

Description of Additional Supplementary Files

File name: Supplementary Data 1

Description: Source data for Figure 4h and 4q. For measuring *ey*s mRNA expression by quantitative RT PCR, Ct values for *ey*s were normalized against those for *gapdh*. Data for SPMO1, SPMO2, and SPMO3 were further normalized against that for ContMO (Figure 4h). Number of cells with mislocalized rhodopsin within a retinal section was counted (Figure 4q).