

TRPM1 is required for the depolarizing light response in retinal ON-bipolar cells

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The ON pathway of the visual system, which detects increases in light intensity, is established at the first retinal synapse between photoreceptors and ON-bipolar cells. Photoreceptors hyperpolarize in response to light and reduce the rate of glutamate release, which in turn causes the depolarization of ON-bipolar cells. This ON-bipolar cell response is mediated by the metabotropic glutamate receptor, mGluR6, which controls the activity of a depolarizing current. Despite intensive research over the past two decades, the molecular identity of the channel that generates this depolarizing current has remained elusive. Here, we present evidence indicating that TRPM1 is necessary for the depolarizing light response of ON-bipolar cells, and further that TRPM1 is a component of the channel that generates this light response. Gene expression profiling revealed that TRPM1 is highly enriched in ON-bipolar cells. In situ hybridization experiments confirmed that TRPM1 mRNA is found in cells of the retinal inner nuclear layer, and immunofluorescent confocal microscopy showed that TRPM1 is localized in the dendrites of ON-bipolar cells in both mouse and macaque retina. The electroretinogram (ERG) of TRPM1-deficient (TRPM1^{-/-}) mice had a normal a-wave, but no b-wave, indicating a loss of bipolar cell response. Finally, whole-cell patch-clamp recording from ON-bipolar cells in mouse retinal slices demonstrated that genetic deletion of TRPM1 abolished chemically simulated light responses from rod bipolar cells and dramatically altered the responses of cone ON-bipolar cells. Identification of TRPM1 as a mGluR6-coupled cation channel reveals a key step in vision, expands the role of the TRP channel family in sensory perception, and presents insights into the evolution of vertebrate vision.

retinal neurobiology | transient receptor potential channel | visual ON-pathway

At the first retinal synapse, the postsynaptic bipolar cells respond with opposite polarities to light-induced changes in synaptic glutamate released from photoreceptor terminals (1). ON-bipolar cells depolarize and OFF-bipolar cells hyperpolarize in response to increases in light intensity. The polarity of the bipolar cell response is determined by the expression of distinct glutamate receptors. The OFF response is mediated by ionotropic glutamate receptors (iGluRs). In contrast, ON-bipolar cells express a distinctive metabotropic glutamate receptor (mGluR), mGluR6, which is localized in the tips of ON bipolar cell dendrites (2). In darkness, activation of this receptor by synaptic glutamate causes the closure of a nonselective cation channel. Deactivation of mGluR6, subsequent to the light-induced decrease in synaptic glutamate, causes these channels to open, thereby generating the depolarizing light response of ON-bipolar cells (3–5). Despite intensive research over the past two decades, the molecular identity of the channel that generates this depolarizing current has remained elusive.

Transient receptor potential (TRP) channels were first discovered in *Drosophila*, where they generate the photoreceptor light response (6–8). Since that time, >30 mammalian TRP channel homologs have been discovered, which serve a wide variety of sensory functions, ranging from thermosensation and

nociception, to gustation and mechanosensation (9). The founding member of the TRPM family, TRPM1, was discovered by differential display as a potential suppressor of tumor metastasis [and originally named melastatin because it is down-regulated in metastatic melanoma (10)]. TRPM1 is the product of a complex gene, spanning 58 kb and 27 exons, and exists as several isoforms, produced by mRNA splice variants (11–13). Little is known about the physiological role of TRPM1, but recently it has been reported to form constitutively active cation channels in melanocytes, where it has been proposed to function in melanin trafficking (13). The basic properties of the channel, including ion selectivity and current-voltage relationship, are similar to those reported for the mGluR6-coupled ion channel in retinal ON-bipolar cells. In the Appaloosa horse, a reduction in TRPM1 has been correlated with the loss of the ERG b-wave (14), and a recent study by Shen et al. (15) showed that the ERG b-wave is lacking in the TRPM1^{-/-} mouse implicating this channel in photoreceptor to ON-bipolar cell synaptic transmission. Here, we present molecular, immunohistochemical, and electrophysiological evidence that TRPM1 is the mGluR6-coupled cation channel in rod-bipolar cells, and provide evidence that TRPM1 is essential for the normal response of cone ON-bipolar cells.

Results

TRPM1 Is Expressed in Retinal Bipolar Cells. Gene expression profiling studies [*SI Text* and *Fig. S1* (16, 32)], demonstrated that TRPM1 is enriched in ON-bipolar cells, suggesting TRPM1 as a potential candidate for the long sought after channel generating the light response in these cells. We verified that the TRPM1 mRNA was expressed in ON-bipolar cells by in situ hybridization (*Fig. 1A Left*). By using an anti-sense probe directed against the mouse TRPM1 mRNA (bp 2,383–3,514, GenBank accession NML001039104), a strong signal was obtained in many cells of the distal inner nuclear layer (INL), where the majority of ON-bipolar cells are located (*Fig. 1A*, arrows). Unlabeled cells in this region of the INL likely correspond to horizontal cells (*Fig. 1A*, white arrowhead is a putative horizontal cell). A control hybridization with a sense probe directed against the same region of the channel cDNA showed no signal above background in any part of the retina (*Fig. 1A Right*).

Immunofluorescence confocal microscopy demonstrated that the TRPM1 channel is localized to ON-bipolar cells. As shown in *Fig. 1B*, immunolabeling of vertical sections of mouse retina with an anti-TRPM1 antibody revealed punctate staining in the

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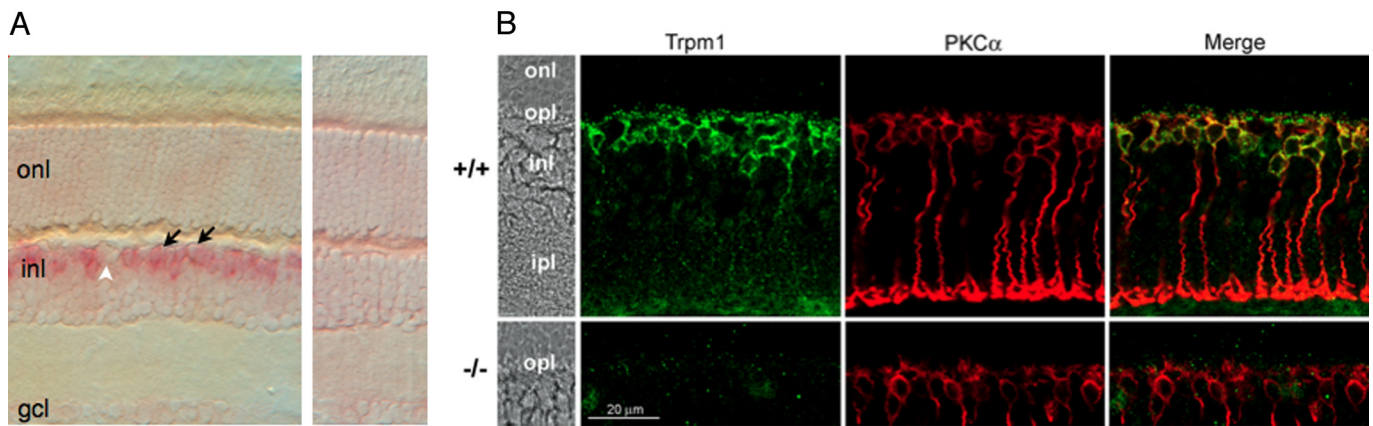


Fig. 1. TRPM1 is expressed by ON-bipolar cells in the mouse retina. (A) In situ hybridization of vertical sections of mouse retina with antisense (Left) and sense control (Right) TRPM1 probes. A hybridization signal is detected in many cell somata in the INL, where bipolar cell nuclei and somata are located (black arrows). Occasional unlabeled cell somata are likely horizontal cells (white arrowhead). (B) Vertical sections from a wild-type (Top) and TRPM1^{-/-} (Bottom) retina were immunofluorescently labeled by antibodies directed against TRPM1 (green) and PKC α (red). Areas of colocalization appear yellow in the merged images (Right). onl, Outer nuclear layer; opl, outer plexiform layer; inl, inner nuclear layer; ipl, inner plexiform layer.

outer plexiform layer (OPL), as well as membrane-associated and possible intracellular staining of bipolar cell bodies. The specificity of the TRPM1 antibody was demonstrated by the absence of staining in the TRPM1^{-/-} mouse (Fig. 1B, lower panels), as well as by labeling of HEK cells transiently transfected with TRPM1 cDNA (Fig. S2). Most of the TRPM1 labeling in the mouse retina was coincident with that for PKC α , indicating that TRPM1 is expressed in rod bipolar cells (Fig. 1B). The TRPM1-positive cells extended dendrites into the OPL, the tips of which could be labeled for mGluR6 (Fig. S3A–C) and are apposed to labeled synaptic ribbons (Fig. S3D–F). Immunofluorescent labeling indicated that mGluR6 is still present in the ON-bipolar cell dendrites in the TRPM1^{-/-} retina (Fig. S4A), and that tips of TRPM1^{-/-} rod bipolar cell dendrites are apposed to photoreceptor synaptic ribbons (Fig. S4B), suggesting that the morphology of the OPL is not grossly disrupted in the TRPM1^{-/-} retina.

Because the TRPM1 antibody was raised against a human polypeptide, TRPM1 immunostaining was even more robust in the macaque retina (Fig. 2). In vertical sections of macaque retina, TRPM1 staining appeared as puncta in the OPL and was associated with cell plasma membranes in the distal INL. In many instances, this labeling was clearly associated with PKC α , indicating that these were rod bipolar cells (Fig. 2B). Similar to mouse, however, not all TRPM1-positive cells were immunoreactive for PKC α (arrowheads, Fig. 2B). This finding is more apparent in a horizontal optical section through the INL of a macaque retinal whole mount shown in Fig. 2C and D (cells marked with asterisks). These cells are most likely cone ON-bipolar cells since all TRPM1 positive cells can be colabeled with Gao (Fig. S5). If this is the case, their dendritic processes should extend to cone photoreceptor pedicles in the OPL. Indeed, immunolabeling for TRPM1 is clearly associated with cone pedicles, as well as with rod terminals, shown by double labeling with the anti-TRPM1 antiserum and the cone marker, Alexafluor peanut agglutinin (PNA) (Fig. 2E–G). Thus, TRPM1 is localized to the dendrites of both rod and cone ON-bipolar cells, where it is optimally positioned to respond rapidly to light-induced changes in the synaptic glutamate levels.

Vision Is Impaired in the TRPM1^{-/-} Mouse. The expression of TRPM1 in retinal bipolar cells implies that the TRPM1 channel is important for vision. To test this hypothesis, we measured spatial frequency and contrast sensitivity thresholds of the optokinetic response (OKR) in TRPM1^{-/-} mice (17). The

spatial frequency threshold of TRPM1^{-/-} mice was reduced 10% compared to wild-type (0.359 ± 0.004 cycles/degree for TRPM1^{-/-}, and 0.400 ± 0.010 cycles/degree for wild-type, $P < 0.05$), and contrast sensitivity was reduced 3-fold (4.61 ± 0.23 for TRPM1^{-/-}, and 14.99 ± 3.85 for wild-type, $P < 0.001$, measured at a spatial frequency of 0.150 cycles/degree). The OKR measurements indicate that the TRPM1^{-/-} mice are visually im-

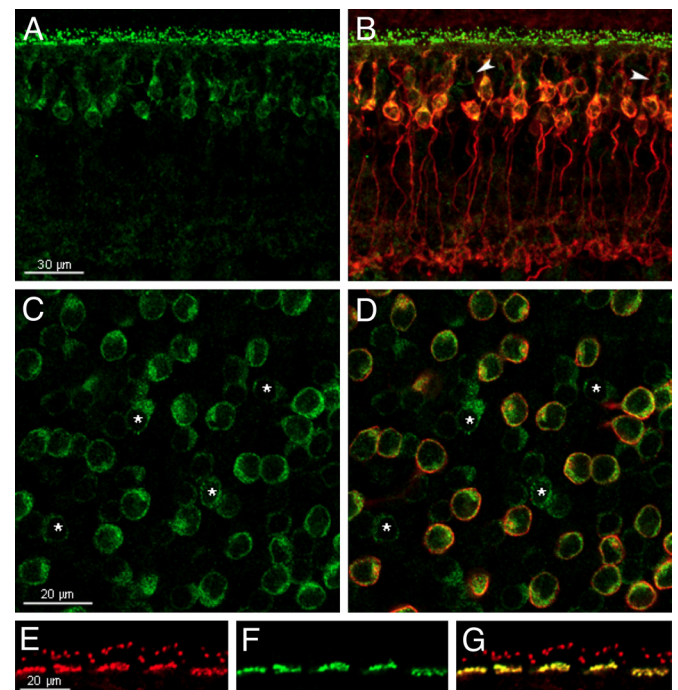


Fig. 2. TRPM1 is associated with both rod and cone ON-bipolar cells in the primate retina. (A and B) Vertical section of macaque retina double labeled for TRPM1 (green, A and B) and PKC α (red, B). Arrowheads indicate putative cone ON-bipolar cells (TRPM1 positive, PKC α negative cells). (C and D) Horizontal optical section in the plane of the inner nuclear layer from a whole mount macaque retina double labeled for TRPM1 (green, C and D) and PKC α (red, D). Asterisks indicate putative cone ON-bipolar cells. (E–G) Vertical section through the outer plexiform layer of the macaque retina double labeled with an antibody against TRPM1 (red, E and G) and the cone marker, peanut agglutinin-Alexafluor (green, F and G).

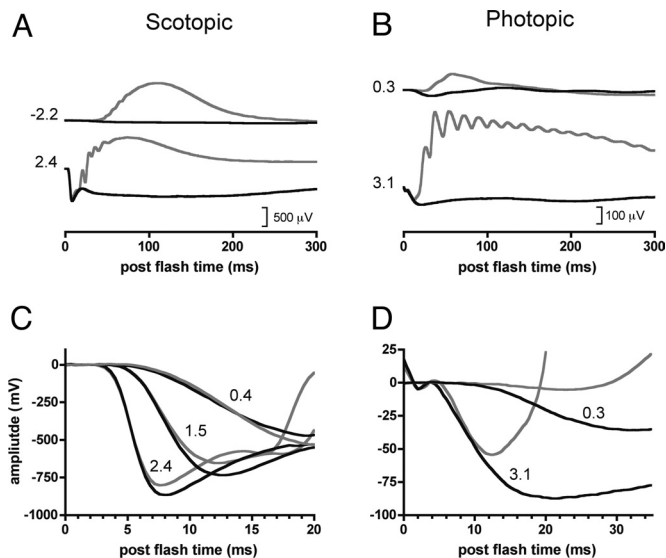


Fig. 3. ERG b-waves and oscillatory potentials are absent in TRPM1-deficient mice. Rod isolated ERGs (A and C) and photopic cone-mediated ERGs (B and D) from combined wild-type and TRPM1^{+/+} mice ($n = 4$; gray lines) compared with TRPM1^{-/-} mice ($n = 5$; solid black lines). Numbers indicate scotopic (A and C) and photopic (B and D) flash intensities (log candela-seconds/meter²). Note that scales vary between scotopic and photopic graphs.

paired, although not profoundly so, similar to the complete congenital stationary night blindness phenotype.

Electroretinogram b-Wave Is Absent in Mice Lacking TRPM1. To investigate the physiological role of TRPM1 in the retina, we recorded the electroretinogram (ERG) from control and TRPM1^{-/-} mice. The ERG responses from TRPM1^{+/+} and wild-type mice were indistinguishable and have been combined to form the control group for analysis. Figure 3 shows representative ERGs from control (gray traces) and TRPM1^{-/-} mice (black traces). The b-wave and oscillatory potentials, both of which are generated downstream from the photoreceptors, were absent from both rod and cone-mediated (both scotopic and photopic) ERGs of the TRPM1^{-/-} mice (Fig. 3A and B). Figure 3C shows an expanded view of the beginning of the rod-isolated ERGs recorded for several flash intensities. The superimposed traces show that the negative-going rod-isolated ERG a-waves were essentially identical between TRPM1-deficient and control

mice. A phototransduction model was ensemble fit to the rising phases of the rod-isolated a-waves. The derived rod phototransduction parameters (mean \pm SE.) were not significantly different between control { $R_{maxP3} = -806 \pm 35 \mu V$; $S = 1,158 \pm 166 [(cd-s/m^2)^{-1} s^{-2}]$; $td = 3.9 \pm 0.05 ms$ } and TRPM1^{-/-} mice { $R_{maxP3} = -877 \pm 30 \mu V$; $S = 1,221 \pm 120 [(cd-s/m^2)^{-1} s^{-2}]$; $td = 4.1 \pm 0.1 ms$ }. Figure 3D shows an expanded view of the cone-isolated photopic ERGs. Rodents have little or no cone mediated photopic ERG a-wave. In our study, the initial negative going photopic response from control mice only tracked the ERG from the TRPM1^{-/-} mice for very bright flash intensities ($>3 \log ph \text{ cd-s/m}^2$) (Fig. 3D). The absence of the ERG b-wave in the TRPM1^{-/-} mice indicates a block in signal transmission between photoreceptors and ON-bipolar cells in these animals. The ERG, however, does not indicate whether this block resides postsynaptically, in the bipolar cell dendrites, or presynaptically, in the photoreceptor terminals. The in situ hybridization and immunofluorescence results (Figs. 1 and 2) support a postsynaptic locus for the block, but to rule out any contribution from defective glutamate release by photoreceptors in the TRPM1^{-/-} retina, we recorded directly from ON-bipolar cells using a technique that bypasses the photoreceptors entirely.

Rod Bipolar Cell Response Is Absent in TRPM1^{-/-} Mice. To examine the role of TRPM1 in generation of the depolarizing light response in ON-bipolar cells, we used the whole-cell patch clamp technique to record chemically simulated light responses from ON-bipolar cells in mouse retinal slices. For these experiments, the slice was bathed at all times in the mGluR6 agonist, L-2-aminophosphonobutyrate (L-AP4) to simulate darkness; pressure application of the mGluR6 antagonist, α -cyclopropyl-4-phosphonophenylglycine (CPPG), to the OPL was used to simulate a step of light. During recording, cells were filled with the dye, Alexa-488 hydrazide, which permitted morphological identification of rod and cone bipolar cells at the termination of the experiment (Fig. 4A and C). As shown in Fig. 4B, application of CPPG to the wild-type mouse retina activated a robust current with a waveform typical of rod bipolar cells (18). These currents had a peak amplitude at approximately 120 ms of $-129 \pm 15 pA$ (SEM; $n = 21$), which decayed rapidly to a plateau. These currents were abolished in the TRPM1^{-/-} retina. In the majority of morphologically identified rod bipolar cells, CPPG application activated no measurable currents ($-2.7 \pm 0.55 pA$; SEM; $n = 25$); a few cells exhibited a slowly activating current of $<5 pA$.

Cone ON-Bipolar Cell Response Is Dramatically Altered in TRPM1^{-/-} Mice. In wild-type retina, cone ON-bipolar cells responded to CPPG with a sustained current of $-75 \pm 8.6 pA$ ($n = 13$) (Fig. 4D).

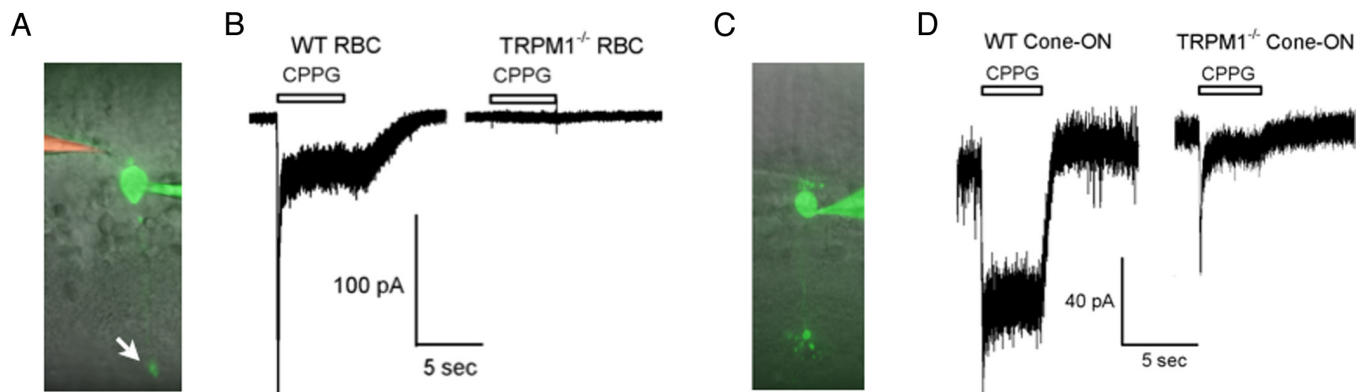


Fig. 4. Chemically simulated ON-bipolar cell light responses from wild-type and TRPM1^{-/-} retina. Patch clamp recording of a rod (A) and a cone (C) ON-bipolar cell in a mouse retinal slice. The patch electrode was filled with internal solution containing Alexa488 hydrazide, and the puffer pipette (visible in A) was filled with Ames media containing 600 μM CPPG and Alexa-594. Rod bipolar cell synaptic terminal is indicated by an arrow in A. Current traces at $-60 mV$ from rod (B) and cone (D) ON-bipolar cells from wild-type and TRPM1^{-/-} retinas; CPPG was applied to the outer plexiform layer as indicated.

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