

# The Neural Code

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Problem Set 5

—STC/GLM—

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1. (STC) (Python exercise) Consider the two filters:

$$a(t) = \exp\left(-\left(\frac{t - 10 \text{ ms}}{\sqrt{10} \text{ ms}}\right)^2\right) \cos\left(2\pi \frac{(t - 10 \text{ ms})}{10 \text{ ms}}\right)$$
$$b(t) = \exp\left(-\left(\frac{t - 10 \text{ ms}}{\sqrt{10} \text{ ms}}\right)^2\right) \cos\left(2\pi \frac{(t - 10 \text{ ms})}{10 \text{ ms}} + \pi/2\right)$$

- (a) Create the filters from 0ms to 20ms with a bin size of 1ms. Also, normalize the filter amplitudes by `numpy.linalg.norm`.
- (b) Generate a white noise stimulus  $x(t)$  with mean 0 and variance 1 for 100s with a bin size of 1ms.
- (c) Consider a neural system which takes  $x(t)$  as input and produce an output  $r(t)$  with a squaring nonlinearity by

$$r(t) = 2[x * a](t)^2 + 1.5[x * b](t)^2$$

Compute  $r(t)$ . To produce a better result, pad an array of zeros in front of the stimulus  $x(t)$  before convolving with the filters with `numpy.convolve(..., mode='valid')`. The zero-padding array should have the size of  $(N_{\text{filter}} - 1)$ , where  $N_{\text{filter}}$  is the size of your filter array.

- (d) The output  $r(t)$  is firing rate. Consider an inhomogeneous poisson process with instantaneous firing rate of  $r(t)$ . Sample a spike train from this process until 100s.
- (e) Compute the spike-triggered average (STA) up to  $N_{\text{filter}}$  time bins before the spikes.

$$\text{STA}(t) = \frac{1}{F} \sum_f x(t_f - t),$$

where  $F$  is the total number of spikes. The STA should look noisy and lack interpretable structure.

- (f) Compute the spike-triggered covariance (STC).

$$\text{STC} = \frac{1}{F} \sum_f \xi_f \xi_f^T$$
$$\xi_f = x(t_f - t) - \text{STA}(t)$$

As a shortcut, you could also utilize the function `numpy.cov( $\xi$ )` to compute the covariance matrix.

- (g) Obtain the eigenvectors and eigenvalues of the STC. (You could use the function `numpy.linalg.eig`.) Plot the eigenvalues from high to low. You should see two eigenvalues which are particularly higher than the rest.
- (h) Plot the corresponding eigenvectors of these two eigenvalues. They should look similar to the filters  $a(t)$  and  $b(t)$ . The signs can be opposite (the eigenvectors have negative values instead of positive). Why?
- (i) How would the model generalize to two pixels (a whitenoise vector  $\vec{x}$  with two entries)? How many kernels could there be for  $n$  pixels?

## 2. (GLM-based spike fitting)

The exponential family distribution

$$p(y) = h(y) \exp[\theta y - A(\theta)]$$

and the (invertible) canonical link function  $\langle y \rangle = g^{-1}(\theta)$ , with  $\theta = \vec{w} \cdot \vec{x}$ , define a generalized linear model.

- (a) Show that  $\vec{w} \cdot \vec{x} = g[\partial_{\theta} A(\theta)]$ .
- (b) Find the canonical link function for the Poisson count distribution  $p(y) = \frac{(\lambda)^y}{y!} e^{-\lambda}$ .  
*Hint:* First, identify  $h(y)$ ,  $\theta$ , and  $A(\theta)$ .
- (c) Let us now consider the following dynamical system (L-NL cascade)

$$\lambda_t = \exp[(s * \phi)_t - (y * \eta)_t] \Delta t ,$$

in which  $\lambda_t$  is the density of an inhomogeneous poisson process that generates spike counts  $y_t$ , and  $\Delta t = 0.1$  ms.  $s_t$  is a stimulus,  $\phi_t$  and  $\eta_t$  are the feedforward and feedback filters, respectively. We now consider the system in discrete time ( $\lambda_t$  is constant in the  $t$ -th time bin). Show that the log-likelihood of an output spike (count) train  $y_t$  equals

$$\mathcal{L} = \sum_t (y_t \ln \lambda_t - \lambda_t).$$

- (d) (Python exercise) Let us consider the two specific kernels

$$\phi_t = a t \exp[-(tb)^2/2] \quad \eta_t = d/c \exp[-tc] \mathcal{H}(t)$$

with  $\mathcal{H}$  denoting the Heaviside function. Plot the kernels and give biological motivations.

- (e) (Python exercise) Load the stimulus  $s$  and the measured spike train  $y$  from `y.npy` and `s.npy` (Sampling rate 10 kHz) and fit the model parameters by maximizing the likelihood (use `scipy.optimize.fmin`)
- (f) Read the paper `Pillow2008.pdf`.

**3. (Logistic Regression)** Logistic regression is (in its simplest form) a binary classification method ( $y = \pm 1$ ) in which, one models the probability  $q$  that an input  $x$  is in class  $y = +1$  by the logistic function

$$q(x) = [1 + \exp(-wx)]^{-1} .$$

The log-likelihood can be written as

$$\ln p(y|x) = \frac{1+y}{2} \ln q(x) + \frac{1-y}{2} \ln [1 - q(x)]$$

- (a) Explain the log-likelihood formula.

(b) For multiple observations  $(y_i, x_i), i = 1, \dots, \ell$  the log-likelihood sums up

$$\ln p(\{y_i\}|\{x_i\})_{i=1,\dots,\ell} = \sum_i \frac{y_i + 1}{2} \ln q(x_i) + \frac{1 - y_i}{2} \ln[1 - q(x_i)]$$

Show that the gradient can be expressed as

$$\partial_w \ln p(\{y_i\}|\{x_i\})_{i=1,\dots,\ell} = \sum_i x_i [z_i - q(x_i)]$$

with  $z_i = (1 + y_i)/2$ .

(c) Explain the relation of binary logistic regression to GLMs.