Awake responses suggest inefficient dense coding in the mouse retina

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Key words: retinal ganglion cells; in vivo recordings; awake; anesthesia; efficient coding.

Acronyms: CMOS, complementary metal-oxide semiconductor; DS, direction-selectivity; dLGN, dorsal lateral geniculate nucleus; DLP, digital light processing; FMM, fentanyl, medetomidine and midazolam; GABA, gamma-aminobutyric acid; LED, light emitting diode; OS, orientation-selectivity; OT, optic tract; PCA, principal component analysis; PFA, paraformaldehyde; RGC, retinal ganglion cells; SC, superior colliculus; STA, spike-triggered average; UV, ultraviolet.

Abstract

The structure and function of the vertebrate retina have been extensively studied across species with an isolated, *ex vivo* preparation. Retinal function *in vivo*, however, remains elusive, especially in awake animals. Here we performed single-unit extracellular recordings in the optic tract of head-fixed mice to compare the output of awake, anesthetized, and *ex vivo* retinas. While the visual response properties were overall similar across conditions, we found that awake retinal output had in general 1) faster kinetics with less variability in the response latencies; 2) a larger dynamic range; and 3) higher firing activity, by ~20 Hz on average, for both baseline and visually evoked responses. Our modeling analyses further showed that such awake response patterns convey comparable total information but less efficiently, and allow for a linear population decoder to perform significantly better than the anesthetized or *ex vivo* responses. These results highlight distinct retinal behavior in awake states, in particular suggesting that the retina employs dense coding *in vivo*, rather than sparse efficient coding as has been often assumed from *ex vivo* studies.

Introduction

The vertebrate retina is one of the best characterized parts of the central nervous system (Gollisch and Meister, 2010; Masland, 2012). It consists of ~100 cell types in total (Shekhar et al., 2016; Yan et al., 2020), including ~30 types of retinal ganglion cells (RGCs) that send the retinal output signals to the brain via the optic nerve (Sanes and Masland, 2015; Baden et al., 2016). Each of these RGC types forms distinct neural circuits within the retina to extract specific features of the visual image coming into the eye, such as color, contrast, and motion. The retina thus performs parallel and dynamic processing as the first stage of the visual pathway.

Most of our knowledge on retinal function has been obtained from *ex vivo* studies because isolated retinal tissues are nevertheless functional, i.e., responsive to light (Barlow et al., 1964). While powerful, however, *ex vivo* physiological approaches have certain limitations.

First, one cannot perform long-term recordings (Meister et al., 1994). Second, one cannot avoid artifacts due to retinal dissection, such as the effect of cutting RGC axons (Vidal-Sanz et al., 2017) or retinal epithelial detachment (Strauss, 2005). Neuromodulatory effects of retinopetal pathways are also difficult to study in an isolated retinal preparation (Repérant et al., 2006; Esposti et al., 2013). *In vivo* studies are thus indispensable for understanding to what degrees our knowledge on the retina can be translated from *ex vivo* to fully physiological *in vivo* conditions and clarifying retinal function thoroughly.

Previous physiological studies on the retina *in vivo* were conducted mostly under anesthetized—and often paralyzed—conditions. To monitor RGC activity *in vivo*, for example, single-unit recordings were made directly from the retina (Kuffler, 1953), at the optic nerve or tract fibers (Hartline, 1938; Lettvin et al., 1959; Enroth-Cugell and Robson, 1966; Mastronarde, 1983, 1985; Sagdullaev and McCall, 2005), or in the form of "slow-potential" in the dorsal lateral geniculate nucleus (dLGN; Bishop et al., 1962; Kaplan and Shapley, 1984). Optical methods were also developed to image the activity of retinal neurons directly through the pupil of a live animal under anesthesia (Geng et al., 2012; Yin et al., 2013, 2014). In contrast, thus far only a handful of studies have reported awake recordings from the retina (Esposti et al., 2013; Hong et al., 2018) or its outputs (Weyand, 2007; Liang et al., 2018, 2020; Schröder et al., 2020; Sibille et al., 2022). Thus, despite a long history of research on the retina, it still remains unclear what exactly the eye tells the brain in awake animals.

As recordings from awake behaving animals became routine for many brain areas (Dombeck et al., 2007; Jun et al., 2017), growing attention has been paid to the roles of an animal's behavior and internal brain states in the function of the sensory systems (Niell and Stryker, 2010; Lee and Dan, 2012). For example, systematic studies on the early visual pathway showed higher firing activity and faster response dynamics in both dLGN (Durand et al., 2016) and the superior colliculus (SC; De Franceschi and Solomon, 2018) of awake animals than those under anesthesia. A critical question that was left unanswered is to what extent such differences originate in the retina.

To clarify differences in the retinal visual response properties between awake and

anesthetized conditions, here we employed single-unit extracellular recording techniques from

head-fixed mice. Specifically, the electrodes were placed in the optic tract (OT), a bundle of

nerve fibers composed of RGC axons as they project from the eye to their main targets: dLGN

and SC (Ellis et al., 2016; Rosón et al., 2019). These recordings are superior to direct *in vivo*

retinal recordings (e.g., with epiretinally implanted mesh electrodes; Hong et al., 2018)

because the retinal circuits and the eye optics remain intact, and also to those extracellular

recordings in SC (Sibille et al., 2022) or dLGN (Weyand, 2007) because RGC axonal signals

do not need to be disambiguated from those of local axons or somata. This advantage also

exists for calcium imaging recordings of the retina directly in the eye of immobilized zebrafish

larvae (Esposti et al., 2013) or those of RGC axon terminals in dLGN (Liang et al., 2018, 2020)

or SC (Schröder et al., 2020; Molotkov et al., 2022). These imaging approaches, however,

lack the temporal precision as in the electrophysiology, which is an important aspect of the

information processing in the retina (Gollisch and Meister, 2008).

From our OT recordings, we examined the visual responses to a set of visual stimuli

widely used for probing retinal function *ex vivo* (Baden et al., 2016; Jouty et al., 2018),

including moving gratings, white-noise stimuli, and full-field flickering stimuli at different

temporal frequencies. We used two different anesthetics that are commonly used in

neuroscience research: isoflurane gas and an intraperitoneal combination of fentanyl,

medetomidine and midazolam (FMM). In both cases, we found that the temporal dynamics of

the retinal outputs were slower than in awake recordings, consistent with the previous studies

on the effects of anesthesia in retinorecipient areas (dLGN, Durand et al., 2016; SC, De

Franceschi and Solomon, 2018). We also found that the retinal outputs in an awake condition

had higher baseline firing rates and a larger dynamic range than in anesthetized or *ex vivo*

conditions. Consequently, while information was conveyed at a comparable rate in bits per

second, the amount of information per spike was lower in the awake responses. Furthermore,

we found that a linear decoder performed significantly better with awake population responses.

Thus, our results do not support sparse efficient coding (Attneave, 1954; Barlow and

Rosenblith, 1961) or minimum energy principles (Laughlin, 2001) that have been used as a

model of the early visual processing (Atick and Redlich, 1990; Gjorgjieva et al., 2019; but see

Schwartz, 2021). We instead suggest that the retina may employ dense coding principles *in*

vivo.

Results

Characterization of retinal output responses *in vivo*

To monitor retinal output *in vivo*, we established extracellular single-unit recording methods

from axons of retinal ganglion cells (RGCs) in the optic tract (OT) of head-fixed mice (Figure

1). In total, we made 17 chronic (75 cells) and 80 acute recordings (207 cells without

anesthesia; 325 cells with isoflurane; 103 cells with FMM), where a standardized set of visual

stimuli were presented to the subject animal to characterize the visual response properties of

the recorded cells (for ~1 hour; Figure 1A,B; see Methods for details). Isolated units typically

had a triphasic spike waveform (e.g., Figure 1F-H) as expected from axonal signals (Barry,

2015); and most of them had a minimum interspike interval above 2 ms (e.g., Figure 1I-K;

93%, awake; 94%, isoflurane; 99%, FMM). Histological verification of the electrode location

(e.g., Figure 1E) further ensured a successful optic tract recording *in vivo*.

Figure 1: *in vivo* extracellular recordings from the mouse optic tract. *A,B*: Schematic diagram of the experimental set-up. We presented visual stimuli to a head-fixed mouse using a digital light processing (DLP) device projecting images onto a spherical screen placed laterally to the subject animal (A, front-view; B, side-view). See Methods for details and specifications. *C,D*: Schematic diagram (C, top-view; D, side-view) of the brain and electrode location to target the optic tract (OT). *E*: Histological image of a representative brain sample (coronal section, 150 μm thick) showing the electrode trace (red, DiI stain deposited on the electrode). F-K: Spike waveform (F-H; black, individual trials; white, mean) and autocorrelogram (I-K; bin size, 1 ms) for three representative units recorded from the optic tract of an awake mouse.

109  We performed physiological classification of the recorded cells (Figure 2; see Methods

110 for details). Specifically, we first identified reliably responsive cells based on the visual

111 responses to full-field contrast-inverting stimuli (Figure 2A,B), and categorized their response

112 polarity into ON, ON/OFF, OFF types (Figure 2C). In both anesthetized and awake conditions,

113 we found around 20-50% of ON cells, 10-40% of ON/OFF cells, and 20% of OFF cells in our

114 data sets. We then further classified the cells from the viewpoint of orientation selectivity (OS)

115 and motion-direction selectivity (DS), based on the responses to moving gratings in eight

116 different directions (Figure 2D-F). In the anesthetized conditions, about a half of the cells (42-

117 54%) were either OS or DS or both regardless of their response polarities or anesthetics. This

118 is consistent with a previous *in vivo* study (Hong et al., 2018), but rather higher than one would

119 expect from anatomical and physiological studies *ex vivo* (about a third; Sanes and Masland,

120 2015; Baden et al., 2016; Jouty et al., 2018). In contrast, we found much less DS/OS cells

121 (17%) in the awake condition. This discrepancy in the fraction of DS/OS cells cannot be

122 explained by a sampling bias *per se*, if any, because we used the same experimental set up

123 between the anesthetized and awake recordings. Instead, it likely arose from a difference in

124 the baseline firing rate (see below and Figure 3). Since DS/OS index is defined as a

125 normalized population vector (Eq.(3). In Methods; Mazurek et al., 2014), higher firing in the

126 awake condition generally leads to a larger denominator, hence a lower DS/OS index value.

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Figure 2: Physiological classification of retinal output responses *in vivo*. *A*:
Representative retinal output responses to full-field contrast inverting stimuli: top, stimulus;
middle, raster graph over trials; bottom, peri-stimulus time histogram (black, mean; gray,
variance; signal-to-noise ratio (SNR), Eq.(1) in Methods; ON-OFF index, Eq.(2)). *B*: SNR of
the retinal output responses in different recording conditions. We set a threshold at 0.15 to
identify reliably responsive cells (black) and low-quality unclassifiable cells (gray). *C*: ON-OFF
index distributions from the reliably responsive cells. While no apparent clusters were
identified, we set a threshold at ±0.25 to categorize the response polarity into ON, ON/OFF,
and OFF cells. Within the ON cells, we further identified those with an "OFF-suppressive"
response to the full-field flickering stimulus (Figure 3). *D-F*: Distribution of DS/OS indices
(Eq.(3)) in each recording condition (D, awake; E, isoflurane; F, FMM). We set a threshold at
0.15 (with *p*<0.2) to identify whether cells are OS/DS (black) or not (gray). *G-I*: Fraction of
identified response classes *in vivo*: ON (blue; OFF-suppressive in green), OFF (red), ON/OFF
(orange), and the rest unclassifiable cells ("N/A", gray). Cells in each category were further
divided based on the OS/DS properties (hued). The OFF-suppressive ON cells were
prominent in the awake condition (G, 36/282 cells), but rarely observed under anesthesia (H,
isoflurane, 2/325 cells; I, FMM, 1/103 cells).

A distinguishing feature of awake recordings is an ongoing behavior of the animal

during recordings (Supplementary Figure 1). Behavioral states affect sensory processing (Neil

and Stryker, 2010), even at the early stages including the retina (Schröder et al., 2020). This

drove us to examine the effects of eye movements, pupil dynamics, as well as locomotion on

the retinal output *in vivo*.

First, we found that eye movements were not uniformly distributed in all directions

(Supplementary Figure 1H). Specifically, saccades were primarily in the horizontal directions

and much less in the vertical directions. Nevertheless, as reported previously (Samonds et al.,

2018; Miura and Scanziani, 2022), head-fixed awake mice did not frequently make eye

movements (0.06 ± 0.03 Hz; mean ± standard deviation, 19 animals) and thus their effects on

RGC activity were considered to be negligible in this study. The effects of microsaccades were

also negligible here as we mainly used full-field stimulation.

We next examined the relationship between pupil size and the retinal output dynamics.

They were both dependent on the light intensity level as evident by the pupillary reflexes

(Hatter et al., 2003) and the visual response polarities (Figure 2C), respectively. To analyze

their relationship, we thus focused on the recording period where the light intensity was

relatively stable over time (see Methods), and found that pupil size was correlated with RGC

firing in a response-polarity-dependent manner (Supplementary Figure 2A,B; Pearson's

R=0.38, p<0.001), likely mediated by the amount of light passing through the pupil.

Lastly, as reported previously (Schröder et al., 2020), we found that the activity of some

cells was significantly correlated with an animal's locomotion speed (Supplementary Figure

3C,D). While there was a positive correlation between pupil size and locomotion speed (0.34

± 0.20 from 19 animals; see also Erisken et al., 2014), unlike the pupil size effects, the

locomotion effects on the retinal output *in vivo* was not dependent on the visual response

polarity of the cells (Supplementary Figure 3D; Pearson's R=0.11, p=0.3). Thus, the

correlation between locomotion and RGC firing dynamics cannot be simply explained by the

light intensity fluctuation due to the associated pupil size dynamics.

Supplementary Figure 1: Eye motion and behavioral data analysis. *A*: Example eye-
tracking frame of an awake mouse eye. Cyan ellipse shows the pupil edge, while magenta dot
shows its center location. *B-G*: Representative dynamics of simultaneously recorded data in
an awake head-fixed condition (B, the presence and direction of the moving gratings stimulus;
C, firing rate heat map of simultaneously recorded seven RGCs; D and E, horizontal and
vertical coordinates of the pupil center location in each frame, respectively; F, pupil size
defined as a mean of the major and minor axis of the ellipse fitted to the pupil edge in each
frame; and G, running speed of the mouse). Red shades indicate eye blinks, whereas the blue
vertical lines indicate saccades. Note that blinks and saccades occurred only once in a while
(0.06 ± 0.03 Hz; mean ± standard deviation, 19 animals); and that the pupil size (F) is positively
correlated with the running speed (G) in general (0.34 ± 0.20; 19 animals). *H-J*: Probability
density of pupil center location (H), pupil size (I), and running speed (J) across different visual
stimulation periods (from left to right, full-field sinusoidally flickering stimuli; randomly flickering
full-field stimuli; and moving gratings; 19 animals).

Supplementary Figure 2: Retinal output was correlated with an animal's behavioral states. *A,B*: Peak cross-correlation values between retinal output and the animal's pupil size, plotted against the *p*-values calculated by a shuffling analysis (1,000 repetitions, see Methods; A) or the ON-OFF index (B): black, correlation with *p*<0.05; gray, correlation with *p*>0.05. The pupil size effects on RGC firing were significantly correlated with the ON-OFF index (B; Pearson's *R*=0.38, *p*<0.001). *C,D*: Corresponding plots for the peak cross-correlation values between retinal output and the animal's running speed. The locomotion effects on RGC firing were not significantly correlated with the ON-OFF index (D; Pearson's *R*=0.11, *p*=0.3).

Higher firing in awake than in anesthetized mice

For characterizing the retinal output properties *in vivo*, we first analyzed the responses to full-

field sinusoidally flickering stimuli that linearly changed the amplitude over time (from 0 to

100% contrast for 10 s; see Figure 3A-C for representative responses). We fitted to the

response an even-power of sine function, weighted with a sigmoid envelope (Eq.(6) in

Methods), and examined the curve-fit parameter values to analyze the response properties in

different recording conditions.

First, we found significantly higher baseline firing rates in awake mice ($B = 20 \pm 27$

Hz; median ± interquartile range) than in those under anesthesia (isoflurane, 2 ± 6 Hz, $p <$

0.001, *U*-test; FMM, 2 ± 4 Hz, $p < 0.001$; Figure 3D). Awake ON cells had particularly high

baseline activity (up to ~100 Hz) and showed a prominent reduction in firing rates in response

to light decrements (see Figure 3B for example). In contrast, due to this high baseline activity,

many of them showed virtually no responses to light increments ($N = 36$ out of 84), except for

a sharp rebound response to a full-contrast inversion ($N = 26$ out of 36; see Figures 2A and

3B for example). These ON cells had negative amplitudes in the curve fit (Figure 3E, green)

with the same response phase as OFF cells (Figure 3F; see Figure 3A for a representative

OFF cell's responses). This indicates that they primarily encode light decrements by

decreasing firing from their high baseline firing rates (62 ± 32 Hz, mean ± standard deviation),

rather than light increments by increasing firing as conventional ON cells do (Masland, 2012;

Sanes and Masland, 2015; Baden et al., 2016; Jouty et al., 2018). We thus categorized these

cells independently as an "OFF-suppressive" class (Figure 2G). This is likely an emerging

property of the cells due to high baseline firing in awake mice. Indeed, such responses were

barely observed in anesthetized animals (isoflurane, 2/325 cells; FMM, 1/103 cells; Figure

2H,I) where the baseline firing rate was generally low (Figure 3D). As expected from previous

ex vivo studies (Masland, 2012; Sanes and Masland, 2015; Baden et al., 2016; Jouty et al.,

2018), anesthetized ON cells increased firing upon light increments (see Figure 3C for a

representative response). Positive amplitudes in the curve fit (Figure 3E) with an opposite

response phase from OFF cells (Figure 3F) further support that these ON cells encode light

increments. These data suggest that the retinal coding depends on the baseline firing level of

RGCs.

Second, we found that the awake responses had a larger dynamic range than those

under anesthesia. Specifically, the absolute amplitude of the evoked responses from the

baseline was larger for the awake condition (Figure 3E; awake, $|A| = 45 \pm 55$ Hz, median ±

interquartile range; isoflurane, 30 ± 31 Hz, *p*<0.001, Mann-Whitney *U*-test against awake data;

FMM, 27 ± 33 Hz; *p*<0.001). Moreover, the sigmoid envelope had a higher midpoint (awake,

$t_0 = 70 \pm 61$ % contrast; isoflurane, 45 ± 32 % contrast, *p*<0.001; FMM, 45 ± 44 % contrast,

p<0.001) and a lower steepness (awake, $\lambda = 0.44 \pm 0.53$ /s; isoflurane, 0.67 ± 0.79 /s,

p<0.001; FMM, 0.69 ± 0.62 /s, *p*<0.001) for the awake responses than for the anesthetized

ones.

We next examined the sensitivity to contrast. At low stimulus contrast, we found a

higher sensitivity in the awake condition (estimated evoked response at 10% contrast = 7 ±

13 Hz, median ± interquartile range) than under anesthesia (isoflurane, 4 ± 12 Hz, *p*<0.001,

Mann-Whitney *U*-test; FMM, 3 ± 10 Hz, *p*<0.001). However, the maximum sensitivity to a

change in contrast – i.e., the slope size at the midpoint of the sigmoid – was comparable

between the awake and anesthetized conditions ($|A\lambda/4| = 5 \pm 11$ Hz / %contrast; isoflurane,

6 ± 9 Hz / %contrast; FMM, 5 ± 7 Hz / %contrast; p = 0.1, Kruskal-Wallis test).

We also found a significant phase shift in the responses between the recording

conditions (Figure 3F). This phase parameter in the curve fit represents the position of the

response peak relative to the sinusoidal stimulus intensity patterns (see Methods for details),

and we identified two clusters in each data set: the one with negative phase for ON cells

(Figure 3F, blue) and the other with positive phase for OFF and OFF-suppressive ON cells

(Figure 3F, red and green, respectively). ON/OFF cells were found in either cluster, depending

on the relative strength of their responses to light increments versus light decrements (Figure

3F, orange). In both clusters, we found that the phase was on average smallest for the awake

condition (-1.9 ± 0.6 and 1.7 ± 0.6 radian, respectively; mean ± standard deviation; isoflurane,

-0.9 ± 0.6 and 2.0 ± 0.5 radian; $p < 0.001$ for both cases, *t*-test), while largest for the FMM

anesthesia (-0.5 ± 0.8 and 2.3 ± 0.5 radian; *p*=0.007 and *p*=0.044 against corresponding

isoflurane data, respectively). This implies that the response dynamics are faster in awake

mice than those under anesthesia, especially with FMM.

Finally, we analyzed the retinal output properties between different visual response

polarities by combining data across recording conditions (ON, 199 cells in total; OFF, 121

cells; and ON/OFF, 170 cells). We found that the baseline firing rate was higher for ON/OFF

cells (17 ± 24 Hz; median ± interquartile range) than for ON or OFF cells (5 ± 24 and 6 ± 10

Hz, respectively; *p*<0.001, Kruskal-Wallis test). Otherwise we did not find any statistically

significant dependency on the response polarity at the population level.

Figure 3: Many ON retinal ganglion cells showed suppressive OFF responses in awake condition. *A-C*: Mean firing rate of representative cells in response to a sinusoidally-flickering stimulus with increasing contrast in the awake (A, OFF; B, OFF-suppressive ON, the same cell as in Figure 2A) or anesthetized conditions (C, ON). Overlaid with the peri-stimulus time histogram (gray) is the model fit (black, Eq.(6) in Methods). The number on top indicates the fit quality (explained variance R^2 in Eq.(5)). *D-F*: population data of the model parameters (D, baseline B; E, amplitude A; F, phase ϕ) across different conditions: isoflurane anesthesia ($N = 147$), FMM anesthesia ($N = 95$) and awake ($N = 247$ in total). The sinusoidal stimulus pattern relative to the response peak is also indicated at the bottom of F. Cell types are color-coded as in Figure 2 (blue, ON; green, OFF-suppressive ON; red, OFF; orange, ON/OFF). Note high baseline with negative amplitude and positive phase for the OFF-suppressive ON cells, which were predominantly found in the awake condition: ***, $p < 0.001$; **, $p < 0.01$; *, $p < 0.05$; ns, non-significant (D, *U*-test; E, *U*-test on the absolute values; F, *t*-test).

Faster response kinetics in awake than in anesthetized mice

To better characterize the kinetics of retinal output *in vivo*, we next analyzed the responses to full-field sinusoidal flickers at different temporal frequencies (Figure 4). In awake mice, most cells responded faithfully to all the stimulation frequencies we tested from 1.875 to 15 Hz (see Figure 4A for example). Some awake cells showed the largest responses to the 15 Hz stimulation (Figure 4A; curve fit quality with Eq. (4)), suggesting that they were possibly tuned to even higher frequencies. In the anesthetized conditions, in contrast, retinal output responses typically followed the stimulation frequencies up to 7.5 Hz, but failed to do so at 15 Hz (see Figure 4B for example). For quantification, we fitted an even-power of sine function to the responses (Eq.(4) in Methods) and set a threshold of the curve fit quality (defined as the explained variance; Eq.(5)) at 0.2 to select cells with robust responses (Figure 4C,D). In the awake condition, a larger number of cells responded well at medium frequencies (3.75 Hz, 63%; 7.5Hz, 71%) than at a low frequency (1.875 Hz, 59%); and the majority of the awake cells remained responsive to a high frequency stimulus (15 Hz, 47%). Under anesthesia, in contrast, the fraction of responsive cells was the largest at a low frequency (1.875 Hz; isoflurane, 77%; FMM, 67%). Fewer cells responded robustly at faster flicker rates (3.75 Hz and 7.5 Hz; isoflurane, 69% and 67%, respectively; FMM, 63% and 59%), and only a small fraction of the anesthetized cells were able to follow the stimulation frequency at 15 Hz (isoflurane, 13%; FMM, 3%). Consistent with the analysis of the response phase (Figure 3), these outcomes indicate that the retinal output dynamics are faster in awake animals than those under anesthesia, especially for FMM.

Across the response polarities, we found that ON/OFF cells (N=170 in total) were most responsive at a medium frequency (7.5Hz, 59%), while ON or OFF cells (N=199 and 121, respectively) were generally responsive from low to medium frequencies (1.875-7.5 Hz; 68-73% and 79-82%, respectively). Indeed, the fraction of responsive cells under our criteria was significantly lower for ON/OFF cells (1.875 Hz, 46%; 3.75 Hz, 49%) than for ON or OFF cells at low frequencies (*p*<0.001 for both frequencies, two-proportion *z*-test). No dependency on

302   the response polarity was found for a high frequency stimulation (15 Hz; ON, 25%; OFF, 29%;

303   ON/OFF, 31%).

Figure 4: Retinal output showed higher temporal frequency sensitivity in awake than in
anesthetized mice. *A*: Representative retinal output (gray, mean firing rate over 10 trials) in
an awake condition in response to full-field sinusoidally flickering stimuli at different temporal
frequencies (1.875, 3.75, 7.5 and 15 Hz), following full-field contrast inversions. Overlaid is
the curve fit (Eq.(4) in Methods; black). The number on top is the explained variance of the
curve fit (R^2, Eq. (5) in Methods), representing the fit quality. *B*: Representative retinal output
responses under isoflurane anesthesia (shown in the same format as in A). *C*: Population data
of the fit quality at four different stimulus frequencies in the awake ($N = 248$) or anesthetized
conditions (isoflurane, $N = 147$; FMM, $N = 95$), color-coded for the responses class as in
Figure 2. The fit quality threshold was set to be 0.2 (black, $R^2 \geq 0.2$; gray, $R^2 < 0.2$). *D*:
Fraction of the cells with the fit quality above the threshold across different conditions (awake,
black line with circles; isoflurane, gray line with vertical crosses; FMM, gray line with diagonal
crosses), representing the frequency tuning of the retinal output at the population level. A
significantly larger fraction of cells was responsive at 15 Hz in the awake condition than in the
anesthetized conditions (***, $p < 0.001$ for both isoflurane and FMM; two-proportion *z*-test).

Comparison of retinal output properties between *in vivo* and *ex vivo*

We have thus far focused on the retinal output properties *in vivo*, and showed higher baseline firing rates (Figure 3) and faster response kinetics (Figure 4) in awake mice than in those under anesthesia. We next asked if RGC responses differ between *in vivo* and *ex vivo* conditions. This is a critical comparison because retinal physiology has been mostly studied and best characterized in an isolated preparation (Gollisch and Meister, 2010; Sanes and Masland, 2015), but little is known about it in awake animals (Weyand, 2007; Liang et al., 2018, 2020; Schröder et al., 2020; Sibille et al., 2022). To what extent can we translate our knowledge on the retina from *ex vivo* to *in vivo*? Here we exploited stimulus ensemble statistical techniques ("reverse correlation"; Meister et al., 1994; Chichilnisky, 2001) to systematically characterize the visual response properties and make a direct comparison across different recording conditions in the linear-nonlinear cascade model framework (see Methods for details). Specifically, using full-field white-noise stimuli, we analyzed 1) the linear temporal filter (Figure 5), estimated by the spike-triggered average (STA) stimulus, i.e., the mean stimulus that triggered spiking responses; and 2) the static nonlinear gain function (Figure 6), i.e., an instantaneous mapping of the STA output to the neural responses (Eq. (7) in Methods). Here we reanalyzed the existing data sets for *ex vivo* recordings (696 cells from 18 isolated mouse retinas recorded at 37 °C; Vlasiuk and Asari, 2021).

Faster response kinetics in awake condition than *ex vivo*

We identified a good quality STA in more than two-thirds of the cells recorded *in vivo* (e.g., Figure 5A). To compare the temporal dynamics of the STAs across different recording conditions, we used the following two measures: 1) the first peak latency of the STA estimated from a difference-of-Gaussian curve fit (Figure 5A); and 2) spectral peak frequency calculated by the Fourier transform of the fitted curve (Figure 5B). Consistent with the results measured by the flickering stimuli at different frequencies (Figure 4), we found faster kinetics in awake

animals than in anesthetized ones (Figure 5C,D): i.e., significantly shorter peak latencies

(awake, 47 ± 18 ms, mean ± standard deviation; isoflurane, 77 ± 19 ms, $p < 0.001$, *t*-test;

FMM, 87 ± 24 ms, $p < 0.001$; Figure 5C) and higher peak frequencies (awake, 8.5 ± 2.9 Hz;

isoflurane, 4.6 ± 1.6 Hz, $p < 0.001$; FMM, 3.7 ± 1.2 Hz, $p < 0.001$; Figure 5D). Interestingly,

we identified that the *ex vivo* data lied in-between. The *ex vivo* STAs had significantly longer

peak latencies (64 ± 28 ms, $p < 0.001$) and lower peak frequencies (4.3 ± 1.8 Hz, $p < 0.001$)

than the awake ones. In contrast, the peak latencies were significantly shorter in an isolated

retinal preparation than in the anesthetized conditions (isoflurane, $p < 0.001$; FMM, $p < 0.001$),

while the peak frequencies were comparable between these conditions (isoflurane, $p = 0.3$;

FMM, $p = 0.052$).

We further analyzed the STA dynamics across different cell classes. For clustering the

STAs obtained in each recording condition, we used the principal component analysis (PCA;

see Methods for details). As in previous *ex vivo* studies (Gollisch and Meister, 2008; Asari and

Meister, 2014), the first two principal components accounted for most of the variance (78-86%,

collectively), and the four quadrants of the PCA biplot generally represented distinct filter

shapes, corresponding to monophasic OFF (mono-OFF), biphasic OFF (bi-OFF), monophasic

ON (mono-ON), and biphasic ON (bi-ON) response classes, respectively. In all recording

conditions, no apparent cluster was found in this feature space, leading to a continuum of the

STA shape patterns across populations (Figure 5E-H). Nevertheless, we identified two

features that were different between *ex vivo* and *in vivo*, especially distinct in the awake

condition. First, while the monophasic cells were generally slower than the biphasic ones,

differences in the peak latencies were much larger in the *ex vivo* (Figure 5G; mono-ON $83 \pm$

24 ms versus bi-ON 58 ± 31 ms, $p < 0.001$, *t*-test; mono-OFF 67 ± 29 ms versus bi-OFF $48 \pm$

8 ms, $p < 0.001$) or anesthetized conditions (Figure 5E, isoflurane: mono-ON 91 ± 22 ms, bi-

ON 67 ± 6 ms, $p < 0.001$; mono-OFF 89 ± 14 ms, bi-OFF 68 ± 8 ms, $p < 0.001$; Figure 5F,

FMM: mono-ON 97 ± 30 ms, bi-ON 78 ± 11 ms, $p = 0.051$; mono-OFF 103 ± 13 ms, bi-OFF

70 ± 11 ms, $p < 0.001$) than in the awake condition (Figure 5H; mono-ON 51 ± 20 ms, bi-ON

44 ± 19 ms, $p = 0.6$; mon-OFF 51 ± 19 ms, bi-OFF 43 ± 12 ms, $p = 0.1$). Second, we found

that ON cells were significantly slower than OFF cells *ex vivo* (mono-ON versus mono-OFF,

$p = 0.002$; bi-ON versus bi-OFF, $p = 0.03$), but not *in vivo* ($p > 0.2$ in all the conditions

examined). Taken together, our results suggest that awake responses are faster and their

kinetics are more similar across different cel classes than *ex vivo* or anesthetized responses.

Figure 5: Retinal ganglion cells showed faster response dynamics in awake condition
than in anesthetized or *ex vivo* conditions. *A*: Temporal filter of a representative awake cell
(gray, spike-triggered average (STA) of the full-field randomly flickering stimulus) and a
difference-of-Gaussian curve fit (black) for estimating the latency of the first peak. *B*: Power
spectra of the example filter in A, based on the curve fit, for estimating the peak frequency.
C,D: Population data of the peak latencies (C) and frequencies (D) across different conditions
(light green, biphasic ON; dark green, monophasic ON; pink, biphasic OFF; violet, monophasic
OFF). Here and thereafter, ***, $p < 0.001$; **, $p < 0.01$; *, $p < 0.05$; ns, non-significant (*t*-test).
The filter types were identified by the quadrants of the PCA biplot (see Methods for details).
E-H: Population data of the temporal filters across different conditions: from top to bottom,
isoflurane anesthesia (E, $N = 238$), FMM anesthesia (F, $N = 69$), *ex vivo* (G, $N = 342$) and
awake (H, $N = 201$). The four filter types are indicated on the right with corresponding colors.

Higher spiking activity in awake condition than *ex vivo*

How does a recording condition affect the firing properties of RGCs? The mean firing rate
during the white-noise stimulus presentation was significantly higher in awake (34.9 ± 24.6 Hz;
mean ± standard deviation) than in all the other recording conditions (isoflurane, 13.0 ± 13.1
Hz; FMM, 10.9 ± 10.7 Hz; *ex vivo*, 6.9 ± 6.6 Hz; all with $p < 0.001$, *t*-test in the logarithmic
scale; Figure 6A). Importantly, this is presumably not due to differences in the stimulus
condition. The maximum projected light intensity was comparable between *in vivo* (~31
mW/m^2 at the eye; see Methods) and *ex vivo* setups (~36 mW/m^2 on the isolated retina;
Vlasiuk and Asari, 2021). Due to absorption by the eye optics, however, photon flux at the
retina was estimated to be ~10 times lower for *in vivo* than for *ex vivo* recordings (on the order
of 10^3 and 10^4 R*/photoreceptor/s, respectively; see Methods). Furthermore, awake mice
show pupillary reflexes (Supplementary Figure 1), leading to a smaller amount of light
impinging on the retina than under anesthesia. Nevertheless, these differences in the lighting
conditions cannot explain the higher firing and faster kinetics in the awake cells (Figure 6A
and Figure 5) because RGCs typically show larger and faster responses to brighter and
higher-contrast stimuli (Baccus and Meister, 2002; Wang et al., 2011).

For a further analysis, we examined the static nonlinear gain function of the recorded

cells (Eq. (7) in Methods). This gain function accounts for nonlinearity associated with spike

generation, such as spike threshold and firing rate saturation, and is generally well explained

by a sigmoid function (see Figure 6B for example). For quantification, we assessed the neutral

firing rate of the cells where the input to the gain function is zero, i.e., the firing rate in the

presence of "neutral" stimuli that are orthogonal to the cell's STA. Consistent with the results

on the baseline firing rates measured by the sinusoidally flickering stimuli (Figure 3), the

neutral firing rate was significantly higher in the awake condition (29.3 ± 22.5 Hz, mean $\pm$

standard deviation) than those under anesthesia (isoflurane, 9.5 ± 11.9 Hz, $p < 0.001$, *t*-test

on a log scale; FMM, 7.5 ± 9.3 Hz, $p < 0.001$; Figure 6C). We also found that the neutral firing

rates *ex vivo* (5.5 ± 7.1 Hz) were as low as those in the anesthetized conditions (isoflurane,

$p = 0.02$; FMM, $p = 0.1$), and significantly lower than those in the awake condition ($p < 0.001$).

We further identified two distinct features in the gain function properties between *ex*

vivo and *in vivo* responses. First, ON cells generally had a higher gain than OFF cells *ex vivo*

(Figure 6F), whereas such cell-type specific differences were not observed *in vivo* (isoflurane,

Figure 6D; FMM, Figure 6E; awake, Figure 6G). Second, the midpoint of the sigmoid fitted to

ex vivo responses (1.6 ± 1.4, median $\pm$ interquartile range) was significantly higher than that

for *in vivo* responses (isoflurane, 1.1 ± 0.9; FMM, 1.1 ± 0.8; awake, 0.8 ± 1.0; all with $p <$

0.001, Mann-Whitney *U*-test; Figure 6H). In agreement with the outcomes obtained with the

sinusoidally flickering stimulus (Figure 3), this suggests that *in vivo* responses are more

linearized to cover a larger dynamic range, especially in the awake condition due to high

neutral responses (Figure 6G). Indeed, while certain batch effects existed across recording

sessions (Supplementary Figure 3), the dynamic range computed from the gain function (see

Methods) was generally largest in the awake condition (56 ± 54 Hz; median $\pm$ interquartile

range) and significantly larger for *in vivo* (isoflurane, 26 ± 30 Hz; FMM, 29 ± 27 Hz) than for

ex vivo (12 ± 12 Hz, all with $p < 0.001$, Mann–Whitney *U*-test).

Figure 6: Retinal ganglion cells showed higher firing activity in awake condition than in anesthetized or *ex vivo* conditions. Cell classes are color-coded as in Figure 5 (light green, biphasic ON; dark green, monophasic ON; pink, biphasic OFF; violet, monophasic OFF). *A*: Population data of the mean firing rates during the stimulus presentation period in four different recording conditions (isoflurane, FMM, *ex vivo*, and awake): ***, $p < 0.001$; **, $p < 0.01$; *, $p < 0.05$; ns, non-significant (*t*-test on the logarithm of firing rates). *B*: Static nonlinear gain function of a representative awake cell (the same one as in Figure 5A), estimated by the stimulus ensemble statistical techniques applied to the responses to a full-field randomly flickering stimulus (gray, Eq.(7) in Methods; black, sigmoid curve fit with the midpoint at 0.71). Note a high neutral stimulus response (40 Hz) defined as the firing rate at zero filter output (i.e., in the presence of stimuli orthogonal to the cell's STA). *C*: Population data of the neutral stimulus responses in each recording condition (in the same format as in A). *D-G*: Population data of the static nonlinear gain function (median for each cell type in corresponding colors; gray, interquartile range of all cells) across different conditions: isoflurane anesthesia (D, $N = 238$), FMM anesthesia (E, $N = 69$), *ex vivo* (F, $N = 342$), and awake condition (G, $N = 201$). *H*: Population data of the midpoint of the sigmoid nonlinearity in each recording condition: ***, p<0.001; Mann-Whitney *U*-test.

Supplementary Figure 3: Batch effects in the datasets. Dynamic range was computed for each cell as the difference between the maximum and minimum firing rates of the static nonlinear gain function (Figure 6D-G), and shown for each recording session in columns (dark green, monophasic ON; light green, biphasic ON; pink, biphasic OFF; violet, monophasic OFF; gray line, median). Some variabilities were present across preparations in each recording condition (isoflurane, *N*=44 recording sessions; FMM, *N*=10; awake, *N*=34; or *ex vivo*, *N*=18), but the batch effects were not large enough to eliminate any particular data set.

Retinal outputs in awake mice are less efficient but more linearly decodable

Our experimental data demonstrated that retinal outputs in awake mice had distinct characteristics compared to those under anesthesia or *ex vivo*, such as higher baseline firing rates (Figures 3 and 6), faster response kinetics (Figures 3-5), and a larger dynamic range (Figuress 3 and 6; Supplementary Figure 3). What are the implications on retinal coding? To address this question, we quantitatively analyzed our data in three ways.

First, we took an information theoretic approach and calculated the information conveyed by the retinal output spike trains recorded under different conditions (Figure 7; see Methods). Overall, we found that *in vivo* responses had a slightly but significantly higher information rate in bits per second (Figure 7A; isoflurane, 8.6 ± 5.9 bits/s; FMM, 7.6 ± 3.6 bits/s; awake, 7.5 ± 2.2 bits/s; median ± interquartile range) than *ex vivo* responses (6.2 ± 1.9 bits/s; *p*<0.001, *p*=0.004, *p*<0.001, respectively; Mann-Whitney *U*-test), but a lower information

rate in bits per spike (Figure 7B; isoflurane, 1.2 ± 1.6 bits/spike; FMM, 1.3 ± 1.4 bits/spike; and

awake, 0.3 ± 0.3 bits/spike; against *ex vivo*, 2.3±1.9 bits/spike; all with p<0.001). In particular,

the amount of information conveyed by a single spike in awake cells was the lowest due to

high average firing rates (34 ± 25 Hz; mean ± standard deviation). This indicates that the

retinal coding *in vivo* is less efficient than that *ex vivo* (Schwartz, 2021).

Figure 7: Retinal responses *in vivo* have a lower information rate in bits per spike but a

higher rate in bits per second than those *ex vivo*. Information rate conveyed by retinal

output spike trains in bits per second (A) and in bits per spike (B; Eq.(8) in Methods) in

response to randomly flickering full-field stimuli under different recording conditions (from left

to right: isoflurane, *N*=231; FMM, *N*=67; awake, *N*=154; *ex vivo*, *N*=328). Cell classes are

color-coded as in Figures 5 and 6 (dark green, monophasic ON; light green, biphasic ON; pink,

biphasic OFF; violet, monophasic OFF). ***, *p*<0.001; **, *p*<0.01 (Mann-Whitney *U*-test).

We next explored which neural coding scheme is favorable for awake retinal responses

(Supplementary Figure 4; see Methods). Specifically, we first simulated the retinal output

responses to a step change in light intensity (e.g., Supplementary Figure 4A,B) using a linear-

nonlinear cascade model with the observed temporal filter (STA; Figure 5) and static nonlinear

gain function (Figure 6). We then calculated the information based on the simulated peak firing

rate ("rate code"; e.g., Supplementary Figure 4C) or response latency ("temporal code"; e.g.,

Supplementary Figure 4D). We found that the information estimated with the rate code was

significantly larger for simulated awake responses (3.4 ± 1.2 bits; median ± interquartile range)

than for anesthetized (isoflurane, 3.0 ± 0.8 bit, *p*<0.001, Mann–Whitney *U*-test; FMM, 3.0 ± 0.8 bit, *p*=0.011) or *ex vivo* responses (2.4 ± 1.3 bit, *p*<0.001; Supplementary Figure 4E). In contrast, information in the latency-based temporal code of the simulated awake responses (1.9 ± 1.1 bits) was comparable to that of *ex vivo* responses (1.7 ± 1.3 bits, *p* = 0.6), but significantly lower than those calculated from the simulated anesthetized responses (isoflurane, 2.3 ± 0.6 bits; FMM, 2.4 ± 0.6 bits; both with *p*<0.001; Supplementary Figure 4F). Here we cannot directly compare the information between the rate and temporal codes as it depends on the assumptions made for the calculation (Eq.(9) in Methods). Nevertheless, information contained in a given coding scheme was calculated under the same assumptions across different recording conditions. Thus, our results suggest that the rate code is in favor of awake responses, while the temporal code is not.

505 **Supplementary Figure 4: Rate coding was more preferable than latency coding for**

506 **simulated awake retinal responses.** *A-D*: Simulated firing rates of a representative awake

507 cell (B) in response to a change in the stimulus contrast at 20 different levels (A, in

508 corresponding colors). The responses were computed from the cell's temporal filter and static

509 nonlinear gain function (see Figures 5 and 6, respectively). The vertical dotted line in B

510 indicates the peak response latency, from which the conditional probability distribution of the

511 peak firing rate, *P*(rate|stimulus), was calculated under the assumption that the standard

512 deviation of the Gaussian distribution follows 2.5% of the peak rates (C). The horizontal lines

513 in B indicate ±10% from the baseline firing rate as the detection threshold to compute the

514 conditional probability distribution of the response latency (D), *P*(latency|stimulus), where the

515 standard deviation of the Gaussian was given by 2.5% of the baseline firing rate (gray shade

516 in B). See Methods for details. *E*: Information in peak rate, $I_{\text{peak rate}}$, in bits computed from

517 *P*(rate|stimulus) as in C for each cell (from left to right: isoflurane, *N*=234; FMM, *N*=67; awake,

518 *N*=195; *ex vivo*, N=328). Cell classes are color-coded as in Figures 5-7 (dark green,

519 monophasic ON; light green, biphasic ON; pink, biphasic OFF; violet, monophasic OFF). ***,

520 *p*<0.001; **, *p*<0.01; *, *p*<0.05 (Mann-Whitney *U*-test). Simulated awake cells have the highest

521 information. *F*: Information in response latency, I_{latency}, computed from *P*(latency|stimulus) as

522 in D for each cell (shown in the same format as E). Given that both increase (red traces in D)

523 and decrease (blue traces in D) of the firing rates are detectable, simulated awake cells have

524 comparable information to cells *ex vivo*, but lower information than those under anesthesia.

525   Finally, to assess population coding properties, we performed a linear decoding

526 analysis using simultaneously recorded population activity in response to the full-field

527 randomly flickering stimulus (Warland, et al., 1997; see Methods). For each recording session,

528 the performance of the optimized linear decoder was evaluated by Pearson's correlation

529 coefficient, R, between the presented and estimated stimuli with 10-fold cross-validation. For

530 each recording condition, the decoding performance was then examined as a function of the

531 number of simultaneously recorded cells by fitting an exponential saturation function (Eq.(10)

532 in Methods; Figure 8). We found that single-cell linear decoding performance was comparable

533 across recording conditions (awake, $R = 0.20 \pm 0.13$, curve fit parameter a with 95%

534 confidence interval; isoflurane, 0.24 ± 0.04; FMM, 0.24 ± 0.06; *ex vivo*, 0.20 ± 0.10). At the

535 population level, however, Pearson's R reached 0.6 in about a third of the cases in the awake

536 recordings with less than 10 cells to be considered, but none in the anesthetized recordings

with less than 20 cells; and at least ~30 cells were required for the *ex vivo* recordings (Figure

8). Consistent with this observation, the exponential saturation constant was smaller for the

awake responses ($b = 8 \pm 4$ from Eq.(10); an estimate with 95% confidence interval) than for

the anesthetized (isoflurane, 23 ± 7; FMM, 21 ± 6) or *ex vivo* responses (72 ± 36). This

indicates that retinal population responses in awake mice are more linearly decodable than in

the other recording conditions, even though an image reconstruction is not necessarily the

objective function of the retina.

Figure8: Linear decoding of retinal population responses worked best with awake

responses. For each recording session, we performed linear decoding of the simultaneously

recorded population activity (333 ms window) in response to the full-field randomly flickering

stimulus (16.7 ms bin). As a measure of the decoding performance, we then calculated

Pearson's correlation coefficient R between the presented and estimated stimuli with 10-fold

cross-validation (from left to right: isoflurane, *N*=44; FMM, *N*=10; awake, *N*=33; *ex vivo*, *N*=18).

The black line shows the exponential curve fit (Eq.(10) in Methods; isoflurane, $a = 0.23 \pm 0.04$,

$b = 23 \pm 7$, estimate with 95% confidence interval; FMM, $a = 0.24 \pm 0.06$, $b = 21 \pm 6$;

awake, $a = 0.20 \pm 0.13$, $b = 8 \pm 4$; *ex vivo*, $a = 0.20 \pm 0.10$, $b = 72 \pm 36$).

554 **Discussion**

Here we established optic tract recordings in head-fixed mice (Figure 1) and systematically

analyzed the retinal output properties *in vivo* using a standardized set of visual stimuli for

characterizing retinal functions in a high-mesopic / low-photopic regime (Baden et al., 2016;

Jouty et al., 2018). We found that awake response properties were overall similar to those

under isoflurane or FMM anesthesia (Figures 2-4) or *ex vivo* (Figures 5 and 6); however, we

identified three distinct features in the awake condition besides behavioral effects

(Supplementary Figures 1 and 2; Schröder et al., 2020): higher firing rates, a larger dynamic

range (Figures 3 and 6; Supplementary Figure 3) and faster response dynamics (Figures 4

and 5). These properties made the awake responses less efficient from an information

theoretic viewpoint (Figure 7 and Supplementary Figure 4) but more linearly decodable at the

population level (Figure 8). Our results highlight that our knowledge on retinal physiology

cannot be simply translated from *ex vivo* to *in vivo* conditions, and that *in vivo* studies are

indispensable to better understand retinal computation.

In awake animals, both the baseline and visually-evoked firing rates of RGCs were

much higher—on average by ~20 Hz—than in those under anesthesia or in an isolated

preparation (Figures 3 and 6). To our knowledge, spiking activity of RGCs in awake mice has

been reported only in some recent studies with a low sample size and none of them provided

any descriptive statistics (Hong et al., 2018; Schröder et al., 2020; Sibille et al., 2022).

Nevertheless, the reported examples (Fig. 3C, Schröder et al., 2020; Fig. S7h, Sibille et al.,

2022) show high baseline activity (>10 Hz), consistent with our observation. These two studies

employed high-density electrodes (NeuroPixels) to record axonal signals of RGCs in

C57BL/6J mice, and are thus arguably comparable to our study from a methodological

viewpoint. In contrast, using mesh electrodes directly injected into the mouse eye, Hong et al.

(2018) reported relatively low light-evoked RGC responses (<10 Hz) in albino CD1 strain. This

is likely not due to the difference in the recording methods, but due to the mouse strain. Albino

mice generally have poor vision with a reduced number of rod photoreceptors (Abdeljalil et al.,

2005; Brown et al., 2007), suggesting a certain level of deficits in their retina. High firing activity

in awake animals has been widely observed in many brain areas across species (Greenberg

et al., 2008; Sellers et al., 2015; Durand et al., 2016; Wright et al., 2017; De Franceschi and

Solomon, 2018; Chen and Song, 2019). Therefore, it could be a general feature of the brain

in an awake state, including the retina.

This high firing activity had three consequences on the retinal output properties. First,

by definition it leads to a lower DS/OS index value calculated as a normalized population

vector (Eq.(3) in Methods). Indeed, DS/OS cells were rather underrepresented in our awake

recordings with high baseline activity (Figures 2 and 3), while overrepresented in our

anesthetized recordings as well as in Hong et al. (2018) with low baseline activity. Second,

the baseline activity level affected the response kinetics (Figures 4 and 5). The higher the

baseline firing is, the higher the conductance of the cell is, hence the shorter the response

latency becomes (Zohar et al., 2011; Wang et al., 2014; Durand et al., 2016). Third, the

dynamic range was also altered (Figures 3 and 6; Supplementary Figure 3). Indeed, many

awake ON cells exhibited an OFF-suppressive response, an emerging feature observed only

in those cells with high baseline firing rates (~60 Hz). Here we categorized these cells

separately from the other ON cells in our classification analysis (Figure 2). It is, however,

unlikely that they form a distinct "cell type" in the retina, even though we cannot deny a

possibility that they were previously overlooked due to a different in the recording modality

(calcium imaging; Baden et al., 2016) or clustering methods (Jouty et al., 2019). Instead, these

cells likely belong to a known cell type, and altered their response properties due to an

increased baseline activity in the awake condition. Possible candidate cell types are: transient

ON alpha cells (although they have a low baseline firing rate *ex vivo*; Krieger et al., 2017), ON

contrast suppressive cells (although a sharp rebound response is missing in the calcium

dynamics *ex vivo*; Baden et al., 2016), or suppressed-by-contrast cells (although their high

baseline activity is suppressed by both ON and OFF stimuli; Mastronarde, 1985; Tien et al.,

2015; Jacoby and Schwartz, 2018). It is interesting to identify the exact cell type of these

RGCs and further characterize their function *in vivo* in future studies.

What are the mechanisms underlying such differences between *in vivo* and *ex vivo*

retinal responses? One possibility is the difference in the physiological states of the retina.

Even if *ex vivo* recordings are made under a proper temperature (Vogel et al., 2016; Vlasiuk

and Asari, 2021) using a perfusion medium specifically formulated to support isolated retinal

tissues (Ames and Nesbett, 1981), there are certain unavoidable differences to the

physiological condition *in vivo*. In particular, the retinal pigment epithelium, a key player for the

visual cycle and spatial buffering of ions, is often removed in an isolated retinal preparation. A

lack of retinal supply can then alter the physiological states of the retinal neurons, potentially

leading to a change in their visual response properties.

Differences in the input stimulus properties, in contrast, cannot explain our results,

even though retinal responses depend a lot on stimulus conditions. For example, the higher

the light intensity level is, the higher the temporal frequency sensitivity becomes for *ex vivo*

RGC responses (threshold at around 20-30 Hz; Wang et al., 2011) and so does the critical

flicker-fusion frequency at the behavioral level (15-40 Hz; Umino et al., 2018; Nomura et al.,

2019). Response latencies depend also on the spatial patterns of stimuli as well as their

contrast (Bolz et al., 1982; Baccus and Meister, 2002; Sagdullaev and McCall, 2005; Pearson

and Kerschensteiner, 2015; Tikidji-Hamburyan et al., 2015). Here we used full-field stimuli at

a somewhat lower light intensity level for *in vivo* recordings than for *ex vivo* ones (see Methods

for details). Pupillary reflexes in awake animals further decrease the dynamic range of the

stimulus impinging on the retina (Hatter et al., 2003). Therefore, higher baseline activity and

faster dynamics observed in awake animals cannot be simply attributed to the difference in

the stimulus conditions.

Finally, when comparing retinal responses across different studies, one cannot ignore

a possible effect of sampling bias because visual response dynamics are cell-type specific

(van Wyk et al., 2009; Krieger et al., 2017; Ravi et al., 2018; Tengölics et al., 2019). In an

extreme case, delayed ON cells have the response latency slower than other RGC types by

hundreds of milliseconds in an isolated mouse retina (Mani and Schwartz, 2017). Response

dynamics are also species-specific. For example, ON RGCs show higher baseline activity and

faster responses than OFF cells in the macaque (Chichilnisky and Kalmar, 2002) and guinea

pig (Zaghloul et al., 2003) retinas, whereas slower in the salamander retina (Gollisch and

Meister, 2008), and such asymmetry between ON and OFF dynamics is pathway-specific in

the rat retina (Ravi et al., 2018). Moreover, kinetics can vary even for the same cell type,

depending on the retinal location (Warwick et al., 2018) or light-adaptation state (Chang and

He, 2014; Tengölics et al., 2019). Sampling bias can then skew the results in many different

ways. Given that most RGCs project their axons through the optic tract (Ellis et al., 2016;

Rosón et al., 2019), we expect that the optic tract recordings employed in this study should

have access to most cell types, if not all. Moreover, here we observed a wide range of visual

responses (Figure 2), and no apparent difference was found in the STA distributions across

different recording conditions (Figure 5). We thus expect that the effects of sampling bias

should be minimal on the observed differences between *in vivo* and *ex vivo* retinal response

characteristics.

Taken together, our findings indicate that clear differences exist in retinal physiology

between *in vivo* and *ex vivo*. This highlights the importance of studying retinal function *in vivo*,

especially in the fully physiological awake condition.

Implications on retinal coding *in vivo*

The optic nerve fiber forms an information bottleneck in the early visual system. The human

retina, for example, contains ~10^8 input neurons (photoreceptors) but only ~10^6 output

neurons (RGCs) whose axons make up the optic nerve (Masland, 2012; Sanes and Masland,

2015). The retina has then been assumed to optimize the channel capacity by compressing

the visual information as much as possible and conveying signals to the brain using as few

spikes as possible (Attneave, 1954; Barlow and Rosenblith, 1961; Laughlin, 2001). This so-

called efficient coding hypothesis is supported by both *ex vivo* experimental data and

theoretical analyses. Under *ex vivo* conditions, while some RGCs exhibit high firing under

certain conditions (Ke et al., 2014; Smeds et al., 2019), many RGCs are silent most of the

time and fire spikes at high rates only when their selective stimulus features are presented

(Gollisch and Meister, 2010; Baden et al., 2016; Jouty et al., 2018). This sparse RGC response,

ensured by low spontaneous activity and strong nonlinearity (Pitkow and Meister, 2012), helps

achieve high efficiency and low redundancy in the visual representation of the retina (Doi et

al., 2012). Furthermore, efficient coding principles can explain various physiological properties

of the retina, such as separation of retinal outputs into multiple cell types (Gjorgjieva et al.,

2019), even though it cannot fully account for the retinal function (Schwartz, 2021).

The retinal output characteristics we observed *in vivo*, however, provide a different

view on the retinal code. First, unlike *ex vivo* conditions, RGCs in awake mice generally

showed a high baseline activity with more linearized responses (Figures 3 and 6).

Consequently, many awake ON cells primarily encoded light decrements by suppression from

high baseline, rather than light increments by activation from low baseline. These cells convey

the same information as conventional ON cells do—the more spikes, the higher light

intensity—but using much more spikes. Thus, the amount of information transmitted per spike

is much lower *in vivo* than *ex vivo* (Figure 7), violating the efficient coding principles. Second,

we found less variability in the visual response dynamics across different RGC classes *in vivo*

(Figure 5E-H). This suggests that temporal coding framework may not be readily applicable *in

vivo* as it requires noticeable differences in spike timing patterns or response latencies

between two or more channels, such as ON and OFF pathways converging onto ON-OFF

RGCs (Gollisch and Meister, 2008). Our modeling analysis indeed demonstrated that awake

responses were in favor of a rate code but not a latency-based temporal code (Supplementary

Figure 4), and that a linear decoder worked significantly better with the population responses

in awake mice (Figure 8), even though an image reconstruction is not necessarily the retina's

objective function. These results point out a need to reconsider retinal function *in vivo*.

What are the advantages of such energy-inefficient retinal coding? As shown in our

awake recordings, one can gain faster response kinetics (Figure 5) and wider bandwidths

(Figure 6). In addition, redundancy in the dense code helps transmit signals accurately even

with intrinsically noisy spike trains, as demonstrated by a high performance of a simple linear

decoder (Figure 8). These features are all highly beneficial from a behavioral viewpoint—e.g.,

to detect predators robustly and promptly—and thus worth achieving for survival at an expense

of energy cost. Interestingly, dense coding can be a general feature of early sensory

processing, as suggested by high baseline firing in vestibular and cochlear nuclei (Fuchs and

Kimm, 1975; Rhode and Smith, 1986; Warchol and Dallos, 1990; Beraneck and Cullen, 2007),

tonic activity in hair cells (Sachs and Abbas, 1974; Wu et al., 2016), and dark current in

vertebrate photoreceptors (Hagins et al., 1970; Okawa et al., 2008). In fact, the energy cost of

dense retinal coding may not be so problematic with a relatively small number of neurons, as

opposed to the total energy required for operating $\sim 10^9$ neurons in the cortex where sparse

coding has certain advantages from the viewpoints of both energy efficiency and sensory

processing (Olshausen and Field, 1996; Asari et al., 2006).

The retina, however, may employ different coding principles under different light levels

(Wang et al., 2011; Tikidji-Hamburyan et al., 2015; Borghuis et al., 2018). All our *in vivo*

recordings were done in a high-mesopic / low-photopic range (at $\sim 10^3$ R*/photoreceptor/s; see

Methods). In a scotopic condition, in contrast, *ex vivo* studies have reported a high tonic firing

rate for OFF RGC types (~50 Hz, OFF sustained alpha cells in mice; Smeds et al., 2019; ~20

Hz, OFF parasol cells in primates; Ala-Laurila and Rieke, 2014), while a low tonic firing for

corresponding ON cell types (<1Hz). Since these ON cells were shown to be responsible for

light detection by firing in the silent background (Smeds et al., 2019), here the retina seems to

operate with a sparse feature detection strategy. Further characterizations of the retinal code

in vivo and its behavioral relevance will be needed to clarify what exactly the eye tells the brain.

Effects of anesthetics in early visual system

Anesthetics generally lower the excitability of nerve cells: e.g., isoflurane acts on the gamma-aminobutyric acid (GABA) type A receptors to silence the brain (Jenkins et al., 1999). It thus makes sense that the retinal outputs were reduced under anesthesia (Figures 3 and 6) and showed slower dynamics (Figures 4 and 5) than in awake animals. It is, however, difficult to generalize the effects of anesthesia because multiple mechanisms of action are involved, exerting complex effects on the sensory systems in anesthetic- and dose-dependent manners (Populin, 2005). For example, while visual cortical responses are reduced under anesthesia (Haider et al., 2013; Vaiceliunaite et al., 2013), auditory responses are enhanced (Raz et al., 2014; Sellers et al., 2015) and the response latency becomes shorter in the auditory pathway (Ter-Mikaelian et al., 2007).

Our results are nevertheless consistent with previous studies on the effects of anesthesia in the early visual system. Specifically, anesthesia reduces the overall activity of both SC and dLGN neurons, and leads to multiple changes in temporal processing of these two major retinorecipient areas, such as reduced sensitivity to high temporal frequencies and longer response latencies (Zhao et al., 2014; Durand et al., 2016; De Franceschi and Solomon, 2018). While different anesthetics were used in these studies (urethane as opposed to isoflurane and FMM), our consistent findings suggest that such changes in the response dynamics arise largely from the retina. In contrast, the effects of anesthesia on spatial response properties seem more complex. While spatial processing remains largely intact in dLGN (Durand et al., 2016), lower sensitivity to contrast and larger receptive field sizes were reported in SC under urethane anesthesia (De Franceschi and Solomon, 2018). Moreover, SC neurons show increased orientation selectivity under isoflurane anesthesia (Kasai and Isa, 2021). Here we also found a larger fraction of DS/OS cells under both isoflurane and FMM anesthesia than in the awake condition (Figure 2D-F). This, however, could be due to the difference in the baseline firing rate (Figure 3) as the DS/OS index is defined by a normalized

738 population vector in response to moving gratings (Eq.(3). In Methods; Mazurek et al., 2014).

739 It is a future challenge for fully characterizing spatiotemporal processing of the retina *in vivo*.

Materials and Methods

741 No statistical method was used to predetermine the sample size. The significance level was

742 0.05 (with Bonferroni correction where appropriate) in all analyses unless noted otherwise. All

743 experiments were performed under the license 233/2017-PR from the Italian Ministry of Health,

744 following protocols approved by the Institutional Animal Care and Use Committee at European

745 Molecular Biology Laboratory. The data analyses were done in Python.

Animals

747 A total of 55 female C57BL/6J mice were used (chronic, 3; acute, 35; failures, 17, including

748 10 initial unsuccessful attempts to record from the optic chiasm) at around three months of

749 age at the time of the surgery. Mice were kept on a 12-h light / 12-h dark cycle and given water

750 and food *ad libitum*. After the chronic implantation of electrodes, the animals were kept single-

751 housed.

Chronic recordings

For implanting electrodes, animals were anesthetized (induction, 4% isoflurane; maintenance,

1-2%) and placed inside a stereotaxic apparatus (Neurostar). Throughout the surgery,

temperature was maintained at 37 °C using a heating pad (Supertech Physiological), and eye

ointment (VitA-POS) was used to prevent the eyes from drying. After positioning the mouse

head, the scalp skin was disinfected (Betadine) and removed with scissors. Soft tissue was

removed from the skull surface with a round scalpel, and ethanol and acetone were applied to

the skull for disinfection and removal of any residual compounds. The skull was glazed with a

drop of cyanoacrylate adhesive (Loctite 401) to secure in place the surrounding skin, and then

registered in the stereotaxic controller software (NeuroStar). A dental drill (WPI) was used to

leave three marks on the skull to label the entry point for targeting the optic tract: [-1.34, +1.87,

+4.74], [-1.70, +1.87, +4.74], and [-1.82, +2.35, +4.07] in [Anterior-Posterior, Medial-Lateral,

Dorsal-Ventral] coordinates, respectively. A hole was drilled for a reference silver wire (A-M

Systems) above the cerebellum, and the wire was inserted sideways to avoid excessive brain

damage. After covering the hole and wire with Vaseline, the wire was attached to the skull

with cyanoacrylate adhesive and dental cement (Paladur, PALA). Subsequently, a custom-

designed titanium head-plate was attached to the skull with dental cement, followed by a

craniotomy (diameter, 1-2 mm) and durotomy. A chronic silicone probe (Buzsaki32L,

NeuroNexus) was then inserted with the stereotaxic controller (75 µm/min) first to a depth of

2 mm, retracted by 1 mm to release pressure created by the initial brain entry, and then to a

depth of 4.5 mm from the skull surface. After reaching the desired depth, a microdrive (dDrive,

NeuroNexus) was attached to the skull with dental cement, and the probe was removed from

its mount and covered with a protective cap. The cables and connectors were cemented to

the cap and covered with paraffin film (Bemis, Parafilm) to prevent the mouse from damaging

the implant.

After the surgery, the animals were kept on a heating pad (Sera, Thermo comfort mat

S) until they recovered from anesthesia. During the next four days, the mice received an anti-

inflammatory/antibiotic cocktail (Rimadyl/Baytril; 0.5 mg/mL each, 0.01 mL/g). The antibiotic

(Baytril) was given for an additional three days through drinking water (0.17 mg/mL).

After a recovery period of five days, the mice were placed on a custom-made rotary

treadmill (diameter, 20 cm) with their head fixed for the recordings (at most two times a day,

each for <2 hours). During the initial sessions, we moved the probe until visual responses

were observed (Figure 1A-D). The electrophysiology data were recorded at 30 kHz from each

recording site (SmartBox, NeuroNexus) together with synchronization pulses from the visual

stimulation device (see below). In total, we made 17 recording sessions from 3 out of 6 animals,

where up to 10 cells were simultaneously recorded in each session (4.4 ± 2.4 cells/session,

mean ± standard deviation).

After all the recording sessions, the electrode position was verified histologically (e.g.,

Figure 1E). The mice were anesthetized (2.5% Avertin, 16 µL/g, intraperitoneal injection) and

perfused with paraformaldehyde (PFA; 4% in phosphate buffer solution). The animal's head

with the electrode left in position was collected without the skull base, and stored in fixative

solution (4% PFA) at 4°C for at least four days. This helped the brain tissue harden around

the silicon probe, hence leaving a visible mark after removing the probe. Harvested brain

tissue was then coronally sliced with a vibratome (Leica, VT1000S; thickness, 150 µm) and

imaged under a bright-field microscope (Leica, LMD7000).

Acute recordings

Animals were anesthetized and placed inside a stereotaxic apparatus with a heating pad as

described above. A contact lens (diameter, 3 mm; Ocuscience) was used for the target eye to

prevent it from drying. The scalp was removed, registered in the stereotaxic controller, and

the three entry points for targeting the optic tract were marked with the stereotaxic drill. A well

was made with dental cement around the marks later to hold saline solution on top of the brain.

After a craniotomy (diameter, 2 mm), an acute silicone probe (Buzsaki32L, Neuronexus)

coated with a fluorescent dye (DiI stain, Invitrogen, D282) was slowly inserted into the brain

(100 µm/min) with a micromanipulator (Sensapex, SMX) attached to the stereotaxic apparatus.

While approaching the target depth, visual stimuli were presented (full-field contrast-inverting

stimulus at 0.5 Hz; see below). The probe was moved until a maximum number of visually

responsive cells were seen at once (up to 20 cells; 7.7 ± 5.3 cells/session, mean ± standard

deviation; 52 sessions in total from 23 out of 27 animals).

Throughout the recordings, the mouse was kept under anesthesia with 1% isoflurane.

The depth of anesthesia was monitored by the breathing rate (~1 breath/s). Alternatively, we

used a cocktail of fentanyl, medetomidine and midazolam (FMM) during the recordings. In this

case, once the electrode was in position, an FMM solution (fentanyl, 0.05 mg/kg;

medetomidine, 0.5 mg/kg; midazolam, 5 mg/kg; in 0.9% NaCl saline) was intraperitoneally

administered, and the isoflurane dose was progressively decreased to 0%. Buprenorphine

(0.1 mg/kg) was injected 20 minutes after the termination of isoflurane, and the recording

session was initiated 10 minutes after the buprenorphine injection. The depth of anesthesia

was monitored through the heart rate (below around 300 beats per minute) and supplemental

FMM anesthesia was provided when required. The electrophysiology data and the visual

stimulation signals were recorded at 30 kHz/electrode (SmartBox, NeuroNexus).

At the end of the recording session, the electrode location was verified histologically.

After retracting the silicone probe, the animal was perfused as described above. The brain

tissue was harvested and post-fixed overnight in 4% PFA at 4°C. Coronal sections of the brain

tissue (thickness, 150 μm) were then examined under a fluorescence microscope (Leica,

LMD7000 with N2.1 filter cube) to visualize the trace left by the DiI stain on the probe.

For acute awake recordings, we first performed head-plate implantation as described

above. In short, after placing an animal on a stereotaxic apparatus under anesthesia, we 1)

surgically removed the skin and the periosteum above the skull; 2) attached a head-plate to

the skull with dental cement; and 3) applied a thin layer of cyanoacrylate adhesive (Loctite

401) and a thick layer of silicone sealant (Kwik-cast) to provide mechanical and thermal

support. Animals were then allowed to recover for a week, and subsequently habituated over

the course of 10 days to head fixation in the recording setup as described above. On the day

of the recording, we removed the silicone to expose the skull surface of the animal with its

head fixed in the recording setup, and determined the electrode penetration path to the target

area using the robotic stereotaxic system (StereoDrive, NeuroStar). The animal was then

briefly anesthetized (for about 5 minutes) and a hole was drilled around the electrode entry

point on the skull. After the removal of the anesthesia, the electrode attached to the robotic

arm was lowered at 5 μm/s until visual responses were found in the target area. Recordings

and histological verification of the electrode location were performed as described above.

Visual stimulation

Visual stimuli were presented by a custom gamma-corrected digital light processing (DLP) device (Texas Instruments, DLPDLCR3010EVM-LC) where the original green and red light-emitting diodes (LEDs) were replaced with ultra-violet (UV; 365 nm; LZ1-00UV00, LED Engine) and infrared (IR; 950 nm; SFH 4725S, Osram) LEDs, respectively. The UV and blue (465 nm) lights were projected onto a spherical screen (radius, 20 cm) with UV-reflective white paint (waterfowl store) placed ~20 cm to the contralateral side of an animal's eye from the implanted probe (Figure 1A), whereas the IR light was used as synchronization pulses, recorded via a photodiode with a custom transimpedance amplifier. The visual stimuli (1280-by-720 pixels; frame rate, 60 Hz) covered $\omega_{azimuth}$ = 73° in azimuth and $\omega_{azimuth}$ = 44° in altitude from the mouse eye position (Figure 1B). The maximum light intensity at the eye position was 31.3 mW/m^2 (E_{365} = 15.4 mW/m^2 for UV LED and E_{365} = 15.9 mW/m^2 for blue LED; measured with S121C sensor, Thorlabs), leading to mesopic to photopic conditions (see also "Light intensity" section below).

Using this DLP setup and QDSpy software (Franke et al., 2019), we presented the following set of visual stimuli: a randomly flickering full-field stimulus (5 min; 60 Hz), a moving grating stimulus (spatial frequencies of square waves, 3° or 20°; moving speed, 7.5°/s or 15°/s, in eight different directions), a full-field stimulus whose intensity followed a sinusoid (1.5 Hz) with a linearly increasing amplitude from 0% to 100% contrast over 10 s (10 trials; Figure 3), and a full-field sinusoidally flickering stimulus (maximum contrast) at different temporal frequencies (1.875, 3.75 and 7.5 Hz, each for 2 s; 15 Hz for 1 s; 10 trials; Figure 4). The last two full-field stimuli were preceded by a sequence of "OFF-ON-OFF" stimulation at maximum contrast (full-field contrast inversion; 2 s each) and interleaved by a 1-s-long gray screen across trials. These stimuli were equivalent to those used for differentiating ~30 RGC types *ex vivo* (Baden et al., 2016; Jouty et al., 2018).

Light intensity

We calculated the photon flux on the mouse retina in vivo as: $I_\lambda = T_\lambda \cdot F_\lambda \cdot A_{\text{pupil}}/A_{\text{retina}}$, where T_λ

is the transmittance of the eye optics at wavelength λ; $F_\lambda = E_\lambda/(hc/\lambda)$ is the photon flux at the

pupil with Planck's constant h and the speed of light c; $A_{\text{pupil}} \approx \pi \cdot (d_{\text{pupil}}/2)^2$ is the pupil area with

diameter d_{pupil}; and $A_{\text{retina}} \approx (\pi \cdot d_{\text{retina}} \cdot \omega_{\text{azimuth}}/360°) \cdot (\pi \cdot d_{\text{retina}} \cdot \omega_{\text{azimuth}}/360°)$ is the illuminated

retinal area with diameter d_{retina}. Assuming that $T_{365} = 50\%$, $T_{454} = 68\%$ (Jacobs and Williams

2007), $d_{\text{pupil}} = 1$ mm and $d_{\text{retina}} = 4$ mm, we then obtained $I_{365} = 3.81 \times 10^3$ and $I_{454} = 6.64 \times 10^3$

photons/s/μm^2. Given a total photon collecting area $a = 0.2$ and 0.5 μm^2 for cones and rods,

respectively (Nikonov et al., 2006), and a relative sensitivity $s_{365} = 90\%$ and $s_{454} = 0\%$ for S-

cones while $s_{365} = 25\%$ and $s_{454} = 60\%$ for M-cones and rods (Jacobs and Williams 2007), we

estimated the photoisomerization (R*) rate as: $a \cdot \Sigma_\lambda\ I_\lambda s_\lambda = 2.5 \times 10^3$ R*/rod/s, 0.7×10^3 R*/S-

cone/s, and 1.0×10^3 R*/M-cone/s.

In contrast, the maximum light intensity of the *ex vivo* set up was $E = 36$ mW/m^2

(Vlasiuk and Asari, 2021). Assuming that the white light from a cathode-ray tube monitor is

centered around $\lambda = 500$ nm, we estimated the photon flux on an isolated retina as: $I = E/(hc/\lambda)$

$= 9 \times 10^4$ photons/s/μm^2. Given the sensitivity $s_{500} = 2\%$ for S-cones and 40% for M-cones and

rods (Nikonov et al., 2006), we then obtained: $a \cdot I \cdot s_{500} = 4 \times 10^4$ R*/rod/s, 2×10^3 R*/S-cone/s,

and 4×10^4 R*/S-cone/s. Thus, the light intensity level was about an order of magnitude higher

for *ex vivo* data than for *in vivo* data.

Physiology data analysis

In total, we obtained 282 cells in awake conditions (75, chronic; 282, acute), and 325 and 103

cells in acute anesthetized recordings with isoflurane and FMM, respectively. Spike sorting

was performed with SpykingCircus (Yger et al., 2018) for semi-automatic cluster detection and

Phy (Rossant, 2020) for data curation. In brief, spikes were detected from highpass-filtered

data traces (cutoff, 300 Hz) as those events 7 standard deviations away from the mean.

Single-units were then identified by clustering in principal component space, followed by

manual inspection of spike shape as well as auto- and cross-correlograms. As expected from

axonal signals (Barry, 2015), spike waveforms typically had a triphasic shape (e.g., Figure 1F-

H). No putative single-units had the interspike intervals below 1 ms for a refractory period (e.g.,

Figure 1I-K), except for 1 (out of 103) for FMM-anesthetized recordings. Most units had a

minimum interspike interval above 2 ms (93%, awake; 94%, isoflurane; 99%, FMM). Here we

included in the analysis those small numbers of visually responsive cells that had a small

fraction of interspike intervals below 2 ms; removing them did not affect the conclusions. Not

all cells responded to the entire stimulus set, but cells responding to any of the presented

stimuli were included in the analysis.

For *ex vivo* recordings, we reanalyzed the data sets in Vlasiuk and Asari (2021).

Specifically, the *ex vivo* data sets included the activity of 696 cells recorded with a multi-

electrode array (from 18 isolated mouse retinas) in oxygenated Ames' medium at 37 °C in

response to a randomly flickering full-field visual stimulus projected from a gamma-corrected

cathode-ray tube monitor (frame rate, 100 Hz) or a DLP device (60 Hz). Given that the

temperature of the ocular surface is ~37 °C (Vogel et al., 2016), the retina likely operates at

around 37 °C *in vivo*.

Response quality

908   We assessed the cell's response quality based on the trial-to-trial reliability of the response

909     $r(t)$ during the "ON-OFF" period of the OFF-ON-OFF stimulus sequence (see Figure 2A for

910   example). Specifically, the signal-to-noise ratio was calculated for each cell as follows:

911
$$\text{SNR} = \frac{\text{var}[\langle r(t) \rangle]_t}{\langle \text{var}[r(t)]_t \rangle}, \tag{1}$$

912   where $\langle \cdot \rangle$ indicates the mean over trials, and $\text{var}[\cdot]_t$ the variance over time t (bin size, $\Delta t =$

913   16.6 ms). We set a threshold at 0.15 to select reliably responsive cells for further analyses

914   (Figure 2B).

Response polarity

916 To characterize the cell's preference to stimulus polarity, we defined an ON-OFF index using

917 the responses to the full-field contrast inversion:

$$\text{ON-OFF index} = \frac{r_{\text{ON}} - r_{\text{OFF}}}{r_{\text{ON}} + r_{\text{OFF}}}, \tag{2}$$

919 where r_{ON} and r_{OFF} are the mean firing rate during the ON and the second OFF periods of the

920 OFF-ON-OFF stimulus sequence, respectively. Positive and negative ON-OFF index values

921 indicate stronger responses to ON and OFF stimuli, respectively (Figure 2C).

Direction and orientation selectivity

923 Direction-selectivity (DS) and orientation-selectivity (OS) indices were calculated by projecting

924 the responses to the moving grating stimuli onto a complex exponential (Mazurek et al., 2014):

$$\text{DS or OS index} = \left\| \frac{\sum_k e^{-i\alpha\omega_k} \cdot r_k}{\sum_k r_k} \right\|, \tag{3}$$

926 where ω_k and r_k are the angle of the k-th direction and the cell's corresponding responses,

927 respectively; and $\alpha = 1$ and 2 for the DS and OS indices, respectively. Cells were considered

928 DS and/or OS when the corresponding index was higher than 0.15 and $p < 0.2$ calculated by

929 bootstrap methods (1,000 repetitions; Figure 2D-F).

Temporal frequency sensitivity

931 Temporal frequency sensitivity was assessed by fitting an even power of sine function to the

932 responses to the full-field sinusoidally flickering stimulus at different frequencies h (1.875, 3.75,

933 7.5, or 15 Hz; Figure 4):

$$s(t) = A \cdot \sin^q \left(\pi h t - \frac{\phi}{2} \right) + B, \tag{4}$$

where A and $B \geq 0$ are the evoked response amplitude and baseline activity, respectively; and

$q = 2n$ with $n \in \mathbb{N}^+$ is the exponent of the sine function. The phase $\phi \in [-\pi, \pi)$ indicates the

relative position between the response peak and the sinusoidal stimulus patterns. For example,

$\phi = -\pi/2$ and $\pi/2$ are obtained if the response reaches its peak when the stimulus is

brightest and darkest, respectively.

The fit quality was then assessed by the explained variance:

$$R^2 = 1 - \frac{\text{var}[\text{Data} - \text{Fit}]}{\text{var}[\text{Data}]}. \tag{5}$$

942   We set a threshold at 0.2 to select cells with a good fit hence responsive to the stimulus (Figure

943   4C), and compared the proportions of the responsive cells across different recording

944   conditions (two-proportion z-test; Figure 4C,D).

Sensitivity to contrast

946   We used an even power of sine function with a sigmoid envelope to characterize the

947   responses to the flickering stimulus ($h = 1.5$ Hz) with increasing contrast (Figure 3):

$$S(t) = \frac{A}{1 + e^{-\lambda(t - t_0)}} \cdot \sin^q \left(\pi h t - \frac{\phi}{2} \right) + B, \tag{6}$$

where t_0 and $\lambda > 0$ are the midpoint and the steepness of the sigmoid, respectively; and the

other free parameters are the same as $s(t)$ in Eq.(4). The fitted parameter values were then

compared across different recording conditions: Mann-Whitney U-test for the baseline B

(Figure 3D) and the absolute amplitude $|A|$ (Figure 3E), and t-test for the phase ϕ for ON and

OFF responses, respectively (k-means clustering with $k = 2$; Figure 3F). As a measure of the

sensitivity to contrast, we used an estimated response magnitude at 10% contrast from Eq.(6)

as well as the absolute slope size at the midpoint of sigmoid ($|A\lambda/4|$).

Temporal filters and static nonlinear gain functions

For systematically characterizing the response dynamics, we used stimulus ensemble

statistical techniques ("reverse correlation" methods; 500 ms window) to calculate the linear

filter (Figure 5) and static nonlinear gain function (Figure 6) of the recorded cells in response

to a randomly flickering visual stimulus (Meister et al., 1994; Chichilnisky, 2001). First, we

obtained the linear filter of each cell by calculating a spike-triggered average (STA) of the

stimulus with ± 1 being "white" and "black," respectively. As a quality measure, p-value was

computed for each time bin against a null hypothesis that the STA follows a normal distribution

with a mean of zero and a variance of $1/C$, where C is the total number of spikes. As a

measure of the cell's temporal frequency tuning, we then estimated the peak latency by fitting

a difference-of-Gaussian curve to the linear filter (e.g., Figure 5A); and the spectral peak

frequency by the Fourier analysis on the fitted curve (e.g., Figure 5B). The curve fitting quality

was assessed by the explained variance R^2 as in Eq. (5). We discarded the cells if $p > 10^{-18}$

for all time bins or $R^2 < 0.8$. We ran a t-test to compare the temporal frequency tuning

properties at the population level across recording conditions (Figure 5C,D) or across different

cell types in each recording condition (Figure 5E-H). Spatial response properties were not

examined in this study because complex nonlinear kinematics of eye movements in the awake

condition precluded the analysis at a high enough spatial resolution; and because it has been

shown that anesthesia has no effect on the spatial processing in dLGN, a direct downstream

of the retina (Durand et al., 2016).

Static nonlinear gain function $P(\text{response}|\text{stimulus})$ of each cell was computed by

taking the ratio between the distribution of spike-triggered stimulus ensembles

$N(\text{stimulus}|\text{response})$ projected onto the $L2$-normalized STA (bin size, 0.1) and that of the

entire stimulus ensembles $N(\text{stimulus})$ (e.g., Figure 6B):

$$P(\text{response}|\text{stimulus}) = \frac{N(\text{stimulus}|\text{response})}{N(\text{stimulus})}/\Delta t. \tag{7}$$

A sigmoid function was fitted to $P(\text{response}|\text{stimulus})$ for smoothing. The neutral stimulus

response was then defined as the vertical-intercept of the sigmoid function at the horizontal

axis value of zero: i.e., the firing rate when the stimulus is orthogonal to the STA. We ran a t-

test to compare the neutral stimulus responses and the average firing rates during the stimulus

presentation at the population level across conditions (Figure 6A,C); and Mann-Whitney U-

test to compare the midpoint of the sigmoid (Figure 6H). The dynamic range was computed

for each cell as the difference between the maximum and minimum firing rates of the static

nonlinear gain function (Supplementary Figure 3).

Response classification

In this study, we were not able to perform a morphological analysis of individual cells as we

employed blind *in vivo* recording methods (Figure 1). We thus focused on the visual response

properties for categorizing cell-classes in the following two ways.

First, we classified cells *in vivo* using the response quality, response polarity, and

motion sensitivity (Figure 2). Specifically, we first divided the cells into two groups: reliably

responsive ones with $\mathrm{SNR} > 0.15$ (Eq. (2)) and the other low quality ones ("N/A" type; Figure

2B). We then set thresholds at ± 0.25 for the ON-OFF index (Eq. (3)) to identify ON, OFF, and

ON/OFF cells within the reliably responsive cells that increased firing in response to light

increments, light decrements, and both, respectively (Figure 2C). For some cells where these

measures were not available, we used the responses to the full-field sinusoidally flickering

stimuli to calculate the response quality and polarity in a similar manner. Within the ON cells,

we further identified an "OFF-suppressive" class if they had a significant negative amplitude

in Eq. (6) with 95% confidence intervals in the parameter estimation. Independently, we also

labeled cells as motion sensitive or not, based on the DS/OS indices as described above

(Eq.(3); Figure 2D-F).

Second, because these measures are not available for the *ex vivo* data sets (Vlasiuk

and Asari, 2021), we performed classification using the temporal filter properties to make a

fair comparison between the *ex vivo* and *in vivo* data sets (Figures 5 and 6). Specifically, we

first ran a principal component analysis (PCA) on the temporal filters ($L2$-normalized STA)

obtained in each condition (Gollisch and Meister, 2008; Asari and Meister, 2014). The first two

principal components (PC1, monophasic filter; and PC2, biphasic filter) were largely sufficient

to fit all the temporal filter dynamics, accounting for 78-86% of the total variance in each

condition. Each temporal filter is a point in the PCA biplot (i.e., the two-dimensional space

spanned by PC1 and PC2), and we grouped its shape into four subtypes based on its position:

monophasic OFF, biphasic OFF, monophasic ON, and biphasic ON from the first to the fourth

quadrants, respectively.

Here we did not aim to fully differentiate all functional cell-types (~35 RGC types from

ex vivo studies; Baden et al., 2016; Jouty et al., 2018) for three reasons. First, our data sets

are not large enough. Second, the presented stimulus sets were not diverse enough. Finally,

some cells may have not been sampled, such as those projecting to the ipsilateral side of the

brain, or those projecting exclusively via the retinohypothalamic tract (e.g., to the medial

terminal nucleus of the accessory optic system; Yonehara et al., 2009; Gauvain and Murphy,

2015). Thus, here we did not use the term "cell-type" to avoid confusion.

Behavior data analysis

We monitored the behavioral states of an animal during *in vivo* recordings in two ways

(Supplementary Figure 1). First, we monitored the running speed based on the turning speed

of the custom-made rotary treadmill. Second, we videotaped the animal's eye, facing towards

the visual stimulation screen. Eye tracking was done by passing individual frames of the eye

videos to Mask R-CNN, an object detection and segmentation neural network (He et al., 2017;

Abdulla, 2017). Mask R-CNN was fine-tuned with 500 manually annotated frames of our

recordings, covering the range of pupil size and background illumination found in our setup.

The detected masks were fitted with an ellipse, retrieving the center coordinate, semi-major

and -minor axes, and rotation angle (e.g., Supplementary Figure 1A).

Pairwise cross-correlation (100 ms bin) was calculated between the RGC firing rate,

running speed and pupil size (Supplementary Figure 2) for the recording period where the

mean light intensity was relatively stable: i.e., randomly flickering full-field stimulation and

moving grating stimulation periods. The peak value was then selected within 5 s lag while the

p-value was calculated from the data outside the range. The peak cross-correlation was

similarly calculated between the pupil size and the light intensity for the full-field sinusoidally

flickering stimulus and the full-field contrast inversion stimulation periods.

Model analysis

Information theoretic analysis

1042 We took an information theoretic approach to quantify the amount of information carried by

1043 the retinal output spike trains (Figure 7; Rieke et al., 1997), focusing on the randomly flickering

1044 full-field stimulation period (T = 300 s). Specifically, we calculated the information conveyed

1045 by a single spike, I_{spike} bits/spike, using the following formula:

$$ I_{\text{spike}} = \frac{1}{T} \int_0^T dt \left[\frac{r(t)}{\bar{r}} \right] \log_2 \left[\frac{r(t)}{\bar{r}} \right], \tag{8} $$

where $r(t)$ is the instantaneous firing rate of a target cell using small time bins Δt, and $\bar{r}$ is the

average firing rate over time. To avoid the estimation bias due to the finite data size, we

computed I_{spike} in Eq.(8) as a function of bin size Δt, performed a linear regression to the well-

behaved linear portion of these measurements, and extrapolated to the limit $\Delta t \to 0$ (i.e. the

y-intercept). The information rate conveyed by a target cell was then given as: $I_{\text{rate}} = \bar{r} \cdot I_{\text{spike}}$

bits/s.

Rate and latency coding models

We took a modeling approach to examine how retinal coding is affected by the response

characteristics observed under different recording conditions (Supplementary Figure 4;

Gollisch and Meister, 2008). Specifically, using the temporal filter (STA; Figure 5) and static

nonlinear gain function (Eq.(7); Figure 6) in the linear-nonlinear cascade model framework,

we first simulated the responses of each cell to a step change in light intensity at 10 different

levels of contrast increase and decrease, respectively (n_{level} = 20 in total; e.g., Supplementary

Figure 4A,B). To estimate the information contained in a given characteristic x of the simulated

responses, we then used the following formula:

$$I = \sum_s \int dx \, p(s) \, p(x|s) \, \log_2 \frac{p(x|s)}{p(x)}, \qquad (9)$$

where $p(s) = 1/n_{\text{level}}$ is the probability of the stimulus s; $p(x|s)$ is the conditional probability density of x given stimulus s; and $p(x) = \sum_s p(s) \, p(x|s)$ is the total probability density of x, obtained as the normalized sum of $p(x|s)$. For rate coding (Supplementary Figure 4E), we focused on the peak firing rate of the simulated responses, and assumed that $p(x = \text{"peak rate"} \mid s)$ follows a Gaussian distribution with the mean and standard deviation of the peak firing rate and its 2.5% value, respectively (e.g., Supplementary Figure 4C). For temporal coding (Supplementary Figure 4F), we focused on the response latency, where we assumed that the detection threshold was given by 10 ± 2.5% deviation from the baseline firing rate (mean ± standard deviation of a Gaussian distribution), from which $p(x = \text{"latency"} \mid s)$ was calculated (e.g., Supplementary Figure 4D). For stimuli that did not trigger strong enough responses to cross the detection threshold, the response latency was considered to be infinity.

Linear decoding model

We used a linear decoding model to investigate how population coding of the retina depends on the recording conditions (Figure 8). Specifically, we first computed the optimal linear decoder (333 ms window with a bin size of 16.7 ms) from the simultaneously recorded population activity in response to the full-field randomly flickering stimulus (Warland, et al., 1997). The performance of the linear decoder was measured by Pearson's correlation coefficient between the presented and estimated stimuli with 10-fold cross-validation. For each recording condition, the decoding performance as a function of the number of input cells, n, was then assessed by fitting the following exponential saturation function with a plateau at 1:

$$f(n) = 1 - (1 - a)e^{-(n-1)/b}, \qquad (10)$$

where $a = f(1) \in [0, 1]$ represents the single-cell decoding performance; and $b > 0$ is the saturation constant, hence a larger value of b indicates that the linear decoding requires a larger population to improve its performance.

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