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β -catenin is essential for lamination but not neurogenesis in mouse retinal development

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Abstract

During vertebrate retinal development, the seven retinal cell types differentiate sequentially from a single population of retinal progenitor cells (RPCs) and organize themselves into a distinct laminar structure. The purpose of this study was to determine whether β -catenin, which functions both as a nuclear effector for the canonical Wnt signaling pathway and as a regulator of cell adhesion, is required for retinal neurogenesis or lamination. We used the Cre-loxP system to either eliminate β -catenin or to express a constitutively active form during retinal neurogenesis. Eliminating β -catenin did not affect cell differentiation, but did result in the loss of the radial arrangement of RPCs and caused abnormal migration of differentiated neurons. As a result, the laminar structure was massively disrupted in *β -catenin*-null retinas, although all retinal cell types still formed. In contrast to other neural tissues, eliminating β -catenin did not significantly reduce the proliferation rate of RPCs; likewise, activating β -catenin ectopically in RPCs did not result in overproliferation, but loss of neural retinal identity. These results indicate that β -catenin is essential during retinal neurogenesis as a regulator of cell adhesion but not as a nuclear effector of the canonical Wnt signaling pathway. The results further imply that retinal lamination and retinal cell differentiation are genetically separable processes.

Keywords

retina; retinal development; β -catenin; retinal lamination; cell adhesion; cell differentiation

Introduction

Tissue formation requires not only the specification and differentiation of functional cell types but also their correct organization. Although much has been learned about the molecular events regulating the differentiation of various cell types, far less is known about the events required to organize cells into tissues and how cellular organization is coordinated with cell-type differentiation.

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The vertebrate retina is a highly representative neural tissue consisting of well-defined cell types that are organized into a distinct multilayered laminar structure. The retina has six neuronal cell types (RGCs, amacrine cells, bipolar cells, horizontal cells, and rod and cone photoreceptors) and one glial cell type (Müller glial cells). The nuclei of the different cell types are located in three layers: an outer nuclear layer (ONL), containing rod and cone photoreceptors; an inner nuclear layer (INL), containing horizontal cells, bipolar cells, Müller glial cells, and amacrine cells; and a ganglion cell layer (GCL) containing RGCs and displaced amacrine cells. In the mouse, retinal neurogenesis starts at embryonic day 11.5 (E11.5), after the optic cup has formed, and continues until postnatal day 11 (P11) (Young, 1985). All seven cell types derive from a single population of naïve RPCs following a distinct temporal birth order (Livesey and Cepko, 2001). Specification and differentiation of the individual cell types requires specific regulatory factors that act singly or in combination to determine cell fate (Hatakeyama and Kageyama, 2004; Mu and Klein, 2004).

In addition to adopting a specific cell fate, a differentiating retinal cell must also establish its spatial identity by migrating to the correct position within the retina (Hinds and Hinds, 1974; Hinds and Hinds, 1979; Hinds and Hinds, 1983). RPCs are oriented radially along the apical/basal axis, and their processes span the full thickness of the retina, with their end feet anchored at both sides (Hinds and Hinds, 1974). Cell differentiation starts with the detachment of the end foot from the apical surface so that the differentiating cell can migrate to the appropriate layer. For example, newly formed RGCs, the earliest cell type to differentiate (Mu and Klein, 2004), begin to migrate toward the basal (inner) side of the neural retina to form the future GCL (Hinds and Hinds, 1974). Similar migration of other newly formed cell types are also required to produce a mature, functional retina (Hinds and Hinds, 1979; Hinds and Hinds, 1983). Fundamental questions concerning retinal cell migration remain unanswered. For example, the basis by which the different cell types assume their appropriate location once their cell fate has been determined is unknown. Furthermore, the relationship between retinal cell-type specification and cell migration is not understood.

It has long been suggested that differential cell adhesion plays a key role in sorting different cell types into a specific cellular organization during development (Steinberg, 1970; Steinberg, 1978). This view has been supported by more recent studies. For example, genetic screens in zebrafish have identified mutations that have severe retinal lamination defects (Pujic and Malicki, 2004). The genes associated with several of these mutations have been identified, and the proteins they encode are invariably involved in cell adhesion (Erdmann et al., 2003; Horne-Badovinac et al., 2001; Jensen and Westerfield, 2004; Malicki et al., 2003; Masai et al., 2003; Wei and Malicki, 2002). Two of them, *glass onion* and *parachute*, are mutations in the coding region of the type I cadherin gene, *N-cadherin* (Erdmann et al., 2003; Malicki et al., 2003; Masai et al., 2003). Type I cadherins form adherens junctions mostly through homotypic interactions (Wheelock and Johnson, 2003). Cadherin molecules in cadherens junctions are linked to the actin cytoskeletal network by β -catenin through α -catenin. Adherens junctions function in various biological processes, such as cell migration, cell sorting, and boundary formation. β -catenin therefore plays a pivotal role, physically bridging adherens junctions to the cytoskeletal network.

β -catenin is also an effector of the canonical Wnt signaling pathway that is activated by binding of the Wnt ligands to their seven-transmembrane receptors, the frizzled proteins (Logan and Nusse, 2004). In the absence of Wnts, cytoplasmic β -catenin is phosphorylated by glycogen synthase kinase-3 β (GSK-3 β) and casein kinase I (CKI) bound to the adenomatous polyposis coli (APC) complex and is targeted for degradation through the ubiquitination pathway. Activation of the Wnt pathway leads to the inhibition of GSK-3 β , resulting in the stabilization and cytoplasmic accumulation of β -catenin. The stabilized β -

catenin then interacts with Tcf/Lef transcription factors in the nucleus and activates downstream genes. In the nervous system, the Wnt- β -catenin pathway is involved in progenitor cell proliferation, cell-fate determination, and axon pathfinding (Charron and Tessier-Lavigne, 2005; Logan and Nusse, 2004).

In the developing mouse retina, many components of the Wnt pathway are expressed, and analysis of transgenic mice harboring a reporter transgene that responds to Wnt- β -catenin signaling, *TCF/Lef-LacZ*, suggests that the canonical pathway is activated during the early stages of retinogenesis (Liu et al., 2003). Other studies indicate that the Wnt pathway functions to promote RPC proliferation in *Xenopus* (Van Raay et al., 2005) and to prevent differentiation in the chick ciliary marginal zone (CMZ) (Kubo et al., 2003; Kubo et al., 2005). However, a definitive role of the Wnt- β -catenin pathway during retinal neurogenesis has not been established.

In this study, we directly determine the requirement for β -catenin in mouse retinal neurogenesis by specifically deleting the *β -catenin* gene in the developing retina. In our study, eliminating β -catenin led to a disorganized retina but did not affect the formation of any of the retinal cell types. In contrast, activating the nuclear function of β -catenin in the retina did not lead to overproliferation of RPCs, suggesting that the Wnt- β -catenin signaling pathway is not required for RPC proliferation during neurogenesis. Our results support the view that lamination and cell differentiation are genetically separable processes, and that during retinal neurogenesis, β -catenin is required solely for cell adhesion.

Materials and methods

Mouse lines

Mice harboring a conditional *β -catenin* allele (here referred to as *catnb^{lox(e2-6)}*) (Brault et al., 2001) were obtained from The Jackson Laboratory (Bar Harbor, ME) and mice with a stabilized *β -catenin* allele (*catnb^{lox(e3)}*) were generously provided by Makoto M. Taketo (Harada et al., 1999). *α -Cre* mice, originally generated by Marquardt et al. (Marquardt et al., 2001), were obtained from Guillermo Oliver (St. Jude Children's Research Hospital, Memphis, TN). *Six3-Cre* mice (Furuta et al., 2000; Mu et al., 2005) were kindly provided by Yas Furuta (M. D. Anderson Cancer Center, Houston, TX). These mice were bred into a 129/BL6 background. In all experiments using mice, the U. S. Public Health Service Policy on Humane Care and Use of Laboratory Animals was followed.

Genotyping

Genotyping for the *β -catenin* allele was performed by PCR following a previously described protocol (Brault et al., 2001). Genotyping for the *α -Cre* and *six3-Cre* mice was performed using PCR to detect the presence of the *Cre* sequence. The *Cre* primers used were 5'-CCT GAT CCT GGC AAT TTC GGC TA-3' and 5'-TCC AAT TTA CTG ACC GTA CAC CAA-3'. The PCR conditions used were: pre-denaturing (94°C, 5 min); 35 cycles of denaturing (94°C, 30 sec), annealing (55°C, 30 sec), and extension (72°C, 40 sec); followed by 2 min of extra extension at 72°C.

Histology and in situ hybridization

For morphological assessment of retinal phenotypes, embryos or eyes from different stages were fixed, paraffin-embedded, and sectioned at 10 μ m. After de-waxing, the sections were stained with hematoxylin and eosin (H&E) as previously described (Mu et al., 2005).

In situ hybridization with 7- μ m paraffin-embedded retinal sections was performed following a previously described protocol (Mu et al., 2004).

Fluorescence staining

Immunofluorescence staining on optimum cutting temperature medium (OCT)-embedded, 16 μm frozen retinal sections was carried out as previously described (Mu et al., 2005). The primary antibodies and the dilutions used in this study were as follows: anti-POU4f2 (previously called Brn3b) (Santa Cruz Biotechnology, Santa Cruz, CA, 1:50), anti- β -catenin (Cell Signaling, Beverly, MA, 1:500), anti-N-cadherin (BD Biosciences, San Jose, CA, 1:400), anti-Pax6 (Development Studies Hybridoma Bank, Iowa City, IA, 1:200), anti-Chx10 (Exalpha Biologicals, Maynard, MA, 1:400), anti-rhodopsin (Sigma-Aldrich, 1:400), anti-neurofilament 160 (NF160) (Sigma-Aldrich, 1:400), anti-PKC α (Sigma-Aldrich, 1:400), anti-Cralbp (Abcam, Cambridge, MA, 1:200), anti-cone arrestin (CAR) (Dr. Craft, UCLA, 1:300), anti-Par3 (Santa Cruz Biotechnology, 1:150), anti-Par6 (Santa Cruz Biotechnology, 1:150), anti-aPKC λ (Santa Cruz Biotechnology, 1:150). Fluorescent secondary antibodies were obtained from Molecular Probes (Eugene, OR) and were used at 1:800 dilution. Filamentous (F)-actin was stained with Alexa-488 conjugated phalloidin (Molecular Probes) at 1:1,000 dilution. Cell counting of individual cell types was performed as previously described (Mu et al., 2005). Briefly, total cells (based on propidium iodide staining) and positive cells for a specific marker were counted in the central retinal region within an arbitrary length unit and the percentage for each cell type was calculated. Effort was made to count cells from equivalent regions for different genotypes. Six sections from two different animals were counted for each marker and genotype. Statistical analysis was performed by two-sample *t* test with unequal variances.

BrdU labeling and TUNEL assay

For BrdU labeling, 100 μg BrdU per gram body weight was injected intraperitoneally into pregnant mice 1.5 hr before euthanization. Embryos were embedded in paraffin and then sectioned at 7 μm . The sections were de-waxed, treated as previously described (Mu et al., 2005), and stained with anti-BrdU primary antibody (Abcam, Cambridge, MA, 1:200) and Alexa 488-conjugated anti-rat IgG secondary antibody (Molecular Probes). The sections were then counter-stained with propidium iodide. The proportion of BrdU positive cells was calculated by counting the numbers of BrdU positive cells and total cells (based on propidium iodide staining) in an arbitrary length unit in the central region of the retinal sections and dividing by the total number of cells. For each genotype, six sections from two embryos were counted.

TUNEL assays were performed with the ApopTag plus fluorescein in situ apoptosis detection kit (Chemicon, Temecula, CA) following the manufacturer's instructions.

Results

Dynamic expression patterns of β -catenin in retinal development

We and others previously showed that *β -catenin* is abundantly expressed in the developing retina (Liu et al., 2002; Mu et al., 2001). To better understand the function of β -catenin in the retina, we determined the spatiotemporal expression pattern of β -catenin protein throughout retinogenesis. Consistent with previous reports (Liu et al., 2002), we observed distinct spatial expression patterns for β -catenin at all developmental stages. At E12.5, when only a few differentiated cells are present, β -catenin was expressed in uniform levels across the retina (Fig. 1A). At E14.5 and E17.5, although all cells still expressed β -catenin, higher expression was observed in RGCs and their axons (Fig. 1B, C). In mature P16 retinas, expression of β -catenin was highest in the outer plexiform layer (OPL), inner plexiform layer (IPL), and optic fibers (Fig. 1D). It was also expressed in the INL and GCL, but not in the ONL (Fig. 1D). Moreover, at all the stages examined, β -catenin was localized mostly to the plasma membrane; nuclear staining was minimal. Neuronal cell processes had high

levels of β -catenin, and from E12.5 to E17.5, β -catenin was strongly expressed at the apical and basal surfaces of the retina, where the end feet of RPCs are anchored (Fig. 1A-C).

Loss of β -catenin disrupts retinal structure

We used mice homozygous for a floxed β -catenin allele (*catnb*^{lox(e2-6)}) (Brault et al., 2001) to specifically delete β -catenin in the developing retina. In this allele, loxP sites flank exons 2 to 6 of the β -catenin gene. The mice were bred to either *six3-Cre* or α -*Cre* transgenic mice. *Six3-Cre* is expressed throughout the developing retina, with highest expression in the central region (Furuta et al., 2000), whereas α -*Cre* is expressed only in the anterior-peripheral retina (Marquardt et al., 2001). The expression of *Cre* in both transgenic lines starts at around E9, when the optic vesicle has formed but neurogenesis has yet to begin. The two *Cre* transgenes allowed us to determine the effects of β -catenin's absence at the neurogenesis stage, starting at around E11. Both *catnb*^{lox(e2-6)/lox(e2-6);six3-Cre} and *catnb*^{lox(e2-6)/lox(e2-6); α -Cre} mice were viable and fertile, but their eyes were smaller compared to those of either *catnb*^{lox(e2-6)/lox(e2-6)} or wild-type mice, and the small-eye phenotype was apparent as early as E14.5 (Fig. 1E-G; for *catnb*^{lox(e2-6)/lox(e2-6); α -Cre} mice, data not shown).

We first analyzed the cellular structure of β -catenin-deleted retinas at P16 when retinogenesis was completed and the multilayered mature retina had fully formed in the wild-type and *catnb*^{lox(e2-6)/lox(e2-6)} retinas. In both *catnb*^{lox(e2-6)/lox(e2-6);six3-Cre} and *catnb*^{lox(e2-6)/lox(e2-6); α -Cre} retinas, the normal laminar structure was disrupted (Fig. 2). This disruption occurred in the central region in *catnb*^{lox(e2-6)/lox(e2-6);six3-Cre} retinas (Fig. 2A-D) and in the peripheral region in *catnb*^{lox(e2-6)/lox(e2-6); α -Cre} retinas (Fig. 2E-H). Cells in these regions formed either rosette structures or were randomly distributed (Fig. 2B, D, F, H). Despite the massive disruption, some localized patterning was observed within the rosette structures (Fig. 2D, H). The rosettes were typically comprised of an inner and outer layer separated by a thin non cellular layer. The inner layer resembled the ONL and outer layer was similar to the INL. Throughout the β -catenin-deleted region, patches of non cellular structures resembling the IPL of the mature wild-type retina could be observed (Fig. 2D, H).

Identical defects were observed in the central region in *catnb*^{lox(e2-6)/lox(e2-6);six3-Cre} retinas and the peripheral region in *catnb*^{lox(e2-6)/lox(e2-6); α -Cre} retinas. We therefore performed further analyses solely with the *catnb*^{lox(e2-6)/lox(e2-6);six3-Cre} and *catnb*^{lox(e2-6)/lox(e2-6)} mice.

To determine when defects in retinal lamination first occurred, *catnb*^{lox(e2-6)/lox(e2-6);six3-Cre} retinas were examined at E12.5, 14.5, and 17.5 (Fig. 3). At E12.5, cellular defects were already obvious. These defects were restricted to the central retinal region, consistent with *six3-Cre* activity. In control retinas and in the peripheral region of *catnb*^{lox(e2-6)/lox(e2-6);six3-Cre} retinas, RPCs had oval-shaped nuclei arranged radially along the apical-basal axis (Fig. 3A). In contrast, the nuclei in the central region of *catnb*^{lox(e2-6)/lox(e2-6);six3-Cre} retinas lost their normal orientation and were round in shape (Fig. 3B). The apical surface of *catnb*^{lox(e2-6)/lox(e2-6);six3-Cre} retinas was rough and folded rather than smooth and curved as seen in control retinas. These defects were more pronounced at E14.5 and E17.5 (Fig. 3C-F). Additionally, at E14.5, the GCL of *catnb*^{lox(e2-6)/lox(e2-6);six3-Cre} retinas was greatly reduced in thickness or entirely absent in the central retinal region (Fig. 3C, D). At E17.5, the rosette structures were obvious (Fig. 3 E, F). Because the first sign of anomaly was the defective orientation and arrangement of RPCs, this could be the initial cause of the disorganized structure in the β -catenin-deleted retinas.

Loss of β -catenin leads to abnormal localization of N-cadherin, F-actin and the aPKC λ /Par3/Par6 complex during retinal development

Because β -catenin-mediated cell adhesion plays critical roles in cellular organization, we sought to determine whether cell adhesion was affected in β -catenin-deleted retinas. β -catenin physically interacts with N-cadherin and links it to F-actin. We therefore examined the expression of N-cadherin and F-actin during retinal development. In control retinas at E12.5, N-cadherin and F-actin were expressed in identical patterns as that of β -catenin (Fig. 4A, C, E). β -catenin, N-cadherin, and F-actin were expressed in all retinal cells, but expression was significantly higher at the apical and basal surfaces, and in the GCL (Fig. 4A, C, E). Expression of all three proteins was minimal in RPCs. The elevated expression of these proteins at both the apical and basal surfaces supports the notion that they have essential roles in maintaining the radial orientation and elongated morphology of RPCs. Moreover, their enhanced expression in the GCL suggests that differential cell adhesion is essential for proper migration of RGCs.

In *catnb^{lox(e2-6)/lox(e2-6);six3-Cre}* retinas, β -catenin was efficiently deleted in the central region by E12.5, as shown by the virtual absence of β -catenin expression (Fig. 4B). Consequently, we observed abnormal expression patterns for N-cadherin and F-actin in this region (Fig. 4D, F). High-level expression of both proteins at the apical surface was largely lost. Instead, we observed only patchy staining in these regions (Fig. 4D, F). The abnormal expression patterns of N-cadherin and F-actin continued at later developmental stages (e.g. E14.5, data not shown). The disruption of normal expression patterns of N-cadherin and F-actin at as early as E12.5 was consistent with the observed cellular defects. These results suggest that β -catenin is essential for maintaining proper cell adhesion and linkage to the cytoskeletal network during retinal development. If so, abnormal cell adhesion might cause the aberrant cell shape and organization in β -catenin-deleted retinas.

Atypical protein kinase C (aPKC) forms a complex with Par3 and Par6 and regulates cell polarity in various biological processes (Henrique and Schweisguth, 2003; Ohno, 2001). Two components of the complex, aPKC λ and Par3 (Pard3 in zebrafish) have been implicated in the establishment and maintenance of cell polarity in the developing neural retina (Koike et al., 2005; Wei et al., 2004). There is also evidence suggesting that the aPKC λ /Par3/Par6 complex interacts with cell adherens junctions in regulating cell polarity (Koike et al., 2005). We therefore examined how the expression of this complex was affected by the deletion of β -catenin. At E12.5 in the control retinas, aPKC λ , Par3, Par6 are expressed in all retinal cells with the highest level at the apical and basal surface (Fig. 4G, I, K). In the β -catenin deleted retinas, although the overall level of these proteins did not change, their localization at the apical surface disappeared (Fig. 4H, J, L), suggesting that the function of the aPKC λ /Par3/Par6 complex is dependent on the cell adherens junctions in retinal development.

β -catenin is not required for the initiation of differentiation but is essential for the normal migration of differentiated cells

Retinal cell differentiation starts at E11.5 with the birth of the first RGCs (Young, 1985). RGC differentiation is preceded by expression of the proneural bHLH gene *math5* in RPCs (Brown et al., 2001; Wang et al., 2001) and marked by expression of *pou4f2* (*brn3b*), a gene encoding a class IV POU domain transcription factor (Xiang et al., 1995). RGC development starts at the central retina and propagates toward the peripheral region as development proceeds (Gan et al., 1999; Gan et al., 1996). The expression of both *math5* and *pou4f2* follows a central-to-peripheral pattern in control retinas as shown by in situ hybridization at E12.5 and E14.5 (Fig. 5A, C, E, G). In *catnb^{lox(e2-6)/lox(e2-6);six3-Cre}* retinas, despite the abnormal cellular organization, expression of both *math5* and *pou4f2*

initiated normally in the central retina (Fig. 5B, F) and propagated toward the periphery (Fig. 5D, H). In addition, the level of expression of the two genes in the central region of *catnb^{lox(e2-6)/lox(e2-6);six3-Cre}* retinas was comparable to that in the peripheral region and in control retinas. These results suggest that the normal RPC orientation and organization was not required for the initiation of RGC differentiation, and hence neurogenesis.

However, the spatial expression pattern of *pou4f2* was not normal in the β -catenin-deleted region. Unlike in control retinas, where *pou4f2* expression was confined to the GCL on the basal side (Fig. 5E, G), *pou4f2* mRNA was found in patches of cells that were located at different positions throughout the retina (Fig. 5F, H). This expression pattern was confirmed using an anti-POU4f2 antibody. At E14.5, in *catnb^{lox(e2-6)/lox(e2-6)}* retinas, the majority of RGCs, as defined by POU4f2 protein expression, were in the GCL; only a few newly born RGCs were in the proliferation layer, and none of them were located on the apical side (Fig. 5I). In contrast, in *catnb^{lox(e2-6)/lox(e2-6);six3-Cre}* retinas, RGCs were found throughout the retina, including the apical side, and the boundary between the proliferation zone and the GCL was lost (Fig. 5J). Because newly differentiated RGCs were not properly located to the GCL in the absence of β -catenin, it is possible that their cell adhesion properties were affected and, hence, that their directional migration was disrupted. Newly born RGCs either failed to migrate, or migrated randomly.

All retinal cell types form in the absence of β -catenin

The fact that the absence of β -catenin did not affect RGC differentiation but did affect their location in the retina prompted us to examine whether the other retinal cell types would also differentiate but be mislocalized. Antibodies corresponding to other retinal cell types were used to determine the extent of differentiation in β -catenin-deleted retinas. At P16, all cell types were observed in the central region of *catnb^{lox(e2-6)/lox(e2-6);six3-Cre}* retinas, despite the fact that the retinal structure was severely disrupted in this region (Fig. 6A-P). Compared to *catnb^{lox(e2-6)/lox(e2-6)}* retinas, the different cell types were mislocalized and observed across the thickness of the retina in the β -catenin-deleted region (Fig. 6A-P). However, mislocalization was not completely random. Consistent with our histological observations, cell types within the rosettes were partially laminated, with an organization similar to that of control retinas along the apical/basal axis. The rosette centers resembled the outer segment of photoreceptors and stained heavily for rhodopsin (Fig. 6H). Adjacent to these centers were the rod and cone photoreceptor nuclei (rhodopsin-positive cells) that resembled the ONL (Fig. 6G, H). At least two types of intermediate neurons, bipolar cells (Fig. 6O, P; Chx10 and PKC α) and amacrine cells (Fig. 6M, X; Pax6), were more distally located, forming INL-like structures. In areas where cells did not form rosettes, the different cell types were randomly distributed. The discontinuous non cellular structures stained positively for PKC α (Fig. 6P), Syntaxin (data not shown) and Cralbp (Fig. 6N), proteins normally expressed in the processes of rod bipolar cells, amacrine cells, and Müller glial cells, respectively. This suggests that these non cellular structures were indeed disorganized IPL. To determine whether the relative proportion of retinal cell types was altered in *catnb^{lox(e2-6)/lox(e2-6);six3-Cre}* retinas, cell types were quantified using markers that allowed for distinction of individual cells because they were expressed predominantly in cell bodies. These included cones (CAR), bipolar cells (Chx10), amacrine cells (Pax6), and RGCs (Pou4f2). No significant changes were detected in any of these cell types in *catnb^{lox(e2-6)/lox(e2-6);six3-Cre}* retinas (Fig. 6).

We also examined the possibility that hybrid cell types containing different cell markers might be formed in the absence of β -catenin, which would indicate abnormal differentiation. We used antibodies against five pairs of markers including CAR/Chx10 (Fig. 7A-C, G-I), Cralbp/Chx10 (Fig. 7D-F, J-L), Cralbp/CAR (Fig. 7M-O, S-U), Pax6/Chx10 (Fig. 7P-R, V-X), and PKC α /Chx10 (data not shown). For each pair, the staining patterns of individual

markers were different in both control and β -catenin–deleted retinas (Fig. 7). Although there was overlap of staining for some pairs (Fig. 7F, L, U), this was caused by overlapping neighboring cells or their processes, but not by staining of a single cell by both markers (inlets, Fig. 7 D-F, J-L, M-O, S-U). We observed no cells that were positive for two different cell type markers in either the control or β -catenin-deleted retinas.

These results indicate that β -catenin was not required for the development of any of the retinal cell types. Furthermore, the partially laminated rosettes observed in the absence of β -catenin indicate that localized areas of retinal lamination still form in retinas lacking β -catenin.

Cell proliferation and apoptosis in the absence of β -catenin

Both $catnb^{lox(e2-6)/lox(e2-6);six3-Cre}$ and $catnb^{lox(e2-6)/lox(e2-6);\alpha-Cre}$ mice had smaller eyes than those of wild-type and $catnb^{lox(e2-6)/lox(e2-6)}$ controls (Fig. 1). The β -catenin-deleted regions in the retinas of the mutant mice were thicker than the corresponding regions of the control retinas (Fig. 2), but this alone could not account for the reduction in retinal size, because the total number of cells in the β -catenin-deleted retinas was also reduced. To determine the potential responsible mechanism, we examined the expression of *cyclinD1*, a gene expressed in RPCs and essential for their proliferation (Fantl et al., 1995; Sicinski et al., 1995). At E12.5 (data not shown) and E14.5 (Fig. 8A, B), no detectable changes in *cyclinD1* expression were observed in $catnb^{lox(e2-6)/lox(e2-6);six3-Cre}$ retinas compared to control retinas. Additionally, no significant changes in cell proliferation were observed in the β -catenin deleted retinas when we used BrdU to chase-label S-phase RPCs at E14.5 (Fig. 8C, D) and compared the proportions of BrdU-positive cells in $catnb^{lox(e2-6)/lox(e2-6);six3-Cre}$ and control retinas (cell counting results not shown).

We also examined whether retinal cells in β -catenin-deleted retinas underwent enhanced apoptosis at E12.5 and E14.5, because it is known that in wild-type retinas, little apoptosis occurs at these stages (Mu et al., 2005). TUNEL analysis showed that enhanced apoptosis was not observed in $catnb^{lox(e2-6)/lox(e2-6);six3-Cre}$ retinas at E12.5 (data not shown), but at E14.5, an increase in the number of apoptotic cells was clearly evident and these cells were restricted largely to the area where β -catenin was deleted (Fig. 8F). It was therefore likely that the reduced size of β -catenin-deleted retinas was caused in large part by enhanced apoptosis. Notably, enhanced apoptosis was also observed in retinas from zebrafish *glass onion* mutants (Pujic and Malicki, 2001), indicating that the loss of N-cadherin and the resulting defects in cell adhesion lead to a decrease in cell survival.

Ectopic activation of the canonical Wnt signaling pathway is detrimental to retinal development and leads to loss of neural retinal identity

The defects observed in β -catenin-deleted retinas during neurogenesis point to a role for β -catenin in cell adhesion. However, this fact does not rule out the possibility that the nuclear function of β -catenin is also required. To determine whether this was the case, we activated the nuclear function of β -catenin in the developing retina by breeding mice carrying the $catnb^{lox(e3)}$ allele to *six3-Cre* or *α -Cre* transgenic mice. The $catnb^{lox(e3)}$ allele contains floxed exon 3 sequences that encode the phosphorylation sites required for β -catenin degradation; deletion of exon 3 by Cre recombinase leads to the production of stabilized β -catenin and activation of the canonical Wnt pathway (Harada et al., 1999). This allele has been used widely to address the consequences of ectopic Wnt signaling in various tissues, and the ectopic stabilization of β -catenin often results in enhanced cell proliferation (Akiyama et al., 2004; Bierie et al., 2003; Harada et al., 1999; Zechner et al., 2003).

Ectopic activation of the Wnt- β -catenin pathway in the retina did not lead to an overproliferation of retinal cells; at E14.5, *catnb^{lox(e3)/+};six3-Cre* and *catnb^{lox(e3)/+};a-Cre* retinas were the same size as control retinas (Fig. 9A, E, data not shown). However, the choroid fissures in β -catenin-activated retinas failed to close (data not shown), and at later stages, the mutant eyes that were smaller than those of wild-type controls (Fig. 9F). β -catenin-activated retinas lost their normal structure as early as E14.5 in regions where *Cre* was expressed (Fig. 9E), and this defect became more severe at later stages (Fig. 9F). Furthermore, pigmented cells were abnormally intermingled with neural retinal cells in both *catnblox^{(e3)/+};six3-Cre* (inlet, Fig. 9F) and *catnb^{lox(e3)/+};a-Cre* retinas (data not shown), suggesting that ectopic activation of the Wnt- β -catenin pathway leads to a transdifferentiation of neural retinal cells into pigmented cells. To further examine the possibility that transdifferentiation had occurred, we determined the expression patterns of *Chx10* and *Mitf*, two transcription factors required for the development of the neural retina and pigmented epithelium, respectively (Fig. 9C, D). In the central region of *catnb^{lox(e3)/+};six3-Cre* retinas, no *Chx10* expression was detected indicating that neural retina development was compromised (Fig. 9G). Instead, the central region contained a sparse number of dispersed cells expressing *Mitf* (Fig. 8H). This result provided additional evidence that forced Wnt signaling altered the developmental pathway of retinal progenitor cells. These results suggest that activation of the canonical Wnt pathway does not facilitate neurogenesis, as might be expected, but rather compromises neural retinal integrity and identity.

Discussion

Cell adhesion function of β -catenin in retinal development

In our study, we showed that retinogenesis is profoundly affected by the loss of β -catenin. These results provide direct *in vivo* evidence that β -catenin has essential functions in retinal development and that these functions are associated with retinal lamination rather than cell proliferation or cell differentiation. During retinal development, β -catenin, N-cadherin, and F-actin have virtually identical spatiotemporal expression patterns (Xu et al., 2002). The absence of β -catenin caused disruption in the normal expression pattern of *N-cadherin*, indicating that β -catenin is essential in maintaining normal cell adhesion during retinal development. Consistent with this, deletion of *β -catenin* led to retinal defects similar to those observed in *N-cadherin* mutations in zebrafish (Erdmann et al., 2003; Malicki et al., 2003; Masai et al., 2003) and N-cadherin loss-of-function perturbation in chick embryos (Matsunaga et al., 1988). In these cases as well as in the *β -catenin*-deleted retinas described here, cell shape and radial alignment of RPCs was disrupted, and the laminar structure of the mature retina failed to form. The disruption of RPC orientation and alignment occurs early, before neurogenesis begins, suggesting that N-cadherin- β -catenin-mediated cell adhesion is essential in establishing and maintaining the orientation, polarity, and alignment of RPCs in the vertebrate retina. As development proceeds, newly differentiated cells appear to rely on the pre-existing apical-basal orientation of RPCs for their migration. Although the precise mechanism that controls the directional migration of newly differentiated retinal cells is unknown, differences in cell adhesion between RPCs and differentiated retinal cells may be essential. This hypothesis is supported by the fact that there is higher expression of N-cadherin and β -catenin in newly differentiated RGCs than there is in overlying RPCs. The defects in cellular organization seen in *β -catenin*-deleted retinas may be caused by changes in adhesion properties in both RPCs and newly differentiated retinal cells. Because localized cellular organization still occurred in *β -catenin*-deleted retinas, N-cadherin is probably not the only cell-adhesion molecule involved in retinal lamination. This is supported by the fact that *β -catenin* is not expressed in photoreceptors, which remain well-organized in *β -catenin*-deleted retinas. Indeed, many cell-adhesion molecules are expressed in the retina besides N-

cadherin (Honjo et al., 2000), suggesting that non- β -catenin cell adhesion mechanisms (such as gap junction) are also important in the formation and maintenance of laminar retinal structures. Moreover, retinal lamination is an actively regulated process requiring both intra- and intercellular signaling. Mutations in genes for many molecules, including aPKC λ (Horne-Badovinac et al., 2001; Koike et al., 2005), Moe (a FERM domain protein) (Jensen and Westerfield, 2004), and Par3 (a PDZ domain protein) (Wei et al., 2004), cause defective lamination. Par3 and aPKC, together with Par6, function as a complex in the regulation of cell polarity in a variety of cell types in different species (Henrique and Schweisguth, 2003; Ohno, 2001). We found that the expression patterns of aPKC λ , Par3, and Par6 were disrupted in the absence of β -catenin, suggesting this complex may function through adherens junctions in the developing retina. At present, however, it is not understood how these molecules act together to coordinate cell orientation, cell polarity, cell migration, and lamination in the retina.

The function of β -catenin may not be limited to just lamination formation during retinal neurogenesis. There is much evidence suggesting that adherens junctions are also required for neurite extension (Elul et al., 2003; Riehl et al., 1996; Stone and Sakaguchi, 1996) and the canonical Wnt pathway inhibits this process (Ouchi et al., 2005).

Cell adhesion and cell differentiation are genetically separable processes in the developing retina

In the absence of β -catenin, RPCs lost their radial orientation and arrangement, but neurogenesis was not affected. This indicates that proper RPC arrangement is not a prerequisite for the differentiation of the retinal cell types. Similar observations have been made with zebrafish mutants with retinal lamination defects (Erdmann et al., 2003; Jensen and Westerfield, 2004; Koike et al., 2005; Malicki and Driever, 1999; Masai et al., 2003). Despite the fact that RGCs, the first retinal cells to differentiate, did not migrate to their proper position in the GCL, all later cell types still developed. These results imply that the sequential development of the different retinal cell types is not dependent on the proper cellular organization. This is consistent with accumulating evidence suggesting that intrinsic mechanisms play a major role in retinal cell differentiation (Cayouette et al., 2003; Kay et al., 2005; Mu et al., 2005).

Although the loss of proper polarity and arrangement of RPCs was the earliest defect observed, it is possible that in the absence of β -catenin, the later aberrant migration of RGCs and other retinal cell types was not solely caused by polarity and arrangement defects in RPCs, but also by defective cell adhesion properties of the differentiated neurons. Migration may require proper adhesion between differentiated neurons and RPCs, as well as their interaction with the extracellular matrix or other substrates. In the developing cortex, differentiated neurons migrated along the processes of radial glial cells, which are the neural progenitor cells (Miyata et al., 2001; Noctor et al., 2001; Noctor et al., 2002). A similar scenario might exist in the developing retina in which the differentiated neuronal cell types rely on the oriented processes of RPCs for their proper unidirectional migration.

Nuclear function of β -catenin and the Wnt signaling pathway in retinal development

Several studies have suggested that the Wnt signaling pathway is important in retinal development in chickens (Kubo et al., 2003; Kubo et al., 2005), *Xenopus* (Van Raay et al., 2005), zebrafish (Cavodeassi et al., 2005; van de Water et al., 2001), and mice (Maretto et al., 2003). In the chick, Wnt2b (also known as Wnt13) is expressed at the tip of the ciliary marginal zone (CMV) of the retina. It promotes RPC proliferation and prevents precocious differentiation by suppressing the expression of proneural and neural genes (Kubo et al., 2003; Kubo et al., 2005). In *Xenopus*, Frizzled5 (Xfz5) has been shown to regulate the

proliferation and neural potential of RPCs. In the mouse, knockout of the Wnt co-receptor *Lrp6* gene leads to microphthalmia and retinal defects (Maretto et al., 2003). These findings combined with the fact that many components of the Wnt pathway are expressed in the developing mouse retina suggest that Wnt signaling is an important process in retinogenesis, although its specific role may vary among different vertebrate species. However, in all the above-mentioned studies, the role of the canonical Wnt pathway in retinal development was investigated by inactivating the pathway at stages prior to cell differentiation, thereby preventing an assessment of its role in cell type specification and differentiation. Transgenic analysis, using the *TCF/Lef-LacZ* transgene, which responds to canonical Wnt- β -catenin signaling, indicates that the Wnt- β -catenin pathway is largely inactive during neurogenesis (Liu et al., 2003). Although some residual LacZ-positive cells were still present at E14.5 in the retina of *TCF/Lef-LacZ* mice, this probably reflected the stability of LacZ rather than the activation of the canonic Wnt pathway. Therefore, Wnt- β -catenin signaling appears only to be required at early stages of retinal development to prevent precocious differentiation and is inactive during neurogenesis. Consistent with this idea, we found that β -catenin functions mainly in cell adhesion during retinal neurogenesis and not in the specification and differentiation of retinal cell types because these events were not affected in the absence of β -catenin. Moreover, we found that ectopic activation of the Wnt- β -catenin pathway did not lead to overproliferation, and was in fact detrimental to neurogenesis, consistent with previous reports both in zebrafish (van de Water et al., 2001) and mice (Ouchi et al., 2005). Retinal neurogenesis may require inactivation of this pathway, and this may be achieved by inhibitory molecules such as Axin (van de Water et al., 2001), Sfrp1, Sfrp2 (Liu et al., 2003; Mu et al., 2001) and WIF-1 (Hunter et al., 2004), which are all expressed in the developing retina. This differs from other neural tissues in which the canonical Wnt- β -catenin pathway is required for cell proliferation (Chenn and Walsh, 2002; Zechner et al., 2003) and cell-fate determination (Hari et al., 2002; Lee et al., 2004). Notably, in *Xenopus*, overexpression of Sfrp2 and Frzb, two inhibitory molecules of the Wnt pathway, also cause retinal lamination defects (Ladher et al., 2000). However, because these genes were overexpressed very early, the defects observed may be indirect. It is also possible that Wnt ligands act through non-canonical signaling pathways in the retina (Esteve et al., 2003; Yu et al., 2004) and that non-canonical Wnt signaling and cell adherens junctions are both required for retinal lamination.

The canonical Wnt- β -catenin pathway may be required in the separation of the neural retinal and pigmented epithelial fates. Chx10 and Mitf function to establish and maintain these two fates respectively (Horsford et al., 2005; Rowan et al., 2004). Our observation that activated β -catenin leads to loss of Chx10 expression and gain of ectopic Mitf expression, and hence an apparent change in the pathway of neural retinal cells into pigmented cells, suggests that the Wnt- β -catenin pathway has a role in pigmented epithelial fate.

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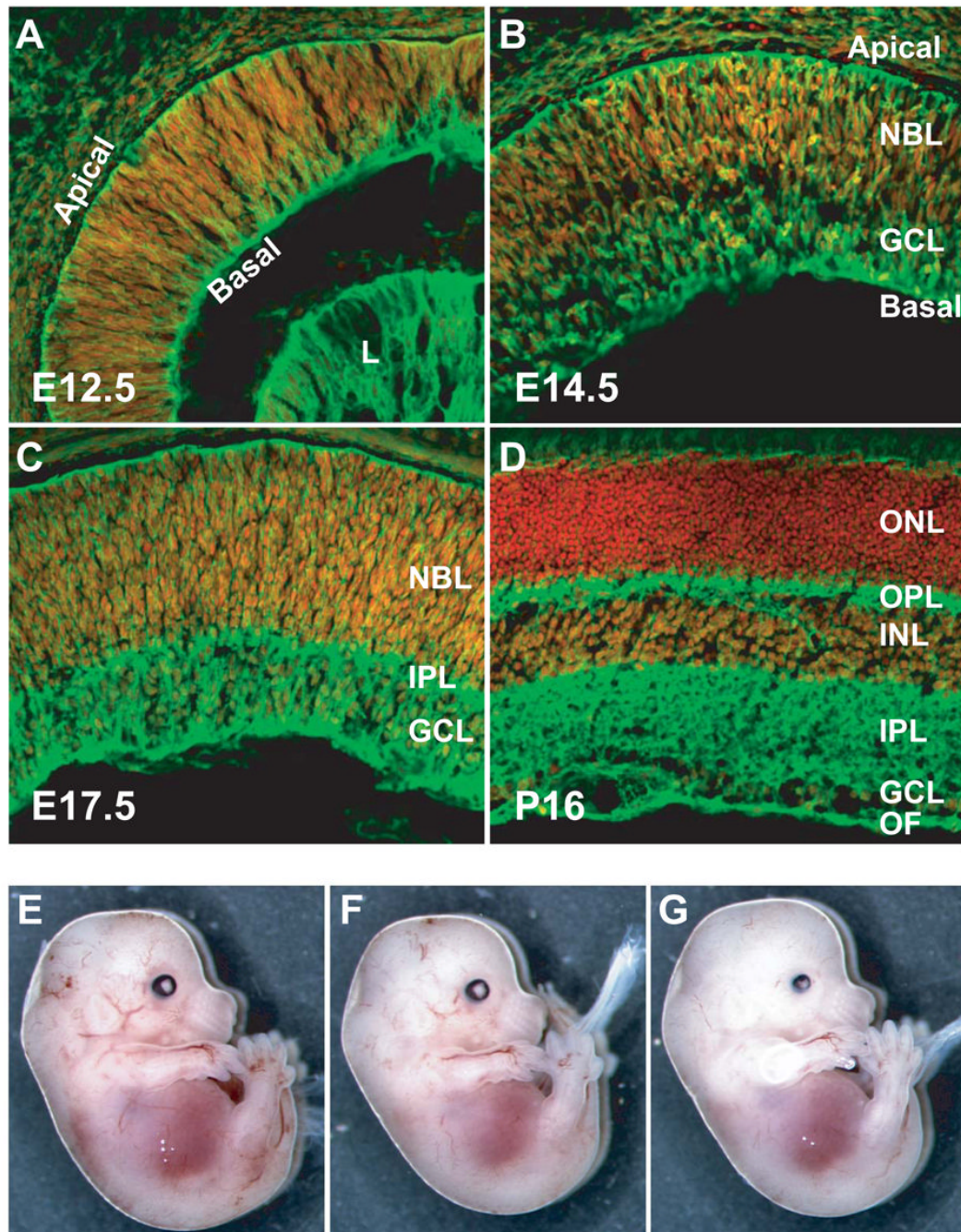


Fig. 1. Expression patterns and conditional deletion of β -catenin in the developing retina. (A-D) Merged confocal images of β -catenin protein expression at E12.5 (A), E14.5 (B), E17.5 (C), and P16 (D) detected by an anti- β -catenin antibody (green). Nuclei were counter-stained red with propidium iodide. (E-G) Three E14.5 embryos of wild-type (E), $catnb^{lox(e2-6)/lox(e2-6)}$ (F), and $catnb^{lox(e2-6)/lox(e2-6);six3-Cre}$ (G). E and F have normal-sized eyes but G has smaller eyes. NBL, neural blast layer; GCL, ganglion cell layer; ONL, outer nuclear layer; OPL, outer plexiform layer; INL, inner nuclear layer; IPL, inner plexiform layer; OF, optic fiber.

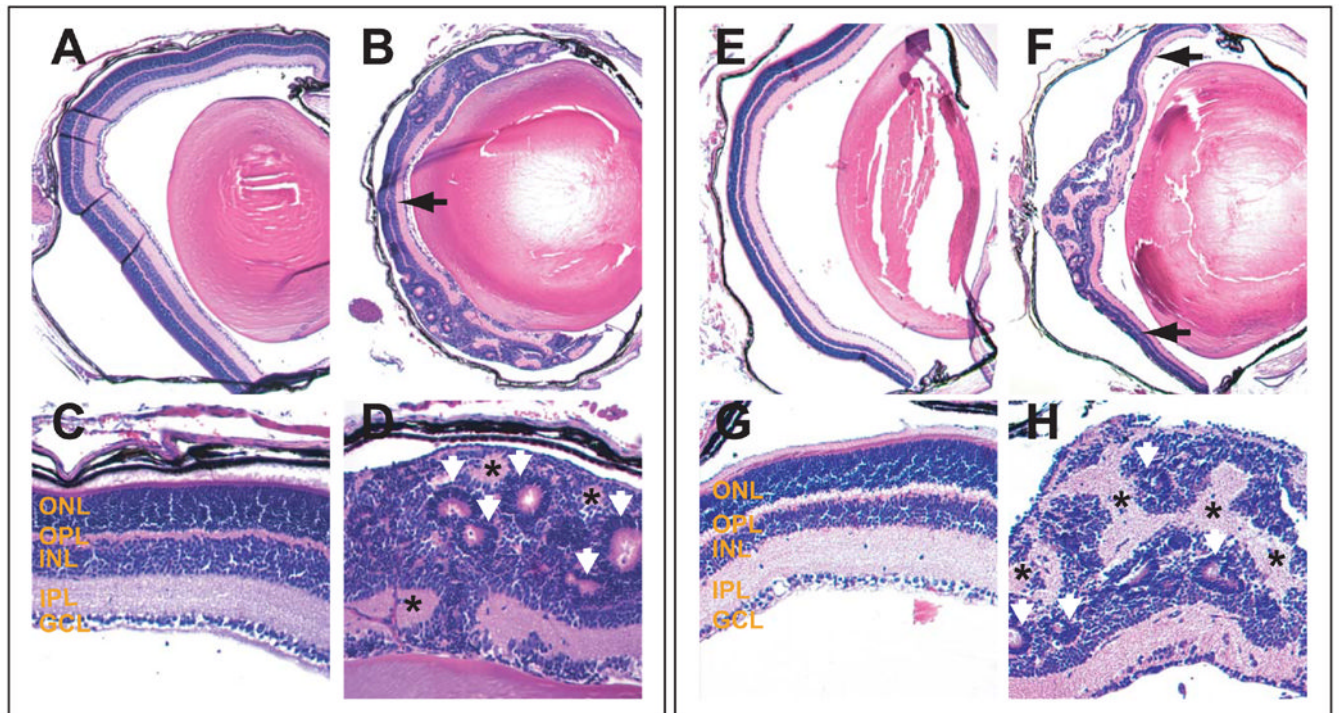


Fig. 2. Deletion of β -catenin in P16 retinas. Retinal sections of different genotypes were analyzed by H&E staining. (A) Control $catnb^{lox(e2-6)/lox(e2-6)}$ retina. (B) $catnb^{lox(e2-6)/lox(e2-6);\alpha-Cre}$ retina. (C and D) Higher magnification of A and B, respectively. (E) Control $catnb^{lox(e2-6)/lox(e2-6)}$ retina. (F) $catnb^{lox(e2-6)/lox(e2-6);six3-Cre}$ retina. (G and H) Higher magnification of E and F respectively. Black arrows point to the normal laminated structures in the central region of the $catnb^{lox(e2-6)/lox(e2-6);\alpha-Cre}$ retina (B) and the peripheral region of the $catnb^{lox(e2-6)/lox(e2-6);six3-Cre}$ retina (F). White arrowheads in D and H point to rosette structures with inner and outer cell layers. Asterisks indicate the non cellular structures in the β -catenin-deleted retinas.

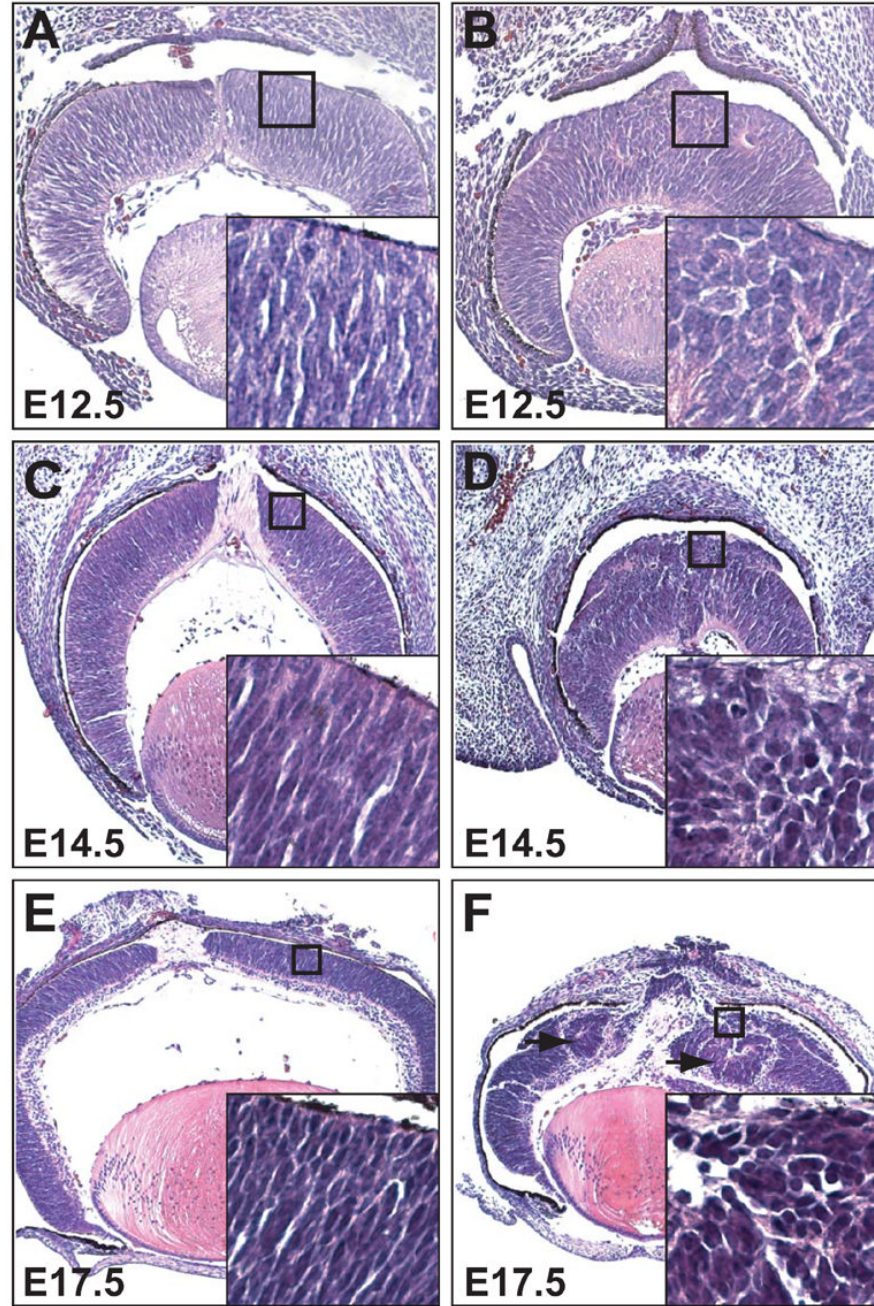


Fig. 3. Progressive loss of radial orientation and cellular organization of RPCs during development. Shown are H&E-stained sections of E12.5 (A, B), E 14.5 (C, D), and E17.5 (E, F) retinas. (A, C, E) Control *catnb^{lox(e2-6)/lox(e2-6)}* retinas showing RPCs with elongated nuclei and aligned radially along the apical-basal axis. (B, D, F) *catnb^{lox(e2-6)/lox(e2-6);six3-Cre}* retinas. Insets at the lower right of each image show a higher magnification of the boxed region. Note that at as early as E12.5 (B), RPCs from β -catenin-deleted retinas lose their normal organization, and their nuclei are round. These defects become more severe at E14.5 (D) and E17.5 (F), and rosette structures begin to appear (arrowheads).

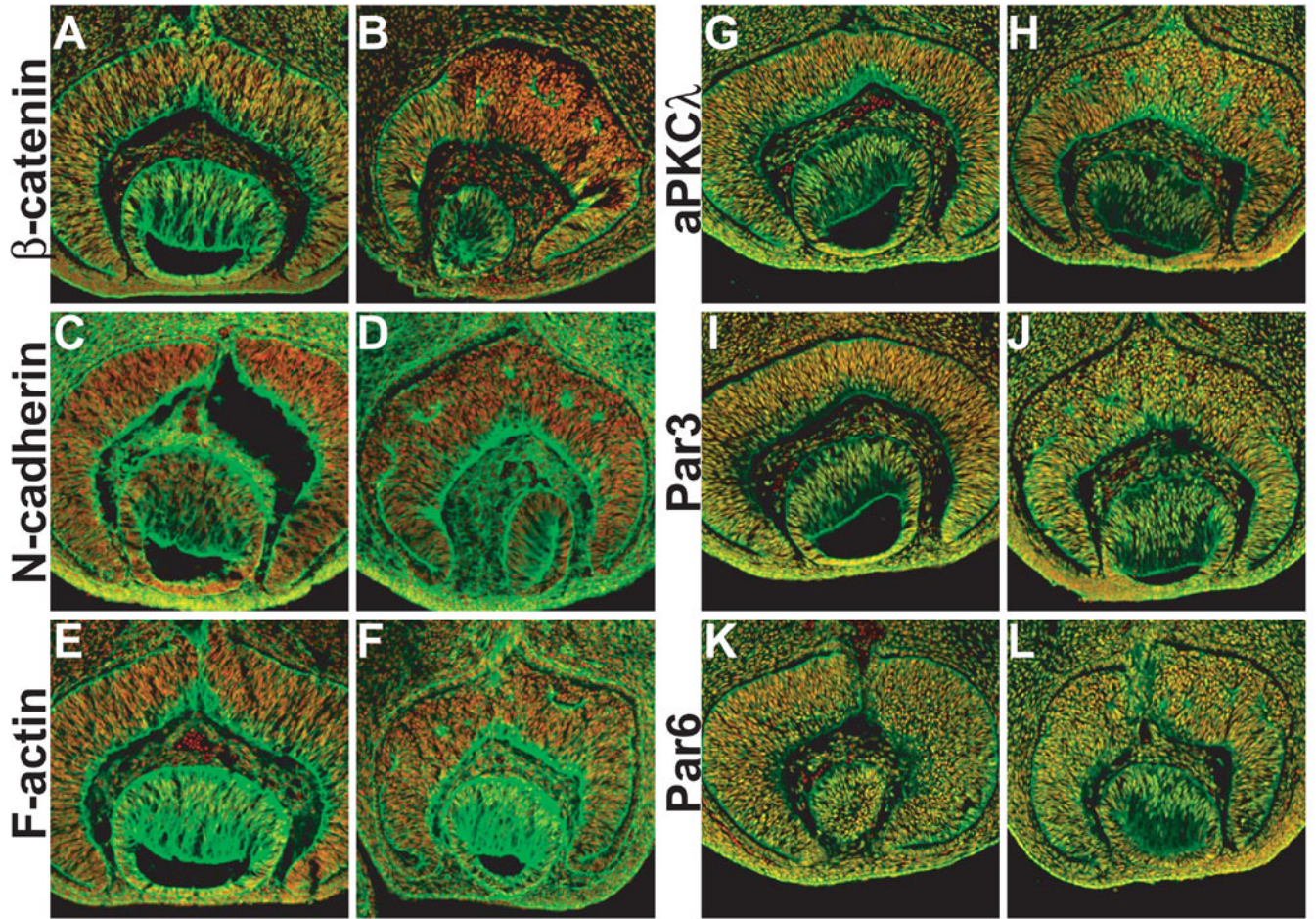


Fig. 4.

Expression of β -catenin, N-cadherin, F-actin, aPKC λ , Par3, and Par6 proteins in E12.5 retinas. Antibody staining is green, and nuclei are stained red with propidium iodide. (A, C, E, G, I, K) β -catenin (A), N-cadherin (C), F-actin (E), aPKC λ (G), Par3 (I), and Par6 (K) expression in *catnb*^{lox(e2-6)/lox(e2-6)} control retinas. (B, D, F, H, J, L) Corresponding expression in *catnb*^{lox(e2-6)/lox(e2-6);six3-Cre} retinas. In A, C, E, G, I, and K, high expression levels are observed at the apical and basal surfaces and the GCL. In the central region of *catnb*^{lox(e2-6)/lox(e2-6);six3-Cre} retinas, β -catenin was not detected, and the normal N-cadherin, F-actin, aPKC λ , Par3, and Par6 expression patterns are disrupted. Note that the staining for N-cadherin, F-actin, aPKC λ , Par3, and Par6 on the apical surface in the central region of *catnb*^{lox(e2-6)/lox(e2-6);six3-Cre} retinas disappears.

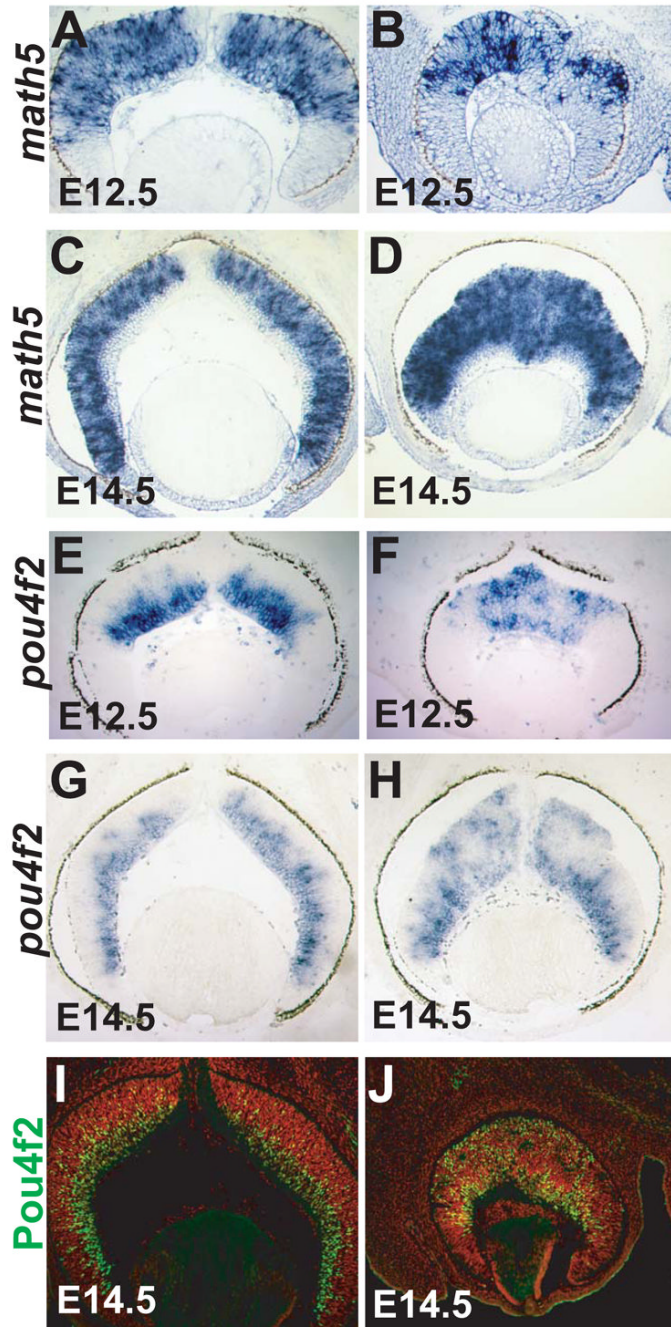


Fig. 5. *Math5* and *pou4f2* expression in control and β -catenin-deleted retinas. (A, C, E, G, I) Control $catnb^{lox(e2-6)/lox(e2-6)}$ retinas. (B, D, F, H, J) $catnb^{lox(e2-6)/lox(e2-6);six3-Cre}$ retinas. *Math5* and *pou4f2* transcript expression are shown in A-H, and POU4f2 protein expression (green) is shown in I and J. Nuclei in I and J are stained red with propidium iodide. Similar to $catnb^{lox(e2-6)/lox(e2-6)}$ control retinas (A, C, E, G), expression of *math5* and *pou4f2* initiates normally at the center of $catnb^{lox(e2-6)/lox(e2-6);six3-Cre}$ retinas (B, F), and expands toward the periphery (D, H). Unlike the $catnb^{lox(e2-6)/lox(e2-6)}$ control retinas (E, G, I), newly formed RGCs, marked by *pou4f2* expression, did not migrate normally to the GCL in β -catenin-deleted retinas (F, H, J).

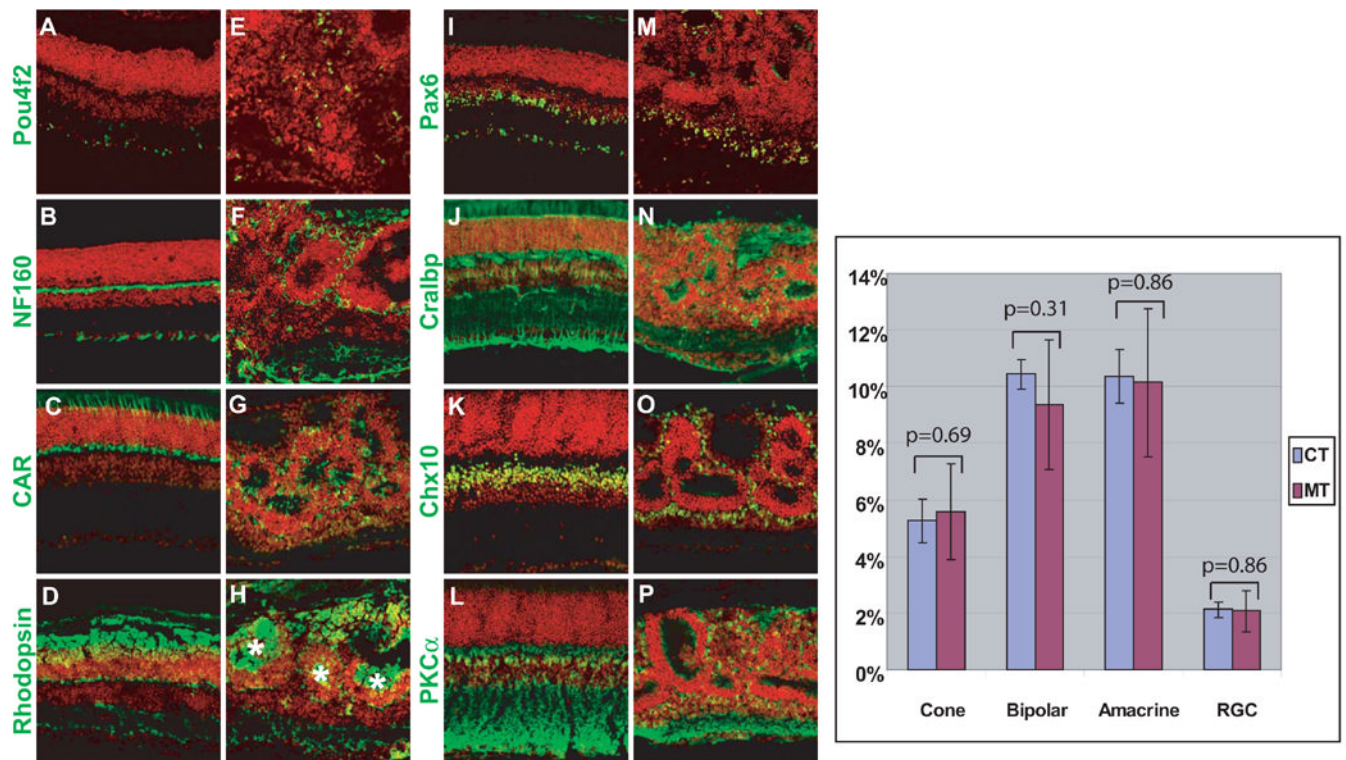


Fig. 6. Differentiation of retinal cell types in control and β -catenin-deleted retinas at P16. Sections of P16 control *catnb^{lox(e2-6)/lox(e2-6)}* (A-D, I-L) or *catnb^{lox(e2-6)/lox(e2-6);six3-Cre}* (E-H, M-P) retinas were stained with individual antibodies against different retinal cell-type markers (green) and counter-stained red with propidium iodide. Pou4f2 (A, E): RGCs; NF160 (B, F): horizontal cells and RGCs; CAR (C, G): cones; rhodopsin (D, H): rods; Pax6 (I, M): amacrine cells and RGCs; Cralbp (J, N): Müller cells; Chx10 (K, O), bipolar cells; and PKC α (L, P), rod bipolar cells. Note that although all cell types are present, they lose their normal positions in *catnb^{lox(e2-6)/lox(e2-6);six3-Cre}* retinas (E-H, M-P) compared to control retinas (A-D, I-L). Localized lamination of varying extents is observed around the rosettes with heavy staining of rhodopsin at the center (asterisks, H). Right panel: Quantification of four cell types (cone, bipolar, amacrine and RGC) in control (CT) and β -catenin-deleted (MT) retinas. Y axis is the percentage of individual cell types within an arbitrary length unit of retinal sections (see Material and methods). Standard deviations (error bars) are shown for each sample (n=6). There is no significant difference for any of the cell types between control and mutant retinas as indicated by the p values.

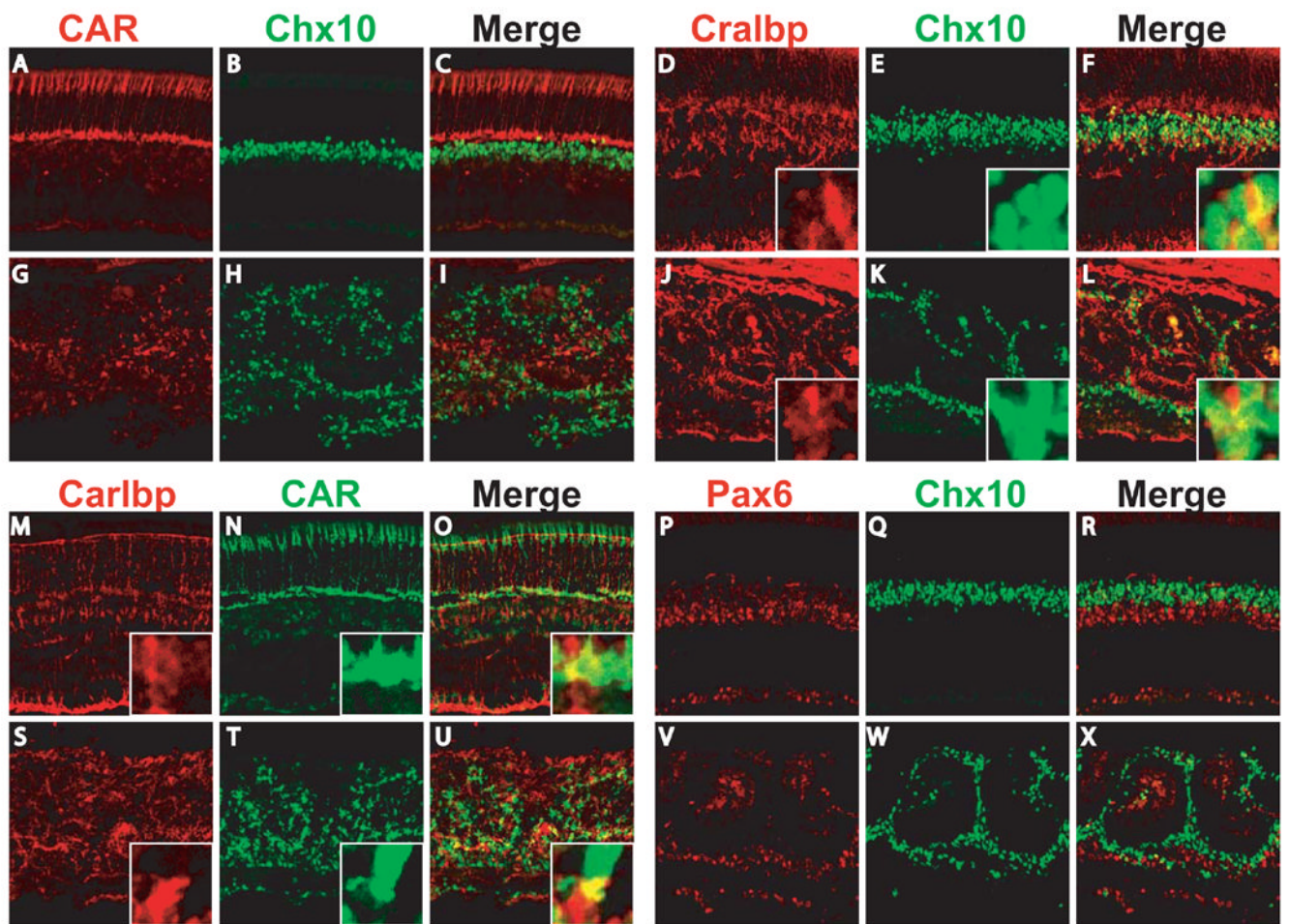


Fig. 7.

No hybrid retinal cell types were formed in the absence of β -catenin. P16 control *catnb^{lox(e2-6)/lox(e2-6)}* (A-F, M-R) or *catnb^{lox(e2-6)/lox(e2-6);six3-Cre}* (G-L, S-X) retinal sections were double stained with markers for two different cell types as indicated. For each pair of markers, images of individual and merged signals were shown. Insets (D-F, J-L, M-O, and S-U) are high magnification of regions that appear yellow in the merged images. The yellow color was resulted from neighboring cells or their processes that overlap in the sections, but not from a single cell positive for both markers. No cell was observed double-positive for any pair of markers in either the control or β -catenin-deleted retinas.

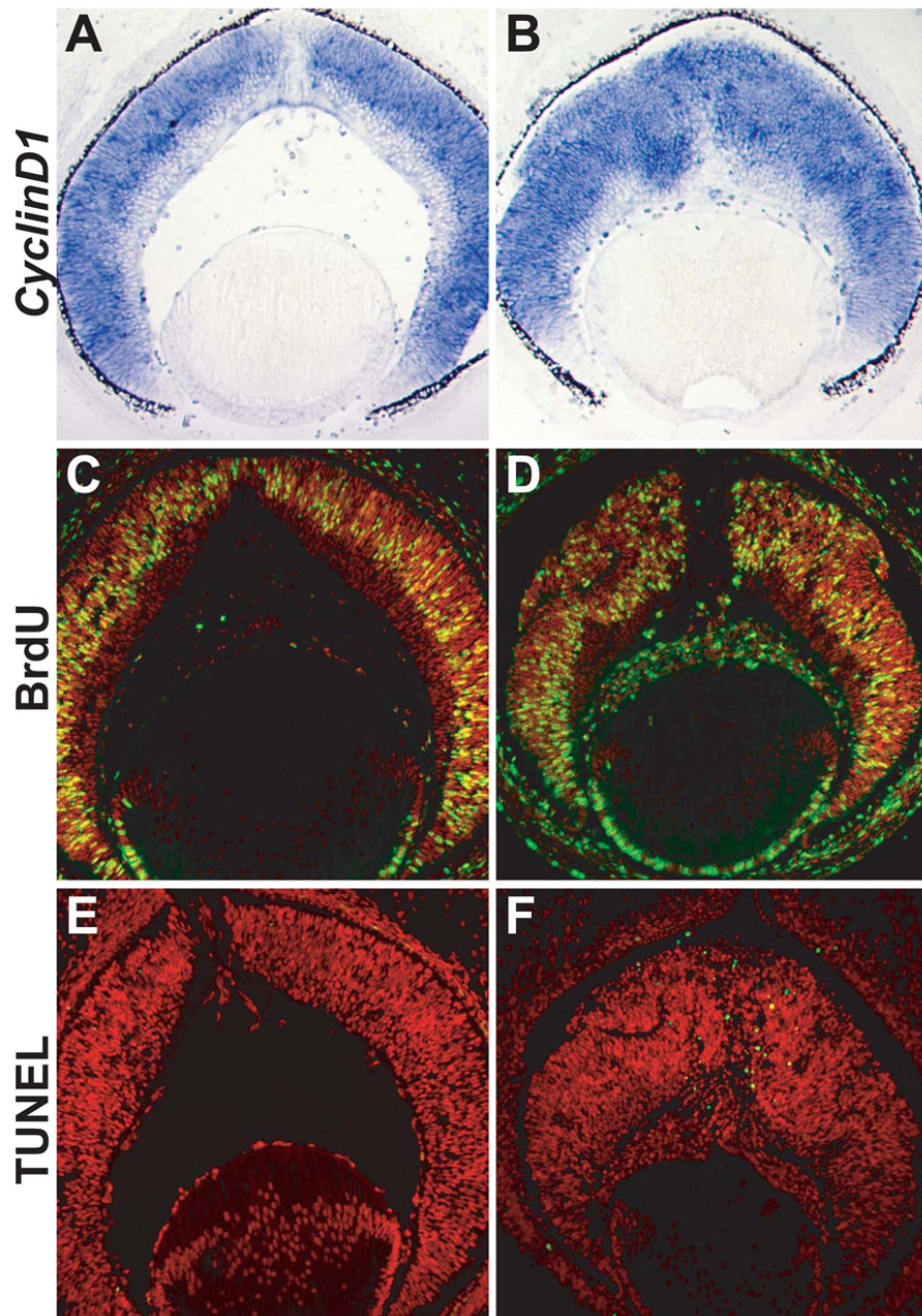


Fig. 8. Cell proliferation and apoptosis in *catnb*^{lox(e2-6)/lox(e2-6)} control (A, C, E) and *catnb*^{lox(e2-6)/lox(e2-6);six3-Cre} (B, D, F) retinas at E14.5. (A, B) Expression of *cyclinD1* transcripts. (C, D) BrdU chase-labeling (green) for S-phase cells. Nuclei are stained red with propidium iodide. (E, F) TUNEL assay for apoptotic cells (green).

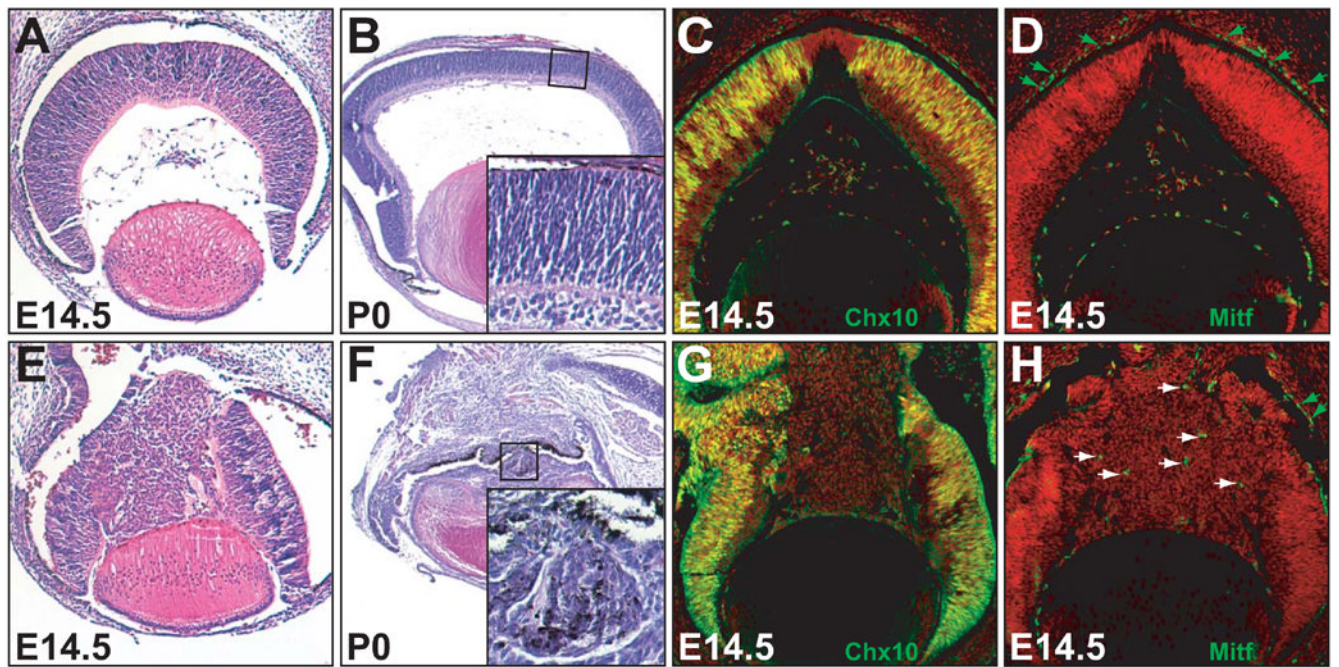


Fig. 9.

Effects of the ectopic expression of stabilized β -catenin on retinal development. (A-D) control *catnb^{lox(e3)/lox(e3)}* retinas. (E-H) *catnb^{lox(e3)/lox(e3);six3-Cre}* retinas. (A, B, E, F) H&E staining of E14.5 and P0 retinas shows cellular disorganization in stabilized β -catenin-activated (*catnb^{lox(e3)/lox(e3);six3-Cre}* retinas. Note the presence of pigmented cells in the *catnb^{lox(e3)/lox(e3);six3-Cre}* neural retina (inlet, F). At later stages (P0), *catnb^{lox(e3)/lox(e3);six3-Cre}* eyes (F) are smaller than the control (B). (C, G, D, H) immunofluorescence staining (green) for Chx10 and Mitf of E14.5 control (C, D) and *catnb^{lox(e3)/lox(e3);six3-Cre}* (H, G) retinas. In the central regions of *catnb^{lox(e3)/lox(e3);six3-Cre}* retinas, Chx10 expression is lost, and some cells (white arrowheads) express Mitf ectopically. Green arrowheads point to normal Mitf-expressing cells in the pigmented epithelia.