1	Title: A critical role for MSI1 and MSI2 in photoreceptor morphogenesis
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3	Authors: Jesse Sundar ¹ , Fatimah Matalkah ¹ , Bohye Jeong ¹ , Peter Stoilov ¹ *, and Visvanathan
4	Ramamurthy ^{1,2,3,*}
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6	Affiliations: Departments of Biochemistry ¹ , Ophthalmology and Visual Sciences ² , and
7	Neuroscience ³ ; Robert C. Byrd Health Sciences Center, West Virginia University; Morgantown,
8	West Virginia, USA, 26505;
9	
10	Address for correspondence:
11	Peter Stoilov, Department of Biochemistry, West Virginia University School of Medicine; 1
12	Medical Center Dr. Morgantown, WV, USA, 26505; Email: pstoilov@hsc.wvu.edu ; Telephone:
13	304-293-6334; Fax: 304-293-6846;
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15	Visvanathan Ramamurthy, Department of Biochemistry, West Virginia University School of
16	Medicine; 1 Medical Center Dr. Morgantown, WV, USA, 26505; Email:
17	ramamurthyv@hsc.wvu.edu; Telephone: 304-293-2479; Fax: 304-293-6846;
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ABSTRACT

We previously proposed a role for the Musashi proteins, MSI1 and MSI2, in photoreceptor cell development that is mediated by their ability to control alternative splicing. Photoreceptors with simultaneous deletion of Msi1 and Msi2 did not respond to light, displayed severely disrupted OS morphology and axonemal defects. At postnatal day 5, we observed an increase in proliferating retinal progenitor cells in the knockout animals, suggesting delay in photoreceptor development. The loss of Musashi prevented the use of photoreceptor-specific exons in transcripts important for OS morphogenesis, ciliogenesis and synaptic transmission. However, deletion of the photoreceptor-specific exons in Ttc8, Cc2d2a, Cep290, Cacna2d4, and Slc17a7 did not impair retinal development or visual function. We demonstrate a critical role for Musashi in the morphogenesis of terminally differentiated photoreceptor neurons. This role is in stark contrast with the canonical function of the two proteins in maintenance and renewal of stem cells.

INTRODUCTION

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In eukaryotes, alternative splicing of pre-mRNA is a process that increases protein diversity and controls gene expression. Diversification of proteomes through alternative splicing is a defining characteristic of metazoans and was expanded dramatically in bilaterians¹. Alternative splicing is particularly prevalent in vertebrate neurons and is critical for the development and function of vertebrate nervous systems^{2–7}. We previously showed that photoreceptor neurons exploit a unique splicing program⁸. Motif enrichment analysis suggested that Musashi-1 (MSI1) and Musashi-2 (MSI2), promote the use of photoreceptor specific exons⁸. We further showed that MSI1 is critical for utilization of photoreceptor specific exon in Tetratricopeptide repeat domain protein-8 (Ttc8)⁸. In addition, Musashi promotes the splicing of several photoreceptor specific exons when over-expressed in cultured cells⁸. Recently, analysis of a comprehensive gene expression data set that spanned multiple tissues and cell types from mice and human proved that photoreceptors utilize a unique set of alternative exons that are primarily regulated by MSI1 and MSI29. Furthermore, the work by Ling et. al. demonstrated that Msi1 transcript levels are upregulated in the developing rod photoreceptors and reach exceptionally high levels compared to all other cell types or tissues in the data set. The MSI1 and MSI2 proteins have two highly conserved RNA binding domains (RBDs) in the N-terminal region which show close to 90% sequence identity and recognize a similar UAG motif in RNA¹⁰. The two RBDs of MSI1 and MSI2 are followed by a less conserved Cterminal region which shows approximately 70% sequence identity¹¹. The high degree of sequence identity between the MSI1 and MSI2 results in functional redundancy between the two proteins ^{12,13}. The canonical function of the Musashi proteins is to control mRNA translation in

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the cvtosol^{14,15}, where they can either block or enhance translation of mRNA depending on cellular context^{16–21}. Vertebrate photoreceptors are neurons specialized in detecting and transducing light stimuli. Photoreceptors are characterized by segmented morphology which compartmentalizes phototransduction, core cellular functions, and synaptic transmission. The light sensing machinery is confined to the outer segment, a stack of membranes that is elaborated by cell's modified primary cilium. The outer segment is dynamic structure that is remade every 7 to 10 days. Consequently, maintenance of the outer segment requires high rate of transport of membranes and proteins through the connecting cilium²². Interestingly, the predicted splicing targets of Musashi in photoreceptors include premRNAs from ciliary (Ttc8, Cep290, Cc2d2a, Prom1) and synaptic-associated genes (Cacna2d4, Slc17a7)^{23–29}. These genes have been showed to be crucial for photoreceptor development and fucntion^{23–29}. We proposed that production of photoreceptor specific splicing isoforms that is promoted by Musashi is necessary for the development and maintenance of photoreceptor cells in $vivo^8$. To test if Musashi drives photoreceptor development and function, we removed Msi1 and Msi2 in the developing retina and rod photoreceptor cells. We find that Musashi proteins are essential for photoreceptor function, morphogenesis, and survival but not their specification. Specifically, the Musashi proteins are crucial for outer segment (OS) and axoneme development. As expected, disruption of the Musashi genes led to loss of expression of photoreceptor specific splicing isoforms. Surprisingly, deleting the photoreceptor-specific exons of the, Ttc8, Cc2d2a, Cep290, Cacna2d4, and Slc17a7 genes does not produce a detectable phenotype suggesting that

the loss of vision in the Musashi mutants is likely independent of the role of the Musashi proteins

in controlling alternative splicing.

MATERIALS AND METHODS

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Generation of mice and genotyping

Mice carrying floxed alleles for *Msi1* and *Msi2* were provided by Dr. Christopher Lengner from the University of Pennsylvania. Six3-Cre transgene or Nrl-Cre transgenes were used to delete the floxed alleles in developing retina or rod photoreceptors. All mouse lines were devoid of naturally occurring rd1 and rd8 alleles^{30,31}. Males hemizygous for the Six3-Cre transgene or Nrl-Cre transgene and floxed for either Msi1, Msi2, or both Msi1 and Msi2 were mated with females floxed for either Msi1, Msi2, or both Msi1 and Msi2 to obtain experimental knockout mice and littermate control. The offspring of breeding pairs were genotyped using PCR of DNA derived from ear biopsies. The Msi1 wildtype and floxed alleles were identified using following primers: (5'-CGG ACT GGG AGA GGT TTC TT-3' and 5'-AGC TCC CCT GAT TCC TGG T-3')³². The Msi2 wildtype and floxed alleles were identified by using following primers: (5'-GCT CGG CTG ACA AAG AAA GT-3' and 5'-TCT CCT TGT TGC GCT CAG TA-3')³². The presence of the Six3 Cre transgene was determined using following primers: (5'-CCC AAA TGT TGC TGG ATA GT-3' and 5'-CCC TCT CCT CTC CCT CCT-3'). The presence of the Nrl Cre transgene was determined using following primers: (5'-TTT CAC TGG CTT CTG AGT CC-3' and 5'-CTT CAG GTT CTG CGG GAA AC-3'). The presence of Cre recombinase was determined using following primers: (5'-CCT GGA AAA TGC TTC TGT CCG-3' and 5'-CAG GGT GTT ATA AGC AAT CCC-3')³³. The knockout of the photoreceptor specific exons in Ttc8, Cep 290, and Cc2d2a was done using CRISPR/Cas9. Two guide RNAs were used for each alternative exon to target intronic sites upstream and downstream of the alternative exons to cause their deletion. The guide RNAs were commercially synthesized by Synthego and had the following targeting

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sequences: Ttc8-1 GUUCCUGGAAAGCGUUAAGA, Ttc8-2 CAGCACAUUUUCCAAUCUUC, Cep290-1 CCCUAAGGGUAGUUAAAGCU, Cep290-2 CGUUUGUUCUUCAAAAGGAC, Cc2d2a-1 GCAUGUCACUCAUCUACCAU, and Cc2d2a-2 AGCGUUUAACUGAUGACUGC. The guide RNAs and Cas9 (Thermo Fisher) were assembled into RNPs and electroporated into zygotes by the WVU transgenics core facility. The founders were back-crossed to wild type C57 Black 6/J for 5 generations to eliminate off-target mutations. The deleted exons were genotyped using the following primers: Ttc8-CRISP-F ctccccatcctagccaatct, Ttc8-CRISP-R tgtgcacaggcaaaataagc, Cep290-CRISP-F gagcaaacacagcagtcagc, Cep290-CRISP-R tettetgaccgccacetact, Cc2d2a-CRISP-F gctacacacatggctggatg, Cc2d2a-CRISP-R ctctcatgtgacaggcagga. All experiments were conducted with the approval of the Institutional Animal Care and Use Committee at West Virginia University. All experiments were carried out with adherence to the principles set forth in the ARVO Statement for the Ethical Use of Animals in Ophthalmic and Vision Research which advocates the use of the minimum number of animals per study needed to obtain statistical significance. Electroretinography, Immunoblotting, and Reverse Transcriptase PCR Electroretinography, immunoblotting, and reverse transcriptase PCR were conducted using previously described protocol from our laboratory^{8,34,35}. **Immunofluorescence Microscopy** Immunofluorescence microscopy was carried out using a modified procedure in our laboratory^{34,35}. Briefly, eyes were enucleated, and the cornea and lens were discarded. After dissection, eyes were fixed by immersion in 4% paraformaldehyde in PBS for one hour. After washing the eyes in PBS three times for ten minutes each, they were dehydrated by overnight

incubation in 30% sucrose in PBS. Eyes were then incubated in a 1:1 solution of OCT:30% sucrose in PBS for one hour and frozen in OCT (VWR). The frozen tissues were sectioned using a Leica CM1850 cryostat for collecting serial retinal sections of 16µm thickness. The retinal cross-sections were then mounted onto Superfrost Plus microscope slides (Fisher Scientific). Slide sections were then washed and permeabilized with PBS supplemented with 0.1% Triton X-100 (PBST) and incubated for one hour in a blocking buffer containing 10% goat serum, 0.3% Triton X-100, and 0.02% sodium azide in PBS. Retinal sections were then incubated with primary antibody in a dilution buffer containing 5% goat serum, 0.3% Triton X-100, 0.02% sodium azide, and primary antibody at 1:500 dilution in PBS overnight at 4°C followed by three 5 minute washes using PBST. Sections were then incubated in the same dilution buffer containing secondary antibody and DAPI at 1:1000 for one hour. Slides were washed with PBST three times for five minutes each before treating with Prolong Gold Antifade reagent (ThermoFisher) and securing the coverslip. The images were collected using a Nikon C2 Confocal Microscope.

Retinal histology of the mouse models

Following euthanasia, eyes were enucleated using a C-shaped forceps after marking the superior pole and incubated in Z-fixative for >48 hours before shipment and tissue processing by Excalibur Pathology Inc. (Norman, OK)^{34,35}. The embedding, serial sectioning, mounting, and hematoxylin/eosin (H&E) staining were performed by Excalibur Pathology. A Nikon C2 Microscope equipped with Elements software was used to image the slides.

Transmission Electron Microscopy

After euthanasia, a C-shaped forceps was used to enucleate the eye, and the cornea was discarded^{34,35}. Eyes were then incubated in a fixative solution containing 2.5% glutaraldehyde

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and 2% paraformaldehyde in 100mM sodium cacodylate buffer at pH 7.5 for 45 minutes before removal of the lens. After lensectomy, eyes were placed back into fixative for 72 hours before shipment, tissue processing, and imaging at the Robert P. Apkarian Integrated Electron Microscopy Core at Emory University. **Antibodies** The following primary antibodies were used throughout our studies: rat anti-MSI1 (1:1000; Medical and Biological Laboratories, Woburn, MA), rabbit anti-MSI2 (1:2000; Abcam, Cambridge, MA), mouse anti-β-tubulin (1:10,000; Sigma-Aldrich, St. Louis, MO), rabbit antiphospho-histone H3 (1:500; Cell Signaling, Danvers, MA), mouse anti-Ki67 (1:500; BD Biosciences, San Jose, CA), rabbit anti-active Caspase-3 (1:500; Promega, Madison, WI), rhodamine peanut agglutinin (1:1000; PNA: cone OS sheath marker, Vector laboratories, Burlingame, CA), rabbit anti-peripherin-2 (1:2000) was a kind gift by Dr. Andrew Goldberg from Oakland University, rabbit anti-PDE6β (1:2000; ThermoFisher, Waltham, MA), mouse anti-acetylated α-tubulin (1:1000; Santa Cruz, Dallas, TX), guinea pig anti-MAK (1:500; Wako, Richmond, VA), mouse anti-glutamylated tubulin (1:500; AdipoGen Life Sciences, San Diego, CA), mouse anti-Ttc8 (1:1000; Santa Cruz, Dallas, TX), rabbit anti-Ttc8 Exon 2A (1:1000; custom made⁸), mouse anti-GAPDH (1:10,000; Fitzgerald, Acton, MA), and 4',6-diamidino-2phenylindole (DAPI: nuclear counterstain; 1:1000; ThermoFisher, Waltham, MA).

RESULTS

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Validation of the conditional knockout mouse models

We analyzed the expression of Musashi proteins in various tissues from adult mice. Out of all the tissues we tested, retina showed the highest expression of MSI1 and MSI2 proteins (Figure 1A), in line with the previously reported high transcript levels for Msi1 and Msi2 in rod photrecptors⁹. To test the biological significance of Musashi protein expression in the murine retina, we used Cre-LoxP conditional recombination to remove either Msi1, Msi2, or both the Msi1 and Msi2 genes throughout the entire retina and ventral forebrain using the Six3 Cre transgene (Supplementary Figure 1)³⁶. Throughout this work, we refer to *Musashi* floxed mice which are hemizygous for the Six3 Cre transgene as ret-Msi-/- mice. The conditional recombination results in the deletion of Msi1's transcription start site, exon 1, and exon 2 (Supplementary Figure 1) 13 . For Msi2, the transcription start site and the first four exons are removed after cre-mediated recombination (Supplementary Figure 1)¹³. The ablation of MSI1 and MSI2 was confirmed by immunoblotting retinal lysates from knockout mice at postnatal day 10 (PN10) (Figure 1B). Immunofluorescence microscopy of retinal cross sections obtained from the knockout mice also affirmed the absence of MSI1 and MSI2 expression in the retina (Figure 1C). Notably, MSI2 protein levels were moderately but reproducibly upregulated in Msi1 knockout retina (Figure 1B, Supplementary Figure 2). We did not observe the inverse, upregulation of MSI1 protein in *Msi2* knockouts (Supplementary Figure 2). These data show that MSI1 protein regulates the expression of MSI2 and could indicate existence of a homeostatic mechanism for regulation overall Musashi protein levels.

The Musashi proteins are crucial for photoreceptor function

To determine if the Musashi proteins are required for photoreceptor function, we

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performed electroretinographic (ERG) recordings of the Musashi conditional knockout mice at PN16 and monitored for changes in retinal function up to PN180. Figure 2A shows the scotopic and photopic ERG waveforms of the ret-Msi1-/-, ret-Msi2-/-, and ret-Msi1-/-:Msi2-/- mice at PN16 immediately after mice open their eyes³⁷. When both *Musashi* genes are removed, no scotopic or photopic retinal function remains as shown by absence of conspicuous "a"-waves and "b"-waves (Figure 2A). However, significant retinal function remains in the ret-Msi1-/- and ret-Msi2-/- single knockout mice. We characterized the retinal function of the ret-Msi1-/- and ret-Msi2-/- mice further to see if there was a photoresponse deficit at higher light intensities or as the mice aged (Figure 2B-E). In ret-Msi1-/- mice, there was a statistically significant reduction in photoreceptor "a"-wave amplitudes at almost all light intensities (Figure 2B). This reduction in the photoreceptor "a"-wave amplitude persisted in ret-Msi1-/- mice up to PN180 (Figure 2C). On the other hand, ret-Msi2-/- mice at PN16 had normal photoreceptor function at all the light intensities we tested (Figure 2D). The "a"-wave amplitude began to decrease progressively in ret-Msi2-/- mice as they aged, and this became significant at PN120 (Figure 2E). Overall, this data shows that the Musashi proteins essential for photoreceptor function, and the two proteins are partially redundant. Intrinsic expression of Musashi in photoreceptors is crucial for photoreceptor function

We next sought to determine if the phenotype of the ret-Msi-/- mice was due to the absence of Musashi protein expression in photoreceptors or if deletion of Musashi in other retinal cell types or retinal progenitors were contributing to the loss of vision. To this end, we generated rod-specific Musashi conditional knockouts by crossing Musashi floxed mice with mice hemizygous for the Nrl Cre transgene where the Nrl promoter activates Cre expression in rod photoreceptors³⁸. Throughout this work, we refer to the *Musashi* floxed mice that are

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hemizygous for the Nrl Cre transgene as rod-Msi-/- mice. After confirming the loss of Musashi proteins in rod photoreceptors (Supplementary Figure 3), we used ERG to analyze the retinal function of the knockout mice after ablation of the Musashi genes in rods (Supplementary Figure 4A-E). Supplementary Figure 4A shows the scotopic and photopic ERG waveforms of the rod-Msi1-/-, rod-Msi2-/-, and rod-Msi1-/-:Msi2-/- mice at PN16. As observed in the ret-Msi1-/-:Msi2-/- mice, no conspicuous rod function was observed in the rod-Msi1-/-:Msi2-/- mice at PN16 which is demonstrated by absence of conspicuous "a"-wave under scotopic testing conditions (Supplementary Figure 4A). Again, we examined the rod-Msi1-/- and rod-Msi2-/single knockout mice further to see if the photoresponse phenotype was comparable to that obtained from the ret-Msi1-/- and ret-Msi2-/- mice. In rod-Msi1-/- mice at PN16, there was a reduction in photoreceptor "a"-wave amplitudes at multiple light intensities (Supplementary Figure 4B). This reduction in "a"-wave amplitude persisted as these mice aged up to PN180 (Supplementary Figure 4C). Contrarily, PN16 rod-Msi2-/- mice had no changes in photoreceptor function at all the light intensities examined (Supplementary Figure 4D). As observed in the ret-Msi2-/- mice, the "a"-wave amplitude began to decrease progressively as these mice aged, and this decrease became statistically significant at PN90 (Supplementary Figure 4E). The similar phenotypes of the ret-Msi and rod-Msi knockout mice shows that the intrinsic expression of Musashi proteins in photoreceptors is crucial for their function and that deletion of Musashi proteins in other cell types likely does not contribute significantly to the phenotype observed in the ret-Msi-/- mice. Therefore, throughout the rest of our studies, we focus on the ret-Msi1-/-:Msi2-/- mouse model for our experiments since there is a compensation in function occurring between MSI1 and MSI2 in the single knockout mice and to avoid confounding results that might be obtained when Msi1 and Msi2 are deleted only in rod but not cone photoreceptors.

Retinal cell death occurs in the absence of the Musashi proteins

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We next wanted to examine the mechanism behind the photoreceptor dysfunction seen in the ret-Msi1-/-: Msi2-/- mouse model. One of the common causes of a reduced ERG is photoreceptor cell death. Therefore, we performed histological analysis of the ret-Msi1-/-:Msi2-/- mice at PN5, PN10, PN16, and PN180 (Figure 3A-D). In ret-Msi1-/-: Msi2-/- mice at PN5, even before the neural retina has differentiated completely, there is a reduction in the neuroblast layer (NBL) thickness which was quantified across the superior-inferior axis (Figure 3A, left and right panels). There is also a more disordered arrangement of NBL nuclei in ret-Msi1-/-:Msi2-/mice with cells more tightly packed together compared to its littermate control (Figure 3A, left panel). At PN10, the outer nuclear layer (ONL), inner nuclear layer (INL), and ganglion cell layer (GCL) of the retina all form in ret-Msi1-/-:Msi2-/- mice but there is a reduction in the number of layers of photoreceptor nuclei (Figure 3B, left and right panels). At PN16, the number of layers of ONL nuclei continue to decrease suggesting that photoreceptor cell death is occurring (Figure 3C, left and middle panels). However, at this age, there are no statistically significant changes in the number of layers of INL nuclei (Figure 3C, left and right panels). By 6 months of age, the retina of ret-Msi1-/-: Msi2-/- mice was severely degenerated with a complete loss of ONL nuclei in addition to a significant reduction in the number of layers of INL nuclei (Figure 3D, left, middle, and right panels). Changes in proliferation during retinal development in the absence of MSI1 and MSI2 After observing a significant reduction in NBL nuclei in ret-Msi1-/-:Msi2-/- mice even before their retinas had fully differentiated, we wanted to investigate if this phenotype is related to the proliferation or premature death of retinal progenitor cells (RPCs). We collected retinal

cross sections of ret-Msi1-/-: Msi2-/- mice at PN5 and probed with antibodies against phospho-

histone H3 (PHH3) and Ki67, which are two commonly used markers of proliferation^{39–41}. We also examined apoptosis of retinal cells in *ret-Msi1-/-:Msi2-/-* mice by staining with an antibody against the apoptotic marker anti-active Caspase-3 (CASP3)^{40,42}. In *ret-Msi1-/-:Msi2-/-* mice, we witnessed a substantial increase of both PHH3+ and Ki67+ cells within the central retina whereas the littermate control had very few PHH3+ and Ki67+ cells within the central retina (Figure 4A and B). We also noticed a trend toward increase of PHH3+ and Ki67+ cell in the peripheral retina of *ret-Msi1-/-:Msi2-/-* mice (Figure 4A and B). No significant changes in proliferation marker staining were witnessed when comparing the superior and inferior retina of *ret-Msi1-/-:Msi2-/-* mice. CASP3 staining of retinal cross sections from *ret-Msi1-/-:Msi2-/-* mice showed no significant differences in the number of CASP3+ cells in either the central or peripheral retina (Figure 4C). Our data points to altered proliferation and not increased apoptosis at early stages of retinal development as the cause for the reduced number of NBL nuclei at PN5 in the Musashi knockout mice.

The Musashi proteins are required for OS development

Photoreceptor cells are present in the *ret-Msi1-/-:Msi2-/-* as indicated by the well-defined ONL (Figure 3C). We therefore examined the structure of the OS in *ret-Msi1-/-:Msi2-/-* mice at PN16 by immunofluorescence microscopy using three different OS markers, anti-Peripherin-2 (PRPH2: OS marker), anti-Phosphodiesterase-6β (PDE6β: rod OS marker), and peanut agglutinin (PNA: cone OS marker). After staining retinal cross sections from *ret-Msi1-/-:Msi2-/-* mice with PRPH2 and PNA, we observed a severe shortening of the photoreceptor outer segment (Figure 5A). This result was not limited to PRPH2, as staining with the rod OS marker PDE6β demonstrated the same phenotype (Figure 5B). The outer segment of cone photoreceptors also appears to be severely shortened as shown by the abnormal PNA staining (Figure 5A-B). This

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defect is likely independent of degeneration since a similar change was observed at PN10 before significant degeneration (Supplementary Figure 6). Lastly, no mislocalization of PDE6β or PRPH2 is found in the ONL or inner segment of ret-Msi1-/-:Msi2-/- mice suggesting that while the Musashi proteins are required for outer segment formation they are not regulating trafficking or localization of OS-resident proteins (Figure 5B). The Musashi proteins are crucial for photoreceptor outer segment and axoneme development Using transmission electron microscopy, we imaged ultrathin retinal sections from ret-Msi1-/-: Msi2-/- mice at PN10 when the OS begins to elaborate (Figure 6). When examining the OS/IS boundary in ret-Msi1-/-: Msi2-/- mice by electron microscopy, we observed very little, if any, conspicuous OS (Figure 6A). Instead, the IS of the ret-Msi1-/-:Msi2-/- mice appears to come in direct contact with the RPE (Figure 6A-B). At higher magnification, the photoreceptors of ret-Msi1-/-: Msi2-/- mice displayed either no OS or aberrant and undersized OS (Figure 6B left, middle, and right panels). The basal body and connecting cilium (CC) appear to be normal in structure and size (Figure 6B, middle panel). To further examine the structure of the connecting cilium and the axoneme, we stained retinal cross sections from ret-Msi1-/-: Msi2-/- mice at PN10 using antibodies directed against the connecting cilium (glutamylated and acetylated α -tubulin) and axoneme (MAK) markers^{40,43–45}. Probing with glutamylated and acetylated α-tubulin antibodies showed that there were no changes in the length of the CC (Figure 7A, C-D). Contrarily, staining with the anti-MAK antibody showed a substantial reduction in the length of the axoneme accompanied with punctate staining suggesting a severe structural defect of the axoneme (Figure 7A-B). The Musashi proteins promote splicing of photoreceptor specific exons

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Our previous studies suggested that the Musashi proteins are regulating alternative splicing of their target pre-mRNAs in vertebrate photoreceptors⁸. To test if the Musashi proteins are responsible for the inclusion of photoreceptor specific exon, we analyzed the splicing in ret-Msi1-/-: Msi2-/- mice of pre-mRNAs from cilia-and OS-related genes that we previously showed to express photoreceptor specific isoforms (Figure 8). We witnessed a drastic reduction in alternative exon inclusion in ret-Msi1-/-:Msi2-/- mice for all tested transcripts (Figure 8B). We analyzed isoform expression at the protein level for Ttc8 using two different antibodies, a panantibody that recognizes all Ttc8 protein isoforms (Pan-Ttc8) and the other that recognizes the photoreceptor-specific isoform of Ttc8 by binding the epitope encoded by Exon 2A (the photoreceptor-specific exon of *Ttc8*) (Figure 8A, bottom panel). After probing retinal lysates from the ret-Msi1-/-: Msi2-/- mice with the pan-Ttc8 antibody, we observed faster migration of the Ttc8 protein compared to the littermate control suggesting that the Exon 2A was not included (Figure 8C). Concordantly, when probing for the photoreceptor-specific isoform of Ttc8 using the Ttc8 Exon 2A antibody, we saw an absence of this isoform in ret-Msi1-/-:Msi2-/- mice (Figure 8C). Taken together, these results demonstrate that the Musashi proteins are required for the inclusion of photoreceptor specific alternative exons. Alternative exons included in photoreceptor mRNAs are not required for photoreceptor cell development To examine the biological significance of the photoreceptor-specific splicing program that is regulated by the Musashi proteins, we used CRISPR-Cas9 to delete the photoreceptorspecific exons from the Ttc8, Cc2d2a, Cep290, Cacna2d4, and Slc17a7 genes in C57BL6/J mice. These exons were chosen because: (i) they are in genes critical for vision; (ii) the exons are used specifically in photoreceptors where nearly all of the transcripts include the exon; (iii) the exons

with exception of the Slc17a7 exon are conserved in vertebrates. After validating the exon knockout mice (Figure 9A), we collected ERG traces from the exon knockout mice to determine if these exons were necessary for the function of rod and cone photoreceptor cells. Figure 9B shows ERG traces from the *Ttc8* exon knockout, *Cep290* exon knockout, *Cc2d2a* exon knockout, *Cacna2d4* exon knockout, and *Slc17a7* exon knockout compared to wildtype (far left). No significant changes in retinal function were observed in any of exon knockout mice compared to control by five months of age (Figure 9B) suggesting that the inclusion of these photoreceptor-specific exon inclusion is not critical for rod and cone function.

DISCUSSION

MSI1 and MSI2 are required for photoreceptor morphogenesis but not specification

Our data shows the requirement for MSI1 and MSI2 in photoreceptor cells. Double knockout of *Msi1* and *Msi2* in retinal progenitors results in complete loss of vision. Two lines of evidence demonstrate that this loss of vision is due to a defect in photoreceptor morphogenesis, rather than developmental defects due to impairment of the retinal progenitor cells. First, the specification of retinal progenitors to photoreceptor cells was not affected by loss of Musashi. The retina of the knockout mice had clearly defined outer nuclear layer. The rod photoreceptor nuclei retained their characteristic morphology, and the photoreceptor cells expressed cell type specific transcripts such as peripherin and PDE6β. Importantly, knockout of *Msi1* and *Msi2* in rod photoreceptors driven by *Nrl-Cre* caused loss of scotopic photoresponse. Thus, the vision phenotype is not due to impairment of the early stages of retinal development and is caused by a defect specific to photoreceptor cells.

Morphological examination by electron microscopy and immunofluorescence showed that the outer segment of the photoreceptors lacking Musashi is either missing or is stunted and disorganized. The absence of outer segment is accompanied by a shortened axoneme. In contrast, the connecting cilium has normal length and did not have obvious defects. Trafficking of PDE6 and peripherin through the connecting cilium also appears to be normal and the two proteins localize to the stunted outer segment wherever one is present. Taken together our findings demonstrates a requirement for Musashi in the morphogenesis and function of the photoreceptor outer segment that appears not to affect transport along the connecting cilium.

Normal photoreceptor function in photoreceptor-specific exon knockout mice

RT-PCR analysis of alternative splicing in the retina of *Msi1* and *Msi2* knockout mice showed that inclusion of photoreceptor specific exons in the mature transcripts is dependent on the Musashi proteins. Even though MSI1 and MSI2 regulate the splicing of the Ttc8, Cc2d2a, Cep290, Cacna2d4, and Slc17a7 pre-mRNAs, knockouts of the photoreceptor specific-exons in these genes revealed that these exons are not crucial for photoreceptor function. The retina of the exon knockout animals developed normally, and no adverse phenotype was observed up to 5 months after birth. Thus, our data does not support a mechanism by which alternative splicing mediates the phenotype of the Musashi knockouts in photoreceptor cells. This is a surprising result considering that four out of the exons are conserved and the genes that host them are critical for vision. Based on the currently available data we cannot completely rule out role for splicing in shaping the phenotype of the Musashi knockouts. Nevertheless, our results point that other mechanism need to be explored, particularly in the light of the documented role for Musashi in control mRNA translation.

Cell proliferation and survival in the Musashi knockout retina

Morphological examination showed reduced cell number in the neuroblastoid layer at postnatal day 5 in the knockout animals. In the mature retina the outer nuclear layer did not reach the size of the corresponding layers in the wild type animals and progressively degenerated with age. The MSI1 protein was previously reported to be required for photoreceptor survival, but no loss of inner neurons was reported, likely due to the presence of the paralogous MSI2⁴⁶. The reduction of the inner nuclear layer that we observe demonstrate that the Musashi proteins are required for the survival of inner retinal neurons in addition to photoreceptors. Interestingly, the reduction of the neuroblastic layer in the Musashi knockout retina at postnatal day 5 was accompanied by increase in the number of proliferating cells that stained positive for PHH3 and

Ki67. Caspase 3 staining did not show differences in the number of apoptotic cells between the wild type and the knockout retina at that stage. These apparently contradictory observations can be explained with the role of the Musashi proteins in supporting stem cell renewal and proliferation through activation of the Notch pathway^{16,47–50}. We propose that loss of Musashi in the developing retina reduces the numbers and proliferation rates of the neuronal precursors leading to delayed development of the neuroblastoid layer.

Functional redundancy within the Musashi protein family

In vertebrates, the Musashi protein family consists of two paralogues, MSI1 and MSI2, which have high degree of sequence identity, and have arisen from a gene duplication event^{51,52}. The RNA binding domains of MSI1 and MSI2 have approximately 90% sequence identity and recognize the same UAG sequence motif *in vitro* and *in vivo*^{53–56}. The high degree of similarity suggest that the two proteins are likely to be functionally redundant when co-expressed in the same cells. Indeed, we observed only minor reductions in visual function after the loss of either MSI1 or MSI2 alone whereas the combined loss of MSI1 and MSI2 resulted in a complete loss of visual function (Figure 2). Similarly, inclusion of photoreceptor specific exons is promoted by both proteins, and the double knockout produces stronger effect on splicing than the knockouts of either *Msi1* or *Msi2*. The functional redundancy in photoreceptor cells that we observe is in agreement with previous reports of redundancy between MSI1 and MSI2 in other cell types^{12,13}.

The redundancy between the two Musashi proteins in the retina appear to be partial. Loss of MSI1 produce more severe phenotype than loss of MSI2. Specifically, we observe an early visual defect in the *Msi1* knockout mice, which is absent in the *Msi2* knockouts. Loss of MSI1

also produced a stronger effect on splicing compared to MSI2. This partial redundancy may reflect intrinsic functional differences between the two proteins, or simply difference in their expression levels in photoreceptor cells. Interestingly, we noticed moderate but reproducible increased in the MSI2 protein levels after knocking out *Msi1* (Supplementary Figure 2). Such mutual regulation can contribute to the redundancy between the two proteins and be a part of a homeostatic mechanism that maintains the overall Musashi protein levels.

Our work highlights roles for MSI1 and MSI2 in retinal development, retinal cell survival and photoreceptor morphogenesis. An interesting aspect of the function of the Musashi proteins in retina our their apparently mutually exclusive roles at different stages of development. At early stages of development MSI1 and MSI2 support the renewal and proliferation of retinal precursor cells. At late stages of retinal development and in the adult retina MSI1 and MSI2 are required for morphogenesis of the differentiated photoreceptor cells and survival of mature neurons. These roles are likely executed through translational control of Musashi targets, as knockouts of alternative exons regulated by the Musashi proteins did not recapitulate any of the aspects of the phenotype of the Musashi knockout mice. Future studies will be aimed at determining the mechanism(s) by which the absence of Musashi causes a complete loss of vision in mice.

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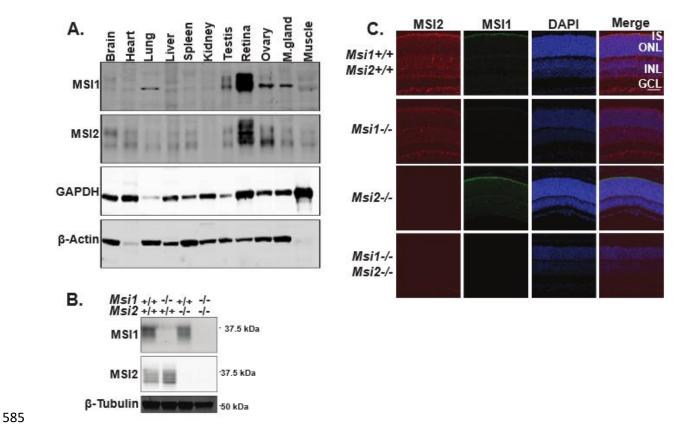
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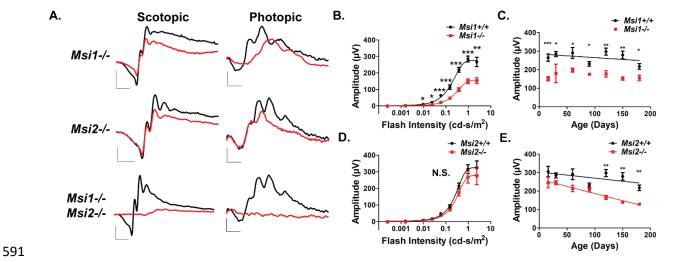
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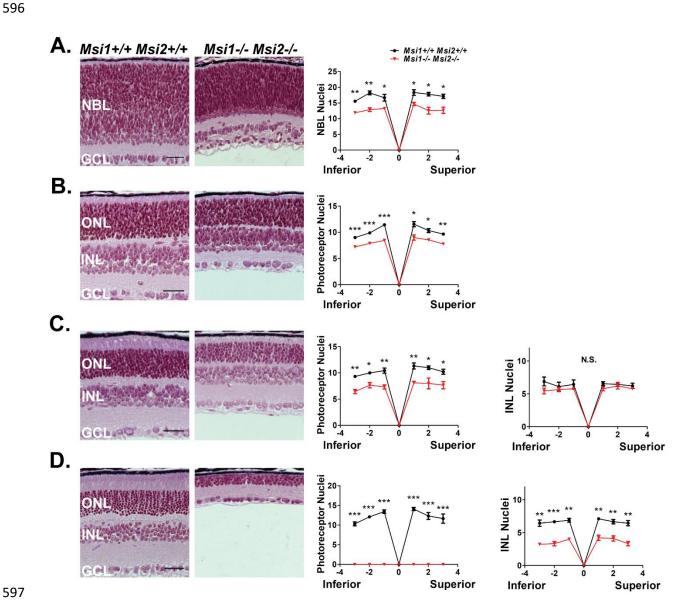
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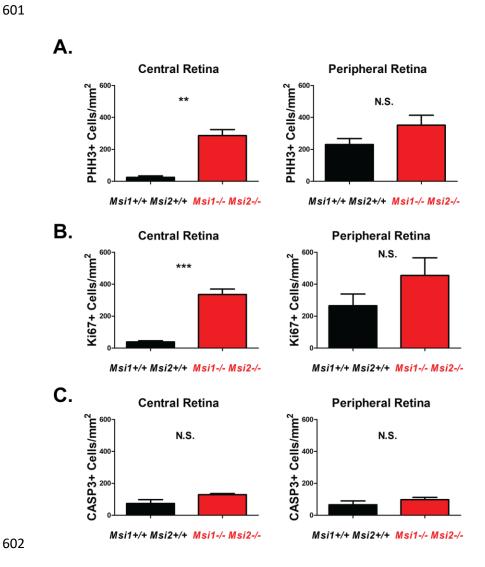
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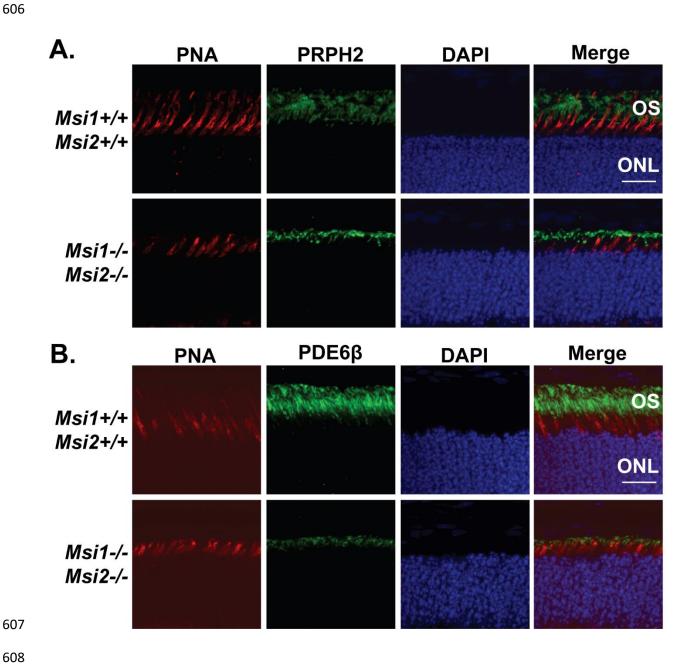


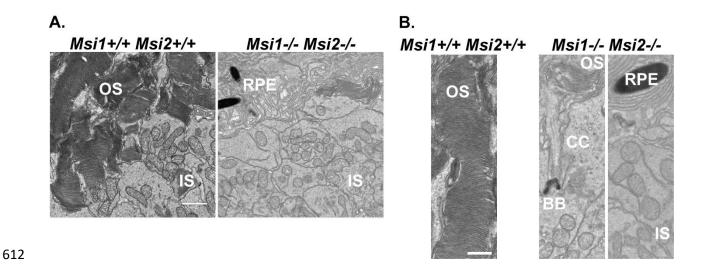


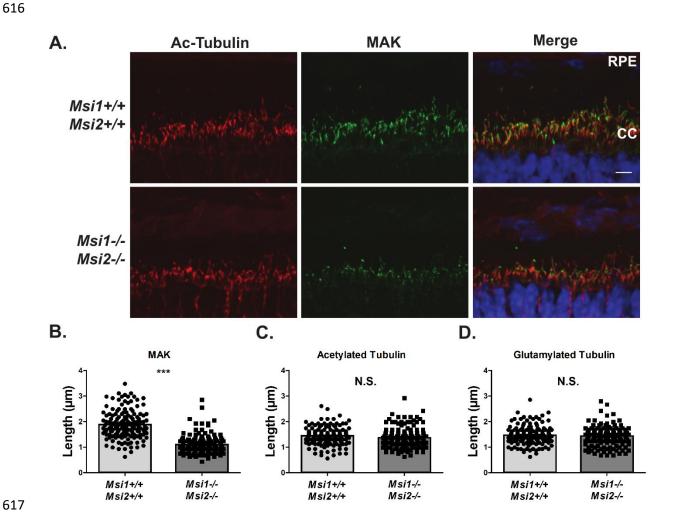


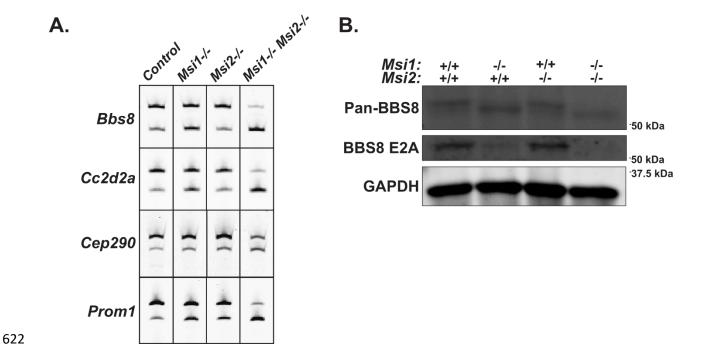


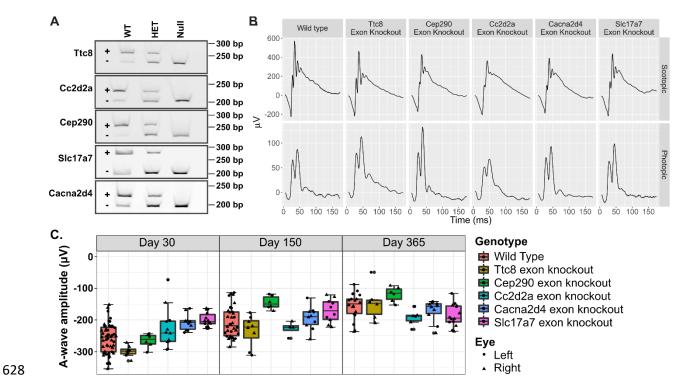
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FIGURE LEGENDS Figure 1: Conditional Musashi knockout mouse models. A. Immunoblot of indicated tissues from adult wildtype mice probed with MSI1 and MSI2 antibodies. GAPDH and β -Actin serve as a loading control. **B.** Western blot analyses of Musashi in retinal lysates from ret-Msi1-/-, ret-Msi2-/-, and ret-*Msi1-/-: Msi2-/-* mice at PN10. β-tubulin levels provide a loading control. C. Retinal sections from ret-Msi1-/-, ret-Msi2-/-, and ret-Msi1-/-: Msi2-/- mice at PN10 probed with MSI1 (Green) and MSI2 (Red) antibodies along with a DAPI nuclear counterstain (Blue). (IS: inner segment, ONL: outer nuclear layer, INL: inner nuclear layer, and GCL: ganglion cell layer). Scale bar = $50\mu m$. Figure 2: The Musashi proteins are crucial for normal visual response. A. Representative scotopic and photopic electroretinograms (ERGs) from the ret-Msi1-/-, ret-Msi2-/-, and ret-Msi1-/-: Msi2-/- mice at PN16. Scotopic ERGs were obtained after overnight dark adaptation using 0.151 cd-s/m² flashes while photopic ERGs were obtained with 7.6 cds/m² flashes under light-adapted conditions using a rod-saturating white background light (Scotopic scale bar: x-axis = 20ms, y-axis = $200\mu V$; Photopic scale bar: x-axis = 20ms, y-axis = $20\mu V$). **B.** Intensity-response plot of the scotopic "a"-wave response from ret-Msi1-/- mice (*=P-value < 0.05; **=P-value < 0.01; ***=P-value < 0.001).

C. Plot of the rod photoreceptor "a"-wave response from ret-Msi1-/- mice against the age of the

mouse during which the ERG was recorded.

D. Intensity-repsonse curve of the scotopic "a"-wave response from ret-Msi2-/- mice (*=P-value 653 < 0.05; **=P-value < 0.01; ***=P-value < 0.001).654 655 **E.** Plot of the rod photoreceptor "a"-wave response from ret-Msi2-/- mice plotted against the age of the mouse during which the ERG was recorded. All data is shown as the mean \pm the SEM, and 656 statistical analyses were carried out using the homoscedastic unpaired student's t-test 657 658 *=P<0.05. 659 660 Figure 3: Retinal cell death occurs in the absence of the Musashi proteins Left: Brightfield microscopic images of H&E stained retinal cross sections from the ret-Msi1-/-: 661 Msi2-/- mice at PN5 (A), PN10 (B), PN16 (C), and PN180 (D). 662 Right: Spider plot of the layer thickness at six regions from the inferior to superior retina in the 663 ret-Msi1-/-: Msi2-/- mice at PN5 (NBL: neuroblast layer, ONL: outer nuclear layer, INL: inner 664 nuclear layer, and GCL: ganglion cell layer) 665 666 All data is shown as the mean \pm the SEM, and statistical analyses were carried out using the homoscedastic unpaired student's t-test (*=P-value < 0.05; **=P-value < 0.01; ***=P-value < 667 0.001). 668 669 Figure 4: Increased proliferation of progenitors in the absence of MSI1 and MSI2 670 671 Quantitation of PHH3+ cells (A), Ki67+ cells (B) and CASP3+ cells (C) in the central (left) and 672 peripheral (right) regions of the retina in PN5 ret-Msi1-/-: Msi2-/- mice and littermate control 673 mice. Figure 5: The Musashi proteins are required for proper OS development 674 675 **A.** Immunofluorescence microscopy images of retinal cross sections from the ret-Msi1-/-: Msi2-

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/- mice at PN16 stained with anti-peripherin-2 antibody (PRPH2: OS marker - Green) and peanut agglutinin (PNA: cone OS marker - Red) along with a DAPI nuclear counterstain (Blue). **B.** Immunofluorescence microscopy images of retinal cross sections from the ret-Msi1-/-: Msi2-/- mice at PN16 stained with anti-phosphodiesterase-6\beta antibody (PDE6\beta: rod OS marker -Green) and peanut agglutinin (PNA: cone OS marker - Red) along with a DAPI counterstain (Blue). (OS: outer segment and ONL: outer nuclear layer). Scale bar = $20\mu m$. Figure 6: Abnormal development of OS in the absence of MSI1 and MSI2 **A.** Low magnification transmission electron microscopy images of ultrathin retinal sections from ret-Msi1-/-: Msi2-/- mice at PN10 visualizing the boundary between the OS and IS showing the lack of typical outer segments in the absence of the Musashi proteins (OS: outer segment, IS: inner segment, and RPE: retinal pigment epithelium). Scale bar = $2\mu m$. **B.** High magnification transmission electron microscopy images of ultrathin retinal sections from ret-Msi1-/-: Msi2-/- mice at PN10 visualizing the boundary between the OS and IS showing that the OS either does not form (far right) or is dysmorphic (middle) in the absence of the Musashi proteins (OS: outer segment, CC: connecting cilium, BB: basal body, RPE: retinal pigment epithelium, and IS: inner segment). Scale bar = 1μ m. Figure 7: The Musashi proteins are crucial for photoreceptor axoneme development **A.** Immunofluorescence microscopy images of retinal cross sections from the ret-Msi1-/-: Msi2-/- mice at PN10 stained with acetylated-α-tubulin antibody (Ac-Tubulin: Red) and male germ cell-associated kinase antibody (MAK: Green) along with DAPI counterstain (Blue) (RPE: retinal pigment epithelium, CC: connecting cilium, and ONL: outer nuclear layer).

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Scatter bar plot showing the distribution of length measurements for the photoreceptor axoneme by MAK staining (B) and connecting cilium by Ac-tubulin staining (C) and glutamylated tubulin staining (**D**). Retinal sections were obtained from PN10 musashi knockouts and littermate controls. Figure 8: The Musashi proteins regulate alternative splicing of their target transcripts **A.** Reverse transcriptase PCR splicing assay using total RNA purified from retinal lysates of ret-Msi1-/-, ret-Msi2-/-, and ret-Msi1-/-: Msi2-/- mice. Ttc8, Cc2d2a, Cep290, and Prom1 are four cilia- and OS-related transcripts shown to have reduced photoreceptor-specific exon inclusion in the absence of MSI1 and MSI2. **B.** Immunoblot of retinal lysates from ret-Msi1-/-, ret-Msi2-/-, and ret-Msi1-/-: Msi2-/- mice. After probing with the pan-Ttc8 antibody (top), a change in the migration of the Ttc8 protein is observed in the absence of MSI1 and MSI2 suggesting that the peptide encoded by Exon 2A was not included. When probing with the Ttc8 E2A antibody (middle), photoreceptor-specific isoform of Ttc8 was not observed in the absence of MSI1 and MSI2. Figure 9: Exon knockout mice have normal photoreceptor function **A.** Analysis of alternative splicing of the *Ttc8*, *Cc2d2a*, *Cep290*, *Slc17a7*, and *Cacna2d4* genes in wild type (WT), heterozygous (Het) and homozygous (Null) exon knockouts by RT-PCR. Isoforms including and skipping the alternative exon are indicated by "+" and "-", respectively. **B.** Representative scotopic and photopic electroretinograms (ERGs) from wildtype, *Ttc8* exon knockout, Cep290 exon knockout, Cc2d2a exon knockout, Cacna2d4 exon knockout, and Slc17a7 exon knockout mouse models at postnatal day 150. Scotopic ERGs were obtained after dark adaptation while photopic ERGs were obtained under light-adapted conditions using a rod-

saturating white background light. No changes in photoreceptor function are observed compared to control.

C. Box blot of a-wave intensities normalized to the median of the wild type a-wave at postnatal days 30, 150 and 356. No significant changes in photoreceptor function of the knockouts of photoreceptor specific exons in the *Ttc8*, *Cep290*, *Cc2d2a*, *Cacna2d4* and *Slc17a7* genes are observed compared to the wild type control.