

SUPPLEMENTAL INFORMATION FOR

An inducible Cre mouse for studying roles of the RPE in retinal physiology and disease

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SUPPLEMENTAL FIGURES, FIGURE LEGENDS, AND TABLES

Figure S1

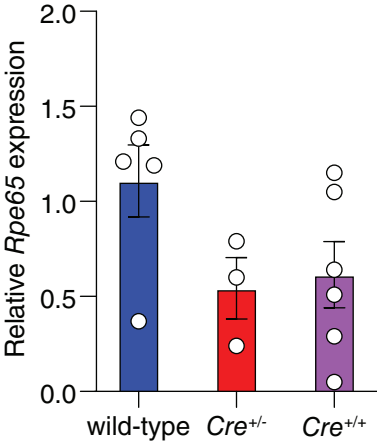


Figure S1. *Quantitative PCR analysis of Rpe65 mRNA expression.* *Rpe65* transcript levels were measured by RT-PCR and normalized to the *Gapdh* transcript levels. *Rpe65* was reduced by ~50% in *Cre*^{+/-} (n = 3) and *Cre*^{+/+} (n = 6) mice as compared to wild-type controls (n = 5), although these differences were not statistically significant as assessed by one-way ANOVA [$F_{(2,11)} = 2.671$, $P = 0.11$]. Each point represents data from a single mouse. Bars indicate mean \pm SEM.

Figure S2

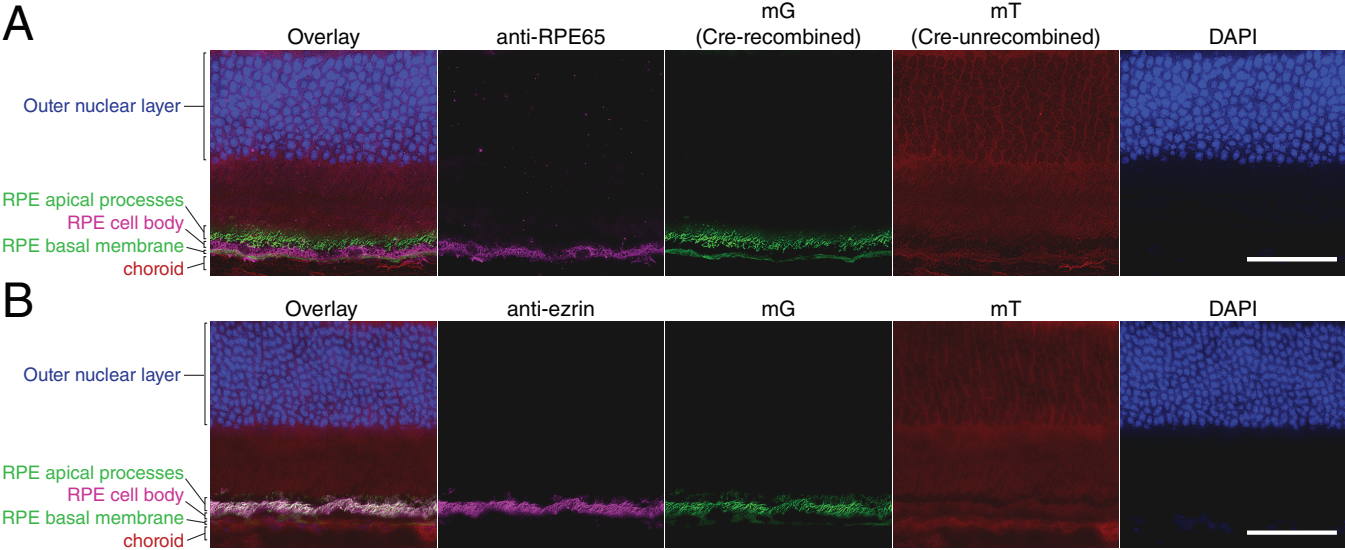


Figure S2. *Relationship of retinal mT and mG fluorescence to RPE cell markers.* **A)** A retina cryosection from a PD21 IP tamoxifen-treated *Cre^{+/-} mT/mG^{+/-}* mouse was co-stained with an anti-RPE65 Ab and DAPI. The leftmost image shows the overlaid fluorescence signals while the images to the right show the individual channels. The RPE65-associated fluorescence is localized to the cell body (1) in-between the mG signals associated with the basal and apical plasma membranes of the RPE. Note that the scattered specs in this image were also present in the secondary antibody-only control. The scale bar on the rightmost panel represents 50 μ m. **B)** A retina cryosection from an PD21 IP tamoxifen-treated *Cre^{+/-} mT/mG^{+/-}* mouse was co-stained with an anti-ezrin antibody and DAPI. The leftmost image shows the overlaid fluorescence signals while the images to the right show the individual channels. The overlap between the mG and ezrin-associated fluorescence signals confirms the localization of mG to the RPE apical membrane (1). The scale bar on the rightmost panel represents 50 μ m.

Figure S3

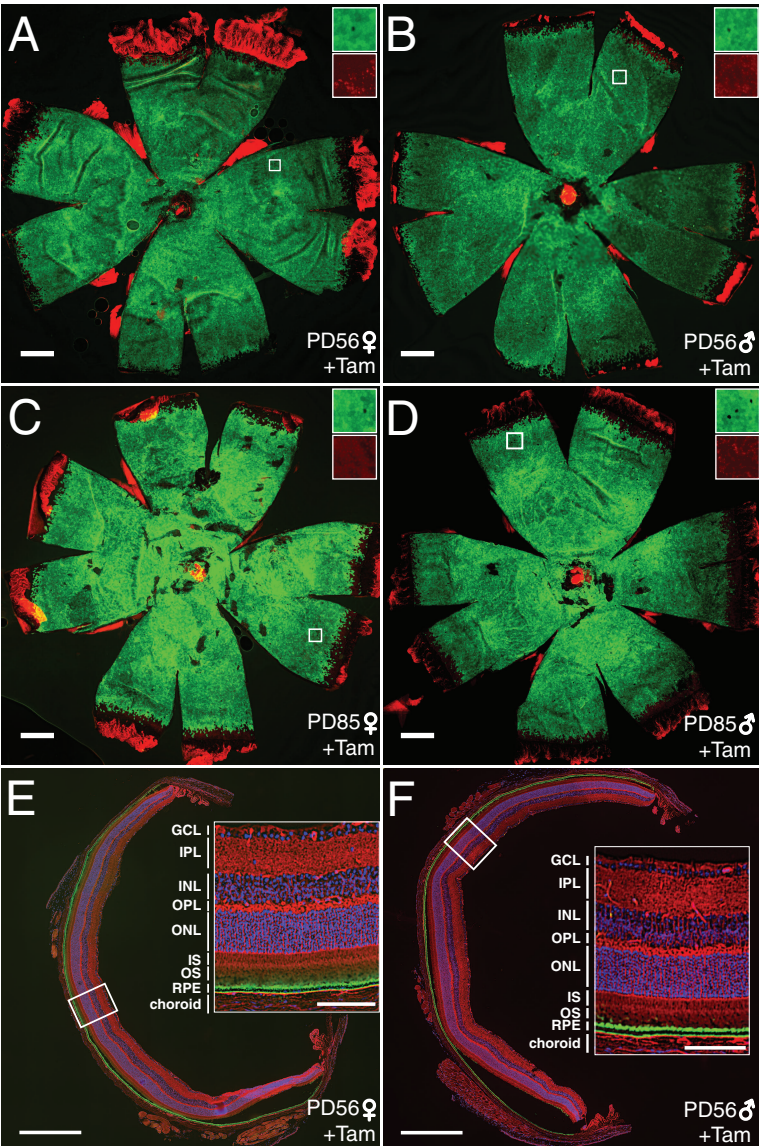


Figure S3. *Cre* recombinase activity in *Rpe65^{CreERT2}* mouse retina assessed with the *mT/mG* reporter. Panels **A** and **B** show RPE flatmounts from *Cre^{+/-}* female and male mice, respectively, that were fed tamoxifen chow diet for three weeks starting on PD21 (n = 2). The flatmounts were obtained two weeks after stopping the tamoxifen chow diet and imaged with a fluorescence microscope. Green fluorescence indicates cells where *Cre*-mediated recombination has occurred, whereas red fluorescence indicates un-recombined cells. Panels **C** and **D** show RPE flatmounts from representative *Cre^{+/-}* female and male mice, respectively, that were fed tamoxifen chow diet for three weeks starting on PD50 (n = 4). The flatmounts were obtained two weeks after stopping tamoxifen chow diet. The insets in panels **A-D** show zoomed areas (marked by white squares) with green and red channels separately displayed to demonstrate the single cell resolution of the imaging method. Panels **E** and **F** show retina cryosections from *Cre^{+/-}* female and male mice, respectively, that were fed tamoxifen chow for three weeks starting on PD21 (n = 2). The cryosections were obtained two weeks after stopping tamoxifen chow diet. The green label represents *Cre*-mediated recombination and is restricted to the RPE. Insets show zoomed views of the areas marked with rectangles with the retinal cell layers labeled. Abbreviations: GCL, ganglion cell layer; INL, inner nuclear layer; IPL, inner plexiform layer; IS, inner segment; ONL, outer nuclear layer; OPL, outer plexiform layer; OS, outer segment; RPE, retinal pigment epithelium. The scale bars in each panel indicate 500 μm while those in the insets in panels **E** and **F** indicate 100 μm . Tam – tamoxifen.

Figure S4

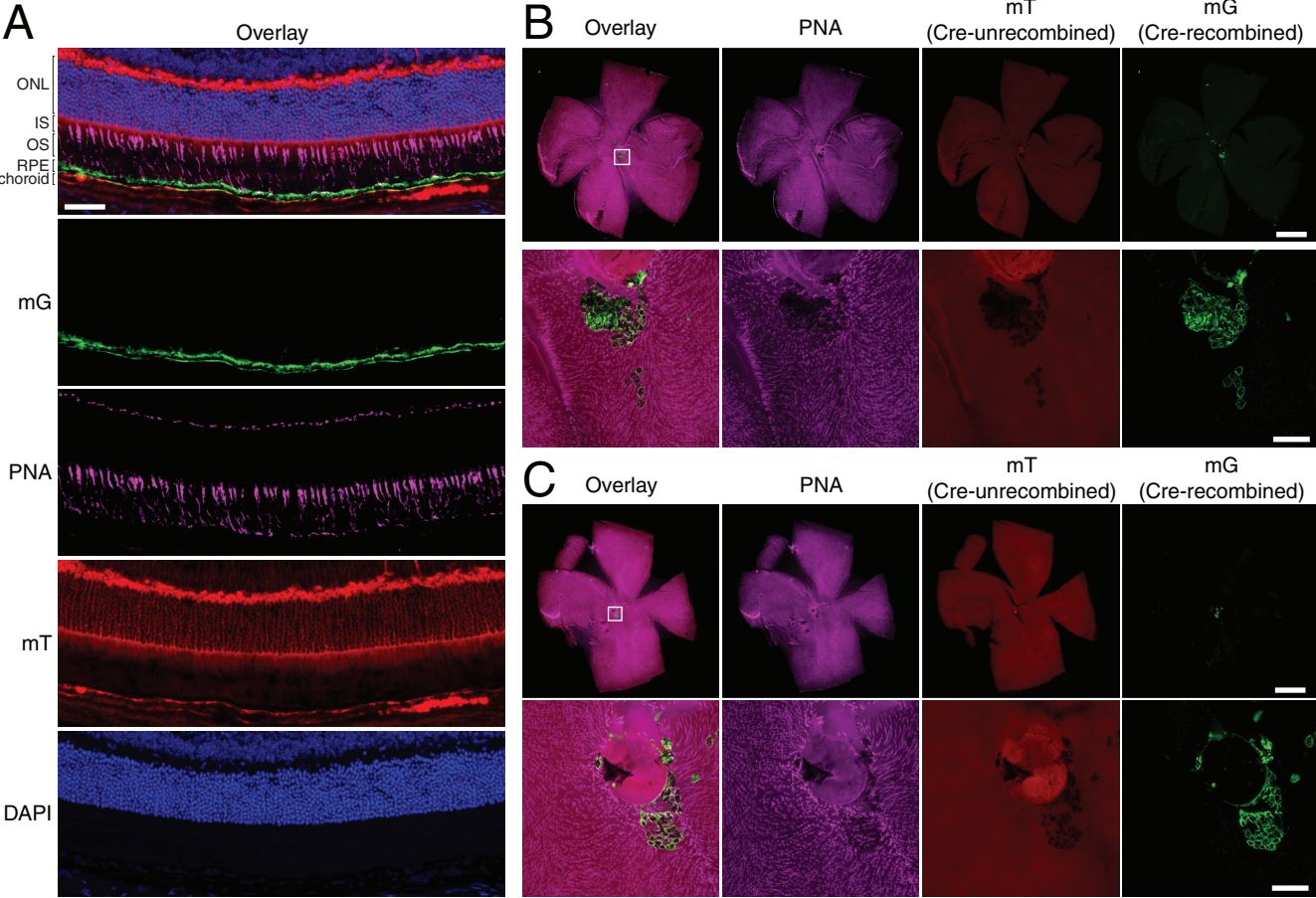


Figure S4. *Lack of mG fluorescence co-localization with cone photoreceptors.* **A)** Retina cryosections from *Cre^{+/-} mT/mG^{+/-}* mice treated with IP tamoxifen starting on PD21 were co-stained with peanut agglutinin, which selectively stains cone photoreceptors, and DAPI. The top image shows the composite fluorescence signal and those below in are the individual channels. No co-localization of the PNA and mG signals was observed. The scale bar represents 50 μm . Whole retina flatmounts from female (**B**) or male (**C**) *Cre^{+/-} mT/mG^{+/-}* mice treated with IP tamoxifen on PD21 were co-stained with PNA to allow visualization of cone photoreceptors. The retinae were imaged with their RPE-associated sides facing the camera. The leftmost images show the composite fluorescence signal while the images to the right show the individual channels. Zoomed images corresponding to the indicated box are shown below each whole-retina image. No mG signal is seen to co-localize with the PNA cone cell marker indicating that Cre recombinase is not expressed in cone photoreceptors in *Rpe65^{CreERT2}* mice. The zoomed images show mG-positive RPE cells that remained adhered to the neural retina during the dissection procedure. The scale bars in the whole retina images represent 1000 μm while those in the bottom zoomed images represent 100 μm .

Figure S5

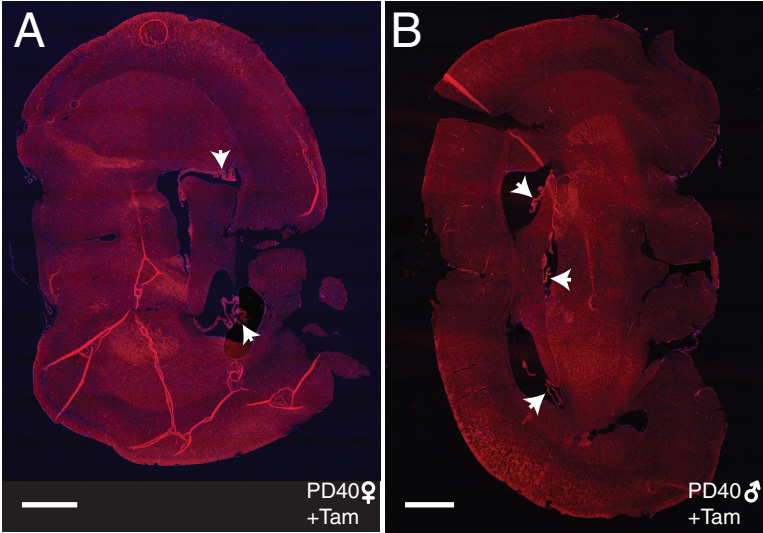


Figure S5. *Lack of Cre recombinase activity in Rpe65^{CreERT2} mouse brain as assessed with the mT/mG reporter.* Panels **A** and **B** show whole brain cryo-sections from *Cre^{+/-}* female (**A**) and male (**B**) mice following Cre induction with IP tamoxifen starting on PD21 (n = 2). Brains were processed 2 weeks after the last dose of tamoxifen. No green fluorescence was evident in the sections, including the choroidal plexus epithelium (white arrows) confirming the absence of Cre-mediated recombination in the brains of *Rpe65^{CreERT2}* mice. The scale bar indicates 1000 μ m.

Figure S6

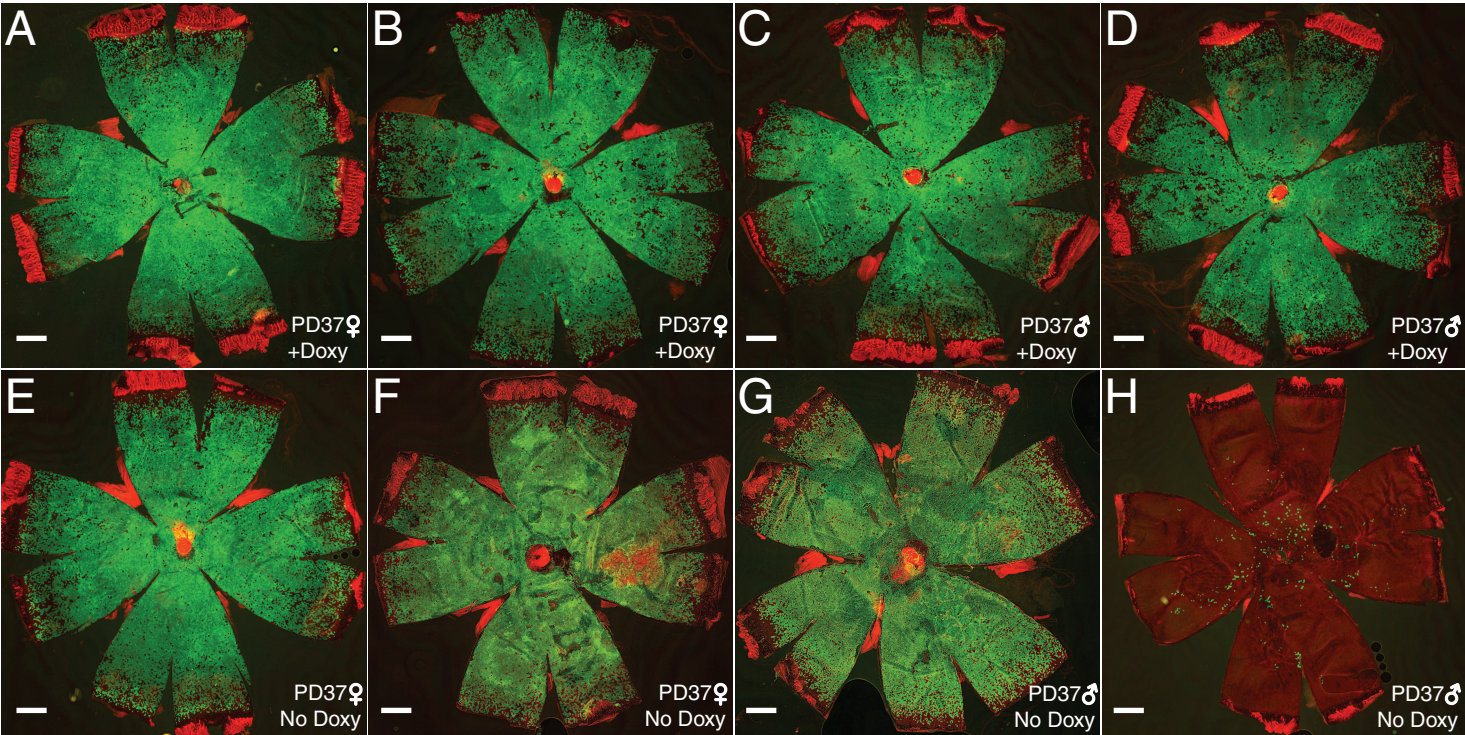


Figure S6. *Cre* recombinase activity in *tet-ON VMD2-Cre* mouse retina as assessed with the *mT/mG* reporter. Panels **A-D** show RPE flatmounts from representative *VMD2-Cre*^{+/-} female (**A, B**) and *VMD2-Cre*^{+/-} male (**C, D**) mice administered 8 mg of doxycycline (Doxy) in water by oral gavage for two consecutive days starting on PD21 (n = 4). The flatmounts were obtained two weeks after the final dose of doxycycline. Green fluorescence indicates cells where *Cre*-mediated recombination had occurred, whereas red fluorescence indicates un-recombined cells. Panels **E-H** show RPE flatmounts from representative *VMD2-Cre*^{+/-} female (**E, F**) and *VMD2-Cre*^{+/-} male (**G, H**) mice administered vehicle (water) by oral gavage for two consecutive days starting on PD21 (n = 4). The flatmounts were obtained two weeks after the final water gavage. Note the variability in doxycycline-independent induction in these mice. The scale bar indicates 500 μm.

Figure S7

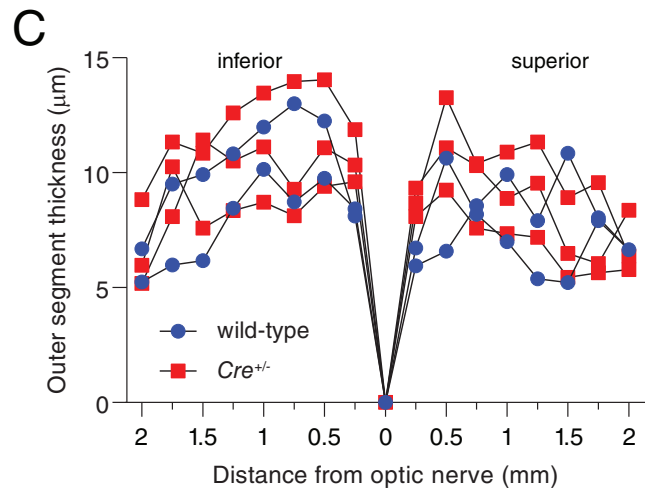
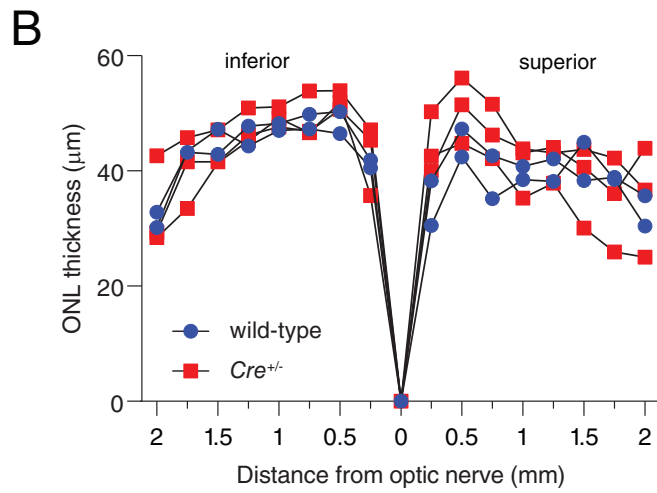
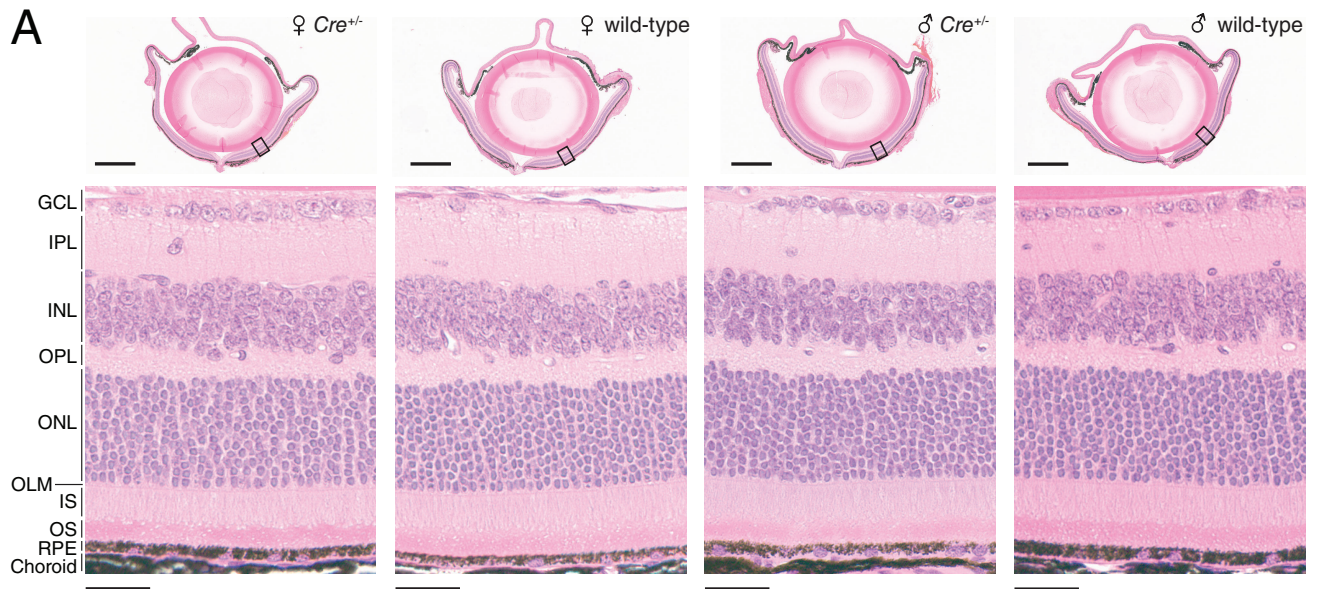


Figure S7. *Retinal histology of 2-month old, IP tamoxifen-treated Cre^{+/-} mice and wild-type littermates.* **A)** Hematoxylin and eosin-stained retinal paraffin sections from Cre^{+/-} mice (n = 3) and wild-type littermate controls (n = 2) that were treated with IP tamoxifen for five days starting on PD21. The upper panels show the complete retinal section. The lower panels show zoomed images corresponding to the areas boxed in black rectangles in the upper panels. The scale bars in the upper and lower panels indicate 600 μm and 25 μm, respectively. Abbreviations: GCL, ganglion cell layer; INL, inner nuclear layer; IPL, inner plexiform layer; IS, inner segment; OLM, outer limiting membrane; ONL, outer nuclear layer; OPL, outer plexiform layer; OS, outer segment; RPE, retinal pigment epithelium. **B)** Quantification of the outer nuclear layer (ONL) thickness as a function of distance from the optic nerve revealed comparable thickness between Cre^{+/-} and wild-type mice. **C)** Quantification of the photoreceptor outer segment thickness as a function of distance from the optic nerve revealed comparable thickness between Cre^{+/-} and wild-type mice.

Table S1. RPE-targeted Cre mouse lines reported to date.

Name	Targeting strategy	Expression driver	Cre recombinase	Characteristics	Reference
<i>Mct3</i> -Cre	random transgene integration	Monocarboxylate transporter 3 (<i>Mct3</i>) promoter	iCreER ^{T2}	<ul style="list-style-type: none"> Tamoxifen inducible Low and mosaic recombination 	(2)
tet-ON VMD2-Cre	random transgene integration	Vitelliform macular dystrophy-2 (<i>Vmd2</i>), promoter	Cre	<ul style="list-style-type: none"> Doxycycline inducible Inducer-independent Cre expression Mosaic recombination 	(3)
<i>Tyr</i> -CreER ^{T2}	random transgene integration	Hsp70 promoter and <i>Tyr</i> enhancer	CreER ^{T2}	<ul style="list-style-type: none"> Tamoxifen inducible Prenatal and post-natal induction possible Mosaic recombination Cre expression in ciliary body 	(4)
<i>Mart1</i> -Cre	random transgene integration	Melanoma-Associated Antigen Recognized by T-cells (MART-1) promoter	Cre	<ul style="list-style-type: none"> Constitutive Cre activity Recombination not specific to the RPE 	(5)
<i>Trp1</i> -Cre	random transgene integration	Tyrosinase-related protein-1 (<i>Trp1</i>) promoter	Cre	<ul style="list-style-type: none"> Constitutive Cre activity Mosaic recombination Recombination not specific to the RPE Intrinsic RPE toxicity 	(6, 7)
<i>Dct</i> -Cre	random transgene integration	dopachrome tautomerase (<i>Dct</i>) promoter	Cre	<ul style="list-style-type: none"> Constitutive Cre activity Recombination not specific to the RPE 	(8)
<i>BEST1</i> -Cre	random transgene integration	Bestrophin-1 (<i>Best1</i>) promoter	Cre	<ul style="list-style-type: none"> Constitutive Cre activity Mosaic recombination Intrinsic RPE toxicity 	(9, 10)
<i>BEST1</i> -CreER ^{T2}	Targeted transgene integration at ROSA26 locus	Bestrophin-1 (<i>Best1</i>) promoter	CreER ^{T2}	<ul style="list-style-type: none"> Mosaic recombination Cre activity in a population of Müller glia 	(11)

Table S2. PCR conditions used for animal genotyping

Locus	Primers (5' – 3')	Cycling parameters		Bands generated (bp)	Notes	Reference
		Time	Temperature (°C)			
<i>Rpe65</i> ^{CreERT2}	<p>RPE65_E14f: CTTCCATGGACTGTTC AAAAGATCC</p> <p>RPE65_E14r: AACTTCCAGGAGTAAG TTCTGTCC</p> <p>RPE65_CreERT2: GCATAACCAAGTGAAC AGCATTG</p>	5 min 45 sec 30 sec 1 min 10 min	94 94 63 x 35 cycles 72 72	449 (<i>Cre</i> ^{+/+}) 449 and 146 (<i>Cre</i> ^{+/-}) 146 (wild-type)		This work
mT/mG	<p>12177: CTTTAAGCCTGCCAG AAGA</p> <p>30297: TAGAGCTTGCGGAACC CTTC</p> <p>30298: AGGGAGCTGCAGTGG AGTAG</p>	5 min 45 sec 30 sec 1 min 10 min	94 94 60 x 35 cycles 72 72	128 (mT/mG ^{+/+}) 128 and 212 (mT/mG ^{+/-}) 212 (wild-type)		(12)
RPE65 (L/M 450 variation)	<p>RPE65f: GCATACGGACTTGGGT TGAATCAC</p> <p>RPE65r: GGTTGAGAAACAAAGA TGGGTTTCAG</p>	5 min 45 sec 30 sec 1 min 10 min	94 94 60 x 35 cycles 72 72	231 (M450)* 231, 142, 89 (M450/L450)* 142, 89 (L450)*	* Following digestion of the PCR product with MwoI. The L450 variant contains an MwoI site at nt 142–151 that is absent in the M450 variant. Digestion with MwoI therefore results in two bands of ~142 and 89 bp for L450, and one band of 231 bp for M450.	(13)
rd8	<p>mCrb1-mF1: GTGAAGACAGCTACAG TTCTGATC</p> <p>mCrb1-mF2: GCCCTGTTTGCATGG AGGAAACTTGAAGAC AGCTACAGTTCTTCTG</p> <p>mCrb1-mR: GCCCCATTTGCACACT GATGAC</p>	5 min 45 sec 30 sec 1 min 10 min	94 94 60 x 35 cycles 72 72	220 (wild-type)* 244 (rd8)*	* The mF1 x mR (wild-type) and mF2 x mR (rd8) reactions are run separately because of the similarity in their product sizes.	(14)
VMD2-Cre	<p>VMD2-Cre_f: GAACCTGATGGACATG TTCAG</p> <p>VMD2-Cre_r: AGTGC GTTCGAACGCT AGAGC</p>	5 min 45 sec 30 sec 1 min 10 min	94 94 60 x 35 cycles 72 72	375 (<i>Cre</i> [*])	* The genotyping protocol does not distinguish between heterozygous and homozygous Cre animals	(15)

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