1	A minimal synaptic model for direction selective neurons in <i>Drosophila</i>
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3	Jacob A. Zavatone-Veth ^{1,2,a} , Bara A. Badwan ³ , Damon A. Clark ^{1,2,4,5,#}
4	1 – Department of Physics
5	2 – Department of Molecular, Cellular, and Developmental Biology
6	3 – School of Engineering and Applied Science
7	4 – Interdepartmental Neuroscience Program
8	5 – Department of Neuroscience
9	Yale University, New Haven, CT 06511, USA
10	
11	a – Present address: Department of Physics, Harvard University, Cambridge, MA 02138, USA
12	# – Corresponding author
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14	Abstract
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16	Visual motion estimation is a canonical neural computation. In Drosophila, recent advances have
17	identified anatomical and functional circuitry underlying direction-selective computations.
18	Models with varying levels of abstraction have been proposed to explain specific experimental
19	results, but have rarely been compared across experiments. Here we construct a minimal,
20	biophysically inspired synaptic model for Drosophila's first-order direction-selective T4 cells
21	using the wealth of available anatomical and physiological data. We show how this model relates
22	mathematically to classical models of motion detection, including the Hassenstein-Reichardt
23	correlator model. We used numerical simulation to test how well this synaptic model could
24	reproduce measurements of T4 cells across many datasets and stimulus modalities. These
25	comparisons include responses to sinusoid gratings, to apparent motion stimuli, to stochastic
26	stimuli, and to natural scenes. Without fine-tuning this model, it sufficed to reproduce many, but
27	not all, response properties of T4 cells. Since this model is flexible and based on straightforward
28	biophysical properties, it provides an extensible framework for developing a mechanistic
29	understanding of T4 neural response properties. Moreover, it can be used to assess the
30	sufficiency of simple biophysical mechanisms to describe features of the direction-selective
31	computation and identify where our understanding must be improved.

Motion estimation is a canonical visual computation that requires integrating information

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33 Introduction

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36 nonlinearly over both time and space. Direction-selective signals are tuned to motion in a 37 preferred-direction (PD), which elicits the strongest responses, while motion in the opposite, 38 null-direction (ND), elicits a weaker response. This directional computation has been described 39 by a wide variety of computational models. Classical models, such as the Hassenstein-Reichardt 40 correlator (HRC) (Hassenstein and Reichardt, 1956) and motion energy model (Adelson and 41 Bergen, 1985), rely on sensing correlations between pairs points separated in time and space. 42 These phenomenological models have provided striking insights into neural and behavioral 43 responses in a variety of species, including in flies (Yang and Clandinin, 2018). 44 45 In the last decade, advances in defining the anatomical and functional connectivity of 46 Drosophila's visual circuits suggest that we should move towards more mechanistic, biophysical descriptions of this computation. Here, we follow previous work (Gruntman et al., 2018; Torre 47 48 and Poggio, 1978) to propose a simple, biophysically-plausible synaptic model for direction-49 selectivity in Drosophila's ON-edge sensitive motion pathway. We compare its predictions to 50 measurements made by several research groups in response to many stimuli, giving us a tool for understanding which features are sufficient to describe different response properties. 51 52 53 The inputs to direction-selective cells have been identified by electron microscopy and through 54 genetic silencing experiments. The most peripheral direction-selective neurons in the Drosophila 55 optic lobe are the T4 and T5 cells, which are sensitive to moving ON-edges (consisting of 56 contrast increments) and OFF-edges (consisting of contrast decrements), respectively (Clark et 57 al., 2011; Joesch et al., 2010; Maisak et al., 2013). Electron microscopy and genetic silencing 58 have identified primary inputs to T4 and T5 cells (Serbe et al., 2016; Shinomiya et al., 2019; 59 Strother et al., 2017; Takemura et al., 2017). These studies suggest that T4 cells receive input 60 from three distinct colinear spatial locations, with the neurons Mi1 and Tm3 both relaying 61 information about the central point, and the neurons Mi9 and Mi4 acting as relays for the two

62 flanking points (Takemura et al., 2017) (Fig. 1A). The neuron T5 appears to have a similar

63 spatial structure, with different input neurons (Shinomiya et al., 2019). Both cell types also

64 receive spatially-localized inputs from other neurons, whose functions remain less well

understood (Shinomiya et al., 2019; Takemura et al., 2017).

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The physiological properties of the inputs to T4 and T5 cells have also been characterized. At 67 68 their receptive field centers, Mi1 and Tm3 cells respond quickly to visual stimuli, and provide 69 excitatory input to T4 (Arenz et al., 2017; Behnia et al., 2014; Gruntman et al., 2018; Strother et 70 al., 2017; Takemura et al., 2017). On the preferred direction side of the receptive field, the cells 71 Mi4 and CT1 are ON cells with slower kinetics, likely inhibiting T4 cells (Arenz et al., 2017; 72 Shinomiya et al., 2019; Takemura et al., 2017). On the null direction side of the receptive field, 73 Mi9 cells are delayed OFF cells, which are likely to provide inhibitory, glutamatergic input to T4 74 cells (Arenz et al., 2017; Salazar-Gatzimas et al., 2018). The inputs to T5 cells similarly appear 75 to be arranged with a fast central input and delayed flanking inputs, but whether these inputs 76 excite or inhibit T5 is less clear (Arenz et al., 2017; Behnia et al., 2014; Shinomiya et al., 2019; 77 Wienecke et al., 2018).

78

79 The functional properties of the T4 and T5 cells and their inputs have been interrogated using 80 many stimulus and measurement modalities. This wealth of data has led to many different 81 models that seek to describe the response properties of T4 and T5 cells (Arenz et al., 2017; 82 Badwan et al., 2019; Behnia et al., 2014; Clark et al., 2011; Creamer et al., 2018; Eichner et al., 83 2011; Gruntman et al., 2018; Haag et al., 2016; Leong et al., 2016; Leonhardt et al., 2016; 84 Salazar-Gatzimas et al., 2018; Salazar-Gatzimas et al., 2016; Serbe et al., 2016; Strother et al., 2017; Wienecke et al., 2018). Many measurements of T4 and T5 have demonstrated 85 86 phenomenology that could not be produced by the classical HRC model. However, proposed 87 models were most often evaluated by how they reproduced the associated dataset, rather than the 88 full the range of phenomena in the literature. Here we ask how a minimal, constrained model 89 reproduces T4 phenomenology (and some T5 phenomenology) from many different experiments. 90 We compare the model to data in response to moving edges, to sinusoids, to apparent motion 91 stimuli, to stochastic stimuli, and to natural scenes.

93 In this minimal model, the spatially-separated inputs to T4 are represented as three linear-94 nonlinear (LN) transformations of the input contrast (Dayan and Abbott, 2001). These model 95 neurons then interact with T4 by altering the conductance of excitatory and inhibitory currents 96 (Gruntman et al., 2018; Torre and Poggio, 1978). This construction is simple enough to allow 97 some algorithmic intuition but incorporates greater biophysical realism than most 98 phenomenological models. We do not fit the model to every dataset. Rather, our goal is to test 99 the sufficiency of a minimal circuit model to account for different measured phenomena in T4 100 cells. This model does not contain any exotic channels or receptors, and it biophysically models 101 the membrane voltage and intracellular calcium concentration in T4 neurons. It does not 102 reproduce all functional properties of T4 cells, but it provides a flexible framework for 103 understanding the sufficiency of simple circuit properties and mechanisms to describe the 104 processing properties of T4 neurons. In cases where this model is insufficient to describe data, 105 we suggest how model parameters might be changed to better describe the data. 106 107 **Methods** 108 109 Constructing an anatomically constrained synaptic model for T4 cells 110 Following proposed synaptic architectures for direction-selective computations (Gruntman et al., 111 2018; Torre and Poggio, 1978), we constructed an elementary motion detector based on the 112 connectome of the Drosophila optic lobe. We simplified this structure to consider three inputs to 113 a T4 cell: a delayed ND-offset OFF inhibitory input representing Mi9, a centered ON excitatory 114 input representing Mi1 and Tm3, and a delayed PD-offset ON inhibitory input representing Mi4

115 (and/or CT1) (**Figure 1A**) (Strother et al., 2017; Takemura et al., 2017).

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We will model these inputs to T4 cells as simple linear-nonlinear (LN) transformations of the input contrast (Behnia et al., 2014). We will further model effects of these synaptic inputs on the membrane potential of the T4 cell by changes in the conductance of excitatory and inhibitory currents (Torre and Poggio, 1978). For notational convenience, we define our model below in continuous space and time, noting as needed where adjustments are made for the discretization used in numerical simulation. We take the inputs to the model to be contrasts. We take each input to the motion detector to have an L_1 -normalized Gaussian spatial acceptance function

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125
$$h(x) = \frac{1}{\sqrt{2\pi\sigma^2}} e^{-\frac{x^2}{2\sigma^2}}$$

126

127 where the spatial parameter σ is related to the full width at half maximum (FWHM) of the 128 acceptance function by FWHM = $2\sqrt{2 \log 2} \sigma$. We fix FWHM = 5.7° to approximately match 129 the spatial acceptance functions of photoreceptors in the fly eye (Stavenga, 2003). To represent 130 the delayed inputs to the motion detector, we use the L_2 -normalized lowpass temporal filter 131

132
$$f(t) = 2 \tau^{-\frac{3}{2}} t \Theta(t) e^{-\frac{t}{\tau}}$$

133

134 where $\Theta(x)$ is the Heaviside step function. To represent the non-delayed central input to the 135 motion detector, we replace the temporal filter f by its derivative \dot{f} . We note that the term 136 resulting from the distributional derivative of $\Theta(t)$ vanishes when \dot{f} is convolved with any signal 137 as it is proportional to $t \,\delta(t)$, where $\delta(x)$ is the Dirac delta distribution. Using these filters, we 138 define the filtered contrast signal s at each point in spacetime:

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- 140 $s(t,x) \coloneqq (fh * c)(t,x)$
- 141

where c(t, x) is the input contrast and * denotes spatiotemporal convolution over the appropriate domain. As taking the temporal derivative of the filtered contrast signal is equivalent to filtering with the derivative of the temporal filter, we will use the notation \dot{s} for the high-pass-filtered signal throughout. For convenient handling of spatial boundary conditions, we numerically simulate the full 360° of visual space, which is a periodic interval.

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We denote the spacing between neighboring inputs as Δ . Here, we use 5° spacing so that the inputs evenly tile 360° of visual space. Then, we define the three inputs to the motion detector as rectified-linear functions of the filtered contrast signal at three points in space, mimicking the polarity-selectivity of the inputs to T4 cells:

153
$$g_1(t,x) \coloneqq g_{\sinh} R \left(-s(t,x-\Delta) \right)$$

154
$$g_2(t,x) \coloneqq g_{\text{exc}} R(\dot{s}(t,x))$$

155
$$g_3(t,x) \coloneqq g_{\rm inh} R(s(t,x+\Delta))$$

156

157 where $R(x) \coloneqq \max\{0, x\}$ is the ramp function and g_{inh} and g_{exc} are parameters scaling the 158 effects of each input on the postsynaptic conductances (**Figure 1A-B**). Thus far, we have 159 represented the conductances as linear-nonlinear (LN) transformations of the input contrast 160 (Dayan and Abbott, 2001).

161

We define the membrane potential $V_{\rm m}$ of the postsynaptic cell such that the reversal potential for leak currents is 0 mV. The cell's membrane voltage dynamics are then given as (Torre and Poggio, 1978)

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$$c_{\rm m}\dot{V}_{\rm m} + V_{\rm m}(g_{\rm leak} + g_1 + g_2 + g_3) = g_1E_{\rm inh} + g_2E_{\rm exc} + g_3E_{\rm inh}$$

167

where $c_{\rm m}$ is the membrane capacitance, $g_{\rm leak}$ is the leak conductance, and $E_{\rm inh}$ and $E_{\rm exc}$ are the reversal potentials for inhibitory and excitatory currents, respectively. Neglecting capacitive currents, we solve for the pseudo-steady-state (Gruntman et al., 2018; Torre and Poggio, 1978).

172
$$V_{\rm m} = \frac{g_1 E_{\rm inh} + g_2 E_{\rm exc} + g_3 E_{\rm inh}}{g_{\rm leak} + g_1 + g_2 + g_3}$$

173

174 Then, we model the transformation from membrane voltage to calcium concentration *C* as a 175 positively rectifying half-quadratic function $R^2(x) \coloneqq (R(x))^2$:

- 176
- 177 $C(t,x) \coloneqq R^2 \big(V_{\rm m}(t,x) \big)$
- 178

179 which qualitatively captures the expansive nonlinear effect of the transformation between

180 voltage and calcium (Kato et al., 2014; Leong et al., 2016) (**Figure 1B**).

182 Visual stimuli

We presented this model with spatiotemporal contrast patterns to mimic a variety of visual stimuli used in the field. Detailed mathematical descriptions of each stimulus are given in Appendix A. Briefly, we presented the model with moving and stationary sinusoidal gratings, with apparent motion stimuli, with stochastic stimuli including those with imposed correlations, and with natural scenes. In each case, we compared how the model responds to the published responses of T4 and T5 neurons.

189

190 Selecting model parameters

191 This model uses a parameter set equal to one that was developed to explain direction-opponency

in T4 cells (Badwan et al., 2019). There, we fixed the filter time constant $\tau = 150$ ms to produce peak responses to PD sinusoidal gratings at ~1 Hz (Badwan et al., 2019; Creamer et al., 2018; Maisak et al., 2013). We fix the excitatory and inhibitory reversal potentials to values of $E_{\text{exc}} =$ 60 mV and $E_{\text{inh}} = -30$ mV, which are plausible based on electrophysiological experiments (Gruntman et al., 2018). As the model membrane potential can be rewritten as

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$$V_{\rm m} = \left(\frac{g_1}{g_{\rm leak}}E_{\rm inh} + \frac{g_2}{g_{\rm leak}}E_{\rm exc} + \frac{g_3}{g_{\rm leak}}E_{\rm inh}\right) \left(1 + \frac{g_1}{g_{\rm leak}} + \frac{g_2}{g_{\rm leak}} + \frac{g_3}{g_{\rm leak}}\right)^{-1}$$

199

200 only the ratios of g_1 , g_2 , and g_3 to g_{leak} , rather than their absolute magnitudes, are relevant. We therefore express the postsynaptic conductances as non-dimensional quantities in units of g_{leak} , 201 leaving $g_{\rm exc}/g_{\rm leak}$ and $g_{\rm inh}/g_{\rm leak}$ as the model's two free parameters. The procedure used to 202 203 select the values of these parameters is described in detail in **Appendix B**. As shown previously 204 (Badwan et al., 2019), there exists a broad region of parameter space for which this model 205 displays responses to sinusoid gratings with a temporal frequency of 1 Hz and a spatial 206 wavelength of 45° consistent with those measured in T4 and T5 cells. We note that our choice of 207 filter normalization, which differs from that in the previous use of this model (Badwan et al., 2019), affects the parameter values chosen, as it scales g_1 , g_2 , and g_3 relative to g_{leak} . Table 1 208 209 summarizes the model parameter values used in all simulations.

Parameter	Value
Photoreceptor spacing Δ	5°
Photoreceptor spatial FWHM	5.7°
Temporal filter time constant τ	0.150 s
Spatial sampling interval Δx	0.5°
Temporal sampling interval Δt	1/240 s
Leak current reversal potential E_{leak}	0 mV
Excitatory current reversal potential E_{exc}	+60 mV
Inhibitory current reversal potential E_{inh}	-30 mV
Excitatory to leak conductance ratio $g_{\text{exc}}/g_{\text{leak}}$	0.1
Inhibitory to leak conductance ratio g_{inh}/g_{leak}	0.3

211 **Table 1:** Parameter values used in all simulations.

212

213 In vivo two-photon calcium imaging in T4 cells

214 Most of our comparisons relate the synaptic model's responses to published data, but we also

215 compare the model to a new dataset of T4 cell responses to glider stimuli. The protocol for two-

216 photon calcium imaging in T4 cells matches published methods (Badwan et al., 2019) and used

217 Psychtoolbox (Brainard, 1997; Kleiner et al., 2007; Pelli, 1997) to present stimuli on a

218 panoramic visual display (Creamer et al., 2019). The glider stimuli presented during these

219 measurements are described in **Appendix A**. Net responses were computed as the difference in

220 responses to stimuli moving in the preferred and null directions of each T4 region of interest, and

then averaged within each fly. Non-parametric two-sided Wilcoxon signed-rank tests were used

to test whether median net responses differed significantly from zero (Hollander et al., 2013).

223 For statistical purposes, each individual fly was considered to be an independent sample.

224

225 Numerical methods

226 Numerical simulations were conducted using Matlab 9.6 (R2019a) (The MathWorks, Natick,

227 MA, USA). For stimuli containing randomly-generated components, responses were averaged

- 228 over 1000 realizations, and bootstrapped 95% confidence intervals for the mean were computed
- using the bias-corrected and accelerated percentile method (Efron, 1987). Results
- 230

231 The synaptic model reduces to HRC-like terms

232 To gain intuition about the operation of the T4 synaptic model, we consider its expansion in the

- small-input limit. To do so, we approximate the ramp function nonlinearity with a smooth
- function that represents a soft rectifier, which can be approximated by a linear function for small

inputs (Fitzgerald and Clark, 2015). In particular, letting $s_1(t) \coloneqq s(t, x - \Delta), s_2(t) \coloneqq s(t, x)$,

236 and $s_3(t) \coloneqq s(t, x + \Delta)$, and defining the non-negative constants $\alpha \coloneqq |g_{\text{inh}}E_{\text{inh}}/g_{\text{leak}}|$ and $\gamma \coloneqq$

237 $|g_{\text{exc}}E_{\text{exc}}/g_{\text{leak}}|$, we have, to lowest order in the inputs,

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239
$$C \approx \frac{1}{16} (\alpha s_1 + \gamma \dot{s}_2 - \alpha s_3)^2$$

240

241 which may be rewritten as

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243
$$C \approx \frac{\alpha^2}{16} (s_1 - s_3)^2 + \frac{\gamma^2}{16} (\dot{s}_2)^2 + \frac{\alpha\gamma}{8} (s_1 \dot{s}_2 - \dot{s}_2 s_3)$$

244

245 This expansion represents a motion-energy approximation of the model. The first term in this 246 expansion is a finite-difference approximation to a spatial derivative, while the second term is a 247 temporal derivative at the center of the model's receptive field. The third term, which is the only 248 direction-selective term, is the subtraction of two offset correlators with opposite directional 249 tuning. This subtraction step provides some intuition for why this model mimics some properties 250 of a fully-opponent HRC model (Badwan et al., 2019). This same direction-selective term also 251 appears in the second-order expansion of the membrane voltage. Because this expansion of the 252 model is only to second-order, it is invariant under contrast inversions, and cannot account for 253 properties like ON-edge selectivity (Clark et al., 2014; Fitzgerald and Clark, 2015; Fitzgerald et 254 al., 2011). Though this simple description does not capture all properties of the synaptic model, it 255 provides intuition for the sensitivity of the model to certain stimulus features. 256

257 The synaptic model is strongly ON/OFF-edge- and direction-selective

T4 and T5 neurons are distinguished by the fact that T4 cells respond to ON-edges while T5 cells respond to OFF-edges (Maisak et al., 2013). We first compared the ON/OFF edge- and direction-selectivity of our T4 synaptic model to responses measured using two-photon calcium imaging in T4 cells sensitive to front-to-back (FTB) motion. Like T4 FTB cells, our synaptic model responded strongly to an ON edge moving in the FTB direction, but displayed little or no response to OFF edges moving in the FTB direction or to edges of either polarity moving in the back-to-front (BTF) direction (**Figure 1C**) (Maisak et al., 2013; Salazar-Gatzimas et al., 2016).

266 The spatiotemporal tuning of the synaptic model is consistent with that of T4 cells

267 Sinusoid grating stimuli are a common tool for characterizing direction-selective computations.

Responses to these stimuli have been used to suggest that the membrane voltage of T5 cells is a

269 nearly linear transformation of the visual input (**Figure 2A**) (Wienecke et al., 2018). In the

270 synaptic model, the membrane voltage is a nonlinear function of the input contrast because the

271 inputs are first rectified and then interact nonlinearly. We applied the same linearity testing

272 protocol to our model membrane voltage, constructing predictions for responses to PD and ND

272 protocor to our moder memorane vorage, constructing predictions for responses to 1 D and 11D

drifting gratings from the responses to counterphase gratings (see Appendix A) (Jagadeesh et al.,
1993; Wienecke et al., 2018). The responses of the T4 synaptic model to drifting gratings were

similar to those predicted by a linear model for membrane voltage (**Figure 2A**). Thus, even a

276 nonlinear system like the T4 synaptic model may appear reasonably linear by this protocol.

277

268

278 T4 and T5 cells display direction-opponent average calcium responses to sinusoid gratings 279 (Figure 2B) (Badwan et al., 2019). This property means that the average response to PD motion 280 is reduced by the addition of ND motion, imposing a strong constraint on models for the 281 direction-selective computation. In particular, it implies that linear-nonlinear models with 282 expansive nonlinearities cannot account for the response properties of these cells (Badwan et al., 283 2019). A variant of this synaptic model was proposed to account for these direction-opponent 284 responses (Badwan et al., 2019). This model reproduces the strong suppression when ND motion is added to PD motion without substantial enhancement when orthogonal-direction (OD) motion 285 286 is added to PD motion (Figure 2B).

288 T4 and T5 cells are tuned to the temporal frequency of sinusoidal stimuli (Figure 2C) (Creamer 289 et al., 2018). This means that the mean neural response is maximal at a single temporal 290 frequency, independent of the wavelength. This property also applies to measurements of fly 291 behavior (Creamer et al., 2018; Kunze, 1961) and is consistent with the classical, fully-opponent 292 HRC. We presented the T4 synaptic model with drifting gratings of different spatial and 293 temporal frequencies to find the mean response to each. The model response was strongly 294 temporal-frequency-tuned (Figure 2C). To quantify the temporal-frequency-tuning, we asked 295 how much of the variance in this surface was accounted for by the product of one function of 296 temporal frequency and one fuction of spatial frequency response (Creamer et al., 2018; Priebe et 297 al., 2006). Such a separable model accounted for 99% of the variance in the response (see 298 Appendix A, Figure 2C). Because of our choice of parameters, the input temporal filters in this 299 model produce peak responses at around 1 Hz, lower than the roughly 2-4 Hz peak measured in 300 these T4 cells.

301

302 T4 and T5 cells respond to static gratings with amplitudes that depend on the grating orientation (Fisher et al., 2015) (Figure 2D). The preferred orientation (defined by the vector normal to the 303 304 edges in a static grating) approximately matches the preferred direction of motion of these cells 305 (Maisak et al., 2013). The convention we use here for defining the orientation of a static grating 306 is rotated 90° relative to that used in the original study, which defined orientation in terms of 307 vectors parallel, rather than normal, to the edges (Fisher et al., 2015) (see Appendix A). When 308 the T4 synaptic model was presented with both static and drifting gratings of many different 309 orientations, it reproduced the orientation tuning observed experimentally for both static and 310 moving gratings (Figure 2D). The model was more selective for both orientation and direction 311 than the T4 cell measurements.

312

313 The synaptic model reproduces the selectivity of apparent motion responses in T4 cells

314 In addition to sinusoid gratings, apparent motion stimuli are a useful tool for investigating

315 direction-selective systems. These stimuli decompose visual motion into summations of simpler

316 spatiotemporal patterns, which can provide strong intuition into the motion computation (Barlow

317 and Levick, 1965).

319 Electrophysiological measurements of T4 cells have shown fast depolarization and delayed, 320 offset hyperpolarization in response to a small flashed white bar placed on a gray background 321 (Gruntman et al., 2018) (Figure 3A). The synaptic T4 model displayed qualitatively consistent 322 responses to the same stimulus (Figure 3A). The positive lobe in the model is narrower than in 323 the electrophysiological recording; this is likely because the true central input to T4 has a wider 324 receptive field than in our model (Behnia et al., 2014; Takemura et al., 2013). Consistent with 325 electrophysiology, the OFF input to the T4 model is not visible under this analysis because it was 326 rectified with a threshold at mean gray (zero contrast).

327

328 Since the synaptic model reproduced T4 cell voltage responses to flashed bars, we sought to 329 characterize its responses to apparent motion stimuli composed of pairs of bars offset in 330 spacetime (Salazar-Gatzimas et al., 2018). These stimuli can induce in humans and in flies the 331 "reverse-phi" motion illusion, in which a reversal of contrast polarity induces a motion percept in 332 the direction opposite the stimulus displacement (Anstis, 1970; Clark et al., 2011; Hassenstein 333 and Reichardt, 1956). We aligned these stimuli so that the temporally-delayed bar is placed at the 334 center of the receptive field (Figure 3B) (Salazar-Gatzimas et al., 2018). T4 cells respond 335 maximally to one phi and one reverse-phi apparent motion stimulus out of eight possible pairings 336 (Salazar-Gatzimas et al., 2018). The synaptic model reproduced this selectivity (Figure 3C-D). 337

Various groups have assessed nonlinear enhancement or suppression of PD and ND apparent
motion stimuli relative to linear decompositions. This analysis can be misleading because it does
not allow one to uniquely characterize the nonlinearity as 'enhancing' or 'suppressing', since
there exist an infinite number of linear decompositions of a given stimulus (Salazar-Gatzimas et
al., 2018). Despite this difficulty, such analyses have been applied as an intuitive way to try to
understand direction-selective computations (Barlow and Levick, 1965; Fisher et al., 2015;
Gruntman et al., 2018; Haag et al., 2016).

345

346 In T4 cells, an analysis of responses to sequential bars has indicated that calcium signals include

347 both PD enhancement and ND suppression relative to a linear prediction from the responses to

individual bars (Haag et al., 2016) (Figure 3E). Our model failed to reproduce this result,

349 showing only suppression of ND motion under this analysis (Figure 3E). This discrepancy could

350 be influenced by the timescale of this stimulus, which is far longer than the 150 ms offset used in

351 the apparent motion stimuli in (Figure 3C-D). Additionally, previous theoretical work has

352 shown that disinhibition can generate PD enhancement in similar models (Borst, 2018; Torre and

- 353 Poggio, 1978); the choice of thresholds in this model did not permit flanking disinhibition with
- 354 ON stimuli.
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356 The synaptic model does not reproduce the fast timescale tuning of T4 cells

357 A third approach to characterizing direction-selective signals has been to apply stochastic stimuli 358 with specified correlation structure. Responses to uncorrelated stimuli can be used to generate an 359 unbiased estimate of a system's linear receptive field (Chichilnisky, 2001). By using reverse-360 correlation and uncorrelated stimuli to extract spatiotemporal receptive fields, T4 cells have been 361 characterized by oriented linear receptive fields with a central excitatory lobe and a delayed, 362 offset inhibitory lobe (Figure 4A) (Leong et al., 2016; Salazar-Gatzimas et al., 2016). The T4 363 synaptic model generates the same shape of receptive field (Figure 4A). However, in the model, 364 the inhibitory lobe lasts longer than that measured in T4 cells, and the tuning of the model was 365 slower overall.

366

367 Responses to stochastic stimuli containing precise pairwise spatiotemporal correlations have 368 revealed fast-timescale tuning in T4 cells (Salazar-Gatzimas et al., 2016). In measurements of T4 369 and T5, the cells could discriminate between spatiotemporal correlations with delays of 0 and 15 370 ms (Figure 4B). We presented the synaptic model with stimuli containing pairwise 371 spatiotemporal correlations at different temporal delays. The model was direction-selective and 372 responded to both positive and negative correlations, as in the cellular measurements. However, 373 the model did not reproduce the fast timescale discrimination between delays (Figure 4B). 374 Furthermore, the synaptic model showed strong suppression of ND-oriented positive correlations 375 and enhancement of ND-oriented negative correlations, which was not observed in the data. 376

377 Behaving Drosophila respond direction-selectively to correlations higher than second-order

378 (Clark et al., 2014; Leonhardt et al., 2016). This cannot be explained by models that compute

379 pairwise correlations in the stimulus, such as the HRC and motion energy model. The sensitivity

380 to higher-order correlations has been assessed using three-point glider stimuli, which contain 381 precise third-order correlations (Hu and Victor, 2010) (Figure 4C). The net responses of T4 cells 382 to these stimuli have previously been inferred from behavioral measurements in *Drosophila* with 383 the synaptic outputs of T5 cells silenced, using gliders updated at 24 Hz (Leonhardt et al., 2016). 384 We used *in vivo* two-photon calcium imaging to measure directly the responses of T4 cells to 385 three-point gliders updated at 5 Hz, and found that the signs of the net responses were consistent 386 with those measured in behavior with T5 cells silenced (Figure 4C, see Methods and Appendix 387 A for details).

388

With an update rate of 24 Hz, the synaptic model correctly predicted the signs of net responses to diverging gliders measured in imaging and behavior, but predicted the wrong converging glider responses (**Figure 4C**). At 5 Hz, the synaptic model correctly predicted the signs of both converging and diverging glider responses, but not the relative magnitudes. Thus, the glider responses in T4 appear relatively insensitive to the glider timescale (24 vs. 5 Hz), but the model's response depends strongly on the input timescales.

395

396 T4 and T5 cells have been shown to display strongly direction-selective responses to rigidly-397 translating stimuli consisting of black and white squares placed at random on a gray background 398 (see Appendix A) (Badwan et al., 2019). When two such stimuli that move in opposite 399 directions are superimposed, they generate transparent motion percepts in primates (Qian et al., 400 1994), and they generate responses in T4 that are reduced compared to presenting PD stimuli 401 alone (Badwan et al., 2019). The synaptic model qualitatively reproduced these responses 402 (Figure 4D). In particular, the responses of T4 cells are suppressed more strongly under the 403 addition of ND motion than under the addition of OD motion, a feature that is reproduced by the 404 synaptic T4 model (Figure 4D). Therefore, as in T4 cells, the selective direction-opponency 405 observed in the model persists even with stimuli containing multiple spatiotemporal frequencies. 406

407 The T4 synaptic model provides decorrelated channels for naturalistic motion

408 Beyond artificial stimuli, natural scenes have been used to investigate the performance of

409 direction-selective signals generated by models, behavior, and neurons (Badwan et al., 2019;

- 410 Chen et al., 2019; Dror et al., 2001; Fitzgerald and Clark, 2015; Leonhardt et al., 2016; Salazar-
- 411 Gatzimas et al., 2018; Straw et al., 2008). We therefore sought to investigate the performance of

412 the T4 synaptic model in natural motion processing. To do so, we presented it with rigidly-413 translating scenes from a database of natural images (see Appendix A, Figure 5A) (Meyer et al., 414 2014) (Badwan et al., 2019; Chen et al., 2019; Fitzgerald and Clark, 2015; Meyer et al., 2014; 415 Salazar-Gatzimas et al., 2018). Though the structure and properties of inputs to T5 cells are 416 known to differ from the inputs to T4 (Serbe et al., 2016; Shinomiya et al., 2019), to make an 417 OFF-edge selective channel we created a 'T5' model by simply inverting the ON/OFF selectivity 418 of the inputs to our T4 synaptic model. This is intended merely to be an OFF-selective channel 419 for the purposes of comparing T4 and potential T5 cell responses. The resulting four channels 420 displayed strongly direction-selective average responses to translating natural scenes (Figure 421 **5B**). 422 423 Measured responses of T4 and T5 cells to translating natural scenes are decorrelated, so that only

424 one channel is active at once (Salazar-Gatzimas et al., 2018). The synaptic models of T4 and

425 'T5' also generated highly decorrelated responses, with the coactivation matrix of the four

426 channels being nearly diagonal (Figure 5C). Such decorrelated parallel channels may provide a

427 convenient representation of motion signals (Salazar-Gatzimas et al., 2018).

428

429 **Discussion**

430

431 An anatomically constrained synaptic model suffices to reproduce many, but not all, of the 432 properties of *Drosophila* T4 cells. This model reproduces the direction-opponency, temporal-433 frequency-tuning, orientation-tuning, and phi/reverse-phi selectivity measured in T4 cells 434 (Figures 2-4). When applied to a naturalistic velocity estimation task, it produces decorrelated 435 signals similar to those measured in T4 and T5 neurons (Figure 5). However, it fails to 436 reproduce the PD enhancement and fast-timescale tuning observed in T4 cells (Figures 3-4). 437 Moreover, though it is sensitive to triplet correlations in its input, it fails to reproduce them on 438 the same timescales as observed in the data (Figure 4). In short, this simple synaptic model is 439 sufficient to reproduce several distinct properties of T4 cells, but cannot account for several 440 observations.

442 Minimal models and levels of understanding

Here, we asked whether a minimal synaptic model could qualitatively reproduce features of T4 cell responses. The minimal model required no exotic neurotransmitter receptors or interactions, and was based on simple synaptic conductances. The simplifications sufficient to explain different phenomena will depend strongly on the features one seeks to reproduce, and on the desired level of fidelity. However, minimal models are useful precisely because they can be relatively straightforward to analyze.

449

450 Marr famously proposed different levels of understanding neural circuitry, including an

451 algorithmic level and a mechanistic level (Marr and Poggio, 1976). As we drive towards a deeper

452 understanding of the visual motion circuit in the fly, the levels of algorithm and mechanism can

453 appear increasingly blurred. It is hard to define what distinguishes the mechanistic circuit

description presented here from a detailed algorithm-level description of the computation.

455 However, it remains important to connect proposed mechanistic models to high-level

456 descriptions of the system such as the HRC. This is because the high-level descriptions of

457 computations provide a level of intuition for the behavior of the system that a more intricate

458 model cannot. Moreover, the HRC explains a wide variety of neural and behavioral data in flies

459 (Borst and Egelhaaf, 1989; Yang and Clandinin, 2018), so an HRC-like algorithm must be a

460 limiting case of any proposed mechanistic model (Potters and Bialek, 1994).

461

462 Sufficiency of models

463 Many details of the function of the early visual system were neglected in this model. For 464 instance, the filter shapes in neurons leading into the model T4 cell have been well-characterized 465 (Arenz et al., 2017; Behnia et al., 2014), but this model used simple exponential filters. Lateral 466 inhibition is widely documented in the early fly visual system (Arenz et al., 2017; Freifeld et al., 467 2013; Meier et al., 2014), but this model used simple Gaussian spatial acceptance functions 468 without lateral inhibition. The synapses that feed into the medulla neurons that synapse onto T4 469 are likely to have complex, nonlinear processing properties (Yang et al., 2016), yet we modelled 470 the entire input pathway as a purely linear filter. The rectifications of neural responses upstream 471 of T4 are imperfect (Behnia et al., 2014; Salazar-Gatzimas et al., 2018), but this model used 472 simple threshold-linear rectifiers. The fly eye possesses neurons that feedback onto earlier stages

473 and create reciprocal interactions between neurons (Takemura et al., 2013; Takemura et al.,

- 474 2017), but this model is entirely feedforward.
- 475

476 Despite these approximations, the synaptic T4 model presented here is sufficient to qualitatively

477 match a variety of T4 neuron responses. Adding some of these neglected details into a model

478 may make it sufficient to reproduce other features of T4 responses. This provides a method for

- 479 understanding which details of processing are related to which response features in T4 cells: one
- 480 may ask how different details of the system affect the sufficiency of a model to reproduce
- 481 specific downstream response properties. As the field acquires more and more detailed

482 information about the motion detection circuitry, this sort of analysis will be critical to

483 understand the functional role of different properties.

484

One might naturally ask whether the synaptic model presented here might be further simplified without sacrificing its ability to account for the response properties of T4 cells. As described in **Appendix C**, a simplified linear-nonlinear cascade (LNLN) representing the numerator of the biophysical nonlinearity can generate some, but not all, of the properties of the full model.

490 Flexibility in extending this minimal synaptic model

491 In selecting parameter values for this synaptic model, we sought to reproduce only a few 492 properties of T4 cells: a temporal frequency maximum of 1 Hz and a direction-opponent average 493 responses to sinusoid gratings with a temporal frequency of 1 Hz and a spatial wavelength of 45° 494 (Appendix B) (Badwan et al., 2019). To capture a larger subset of the measured properties of T4 495 cells, one could optimize the parameters of the model capture many response properties (Deb, 496 2014). Such a solution would provide information about the maximal ability of this synaptic 497 model to reproduce the properties of T4 cells, but it seems unlikely to provide insight into the 498 predictive power of the core features of the model.

499

500 The organization of this model allows for several clear tuning mechanisms. First, the temporal

501 filters could be modified to better match measured filters (Figure 2). Second, the degree to

502 which inhibition is shunting or hyperpolarizing can be adjusted by changing the reversal

503 potential of inhibitory currents. This could effectively hide inhibition under some stimuli and

504 measurements. Third, it is clear that to better represent preferred direction enhancement, the 505 threshold for the OFF-inhibitory input could be changed (**Figure 3**) (Borst, 2018). This would 506 allow disinhibition of Mi4 to change the gain for the central input.

507

508 In the model analyzed here, we chose all thresholds of the input LN models to be zero. This

509 effectively ignores contrast asymmetries in the natural world (Geisler, 2008), which have been

510 used to understand many functional properties of motion detectors in flies (Chen et al., 2019;

511 Clark et al., 2014; Fitzgerald and Clark, 2015; Fitzgerald et al., 2011; Leonhardt et al., 2016;

512 Salazar-Gatzimas et al., 2018). Changing these thresholds to optimize for natural scene motion

513 estimation might also generate a parameter set that better captures responses to triplet

514 correlations (Figure 4) (Fitzgerald and Clark, 2015). In short, the synaptic model presented here

515 is highly flexible and extensible, and uses only simple, known biophysical mechanisms.

516

For the sake of simplicity, we have used single delay and non-delay lines in this work (**Figure 1A**). However, T4 cells receive fast excitatory input at the center of their receptive fields from both Mi1 and Tm3 cells, and delayed OFF inhibitory input offset in the PD from both Mi4 and CT1 cells (Shinomiya et al., 2019; Takemura et al., 2017). Dissecting how information from these parallel channels is used, particularly if it is nonlinearly combined, will be important in developing a full understanding of the direction-selective computation performed by T4 cells.

523

524 *Modelling temporal processing*

525 The model presented here failed to capture some of the fast-timescale tuning measured in T4, 526 including in its responses to pairwise and triplet spatiotemporal correlations (Figure 4). In this 527 minimal model, we represented all temporal processing by linear filters. However, the temporal 528 processing upstream of T4 cells involves nonlinear and adaptive mechanisms, which can affect 529 temporal response properties (Howard et al., 1984; Zheng et al., 2006). Thus far, the study of 530 nonlinear mechanisms in the fly visual system has focused on static nonlinear effects such as 531 rectification (Behnia et al., 2014; Yang et al., 2016) and on nonlinear interactions between 532 linearly filtered signals (Borst et al., 2005; Fitzgerald and Clark, 2015). The inclusion of 533 nonlinear effects on the dynamics themselves may be necessary to accurately capture the temporal processing upstream of T4 cells. As a first step towards experimentally understanding 534

535 adaptation in this circuit, one might characterize the temporal kernels of the inputs to T4 cells

536 with high resolution (Mano et al., 2019; Yang et al., 2016) and study how their properties depend

537 on stimulus statistics and history (Baccus and Meister, 2002; Kim and Rieke, 2001; Rieke,

538 2001). Only a few models have focused on these sorts of changes in processing dynamics (Clark

t al., 2013). Though the analysis of dynamic temporal nonlinearities is complex, incorporating

- 540 them into models may provide insight into how fast timescale tuning of T4 cells arises.
- 541

542 A T5 synaptic model

543 In this work, we used a sign-inverted version of our T4 synaptic model to represent the OFF-

544 edge-selective T5 cells. This representation would correspond to a first-order direction-selective

545 cell that receives OFF excitatory input at the center of its receptive field, delayed OFF inhibitory

546 input offset in its preferred direction, and delayed ON input offset in the null direction. Such a

547 model would correctly predict the selective responses of T5 cells to phi and reverse-phi apparent

548 motion stimuli (Salazar-Gatzimas et al., 2018). However, the functional and anatomical structure

549 of the inputs to T5 cells suggests that it receives only OFF inputs (Serbe et al., 2016; Shinomiya

et al., 2019). Somehow, however, signals in T5 cells are sensitive to both contrast increments and

decrements (Salazar-Gatzimas et al., 2018; Wienecke et al., 2018). Further study of the physical

and functional connectome of the OFF-edge motion pathway, will be required to elucidate how

- 553 the direction-selective computations in T4 and T5 cells differ.
- 554

555 Relationships to mammalian visual systems

556 The organization of the fly's visual motion detection circuits bear striking similarities to those in

557 mammalian retina in their anatomy, circuitry, and algorithmic processing (Borst and

Helmstaedter, 2015; Clark and Demb, 2016; Sanes and Zipursky, 2010). In mammalian retina,

the earliest direction-selective signals are generated by starburst amacrine cells (SACs), which

are also tuned to ON- and OFF-edges (Euler et al., 2002; Famiglietti Jr, 1983). It appears that

561 SACs may receive inputs that are differentially delayed (Fransen and Borghuis, 2017; Kim et al.,

562 2014), similar to the inputs to T4 cells. It would be interesting to investigate how much SAC

563 phenomenology that mechanism alone could account for, when linked to simple biophysical

564 mechanisms. As in this study, it could provide insight into where the circuit understanding is

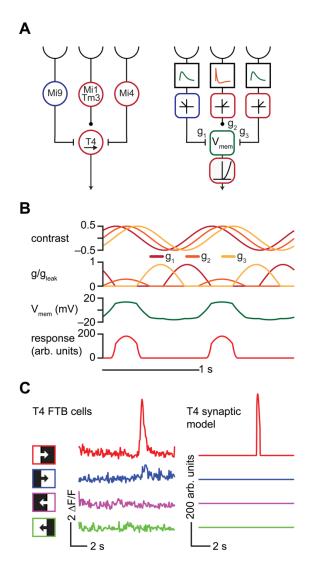
565 lacking, especially when complex stimuli are used to probe SAC function (Chen et al., 2016).

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It is notable that the ON-ON-OFF spatial organization of T4 inputs (Fig. 1A) is almost identical
to a model proposed to explain cortical responses to pairwise correlations (Mo and Koch, 2003).
This suggests there may be deep parallels between T4 and T5 and cortical motion processing
steps. Models for fly and cortical direction-selectivity have traditionally differed in whether they
assume discrete inputs (fly, HRC-like models) or more continuous inputs (cortex, motion-
energy-like spatiotemporal filtering). If synaptic interactions are considered, then continuous
linear filters cannot be applied, and models must incorporate the discrete receptive fields of the
inputs to a cell. It would be interesting to ask how such conductance models fare in predicting
cortical responses; the statistical nature of cortical connections make it more difficult to make a
general model of this type.
In this synaptic model of T4 cell function, we have paired known connectivity with measured
physiology and simple biophysics to predict many circuit processing properties. This allows us to
define where such a model succeeds and where it fails. This represents progress towards the
ultimate goal of understanding this circuit at all levels, from utility to algorithm to mechanism.
Author contributions
JAZV and DAC conceived of numerical experiments. JAZV performed numerical simulations
and analyzed the model. BAB acquired calcium imaging data. JAZV and DAC wrote the paper.
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592 Figure Legends



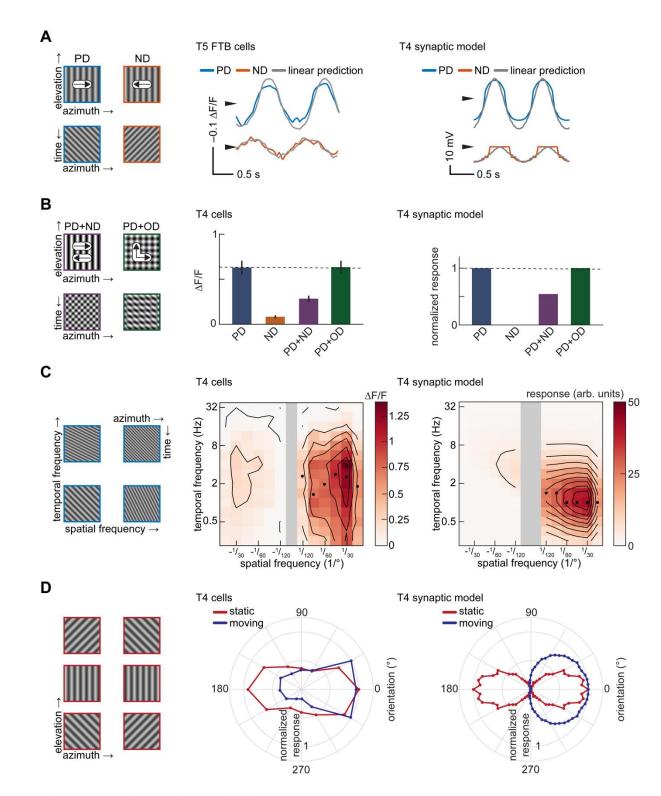


594 Figure 1, Zavatone-Veth et al. (2019)

595 Figure 1: An anatomically constrained synaptic model for T4 cells.

A. *Left:* Diagram of proposed inputs to *Drosophila* T4 first-order direction-selective cells
based on anatomical and physiological measurements. Mi1 and Tm3 cells provide ON
excitatory input at the center of the receptive field of each T4 cell, while Mi9 provides
delayed OFF inhibitory input offset in the null direction, and Mi4 provides delayed ON
inhibitory input offset in the preferred direction. *Right:* Synaptic model based on the
anatomical structure shown at left.

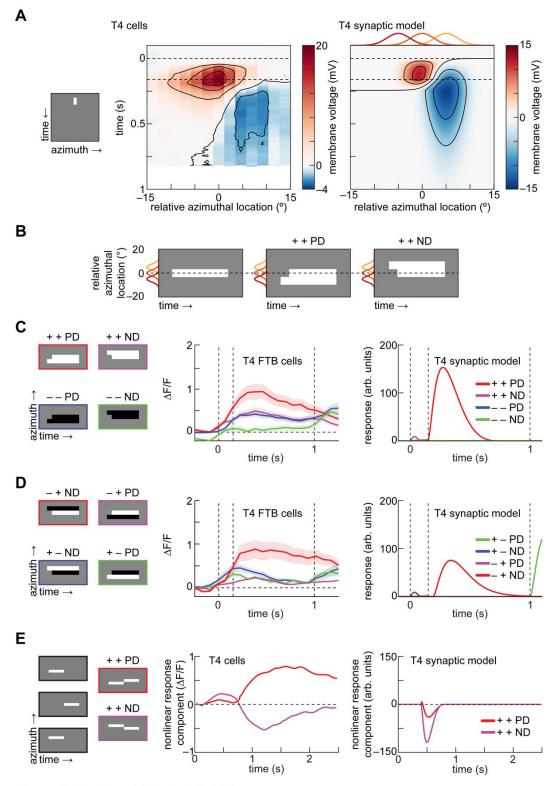
602 B. Responses of each component of the synaptic model to a 1 Hz, 45° sinusoidal grating 603 drifting in the preferred (rightward) direction. Top: Input contrasts to each of the three 604 presynaptic units of the model. Upper middle: Conductances of excitatory and inhibitory 605 currents corresponding to each input in response to the sinusoidal stimulus. Lower 606 middle: Membrane voltage. Bottom: Calcium signal. 607 C. Left: Responses of T4 cells sensitive to front-to-back (FTB) motion to ON and OFF 608 edges moving FTB and back-to-front (BTF) at 30%, measured using two-photon calcium 609 imaging (data from (Salazar-Gatzimas et al., 2016)). Right: As at left, but for the T4 610 synaptic model. 611



612 **Figure 2**, Zavatone-Veth et al. (2019)

Figure 2: The T4 synaptic model reproduces sinusoidal grating responses measured in T4 cells.

615	A. Left: Images and kymographs of sinusoid gratings drifting in the prefer	red (PD) and null
616	(ND) directions. Center: Membrane voltage of T5 FTB cells to 1 Hz, 22	5° drifting gratings
617	compared with linear predictions from contrast-modulated counterphase	e gratings,
618	measured using voltage indicators (data from (Wienecke et al., 2018)).	<i>Right:</i> As at
619	center, but for voltage responses of the T4 synaptic model. The model h	as coefficients of
620	determination for PD and ND of 0.92 and 0.82.	
621	B. Mean responses to 1 Hz, 45° sinusoid gratings. Left: Images and kymog	raphs of
622	composite sinusoid gratings containing PD and ND motion or PD and o	orthogonal
623	direction (OD) motion. Center: Mean responses of T4 cells to drifting g	gratings, measured
624	using a calcium indicator (data from (Badwan et al., 2019)). Error bars	indicate ±1 SEM.
625	Right: As at center, but for calcium responses of the T4 synaptic model	
626	C. Spatiotemporal frequency tuning. Left: Kymographs of sinusoid grating	s with varying
627	spatiotemporal frequency content. Center: Spatiotemporal frequency tu	ning of T4 cells
628	(data from (Creamer et al., 2018)). Black circles indicate the temporal f	requency at which
629	the maximum response at a given spatial frequency is attained. Right: A	s at center, but
630	for the T4 synaptic model.	
631	D. Orientation and direction tuning. Left: Images of oriented sinusoid grati	ngs. Center:
632	Orientation tuning of T4 and T5 cells with static gratings (data from (Fi	sher et al., 2015))
633	and direction tuning of T4 cells with drifting gratings (data from (Maisa	ak et al., 2013)).
634	The orientation of a static grating is defined by the vector normal to the	apparent edges,
635	the same definition as for moving gratings (see Appendix A). Right: A	s at center, but for
636	the T4 synaptic model.	



638 Figure 3, Zavatone-Veth et al. (2019)

Figure 3: The T4 synaptic model reproduces the spatial organization and selectivity of apparent motion responses in T4 cells.

- A. Responses to a single white bar flashed at different spatial locations. *Left:* Kymograph of
 2° white bar presented for 160 ms. *Center:* Membrane voltage of T4 cells to flashed white
 bars, measured using electrophysiology (data from (Gruntman et al., 2018)). *Right:* As at
 center, but for the T4 synaptic model. Red, orange, and yellow lines indicate the spatial
 acceptance functions of the three model inputs.
- B. Apparent motion stimuli are aligned such that the lagging bar is located at the center of
 the receptive field (Salazar-Gatzimas et al., 2018). The leading bar is presented at time
 zero and lasts for 1 second, and the lagging bar is presented 150 ms later. Each bar
 subtends 5° of visual angle. Red, orange, and yellow lines indicate the spatial acceptance
 functions of the three inputs.
- C. Responses to phi apparent motion stimuli, aligned as in (B). *Left:* Kymographs of all four
 possible phi apparent motion stimuli. *Center:* Responses of T4 FTB cells to all four phi
 apparent motion stimuli, measured using two-photon calcium imaging (data from
 (Salazar-Gatzimas et al., 2018)). Error patches indicate ±1 SEM. *Right:* As at center, but
 for the T4 synaptic model.

D. As in (C), but for reverse-phi apparent motion stimuli, in which the sequentially presented bars have opposite contrasts.

E. Assessing PD enhancement and ND suppression. *Left:* Kymographs of linear
decomposition of flashed apparent motion stimuli, with 4.5°-wide white bars presented
sequentially for 400 ms each. *Center:* Nonlinear response component, defined as the
residual of the linear prediction, measured using a calcium indicator (data from (Haag et
al., 2016)). *Right:* As at center, but for the T4 synaptic model.

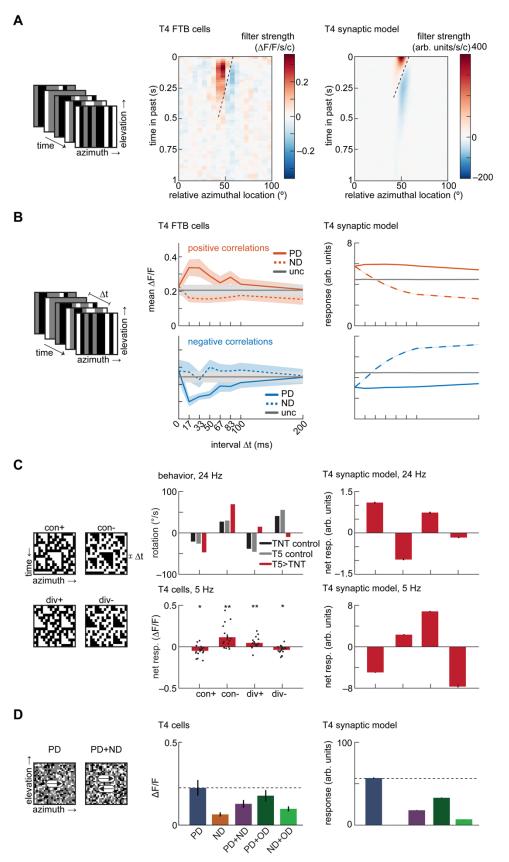
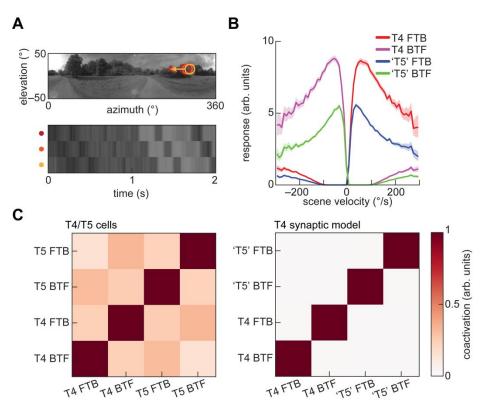


Figure 4, Zavatone-Veth et al. (2019)

Figure 4: The minimal T4 synaptic model is not sufficient to reproduce the fast-timescale tuning of T4 cells.

667 A. Linear receptive field measurements. Left: Schematic depiction of binary, uncorrelated spatiotemporal noise. Center: Linear receptive field of T4 FTB cells (data from (Salazar-668 669 Gatzimas et al., 2016)). Right: As at center, but for the T4 synaptic model. 670 B. Correlation interval receptive field measurements. *Left:* Schematic depiction of ternary 671 noise containing pairwise correlations at a specified interval Δt . Center: Responses of T4 672 FTB cells to positive and negative correlations, measured using a calcium indicator (data 673 from (Salazar-Gatzimas et al., 2016)). Error bars indicate ±1 SEM. *Right:* As at center, 674 but for the T4 synaptic model. Error patches, which are barely visible, indicate 95% 675 confidence intervals of the mean, which is variable due to the stochastic stimulus. 676 C. Triplet correlation sensitivity. Left: Kymographs of three-point glider stimuli containing 677 positive and negative triplet correlations. *Center top:* Turning behavioral responses to 678 three-point gliders updated at 24 Hz of flies with the synaptic outputs of T5 cells silenced 679 (data from (Leonhardt et al., 2016)). Positive rotations correspond to the direction of the 680 displacement of the spatial mean location of each triplet. Center bottom: Net responses of 681 T4 cells to three-point gliders updated at 5 Hz, measured using a calcium indicator (see 682 **Methods**). Asterisks indicate that median net response differs from zero at the p < 0.05683 (*) or p < 0.01 (**) level by a Wilcoxon signed-rank test with N = 16 flies. Exact p-684 values are p = 0.0174, 0.0061, 0.0097, and 0.0131 for con+, con-, div+, and div-, 685 respectively. Error bars indicate ± 1 SEM over flies, and black circles indicate individual 686 per-fly means. *Right:* As at center, but for the T4 synaptic model. Error bars indicate 95% 687 confidence intervals of the mean. 688 D. Responses to rigidly translating stimuli with stochastic checkerboard patterns. Left: 689 Images of random checkerboard stimuli. Center: Mean responses of T4 cells to 690 checkerboard stimuli translating at 100%, measured using a calcium indicator (data from 691 (Badwan et al., 2019)). Error bars indicate ± 1 SEM. *Right:* As at center, but for the T4 692 synaptic model. Error bars indicate 95% confidence intervals of the mean.



⁶⁹⁴ Figure 5, Zavatone-Veth et al. (2019)

Figure 5: The T4 synaptic model generates decorrelated signals in response to naturalistic motion.

- A. A rigidly rotating panoramic natural scene, with three spatially offset input signals as afunction of time.
- B. Average responses of the T4 synaptic model and a 'T5' variant to naturalistic motion
 constructed by rigidly translating natural scenes at a variety of velocities. Error patches
 indicate 95% confidence intervals of the mean. 'T5' cells were constructed by signinverting the inputs to the minimal T4 synaptic model.
- C. Decorrelation of channels with naturalistic motion. *Left:* Coactivation matrix of T4 and
 T5 cells in response to rigidly translating natural scenes (data from (Salazar-Gatzimas et al., 2018)). *Right:* As at left, but for the T4 synaptic model and 'T5' variant. Coactivation
 was computed for an ensemble of (image, velocity) pairs, in which the velocity was
 chosen from a Gaussian distribution with zero mean and 100°/s standard deviation.
- 708

709	Appendix A: Visual stimuli used in simulations and imaging experiments
710	
711	In this appendix, we describe in detail all stimuli used in this work.
712	
713	ON and OFF edges (Figure 1)
714	ON and OFF edges were constructed by placing white (respectively black) edges on a gray
715	background. All edges translated at 30°/s.
716	
717	Sinusoid grating stimuli (Figures 1 and 2)
718	Sinusoid grating stimuli were constructed as in (Badwan et al., 2019; Creamer et al., 2018;
719	Maisak et al., 2013). Briefly, rightward- and leftward-drifting gratings were constructed as
720	
721	$c(t,x) = c_0 \sin(\omega t \mp \kappa x)$
722	
723	where c_0 is the input contrast, ω is the temporal frequency in units of radians per second, κ is the
724	spatial frequency in units of radians per degree, and the negative sign is taken for rightward-
725	drifting gratings. To assess whether our model is temporal-frequency-tuned, we computed the
726	fraction of the total variance in a spatiotemporal frequency sweep of its responses accounted for
727	by a separable approximation resulting from its singular value decomposition (Creamer et al.,
728	2018). Counterphase gratings were constructed as
729	
730	$c(t,x) = c_0 \sin(\omega t + \kappa x + \phi_1) + c_0 \sin(\omega t - \kappa x + \phi_2)$
731	
732	where ϕ_1 and ϕ_2 are uniformly sampled phase offsets, over which we average in all analyses.
733	Gratings containing preferred- and orthogonal-direction motion were constructed as
734	
735	$c(t, x) = c_0 \sin(\omega t + \kappa x + \phi_1) + c_0 \sin(\omega t + \phi_2)$
736	
737	The linearity analysis in Figure 2A was applied to T5 cells (Wienecke et al., 2018), following a
738	previously developed protocol (Jagadeesh et al., 1993). This analysis relies upon the fact that a
739	drifting sinusoid grating may be decomposed into a sum of counterphase gratings as

740

741
$$c(t,x) = c_0 \sin(\omega t \mp \kappa x) = \frac{c_0}{4} \sum_{n=0}^7 \sin\left(\omega t + \frac{n\pi}{8} \mp \frac{\pi}{2}\right) \sin\left(\kappa x \pm \frac{n\pi}{8}\right)$$

742

Therefore, if a system is linear, its scaled, summed response of a linear system to counterphase
gratings with these phase shifts will be equal to its response to the corresponding drifting grating.
By comparing the linear prediction of the drifting grating response to the actual response, one
may assess a system's linearity.

747

To assess the orientation- and directional-tuning of the model with sinusoid gratings in Figure

749 2D, we defined a two-dimensional grating

750

 $c(t, x, y) = c_0 \sin(\omega t - \kappa(x \cos \theta + y \sin \theta))$

752

where the angle θ defines its orientation. In this analysis, we assume that the ring of detectors is located at y = 0, and that the Gaussian spatial filter is symmetric in x and y. Static gratings were formed by setting $\omega = 0$. We note that our convention for the orientation of a static grating differs from the original manuscript (Fisher et al., 2015); we define the orientation as the angle between the normal to the apparent edge and the preferred direction rather than the angular position of the edge itself. Therefore, in our convention the preferred orientations and directions align.

760

761 Apparent motion stimuli (Figure 3)

Single-bar stimuli were constructed as previously published (Gruntman et al., 2018; Salazar-Gatzimas et al., 2018). Briefly, 5° (respectively 2°) black or white bars were placed on a gray background, and presented for one second (respectively 160 ms) to match (Salazar-Gatzimas et al., 2018) (respectively (Gruntman et al., 2018)). Bar pair apparent motion stimuli were constructed as in (Salazar-Gatzimas et al., 2018). Briefly, 5° black or white bars were placed on a gray background and presented for one second. To create apparent motion, a second black or white bar was added 150 ms after the onset of the first bar at a neighboring spatial location.

Responses to these bar pair apparent motion stimuli were aligned such that the location of the

170 lagging bar matched the location of peak single-bar responses, as in (Salazar-Gatzimas et al.,

2018). Flashed apparent motion stimuli were constructed similarly to those presented to T4 and

- T5 cells in (Gruntman et al., 2018; Haag et al., 2016). Briefly, 4.5° white bars were placed on a
- gray background and were presented for 400 ms in sequential spatial positions.
- 774

775 Noise stimuli and linear receptive field extraction (Figure 4)

As previously published (Salazar-Gatzimas et al., 2016), we extracted linear receptive fields
from responses to uncorrelated binary stimuli composed of 5° black or white bars, updated at 60
Hz. We estimated the linear receptive field from these responses using reverse correlation
(Chichilnisky, 2001). Ternary noise stimuli with pairwise correlations were constructed as in
(Salazar-Gatzimas et al., 2016). Briefly, the contrast of the correlated noise stimulus was given

781

as

783
$$c(t,x) = \frac{1}{2} \left(B(t,x) \pm B(t+\delta t, x+\delta x) \right)$$

784

where B(t, x) is an uncorrelated binary stimulus composed of 5° black or white bars, and addition (respectively subtraction) generates positive (respectively negative) correlations. The stimulus was updated at a fixed rate, and the temporal offset δt was taken to be one cycle, with its sign determining whether the stimulus was oriented in the preferred or null direction. The spatial offset δx was fixed to be one bar width. As shown in (Salazar-Gatzimas et al., 2016), the autocorrelation function of this stimulus, with spacetime discretized by the bar width and sampling rate, is

792

793
$$\langle c(t,x)c(t+\tau,x+\rho)\rangle = \frac{1}{2}\delta_{\tau,0}\delta_{\rho,0} + \frac{1}{4}\left(\delta_{\tau,\delta t}\delta_{\rho,\delta x} + \delta_{\tau,-\delta t}\delta_{\rho,-\delta x}\right)$$

794

795 where $\delta_{i,j}$ is the Kronecker delta.

796

797 Three-point glider stimuli (Figure 4)

As in previous studies (Clark et al., 2014; Fitzgerald and Clark, 2015), we constructed three-

point glider stimuli following (Hu and Victor, 2010). Briefly, these binary stimuli enforce

800 correlations over space and time among triplets of pixels. Three-point gliders may be categorized

801 into four types: converging gliders with positive parity (con+), converging gliders with negative

802 parity (con-), diverging gliders with positive parity (div+), and diverging gliders with negative

803 parity (div-). Letting ρ be the pixel spacing and δ be the frame duration (the inverse of the

update rate), the update rules for each of the four three-point glider types are (see kymographs inFigure 4):

806

807	$c_{\rm con+}(t,x)c_{\rm con+}(t,x+\rho)c_{\rm con+}(t+\delta,x+\rho) = +1$
808	$c_{\rm con-}(t,x)c_{\rm con-}(t,x+\rho)c_{\rm con-}(t+\delta,x+\rho) = -1$
800	$a (t, y) = a (t + \delta y) = a (t + \delta y + a) = a$

809
$$c_{\text{div}+}(t,x)c_{\text{div}+}(t+o,x)c_{\text{div}+}(t+o,x+\rho) = +1$$

810
$$c_{\text{div}-}(t,x)c_{\text{div}-}(t+\delta,x)c_{\text{div}-}(t+\delta,x+\rho) = -1$$

811

812 The direction of the displacement of the spatial mean location of each triplet is inverted by 813 inverting the sign of the pixel spacing. Starting from an initial seed state, the values of each pixel 814 at each timepoint are determined by these update rules using the surrounding pixels' values. As 815 we simulate the full 360° of visual space, we use periodic boundary conditions to avoid 816 undetermined edge pixel values. As in previous studies (Clark et al., 2014; Fitzgerald and Clark, 817 2015; Leonhardt et al., 2016), the pixel spacing was taken to be 5° in both imaging experiments 818 and numerical simulations. In imaging experiments, visual stimuli were generated and presented 819 as described in previous studies (Badwan et al., 2019).

820

821 Random checkerboard stimuli (Figure 4)

Random checkerboard stimuli were constructed as in (Badwan et al., 2019). Briefly, 5° black or
white bars were placed at random with a density of 40% on a gray background. The resulting
checkerboards were then rigidly translated at a velocity of 100°/s. When combining rightwardand leftward-moving stimuli, summation was defined such that two white bars summed to white,
two black bars summed to black, and one white and one black bar summed to gray. Therefore,
the contrast of the composite stimulus matched that of the individual components, though its
density rose to 64%.

830 Natural scene stimuli (Figure 5)

831 Following prior work (Chen et al., 2019; Clark et al., 2014; Fitzgerald and Clark, 2015), we 832 generated a left-right symmetric ensemble of natural scenes by drawing independent row and 833 column samples from the database gathered by (Meyer et al., 2014). In this ensemble, scenes 834 were rigidly-translated at velocities sampled from a Gaussian distribution with a standard 835 deviation of 100%, which roughly matches typical rotational velocities of walking flies 836 (DeAngelis et al., 2019; Katsov and Clandinin, 2008). To convert the scenes to contrast signals, 837 we spatially filtered each image with the photoreceptor kernel to generate blurred images $I_{\rm blur}$, 838 and then used a Gaussian kernel with a standard deviation of 20° to estimate locally-averaged 839 images I_{mean} (Chen et al., 2019). The contrast signal was then defined as 840 $I_{\rm blur}(x,y) - I_{\rm mean}(x,y)$

841
$$c(x,y) \coloneqq \frac{I_{\text{blur}}(x,y) - I_{\text{mean}}(x,y)}{I_{\text{mean}}(x,y)}$$

842

As in previous studies of coactivation (Salazar-Gatzimas et al., 2018), the coactivations in Figure
5C were computed as normalized inner products of response timeseries. For all analyses in

Figure 5, we used an ensemble with 10^6 elements.

847 Appendix B: Parameter value selection

848

- 849 In this appendix, we briefly describe how we selected values of the weighting parameters
- 850 $g_{\rm exc}/g_{\rm leak}$ and $g_{\rm inh}/g_{\rm leak}$. We evaluated the model solely based on its ability to produce
- direction-opponent average responses to 1 Hz, 45° sinusoid gratings similar to those measured in
- T4 cells (Badwan et al., 2019). To do so, we considered the direction selectivity index and

analogous indices of direction-opponency and orthogonal direction enhancement, defined as

854

855
$$\mathrm{DSI} \coloneqq \frac{r(PD) - r(PD)}{r(PD) + r(PD)},$$

856

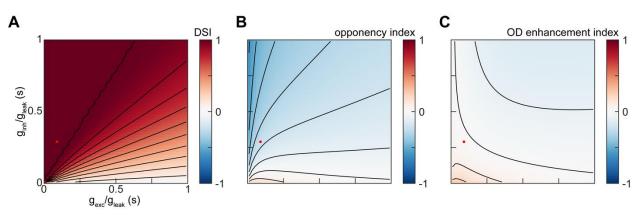
857
$$I_{\rm PD+ND} \coloneqq \frac{r(PD+ND) - r(PD)}{r(PD+ND) + r(PD)}$$

858 and

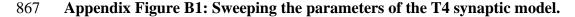
859
$$I_{\text{PD+OD}} \coloneqq \frac{r(PD+OD) - r(PD)}{r(PD+OD) + r(PD)}$$

860

As shown in Figure B1, there exists a broad region of parameter space for which the model
produces responses with a similar degree of direction-opponency to that measured in T4 cells
without significant PD+OD enhancement. We therefore made a simple choice of round-number
values within that region.



866 Appendix Figure B1, Zavatone-Veth et al. (2019)



- A. The direction-selectivity index of the T4 synaptic model's responses to drifting sinusoid
- gratings as a function of $g_{\text{exc}}/g_{\text{leak}}$ and $g_{\text{inh}}/g_{\text{leak}}$. Red dot indicates the selected values
- 870 of $g_{\text{exc}}/g_{\text{leak}} = 0.1$ and $g_{\text{inh}}/g_{\text{leak}} = 0.3$.
- B. As in (A), but for the opponency index.
- 872 C. As in (A), but for the OD enhancement index.
- 873

874 Appendix C: LNLN cascade factorization of the T4 synaptic model

875

876 In this appendix, we show how our T4 synaptic model may be factorized as a product of linear-

877 nonlinear-linear-nonlinear (LNLN) cascades representing the numerator and denominator of the

biophysical nonlinearity. The response C of the full model at each point in spacetime is given in

879 terms of the filtered contrast signal *s* as

880

881
$$C = R^2 \left(\frac{\tilde{g}_1 E_{\text{inh}} + \tilde{g}_2 E_{\text{exc}} + \tilde{g}_3 E_{\text{inh}}}{1 + \tilde{g}_1 + \tilde{g}_2 + \tilde{g}_3} \right)$$

882

883 where we have defined $\tilde{g}_i \coloneqq g_i/g_{\text{leak}}$ for brevity. Noting that the denominator of this expression 884 is always positive, we may re-express the response as

885

886
$$C = \frac{R^2 (\tilde{g}_1 E_{\text{inh}} + \tilde{g}_2 E_{\text{exc}} + \tilde{g}_3 E_{\text{inh}})}{(1 + \tilde{g}_1 + \tilde{g}_2 + \tilde{g}_3)^2}$$

887

hence the full EMD model admits a factorization into a product of LNLN models as

890 C(t,x) = N(t,x) D(t,x)

891 where

892 893

 $N(t, x) \coloneqq R^2(\tilde{g}_1 E_{\rm inh} + \tilde{g}_2 E_{\rm exc} + \tilde{g}_3 E_{\rm inh})$

- 894
- 895 and
- 896
- 897 $D(t, x) \coloneqq (1 + \tilde{g}_1 + \tilde{g}_2 + \tilde{g}_3)^{-2}$
- 898

899 which is bounded as $D(t, x) \le 1$. Because $D(t, x) \le 1$, $C(t, x) \le N(t, x)$.

900

901 The denominator LNLN cascade *D* is the result of applying a convex function (x^{-2} for x > 0) to

902 a non-negative linear combination of LN models with convex nonlinearities. Therefore, it cannot

 $R_{\beta}(x) \coloneqq \beta^{-1} \log(1 + \exp(\beta x))$

903 generate direction-opponent (DO) average responses to sinusoid gratings. The proof of this

904 proposition is a minor extension of our previous results on LNLN models with continuously-

905 differentiable convex nonlinearities and non-negative secondary linear filters (Badwan et al.,

- 906 2019). We define the soft ramp function
- 907
- 908
- 909

910 which is a continuously differentiable, monotone increasing, non-negative, and convex function of x for all positive β . As $\beta \to \infty$, $R_{\beta}(x) \to R(x)$ pointwise. By continuity, defining $D_{\beta}(t, x)$ 911 using R_{β} , we have $0 \le D_{\beta}(t, x) \to D(t, x) \le 1$ as $\beta \to \infty$. We denote the nonlinear functional 912 corresponding to the spacetime average of $D_{\beta}(t, x)$ for some input stimulus f as $D_{\beta}[f]$. As we 913 914 have the integrable constant dominating function 1, by the Lebesgue dominated convergence 915 theorem we have $0 \le D_{\beta}[f] \to D[f] \le 1$ as $\beta \to \infty$ (Stein and Shakarchi, 2009). By the result of (Badwan et al., 2019), we know that $D_{\beta}[PD + ND] \ge D_{\beta}[PD]$ and $D_{\beta}[PD + ND] \ge D_{\beta}[ND]$, 916 where $D_{\beta}[PD]$, $D_{\beta}[ND]$, and $D_{\beta}[PD + ND]$ are the average responses to PD, ND, and PD+ND 917 918 sinusoid gratings, respectively. As these inequalities hold pointwise for all positive β , by taking 919 $\beta \to \infty$ we may obtain $D[PD + ND] \ge D[PD]$ and $D[PD + ND] \ge D[ND]$. Therefore, the 920 denominator LNLN cascade cannot generate DO average responses to sinusoid gratings. 921 922 However, as the numerator LNLN model is the result of applying a convex function to a non-923 convex linear combination of LN models with convex nonlinearities, we cannot analytically 924 exclude the possibility that it could generate DO average responses to sinusoid gratings using the 925 results of (Badwan et al., 2019). In fact, numerical simulation shows that it can generate DO 926 average responses to sinusoid gratings, though it generates strong PD+OD enhancement (Figure 927 **C1**). It also generates DO responses over a smaller region in spatiotemporal frequency space 928 than the full model. If one replaced the infinitely sharp ramp functions with more biophysically 929 plausible soft rectifiers, the numerator LNLN cascade would be well-approximated for small

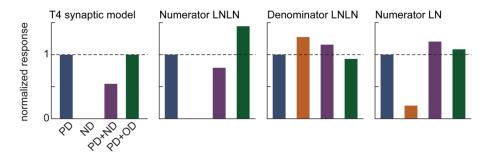
930 input contrasts by a LN model with a quadratic nonlinearity. Therefore, it could not generate DO

average responses for sufficiently small input contrasts. However, even in the limit in which both

932 the numerator and denominator are represented as LN models with quadratic nonlinearities, the

- 933 full model could likely generate DO average responses. In particular, this limiting construction
- 934 would resemble a type of adaptive gain model which was previously shown to generate DO
- 935 average responses (Badwan et al., 2019).





937 Appendix Figure C1, Zavatone-Veth et al. (2019)

938 Appendix Figure C1: Sinusoid grating responses of different components of the LNLN

939 factorization.

- 940 From left to right: Average responses of the full T4 synaptic model, the numerator LNLN
- 941 cascade, the denominator LNLN cascade, and the numerator LN cascade to 1 Hz, 45° sinusoid
- 942 gratings. All responses are normalized by the response of the given component to a grating
- 943 drifting in the PD of the full model.
- 944

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