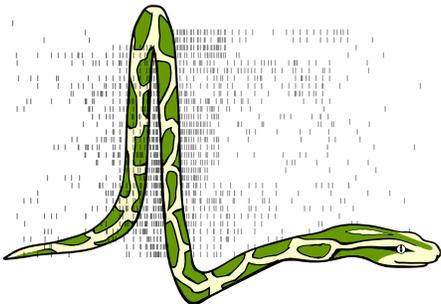


Python in Electrophysiology



Find the community @ <http://neuralsemble.org>

Data I/O

Neuroshare Tools

neuroshare

<http://g-node.org/neuroshare-tools>

Neuroshare is a standardized API for accessing neurophysiology data stored in vendor-specific binary formats in a vendor-neutral way.

- High-level Python library to access Neuroshare compatible data-files
- Automatically detects file types and loads the corresponding vendor library
- Support for GNU/Linux, MacOS X, and Windows
- Neuroshare-WineProxy enables the use of vendor libraries for Windows under GNU/Linux and MacOS X
- Comes with a tool to convert any data file supported by Neuroshare to the HDF5 format

Neo

<http://packages.python.org/neo>

neo

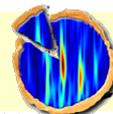
Neo provides a common model for representing electrophysiology data in Python. It provides I/O for reading a wide range of neurophysiology file formats (Spike2, NeuroExplorer, AlphaOmega, Axon, Blackrock, Plexon, Tdt) and for writing to a subset of these formats plus non-proprietary formats including HDF5.

Neo implements a hierarchical data model well-adapted to intracellular and extracellular electrophysiology and EEG data with support for multi-electrodes (e.g., tetrodes). Neo's data objects build on the [quantities](#) package, which in turn builds on [NumPy](#) by adding support for physical dimensions. Thus Neo objects behave like normal NumPy arrays but with additional metadata, checks for dimensional consistency and automatic unit conversion.

A project with similar aims but for neuroimaging file formats is [NiBabel](#).

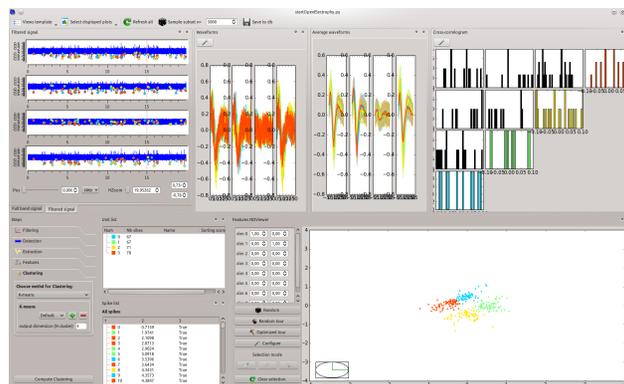
OpenElectrophy

<http://packages.python.org/OpenElectrophy>



OpenElectrophy is build on top of neo. It provides

- Powerful GUI
- Collection of methods for spike sorting
- Wavelet method for LFP transient oscillations analysis
- Customisable database to organize datasets



Data Management

G-Node Data API Python Client

<http://g-node.github.com>



G-Node provides a free cloud-based service neuroscientists can use for storage, management and sharing of data (<https://portal.g-node.org/data/>). An API for data access is provided (<http://g-node.github.com/g-node-portal/>), allowing developers to code their own clients. A client for Python is currently under development:

- Compatibility with NEO
- Smart lazy loading and caching for frugal bandwidth and memory usage
- Possibility to work in a mixed workflow: work on the same data in Python and MATLAB[®]

Metadata Management

odML libraries & Editor

<http://www.g-node.org/odml/>



Use the open metadata Markup Language to annotate data with information about the stimulus, data acquisition, and experimental conditions.

- Developer friendly libraries for Python and Java
- Fully functional graphical editor for Linux, Windows, and MacOS X

Name	Name	Value	Definition	Type	Unit	Comment	Encoder	Checksum	Reference
Electrode	Author	Christian Kellner	person						
	Date								
	Description								
	Method								
	DatasetFileURL								

Document: Analysis
Attribute Value
definition
include http://portal.g-node.org/odml/terminologies/v1.0/analysis/analysis.xml
link

Grewe J, Wachtler T, Benda J (2011) A bottom-up approach to data annotation in neurophysiology. *Front. Neuroinform.* 5:16. doi: 10.3389/fninf.2011.00016

Simulation Tools

LFPy

<http://compneuro.umb.no/LFPy>



LFPy is a Python module for simulation of extracellular electrical potentials evoked by activity of multi-compartment model neurons.

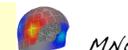
LFPy provides:

- A forward modeling scheme for calculating extracellular potentials from compartmental membrane currents in an infinite homogeneous linear extracellular medium
- Scripting capabilities thanks to NEURON and the Python programming environment
- Simultaneous simulation of the model cell responses and extracellular potentials based on a biophysically detailed neuronal model
- Support for common formats for reconstructed neuronal morphologies, allowing use of publicly available 3D-reconstructions (e.g., <http://www.neuromorpho.org>).

Analysis

MNE-Python

<http://www.martinos.org/mne>

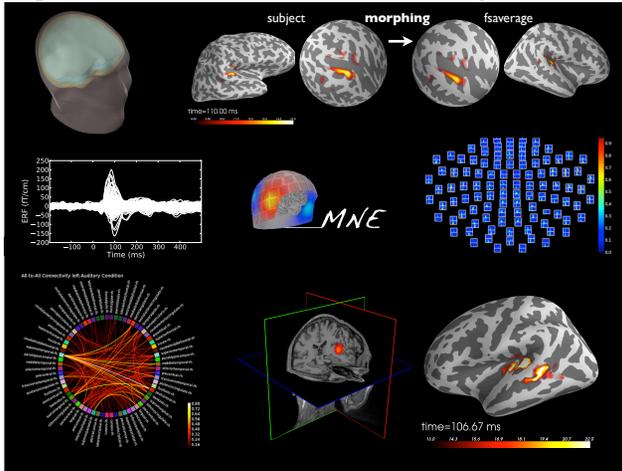


MNE is a software package for processing magnetoencephalography (MEG) and electroencephalography (EEG) data. (<http://www.martinos.org/mne>)

- Data conditioning and data conversion utilities
- Interactive and batch-mode modules for the forward and inverse calculations
- Cortically-constrained source estimates
- Associated dynamic statistical parametric maps, option-

ally constrained by fMRI

- Frequency-domain and time-frequency analyses, non-parametric statistics, multivariate decoding



A. Gramfort, *et al.* MNE software for processing MEG and EEG data, *NeuroImage*, Volume 86, 1 February 2014, Pages 446-460, ISSN 1053-8119, <http://dx.doi.org/10.1016/j.neuroimage.2013.10.027>

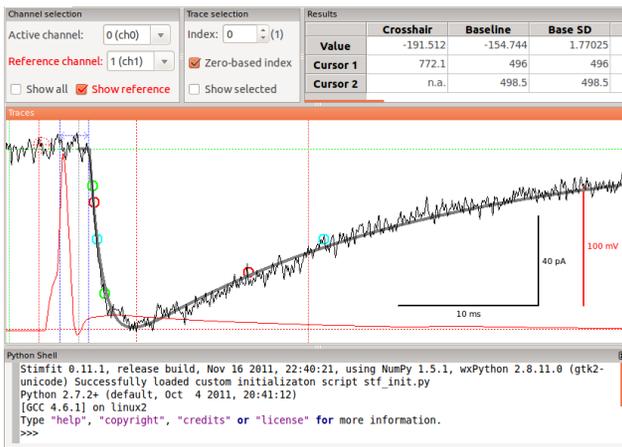
Stimfit

<http://www.stimfit.org>



Visualize and quantify electrophysiological data.

- With a focus on patch-clamp recordings
- Supports most standard patch-clamp file types
- Embedded Python shell
- Measures action potential, EPSC and EPSP kinetics
- Extracts spontaneous and evoked events
- Successfully used in many publications for > 5 years



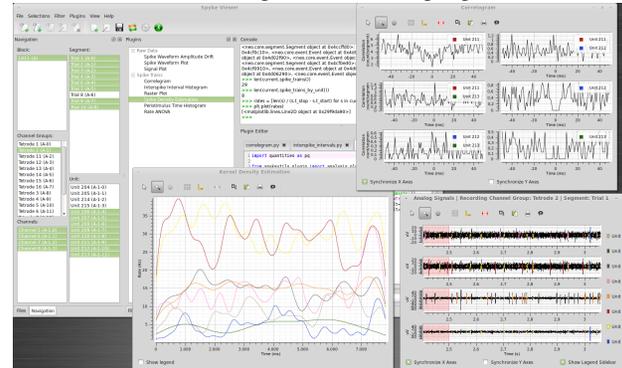
Spyke Viewer

<http://spyke-viewer.readthedocs.org>



Navigate, visualize and analyze electrophysiological data. Designed to be flexible and extensible.

- Based on Neo for easy data and algorithm sharing
- Flexible: Work with selected data using the embedded Python console
- Easily extensible: Create useful plugins in minutes
- Use supplied plugins for common plots such as Raster Plot, PSTH, Correlogram and analog signals



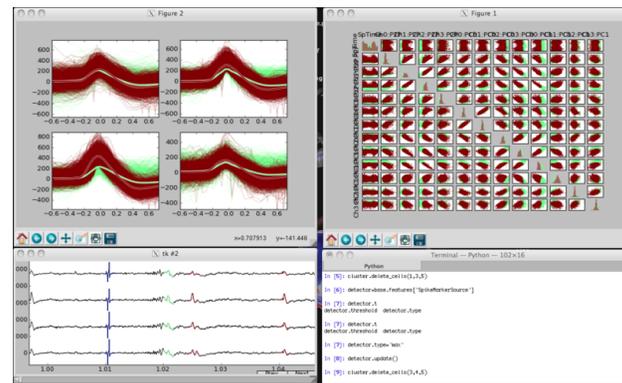
SpikeSort

<http://spikesort.org>



SpikeSort is a flexible spike sorting framework implemented completely in Python. It features manual and automatic clustering, many data formats and it is memory-efficient. Truly Open Source, BSD-licensed.

- Interactive command-line interface in Python
- GUI and visualization widgets
- Support for multi-channel data



Data Acquisition

ACQ4

<http://acq4.org>



ACQ4 is a complete software system for data acquisition and analysis in neurophysiology research. It is currently used for patch clamp electrophysiology, multiphoton imaging, photostimulation mapping, calcium imaging, and intrinsic imaging.

- Easy to design protocols synchronizing multiple devices
- Live camera viewing for patching and online analysis of calcium imaging
- Modular and scalable design – easy to expand support for new hardware and experiments
- Integrated data manager for hierarchical data storage with annotations and other metadata

