Title

More than expected: extracellular waveforms and functional responses in monkey LGN

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Abstract

Unlike the exhaustive determination of cell types in the retina, key populations in the lateral geniculate nucleus of the thalamus (LGN) may have been missed. Here, we have begun to characterize the full range of extracellular neuronal responses in the LGN of awake monkeys using multi-electrodes during the presentation of colored noise visual stimuli to identify any previously overlooked signals. Extracellular spike waveforms of single units were classified into seven distinct classes, revealing previously unrecognized diversity: four negative-dominant classes that were narrow or broad, one triphasic class, and two positive-dominant classes. Based on their mapped receptive field (RF), these units were further categorized into either magnocellular (M), parvocellular (P), koniocellular (K), or non-RF (N). We found correlations between spike shape and mapped RF and response characteristics, with negative and narrow spiking waveform units predominantly associated with P and N RFs, and positive waveforms mostly linked to M RFs. Responses from positive waveforms exhibited shorter latencies, larger RF sizes, and were associated with larger eccentricities in the visual field than the other waveform classes. Additionally, N cells, those without an estimated RF, were consistently responsive to the visually presented mapping stimulus at a lower and more sustained rate than units with an RF. These findings suggest that the LGN cell population may be more diverse than previously believed.

Significance statement

This study uncovers evidence for an intricate diversity of neuronal responses within the lateral geniculate nucleus (LGN), challenging conventional classifications and revealing

previously overlooked populations. By characterizing extracellular spike waveforms and revising receptive field classifications, we provide novel insights into LGN function. Our findings have significant implications for understanding early visual processing mechanisms and interpreting extracellular signals in neural circuits. Furthermore, we identify non-receptive field units, prompting exploration into their functional roles and broader implications for visual and non-visual computations. This study not only advances our understanding of LGN organization but also highlights the importance of considering recording biases in electrophysiological studies. Overall, our work opens new avenues for interdisciplinary research and contributes to advancing our knowledge of neural dynamics in the visual system.

Introduction

The lateral geniculate nucleus (LGN) of the thalamus serves as a pivotal hub for processing visual information in mammals, acting as the primary relay station between the retina and the visual cortex (Sherman & Guillery, 2006; Solomon & Lennie, 2007). In primates, the LGN is anatomically segregated into distinct layers, namely magnocellular (M), parvocellular (P), and koniocellular (K). Electrophysiological studies in primates have delineated the functional characteristics of these layers: M neurons exhibit rapid temporal dynamics, well suited for motion detection; P neurons are sensitive to chromatic dynamics, well suited for color and form processing; and K neurons are responsive to short-wavelength ("blue") photoreceptor inputs (Wiesel & Hubel, 1966; Schiller & Malpeli, 1978; Kaplan & Shapley, 1982; Derrington & Lennie, 1984; Hubel & Livingstone, 1990; Maunsell *et al.*, 1999; Reid & Shapley, 2002; Tailby *et al.*, 2008).

Despite the extensive characterization of these divisions (with K neurons being

comparatively less studied), much of our understanding of LGN function stems from

electrophysiological recordings employing single-channel electrodes to discern neuronal

responses in the form of extracellular spikes. Typically, these signals are biphasic with a

dominantly negative voltage excursion, allowing for tentative classification of neuronal cell

types: broader waveforms are indicative of excitatory neurons, while narrower waveforms

suggest inhibitory neurons (Henze *et al.*, 2000). However, recent advancements in dense

multi-electrode arrays and sophisticated spike-sorting algorithms have unveiled a spectrum of

waveform shapes across various species and brain regions, diverging from the traditional

biphasic and negative waveforms.

These waveform variations include triphasic-spiking waveforms found in rat

hippocampus (Barry, 2015) and superior colliculus (Sibille *et al.*, 2022); positive-spiking and

long-broad negative waveforms found in the cat visual cortex (Gold *et al.*, 2009; Sun *et al.*,

2021) and human prefrontal cortex (Paulk *et al.*, 2022); and doublet-spiking waveforms

consisting of two short downward deflections found in ferret LGN (Murphy *et al.*, 2020).

Such discoveries challenge the conventional understanding of extracellular signals and

underscore the limitations of single-electrode sampling biases (Olshausen & Field, 2005;

Talebi & Baker, 2016), potentially leading to the oversight of crucial neuronal populations.

Identifying the gamut of extracellular signals from the LGN may help in understanding the

processing in the early visual pathway.

Thus, in this study, we agnostically survey the extracellular space in the LGN of

rhesus macaques through recordings that were not optimized for single-unit isolation,

employing a variety of stimulus and electrode properties. For each recorded LGN unit, we

characterize its receptive field (RF) and extracellular spike waveform classes, alongside

several response metrics, to elucidate potential relationships between spike shape and

neuronal identity. Our investigation also unveils a subset of units lacking an estimated RF

(termed non-RF, or N units), an occurrence rarely reported in existing literature, with most of

these units exhibiting consistent responsiveness to the visual stimulus at a lower and more

sustained response rate to units with an RF. The presence of N cells suggests a more nuanced

complexity in LGN processing beyond the classical M, P, and K divisions, and hints at

undiscovered computational mechanisms within the visual system. Our findings challenge the

conventional paradigm of LGN organization and underscore the necessity for a fuller

understanding of neural populations within this critical visual relay station. By delineating the

LGN's neural landscape beyond traditional classifications, our study contributes to advancing

knowledge of early visual processing mechanisms and the interpretation of extracellular

signals.

Methods

Ethics

Recordings were made from three awake adult rhesus monkeys (*Macaca mulatta*, 3M, 19–20 kg). The animals were maintained in the AAALAC accredited animal facility at the Massachusetts General Hospital. All research procedures were approved by the Massachusetts General Hospital Institutional Animal Care and Use Committee (IACUC) and were carried out according to the NIH Guide for the Care and Use of Laboratory Animals.

Animal preparation

The animals were surgically implanted with custom titanium head-holding posts and recording cylinders that allowed bilateral access to the LGN. Animals were trained to sit in a primate chair (B & M Plastics, Inc.), and placed in a shielded recording chamber (Crist Instrument Co.) during recordings. The animals were positioned so that their eyes were 43 cm from a 22-inch CRT monitor (Viewsonic P220f) with neutral gaze position near the center of the monitor.

Gaze location was monitored by an infrared video camera at a 500 Hz sampling rate (High-Speed Primate, SensoMotoric Instruments GmbH), with infrared illuminators stacked to yield a single corneal reflection. Software (iView X, SensoMotoric Instruments) was used to adjust the gains and offsets of the pupil-corneal reflection distance. Gaze position signals were then sent via analog signals to a behavioral control computer, where additional calibration was performed (2D quadratic fit to nine calibration points). See Figure 1A for a schematized depiction of the recording cylinder, monitor positioning and eye tracking.

Figure 1. Experimental design. **(A)** Animals were placed in front of a CRT monitor, with neutral gaze position near the center of the monitor. Gaze location was monitored through an infrared eye tracker. Multi-channel electrodes were inserted through surgically-implanted titanium recording cylinders which maintained chronic access to the dura. **(B)** The trial-based mapping task. The animal was required to fixate on a small spot in the center of the screen for 0.4 sec, before the mapping stimulus commenced, and to maintain fixation on the spot for the duration of the mapping stimulus. Mapping stimuli were presented for 2.5 to 5.0 seconds. **(C)** Schematic of extracellular spike waveform extraction. For each single unit (u), spike waveforms were extracted from the raw signal (W_u^k) in 4.5 ms windows and then averaged to obtain the mean waveform ($\overline{W_u}$).

Behavioral task

A trial-based task was used to map the response fields of neurons (Figure 1B). To begin each

trial, the animal was required to fixate on a centrally presented small, circular target (0.2°

diameter white spot) for 400 ms. The visual mapping stimulus then began with the fixation

point overlaid on the animated noise so that it continued to be visible, and the animal was

required to continue maintaining fixation within a 2.0° diameter circular window for the

duration of the stimulus (2.5–5.0 seconds, mode of 2.5 seconds; typical fixation performance

was much tighter than this window; see Results). Successful fixations through the entire

stimulus were rewarded with drops of sweet liquid and pleasant audio feedback. Data from

successful trials were used for mapping analysis (4–215 trials per experiment, median of 39

trials), but data from all trials were used for the spike waveform analysis.

Visual Stimuli

Custom software (PLECS, Pezaris Lab Experiment Control System) was used to display

noise stimuli that spanned both luminance and chrominance spaces, in addition to controlling

behavioral state and data collection (Killian *et al.*, 2016). The visual stimuli were presented

on a CRT display (Viewsonic P220f) at 160 Hz refresh rate and 8 bits per color channel on a

display area of 40 by 30 cm (400 by 300 pixels) that subtended 53° by 40° in visual field.

Two classes of noise stimuli were presented: high-resolution 400 by 300 pixel sets with

power spectra proportional to one over frequency (five sets, described below) and low-

resolution 80 by 60 pixels with a white power spectrum (one set). Each recording used a

single mapping stimulus from this collection.

The high-resolution noise stimuli were generated through computations in the

frequency domain:

$$\xi = \left(\xi_x, \xi_y, \xi_t\right) \tag{1}$$

$$170 \qquad\qquad\qquad S(\xi) = \frac{1}{\|\xi\|^\beta} \qquad\qquad\qquad (2)$$

$$171 \qquad\qquad\qquad |(Ff)(\xi)|^2 = S(\xi) \qquad\qquad\qquad (3)$$

$$172 \qquad\qquad\qquad Ff(\xi) = \left(S(\xi)\right)^{1/2} e^{-i2\pi\varphi} \qquad\qquad\qquad (4)$$

$$173 \qquad\qquad\qquad \varphi = rand[0,1] \qquad\qquad\qquad (5)$$

$$174 \qquad\qquad\qquad f(x,y,t) = F^{-1}(Ff)(x,y,t) \qquad\qquad\qquad (6)$$

where ξ is the spatiotemporal frequency vector represented in space (x, y) and time (t), $S(\cdot)$ is

the power spectra, β is a set variable of 0, 1.5, 2.5, 3.14, 3.5, or 4.0; F is the Fourier

transform; and $f(x,y,t)$ in the value of each pixel in space and time. Increasing values of β

decrease the slope of the power spectrum, where a β value of 0 yields a white spectrum (used

here for 80 by 60-pixel stimuli), and higher values of β yield more naturalistic spectra with

higher spatial and temporal correlations (Simoncelli & Olshausen, 2001). See Extended

Figure 1-1 for illustrated examples.

Electrode implantation

LGN recordings were made with acutely inserted electrodes (n = 99 recordings; 77

multichannel recordings, 22 single-channel recordings; three animals) or chronically

implanted electrodes (n = 12 recordings; one animal). Acute electrodes were either a custom

single linear 16-electrode array with 200 µm or 260 µm channel spacing, a linear 16-

electrode tetrode configuration array (four groups of four contacts) with 50 µm intra-group

spacing and 450 µm inter-group spacing (Plexon U-Probe), or a single traditional tungsten

electrode (FHC, Inc.). Chronic electrodes were custom 64-channel microwire bundles

designed to splay at depth.

Before acute penetration, the recording chamber was opened and cleaned, the

micromanipulator attached (Kopf Instruments Model 650), and used to advance the electrode

to the area of interest. Neural responses were recorded by a data acquisition system (Power

1401, Cambridge Electronic Design) and software (Spike 2, Cambridge Electronic Design)

that sampled at 40 kHz for all electrodes and channels, to 16-bit resolution. Unless stated

otherwise, the data presented in this project are from electrodes with multiple channels

(single-channel electrode data from traditional tungsten electrodes are shown only in Figures

4D and 5C).

Spike sorting

Extracellular recordings can have spikes from multiple nearby neurons recorded on the same

channel, and for multi-electrode channels, spikes from the same neuron may appear on

multiple channels. To distinguish spikes from different neurons, extracellular signals were

automatically sorted through KiloSort 2.5 (Pachitariu *et al.*, 2016). The output was then

manually curated with *phy* (Rossant *et al.*, 2016). During manual curation, clusters were

cleaned by drawing a boundary in the principal component analysis (PCA) space to remove

any abnormal waveforms. The clusters were then assigned a label of *good* or *noise. Good*

clusters showed evidence of a refractory period (longer than 1 ms) in the autocorrelogram

with a dip to approximately zero, and clear isolation from other clusters in PCA space.

Merges were made between *good* clusters if they had similar waveforms (overlapping PCAs),

a shared refractory period (a central dip in their crosscorrelogram similar to that from their

autocorrelograms), and possibly if there was evidence of drift (one cluster halts spiking when

the other starts at an adjacent channel). The lack of an autocorrelogram-like shape in the

crosscorrelogram would disqualify a merge. All *good* clusters were identified as single units

(SU). The remaining *noise* clusters were not included in additional analysis. See Extended

Figure 2-1 for spike sorting examples.

Extracellular spike waveform analysis

We will represent a single unit's extracellular signal by its mean extracellular spike

waveform, which is then analyzed and compared to the mean waveform of other single units.

To obtain the mean waveform of each single unit, u, up to 2,000 spike waveforms, k, were

extracted from the raw signal, randomly selected from identified firing times for that neuron

and collected as a set W_u^k (Figure 1C). Each waveform was extracted from a window of 60

samples (1.5 ms) before to 120 samples (3 ms) after the selected spike time; extracted single

waveforms were then averaged together to obtain the mean waveform $\overline{W_u}$. If a single unit

produced less than 2,000 spikes, then all spikes were used (this limitation occurred with 44%

of units). For multi-channel electrode recordings, all channels were sampled and averaged,

with the single channel with the largest mean spike waveform amplitude selected as the mean

waveform for each of those cells. Through the remainder of this report, we will use *spike*

waveform to refer to the mean extracellular waveform of a single unit, $\overline{W_u}$ from the channel

with the largest signature. The recording quality of single units was quantified by a signal-to-

noise ratio (SNR; Kelly et al., 2007); the population median of these values was 2.85 ($n =$

303).

Spike waveform classification

Mean waveforms, $\overline{W_u}$, were automatically classified using WaveMAP (Lee *et al.*, 2021),

which uses Uniform Manifold Approximation and Projection (UMAP; McInnes *et al.*, 2020)

for dimensionality reduction and the Louvain method for cluster identification (Blondel *et al.*,

2008). As in Lee *et al.* (2021), before WaveMAP classification, the mean waveform of each

single unit, $\overline{W_u}$, was normalized so that the maximum of its absolute value was 1.

Receptive Field Analysis

Spike-triggered averaging

Each single unit's chromospatiotemporal (CSTRF, or just RF) was estimated by computing a

spike-triggered average (STA; Schwartz *et al.*, 2006), which averages the visual stimulus

around each spike time by the following formula:

$$STA = \frac{1}{N}\sum_i s_i \qquad\qquad \textit{Eq. 1}$$

where s_i represents the stimulus in a fixed-size temporal window spanning the time of the ith

spike, and N is the total number of spikes. The STA was computed using 25 stimulus frames

before to 10 frames after each spike (i.e., time lags of –25 to 10 frames, or –156.25 ms to

62.5 ms in steps of 6.25 ms). The entire STA was then whitened by multiplying it with the

stimulus covariance matrix (Sharpee, 2013). The resulting chromospatiotemporal receptive

field is in four dimensions: time, position in the horizontal dimension, position in the vertical

dimension, and color by phosphor (red, R; green, G; blue, B).

RF position and latency

The position of each RF was determined by the location of the maximum excursion across the RF. The process of identifying RF position was performed iteratively: a first pass was used to find the absolute maximum; a second pass, to determine the background noise after clipping out a rectangular region 4 degrees on a side centered spatially at the maximum and spanning all time; and a third pass, to enforce statistical selection constraints based on z-score. The localization of the peak value was performed in luminance space after converting the CSTRF by taking the vector length of the RGB components for each pixel:

$$Lum_{x,y,t} = \sqrt{R_{x,y,t}^2 + G_{x,y,t}^2 + B_{x,y,t}^2} \qquad \text{Eq. 2}$$

$$RF_{location} = argmaxLum_{x,y,t} \qquad \text{Eq. 3}$$

where R, G, and B are the intensity of the colored phosphors red, green, and blue, respectively, and x, y, and t are locations in the CSTRF matrix in x-space, y-space, and time, respectively. The luminance matrix, Lum, was determined by the magnitude of the colored phosphors. From Lum, the 3-tuple representing the position of its maximum was recorded as the spatiotemporal location of the RF. Throughout this report, *RF position* refers to the x and y spatial components of $RF_{location}$, *RF latency* refers to its temporal component, and RF_{max} to the spatial plane at the latency point. Lum_{max} is the luminance value of RF_{max}.

RF size and RF spatial window

The RF size and spatial window were measured from the spatial envelope of Lum_{max}. First, a 2D Gaussian fit was applied to Lum_{max}, and then the length and width of the RF were defined as the full width at 60% maximum (i.e., one standard deviation) along the vertical

and horizontal directions, respectively, and corrected for cosine error. The RF size was

defined as the average of length and width, and the RF spatial window as the area length

times width. All data referring to RF size in this study are from receptive fields from which a

clean spatial envelope could be extracted when mapped using high-resolution stimuli.

RF temporal window

A temporal window was used to select frames around the peak by fitting a two-term Gaussian

model to $Lum_{x,y}$ at the RF position, through time. If the polarity of RF_{max} was *off*, pixels

were multiplied by -1 before fitting. Frames where this value at least one standard deviation

above the noise level and were of the same polarity as RF_{max} were considered the RF

temporal window.

Z-score

To evaluate the strength of each RF and impose statistical constraints, we computed its *z*-

score, its amplitude *a* divided by the noise level η. The amplitude was computed as an un-

normalized peak value by averaging the CSTRF pixels within the spatial and temporal

windows. This average value, an RGB triplet, was then transformed to luminance space to

obtain the RF amplitude *a*. The noise floor η was determined by the standard deviation of

amplitude computed in the same manner with an equivalent number of pixels, but taken

randomly from RF_{noise}, the acausal frames (frames after the spike, thus positive latencies) of

the CSTRF, and repeated without replacement until all pixels in RF_{noise} were used.

Normalizing the amplitude by the noise floor yielded the *z*-score for the RF: $z = a / \eta$.

Cone Weights Calculation

We used the method described in Horwitz & Albright (2005) to obtain the RF color weights

(RGB; red, green, and blue phosphor intensities, also known as *gun values*) and then

converted those values to cone weights (LMS; corresponding to sensitivities for long,

medium, and short wavelength cones). The color weights were obtained by computing the

mean STRF value within the spatial and temporal windows and then converting to cone

weights by the following formula:

$$[LMS] = A^{-T} * [RGB]^{T}$$

Eq. 4

where A^{-T} is the inverse transpose of the three-by-three transformation matrix, A. To obtain

the transformation matrix, RGB gun values from the CRT were computed for each color

channel in independent trials. Spectral radiance values were measured (PR650 SpectraScan

Colorimeter, Photo Research, Inc), and the transformation matrix was obtained from pairwise

inner products of the monitor spectral radiance values and 10° cone fundamentals (Stockman

& Sharpe, 2000). The LMS cone weights were then normalized by dividing each cone weight

by the sum of their absolute weights.

Cone weight noise

For each of the three cone weights in an RF, a corresponding noise level was measured to

determine significance. The noise levels were calculated analogously from the acausal pixels

presumed not to contain signal as described for η above, except, instead of transforming the

averaged pixels into luminance space, those pixels were transformed into cone space using

Equation 4. The noise threshold for each cone weight was then taken as one standard

deviation of the corresponding cone space values derived from acausal pixels.

Response metrics

PSTH responsiveness and response latency

The peristimulus time histogram (PSTH) was used to determine responsiveness and response

latency for each cell. The PSTH was computed during the time range 600 ms before stimulus

onset through the end of stimulus presentation, binned to 1 ms. Responsiveness was

determined by comparing bins that were 600–0 ms before to those 200–800 ms after stimulus

onset (paired Student's t-test, $p < 0.05$). When estimating the stimulus response latency, the

PSTH was smoothed by applying a running average over a sliding window of length 5 ms,

and the response latency was determined by the delay between stimulus onset and when the

firing rate crossed 50% between baseline and maximum value during stimulus presentation.

Burst index

To describe the firing patterns of LGN cells, we used a burst index which measured how

bursty (bunched together in time) or tonic (evenly spread out) a spike train is. Bursts were

defined by an inter-spike interval of greater than 100 ms, followed by a string of inter-spike

intervals of less than 4 ms. The burst index was then computed as the ratio of spikes

occurring in a burst over all spikes in the spike train (Sherman, 2001; Wang *et al.*, 2006).

Single Unit Screening

To remain agnostic about cellular responses while maintaining confidence that recorded cells were from within LGN, we used a screening criterion that started with each well-isolated cell for which an RF was found ($n = 228$; RF units) and then included all other well-isolated cells appeared on the same channel ($n = 75$; non-RF units). This screening method resulted in 303 units from 89 experiments spanning three animals.

Statistical Analysis

Several statistical analyses and significance comparisons are made throughout this study. All results are shown as the mean with standard deviation (SD), except for Figure 4B, which uses mean with standard error of the mean (SEM). Significance tests involving the mean were done with unpaired Student's *t*-tests unless stated otherwise. All statistical analyses were performed using MATLAB.

Results

Extracellular spike waveforms

A total of 303 single units (SUs) were recorded from 89 electrode penetrations in the LGN of three macaques (M_{CH}, $n = 104$; M_{ST}, $n = 51$; M_{VG}, $n = 148$) and were classified using WaveMAP (Lee *et al.*, 2021), yielding seven distinct spike waveform classifications. Four exhibited typical negative-spiking waveforms (Figure 2 left side): a dominant negative

deflection followed by a smaller and slower positive deflection. The differences between

these four groups are their width (i.e., trough-to-peak duration) and the magnitude of positive

deflection (trough-to-peak ratio). Negative-spiking waveforms are often classified into broad-

and narrow-spiking classes, which have been associated with excitatory and inhibitory

neurons, respectively (Barthó *et al.*, 2004). Therefore, we defined these four classes as

Narrow 1 (N1, $n = 39$, 13%, shown as dark blue in Figures 2 and 5), *Narrow 2* (N2, $n = 67$,

22%, purple), *Broad 1* (B1, $n = 49$, 16%, orange), and *Broad 2* (B2, $n = 19$, 6%, light blue).

389   In addition to the typical negative-spiking waveforms, a subset of units exhibited

atypical spike waveforms (visualized in the insets under the grouped waveforms within

Figure 2). For example, within the four negative-spiking classes, a small population were

doublet-spiking waveforms (8%, $n = 14/174$; see right insets of N2 and B2 in Figure 2),

where the spike waveform consisted of two distinct negative deflections (examined

qualitatively). These doublet-waveforms were predominantly observed in N2 units (67%, $n =$

$8/12$), with fewer instances in B1 (8%, $n = 1/12$, and B2 units (25%, $n = 3/12$), while no

occurrences were noted in N1 units. See Discussion for possible justifications of doublet

waveforms along with all other spike waveforms mentioned in this study.

398   The fifth waveform class, Triphasic (T, $n = 38$, 13%, green), was also negative

399 spiking but had the addition of an initial positive deflection preceding the primary, negative

400 deflection, and subsequent positive deflection. Even though we defined this group as

401 triphasic, not all units had three distinct phases when examined qualitatively (76%

402 qualitatively triphasic, $n = 29/38$; see Figure 2 insets for examples).

403   The remaining two classes were positive spiking, which we defined as *Positive 1* (P1,

n = 47, 16%, black) and *Positive 2* (P2, $n = 44$, 15%, red). We noticed two main differences

between these two positive classes. Firstly, P1 waveforms featured a larger negative

deflection than those from P2; secondly, P2 as a class had a higher proportion of waveforms

with two prominent positive peaks surrounding a negative deflection (i.e., M-shaped; see

examples in Figure 2 under the grouped waveforms). The P1 and P2 classes had 9% (*n* =

4/47) and 45% (*n* = 20/44) M-shaped positive spikes, respectively, when qualitatively

examined. These positive and triphasic-shaped waveforms, although less commonly reported,

have been linked to action potentials propagating through axonal fibers (Meeks *et al.*, 2005;

Gold *et al.*, 2006; Lewandowska *et al.*, 2015) and afferent fibers originating from preceding

processing areas (Barry, 2015; Sun *et al.*, 2021; Sibille *et al.*, 2022), i.e., from the retina in

our case here.

Figure 2. Extracellular mean spike waveforms are classified using WaveMAP (Lee *et al.*,

2021). In the middle, scatter plot visualization of each single unit's mean spike waveform in

UMAP space, where the horizontal and vertical axes represent the dimension-reduced UMAP

coefficients. The data points are colored by the seven waveform classifications from

WaveMAP: *Narrow 1* (N1, blue); *Narrow 2* (N2, purple); *Broad 1* (B1, orange); *Broad 2*

(B2, light blue); *Triphasic* (T, green); *Positive 1* (P1, black); and *Positive 2* (P2, red).

Surrounding the scatter plot are the mean spike waveform traces for each single unit (light

lines) with the group mean waveform overlayed on top (dark and bold lines), for each class of

waveforms. Negative-spiking groups are normalized to the negative peak, and positive-

spiking classes (P1 and P2) are normalized to the positive peak. The smaller individual

waveforms illustrated below the grouped waveforms are two examples to show the variation

within each class. Gray lines within the illustrated waveforms represent baseline voltage. All

waveforms were plotted within a 2.5 ms window. See Extended Figure 2-2 for illustrations of

all extracellular spike waveforms.

Receptive fields: RF and non-RF units

All 303 SUs were presented with chromatic noise visual stimuli (see Methods) to estimate

their chromospatiotemporal receptive field using spike-triggered averaging. SUs with

successful CSTRF, defined by a z-score of greater than three (see Extended Figure 3-1A for

z-score distribution), were categorized as *RF units* ($n = 228/303$, 75%), and nearly all

CSTRFs had a lower spatial variance than the gaze location ($n = 141/142$, 99%; Extended

Figure 3-1B). The remaining SUs without successful CSTRF mapping are termed *non-RF*

units or *N units* ($n = 75/303$, 25%). Both RF and N units were analyzed in this study.

Two stimulus resolutions, coarse and fine, were used during RF mapping (no cell was

presented with both). The first two rows in Figure 3 illustrate two example RFs, one from a

coarse stimulus (RF 1) and one from a fine stimulus (RF 2). Both examples feature an *on-*

center (responding to RGB increments) accompanied by an *off* rebound (responding to RGB

decrements). These example cells demonstrate how fine stimuli allowed measurements of

RFs with higher spatial resolution. However, we found that this advantage came with the cost

of a lower success rate at observing significant RFs than when using coarse stimuli (fine:

70%, n = 158/226; coarse: 91%, n = 70/77), likely due to the higher contrast in the coarse

stimuli than the fine stimuli. This is evident in the surround estimated in RF 1 at –31.25 ms

compared to the weak indication of a surround in RF 2. Since RF 1 was one of the few RFs

that had a clear surround in our measurements, we will concentrate on RF centers.

Figure 3. Five example chromospatiotemporal receptive fields (RFs), one per row, spanning

12 stimulus frames, one per 6.25 ms. RF 1 was measured using coarse-resolution stimuli, and

RF 2–5 were measured using fine-resolution stimuli. All plots span 2.0° of the visual field.

Bolded black outlines indicate the frame in an RF with the largest magnitude (RF_{max}). On the

right are the cone weights for each example, indicated as long- (L, magenta), medium- (M,

green), and short-cone wavelengths (S, blue) on the x-axis and the normalized response on

the y-axis. See Extended Figure 3-2 for the RF_{max} and cone weights for all RF units.

M, P, and K classifications of RF units

The 228 mapped RFs were classified into magnocellular (M), parvocellular (P), or

koniocellular (K) based on their relative responses to long-, middle- and short-wavelengths

(LMS cone weights; Eq. 4). Firstly, RFs were classified into koniocellular if the S-cone

weight exhibited the largest magnitude (K, $n = 21$, 9%; see RF 5 in Figure 3 for an example);

secondly, parvocellular if the L- and M-cone weights displayed opposing and significantly

above noise responses (P, $n = 69$, 30%; RF 3 and 4 in Figure 3 for examples; see Discussion

for justification of this definition of parvocellular response); and lastly, all remaining RFs

were classified as magnocellular (M, $n = 138$, 61%; RF 1 and 2 for examples), i.e., L- and M-

cone weights that were of equal sign. It is important to note that this LGN cell classification

is based on functional considerations, which are closely related to the anatomical

classifications documented in prior LGN studies (Wiesel & Hubel, 1966; Derrington *et al.*,

1984; White *et al.*, 1998; Reid & Shapley, 2002).

M, P, and K cone response characteristics

The normalized LMS cone weights of all 228 RFs are visualized in the diamond plot of

Figure 4A, a method employed in previous studies comparing LGN neuron cone input

(Derrington *et al.*, 1984; Reid & Shapley, 2002; Horwitz *et al.*, 2007). M RFs exhibited an

even distribution across the LM-on and LM-off planes (Fig 4., top-right and bottom-left

lines, respectively), indicating varying contributions from both L- and M-cones. In contrast, P

RFs are clustered around the midpoint along the L-on and M-off plane (bottom-right line) but

are more evenly spread along the L-off and M-on plane (top-left line). K RFs predominantly

displayed S-cone *on* responses ($n = 15/21$, indicated by the high proportion of circular purple

data points), and half of the K RFs exhibiting an opposing contribution from L- and M-cones

(i.e., S-on and LM-off, = 10/21) are clustered in the bottom-left quadrant of Figure 4A. The

observed 2.5:1 ratio of S-*on* to S-*off* RFs aligns with previous studies in marmoset LGN

(Martin & Lee, 2014; Pietersen *et al.*, 2014) and macaque retinal ganglion cells (De

Monasterio *et al.*, 1975; De Monasterio & Gouras, 1975).

The *temporal weighting function* (TWF), LMS response over time for the RF center

(Reid & Shapley, 2002), revealed distinct differences among RF classes. M and P RFs

exhibited contrasting responses in the L- and M-cone space (M were chromatic non-opponent

while P were chromatic opponent), aligning with their role in color vision, with both M and P

units showed minimal to no S-cone response (11% and 6% relative to the peak cone response,

respectively), also consistent with prior work in macaque LGN (Callaway, 2005) and retinal

ganglion cells (Sun *et al.*, 2006). Additionally, M units displayed two clear opposing phases

for computing temporal changes, while P units were monophasic for computing ordinary

intensity. The TWF of K RFs were monophasic and chromatic opponent, with a moderate

contribution from L- and M-cones (29% and 28% relative to peak, respectively).

Figure 4. Receptive field classification and analysis ($n = 228$). **(A)** Normalized cone weights

($|L| + |M| + |S| = 1$) for all RFs plotted on the LM plane. The horizontal and vertical axes

represent L- and M-cone weights, respectively. The magnitude of the S-cone weight is

represented as the distance from the diamond lines. The circular and diamond-shaped data

points with positive and negative S-cone weights, respectively. The data points colored in

green, light-blue, and purple represent magnocellular (M), parvocellular (P), and

koniocellular (K) RFs, respectively (also in subplots C and D). **(B)** Mean temporal weighting

functions (TWFs), describing the average cone response over time for M (top-left), P (top-

right), and K (bottom) RFs. The grey line indicates spike onset. For M and K RFs, weighting

functions were multiplied by –1 for *off* RFs, and for P RFs, responses were flipped if the

maximum L-cone response was negative. **(C)** Intensity visualization summarizing the mean

value of a response metric for each RF class separated into M, P, and K on the left and RF

and non-RF (N) units on the right. The area of the inner and outer circles represents the mean

and standard deviation, respectively. The notations *, ** and *** represent $p < 0.05$, $p < 0.01$

and $p < 0.001$ (unpaired t-test), respectively. Note that response latency (indicated by the

dagger symbol, †) was calculated differently for the left (from the RF_{max}) and right datasets

(from the PSTH). Since differences in stimulus statistics have been known to affect spiking

metrics (Almasi *et al.*, 2022; Sanchez *et al.*, 2023), the RF size, spike-rate, burst index, and

response latency mentioned in this and following sections were from units stimulated with

fine-stimulus only. |Latency| refers to the absolute response latency. **(D)** Percentage of units

recorded from multi-channel electrodes versus single-channel electrodes (left) and units

recorded with the fine stimuli versus coarse stimuli (right) for each RF class.

M, P, and K spiking characteristics
528

Within the RF classes, distinct response metrics differentiate between M, P, and K neurons.

First, within 5° of central vision ($n = 75$), the mean RF sizes of M (0.17° ± 0.06°) and P RFs

(0.18° ± 0.07°) were significantly smaller than K RFs (0.32° ± 0.06°; $p < 10^{-6}$). We also

observed that although the rate of activity of M, P, and K units was not significantly different

from each other ($p > 0.27$; M, 21.8 ± 15.6 spikes/s; P, 19.5 ± 13.3 spikes/s; K, 17.0 ± 17.1

spikes/s), the spiking activity of M units were more frequent in bursts (burst index of 0.046 ±

0.035) than the more tonic P (0.023 ± 0.023, $p < 10^{-4}$) and K units (0.021 ± 0.031, $p = 0.012$).

Not only did M units spend more time in bursts, but they also had shorter mean unsigned

response latency (31.8 ± 7.8 ms) than P (42.1 ± 11.8 ms, $p < 10^{-8}$) and K units (47.5 ± 5.2 ms,

$p < 10^{-10}$). These findings are supported by previous LGN studies regarding RF size and

response latency (Maunsell *et al.*, 1999; Pietersen *et al.*, 2014; Eiber *et al.*, 2018) but present

novel insights regarding burst spikes across RF classes (Ruiz *et al.*, 2006; Pietersen *et al.*,

2017).

542   We also found two unexpected differences between cells classified as M and P by

RFs. Due to their larger cell bodies, we expected M units to have had the largest recorded

waveform amplitude out of the three classes. This was not the case with the M units in the

current study ($p > 0.37$) (Figure 4C fifth row), likely because we did not optimize electrode

position for each cell, and thus units were recorded at varying distances from the electrode

tips (as amplitude is inversely proportional to source-to-site distance: Holt & Koch, 1999;

Gold *et al.*, 2006). Second, the mean estimated depths for M and K units were significantly

more superficial than P units ($p < 0.02$) (Figure 4C sixth row), contrary to expectations based

on LGN anatomy. It is important to note that the estimated depth is not a complete and

accurate measurement of somatic depth (see Discussion for further details).

Non-receptive field (N) units

A substantial portion (25%, $n = 75/303$) of our recorded population, termed N units, did not

exhibit measurable receptive fields despite being recorded alongside units with identifiable

RFs. These N units displayed significantly lower spiking activity than RF units (N, 7.3 ± 10.2

spikes/s; RF, 20.6 ± 15.0 spikes/s; $p = 10^{-9}$; right side of Fig. 4D), suggesting that that the

visual stimulus did not activate these units, however more than half of N units had a

significantly elevated firing rate during stimulus onset versus immediately prior (65% of N

units; paired t-test; see Methods; 82% for RF units). The lower spiking activity of N units

may be due in part to insufficient data for a significant receptive field to be estimated, as RF

quality as assessed by z-score was moderately correlated with number of spikes in RF cells (r

$= 0.58$).

Additionally, N units exhibited more sustained spiking activity than RF units (burst

index of; N, 0.015 ± 0.013; RF, 0.036 ± 0.033; $p = 10^{-6}$), which was moderately correlated

with RF z-score (r = 0.45), and were estimated at a depth more superficial than RF units (N,

50.9 ± 2.4 mm; RF, 51.6 ± 2.7 mm; $p = 0.049$), suggesting that anatomical location may be

relevant to the recording of non-RF units. For the remaining response metrics, N units had no

significant differences in response latency compared to RF units (N, 47.2 ± 20.4 ms; RF, 50.7

± 25.9 ms; $p = 0.53$; estimated from PSTHs: see Methods), and no significant differences in

waveform amplitude ($p = 0.80$), suggesting that the timing and quality of the spikes were

irrelevant to the recording of non-RF units.

We observed two experimental factors that may have influenced the N unit

population. By including the single-channel units that were initially excluded from the dataset

($n = 48$) because of potential technologically-driven sampling biases (Talebi & Baker, 2016),

N units were more commonly recorded with single-channel electrodes than multi-channel

electrodes (44%, $n = 21/48$, versus 25%, $n = 76/303$; Figure 4E left panel). This discrepancy

is likely due to a difference in electrode technology, as the single-channel electrodes used

were sharper and had higher impedance than the multi-channel electrodes, reinforcing our

initial concern to exclude single-channel units (see Discussion for further details). Moreover,

the proportion of N units recorded with coarse stimuli was much lower (9%, $n = 7/77$) than

with fine stimuli experiments (30%, $n = 68/226$; Figure 4D right panel), likely due to the

stimulus' larger pixel size and higher contrast: driving neurons more strongly and perhaps

driving a larger proportion of the population than the fine stimuli.

Overall, our findings reveal a population of non-RF units with distinct response

characteristics and dependencies on electrode type and stimulus type, shedding light on

previously unreported aspects of neural activity in the LGN.

Correlations

Correlating spike waveform classifications (N1, N2, B1, B2, T, P1, and P2), RF

classifications (M, P, K, and N), and various response characteristics (e.g., spike-rate,

response latency, RF size) revealed intriguing patterns and insights into LGN neural

processing. When examining spike waveform classifications alongside RF classifications,

several notable observations emerged: (*i*) the narrow negative-spiking classes (N1 and N2)

were predominantly associated with P RFs (36–46%) with approximately a 2-to-1 ratio to M

RFs for N1 units and a 1-to-1 ratio for N2 units; (*ii*) the broad negative-spiking classes (B1

and B2) had the largest proportion of N units (37–47%); and (*iii*) the positive-spiking classes

(P1 and P2) were mostly linked to M RFs (66–68%) with a 5-to-1 ratio to P RFs and the least

proportion of K RFs (0–4%). This heterogeneity indicates there may be functional differences

between waveform classes in the LGN but may also reflect sampling biases inherent in the

recording process (Towe & Harding, 1970). For example, the higher proportion of M RFs

from the positive-spiking units may indicate the presence of more axonal projections from the

deeper M layers (passing upward through the P layers as they flow toward the optic tract)

than the P and K projections in our recordings.

Figure 5. Correlations and analyses of spike waveform classes against RF classes and

response characteristics. **(A)** Bar plot of waveform class population to their RF type (M, P, K,

N) as number of units ($n = 303$). The percentage above each bar indicates the proportion of

each RF type within that waveform class. **(B)** Intensity visualization summarizing the mean

value of a response metric for each waveform class, like in Figure 4C. The area of the inner

and outer circles represents the mean and standard deviation, respectively. The notations *,

** and *** represent $p < 0.05$, $p < 0.01$ and $p < 0.001$ (unpaired *t*-test), respectively. RF

population for each dataset is stated below each tick label along the vertical axis. Spike-rate,

burst index, latency, and RF size were estimated from units presented with fine-stimuli. **(C)**

The percentage of units recorded from multi-channel versus single-channel electrodes for

each spike waveform class. The percentages on the right of each bar denote the proportion of

each waveform class, and the number above denotes the total number of units within that bar.

Further comparative analysis of response characteristics reinforced these findings.

(Figure 5B). The two positive-spiking classes (P1 and P2) responded to the visual stimuli

with a significantly shorter response latency than the four negative-spiking classes (N1, N2,

B1 and B2) ($p < 0.004$); the RF size of narrow negative-spiking classes (N1 and N2) were

significantly smaller than all other classes ($p < 0.03$); and positive and triphasic (P1, P2 and

T) classes had RFs at significantly larger eccentricities the negative-spiking classes ($p <$

0.003). These observations are particularly significant because the shorter response latencies

of the non-classical positive-spiking classes support the notion that these waveforms originate

from axons in the retinothalamic projection, and, if the recorded positive waveforms were

axonal, then it accounts for their higher eccentricities as eccentricity is positively correlated

with axon diameter (Walsh *et al.*, 2000). Therefore, our observations show that the non-

classical positive-spiking waveforms recorded in the LGN have distinct functional

characteristics to the classical negative-spiking classes, similar to previous studies (cat V1:

Sun *et al.*, 2021).

Moreover, we also noted two additional metrics that may be of interest (Figure 5B last

two rows): overall spike size (amplitude; trough-to-peak magnitude) and recording position

(estimated depth). N1 waveforms exhibited the largest mean amplitude of all other classes (p

< 0.001), signifying that N1 may represent recordings from large neurons or recordings of

cells that were near the electrode (Holt & Koch, 1999; Gold *et al.*, 2006). In terms of the

anatomical recording positions, three out of the four classical negative-spiking classes (N2,

B1, and B2) were recorded significantly deeper than the positive-spiking and triphasic classes

(T, P1, and P2) ($p < 0.035$), indicating differences in the anatomical origin of these

waveforms. The more superficial position of positive and triphasic units suggests they were

from thalamocortical or corticothalamic axonal fibers traveling to or from the LGN, with

thalamocortical projections being more frequent due to the significantly faster response

latencies of the positive-spiking groups.

Methodological factors, such as electrode type and stimulus type, also influenced the

distribution of waveform types (Figure 5C). Single-channel electrodes predominantly

recorded negative-spiking units (96%, $n = 46/48$), while multi-channel electrodes captured a

more diverse range of waveform types, perhaps due to the dissimilar electrode properties or

selection bias for strong signals when using single-channel electrodes as opposed to multi-

channel electrodes (Talebi & Baker, 2016). Additionally, coarse stimuli resulted in a higher

proportion of positive-spiking units compared to fine stimuli (46% versus 24%), highlighting

a dependency between stimulus type and recorded waveform types.

Discussion

Extracellular spike waveforms

Our investigation into extracellular spike waveforms within the LGN reveals a diverse array of classes, each with distinct characteristics. The negative-dominant waveform classes (N1, N2, B1, B2) exhibit variations in spike duration, attributed by the opening and closing of sodium and potassium channels, and occur when the recording contact is near the neuron's soma with minimal influence from other structures (Holt & Koch, 1999; Gold *et al.*, 2006). The variations is spike duration has been linked to neuronal type, with broad-spiking waveform associated with excitatory neurons and narrow-spiking associated with inhibitory neurons (Henze *et al.*, 2000; Barthó *et al.*, 2004; Sukman & Stark, 2022). We have not made this assumption for our dataset because excitatory cells have exhibited narrow-spiking extracellular waveforms (Vigneswaran *et al.*, 2011), and inhibitory interneurons in mouse cortex have exhibited broad-spiking waveforms, intracellularly (Gentet *et al.*, 2010).

The upward deflection in triphasic- and positive-spiking waveforms has been described as the mixed-ion capacitive current from surrounding sources for recordings distal to the soma as the spike propagates through an isolated unmyelinated axon (Clark & Plonsey, 1968; Raastad & Shepherd, 2003; Heinricher, 2004; Gold *et al.*, 2006; Lewandowska *et al.*, 2015). Furthermore, previous studies have shown that only triphasic- and positive-spiking waveforms were recorded from silenced brain regions (Barry, 2015; Sun *et al.*, 2021; Sibille *et al.*, 2022), demonstrating that these waveforms are of afferent fibers originating from previous processing areas. Our findings are consistent with these reports, as the positive waveforms (P1, P2), and to a lesser extent the triphasic waveforms (T), had shorter response

latencies than the negative-spiking classes, suggesting they reflect axonal fibers of passage of

retinal ganglion cells.

Although the waveform clustering algorithm used here provided well-defined

waveform groupings, there were multiple variations within each waveform class. Within the

negative classes, a small population were doublet-spiking waveforms that may reflect retinal

spikes impinging on LGN neurons (S-potentials: Sincich *et al.*, 2007), or intermediate LGN

neurons like those found in ferret LGN (Murphy *et al.*, 2020). Within the positive classes, we

noticed several M-shaped waveforms (double positive peaks), which could also be from S-

potentials or from distal neuronal activity such as from dendrites (Holt & Koch, 1999; Gold

et al., 2006). Due to their rarity, we did not investigate if these unusual waveforms

functionally differed from others. Such subpopulations are rarely reported in the literature

and indicate potential complexities in extracellular action potentials that warrant further

investigation.

RF classification

Our study revises the traditional RF classification method (Wiesel & Hubel, 1966; Reid &

Shapley, 2002), to account for experimental uncertainties: i.e., for a RF to be considered P,

both opposing L- and M-cones had to be significantly above noise. This refinement is

important because, when classifying our RFs using the traditional method (sign only), we

found several RFs that qualitatively appeared to be magnocellular, but would have been

labeled parvocellular by classical criteria due to L- or M-cone responses within the noise

floor, resulting in an inaccurate classification (see Extended Figure 3-2 for examples). Our

new classification is more robust in the face of experimental uncertainty while preserving the

functional definitions.

We reported above (Figure 4C) that magnocellular units (sorted spike trains leading to

M RFs) were unexpectedly found more superficial than parvocellular units (those leading to

P RFs), which is not what is traditionally held. Bearing in mind that we likely recorded

signals of both somatic and non-somatic origin, we found magnocellular units consisting of

both negative-spiking and positive-spiking units, and that these sub-classes differed in depth.

The positive-spiking magnocellular units were recorded at estimated depths significantly

more superficial to those with negative waveforms, suggesting that magnocellular units with

positive waveforms were axonal activity traversing through the LGN. In fact, when

considering only the negative-waveform magnocellular units, there were no significant

differences in depth to those that lead to P or K RFs.

Correlations

Distinct correlations between extracellular spike shape and RF characteristics were observed.

First, the narrow-spiking classes (N1 and N2) had a 1.5-to-1 ratio of P-to-M RFs, as often

seen in LGN extracellular recordings (Pietersen *et al.*, 2014). Even though this ratio is much

lower than the 8-to-1 ratio of the parvocellular-to-magnocellular population in the LGN

(Prasad & Galetta, 2011), the extracellular ratio may be due to a sampling bias of

extracellular recordings towards the larger magnocellular neurons ("the larger the fish, the

higher the probability of a hit" — Towe & Harding, 1970).

Second, positive-spiking waveforms (P1 and P2) had a 1-to-5 ratio of P-to-M RFs. If

we assume positive spikes are afferent axons from the retina, and the LGN has a 1-to-1 ratio

to retinal ganglion cell input (Spear et al., 1996), then why are the ratios of RF classes in the

positive and negative classes unequal? One might expect that negative and positive spikes are

functionally different, but we think that sampling biases were involved when recording from

axonal activity. In particular, factors including the generally larger diameter axons from

parasol/magnocellular neurons compared to those of midget/parvocellular and koniocellular

cells (Walsh *et al.*, 2000), the larger magnocellular-to-parvocellular ratio as eccentricity

increases (Livingstone & Hubel, 1988), differential degrees of axon myelination (Holt &

Koch, 1999), and the prevalence of magnocellular LGN axons in upper layers as they course

through the interior of the area toward the primary visual cortex, could have biased our

recording of axonal (positive spike) signals toward the magnocellular population.

Third, N RFs (those without an estimated RF) were found with higher proportion in

the broad-spiking waveform classes (B1 and B2) compared to all other waveform classes. We

observed that N RFs had many response characteristics that differed from units with an

estimated RF (discussed further below).

Non-RF units

The presence of N units raises intriguing questions regarding their identity and functional

significance. We found that almost two-thirds of N units responded reliably to the stimulus

albeit at a reduced spike and burst rate than RF units. The shortage in spiking activity could

be from the electrode channels losing contact with the source, which may explain the high

proportion of N units in the broad-spiking population, as contact distance is proportional to

spike waveform width (Gold *et al.*, 2006). The shortage of burst activity for N units may be

associated with a lack of stimulus contrast processing (Sanchez *et al.*, 2023), higher label

information processing (Butts *et al.*, 2010), or a lack of retinothalamic transmission of visual

information (Alitto *et al.*, 2019). Together, this evidence from the literature can suggest that

some of the N units may have been corticothalamic projections (evident in the more

superficial recorded depth), such as those that influence response gain (Murphy *et al.*, 2021).

For the remaining N units that did not reliably respond to the stimulus (43%), what do

these units do if they are not being visually stimulated? As here, Vries *et al.* (2020) found

23% of mouse visual cortex neurons (~60,000 total) did not reliably respond to visual stimuli,

with the proportion of non-responsive units increasing in higher visual areas. They speculated

that these units might be involved in specific natural features from hierarchical processing,

modulated by multimodal senses (Stringer *et al.*, 2019), or non-visual computation, such as

motion, identified in the dLGN of mice (Orlowska-Feuer *et al.*, 2022), which may be true for

our N units. It is also possible that these N units may explain the retinal ganglion cell classes

projecting to the LGN that are not associated with classical thalamic responses (~20%:

Dacey, 2004). While it may be simply that there was insufficient activity for a significant RF

to be recovered here, gaining insight into non-responsive units in future studies is integral to a

complete understanding of visual processing (Olshausen & Field, 2005).

Evidence for recording bias

Differences in populations between our multi-channel and single-channel recordings is likely

due to technologically-driven sampling bias. Talebi & Baker (2016) postulated that manually

controlled single-channel electrodes in combination with a search stimulus to identify

responsive neurons, largely avoided in our multi-channel recordings, creates a selection bias

toward user-preferred neurons. Similarly, previous studies utilizing single-channel electrodes

may have overlooked atypical positive-spiking units when searching for neurons, especially

when using a negative trigger threshold when searching for neurons or during off-line spike-

sorting.

It is important to note that the current study should be viewed as a partial survey of

the LGN, rather than an exhaustive one. We searched while listening for spikes along with

looking for them, often using a negative trigger threshold. This strategy may have created a

selection bias on the channel being monitored despite our attempts to mitigate those effects,

and the true proportion of positive-spiking waveforms may be higher than what we have

reported. For example, Paulk *et al.* (2022) recorded more positive-spiking than negative-

spiking units in human cortex using dense multi-channel electrodes. Several other

experimental biases are difficult to avoid in extracellular recordings, such as preference for

large extracellular action potentials, units with high firing rates, or units that are visually

responsive to the stimulus (Olshausen & Field, 2005). Thus, future studies need to consider

the possible biases when recording extracellularly to overcome the difficulties involved in

obtaining a completely objective study of a neuronal population.

Conclusion and implications

In conclusion, our study sheds light on the complexity of LGN neuronal populations,

challenging traditional classifications and highlighting the presence of previously

undocumented LGN units such as non-RF units. These findings expand our understanding of

LGN function and offer significant implications for future research, particularly elucidating

the intricacies of extracellular signals, addressing recording biases, and providing a deeper

understanding of receptive field data in the awake macaque LGN, for which there is limited

information at the high spatiotemporal resolution presented in this study. Furthermore, our

work opens avenues for interdisciplinary exploration, with potential applications in broader

neuroscience research (e.g., comparing function to principal cell types through

transcriptomics: Bakken *et al.*, 2021). Overall, our study contributes valuable insights into

LGN function and underscores the need for continued investigation into its neural dynamics.

SUPPLEMENTARY MATERIAL

SUPPLEMENTARY MATERIAL

Appendix S1: Glossary

SU, single unit: isolated extracellular recording from an individual neuron.

CSTRF, chromospatiotemporal receptive field: the receptive field of an LGN cell measured

across color and visual spaces, and through time.

RF, receptive field: the response of the optimal stimulus for a given cell, used

interchangeably with the CSTRF.

RF unit: a single unit with a significant receptive field.

STA, spike triggered average: the averaged stimulus image conditioned on spike time, here

used to calculate the CSTRF.

R, G, B; red, green, blue: the three phosphor colors used in computer monitors and, here, to

generate visual stimuli.

Lum, luminance matrix: determined by the magnitude of the colored phosphors.

$RF_{location}$: the location of CSTRF with the maximum, from *Lum*, in *x*-space, *y*-space, and

time.

RF position: the *x* and *y* spatial components of $RF_{location}$.

RF latency: the temporal component of $RF_{location}$.

RF_{max}: the spatial plane at RF latency.

Lum_{max}: the luminance matrix of RF_{max}.

RF_{noise}: the acausal frames of the CSTRF.

η: noise in a signal; used to select RFs with significant amplitudes.

L, M, S; long, medium, short: retinal photoreceptor cones classified by wavelength of peak

sensitivity.

M, magnocellular: RFs with contribution from *L*- and *M*-cone weights that were non-

opposing.

P, parvocellular: RFs with contribution from *L*- and *M*-cone weights that were opposing and

significantly above noise.

K, koniocellular: RFs with the largest contribution from *S*-cone.

N, non-RF unit: cells that do not have significant receptive fields.

TWF, temporal weighting function: the LMS response of a given pixel, or average pixel, over

time.

Narrow 1, 2: narrow, negative-going spike waveform classifications (see Fig. 2).

Broad 1, 2: broad, negative-going spike waveform classifications (see Fig. 2).

Triphasic: negative-going spike waveform classification but typically with an addition of an

initial positive deflection (see Fig. 2).

Positive 1, 2: positive-going spike waveform classifications (see Fig. 2).

Coarse stimulus: 80 x 60 colored noise stimulus.

Fine stimulus: 400 x 300 colored noise stimulus.

Extended Figures

Figure 1-1. Visualizing the range of spatiotemporal correlations created by increasing values

of β. (A) Example single frames from stimulus sets for β = 1.5, 2.0, 2.5, 3.0, 3.5, and 4.0. As

β increases, so does the spatiotemporal correlation. (B) Example frames from two

independent stimulus sets with β = 2.5. Each row depicts the central 2° for six consecutive

frames from a given set.

Figure 2-1. Spike sorting examples. **(A-C)** Three example single-channel recordings, each

with multiple isolated units. The top row illustrates the PCA feature space of the spike

waveforms (limited in this figure to 200 spikes). The grey cluster is noise. The second row is

the mean waveform of each single unit sorted on that recording channel. The third row is the

auto- and cross-correlograms between the same example units. **(D)** Five example single units

from additional recordings with only one single sorted unit, with the same information as in

A-C.

Figure 2-2. All extracellular spike waveforms plotted in **(A)** UMAP space, and **(B)** mean

spike waveform traces for each single unit with their corresponding UMAP coordinates. The

seven colors refer to the seven waveform classifications from WaveMAP.

868 **Figure 3-1.** Receptive field statistics. **(A)** Distribution of z-scores across all single units. The

869 histogram is binned in size of 0.5 z-score. The black and grey bins refer to RF and non-RF

870 units, respectively. **(B)** Standard deviation of RF size (°) plotted against standard deviation of

871 gaze position (°). The identity line is plotted in red. Nearly all CSTRFs had a lower spatial

872 variance than the gaze location.

Figure 3-2. RF_{max} and corresponding cone weight from all RF units. RFs are visualized in

RGB, and the three cone weights correspond to L, M, and S. The three groups of RFs are M

RFs (green border), P RFs (blue border), and K RFs (purple border). All RF plots span 2° in

visual space.

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