Histamine interferes with the early visual processing in mice

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Acronyms: AAV, adeno-associated virus; ANOVA, analysis of variance; DLP, digital light processing; DRN, dorsal raphe nucleus; HDC, histidine decarboxylase; GABA, γ-aminobutyric acid; IQR, interquartile range; IR, infrared; LED, light emitting diode; LGN, lateral geniculate nucleus; OT, optic tract; PFA, paraformaldehyde; PSAM/PSEM, pharmacologically selective actuator/effector module; RF, receptive field; RGC, retinal ganglion cell; STA, spike-triggered average; tSNE, t-distributed Stochastic Neighbor Embedding; UV, ultraviolet.

1   Abstract

Sensory processing is dynamically modulated by an animal's behavior and internal states.

Growing evidence suggests that such modulation starts from early stages, even at the retina,

but the underlying mechanisms remain elusive. Combining pharmacological and

chemogenetic tools with single-unit extracellular recordings in awake head-fixed mice, here

we identified that the visual responses of retinal ganglion cells and the lateral geniculate

nucleus were both made weaker and slower by histaminergic projections from the

tuberomammillary nucleus of the posterior hypothalamus. The observed changes in the visual

responses were, however, not directly linked with histaminergic modulation of pupil dynamics

or locomotion behavior. Our computational modelling analysis instead suggests neuronal

circuit effects, such as gain modulation via the histamine H1 receptors in the retina. As

nocturnal animals, facilitation of visual processing at low histamine level is ethologically

beneficial for mice to respond faster to visual threats when animals are less active during

daytime.

15   Introduction

In the mammalian brain, histaminergic neurons are exclusively located in the

tuberomammillary nucleus (TMN) and surrounding areas of the posterior hypothalamus (Haas

and Panula, 2003). They express histidine decarboxylase (HDC) and innervate widely across

the brain (Lin et al., 2023), including the retina (Gastinger et al., 1999; Repérant et al., 2007;

Warwick et al., 2024). These neurons remain silent during sleep, but increase their firing during

wakefulness and reach their maximal activity at the state of high vigilance, much as other

monoaminergic systems do, such as the serotonergic neurons in the dorsal raphe nucleus

(DRN) of the brain stem (Lin et al., 2023; Takahashi et al., 2006; Thakker, 2011). Brain

histamine exerts its effects via three types of receptors: the H1 and H2 receptors mediate

excitation of postsynaptic cells, while the H3 receptors serve as an auto-receptor to

presynaptically inhibit histamine release (Haas and Panula, 2003). The central histaminergic

system has been implicated in regulating an animal's state and a broad range of behavior,

including the sleep-wake cycle (John et al., 2004; Mochizuki et al., 1992), arousal (Parmentier

et al., 2002; Scammell et al., 2018), suppression of food intake (Khouma et al., 2023; Valdés

et al., 2010; Wang et al., 2015) and learning (Kamei et al., 1993; Klapdor et al., 1994; Orsetti

et al., 2001; Rubio et al., 2002). Histaminergic neurons, however, corelease γ-aminobutyric

acid (GABA) and interact with other neuromodulatory systems, such as the cholinergic and

serotonergic systems (Brown et al., 2002; Khateb et al., 1995; Lin et al., 2023). This makes it

difficult to dissect the effects of histamine by itself on neural circuit functions. In particular,

histaminergic modulation of sensory processing remains largely elusive (Lee and Dan, 2012;

McCormick et al., 2020).

The retina performs the first stage of visual processing, and conveys visual information

to the brain via spike trains of retinal ganglion cells (RGCs). Growing evidence supports that

the activity of RGCs is correlated with various behavioral measures, such as locomotion and

pupil size (Schröder et al., 2020; Liang et al., 2020; Reggiani et al., 2023; Boissonnet et al.,

2023), pointing to a more dynamic role of the retina than is assumed for a deterministic

processor. The vertebrate retina is part of the central nervous system equipped with both

intrinsic and extrinsic neuromodulation mechanisms, much as other brain regions are (Lee

and Dan, 2012; McCormick et al., 2020). For instance, dopamine is synthesized and released

by a group of retinal amacrine cells under the control of circadian rhythm, and contributes to

adjusting the retinal circuitry for rod or cone vision by modulating gap-junction couplings

(Bloomfield and Völgyi, 2009). Such retinal intrinsic mechanisms of neuromodulation are,

however, much slower than needed to modulate retinal function to follow rapid behavioral

changes. To this end, the retina seems to exploit the centrifugal system to receive inputs from

a variety of brain regions in a species-specific manner (Repérant et al., 2007). For example,

histaminergic and serotonergic projections from TMN and DRN, respectively, have been

anatomically described in the mammalian retina (Gastinger et al., 1999; Greferath et al., 2009;

Labandeira-Garcia et al., 1990; Warwick et al., 2024). However, compared to the isthmo-optic

pathway in birds (Wilson and Lindstrom, 2011) or the olfacto-retinal circuit in fish (Esposti, et

al., 2013), little is known about the function of the mammalian centrifugal system.

Combining in vivo single-unit extracellular recordings from either the lateral geniculate

nucleus (LGN) or the optic tract (OT) with pharmacological and chemogenetic tools, here we

explored how histamine affects the early visual processing in awake head-fixed mice. During

the recording sessions, we simultaneously monitored the subject animal's behavior, such as

pupil dynamics and locomotion, to better correlate the effects of histamine on the visual

response properties with those on the behavioral measures. We also took a computational

approach to clarify the net effect of histamine on the visual responses from functional

viewpoints. Because visual processing in the brain fully relies on the signals from the retina,

characterization of the retinal operation in vivo is indispensable to better understand how the

visual system works.

Results

Histamine leads to weaker and slower visual responses in the mouse lateral geniculate nucleus

To investigate how histamine influences the mouse visual system, we first performed in vivo

single-unit extracellular recordings from the lateral geniculate nucleus (LGN) in awake head-

fixed mice, and compared the visual response properties of the recorded cells before and after

perturbing the central histaminergic system (Figure 1A; see Methods for details). Stimulus

ensemble statistical techniques ("reverse correlation"; Chichilnisky, 2001) were employed to

systematically characterize the visual response properties of the recorded cells in the

framework of a linear-nonlinear cascade model (Boissonnet et al., 2023). In particular, using

full-field white-noise stimuli, we analyzed histaminergic effects on 1) the linear temporal filter,

estimated by the spike-triggered average (STA) stimulus, i.e., the mean stimulus that triggered

spiking responses; and 2) the static nonlinear gain function, i.e., an instantaneous mapping of

the STA output to the neural responses (Figure 1B). We also functionally mapped the cells'

receptive field (RF) by calculating the linear spatial filter from the responses to random binary

"checkerboard" stimuli, where the contrast of each spatial location was randomly and

independently inverted over time. To manipulate the histaminergic system, we used two

different approaches (Supplementary Figure 1). The first was the chemogenetic activation of

the central histaminergic system (Magnus et al., 2011). We injected cre-dependent adeno-

associated viruses (AAVs) carrying pharmacologically selective actuator module (PSAM)-

5HT3HC channels into the tuberomammillary nucleus (TMN) of the posterior hypothalamus in

histidine decarboxylase (HDC)-cre transgenic mice, and then achieved selective activation of

the target neurons by systemic administration of the pharmacologically selective effector

molecule (PSEM). The second approach was the use of pharmacological tools in wild-type

mice (Thakkar, 2011). Specifically, we used chlorphenamine (5 mg/kg), an antagonist of a

postsynaptic histamine receptor H1; and ciproxifan (12 mg/kg), an antagonist of the H3 auto-

receptor that presynaptically inhibits histamine release. These drugs effectively down- and up-

regulate the action of histamine in the nervous system, respectively. Thus, resulting changes

in visual processing, if any, were expected to be in opposite directions.

Upon chemogenetically activating HDC+ neurons in TMN, we found that LGN neurons

generally showed weaker and slower visual responses (Figure 1C,F; N=58 from 3 animals).

The mean firing rate during the visual stimulation significantly reduced (median, -5.0 Hz;

interquartile range (IQR), 9.9 Hz; p(U-test)<0.001); and the response gain also significantly

decreased (median, -41%; IQR, 54%; p(U-test)<0.001) that we computed as the ratio of the

estimated static nonlinearity functions before and after PSEM injection. The temporal filter

became elongated regardless of their functional cell types (Supplementary Figure 2), leading

to significantly longer peak latencies (median, +32 ms; IQR, 27 ms; p(U-test)<0.001) and lower

peak frequencies (median, -16%; IQR, 5 %; p(U-test)=0.01).

Consistent results were obtained when histamine release was pharmacologically up-

regulated. Specifically, after the application of ciproxifan (H3 antagonist; Figure 1D,G; N=48

from 2 animals), we found significantly reduced mean firing rates (median, -3.7 Hz; IQR, 5.0

Hz; p(U-test)<0.001) and response gain (median, -30%; IQR, 52%; p(U-test)=0.003), and

significantly longer peak latency (median, +17 ms; IQR, 13 ms; p(U-test)<0.001) and lower

peak frequency (median, -9%; IQR, 14%; p(U-test)<0.001) of the linear temporal filters. These

effects of the pharmacological treatment were less prominent than those of the chemogenetic

manipulation (Figure 1I-L), except for the changes in the receptive field size where the

reduction reached the statistically significant level with ciproxifan (median, -26%; IQR, 38 %;

p(U-test)=0.004, N=28 from 2 animals) but not with the chemogenetic manipulation (median,

-11%; IQR, 32 %; p(U-test)=0.11, N=19 from 2 animals; Supplementary Figure 3A-F).

In contrast, chlorphenamine (H1 antagonist) led to marginal changes in the visual

response properties of LGN neurons in the opposite direction (Figure 1E,H; N=116 from 3

animals). We observed a tendency towards an increased mean evoked firing rate (median,

+1.0 Hz; IQR, 7.2 Hz; p(U-test)=0.09) and response gain (median, +13%; IQR, 87%; p(U-

test)=0.3), and facilitated response kinetics with significantly higher peak frequency of the

linear temporal filter (median, +4%; IQR, 15%; p(U-test)=0.03) but no significant changes in

its peak latency (median, 0 ms; IQR, 5 ms; p(U-test)=0.7) or the receptive field size (median,

+5%; IQR, 34 %; p(U-test)=0.08, N=38 from 3 animals; Figure 1I-L and Supplementary Figure

3A-F). Taken together, these results indicate that histamine reduces the response gain and

slows down the response kinetics in the mouse LGN.

Figure 1: Histamine made the visual responses of LGN weaker and slower in awake head-fixed mice. *A,B*: Schematic diagram of LGN recordings (A) and linear-nonlinear cascade model (B). Reverse-correlation was used to estimate the linear filter and static nonlinearity that represent response kinetics and strength, respectively. *C*: Linear filters of two representative LGN cells before (gray) and after chemogenetic activation of HDC+ cells in TMN (red). *D,E*: Representative LGN linear filters before (gray) and after pharmacological treatment (D, 12 mg/kg ciproxifan, green; E, 5 mg/kg chlorphenamine, orange). *F-H*: Static nonlinearities averaged across LGN population before and after treatment in corresponding colors (F, chemogenetics, N=58 from 3 animals; G, ciproxifan, N=48 from 2 animals; H, chlorphenamine, N=116 from 3 animals): line, median; gray shade, 25 and 75 percentiles. *I-L*: Cumulative distributions of response feature ratio before and after each treatment (in corresponding color): peak latency (I) and frequency (J) of the linear temporal filters, mean evoked firing rate during stimulation (K), and static nonlinearity (i.e., response gain; L). One-way analysis of variance (ANOVA) with Tukey-Kramer correction was used to compute the p-value against saline control (blue, N=93 from 3 animals): *, p<0.05; **, p<0.01; ***, p<0.001.

Supplementary Figure 1: Pharmacological and chemogenetic approaches to manipulate the central histaminergic system in the mouse brain. *A,B*: Viral delivery of PSAM 5HT3HC channel to HDC+ cells in the TMN (marked with dotted lines) of the posterior hypothalamus (A) or those in the anterior hypothalamus (B; control). *C*: Schematic diagram of histamine receptor localizations and their cellular function. Chlorphenamine blocks the postsynaptic receptor H1 (orange) that mediates the action of histamine postsynaptically, while ciproxifan blocks the auto-receptor H3 (green) that inhibits presynaptic histamine release.

Supplementary Figure 2: Functional classification of LGN/RGC visual response types.
A: t-SNE embedding of the STAs of LGN cells. Different markers are used for distinct response
categories (shown in B). *B*: Each panel represents one of the six response categories: slow
off, off, fast off, fast on, on, and slow on. In each panel, each row represents a cell's STA
(color-coded with red and blue indicating positive and negative filter values, respectively); and
the overlaid gray line shows the average STAs in each response type. *C,D*: Corresponding
data for RGCs obtained by in vivo optic tract (OT) recordings methods.

Supplementary Figure 3: Histamine reduced the LGN but not RGC RF size. *A*: Schematic diagram of LGN recordings with white-noise "checkerboard" stimuli. *B*: Estimated spatial filter (receptive field) of two example LGN cells before (top) and after (bottom) chemogenetic activation of HDC+ cells in TMN. Note a decrease of the RF size of this OFF LGN cell (right), while no change was observed for this ON cell (left). *C-F*: Cumulative distribution of the RF size ratio of LGN cells before and after treatment (C; orange, chlorphenamine, N=38 from 3 animals; green, ciproxifan, N=28 from 2 animals; red, PSAM/PSEM for HDC+ cells in TMN, N=19 cells from 2 animals) and corresponding figure panels for the peak latency (D) and frequency (E) of the RF center dynamics as well as for the mean firing rate (F). One-way ANOVA with Tukey-Kramer correction against saline control (N=48 from 3 animals): *, p<0.05; **, p<0.01; ***, p<0.001. *G-L*: Corresponding figure panels for RGCs: chlorphenamine (H1 antagonist), N=6 from 3 animals; ciproxifan (H3 antagonist), N=25 from 4 animals; saline, N=18 from 3 animals.

Histaminergic modulation starts from the mouse retina

Histaminergic neurons are localized solely in the hypothalamus, most densely in TMN, and project their axons widely across the brain (Lin et al., 2023), including the mouse retina (Gastinger et al., 1999; Repérant et al., 2007; Warwick et al., 2024). To test if the observed histaminergic modulation of the visual processing in LGN originates in the retina, we next performed in vivo optic tract (OT) recordings to monitor the retinal output responses in awake head-fixed mice (Boissonnet et al., 2023) before and after manipulating the central histaminergic system (Figure 2).

Consistent with the results in LGN, chemogenetic activation of HDC+ neurons in TMN led to reduced firing (median, -13.0 Hz; IQR, 25.5 Hz; p(U-test)=0.11) with lower response gain (median, -51%; IQR, 50 %; p(U-test)=0.04); and slower response kinetics with elongated linear temporal filters in retinal ganglion cells (RGCs; peak latency change, median, 46 ms; IQR, 22 ms; p(U-test)=0.008; peak frequency change, median, -36%; IQR, 22%, p(U-test)=0.04; N=10 from 2 animals; Figure 2A,D). In contrast, no change in these response parameters was observed when targeting HDC+ populations in the anterior hypothalamus outside TMN for the PSAM/PSEM activation (firing rate change, median, -0.7 Hz, IQR, 16 Hz, p(U-test)=1; gain change, median, 0.5%, IQR, 79%, p(U-test)=1; latency change, median +2 Hz, IQR, 11 Hz, p(U-test)=0.8; frequency change, median -2.9%, IQR, 11%, p(U-test)=0.12; N=23 from 3 animals). This supports that the histaminergic modulation of the early visual system is mediated by the centrifugal projections specifically from TMN (Figure 2G-J and Supplementary Figure 1).

The effects of the pharmacological treatments were overall similar between LGN and RGCs, though certain differences existed. First, pharmacologically blocking the H3 receptors with ciproxifan led to slower response kinetics in both LGN and RGCs (latency change, median, 8 ms; IQR, 14 ms; p(Utest)<0.001; frequency change, median, -7%; IQR, 19%; p(U-test)=0.04; N=40 from 4 animals), but the firing rates and the response gain remained largely unaffected in RGCs (Figure 2B,E; firing rate change, median -0.3 Hz; IQR, 14.2 Hz; p(U-text)=0.8; gain

change, median, -1.3%; IQR, 68%, p(Utest)=0.8). Second, while chlorphenamine (H1

antagonist) drove faster kinetics with marginally higher firing in both LGN and RGCs, the

effects on the temporal filters were primarily on the peak frequencies in LGN (Figure 1E,H)

but on the peak latencies in RGCs (Figure 2C,F; latency change, median, -5 ms; IQR, 10 ms;

p(U-test)=0.004; frequency change, median, +4 Hz; IQR, 19 Hz; p(U-test)=0.7; firing rate

change, median, +2.1 Hz, IQR, 10.4 Hz; p(U-test)=0.14; gain change, median, 5%, IQR, 88%,

p(U-test)=0.2; N=34 RGCs from 3 animals). Finally, RGC receptive fields did not show

significant changes in size (Supplementary Figure 3G-L; ciproxifan, median, -4%, IQR, 54%,

p(U-test)=1, N=25 from 4 animals; chlorphenamine, median, ±0%, IQR, 36%, p(U-test)=1,

N=6 from 3 animals; saline, median, 5%, IQR, 30%, p(U-test)=0.36, N=18 from 3 animals).

Taken together, our electrophysiology data indicate that 1) the mouse early visual processing

is subject to histaminergic modulation controlled by TMN, starting from the retina; 2) histamine

generally suppresses the firing and slows the response kinetics; and 3) response latency and

frequency tuning are likely modulated by different mechanisms.

Figure 2: histamine primarily affected the visual response kinetics of RGCs. *A-C*: Linear filters of two representative RGCs before (gray) and after chemogenetic (A) or pharmacological treatment (B, ciproxifan, green; C, chlorphenamine, orange). Schematic diagram of in vivo optic tract (OT) recordings is displayed on the left. *D-F*: Static nonlinearities averaged across RGC population before and after treatment in corresponding colors (D, chemogenetics, N=10 from 2 animals; E, ciproxifan, N=40 from 4 animals; F, chlorphenamine, N=34 from 3 animals): line, median; gray shade, 25 and 75 percentiles. *G-J*: cumulative distributions of the response feature ratio before and after each treatment (in corresponding colors): peak latency (G) and frequency (H) of the linear temporal filters, mean evoked firing rate during stimulation (I), and static nonlinearity gain function (J). One-way ANOVA with Tukey-Kramer correction (chemogenetics against non-TMN targets, N=23 from 3 animals, dark gray; pharmacology against saline control, N=46 from 4 animals, blue): *, p<0.05; **, p<0.01; ***, p<0.001.

Pupil dynamics and locomotion are irrelevant to histaminergic modulation of early visual processing

Histamine is a potent neuromodulator implicated in various brain functions and behaviors, such as sleep-wake cycles and arousal (Haas and Panula, 2003; Thakkar, 2011). We then cannot deny the possibility that the observed histaminergic effects on the visual response properties in the mouse early visual system are by-products of behavioral modulation. For example, a larger pupil size leads to stronger responses because more photons reach the retina (Boissonnet et al., 2023). To address this issue, we examined the relationship between the histaminergic effects on the RGC/LGN visual responses and simultaneously monitored behavioral measures (Figure 3).

We first analyzed the histaminergic modulation of pupil dynamics (Figure 3A). We found pupil dilation in all 6 mice after chlorphenamine (H1 antagonist) application, but pupil constriction in 7 out of 8 animals when histamine release was enhanced by systemic injection of ciproxifan (H3 antagonist). Chemogenetic activation of the central histaminergic system, however, led to pupil dilation in most animals (4 out of 5 mice). Thus, the polarity of the changes in pupil size was not consistent with that of RGC/LGN visual responses. This suggests that the histamine-driven changes of pupil size cannot explain the modulation of the early visual responses.

We next examined how histamine affects locomotion of the head-fixed animals on the treadmill during recordings (Figure 3B,C). These mice often stayed still and moved fast ("running" at >2 cm/s) only about 20% of the time on average. In general, none of the perturbations of the histaminergic system strongly affected the time spent running in most mice (Figure 3B). When they did, however, we found that the chemogenetic activation of HDC+ neurons in TMN resulted in a substantial increase of no-running periods (2 out of 6 mice), whereas both chlorphenamine and ciproxifan led to a substantial decrease of no-running periods (2 out of 6 and 1 out of 7 mice, respectively). Furthermore, the mean running

speed during the recordings decreased upon the chemogenetic activation, whereas it varied

in both directions after the pharmacological treatments (Figure 3C). Thus, changes in

locomotion driven by histamine were not in agreement with the histaminergic effects on the

RGC/LGN visual responses (Figures 1 and 2).

Taken together, neither pupil dynamics nor locomotion behavior was causally linked

with the histaminergic modulation of the mouse early visual system. We conclude that the

central histaminergic system modulates the mouse vision, effectively leading to slower and

weaker responses.

Figure 3: histaminergic modulation of pupil dynamics and locomotion behavior was not

causally linked with that of RGC/LGN visual responses. *A*: the ratio of the average pupil

size before and after treatment (in corresponding colors). *B*: the ratio of the "no-running"

periods (threshold, 2 cm/s) before and after treatment. *C*: the ratio of the average locomotion

speed during recording sessions before and after treatment.

Response gain modulation is more strongly correlated with peak latency modulation than with peak frequency modulation

What are the mechanisms underlying the histaminergic modulation of early visual processing?

To address this question, we next took a computational approach to examine the relationship

between different aspects of the visual response properties that we identified to be affected

by histamine (Figure 4). In particular, we simulated the activity of an integrate-and-fire neuron

with Poisson spiking and refractoriness in response to white-noise input stimuli (Figure 4A),

and processed the output spike trains in the framework of the linear-nonlinear cascade model

to characterize the simulated response properties in the same way as the experimental data

(Figure 1A). Here we varied two model parameters, gain and resting baseline, to examine the

effects of their changes on the following four aspects of the simulated responses: spike counts,

the peak latency and frequency of the estimated linear temporal filter, and the slope of the

estimated static nonlinearity expressed as half-wave rectification. The rest model parameters

were fixed in all simulations, including the underlying linear filter given by a trigonometric

function with logarithmic scale in time, and the refractory period filter given by an exponential

function. The spike threshold was drawn from a random uniform distribution ranging from 0 to

1 for each time point to approximate the Poisson spiking properties.

When the model neuron had a higher gain or higher resting baseline, subthreshold

signal reached the spike threshold faster and more frequently; hence, total spike counts

became higher (Figure 4B) and the peak latency of the estimated linear temporal filter

appeared shorter (Figure 4C). In contrast, the apparent peak frequency of the simulated

responses was dependent more strongly on the resting baseline than on the gain (Figure 4D).

These changes in the estimated linear temporal filter properties were consistent with the

experimental observation, where the magnitude of the response gain modulation by

histaminergic perturbation was significantly correlated with that of the peak latency modulation

(Figure 4F,H), but much less with that of the peak frequency modulation (Figure 4G,I). This

agreement between the observed and simulated data indicates that, despite the simplicity, our

model captures aspects of neuronal response modulation to a first approximation.

The model analysis supports that the histaminergic effects on RGCs should involve at

least gain and baseline modulations. First, the H1 receptors likely mediate the gain modulation

in RGCs, given that chlorphenamine (H1 antagonist) led to changes in the peak latency but

not the peak frequency, along with changes in the estimated static nonlinearity largely by

scaling (Figure 2). Second, the peak frequency was modulated as well as the peak latency of

RGCs upon chemogenetic activation of HDC+ neurons in TMN, accompanied by a shift of the

estimated static nonlinearity. This implies changes in the baseline resting potential of the cells.

Blocking the H3 receptors, however, led to changes in the RGC response kinetics without

much affecting the mean firing rates. Furthermore, the net effects of histamine perturbations

on LGN responses were more diverse (Figure 1) and not fully explainable by the simple model

(Figure 4). Thus, we expect that mechanisms besides gain and baseline modulations are also

involved, such as neural circuit effects through retinal amacrine cells for example (Horio et al.,

2018).

Figure 4: Gain modulation accompanies the peak latency modulation but not the peak
frequency modulation via histaminergic effects. *A-E*: Using a computational model
outlined in A, here we examined the consequences of varying the baseline and gain levels on
the response features, such as spike counts (B), peak latency (C) and frequency (D) of the
estimated linear filter from the simulated responses, and slope of static nonlinear gain function
of the model neuron (E). See Methods for details. *F,G*: Relationship between the magnitudes
of the gain modulation and response kinetics modulation by perturbing the histaminergic
system for LGN neurons (chemogenetics, red; ciproxifan, green; chlorphenamine, orange;
N=222 from 8 animals). Cell types are indicated by different markers (Supplementary Figure
2). A stronger correlation was found for the peak latencies (F, R = –0.42, p<0.001) than for
the peak frequencies (G, R = 0.17, p=0.01). *H,I*: corresponding figure panels for RGC
responses (N=84 from 9 animals). Significant correlation to the gain modulation was found for
the peak latency modulation (R = –0.25, p = 0.02), but not for the peak frequency modulation
(R= 0.20, p = 0.07).

Discussion

Here we provide in vivo electrophysiological evidence for histaminergic modulation of the

mouse early visual system: the more the histamine released, the slower and weaker the LGN

and RGC visual responses in awake head-fixed mice (Figures 1 and 2). This is opposite from

what one would expect for the effect of histamine, given that the level of brain histamine

typically increases as animals become more aroused (Thakker, 2011). At the behavioral level,

antihistamine drugs lead to drowsiness in humans, impairing sensory processing (van

Ruitenbeek et al., 2009). At the neurophysiological level, it has been shown that histamine

leads to slow depolarization of LGN relay cells in cats and guinea pigs (McCormick and

Williamson, 1991; Uhlrich et al., 2002). Histamine was thus expected to facilitate visual

processing, unlike what our observations indicated. These discrepancies may arise from two

reasons. First, the effects of histamine can be species-specific. In particular, many prey

species often sleep with their eyes open, including mice. Furthermore, as nocturnal animals,

mice are less active during daytime. Thus, facilitating visual responses at low histamine level

may be ethologically beneficial for mice to respond faster to visual threats during daytime,

especially when mice are dormant. Future studies in the context of sleep-wake cycles as well

as in diurnal animals will give further insights into this question.

Second, our unexpected results may be due to differences in experimental conditions.

Electrophysiological studies on neuromodulation were conventionally conducted in vitro or in

anesthetized animals (McCormick and Williamson, 1991; Uhlrich et al., 2002; Warwick et al.,

2024). It has been shown, though, that substantial differences exist in the early visual

processing between awake and anesthetized conditions (Durand et al., 2016), including the

retinal output (Boissonnet et al., 2023). Here we took a computational modelling approach to

better understand the net effect of histamine from functional viewpoints, rather than pursuing

the underlying biophysical mechanisms. We suggest that the H1 receptors mediate gain

modulation in RGCs: chlorphenamine (H1 antagonist) affected the response latency but not

the frequency tuning (Figure 2), a distinctive feature of gain modulation identified by our

computational modelling analysis (Figure 4). This action of histamine does not have to be

directly on RGCs expressing the H1 receptors, but can be mediated indirectly by circuit

mechanisms. Indeed, histamine affects both inward and outward currents of amacrine cells in

the mouse retina (Horio et al., 2018). The effects of histamine can then be more profound than

gain modulation, much as we observed diverse outcomes in RGCs and LGN when different

pharmacological and chemogenetic perturbation tools were employed (see also Warwick et

al., 2024). Furthermore, in awake conditions, the histaminergic system acts in concert with

other neuromodulatory systems (McCormick et al., 2020). For example, serotonergic neurons

in the dorsal raphe nucleus have been suggested to project to the retina (Schütte, 1995;

Repérant et al., 2000; Gastinger et al., 2006). As the brain state is dynamically controlled by

multiple neuromodulators, it will be critical to further characterize neuromodulation of the retina

and downstream visual areas in various behavioral contexts.

What is the biological relevance of histaminergic modulation of the early visual system,

starting at the retina? Growing evidence supports significant correlations between awake

retinal responses and various behavioral measures, such as pupil dynamics and locomotion

(Schröder et al., 2020; Liang et al., 2020; Boissonnet et al., 2023). We, however, show that

resulting changes in pupil size and locomotion behavior by manipulating the histaminergic

system were not always consistent with the effects on LGN or RGC visual responses (Figure

3). Therefore, the observed changes in their visual responses were unlikely to be associated

with behavioral modulation by histamine. This in turn suggests that histamine may not fully

explain the fast modulation of RGCs by arousal or locomotion. Additional mechanisms of

retinal modulation should then exist, such as the centrifugal serotonergic system (Schütte,

1995; Repérant et al., 2000; Gastinger et al., 2006). Further studies on the function of the

retina as an integrated part of the whole organism, but not as an isolated tissue, will be critical

to better understand sensory information processing in awake animals under natural

conditions.

Materials and Methods

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

0.05 (with multiple comparison correction where appropriate) in all analyses unless noted

otherwise. All experiments were performed under the license 233/2017-PR and 220/2024-PR

from the Italian Ministry of Health, following protocols approved by the Institutional Animal

Care and Use Committee at European Molecular Biology Laboratory. The data analyses were

done in Python and Matlab.

Animals

384 Animals were housed on a 12h light-dark cycle, with ad libitum access to water and food. All

385 experiments were performed on female mice. A total of 20 wild-type (C57BL6/J;

386 RRID:IMSR_JAX:000664) and 10 hemizygous HDC-cre (Hdctm1.1(icre)Wwis/J;

387 RRID:IMSR_JAX:021198) mice were used for pharmacological and chemogenetic

388 manipulations, respectively.

Surgical procedures

All surgical procedures were performed in animals from 4 to 19 weeks old (9.2 weeks median

age). Before the surgery, animals were injected with Caprofen (5 mg/Kg) and then

anaesthetized with isoflurane (4% for induction, 1% for maintenance in O_2). Throughout the

surgery, the temperature of the animals was kept stable at 37°C using a heating pad

(Supertech Physiological). Ointment (VitA-Pos, Ursapharm) was applied on both eyes to

prevent them from drying during the surgery, and the animals were placed in a stereotaxic

frame (Stoelting 51625). A portion of the scalp was removed to expose the skull and the

periosteum was scraped away with a round scalpel to increase adherence of the dental

cement. For those mice belonging to the chemogenetic manipulation group, the skull was

aligned in order to level bregma and lambda on the same horizontal plane, and a 100 μm

craniotomy was performed using a motorized drill attached to the stereotaxic arm. Viral

solution (250 nL; AAV9::FLEX-PSAM Y115F,L141F:5HT3 HC; Vector Biosystems, VB773)

was injected with an automatic injector pump at 1nL/sec, at the coordinate of AP: -2.6, ML:

±0.8, DV:-5.3. All animals were then implanted with a custom-made titanium head plate,

cemented on the skull using a mixture of cyanoacrylate (Loctite 401, Henkel) and dental

cement (Paladur, Kulzer). The head plate featured a hole (8 mm diameter), allowing for clear

observation of both bregma and lambda. The skull surface was then glazed with a thick layer

of cyanoacrylate in order to support the skull with mechanical, atmospheric and biological

protection, while still allowing for visual identification of reference points. After the surgery

animals were allowed to recover on a heating pad and then housed in individual cages. During

the following seven days, the mice were administered with painkillers (Caprofen) diluted in the

drinking water at a concentration of 50 mg/mL.

In vivo electrophysiology

After recovery from the surgery, animals were habituated to head fixation by placing them in

the experimental apparatus over the course of one week, twice a day for up to two hours. On

the day of the recording, we first placed the subject animal in the recording set up with its head

fixed, and determined the electrode penetration path 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, an acute silicone probe (Buzsaki32L, Neuronexus; or P2,

Cambridge Neurotech) coated with a fluorescent dye (DiI stain, Invitrogen, D282) was lowered

at 5 µm/s using the robotic arm until visual responses were found in the target area.

After presenting a battery of visual stimuli, we briefly anesthetized the animal and

injected 0.9% saline solution alone or that containing chlorphenamine (H1 antagonist, 5

mg/kg), ciproxifan (H3 antagonist, 12 mg/kg), or PSEM (5 mg/kg). About 20 minutes after the

injection and removal of the anesthesia, we presented the same battery of visual stimuli to the

animal for the second recording session.

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

retracting the silicon probe, the mice were anesthetized (2.5% Avertin, 16 µL/g, intraperitoneal

injection) and perfused with paraformaldehyde (PFA; 4% in phosphate buffer solution). 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.

Visual stimulation

Visual stimuli were presented as described before (Boissonnet et al., 2023). Briefly, visual

stimuli were projected onto a spherical screen (20cm in radius) painted with ultra-violet (UV)

reflective white paint, placed 20 cm from the mouse's left eye. A gamma-corrected digital light

processing device (DLP, Texas Instruments, DLPDLCR3010EVM-LC) was used as a light

source after the green and the red LEDs were replaced with UV (365nm, LZ1-00UV00, LED

Engine) and infrared (IR: 950nm, SFH 4725S, Osram) LEDs respectively. The IR light was

used as a synchronization signal and was recorded with a photodiode (PDA100A2, Thorlabs).

Stimulation was conducted at 60 frames per second (FPS), and covered 73° in azimuth and

44° in altitude from the mouse eye position. The maximum light intensity at the eye position

was 31 mW/m^2 (15.4 mW/m^2 for UV LED and 15.9 mW/m^2 for blue LED; measured with S121C

sensor, Thorlabs), leading to mesopic to photopic conditions. Using QDSpy software (Franke

et al., 2019), we presented the following stimuli: a randomly flickering full-field stimulus (5

minutes) consisting of a random sequence of black and white frames at 60 FPS; and a black-

and-white binary dense noise stimulus (15 minutes), consisting of a 32-by-18 pixels

checkerboard, where each pixel followed an independent random sequence at 60 FPS but the

overall luminance of the frame was kept constant at the mean value.

Electrophysiology data analysis

Raw data from the same animal before and after treatment were concatenated and processed

as a single recording. Spike sorting was performed with Kilosort2 for semi-automatic cluster

detection and Phy for visualization and data curation. In total, we recorded 311 RGCs from 18

animals and 421 cells in LGN from 12 mice.

For systematically characterizing the visual response properties, we used stimulus

ensemble statistical techniques ("reverse correlation" methods; 500 ms window; $\Delta t = 1/60$ s

bin size) to calculate the linear filter and static nonlinear gain function of the recorded cells in

response to white-noise stimuli as described before (Figure 1A; Boissonnet et al., 2023).

Briefly, we first 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 response kinetics, we then estimated the peak latency by fitting a

difference-of-Gaussian curve to the linear filter; and the spectral peak frequency by the Fourier

analysis on the linear filter. Spatial response properties were examined for STAs from

"checkerboard" stimulation. In particular, we fitted a two-dimensional Gaussian envelope to

the spatial filter at the peak latency (e.g., Supplementary Figure 3B,H), and the receptive field

(RF) size was estimated as twice the mean standard deviation (SD) of the long and short axes.

The linear temporal filters from full-field white-noise stimuli were used for broadly

classifying the response types of RGCs and LGN cells, respectively (Supplementary Figure

1). Specifically, we first used t-Distributed Stochastic Neighbor Embedding (t-SNE) to map the

STAs onto two-dimensional space, and then used K-means++ algorithm for heuristically

categorizing the responses into the following six types: fast ON, ON, slow ON, fast OFF, OFF,

and slow OFF.

Static nonlinear gain function of each cell (bin size, 0.1) was computed as: P(response

| stimulus) = N(stimulus | response) / N(stimulus) / Δt, where N(stimulus | response) and

N(stimulus) are the distributions of spike-triggered stimulus ensembles projected onto the L2-

normalized STA and the entire stimulus ensembles, respectively. The ratio of P(response |

stimulus) before and after the treatment was used as a measure of response gain modulation

for each cell.

Behavioral data analysis

For all experiments, the mouse's left eye – i.e., the side presented with visual stimuli – was

recorded using an IR camera (The Imaging Source, DMK23UV024) to monitor its motion and

pupil dynamics at 30-60Hz. Pupil detection was done using a Mask-Region based

Convolutional Neural Network (Mask-RCNN), trained as described previously (Boissonnet et

al., 2023). Two-dimensional ellipse was fit to the pupil, and the large axis diameter was used

as a measure of the pupil size. We then calculated the mean pupil size during the recording

sessions, and used the ratio before and after pharmacologic or chemogenetic treatment as a

measure of the treatment effects on the pupil dynamics (Figure 3A).

We also monitored the running speed of an animal based on the turning speed of the

custom-made rotary treadmill throughout the recordings. We set a threshold at 2 cm/s to detect

running behavior. The animals, however, did not run frequently on the treadmill (17 ± 25%,

mean ± standard deviation; 30 animals). As a measure of the treatment effects, we thus used

a) the ratio of the fraction of the "non-running" period (Figure 3B); and b) that of the average

running speed during the recording session before and after pharmacological or chemogenetic

treatment (Figure 3C).

Model analysis

To explore how different aspects of neuronal intrinsic properties and output responses are

interrelated between each other, we simulated a model neuron's responses to white-noise

stimuli using different model parameter values (Figure 4A), and characterized these simulated

responses using a linear-nonlinear cascade model as we did for the experimental data.

Specifically, we employed an integrate-and-fire neuron model with Poisson spiking and

refractoriness. A model neuron's discrete linear filter (of length $n_T = 25$) was given by a

trigonometric function with logarithmic scale in time: i.e., $L(i) = \sin[2*\pi*((10^{\wedge}t_i-1)/(10^{\wedge}s_t-1))]$,

where $s_t = 1.5$ is a temporal spacing parameter and $t_i = i*s_t/(n_T-1)$ for $i=0,\ldots,n_T-1$. The spike-

generating function $V(i)$ was then modelled by first projecting the white-noise Gaussian

stimulus $S(i)$ (of length 1,000,000; normalized to have a maximum value of 1) onto $L(i)$,

followed by the addition of baseline B and the multiplication by gain G: i.e., $V(i) = G*(\Sigma_j L(j)*S(i-$

$j) + B)$. For each simulation, we chose the value of B from -0.45 to 0.45 in steps of 0.15; and

that of G from $1.2^{\wedge}g$ with g = -3, -2, -1, 0, 1, 2, 3. The spiking response was then generated

by thresholding $V(i)$ (at a random threshold for every time point derived from a uniform

distribution from 0 to 1) and refractoriness (given by an exponential filter: $-\exp[-i/s_h]$ for i =

$0,\ldots,5$ with $s_h = 1$). The output spike trains were then subject to the reverse-correlation analysis

as described above to characterize the apparent linear filter and static nonlinearity of the

model neuron, much as we did for the experimentally recorded RGCs and LGN cells.

Specifically, we focused on the peak latency and frequency of the estimated linear filters and

the slope of the estimated static nonlinear gain function, and examine how these features

depend on the baseline B and the intrinsic gain G of the model neuron (Figure 4B-E).

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