The MNE package for M/EEG data processing

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Health Sciences 8

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The MNE Software Vision

- State-of-the-art methods, many examples, documented and tested
- ■Open development: collaboration between several centers
- ■Share the best practices, promote reproducible research

Software Features

Preprocessing

- Review raw data, filter, correct ECG / EOG with SSPs, ICA Forward & inverse modeling
- FreeSurfer structural data: Automatic forward modeling
- ■MNE dSPM sLORETA (TF-)MxNE LCMV

Statistics (sensor and source spaces)

- Time-Frequency (Phase-Locking, Induced Power)
- Parametric and non-parametric stats, with clustering Connectivity (sensor and source spaces)
- ■Functional and effective connectivity measures

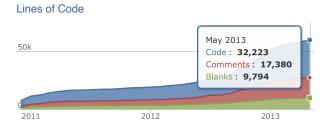
http://martinos.org/mne



A. Gramfort, M. Luessi, E. Larson, D. Engemann, D. Strohmeier, C. Brodbeck, L. Parkkonen, M. Hämäläinen MNE software for processing MEG and EEG data, Submitted.

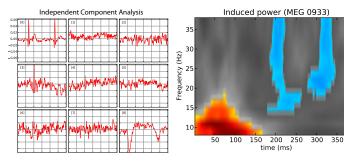
MNE-Python

- Python: general-purpose, high-level language
- Free: can run on a cluster without license problems
- Permissive BSD license: allows use in commercial products
- Many third-party packages easily integrated, e.g., ML
- ■Open, 29 contributors so far: ≈ 8 person years of effort

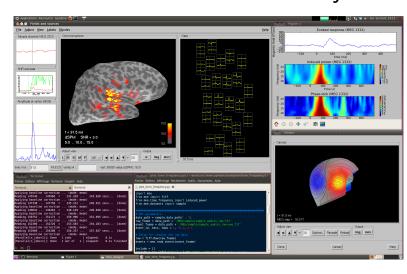


Learn more

- Mailing list: mne_analysis@nmr.mgh.harvard.edu
- ■http://martinos.org/mne/ (general doc)
- http://martinos.org/mne/python_tutorial.html
- http://martinos.org/mne/auto_examples/ (> 70 demos)
- ■http://mne-tools.github.com/mne-python-intro-slides



The MNE Software Family MNE-C – MNE-Matlab – MNE-Python



http://github.com/mne-tools D. Strohmeier, C. Brodbeck, L. Parkkonen, M. Hämäläinen

From raw to dSPM in < 30 lines of code

morph it to average brain for group study
stc_avg = mne.morph_data('sample', 'fsaverage', stc, 5, smooth=5)

