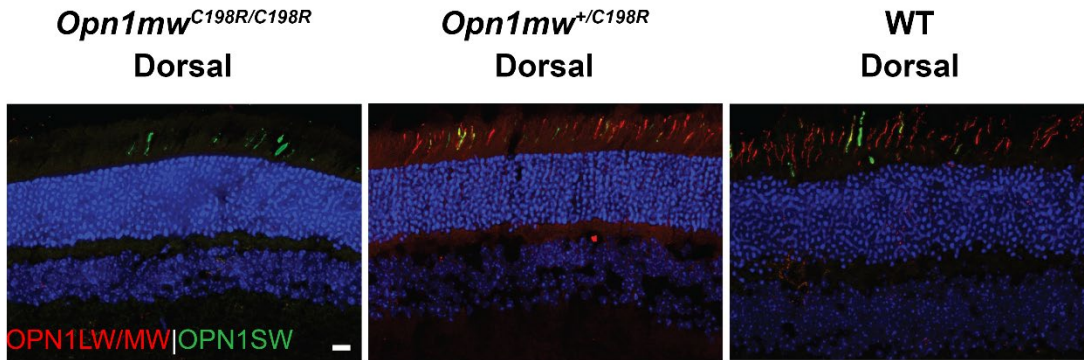


1 SUPPLEMENTAL MATERIALS

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5 **Supplemental Figure 1. M- and S-opsin localization in homozygous and heterozygous**

6 ***Opn1mw^{C198R}* mice.** Representative IHC images of the dorsal region of WT, heterozygous

7 *Opn1mw^{+/C198R}* female, and homozygous *Opn1mw^{C198R}* cross-sections collected at P30 and

8 stained with antibodies against S-opsin (green) and M-opsin (red). Scale bar = 20 μ m.

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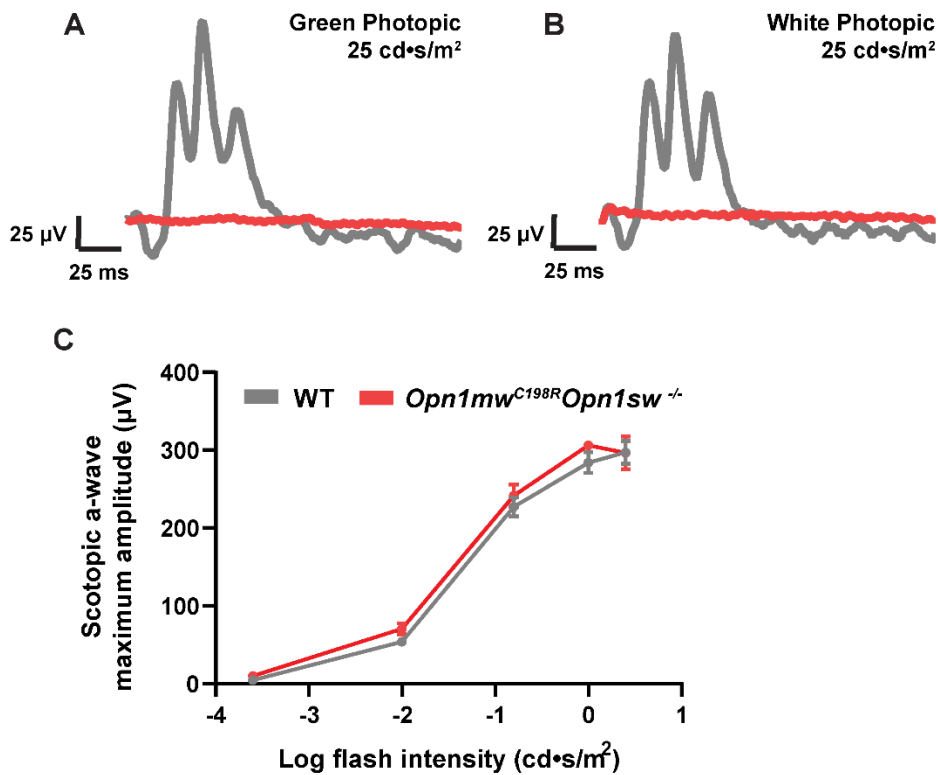
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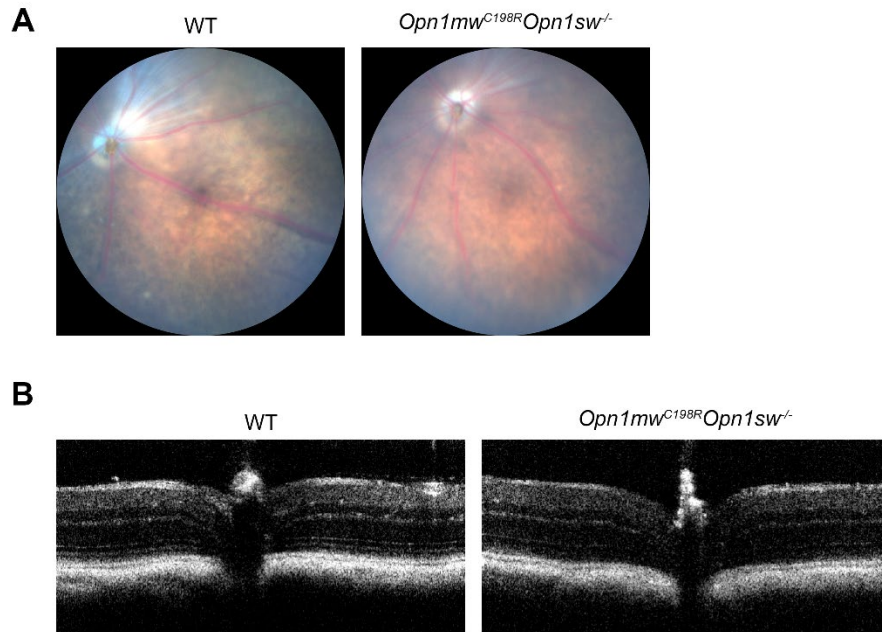
20 **Supplemental Figure 2. *Opn1mw^{C198R}Opn1sw^{-/-}* cones are non-functional while rods are**
21 **normal. (A-B)** Representative waveforms of P30 WT (grey) and *Opn1mw^{C198R}Opn1sw^{-/-}* mice
22 (red) following ERG recordings at 25 cd•s/m² under (A) middle-wavelength (green) and (B)
23 photopic (white) light. (C) Sensitivity curve demonstrating the scotopic a-wave maximum
24 amplitude of WT (grey) and *Opn1mw^{C198R}Opn1sw^{-/-}* mice (red) at P30 following ERG recordings
25 under dim light conditions at various light intensities (0.0003, 0.01, 0.2, 1.0, and 2.5 cd•s/m²).
26 Data is represented as mean ± SEM (n=6).

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33 **Supplemental Figure 3. Normal morphology of *Opn1mw^{C198R}Opn1sw^{-/-}* retinas. (A)**

34 Representative bright field fundus images and **(B)** OCT images from 10-month old WT and

35 *Opn1mw^{C198R}Opn1sw^{-/-}* mice revealed no gross morphological abnormalities (n=3).

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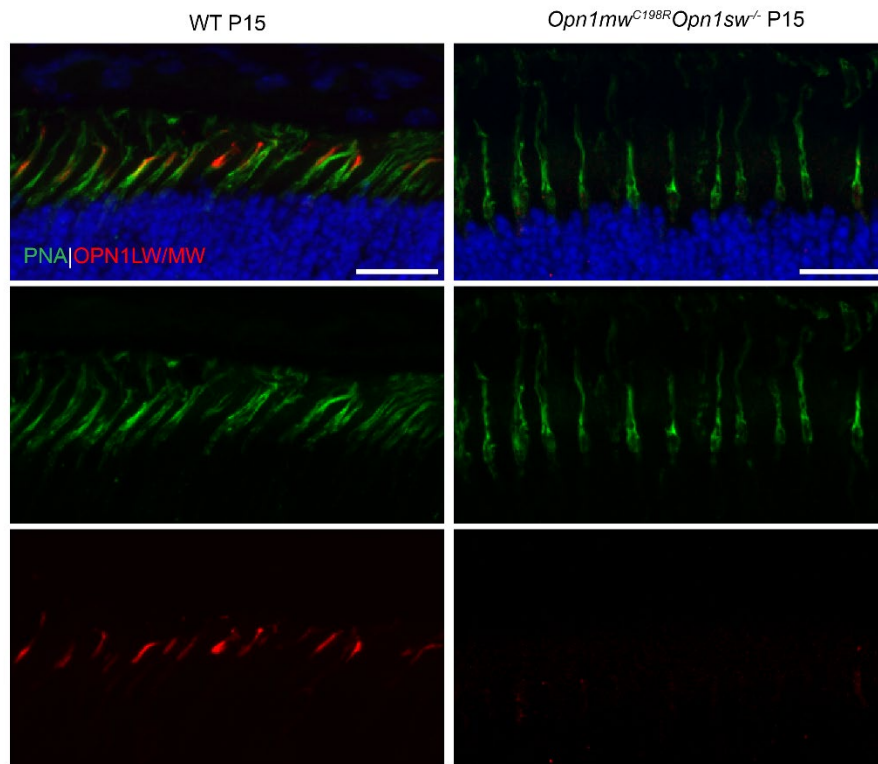
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44 **Supplemental Figure 4. *Opn1mw^{C198R}Opn1sw^{-/-}* cones lack L/M-opsin expression at P15.**

45 Representative IHC images of WT (left) and *Opn1mw^{C198R}Opn1sw^{-/-}* (right) cross-sections stained
46 with an antibody against L/M-opsin and with PNA. Scale bar = 20 μ m.

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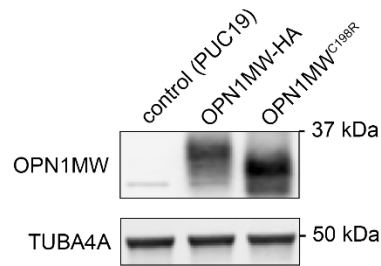
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58 **Supplemental Figure 5. Antibody against L/M-opsin recognizes mutant OPN1MW^{C198R}.**

59 Western blot analysis of HEK293T cell lysates following single-plasmid transfection with PUC19
60 (left lane), WT OPN1MW-HA (middle lane), or mutant OPN1MW^{C198R} protein (right lane), showing
61 that the antibody against L/M-opsin recognizes both WT and mutant OPN1MW^{C198R} (OPN1MW).
62 TUBA4A was used as a loading control.

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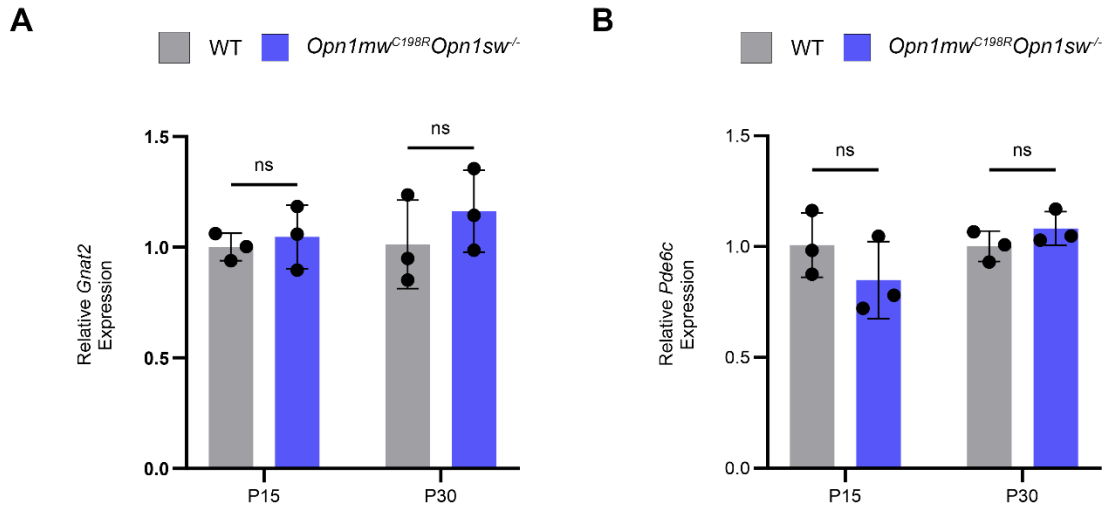
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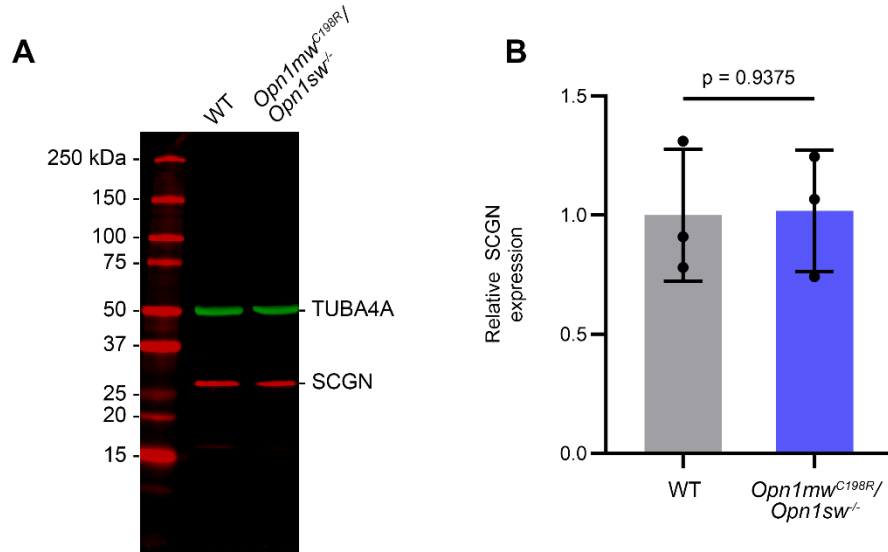
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77 **Supplemental Figure 6. Gene expression of *Pde6c* and *Gnat2* is comparable between**
78 ***Opn1mw^{C198R}Opn1sw^{-/-}* and WT at P15 and P30.** Real-time qPCR of **(A) *Pde6c*** and **(B) *Gnat2***
79 mRNA levels in *Opn1mw^{C198R}Opn1sw^{-/-}* retinas at P15 and P30 relative to age-matched WT
80 controls. Data represented as mean \pm SD, 2-way ANOVA ($n = 3$; $*p \leq 0.05$, $**p < 0.002$, $***p <$
81 0.001).

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90 **Supplemental Figure 7. *Opn1mw^{C198R}Opn1sw^{-/-}* cones exhibit normal protein levels of the**
91 **cone bipolar cell marker secretagogin compared to age-matched WT controls. (A)** Western
92 blot and **(B)** quantification of 10-month old WT (left lane) and *Opn1mw^{C198R}Opn1sw^{-/-}* (right lane)
93 retinal lysates with an antibody labeling the cone bipolar cell marker secretagogin (SCGN, red).
94 TUBA4A (green) was used as a loading control. Data represented as mean ± SD, unpaired 2-
95 tailed t-test with Welch's correction (n=3; $p = 0.9375$).

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