Control of lens development by Lhx2-regulated neuroretinal FGFs
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ABSTRACT
Fibroblast growth factor (FGF) signaling is an essential regulator of lens epithelial cell proliferation and survival, as well as lens fiber cell differentiation. However, the identities of these FGF factors, their source tissue and the genes that regulate their synthesis are unknown. We have found that Chx10-Cre;Lhx2lox/lox mice, which selectively lack Lhx2 expression in neuroretina from E10.5, showed an early arrest in lens fiber development along with severe microphthalmia. These mutant animals showed reduced expression of multiple neuroretina-expressed FGFs and canonical FGF-mediated microphthalmia. These mutant animals showed reduced expression of multiple neuroretina-expressed FGFs and canonical FGF-mediated microphthalmia. These mutant animals showed reduced expression of multiple neuroretina-expressed FGFs and canonical FGF-regulated genes in neuroretina. When FGF expression was genetically restored in Lhx2-deficient neuroretina of Chx10-Cre; Lhx2lox/lox mice, we observed a partial but nonetheless substantial rescue of the defects in lens cell proliferation, survival and fiber differentiation. These data demonstrate that neuroretinal expression of Lhx2 and neuroretina-derived FGF factors are crucial for lens fiber development in vivo.

KEY WORDS: FGF, Genetics, Lens, Mouse, Retina, Transcription factor

INTRODUCTION
Vertebrate lens development has long been a model system for the study of the role of inductive signaling in tissue patterning and cell specification (Gunhaga, 2011). During embryogenesis, the surface ectoderm adjacent to the optic vesicle thickens and invaginates to give rise to the lens vesicle. Subsequently, cells in the anterior lens vesicle become a monolayer of lens epithelial cells, while cells in the posterior half of the vesicle elongate and differentiate to form primary lens fibers, which fill the lens vesicle. After this distinctive architecture has been established, lens growth continues in a spatially restricted manner maintaining lens polarity. Lens epithelial cells proliferate in the germinative zone, the region just above the lens equator. The progeny then migrate or are displaced below the equator into the transitional zone where they give rise to the lens fiber. FGFs have long been a top candidate among these, following landmark studies demonstrating that treatment of cultured lens epithelial cells with FGF1 and FGF2 was sufficient to induce differentiation into lens fiber cells (Chamberlain and McAvoy, 1987, 1989). Subsequent studies have provided compelling evidence that FGFs promote lens epithelial cell proliferation and fiber cell differentiation in a dose-dependent manner (Lovicu and McAvoy, 2005; McAvoy and Chamberlain, 1989; Schulz et al., 1993). Although this stimulated speculation that FGFs might be the long-sought neuroretina-derived signal, targeted mutation studies carried out over recent decades have yet to identify any specific FGFs that are necessary, either individually or in combination, for lens cell proliferation or differentiation (Robinson, 2006). More recently, targeted deletion of three FGF receptor genes (Fgfr1, Fgfr2 and Fgfr3) in lens pit/vesicle have been shown to disrupt lens cell survival and fiber cell differentiation. However, individual or pairwise deletion of FGF receptor genes still resulted in grossly normal lens formation (Zhao et al., 2008). This functional redundancy of FGF receptors suggests that multiple FGFs act in concert to promote lens fiber differentiation. The identities and source of these factors, however, remain obscure. 

Lhx2 is a LIM-homeodomain transcription factor essential for eye development. A previous study has shown that lens development arrests at, or just prior to, lens placode formation in Lhx2 germline mutant animals. FGF15, BMP4, BMP7 and phosphorylated SMAD1, SMAD5 and SMAD8 (pSMAD1/5/8), an indicator of BMP signaling, are all downregulated in the Lhx2+/− optic neuroepithelium and lens-forming region of the surface ectoderm. Restoration of BMP4 and BMP7 led to upregulation of FGF15 in the optic neuroepithelium and induction of the lens placode marker Sox2 in the surface ectoderm of Lhx2−/− animals. But BMP treatment does not rescue the morphology of Lhx2−/− mutant optic vesicle or surface ectoderm. These findings suggest that BMP signaling partially mediates Lhx2-dependent regulation of optic vesicle and lens vesicle induction (Yun et al., 2009). However, whether Lhx2 is required for lens development beyond lens placode formation is not known.

We recently observed that selective deletion of Lhx2 in neuroretina after lens placode induction led to severe microphthalmia and a near-total loss of the expression of many retinal progenitor-specific genes (Roy et al., 2013). Interestingly, we also noticed severe disruptions of lens development in these animals. This suggested that Lhx2 may regulate the expression of neuroretina-derived diffusible factors that are necessary for lens development beyond the lens placode stage. To address this possibility, we set out to identify candidate factors whose expression was disrupted in Lhx2-deficient neuroretina, and to determine whether restoring the activity of these factors was sufficient to rescue lens cell proliferation and fiber differentiation.
RESULTS
Selective deletion of Lhx2 in embryonic neuroretina disrupts lens development

We observed that selective deletion of Lhx2 in neuroretina using Chx10-Cre (Vix2-Cre) resulted in severe microphthalmia in adult mice (Fig. 1A,B). Close examination of Chx10-Cre;Lhx2lox/lox eye sections revealed severe defects in lens development at embryonic day (E)13.5, as well as postnatal day (P0.5) (Fig. 1C-F). The lens was either missing altogether, or detectable as only a vestigial lumen rudiment (Fig. 1D, F). Lens size was reduced to a substantially greater extent (740-fold smaller) than retinal size (7.3-fold smaller) at P0.5 (Fig. 1G). Analysis of Chx10-Cre;Ai9Rosa26Sl-tdTomato mice confirmed previous reports (Rowan and Cepko, 2004) that this Cre line is selectively active in neuroretina and not active in other ocular tissues, including lens (Fig. S1A-D). Immunostaining for Lhx2 showed selective deletion of Lhx2 in neuroretina starting at E10.5 with the expression almost entirely lost in neuroretina by E11.5 (Fig. S1E-L). Even though some Lhx2 staining was observed in both control and mutant lenses, signals were never detected in the nuclei of lens cells. Moreover, Lhx2 mRNA was also never detected in either control or Chx10-Cre;Lhx2lox/lox lenses (insets in Fig. 2B-G). As previous work has shown that Lhx2 expression is absent in the lens at all stages of ocular development (Hägglund et al., 2011), we conclude that the Lhx2 signal observed in the lens here represents non-specific background staining. Furthermore, as the Cre is active only in the neuroretina of our animals, we conclude that the disruption of lens development observed in our Chx10-Cre;Lhx2lox/lox animals must have resulted from disrupted signaling by secreted retinal-derived factors.

FGF signaling is downregulated in Lhx2-deficient eyes

To identify candidate retinal-derived factors whose expression was altered in Lhx2-deficient neuroretina, we analyzed previously reported microarray data obtained from retinas of control and Chx10-Cre;Lhx2lox/lox mice at E13.5 (Roy et al., 2013; Accession number GSE87707). These data indicated that multiple different FGFs and FGF-regulated genes showed altered expression in Chx10-Cre;Lhx2lox/lox retina. Fgf3, Fgf9 and Fgf15 – all of which have previously been reported to be expressed in embryonic neuroretinal progenitors (Colvin et al., 1999; Kurose et al., 2004; Wilkinson et al., 1989) – were dramatically downregulated (~5.0-, ~3.2- and ~11.9-fold, respectively). The FGF-regulated genes En5 and Spry2 were also downregulated (~11.4- and ~2.2-fold, respectively), whereas the FGF receptor gene Fgfr3 was upregulated (3.1-fold). Loss of Fgf3, Fgf9 and Fgf15 expression in Chx10-Cre;Lhx2lox/lox retinas at E13.5 was confirmed using qRT-PCR and in situ hybridization analysis (Fig. 2). In situ hybridization analysis at E13.5 showed a reduction in the expression of the FGF-regulated genes En5, En1 and Spry2 in the lenses of Chx10-Cre;Lhx2lox/lox mice (Fig. 3I,K,M,O,Q,S). Taken together, these data suggested that loss of expression of multiple different FGF genes in Chx10-Cre;Lhx2lox/lox retinas resulted in a global loss of FGF signaling in the lenses.

Forced expression of FGF10 in neuroretina induces FGF-regulated genes in the lens

As FGFs typically signal through multiple different FGF receptors (Ornitz and Itoh, 2015), and previous work had shown considerable redundancy in FGF receptor action in control of lens fiber development (Lovicu and Overbeek, 1998; Zhao et al., 2008), we hypothesized that selectively activating the expression of any individual FGF in Lhx2-deficient retina might rescue lens development. We accomplished this by using pMes-Fgf10 transgenic mice, a previously described line in which Cre-dependent excision of a transcriptional stop cassette leads to expression of full-length Fgf10 under the control of the chick β-actin promoter (Song et al., 2013). Using this line, we aimed to induce Fgf10 expression selectively in neuroretina of both control and Chx10-Cre;Lhx2lox/lox mice (Fig. 3B,D). We confirmed that Fgf10 expression could be robustly and selectively induced in neuroretina of E13.5 Chx10-Cre;Lhx2lox/lox;pMes-Fgf10 mice as well as Chx10-Cre;Lhx2lox/lox;pMes-Fgf10 mice (Fig. S2A; Fig. 3F,H). Furthermore, we observed that the FGF target genes En5, En1 and Spry2 were substantially upregulated in lenses of Chx10-Cre;Lhx2lox/lox;pMes-Fgf10 mice relative to Chx10-Cre;Lhx2lox/lox controls (Fig. 3). Strikingly, we also observed robust induction of these same FGF target genes in the lenses of Chx10-Cre;Lhx2lox/lox;pMes-Fgf10 mice relative to Chx10-Cre;
expression levels in Data were analyzed using unpaired two-tailed t-test; n=3; ***P<0.001; ****P<0.0001. (B-G) In situ hybridization of Fgf3, Fgf9 and Fgf15 mRNA expression levels in Chx10-Cre;Lhx2lox/lox retinas compared with Chx10-Cre;Lhx2lox/lox controls at E13.5. Data represent mean normalized to Gapdh values±s.e.m. (C,E,G) eyes at E15.5. (C,E) Dotted red circles mark the lenses shown in right-hand insets. (B-G) Lens Lhx2 expression in adjacent sections is shown in right-hand insets. Lhx2 expression in RPE is maintained and outlines the neural retina in unpigmented animals (right-hand insets in C,E). Scale bar: 100 µm.

Retinal overexpression of Fgf10 leads to the persistence of lens stalk
Overexpression of Fgf10 on a control background did not lead to any gross defects in lens development. The morphology and marker expression of Chx10-Cre;Lhx2lox/pMes-Fgf10 lenses were comparable with those of Chx10-Cre;Lhx2lox/+ controls (Figs 4, S5). However, Chx10-Cre;Lhx2lox/+;pMes-Fgf10 lenses were slightly but significantly larger than control lenses at all time points examined (Fig. S2B). In addition, the anterior pole of the lens remained tethered to the surface ectoderm (Fig. S4B,D,F,H) and this persistent lens stalk led to the development of corneal opacification in some adult mice (Fig. S4J).

Forced expression of FGF10 in Lhx2-deficient neuroretina rescues lens fiber differentiation
To investigate whether restoration of retinal FGF expression was able to rescue lens development in Lhx2 knockout animals, we conducted a detailed characterization of the expression of lens markers at multiple time points. At E11.5, prior to the formation of distinct lens fiber cells, no clear difference in lens size or lens fiber differentiation became more prominent as development proceeded. By E15.5, Chx10-Cre;Lhx2lox/pMes-Fgf10 mice showed well-defined lenses filled with β-crystallin-positive lens fiber cells (Fig. 4P). By P0.5, more than half of all Chx10-Cre;Lhx2lox/pMes-Fgf10 mice no longer had a visible lens rudiment. When a lens rudiment was detectable, it was extremely small (Fig. 4W; Fig. S2B). However, all Chx10-Cre;Lhx2lox/pMes-Fgf10 mice examined at P0.5 showed well-defined, though often small, lenses that expressed Prox1 and β-crystallin. Comparisons of eyes for the presence or absence of lenses in Chx10-Cre;Lhx2lox/pMes-Fgf10 and Chx10-Cre;Lhx2lox/pMes-Fgf10 mice at P0.5 indicated that the maintenance of lenses from FGF10 overexpression in Chx10-Cre;Lhx2lox/pMes-Fgf10 mice was statistically significant (Fig. S2C). Interestingly, at no point did overexpression of Fgf10 rescue defects in neuroretinal size (Fig. S2D). Expression of β-crystallin in lens epithelial cells through P0.5 was observed in lens epithelial cells of both Chx10-Cre;Lhx2lox/pMes-Fgf10 and Chx10-Cre;Lhx2lox/pMes-Fgf10 mice (Fig. 4G,H,K,P,T,X). By contrast, β-crystallin expression was restricted to lens fiber cells in Chx10-Cre;Lhx2lox/+ controls and Chx10-Cre;Lhx2lox/+;pMes-Fgf10 by E12.5 (Fig. 4E,F).

To further characterize the rescue of lens development observed following neuroretinal-specific overexpression of Fgf10 in Lhx2-deficient mice, we analyzed the expression of additional lens markers, a selective marker for lens epithelial cells, E-cadherin, and two markers for lens fiber cells, N-cadherin and Aquaporin-0 (originally known as main intrinsic polypeptide, MIP) (Xu et al., 2002; Yancey et al., 1988). We observed that Chx10-Cre;Lhx2lox/+ animals initially formed E-cadherin-positive lens vesicles but failed to differentiate to form lens fiber cells that are positive for N-cadherin or Aquaporin-0 at E12.5 (Fig. S5G; Fig. 5G). As development progressed, these E-cadherin-positive lens vesicles became smaller and were no longer detectable in most animals by E18.5 (Fig. S5S). By contrast, Chx10-Cre;Lhx2lox/+;pMes-Fgf10 mice continued to express E-cadherin robustly at E13.5, and began to show a progressive increase in N-cadherin expression from this...
point through P0.5 (Fig. S5L,P,T,X). We also observed that by E15.5, Chx10-Cre;Lhx2lox/lox;pMes-Fgf10 mice showed robust expression of Aquaporin-0, a mature lens fiber cell marker, although the onset of Aquaporin-0 expression was delayed relative to controls (Fig. 5). However, the Chx10-Cre;Lhx2lox/lox; pMes-Fgf10 rescue lens failed to develop normal lens polarity. E-cadherin and Prox1-positive cells persisted in the posterior pole of the lens at all time points examined (Figs 4, 5; Fig. S5).

Fig. 3. Cre-mediated induction of Fgf10 expression in Chx10-Cre;Lhx2lox/lox;pMes-Fgf10 retinas restored expression of FGF-regulated genes in lens. (A-D) Mouse genetics and anticipated retinal-derived FGF expression for (A) Chx10-Cre;Lhx2lox/+, (B) Chx10-Cre;Lhx2lox/++;pMes-Fgf10, (C) Chx10-Cre;Lhx2lox/lox, and (D) Chx10-Cre;Lhx2lox/lox;pMes-Fgf10 eyes. (E-T) In situ hybridization demonstrates Fgf10 induction in neuroretina (E-H), and induction of expression of FGF-regulated genes, including Etv5 (I-L), Etv1 (M-P) and Spry2 (Q-T). Right-hand insets indicate Lhx2 expression in adjacent sections. Lhx2 expression in RPE is maintained and outlines the neural retina in unpigmented animals (insets in G,K,O,S,H,T). Dotted red circles mark the lenses shown at higher magnification in the left-hand insets (G-T). Scale bar: 100 µm.

Fig. 4. Overexpression of Fgf10 in Chx10-Cre;Lhx2lox/lox;pMes-Fgf10 animals rescued lens fiber development. Developmental time-course of immunohistochemical staining for Prox1 (green) and β-crystallin (red) in lenses at E11.5 (A-D), E12.5 (E-H), E13.5 (I-L), E15.5 (M-P), E18.5 (Q-T) and P0.5 (U-X) of Chx10-Cre;Lhx2lox/+(A,E,I,M,Q,U), Chx10-Cre;Lhx2lox/++;pMes-Fgf10 (B,F,J,N,R,V), Chx10-Cre;Lhx2lox/lox (C,G,K,O,S,W) and Chx10-Cre;Lhx2lox/lox; pMes-Fgf10 (D,H,L,P,T,X) animals . Nuclei are counterstained with DAPI (blue). Scale bars: 100 µm.
FGF10 overexpression in Lhx2-deficient neuroretina promotes lens cell proliferation and survival

Exogenous FGFs have been reported to both promote the proliferation and survival of lens epithelial cells, in addition to driving lens fiber cell differentiation. To determine whether lens epithelial cell proliferation and survival were altered by overexpression of Fgf10, we examined the expression of Ki67 (Mki67 – Mouse Genome Informatics) and activated caspase 3 (c-caspase 3), respectively, at E12.5 (Fig. 6). We observed a significant increase in lens epithelial cell proliferation in Chx10-Cre;Lhx2lox/lox;pMes-Fgf10 relative to Chx10-Cre;Lhx2lox/lox mutant lens (Fig. 6E). Cell death was increased in both Chx10-Cre;Lhx2lox/lox mutants and Chx10-Cre;Lhx2lox/lox;pMes-Fgf10 rescue mice relative to the Chx10-Cre;Lhx2lox/lox controls, but was decreased in rescue mice relative to mutants (Fig. 6F). We thus conclude that overexpression of Fgf10 in Lhx2-deficient neuroretina promotes lens cell proliferation, survival and differentiation.

Loss of Lhx2 expression in neuroretina also disrupts BMP signaling in the lens

A previous study has shown that Lhx2 regulates lens placode formation through BMP signaling (Yun et al., 2009), whereas other studies have shown an essential role of BMP signaling in lens induction as well as fiber differentiation (Boswell et al., 2008; Furuta and Hogan, 1998; Jarrin et al., 2012; Muruli et al., 2005; Wawersik et al., 1999). To investigate whether BMP signaling was altered in either Lhx2-deficient or Fgf10-overexpressing eyes, we performed in situ hybridization for Bmp4 and Bmp7, and immunostaining for phosphorylated SMAD1, SMAD5 and SMAD9 (pSmad1/5/9) at E13.5 (Fig. 7). In both Chx10-Cre;Lhx2lox/+ and Chx10-Cre;Lhx2lox/lox;pMes-Fgf10 controls, broad Bmp4 expression was observed in neuroretina and RPE, and in lens epithelial and differentiating fiber cells, whereas more restricted Bmp7 expression was detected in a subset of neuroretinal cells, RPE and lens epithelial cells (Fig. 7A,B,E,F). Loss of neuroretinal Lhx2 expression led to a dramatic reduction of Bmp4 expression in Chx10-Cre;Lhx2lox/lox mutant lens, whereas forced Fgf10 expression partially rescued Bmp4 expression in Chx10-Cre;Lhx2lox/lox;pMes-Fgf10 neuroretina and lens (Fig. 7C,D). Bmp7 expression was not altered in both Chx10-Cre;Lhx2lox/lox mutants and Chx10-Cre;Lhx2lox/lox;pMes-Fgf10 rescue mice in agreement with a previous study by Hägglund et al. (2011) (Fig. 7G,H). Robust pSmad1/5/9 expression was observed in the neuroretina of both Chx10-Cre;Lhx2lox/lox and Chx10-Cre;Lhx2lox/lox;pMes-Fgf10 controls in the dorsoventral gradient. The expression was also observed in the nuclei of differentiating lens fiber cells (Fig. 7M,N). By contrast, even though some pSmad1/5/9 staining was observed the neuroretina of both Chx10-Cre;Lhx2lox/lox and Chx10-Cre;Lhx2lox/lox;pMes-Fgf10 animals, pSmad1/5/9 expression was not detected in the lens of both lines (Fig. 7O,P). We conclude that loss of neuroretinal Lhx2 disrupts Bmp4 expression and severely reduces BMP signaling in the lens.

DISCUSSION

In this study, we showed that Lhx2 regulated the expression of Bmp4 and multiple FGF genes – Fgf3, Fgf9 and Fgf15 – in neuroretina during early stages of retinal neurogenesis. The downregulation of these retinal-derived FGFs in Lhx2 knockout animals contributed to severe defects in lens cell proliferation, survival and differentiation. Restoring FGF expression in neuroretina, however, led to a partial rescue of lens development defects, despite continued retinal

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Fig. 5. Chx10-Cre;Lhx2lox+/pMes-Fgf10 animals expressed the mature lens fiber marker Aquaporin-0. Developmental time-course of immunohistochemistry for Prox1 (green) and Aquaporin-0 (red) expression in lenses at E11.5 (A-D), E12.5 (E-H), E13.5 (I-L), E15.5 (M-P), E18.5 (Q-T) and P0.5 (U-X) of Chx10-Cre;Lhx2lox+/pMes-Fgf10 (B,F,J,N,R,V), Chx10-Cre;Lhx2lox+/pMes-Fgf10 (C,G,K,O,S,W) and Chx10-Cre;Lhx2lox+/pMes-Fgf10 (D,H,L,P,T,X) animals. Nuclei are counterstained with DAPI (blue). Scale bars: in N, 100 µm for M,N; in L, 100 µm for A-L; in P, 100 µm for O,P; in X, 100 µm for Q-X.
development defects. These findings demonstrate the central role of Lhx2 in regulating lens development, and strongly suggest that FGFs do indeed constitute a major component of the long-sought retinal-derived signals that control lens maturation.

Previous work has shown that Lhx2 expression in the optic neuroepithelium is essential for lens placode specification (Yun et al., 2009). Our findings demonstrate the continued requirement of Lhx2 in the neuroretina past lens vesicle formation in regulating lens cell proliferation, survival and differentiation. The present study examined later stages of lens development than this earlier work, and identified a number of key differences in the relative contribution of BMP and FGF signaling. The previous study showed that loss of Lhx2 from optic vesicle severely disrupted Bmp4/7 expression, but only modestly reduced FGF signaling, as evidenced by continued expression of Fgf8, Etv5 (Erm) and phosphorylated ERK1/2 (pERK) in the optic vesicle and surface ectoderm. This same study showed that adding BMP7 or the combination of BMP4 and BMP7 to Lhx2-/- head cultures was sufficient to induce Sox2 expression in surface ectoderm (Yun et al., 2009). Other studies have also demonstrated an essential role for BMP4/7 signaling in early stages of lens development (Furuta and Hogan, 1998; Murali et al., 2005; Pandit et al., 2015; Wawersik et al., 1999). By contrast, our study indicated FGF signaling as perhaps the major mediator in Lhx2 regulation of lens maturation. Loss of neuroretinal Lhx2 led to reduced retinal Bmp4 expression, and severely disrupted BMP signaling in the lens; these changes were not rescued by overexpression of Fgf10 (Fig. 7). Nonetheless, lens fiber development was substantially, though not completely, rescued, suggesting that this process is at least partially independent of BMP signaling. This difference in the relative contribution of FGF and BMP signaling may reflect differences in the roles of neuroretina-derived BMP and FGF signaling during earlier and later stages of lens development. However, the lack of complete rescue may thus result from the well-documented synergistic role of BMP and FGF signaling in promoting lens fiber differentiation (Boswell et al., 2008; Boswell and Musil, 2015; Jarrin et al., 2012). The extent to which restoring BMP signaling can rescue lens development in Lhx2-deficient mice awaits further investigation.

Our finding that Lhx2 regulates neuroretinal FGFs is supported by a previous study from our lab that showed direct binding of Lhx2 to cis-regulatory regions of Fgf15 in neonatal mouse retina (de Melo et al., 2016). Although neuroretina-specific loss of function of other retinal progenitor-expressed transcription factors disrupts retinal progenitor cell proliferation and differentiation, and leads to microphthalmia (Burmeister et al., 1996; Marquardt et al., 2001; Taranova et al., 2006), only loss of Lhx2 significantly disrupts lens development. It is likely that these other mutant lines maintain neuroretinal expression of one or more FGFs, leading to normal lens development even in the face of a hypoplastic neuroretina.

Previous cell culture-based studies demonstrate that FGF signaling regulates multiple aspects of lens cell development and maturation (McAvoY and Chamberlain, 1989; Schulz et al., 1993). This study provides in vivo evidence for the importance of FGF
signaling in lens cell proliferation, survival and differentiation. Our findings identify FGF3, FGF9 and FGF15 as the three neuroretinal-derived FGFs that regulate lens development. Previous studies show that FGF3 can activate FGFR2b and FGFR1b; FGF9 can activate FGFR3c, FGFR2c, FGFR1c, FGFR3b and FGFR4; and FGF15 can activate FGFR1c, FGFR2c, FGFR3c and FGFR4 (Ornitz et al., 1996; Zhang et al., 2006). Neuroretinal expression of these three different FGFs, each of which can activate different subsets of FGF receptors, likely accounts for the observation that only Fgfr1/Fgfr2/Fgfr3 triple mutants produced severe defects in lens development (Zhao et al., 2008). This also likely accounts for the failure thus far to observe defects in lens fiber development in individual or pairwise combinations of targeted Fgf gene mutations (Robinson, 2006). Our ability to rescue lens fiber development by overexpression of Fgf10, which is expressed at only low levels in developing retina, agrees with previous findings of redundant roles for FGF3 and FGF10 in regulating cardiovascular and inner ear development (Alvarez et al., 2003; Urness et al., 2011).

Retinal overexpression of Fgf10 in animals that are not deficient for Lhx2 leads to downregulation of BMP signaling in the neuroretina and lens. (A-L) In situ hybridization of Bmp4 (A-D), Bmp7 (E-H) and Lhx2 (I-L) mRNA expression levels in Chx10-Cre;Lhx2lox+/+(A,E,I), Chx10-Cre;Lhx2lox-/+(B,F,J), Chx10-Cre;Lhx2lox+/pMes-Fgf10(K) and Chx10-Cre;Lhx2lox-/pMes-Fgf10 (D,H,L) eyes at E13.5. Dotted red circles mark the lenses (C,D,G,H,L). (M-P) Immunohistochemical staining for Pax6 (green) and pSmad1/5/9 (red) in lenses of Chx10-Cre;Lhx2lox+/-(M), Chx10-Cre;Lhx2lox+/pMes-Fgf10 (N), Chx10-Cre;Lhx2lox+/pMes-Fgf10 (O) and Chx10-Cre;Lhx2lox-/pMes-Fgf10 animals (P) at E13.5. Nuclei are counterstained with DAPI (blue). The images of pSmad1/5/9 staining in single channel and at higher magnification are included underneath. Dotted white circles mark the lenses (O,P). Scale bars: 100 µm.
overexpression of Fgf10 in neuroretina did not rescue BMP signaling in the lens in Lhx2-deficient eyes. A schematic figure summarizing our results is shown in Fig. 6G.

MATERIALS AND METHODS

Animals

Lhx2loxloxplox (Mangale et al., 2008) mice were obtained from Dr Edwin Monuki (University of California, Irvine, CA, USA). Chx10-Cre (Rowan and Cepko, 2004) mice were a gift from Dr Connie Cepko (Harvard, Boston, MA, USA). pMes-Fgf10 (Song et al., 2013) mice were generously provided by Dr Yang Chai (University of Southern California, Los Angeles, CA, USA). A9 (R26-CAG-lox-stop-lox-sMMP20) (Madin et al., 2010) mice were a gift from Dr Xinzhong Dong (Johns Hopkins, Baltimore, MD, USA). Chx10-Cre; Lhx2lox/lox, Chx10-Cre;Lhx2lox/lox-tdTomato, pMes-Fgf10, Chx10-Cre;Lhx2loxloxplox, Chx10-Cre;Lhx2loxloxplox;pMes-Fgf10 and Chx10-Cre;A9 mice were generously provided by Dr Yang Chai (University of Southern California, Los Angeles, CA, USA). Anti-Pax6 (1:200; AB_528427, DSHB), rabbit anti-β-crystallin (1:500; a generous gift from Dr Jeremy Nathans, Johns Hopkins University, Baltimore, MD, USA), mouse anti-Prox1 (1:200; MAB5654, Millipore), rabbit anti-E-Cadherin (1:200; 3195, Cell Signaling), mouse anti-N-Cadherin (1:200; 333900, Novex), rabbit anti-Aquaporin 0 (1:200; AB3071, Millipore), rabbit anti-c-caspase 3 (1:200; 9664, Cell Signaling), mouse anti-Ki67 (1:200; 550609 (Clone B56), BD Pharmingen), rat anti-RFP (1:1000; ABIN334653, Chromotek) and rabbit anti-pSmad1/5/9 (1:1300; 18920, Cell Signaling). Secondary antibodies used were as follows: FITC-conjugated donkey anti-mouse IgG (1:500; 715-095-150, Jackson ImmunoResearch), Alexa Fluor 488-conjugated goat anti-mouse IgG, Fcy subclass 1 specific (1:500; 115-545-205, Jackson ImmunoResearch), FITC-conjugated goat anti-mouse IgG (subclasses 1+2a+3b+3c+3d; 1:500; 115-095-164, Jackson ImmunoResearch), Alexa Fluor 594-conjugated donkey anti-rabbit IgG (1:500; 711-585-152, Jackson ImmunoResearch) and Alexa Fluor 555-conjugated goat anti-rat IgG (1:500; A-21434, Invitrogen). Hematoxylin and Eosin sections were imaged using a Zeiss Axioskop 2 Mot Plus microscope. Immunohistochemical sections were imaged on a Zeiss Meta 510 LSM confocal microscope.

Quantitative RT-PCR

The cDNAs of E13.5 dissected retinas were synthesized from total RNAs using SuperScript III First-Strand Synthesis System (Invitrogen). Quantitative RT-PCR was performed using Brilliant III Ultra-Fast SYBR Green PCR Master Mix (Agilent) on a 7300 Real-Time PCR System (Applied Biosystems) following the manufacturers’ recommended protocols. Primer sets for genes examined were as follows: fibroblast growth factor 3 (Fgf3) forward, TGGCCGTACCAATGACACC and Fgf3 reverse, CACCAGCAATCTACAGGAT; Fgf9 forward, ATGGCTCCCCATAGGAGT and Fgf9 reverse, TGGCCGTAAGAATCCCCCTT; Fgf10 forward, GAGCGGCAATGTTGCCGCA; Fgf10 reverse, GTGTTGCAATGCTTACA; Fgf15 forward, GCGCCGTACCAATGACACC and Fgf15 reverse, CTTGTTGCAATGCTTACA; Gapdh forward, AGGTCTGCGGCTACA; Gapdh reverse, GTGGAAGAGGATGACGT and Gapdh fragment (1:500; 115-095-164, Jackson ImmunoResearch). Lethal retinal sizes were measured using ImageJ (n>4 for E11.5; n≥9 for E12.5; n≥6 for E13.5; n≥7 for P0.5) All cell counts and measurements were repeated blindly by a student researcher.

Statistical analysis

Data are presented as mean±s.e.m. Statistical analysis was performed using GraphPad Prism Software. Comparisons of means between two groups were evaluated using Student’s t-test. Comparison of means among multiple groups was performed by one-way ANOVA followed by Tukey’s and Sidak’s multiple comparison tests. Categorical data was analyzed using Fisher’s exact test. 

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Competing interests

The authors declare no competing or financial interests.

Author contributions


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Supplementary information

Supplementary information available online at http://dev.biologists.org/lookupdoi/10.1242/dev.137760.supplemental.

Histology and immunohistochemistry

Haematoxylin and Eosin staining and fluorescence immunohistochemistry were performed as previously described (de Melo et al., 2012; Roy et al., 2013). The following primary antibodies were used: rabbit anti-Lhx2 (1:1000; generated for our lab by Covance (de Melo et al., 2012)), mouse anti-Pax6 (1:200; AB_528427, DSHB), rabbit anti-β-crystallin (1:500; a generous gift from Dr Jeremy Nathans, Johns Hopkins University, Baltimore, MD, USA), mouse anti-Prox1 (1:200; MAB5654, Millipore), rabbit anti-E-Cadherin (1:200; 3195, Cell Signaling), mouse anti-N-Cadherin (1:200; 333900, Novex), rabbit anti-Aquaporin 0 (1:200; AB3071, Millipore), rabbit anti-c-caspase 3 (1:200; 9664, Cell Signaling), mouse anti-Ki67 (1:200; 550609 (Clone B56), BD Pharmingen), rat anti-RFP (1:1000; ABIN334653, Chromotek) and rabbit anti-pSmad1/5/9 (1:1300; 18920, Cell Signaling). Secondary antibodies used were as follows: FITC-conjugated donkey anti-mouse IgG (1:500; 715-095-150, Jackson ImmunoResearch), Alexa Fluor 488-conjugated goat anti-mouse IgG, Fcγ subclass 1 specific (1:500; 115-545-205, Jackson ImmunoResearch), FITC-conjugated goat anti-mouse IgG (subclasses 1+2a+3b+3c+3d; 1:500; 115-095-164, Jackson ImmunoResearch), Alexa Fluor 594-conjugated donkey anti-rabbit IgG (1:500; 711-585-152, Jackson ImmunoResearch) and Alexa Fluor 555-conjugated goat anti-rat IgG (1:500; A-21434, Invitrogen). Hematoxylin and Eosin sections were imaged using a Zeiss Axioskop 2 Mot Plus microscope. Immunohistochemical sections were imaged on a Zeiss Meta 510 LSM confocal microscope.

Cell quantification

The total number of DAPI+, Ki67- and c-caspase 3-positive cells on each lens section was quantified and the percentage of Ki67- and cleaved caspase 3-positive cells was calculated (n=6 for Chx10-Cre;Lhx2lox/lox and Chx10-Cre;Lhx2lox/lox-tdTomato; n=18 for Chx10-Cre;Lhx2lox/lox and Chx10-Cre;Lhx2lox/lox+pMes-Fgf10). Lens and retinal sizes were measured using ImageJ (n≥4 for E11.5; n≥9 for E12.5; n≥6 for E13.5; n≥7 for P0.5) All cell counts and measurements were repeated blindly by a student researcher.

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References


