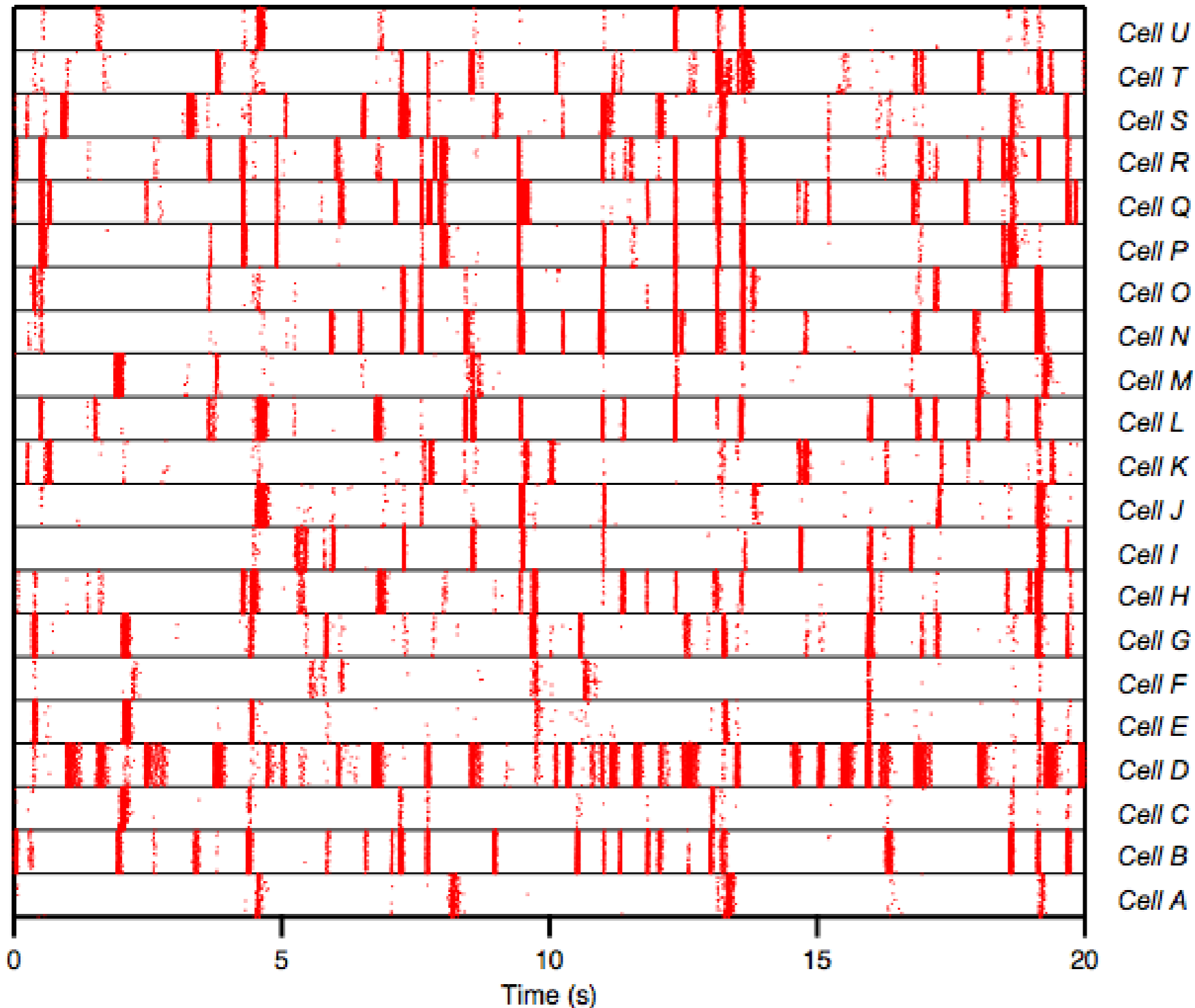


What is the neural code?



Single neurons

Populations

Encoding and decoding

$P(\text{response} \mid \text{stimulus})$

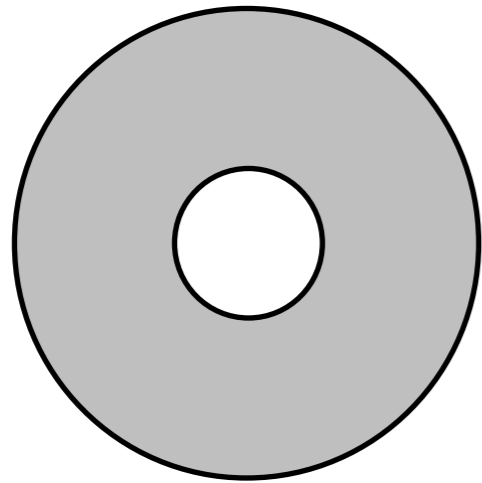
encoding

$P(\text{stimulus} \mid \text{response})$

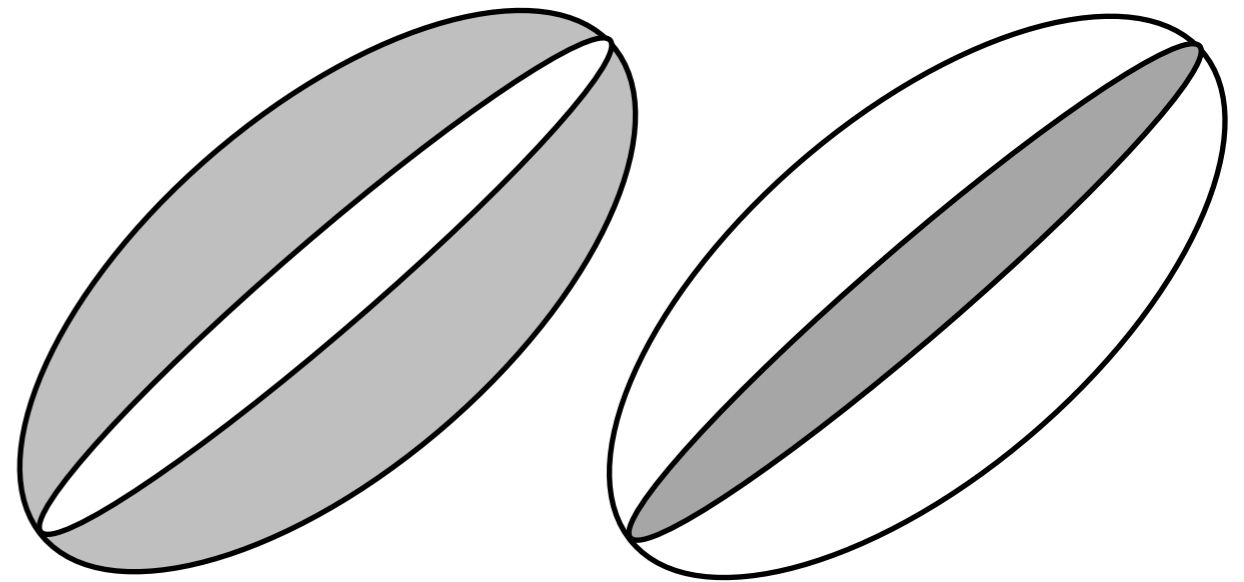
decoding

- What is response?
- What is stimulus?
- What is the function P?

Basic coding model: linear filtering



retina



Visual cortex

Spatial filter: $r = \iint f(x,y) I(x_0-x, y_0-y) dx dy$

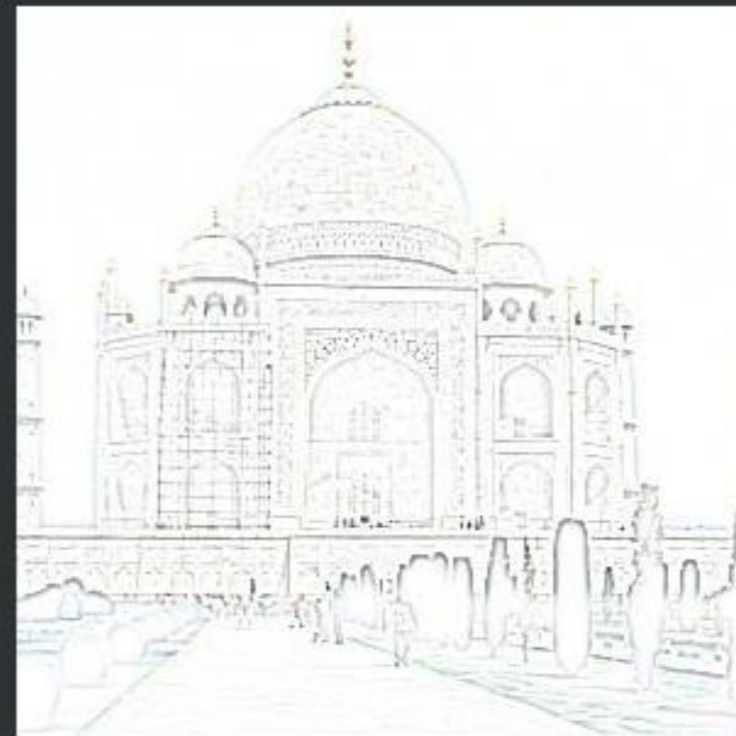
Spatial filtering

7.2.1. Overview

Figure 16.136. Applying example for the “Difference of Gaussians” filter

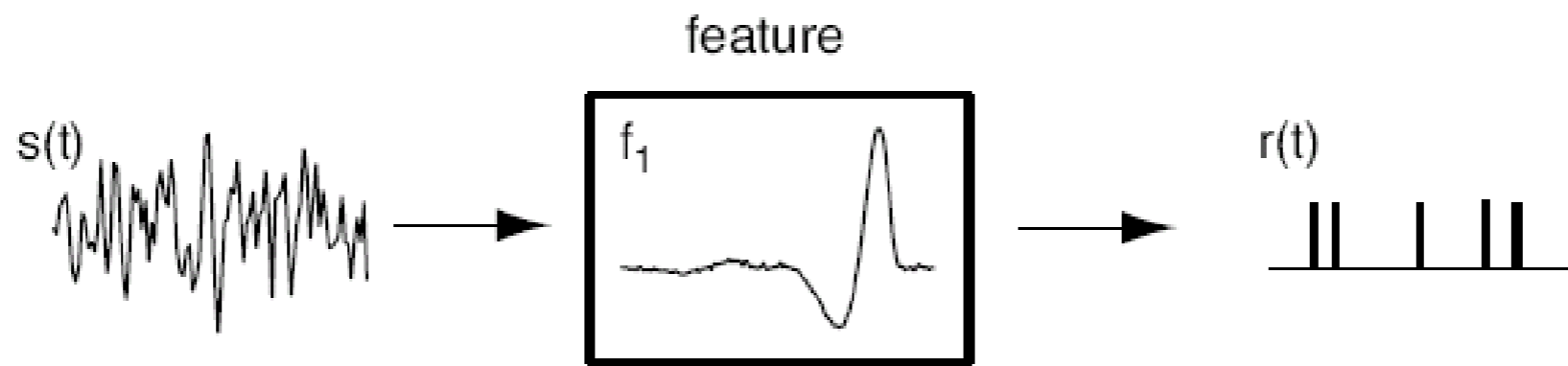


Original image



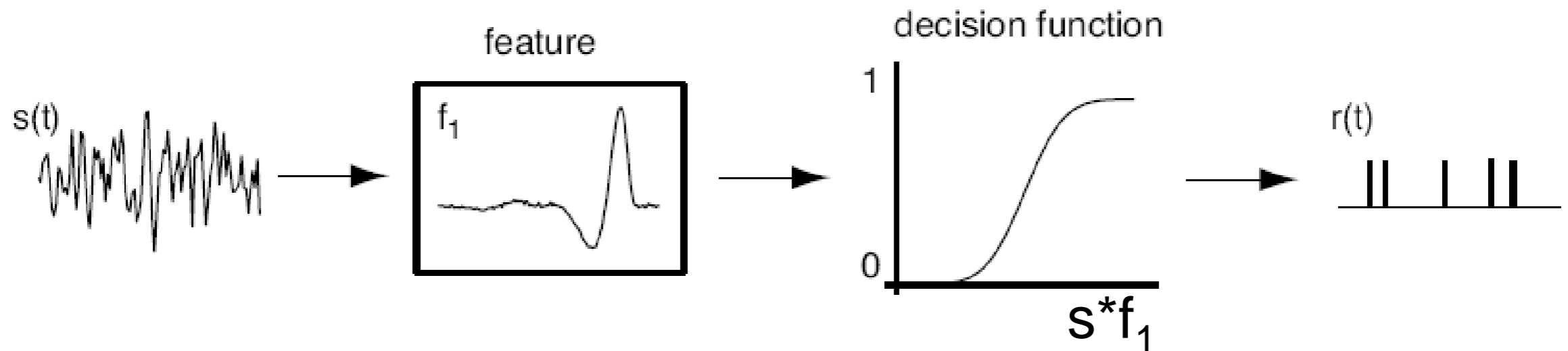
Filter “Difference of Gaussians” applied

Basic coding model: temporal filtering



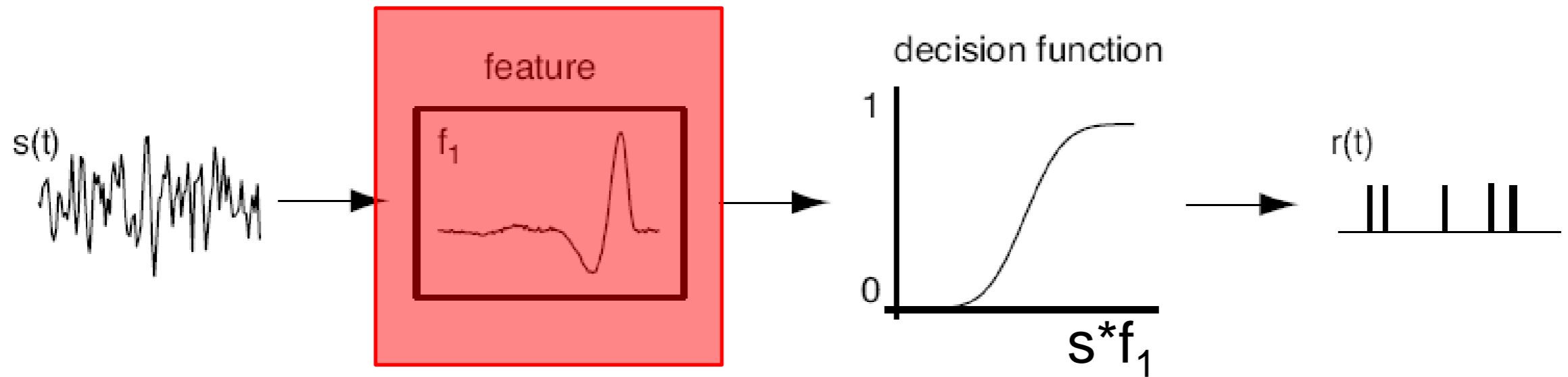
Linear filter:
$$r(t) = \int s(t-\tau) f(\tau) dt$$

Next most basic coding model

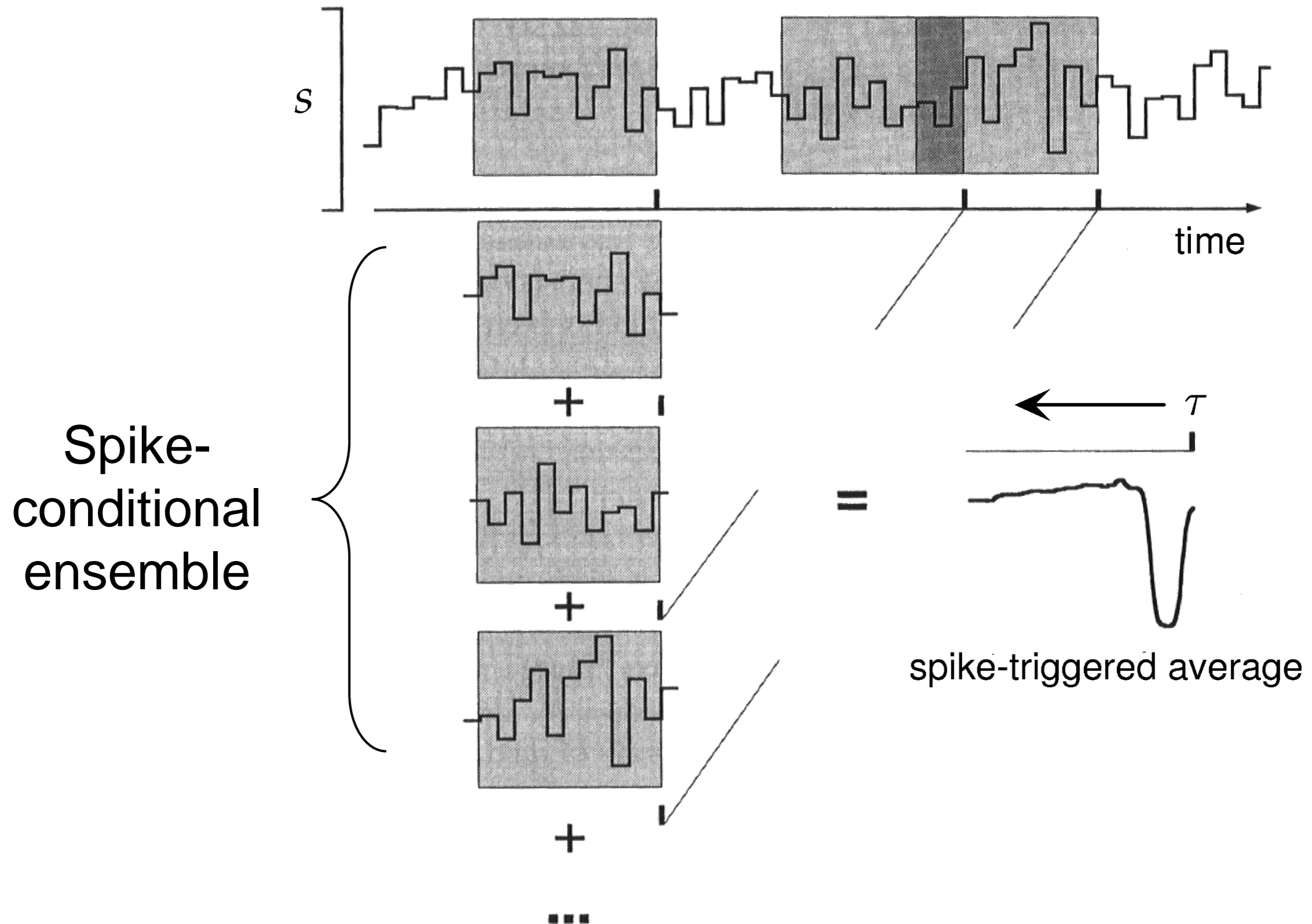


Linear filter & nonlinearity: $r(t) = g\left(\int s(t-\tau) f(\tau) d\tau\right)$

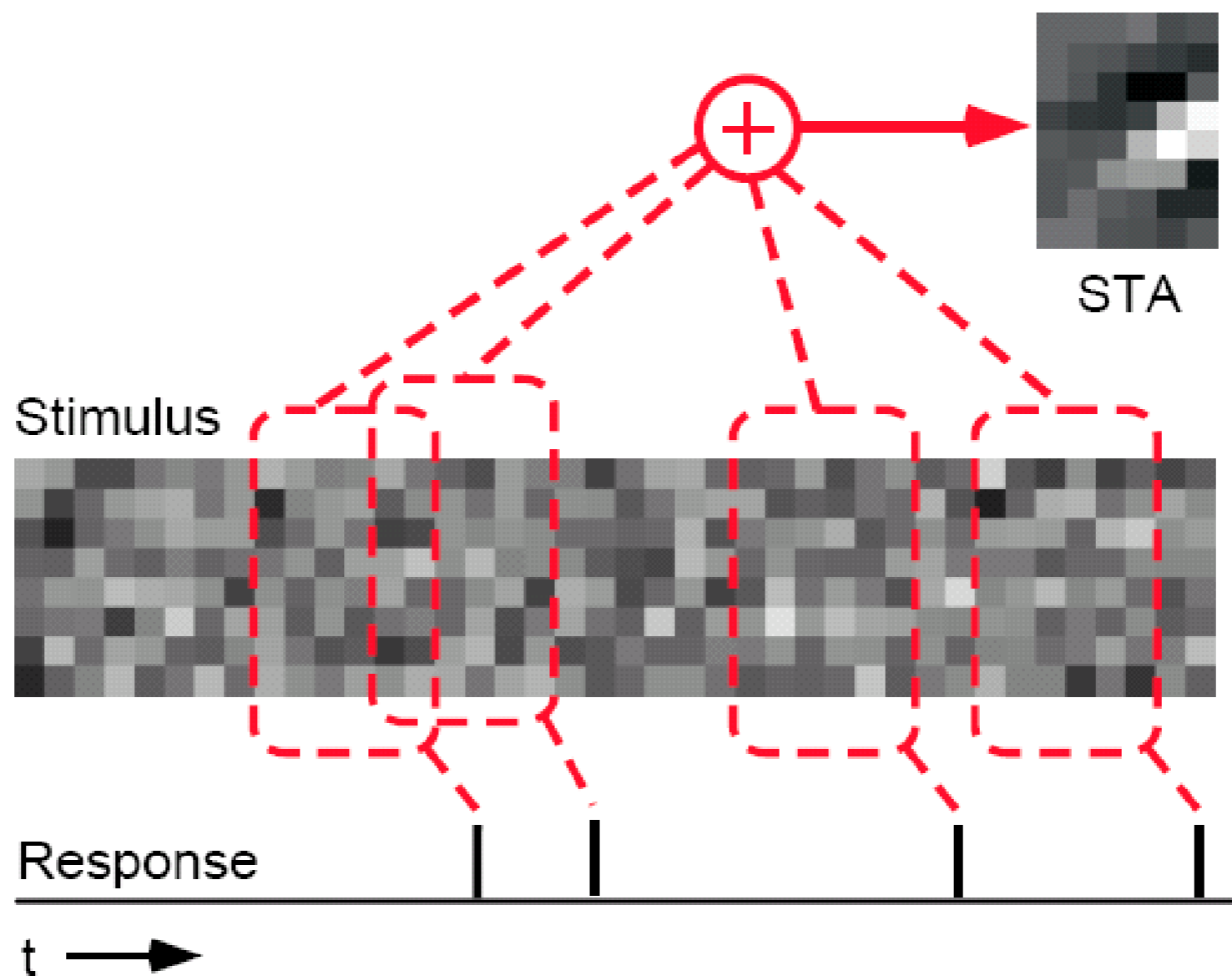
How to find the components of this model



Reverse correlation: the spike-triggered average



The spike-triggered average

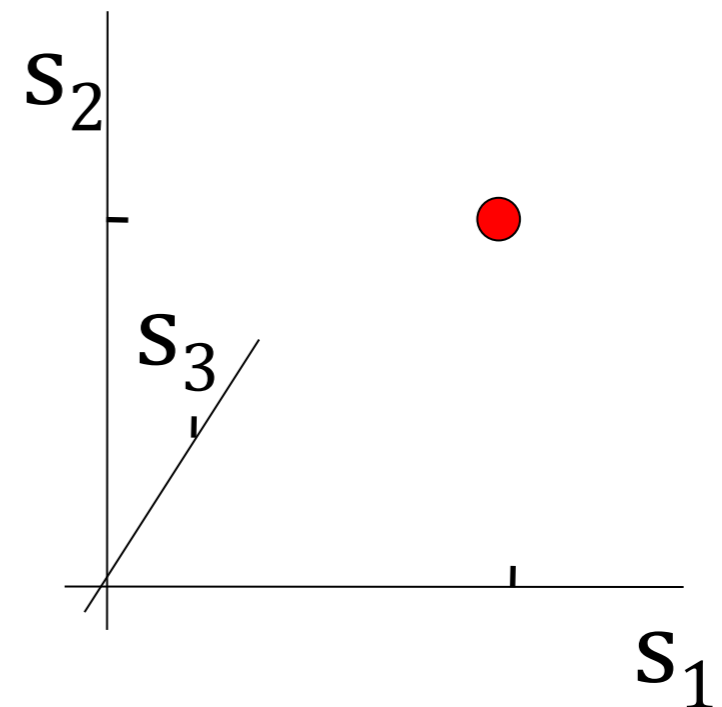
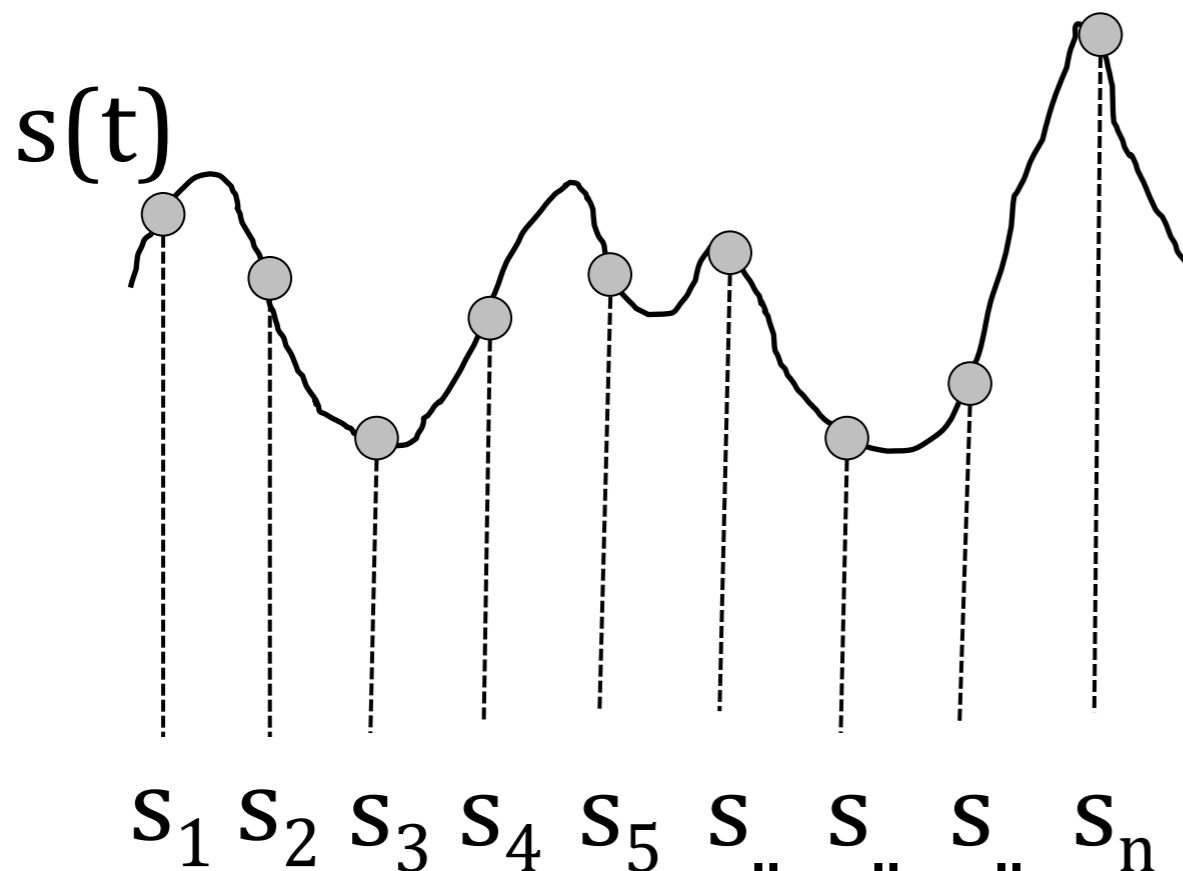


Dimensionality reduction

More generally, one can conceive of the action of the neuron or neural system as *selecting a low dimensional subset* of its inputs.

$$P(\text{response} \mid \text{stimulus}) \rightarrow P(\text{response} \mid s_1, s_2, \dots, s_n)$$

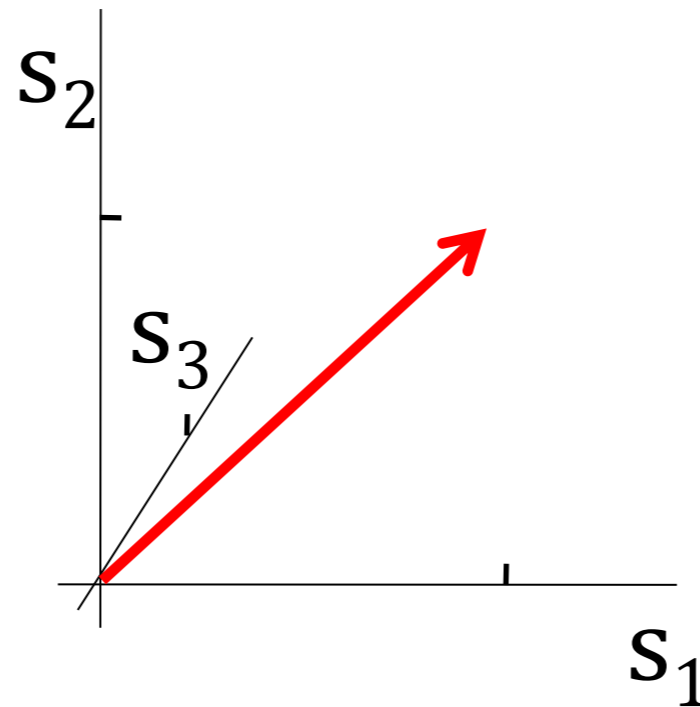
Start with a very high dimensional description (eg. an image or a time-varying waveform) and pick out a small set of relevant dimensions.



$$S(t) = (s_1, s_2, s_3, \dots, s_n)$$

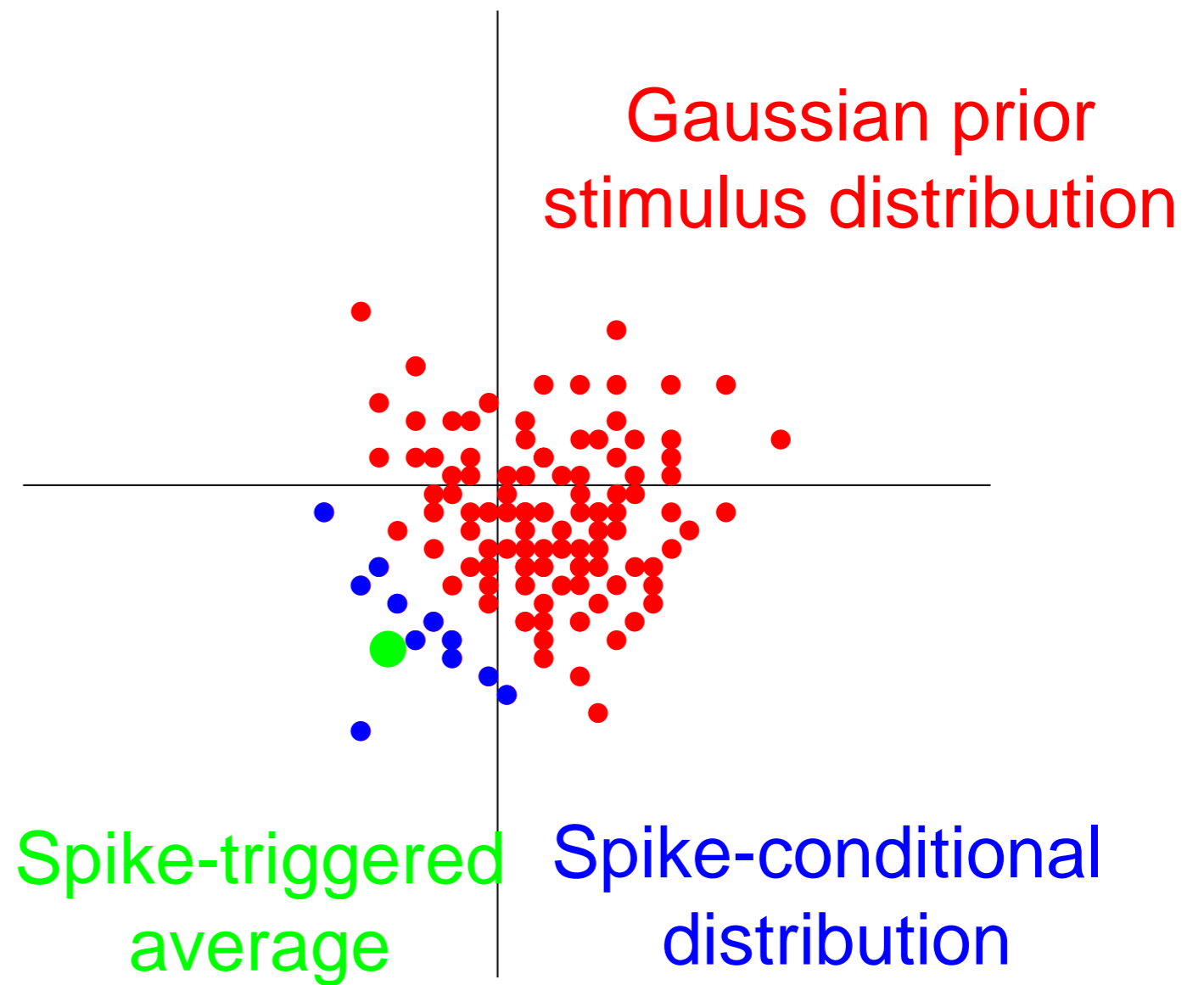
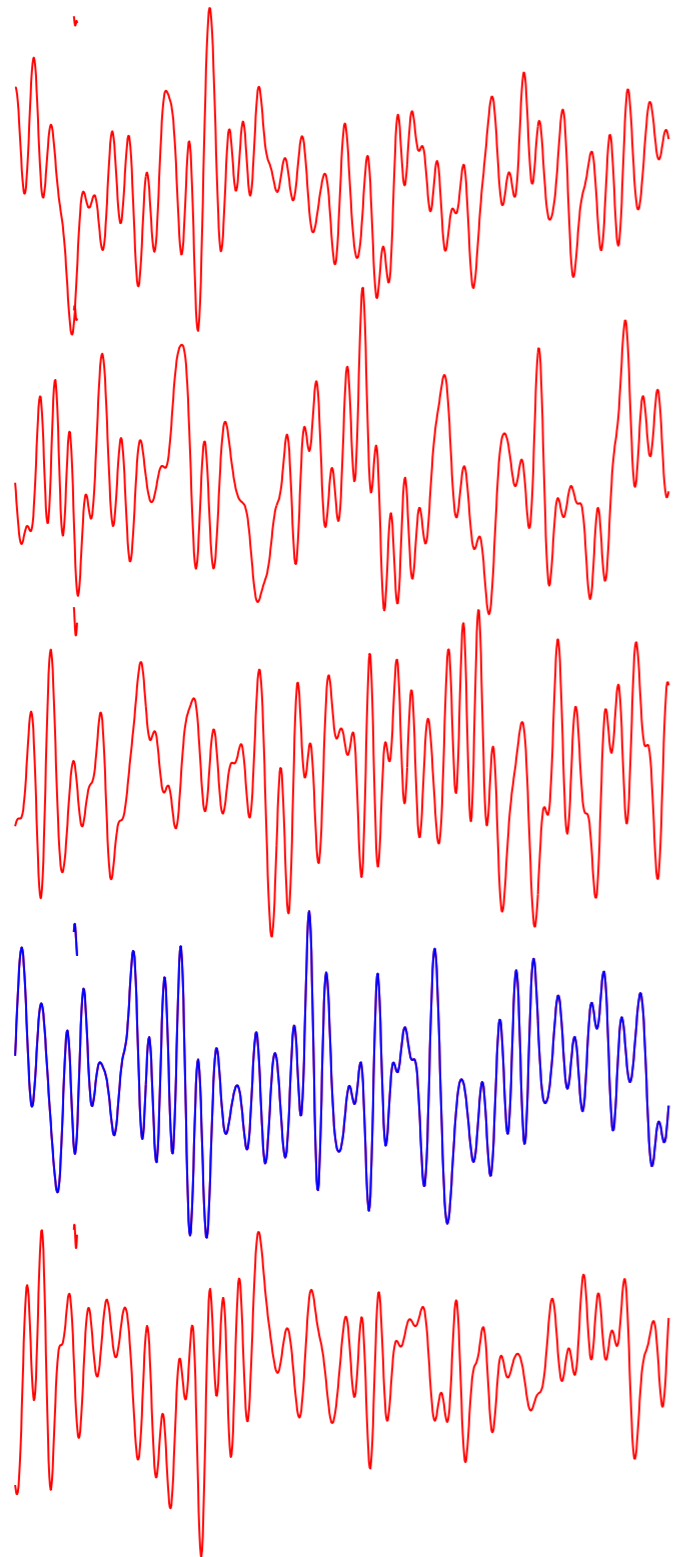
Linear filtering

Linear filtering = convolution = projection

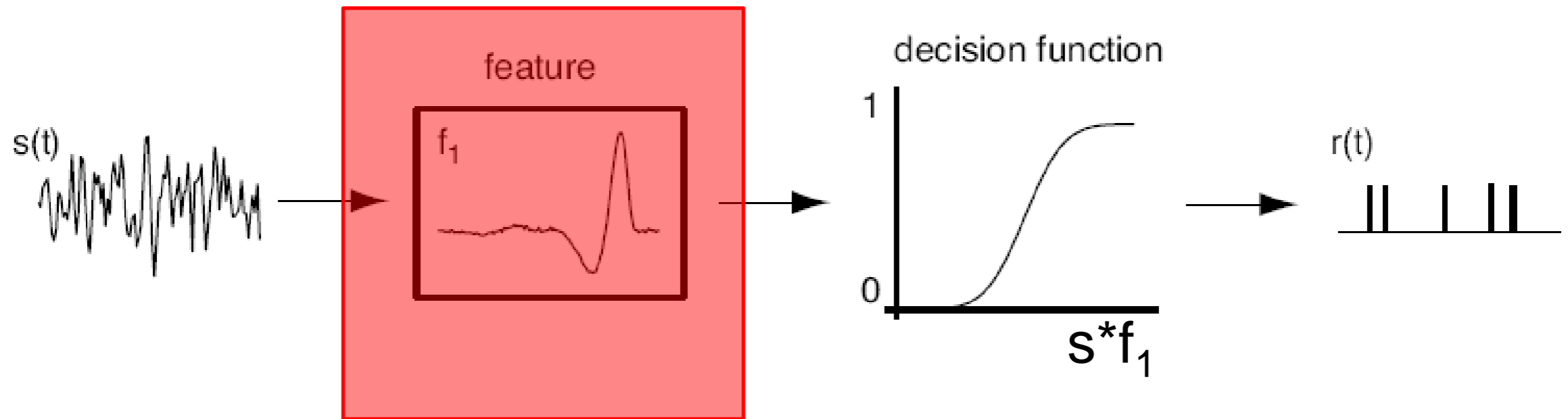


Stimulus feature is a vector in a high-dimensional stimulus space

Determining linear features from white noise



How to find the components of this model



Determining the nonlinear input/output function

The input/output function is:

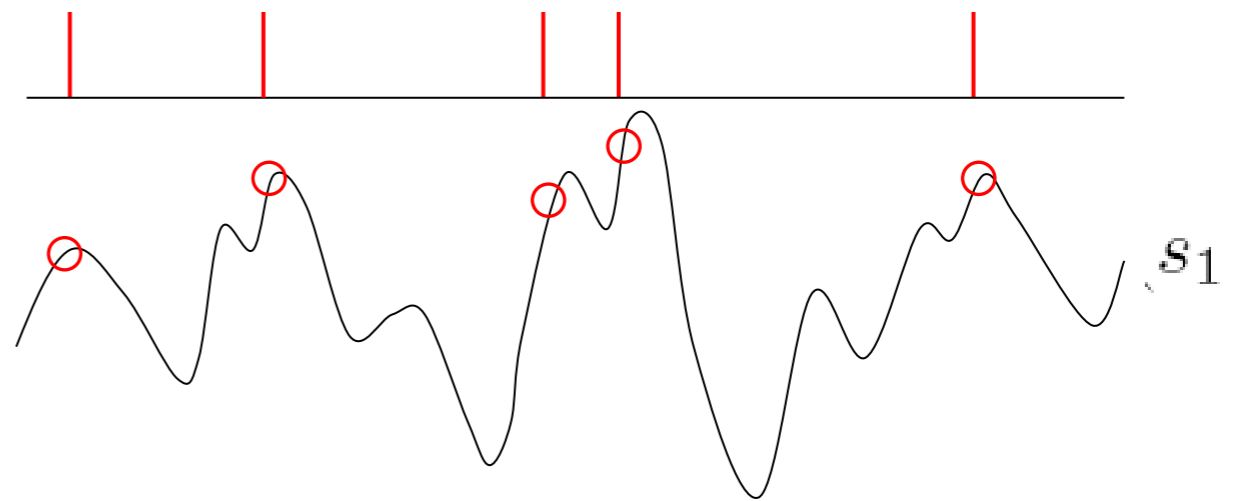
$$P(\text{spike}|\text{stimulus})$$

which can be derived from data using Bayes' rule:

$$P(\text{spike}|s_1) = \frac{P(s_1|\text{spike})P(\text{spike})}{P(s_1)}$$

$P(s_1)$

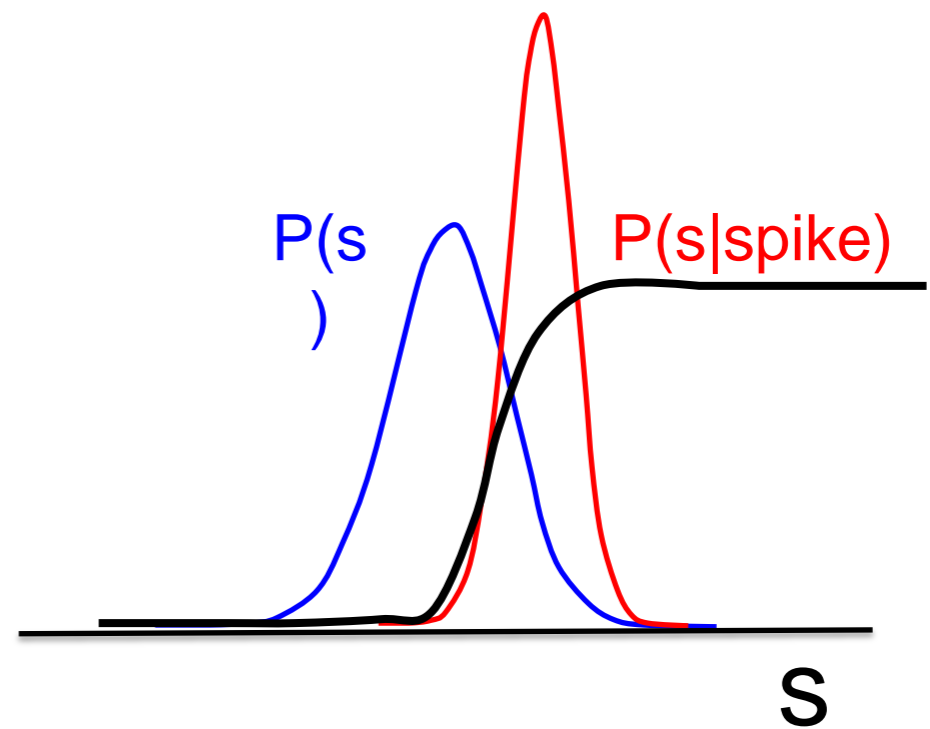
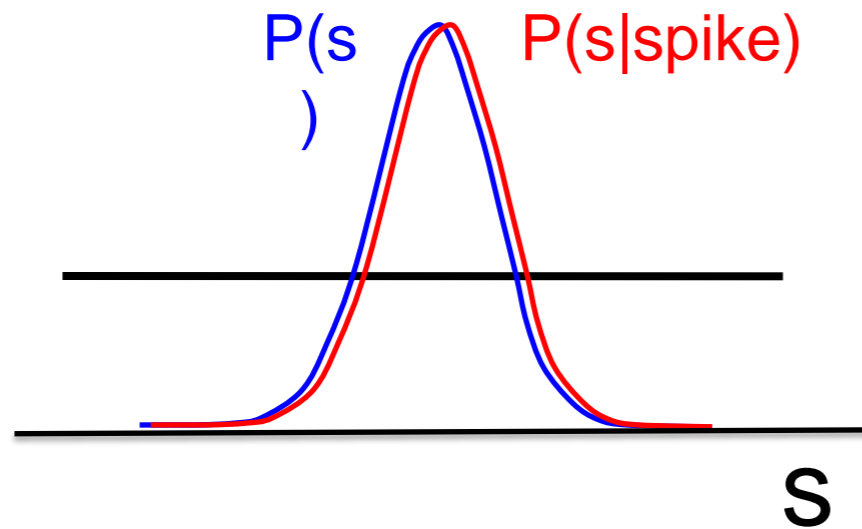
$P(s_1|\text{spike})$



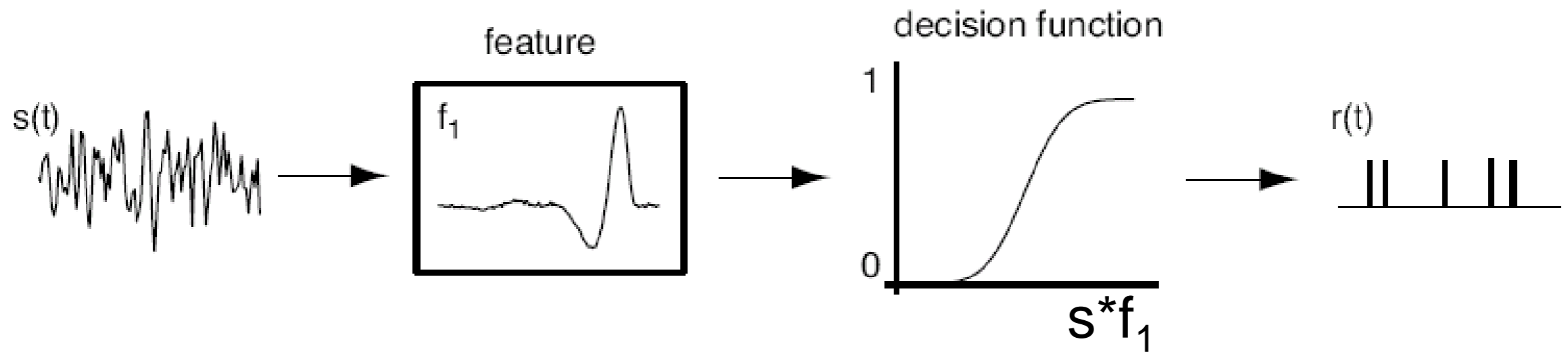
Nonlinear input/output function

Tuning curve:

$$P(\text{spike}|s) = P(s|\text{spike}) P(\text{spike}) / P(s)$$



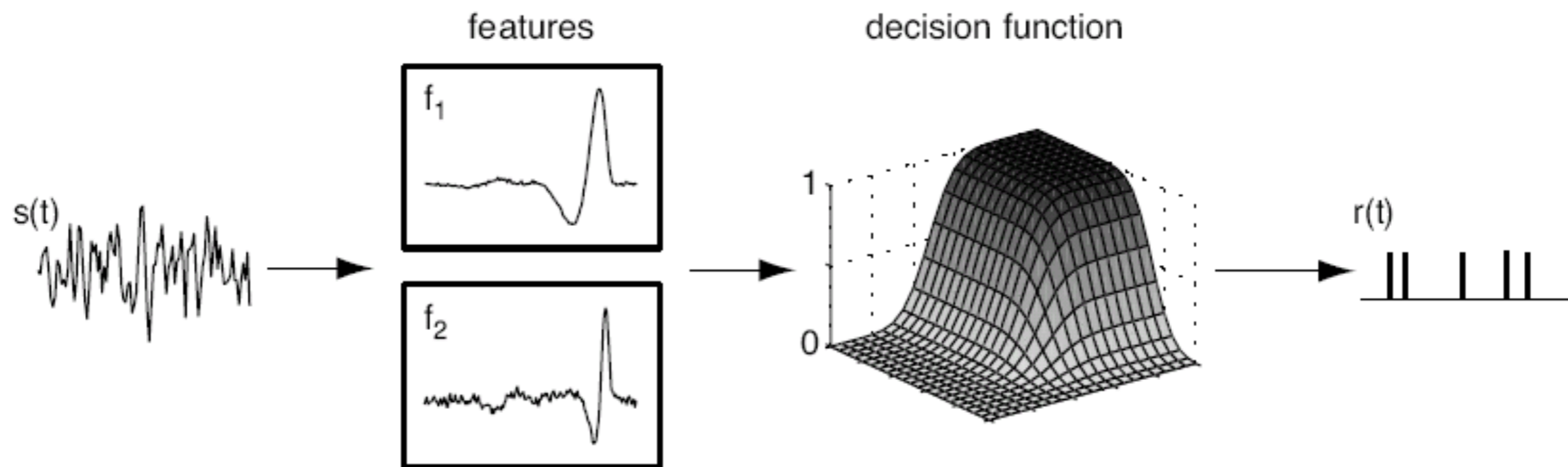
Next most basic coding model



Linear filter & nonlinearity: $r(t) = g\left(\int f(t-\tau) s(\tau) d\tau\right)$

...shortcomings?

Less basic coding models



Linear filters & nonlinearity: $r(t) = g(f_1 * s, f_2 * s, \dots, f_n * s)$

A bit of linear algebra

$$\mathbf{u} \bullet \mathbf{v} = \sum u_i v_i$$

$\mathbf{u} \bullet \mathbf{v}$ = projection of u onto v

$\mathbf{u} \bullet \mathbf{u}$ = length of u

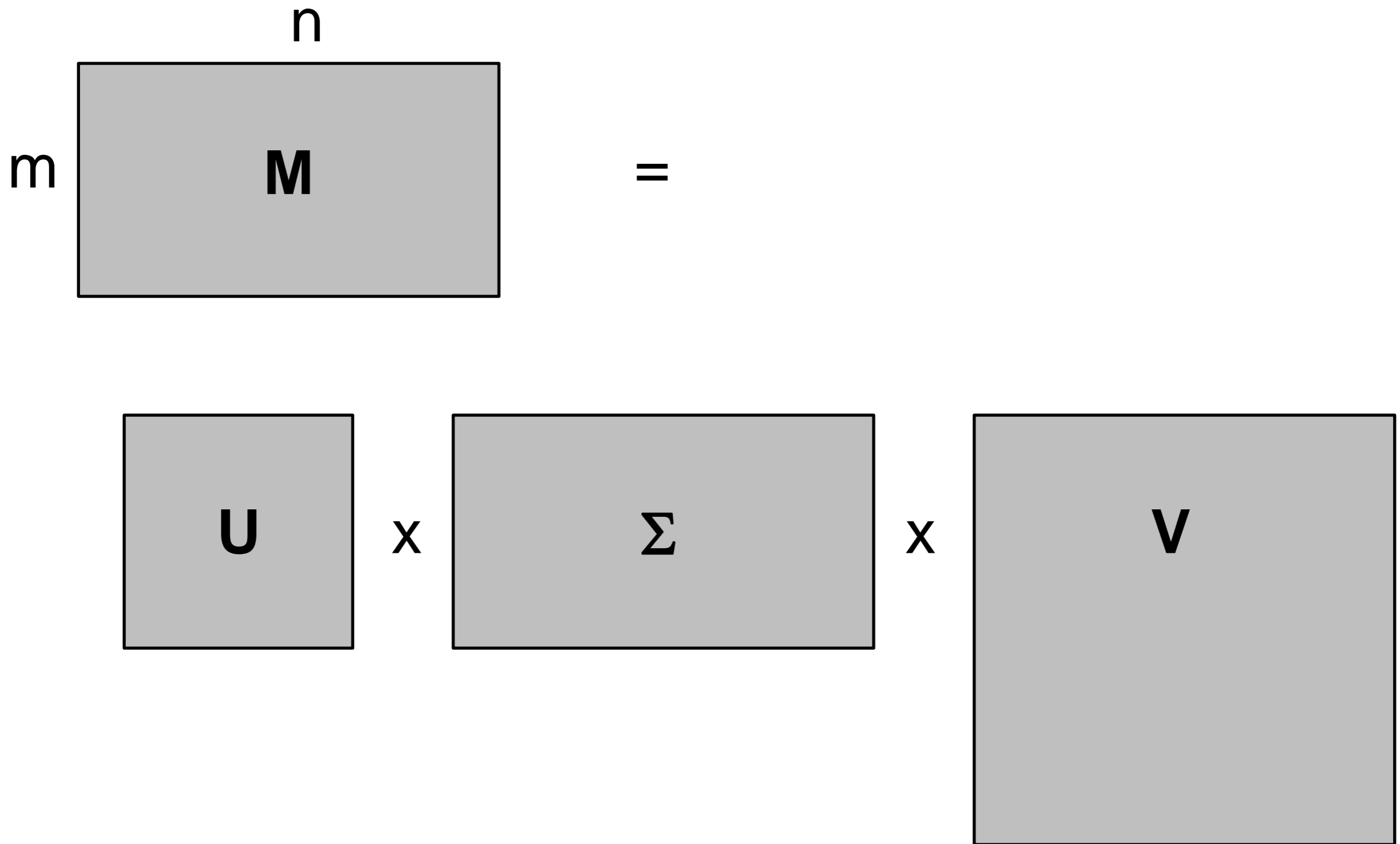
Linear filtering: $r(t) = \int f(t-\tau) s(\tau) d\tau$

$$\mathbf{u} \times \mathbf{v} = \mathbf{M} \quad \text{where } M_{ij} = u_i v_j$$

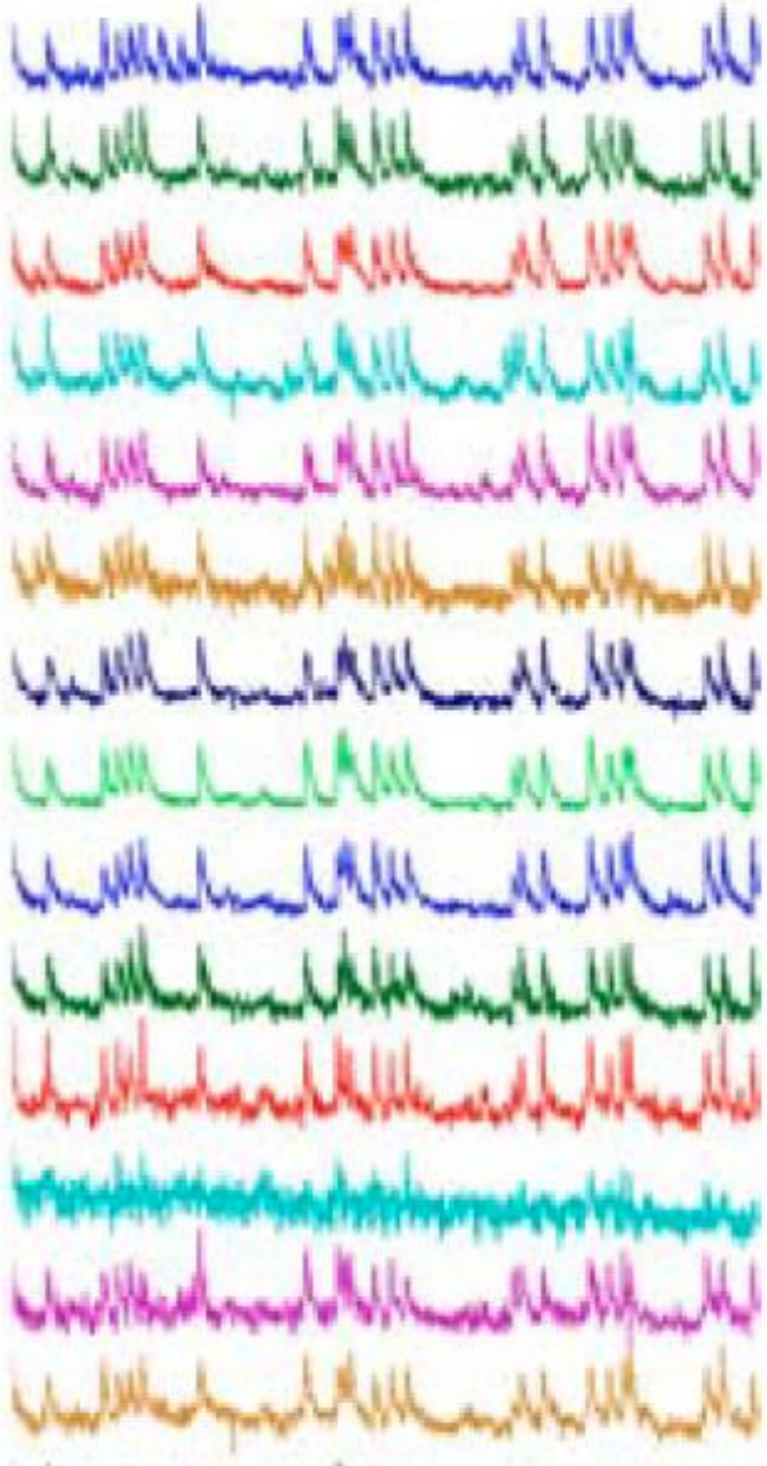
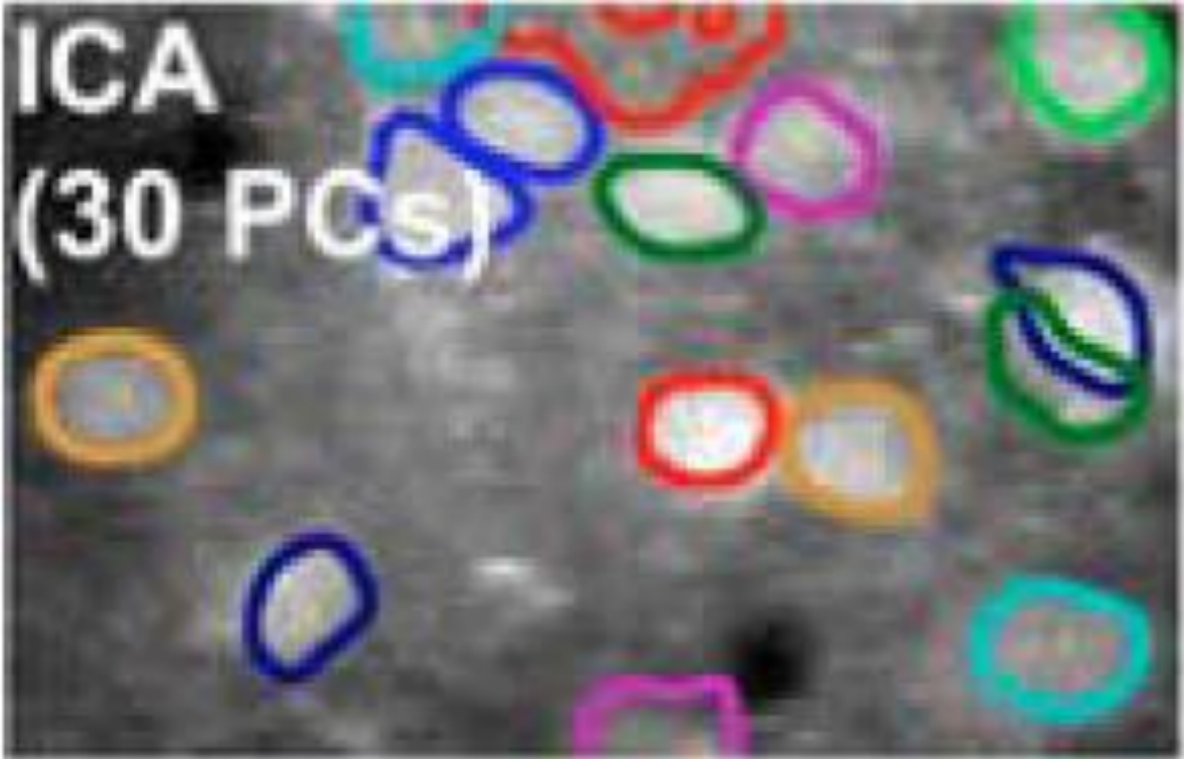
Eigenvalues and eigenvectors

$$\mathbf{M} \mathbf{u} = \lambda \mathbf{u}$$

Singular value decomposition



Identifying dynamical processes

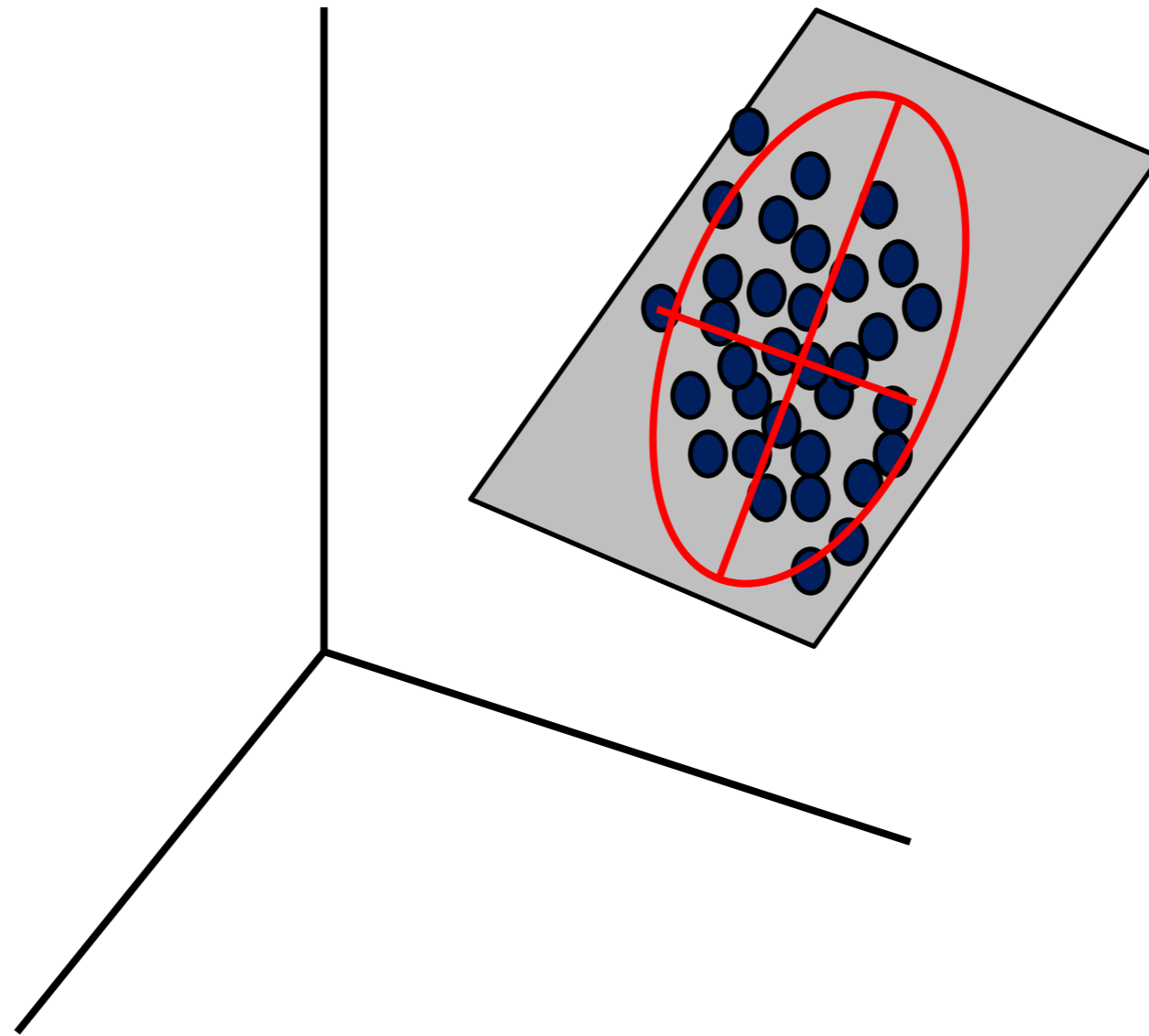


Principal component analysis

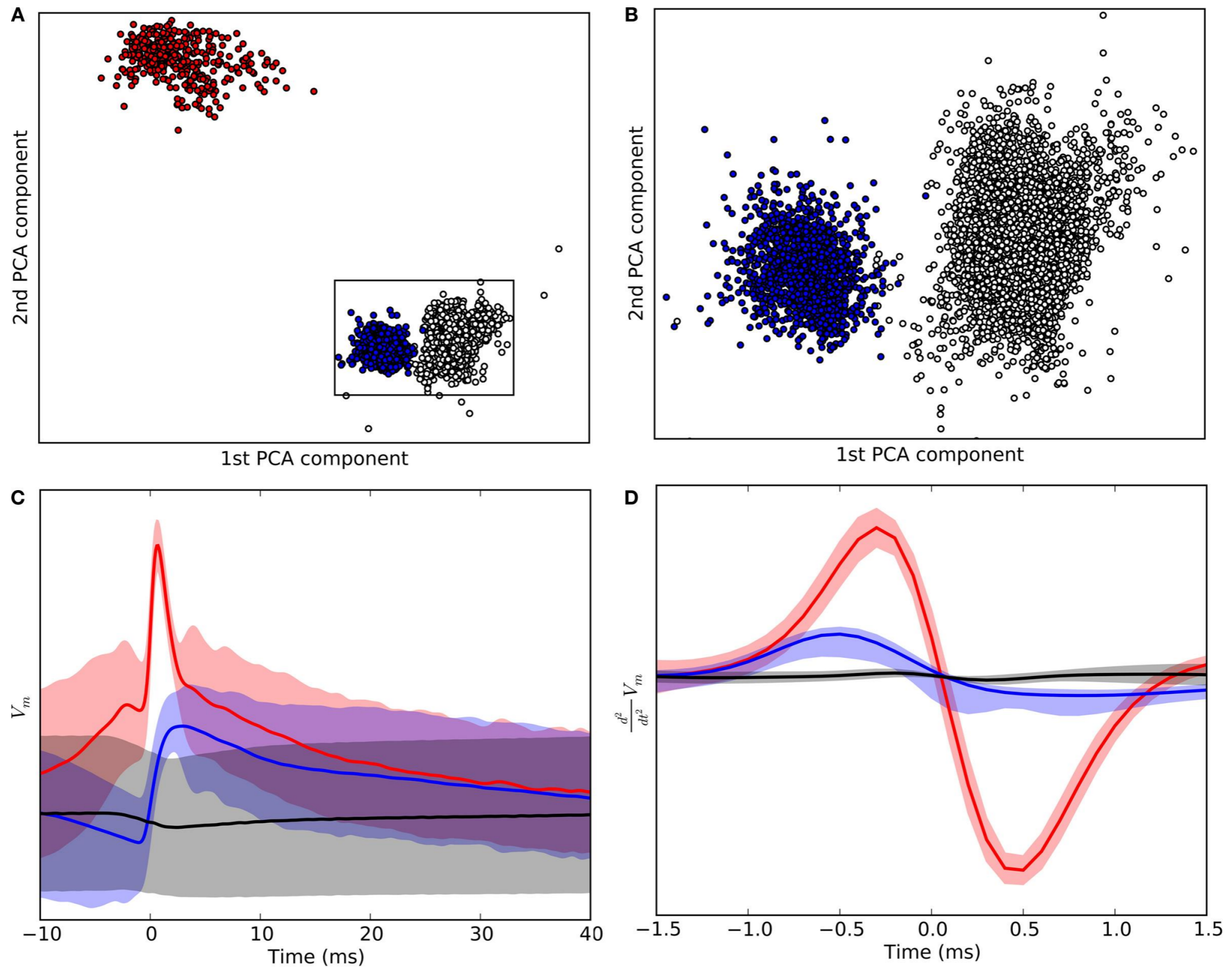
$$\begin{aligned} M M^* &= (U \Sigma V)(V^* \Sigma^* U^*) \\ &= U \Sigma (V V^*) \Sigma^* U^* \\ &= U \Sigma \Sigma^* U^* \end{aligned}$$

$$C = U \Lambda U^*$$

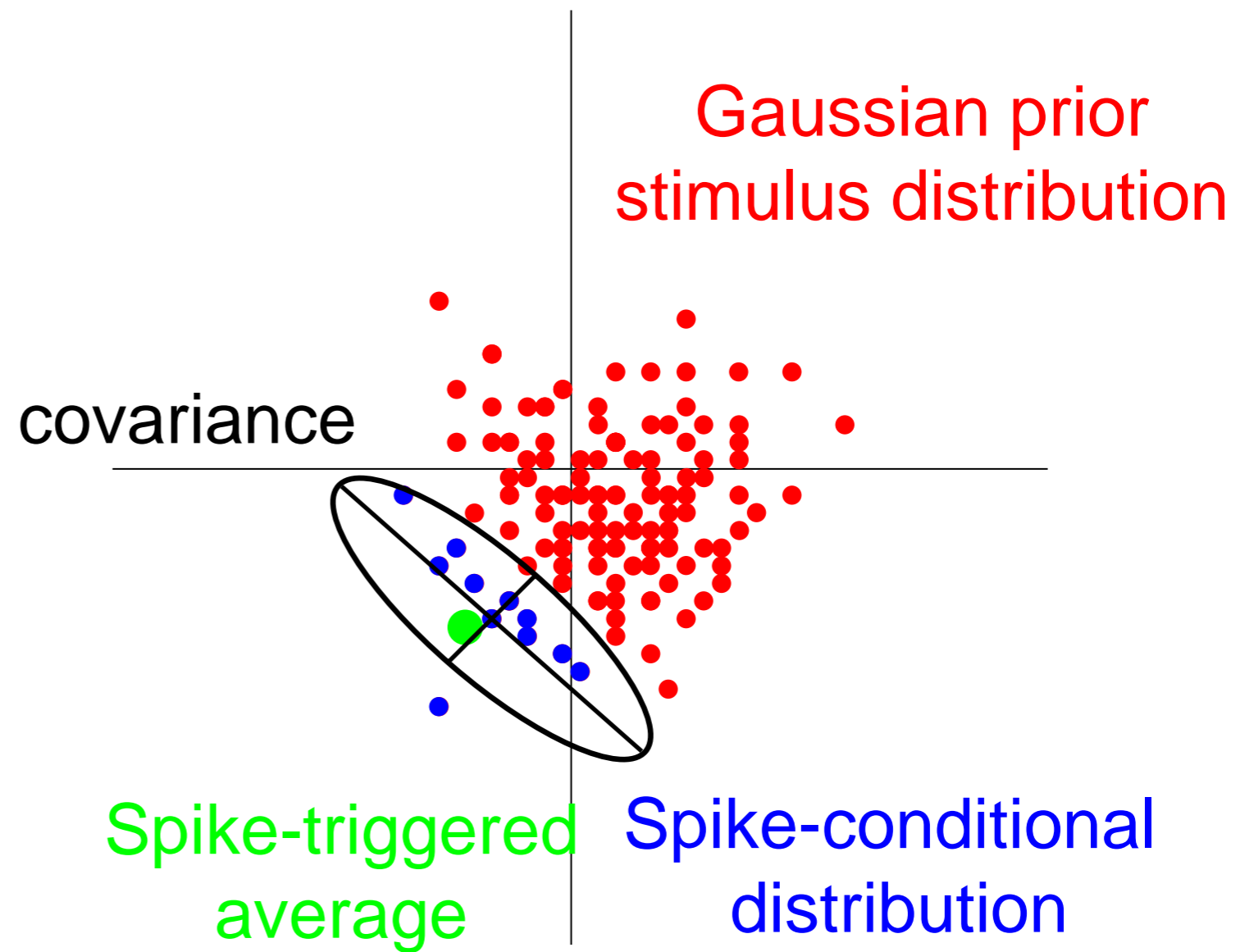
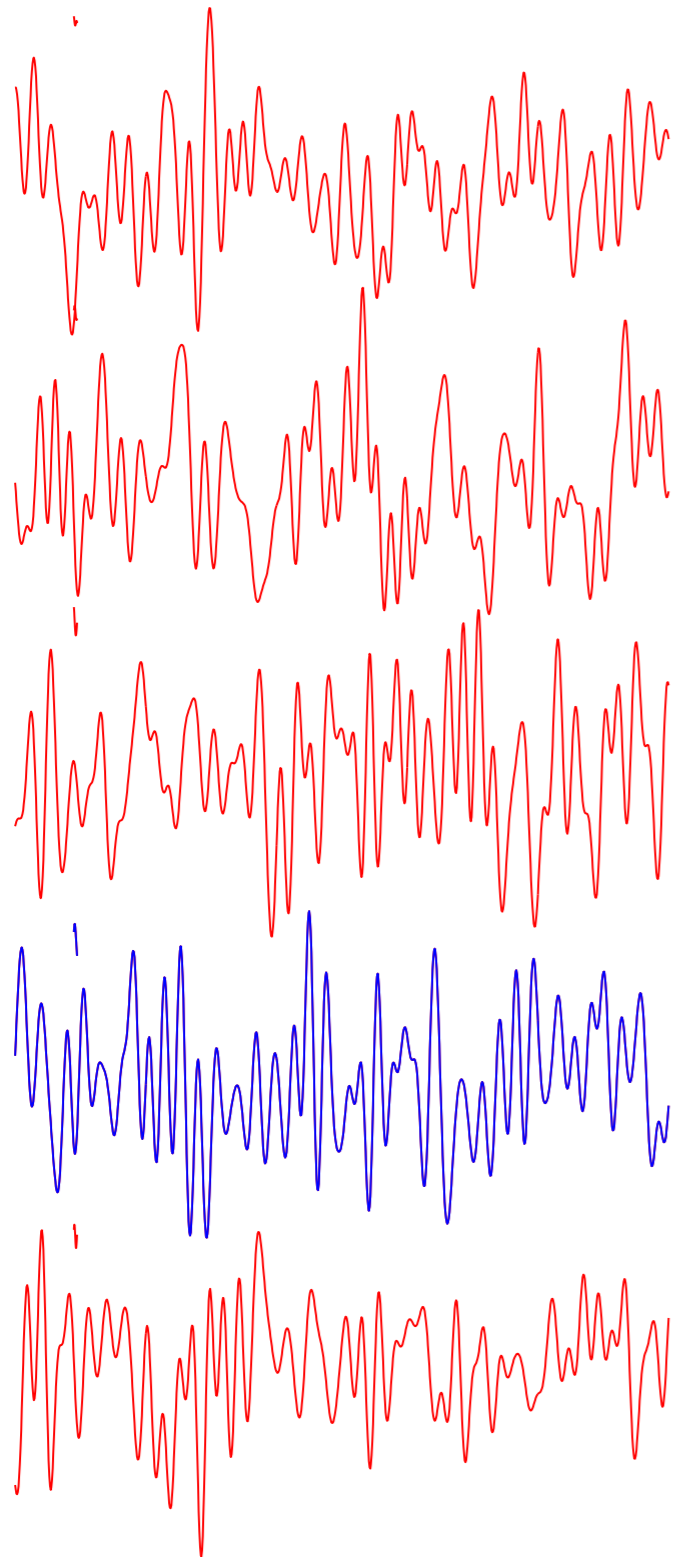
Principal component analysis



Principal component analysis



Determining linear features from white noise



Identifying multiple features

The covariance matrix is

$$C_{ij} = \langle S(t_{\text{spike}} - t_i) S(t_{\text{spike}} - t_j) \rangle - \bar{S}(t_i) \bar{S}(t_j) - \langle I(t - t_i) I(t - t_j) \rangle$$

↑
Spike-triggered stimulus
correlation

↑
Spike-triggered
average

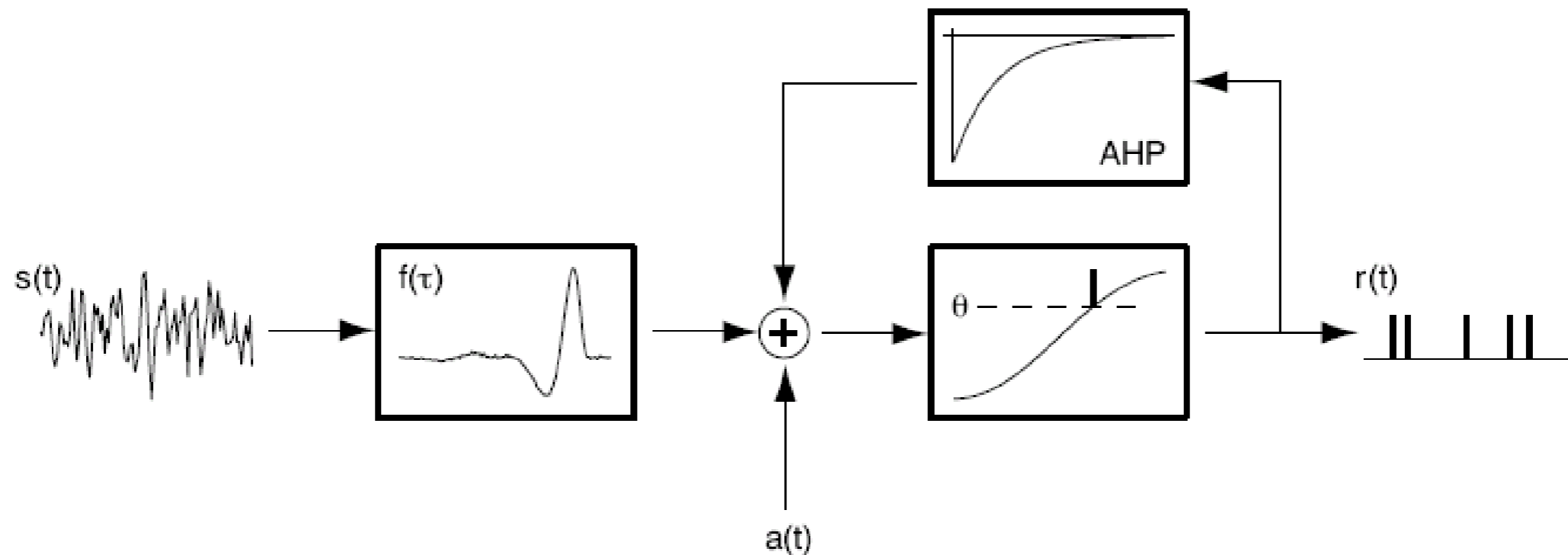
↑
Stimulus prior

Properties:

- The number of eigenvalues significantly **different from zero** is the number of relevant stimulus features
- The corresponding eigenvectors are the relevant features (or span the *relevant subspace*)

A toy example: a filter-and-fire model

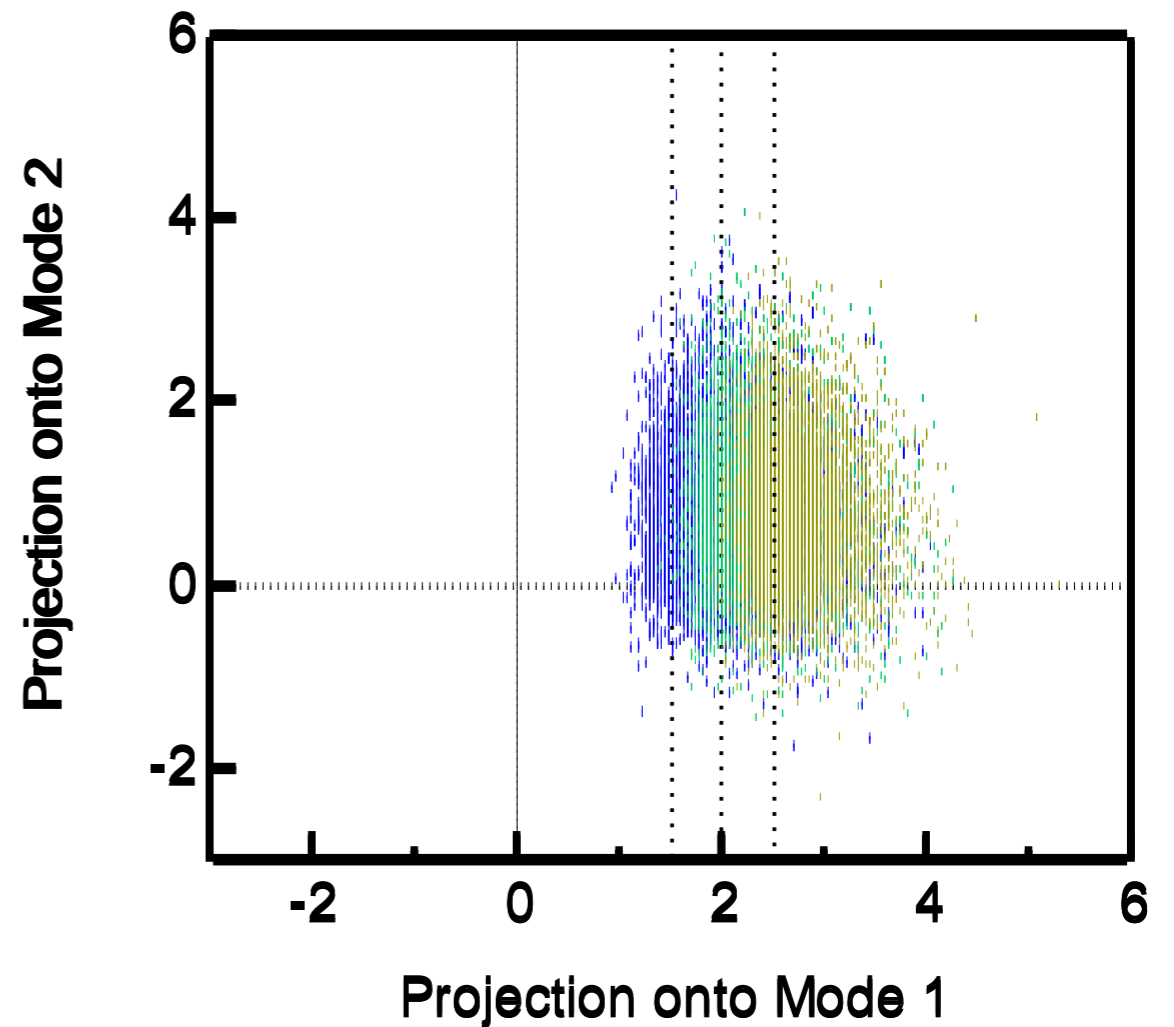
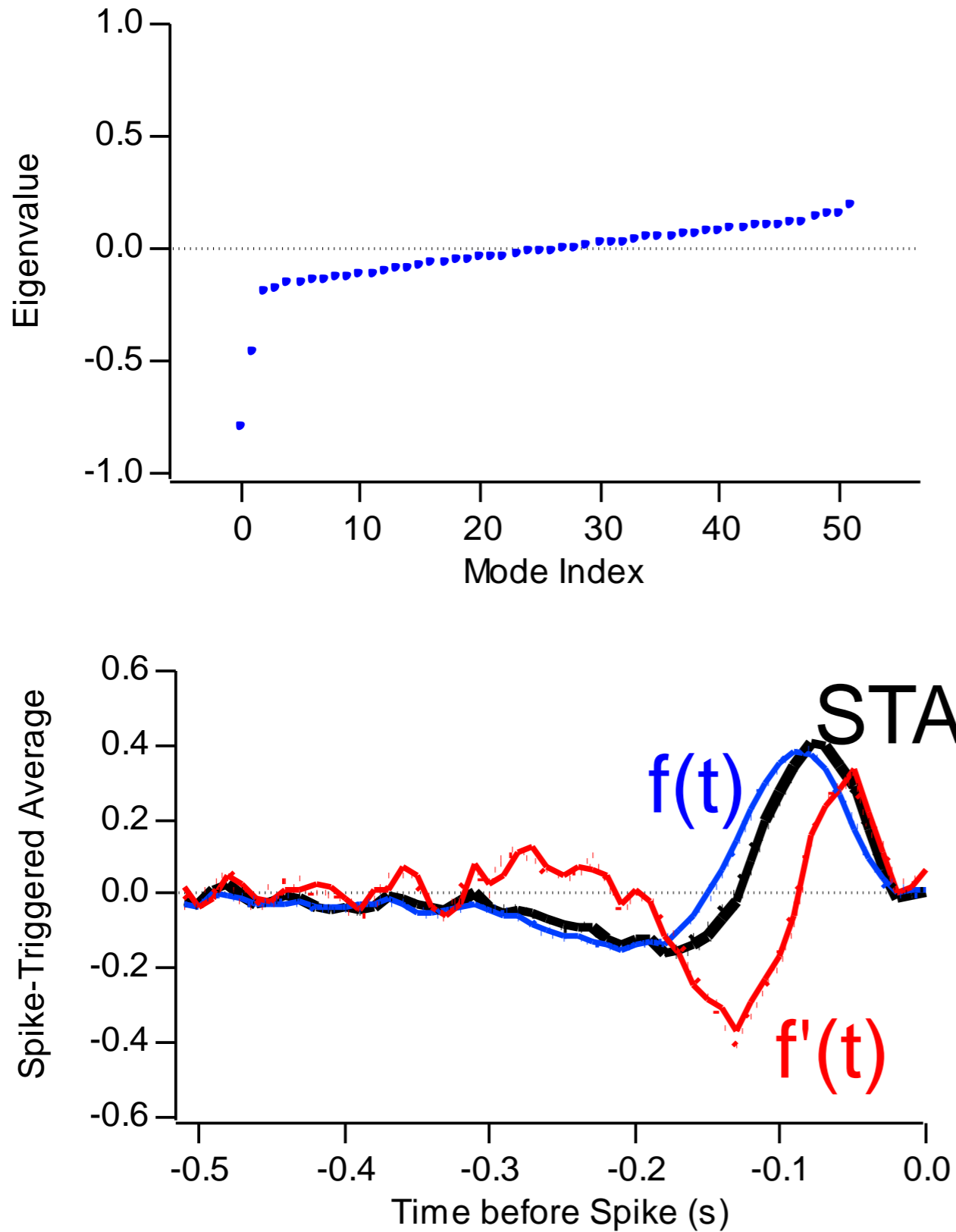
Let's develop some intuition for how this works: a filter-and-fire threshold-crossing model with AHP



Keat, Reinagel, Reid and Meister, Predicting every spike. Neuron (2001)

- Spiking is controlled by a single filter, $f(t)$
 - Spikes happen generally on an upward threshold crossing of the filtered stimulus
- ➔ expect 2 **relevant features**, the filter $f(t)$ and its time derivative $f'(t)$

Covariance analysis of a filter-and-fire model



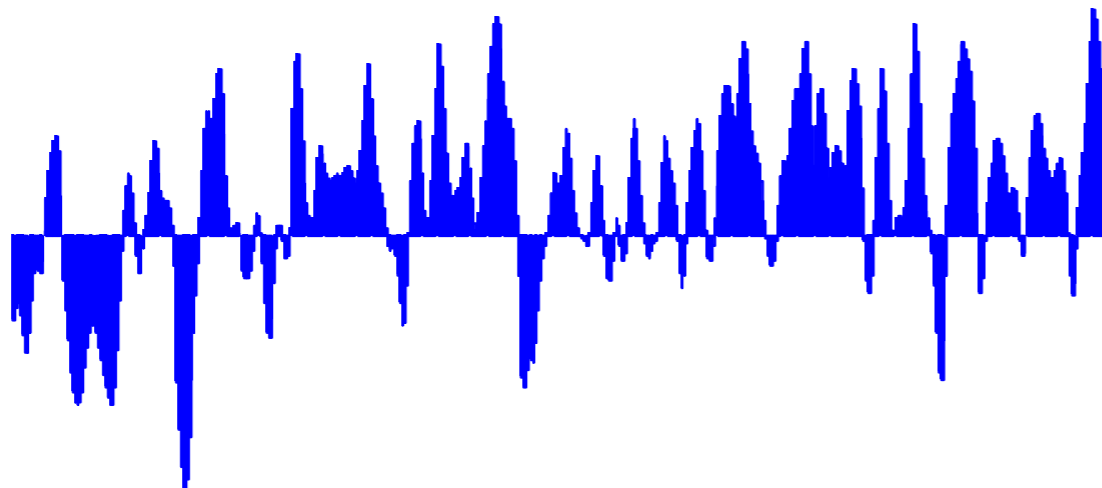
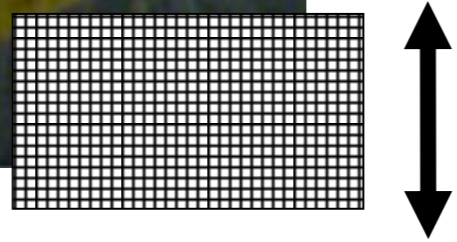
Let's try it

Example: rat somatosensory (barrel) cortex

Rasmus Petersen and Mathew Diamond, SISSA

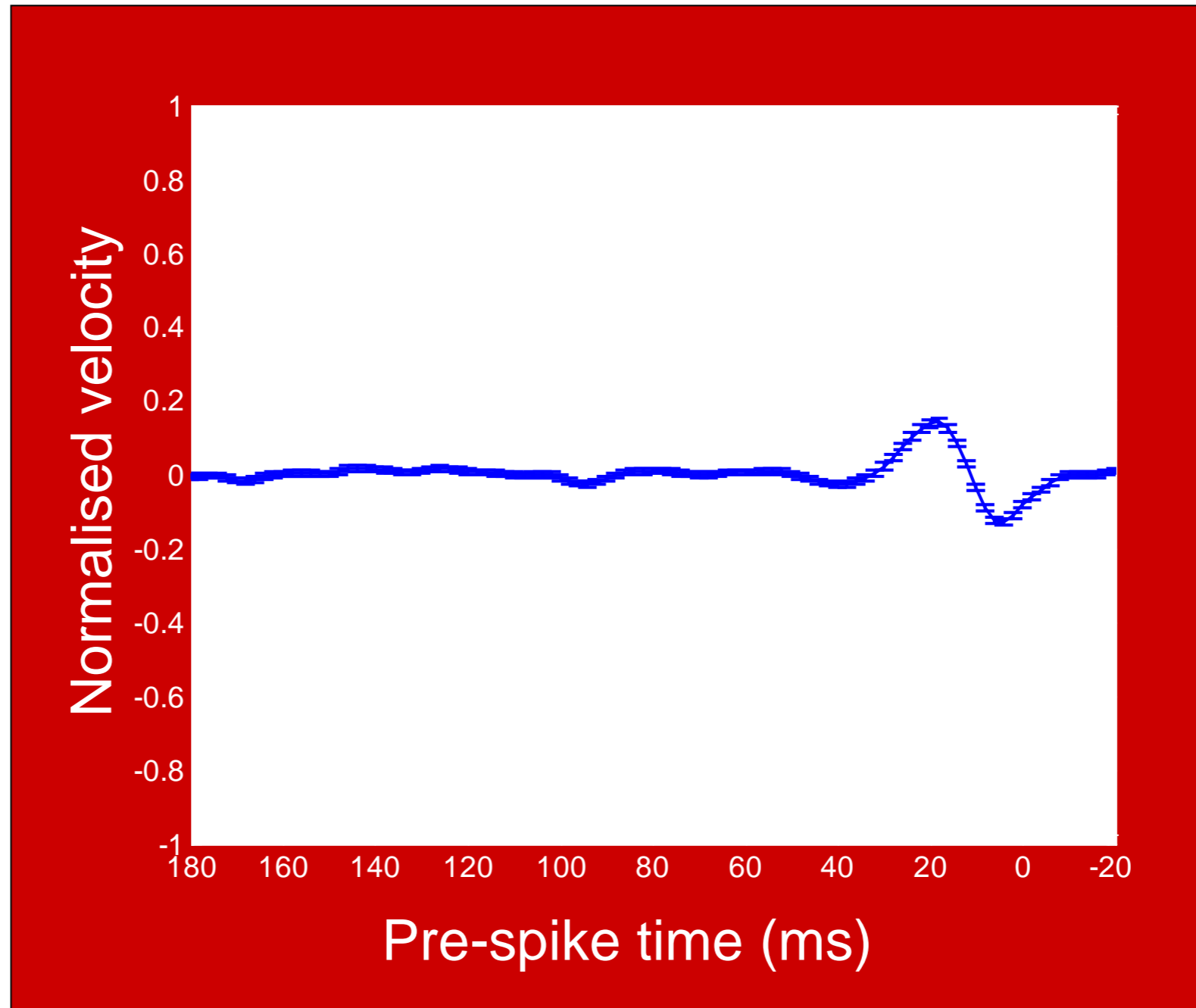


Record from single units in barrel cortex



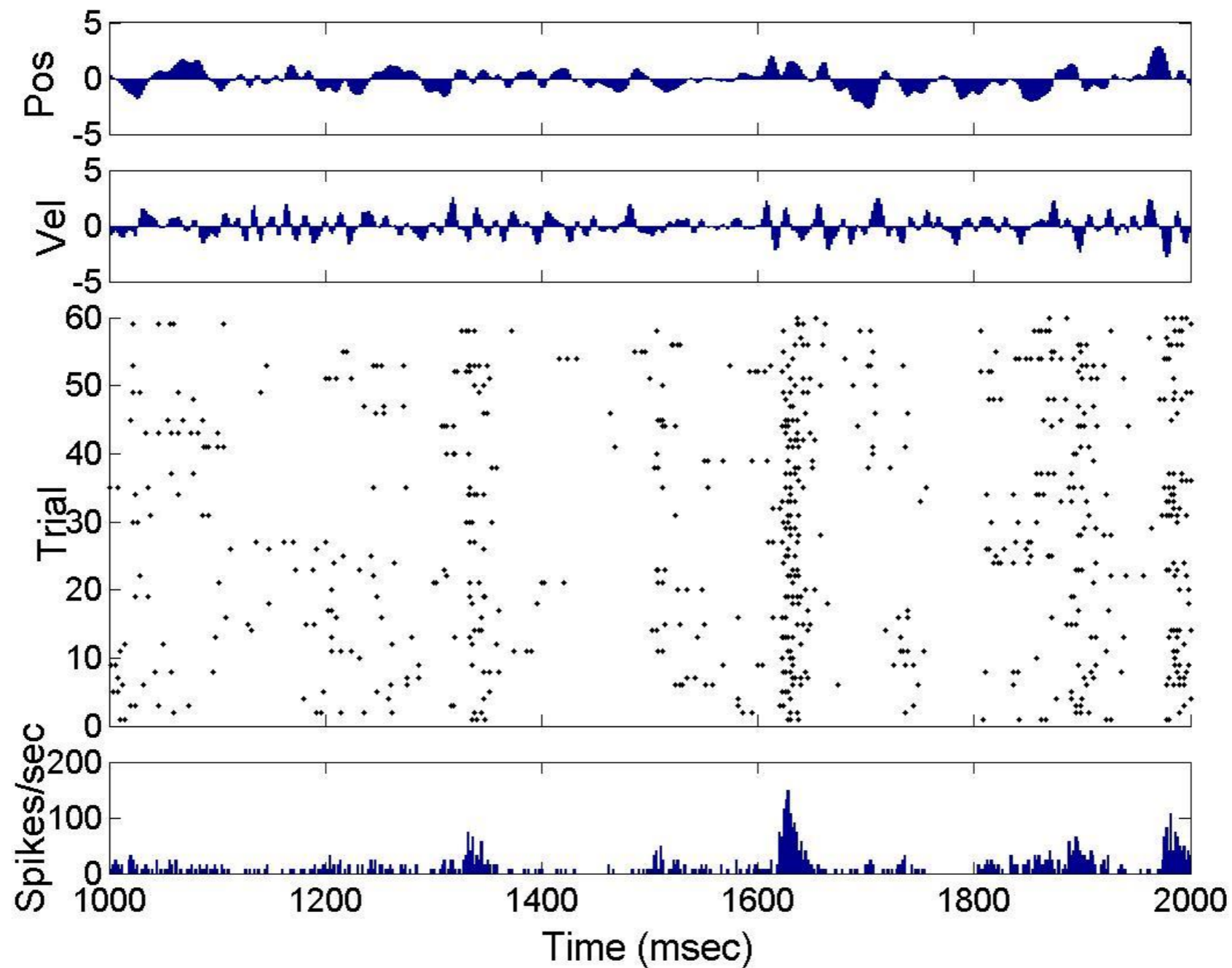
White noise analysis in barrel cortex

Spike-triggered average:



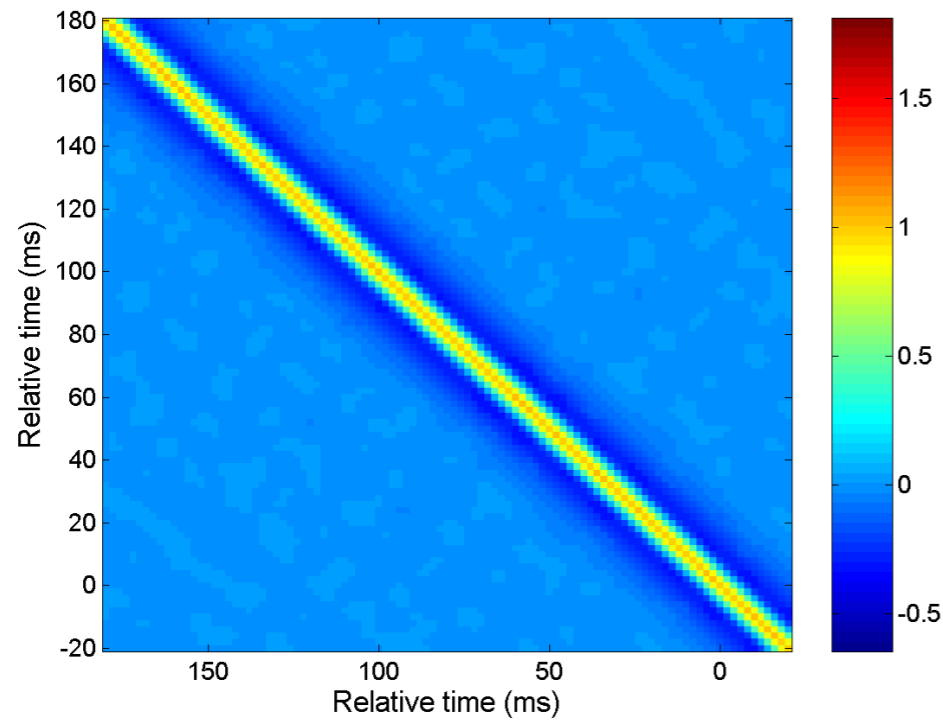
White noise analysis in barrel cortex

Is the neuron simply not very responsive to a white noise stimulus?

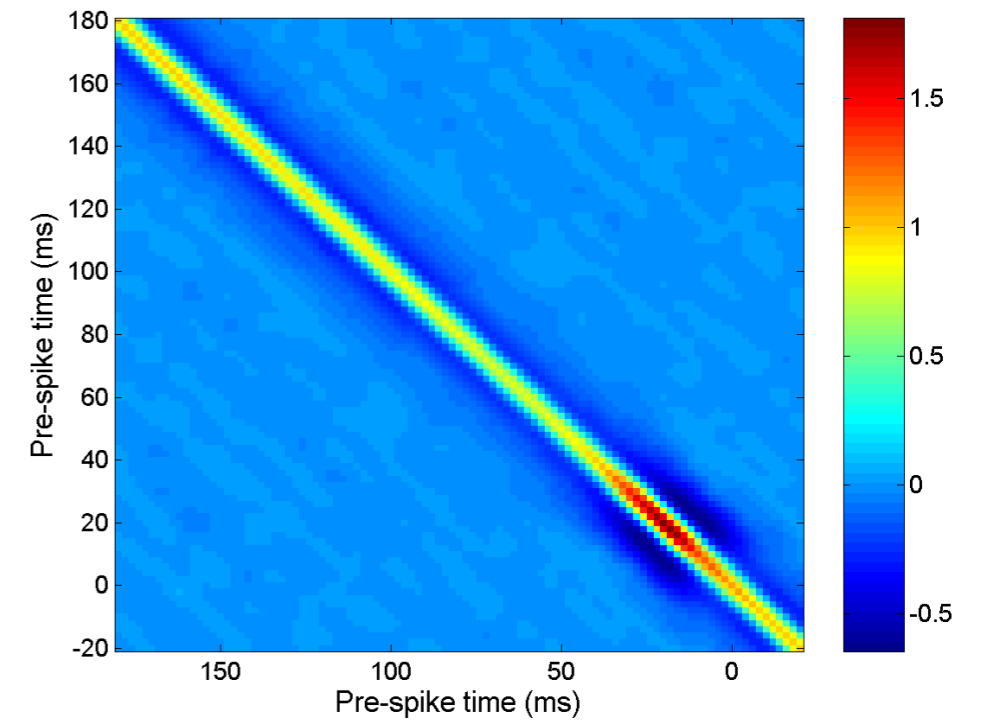


Covariance matrices from barrel cortical neurons

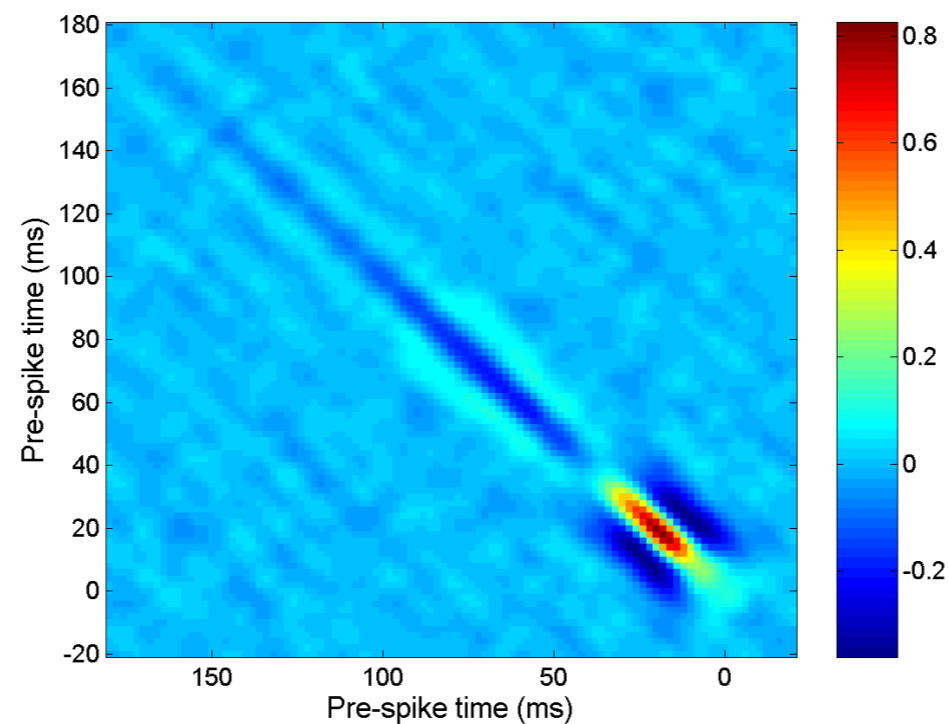
Prior



Spike-triggered

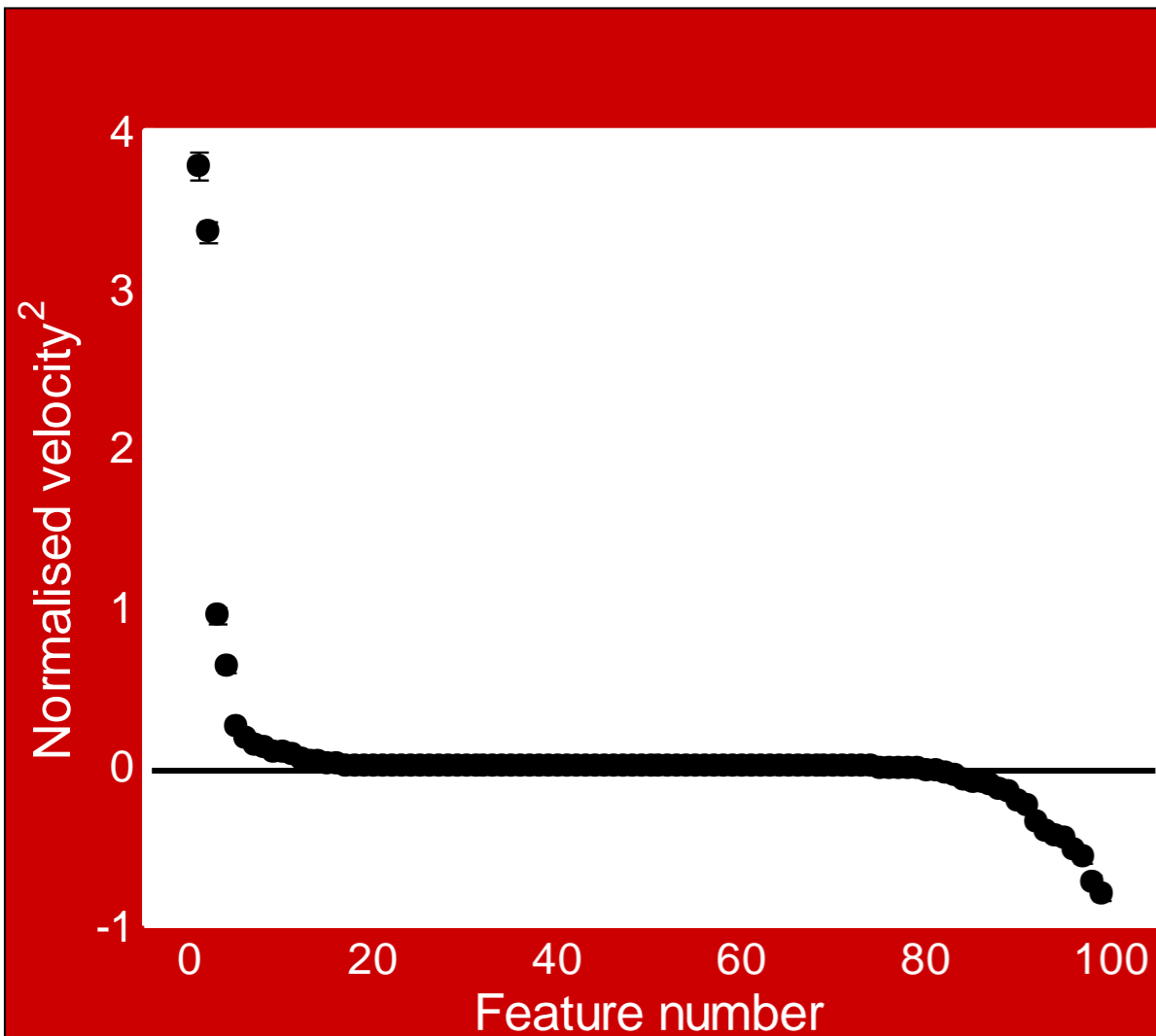


Difference

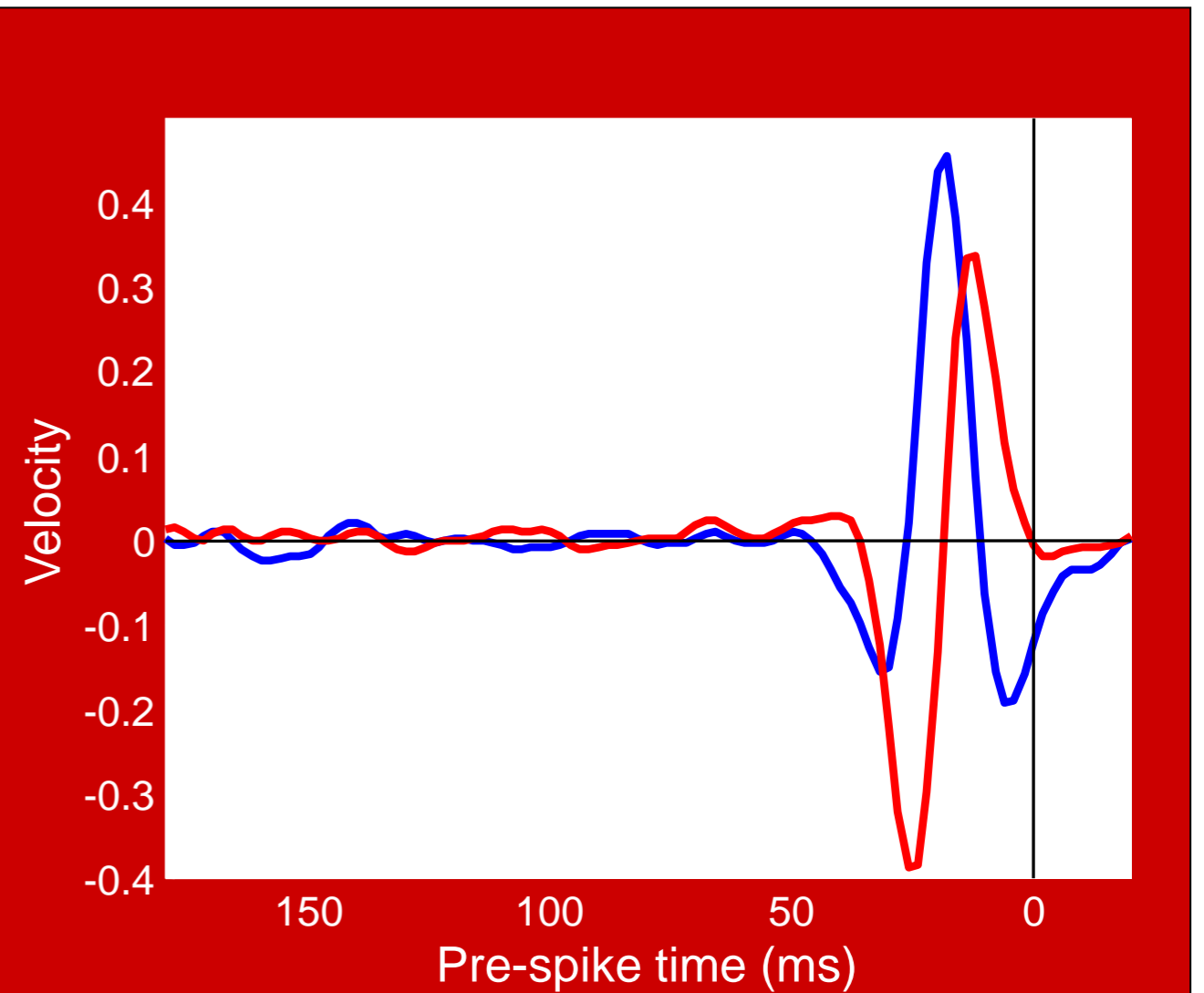


Eigenspectrum from barrel cortical neurons

Eigenspectrum

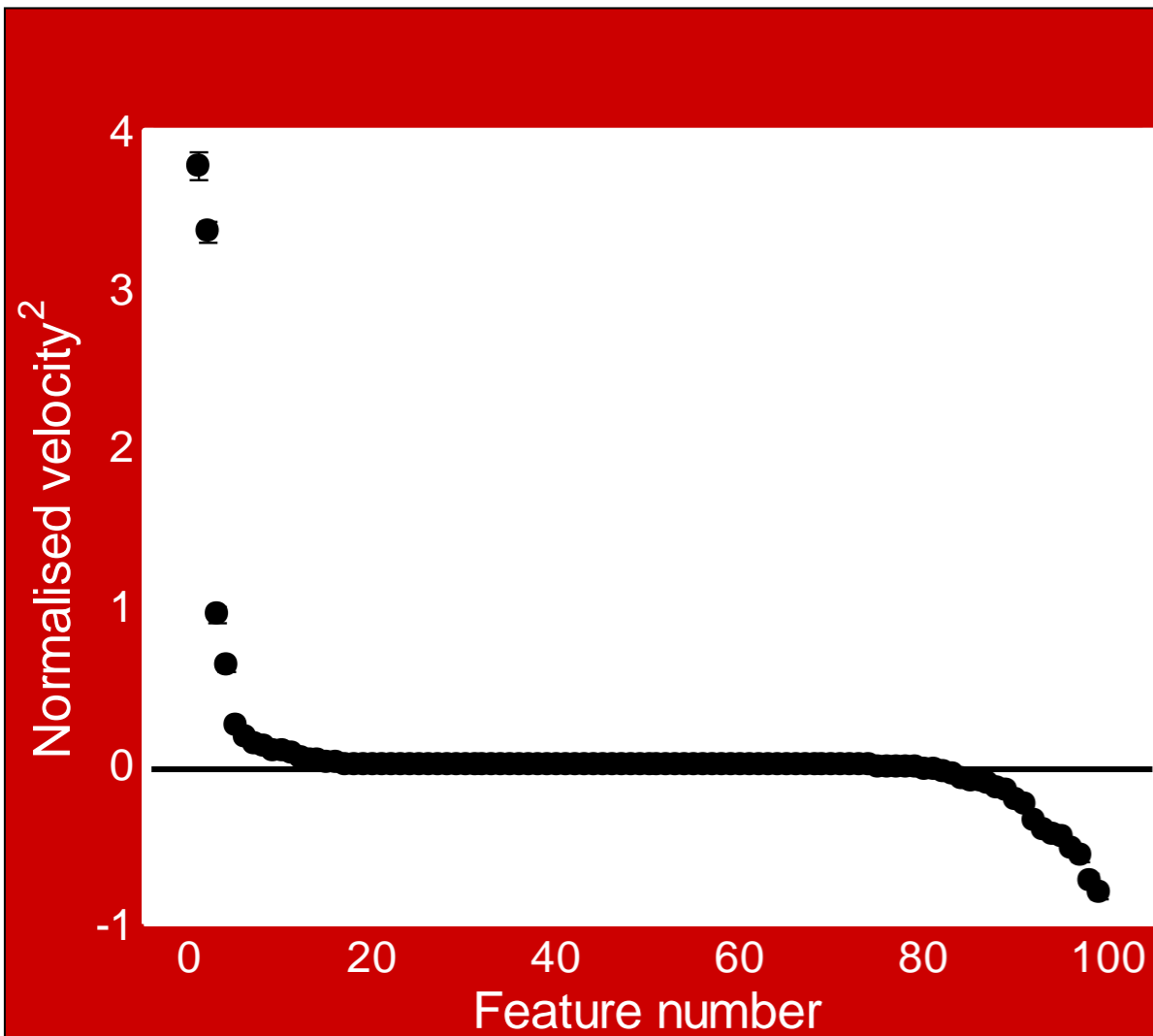


Leading modes

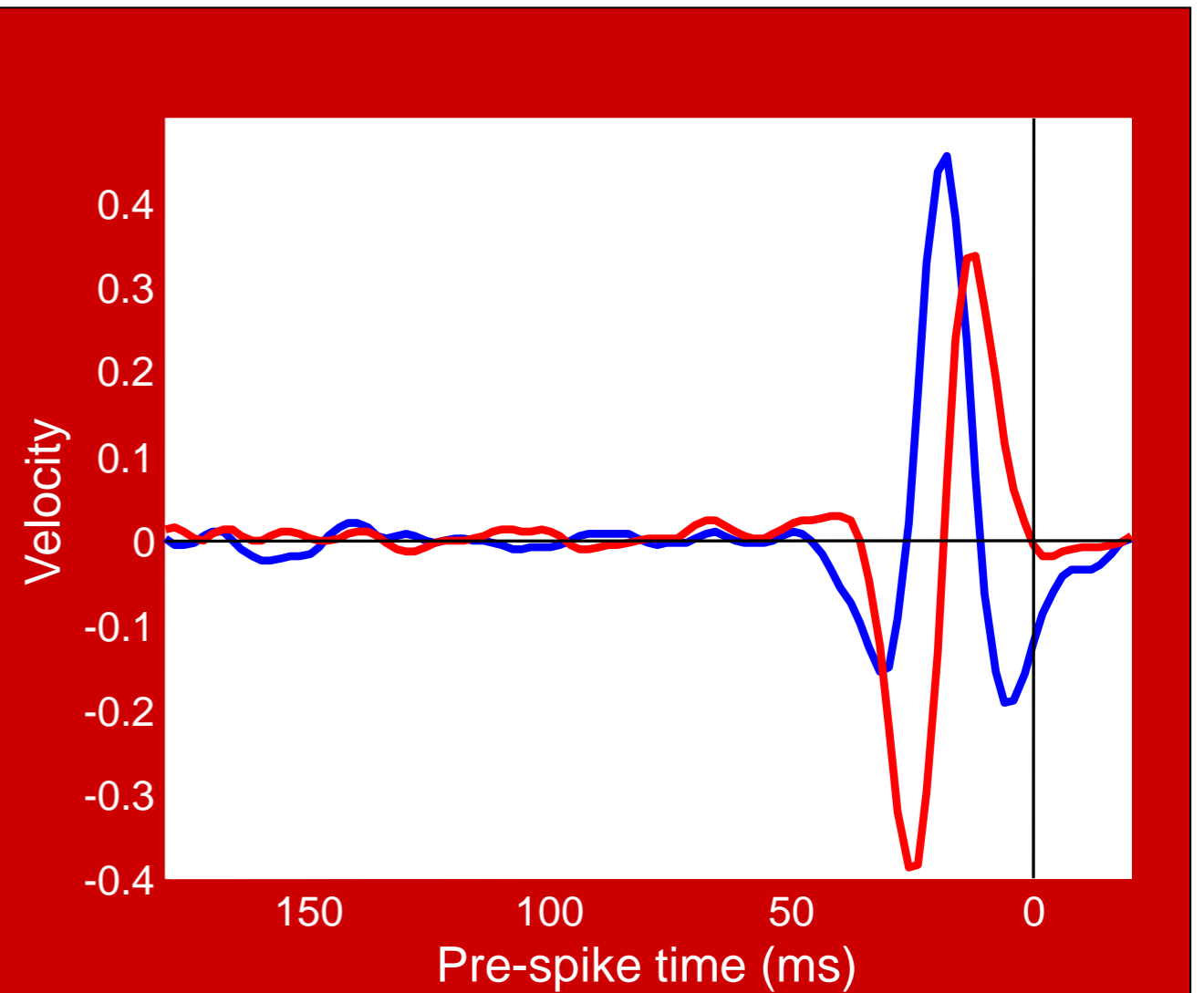


Eigenspectrum from barrel cortical neurons

Eigenspectrum

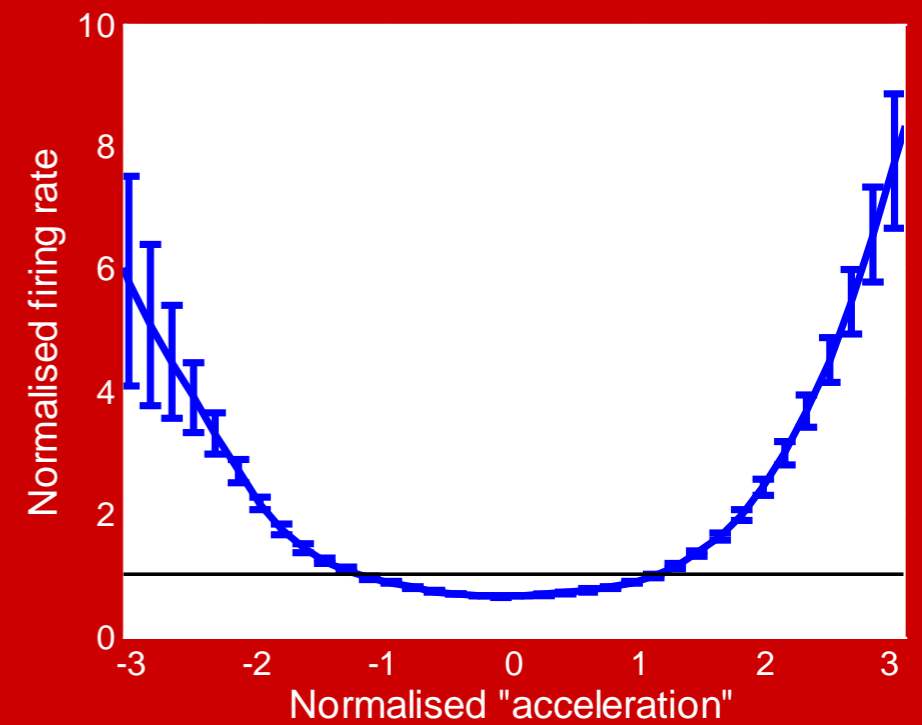
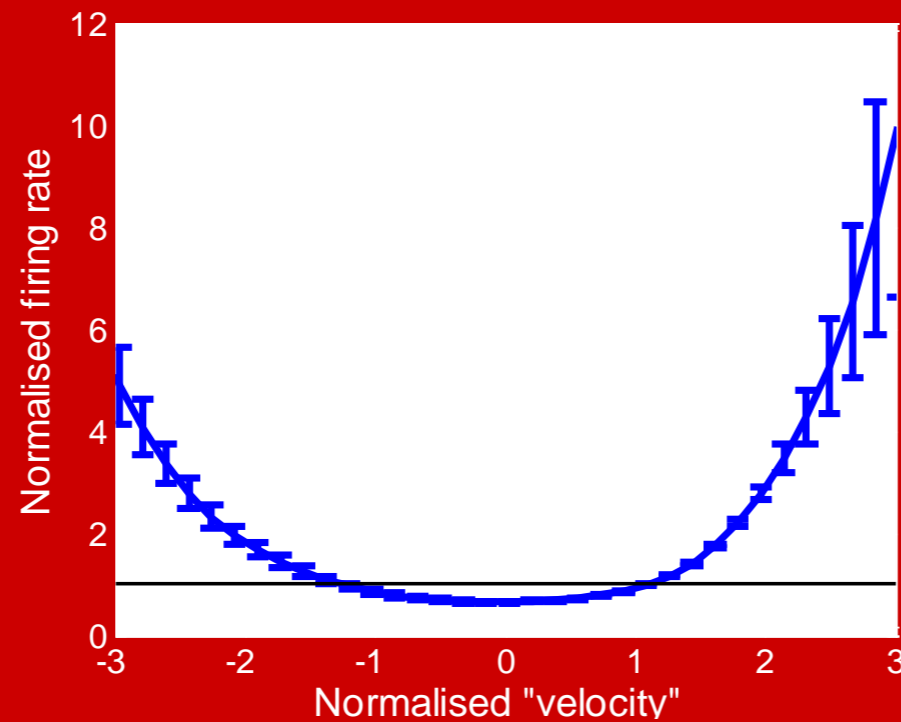


Leading modes

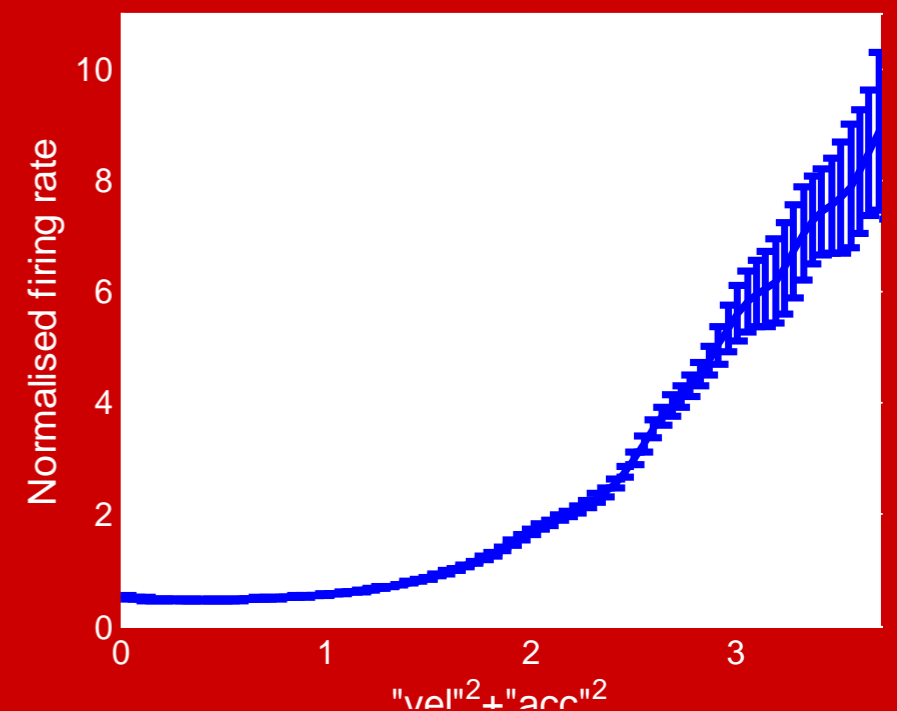
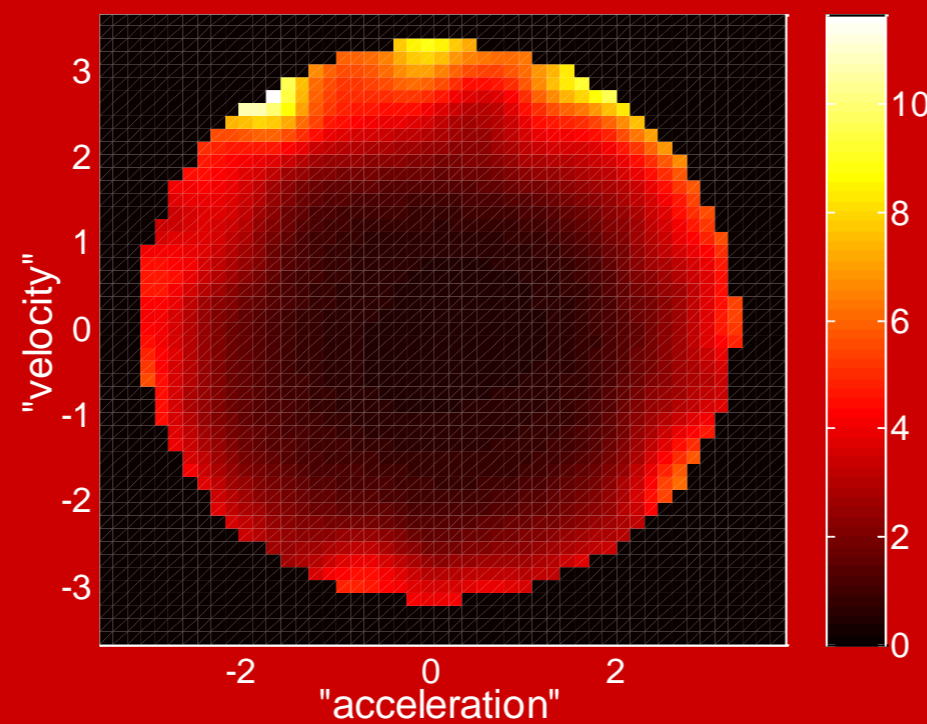


Input/output relations from barrel cortical neurons

Input/output relations wrt first two filters, alone:



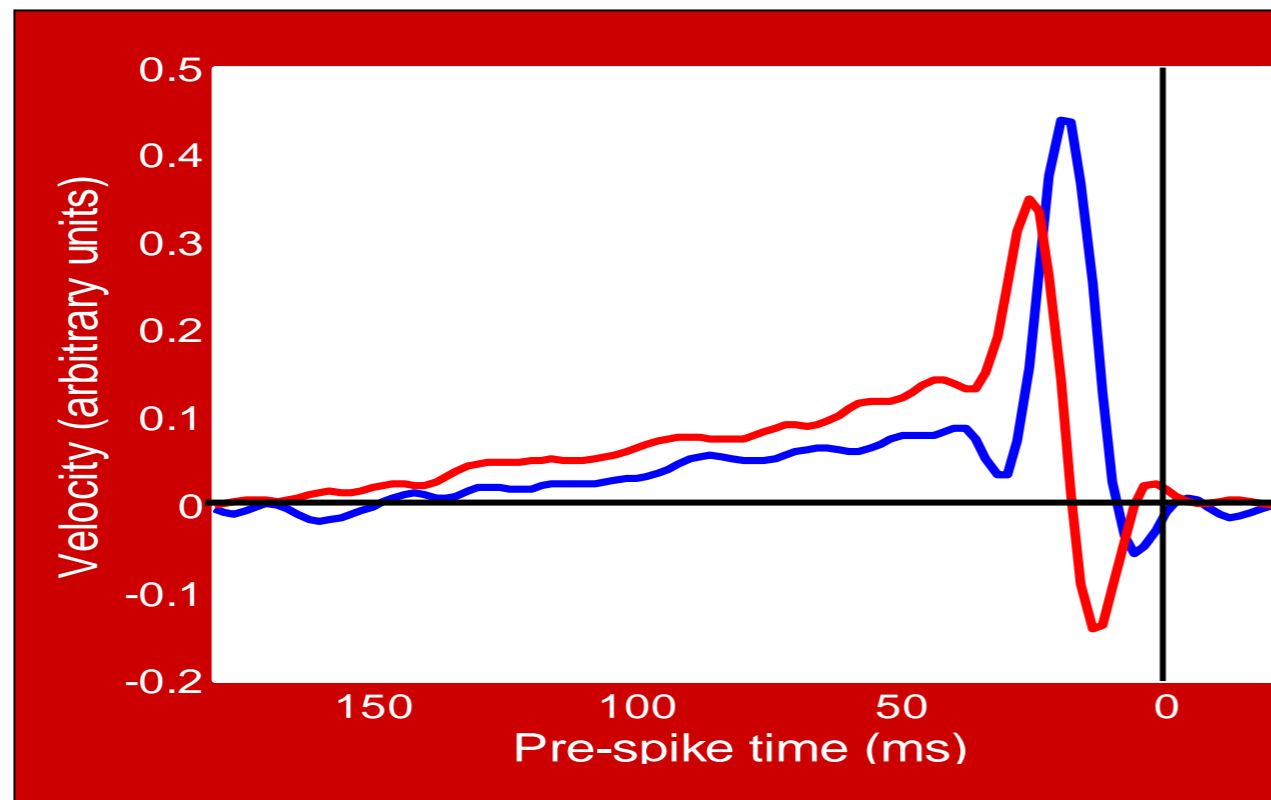
and in quadrature:



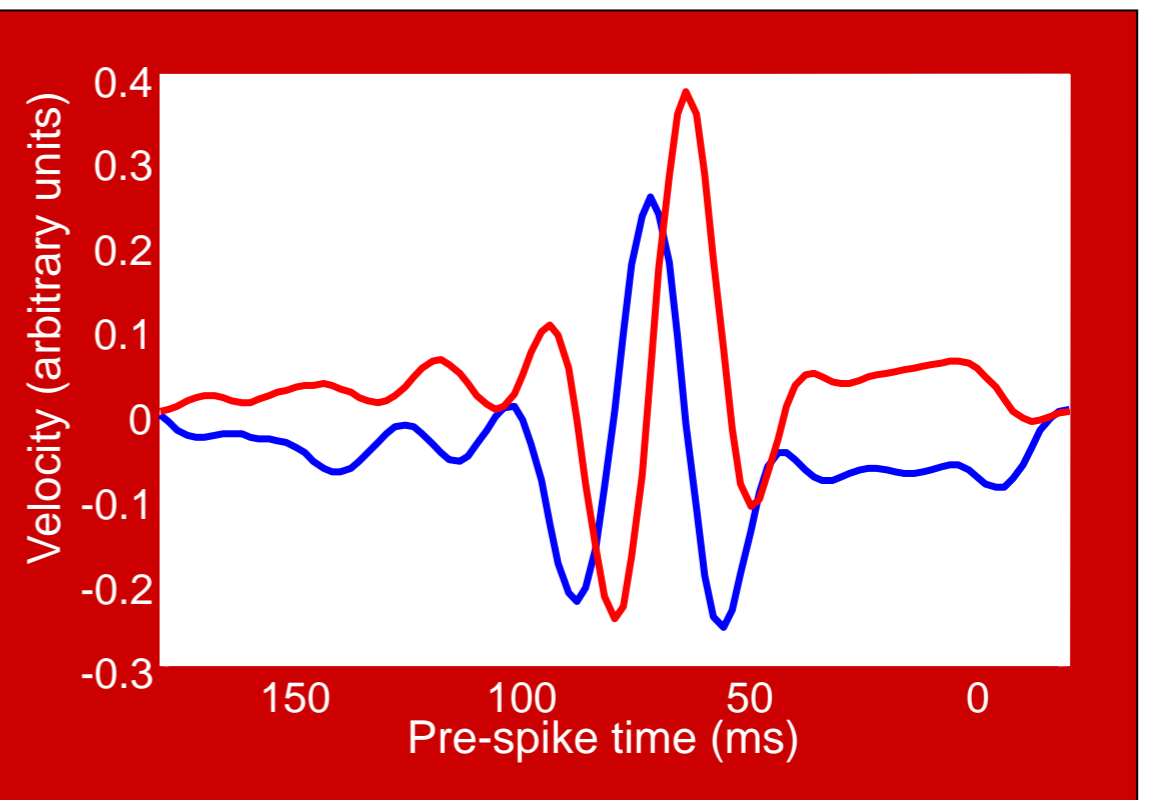
Less significant eigenmodes from barrel cortical neurons

How about the other modes?

Next pair with +ve eigenvalues

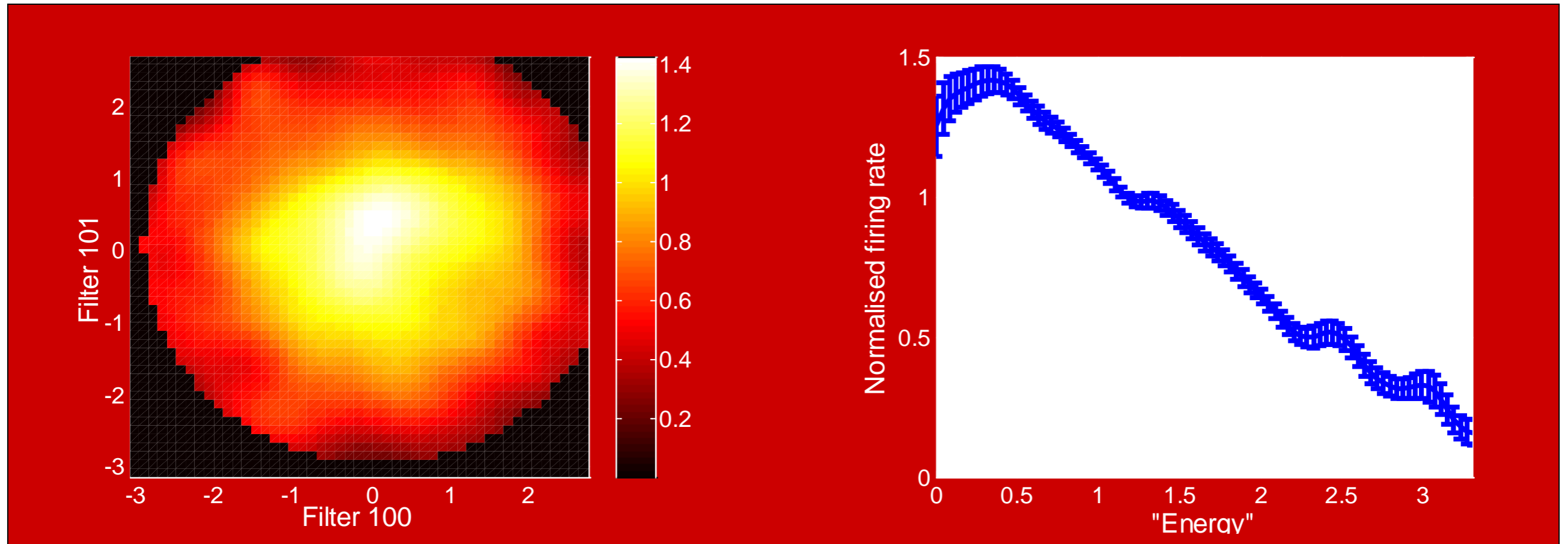


Pair with -ve eigenvalues



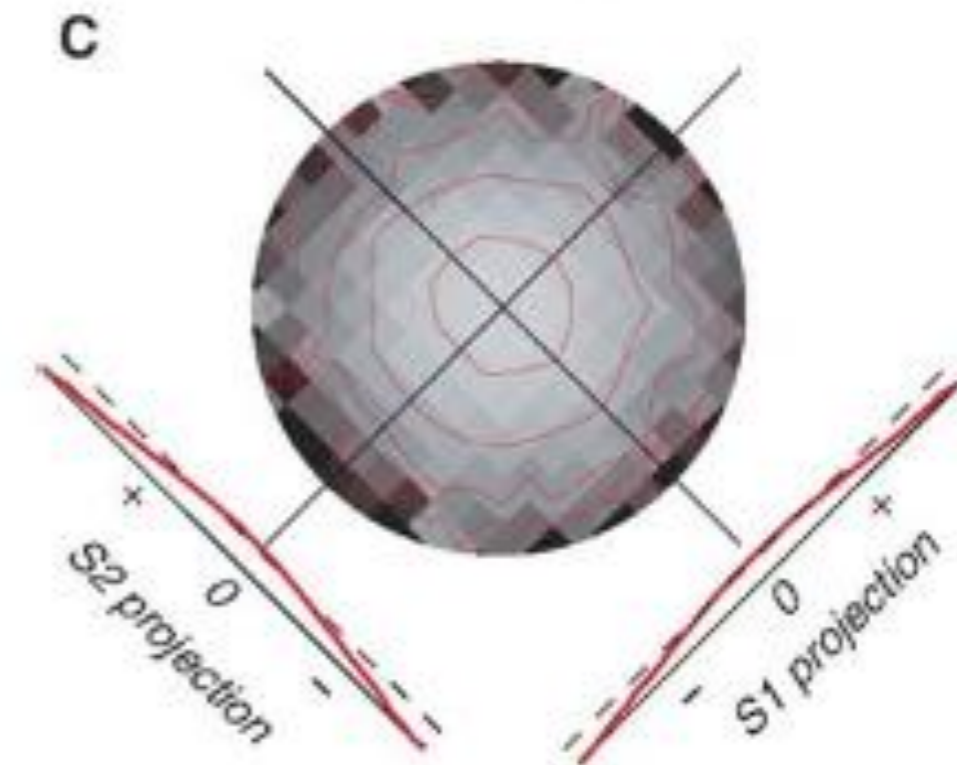
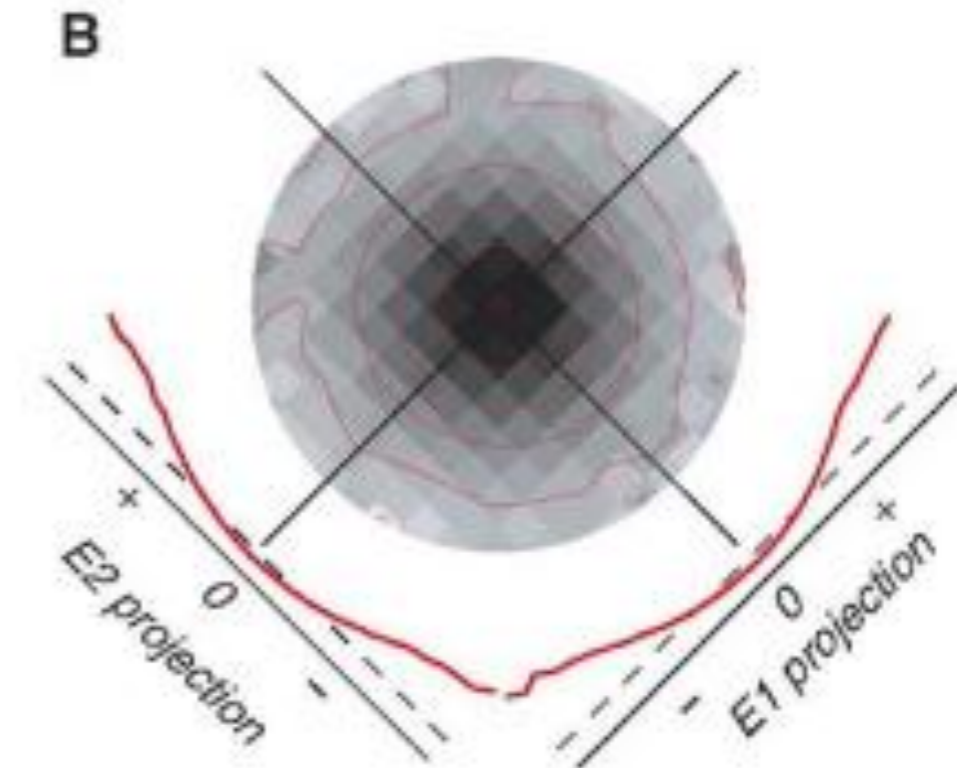
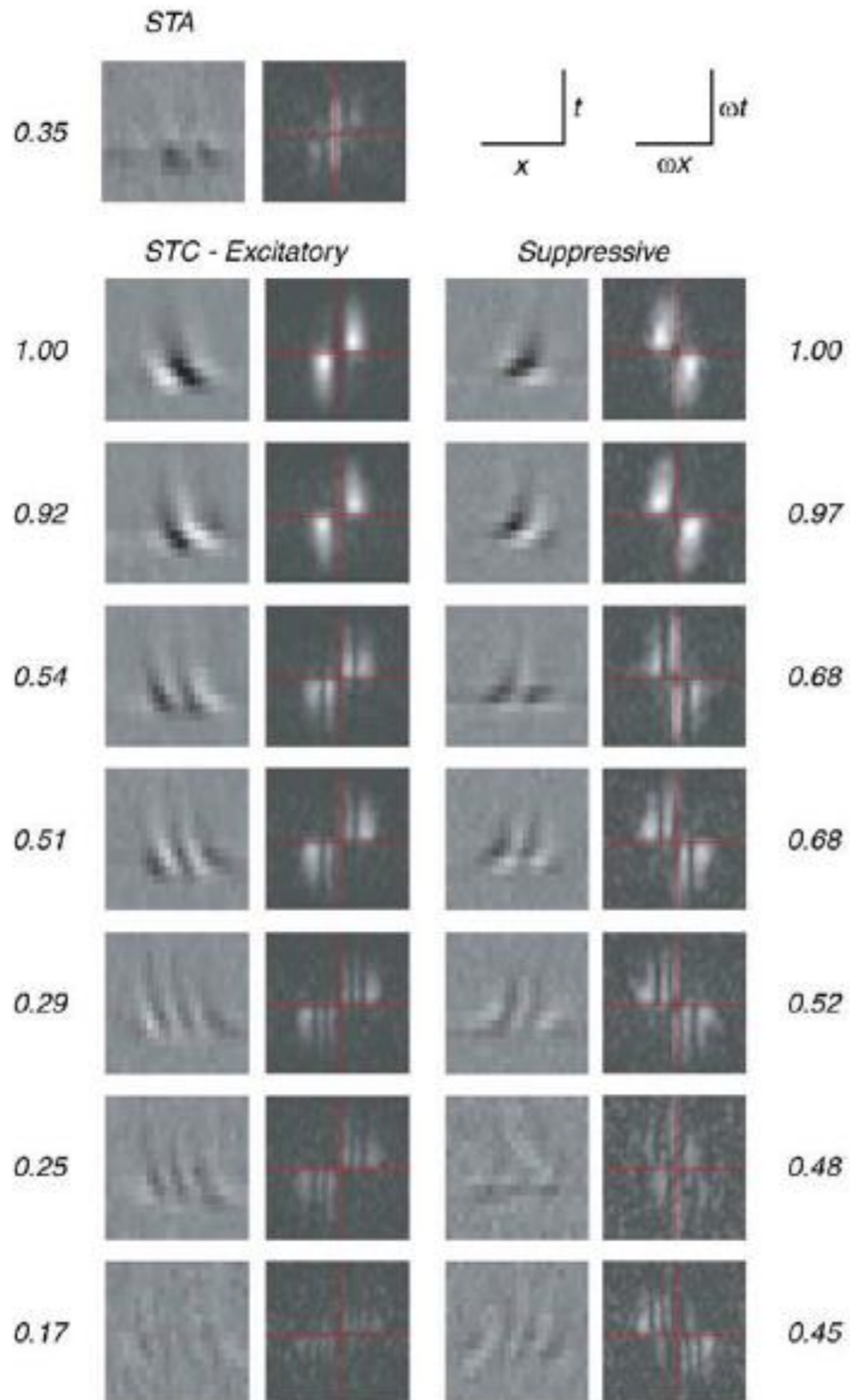
Negative eigenmode pair

Input/output relations for negative pair

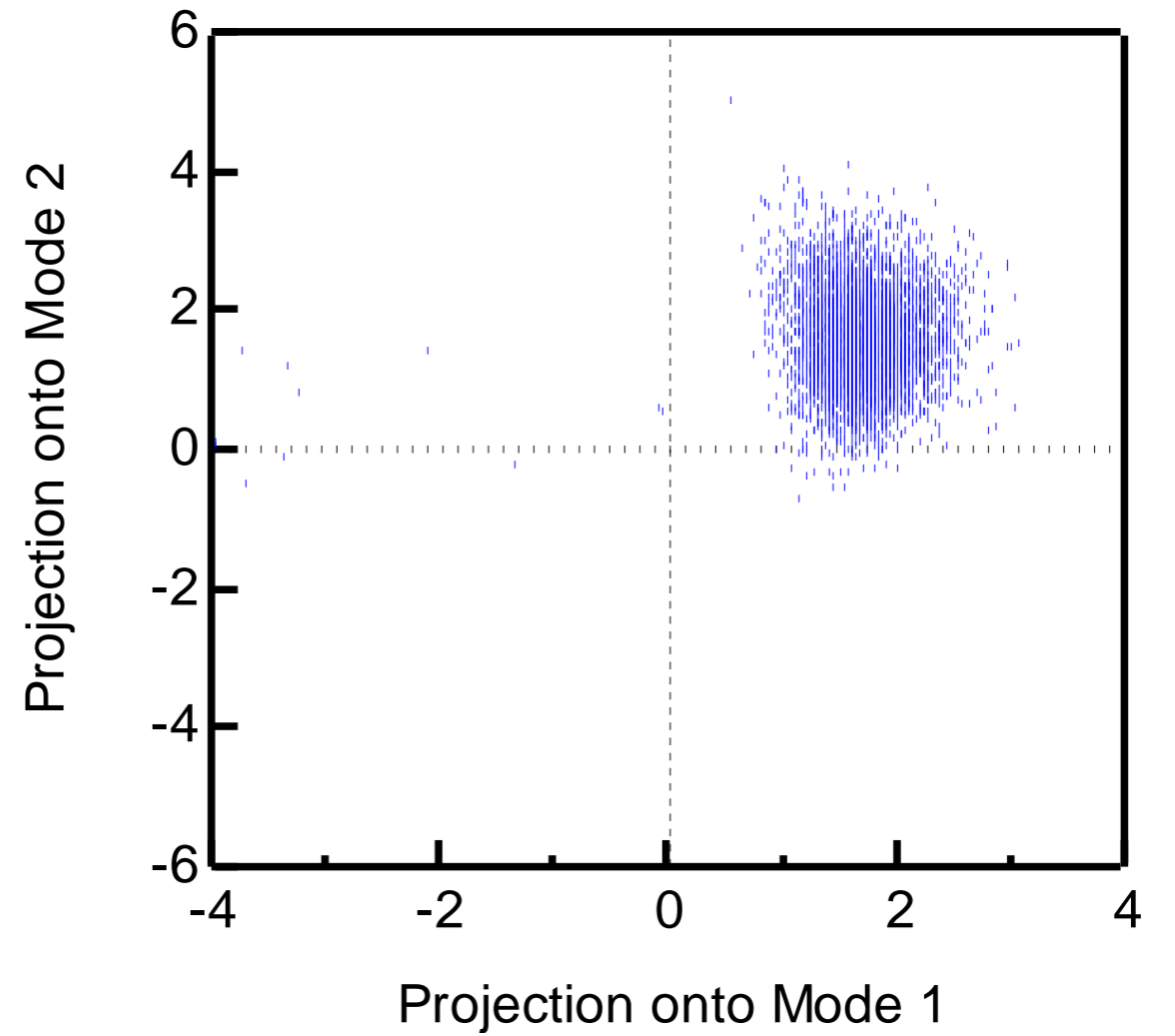
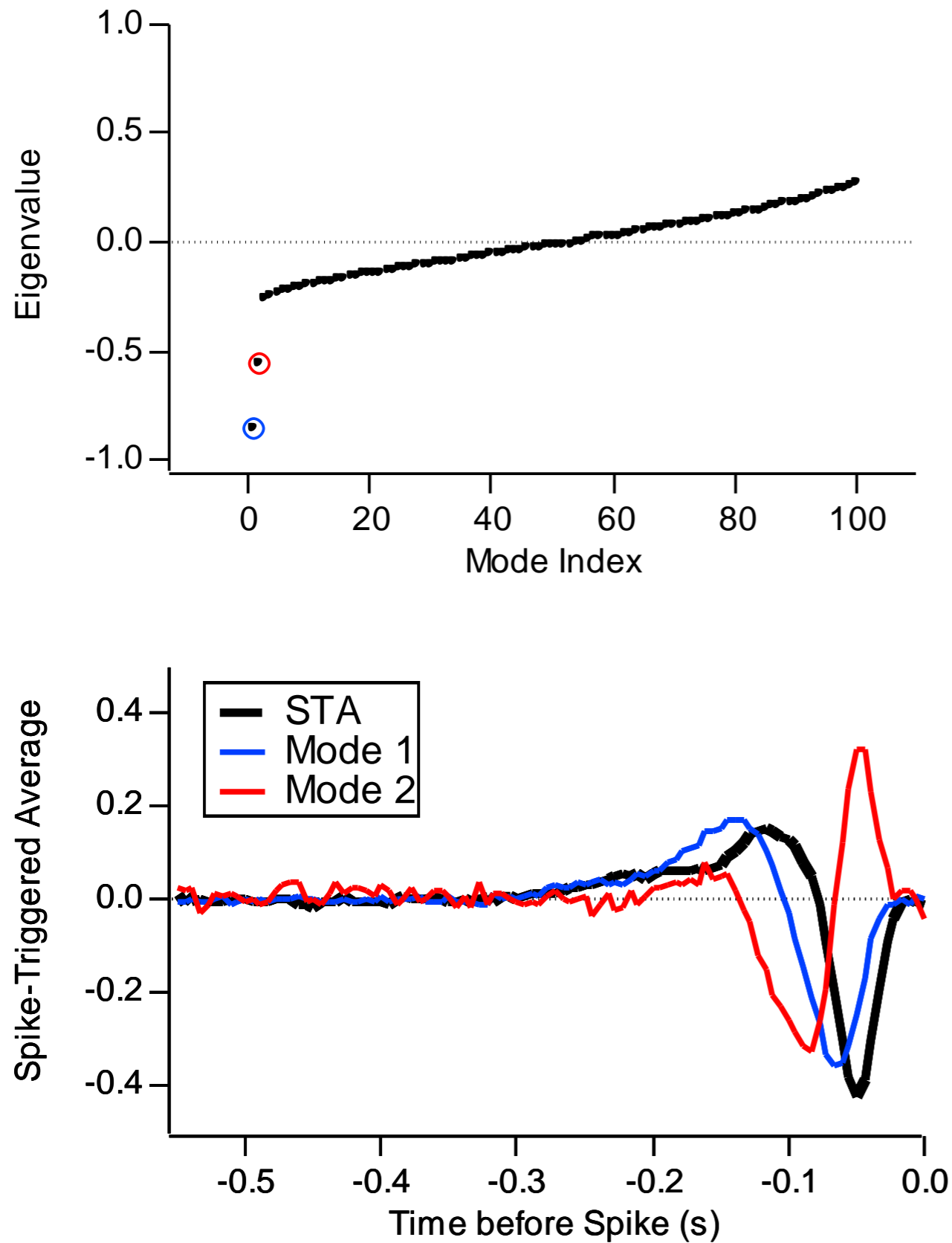


Firing rate *decreases* with increasing projection:
suppressive modes

Complex cells in V1

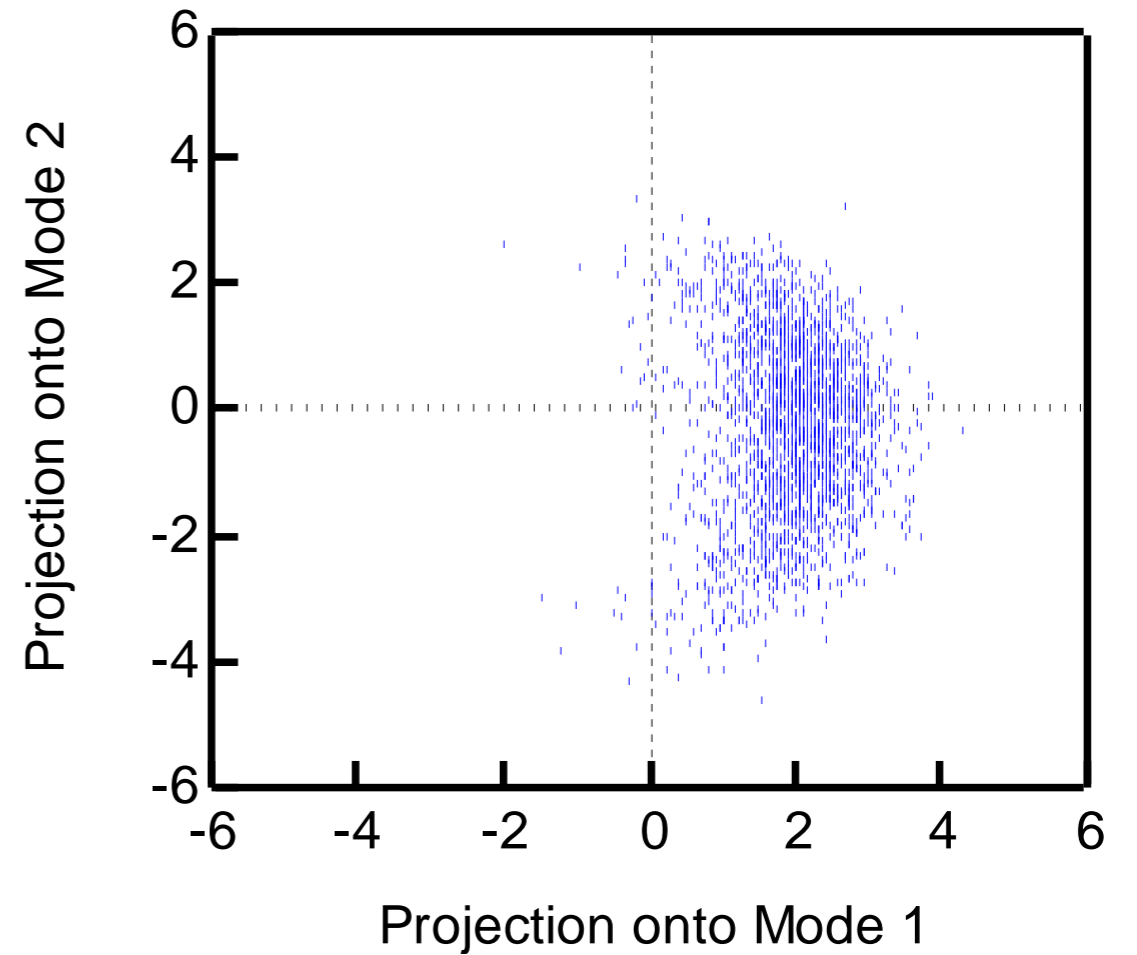
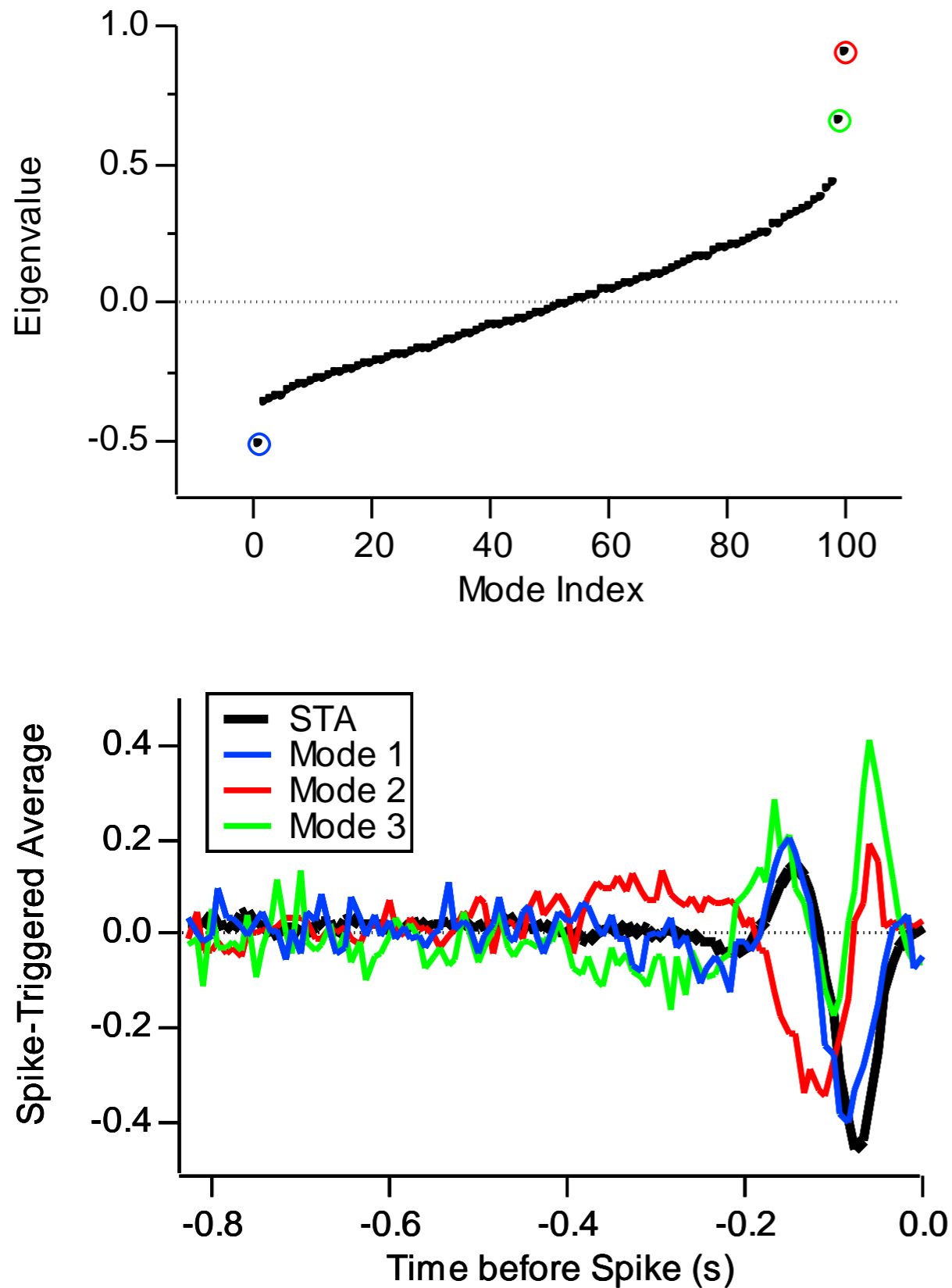


Salamander retinal ganglion cells perform a variety of computations

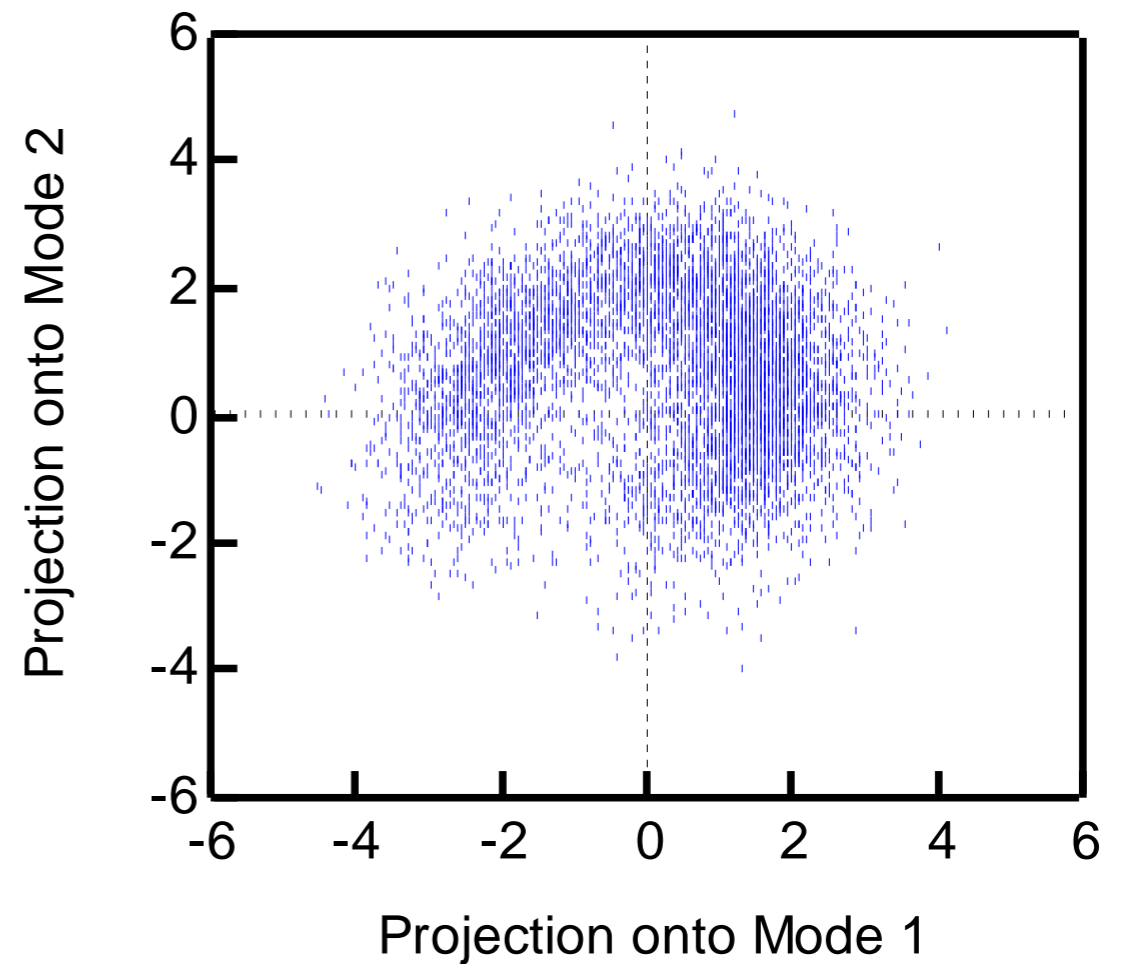
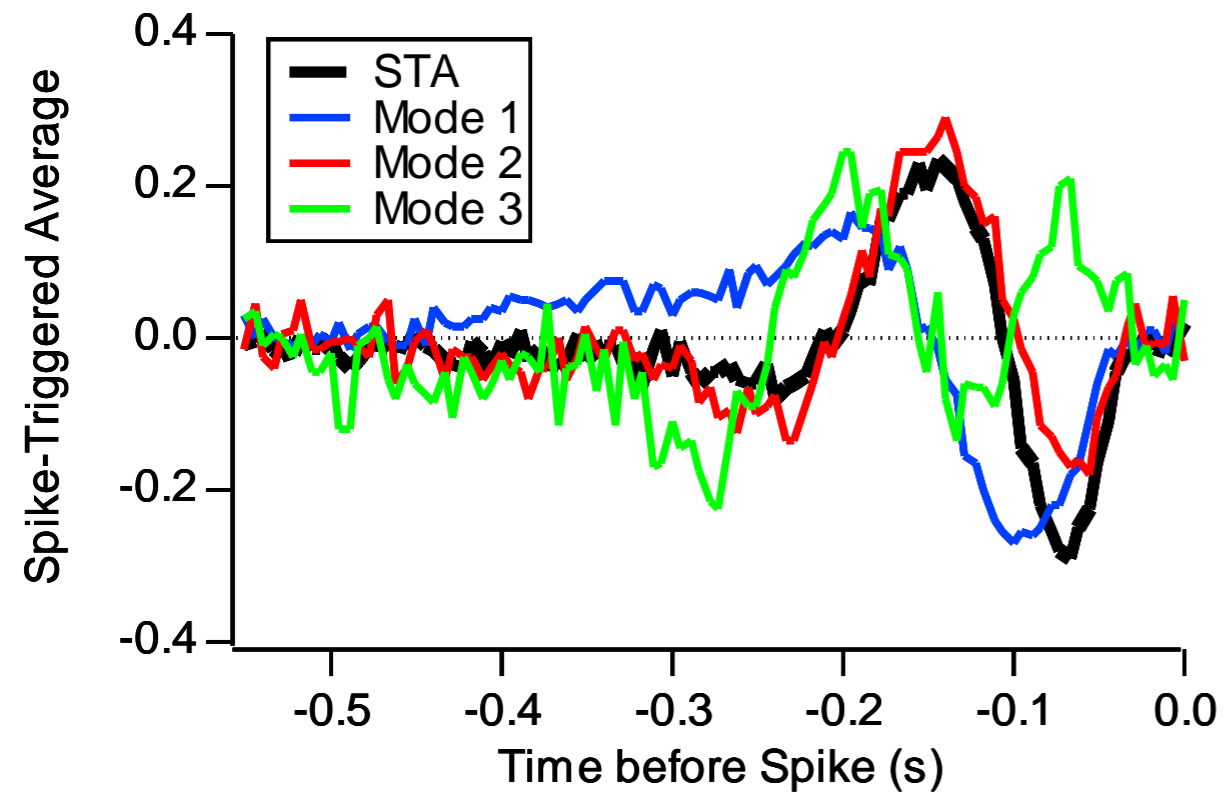
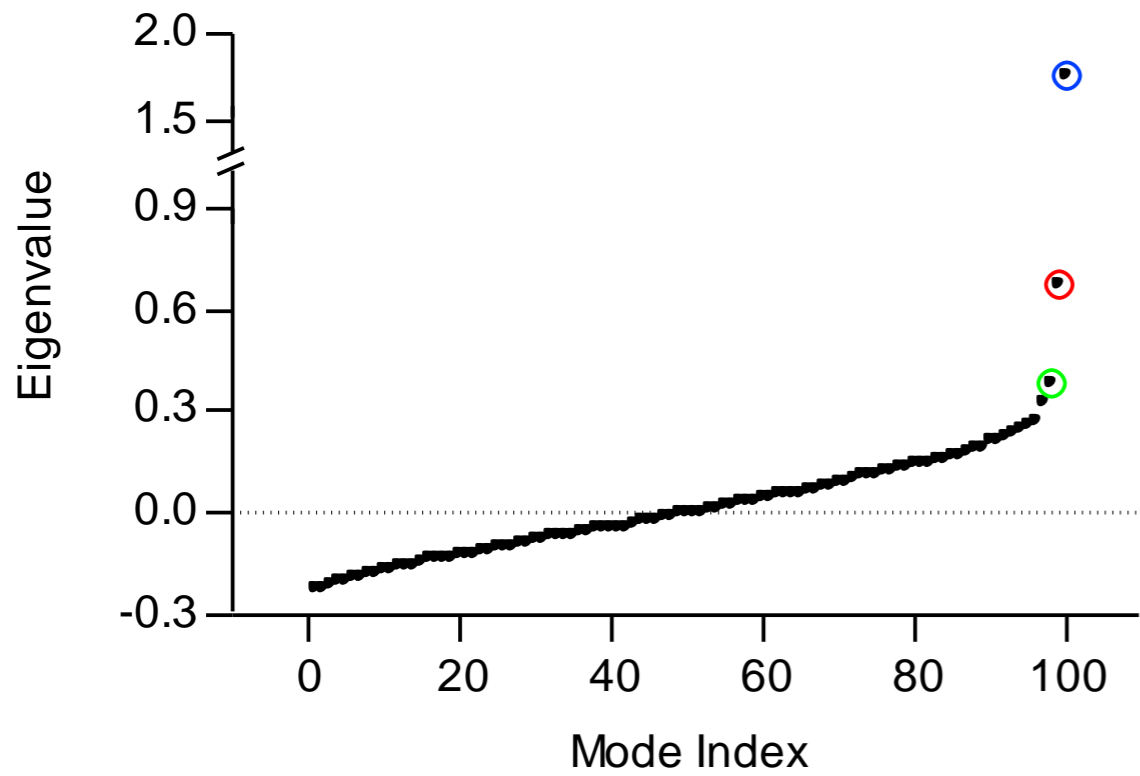


Michael Berry

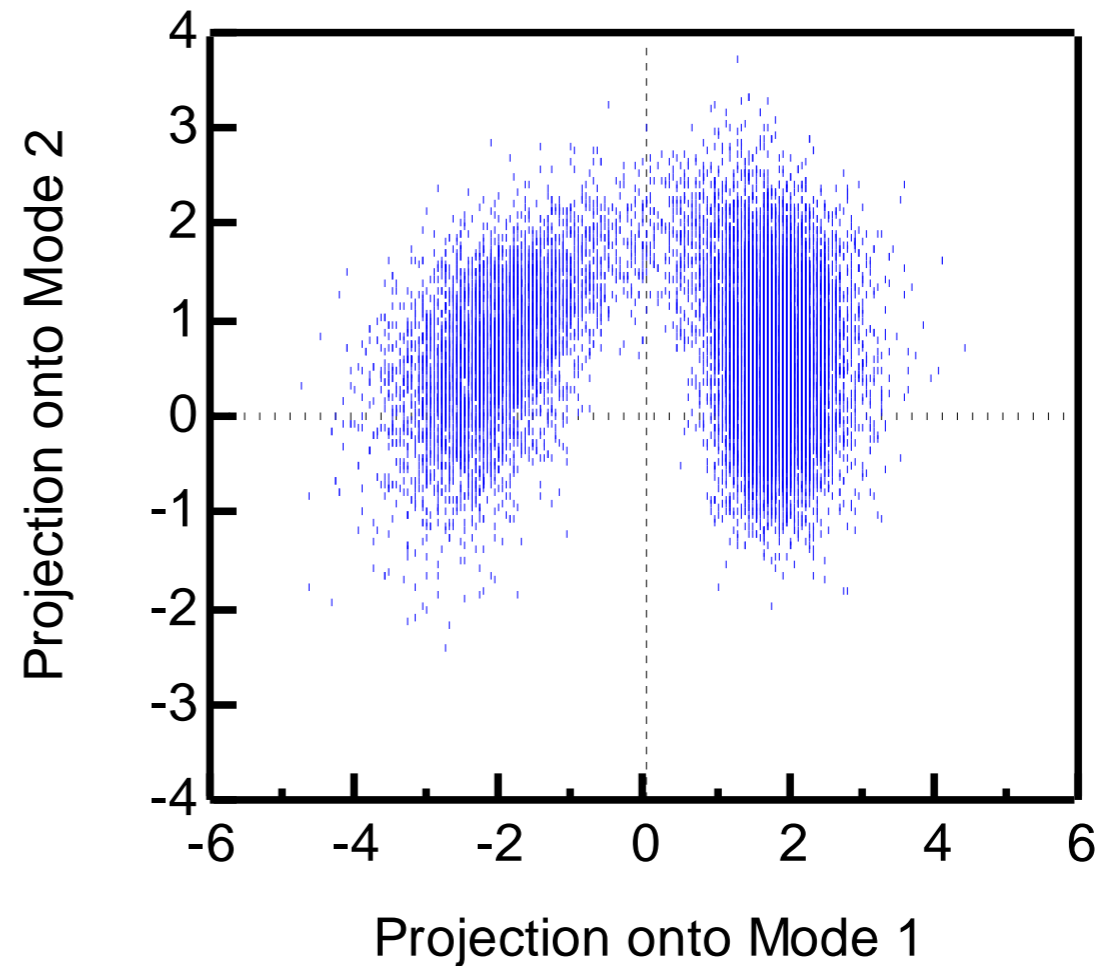
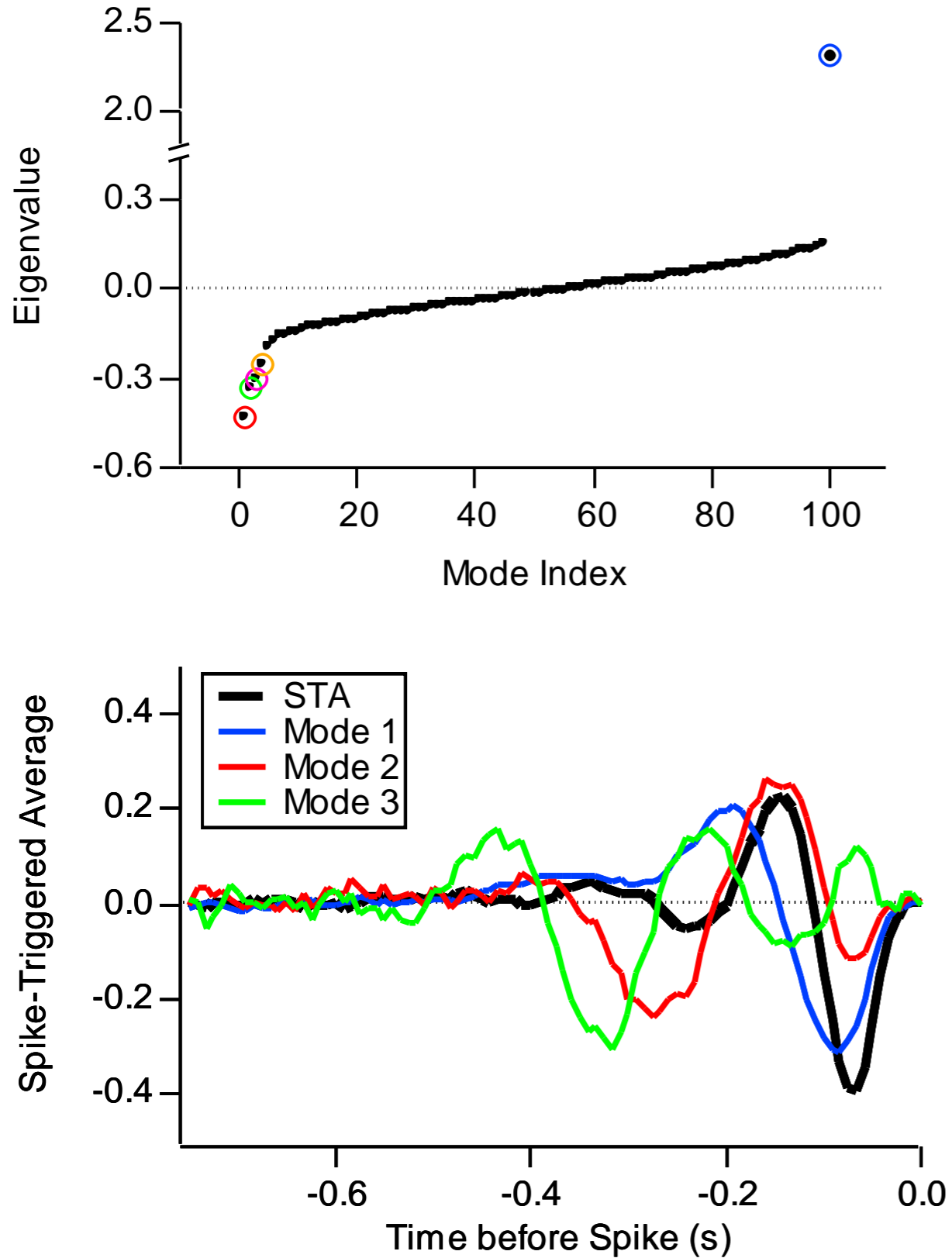
Not a threshold-crossing neuron



Complex cell like



Bimodal: two separate features are encoded



Basic types of computation

- **integrators**
H1, some single cortical neurons
- **differentiators**
Retina, simple cells, HH neuron, auditory neurons
- **frequency-power detectors**
V1 complex cells, somatosensory, auditory, retina

When have you done a good job?

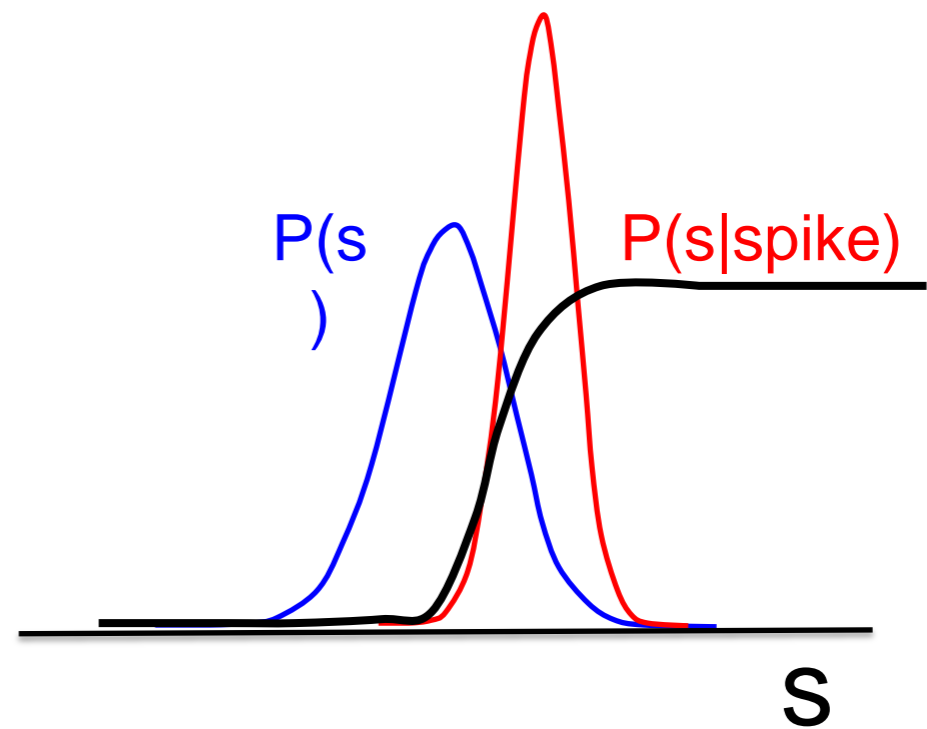
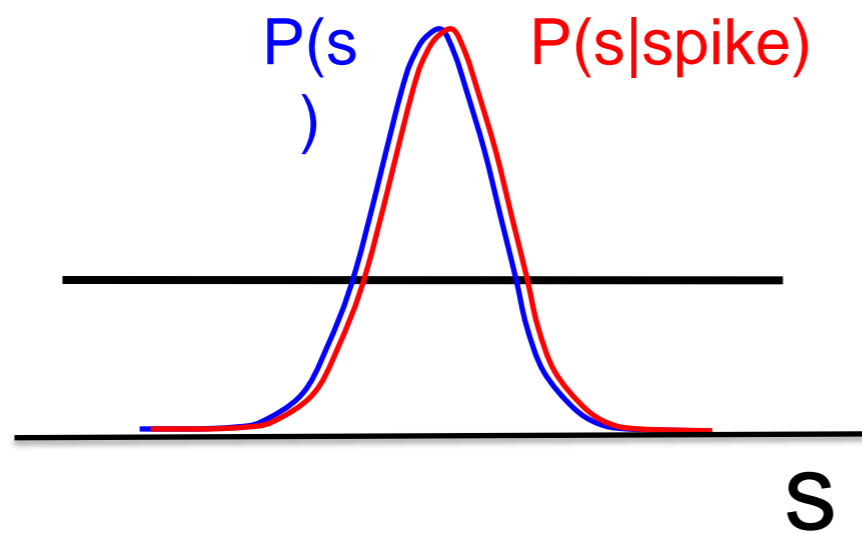
- When the tuning curve over your variable is *interesting*.
- How to quantify interesting?

When have you done a good job?

Tuning curve: $P(\text{spike}|s) = P(s|\text{spike}) P(\text{spike}) / P(s)$

Boring: spikes unrelated to stimulus

Interesting: spikes are selective

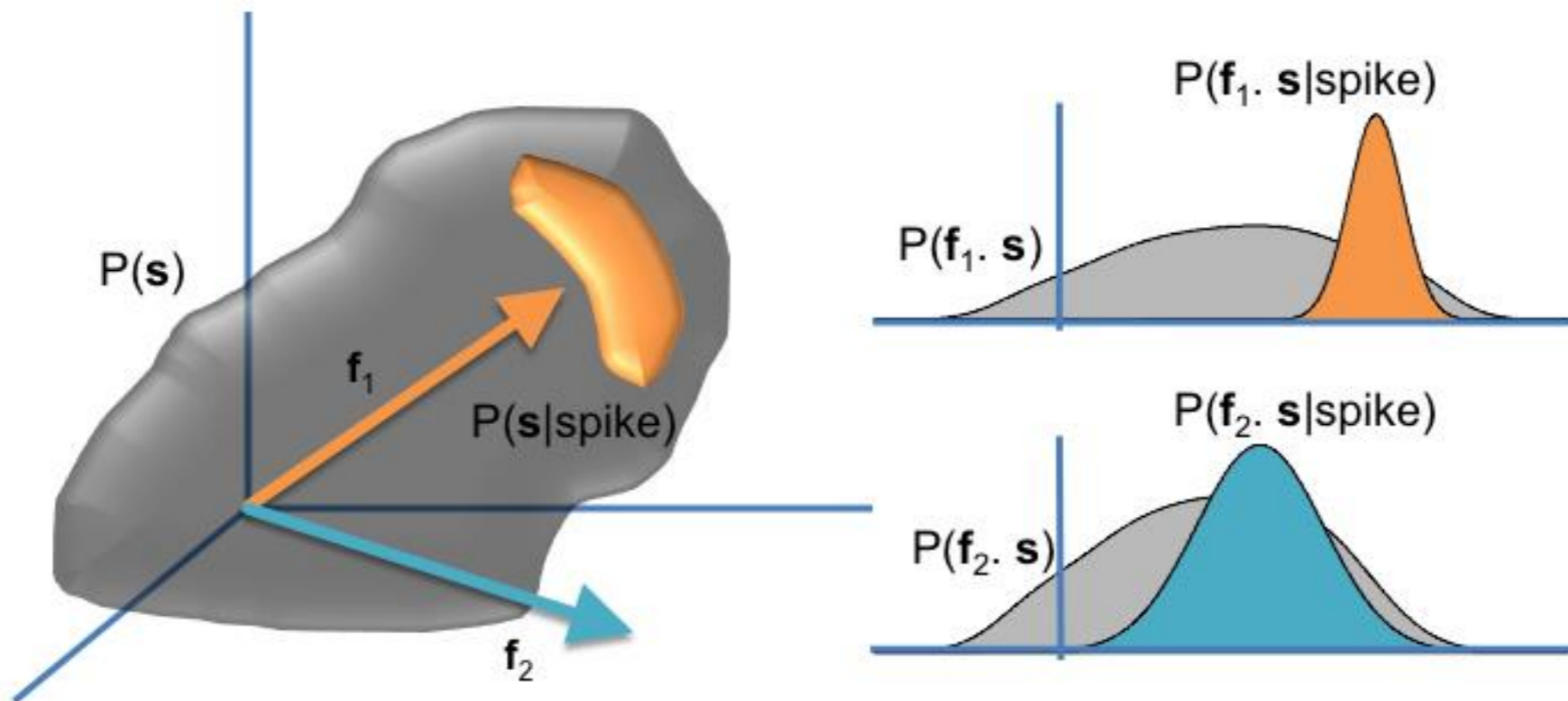


Goodness measure: $D_{\text{KL}}(P(s|\text{spike}) | P(s))$

Maximally informative dimensions

Sharpee, Rust and Bialek, Neural Computation, 2004

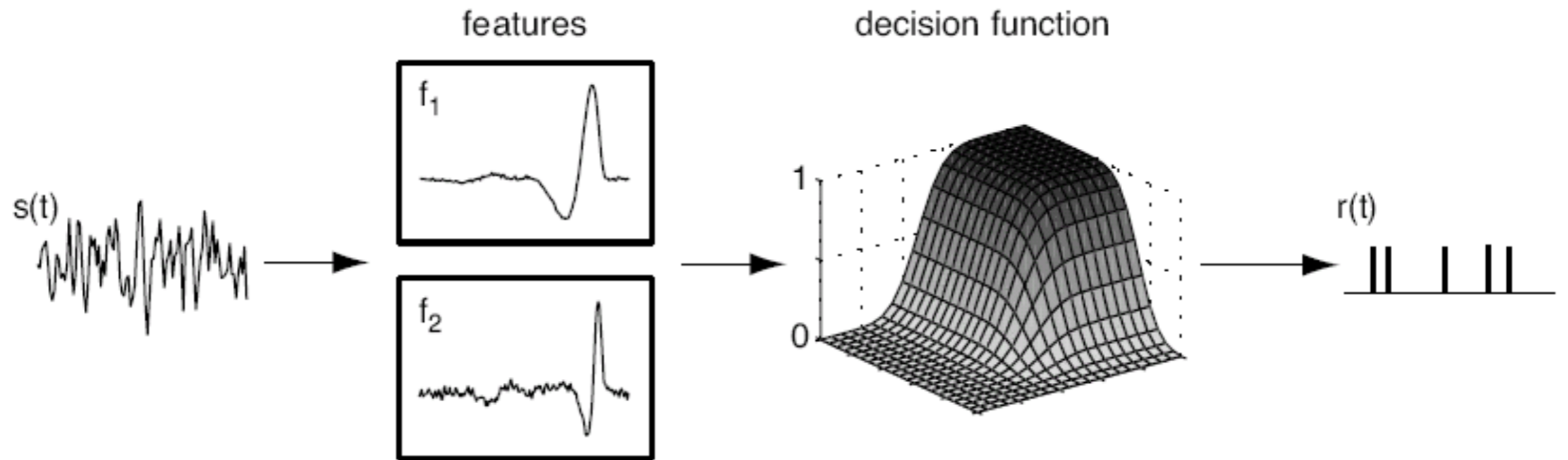
Choose filter in order to maximize D_{KL} between spike-conditional and prior distributions



Finding relevant features

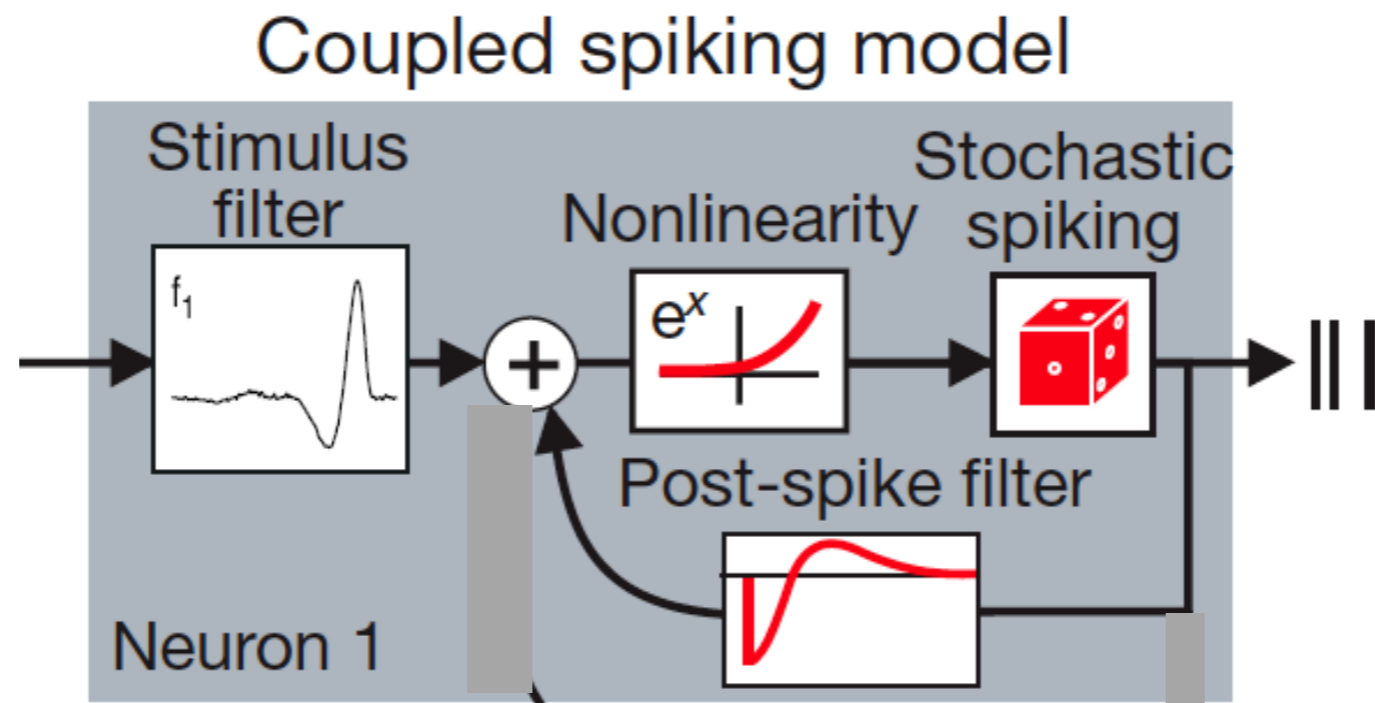
1. Single, best filter determined by the first moment
2. A family of filters derived using the second moment
3. Use the entire distribution: information theoretic methods

Less basic coding models



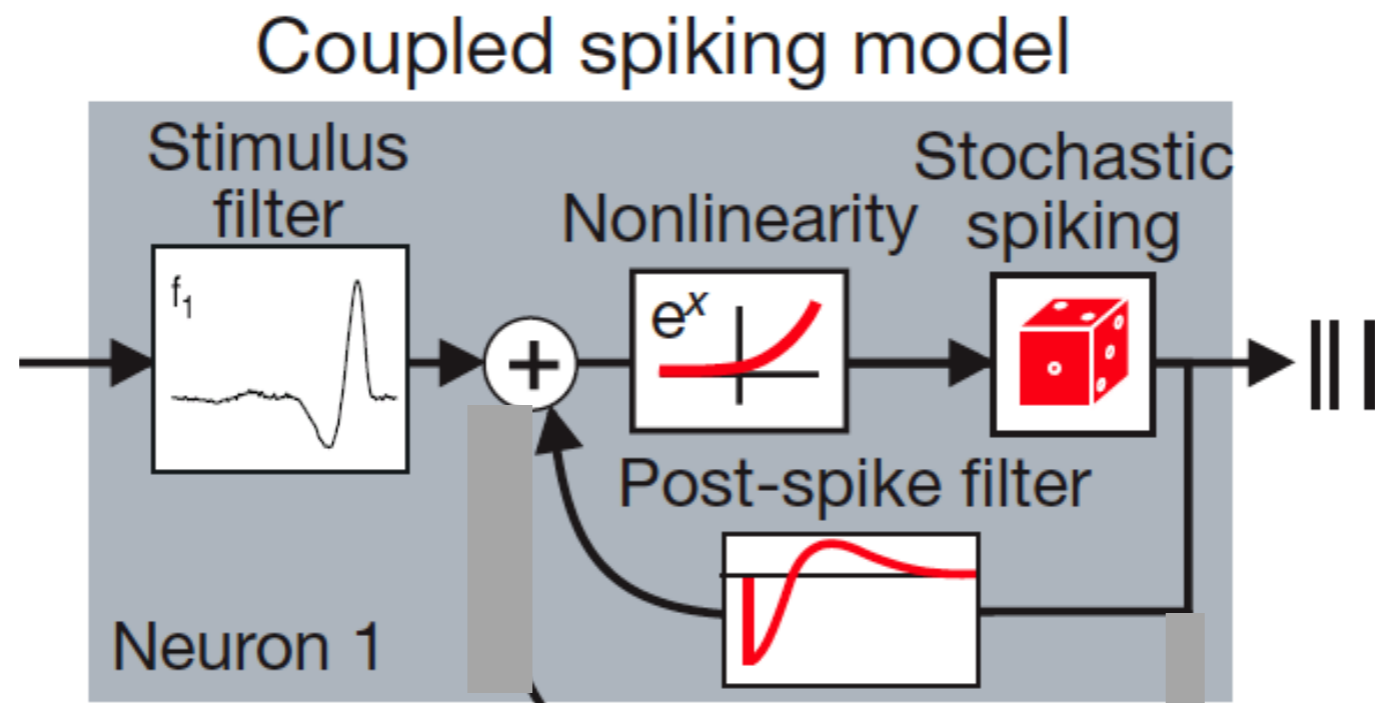
Linear filters & nonlinearity: $r(t) = g(f_1 * s, f_2 * s, \dots, f_n * s)$

Less basic coding models



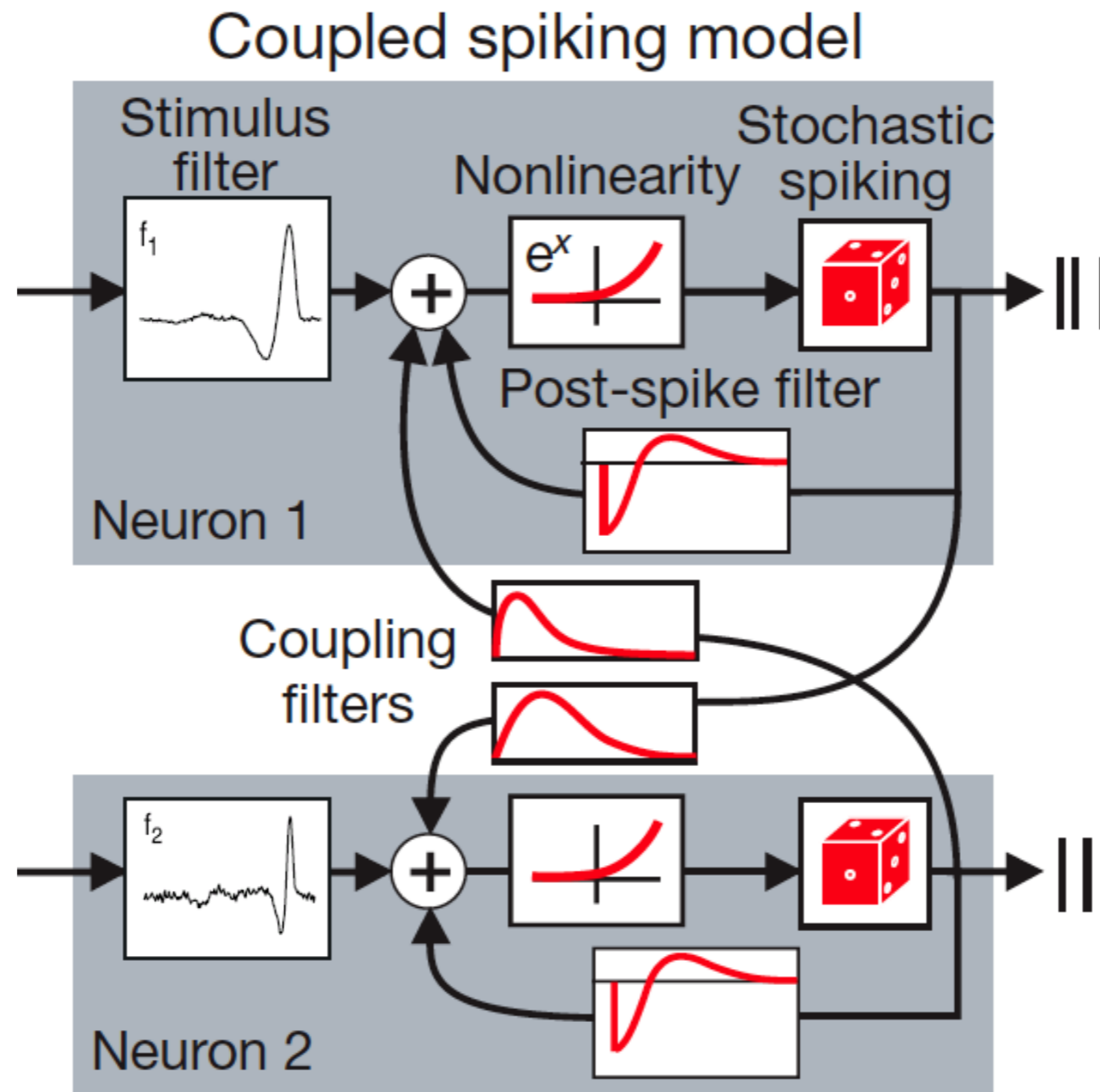
$$\text{GLM: } r(t) = g(f_1 * s + f_2 * r)$$

Less basic coding models



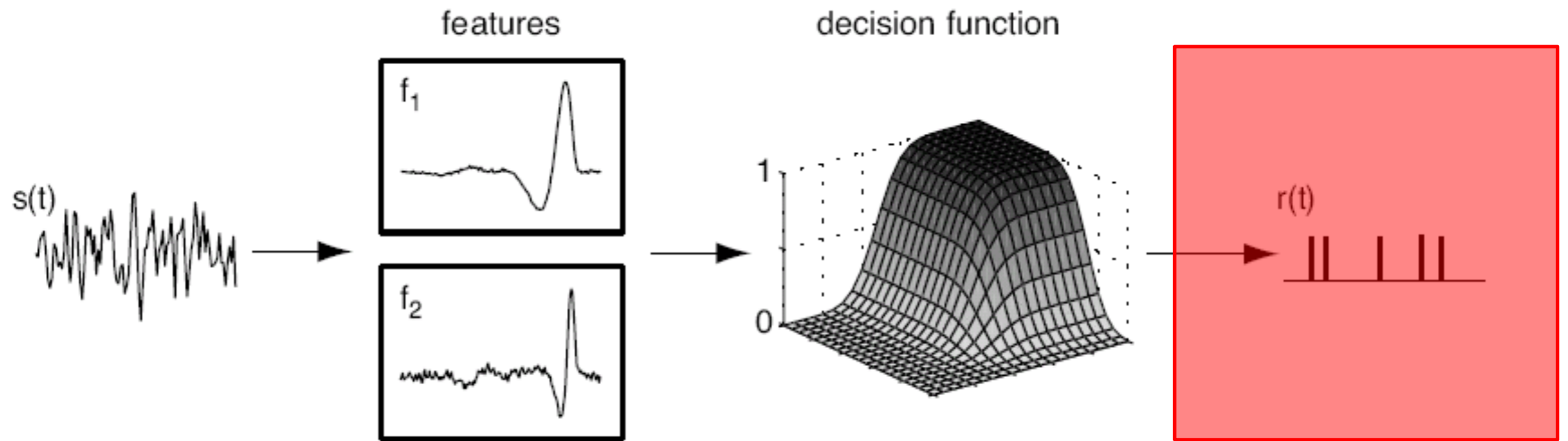
$$\text{GLM: } r(t) = g(f^*s + h^*r)$$

Less basic coding models



$$\text{GLM: } r(t) = g(f_1 * s + h_1 * r_1 + h_2 * r_2 + \dots)$$

Point process model



Binomial spiking

Properties:

Distribution: $P_n[k] = ?$

Mean: $\langle x \rangle = ?$

Variance: $\text{Var}(x) = ?$

Binomial spiking

Properties:

Distribution: $P_T[k] = (rT)^k \exp(-rT) / k!$

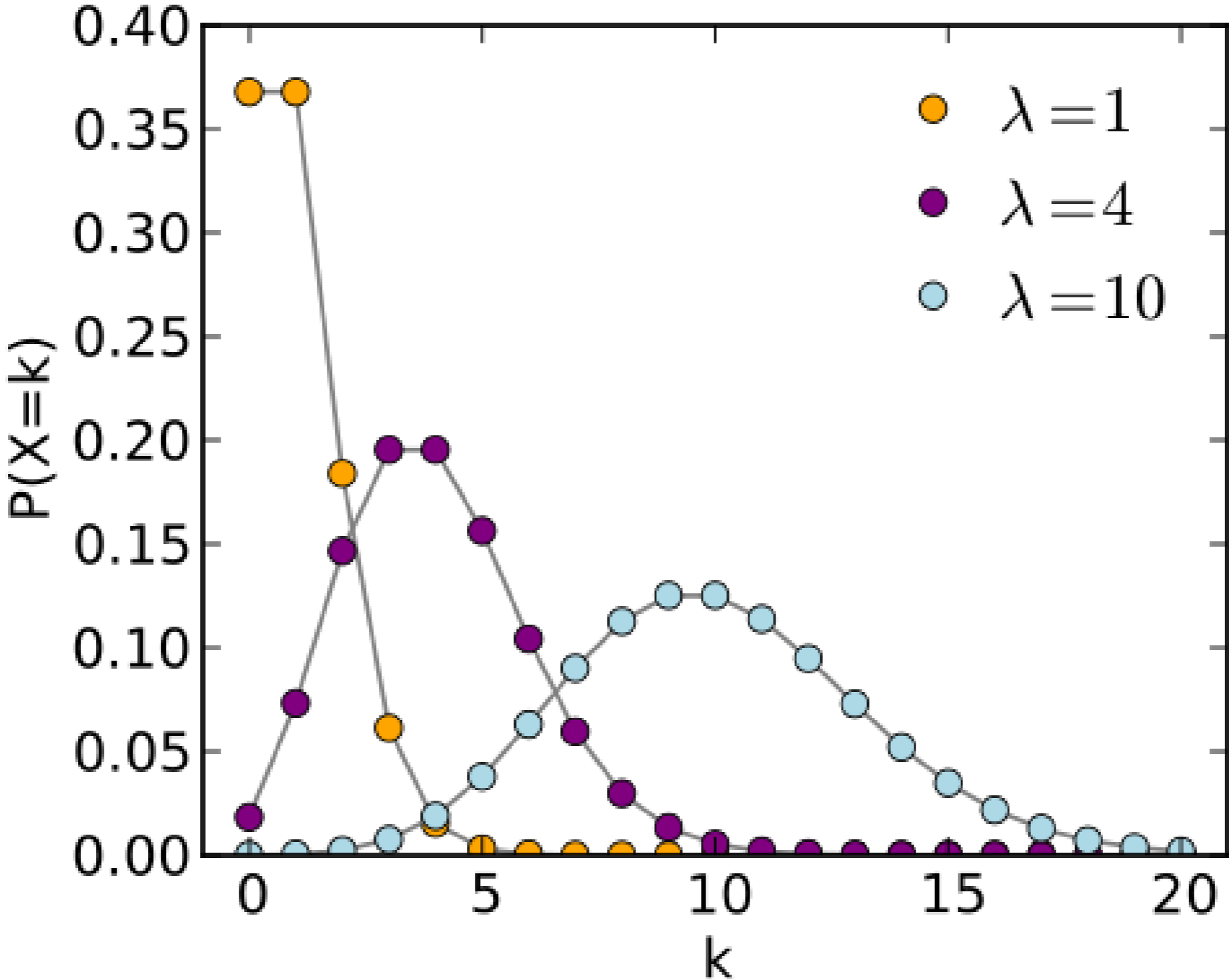
Mean: $\langle x \rangle = rT$

Variance: $\text{Var}(x) = rT$

Fano factor: $F = 1$

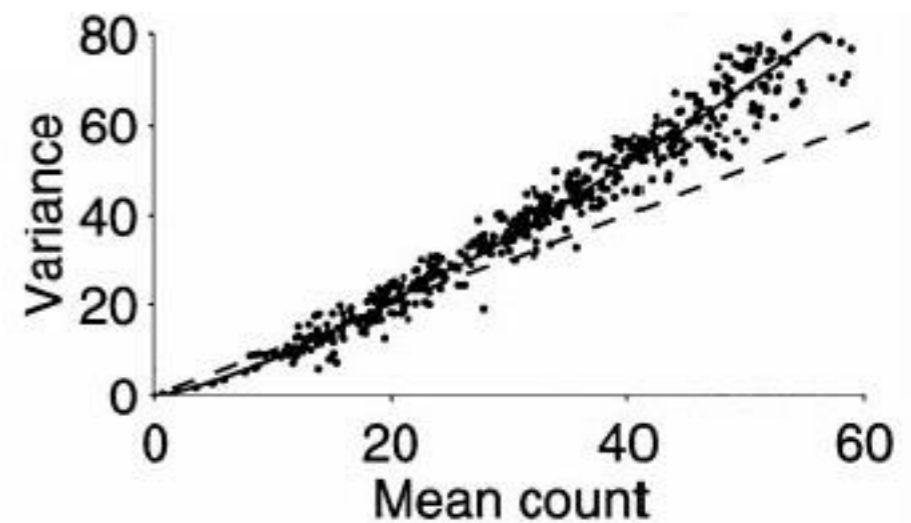
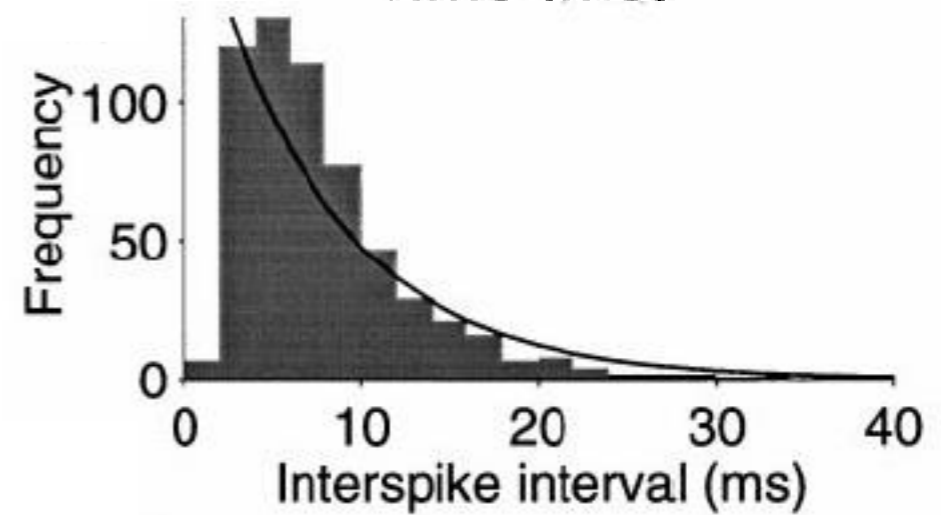
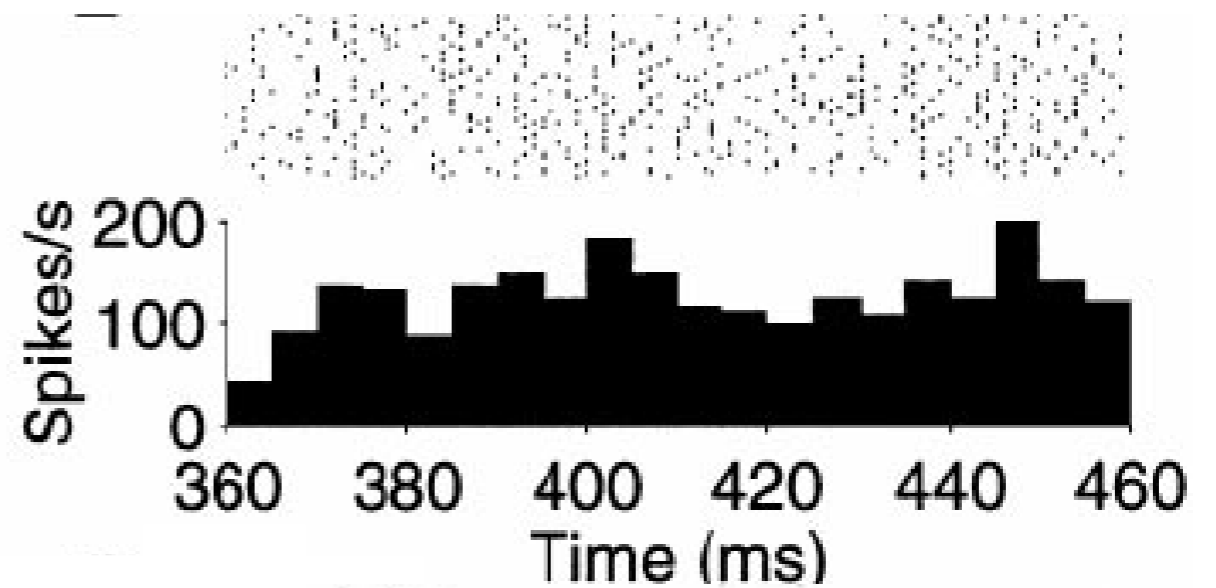
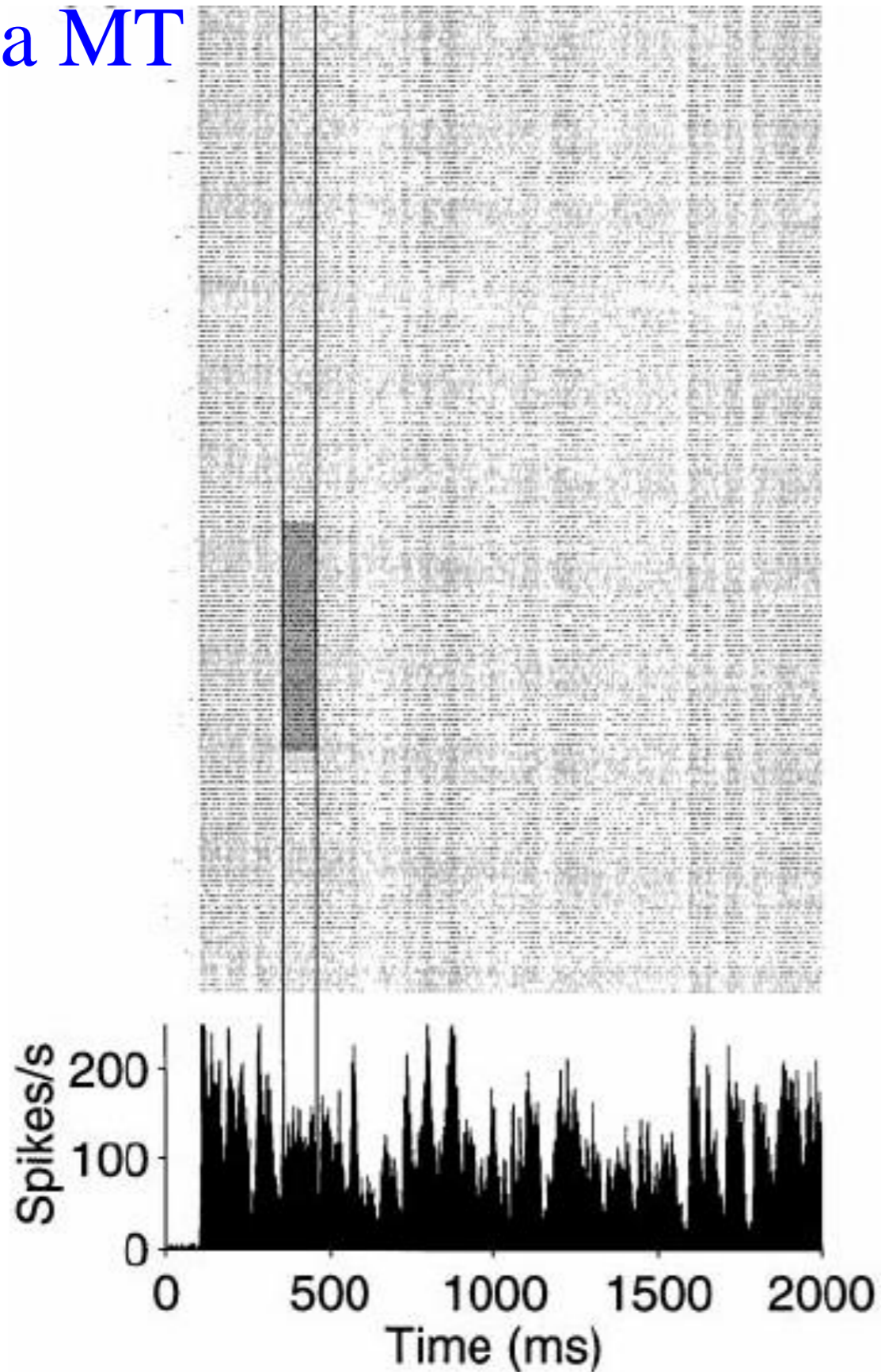
Interval distribution: $P(T) = r \exp(-rT)$

Poisson distribution



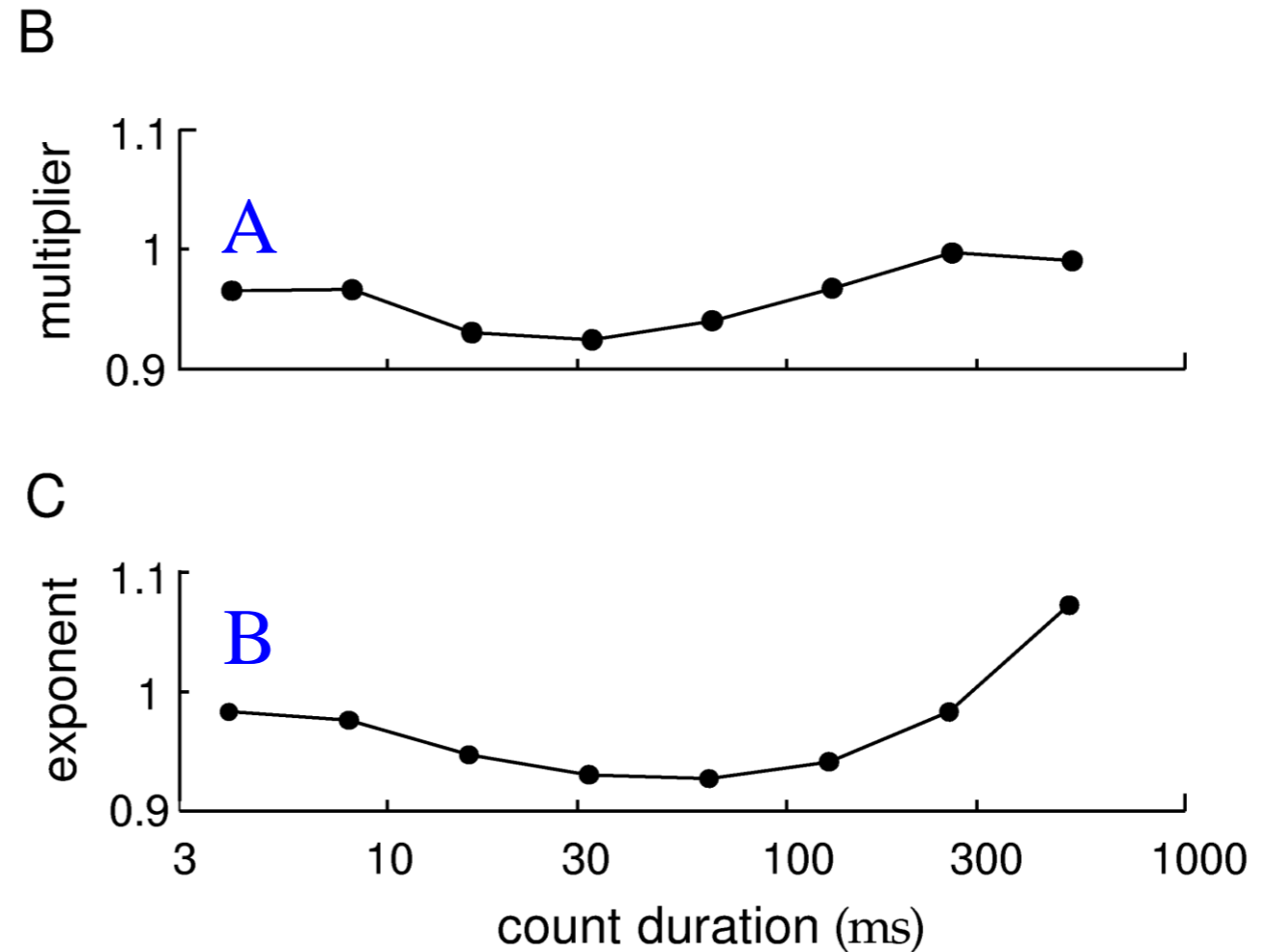
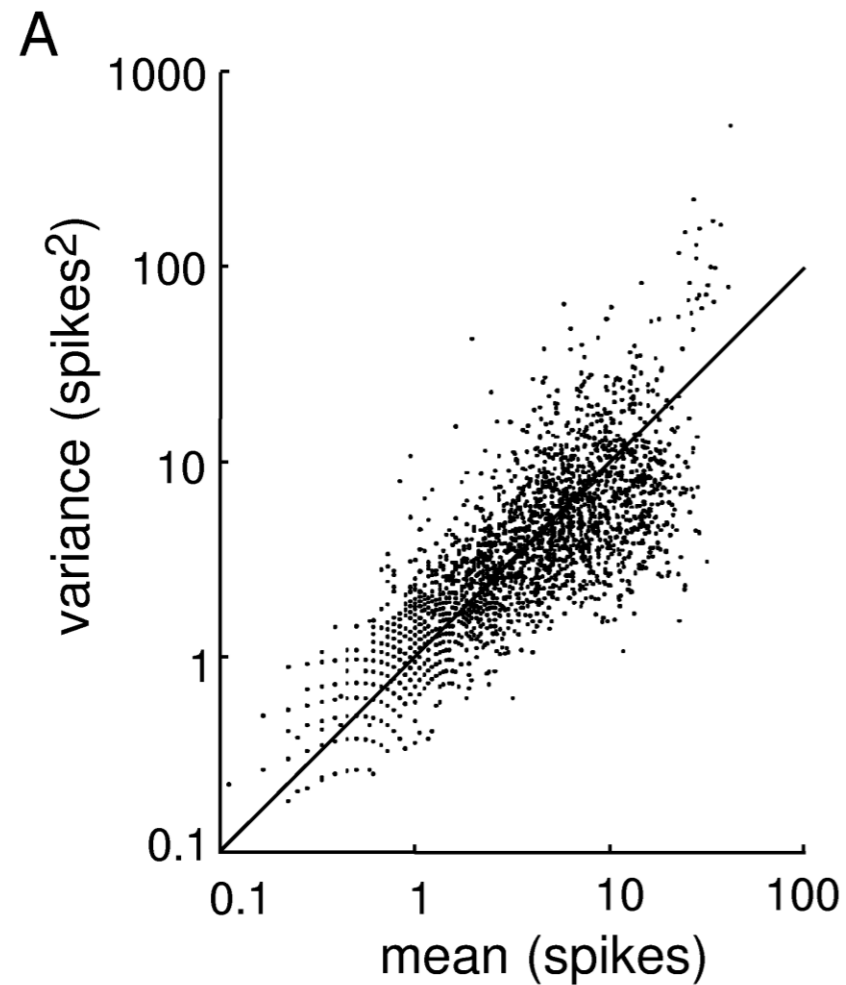
Poisson distribution

Area MT



Poisson spiking in the brain

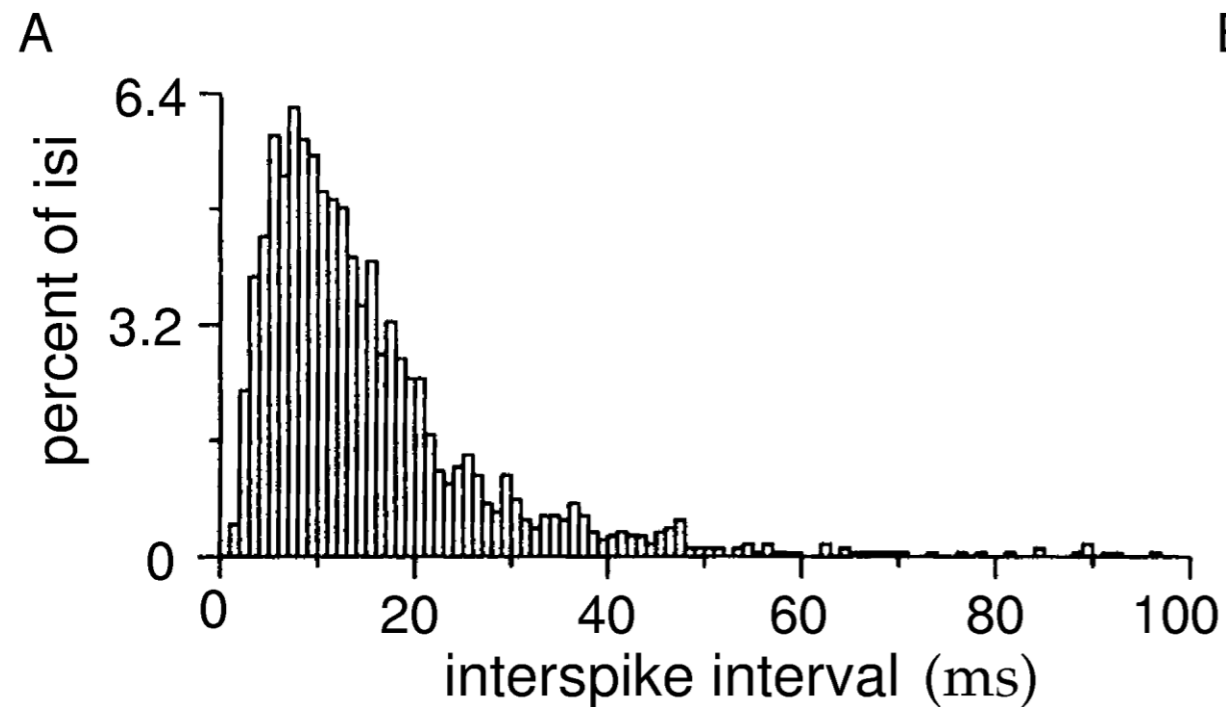
Area MT



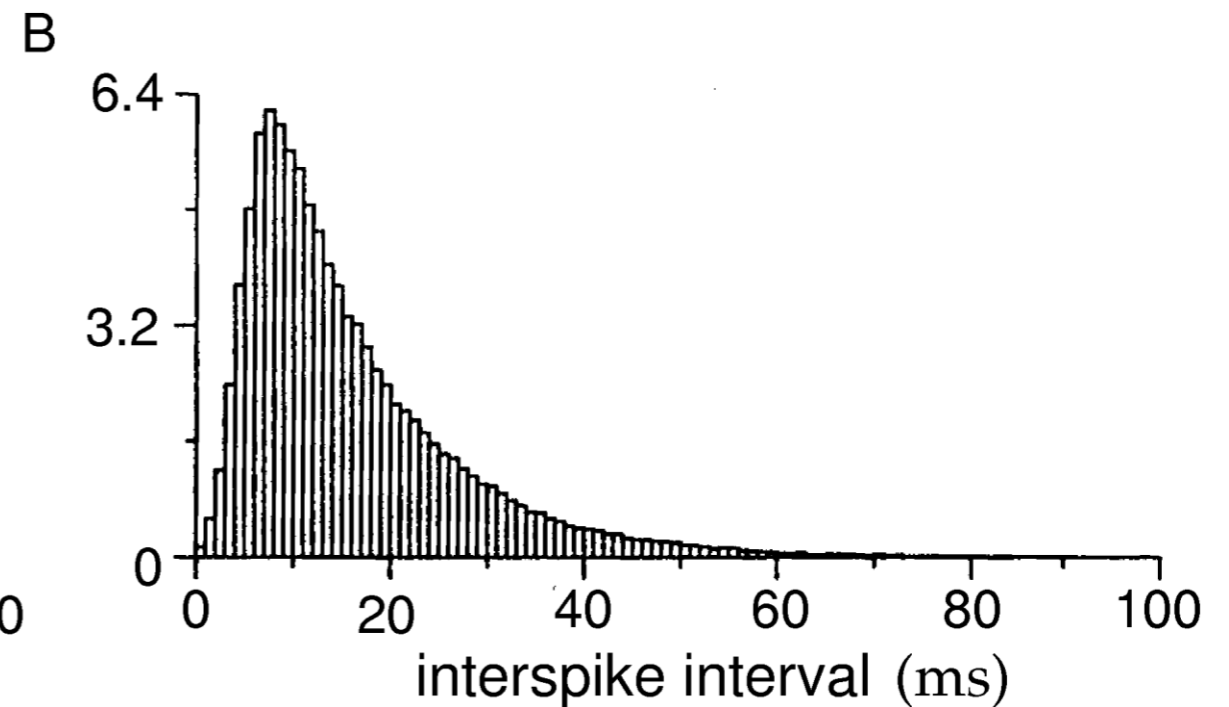
Fano factor

Data fit to:
 $\text{variance} = A \times \text{mean}^B$

Poisson spiking in the brain

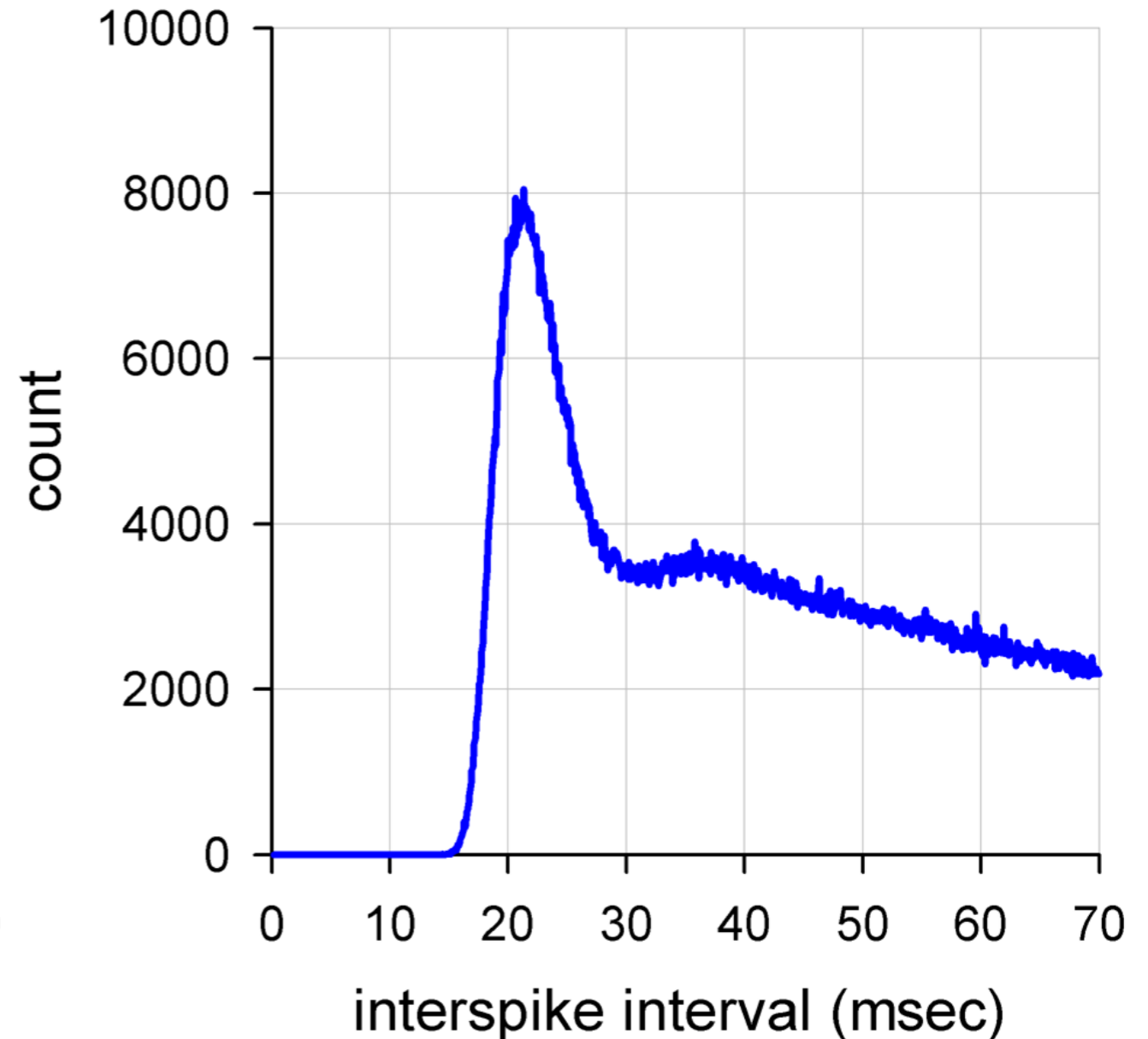
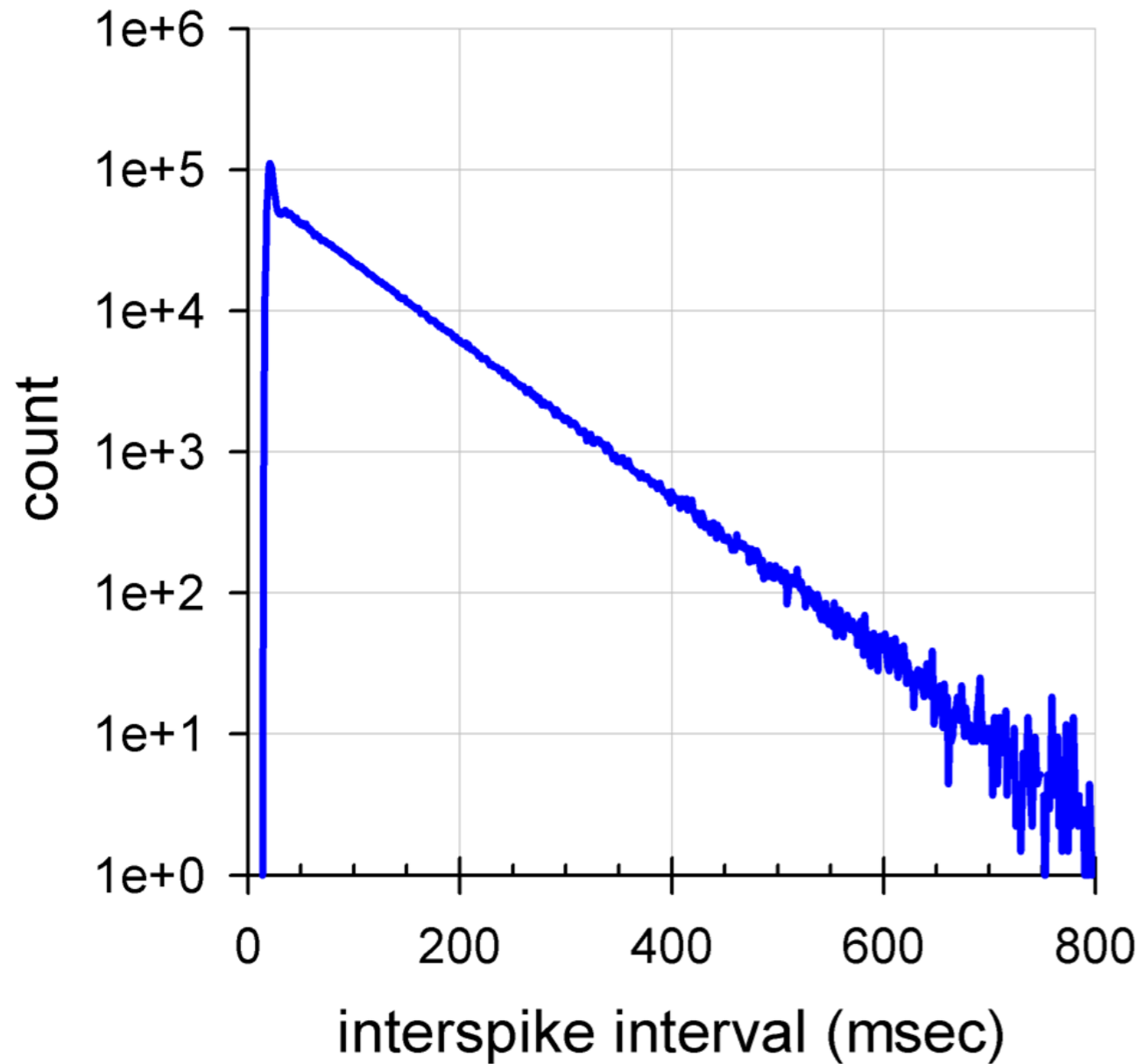


ISI Distribution from an
area MT Neuron

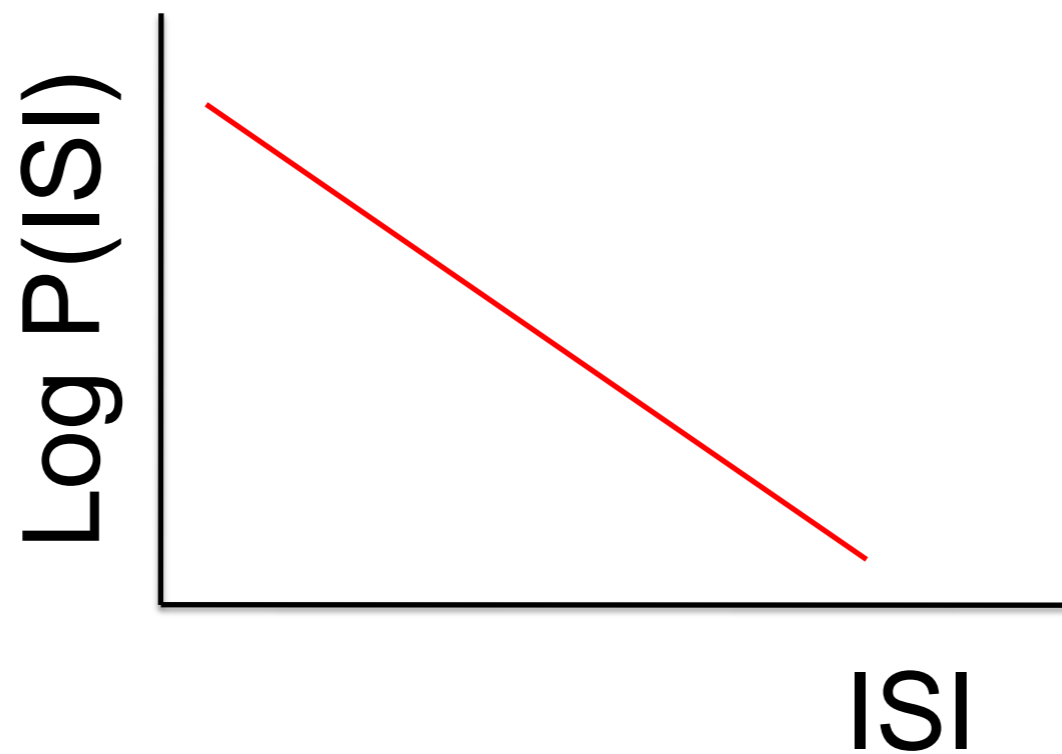
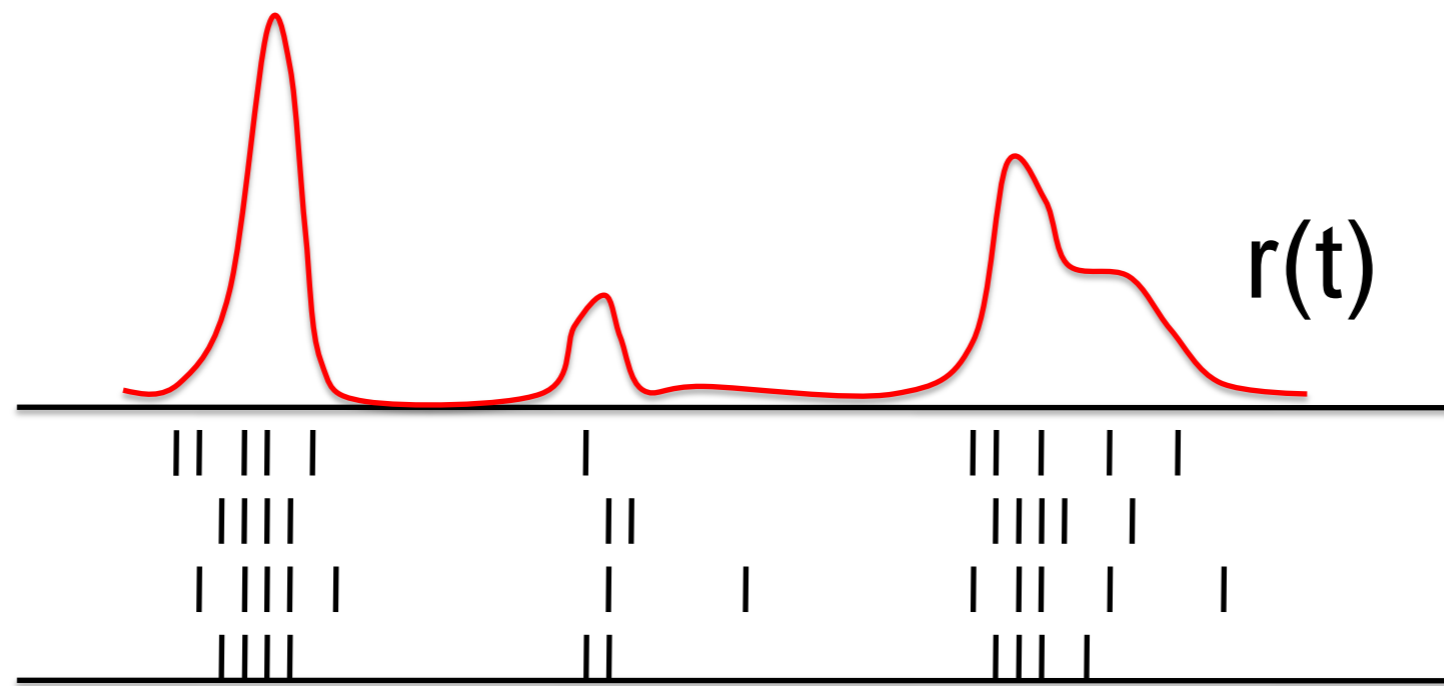


ISI distribution generated from
a Poisson model with a Gaussian
refractory period

Hodgkin-Huxley neuron driven by noise

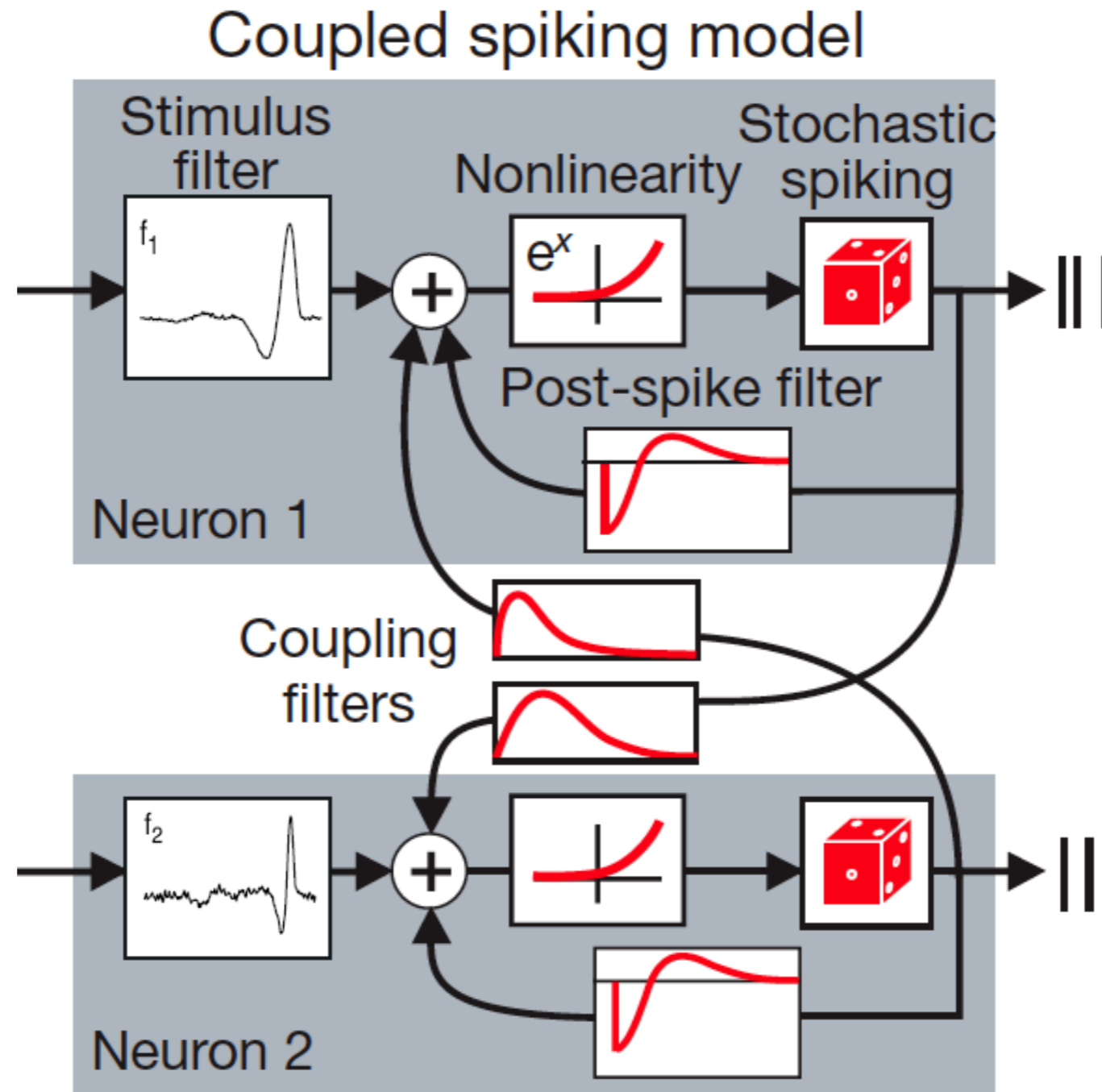


Time-rescaling theorem



Exponential distribution
of *scaled* ISIs:
 $(t_{i-1} - t_i) r(t)$

GLM



$$\text{GLM: } P(\text{spike at } t) = \exp(f_1 * s + h_1 * r_1 + h_2 * r_2 + \dots)$$