### Summary

Modern techniques in optophysiology have allowed neuroscientists unprecedented access to neuronal activity in vivo. The time series datasets generated from these experiments are becoming increasing larger as new technologies allow for faster acquistion rates of raw data. These fluorscent recordings are being made with an ever expanding library of indicators for calcium, voltage, neurotransimitter and neuromodulator activity. These signals generated from these fluorscent bioindicators contain the information of the underlying neuronal activity but all have unique molecular kinetics and inherit signal-noise ratios which must be taken into account during singal processing. The development of pyNeuroTrace, an open-source Python library, was made to aid in the processing of these neurnal signals which must be filtered with these unique aspects in mind before analysis can be completed.

#### Statement of need

Many neuroscience labs that use optophysiological methods, such as two-photon microscopy or fibrephotomotery, frequently must rewrite and maintain common functions and filters needed to analysis the raw recordings. Furthermore, many technique and algorithms for signal processing are scattered throughout the literature and are frequently implemented programing languages other than Python. pyNeuroTrace meets the need of a time series analysis written purely in Python for neuronal activity. Our package is a collection of filters and algorithms implemented in a genralizable manner for time series data in either 1D-arrays or a collection of recording in a 2D-arrays. Additionally, with the increase in aquistion rate of new imaging techniques, we have implementations of these algorithms using GPU compatiable code to increase the speed in which the techniques can be applied to larger datasets collected at kilohertz rates.

# Signal Processing

### DeltaF/F

There are several methods for calculating the DeltaF/F of a fluorscent trace. We implemented the method described by Jia et al which includes several smoothing steps to help with shot noise[@Jia2010]. In short, F0 is calculated by finding a the minmum signal in a window of the rolling average of the raw signal. Then DeltaF is calculated by the difference in the raw signal and F0 divided by F0 (F-F0)/F0. This DeltaF/F signal is optionally smoothed using an exponetially wieghted moving average (ewma) to further remove shot noise.

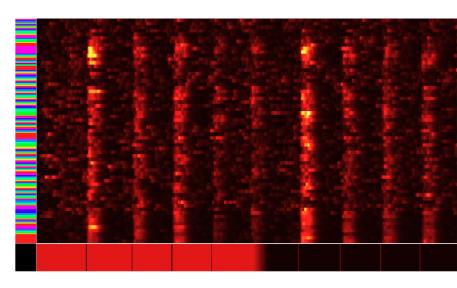
#### Okada Filter

We implement the Okada Filter in Python[@Okada2016]. This filters is designed to filter shot-noise from traces in low-signal to noise paradigms, which is common for calcium imaging with two-photon imaging where the collected photon count is low and noise from PMT can nontrivial.

## Nonnegative Deconvolution

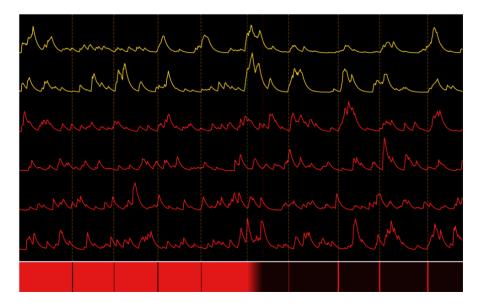
#### **Event Detection**

# Visualization

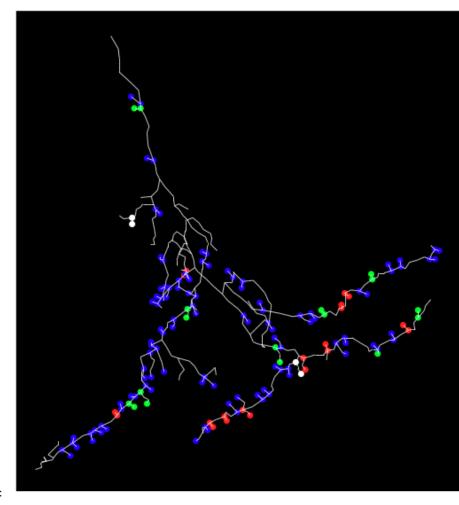


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### **GPU Acceleration**

Several of the filters in pyNeuroTrace have been rewritten to be almost entirerly vectorized in their calculations. The benefit being a noticable difference in the performance for larger time series. These vectorized versions gain further speed by being excuted on a GPU. To excuted these version the filters can be imported from the module, pyneruopyneurotrace.gpu.filters, and a CUDA compatiable graphics card is needed. This functionality is becoming increasingly important as aquistion rates increase for kilohertz imaging of activity can generate arrays hundreds of thousands datapoints in length in just a few minutes. Figure 1 shows the difference in calculating arrays of various sizes using either the CPU or vectorized GPU based approach of the dF/F function. The CPU used in these calcultions was a Intel i5-9600K with six 4.600GHz cores, the GPU was a

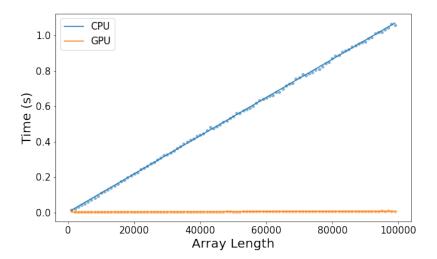


Figure 1: Comparison between dF/F with EWMA calculations for different array sizes.

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Double dollars make self-standing equations:

$$\Theta(x) = \begin{cases} 0 \text{ if } x < 0\\ 1 \text{ else} \end{cases}$$

You can also use plain LATEX for equations

$$\hat{f}(\omega) = \int_{-\infty}^{\infty} f(x)e^{i\omega x}dx \tag{1}$$

and refer to Equation 1 from text.

# Acknowledgements

The development of this software was supported by funds from the Canadian Institutes of Health Research (CIHR) Foundation Award (FDN-148468).

### References