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Pitx3 controls multiple aspects of lens development

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Abstract

The transcription factor *Pitx3* is critical for lens formation. Deletions in the promoter of this gene cause abnormal lens development in the *aphakia* (*ak*) mouse mutant, which has only rudimentary lenses. In this study, we investigated the role of *Pitx3* in lens development and differentiation. We found that reduced expression of *Pitx3* leads to changes in the proliferation, differentiation and survival of lens cells. The genetic interactions between *Pitx3* and *Foxe3* were investigated, as these two transcription factors are expressed at the same time in lens development and their absence has similar consequences for lens development. We found no evidence that these two genes genetically interact.

In general, our study shows that the abnormal phenotype of the *ak* lenses is not due to just one molecular pathway, rather in the absence of *Pitx3* expression multiple aspects of lens development are disrupted.

Keywords

aphakia; *Foxe3*; *Pitx3*; lens

Introduction

During vertebrate lens development, the lens placode undergoes a carefully programmed morphogenetic process that leads to the formation of a lens with relatively undifferentiated, proliferative lens epithelial cells in the anterior of the lens, and highly differentiated, mitotically-inactive fiber cells in the posterior of the lens (for review see (McAvoy et al., 1999; Chow and Lang, 2001; Donner et al., 2006; Cvekl and Duncan, 2007; Medina-Martinez and Jamrich, 2007)). The proliferation and differentiation of the lens cells are controlled by several transcription factors. While the transcription factor Pax6 appears to be the key regulator of lens development, as mutations in this gene lead to eye malformations in humans (Ton et al., 1991; Jordan et al., 1992; Glaser et al., 1994; Hanson et al., 1994), mice and rats (Hill et al., 1991; Fujiwara et al., 1994), several other transcriptional regulators are also critical for lens formation. One of them is the homeodomain-containing factor *Pitx3*. *Pitx3* is expressed during early vertebrate lens development. A double deletion that eliminates the promoter region and a part of the coding region of this gene causes the abnormal lens phenotype in the mouse line *aphakia* (*ak*) (Semina et al., 2000; Rieger et al., 2001) and mutations in *PITX3* lead to the development of autosomal-dominant cataract in humans (Semina et al., 1998). In *ak* mice the lens begins to form, but its development is

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abnormal. Eventually, lens development arrests and the lens disappears (Varnum and Stevens, 1968; Zwaan, 1975; Zwaan and Kirkland, 1975; Grimm et al., 1998). Some aspects of lens development in the *aphakia* mutant are similar to the lens development in *Foxe3* null mice (Medina-Martinez et al., 2005). In both mutants the invaginating lens placode does not separate properly from the head surface ectoderm and the lens remains connected to the surface ectoderm with a stalk (Varnum and Stevens, 1968; Zwaan, 1975; Zwaan and Kirkland, 1975; Grimm et al., 1998; Blixt et al., 2000; Brownell et al., 2000; Medina-Martinez et al., 2005; Blixt et al., 2006; Medina-Martinez and Jamrich, 2007). *Foxe3* is a conserved forkhead transcription factor that is critical for lens development in several vertebrate species (for review see (Medina-Martinez and Jamrich, 2007)). Mutations in this gene, or its altered expression, cause abnormal lens development in mouse, humans and zebrafish (Blixt et al., 2000; Brownell et al., 2000; Semina et al., 2001; Ormestad et al., 2002; Medina-Martinez et al., 2005; Shi et al., 2006; Valleix et al., 2006; Medina-Martinez and Jamrich, 2007; Swindell et al., 2008).

Since the expression of these genes is very similar during early stages of lens development and mutations in these genes result in a similar lens phenotype, we investigated whether these two transcription factors are a part of the same regulatory cascade during mouse lens development. As these experiments provided no evidence for genetic interaction of these two genes, we have compared the expression of several diagnostic markers of lens development and differentiation in the *ak* and wild type lenses. Results of these experiments have provided a better understanding of the requirement for the *Pitx3* function in the mouse lens.

Results

Expression of *Foxe3* and *Pitx3* in wild type and *ak* embryos

Since previous reports indicated that the expression of *Pitx3* and *Foxe3* is similar during early lens development (Semina et al., 1997; Semina et al., 1998; Blixt et al., 2000; Brownell et al., 2000), and since mouse embryos with mutations in these two genes display similar lens phenotypes (Varnum and Stevens, 1968; Blixt et al., 2000; Brownell et al., 2000; Medina-Martinez et al., 2005; Medina-Martinez and Jamrich, 2007), we investigated the expression of *Pitx3* and *Foxe3* in more detail. To obtain better information about the temporo-spatial expression pattern of *Foxe3* and *Pitx3* during early lens development, we analyzed expression of these two genes by whole mount *in situ* hybridization. Using this method, *Foxe3* expression can be easily detected in the developing lens placode at E9.5 (Fig. 1A) (Blixt et al., 2000; Brownell et al., 2000). *Pitx3* expression can be detected by whole mount *in situ* hybridization few hours later, around E9.75 (Fig. 1B). Sections of these embryos show that at E9.75 *Pitx3* is expressed only in a small portion of the lens placode, while *Foxe3* is expressed in the entire lens placode (Fig. 1C, D). However, by the time the lens placode begins to invaginate, both *Foxe3* and *Pitx3* are expressed in the entire lens pit (Fig. 1E,F). Thus, *Foxe3* expression precedes *Pitx3* during the formation of the lens placode, but at later stages their expression appears to be identical.

Since *Foxe3* and *Pitx3* expression is similar in the developing lens, we evaluated whether the expression of these two genes depends on each other. For this purpose, we analyzed the expression of *Foxe3* in the *ak* mutant and the expression of *Pitx3* in the *Foxe3* null mutant by *in situ* hybridization. During normal lens development *Foxe3* is initially expressed in the entire lens vesicle. As lens development proceeds, *Foxe3* remains active in the anterior of the lens, in cells that will form the anterior lens epithelium (Blixt et al., 2000; Brownell et al., 2000). The differentiating lens fiber cells at the posterior of the lens cease to express *Foxe3*. *In situ* hybridizations on E11.5 wild type and *ak* embryos show an altered spatial distribution of *Foxe3* transcripts within the *ak* lens. While in the wild type lens the highest

levels of *Foxe3* expression are found in the anterior of the lens (Fig. 2A), which develops into the lens epithelium, in the *ak* lens most of the *Foxe3* transcripts are found in the posterior of the lens (Fig. 2B). Expression of *Foxe3* in the posterior of the *ak* lens indicates that the primary lens fiber cells in the *ak* embryo do not undergo proper differentiation. Lack of expression of *Foxe3* in the anterior compartment indicates that *Pitx3* activity is necessary for the maintenance of *Foxe3* expression in the anterior lens epithelium. Indeed, the significant downregulation of *Foxe3* expression in the *ak* mutant by E12.5 (Fig. 2C,D) suggests that *Pitx3* is necessary for the maintenance of the *Foxe3* expression. In contrast, *Pitx3* expression is relatively unaffected in the *Foxe3* null embryos (Fig. 2E-H), suggesting that *Foxe3* does not have an important role in the regulation of *Pitx3* transcription during early lens development.

Analysis of genetic interactions between *Foxe3* and *Pitx3*

Since some of the defects in the *Foxe3* and *Pitx3* mutants appear superficially similar, there was a possibility that these two genes genetically interact. For this reason, we compared lens development in *Foxe3* and *Pitx3* compound heterozygous and homozygous animals by interbreeding *Foxe3*^{+/-} heterozygous mice with *+ak* heterozygous mice. As previously reported (Medina-Martinez et al., 2005), the *Foxe3*^{+/-} heterozygous mice display no abnormal phenotype (Fig. 3A), whereas the lenses in *+ak* heterozygous animals are almost normal, occasionally displaying a stalk connecting the lens and the surface ectoderm (Fig. 3B). Lenses in *Foxe3*-deficient animals are smaller than normal and display a persistent connection with the surface ectoderm (Fig. 3C). Lenses in *ak/ak* animals are typically club-shaped and smaller than lenses in *Foxe3*-deficient animals. They also display a persistent connection to the surface ectoderm (Fig. 3D). Analysis of lens development in mice with different genotypic combinations of *Foxe3* and *Pitx3* revealed that in *Foxe3*^{+/-};*+ak* double heterozygous embryos (Fig. 3E), the morphology of the lens is similar to the lens in *ak* heterozygous animals (Fig. 3C). Thus, superposition of *Foxe3*^{+/-} heterozygosity does not exacerbate the *+ak* phenotype suggesting that there is no synergistic genetic interaction between *Foxe3* and *Pitx3*. We next evaluated the lens development in *Foxe3*^{+/-};*ak/ak*, *Foxe3*^{-/-};*+ak*, and *Foxe3*^{-/-};*ak/ak* compound mutants. As expected, the lens in the *Foxe3*^{-/-};*+ak* had the same morphology (Fig. 3F) as in the *Foxe3* null allele (Fig. 3C). The *Foxe3*^{+/-};*ak/ak* lenses (Fig. 3G) resemble those of the *ak* mutants (Fig. 3D). In the *Foxe3*^{-/-};*ak/ak* compound mutants (Fig. 3H), the lenses are similar to lenses in *ak/ak* animals (Fig. 3D). These experiments suggest that there is no synergistic interaction between *Foxe3* and *Pitx3* during lens development.

Expression of crystallins in *ak* embryos

To better understand the developmental state of lens cells in the *ak* mutant, we investigated the expression of crystallins. Crystallins are a family of chaperone proteins that are expressed at very high levels in the vertebrate lens (Wistow and Piatigorsky, 1987; Piatigorsky, 1998; Cvekl and Duncan, 2007). Expression of individual crystallin genes is characteristic of the state of differentiation of individual lens cells. During mouse lens development, α B-crystallin is the first crystallin gene activated. Its transcription starts already in the lens placode (Robinson and Overbeek, 1996; Cvekl and Duncan, 2007). It is followed by α A-crystallin, which is activated at the transition from the lens pit to lens vesicle (Robinson and Overbeek, 1996). The β and γ -crystallins are highly expressed in differentiating fiber cells. We have examined the expression of crystallins in the *ak* lenses using both *in situ* hybridizations and antibodies against different crystallins. *In situ* hybridization shows that at E12.5 α A-crystallin (*Crya1*) and α B-crystallin (*Crya2*) are transcribed in the wild type embryos at high levels (Fig. 4A and C), but their transcription is strongly reduced in the *ak* mutant (Fig. 4B and D). Using antibodies that recognize α A and α B-crystallins, we found that only a few cells are positive for these two proteins in the

aphakia lens (Fig.4F), while the entire lens labels with these antibodies in the wild type embryo (Fig. 4E). Expression of β A3/A1-crystallin (*Cryba1*) is also dramatically altered. In the wild type lens, this gene is strongly transcribed already at E12.5 (Fig. 4G). In the *ak* lens, the β A3/A1-crystallin is only weakly transcribed at E12.5 (Fig. 4H). Immunostaining using antibodies against β A-crystallins shows that in the wild type lens the distribution of β A-crystallins proteins is not uniform. Most of the β A-crystallins can be found in mature fibers (Fig. 4I). In the *ak* mutant, the β A-crystallins are almost undetectable (Fig. 4J). Finally, γ -crystallin transcripts, as well as the γ -crystallin proteins, are present primarily in the mature lens fiber cells of the wild type lens (Fig. 1K, M and O). Interestingly, in the *ak* mutant, relatively high levels of γ -crystallin transcripts can be detected (Fig. 4L, N), but the γ -crystallins proteins are practically undetectable when using antibodies against γ -crystallins (Fig. 4P). Furthermore, the γ -crystallin transcripts show an abnormal distribution within the *ak* lens. In contrast to the wild type lens, where the transcripts are present mostly in the lens fiber cells (Fig. 4M), in the *ak* lens, the γ -crystallin transcripts are more abundant in the anterior of the lens (Fig. 4N). This indicates that the differentiation process of lens cells is abnormal in the *ak* mutant, since all lens cells transcribe γ -crystallin mRNAs at relatively high levels and that the temporo-spatial regulation of γ -crystallin expression is abnormal. In addition, our results suggest that in *ak* lenses, the expression of γ -crystallins and maybe also β A-crystallins, is controlled at a post-transcriptional level.

Expression of key transcriptional regulators in *ak* embryos

Since the altered expression of crystallins was diagnostic of abnormal differentiation of lens cells, we investigated in the *ak* mutant the expression of transcription factors that were previously implicated in the regulation of lens development and differentiation. We evaluated expression of several regulatory genes including *c-maf*, *Maf-B*, *Sox1*, *Sox2*, *Six3*, *Prox1* and *Hsf-4* by *in situ* hybridization. At E12.5, we have observed only slight reduction in the intensity of expression of *c-maf* in the *ak* mutant when compared to the *+ak* embryos (Fig. 5A, B). Expression of *Maf-B* (Fig.5C, D), *Hsf4*, *Sox1* and *Sox2* (data not shown) in the *ak* lenses was not significantly altered with the caveat that the **area** of expression was smaller than in *+ak* lenses. The most dramatic reduction was observed in the expression of *Prox1* (Fig. 5E, F). *Prox1* is involved in the transition of lens epithelium to the lens fiber cells. *Prox1*-deficient lenses do not develop fiber cells and remain arrested at the lens vesicle stage (Wigle et al., 1999). *Prox1* promotes exit of cells from the cell cycle and *Prox1*-deficient cells have been shown to be less likely to stop dividing (Wigle et al., 1999; Dyer, 2003; Dyer et al., 2003).

Since expression of *Prox-1* was affected in *ak* mutants, we investigated whether the expression of the cell cycle regulator *p57^{KIP2}* is also affected. Expression of *p57^{KIP2}* is known to be controlled by *Prox-1* (Wigle et al., 1999) and reduction in *p57^{KIP2}* expression would indicate that cell cycle control in this mutant is affected. We found that *p57^{KIP2}* is not expressed in the *ak* lens (Fig. 5H), an observation that is consistent with the absence of *Prox1* expression in the *ak* lens.

Apoptosis and proliferation in *ak* embryos

To examine whether proliferation and apoptosis is affected in *ak* lenses, we analyzed cell proliferation by using antibodies against phosphohistone H3 (Wei et al., 1999). Phosphohistone H3 (pH3) is an indicator of cell proliferation, as histone H3 is phosphorylated during the M-phase of the cell cycle. We have examined the distribution of phosphohistone H3 in *+ak* and *ak* embryos ranging from E9.75 to E12.5. We found that at all stages there are fewer lens cells labeling in the *ak* mutant than in the *+ak* lenses. Already at E9.75 and E11.5 the presumptive lens placode and the lens vesicle shows fewer pH3 positive cells than the *+ak* counterparts (Fig. 6A-D). At E12.5, in *+ak* embryos the

antibodies against pH3 label many proliferating cells in the anterior lens epithelium (Fig. 6E, G). In the *ak* lenses, only occasional cells can be found displaying a positive reaction using these antibodies (Fig. 6F, H). These cells are typically not in the anterior of the lens, but rather in the posterior compartment. This indicates that in *ak* embryos cells in the anterior of the lens stopped proliferation. Only aberrant proliferation of cells in the posterior of the lens is present. A TUNEL assay demonstrates that at the same time as the proliferation of lens cells is reduced in the *ak* mutant, the cell death is increased when compared to the wild type lens (6I, J).

Discussion

Lack of Pitx3 activity causes reduced proliferation and increased apoptosis in the *ak* lenses

In this paper, we report studies of gene expression, proliferation and differentiation of lens cells in the *aphakia* mouse mutant (Semina et al., 2000). In this mutant, the activity of *Pitx3* is reduced by about 90% (Semina et al., 2000; Rieger et al., 2001). As a result of this reduced *Pitx3* activity, in the *ak* animals the lens is abnormally small and club-shaped. It remains connected to the surface ectoderm and eventually it disappears. In order to understand better the causes for the abnormal lens phenotype, we investigated apoptosis, proliferation and gene expression in the *ak* lenses. We found that the proliferation in the developing lens is reduced in the *ak* mutant as early as E9.75. The reduced proliferation is concomitant with the abnormal expression of the transcription factor *Foxe3*. Since the proliferation of lens cells is also affected in *Foxe3* mutants (Blixt et al., 2000; Brownell et al., 2000; Medina-Martinez et al., 2005), we investigated whether these two transcription factors cooperate in the regulation of proliferation and whether the reduced proliferation in the *ak* lens can be explained by the abnormal expression of *Foxe3*. Genetic analysis showed no evidence of synergistic interaction between these two genes, as the lenses in the compound heterozygous animals did not have a more severe phenotype than the lenses in *ak* heterozygous animals. A comparison of expression of *Pitx3* and *Foxe3* in wild type, *ak* and *Foxe3* mutants showed that *Foxe3* is activated in the lens placode before *Pitx3* and therefore *Pitx3* is unlikely to have a role in the activation of *Foxe3*. However, *Foxe3* expression in the *ak* mutant is not maintained past E12.5, suggesting that *Pitx3* is directly or indirectly required for the maintenance of *Foxe3* transcription. Since *Foxe3* was shown to play a role in the proliferation of the anterior epithelium (Blixt et al., 2000; Brownell et al., 2000; Medina-Martinez et al., 2005; Medina-Martinez and Jamrich, 2007), it is possible that the decreased proliferation of anterior lens cells in the *ak* mutant is due to the reduced activity of *Foxe3*. In contrast to the wild type lens, in the *ak* lens the *Foxe3* is no longer expressed in the anterior lens cells. Reduction in *Foxe3* expression is known to reduce the proliferation of anterior lens cells. Interestingly, *Foxe3* transcripts can be found in the posterior lens cells. This must be detrimental to the differentiating lens fiber cells since a persistent expression of *Foxe3* blocks cytoskeletal remodeling during the differentiation of lens fiber cells (Landgren et al., 2008). This block in cytoskeletal remodeling might explain while we never see true fiber cells in the *ak* lens. However, the reduction in *Foxe3* expression cannot account entirely for the lack of proliferation in *ak* lenses. This is because in *Foxe3*-deficient embryos, the proliferation of the lens epithelium is less affected than in the *ak* mutants, which have at least some *Foxe3* activity. However the reduction in proliferation in conjunction with the observed apoptosis of the lens cells is likely to be the cause of smaller lenses in the *ak* mutant.

Lack of Pitx3 activity causes abnormal differentiation in the *ak* lenses

While some differentiation of lens cells takes place in the *ak* mutant, expression of crystallins, markers of lens differentiation, is abnormal. In contrast to the wild type lens, the

expression of α and β -crystallins is rare in the posterior lens cells. γ -crystallins are transcribed, but they are not efficiently translated. Furthermore, their transcription is spatially dysregulated. While typically γ -crystallins transcripts are found in the differentiating lens fiber cells of the wild type lens, but not in the cells of the anterior lens epithelium, in the *ak* lens, high levels of γ -crystallins transcripts can be found in the anterior lens cells indicating that γ -crystallins are transcribed inappropriately early during lens cell differentiation. Our observations taken together with the reports that the lens cells that move posteriorly do not assume typical fiber shape (Varnum and Stevens, 1968; Zwaan, 1975; Malinina and Koniukhov, 1981; Semina et al., 1997; Grimm et al., 1998; Semina et al., 2000; Rieger et al., 2001) show that the differentiation of lens fiber cells is grossly abnormal in the *ak* mutants. The deregulation of crystallin expression might be important for the survival of lens cells. The reduction of expression of α A-crystallin and α B-crystallin might contribute to the observed apoptosis of lens cells in *ak* animals. α A/ α B-crystallins were shown to inhibit apoptosis, as lens fibers disintegrate in the α A/ α B-crystallin double knockout (Morozov and Wawrousek, 2006).

The downregulation of crystallin expression in the *ak* mutant might be due to the downregulation of *Prox1* expression. *Prox1* has been shown to regulate expression of crystallin genes (Cui et al., 2004; Chen et al., 2008). On the other hand, almost normal crystallin expression in the *Prox1*-deficient embryos argues against this possibility (Wigle et al., 1999).

In general, the development and differentiation of the lens in the *ak* mutants is strongly disrupted. *Pitx3* is clearly essential for the normal lens development and differentiation. In the *ak* mutant, the developing lens does not undergo a typical development in which an anterior lens epithelium is formed. The lens cells do not undergo a proper differentiation and fiber cells are not formed. The proliferation of lens cells is reduced and the apoptosis is increased. The temporo-spatial expression of genes diagnostic of early and late stages of lens development is abnormal. Based on these results we conclude that the abnormal phenotype of the *ak* lenses is not due to just one molecular pathway or a specific interaction, rather in the absence of *Pitx3* expression, the apoptosis, proliferation and differentiation of lens cells is severely disrupted.

Experimental Procedures

Mouse genetics

Foxe3-deficient mice were previously described (Medina-Martinez et al., 2005). The *ak* mice that were generously provided by Dr. Paul Overbeek (Semina et al., 2000; Varnum and Stevens, 1968) arose as a spontaneous mutation in the C57BL/6 genetic background, but our mutants have a mixed genetic background. In order to generate animals with different genotypic combinations of *Foxe3* and *Pitx3*, *Foxe3*^{+/-} heterozygous mice were backcrossed at least two generations with *+/ak* heterozygous mice. The *ak* mice were crossed to *Foxe3*^{-/-} (C57BL/6X 129 SvEv) mice. Double heterozygous animals (*Foxe3*^{+/-}/*+/ak*) were interbred to obtain the desired embryos. Three embryos were analyzed for each genotype.

Histology, in situ hybridization and nomenclature

For histological analysis, embryos were fixed in 10% formalin, dehydrated in an ethanol series. They were embedded in paraffin, sectioned and stained with hematoxylin and eosin. Whole mount in situ hybridizations were performed according to standard procedures (Wilkinson, 1992).

In this paper, we refer as α A-crystallin and α B-crystallin to the mouse genes *Crya1* and *Crya2*. β A3/A1-crystallin refers to the mouse gene *Cryba1*.

Immunohistochemistry

Embryos were fixed in 10% formalin overnight, dehydrated, embedded in paraffin and cut into 8mm serial sections. Sections were treated with different anti-crystallin antibodies (Yoshimoto et al., 2005) or with a phosphohistone H3 antibody (Hendzel et al., 1997) from Upstate Biotechnology.

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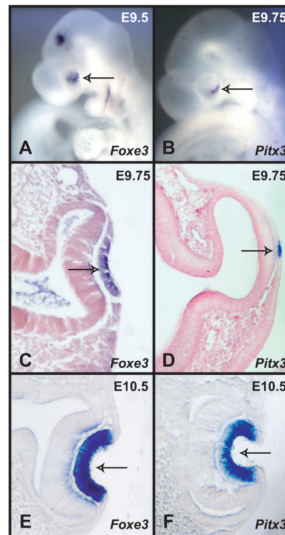


Figure 1.

In situ hybridizations of *Foxe3* and *Pitx3* probes to wild type mouse embryos. (A) Whole mount *in situ* hybridization of a *Foxe3* probe to an E9.5 embryo showing expression in the lens placode (arrow). (B) Whole mount *in situ* hybridization of *Pitx3* probe to an E9.75 embryo showing expression in a part of the lens placode (arrow). (C) A section showing the expression of *Foxe3* in the lens placode (arrow). (D) A section through the embryo in B showing expression of *Pitx3* in a few cells of the lens placode (arrow). (E) A section of an E10.5 embryo hybridized with *Foxe3* probe showing strong expression in the lens pit (arrow). (F) A section of an E10.5 embryo hybridized with *Pitx3* probe showing strong expression in most of the cells of the lens pit (arrow).

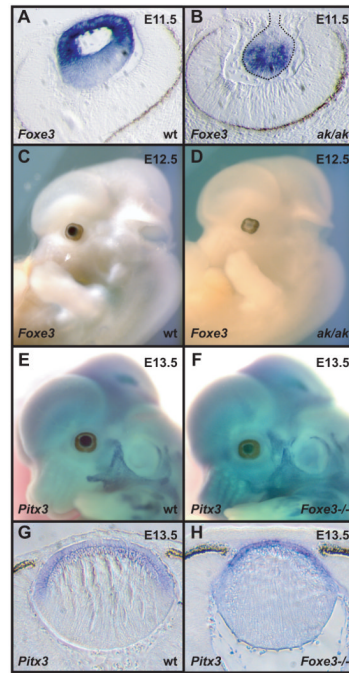


Figure 2.

In situ hybridizations of *Foxe3* and *Pitx3* to wild type, *ak* and *Foxe3*-deficient embryos. (A) A section of an E11.5 wild type embryo hybridized with *Foxe3* probe showing strong labeling in the developing lens. (B) A section of an E11.5 *ak* embryo hybridized with a *Foxe3* probe showing expression of this gene in the abnormal lens. (C) Whole mount *in situ* hybridization of *Foxe3* probe to an E12.5 wild type embryo showing expression in the lens. (D) Whole mount *in situ* hybridization of *Foxe3* probe to an E12.5 *ak* embryo showing dramatically reduced expression of this gene in the lens. (E) Whole mount *in situ* hybridization of *Pitx3* probe to an E13.5 wild type embryo showing expression of this gene in the lens. (F) Whole mount *in situ* hybridization of *Pitx3* probe to an E13.5 *Foxe3*-deficient embryo showing only slightly reduced levels of expression of this gene in the lens. (G) A section through the embryo in (E) showing the expression of *Pitx3* in wild type embryo. (H) A section through the embryo in (F) showing the expression of *Pitx3* in *Foxe3*^{-/-} embryo.

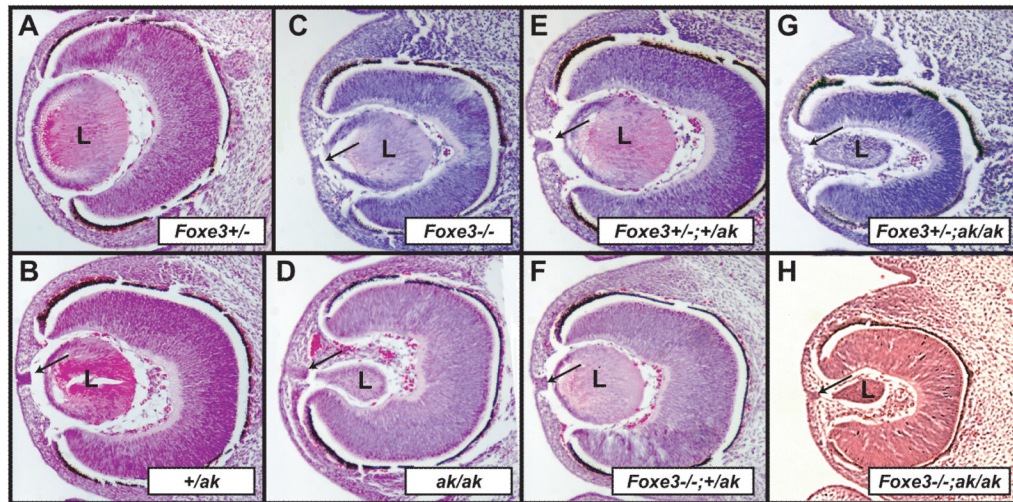


Figure 3.

Morphological analysis of genetic interactions between *Foxe3* and *Pitx3*. (A-H) Hematoxylin and eosin (H&E) stained sections of a *Foxe3*^{+/-} eye (A), an eye from *+/ak* embryo (B), an eye from a *Foxe3*^{-/-} embryo (C), an eye from *ak/ak* embryo (D), an eye of an embryo heterozygous for *Foxe3* and *Pitx3* (E), an eye from an embryo deficient for *Foxe3*^{-/-} and heterozygous for *Pitx3* (F), an embryo heterozygous for *Foxe3* and deficient for *Pitx3* (G) and an eye of an embryo deficient for *Foxe3* and *Pitx3* (H). Notice the club-shaped lens in the *ak/ak* embryos. Arrows in sections point to the connections between the lens and surface ectoderm that broke off during the preparations of sections. L- lens

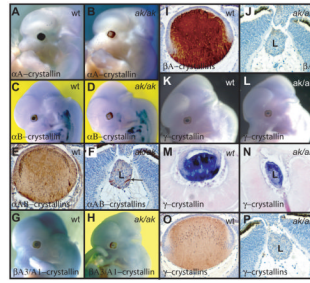


Figure 4.

Analysis of crystallin expression in E12.5 wild type and *ak* embryos. (A, B) Whole mount in situ hybridizations of an α A-crystallin (*Crya1*) probe to a wild type embryo (A) and to an *ak* embryo (B). (C, D) Whole mount in situ hybridizations of an α B-crystallin (*Crya2*) probe to a wild type embryo (C) and to an *ak* embryo (D). (E, F) Immunostaining of lens sections from a wild type embryo (E) and an *ak* embryo (F) with antibodies directed against α A and α B-crystallins. Only few cells show expression in the *ak* lens (arrow). (G, H) Whole mount in situ hybridizations of β A3/A1-crystallin (*Cryba1*) probe to a wild type embryo (G) and an *ak* embryo (H). (I, J) Immunostaining of lens sections from a wild type embryo (I) and an *ak* embryo (J) with antibodies directed against β A-crystallins. (K, L) Whole mount in situ hybridizations of γ -crystallin probe to a wild type embryo (K) and to an *ak* embryo (L). (M) A section of the embryo in (K) showing high levels of γ -crystallin expression in the posterior of the lens. (N) A section of the embryo in (L) showing γ -crystallin expression in the anterior part of the lens. (O, P) Immunostaining of lens sections from a wild type embryo (O) and an *ak* embryo (P) with antibodies direct against γ -crystallins.

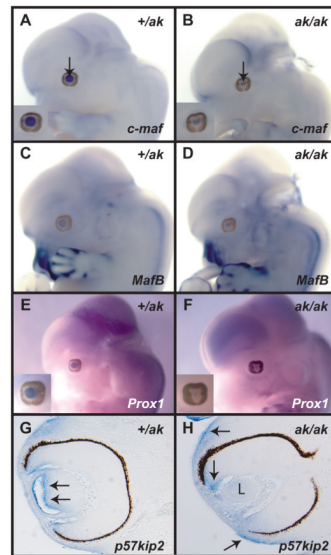


Figure 5.

Analysis of expression of *c-maf*, *MafB*, *Prox1* and *p57^{KIP2}* in *Pitx3* heterozygous and deficient embryos. (A, B) Whole mount in situ hybridizations of a *c-maf* probe to a +/ak embryo (A) and an *ak/ak* embryo (B). Arrows point to the lens. Insets provide a high magnification of the eye region. (C, D) Whole mount in situ hybridization of a *MafB* probe to a +/ak embryo (C) and an *ak* embryo (D). (E, F) Whole mount in situ hybridizations of a *Prox1* probe to a +/ak embryo (E) and an *ak* embryo (F). Insets provide a high magnification of the eye region. Notice the virtual absence of *Prox1* expression in F. (G) A section of a +/ak eye hybridized with *p57^{KIP2}* probe. Arrows point to the expression of this gene in the lens. (H) A section of an *ak* eye hybridized with the *p57^{KIP2}* probe. There is no expression detected in the lens (L), but the expression in the retina and surface ectoderm remains (arrows).

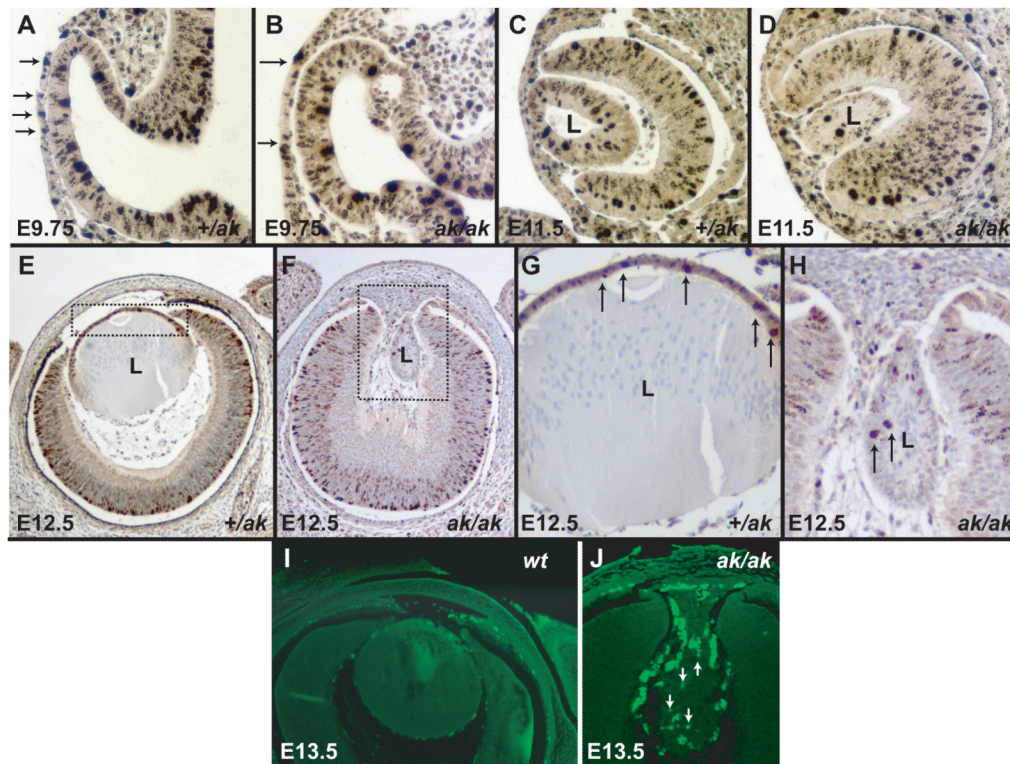


Figure 6.

Proliferation and apoptosis in *Pitx3* heterozygous and *Pitx3*-deficient embryos. (A) A section of an E9.75 *+ak* eye reacted with antibody against phosphohistone H3. Arrows point to the pH3 positive cells. (B) A section of an E9.75 *ak/ak* eye showing reaction with an antibody against phosphohistone H3. Arrows bracket the presumptive lens placode that shows a reduced labeling with antibodies against pH3. (C) A section of an E11.5 *+ak* eye reacted with antibody against phosphohistone H3. (D) A section of an E11.5 *ak/ak* eye showing reaction with the antibodies against phosphohistone H3. (E) A section of an E12.5 *+ak* eye reacted with antibody against phosphohistone H3. The boxed region is magnified in G. (F) A section of an E12.5 *ak/ak* eye showing reaction with the antibody against phosphohistone H3. The boxed region is magnified in H. Arrows indicate phosphohistone H3 positive cells. (I, J) TUNEL assay on the wild type eye (I) and on the *ak* eye (J). Apoptosis is not observed in the wild type lens, while TUNEL positive cells (fluorescent nuclei) are seen in the *ak* lens (arrows). Fluorescent cells surrounding the lens are likely to be of neural crest origin.