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BCL2L1 (BCL-x) promotes survival of adult and developing retinal ganglion cells

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Abstract

The Bcl-2 family is responsible for regulating cell death pathways in neurons during development, after injury and in disease. The activation of the pro-death family member BAX is often the final step before cell death in neurons. Pro-survival family members such as BCL-X (BCL2L1) act to inhibit BAX activation. Overexpression studies have suggested that BCL-X could play an important physiological role in mediating neuronal viability. Loss-of-function studies performed *in vivo* have implicated BCL-X as a mediator of neuronal survival during the early stages of neurodevelopment. To assess whether BCL-X is needed to promote the survival of neurons in the central nervous system throughout life, *Bcl-x* was conditionally removed from the optic cup or throughout the adult mouse. During development BCL-X was required for the survival of differentiating retinal ganglion cells (RGCs) leading up to their normal window of developmental death. Despite its expression in adult RGCs, BCL-X was not required for maintaining RGC viability in adult retinas. However, the loss of BCL-X in adult RGCs did significantly increase the rate of death of RGCs after axonal injury. Thus, in developing and injured RGCs there appears to be an active cell survival program preventing neuronal death.

Introduction

The Bcl-2 family of genes mediates the intrinsic pathway of apoptosis, which significantly contributes to neuronal death during development, after injury, and in disease. For instance, the pro-death Bcl-2 family member BAX is required for retinal ganglion cell (RGC) death during development, after acute axonal injury, and in ocular hypertensive glaucoma (Li et

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al., 2000; Libby et al., 2005; Mosinger Ogilvie et al., 1998; Qin et al., 2004; White et al., 1998). BAX activation is controlled by the opposing actions of pro-death and pro-survival members of the Bcl-2 family. During development and after injury RGC apoptosis requires upstream pro-death Bcl-2 family members (Harder and Libby, 2011; McKernan and Cotter, 2007). The physiological role of the pro-survival Bcl-2 family members is less well understood than their pro-death counterparts. Importantly, while *Bcl2* was shown to not have a role in maintaining RGC survival after axonal injury (Dietz et al., 2001), it does help maintain RGC viability in maturing RGCs (Cellerino et al., 1999). Thus, pro-survival Bcl-2 family members can play critical roles in maintaining RGC viability.

There are five pro-survival members of the Bcl-2 family (*Bcl2*, *Bcl-x*, *Bcl-w*, *Bcl2a1a*, and *Mcl1*). Their main function is to prevent BAX (or BAK1) activation and, therefore, cell death (Puthalakath and Strasser, 2002; Strasser, 2005; Willis et al., 2005; Willis et al., 2007). Most of the pro-survival Bcl-2 family members have important roles in neuronal development (Arbour et al., 2008; Crosio et al., 2006; Lukiw et al., 2005; Middleton et al., 2001; Mori et al., 2004; Motoyama et al., 1995; Shacka and Roth, 2006). In fact, antagonization of pro-survival Bcl-2 family members with small molecule inhibitors is enough to trigger neuronal death *in vitro* (Young et al., 2010). BCL-X has been specifically implicated as an important pro-survival factor in neuronal development and disease. Germline deletion of *Bcl-x* leads to death of neurons in the developing central nervous system and embryonic lethality (Motoyama et al., 1995). Conditional deletion of *Bcl-x* in dopaminergic neurons showed that *Bcl-x* is required for the survival of all but a few catecholaminergic cells in the developing substantia nigra (Savitt et al., 2005). Numerous neuroprotective treatments are reported to increase the intracellular ratio of BCL-X to pro-apoptotic members (Kilic et al., 2005; Koh, 2009; Ma et al., 2005; Pike, 1999; Wang et al., 2000) and in injured neurons overexpressing BCL-X can increase survival and sustain neuronal function (Garrity-Moses et al., 2005; Parsadanian et al., 1998; Wiessner et al., 1999). In RGCs *Bcl-x* transcript and protein expression is regulated after injury (Isenmann et al., 1997; Levin et al., 1997; McKernan and Cotter, 2007; Pelzel et al., 2010) and overexpression of BCL-X or BCL2 protects RGCs after axonal injury (Bonfanti et al., 1996; Cenni et al., 1996; Chierzi et al., 1999; Malik et al., 2005). Together these studies suggest that BCL-X may play a necessary physiological role in maintaining survival of adult and developing neurons. However, despite the importance of apoptotic cell death during development and in disease, to date there is limited knowledge of how critical physiological levels of pro-survival Bcl-2 family members are in maintaining neuronal survival throughout life (Isenmann et al., 2003). To test the function of an endogenous pro-survival Bcl-2 family member in the central nervous system, the role of *Bcl-x* (*Bcl2l1*) was assessed in developing and adult RGCs *in vivo*.

Methods

Animals

A floxed allele of *Bcl2l1^{tm1.1Mam}* (*Bcl-xfl*; Rucker et al., 2000) was removed from the developing retina using the *Six3-cre* allele (Furuta et al., 2000) and from the adult mouse using a ubiquitously expressed, tamoxifen inducible cre (*Cre-ERTM*; Hayashi and McMahon, 2002). A tamoxifen dose equivalent to 5mg/40g mouse was administered by intraperitoneal injection to 45-75 day old mice for 5 consecutive days. Experiments were performed either 15 days (controlled optic nerve crush) or 60 days (assessing long term survival in the absence of *Bcl-x*) following the first tamoxifen injection. Control mice were either heterozygous or wild type for the floxed allele. In addition, control mice containing cre were compared to control mice without cre to rule out effects of cre toxicity or *Bcl-x* heterozygosity in the retina using at least 3 mice with and without cre for comparison. No differences were noted between any control genotype (*Bcl-x^{+/?} Six3-cre[?]* or *Bcl-x^{+/?} Cre-*

ER^{TM2}) and control genotypes were combined and are referred to throughout the manuscript as control. Note that the \times in the control genotypes is to denote that two types of genotypes are combined to make the control group: for the *Bcl-x* gene the \times reflects the fact that that allele could be wildtype or floxed; for Cre locus the \times reflects the fact that the allele could be present or absent. Morning vaginal plug checks were used to establish age E0.5 for embryonic stages. All experiments were conducted in accordance with the Association for Research in Vision and Ophthalmology's statement on the use of animals in ophthalmic research and were approved by the University of Rochester's University Committee on Animal Resources. Significant differences were determined by comparing control and *Bcl-x* knockout groups with Student t test at each time point during development and after CONC. At least 4 animals are used for each group in each comparison. Specific animal numbers for each experiment are detailed in the relevant figure legend.

Histology and Immunocytochemistry

For retinal whole mounts and sectioning (plastic and frozen), tissue was processed as previously described (Fernandes et al., 2012; Harder and Libby, 2011). Plastic sections were used for retinal cross-section thickness measurements. Four areas were averaged per retina, each within 50 μ m of the optic nerve. The measurement was taken from the nerve fiber layer to the tips of the photoreceptor outer segments. For retinal area measurements, whole retina images were reconstructed from 5X images of flat mounted retinas and area measurements were made using Image J. For immunohistochemistry, the following primary antibodies and dilutions in blocking solution were used: rabbit anti-cCASP3 marks cells with activated caspase 3, R&D Systems, 1:1000; goat anti-POU4F2, also known as BRN3B, labels RGCs (Gan et al., 1999), Santa Cruz, 1:300; mouse anti-TUJ1 immunolabels β III-tubulin which is expressed in RGCs (Cui et al., 2003), Covance, 1:1000; rabbit anti-BCL-X, Cell Signaling, 1:500. Alexafluor-conjugated secondary antibodies (Invitrogen) were used at a dilution of 1:1000. For all cell counts identical retinal areas were assessed. For adult anti-CASP3 and anti-TUJ1 counts flat mounted retinas were counted over 20X fields and 40X fields respectively. 8 peripheral fields, 2 from each quadrant, were counted per eye. For embryonic cell counts four comparable central horizontal sections were counted per eye (cCASP3+ cells were counted along the entire section and POU4F2+ cells were counted per section at E12.5 and over 250 μ m at E18.5).

Controlled optic nerve crush (CONC)

CONC was performed 15 days after the first tamoxifen injection, as previously described (Fernandes et al., 2012; Harder and Libby, 2011). In anesthetized mice, the optic nerve was exposed and clamped using self-closing forceps (Roboz RS-5027) for 3-4 seconds just behind the eye. Unmanipulated or sham surgery (all procedures performed except clamping the nerve) eyes were used as controls.

Results

Immature RGCs require BCL-X for survival

BCL-X is expressed throughout the neuroblastic retina (Fig. 1A). At E12.5, RGCs are being born at the ventricular surface (VS) and migrating to the presumptive RGC layer at the inner surface of the retina. BCL-X and the RGC marker β III-tubulin (TUJ1; Cui et al., 2003) are coexpressed in the GCL at E12.5 and E18.5. The expression pattern of BCL-X indicates it may affect RGC birth and survival. Since germline deletion of *Bcl-x* results in embryonic lethality (Motoyama et al., 1995; Savitt et al., 2005), Six3-cre was used to delete a floxed allele of *Bcl-x* (*Bcl-x^f*) from the developing optic cup. RGCs are among the earliest retinal neurons to get specified, represent a large proportion of the first wave of differentiation (Gan et al., 1999), and undergo a significant amount of programmed cell death during

development (Mosinger Ogilvie et al., 1998). Based on POU4F2 immunolabeling (an early marker of RGC differentiation; Gan et al., 1999), *Bcl-x* deletion does not alter RGC generation at E12.5 (Fig. 2A,B). Also, the loss of BCL-X does not significantly increase cell death at E12.5, suggesting that both retinal progenitors and newly born RGCs do not require BCL-X for survival (Fig. 2C). Substantial naturally occurring developmental death of RGCs begins around E18.5 (Pequignot et al., 2003). However, without BCL-X large numbers of RGCs are prematurely lost between E12.5 and E18.5. During this time period, ectopic CASP3 activation and thinning of the retina indicate cell death is coincident with the loss of RGCs (Fig. 2C,D). These data indicate that in differentiating RGCs BCL-X is required to prevent apoptotic cell death.

In addition, other types of retinal neurons appear to be susceptible to apoptotic death in the absence of *Bcl-x*. At E18.5, the few surviving RGCs are primarily cells in which *Bcl-x* was not deleted (Fig. 2A arrow), which is consistent with known mosaic expression of *Six3-cre*, particularly in the retinal margin (Cai et al., 2011; Fuhrmann et al., 2009; Poche et al., 2008). Thus, the large increase in cell death in *Bcl-x^{fl/fl}* *Six3-cre⁺* retinas at E18.5 is likely also associated with abnormal cell death of other types of retinal neurons.

The increase in cell death during development produced a smaller adult retina in the *Bcl-x^{fl/fl}* *Six3-cre⁺* mice compared to wild type. The *Bcl-x* knockout retina was reduced in thickness (Fig. 3A; *Bcl-x^{+/?}* *Six3-cre⁺* 180±7 μm, *Bcl-x^{fl/fl}* *Six3-cre⁺* 96±7 μm; P<0.001, N=5 for each genotype) and surface area (Fig. 3B; *Bcl-x^{+/?}* *Six3-cre⁺* 18±1 mm², *Bcl-x^{fl/fl}* *Six3-cre⁺* 13±2 mm²; P<0.001, N=5 for each genotype). The *Bcl-x^{fl/fl}* *Six3-cre⁺* retina consisted of all major cell types, albeit reduced in number, and retained normal gross morphology (Fig. 3A and data not shown). In wild type adult retinas all RGCs express BCL-X, as determined by colabeling with the RGC marker TUJ1 (Fig. 3C). In 6 out of 6 *Bcl-x* knockout retinas all surviving RGCs expressed BCL-X (the *Six3-cre* is not a complete retinal deleter (Cai et al., 2011; Fuhrmann et al., 2009; Poche et al., 2008)). These results indicate that RGCs require BCL-X for survival during development and suggest that the survival of RGCs in the *Bcl-x^{fl/fl}* *Six3-cre* retinas is the result of incomplete deletion of *Bcl-x* in the developing retina.

***Bcl-x* is not required for survival of adult RGCs**

The continued expression of BCL-X in adult RGCs and its role as a required survival factor during development raises the question of whether adult RGC viability is also dependent on BCL-X. To address the importance of BCL-X in adult RGCs, *Bcl-x^f* was removed in adult mice using a ubiquitously expressed, tamoxifen inducible cre-recombinase (Cre-ERTM). Tamoxifen-induced recombination was highly effective, reducing the number of BCL-X positive cells in the RGC layer to 2% of control by 15 days following treatment (Fig 4A). Loss of BCL-X produced no noticeable change in retinal architecture and did not induce immediate cell death (Fig 4B, 5A,B) or signs of glial activation (data not shown). In fact, two months after *Bcl-x* deletion there was still normal retinal architecture (Fig 4B) and normal numbers of RGCs (Fig. 4C).

BCL-X is a pro-survival factor in injured RGCs

Adult RGCs die by apoptotic pathways regulated by the Bcl-2 family after axonal injury (either mechanical or glaucomatous axonal injury; Bahr, 2000; Isenmann et al., 2003; Li et al., 2000; Libby et al., 2005). This cell death involves activation of BAX by other pro-death Bcl-2 family members. Over-expression of BCL-X can protect against cell death following axonal injury (Malik et al., 2005), but it is unknown whether endogenous BCL-X prevents death in injured adult neurons. After acute axonal injury there is near immediate pro-death injury signaling in RGCs (Agudo et al., 2009; Fernandes et al., 2012; Lukas et al., 2009), but RGC death does not begin until three days after injury and RGC dropout continues over the

course of weeks. *Bcl-x* expression has been reported to transiently increase in retinas after axonal injury by several groups (Isenmann et al., 1997; Levin et al., 1997; Pelzel et al., 2010). Thus there is evidence supporting the possibility of an active pro-survival pathway in adult RGCs.

To test the importance of BCL-X in injured adult neurons, RGC axons were mechanically injured using a controlled optic nerve crush (CONC, a standard technique used to induce axonal injury in RGCs) injury. Similar to wild type, the BCL-X negative retina had no observable cell death at one day following axonal injury. In distinct contrast to the negligible number of apoptotic RGCs at two days in the injured wild type retina, the BCL-X negative retina had a significant increase and an immediate peak of apoptotic death (Fig 5A,B). Cell death did not peak in the wild type retina until at least 5 days (Fig. 5B and Harder and Libby, 2011) and this peak was significantly less than the amount of death observed at 2 days in the BCL-X negative retina. The early loss of RGCs in *Bcl-x* deficient retinas was confirmed by counting surviving RGCs 5 days after injury (Fig. 5C,D). This striking pattern of early cell death in the mutant indicates that BCL-X acts as a survival factor allowing many RGCs to withstand the initial cell death signaling that occurs following axonal injury.

Discussion

Pro-survival Bcl-2 family members are powerful mediators of cell survival (Adams, 2003; van Delft and Huang, 2006; Willis et al., 2007). In fact, antagonizing pro-survival Bcl-2 family members can initiate the mitochondrial cell death pathway in at least one type of cultured neuron (Young et al., 2010). In addition pro-survival Bcl-2 family members prevent pro-apoptotic BH3-only proteins from activating BAX (or its subfamily members BAK1 and BOK) (Puthalakath and Strasser, 2002; Strasser, 2005; Willis and Adams, 2005). Different cell types, including neurons, appear to vary in their latent potential for BAX activation in the absence of pro-survival Bcl-2 family members (Young et al., 2010). Overexpression of both BCL-X and BCL2 in RGCs has been shown to protect RGCs from death during development and after insult in the adult (Bonfanti et al., 1996; Cenni et al., 1996; Chierzi et al., 1999; Malik et al., 2005). Different forms of stress can also variably alter the expression of pro-survival and pro-apoptotic Bcl-2 family members and induce cell death. Despite a potentially critical role in survival, the physiological function of pro-survival Bcl-2 family members in neurons throughout life is not well understood. Here, loss-of-function studies were used to determine at what stages RGCs require pro-survival signaling of BCL-X for survival.

Developing RGCs require *Bcl-x* for survival

Proper retinal development requires a significant amount of programmed cell death and this death is dependent on pro-apoptotic Bcl-2 family members (BAX and BBC3; Harder and Libby, 2011; Mosinger Ogilvie et al., 1998; White et al., 1998). The pro-survival Bcl-2 family member BCL-X is expressed in the retina throughout development and is present in RGCs suggesting it may be required to counteract pro-apoptotic signaling in developing RGCs. In the absence of *Bcl-x*, widespread ectopic cell death in the developing nervous system has been reported (Motoyama et al., 1995), although embryonic lethality at E13.5 precluded assessing the retinal phenotype or the absolute requirement for BCL-X for neuronal survival. Deleting *Bcl-x* with a retinal specific cre at the beginning of retinal development (Six3-cre; Furuta et al., 2000) led to ectopic death of developing RGCs. In the developing brain, based on cell position in the developing tissue, it has been suggested that the general role of *Bcl-x* is in maintaining survival of differentiating neurons (Motoyama et al., 1995; Shindler et al., 1997). Using an early, specific marker for RGCs, POU4F2 (Badea et al., 2009; Wang et al., 2002), it is clear that BCL-X is not required for RGC survival in the earliest stages of differentiation. RGCs were initially generated in normal numbers and

migrated to/toward their proper position in the retina. Normal developmental death of RGCs occurs between E18.5 and P7 after RGC generation is complete and results in about half of all RGCs dying (Pequignot et al., 2003). Surprisingly, in the *Bcl-x* deficient retina, RGC death occurred prematurely between E14.5 and E18.5, corresponding with a period when few wild type RGCs die. The data presented here demonstrate that an active pro-survival factor is required for RGCs to survive soon after differentiation begins. This result implies that a pro-survival pathway that includes BCL-X antagonizes an active cell death signal that is normally present during the differentiation process. It will be interesting to determine if the downregulation of pro-survival factors contributes to RGC death during the normal death window because this change appears to be sufficient to induce apoptosis. Defining the extent and regulation of both the pro-survival and pro-death pathways will be needed to fully understand neuronal developmental death.

Based on the adult retinal morphology including decreased retinal size and decreased thickness of all retinal layers, other retinal neurons may also require BCL-X during development. Consistent with increased death of other developing neurons, after most RGCs have died at E18.5, there are significantly more activated CASP3+ cells in the *Bcl-x* deficient retinas. However, early born RGCs can help control later retinal neuron production by affecting retinal progenitor proliferation (Mu et al., 2005; Wang et al., 2005). Thus, the loss of RGCs may also contribute to the decrease in other cell types. The use of other cres will be needed to test the importance of *Bcl-x* in the development of later born retinal neurons.

Endogenous BCL-X delays adult RGC death after acute axonal injury

Similar to developing RGCs, adult RGCs robustly express BCL-X, raising the possibility that adult RGCs also require BCL-X in order to survive. To rule out developmental effects, *Bcl-x* was conditionally disrupted in the adult and RGC survival was assessed. Gross retinal morphology, including the number of RGCs, was still normal 60 days after *Bcl-x* deletion. Thus at some point during development RGCs lose their requirement for BCL-X as a survival factor. Other neurons in the central nervous system appear to exhibit a similar phenotype. For example, although BCL-X is normally expressed in adult catecholaminergic neurons, Savitt et al. observed a few surviving catecholaminergic neurons not expressing BCL-X in the adult (Savitt et al., 2005). One caveat of their observation was the possibility of developmental effects due to *Bcl-x* deletion since *Bcl-x* was conditionally deleted in these neurons during embryonic development. However in combination with results presented here, it appears that BCL-X is not required for the survival of normal adult neurons despite its wide expression pattern in the central nervous system. Interestingly, using a *Bcl2* knockout mouse Cellerino and colleagues showed that *Bcl2* was involved in RGC survival after the normal window of RGC death (Cellerino et al., 1999). These data suggest different *Bcl-2* family members play key roles in maintaining RGC viability at various stages of maturation.

RGC death following axonal injury, an important insult in glaucoma (Anderson and Hendrickson, 1974; Howell et al., 2007; Howell et al., 2012; Quigley et al., 1983; Schlamp et al., 2006), is another form of apoptotic death that requires pro-apoptotic *Bcl-2* family members. The activation of an apoptotic death pathway suggests that endogenous pro-survival *Bcl-2* family members may determine an RGC's susceptibility to death after an axonal insult. In fact, RGC death occurred earlier after mechanical optic nerve injury in the *Bcl-x* deficient retina compared to controls. By 2 days following injury, an RGC's resistance to apoptosis appears to depend in part on BCL-X. This may be attributed to its ability to interact with BAX (Adams, 2003; van Delft and Huang, 2006) and its role in linking cellular metabolism to apoptotic sensitivity (Yi et al., 2011). Endogenous expression levels of BCL-X have been reported to be upregulated and downregulated after axonal injury in RGCs

(Iseman et al., 1997; Levin et al., 1997; McKernan and Cotter, 2007; Pelzel et al., 2010) suggesting that to fully unravel the process by which RGCs die. Thus, RGCs need to be examined on an individual cell basis in order to isolate surviving versus dying cells. In pathological conditions, RGC death is likely achieved by the complex interplay between the pro-survival and pro-death Bcl-2 family members (Nickells, 2010). BCL-X clearly has a major role in this process, but pro-death members BAX, BIM, BBC3, and potentially others are required to activate the intrinsic pathways of apoptosis in RGCs after axonal injury (Harder and Libby, 2011; Li et al., 2000; Libby et al., 2005; McKernan and Cotter, 2007; Qin et al., 2004). The relative level of expression of all these molecules is likely critical to determining how much of an insult a cell can withstand before undergoing apoptosis. Finally, due to the specific importance of BCL-X in resisting pro-apoptotic signaling in RGCs, fluctuations in BCL-X expression may also affect susceptibility to chronic or variable insults. Thus it may be important to identify how BCL-X and other Bcl-2 family members are regulated in adult neurons. Manipulating these endogenous pathways may increase a neurons ability to withstand an insult and therefore, be a potentially powerful therapeutic target.

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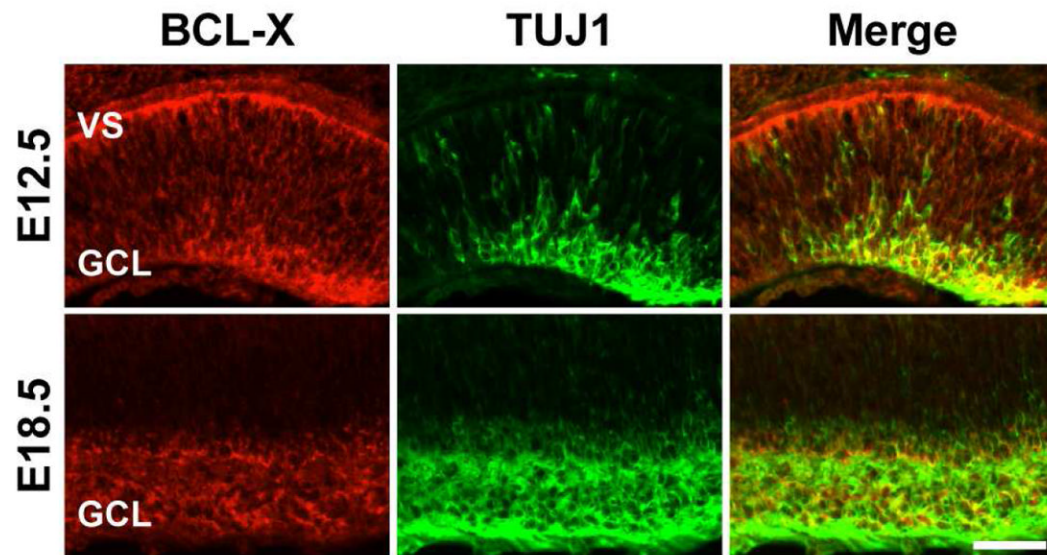


Figure 1. BCL-X is expressed in differentiating RGCs

BCL-X (red) is expressed by differentiating RGCs (TUJ1+, green) in the retina. At E12.5, RGCs are being born at the ventricular surface (VS) and migrating to the presumptive RGC layer (GCL). BCL-X and TUJ1 are coexpressed (yellow) in the GCL at both E12.5 and E18.5. 4 retinas were examined for each genotype and time point.

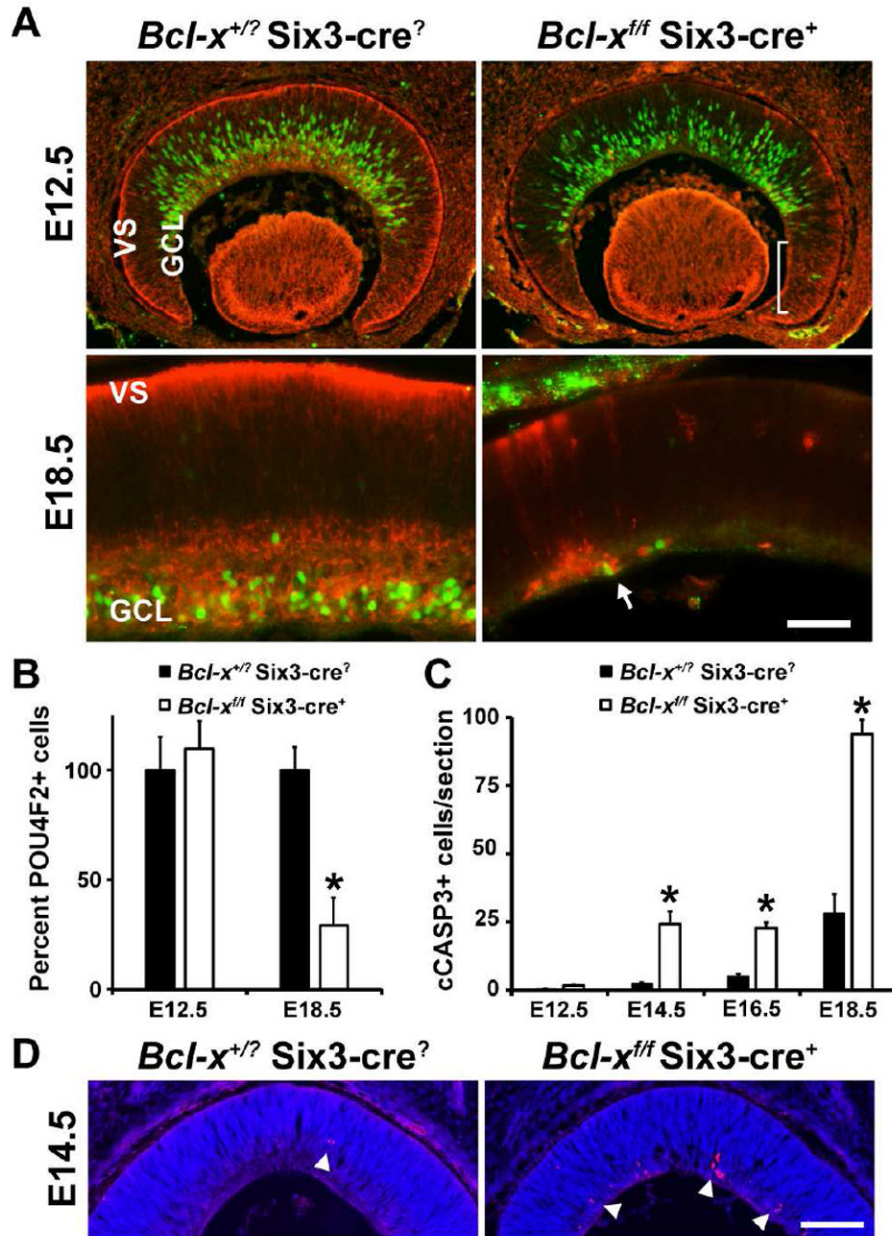


Figure 2. BCL-X is required for the survival of differentiating RGCs

(A, B) Control (*Bcl-x^{+/?} Six3-cre[?]*) RGCs express the RGC specific marker POU4F2 (green) soon after undergoing fate determination at the ventricular surface of the retina. Loss of BCL-X (red) in the retina (*Bcl-x^{fl/fl} Six3-cre⁺*) does not alter the production or migration of RGCs as judged by their distribution and number. By E18.5, a time when all RGCs are born and reside in the GCL, there is a significant reduction in the number of RGCs ($P < 0.001$), suggesting RGCs die during the differentiation process. Note, there is an incomplete deletion of *Bcl-x* by the *Six3-cre* allele (bracket) and surviving RGCs still express BCL-X (arrow). (C) The loss of RGCs by apoptosis is supported by an increase in the number of cleaved caspase 3 (cCASP3+) cells. This increase was statistically significant at each time point assessed (*, $P < 0.01$) except for E12.5 ($P = 0.78$). (D) Representative images of anti-cCASP3 immunostaining (red) with a nuclear counter stain (DAPI, blue) at E14.5 in the

retinas of control and *Bcl-x^{fl/fl}* Six3-cre⁺ mice. At least 4 different animals at each time point and for each genotype were assessed. Scale bar, **A**, 50μm; **D**, 100μm.

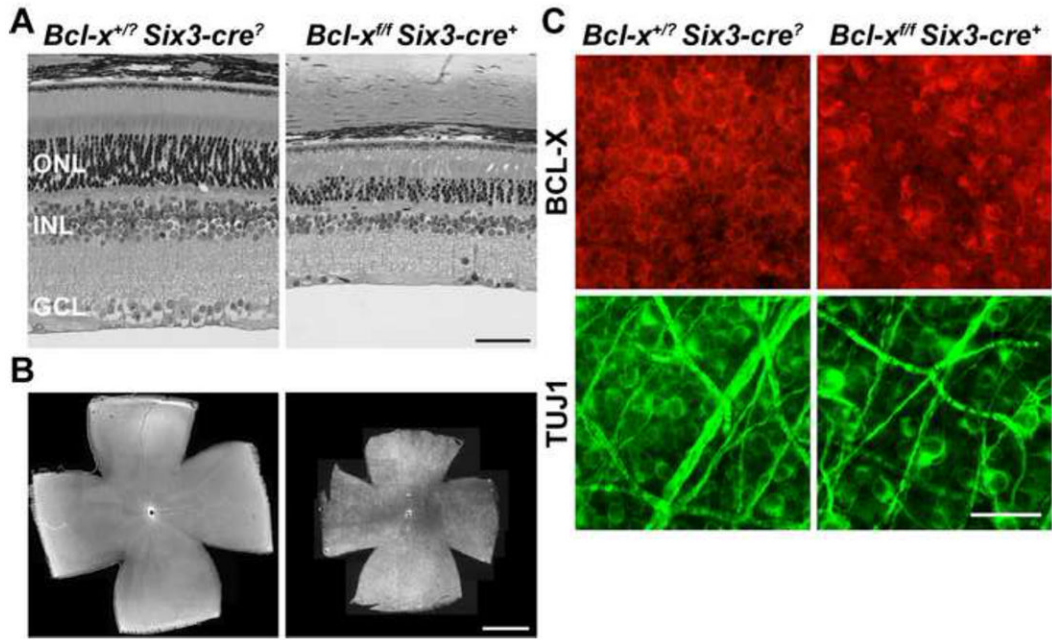


Figure 3. BCL-X is required for retinal neuron survival during development

(A) *Bcl-x^{f/f} Six3-cre⁺* retinas at P45 have normal retinal layer organization, but a significant loss in retinal neuron number compared to control retinas (both thickness and surface area were significantly smaller; $P < 0.001$ for each measurement; $N = 5$ for each genotype). (B) Whole mount retinas, ganglion cell layer up, from control animals shows that BCL-X is expressed across the retina. In the smaller *Bcl-x^{f/f} Six3-cre⁺* retinas BCL-X appears to still be expressed in the ganglion cell layer ($N = 5$ for each genotype). (C) Colabeling of the RGC marker TUJ1 and BCL-X shows that all surviving RGCs contained BCL-X (6 retinas, with at least 8, 40X fields examined per retina). Thus, BCL-X is required for RGC survival during development and the remaining RGCs in *Bcl-x^{f/f} Six3-cre⁺* eyes likely survive because *Bcl-x* is still present. ONL, Outer Nuclear Layer; INL, Inner Nuclear Layer; GCL, Ganglion Cell Layer; Scale bar: A, 50 μ m; B, 1mm; C, 25 μ m.

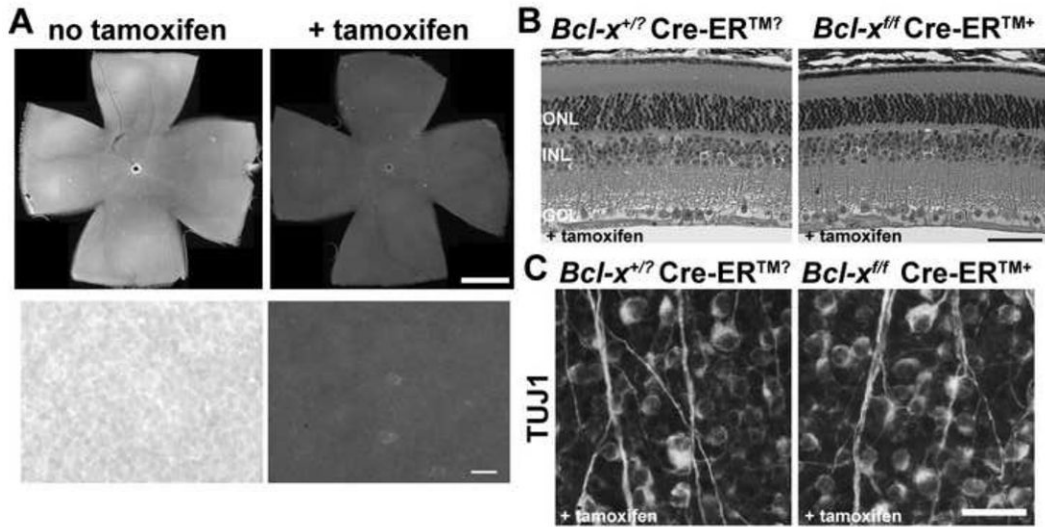


Figure 4. Adult RGCs do not require BCL-X for survival

Bcl-x^f was deleted in the adult using a ubiquitously expressed, tamoxifen inducible cre (Cre-ERTM). (A) Assessment of BCL-X expression in *Bcl-x^{fl/fl}* Cre-ER^{TM+} mice with and without tamoxifen administration in retinal flat mounts (ganglion cell layer facing up) showed that *Bcl-x* was efficiently deleted in the adult using this Cre-ERTM (Top, low power composite image of entire retinas immunolabeled for BCL-X; Bottom, high power image showing individual cells). (B) There was no apparent loss of retinal cells up to 60 days after deletion of *Bcl-x* (4 animals were assessed for each genotype). (C) Quantification of RGCs using the RGC specific marker TUJ1+ (Cui et al., 2003) confirmed that there was no loss of RGCs 60 days after deleting BCL-X (RGCs/mm² in controls: 2828±203; *Bcl-x^{fl/fl}* Cre-ER^{TM+}: 2695±221; P=0.67; N=4 for each genotype). ONL, Outer Nuclear Layer; INL, Inner Nuclear Layer; GCL, Ganglion Cell Layer. Scale bar: A top, 1mm, bottom, 25µm; B, 50µm; C, 25µm.

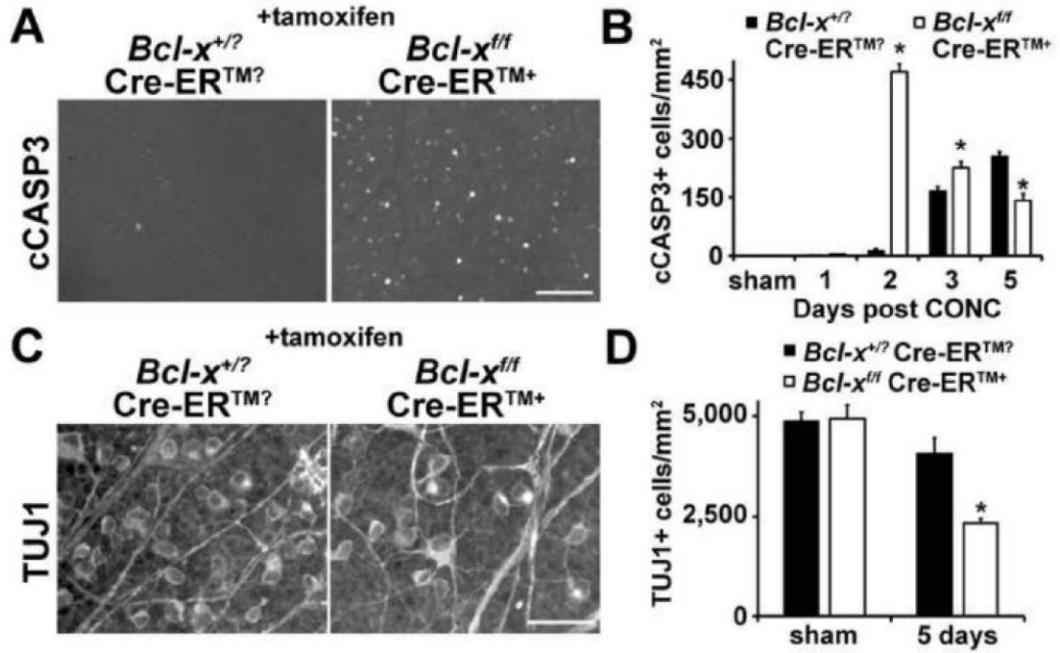


Figure 5. BCL-X is a pro-survival factor in injured RGCs

(A,B) After axonal injury (controlled optic nerve crush, CONC), RGC death begins around 2 days after injury, with only occasional cleaved caspase 3+ cells (cCASP3) being detected in wild type animals and the peak of RGC death occurring 5 days after injury (Harder and Libby, 2011). In the absence of *Bcl-x* (deleted in 45-75 day old mice using a ubiquitously expressed tamoxifen inducible cre, Cre-ERTM) RGC cells undergo death at a much faster rate. In fact, RGC death peaks at 2 days after injury. Though by 5 days after injury, wild type mice have more cCASP3+ cells than *Bcl-x* knockout mice, presumably because there are less RGCs remaining to die in the mutants. (C,D) The increase in RGC death was confirmed by counting RGCs 5 days after injury. Eyes deficient in *Bcl-x* (*Bcl-x*^{fl/fl} Cre-ER^{TM+}) had a significantly reduced number of TUJ1+ RGCs (TUJ1+ is an RGC specific marker; Cui et al., 2003). At least 5 different animals at each time point and for each genotype were assessed. *, P<0.01; Scale bar: A, 50μm; C, 25μm.