RESEARCH ARTICLE

Classical Photoreceptors Are Primarily Responsible for the Pupillary Light Reflex in Mouse

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Abstract

Pupillary light reflex (PLR) is an important clinical tool to assess the integrity of visual pathways. The available evidence suggests that melanopsin-expressing retinal ganglion cells (mRGCs) mediate PLR—driven by the classical photoreceptors (rods and cones) at low irradiances and by melanopsin activation at high irradiances. However, genetic or pharmacological elimination of melanopsin does not completely abolish PLR at high irradiances, raising the possibility that classical photoreceptors may have a role even at high irradiances. Using an inducible mouse model of photoreceptor degeneration, we asked whether classical photoreceptors are responsible for PLR at all irradiances, and found that the PLR was severely attenuated at all irradiances. Using multiple approaches, we show that the residual PLR at high irradiances in this mouse was primarily from the remnant rods and cones, with a minor contribution from melanopsin activation. In contrast, in rd1 mouse where classical photoreceptor degeneration occurs during development, the PLR was absent at low irradiances but intact at high irradiances, as reported previously. Since mRGCs receive inputs from classical photoreceptors, we also asked whether developmental loss of classical photoreceptors as in rd1 mouse leads to compensatory takeover of the high-irradiance PLR by mRGCs. Specifically, we looked at a distinct subpopulation of mRGCs that express Brn3b transcription factor, which has been shown to mediate PLR. We found that rd1 mouse had a significantly higher proportion of Brn3b-expressing M1 type of mRGCs than in the inducible model. Interestingly, inducing classical photoreceptor degeneration during development also resulted in a higher proportion of Brn3b-expressing M1 cells and partially rescued PLR at high irradiances. These results suggest that classical photoreceptors are primarily responsible for PLR at all irradiances, while melanopsin activation makes a minor contribution at very high irradiances.
Introduction

Broadly, classical photoreceptors (rods and cones) produce image-forming vision, while melanopsin-expressing retinal ganglion cells (mRGCs) are considered to be responsible for non-image-forming vision. However, this distinction has blurred with several recent discoveries. For example, classical photoreceptors are also involved in non-image-forming functions, such as circadian activity and pupillary light reflex (PLR), and mRGCs also contribute to pattern vision [1–9]. The majority of the work designed to unravel the differential roles of classical photoreceptors and mRGCs has been carried out using genetically-engineered mice. Since genetic alterations—especially when introduced during development—can result in a variety of compensatory biological responses, the results from using such animal models may be difficult to interpret [5,10–15]. For example, knocking out TRβ (thyroid hormone receptor) gene, which is involved in the development of m-cones, results not only in lack of m-opsin but also in upregulation of s-opsin and melanopsin [15]. Since mRGCs are known to receive signals from photoreceptors [16–20], loss of photoreceptors during development could potentially cause compensatory changes in mRGCs, making it difficult to understand their specific contributions to vision.

Loss of photoreceptors during development, caused by genetic mutation (eg, in PDE6βrd1, rd/rd-cl or gnat–/–;cnga3–/– mice), results in elimination of PLR at low irradiances but leaves it intact at high irradiances [1,7,21–23]. This suggested that photoreceptors are responsible for PLR only at low irradiances. Knocking out the melanopsin gene in these mice (PDE6βrd1; Opn4–/– or gnat–/–;cnga3–/–;Opn4–/–) eliminates PLR at all irradiances, suggesting that melanopsin is responsible for PLR at high irradiances [1,22]. However, this is inconsistent with the observations that knocking out the melanopsin gene (Opn4–/–) or acutely blocking the melanopsin activation with a specific antagonist in an otherwise normal mouse causes only 10%–15% reduction in the high-irradiance PLR [1,21,22,24].

Here, we explored the possibility that classical photoreceptors are responsible for PLR at all irradiances. Since classical photoreceptors provide input to mRGCs, it was possible that loss of classical photoreceptors during development leads to a compensatory response in mRGCs which then take over the PLR function at high irradiances. To test these hypotheses, we measured PLR in adult mouse in which classical photoreceptor degeneration was induced pharmacologically after normal retinal development; we predicted that unlike in rd1 mouse the PLR would be attenuated at all irradiances. To look for the potential compensatory cellular response in rd1 mouse, we studied Brn3b-expressing mRGCs which have been shown to mediate PLR [25].

Materials and Methods

Animals and Drug Injections

All experiments reported here were approved by the Institutional Animal Ethics Committee of the National Brain Research Centre, India. Adult (8–12 weeks) wild-type (C57BL/6J) and rd1 mice (CBA/J) were used. In addition, wild-type pups (P-8 to P-12) and pregnant mice were used for some experiments (see more details in Results). Mice were obtained from Jackson Laboratory (Bar Harbor, USA) and bred locally at the animal facility of the National Brain Research Centre, India. All animals were maintained on a 12-hour light:dark cycle with an average ambient daylight of approximately 200 lux (measured with an IL1400 photometer, International Light Technologies, Peabody, MA).

Some adult male mice (both wild-type and rd1) received a single injection of N-methyl-N-nitrosourea (MNU; 65 mg/kg in 0.05% acetic acid; i/p; Sigma-Aldrich), sodium iodate (NaIO3;
50 mg/kg; i/v; Sigma-Aldrich), or both MNU and NaIO₃ [26,27]. MNU was also injected in some wild-type mice during development, either postnatally (P-8 to P-12; 60 mg/kg; i/p) or prenatally (in pregnant mothers at E-16; 5 mg/kg; i/p) [28]. Similarly, NaIO₃ was injected in some wild-type pups at P-8 (60 mg/kg; i/p) [26].

To block melanopsin activity, we injected in some animals a specific melanopsin antagonist (1-(2,5-dichloro-4-methoxy-benzenesulfonyl)-piperidine (AA41612; 10 mg/kg, i/p; Princeton BioMolecular Research, Inc., Monmouth, NJ) [24]. The drug was prepared in dimethyl sulfoxide (Sigma-Aldrich) and phosphate-buffered saline (PBS) containing 18.2% hydroxypropyl-beta-cyclodextrine and 0.009M methane sulphonic acid (Sigma-Aldrich).

Pupillary Light Reflex

The PLR was recorded between Zeitgeber Time 3 (ZT3) and ZT9 (ZT0: light turning on at 6 am) [7]. A mouse was dark adapted for 1 hour, lightly anesthetized with ketamine (50 mg/kg) and xylazine (5 mg/kg), and placed on a platform in a dark room. The light sedation was given to avoid using restraints, which can cause variable magnitude of anxiety and therefore variability in the measured PLR [7,29]. Consensual PLR was recorded using an infrared video camera (frame rate 30 Hz; DCR-HC96, Sony, Japan). White light from a 100 W halogen bulb was shone in one eye through a light guide connected to the lamp source (color temperature, 3100K; LG-PS2, Olympus, Japan) while recording the pupil response from the other eye. A custom-made barrier blocked the light from falling on the recorded eye. Neutral density filters (Olympus, Japan) were used to cover approximately 8 log units of irradiance (10⁻⁴ μW/cm² to 2.8 x 10⁴ μW/cm²). Following the baseline recording for 60 sec in dark, PLR was recorded at different light intensities, presented in increasing order of intensity. Each stimulus was presented for 20 sec, which was followed by 100 sec of dark adaptation. The PLR in mice which received a melanopsin antagonist was recorded before the drug injection and again between 20 and 30 min after the injection [24]. For these animals, since the 10-min window was not sufficient to record the PLR at all irradiances, some of the irradiances were skipped, but we covered the full range of irradiances used for other animals.

The video frames were analyzed using imageJ software (NIH); the pupil boundary was marked using the circle tool in imageJ and the area within the boundary computed. The pupil constriction at a given light intensity was calculated as percentage change from the baseline pupil area. The initial constriction, which occurred at 2–3 sec after the stimulus onset, was taken as the transient pupil response. The constriction during the last 5 seconds of stimulus presentation (mean of 5 values at 1-second intervals during 16th to 20th second after the stimulus onset) was taken as sustained pupil response. The sustained pupil constriction is presented as PLR, unless noted otherwise.

Characterization of primary antibodies

The primary antibodies used here are described in Table 1. The Brn3a antibody recognizes a 47 kDa band in Western blot (manufacturer’s specification). The specificity of the Brn3b antibody was confirmed by preadsorption with the appropriate Brn3b peptide (SC-6026P, Santa Cruz Biotechnology) (not illustrated). The melanopsin antibody has been extensively used previously. The specificity of this antibody has also been verified by the lack of immunoreactivity in melanopsin knockout mouse [30,31]. The melanopsin antibody used here for Western blotting (Table 1) detects two bands: one for glycosylated melanopsin at 85 kDa and the other for unglycosylated melanopsin at 53 kDa (manufacturer’s specifications) [32]. The α-tubulin antibody, which was used for loading control in Western blotting, detects a single band at 55 kDa (manufacturer’s specifications) [33].
To measure the extent and pattern of rod and cone loss, retinal flatmounts were immunolabeled for rhodopsin (rods), m-opsin (m-cones), or s-opsin (s-cones). The antibody against rhodopsin showed a strong labeling of the outer segments and weak labeling of the inner segments and the cell bodies of rods [27], confirming the antibody specificity. The antibodies against m-opsin and s-opsin detect 40 kDA and 39 kDa bands, respectively, in Western blotting (manufacturer’s specification). Since long wavelength (red) cones are absent in mouse, the red/green opsin antibody labeled only green cones.

### Immunohistochemistry

Retinal flatmounts were prepared from the mice used for PLR measurements and immunolabeled as described previously [9]. Briefly, eyes were removed after cervical dislocation, and eyecups were fixed in 4% paraformaldehyde (PFA; pH 7.4) at 4°C for 1 hour. Retina was isolated from the eyecup, incised radially or cut into 3–4 radial pieces, and flattened on a membrane filter (Millipore, Billerica, MA). For double labeling with melanopsin and Brn3b antibodies, the flatmounts were first incubated for 1 hour in a blocking solution containing 10% normal donkey serum (NDS), 3% bovine serum albumin (BSA), and 0.3% Triton X-100 in PBS (pH 7.4). The samples were then incubated in a mixture of antibodies against melanopsin and Brn3b at 4°C for 3 days. After washing in PBS for 5 x 5-min, the samples were incubated in the appropriate secondary antibody (donkey anti-rabbit conjugated with Alexa Flour 488 [1:500] for melanopsin and donkey anti-goat conjugated with Alexa Flour 594 [1:500] for Brn3b; both from Molecular Probes, Eugene, OR) for 1 hour at room temperature. The samples were washed again in PBS for 5 x 5-min and mounted in Vectashield containing 4-,6-diamidino-2-phenylindole (DAPI; Vector Laboratories Inc., Burlingame, CA).

To assess the effect of photoreceptor loss on density of non-mRGCs, retinal flatmounts of adult wild-type, rd1 and MNU-injected mice were immunolabeled for Brn3a [9]. The blocking solution was same as mentioned above, and the secondary antibody was donkey anti-mouse conjugated with Alexa Flour 488 (1:500).

To quantify the number of rods, m-cones and s-cones, we used rhodopsin, m-opsin and s-opsin antibodies, respectively. The retinal wholemounts were incubated for 1 hour in a blocking solution containing 3% NDS, 1% BSA and 0.3% Triton X-100 in PBS (pH 7.4). The samples

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**Table 1. List of primary antibodies.**

| Antigen   | Immunogen                                                                 | Source                                  | Species, type | Dilution  |
|-----------|---------------------------------------------------------------------------|-----------------------------------------|---------------|-----------|
| Brn3a     | 1–109 N-terminal amino acids of mouse Brn3a                               | Santa Cruz Biotechnology; #sc-8429      | Mouse, monoclonal | 1:250     |
| Brn3b     | 397–410 C-terminal amino acids (MDSPGPRVLSLTLQ) of human Brn3b            | Santa Cruz Biotechnology; #sc-6026      | Goat, polyclonal | 1:250     |
| Melanopsin| a synthetic peptide consisting of 1–15 N-terminal amino acids of mouse melanopsin | Advanced Targeting Systems; #UF006 | Rabbit, polyclonal | 1:7500 (immunohistochemistry) |
| Melanopsin| A synthetic peptide consisting of amino acids 1–19 of rat melanopsin      | Abcam; ab19306                          | Rabbit, polyclonal | 1:2000 (Western blotting) |
| M-Opsin   | Recombinant human red/green opsin (last 38 C-terminal amino acids)       | Chemicon; # AB5405                     | Rabbit, polyclonal | 1:500     |
| S-Opsin   | Recombinant human blue opsin (last 42 C-terminal amino acids)             | Chemicon; #AB5407                      | Rabbit, polyclonal | 1:1000    |
| Rhodopsin | Bovine rhodopsin (clone: Rho 1D4; last 9 C-terminal amino acids)          | Chemicon; # MAB5356                    | Rabbit, polyclonal | 1:5000    |
| β-Tubulin | Tubulin derived from rat brain                                            | Sigma; #T4026                          | Mouse, monoclonal | 1:16000   |

To quantify the number of rods, m-cones and s-cones, we used rhodopsin, m-opsin and s-opsin antibodies, respectively. The retinal wholemounts were incubated for 1 hour in a blocking solution containing 3% NDS, 1% BSA and 0.3% Triton X-100 in PBS (pH 7.4). The samples...
were then incubated overnight in respective primary antibodies at 4°C. After washing in PBS for 5x 5-min, the samples were incubated in the appropriate secondary antibody (donkey anti-mouse conjugated with Alexa Fluor 488 (1:500) for rhodopsin; donkey anti-rabbit conjugated with Alexa Fluor 488 (1:500) for m-opsin; donkey anti-rabbit conjugated with Alexa Fluor 488 (1:500) for s-opsin (all from Molecular Probes, Eugene, OR) for 1 hour at room temperature.

**Imaging and morphometric analyses**

The imaging and analyses were performed as described previously [9]. Briefly, the images of immunolabeled flatmounts were captured on an epifluorescent microscope (AxioImager.Z1; Carl Zeiss, Gottingen, Germany) fitted with a CCD camera (AxioCam MRm). The microscope was equipped with ApoTome grid projection system, which allowed capturing images at higher contrast and enhanced optical resolution in z-axis. M1 and non-M1 type of mRGCs were identified based on their dendritic stratification in the IPL. For tracking the dendrites of melanopsin cells, several 1-μm thick serial optical sections were captured in z-axis to cover the GCL and the IPL. The images were taken such that the relatively bright (likely M1) cells were near the center of the field. If the dendrites of these cells stratified in outer IPL, we then determined whether any of the other dendrites stratified in inner IPL, to differentiate between M1 and M3 cells. The remaining cells were identified as M2 or other non-M1 cells. Once a melanopsin cell was identified as M1 or non-M1 type, we examined each cell for Brn3b expression. For this, we enhanced the image contrast to be able to detect even the very faint Brn3b expression [9]. Total mRGCs, M1 and non-M1 mRGCs, and Brn3b-expressing mRGCs were counted and density was computed.

To quantify the number of rods and cones, images of labeled retinas were captured using a confocal (LSM 510 Meta, Carl Zeiss) or the epifluorescent microscope. For cones, overlapping images were captured, and stitched using Photostitch software (Canon, USA) to generate a montage of the whole retina. The m-cones and s-cones were counted in these montages using ImageJ. For rods, we captured non-overlapping images to cover a large part of a retina, and measured the labeling intensity. All z-sections with any rhodopsin signal in a frame were merged while averaging the labeling intensity at each pixel. The integrated density of rhodopsin labeling was computed by summing intensity values of all the pixels after subtracting the background.

**Western blotting**

Western blotting was performed as described earlier [34]. Briefly, a 20 μg of each tissue sample was electrophoresed in 9% SDS-PAGE and then transferred to a polyvinylidene fluoride (PVDF) membrane. The blots were incubated for 1 hour at room temperature in a blocking solution (5% BSA in Tris-buffered saline with 0.1% Tween-20 (TBST), pH 7.4) and then in melanopsin antibody (1:2000) overnight at 4°C. β-tubulin was used as loading control. This was followed by 3x 10-min washes in TBST, followed by incubation in the secondary antibody at room temperature for one hour. The secondary antibodies used were horseradish-peroxidase-conjugated anti-rabbit and anti-mouse (Vector Laboratories, Burlingame, USA) for melanopsin and β-tubulin, respectively (both 1:4000). Blots were washed again 3x 10-min in TBST. The signals were developed using enhanced chemiluminescence (ECL; Millipore, Billerica, USA) and captured on GelDoc (Syngene Chemi Genius 2 Bio imaging system, Cambridge, UK). Densitometric analysis was performed using ImageJ software. The grayscale image was “inverted” and the background was subtracted using the rolling ball radius method in imageJ. The data was expressed as optical density (OD).
Quantitative Real-Time PCR

Adult wild-type (n = 11), rd1 (n = 11) and MNU-injected mice (14 days post-injection; n = 11) were used to study the melanopsin expression levels. Animals were sacrificed by cervical dislocation at ZT 14, eyes removed, hemisected, and retina isolated and processed for RNA isolation. Retinas were collected in 250 μl of Trizol (Sigma-Aldrich) and homogenized using a homogenizer (IKA-Werke). The RNA was then isolated using the phenol-chloroform extraction method. The cDNA was synthesized from 1μg of total RNA, using Transcriptor First Strand cDNA Synthesis Kit (Roche, Cat.No: 04896866001) as per kit protocol. The primers for melanopsin and 18S ribosomal RNA were designed using NCBI and obtained from Sigma-Aldrich. The annealing temperature was standardized using end point Reverse Transcriptase PCR. The primer sequences are as follows:

18S ribosomal RNA:
Forward: GAGGGAGCCTGAGAAACGG
Reverse: GTCGGGAGTGGGTAATTTGC

Melanopsin:
Forward: CCCAGCACCTGCCTTGCCTT
Reverse: TTCTGTGTCTGTCCAGCCCACT

The Real Time PCR reactions were performed in PRISM 7500 Sequence Detection system (Applied Biosystems, CA, USA) using Power SYBR Green PCR Master Mix (Applied Biosystems, CA, USA) as per the protocol. Each reaction was done in duplicate and 18S r-RNA was used as a loading control. 500ng of cDNA was used as the starting material for all the samples. The PCR conditions were: 95°C for 10 min (1 cycle) for activation of Taq polymerase enzyme, followed by 40 cycles of 95°C for 20s (denaturation), 58°C for 30s (annealing temperature) and 72°C for 40s (extension). The data collection was done at extension phase. To verify that the efficiency of primer pairs, without homo- and hetero-dimer formation and producing only a single product, a dissociation protocol was performed. The data was collected using sequence detector software (SDS) and analyzed using ΔΔCT method (relative quantification).

Statistical Analyses

Mixed design analysis of variance (ANOVA) with repeated measures, two-way ANOVA with repeated measures, one-way ANOVA with or without repeated measures were employed to analyze the data in Sigmaplot (Systat Software Inc., CA, USA) or SPSS softwares (IBM, NY, USA). A p-value of less than 0.05 was considered statistically significant. The data on proportion of M1 cells expressing Brn3b were transformed for statistical comparison using arcsine transformation. A two-tailed (paired or unpaired) t-test was used for some comparisons.

Results

PLR was severely attenuated at all irradiances in mice in which classical photoreceptor degeneration was induced during adulthood

In wild-type mouse, the sustained pupil constriction increased with irradiance, first rapidly (~25% constriction at 0.1 μW/cm² and ~45% at ~1 μW/cm²) and then slowly (~55% constriction at 10 μW/cm² and ~70% at ~10⁴ μW/cm²) (Fig 1A and 1B). The rd1 mouse showed negligible pupil constriction at irradiances up to 0.1 μW/cm² and ~5% constriction at 1 μW/cm², but the constriction increased rapidly beyond 1 μW/cm² to approach the levels similar to those in wild-type. These observations in wild-type and rd1 mice are similar to earlier reports [7,21,22,35], and imply that classical photoreceptors are required for PLR at low irradiances.
In mouse in which classical photoreceptor degeneration was induced with MNU during adulthood, the pupil constriction at 7 days post-injection (dpi) was similar to rd1 mouse at lower irradiances (Fig 1A and 1B). However, unlike rd1 mouse, pupil constriction in MNU-injected mouse did not increase rapidly beyond 1 μW/cm². Even at the highest irradiance used here (2.8 x 10^4 μW/cm²), the pupil constriction in MNU-injected mouse was only ~30% (Fig 1B). We measured PLR in these mice for up to 6 months after MNU injection, and found that most of the reduction occurred during the first week (data not illustrated). These results implied that classical photoreceptors have a crucial role in PLR, not just at low, but also at high irradiances.

Attenuation of PLR in MNU-injected mouse was attributable to loss of classical photoreceptors

One concern was that the attenuation in PLR in MNU-injected mouse could come from the drug directly affecting other retinal neurons, particularly the mRGCs. We addressed this with two experiments. First, we injected in some animals NaIO₃, which is also known to induce classical photoreceptor degeneration [26]. The NaIO₃-induced photoreceptor loss in adult mouse also led to severely attenuated PLR at all irradiances (Fig 2A). Second, we injected MNU in adult rd1 mouse. If MNU affected the mRGCs or other retinal neurons, the PLR in the MNU-injected rd1 mouse should also be significantly reduced at high irradiances. However, the PLR in rd1 mouse before and 7 days after MNU injection was similar at all irradiances (Fig 2B). Together, these results implied that the severely attenuated PLR in mice that were injected with MNU (or NaIO₃) after normal retinal development was attributable to loss of classical photoreceptors, and not to any direct effect of the drug on mRGCs or other neurons in retina or other parts of the brain.
Residual PLR at high irradiances in MNU-injected mouse was attributable primarily to remnant classical photoreceptors

If classical photoreceptors play a critical role in PLR at all irradiances, we wondered why there was any residual PLR in MNU-injected mouse at high irradiances (Fig 1B). One possibility was that some photoreceptors escaped MNU-induced degeneration and were responsible for the residual PLR. Alternatively, or additionally, the residual PLR could have originated from activation of melanopsin by high-irradiance stimuli. Using several approaches, we explored these possibilities.

First, we studied the transient pupil response in rd1 and MNU-injected mice. The transient pupil response is considered to originate from classical photoreceptors [7,18]. We found that wild-type and MNU-injected mice, but not rd1 mice, showed transient pupil response at approximately 2 sec after the stimulus onset (Fig 3A; arrow). Furthermore, MNU-injected mice appeared to show more rapid recovery at the stimulus offset than rd1 mice (Fig 3A). These observations suggested that the MNU-injected mouse had more remnant photoreceptors than rd1 mouse. This was also consistent with our finding that injecting MNU in rd1 mouse did not further reduce its PLR (Fig 2B).

We also compared the status of remnant photoreceptors in rd1 and MNU-injected mouse retinas directly by immunolabeling them for s-opsin, m-opsin, or rhodopsin. The number of puncta labeled for s-opsin in MNU-injected mouse (mean ± SE; 2620 ± 1092 per retina; n = 2 retinas) was approximately 5-fold higher than in rd1 mouse (521 ± 189; n = 2 retinas). However, the number of somas and puncta labeled for m-opsin in MNU-injected mouse (5744 ± 2269; n = 4 retinas) was generally similar to that in rd1 mouse (4247 ± 235; n = 2 retinas). In rd1 mouse, we found 2735 ± 790 (n = 2 retinas) rhodopsin-positive cell somas per

Figure 2. Attenuation of PLR in MNU-injected mice was attributable to loss of photoreceptors. A) NaIO3-induced photoreceptor loss led to severely attenuated PLR at all irradiances used here. At 7 days after NaIO3 injection, pupil constriction at the highest irradiance was ~35% (n = 6 mice). PLR data for wild-type and MNU-injected animals are replotted from Fig 1B for comparison. B) MNU did not affect PLR in rd1 mouse. PLR was recorded before and 7 days after MNU injection (two-way ANOVA with repeated measures; F1,8 = 0.02, P = 0.9; n = 5 mice).

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Fig 3. Residual PLR in MNU-injected mice was attributable primarily to remnant photoreceptors. A) Temporal profile of PLR (mean ± SE; n = 5 mice) at a high irradiance (6x10^3 μW/cm^2) for three groups of mice. MNU-injected (7 dpi) mice showed transient response ~2 sec after the stimulus onset (arrow), which was qualitatively similar to that in wild-type mice, whereas rd1 mice did not show any such transient response. B) Representative images of flatmounted rd1 (left column) and MNU-injected (right) mouse.
peripheral retinas, showing immunolabeling for s-opsin (top row; inferior retina), m-opsin (middle; superior retina) and rhodopsin (bottom; superior retina). S-opsin labeling was present as bright puncta (arrows), possibly representing remnant s-cone outer segments. M-opsin and rhodopsin labeling was present as puncta (arrows), as well as in cell somas (arrowheads). Scale bar: 20 μm. C) Schematics or montages of wholemount retinas labeled for s-opsin (top row), m-opsin (middle) and rhodopsin (bottom) showing the extent and pattern of photoreceptor loss in rd1 (left) and MNU-injected (right) mice. Each black dot in the schematics (top 2 rows) represents a remnant cone. To ensure that cones could be seen as distinctly as possible, we used smaller dot sizes where their density was too high. The number of rods in the MNU-injected sample was so high, especially in peripheral retina that we could not generate a schematic, and therefore show here the images of the montage from both rd1 and MNU-injected mice. Scale bar: 1 mm. D) Administration of NaIO3 in MNU-injected mice resulted in further reduction in PLR at high irradiances at 7 dpi. A two-way ANOVA with repeated measures on irradiance showed a significant difference between the two groups (F1,45 = 8.7; P = 0.032; n = 6 mice). A pairwise comparison using post hoc Holm-Sidak test revealed significant differences at irradiances ≥ 10 μW/cm². *P < 0.05.

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retina. These somas were present mostly in the peripheral retina. However, the number of rhodopsin-positive entities in the MNU-injected sample was so high, especially in peripheral retina, that we could not count them reliably. Therefore, we measured the integrated density of rhodopsin labeling in retinal wholemounts, and found that it was approximately 20-fold higher in MNU-injected mouse (2.9 x 10⁶; n = 1 retina) than in rd1 mouse (1.6 x 10⁵; n = 1 retina) (Fig 3B and 3C).

Next, to understand whether the higher numbers of remnant classical photoreceptors in the MNU-injected mouse could account for the residual PLR, we administered NaIO3 (60 mg/kg; i/v) in some MNU-injected mice, expecting that this will lead to additional loss of classical photoreceptors and will further reduce the residual PLR. We found that NaIO3 indeed resulted in further reduction in PLR at high irradiances (Fig 3D). In some animals, we also injected multiple doses of MNU and found further reduction in PLR (data not shown). Together, these results suggested that there were more remnant classical photoreceptors in the MNU-injected mouse than in the rd1 mouse, and that they contributed to the residual PLR in the MNU-injected mouse.

Melanopsin activation made a minor contribution to PLR at high irradiances

Our results from inducible models of classical photoreceptor degeneration suggested that melanopsin contribution to PLR was relatively minor (<30% at the highest irradiance used here; Figs 1B, 2A and 3D). To further investigate whether, and to what extent, melanopsin activation contributed to the high-irradiance PLR, we blocked melanopsin activity in the animals previously injected with MNU and NaIO3, using a recently-discovered melanopsin antagonist (AA41612, 10 mg/kg, i/p) [24]. This resulted in further reduction in PLR at high irradiances (>10³ μW/cm²) (Fig 4). Interestingly, even at the highest irradiance used here, the melanopsin antagonist reduced the PLR by <15% (Fig 4), a value that matches the magnitude of reduction in high-irradiance PLR in melanopsin knockout mouse [22].

A higher proportion of M1 cells expressed Brn3b in rd1 mouse, but not in MNU-injected mouse

If classical photoreceptors are responsible for PLR even at high irradiances, and considering that the extent of photoreceptor loss in rd1 mouse is similar (m-cones) or more (rods, s-cones) than in MNU-injected mouse, it was unclear why the high-irradiance PLR is intact in rd1 mouse. One possibility was that loss of classical photoreceptors during development in rd1 mouse results in a compensatory response in mRGCs which may then take over the PLR.
function at high irradiances. To explore this, we measured melanopsin expression levels (Fig 5) and counted the numbers of mRGCs in wild-type, rd1 and MNU-injected mice (Fig 6).

Using quantitative immunoblotting and real-time PCR, we measured melanopsin protein and mRNA levels respectively in wild-type, rd1 and MNU-injected (14 dpi) mice. We found that melanopsin expression levels (protein and mRNA) in both rd1 and MNU-injected mice were statistically similar to those in wild-type (Fig 5). However, the numbers of mRGCs in both rd1 and MNU-injected mice were significantly higher than in wild-type (Fig 6C). To investigate whether this effect was specific to mRGCs, we also counted the non-mRGCs using Brn3a immunohistochemistry. The transcription factor Brn3a has been shown to be absent in mRGCs [9,36]. We found that the numbers of Brn3a-positive RGCs in rd1 and MNU-injected mice were not statistically different from wild-type (Fig 6D). These results suggested that loss of photoreceptors, irrespective of the etiology and the age of onset, leads to increase in the numbers of mRGCs.

The mRGCs have been classified based on their dendritic stratification into M1 and non-M1 types (Fig 6A and 6B) [9,19,31,37]. Nearly all non-M1 and a small subset (10%–15%) of M1 cells express the transcription factor Brn3b, while the majority of M1 cells do not [9,38]. The Brn3b-expressing mRGCs have been implicated in PLR [25], although it is not clear whether Brn3b-expressing M1 cells, non-M1 cells, or both are involved. To investigate whether the increase in the numbers of mRGCs was specific to a subtype, we counted M1 and non-M1 cells identified based on their dendritic stratification in the IPL [9].

We found that the densities of M1 cells in rd1 (mean ± SD; 45 ± 12 cells/mm²) and MNU-injected mice (43 ± 10) were similar to that in wild-type (41 ± 12) (Fig 6E). However, the density of non-M1 cells was significantly greater in both rd1 (132 ± 28) and MNU-injected mice (150 ± 32) compared with wild-type (93 ± 31) (Fig 6E). Since the number of non-M1 cells, which express Brn3b, increased after loss of classical photoreceptor, we asked if the Brn3b-
expressing M1 cells were also affected in the two models. Interestingly, we found that 42% of the M1 cells (19 ± 7 out of 45 ± 12 cells/mm²; mean ± SD) expressed Brn3b in rd1 mouse, a proportion that was approximately four-fold higher than in wild-type (11%, or 4.6 ± 3 out of 41 ± 12) (Fig 6F). However, the proportion of these cells in MNU-injected mouse (19%, or 8 ± 5 out of 43 ± 10) was statistically similar to that in wild-type (Fig 6F). Even in the mouse...
Fig 6. Both rd1 and MNU-injected mice had higher numbers of mRGCs, but only rd1 mice had a higher proportion of Brn3b-expressing M1 cells than in wild-type. A) Representative images from flatmounted retinas double-labeled for melanopsin (green) and Brn3b (red), showing non-M1 cells (arrows) in wild-type (WT, left), rd1 (middle) and MNU (7 days post injection; right). Top row shows original images, middle row shows dendritic stratification of the marked non-M1 (arrow) cells, and bottom row shows the same images with melanopsin signal digitally removed to more clearly show the Brn3b labeling in the non-M1 cells. The middle images show...
2 thin solid lines marking the boundaries of the IPL and a thin dashed line separating the Off (top) and On (bottom) sublaminas in the IPL. The faintly-labeled non-M1 cells (arrows) stratified in ON sublamina and expressed Brn3b in wild-type, rd1 and MNU-injected mice. Scale bar: 50 μm. B) Representative images from flatmounted retinas double-labeled for melanopsin (green) and Brn3b (red), showing M1 cells (arrows) in wild-type (left), rd1 (middle) and MNU (7 days post injection; right). Top row shows original images, middle row shows dendritic stratification of the marked M1 (arrow) cells, and bottom row shows the same images with melanopsin signal digitally removed to more clearly show the Brn3b labeling in the M1 cells. The middle images show 2 thin solid lines marking the boundaries of the IPL and a thin dashed line separating the Off (top) and On (bottom) sublaminas in the IPL. The brightly-labeled M1 cells (arrows) stratified in OFF sublamina and expressed Brn3b in rd1, but not in wild-type or MNU-injected mice. Scale bar: 50 μm. C) The density of mRGCs (mean ± SE) in wild-type, rd1 and MNU-injected mice. A one-way ANOVA showed significant differences among the 3 groups (F2,33 = 8.18; P = 0.001; n = 8 mice for wild-type and rd1, n = 20 for MNU-injected). A post hoc Holm-Sidak test revealed that both rd1 and MNU-injected mice had significantly higher numbers of mRGCs than wild-type. *P < 0.05; †P < 0.001. D) The numbers of non-mRGCs, identified by presence of Brn3a immunoreactivity, in wild-type, rd1, and MNU-injected mice were statistically similar (F2,5 = 1.16; P = 0.39; one-way ANOVA; n = 3 mice for wild-type and rd1, n = 2 for MNU-injected). E) Densities of M1 type of mRGCs in wild-type, rd1 and MNU-injected mice were similar (F2,33 = 0.32; P = 0.73; one-way ANOVA), whereas the densities of non-M1 cells were different (F2,33 = 9.74, P < 0.001). A post hoc Holm-Sidak test showed that both rd1 and MNU-injected mice had significantly higher numbers of non-M1 cells than in wild-type (n = 8 mice for wild-type and rd1, n = 20 for MNU-injected). The MNU-injected group included animals in which mRGCs were counted at 3, 7, 14 or 28 days after the MNU injection (n = 5 mice each); data from all samples were pooled, because there was no statistical difference among the subgroups (data not shown). *P < 0.05, †P < 0.001. F) There was a significant difference in the proportion of M1 cells that expressed Brn3b among wild-type, rd1 and MNU-injected mice (F2,32 = 13.39, P < 0.001, one-way ANOVA; n = 8 mice for wild-type, 7 for rd1, 20 for MNU). A post hoc Holm-Sidak test showed that the numbers in rd1 mice, but not in MNU-injected mice, were significantly higher than in wild-type. †P < 0.001.

Classical photoreceptor degeneration induced during development resulted in partial rescue of PLR at high irradiances

If the developmental loss of classical photoreceptors was responsible for the intact high-irradiance PLR in rd1 mouse, we expected that photoreceptor degeneration induced with MNU or NaIO3 during development would also result in more M1 cells expressing Brn3b, we injected MNU (60 mg/kg) in wild-type mice at P-8 to P-12, the time when photoreceptors start to degenerate in rd1 mouse. At 2–3 weeks after the injection, only 1–3 cell layers remained in the ONL, similar to age-matched rd1 (P-21)
Fig 7. Photoreceptor loss induced during development resulted in partial rescue of PLR at higher irradiances. A) PLR in adult (P-72) mice that were injected with MNU prenatally (5 mg/kg i/p at embryonic day 16; n = 5 mice). The 2 parentheses in the label show animal ages when the drug was injected and PLR was recorded, respectively. PLRs of mice injected with MNU during adulthood and of adult rd1 mice are replotted from Fig 1B for comparison. A mixed ANOVA with repeated measures on irradiance showed significant differences among the 3 groups ($F_{2,18} = 60.7, P < 0.0001$). A post hoc test using Bonferroni correction showed that the PLR in MNU (prenatal)(P-72) animals was significantly higher than in MNU (adult)(adult) animals ($P = 0.0002$), but significantly lower than the adult rd1 mice ($P = 0.001$). A pairwise comparison at each irradiance revealed that the PLR in MNU (prenatal)(P-72) mice was similar to that in MNU (adult)(adult) at $<1 \mu W/cm^2$, but significantly higher at irradiances $>1 \mu W/cm^2$. The PLR in MNU (prenatal)(P-72) mice was similar to that in adult rd1 mice at $<10^3 \mu W/cm^2$, but significantly lower at irradiances $>10^3 \mu W/cm^2$. * $P < 0.0005$ for MNU (prenatal) versus MNU (adult); † $P < 0.0001$ for MNU (prenatal) versus adult rd1.

B) PLR in adult (P-70) mice that were injected with NaIO$_3$ at P-8 (60 mg/kg; i/p; n = 6 mice). PLRs of mice injected with NaIO$_3$ during adulthood and of adult rd1 mice are replotted from Figs 2A and 1B, respectively, for comparison. A mixed ANOVA with repeated measures on irradiance showed significant differences among the 3 groups ($F_{2,17} = 14.3; P = 0.0003$). A post hoc test using Bonferroni correction showed that the PLR in both rd1 and NaIO$_3$ (P-8)(P-70) mice was significantly higher than that in NaIO$_3$ (adult)(adult) mice ($P = 0.0002$ and 0.038, respectively), while the PLR in rd1 mice was similar to that in NaIO$_3$ (P-8).
mouse (Fig 8A). The proportion of M1 cells that expressed Brn3b in these animals (24% or 13 ± 4 out of 53 ± 18 cells/mm²; mean ± SD) as well as in age-matched rd1 mice (41% or 18 ± 10 cells/mm² out of 44 ± 17) was significantly higher than in age-matched wild-type mouse (14% or 6 ± 2 cells/mm² out of 43 ± 11), while there was no statistical difference between the MNU-injected and rd1 mice (Fig 8B).

Discussion

Primary role for classical photoreceptors in generating PLR

PLR is thought to originate from classical photoreceptors at low irradiances and from melanopsin activation at high irradiances [1,7,21–23]. However, since classical photoreceptors can encode all physiologically-relevant light levels [39], it seemed puzzling that melanopsin activation should contribute to normal PLR. By inducing photoreceptor loss after normal retinal development in mice, we show here that classical photoreceptors are required for PLR at all irradiances. These mice showed no pupil constriction at low irradiances, and a severely attenuated response at high irradiances. Using two experimental approaches we demonstrate that the loss of PLR in these mice was attributable to loss of photoreceptors (Fig 2).

The inducible mouse models for photoreceptor loss used in this study have been well characterized. MNU is one of the most commonly used chemical agents to induce photoreceptor degeneration [27,40,41]. Multiple approaches, such as biochemical assays, electroretinogram measurements, and behavioral tests have demonstrated that both rods and cones are severely impaired and gross vision is lost in both MNU- and NaIO₃-treated animals [27,42–44]. However, the ganglion cell layer is relatively preserved in the MNU mouse model [27] and has not
been investigated in the NaIO₃ model. Our finding that MNU did not alter the PLR in rd1 mouse, further confirms that MNU does not adversely affect the retinal ganglion cells, especially the mRGCs (Fig 2A).

Several observations suggested that the residual PLR at high irradiances in the MNU-injected mouse originated from the remnant classical photoreceptors. First, MNU-injected mouse exhibited a transient PLR response which was absent in rd1 mouse. Second, injecting NaIO₃ in the MNU-injected mouse resulted in additional reduction in PLR. Finally, the numbers of remnant rods and s-cones in MNU-injected mouse were considerably higher than in rd1 mouse. Rods, although they operate at low light levels, have been recently shown to also respond to very bright light [45,46]. Interestingly, s-cones have been shown in mouse to produce a sustained response to bright light in olivary pretectal nucleus (OPN), the brain area responsible for PLR [6]. A previous report, showing that MNU-injected mouse retina retains some light response, provides evidence that remnant photoreceptors in this model can not only respond to light, but also send signals through the retina [47].

A relatively small component (up to ~15%) of the high-irradiance PLR also originated from activation of melanopsin. We concluded this from the extent of PLR reduction observed in the mice with both MNU and NaIO₃ injected and melanopsin pharmacologically blocked (Fig 4). This magnitude of PLR reduction was consistent with previous observations in Opn4⁻/⁻ mouse [21,22]. Even if the antagonist did not block melanopsin activation completely, the melanopsin contribution to total PLR could not have exceeded ~25% at the highest irradiance used here [24]. Furthermore, the high irradiances (≥10⁶ μW/cm²) where melanopsin did contribute to PLR are likely non-physiological for a mouse, a nocturnal animal. Using a genetic mouse model where m-cones respond to a higher wavelength (650 nm), Lall et al. reported that melanopsin contribution to high-irradiance PLR is greater than that of m-cones, but they also showed that 40% of pupil constriction could be achieved at high irradiances in cone-only (Gnat⁻/⁻,Opn4⁻/⁻) mouse [5], suggesting a significant role for cones at high irradiances. The melanopsin contribution to PLR may vary depending on stimulation conditions, such as stimulus wavelength, stimulus duration, and light adaption [5,48,49].

Photoreceptor loss during development results in compensatory rescue of high-irradiance PLR

If classical photoreceptors are responsible for PLR at all irradiances, it was unclear how the PLR is preserved at high irradiances in rd1 mouse. The remnant photoreceptors cannot explain this, considering that rd1 mouse has fewer photoreceptors than MNU-injected mouse. We explored the possibility that the loss of photoreceptors during rd1 mouse development results in some compensatory response in mRGCs.

Ablating the mRGCs abolishes PLR and circadian photoentrainment [50,51]. Ablating the Brn3b-expressing mRGCs results in severely attenuated PLR, implying that PLR is mediated specifically by these cells [25]. However, since Brn3b is expressed by both M1 and non-M1 type of mRGCs, it was not clear which of these cell types mediate PLR. We found that in rd1 mouse, in which the photoreceptors start to degenerate during development, the numbers of the Brn3b-expressing M1 cells were higher than in mice in which photoreceptor loss was induced during adulthood. Since the high-irradiance PLR is preserved in rd1 mouse but not in MNU-injected mouse, this suggested that the increased number of Brn3b-expressing M1 cells is responsible for the preserved PLR. Consistent with this, we found that inducing photoreceptor degeneration with MNU or NaIO₃ during development also resulted in significantly higher proportions of Brn3b-expressing M1 cells and partially rescued PLR. Interestingly, onset of PLR during development in wild-type mouse is marked by the innervation of OPN shell by M1
cells [52]. The higher numbers of Brn3b-expressing M1 cells following developmental loss of photoreceptors (either in rd1 mouse or in inducible models) could potentially cause enhanced innervation of OPN by these cells, leading in turn to the rescued high-irradiance PLR in these mice.

The photoreceptor loss induced during development, however, led to only a partially-rescued high-irradiance PLR. One possibility was that the higher PLR in these mice at high irradiances originated from the remnant photoreceptors (Fig 3) and not from any compensatory response in mRGCs. However, our finding that the PLR in these mice increased over time during development (Fig 7C and 7D) strongly suggested that the rescued high-irradiance PLR was an emergent property that cannot be explained by remnant photoreceptors.

Regulation of mRGCs by photoreceptors

A previous report showed that rd1 mouse has more mRGCs than the congenic wildtype; the reason they suggested was that the mRGCs in rd1 mouse do not undergo apoptosis during development [23]. However, our finding that the numbers of mRGCs was higher even when photoreceptor degeneration was induced with MNU during adulthood suggests alternative/additional mechanisms. Although both M1 and non-M1 type of mRGCs receive signals from the photoreceptors, the latter receive a stronger input [53,54]. We found that loss of photoreceptors resulted in increased numbers of non-M1 cells in both rd1 and MNU-injected mice (Fig 6E), indicating that photoreceptors regulate the expression of melanopsin in RGCs, dynamically and negatively. It should also be noted that the numbers of mRGCs in rd/rd cl mouse have been reported to be similar to its congenic control [55–57], possibly because the loss of photoreceptors occurs at a much faster rate in this model.

Despite the higher number of mRGCs in both rd1 and MNU-injected mice, we did not find any change in melanopsin expression (protein or m-RNA) levels (Fig 5). This is consistent with previous reports showing no change in melanopsin expression in rd/rd cl or rd1 mouse models [23,55]. However, melanopsin levels have been reported to be reduced in rat models of retinal degeneration [32,58,59], raising the possibility that different mechanisms may be involved in different species.

The number of Brn3b-expressing RGCs declines during early development (from E16 to P-5) in normal mouse [60]. It is possible that Brn3b expression by RGCs is also negatively regulated by photoreceptors during development. This was consistent with our observation that the numbers of Brn3b-expressing M1 cells declined from P-14 to adulthood in wild-type, but not in rd1 mouse (data not shown). This could explain the higher numbers of Brn3b-expressing M1 cells in rd1 mice or in mice in which the photoreceptor degeneration is induced during development than in wild-type mice or in mice in which the photoreceptor degeneration is induced during adulthood.

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Author Contributions

Conceived and designed the experiments: VJ NKD. Performed the experiments: VJ IS SP MG SKS. Analyzed the data: VJ IS SP MG NKD. Wrote the paper: VJ MG NKD.
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