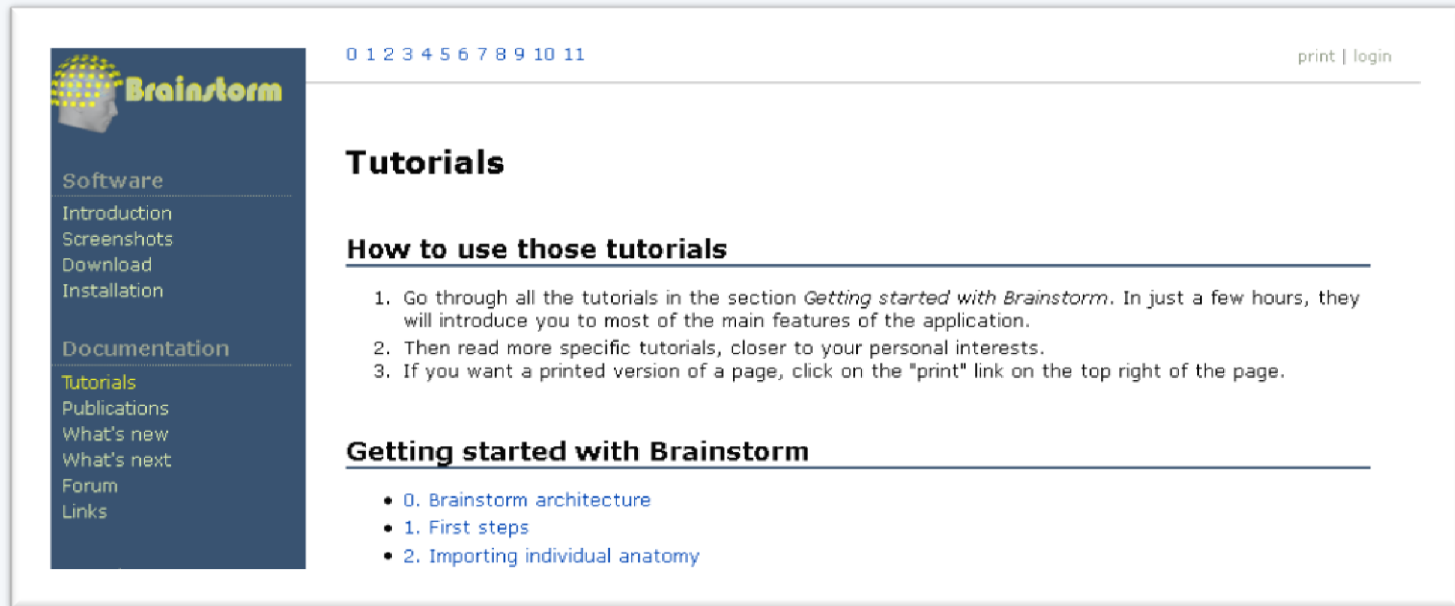


- Objectives: Describe the interaction between two brain regions, identify the brain networks
- Measures:
 - Correlation
 - Coherence
 - Granger causality
 - Phase locking value
- Both at sensor and source levels
- Problem of representation: too many dimensions





- Brainstorm online tutorials and forum:



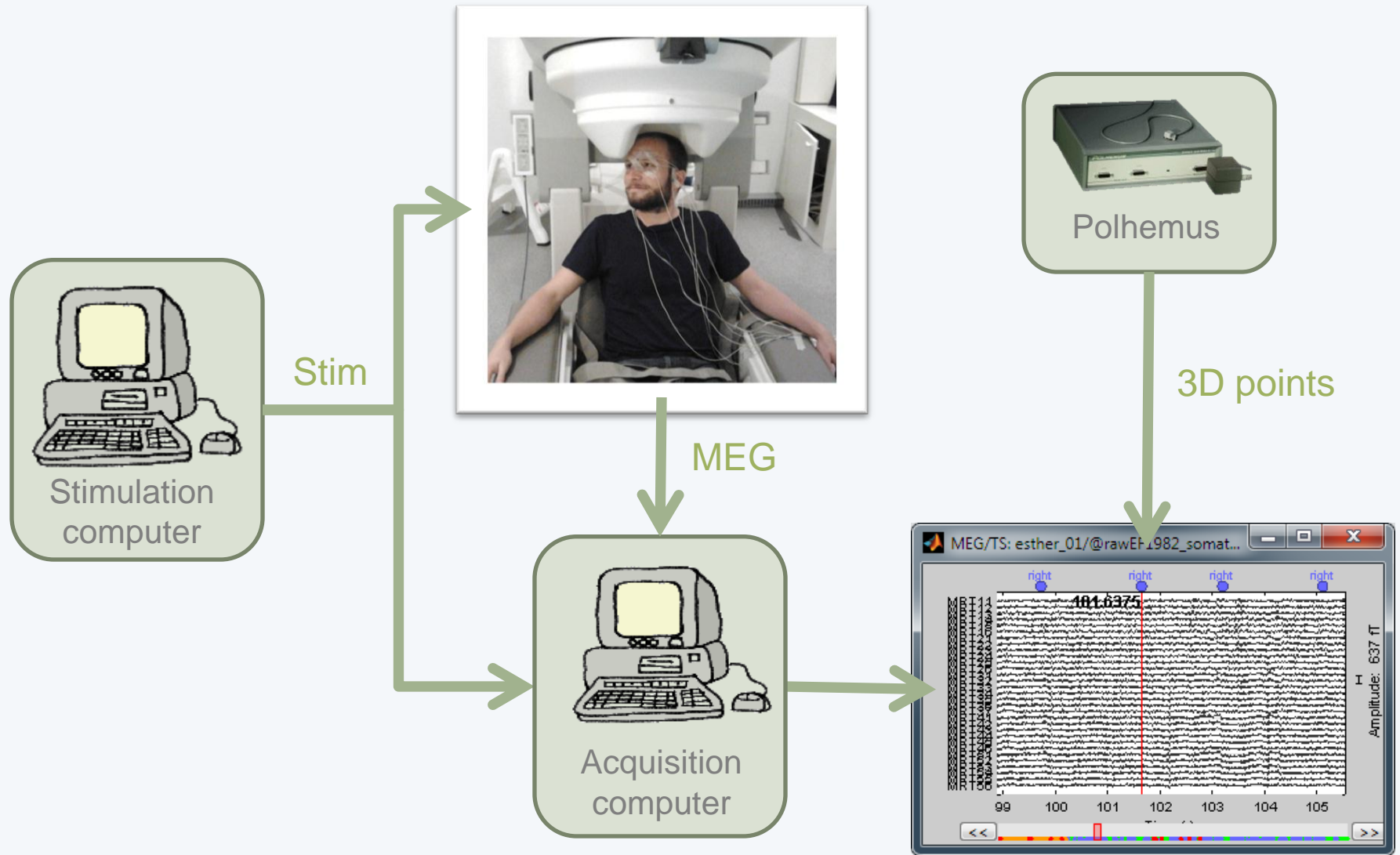
The screenshot shows the Brainstorm website's support page. On the left is a dark blue sidebar with the Brainstorm logo (a yellow dotted brain) and a list of navigation links: Software, Introduction, Screenshots, Download, Installation, Documentation, Tutorials (highlighted in yellow), Publications, What's new, What's next, Forum, and Links. The main content area has a top navigation bar with links 0 through 11, a 'print | login' link, and a 'Tutorials' section. Below this is a section titled 'How to use those tutorials' with three numbered steps: 1. Go through all the tutorials in the section *Getting started with Brainstorm*. In just a few hours, they will introduce you to most of the main features of the application. 2. Then read more specific tutorials, closer to your personal interests. 3. If you want a printed version of a page, click on the "print" link on the top right of the page. Below this is a section titled 'Getting started with Brainstorm' with a bulleted list of links: 0. Brainstorm architecture, 1. First steps, and 2. Importing individual anatomy.

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

- Median nerve stimulation
(Nov 2011, Montreal Neurological Institute, McGill)
 - Random electric stimulation of both arms
 - ~ 100 trials per arm
 - Acquisition at 1200 Hz
 - Recorded on CTF 275 MEG sensors
 - + 26 reference sensors
 - + EOG + ECG + STIM + ... = 302 channels
 - 6 minutes of recordings, 500 Mb

Sample data

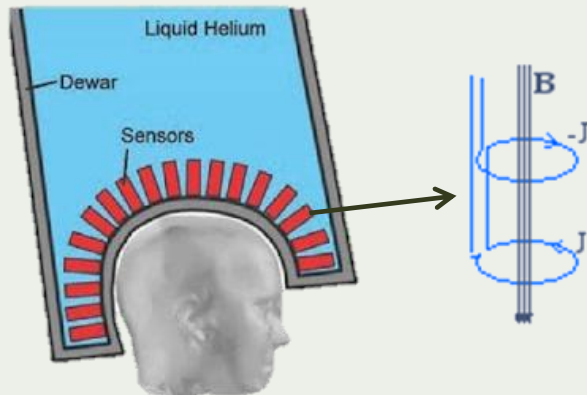
Acquisition setup



CTF vs. Elekta Neuromag

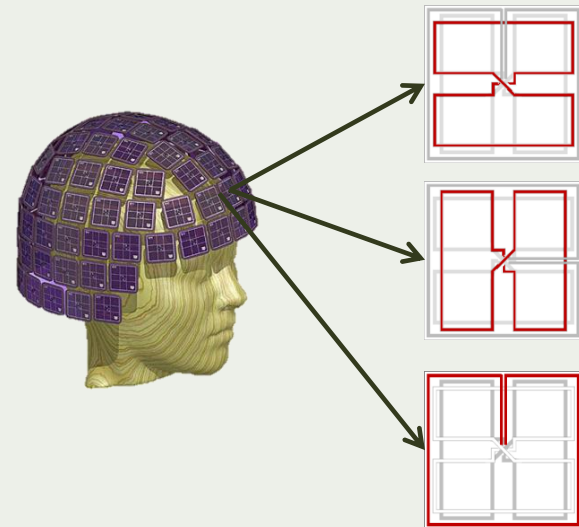
CTF (MNI)

275 axial gradiometers



Neuromag (MIT)

204 planar gradiometers
102 magnetometers



- Create a protocol, with one subject
- Anatomy
 - Import the MRI, define the anatomical landmarks
 - Import the surfaces
 - 2D / 3D display
- Recordings:
 - Review the continuous file
 - Mark cardiac peaks + eye blinks
 - Remove the ocular artifact (SSP)

- Recordings:
 - Import trials: [-100, +300] ms around each stimulus
 - Average the trials (left and right)
 - Explore the average at the sensor level
- Source estimation:
 - Head model
 - Noise covariance matrix
 - Sources time series
 - Review visually the results for left and right stim
 - Create a couple of regions of interest (scouts)

Contributors

Investigators



Sylvain Baillet
MNI



Richard Leahy
USC

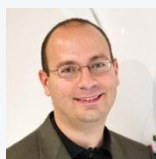


John Mosher
Cleveland Clinic

Key collaborators



Alexandre Gramfort
MGH / INRIA



Dimitrios Pantazis
MIT

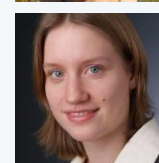


Rey Ramirez
UW

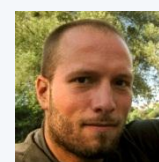
MEG @ McGill



Elizabeth Bock
MNI



Esther Florin
MNI



Francois Tadel
MNI

France

Lucie Charles
Ghislaine Dehaene-Lambertz
Claude Delpuech
Antoine Ducorps
Guillaume Dumas
Line Garnerio
Etienne Labyt
Karim N'Diaye
Lauri Parkkonen
Denis Schwartz
Lydia Yahia-Cherif

USA

Syed Ashrafulla
Sergul Aydore
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Belma Dogdas
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Matti Hamalainen
Sheraz Khan
Esen Kucukaltun-Yildirim
Alexei Ossadtchi
Darren Weber