

# MNE for MEG/EEG data processing: What's up?



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## The MNE Software Vision

- State-of-the-art methods, 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 – DICS

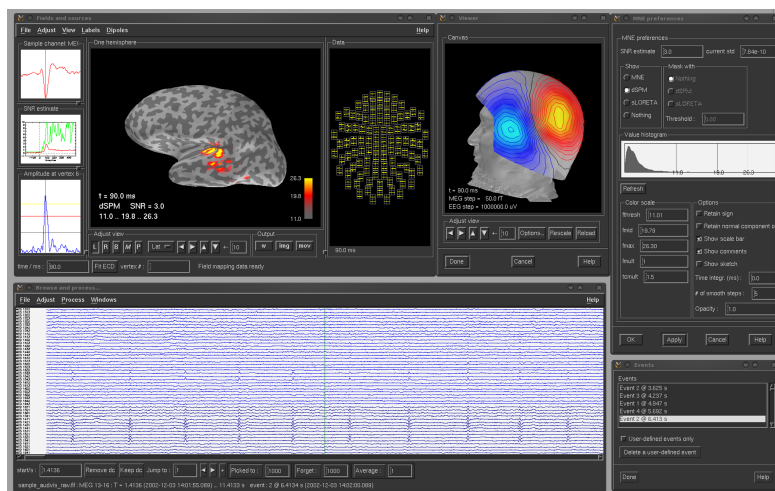
### Statistics (sensor and source spaces)

- Time-Frequency (Phase-Locking, Induced Power)
- Parametric and non-parametric stats, with clustering
- Decoding/MVPA (SVM, ...)

### Connectivity (sensor and source spaces)

- Functional and effective connectivity measures

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



<http://martinos.org/mne>

<http://github.com/mne-tools>



A. Gramfort, M. Luessi, E. Larson, D. Engemann, D. Strohmeier, C. Brodbeck, L. Parkkonen, M. Hämäläinen [MNE software for processing MEG and EEG data](#), Neuroimage, 2014

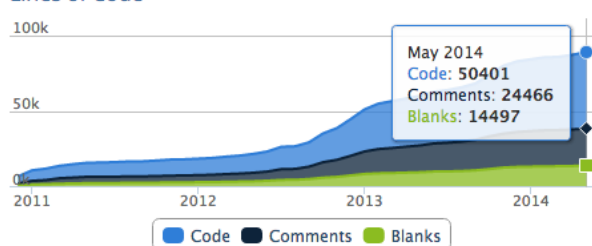


A. Gramfort, M. Luessi, E. Larson, D. Engemann, D. Strohmeier, C. Brodbeck, R. Goj, M. Jas, T. Brooks, L. Parkkonen, M. Hämäläinen, [MEG and EEG data analysis with MNE-Python](#), Frontiers in Neuroscience, 2013

## 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
- 46 contributors in last 12 months  $\approx$  13 person-years of effort

Lines of Code



## Learn more

- Mailing list: [mne\\_analysis@nmr.mgh.harvard.edu](mailto:mne_analysis@nmr.mgh.harvard.edu)
- <http://martinos.org/mne/> (general doc)
- [http://martinos.org/mne/dev/whats\\_new.html](http://martinos.org/mne/dev/whats_new.html)
- <http://mne-tools.github.com/mne-python-intro-slides>
- [http://martinos.org/mne/auto\\_examples/](http://martinos.org/mne/auto_examples/) (> 70 demos)

## From raw to dSPM in < 30 lines of code

```
import mne

fname = 'raw.fif'
raw = mne.io.Raw(fname) # load data
raw.info['bads'] = ['MEG 2443', 'EEG 053'] # mark bad channels

# band-pass filter data in beta band, and save it
raw.filter(13.0, 30.0, filter_length=4096, n_jobs='cuda')
raw.save(fname[:-4] + '_beta.fif')

# extract epochs
picks = mne.pick_types(raw.info, meg=True, eeg=True, eog=True)
events = mne.find_events(raw)
epochs = mne.Epochs(raw, events, event_id=1, tmin=-0.2, tmax=0.5, proj=True,
                    picks=picks, baseline=(None, 0), preload=True,
                    reject=dict(grad=4000e-13, mag=4e-12, eog=150e-6))

# compute evoked response and noise covariance, and plot evoked
evoked = epochs.average()
cov = mne.compute_covariance(epochs, tmax=0)
evoked.plot()

# compute inverse operator
fwd_fname = 'sample_audvis-meg-eeg-oct-6-fwd.fif'
fwd = mne.read_forward_solution(fwd_fname, surf_ori=True)
inv = mne.minimum_norm.apply_inverse(raw.info, fwd, cov, loose=0.2)

# compute inverse solution
stc = mne.minimum_norm.apply_inverse(evoked, inv, lambda2=1 / 3.0 ** 2,
                                     method='dSPM')

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

