1	Chromatic summation and receptive field properties
2	of blue-on and blue-off cells in marmoset lateral
3	geniculate nucleus
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17	Running head: Cone summation in blue-on cells
18	Key words: lateral geniculate nucleus, color vision, primate.

19

#### 20 Abstract

The "blue-on" and "blue-off" receptive fields in retina and dorsal lateral geniculate nucleus 21 (LGN) of diurnal primates combine signals from short-wavelength sensitive (S) cone 22 photoreceptors with signals from medium/long wavelength sensitive (ML) photoreceptors. 23 24 Three questions about this combination remain unresolved. Firstly, is the combination of S 25 and ML signals in these cells linear or non-linear? Secondly, how does the timing of S and ML inputs to these cells influence their responses? Thirdly, is there spatial antagonism 26 within S and ML subunits of the receptive field of these cells? We measured contrast 27 sensitivity and spatial frequency tuning for four types of drifting sine gratings: S cone 28 29 isolating, ML cone isolating, achromatic (S + ML), and counterphase chromatic (S – ML), in 30 extracellular recordings from LGN of marmoset monkeys. We found that responses to stimuli which modulate both S and ML cones are well predicted by a linear sum of S and ML 31 signals, followed by a saturating contrast-response relation. Differences in sensitivity and 32 timing (i.e. vector combination) between S and ML inputs are needed to explain the 33 amplitude and phase of responses to achromatic (S + ML) and counterphase chromatic (S -34 35 ML) stimuli. Best-fit spatial receptive fields for S and/or ML subunits in most cells (> 80%) required antagonistic surrounds, usually in the S subunit. The surrounds were however 36 37 generally weak and had little influence on spatial tuning. The sensitivity and size of S and ML subunits were correlated on a cell-by-cell basis, adding to evidence that blue-on and 38 39 blue-off receptive fields are specialised to signal chromatic but not spatial contrast.

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#### 41 Introduction

Color vision in primates originates in short– (S), medium– (M), and long-wavelength 42 sensitive (L) cone photoreceptors in the retina. The cone signals are processed within the 43 retina to form wavelength-selective afferent channels, including M – L opponent or 44 "red/green" channels, in which M and L signals are differenced, and "blue/yellow" channels, 45 driven by S cones in opposition to combinations of M and L cones. This paper addresses 46 three outstanding questions about S-cone signalling in "blue-on/yellow off" (also known as 47 blue-on, or more properly S-on) cells, and less commonly encountered "blue-off/yellow-on" 48 (also known as blue-off, or more properly S-off) cells. Firstly, is the combination of S and ML 49 50 signals in these cells linear or non-linear? Secondly, how does the timing of S and ML inputs 51 to these cells influence their responses? Thirdly, is there spatial antagonism within S and ML subunits of the receptive field of these cells? Answers to these questions will improve our 52 53 understanding of subcortical signals serving color vision.

54 Blue-on and blue-off cells have been the subject of longstanding study (reviewed by Dacey, 2004; Martin & Lee, 2014), and recordings from dorsal lateral geniculate nucleus 55 (LGN) have associated blue-on and blue-off cells with the koniocellular layers of the LGN 56 (Martin, White, Goodchild, Wilder & Sefton, 1997; Roy, Martin, Dreher, Saalmann, Hu & 57 Vidyasagar, 2009). The small bistratified (SBS) retinal ganglion cell is established as 58 59 exhibiting a blue-on/yellow off receptive field (Dacey & Lee, 1994; Szmajda, Martin & 60 Grünert, 2008; Crook, Davenport, Peterson, Packer, Detwiler & Dacey, 2009) but the anatomical substrate of blue-off cells in retina is less clear (reviewed by Dacey, 2004; Martin 61 & Lee, 2014). 62

63

Firing rates of blue-on cells and blue-off cells are determined by opponent

64 (antagonistic) combination of the S and ML inputs to the receptive field (Yeh, Lee & Kremers, 1995a; Solomon, Lee, White, Rüttiger & Martin, 2005; Field et al., 2007; Tailby, 65 66 Solomon & Lennie, 2008a; Crook et al., 2009). Responses to stimuli which modulate both S 67 and ML cones should therefore be determined by the (signed) sum of the contrast delivered to the S and ML cones. This prediction was confirmed for blue-on cells in macaque retina 68 when tested using combinations of S and ML contrast in spatially uniform fields (Crook et 69 al., 2009). The question as to whether this finding holds across the spatial frequency tuning 70 71 range of blue-on and blue-off cells has however not been specifically tested. Further, most 72 ganglion cells, including blue-on and blue-off cells (Yeh et al., 1995a), show some degree of response saturation for high contrast stimuli (Shapley & Victor, 1978; Lee, Virsu & Elepfandt, 73 1983; Kaplan & Shapley, 1986; Solomon et al., 2005) The specific question as to whether 74 response saturation influences responses of blue-on and blue-off cells to combined S and 75 ML contrast has also not been addressed. 76

77 Lee, Valberg and colleagues showed linear combination of saturating cone inputs can account for responses of blue-on ganglion cells to high-intensity narrow-band lights (Lee et 78 79 al., 1983; Lee, Valberg, Tigwell & Tryti, 1987). In their model, illustrated schematically in 80 Figure 1, nonlinearities in contrast response are present in the S cone and ML cone signals prior to their combination (Lee et al., 1983; Schnapf, Nunn, Meister & Baylor, 1990). If this is 81 the case, then differences in gain and/or saturation between S and ML inputs could yield 82 non-monotonic "supersaturating" responses to stimuli which activate both inputs. An 83 84 alternative (but not mutually exclusive) possibility is that linear S cone and ML cone inputs may be first summed, then subject to nonlinear distortion. In this case, the integrated 85 86 output must be monotonic with increasing contrast. These alternative models were not

explicitly compared in previous studies of cone inputs to macaque LGN cells (Lee et al.,
1983; Lee et al., 1987). Therefore, the first question we address in this paper is: how linear is
cone summation in blue-on and blue-off cells?

90 In addition to assuming linear combination of S and M inputs, most studies among those cited above had assumed that the opponent inputs arrive at the same time at the 91 92 integrating site, and therefore can be modelled as a simple difference of the input functions. More recent studies, however, indicate that there is a variable degree of delay between S 93 and ML inputs (~5 – 10 ms) to blue-on cells (Chichilnisky & Baylor, 1999; Field et al., 2007; 94 Tailby et al., 2008a; Pietersen, Cheong, Solomon, Tailby & Martin, 2014). Such delays could 95 96 help account for direction-selective achromatic responses observed in blue-on cells in 97 macague and marmoset LGN (Tailby et al., 2008a; Tailby, Dobbie, Hashemi-Nezhad, Forte & 98 Martin, 2010), but responses to combined S and ML contrast have not been systematically analysed. This is the second question we address in this paper. 99

100 The spatial distribution of cone inputs to S cone-signaling cells in the LGN was first described as "Type II", consisting of spatially overlapping S and ML subfields, in early work 101 102 by Wiesel and Hubel (1966). Later measurements showed that the ML-off subfield is slightly larger than the S-on subfield (Solomon et al., 2005; Field et al., 2007; Crook et al., 2009). 103 104 Crook et al. (2009) additionally provide evidence that inputs to the S-on subfield to blue-on 105 SBS cells have centre-surround structure (S-on, ML-off), and blue-on cells showing band-pass response to S cone gratings have been recorded in macaque and marmoset LGN 106 107 (Tailby et al., 2008a; Tailby, Szmajda, Buzás, Lee & Martin, 2008b). A population survey of 108 the spatial transfer properties of S cone and ML cone receptive field subunits has not been made, and the degree to which blue-on and blue-off cells are specialised to transmit 109

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- 110 chromatic versus spatial signals remains uncertain. Therefore, the third question we address
- in this paper is: are blue-on cells really Type II cells?

#### 112 Methods

113 Ethical approval.

114 Procedures conformed to the Australian National Health and Medical Research Council

115 (NHMRC) code of practice for the use and care of animals and institutional animal care and

116 ethics committee at the University of Sydney. Procedures also conform to the code of ethics

117 of the World Medical Association (Declaration of Helsinki).

#### 118 Animal preparation.

119 Details of our animal preparation, recording technique, and visual stimulation environment

have been published previously (Tailby et al., 2010; Pietersen et al., 2014). To summarize,

121 extracellular recordings of single units were performed in the lateral geniculate nucleus

122 (LGN) of common marmosets Callithrix jacchus. Animals were sedated with an

intramuscular injection of Alfaxan (12 mg kg<sup>-1</sup>, Jurox, NSW, AUS) and Diazepam (3 mg kg<sup>-1</sup>,

124 Roche, NSW, AUS), and anesthesia was maintained by continuous intravenous delivery of

125 Sufentanil citrate (6 – 30 μg kg<sup>-1</sup> h<sup>-1</sup>; Sufenta Forte, Janssen Cilag, Beerse, BEL). Depth of

- anesthesia was monitored by continuous electroencephalography and pulse oximetry
- 127 (SurgiVet, OH, USA). The animal was artificially respired with a 70%–30% mixture of

128 NO<sub>2</sub>-Carbogen (5% CO<sub>2</sub> in O<sub>2</sub>) and head-fixed in a stereotaxic frame. A durotomy was made

- above the LGN and a guide tube containing the recording electrode was inserted into the
- 130 brain. Action potential waveforms of single cells were discriminated by principal component
- analysis of amplified voltage signals from single microelectrodes (5 11 M $\Omega$ , FHC Inc.,

Bowdoin, Maine, USA). The position of each cell relative to the brain surface was recorded
from a hydraulic microdrive (David Kopf Model 640). Electrolytic lesions (3–6 μA x 3–6 s,
electrode positive) were made to assist in track reconstruction. At the conclusion of
recordings the animal was killed with an overdose of pentobarbitone sodium (80–150 mg
kg<sup>-1</sup>, i.v.). The position of recorded cells was reconstructed histologically as described in
detail in our previous publications (White, Goodchild, Wilder, Sefton & Martin, 1998;
Cheong, Tailby, Solomon & Martin, 2013).

139 Visual Stimuli

140 Stimuli comprised drifting sine gratings and sine-modulated flashing dots, displayed on a 141 stimulus monitor (refresh rate 100 or 120 Hz) against a grey background (mean luminance  $50 \text{ cd/m}^2$ ) which was centred on each receptive field using a front-silvered gimbaled mirror. 142 The driving voltage of the red, green, and blue phosphors of the stimulus monitor were 143 144 adjusted to produce cone selective ("silent substitution") gratings, using the spectral radiance distribution of the monitor phosphors, the sensitivity distribution of the marmoset 145 146 cone photoreceptors, and knowledge of the spectral absorbance of the optic media and 147 macular pigment (Brainard, 1996; Blessing, Solomon, Hashemi-Nezhad, Morris & Martin, 2004; Tailby et al., 2008a). Visual stimuli were generated using custom software which also 148 149 collected and sorted recorded spike waveforms and times to within 0.1 ms. (EXPO; 150 P. Lennie, University of Rochester, Rochester, NY). The S cone selective stimulus produced 60-80% contrast in S cones and less than 5% contrast in ML cones. The ML cone selective 151 152 stimulus produced 60-80% contrast in ML-class cones and less than 5% contrast in S cones, 153 relative to the nominal maximum achievable Michelson contrast of 100% for an achromatic (S + ML) stimulus, which modulated both cone classes equally. For brevity, the 154

cone-isolating stimuli are referred to as S cone and ML cone stimuli hereinafter. An

156 "optimal" mixed (S – ML) stimulus which drove the S and ML cones in counter-phase at half

the RMS cone contrast of the S cone and ML cone stimuli was also used. Drifting gratings

typically had a temporal frequency of 5 Hz (range 2-15 Hz). Stimulus aperture was typically

4° (range  $1^{\circ} - 12^{\circ}$ ), adjusted to ensure stimulation of both centre and surround.

Spatial frequency tuning curves were collected with drifting gratings ranging from
0.1 – 12.8 cycles/degree (cyc/deg) at 50% contrast. Contrast response curves were collected
with drifting gratings ranging from 2% – 70% contrast relative to the maximum achievable
achromatic contrast, typically at the identified peak spatial frequency (range 0 – 2.7
cyc/degree). The amplitude and phase of the first Fourier harmonic of the stimulus
frequency were taken as the primary measure of the cells' response to stimulation.

# 166 Analysis of contrast sensitivity

For the purposes of comparing S, ML, S + ML, and S – ML contrast response functions, the amplitude of the cell's response as a function of contrast were fit to saturating hyperbolic functions (Naka & Rushton, 1966; Sclar, Maunsell & Lennie, 1990) of the form

170 
$$K = \frac{M(c^n)}{c^n + c_{50}^n} + b$$
 (Eq. 1)

171 In which the spike rate K is a function of stimulus contrast c, theoretical maximum spike 172 rate M, semisaturation contrast  $c_{50}$ , exponent n, and noise-derived discharge at zero 173 contrast b. The semisaturation contrast  $c_{50}$  is the contrast at which the response is at half 174 of maximum. The exponent term n controls the steepness and curvature of the curve at 175 intermediate contrasts. Fits were performed in MATLAB (Mathworks, Natick MA) using 176 constrained non-linear least-squares minimization in which all values were constrained to be positive, the values of  $c_{50}$  were constrained to fall between 0 and 200%, and the values of the exponent term were constrained to be less than 3. These constraints ensured that the gain of the cell can be estimated even for low amplitude responses. When the fitted semisaturation constant was greater than the maximum contrast, it indicated a non-saturating response; when semisaturation constants were not constrained they could grow without bound without substantially improving the model fit to the data. The contrast gain is given by the derivative of Eq. 1,

184 
$$K' = \frac{n M c^n c_{50}^n}{c(c^n + c_{50}^n)^2}$$
 (Eq. 2)

and is computed at the semisaturation point  $c_{50}$  for saturating cells ( $c_{50} < 100\%$ ) and at maximum contrast for non-saturating cells. In prior work where the value of n was fixed to 1 (e.g. Kaplan and Shapley, 1986), the contrast gain was instead computed at low contrast (c = 0). It can be determined from Eq. 2 that at c = 0, K' exists and is non-zero if and only if n = 1.

For the purpose of model identification, data were also fit to a simpler version of Eq. 1 in which the exponent was fixed at 1, and to an elaborated version of Eq. 1. (Peirce, which permits non-monotonic supersaturating responses:

193 
$$K = \frac{M(c^{n_1})}{c^{n_2} + c_{50}^{n_2}} + b.$$
 (Eq. 3)

In this model there are two exponent terms n<sub>1</sub> and n<sub>2</sub>, and variables are otherwise as
Eq. 1. To test the hypothesis that this equation with its increased number of free
parameters provided a significantly better fit to the data, we calculated the residual errors
for each model, following Buzás, Kóbor, Petykó, Telkes, Martin & Lénárd (2013). If the more
complex model significantly reduced the residual error over its predecessor, as assessed by

a 1-sided 2-sample F test for equal variances (*p* < 0.05), then the more complex model was</li>
adopted. The data were also fit to a model which implements a thresholding rather than an
expansive nonlinearity:

$$K = \max\left(\frac{M(c-c_0)}{c-c_0+c_{50}}, 0\right) + b. \text{ (Eq.4)}$$

In this model the exponent term is fixed to 1, but a constant value is added to the contrast. At contrasts below  $c_0$ , the predicted response is the noise-derived discharge at zero contrast *b*.

# 205 Analysis of chromatic summation

Two extended models incorporating alternate hypothetical mechanisms for the summation of S and ML inputs were fit to the amplitude and phase of each cell's responses. The first model, illustrated in Figure 1A, assumes separable contrast mechanisms for S cone and ML cone input and is given by

$$K = \frac{M_S c_s |c_s|^{n_s - 1}}{|c_s|^{n_s} + c_{S,50}^{n_s}} + \frac{M_{ML} c_{ML} |c_{ML}|^{n_{ml} - 1}}{|c_{ML}|^{n_{ml}} + c_{ML,50}^{n_{ml}}} + b \text{ (Eq. 5)}$$

210 In which K is the complex spike rate,  $c_S$  is S cone contrast and phase,  $M_S$  is the theoretical saturation spike-rate (and relative phase) for S cone contrast,  $n_s$  is the 211 exponent of the S cone contrast function, and  $c_{S,50}$  is the semisaturation contrast for the S 212 cone contrast function. The notation for ML cone contrast mirrors that for S cone contrast. 213 The complex spike rate K is the complex sum of one Naka-Rushton curve representing S 214 215 cone input  $c_S$  and a second Naka-Rushton curve representing ML cone input  $c_{ML}$ . Prior 216 evidence (Yeh et al., 1995a; Solomon & Lennie, 2005) suggests that phase advance is not 217 prominent for blue-on cells, and so this expression ignores the phenomenon of phase

advance (Shapley & Victor, 1978; Benardete & Kaplan, 1999). In other words, all responses
to S cone stimuli share a single phase, as do all responses for ML cone stimuli. This model
assumes that the nonlinearities in the contrast response are present in the S and ML signals
prior to their combination, and also predicts that the overall contrast response to stimuli
having both S and ML contrasts need not be monotonic where there is a mismatch between
S and ML parameters.

224 An alternative model, illustrated in Figure 1B, assumes linear summation of cone 225 inputs followed by nonlinear distortion, and is given by

$$K = \frac{(M x)|x|^{(n-1)}}{(|x|^n + c_{50}^n)} + b,$$
$$x = w c_{ML} e^{i\phi_{ML}} + (1 - w)c_S e^{i\phi_S}$$

226 In which the complex spike-rate K is a function of the complex drive x, which itself is the weighted complex sum of the S cone and ML cone contrasts. As in Eq. 1, M is theoretical 227 228 saturation spike-rate,  $c_{50}$  is semisaturation contrast, n is an exponent term, and the tonic 229 discharge rate is b. The ML cone fraction w varies from 1 when the cell receives exclusively ML cone input to 0 when the cell receives exclusively S cone input, and the 230 231 response phase of the S cone and ML cone inputs are given by  $\phi_{ML}$  and  $\phi_{S}$ , respectively, for S cone contrast  $c_S$  and ML cone contrast  $c_{ML}$ . This model assumes that a linear 232 summation of approximately linear S cone and ML cone signals occurs, and that the output 233 234 of that summation is distorted by nonlinearity either at the ganglion cell level, at the ganglion cell to LGN synapse, or at the point of spike generation in the LGN. 235

# 236 Analysis of spatial distribution of cone inputs

We employed a straightforward extension of the standard difference-of-Gaussians (DoG) model of Rodieck & Stone (1965) and Enroth-Cugell & Robson (1966), which accounts for response phase by taking the complex sum of Gaussian kernels (Frishman, Freeman, Troy, Schweitzer-Tong & Enroth-Cugell, 1987). Each kernel  $G_n$  is given by

$$G_n(\omega) = \pi r_n^2 e^{2\pi i (\phi_n + \delta_n \omega) - (\pi r_n \omega)^2} (\text{Eq. 7})$$

in which the stimulus spatial frequency is  $\omega$ , the radius of the Gaussian kernel is  $r_n$ , the displacement of the field from the origin is  $\delta_n$ , and the response phase to a spatially uniform stimulus is  $\phi_n$ , which is determined from the cell's dominant (S or ML) input and response latency. The overall response spike-rate K is the complex sum of a variable number of Gaussian kernels, each having a magnitude  $M_n$ ,

$$K = \sum_{n=1}^{N} M_n G_n(\omega) . \text{ (Eq.8)}$$

This formulation is able to support spatial-frequency-dependent changes in response phase (Enroth-Cugell, Robson, Schweitzer-Tong & Watson, 1983). Fits were performed in MATLAB again using a constrained non-linear least-squares algorithm. The volume  $V_n$  of a receptive subfield, assuming circular receptive fields (but see also Chichilnisky & Baylor, 1999; Field et al., 2007; Tailby et al., 2010), is given by

$$V_n = \pi r_n^2 M_n$$
, (Eq. 9)

and can be thought of as the expected response of the cell if all other excitatory or
inhibitory subfields were blocked and the entire subfield were to be stimulated by its
preferred stimulus.

Not every cell for which spatial frequency tuning was available also had chromatic contrast
response data, and so in the spatial model contrast nonlinearities are ignored; instead,

256 response amplitudes were normalized by contrast. For each cell, a sequence of

Model Type	N =	Constraints	Free parameters
Classic type II RF	2 (S,ML)	$r_{S} = r_{ML}; \ \delta_{s} = \delta_{ML}; \\ \delta_{n} < 2 \ r_{n}$	6
Simple Opponent Cell	2 (S,ML)	$\delta_n < 2 r_n$	8
S cone	3	$r_{Sc} < r_{Ss}; \ \delta_{Sc} = \delta_{Ss};$	10
Centre-Surround	(Sc, Ss, ML)	$\phi_{Sc} = \phi_{Ss} + \pi; \ \delta_n < 2 \ r_n$	10
ML cone Centre-Surround	3 (S,MLc,MLs)	$r_{MLc} < r_{MLs}; \ \delta_{MLc} = \delta_{MLs};$ $\phi_{MLc} = \phi_{MLs} + \pi; \ \delta_n$ $< 2 \ r_n$	10
Double-Opponent Centre-Surround	4 (Sc, Ss, MLc, MLs)	$r_{Sc} < r_{Ss}; \ \delta_{Sc} = \delta_{Ss};$ $r_{MLc} < r_{MLs}; \ \delta_{MLc} = \delta_{MLs};$ $\phi_{Sc} = \phi_{Ss} + \pi;$ $\phi_{MLc} = \phi_{MLs} + \pi;$ $\delta_n < 2 r_n$	12

257 progressively more elaborate models was fit, as described in Table 1.

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For each model, residual errors were calculated and compared using 1-sided 2-sample F tests for equal variances (p < 0.1). The simplest model which significantly reduced the variance of the data compared to its predecessor was adopted, as described above for contrast responses.

263 Dataset

The dataset comprised recordings from 155 cells (51 blue-on cells, 6 blue-off cells, 74 P cells and 24 M cells) from 27 animals. Responses of 68 cells to parts of the stimulus set were previously described (Tailby et al., 2008b; Tailby et al., 2010; Pietersen et al., 2014; Cheong & Pietersen, 2014). What is new here is our analyses of datasets comprising responses to ML isolating stimuli and mixed chromatic (S – ML) stimuli as well as S and achromatic (S + ML) stimuli. Most receptive fields (63.3%) were located between 2° and 12° eccentricity; 14.8% were located within 2° of the fovea and 21.9% were located at more than 12°

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eccentricity. No systematic differences in responses to S cone stimulation were found
between trichromatic female animals (identified by the presence of red-green opponent
parvocellular cells), and the other dichromatic animals, all of which had a visual phenotype
consistent with the presence of S cones and one cone type with peak sensitivity close to 543
nm, 556 nm, or 563 nm (ML cones), and so data were pooled for analysis. Not every cell had
both a contrast and a spatial frequency measurement recorded.

277 Cells were classified as blue-on, blue-off, parvocellular (P), or magnocellular (M) by 278 the response to brief (200 or 500 ms) temporal square-wave stimuli (Pietersen et al., 2014) 279 supplemented by measurements of spatial and temporal frequency sensitivity. Examples of 280 a typical blue-on, P, and M response are shown in Figure 2.

Anatomical locations of 41%, or 64/155 cells (22/51 blue-on, 2/6 blue-off, 32/74 P, 281 and 8/24 M) were confirmed offline with histology as described above. In cases where track 282 283 location was not determined, the receptive field properties, eye dominance, encounter position, and response characteristics were used as criteria. Based on the combined 284 anatomical and physiological criteria, one blue-on cell was located in the (ventral-most) 285 286 koniocellular layer K1, 1 cell was in K2 (between the M layers), 17 cells were in K3 (between the M and P layers), 9 were in K4 (between the internal and external parvocellular layers), 287 288 and 4 cells were in K6 (dorsal to the external parvocellular layer). One blue-on cell was 289 located 'ectopically' in the ipsilateral M layer. The laminar location of 18 blue-on cells could 290 not be determined unequivocally. Two blue-off cells were located in K3, two blue-off cells 291 were located 'ectopically' in a P layer, and two blue-off cell locations could not be 292 determined unequivocally. No systematic differences in blue-on or blue-off cell properties were apparent on comparing receptive fields from different layers. 293

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#### 294 Results

#### 295 Chromatic contrast response functions

Figure 3A shows raster plots of a typical blue-on cell for 2 s presentations of 5 Hz S cone, ML 296 297 cone, achromatic (S + ML) and mixed chromatic (S – ML) drifting gratings, arranged by 298 stimulus contrast. At intermediate and high contrast the cell clearly responds in a phase-locked manner to each stimulus cycle, and the S cone response appears at close to 299 opposite phase of the ML response. Response phase for S cone and ML cone remains 300 301 constant across contrast. The lack of contrast-dependent phase advance agrees with previous data from blue-on cells in LGN of macaque and marmoset (Solomon & Lennie, 302 303 2005; Tailby et al., 2008b) and retina of macaque (Yeh et al., 1995a) and capuchin monkey Cebus apella (Silveira et al., 1999). The cell responds only feebly to achromatic (S + ML) 304 stimulus, but responds vigorously to the chromatic (S – ML) stimulus. Figure 3B shows the 305 306 average firing rate and first 5 Fourier components of the response at maximum contrast; the zeroth and first harmonic contain the majority of the response power. The power in the 307 308 higher harmonics chiefly arises from rectification of the response; when rectification is 309 addressed by fitting a rectified sine wave to the response, the residual harmonic distortion ratio (HDR; the ratio of the sum of the squares of the 2<sup>nd</sup> -5<sup>th</sup> harmonics to the square of the 310 311 fundamental) decreased by more than 20-fold. Across the population of recoded blue-on cells, correcting for rectification in the response results on average in an 8-fold decrease in 312 HDR (data not shown). Before correcting for rectification, median values of HDR for blue-on 313 314 cells are 0.57 (90% range 0.13-3.2) for S responses and 0.46 (90% range 0.12-2.0) for ML 315 responses. After correction, these ranges decrease to 0.17 (90% range 0.02-0.88) and 0.16 (90% range 0.01-1.04), respectively. 316

317 Figure 3C shows the amplitude of the first-harmonic responses with fitted Naka-Rushton curves, as given by Eq. 1. The response to S cone stimulation is expansive at 318 319 low contrast and shows mild saturation at high contrast, yielding a sigmoid shape. Expansive 320 nonlinearity at low contrast is less evident in the off-phase response to ML cone stimulation (Fig. 3B), but are also evident for the response to a mixed chromatic stimulus. It is 321 322 interesting to note that blue-on ganglion cells in macaque retina show little sign of expansive nonlinearity (Yeh et al., 1995a; Crook et al., 2009); we return to this point in a 323 324 later section.

Figure 3D shows the responses plotted in the complex plane, with time passing in clockwise direction. Plotting the data this way reveals that the ML cone response is in close-to-opposite phase but ~10 deg (~5 ms at 5 Hz) slower than the S response, as previously reported (Tailby et al., 2010; Pietersen et al., 2014). Figure 3E-H show example responses for a blue-off, yellow-on cell. This cell has a lower baseline firing rate and overall activity, but clear phase-locking and opposite-phase responses can also be observed. The Sand ML cone response phases are opposite to those observed for the blue-on cell.

332 For blue-on cells, there were no significant differences between the distributions of the fitted Naka-Rushton semisaturation constants for curves fitted to the first-harmonic 333 334 response, as compared to curves fitted to a rectification-corrected harmonic response, for 335 either S, ML, achromatic, or mixed S – ML stimulation (p > 0.6 for each case, paired Wilcoxon test). All values reported above were fitted to unmodified first-harmonic 336 337 responses, as is typically reported. Figure 4 shows distributions of the fitted semisaturation 338 constants (Fig. 4A), maximum gains (Fig. 4C), and exponent parameters (Fig. 4E) fitted using Eq. 1 to S cone and ML cone stimuli for blue-on and blue-off cells, as well as the 339

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340	distributions of these fitted parameters for P and M cells (Fig. 4B, D, and F). Summary
341	statistics for the data shown in Figure 4 are given in Table 2. As expected (Kaplan & Shapley,
342	1986; Yeh et al., 1995b), the M cell semisaturation constants were lower, and M cell gain
343	was significantly higher, than those of P or blue-on cells (p < 0.01 for all comparisons, 3-way
344	Kruskal-Wallis test [KW]). The P cell exponent terms were significantly lower than those of
345	blue-on or M cells (p < 0.01, KW). For blue-on cells, S cone gain was correlated with ML cone
346	gain (Fig. 4C, $r = 0.77$ , $p < 0.01$ ) and significantly greater than ML gain ( $p = 0.01$ , 2-way KW),
347	suiting these cells' description as blue-on as opposed to yellow-off. Similarly, semisaturation
348	constants fitted to S cone responses were significantly lower than those fit to ML cone
349	responses (p < 0.01, 2-way KW). Of the 33 blue-on cells within 10° of the fovea for which a
350	contrast response function could be fit, 26 had a clear half-maximal response nonlinearity
351	( $c_{50} < 100\%$ ) in their S response and 20 had a clear nonlinearity in their ML response.

# 352

Table 2
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	Semisaturation (%)	Contrast gain (imp/s % <sup>-1</sup> )	Exponent
Blue-on cells S cone	63.4 ± 45.2	0.60 ± 0.42	2.4 ± 0.6
Blue-on cells ML cone	87.3 ± 46.7	0.40 ± 0.30	2.3 ± 0.7
Blue-off cells S cone	29.4 ± 17.9	0.55 ± 0.36	2.3 ± 0.8
Blue-off cells ML cone	65.6 ± 58.5	0.32 ± 0.22	2.2 ± 1.0
P cells	92.9 ± 51.0	0.35 ± 0.34	1.9 ± 0.7
M cells	22.1 ± 29.6	2.17 ± 1.34	2.4 ± 0.6

353

All reported values mean ± standard deviation

The scatterplot in Figure 4A raises the impression of two populations of blue-on cells with respectively high and intermediate semisaturation constants. We found however that (apart from the correlation of S and ML gain noted above) the joint distributions of the fitted Naka-Rushton parameters for blue-on cells did not hold significant correlations, nor was there evidence of clustering in the parameter space (data not shown). A low semisaturation

359	constant for S cone contrast neither predicted nor was predicted by a low semisaturation
360	constant for ML cone contrast ( $chi^2 = 0.04$ , p = 0.833). Contrast gain was mildly correlated
361	with eccentricity for both S cone and ML cone (r = 0.49 and r = 0.55, respectively); however,
362	no other parameter was correlated with eccentricity. This result stands in contrast to the
363	result for P cells in which semisaturation, was inversely correlated with eccentricity (r =
364	-0.36; p = 0.018 vs. r = 0.17; p = 0.273) as previously observed (Solomon, White & Martin,
365	1999). Blue-off cells (green cross symbols, Fig. 4A,C,E) had similar characteristics to blue-on
366	cells but the number of cells is too small to make statistical comparisons. In comparison to a
367	simplified model in which the exponent term is held constant (see Methods section), the
368	expansive nonlinearity of Eq. 1 was necessary to explain the responses for a majority (29/44,
369	65%) of blue-on cells. Interestingly, the addition of an expansive term also improved the fits
370	for a substantial fraction (26/53, 49%) of P cells. On the other hand, the use of a
371	supersaturating Naka-Rushton curve (Eq. 3) did not significantly improve the fit to the data
372	for any cell, and non-monotonic responses were never observed in blue-on cells. Either an
373	expansive or a thresholding nonlinearity (Eq. 4) could adequately fit the data overall.
374	However, only the expansive nonlinearity model (Eq. 1) could consistently predict responses
375	at the contrast threshold. For blue-on cells, the overall RMS error of the expansive
376	nonlinearity model was not significantly different to the overall RMS error of the threshold
377	model (3.7 $\pm$ 1.4 imp/s vs. 3.9 $\pm$ 1.5 imp/s, respectively, p = 0.34, paired Wilcoxon test).
378	Response saturation is customarily associated with contrast gain control, which is
379	also manifest as advance in response phase with increasing contrast (Shapley & Victor,
380	1981; Kaplan & Benardete, 2001). On comparing response phase at half-maximum and
381	maximum contrast for preferred chromatic direction we found substantial phase advance in

382 MC cells (37.4 ± 18 degrees, n = 22), as would be expected (Yeh et al., 1995a; Levitt, Shumer, Sherman & Spear, 2001; Solomon, Lee & Sun, 2006). Phase advance in MC cells was 383 384 significantly greater (p < 0.02, Kruskal-Wallis test) than the (negligible) phase advance in PC 385 cells (6.0 degrees  $\pm$  13, n = 46) and blue-on cells (3.3  $\pm$  9, n = 44); there was no difference between PC cells and blue-on cells (p = 0.95, Kruskal-Wallis test). There was no clear sign of 386 correlation between semisaturation constant and phase advance in blue-on cells ( $r^2 < 0.01$ , 387 p = 0.82). These data suggest that blue-on cells do not show contrast gain control to any 388 389 great extent. A more extensive analysis of response timing may be of interest for future 390 work.

# 391 Achromatic responses arise from the summation of chromatic signals

Blue-on cells integrate S cone signals with ML cone signals, and it is clear from Figure 4 that 392 there is variation between the S cone and ML cone responses within individual cells. We 393 394 showed above that this variation may be due to S and ML signals having different contrast response characteristics at the level of the outer retina. If this is the case, then the 395 integrated responses to co-stimulation of S and ML cones is given by the sum of the 396 397 responses to cone stimuli, which could lead to non-monotonic achromatic contrast responses (Fig. 1A, Eq. 5). Alternatively, the S cone and ML cone inputs to the blue-on cell 398 399 may be approximately linear, in which case the integrated output cannot be expressed 400 exactly as the sum of the component responses, but must be monotonic with increasing 401 contrast (Fig. 1B, Eq. 6).

Figure 5 shows a particularly illustrative example of a blue-on cell's responses where the predictions made by these two models diverge. Figure 5A shows the cell's responses as a function of contrast with fitted Naka-Rushton curves, as given by Eq. 6, for S and ML cone 405 stimuli. Chromatic contrast is expressed relative to stimulus cone contrast (80% for S and ML cones). Figure 5B shows the measured achromatic contrast response function, along 406 407 with predicted contrast response functions given by fitting the two competing models to the 408 S and ML cone response data. The model with separable contrast mechanisms (dashed line, Fig. 5B) predicts a non-monotonic achromatic contrast response. The non-monotonic 409 410 prediction arises because the expansive nonlinearity for ML-off is weaker than that for S-on, predicting response cancellation at high but not intermediate contrast. The measured data, 411 412 however, falls on the curve predicted by the summation of linear cone inputs which are 413 distorted by a nonlinearity post-integration. Figure 5C shows the amplitude and phase of 414 the responses in the complex plane.

415 Overall, the weight of experimental evidence favors the model where summation of 416 linear cone inputs occurs prior to non-linear distortion, as described in Eq. 6 and illustrated in Figure 1B. Across the population of blue-on cells, non-monotonic achromatic contrast 417 responses were never observed. Most cells could be reasonably fit by either model, as 418 419 would be expected if the cells are operating in the linear regime of their contrast response 420 functions. Figure 6 shows measured vs. predicted responses to achromatic (S + ML) and 421 mixed chromatic (S - ML) stimuli for the population of cells tested. Figure 6A shows the 422 predictions of Eq. 6 for achromatic (S + ML) stimulation (median RMS error 8.1 imp/s), Figure 6B shows the predictions of Eq. 6 for mixed chromatic (S – ML) co-stimulation 423 (median RMS error 5.4 imp/s), Figure 6C shows the predictions of Eq. 5 for achromatic (S + 424 425 ML) stimulation (median RMS error 8.3 imp/s), and Figure 6D shows the predictions of Eq. 5 426 for S – ML stimulation (median RMS error 11.0 imp/s). The two models perform similarly in 427 prediction of achromatic responses (p = 0.66, 2-way KW), but the linear inputs model clearly 428 outperforms the separable contrasts model for chromatic (S – ML) stimulation (p < 0.01,</li>
429 2-way KW).

#### 430 Amplitude and timing of opponent inputs to blue-on and blue-off cells

431 Figure 7 shows, for the data collected using 5 Hz stimulation, the amplitude and phase of 432 responses to maximum-contrast S cone (Fig. 7A), ML cone (Fig. 7B), achromatic (Fig. 7C), and S – ML stimulation (Fig. 7D) in the complex plane. Figures 7C and 7D also show (square 433 434 symbols) the distributions of achromatic and S – ML responses predicted by a vector sum of 435 S and ML inputs. There are four main points to take from these data. Firstly, as expected, the S and ML responses cluster around close-to-opposite phase, consistent with S-on and 436 437 ML-off excitation. The mean phase delay between S and ML was 187 deg, equivalent to an addditional 3.6 ms at 5 Hz. Secondly, there is considerable variability in response phase 438 across the population for all stimuli. As the data of figure 7 were collected at varying spatial 439 440 frequencies, some variation in the observed response phases is due to offset of the centre of the receptive field from the centre of the stimulus patch. The true response phases to 441 442 uniform stimulus are less variable than this figure implies – the circular standard deviation 443 of the response phases to spatially uniform 5 Hz S cone stimulation is 22.4° (n = 40), equivalent to  $\pm 12.4$  ms at 5 Hz. Thirdly, for most (but not all) blue-on cells the phase of the 444 response to achromatic modulation is close to that for S cone modulation (circular r = 0.42, 445 p < 0.02). Finally, the fit predictions for achromatic and S – ML stimuli show heavy overlap in 446 phase space with the measured values, most evidently for S – ML stimuli (Fig. 7D). The 447 448 reader should note that these data are also consistent with the "sum-then-distort" contrast 449 integration model described above (Eq. 6, Fig 1B, Fig. 6).

450 Figure 7E shows on a per-cell basis the amplitude and phase of responses to

451 maximum-contrast S cone, ML cone, together with measured achromatic (S + ML) response (black vector) and the vector sum prediction (magenta square symbols). Responses are 452 453 normalized to the maximum response amplitude, shown at bottom of each sub-plot. Here 454 we see that the broad distribution of response phase across the population belies some hidden regularity, by revealing that the S and ML response phases are correlated on a cell by 455 cell basis (circular r = 0.61, p < 0.01). Responses to achromatic stimuli are attributable to 456 differences in amplitude and/or timing of the S and ML inputs. Two example cells are 457 458 marked in Figure 7E. The cell in the first example (open arrowhead, Fig. 7E [68, bottom left]) 459 shows identical amplitude for S and ML inputs, but the ML input lags the S input by 209°, predicting a vigorous on-type achromatic response. For the second, [54, upper right] the 460 response phase difference is 187° but the S input amplitude is greater than the ML input 461 amplitude, again predicting a (weak) achromatic on-type response. These data show that 462 the gain and timing of S and ML inputs jointly determine responses to non-cone-isolating 463 464 stimuli.

#### 465 Spatial properties of S and ML inputs to blue-on cells

466 A variety of receptive field structures was found for blue-on cells, as illustrated in Figure 8. The simplest model (spatially coextensive S and ML subfields consistent with classical "type 467 468 II" organization), best fit the responses of 15 of 47 cells (32%). An example of the receptive field structure of a type II cell is shown in Figure 8A. The associated S cone, ML cone, and 469 470 achromatic tuning curves and fitted Gaussians are shown in Figure 8B, and the S and ML 471 cone responses are plotted in the complex plane in Figure 8C. These cells rarely showed 472 achromatic responses at any spatial frequency; however, spatially low-pass achromatic responses were observed in 5 of the 15 type II cells. In 5 cells (11%), a S-on subfield and a 473

474 ML-off subfield of different sizes (which were not necessarily concentric) were sufficient to explain the cell's response, consistent with the receptive field organization described in 475 476 Tailby et al. (2010). An example of this receptive field structure is shown in Figure 8D, along 477 with the associated fitted S cone, ML cone, and achromatic tuning curves (Fig. 8E) and location in the complex plane (Fig. 8F). Note the band-pass response to achromatic 478 479 stimulation. In 13 cells (28%), the S cone input had spatial bandpass tuning without evidence of a corresponding bandpass characteristic for ML cone input. Responses of such 480 481 cells are consistent with the presence of an S cone surround. An example of such a cell is 482 shown in Figure 8G, along with the associated fitted S cone, ML cone, and achromatic tuning curves (Fig. 8H) and location in the complex plane (Fig. 8I). Both the achromatic and the S 483 484 cone spatial frequency tuning curves show response roll-off at low spatial frequencies. Bandpass achromatic responses were present in 6 of 13 cells, with low-pass responses 485 486 observed in a further 2 cells. The remainder (14 out of 47 cells, 30%) required both an S-off 487 surround and a weak ML-on surround to explain the cell's response. An example of such a receptive field is shown in Figure 8J, along with the associated fitted S cone, ML cone, and 488 achromatic tuning curves (fig. 8K) and location in the complex plane (fig. 8L). As we show 489 490 below, overall the ML surrounds were weaker than S surrounds, and made only a small but 491 significant improvement in fit quality. Spatial bandpass achromatic responses were more frequent in this population, being present in 9 out of 14 cells. 492

Figure 9 shows radius and volume of S cone and ML cone subunits for blue-on and blue-off cells, and compares these parameters with ML cone inputs to P cells. Summary statistics for the data shown in Figure 9 (A, B, C, E) are given in Table 3. For blue-on cells, the S-on subfield has a marginally smaller radius than the ML-off subfield (Fig. 9A, p = 0.10,

497	paired Wilcoxon test), but has, on average, significantly larger volume (Fig. 9B, p < 0.01,
498	paired Wilcoxon). The S-on subfield radii increased with foveal eccentricity (Kendall's tau =
499	0.31, p < 0.01) as has been reported previously (Tailby et al., 2008b). Surprisingly, the
500	eccentricity–dependence was less pronounced for the ML-off subfields (tau = 0.15, $p$ =
501	0.17). The reason for this difference is not clear. The reported receptive field radii for the
502	S-on and ML-on subunits are within the range which would be expected based on the
503	dendritic arbor of the SBS RGC: 0.50° $\pm$ 0.22° for S cones and 0.26° $\pm$ 0.15° for ML cones (see
504	also Tailby et al., 2010).

505

# Table 3

Blue-on cells	Radius (deg)	Volume (imp/s % <sup>-1</sup> )	Frequency
S-cone on-polarity centre	0.38° ± 0.37°	1.58 ± 1.15	100% (47/47)
ML-cone off-polarity centre	0.40° ± 0.33°	1.29 ± 0.98	100% (47/47)
S-cone off-polarity surround	1.08° ± 0.69°	1.09 ± 1.13	55% (26/47)
ML-cone on-polarity surround	2.41° ± 5.44°	1.33 ± 1.56	30% (14/47)
P cell centre (ML-cone)	0.10° ± 0.10°	1.19 ± 0.48	100% (72/72)
P cell surround (ML-cone)	0.80° ± 0.88°	0.91 ± 0.40	85% (61/72)

506 All values mean ± SD

507	Figure 9C compares ML centre and surround radius for P cells against the S-on, S-off,
508	ML-off, and ML-on subunits (where detectable) of blue-on cells. Figure 9D shows the ratios
509	of these quantities. As expected (for review, see Martin & Lee, 2014), P cell centre radii are
510	significantly smaller than S-on and ML-off radii in blue-on cells (p < 0.01 for both
511	comparisons, independent Wilcoxon). Also as expected, the S-on/ML-off ratio for blue-on
512	cell RF radii (0.993 $\pm$ 0.408) is greater than the ratio of P cell centre/surround radii (0.203 $\pm$
513	0.173, p < 0.01, n-way KW), reflecting their distinct origins in retinal wiring (Field et al.,
514	2007; Crook et al., 2009). Where measurable, the blue-on cell S-on/S-off and ML-off/ML-on
515	radius ratios (0.347 $\pm$ 0.218 and 0.365 $\pm$ 0.186, respectively) were also distinguishable from

P cells (p < 0.01 and p = 0.04, respectively); perhaps unsurprisingly, the distribution of</li>
ML-cone centre/surround ratios was more similar to the distribution of P cell
centre/surround ratios.

519 Figure 9E compares ML centre and surround volumes for P cells against the S-on, S-off, ML-on and ML-off subunits (where detectable) for blue-on cells. Figure 9F shows the 520 521 ratios of these quantities. In contrast to the marked differences in radii, P cell centre volumes were not significantly different from S or ML subunit volumes in blue-on cells (p = 522 0.09 and p = 0.28 for S cone and ML cone subfields, respectively, independent Wilcoxon). 523 Correspondingly, the S-on/ML-off volume ratio in blue-on cells (1.470 ± 1.037) was close to 524 the centre/surround volume ratio in P cells ( $1.596\pm0.860$ , p = 0.414, independent Wilcoxon). 525 This result may help explain why P cells and blue-on cells both respond weakly to 526 527 achromatic contrast at low spatial frequencies: the antagonistic subunit sizes are very different in P cells and blue-on cells, yet the integrated sensitivities are similar and 528 well-matched. On the other hand, where detectable, the volume ratios S-on/S-off and 529 530 ML-off/ML-on in blue-on cells were higher  $(2.613 \pm 1.952 \text{ and } 2.286 \pm 1.070, \text{ respectively})$ . 531 In other words, the spatially antagonistic subunits in blue-on cells contribute only weakly to 532 shaping the cells' overall response.

#### 533 Discussion

We show that when stimulated with low-to-moderate S cone or ML cone contrast, blue-on cells show predominantly linear contrast response functions. Mild response saturation at high contrast and some expansive nonlinearity at low contrasts are evident.

537 The degree of response saturation in blue-on cells shows heavy overlap with that of

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538 PC cells and is much weaker than that of MC cells (Fig 4A, B). The expansive nonlinearity could potentially sharpen blue-on cells' chromatic selectivity as proposed for cortical 539 540 neurons (Heeger, 1992; DeValois, Cottaris, Elfar, Mahon & Wilson, 2000; Solomon, Peirce & Lennie, 2004; Solomon & Lennie, 2005), but our stimulus set was too restricted to address 541 this possibility. The fact that blue-on ganglion cells show little sign of expansive nonlinearity 542 (Yeh et al., 1995a; Crook et al., 2009) suggests that this distortion occurs either at the 543 ganglion cell to LGN cell synapse or at the LGN spike output stage. Expansive nonlinearity at 544 545 low contrasts is also a feature of PC cell responses in LGN of awake and anesthesthetised 546 macaque monkeys (Alitto, Moore, Rathbun & Usrey, 2011), thus its presence in the PC cells and blue-on cells we recorded is not simply attributable to a species difference, or 547 sufentanil anesthesia, or a switch of the LGN into a non-responsive "burst" mode (Sherman, 548 1996). Is it important to note however that the effects we see in LGN are much milder than 549 the nonlinearities in cortical cell responses reported in the studies cited above and 550 elsewhere. 551

We found that responses to simultaneous recruitment of both S cones and ML cones 552 553 are consistent with integration of linear S cone and ML cone signals prior to non-linear 554 distortion of the integrated signal. This result is consistent with measurements of linear S cone and ML cone synaptic inputs to SBS ganglion cells (Chichilnisky & Baylor, 1999; Field et 555 al., 2007; Crook et al., 2009). In common with these previous reports, in our experimental 556 setup we could only accesss a limited range of intensities (maximum 120 cd m<sup>-2</sup>), where the 557 558 cone signals are likely operating in a linear range. Lee, Valberg and colleagues (1983, 1987) used narrow band spectral lights to show saturating cone inputs are needed to account for 559 responses of blue-on cells at high intensities (> 500 cd  $m^{-2}$ ). The presence of response 560

saturation at a pre-cortical level may also help to explain some features of cortical contrast
response (Solomon & Lennie, 2005) and fits within some models of the LGN's computational
contribution to vision (Dan, Atick & Reid, 1996; Mante, Frazor, Bonin, Geisler & Carandini,
2005).

565 Our observation of blue-on LGN cells with varying receptive field structures may also 566 have implications for current theories on the retinal circuitry underpinning blue-on cells. One current theory is that the SBS ganglion cell receives convergent input from a S 567 cone-specific bipolar cell (BB) as well as one or more classes of diffuse off-type bipolar cell 568 (DB) (Crook et al., 2009; Dacey, Crook & Packer, 2013). The S cone signal from the BB input 569 570 is expected to carry an off-polarity ML cone surround (Packer, Verweij, Li, Schnapf & Dacey, 571 2010) and the ML cone signal from the DB cells carries an on-polarity ML cone surround 572 (Dacey, Diller, Verweij & Williams, 2000). Crook et al. (2009) proposed that mutual 573 anhiallation of these ML surround signals generate spatially co-extensive S cone and ML cone receptive fields (Crook et al., 2009, Dacey et al., 2014). Cell-specific imbalances in this 574 mechanism could be expected to produce a distribution of ML cone surrounds, which could 575 576 partially explain the variation in receptive field structure observed in blue-on cells in the 577 LGN. The convergent BB+DB input theory, however, fails to provide an explanation for the off-polarity S cone surround that we observed in 57% of cells, nor why we did not observe 578 blue-on cells with ML cone, but not S cone, surrounds. It is possible that the S cone surround 579 is mediated by inhibitory amacrine cell input onto SBS ganglion cells (Ghosh & Grünert, 580 581 1999); however, if this were indeed the case, why have inhibitory S cone surround 582 responses not been reported in SBS ganglion cells? An alternative possibility is that the S 583 cone off-polarity surround is a consequence of lateral inhibition in the LGN.

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584 In summary, the image of blue-on and blue-off cells which emerges from these data is largely compatible with the blue-yellow color opponent channel originally hypothesized by 585 Hering (Hering, 1878; Jameson & Hurvich, 1955). The (lilac-lime) axis of maximum sensitivity 586 for blue-on cells is not exactly aligned with the perceptual blue-yellow axis proposed by 587 Hering. But other properties of blue-on cells conform nicely to requirement of a chromatic 588 opponent channel. Blue-on cells give responses of opposite polarity to short-wave and 589 medium-wave regions of the spectrum. These responses show mutual antagonism, yielding 590 591 weak responses to achromatic (S + ML) stimuli and strong responses to mixed chromatic 592 (S–ML) stimuli. Further, the spatial properties of the antagonistic S-on and ML-off subunits are well-matched, yielding vigorous low-pass responses to chromatic contrast. Spatial 593 centre-surround structure in blue-on receptive fields, where detectable, was weak and 594 variable. One interpretation of this variability is that central mechanisms may not prioritize 595 spatial contrast enhancement through blue-on cells. In this case the question whether 596 597 blue-on cells are strictly Type II or not becomes largely academic. As a population these cells are clearly specified to favour chromatic over spatial contrast. 598

599

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747

#### 748 Figure Legends

749 Figure 1

750 Prospective models for cone opponency. A: nonlinear inputs, linearly separable output

751 model corresponding to Eq. 5 B: linear input, nonlinear output model corresponding to Eq. 6

752 Figure 2

753 Cell Type classification. Example response raster plots and PSTHs showing responses to brief 754 (0.2s) increments and decrements of ML- and S cone contrast. Representative blue-on 755 koniocellular (A), parvocellular (B), and magnocellular (C) cells are shown. D: Response of cell (A) to S- and ML cone-isolating drifting gratings (1 cyc/deg, 5 cyc/s, 4° aperture) as a 756 757 function of cone contrast. Error bars, in some cases smaller than the data point, are ± std. 758 error. Arrowhead shows the contrast used for the data shown in (E). E: Response of cell (A) to S- and ML cone-isolating drifting gratings (50% contrast, 5 cyc/s) as a function of grating 759 760 spatial frequency. F, G: Responses of cell (B) to achromatic flashing dot and drifting gratings 761 (7.1 cyc/s, 2° aperture (F) and 8° aperture (G)). H, I: Response of cell (C) to achromatic flashing dot and drifting gratings (5 cyc/s, 1° aperture). 762

763

#### 764 Figure 3

765 Typical contrast responses for a blue-on and blue-off cell. A: Raster plots for 2 s

766 presentations of 5 Hz S cone-isolating, ML cone-isolating, achromatic S+ML, and mixed S-ML

<sup>767</sup> 2° flashing dot, organized by stimulus contrast, for a typical blue-on cell. B: Mean firing rate

- and first 5 Fourier components of the response at maximum contrast. The zeroth and first
- harmonic contain the majority of the response power. C: First-harmonic contrast response
- curves for the data presented in (A), showing response amplitude. Error bars, typically

smaller than the data point, are ± std. error. Smooth curves show fits of Eq. 1 to the data. D:
Complex plane plot for the data presented in (A), showing the amplitude and phase of the
first-harmonic response. This cell has an on-phase response to S cone stimulation and an
off-phase response to ML cone stimulation. E, F, G, H: Responses of a blue-off cell to a 4Hz
drifting grating (0.1 cyc/deg, 8° aperture) of the above chromaticities. This cell has an
off-phase response to S cone stimulation and an on-phase response to ML cone stimulation.

778 Figure 4

779 Scatterplots and distributions of fitted Naka-Rushton equation parameters. A: Scatterplot of 780 fitted semi-saturation constants for S- and ML cone-isolating stimuli for blue-on and blue-off 781 cells. Lines of unity shown in grey. Values over 100% indicate non-saturating responses. B: 782 Distributions of fitted semi-saturation constants for ML cone-isolating stimuli for blue-on, P, 783 and M cells. Hatching indicates truncation of plotted distributions. C: Scatterplot of contrast gains for S- and ML cone-isolating stimuli for blue-on and blue-off cells. D: Distributions of 784 contrast gains for ML cone-isolating stimuli for blue-on, P, and M cells. E: Scatterplot of 785 786 fitted exponents for S- and ML cone-isolating stimuli for blue-on and blue-off cells. F: Distributions of fitted exponents for ML cone-isolating stimuli for blue-on, P, and M cells. 787 788 Figure 5 789 Illustrative example of difference in predictions made by Eq. 5 and Eq. 6 for a selected 790 sample cell. A: First-harmonic S- and ML cone contrast response functions. Error bars, in 791 most cases smaller than the data point, are ± std. error. Smooth curves show fits of Eq. 6 to 792 the data. Open circles are S cone responses; filled circles are ML cone responses; black 793 diamonds are achromatic responses. B: Measured and predicted first-harmonic response to 794 achromatic stimulation. Dashed curve plots the achromatic response predicted by Eq. 5,

solid curve plots the achromatic response predicted by Eq. 6. C: Complex plane plot for the
data presented in (A) and (B), showing the amplitude and phase of the first-harmonic
response.

798 Figure 6

799 Predicted vs. measured response amplitudes for achromatic S+ML and mixed S – ML

stimulation. A,B: Scatterplot of predicted vs. measured achromatic (A) and "optimal" S – ML

801 (B) response amplitudes across all contrasts, excluding contrasts at which either S cone or

ML cone stimulation did not evoke a response > 5 imp/s. The model assuming linear inputs,

803 corresponding to fig. 1B, is used to generate predictions for each cell. C,D: Scatterplot of

804 predicted vs. measured achromatic (C) and "optimal" S – ML (D) response amplitudes across

contrasts. The model assuming separable nonlinear inputs, corresponding to fig. 1A, is used
to generate predictions for each cell.

807 Figure 7

808 Predicted vs. measured complex achromatic response across all blue-on cells measured at 5 809 Hz. A-D: Population summary of aggregate F1 response in the complex plane, for S cone-810 isolating (B), ML cone-isolating (C), achromatic S + ML (D), and counter-phase chromatic S – 811 ML (E) stimuli. Open squares show predictions by Eq. 6 for (D) and (E). E: aggregate S 812 cone-isolating, ML cone-isolating, achromatic, and predicted achromatic responses for each 813 cell, arranged from most foveal (top left) to most peripheral (bottom right) and normalized 814 to each cell's response (in imp/s), given in bottom right of each plot. Clockwise motion on 815 each dial indicates increasing phase lag. Off-phase responses fall in the shaded region of 816 each dial. Markers are as used in (B-E). Open arrowhead (lower left) indicates a cell in which phase lag dominates the predicted achromatic response. Closed arrowhead (upper right) 817 818 indicates a cell in which response gain dominates the predicted achromatic response.

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Examples of observed receptive field organization for blue-on cells, corresponding to the 820 model types listed in table 1. A: Example classical "type II" spatially coextensive receptive 821 822 field for a blue-on cell at 2.8° eccentricity. B: S- and ML cone contrast response functions and measured achromatic contrast responses for the cell receptive field shown in (A). Error 823 bars, in some cases smaller than the data point, are ± std. error. Smooth curves show fits of 824 Eq. 7 to the data. Open circles are S cone responses; filled circles are ML cone responses. 825 826 Dashed curve shows the predicted achromatic response. C: Complex plane plot for the data 827 presented in (A) and (B), showing the amplitude and phase of the first-harmonic response. Smooth curve shows the amplitude and phase of the fitted model. D-F: Example simple 828 opponent blue-on cell at 0.7° eccentricity. Filled red circle shows the spatial extent of the 829 830 ML cone receptive field; open blue circle shows the spatial extent of the S cone receptive field. G-I: Example cell with centre-surround organization of S cone inputs at 4.8° 831 832 eccentricity. Dashed blue circle shows the spatial extent of the S cone surround. J-L: 833 Example cell with centre-surround organization of both S cone inputs and ML cone inputs at 3.9° eccentricity. Filled dashed circle shows the spatial extent of the ML cone surround. 834 835

Scatterplots and distributions of fitted difference-of-Gaussians equation parameters. 837 A: Scatterplot of fitted receptive field radius for S- and ML cone inputs to blue-on and 838 blue-off cells. Cells with classical "type II" coextensive receptive fields fall on the line of 839 unity, shown in grey. B: Scatterplot of fitted receptive field volumes for S- and ML cone 840 inputs for blue-on and blue-off cells. C: Scatterplot of centre and surround radius for S cone 841 input to blue-on cells, ML cone input to blue-on cells, P on cells, and P off cells. Marker 842 843 legend per (F). D: Distributions of radius ratios. From left to right: ratio of S subfield to ML 844 subfield radius for blue-on cells; ratio of ML centre radius to ML surround radius for P on cells, ratio of ML centre radius to ML surround radius for P off cells; ratio of S cone centre 845 radius to S cone surround radius for blue-on cells; ratio of ML centre radius to ML surround 846 radius for blue-on cells off cells. E: Scatterplot of centre and surround volume for S cone 847 input to blue-on cells, ML cone input to blue-on cells, P on cells, and P off cells. 848 849 F: Distributions of volume ratios, in the same format as panel D.

850

Α

В

Separable inputs



Linear inputs  $S \longrightarrow f \rightarrow f \rightarrow Response$  $ML \longrightarrow Response$ 





Figure 3

Figure 4









Е

Foveal >

n < ф D, <mark>6</mark>0  $\Box$ ► Peripheral imp/s



Figure 9



Α