

Nitime: an open-source package for time-series analysis of neuroscience data



for Theoretical Neuroscience



Neuroimaging In PYthon

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Introduction

>> The analysis of time-series data is central to many research methods in neuroscience, ranging in temporal and spatial scale from single-cell recordings to BOLD fMRI. Nitime aims to provide a common programming interface for the representation of time-series data from different modalities and to provide algorithms and visualization tools for the analysis of these data.

>> Nitime is a part of the NIPY project (http://nipy.org), which is a development community of open-source tools for neuroimaging research (see posters #2990, #3429, #3841, #2927). Nitime is written in Python using Scipy (http://scipy.org).

Software design and features (see also: http://nipy.org/nitime/users/overview.html):

>> Container objects for representation of time and for representation of time-series data:

- ^x TimeArray: Time-points with time-unit handling (time-unit display, conversion, etc).
- x UniformTime: Uniformly sampled time-points, with the associated sampling rate/interval and time-unit; allows indexing with time information.
- ^x TimeSeries: Uniformly sampled time + data; indexing with time information.
- ¤ Events: Time-points + data.
- ¤ Epochs: intervals in time (start-time, stop-time, offset).
- >> General purpose algorithm library (not dependent on the design of the time-series objects):
 - Spectral decomposition, coherency, wavelet transforms, FIR, etc.
- >> 'Analyzer' objects bridge between the time-series represenation and the algorithms, providing an easy-to-use interface
- >> Lazy initialization: intense computations in the analyzer objects are done on a need-to-know basis. Once a computation is done, the results are cached for reuse.
- >> Visualization of time-series data and results of analysis (see examples below).

Examples (see also: http://nipy.org/nitime/examples/index.html):

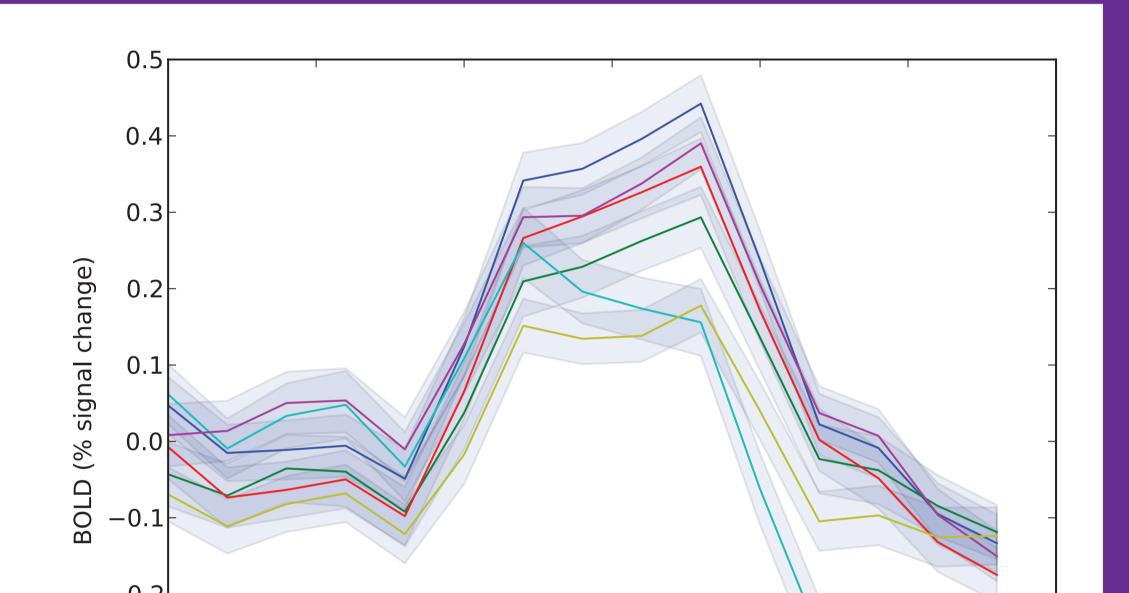
Event related analysis:

Visualize result:

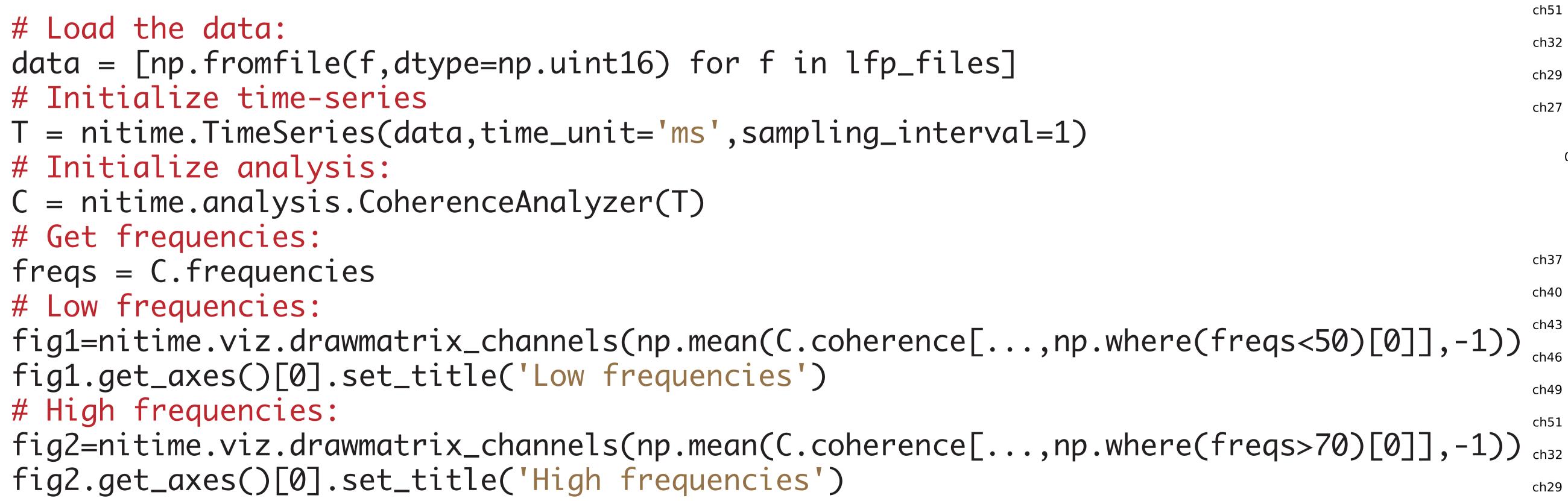
BOLD fMRI in human visual system in response to different directions of motion:

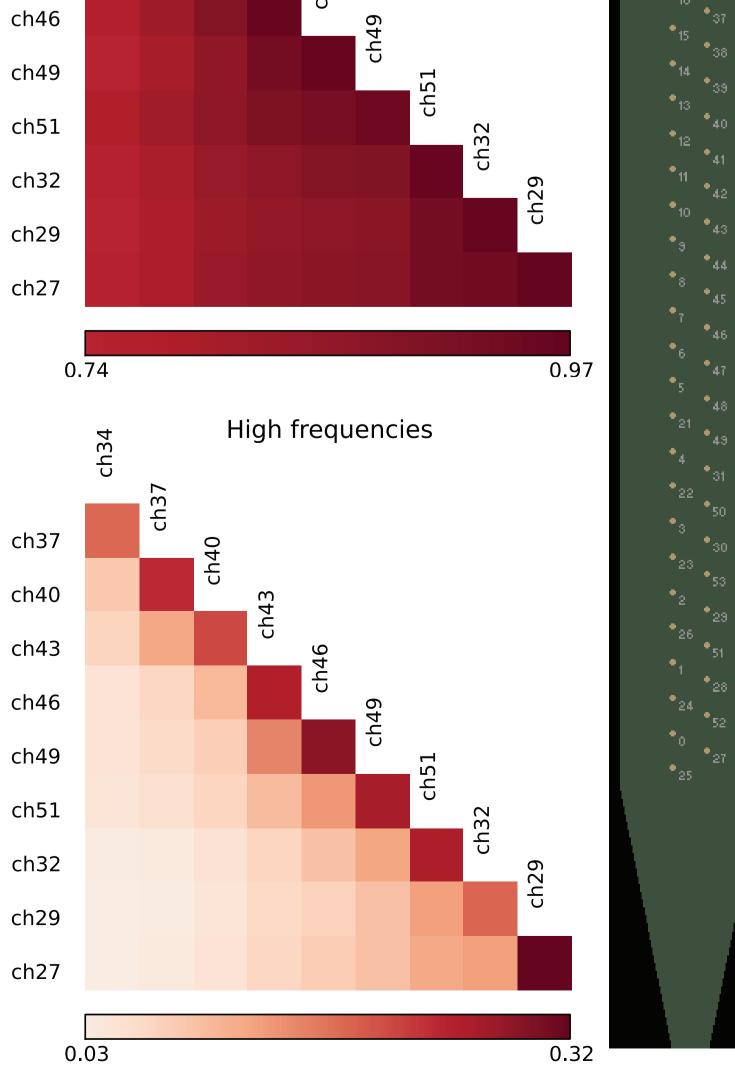
nitime.viz.plot_tseries(event_related.eta,ylabel='BOLD (% signal change)',

data = numpy.recfromcsv('data/event_related_fmri.csv') # Initialize time-series, events objects: T1 = nitime.TimeSeries(data.bold,sampling_interval=2) T2 = nitime.TimeSeries(data.stimulus,sampling_interval=2) # Initialize 'analyzer' object: event_related = nitime.analysis.EventRelatedAnalyzer(T1,T2,15,offset=-5)



yerror=event_related.ets) 15 -5 5 10 0 Time (s) Intra-cellular recordings of spike-times in the grasshopper auditory system in response to an auditory stimulus (data available on the CRCNS data-sharing site: http://crcns.org): $_{a}$ # Load data: stimulus = numpy.loadtxt('data/grasshopper_stimulus1.txt') # Initialize time-series object: T = nitime.TimeSeries(data=stimulus,sampling_interval=50,time_unit='us') (dB SPL) # Load the spike-times from the data file: spike_times = numpy.loadtxt('data/grasshopper_spike_times1.txt') # Initialize the Event object holding the spike-times: E = nitime.Events(spike_times,time_unit='us') # Initialize the analysis object: event_related = nitime.analysis.EventRelatedAnalyzer(T,E,200,offset=-200) # Visualize the results: nitime.viz.plot_tseries(event_related.eta,ylabel='Amplitude (dB SPL)', 54 - 10-8 -2 -6 time_unit='ms') Time (ms) Low frequencies Coherence analysis: Concernce analysis. Coherence is a spectral analogue of correlation: $R_{xy}(\lambda) = \frac{f_{xy}(\lambda)}{\sqrt{f_{xx}(\lambda) \cdot f_{yy}(\lambda)}}$ ch37 ch40 ch43 ch46 LFP data acquired with a polytrode in cat visual cortex (data available on CRCNS): ch49





Future developments:

>> Functional connectivity with Granger casuality (Kayser et al. 2009) >> Readers for common file-formats