SUPPLEMENTARY MATERIAL

Table S1. PCR primer sequences for genotyping

Transcript	Forward primer sequence (5'-3')	Reverse primer sequence (5'-3')
bsx	ATTGCAAAAGGAATGCAGATG	ATTGTCGTCCAGCGTGTATCT
gfp	TTTTCAAGAGTGCCATGCCC	CCATGTGTAATCCCAGCAGC
egfp	CAGATGAACTTCAGGGTCAG	CTGCCGTCCTCGATGTTGTG

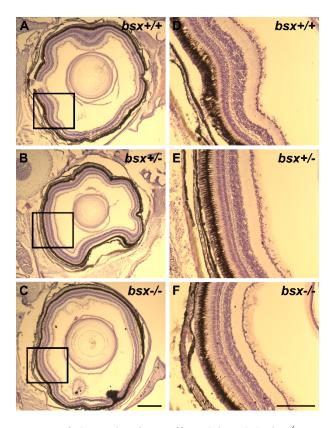


Figure S1. Morphological structure of the retina is unaffected in adult $bsx^{-/-}$ mutant zebrafish. Representative images of sagittal sections of $bsx^{-/-}$ mutant zebrafish eyes (C, F) stained in cresyl violet. Images on the right (D-F) represent the boxed regions on the left (A-C) at higher resolution. Wildtype $(bsx^{+/+})(A, D)$ and heterozygote $(bsx^{+/-})(B, E)$ controls are shown for comparison. For examined eyes, n=4 per genotype. Scale bars, 500 µm (left) and 200 µm (right).

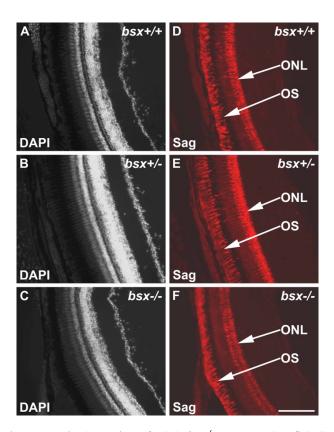


Figure S2. S-antigen protein is present in the retina of adult $bsx^{-/-}$ mutant zebrafish. Immunofluorescent staining for S-antigen (Sag) on sagittal sections of adult $bsx^{-/-}$ mutant zebrafish retina (**F**); sections were also stained in DAPI (**C**). Wildtype $(bsx^{+/-})(\mathbf{A}, \mathbf{D})$ and heterozygote $(bsx^{+/-})(\mathbf{B}, \mathbf{E})$ controls are shown for comparison. Scale bar, 100 μ m. ONL, outer nuclear layer, OS, photoreceptor outer segments.

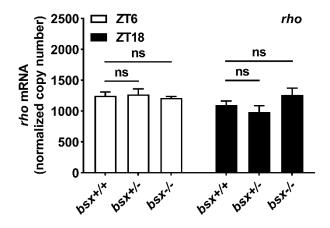


Figure S3. Abundant *rho* transcript levels are maintained in whole eyes of adult $bsx^{-/-}$ mutant zebrafish. Analyses of transcript levels of rhodopsin (rho) at daytime (ZT6) and nighttime (ZT18) as determined by qRT-PCR. Copy numbers were normalized to the geometric means of copy numbers of eef1a1 and actb2, which were detected at high and stable levels in all samples; copy numbers were further divided by 10000 to account for abundance of transcripts. Two-way ANOVA did not detect significant effects of genotype or time of day. Transcript levels of homozygous ($bsx^{-/-}$) and heterozygous ($bsx^{+/-}$) mutants were compared to those of wildtype ($bsx^{+/+}$) controls. Values on the graphs represent mean \pm SEM; n=4, where each sample is pooled tissue from 3 animals. P-values as determined by Dunnett's multiple comparisons test: ns, not significant. ZT, Zeitgeber time.