Targeting K_v channels rescues retinal ganglion cells in vivo directly and by reducing inflammation

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Abbreviations: RGC, retinal ganglion cell; GCL, ganglion cell layer; NFL, nerve fiber layer; IPL, inner plexiform layer; INL, inner nuclear layer; ONL, outer nuclear layer; CNS, central nervous system; BDNF, brain derived neurotrophic factor; CNTF, ciliary neurotrophic factor; GDNF, glial-cell line derived neurotrophic factor; NGF, nerve growth factor; bFGF, basic fibroblast growth factor; IGF, insulin-like growth factor; EGF, epidermal growth factor; NRTN, neurturin; iNOS, inducible nitric oxide synthase; IL-1 β , interleukin-1-beta; IL-1ra, interleukin-1 receptor antagonist; TNFα, tumor necrosis factor-alpha; TGFβ, transforming growth factor-beta; CR3, complement receptor 3; MHC Class II, major histocompatibility complex class 2; GFAP, glial fibrillary acidic protein; HPRT-1, hypoxanthine phosphoribosyltransferase 1; AgTx-2, agitoxin-2; MgTx, margatoxin; α-DTX, alpha-dendrotoxin; K_v, voltage gated potassium channel subunit; K_v1.1, voltage gated potassium channel family 1 member 1; K_v1.2, voltage gated potassium channel family 1 member 2; K_v1.3, voltage gated potassium channel family 1 member 3; Caspase-3, cysteine-aspartic protease-3; Caspase-9, cysteine-aspartic protease-9; Bad, Bcl-2-associated death promoter; Bclx, B-cell lymphoma-extra large

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Retinal ganglion cell (RGC) degenvisual impairment, and results in part from microglia-mediated inflammation. Numerous experimental studies have focused on identifying drug targets to rescue these neurons. We recently showed that K_v1.1 and K_v1.3 channels are expressed in adult rat RGCs and that siRNA-mediated knockdown of either channel reduces RGC death after optic nerve transection. Earlier we found that K_v1.3 channels also contribute to microglial activation and neurotoxicity; raising the possibility that these channels contribute to neurodegeneration through direct roles in RGCs and through inflammatory mechanisms. Here, RGC survival was increased by combined siRNA-mediated knockdown of K_v1.1 and K_v1.3 in RGCs, but survival was much greater when knockdown of either channel was combined with intraocular injection of a K_v1.3 channel blocker (agitoxin-2 or margatoxin). After axotomy, increased expression of several inflammation-related molecules preceded RGC loss and, consistent with a dual mechanism, their expression was differentially affected when channel knockdown in RGCs was combined with K_v1.3 blocker injection. K_v1.3 blockers reduced activation of retinal microglia and their tight apposition along RGC axon fascicles after axotomy, but did not prevent their migration from the inner plexiform to the damaged ganglion cell layer. Expression of several growth factors increased after axotomy; and again, there were differences following blocker

injection compared with RGC-selective channel knockdown. These results provide evidence that $K_v 1.3$ channels play important roles in apoptotic degeneration of adult RGCs through cell-autonomous mechanisms mediated by channels in the neurons, and nonautonomous mechanisms mediated by microglia and inflammation.

Introduction

Information is encoded by retinal ganglion cells (RGCs) before transmission to higher brain centers. RGC death underlies several important visual neuropathies (e.g., glaucoma, diabetic retinopathy) and is thus an important therapeutic target. The optic nerve transection model in adult rats has been used extensively to assess the potential of targeting growth factors and apoptosis-related molecules (reviewed in ref. 1-4). Using this model, we recently showed that depleting either K_v1.1 or K_v1.3 channels from RGCs by injecting channel-specific siRNAs into the cut optic nerve stump greatly reduced RGC degeneration.⁵ While this approach ensures that only the channels in RGCs are affected, it is not clinically feasible. In contrast, when drugs are injected into the vitreous chamber of the eye, they exit by hydraulic and vascular clearance, allowing sustained drug entry into the retina and actions on multiple cell types.⁶

Inflammatory processes associated with activated microglia are important contributors to RGC death.⁷⁻⁹ While microglia are in a relatively 'resting' state in the healthy adult retina, they become 'activated' following damage by mechanical injury, ischemia, infection or in autoimmune disorders. Microglial activation is complex; some functions such as phagocytosis and production of growth factors can contribute to repair after CNS injury, while others (e.g., overproduction of pro-inflammatory and oxidative molecules) can exacerbate damage. Consequently, microglial activation is an important, albeit complicated target for reducing neurodegeneration, including RGC loss in retinal neuropathies.

One potential therapeutic target is $K_v 1.3$; a voltage-gated K⁺ channel that is highly expressed in microglia, where it contributes to superoxide production and their neurotoxic capacity in vitro.10 Ky1.3 channels are also expressed in RGCs, and we recently found that siRNA-mediated depletion of $K_v 1.3$ (or $K_v 1.1$) from RGCs substantially increases their survival after optic nerve transection.5 K₁1.1 knockdown increased the anti-apoptotic gene, Bcl-X, while K, 1.3 knockdown reduced the pro-apoptotic genes, caspase-3, caspase-9 and Bad. The present study was designed to address the possibility that RGC rescue involves a dual mechanism; direct protection of RGCs, and effects on growth factors and inflammation.

Results

Inflammation and the microglial response after optic nerve transection. Figure 1 illustrates the structure of the healthy adult rat retina (Fig. 1A), and the distribution of retinal ganglion cells and microglia before and after optic nerve transection. Microglia were very sparse in the outer retina (e.g., in the outer plexiform layer; Fig. 1B), and were absent from the outer nuclear and inner nuclear layers (not shown). The inner plexiform layer (Fig. 1C) contained a dense array of microglia with the highly ramified morphology typical of resting cells. Immediately below the ganglion cell layer (Fig. 1D), ramified microglia were distributed between the many somata of Fluorogold-labeled retinal ganglion cells (RGCs). In contrast, the healthy nerve fiber layer (Fig. 1E) contained only sparse OX-42 labeled cells with a round or amoeboid morphology. In these images of the healthy retina, numerous RGC cell bodies can be seen, along with several Fluorogold-labeled axon bundles running radially toward the optic disc.

By 14 days after axotomy, the distribution of microglia had dramatically changed (Fig. 1F). In retinas that had been injected with saline at the time of axotomy and four days later (controls), microglia had migrated to the nerve fiber layer (NFL). They were less ramified than in the healthy retina, and most were aligned with the fascicles of degenerating RGC axons. Very few Fluorogoldlabeled RGCs remained (see below), which is consistent with the time-course of neurodegeneration previously observed in this model.¹¹⁻¹³ Phagocytic microglia containing apoptotic Fluorogold labeled RGCs were also clearly visible under these conditions (Fig. 1F, inset).

We next examined effects on microglia of two potent K_v1.3-blocking peptides (agitoxin-2, AgTx-2; margatoxin, MgTx), which we previously found inhibits microglial activation in vitro¹⁰ and RGC degeneration in vivo.5 We note that K_v channel blockers are not as selective as siRNA-mediated channel knockdown; AgTx-2 blocks K_v1.1 and K_v1.3, while MgTx blocks K_v1.2 and K_v1.3.¹⁴ For comparison, we also used α -dendrotoxin (α -DTx), a potent blocker of K_v1.2 channels (and less effective blocker of K_v1.1 and K_v 1.3), which failed to inhibit microglial activation or rescue RGCs in our earlier studies.5,10 Consistent with our recent quantitative study,⁵ the images in Figure 1 show more RGC survival after intraocular injection of either AgTx-2 (Fig. 1G) or MgTx (Fig. 1H). There were several morphological indications of reduced microglial activation; they were more scattered, more ramified and less aligned with the axon bundles. Of note, many microglia were present in the NFL; thus, blocking K_v1.3 channels did not prevent microglial migration into the damaged layer, but reduced their association with damaged axons. α -dendrotoxin (α -DTx) did not affect microglial morphological changes or alignment with RGC axon bundles (compare Fig. 1F and I). As a further control, AgTx-2 injection into the naïve retina did not affect the microglial appearance or distribution (not shown).

Expression of several genes was quantified by real-time RT-PCR to monitor the retinal response to optic nerve transection. Several markers of microglial and

Figure 1 (See opposite page). Inflammation and the microglial response after optic nerve transection. (A) Schematic showing the layers and neural types in the normal adult retina. The nerve fiber layer (NFL) is comprised of axons of retinal ganglion cells (RGCs); the ganglion cell layer (GCL) contains RGC somata; the inner plexiform layer (IPL) includes amacrine cells and processes of RGCs; the inner nuclear layer (INL) contains bipolar cells; the outer plexiform layer (OPL) contains horizontal cells; and the outer nuclear layer (ONL) contains photoreceptors. (B-E) Representative confocal images from flat mounts of healthy retinas (n = 4 for each treatment), taken from the mid-periphery (see inset in Fig. 4E). RGCs and their axons were retrogradelabeled with Fluorogold (false-colored green); microglia were labeled with OX-42 antibody (red). The level at which each image was taken is indicated by corresponding red letter in (A). Microglia were very sparse in the outer plexiform layer (B). Highly ramified microglia were densely distributed in the inner plexiform layer (C), and adjacent to the GCL (D), where numerous RGC cell bodies can be seen. In the nerve fiber layer (E), only a few amoeboid microglia are seen among the RGC axon fascicles. Scale bar, 50 µm; applies to all parts. (F) Microglia in the NFL at 14 days after axotomy in a saline-injected retina. Some microglia (yellow in the overlay) are labeled with Fluorogold, as a result of phagocytosing apoptotic RGCs (green); the higher-magnification inset shows phagocytosed RGCs inside two microglia. Very few RGC cell bodies remain and their axons are no longer visible. Instead, there are large numbers of microglia, which are much less ramified than in the healthy retina and mainly organized in rows, similar to the RGC axon fascicles in the healthy retina (E). (G–I) Microglia in the NFL (14 days after axotomy), following intraocular K, blocker injections (50 μM) at the time of axotomy and 4 days later. The blockers were: agitoxin-2 (G), margatoxin (H) and lpha-dendrotoxin (I). (J) Changes in gene expression at 7 days after optic nerve transection, measured by quantitative real-time RT-PCR (for primer sequences, see Table 1). One group of genes represents responses of microglia, complement receptor-3 (CR3) and MHC class II; astrocytes, glial fibrillary acidic protein (GFAP); and neurons, synaptophysin (SYP). The second includes cFos and the inflammation-related genes, TNFα, IL-1β, iNOS, IL-1 receptor antagonist (IL-1ra). For each gene, the relative expression was normalized to the housekeeping gene, HPRT-1, and values are shown as mean ± SEM (n = 6 retinas each), with significant differences (*p < 0.05; **p < 0.01) determined by ANOVA, followed by Fisher's test.



Figure 1. For figure legend, see p. 338



Figure 2. K_v block and RGC-specific knockdown differentially affect growth factor expression. Retinal expression of gene transcripts was compared by quantitative real-time RT-PCR at seven days after optic nerve transection. Expression of each gene was normalized to the value in untreated retinas after optic nerve transection (axotomized). Data are expressed as mean \pm SEM (6 retinas each); the statistical significance was determined by ANOVA, followed by Fisher's test. (A) Effects of siRNA-mediated depletion of either K_v1.1 or K_v1.3 from RGCs only. Statistical comparisons between healthy retinas (naïve) and axotomized retinas are indicated (*p < 0.05), as are differences from retinas treated with control siRNA directed against firefly luciferase (*p < 0.05). (B) Two intraocular injections of 50 μ M AgTx-2 or MgTx were given: one at the time of axotomy and another four days later. Comparisons are between the K_v channel blockers and control (saline-injected) retinas (*p < 0.05).

astrocyte activation, and neuron remodelling were significantly upregulated at seven days after axotomy (Fig. 1J); a time chosen to correspond with the maximal rate of RGC apoptosis in this model.¹¹⁻¹³ Indications of microglial activation were the ~10-fold increase in expression of MHC class II-a molecule that increases when antigen-presenting cells (including microglia) are activated; and ~2-fold increase in complement receptor 3-a surface receptor for immune complexes that is labeled by the OX-42 antibody. Consistent with the observed constitutive expression of CR3, microglia in both the healthy and damaged retina were labelled with OX-42 (see below). After axotomy, there was a ~ 3 fold increase in expression of glial fibrillary acidic protein (GFAP), which is upregulated in reactive astrocytes. The ~1.7 fold increase in the synaptic protein, synaptophysin, is consistent with an earlier observation that surviving RGCs have expanded dendritic trees.¹⁵ Further indications of inflammation (**Fig. 1J**) were the increased expression of TNF α and the IL1-receptor antagonist (IL-1ra); and cFos, which can be upregulated by TNF α in the injured CNS.¹⁶

 K_v blockers and RGC-specific knockdown differentially affect growth factor expression. While RGC-specific channel knockdown addresses the cellautonomous role of K_v channels in these neurons, intraocular injection of drugs can affect channels in other cell types. In Figure 2, transcript expression was quantified for several genes known to be involved in retinal responses to damage.^{3,17-19} When mRNA levels were monitored in whole-retina samples at seven days (Fig. 2A), most axotomy-induced changes were small, but significant increases were seen in nerve growth factor (NGF), brain-derived neurotrophic factor (BDNF), epidermal growth factor (EGF) and ciliary neurotrophic factor (CNTF). As before,⁵ channel-specific siRNAs were compared with a control siRNA directed against firefly luciferase. The outcome of RGC-specific K_v1.1 knockdown differed from K_v1.3 knockdown. K_v1.3 siRNA treatment decreased expression of glial-derived neurotrophic factor (GDNF) and basic fibroblast growth factor (bFGF; also known as FGF2 or FGF- β), while K_v1.1 knockdown decreased EGF and increased CNTF. In addition, effects of the two channel blockers differed from each other (Fig. 2B) and from channel knockdown in RGCs (compare with Fig. 1A). MgTx significantly increased (bFGF)



Figure 3. Combining channel knockdown with K_v 1.3 blockade differentially affects inflammatory gene expression. Retinal expression of gene transcripts was assessed by quantitative real-time RT-PCR at seven days after optic nerve transection. **Table 1** shows the primer sequences used to detect these inflammation-related genes. Freshly cut nerve stumps were immediately injected with siRNA against K_v 1.1 or K_v 1.3, and compared with a control siRNA against firefly luciferase (as for **Fig. 2A**). When used, two intraocular injections of 50 μ M AgTx-2 were given: one at the time of axotomy and another four days later (as for **Fig. 2B**). Expression of each gene was normalized to the value in axotomized retinas injected with the control siRNA. Data are expressed as mean ± SEM (six retinas each). The statistical significance was determined by ANOVA, followed by Fisher's test: *p < 0.05 indicates that the combination of the siRNA and AgTx-2 was significantly more effective than the siRNA alone.

expression several-fold and decreased neurturin (NRTN) levels, while AgTx-2 did not affect expression of any of the growth factors examined. These disparate effects might be mediated through the additional block of K_v 1.2 channels by MgTx.

Combining channel knockdown with K_v1.3 blockade differentially affects inflammatory gene expression and RGC survival. Expression of several inflammatory genes (Fig. 3) was compared after siRNA-mediated channel knockdown in RGCs alone, versus combined knockdown and intraocular AgTx-2 injection. After K_v1.1 knockdown, there was an increase in retinal TNF α , IL-1 β and TGF β regardless of whether AgTx-2 was injected; however, only the combined treatment increased cFos and IL-1ra. K_v1.3 knockdown from RGCs did not affect expression of the inflammatory genes; whereas, combining this with AgTx-2 injection increased TNF α , IL-1ra and TGF β . The most striking change was a 4- to 5-fold increase in expression of the anti-inflammatory molecule, IL-1ra, after intraocular injection of AgTx-2, but no change after cell-specific knockdown of either K_v 1.1 or K_v 1.3 from RGCs.

If RGC degeneration involves both cellautonomous and non-autonomous mechanisms, then additive neuroprotection is expected following combined treatment with a K_v1.3 blocker and channel knockdown. Results in Figure 4 show additive or supra-additive rescue of RGCs. For comparison, in the healthy rat retina, RGC densities (cells/mm²; n = 6) were: 2,446 ± 68 (inner eccentricity), $2,501 \pm 42$ (middle) and 2,208 ± 109 (outer eccentricity). At 14 days after optic nerve transection, fewer than 10% of the RGCs remained; i.e., 198 ± 12 cells/mm² at the inner eccentricity, 206 \pm 14 in the middle and 178 \pm 12 at the outer eccentricity. Representative confocal images of surviving RGCs in flatmounted retinas are shown (Fig. 4A-D), along with the summarized RGC densities measured at inner, middle and outer retinal eccentricities (Fig. 4E). By 14 days after axotomy, >90% of the RGCs had degenerated if control siRNA was injected into the cut nerve stump (Fig. 4A and E). Of note, all the treatments significantly increased RGC survival. Combined knockdown of K_v 1.1 and K_v 1.3 from RGCs (Fig. 4B and E) increased survival ~3-fold to 600-700 cells/mm²; however, this is only slightly higher than survival following knockdown of either channel alone;5 i.e., ~550 cells/ mm^2 after $K_{\rm v}1.1$ depletion (K_v1.1-922 siRNA) and ~600 cells/mm² after K_v1.3 depletion (K_v1.3-1169 siRNA). Much more survival was seen when injection of a K_v blocker was combined with either K_v 1.1 or K_v1.3 knockdown. For instance, AgTx-2 injection combined with Ky1.1-922 or K_v1.3-1169 siRNA (Fig. 4C-E) increased RGC survival to ~1,400 cells/mm² (Fig. 4E); ~2-fold higher than combined knockdown of both channels. This result is also notable because AgTx-2 blocks K_v1.1 and K_v1.3 channels, and it demonstrates that AgTx-2 is not just acting on RGCs.



Figure 4. Supra-additive RGC rescue by combined K_v block and RGC-specific K_v knockdown. (A–D) Representative confocal micrographs from flatmounted retinas of adult rats at 14 days after axotomy. RGCs were retrograde labeled with Fluorogold applied to the freshly cut optic nerve stump (see Methods). In all examples, siRNAs were injected into the cut nerve stump immediately after optic nerve transection; scale bar, 50 µm in all panels. Few RGCs survived at this time following injection of control siRNA directed against firefly luciferase (A). (B) shows increased RGC survival after injecting combined siRNAs directed against K_v1.1 (K_v1.1-922) and K_v1.3 (K_v1.3-1169). Many more RGCs were seen after intraocular injection (on Day 4) of 50 µM agitoxin-2 (AgTx-2) was combined with injection of an siRNA directed against either K_v1.1 (K_v1.1-922; (C) or K_v1.3 (D); K_v1.3-1169). (E) Summary of RGC densities 14 days after axotomy. The inset cartoon shows the approximate locations of the 12 fields that were counted from the inner, midperiphery and outer retinal eccentricities (four fields per site). The values for RGC density are mean ± SEM of four retinas in each experimental group. From left to right, the bars show the treatments: control siRNA against firefly luciferase (as above); an siRNA directed against K_v1.1, combined with two different siRNAs directed against K_v1.1 siRNA combined with intraocular injection of 50 µM margatoxin (MgTx) or AgTx-2; one of the K_v1.3 siRNAs combined with intraocular injection of 50 µM margatoxin (MgTx) or AgTx-2; one of the K_v1.3 (C) p < 0.001 with respect to K_v1.1-922 or MgTx or AgTx-2 alone; (D) p < 0.001 with respect to K_v1.3-1169 or MgTx or AgTx-2 alone.

Discussion

There are several salient findings in this study. (i) After optic nerve transection, intraocular injection of agitoxin-2, a potent blocker of K_v 1.3 channels, reduced microglial activation and expression of several inflammatory genes in the damaged retina. This provides the first evidence that identified K_v channels contribute to inflammation in the adult mammalian central nervous system in vivo,

and specifically, in the retina. (ii) Retinal expression of several growth factors was upregulated after axotomy. Intraocular injection of MgTx increased retinal b-FGF; whereas, RGC-specific depletion of K_v1.3 from RGCs decreased GDNF and b-FGF levels. AgTx-2 injection did not affect growth factor levels. (iii) Injecting AgTx-2 increased cFos, TGF β , IL-1 β , IL-1ra and TNF α beyond any effects of K_v1.1 or K_v1.3 channel knockdown in RGCs. (iv) Combining siRNA-mediated knockdown of either $K_v 1.1$ or $K_v 1.3$ with intraocular injection of AgTx-2 or MgTx provided much greater RGC rescue; up to 55% of RGCs survived at 14 days after optic nerve transection. Together with our recent study,⁵ the present results provide evidence that K_v channels are functionally linked to RGC degeneration through two distinct mechanisms: directly through cellautonomous RGC death, and indirectly through non-neuronal cells, most likely by blocking $K_v 1.3$ channels in microglia.

Microglia in the healthy adult mammalian retina are present in a thin, densely packed array in the inner plexiform layer²⁰ (reviewed in ref. 7, 9, 20 and 21). Their processes are in close proximity to RGCs and displaced amacrine cells, and their highly ramified morphology is similar to 'resting' microglia elsewhere in the healthy CNS. In contrast, the normal ganglion cell layer contains only a small number of non-ramified macrophage-like cells; often called 'perivascular microglia'. After optic nerve transection, microglia migrate to the ganglion cell layer, where they contribute to RGC degeneration. At seven days after axotomy, microglial activation and inflammation was apparent, as judged by upregulation of complement receptor-3 (CR3), a molecule involved in phagocytosis; MHC class II, which is necessary for antigen presentation; TNF α , a proinflammatory cytokine; IL-1ra, an antiinflammatory protein that antagonizes the effects of IL-1 β ; and cFos, a transcription factor involved in many intracellular signalling events. TNF α , IL-1 β and IL-1ra have been shown to have direct causative or preventative roles in RGC degeneration following injury.4,22-24 Because the entire retina was sampled, detectable changes imply large and/or wide-spread responses. If cell-specific changes occurred, they would be underestimated because microglia, astrocytes and RGCs each contribute a small proportion of total retinal cells. For instance, RGCs comprise only ~7% of the cells in the healthy retina.²⁵

When microglia activate they undergo dramatic morphological changes; their processes retract, cell bodies appear to enlarge, and they become amoeboid and mobile. Consistent with an earlier study,²⁰ at 14 days after axotomy, microglia were aligned with RGC axon fascicles in the nerve fiber layer, and did not appear to be as ramified as in the healthy retina. We previously found that blocking $K_v 1.3$ channels in LPS-activated microglia reduced their ability to produce superoxide and kill neurons in vitro; the signaling pathway was not identified but p38 MAPK activation did not appear to be involved.10 RGC death was reduced after axotomy when AgTx-2 or MgTx was injected into the eye,5 and the microglia remained ramified, much like resting microglia seen in the present study. The blocker profile in both studies suggests that $K_v 1.3$ is the main molecular target for reducing RGC degeneration. AgTx-2 blocks $K_v 1.1$ and $K_v 1.3$ channels, MgTx blocks $K_v 1.3$ and $K_v 1.2$; but α -DTX, which potently inhibits $K_v 1.2$ channels, had no effect.

Together, our two studies show that RGC rescue after injecting AgTx-2 or MgTx is not solely due to channel block in either microglia or RGCs, and might implicate other glial or neuronal cell types. RGC somata contain K_v1.1 and K_v1.3 proteins, and RGC-specific knockdown of either channel increased survival to ~25% of normal.⁵ Depleting both K_v1.1 and $K_v 1.3$ was about as neuroprotective (700– 750 RGCs/mm² survived; present study) as previously seen after injecting either MgTx (700-750 cells/mm²) or AgTx-2 (~550 cells/mm²).⁵ In contrast, combining channel knockdown with injection of either AgTx-2 or MgTx rescued about twice as many RGCs (~1,400 cells/mm²; present study). Considering that <10% of RGCs survived in untreated retinas at 14 days after axotomy, the survival of ~60% of RGCs with the most effective combination (AgTx-2 injection, plus knockdown of K_v 1.1 or K_v 1.3) is striking.

During the acute phase of inflammation after CNS damage, activated microglia can produce numerous inflammatory molecules (reviewed in ref. 8), but the consequences are not fully understood. For instance, TGF β , IL-1 β and TNF α from activated microglia play prominent roles in processes that are thought to contribute to neuron death (reviewed in ref. 26). Consistent with a role for inflammation in RGC degeneration after injury, administering IL-1ra was neuroprotective following retinal ischemia or optic nerve injury.^{23,24} However, cytokine functions are often pleiotropic, and there is evidence that TNF α and IL-1 β can improve RGC survival after axotomy. In RGCs, TNFa downregulated a K⁺ current,⁴ and IL-1 β altered Na⁺ and K⁺ currents and activated the PI3 kinase pathway.²² Both treatments reduced RGC death, but this rescue was not abolished by IL-1ra22 and it was concluded that IL-1ß exerted direct neuroprotective effects, independent of effects on microglia.27 We monitored change in mRNA levels, and while we do not know to what degree the protein levels changes, our observation that IL-1ra expression was increased by AgTx-2 treatment (but not K₁1.1 or K₁1.3 depletion from RGCs) is consistent with IL-1ra counteracting the pro-inflammatory effects of IL-1 β , while permitting the direct anti-apoptotic effects of IL-1ß in injured RGCs. TGFB regulates RGC apoptosis during development,28 and we found that AgTx-2 injection increased TGFB expression; however, adenoviral-mediated expression of TGF β did not improve RGC survival in an earlier study.29 Elevated nitric oxide can contribute to RGC degeneration in this model.³⁰ We did not observe changes in iNOS expression but it is likely that localized or transient changes were missed when we sampled the entire damaged retina. That is, Mueller glia constitutively express iNOS, which is regulated during phases of light and dark adaptation, and because these cells are large and distributed throughout the inner and outer retina, localized changes in iNOS in the damaged ganglion cell layer might not be detected. In addition, it is possible that elevated expression of IL-1ra prevented the upregulation of iNOS that follows increased IL-1 β expression.

Our finding of additive or supra-additive rescue of RGCs by injecting a K_v blocker along with depleting either K_v1.1 or K_v1.3 in RGCs points to additional indirect mechanisms of neuroprotection. Microglia can produce neurotrophic factors and microglial activation depends on functional K_v1.3 channels.¹⁰ MgTx injection, but not RGC-specific K_v1.1 or K_v1.3 depletion, increased bFGF, a growth factor previously found to rescue RGCs.³¹ Interestingly, AgTx-2, which also blocks K_v1.1 and K_v1.3, did not affect bFGF expression, suggesting that increased bFGF levels might result from the additional block of K_v1.2 by MgTx, independent of $K_v 1.1$ or $K_v 1.3$ channels.

Our observation that $K_v 1.3$ channel blockers reduce RGC degeneration in vivo has broader implications. Not only do $K_v 1.3$ channels contribute to activation of microglia,¹⁰ which are often the first cells to respond to CNS injury or disease, but $K_v 1.3$ is also expressed in some neurons. Thus, $K_v 1.3$ inhibition might be a useful

Table 1. Primers used for real-time quantitative real-time RT-PCR

Gene	GenBank accession #	Primer sequences (5'-3') (F, Forward primer; R, Reverse primer)
HPRT-1	NM_012583	F: CAGTACAGCCCCAAAATGGT (520)
		R: CAAGGGCATATCCAACAACA (646)
CR3	NM_012711	F: TGCTGAGACTGGAGGCAAC (1850)
		R: CTCCCCAGCATCCTTGTTT (1950)
MHC class II	AJ554214	F: CCAACACCCTCATCTGCTTT (376)
		R: AAGCCATCTTGTGGAAGGAA (519)
GFAP	NM_017009	F: CAGCTTCGAGCCAAGGAG (326)
		R: TGTCCCTCTCCACCTCCA (446)
synaptophysin	NM_012664	F: GTGCCAACAAGACGGAGAGT (179)
		R: ATCTTGGTAGTGCCCCCTTT (293)
τνξα	NM_012675	F: GCCCACGTCGTAGCAAAC (277)
		R: GCAGCCTTGTCCCTTGAA (444)
IL-1β	NM_031512	F: GGCTGTCCAGATGAGAGCA (197)
		R: TGCCACAGCTTCTCCACA (300)
IL-1ra	NM_022194	F: GGGAAAAGACCCTGCAAGA (88)
		R: GTGGATGCCCAAGAACACA (261)
iNOS	NM_012611	F: AAGCCCCGCTACTACTCCAT (2886)
		R: AGCTGGAAGCCACTGACACT (3280)
cFos	X06769	F: GGCAGAAGGGGCAAAGTAG (504)
		R: TTGATCTGTCTCCGCTTGG (647)
BDNF	NM_012513	F: TCCCTGGCTGACACTTTTG (427)
		R: GTCCGCGTCCTTATGGTTT (519)
CNTF	NM_013166	F: CGACTCCAAGAGAACCTCCA (291)
		R: CCTTCAGTTGGGGTGAAATG (385)
EGF	NM_012842	F: CACCAACAGCCCAAGAATG (3707)
		R: CCCGAGTTACAGGGAGGAA (3796)
bFGF (FGF2)	NM_019305	F: GTCTCCCGCACCCTATCC (22)
		R: ACAACGACCAGCCTTCCA (142)
NRTN	NM_053399	F: GGCCTACGAGGACGAGGT (501)
		R: GTCGGGGCGTGAGGTAG (610)
GDNF	NM_019139	F: AGAGGGAAAGGTCGCAGAG (359)
		R: AGCCCAAACCCAAGTCAGT (449)
TGFβ	NM_021578	F: TTCCTGGCGTTACCTTGG (943)
		R: CCGTCTCCTTGGTTCAGC (1050)

approach for controlling inflammation in other retinal and CNS disorders, and might also be directly neuroprotective. Some potent peptidyl inhibitors of K_v 1.3 have been tested in animal models, without evidence of toxicity (reviewed in ref. 14), and even poorly selective K_v 1-family blockers (e.g., 4-aminopyridine) are being tested in human clinical trials for multiple sclerosis and spinal cord injury, apparently without intolerable side effects.

Materials and Methods

Optic nerve transection, siRNA-mediated channel knockdown and K_v blocker injection. These procedures were carried out in accordance with guidelines from the Canadian Council on Animal Care. For detailed methods and references please see our recent publication.⁵ In brief, to observe retinal ganglion cells (RGCs) in the normal retina, Fluorogold was injected into the superior colliculus, allowed to undergo retrograde transport for 24 h, and then the retina was fixed. For optic nerve transection, a 1 mm section of the nerve was removed from within 1.5 mm of the back of the eye; RGCs were labeled by applying Fluorogold to the freshly cut optic nerve stump, which was retrogradely transported to their somata. $K_v 1.1$ or $K_v 1.3$ channels were selectively depleted from RGCs by injecting 10 ng of a channel-specific siRNA into the nerve stump. The siRNA sequences and pseudonyms were: 5'-CCU UGU GUA UCA UCU GGU U-3' (K_v1.1-922); and 5'-CCA TGA CAA CTG TCG GTT A-3' (K_v1.3-1169). Control injections consisted of an irrelevant Cy3-labeled siRNA (Upstate, Charlottesville, VA) directed against a non-mammalian protein, firefly luciferase: 5'-CGU ACG CGG AAU ACU UCG A-3'. We previously showed⁵ that these siRNAs were not toxic, and their effectiveness was demonstrated using immunohistochemistry in transverse retinal sections; i.e., there was specific knockdown of the salient K_v1 channel from RGCs without effects on channels in other retinal cells.

K, channel blockers were administered by intraocular injection at the time of axotomy and again at four days, the time at which RGC apoptosis begins in this model (reviewed in refs. 1-3 and 19). Three microliters of saline were injected, containing 50 µM margatoxin (MgTx) or agitoxin-2 (AgTx-2) or α -dendrotoxin (α -DTx). For details about these blockers, see ref. 14. We previously showed that these blocker concentrations had no adverse effects on the rats following intraocular injection.⁵ Control injections consisted of saline alone; i.e., 100 mM NaCl, 10 mM TRIS, 1 mM EDTA, 0.1% BSA, pH 7.5. Rats were briefly anaesthetized with isofluorane, and the vitreous chamber was injected at the time of optic nerve transection and again four days later. Injections were made with a glass micropipette posterior to the limbus, being careful to avoid damaging anterior ocular structures or puncturing the lens, which evokes release of growth factors. To assess combined responses to a channel blocker plus an siRNA, comparisons were made between control siRNA and a channel-specific siRNA delivered to the optic nerve stump.

Immunohistochemistry. Locations and morphologies of microglia were monitored in normal retinas and five days after optic nerve transection. Whole mounts were prepared from fixed retinas and labeled overnight at 4°C with a mouse monoclonal antibody against complement receptor 3 (OX-42; Serotec, Oxford, UK) at a 1:250 dilution in phosphate-buffered saline (PBS) containing 3% normal goat serum and 0.3% Triton X-100. Retinas were then washed with PBS (3x, 15 min each) and incubated for 4 h at room temperature with an affinity-purified Alexa 488-conjugated goat-anti-mouse secondary antibody (Jackson ImmunoResearch, West Grove, PA), diluted in PBS. After washing with PBS (3x, 15 min each), retinas were flat-mounted with 1:1 glycerol:PBS and stored at 4°C. For each assay, the concentration of antibodies, fluorescence intensity gain, and pinhole and neutral density filters were kept constant. There was no staining when the primary antibody was omitted (not shown), and the black regions within each image in Figure 1 show the lack of non-specific OX-42 staining.

quantitative RT-PCR. Real-time Changes in gene transcript expression were monitored by quantitative RT-PCR (qRT-PCR) in whole retinas harvested seven days after axotomy, as before.⁵ Gene-specific primers (Table 1) were designed using the 'Primer3Output' program (HUhttp://frodo.wi.mit.edu/ cgi-bin/primer3/primer3_www.cgiUH). RNA was isolated (RNeasy; Qiagen, Inc., Mississauga, Ontario) after degrading any contaminating DNA with DNaseI (0.1 U/mL, 15 min, 37°C; Amersham Biosciences, Baie d'Urfe, Quebec). Total RNA (2 μ g) was reverse transcribed in 20 µL volume using 200 U of SuperScriptII RNase H-reverse transcriptase, with 0.5 mM dNTPs (Invitrogen, Carlsbad, CA) and 0.5 µM oligo dT (Sigma-Aldrich, St. Louis, MO). Amplification was performed on an ABI PRISM 7900 Sequence Detection System (PE Biosystems, Foster City, CA) at 95°C for 10 min, followed by 40 cycles at 95°C for 15 s, 55°C for 15 s and 72°C for 30 s. For each gene, 'no-template' and 'no-amplification' controls were negative, and melt curves showed a single peak, confirming specific amplification. The threshold cycle (C_T) for each gene was determined and normalized against the housekeeping gene, hypoxanthine guanine phosphoribosyl transferase (*HPRT-1*). Significant differences between experimental treatments were identified by ANOVA, followed by Fisher's Mean Interval Comparisons.

Quantifying RGC survival after axotomy. As before,⁵ we quantified RGC survival at 14 days post-axotomy. Rats were sacrificed by an overdose of anaesthetic (7% chloral hydrate), their eyes were enucleated, the cornea and lens removed, and the remaining eye cups fixed (1.5 h, room temperature) in 4% paraformaldehyde in PBS containing 2% sucrose. The eye cup was washed with PBS for 15 min, and the neural retina was dissected away from the sclera and flat-mounted in 1:1 glycerol:PBS. The density of surviving Fluorogold-labeled RGCs was counted from confocal micrographs (Zeiss LSM 510). Because RGC density varies somewhat across the retina, we used a standard procedure, wherein cells in flat-mounted retinas were counted in a 70,000 µm² area at three retinal eccentricities (inner, middle, outer; see cartoon in Fig. 4E) from each quadrant of the retina, for a total of 12 fields of cells per retina.

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