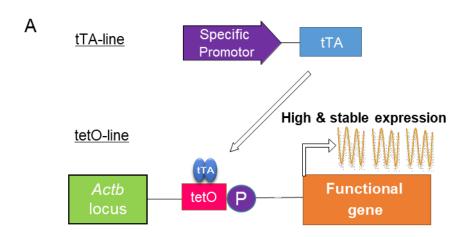
## **Supplemental Information for**

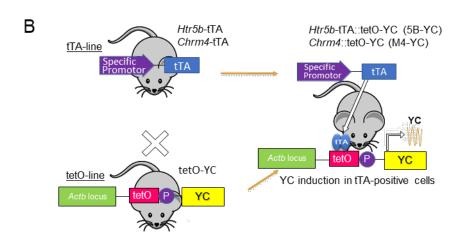
- 2 Starburst amacrine cells amplify optogenetic visual restoration through
- 3 gap junctions in the murine retina
- 4 Katada Y et al.

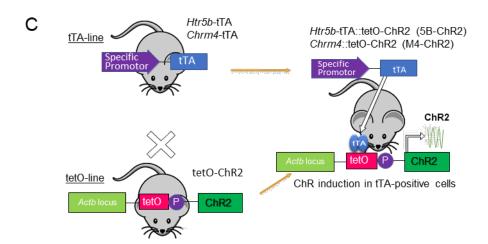
5

## **7 Supplemental Figures**

## 8 Figure S1. KENGE-tet system and tetO-Yellow cameleon

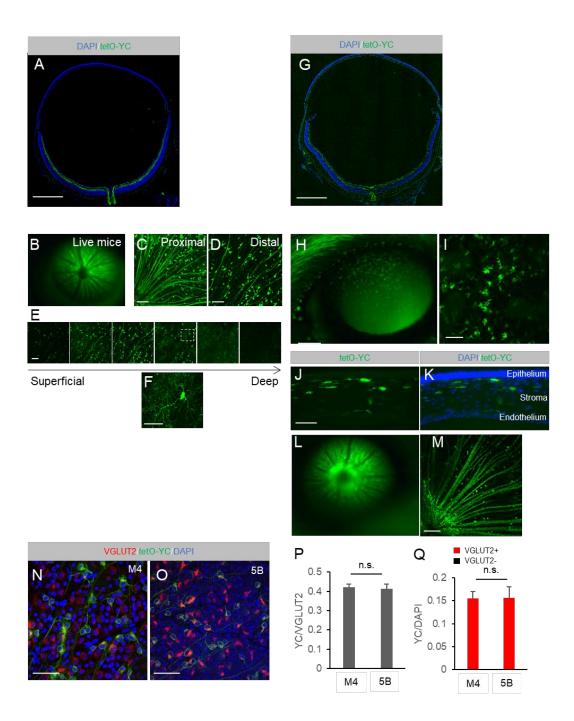






10 (A) KENGE-tet system. tTA is expressed under the control of cell type-specific promoters. tTA transactivates the tetO promoter, and the functional protein is induced in a cell-type-11 specific manner. (B, C) Two different mouse lines are employed that express the gene 12encoding tTA protein under the control of a cell-type-specific promoter, muscarinic 13 acetylcholine receptor M4 or serotonin receptor 5B control region. These mice were further 14 crossed with another transgenic mouse line containing a YC fluorescent gene connected into 15 the downstream of the tetO promoter. The YC gene expression was induced only by the 16 presence of tTA protein in the double transgenic mice (M4-YC or 5B-YC)(B). Next, as a 17 visual restoration model, the tTA line was crossed with tetO-ChR2. (C) tTA drove ChR2 18 expression in RGC. 19

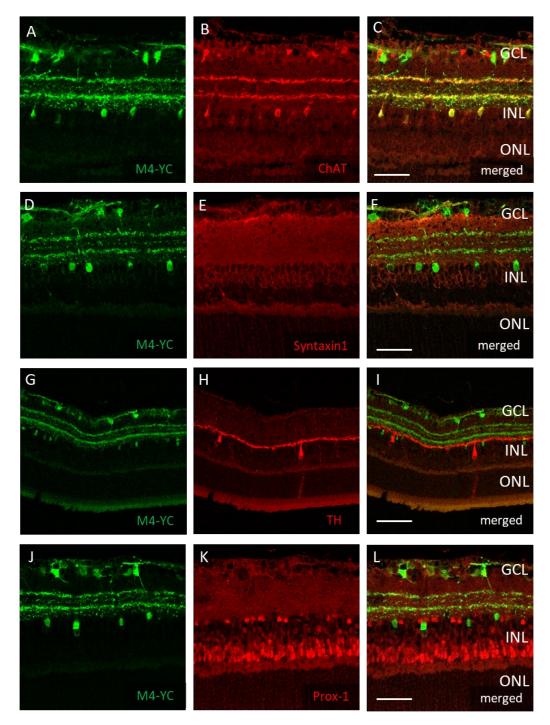
## Figure S2. Expression of Yellow cameleon in M4-YC and 5B-YC



In the M4-YC mouse retina, we identified the expression of YC (green) in RGC and 24amacrine cells within sections (A), with in vivo fluorescence microscopy (B), and with flat 25mounted retina (C-F). In the 5B-YC mouse retina, we identified the expression of YC in 2627 RGC and the corneal stromal layer with in sections (G, J, K), in vivo fluorescence microscopy (H, I, L), and the flat mounted retina (M). Coexpression of the RGC marker 28 VGLUT2 in flat mounted retina of M4-YC (N) and 5B-YC (O). Percentage of YC-positive 29 cells in VGLUT2-positive (P) or DAPI-positive cells (Q) and VGLUT2-positive cells in 30 YC-positive cells (Q) in both lines from confocal flat mounted GCL (n = 3 retinas each). 31 Regions were chosen in each quadrant, and we obtained VGLUT2, DAPI-positive, YC-32positive, and co-labeled cells. Error bars represent the SEM. Scale bar: 50 µm in (F), (N) 33 and (O). 100 µm in (C-E), (I), (J) and (M). 1,000 µm in (A) and (G). 34

## Figure S3. Immunostaining of amacrine cells in M4-YC mice

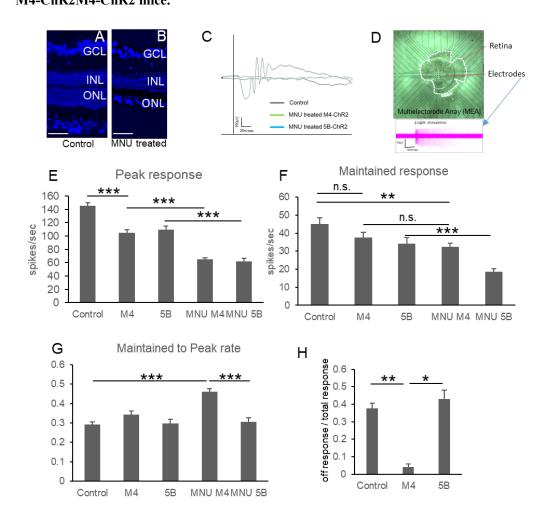
36



Immunohistochemistry on the transverse retinal cryosection of M4-YC mouse labeled with choline acetyltransferase (ChAT), a starburst amacrine marker (A-C), syntaxin1, a pan-amacrine cell marker (D-F), tyrosine hydroxylase (TH), a dopaminergic amacrine

- 41 marker (G-H) and Prox-1, an All amacrine marker (J-L). GCL, ganglion cell layer; INL,
- inner nuclear layer; ONL, outer nuclear layer. Scale bars, 50 μm in A-L.

# Figure S4. The maintained response was retained regardless of photoreceptor degeneration in M4-ChR2M4-ChR2 mice.

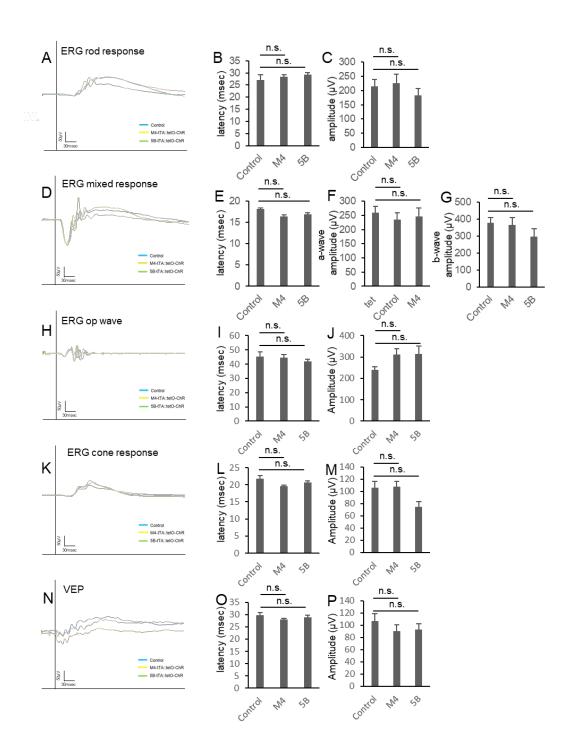


(A, B) Comparison of retinal sections after intraperitoneal administration of MNU. MNU was used to produce a photoreceptor degeneration model. Two weeks after the MNU injection, DAPI nuclear counter-stain is shown in blue. Scale bar: 50 μm. (C) Representative ERG traces from MNU-injected and control mice. White LED light stimulation of 10.0 log cd-s/m² was delivered. (D) Image of MEA. It can measure the extracellular potential of RCGs in contact with the electrode ex vivo. (E-F) Comparison of peak response (E),

maintained response (F) and maintained to peak rate (G) from MEA recordings among control (tetO-ChR2; n = 3 retinas, 112 cells), M4-ChR2M4-ChR2 (n = 3 retinas, 62 cells), 5B-ChR2 (n = 3 retinas, 48 cells), MNU-treated M4-ChR2M4-ChR2 (n = 3 retinas, 164 cells) and MNU-treated 5B-ChR2 (n = 3 retinas, 117 cells) mice. (H) Comparison of ratio of off response to total light response from MEA recording. Error bars represent SEMs. \*p < 0.05, \*\*p < 0.01, \*\*\*p < 0.001. Games-Howell test.

#### 61 Figure S5. Ectopic channelrhodopsin induction had no significant effect on ERG or

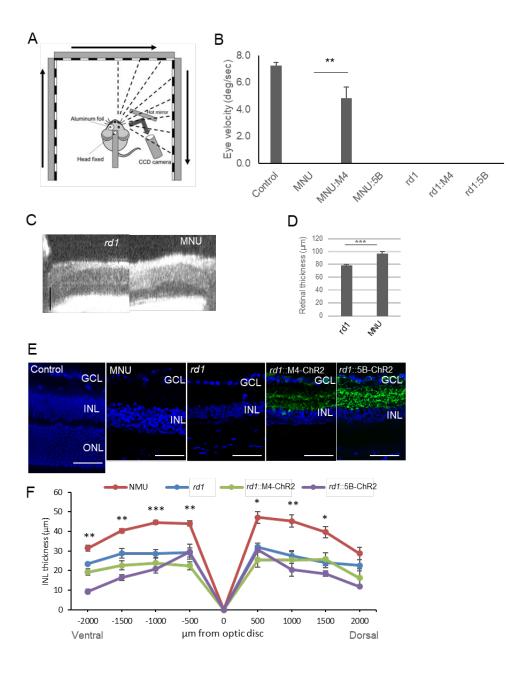
#### **VEP.**



- Representative ERG waveforms from ERG rod response (A), mixed response (D),
- oscillatory potentials (H), cone response (K) and VEPs (N). Quantification of latency
- 66 (B, E, I, L, O) and amplitude (C, F, G, J, M, P) in each protocol in control (tetO-ChR2;
- n = 6), M4-ChR2M4-ChR2 (n = 8), and 5B-ChR2 (n = 8) mice. Error bars represent
- 68 SEMs. n.s., not significant. Games-Howell test.

## Figure S6. Thinning of INL in the genetic model of retinal degeneration attenuating

## the restoration effect.

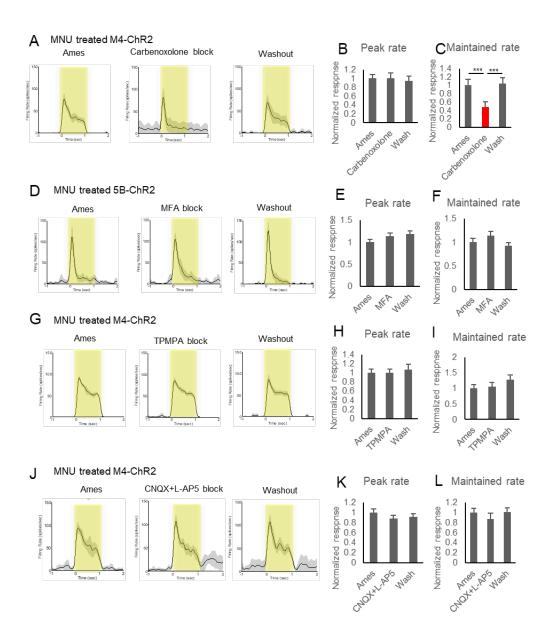


(A) Schematic view of the OKR system. The images of the right or left eyes are captured by a CCD camera placed on the same side. During measurement, the contralateral eyes are

- covered with aluminum foil. Visual stimulation is presented on three LCD monitors around 74the mouse, the head of which is fixed in the middle. (B) The average eye velocities of the 75control tetO-ChR2 mice (n = 3), MNU-injected tetO-ChR2 mice (n = 3), M4-ChR2M4-76 ChR2 mice (n = 9), 5B-ChR2 mice (n = 5), rd1; tetO-ChR2 mice (n = 8), rd1; M4-ChR2M4-77ChR2 mice (n = 10) and rd1;5B-ChR2 mice (n = 5) measured from the OKR system at 10 78weeks of age. (C, D) Retinal structure images of rd1 (n = 9) and MNU (n = 10)-treated mice 79 at 8 weeks of age by optical coherence tomography (OCT). (E) Retinal sections of control 80 (tetO-ChR2), MNU-treated tetO-ChR2, rd1::tetO-ChR2, rd1;M4-ChR2M4-ChR2 and 81 rd1;5B-ChR2 mice at 8 weeks of age. Nuclear counter-staining with DAPI (blue) and 82 ectopic gene induction with YC (green) are shown. (F) The INL thickness quantification 83 from the sagittal sections. Cryosections from MNU-treated tetO-ChR2 (n = 6), and 84 rd1::tetO-ChR2 mice (n = 6) at 8 weeks of age and rd1;5B-ChR2 (n = 3) and rd1;M4-85 ChR2M4-ChR2 (n = 3) mice at 10 weeks of age were used for quantification. 86
- 87 All scale bars: 50  $\mu m.$  All error bars represent the SEMs. \*p < 0.05, \*\*p < 0.01, \*\*\*p <
- 88 0.001. Student's 2-tailed t-test or Tukey's test.

#### Figure S7. Gap junction was involved in the maintained response.

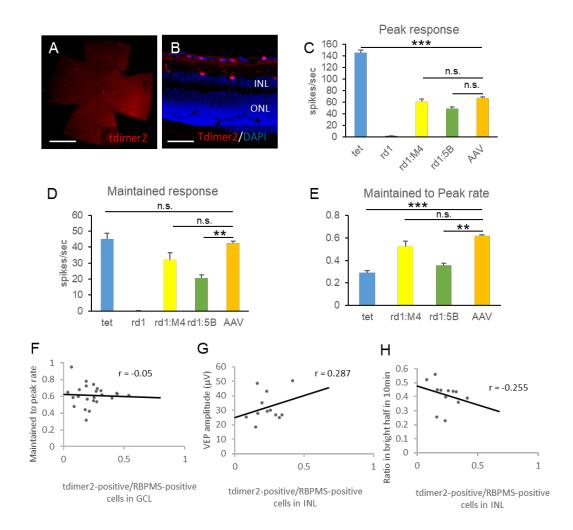
90



92 (A, D, G, J) Mean ± SEM of exemplar cell response firing rate recorded during normal
93 Ames' medium superfusion (left), in synaptic block (middle), and after washout (right).
94 MNU-treated M4-ChR2M4-ChR2 mice with carbenoxolone block (n=3 retinas, 25 cells)
95 (A), MNU-treated 5B-ChR2 mice with MFA block (n=3 retinas, 117 cells) (D), MNU

- 96 treated M4-ChR2M4-ChR2 mice with TPMPA block (n=3 retinas, 51 cells) (G), and MNU
- 97 treated M4-ChR2M4-ChR2 mice with CNQX and L-AP5 block (n=3 retinas, 52 cells) (J).
- 98 The gray areas around the averaged traces represent the SEM. (B, C, E, F, H, I, K, L)
- Averaged normalized peak firing rate and maintained rate. Maintained time frame is 0.4 to
- 1.0 seconds from light stimulation. Light intensity was 13.6 log photons/cm<sup>2</sup>/s.
- All error bars represent the SEMs. \*\*\*p < 0.001. One-way ANOVA and Tukey's test.

#### Figure S8. Visual restoration profile of rAAV model



(A) AAV2–CAG–tdimer2–WPRE intravitreally injected mouse flat mounted retina and its section. (C-E) Comparison of peak response (C), maintained response (D) and maintained to peak rate (E) from MEA recordings among control (tetO-ChR2; n = 3 retinas, 112 cells), rd1::tetO-ChR2 (n = 3 retinas, 86 cells), rd1;M4-ChR2M4-ChR2 (n = 3 retinas, 18 cells), rd1;5B-ChR2 (n = 3 retinas, 17 cells) and rAAV treated rd1::tetO-

110 ChR2 mice (n = 24 retinas, 1,151 cells) at 10 weeks of age. (F-H) Correlation between

111 transfection efficiency into RGCs (tdimer2-positive cells/RBPMS-positive cells in INL)

112 and maintained to peak rate (F) (n = 24), VEP amplitude (G) (n = 24) and % time in

113 bright half in LDT (H) (n = 12). All error bars represent the SEMs. INL, inner nuclear

114 layer; GCL, ganglion cell layer, RGC, retinal ganglion cell, ONL, outer nuclear layer.

115 Scale bars, 1,000 μm in (A), 50 μm in (B), n.s.: not significant, \*p < 0.05, \*\*p < 0.01,

116 \*\*\*p < 0.001. Games-Howell test (C, D, E), Pearson's correlation coefficient (F-H).