

# Coding and computation by neural ensembles in the primate retina

Liam Paninski

Department of Statistics and Center for Theoretical Neuroscience  
Columbia University

<http://www.stat.columbia.edu/~liam>

*liam@stat.columbia.edu*

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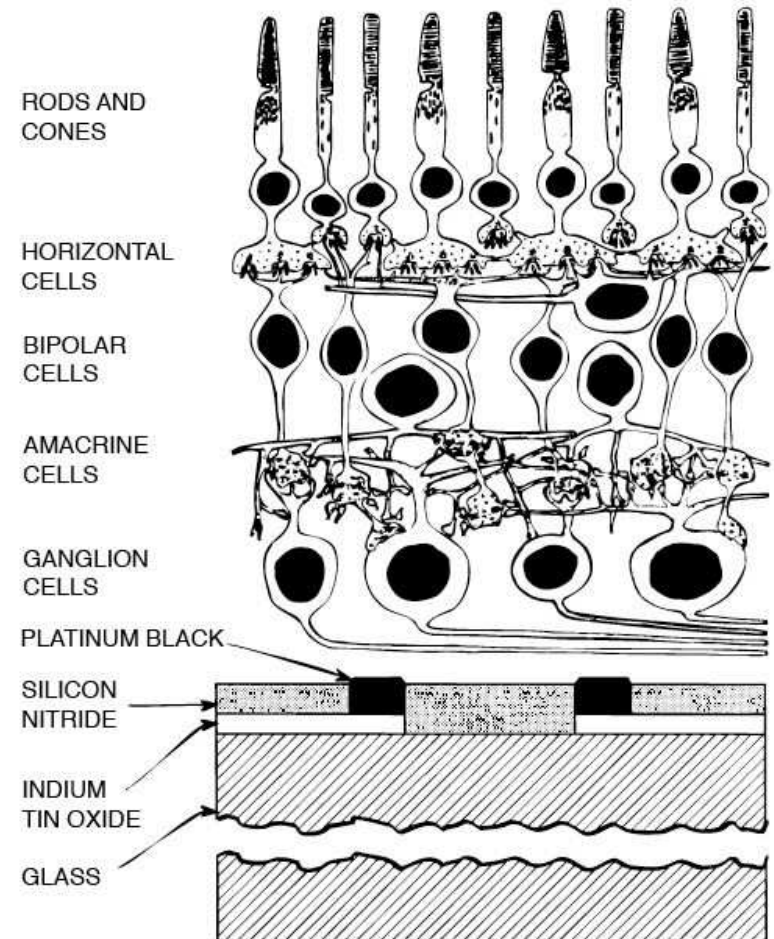
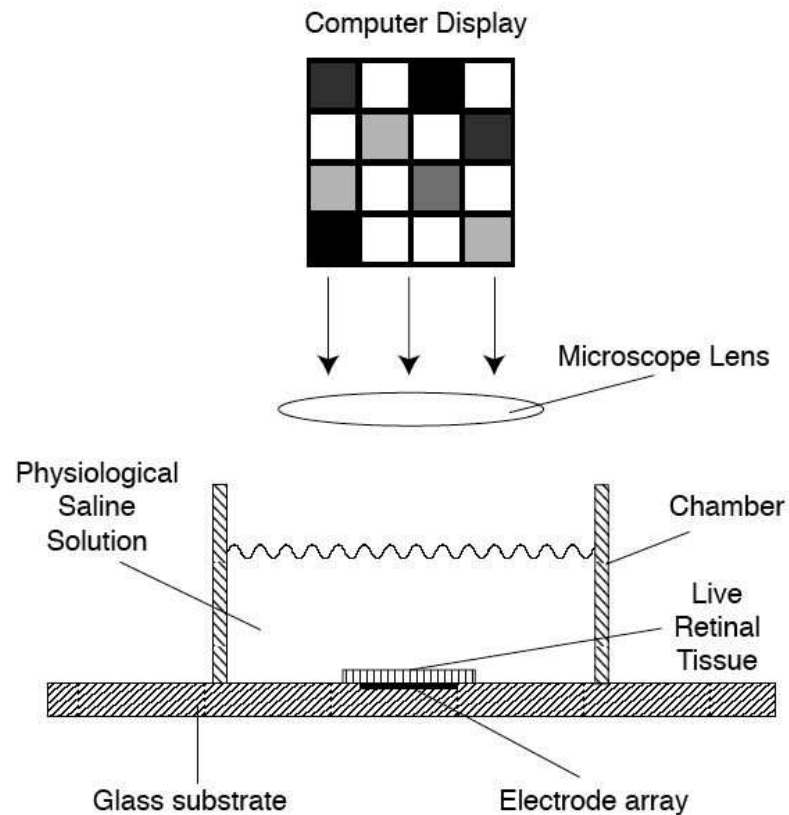
co-PI's: E. Simoncelli (NYU), E.J. Chichilnisky (Salk)

— with J. Pillow (UT Austin), G. Field, J. Gauthier, J. Shlens (Salk), A. Litke (UCSC), E. Lalor (TC Dublin), S. Koyama (CMU), **Y. Ahmadian**, J. Kulkarni, H. Liu, T. Machado, D. Pfau, X. Pitkow, **M. Vidne** (Columbia).

# Retinal ganglion neuronal data

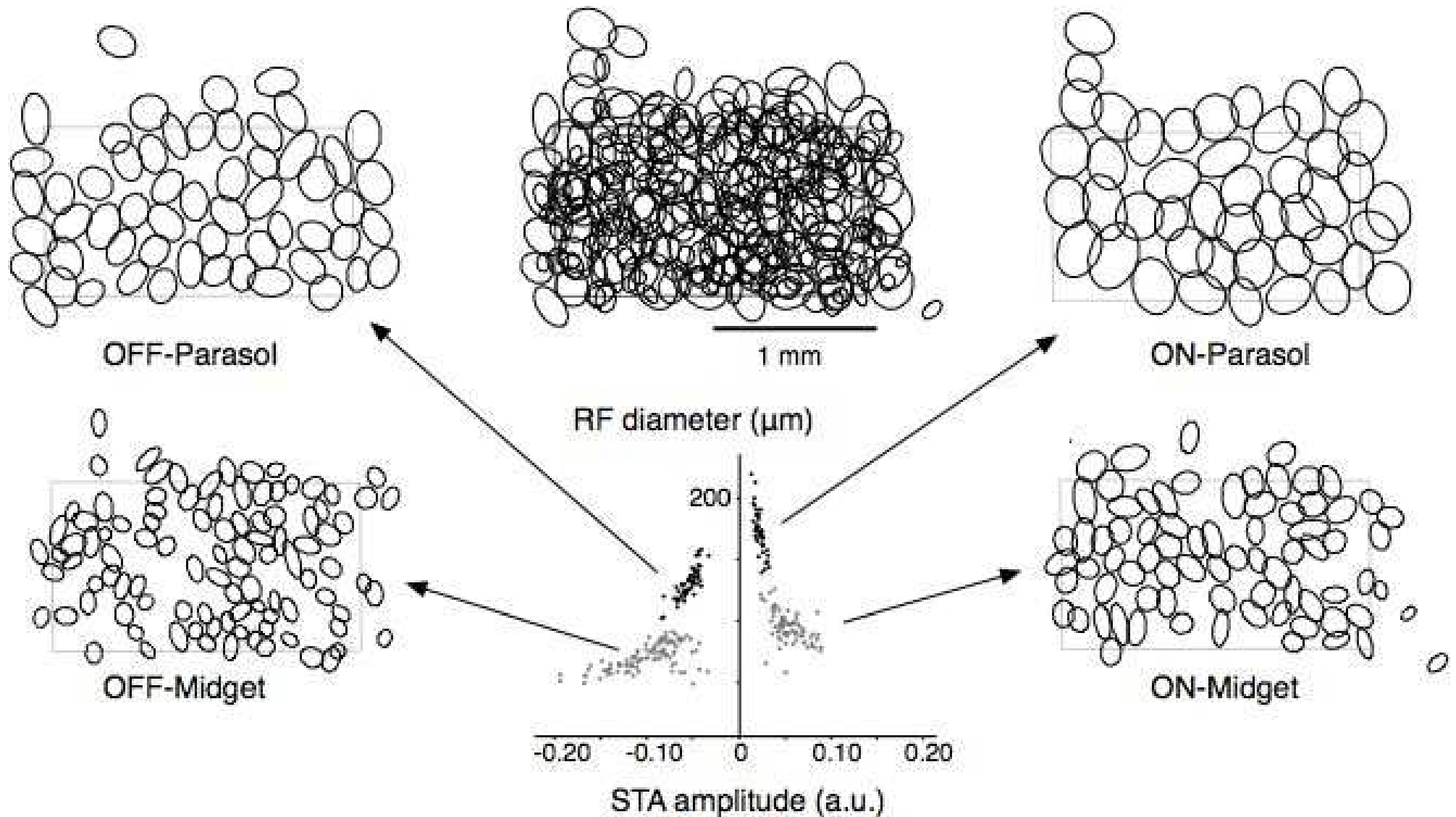
Preparation: dissociated macaque retina

— extracellularly-recorded responses of populations of RGCs

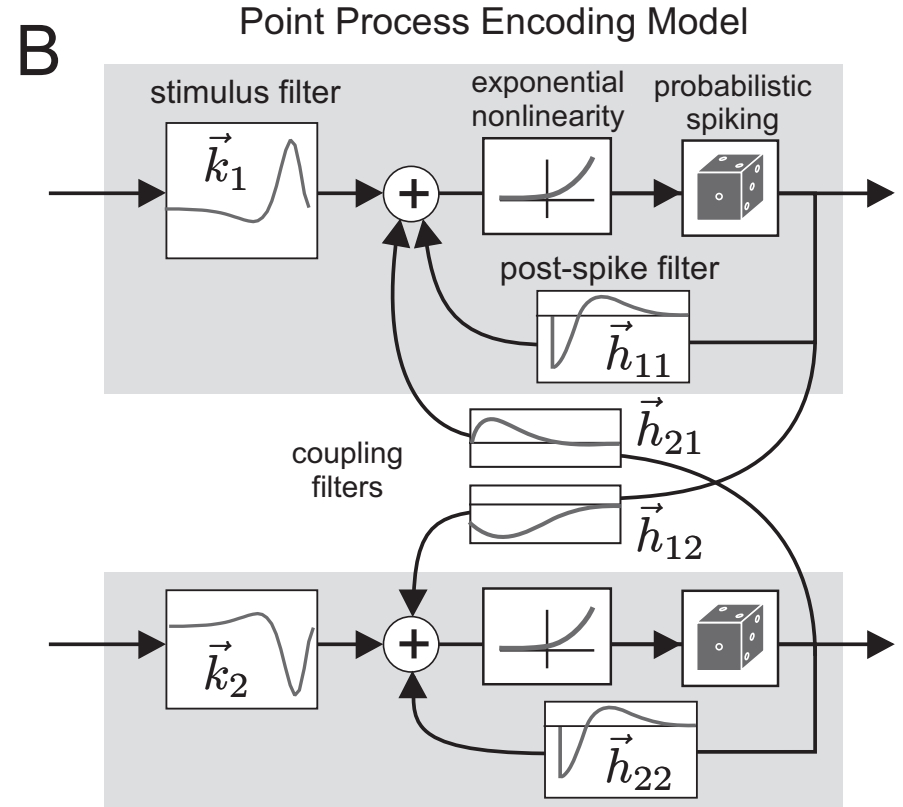
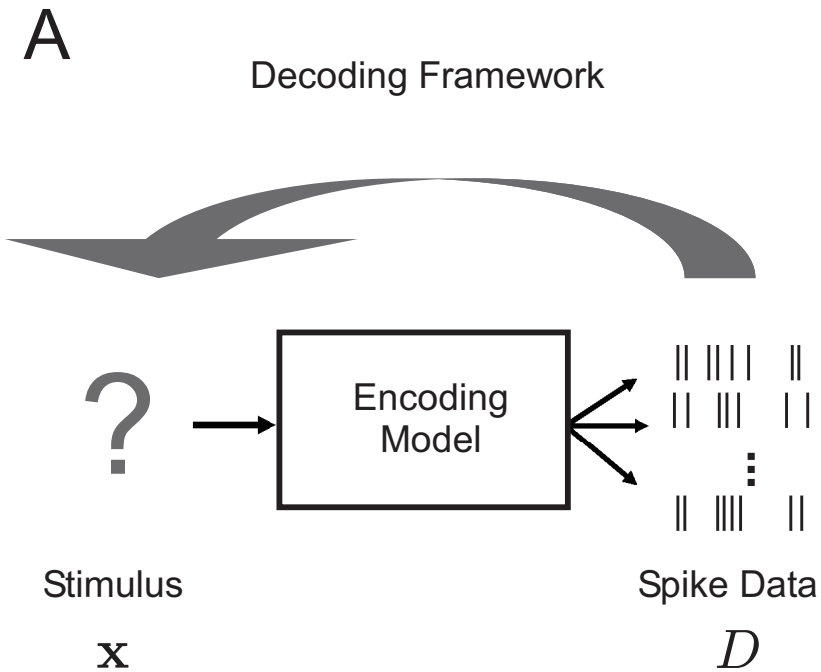


Stimulus: random spatiotemporal visual stimuli (Pillow et al., 2008)

# Receptive fields tile visual space



# Multineuronal point-process model



$$\lambda_i(t) = f \left( b_i + \vec{k}_i \cdot \vec{x}(t) + \sum_{i',j} h_{i',j} n_{i'}(t-j) \right),$$

— likelihood is easy to compute and to maximize (concave optimization)  
(Paninski, 2004; Paninski et al., 2007; Pillow et al., 2008)

— close connections to noisy integrate-and-fire model

# Optimal Bayesian decoding

$$E(\vec{x}|\text{spikes}) \approx \arg \max_{\vec{x}} \log P(\vec{x}|\text{spikes}) = \arg \max_{\vec{x}} [\log P(\text{spikes}|\vec{x}) + \log P(\vec{x})]$$

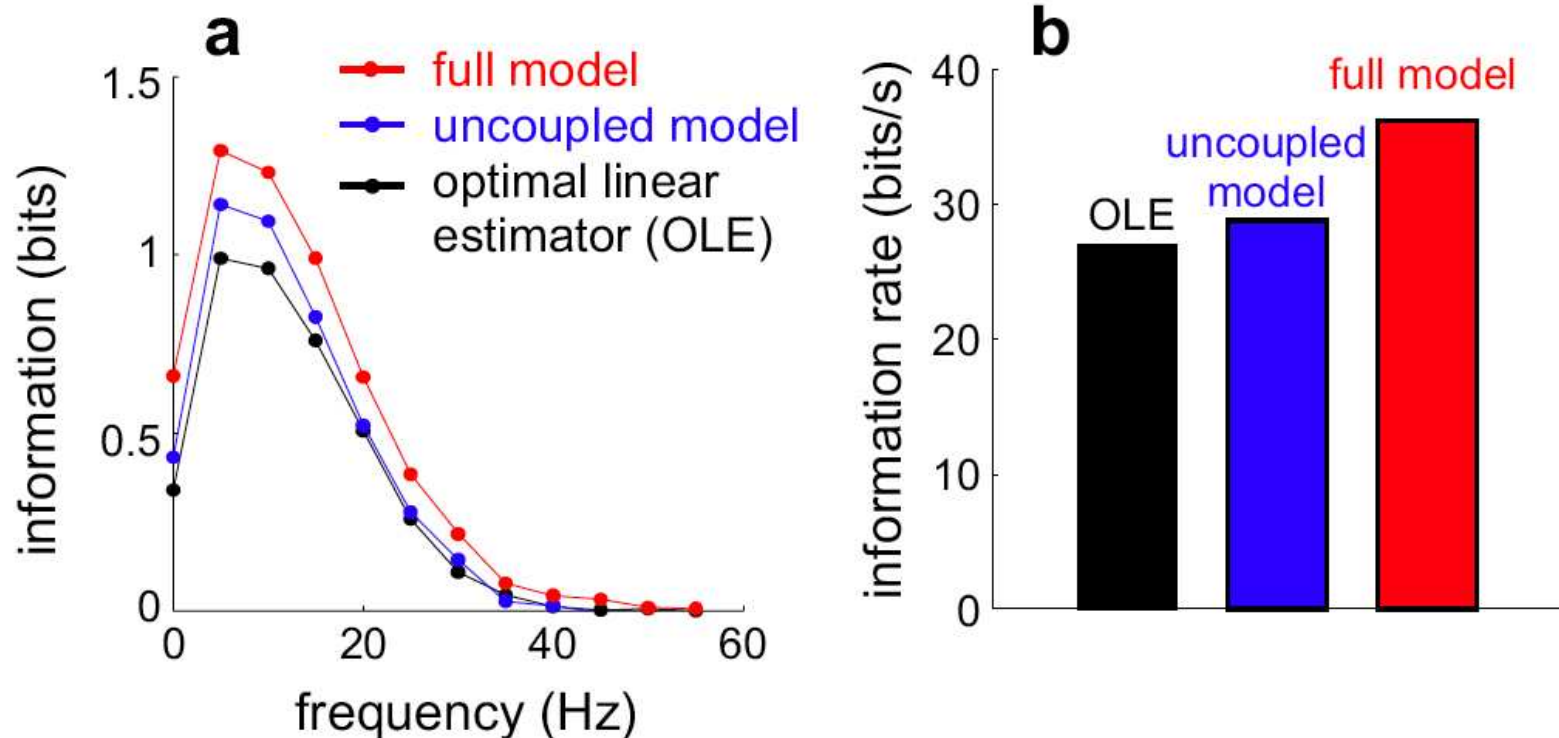
(Loading yashar-decode.mp4)

— Computational points:

- $\log P(\text{spikes}|\vec{x})$  is concave in  $\vec{x}$ : concave optimization again.
- Decoding can be done in linear time via standard Newton-Raphson methods, since Hessian of  $\log P(\vec{x}|\text{spikes})$  w.r.t.  $\vec{x}$  is banded (Pillow et al., 2010; Ahmadian et al., 2010).

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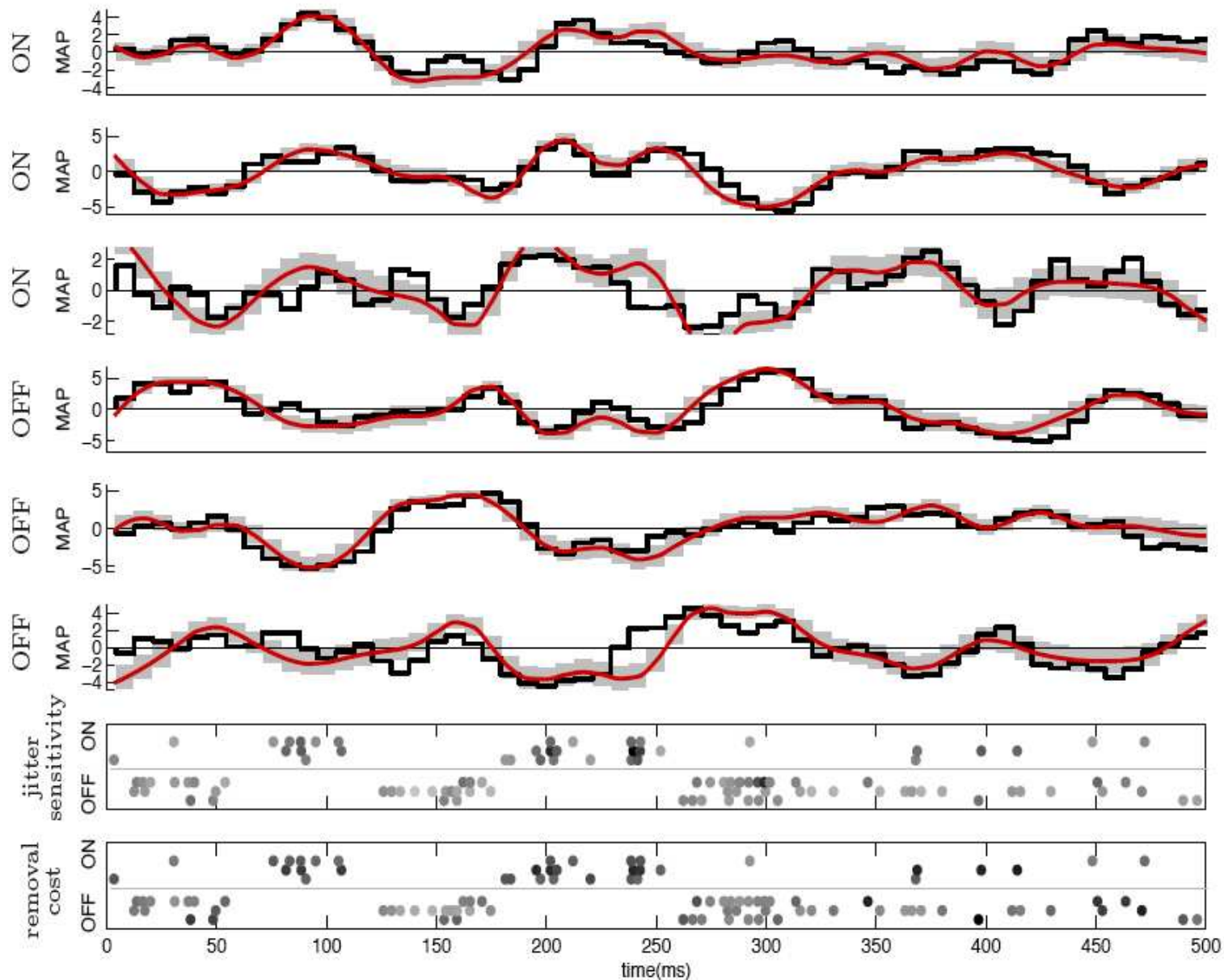


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— Biological point: paying attention to correlations improves decoding accuracy.

# Application: how important is timing?

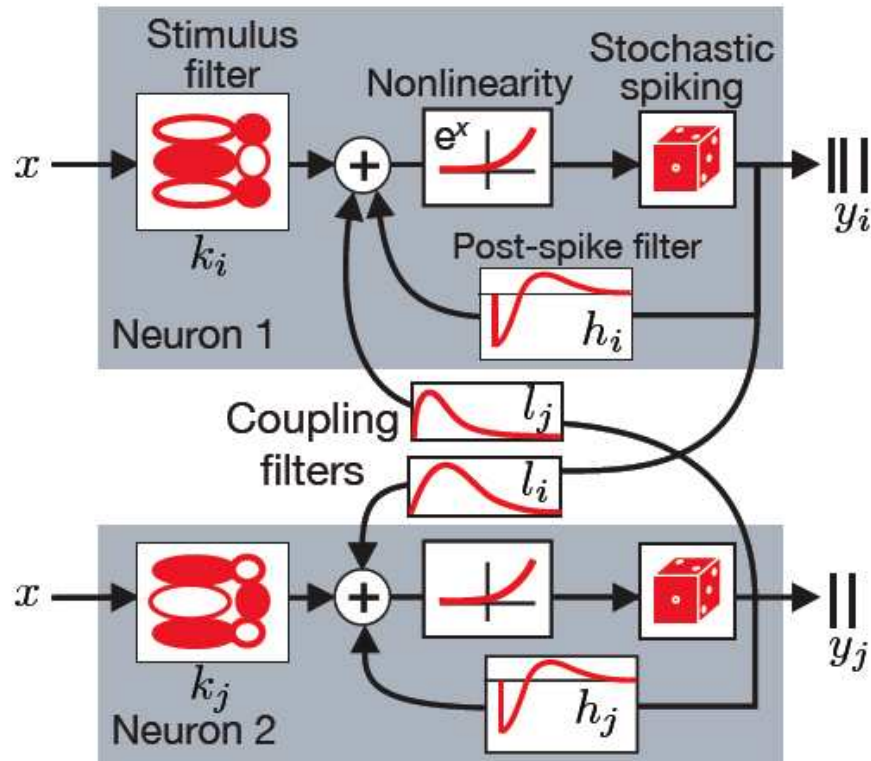


— further applications: decoding velocity signals (Lalor et al., 2009), tracking images perturbed by eye jitter (Pfau et al., 2009)



# Next steps: reconsidering the model

$$\lambda_i(t) = \exp \left( k_i \cdot x(t) + h_i \cdot y_i(t) + \sum_{i \neq j} l_{i,j} \cdot y_j(t) \right)$$



## Pros:

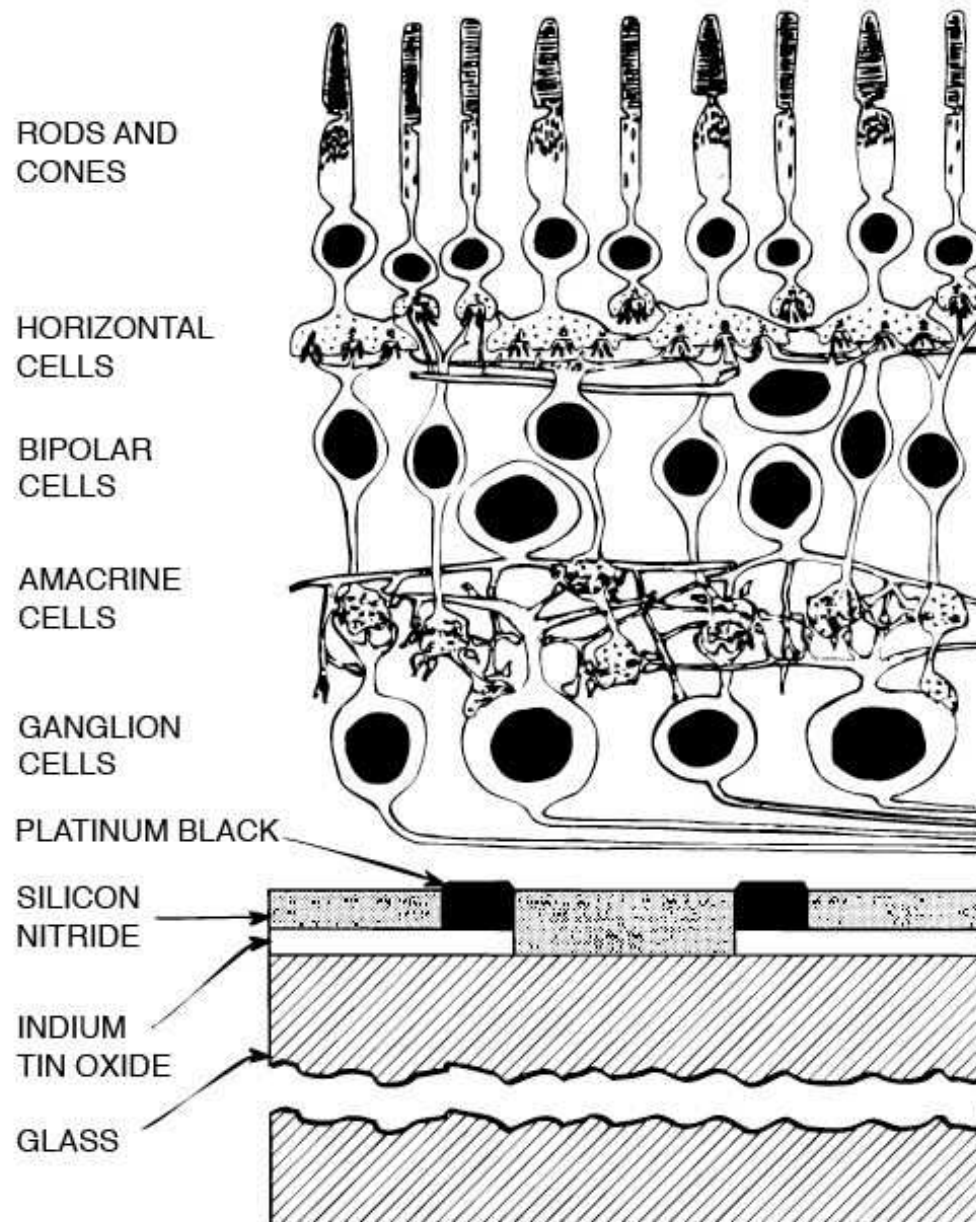
- Tractable model-fitting and optimal decoding
- Captures response statistics

## Cons:

- Instantaneous coupling filters
- No explicit Common Input



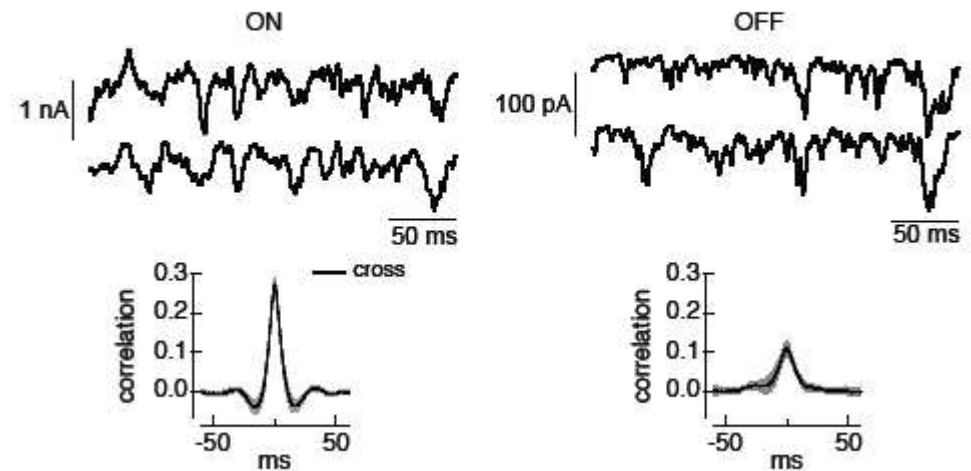
# Considering common input effects



— universal problem in network analysis: can't observe all neurons!

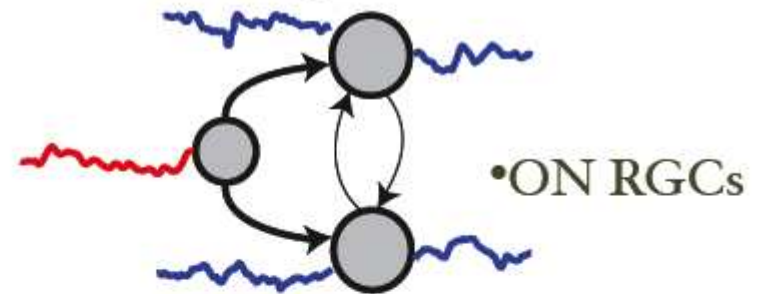
## Intracellular findings:

- RGCs receive strongly correlated synaptic input in the absence of modulated light stimuli

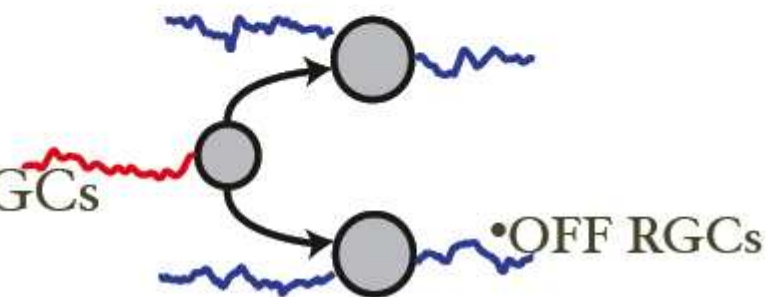


Khuc Trong & Rieke Nature Neuro 2008

- ON RGCs are weakly electrically coupled

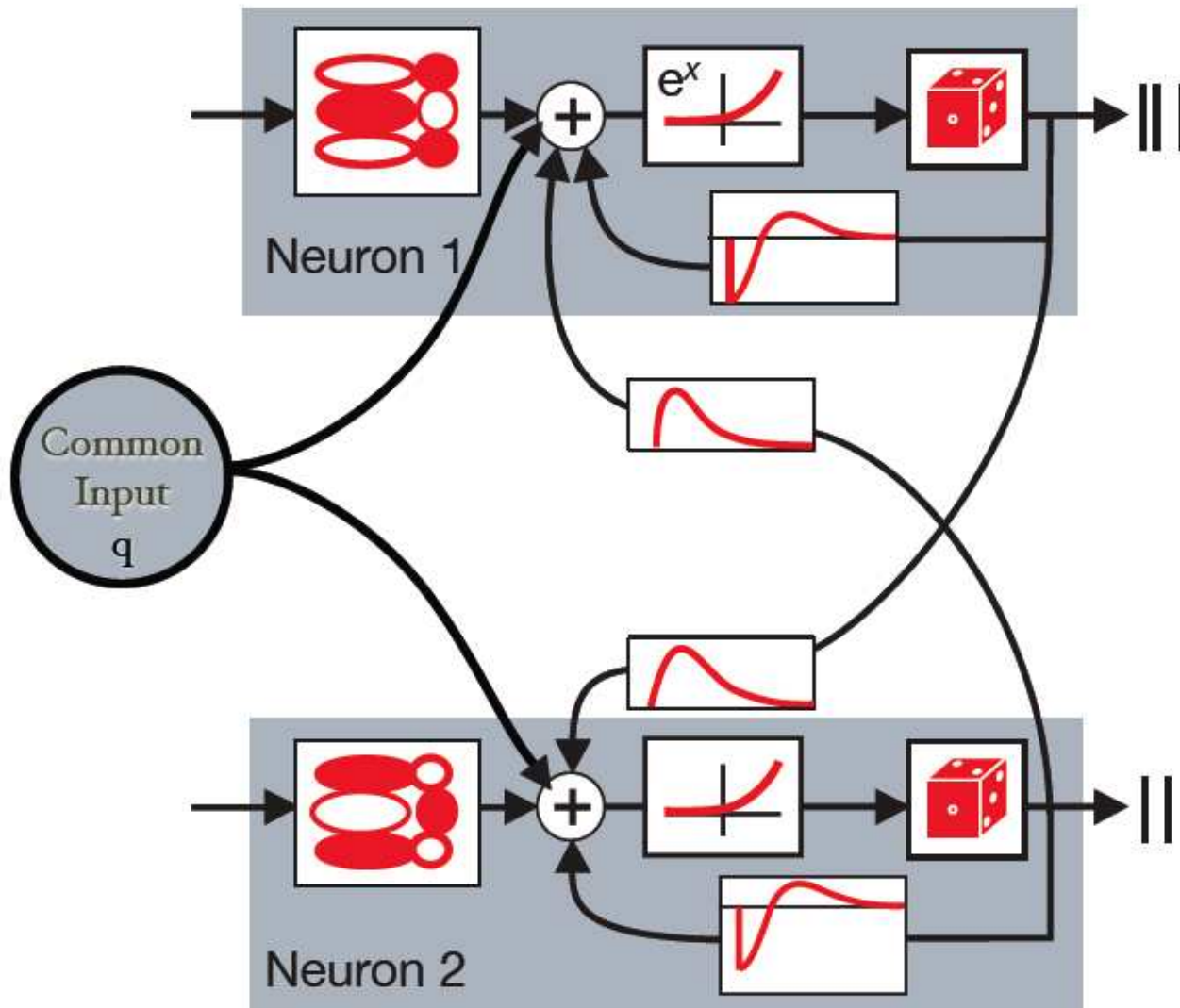


- No electrical coupling seen between OFF RGCs



# Extension: including common input effects

$$\lambda_i(t) = \exp \left( k_i \cdot x(t) + h_i \cdot y_i(t) + \sum_{i \neq j} l_{i,j} \cdot y_j(t) + Lq(t) \right)$$



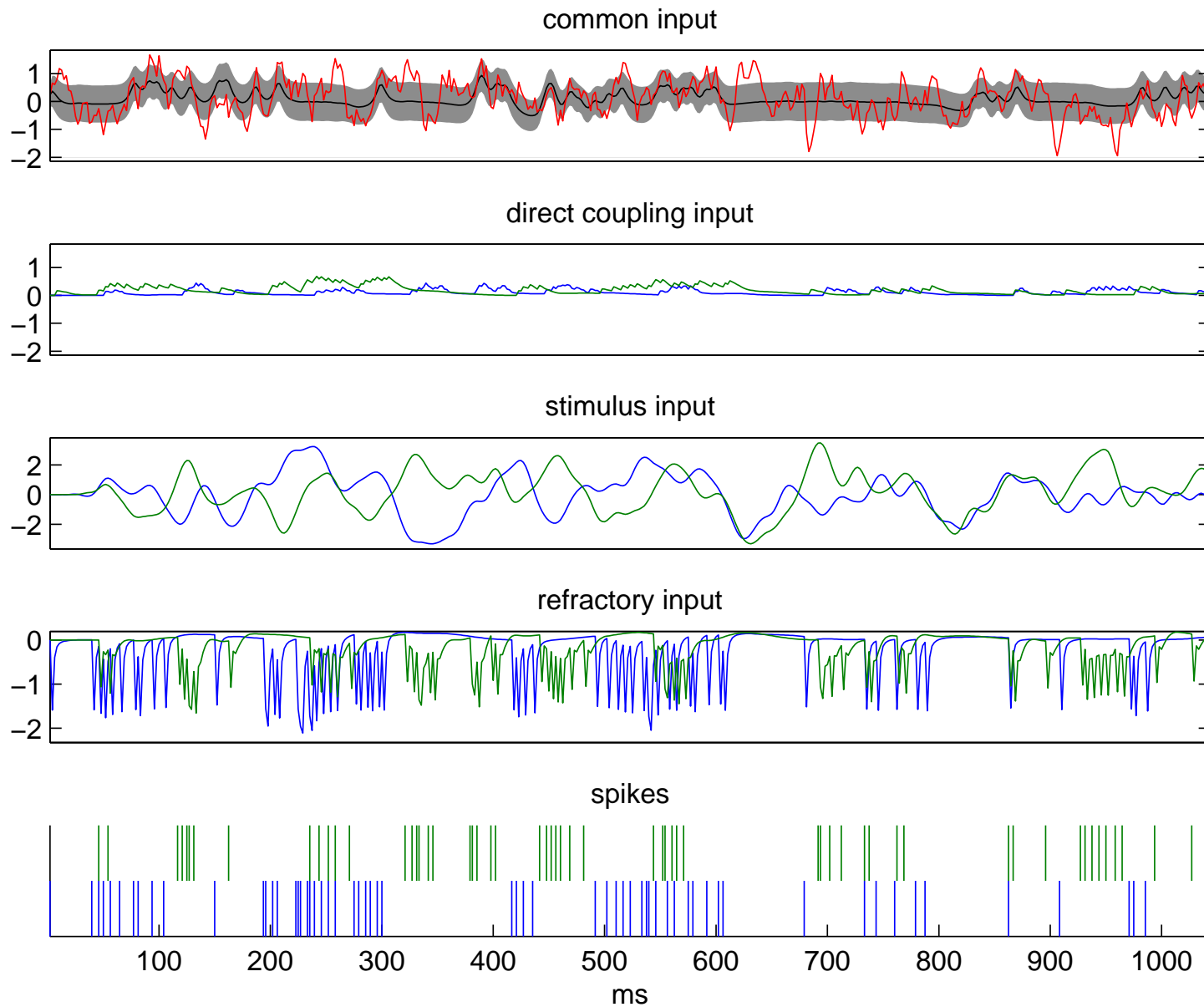
# Direct state-space optimization methods

To fit parameters, optimize approximate marginal likelihood:

$$\begin{aligned}\log p(\textit{spikes}|\theta) &= \log \int p(Q|\theta)p(\textit{spikes}|\theta, Q)dQ \\ &\approx \log p(\hat{Q}_\theta|\theta) + \log p(\textit{spikes}|\hat{Q}_\theta) - \frac{1}{2} \log |J_{\hat{Q}_\theta}| \\ \hat{Q}_\theta &= \arg \max_Q \{ \log p(Q|\theta) + \log p(\textit{spikes}|Q) \}\end{aligned}$$

- $Q$  is a very high-dimensional latent (unobserved) “common input” term. Taken to be a Gaussian process here with autocorrelation time  $\approx 5$  ms (Khuc-Trong and Rieke, 2008).
- correlation strength specified by one parameter per cell pair.
- all terms can be computed in  $O(T)$  via banded matrix methods (Paninski et al., 2010).

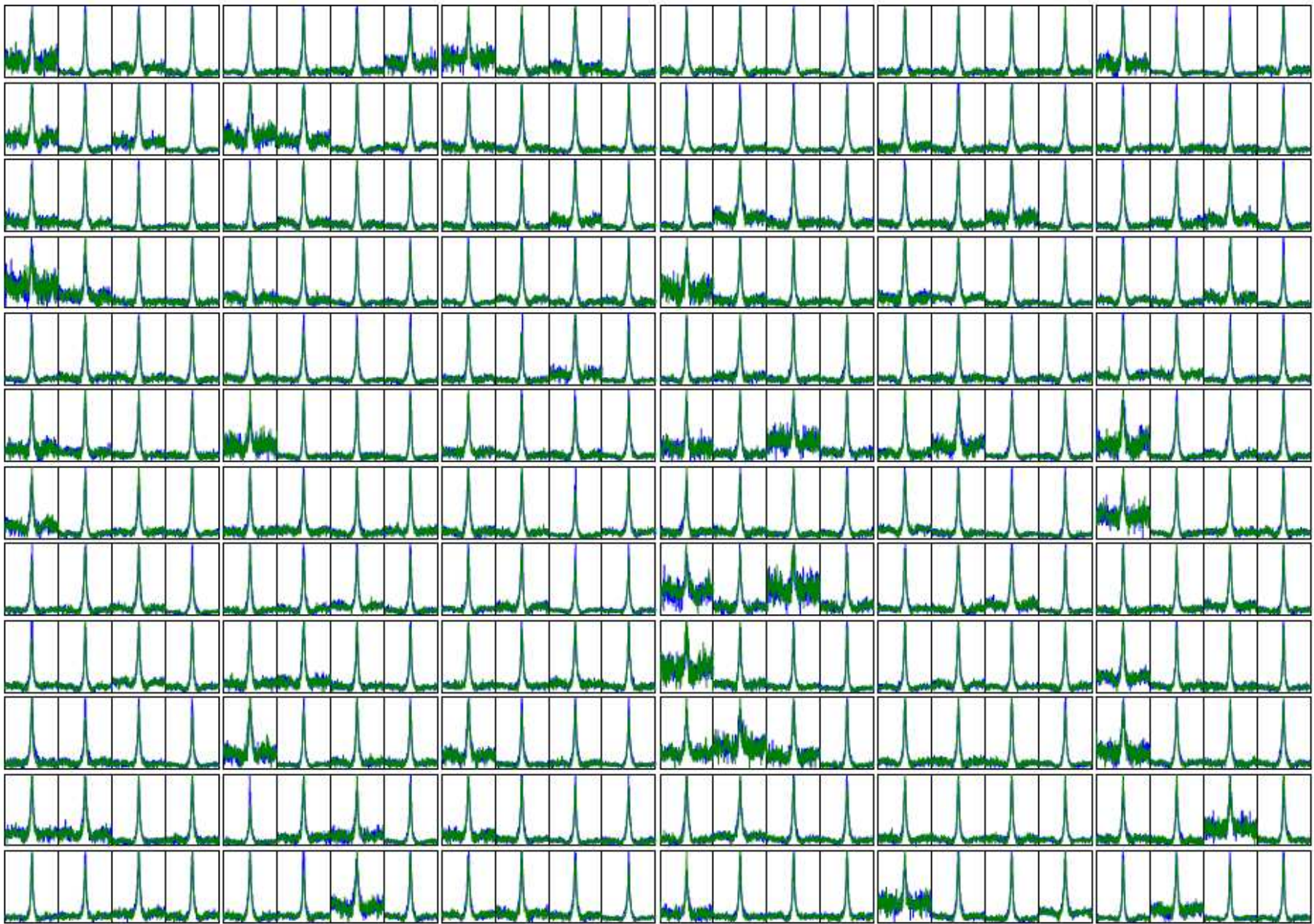
# Inferred common input effects are strong



— note that inferred direct coupling effects are now relatively small.



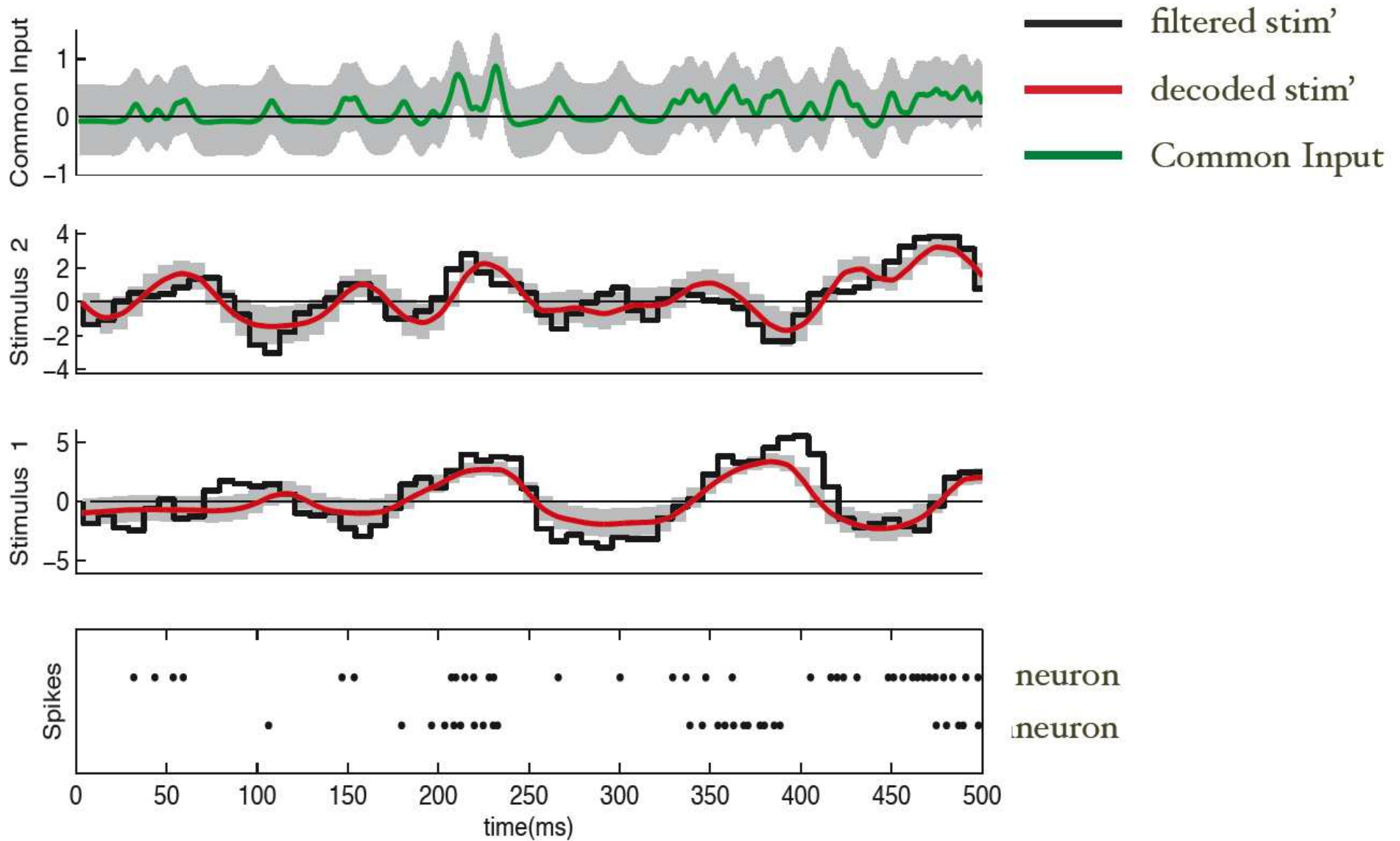
# Common-input-only model captures x-corrs



— single and triple-cell activities captured well, too (Vidne et al., 2009)

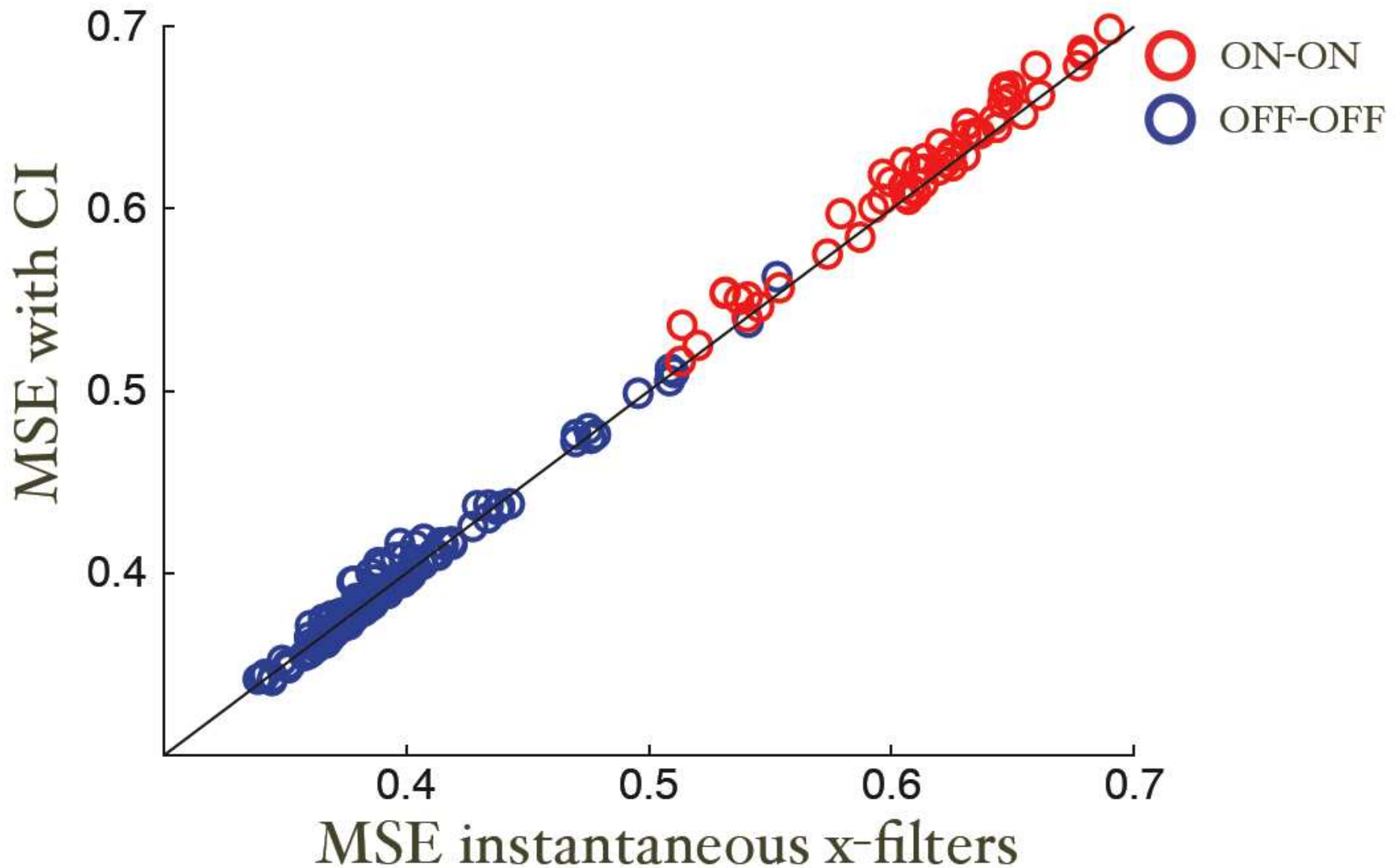


# Decoding the stimulus and hidden input



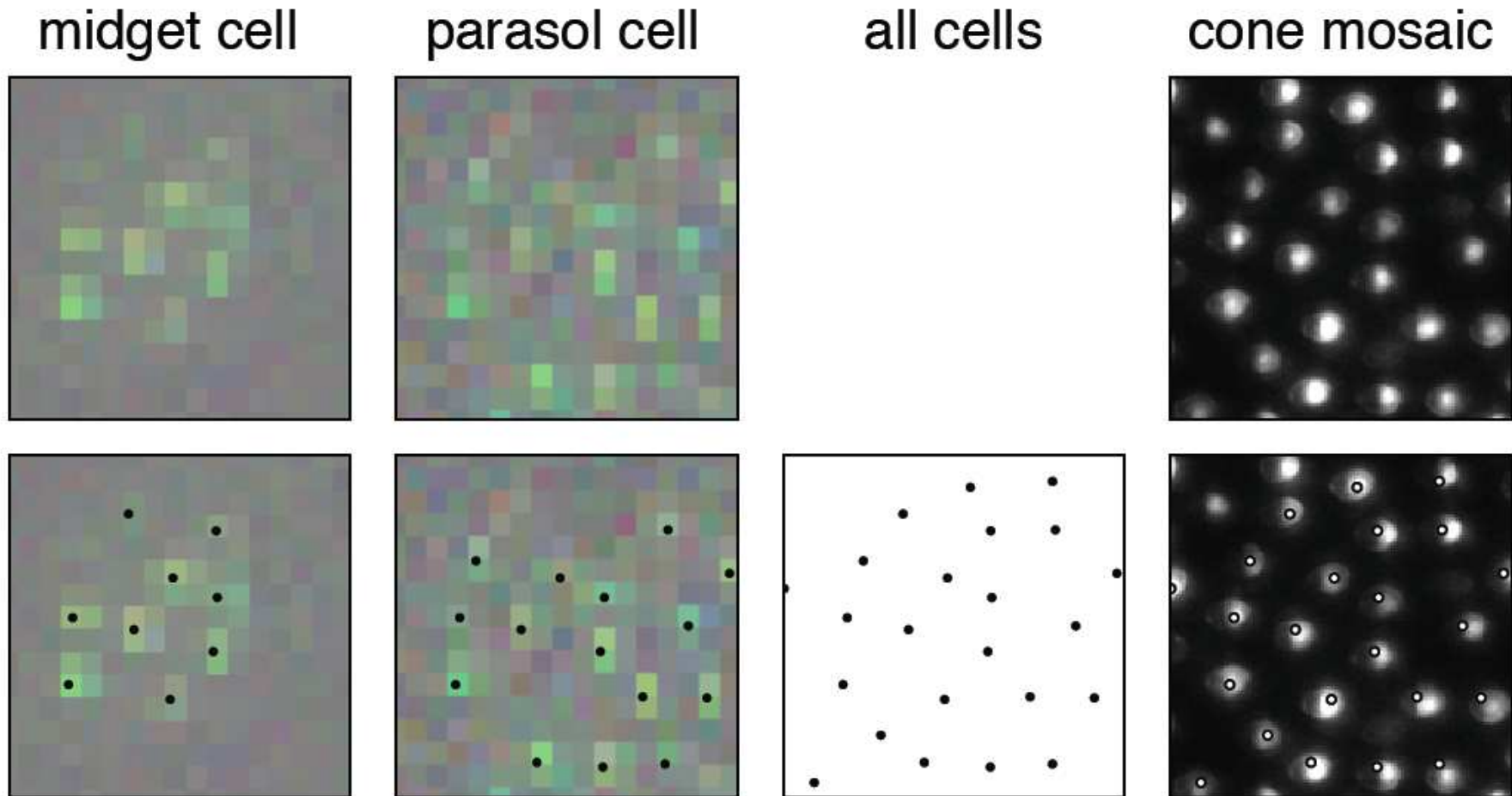
$$\arg \max_{\vec{x}} p(\vec{x}|y, \theta) = \arg \max_{\vec{x}} \int p(\vec{x}, Q|y, \theta) dQ \approx \arg \max_{\vec{x}, Q} p(\vec{x}, Q|y, \theta)$$

# Models lead to similar decoding performance



...but CI model is more robust to spike jitter and deletions (Vidne et al., 2009).

# Next steps: inferring cones



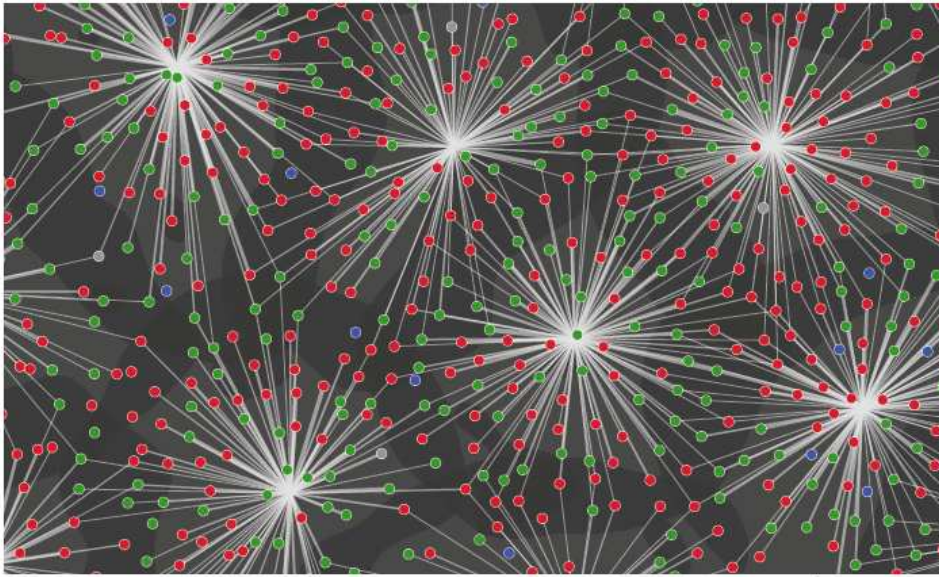
— cone locations and color identity can be inferred accurately with high spatial-resolution stimuli via maximum a posteriori estimates (Field et al., 2010).



# Next steps: inferring circuitry?

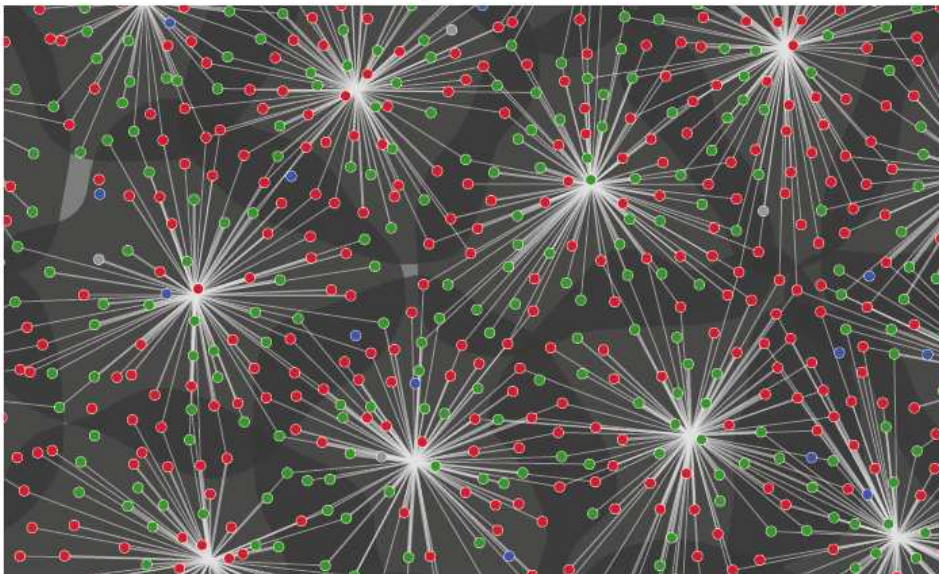
*ON parasol*

*retina 1,*



*OFF parasol*

50  $\mu\text{m}$



RODS AND CONES

HORIZONTAL CELLS

BIPOLAR CELLS

AMACRINE CELLS

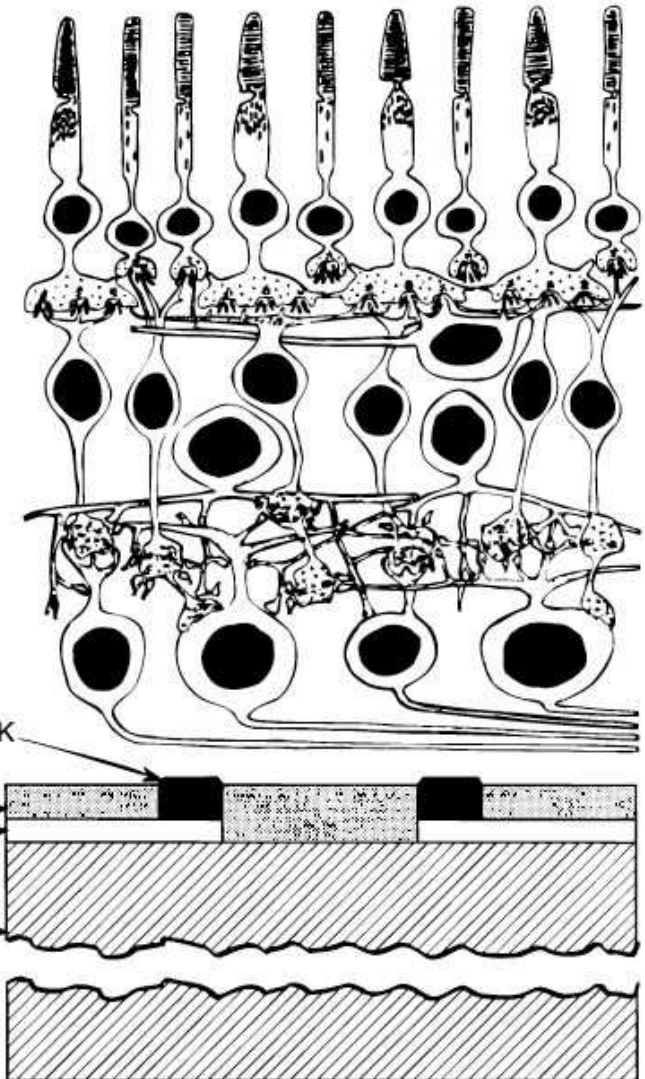
GANGLION CELLS

PLATINUM BLACK

SILICON NITRIDE

INDIUM TIN OXIDE

GLASS



# References

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