Jeremy Freeman, Greg Field, Peter Li, Martin Greschner, Lauren Jepson, Neil Rabinowitz, Eftychios Pnevmatikakis, Deborah Gunning, Keith Mathieson, Alan Litke, EJ Chichilnisky, Liam Paninski, Eero Simoncelli

INTRODUCTION

APPROACH TO FITTING



How do *circuit* elements give rise to *functional* response properties of retinal ganglion cells?

Anatomically, retinal ganglion cells (RGCs) are at the third stage of a hierarchical circuit containing two feed-forward stages of convergence and nonlinear transduction.

Functionally, spatial nonlinearities have been characterized in the responses of many RGC types, and modeled as nonlinear "subunits" within the receptive field (Hochstein and Shapley, 1976; Victor and Shapley, 1979). Subunits are thought to reflect bipolar cell circuitry (Demb et al., 2001; Crook et al., 2008), but it is challanging to directly record from this interneuron layer.

Computationally, the roles of many RGC types are influenced by subunits – e.g. detection of fine-grained patterns and spatial invariance – but quantitative models of subunit structure in terms of the elements of the circuit, especially at the population level, are lacking.

We measured spiking responses in large populations of midget RGCs by recording from isolated retinas using a multi-electrode array. We presented high-resolution visual noise stimuli, and fit a hierarchical model to describe the measured RGC responses. Given the complete input and output of the circuit, we infer properties of the interneuron computation, and our findings are consistent with properties of bipolar cells.

MODEL FITS

Single ON midget RGC





Punctate spots of light within the receptive field indicate cone locations, and allow us to transform the stimulus from pixel space to cone space. For most midget RGCs, the model recovers an accelerating subunit nonlinearity, with subunits of 1, 2, or 3 cones. Cross-validated improvement in explained variance is ~10-20%. If we only consider stimuli for which the models differ in their predictions, it exceeds 50%. Improvement is driven more by the subunit nonlinearity than by the cone-to-subunit assignments.

SPATIAL STRUCTURE AND ORGANIZATION OF NONLINEAR SUBUNITS IN PRIMATE RETINA

Model choices

- Subunits do not overlap
- Cone weights are positive
- Nonlinearities are splines
- Poisson spiking

Model fitting

 Block coordinate ascent on ikelihood to estimate cone weights, subunit weights, and nonlinearities • Greedy search over assignment matrices

baseline level of response. Paired to cones across subunits



subunits







Single retina summary



Midget RGC responses to contrast-modulated sinusoidal gratings exhibit "frequency doubling". The subunit model predicts frequency oubling but the LN model does not. Subunit model in greatest at high spatial frequencies.

Example cell



CONFIRMING SUBUNIT STRUCTURE WITH TARGETED STIMULI

Cancellation within a subunit

VALIDATING WITH GRATINGS



VALIDATING WITH NOISE

Example cell

The same 8-second sequence of noise was presented 50 times.



The subunit model captures more than 50% of the explainable variance on average, and outperforms the LN model by 40-50%.

Population



SUBUNITS VARY WITH ECCENTRICITY









Across multiple retinas, the relative fractions of 1, 2, and 3-cone subunits in the model fits approximately match anatomical predictions, and vary as expected with eccentricity.

