

Dlic1 deficiency impairs ciliogenesis of photoreceptors by destabilizing dynein

Shanshan Kong¹, Xinrong Du¹, Chao Peng¹, Yiming Wu¹, Huirong Li², Xi Jin², Ling Hou², Kejing Deng¹, Tian Xu^{1,3}, Wufan Tao¹

¹State Key Laboratory of Genetic Engineering and Institute of Developmental Biology and Molecular Medicine, School of Life Science, Fudan University, Shanghai 200433, China; ²Developmental Cell Biology and Disease Program, State Key Laboratory Cultivation Base and Key Laboratory of Vision Science of Ministry of Health, Wenzhou Medical College, Wenzhou, Zhejiang 325003, China; ³Howard Hughes Medical Institute, Department of Genetics, Yale University School of Medicine, New Haven, CT 06536, USA

Cytoplasmic dynein 1 is fundamentally important for transporting a variety of essential cargoes along microtubules within eukaryotic cells. However, in mammals, few mutants are available for studying the effects of defects in dynein-controlled processes in the context of the whole organism. Here, we deleted mouse *Dlic1* gene encoding DLIC1, a subunit of the dynein complex. *Dlic1*^{-/-} mice are viable, but display severe photoreceptor degeneration. Ablation of *Dlic1* results in ectopic accumulation of outer segment (OS) proteins, and impairs OS growth and ciliogenesis of photoreceptors by interfering with Rab11-vesicle trafficking and blocking efficient OS protein transport from Golgi to the basal body. Our studies show that *Dlic1* deficiency partially blocks vesicle export from endoplasmic reticulum (ER), but seems not to affect vesicle transport from the ER to Golgi. Further mechanistic study reveals that lack of *Dlic1* destabilizes dynein subunits and alters the normal subcellular distribution of dynein in photoreceptors, probably due to the impaired transport function of dynein. Our results demonstrate that *Dlic1* plays important roles in ciliogenesis and protein transport to the OS, and is required for photoreceptor development and survival. The *Dlic1*^{-/-} mice also provide a new mouse model to study human retinal degeneration.

Keywords: ciliogenesis; *Dlic1*; dynein stability; photoreceptor degeneration; Rab11 vesicles
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Introduction

Photoreceptor cells in the vertebrate retina are highly polarized light-sensory neurons, which can be divided into several morphologically and functionally distinct compartments: the photoreceptor outer segment (OS) and inner segment (IS) continuing into the perikaryon and the synaptic region. The OS is a specialized primary cilium composed of stacks of disk membranes, which links to the IS by the connecting cilium (CC). The disk mem-

branes bear large quantities of phototransduction proteins and undergo renewal throughout life by the assembly of new disks at the base and by the shedding of older disks from the tip [1]. All of the OS proteins including rhodopsin are synthesized in the endoplasmic reticulum (ER) located in the IS and perikaryon, and funneled through the Golgi. They are then apically transported to the base of the connecting cilium in a dynein-dependent manner [2] and finally to the OS [3]. Accumulating evidence indicates that mutations that affect OS development and/or protein transport to the OS generally lead to photoreceptor degeneration [4].

Cytoplasmic dynein 1 is a major microtubule minus-end-directed motor complex and has been implicated in many important processes, including mitosis, membrane trafficking, organelle positioning, and cell migration within eukaryotic cells [5]. Dynein 1 consists of four distinct classes of subunits: dynein heavy chain (DHC,

Correspondence: Wufan Tao^a, Tian Xu^b

^aTel: +86-21-65643949; Fax: +86-21-65642111

E-mail: wufan_tao@fudan.edu.cn

^bTel: 203-767-2623; Fax: 203-767-1762

E-mail: tian.xu@yale.edu

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DYNC1H1), dynein intermediate chain (DIC, DYNC1H1), dynein light intermediate chain (DLIC, DYNC1LI) and dynein light chain (DLC, DYNL) [6]. The DHC harbors the ATPase and microtubule motor domains, whereas the remaining subunits play regulatory functions, such as the selection of cargoes. DLIC is highly conserved among different eukaryotic species. In mammals there are two *Dlic* genes for cytoplasmic dynein 1, *Dlic1* (*Dync1li1*) and *Dlic2* (*Dync1li2*), while in lower eukaryotes there is only one *Dlic* gene. The function of the DLIC protein was first implicated in the control of the mitotic spindle and the assembly of centrioles by the fact that DLIC1, not DLIC2, specifically interacted with pericentrin (PCNT) in Cos-7 cells [7]. Mutations in the gene encoding DLIC or depleting its gene product by RNAi were shown to result in a variety of mitotic defects from yeast to mammals [8-11]. DLIC1 also participates in intracellular vesicle transport via forming a complex with small GTPases Rab4 and Rab11, respectively [12, 13]. However, the functions of the DLIC subunit in keeping the integrity of dynein and Golgi apparatus and the development of neurons are controversial. It was shown that yeast DLIC and mammalian DLIC1 were not required for the stability of dynein complex [9, 11, 14], but depletion or loss of DLIC in *Drosophila* cells and *Aspergillus nidulans* led to destabilization of DHC and DIC [10, 15]. Palmer *et al.* [14] reported that depleting DLIC1 by RNAi in HeLa cells disrupted the Golgi apparatus, but Sivaram *et al.* [11] claimed that it had no effect on the Golgi. Mutating *Dlic* gene blocks the transport of neuronal retrograde cargoes in worms and flies [16-18] and results in dendritic and axonal defects in neurons, such as a reduction in the length and number of dendrite branches in *Drosophila*. However, mice with a point mutation *Dlic1*^{N235Y} display an increase in dendrite length of cortical neurons (with no changes in branching) and the number of dendrite branches of dorsal root ganglia (DRG) neurons, in addition to its increased anxiety-like behavior and altered gait [19]. *Dlic1*^{N235Y} was thought to be a gain-of-function mutation.

To better understand the *in vivo* function of *Dlic1* in mammals, we deleted the *Dlic1* gene in mice. We found that *Dlic1* is not essential for mouse embryonic development or adult survival. The ablation of *Dlic1* in mice results in impaired OS growth and ciliogenesis, and photoreceptor degeneration. *Dlic1* deficiency leads to the ectopic accumulation of OS proteins, impaired ER export, disturbance of Rab11-vesicle trafficking, and the reduced protein level and altered distribution of dynein subunits. Our data supports the notion that *Dlic1* plays an important role in keeping dynein function and is necessary for photoreceptor development and survival.

Results

Establishment of *Dlic1*^{-/-} mice

We first generated a modified *Dlic1* allele (Figure 1A) by homologous recombination in mouse embryo stem (ES) cells. Targeted ES clones containing the modified allele were identified by PCR and verified by Southern blot (Figure 1B). In the modified *Dlic1* allele, a bacterial β -gal reporter gene with a splicing acceptor and a neo expression cassette flanked by FRT sites were inserted after exon 4, and then exon 5 was flanked by loxp sites (Figure 1A). The mosaic mice were created by injecting targeted ES clones into C57BL/6 blastocysts, and then bred with PGK-Cre transgenic mice to generate *Dlic1*^{+/^{galeo}} mice. In the *Dlic1*^{galeo} mutant allele, in which the exon 5 was deleted, the expression of β -gal reporter was under the control of endogenous *Dlic1* promoter and the β -gal reporter was fused in frame with the first 189 amino acids of DLIC1 protein after splicing. Therefore, the expression pattern of *Dlic1* in *Dlic1*^{+/^{galeo}} mice can be examined using β -gal as a reporter. To establish *Dlic1*^{+/-} mice, the *Dlic1*^{+/^{galeo}} mice were crossed with Act-*Flp-e* transgenic mice to remove the β -gal reporter gene and the Neo expression cassette simultaneously (Figure 1A). The *Dlic1* gene has 13 exons coding for 523 amino acids. Deletion of exon 5 from *Dlic1* resulted in a nonsense mutation at the beginning of exon 6, therefore generating a *Dlic1*-null mutation. This was further confirmed by analyzing the expression of DLIC1 proteins in the *Dlic1*^{-/-} mouse brain (Figure 1C).

Retinal degeneration in *Dlic1*^{-/-} mice

Dlic1^{-/-} mice are viable, fertile and show lower body weight (Supplementary information, Figure S1A and S1B). To help to study the physiological role of *Dlic1* in mice, we examined the expression pattern of *Dlic1* in adult *Dlic1*^{galeo/galeo} mice using the inserted β -gal as a reporter. We found that *Dlic1* was highly expressed in the outer nuclear layer (ONL) of mouse retinas, cerebella and hippocampi (Supplementary information, Figure S1C). The results also revealed that the ONL thickness of *Dlic1*^{-/-} retinas was reduced compared with the controls (Supplementary information, Figure S1C, top panels), suggesting that the *Dlic1* gene may be involved in the development or maintenance of photoreceptor cells.

To further investigate the roles of *Dlic1* in the development of mouse retinas, we performed a pathological analysis of *Dlic1*^{-/-} retinas. As the body weight and the ONL thickness of *Dlic1*^{+/-} mice are not obviously different from those of wild types, *Dlic1*^{+/-} mice were used as controls. The results showed that the *Dlic1*^{-/-} retina displayed a gradual loss of photoreceptor cells.

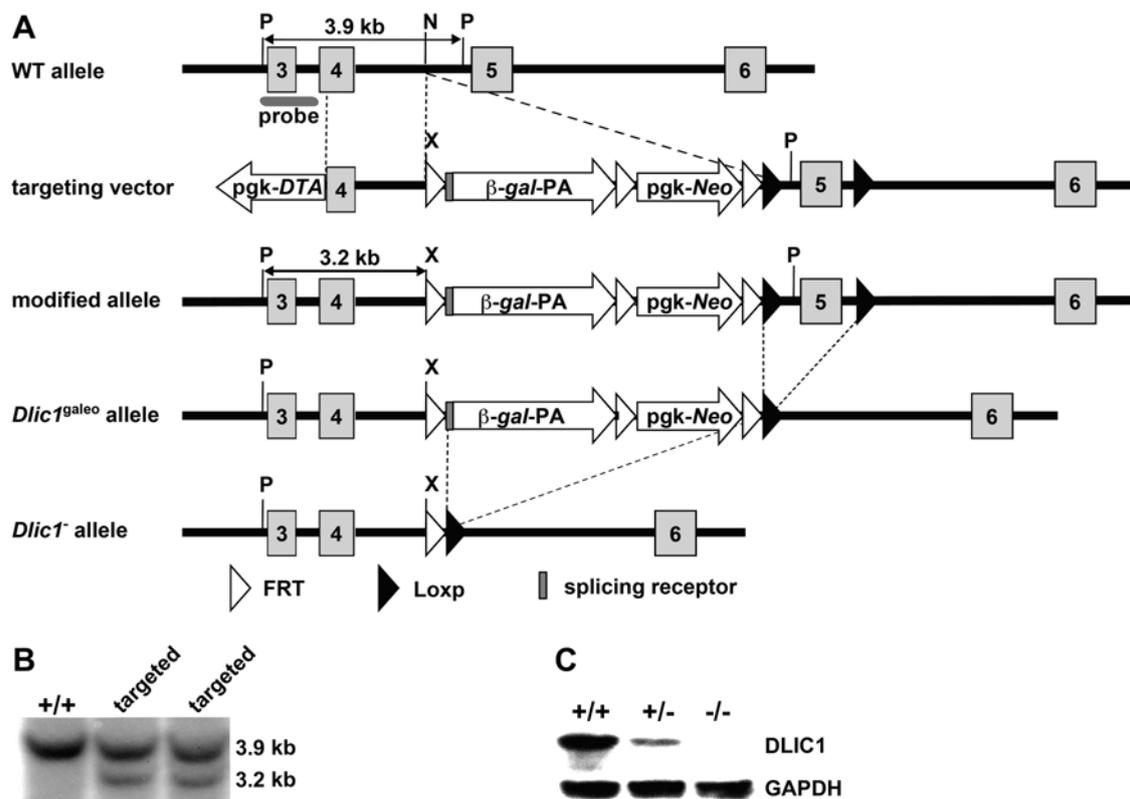


Figure 1 The generation of *Dlic1* knockout mice. **(A)** Schematic strategy to generate *Dlic1*^{-/-} mice. WT allele: genomic DNA fragment of the WT *Dlic1* gene containing exons 3-6. Targeting vector: schematic structure of the *Dlic1* targeting vector. Modified allele: genomic structure of the *Dlic1* modified allele after homologous recombination. *Dlic1^{galeo}* allele: the *Dlic1* null allele carrying the *β-gal* reporter fused in frame to exon 4 of *Dlic1* after removing a fragment containing exon 5 by CRE. *Dlic1⁻* allele: the *Dlic1* null allele derived from the *Dlic1^{galeo}* allele by removing *β-gal* and *Neo* expression cassette by FLP. *pgk-DTA* and *pgk-Neo* represent the diphtheria toxin A and the neomycin expression cassettes, respectively. *β-gal* represents a modified *E. coli* *β*-galactosidase gene containing a nuclear localization signal at its N-terminus. The arrows indicate the transcriptional direction of *DTA*, *β-gal* and *Neo* genes. Exons are numbered and depicted by grey boxes. FRT and loxP sequences are represented by white and black arrowheads, respectively. P, N and X indicate the *Puv* II, *Nhe* I and *Xho* I restriction sites, respectively (Note: *Puv* II and *Nhe* I are not unique sites). The gray ellipse under exon 3 in the WT allele depicts the probe used for Southern blot analysis. The expected sizes of the restriction fragments hybridized with the probe are indicated in WT and modified alleles. **(B)** Southern blot analysis of genomic DNA extracted from WT and targeted ES cell clones and digested with *Pvu* II and *Xho* I restriction enzymes. The 3.9 kb and 3.2 kb fragments represent WT and modified alleles, respectively. **(C)** Western blot analysis of the DLIC1 protein expression in mouse brains using anti-DLIC1 antibody. GAPDH was used as loading control. ^{+/+}, WT; ^{+/-}, *Dlic1* heterozygote; ^{-/-}, *Dlic1* homozygote.

By 12 days after birth (P12), the thickness of the ONL in *Dlic1*^{-/-} and control retinas was comparable at 11-13 rows of nuclei (Figure 2A and 2B left panel). However, the thickness of ONL of one month (1M) *Dlic1*^{-/-} retinas was significantly reduced to ~6 rows of nuclei (Figure 2A and 2B left panel), suggesting a dramatic loss of photoreceptor cells in *Dlic1*^{-/-} retinas at age 1M. This loss of cells continued as the *Dlic1*^{-/-} mice aged, but at a much slower rate (Figure 2A and 2B left panel).

Further TUNEL analysis showed that many TUNEL-positive signals were detected in the ONL of P20 (post-

natal day 20) *Dlic1*^{-/-} retinas, while very few to no TUNEL-positive signals were found in the ONL of P12 *Dlic1*^{-/-} retinas (Figure 2C and 2D). Consistent with the reduced loss of photoreceptor cells in older *Dlic1*^{-/-} mice, TUNEL signals in the ONL of *Dlic1*^{-/-} retinas were gradually reduced after one month of age, and only a few TUNEL signals were still detected in 10-month *Dlic1*^{-/-} retinas (Figure 2C and 2D). These data demonstrate that the gradual loss of the photoreceptors in the *Dlic1*^{-/-} retina is mainly caused by apoptosis. Furthermore, we also found that the expression of glial fibrillary acidic protein

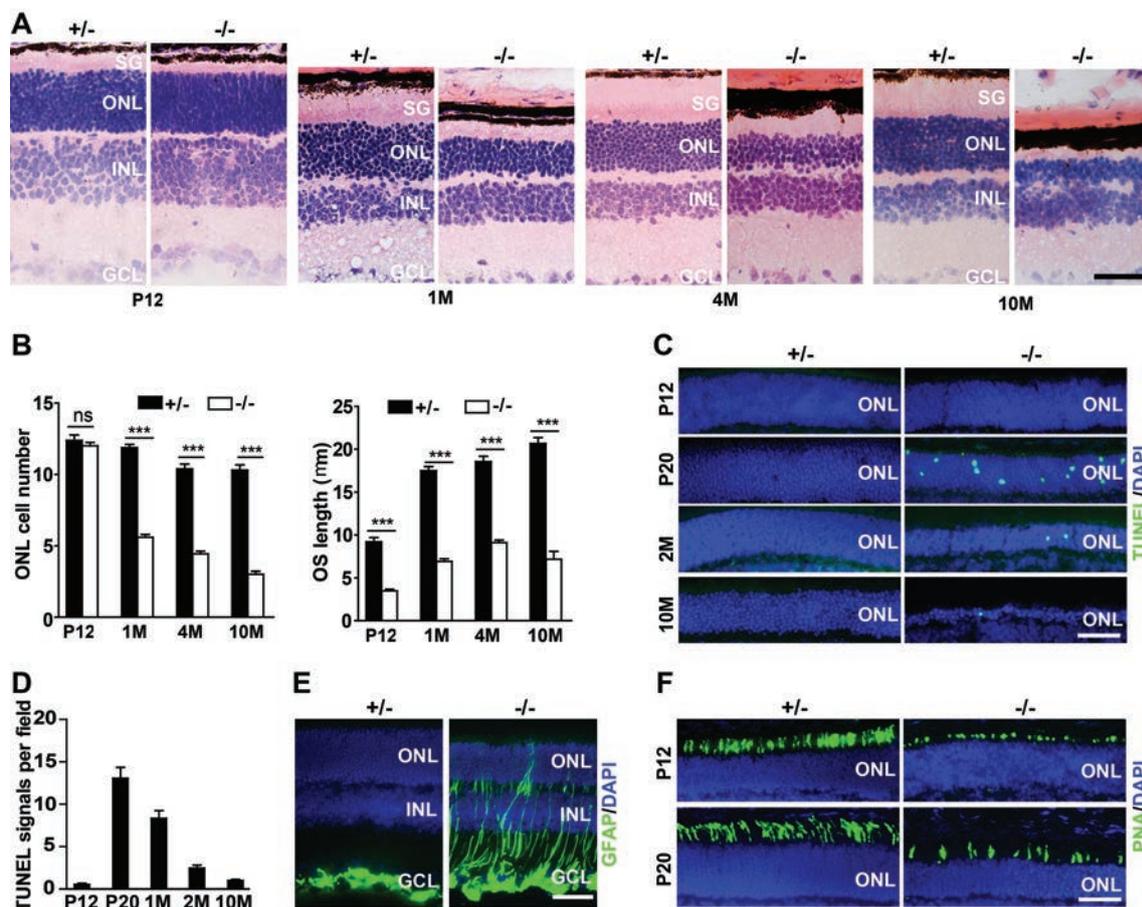


Figure 2 Photoreceptor degeneration and impaired OS development in the *Dlic1*^{-/-} mouse retina. **(A)** Images of HE-stained cryosections of *Dlic1*^{+/-} and *Dlic1*^{-/-} mouse retinas at the indicated ages. **(B)** Quantitative analysis of the thickness of the ONL (left) and length of the OS (right) in *Dlic1*^{+/-} and *Dlic1*^{-/-} mouse retinas at the indicated ages. **(C, D)** Representative images of TUNEL assay **(C)** and quantitative analysis **(D)** of apoptotic cells in the ONL of *Dlic1*^{+/-} and *Dlic1*^{-/-} retinas at the indicated ages. **(E)** Representative fluorescent images of cryosections of P20 *Dlic1*^{+/-} and *Dlic1*^{-/-} retinas stained with anti-GFAP show upregulation of GFAP in *Dlic1*^{-/-} Müller glia. **(F)** Representative images of PNA-stained cone cells in *Dlic1*^{+/-} and *Dlic1*^{-/-} retinas at P12 and P20 show the reduced number and impaired OS development of *Dlic1*^{-/-} cone cells. Cell nuclei in **C, E** and **F** were stained with DAPI. Images are representative retina sections from at least three mice per group. SG = OS + IS; ONL, outer nuclear layer; INL, inner nuclear layer; GCL, ganglion cell layer; M, month; ^{+/-}, *Dlic1*^{+/-}; ^{-/-}, *Dlic1*^{-/-}. The values in **B** and **D** represent the means ± SEM of three mice. ns, not significant; ****P* < 0.001. Bar = 50 μm.

(GFAP) was upregulated in *Dlic1*^{-/-} retinas (Figure 2E), indicating that Müller cells were activated for proliferation as a result of retinal damage [20].

All the results above suggest that ablation of the *Dlic1* gene in the mouse results in photoreceptor degeneration.

Ablation of *Dlic1* impairs the development of photoreceptor cells

Detailed pathological analysis also revealed that the length of the OS of P12 *Dlic1*^{-/-} retina was significantly shorter compared with the control although no obvious cell loss in the ONL could be observed in mutant retinas

(Figure 2A most left panel, and 2B). The OS length of the mutant retina in aged mice never reached that of their controls (Figure 2A and 2B, right panel). However, ultra-structure analysis with electron microscopy demonstrated that the disk membranes remained well ordered and tightly stacked in the OS of *Dlic1*^{-/-} photoreceptor cells (Supplementary information, Figure S2). These results suggest that *Dlic1* deficiency impairs the OS growth, but has no obvious effect on the disk membrane structures of the OS in the *Dlic1*^{-/-} retina.

The ONL of mouse retina is composed of ~95% rod and ~5% cone nuclei. Thus, the observations of shorter

OS length and the reduced ONL cell number in the *Dlic1*^{-/-} retina strongly suggest that the OS development and survival of rod cells are affected by *Dlic1* deficiency. However, it is not clear whether cone cells are also affected in the *Dlic1*^{-/-} retina. In order to determine the effect of *Dlic1* deficiency on cone cells, we performed immunofluorescent staining with FITC-conjugated peanut agglutinin (PNA), which specifically labels cone photoreceptors. Similarly to *Dlic1*^{-/-} rod cells, the results showed not only a decrease in the OS length of cone cells, but also a dramatic decrease in the number of cone cells in P12 *Dlic1*^{-/-} retinas (Figure 2F, top panels). The similar phenotypes were also observed in older *Dlic1*^{-/-} retinas (Figure 2F, bottom panels). However, almost no TUNEL signal could be detected in P12 *Dlic1*^{-/-} retina (Figure 2C). Therefore, these results suggest that the development of cone cells is also impaired.

Dlic1 deficiency leads to impairment of primary ciliogenesis

The OS of photoreceptor cells is a specialized cilium. Accumulating evidence indicates that mutations in genes involved in ciliogenesis and/or protein transport to the OS generally affect OS development and finally lead to photoreceptor degeneration [4]. Severe photoreceptor degeneration and shorter OS in *Dlic1*^{-/-} retinas (Figure 2A-2C) suggest the effects of *Dlic1* deficiency on ciliogenesis and/or protein transport to the OS. Therefore, we first verified the effect of *Dlic1* deficiency on ciliogenesis by examining the development of connecting cilia in *Dlic1*^{-/-} photoreceptors at ages of P12 and P20, when photoreceptor OS was growing [21]. Cryosections of *Dlic1*^{-/-} and control retinas were immunofluorescent stained with an antibody against acetylated-tubulin (Ac-tubulin), a marker for primary cilia. Notably, we found that the mean cilium lengths were significantly shorter in P12 and P20 *Dlic1*^{-/-} photoreceptors (Figure 3A-3C, $0.59 \pm 0.02 \mu\text{m}$ ($n = 249$) and $0.45 \pm 0.02 \mu\text{m}$ ($n = 171$) for the control and mutant at P12, respectively; $1.01 \pm 0.05 \mu\text{m}$ ($n = 117$) and $0.84 \pm 0.04 \mu\text{m}$ ($n = 136$) for the control and mutant at P20, respectively). These results indicate that ciliogenesis is impaired in *Dlic1*^{-/-} photoreceptor cells.

To further ascertain the effect of *Dlic1* deficiency on ciliogenesis, we also examined ciliogenesis in *Dlic1*^{-/-} MEFs. *Dlic1*^{-/-} and control MEFs were first starved for serum to induce ciliogenesis, and then the number of cells with cilia and the length of primary cilia were evaluated by immunofluorescent staining with anti-Ac-tubulin. The experiments revealed that both the length of primary cilia and the percentage of ciliated cells in *Dlic1*^{-/-} MEFs were reduced compared with those in the

control (Figure 3D and 3E). Therefore, we demonstrate that *Dlic1* deficiency impairs primary ciliogenesis in both photoreceptor and MEF cells.

Loss of *Dlic1* does not affect apical location of centrioles during ciliogenesis

In mammalian photoreceptor precursor cells, a pair of centrioles complete its migration towards the apical cell membrane by midgestation, and the basal body is developed from the mother centriole and remains in that position [22]. The apical localization of centrioles is thought to be one of the prerequisites for primary ciliogenesis [23, 24]. As dynein plays a key role in the movement of centrosomes in many cellular events, such as cell division and cell differentiation [5], loss of *Dlic1* might impair ciliogenesis by affecting the centriole localization. To understand the mechanisms underlying the impairment of ciliogenesis in *Dlic1*^{-/-} photoreceptor cells, we examined the location of basal bodies in P5 *Dlic1*^{-/-} retinas by immunofluorescent staining with an antibody against PCNT, which is a component of the basal body complexes of the CC in photoreceptors [25]. Our study revealed that the basal bodies in both P5 *Dlic1*^{-/-} and the control retinas were properly localized in the apical region of the IS of photoreceptors (Figure 3F), suggesting that *Dlic1* deficiency does not affect the apical localization of basal bodies in photoreceptor cells before the OS starts to develop.

Disrupted Rab11-vesicle transport from the Golgi to the basal body in *Dlic1*^{-/-} photoreceptors

The Rab11-mediated vesicle transport from the Golgi to the basal body is required for ciliogenesis [26, 27]. Rab11 was linked to cytoplasmic dynein via the direct interaction of Rab11-FIP3 with DLIC1 or DLIC2 [13, 28]. On the basis of these previous studies, we questioned whether *Dlic1* deficiency might impair ciliogenesis by blocking the transport of Rab11-mediated vesicle from the Golgi to cilia due to the reduced interaction between dynein and Rab11. To address the question, we first examined the localization of Rab11 in *Dlic1*^{-/-} retinas. The results showed that large quantities of Rab11 were localized in the ONL layer (perinuclei) of *Dlic1*^{-/-} retina, while they were mainly localized in the IS of control retinas (Figure 3G), demonstrating the blockage of Rab11-vesicle transport from the Golgi to cilia. Next we examined whether the interaction between dynein and Rab11 decreased in *Dlic1*^{-/-} photoreceptor cells by immunoprecipitation and western blot. The analysis showed that the protein level of Rab11 coimmunoprecipitated with dynein was reduced (Figure 3H), indicating that the decreased ability of dynein to capture the Rab11-vesicle

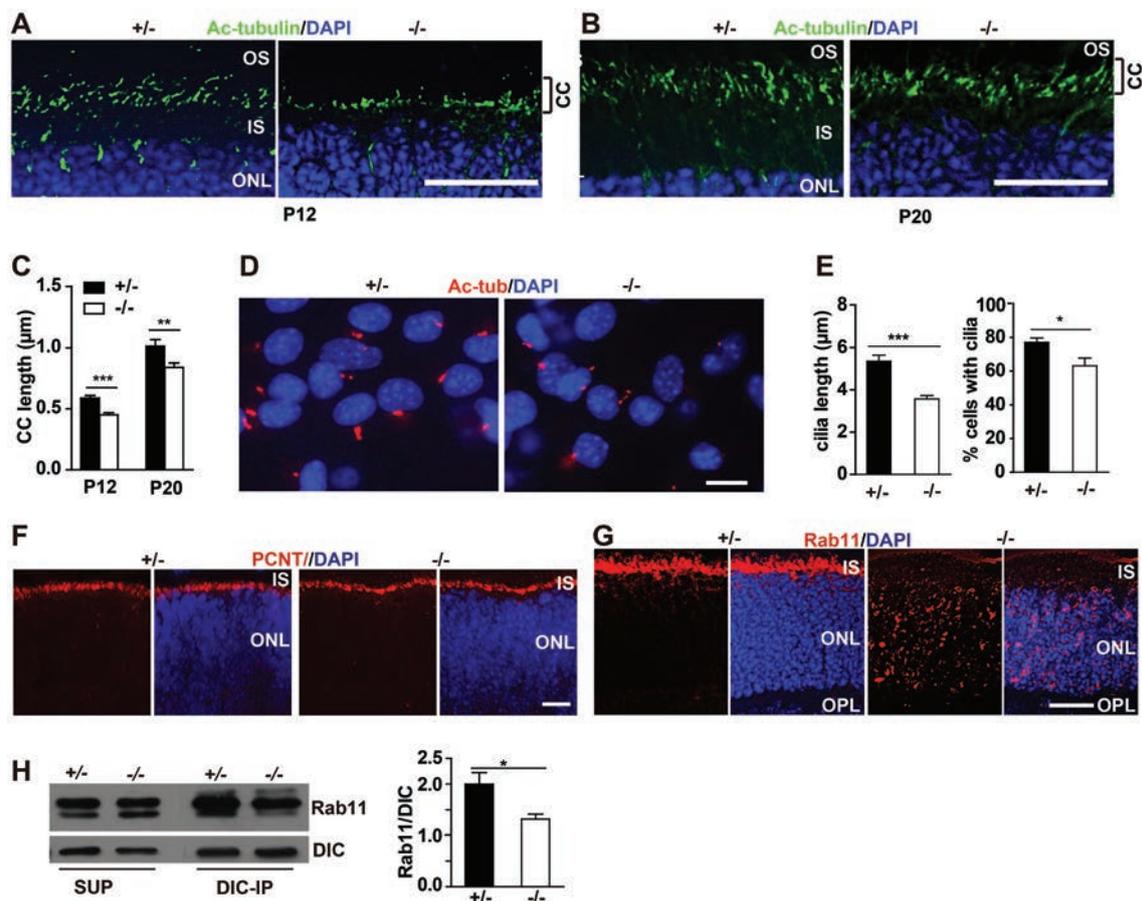


Figure 3 *Dlic1* deletion impairs primary ciliogenesis by disrupting Rab11-vesicle transport. **(A, B)** Impaired development of the connecting cilium in the *Dlic1*^{-/-} photoreceptors. Cryosections of the *Dlic1*^{+/+} and *Dlic1*^{-/-} retinas at age of P12 **(A)** and P20 **(B)** were immunofluorescent stained with anti-Ac-tubulin. **(C)** Statistical analysis of the length of the connecting cilia in *Dlic1*^{+/+} and *Dlic1*^{-/-} mouse photoreceptors at the indicated ages. Length of cilia in five random areas per section were measured and retinas from three mice in each group were used for the analysis. **(D, E)** Impaired ciliogenesis in *Dlic1*^{-/-} MEFs. Primary ciliogenesis of *Dlic1*^{+/+} and *Dlic1*^{-/-} MEFs was induced by serum starvation. Images of primary cilia stained with anti-Ac-tubulin **(D)** and statistical analysis of the cilia length **(E, left panel)** and the percentage of cells with cilia **(E, right panel)** in serum-starved *Dlic1*^{+/+} and *Dlic1*^{-/-} MEF cells ($n = 32$ and 44 for the control and mutant respectively). **(F)** *Dlic1* deletion does not affect the distribution pattern of PCNT in photoreceptor cells. Cryosections of P12 *Dlic1*^{+/+} and *Dlic1*^{-/-} retinas were IF stained with anti-PCNT. **(G)** IF stained cryosections of P12 *Dlic1*^{+/+} and *Dlic1*^{-/-} retinas with anti-Rab11 show the misaccumulation of Rab11 vesicles in the ONL of *Dlic1*^{-/-} retina. **(H)** IP-western blot (left pane) and quantitative analyses (right panel) show that loss of *Dlic1* leads to a decreased interaction between Rab11 and dynein. Cell nuclei are stained with DAPI. CC, connecting cilium. Images are representative retina sections from at least three mice per group. The values in **C, E, H** represent the means \pm SEM. * $P < 0.05$, ** $P < 0.01$, *** $P < 0.001$. Bar = 20 μm .

was due to *Dlic1* deficiency. This result provides at least one explanation for the failure of Rab11-vesicle transport. The results presented above also suggest that *Dlic1* deficiency-mediated blockage of Rab11-vesicle transport from the Golgi to cilia is one cause leading to the impairment of ciliogenesis in *Dlic1*^{-/-} photoreceptor cells.

On the basis of these results, we conclude that *Dlic1* deficiency impairs photoreceptor ciliogenesis by blocking the Rab11-vesicle transport to basal bodies, but not by interfering with the apical localization of basal bodies.

Mislocalization of OS proteins in *Dlic1*^{-/-} photoreceptor cells

The photoreceptor OS contains large amounts of phototransduction proteins. All OS proteins are synthesized in the IS. Impaired transport of OS proteins from the IS to the basal body or insufficient supply of OS proteins will impede OS development, and finally lead to photoreceptor degeneration [4, 29]. In *Drosophila*, rhodopsin normally colocalizes with Rab11 in vesicles, and the loss of Rab11 activity results in the accumulation of

rhodopsin-containing vesicles in the cytoplasm [30]. We found that *Dlic1* deficiency blocks Rab11-vesicle transport from the Golgi to cilia (Figure 3G). Therefore, it is conceivable to postulate that *Dlic1* deficiency causes OS underdevelopment by impeding OS protein transport from the IS to the OS of photoreceptor cells. To test our model, we examined the localizations of the OS proteins in *Dlic1*^{-/-} retinas by immunofluorescent staining. Our results showed that rhodopsin was ectopically accumulated in the cell body especially in the outer plexiform layer (OPL) of P20 *Dlic1*^{-/-} retina in addition to its normal OS localization (Figure 4A and 4B). We also found that the distribution of another OS protein arrestin displayed a similar ectopic accumulation in P20 *Dlic1*^{-/-} retinas (Figure 4C). These results demonstrate that *Dlic1* deficiency impairs protein transport in photoreceptor cells. We suggest that *Dlic1* deficiency-mediated impairment of protein transport provides a mechanism underlying underdevelopment of OS in *Dlic1*^{-/-} retinas.

Dlic1 deficiency does not disrupt the Golgi architecture and may not affect vesicle transport from the ER to Golgi

Rab11 vesicles and rhodopsin were both accumulated in the cytoplasm of *Dlic1*^{-/-} photoreceptor cells (Figures 3G, 4A and 4B). However, we found that their accumulation patterns in the cytoplasm of *Dlic1*^{-/-} photoreceptor cells were not exactly the same, e.g., rhodopsin heavily accumulated in the OPL of P20 *Dlic1*^{-/-} retina, while Rab11 did not. The OS proteins are synthesized in the ER, funneled through the Golgi and then transported to the OS. As Rab11 is implicated in mediating vesicles transport from the Golgi to the basal body [26, 27], the distinction of accumulation patterns between Rab11 and rhodopsin hints that *Dlic1* deficiency may also affect protein transport from the ER to Golgi, in addition to its effect on the protein transport from the Golgi to cilia. Indeed, knocking down *Dlic1* but not *Dlic2* has been reported to disrupt the Golgi architecture and block ER-to-Golgi transport in HeLa cells [14].

On the basis of these previous studies, we assumed that deleting *Dlic1* gene also resulted in impairment of protein transport from the ER to Golgi and disruption of the Golgi architecture. To validate the assumption, we first tried to verify the effects of *Dlic1* deficiency on the Golgi architecture by immunofluorescent staining with an antibody against giantin, a marker for the Golgi. Our studies showed that the Golgi apparatus was normally concentrated in the proximal portion of the IS in control retinas (Figure 5A, left panel). Our studies also revealed that the distribution of Golgi was not changed and the Golgi architecture appeared intact in *Dlic1*^{-/-} retinas (Figure 5A, right panel), suggesting that the Golgi ap-

paratus is not disrupted in *Dlic1*^{-/-} photoreceptor cells. To further confirm the integrity of the Golgi apparatus in *Dlic1*^{-/-} cells, we performed the same experiment with *Dlic1*^{-/-} MEFs. Results showed that the Golgi architecture in *Dlic1*^{-/-} MEFs was not scattered, but assembled normally at the juxtannucleus (Figure 5E, right panel), which was very similar to that in the control (Figure 5E, left panel). These results demonstrate that *Dlic1* deficiency does not disrupt the Golgi architecture in either photoreceptor or MEF cells.

Next, we investigated whether vesicle transport from the ER to Golgi was impaired by examining the localization of 1, 4-galactosyltransferase I (GalT) in *Dlic1*^{-/-} and control retinas. GalT-containing vesicles cycle slowly between the ER and Golgi [31], and GalT localizes at steady state to *trans*-Golgi cisternae [32], thereby, serving as an endogenous indicator of ER-to-Golgi transport. Our immunofluorescent staining with anti-GalT showed that unlike the giantin that was mainly located in the IS of the control retina, GalT was distributed in both the IS and the ONL (perinuclei) of the control retina, indicating that GalT is located at both the ER and Golgi in photoreceptor cells (Figure 5B, left panel). *Dlic1*^{-/-} retina displayed a GalT distribution pattern very similar to that of the control except for the distribution of GalT extended into the OPL (Figure 5B, right panel). These results indicate that (1) the ER is mislocalized in *Dlic1*^{-/-} photoreceptor cells, and (2) GalT is not a proper marker for ER-to-Golgi transport in photoreceptor cells.

To evaluate the effect of *Dlic1* deficiency on ER-to-Golgi transport, we analyzed the distribution of GalT in MEFs by immunofluorescent staining with anti-GalT. Our results showed that GalT in control MEFs displayed a juxtannuclear pattern strongly reminiscent of the Golgi apparatus, and colocalized with giantin (Figure 5E, left panel). These results are consistent with that previously reported in HeLa or other cells [14, 32], suggesting that GalT can serve as a marker for ER-to-Golgi transport in MEFs. However, we surprisingly found that GalT in *Dlic1*^{-/-} MEFs not only displayed a very similar pattern to that of the control, but also colocalized with giantin (Figure 5E, right panel), indicating that *Dlic1* deficiency may not affect normal vesicle transport from the ER to Golgi in MEFs, or have very little effect.

Dlic1 deficiency partially blocks ER export

The extension of GalT distribution into the OPL in *Dlic1*^{-/-} retinas (Figure 5B) strongly indicates a mislocalization of the ER in *Dlic1*^{-/-} photoreceptor cells. To ascertain the mislocalization of the ER in *Dlic1*^{-/-} photoreceptors, we examined the ER distribution in *Dlic1*^{-/-} photoreceptors by immunofluorescent staining with an

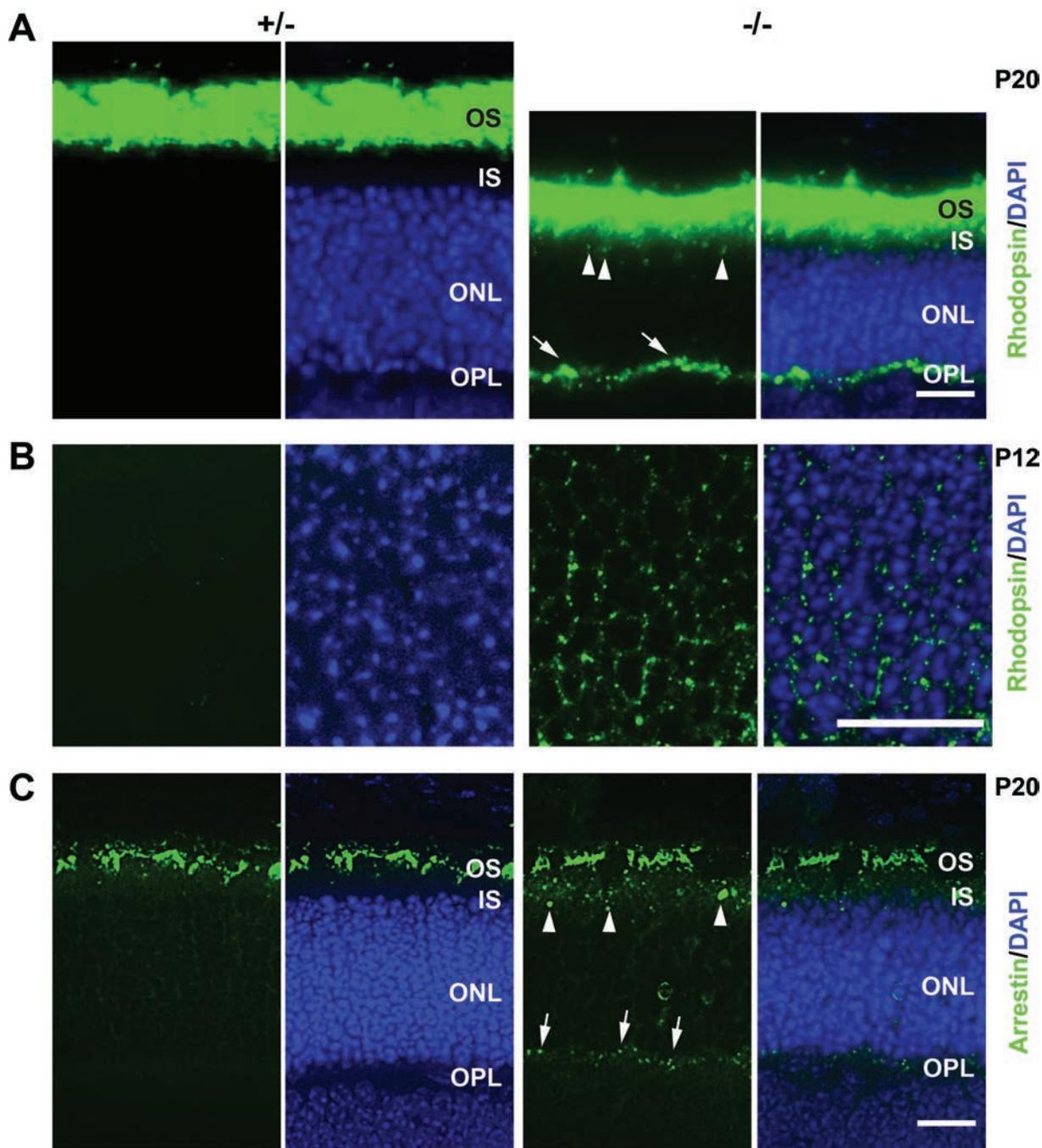


Figure 4 Mislocalization of OS proteins in *Dlic1*^{-/-} photoreceptor cells. **(A)** Immunofluorescent staining of cryosections of P20 *Dlic1*^{+/+} and *Dlic1*^{-/-} retinas with anti-rhodopsin reveals the mislocalization of rhodopsin in *Dlic1*^{-/-} photoreceptor cells. **(B)** Higher magnification of P12 mouse retinas immunofluorescent -stained with anti-rhodopsin. Perinuclear mislocalized rhodopsin is only shown in the *Dlic1*^{-/-} retina. **(C)** Immunofluorescent-stained cryosections of P20 *Dlic1*^{+/+} and *Dlic1*^{-/-} retinas with anti-arrestin. Arrows and arrowheads in **A** and **C** indicate the mislocalization of rhodopsin or arrestin in the OPL and the IS, respectively. Cell nuclei were stained with DAPI. Images are representative retina sections from at least three mice per group. Bar = 20 μm.

antibody against calnexin (an ER-specific protein). The results revealed that the ER was accumulated in the OPL

of the *Dlic1*^{-/-} retina in addition to its distribution in the IS and the ONL (perinuclei), while it was only scat-

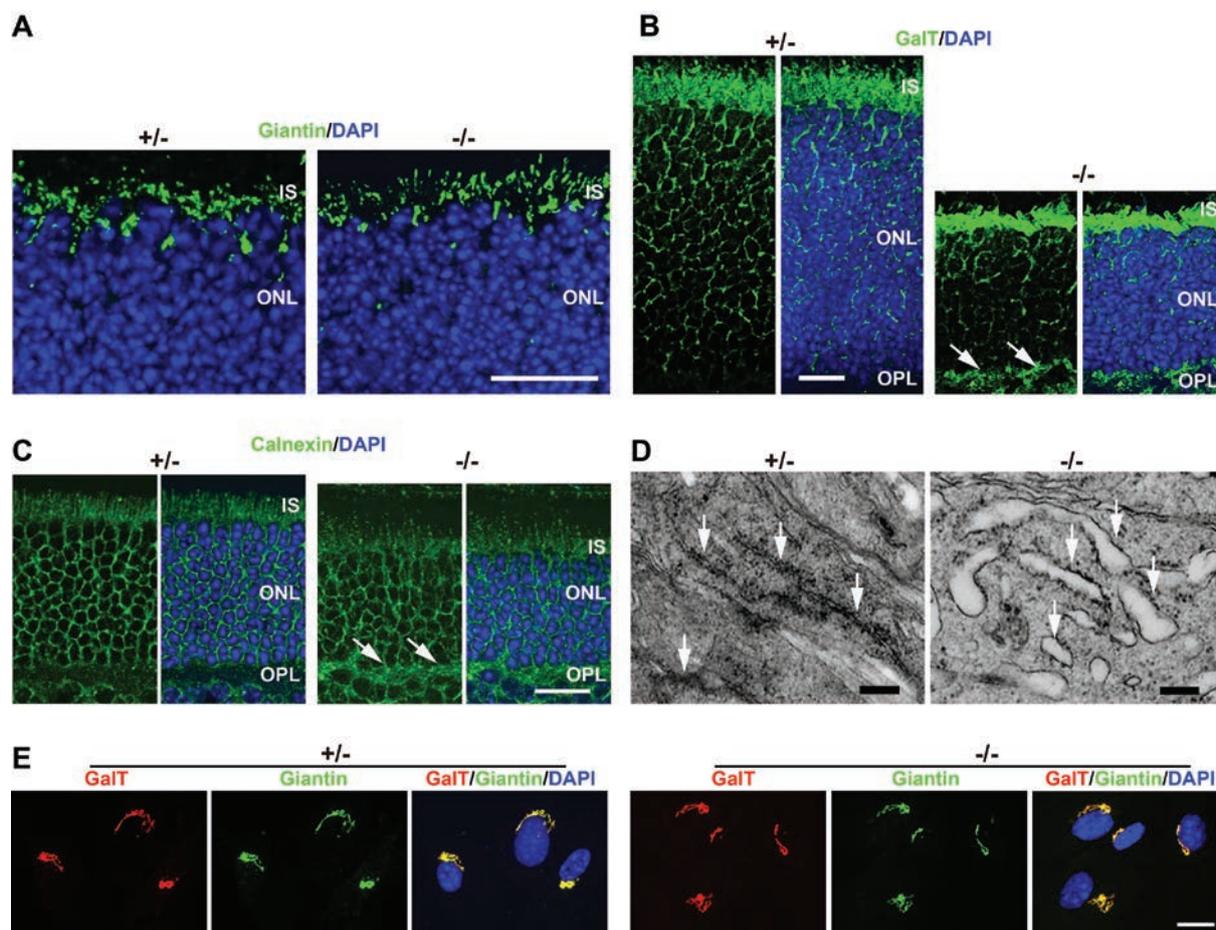


Figure 5 The effect of *Dlic1* deficiency on the ER-to-Golgi transport in photoreceptors. **(A–C)** *Dlic1* deficiency does not affect the Golgi structure and distribution, but leads to the mislocalization of GalT and calnexin in the OPL. Cryosections of P20 *Dlic1*^{+/+} and *Dlic1*^{-/-} retinas were stained with anti-giantin **(A)**, anti-GalT **(B)** and anti-calnexin **(C)** separately. Arrows in **B** and **C** indicate the mislocalization of GalT and calnexin in the OPL, respectively. **(D)** Electron micrographs of photoreceptors in P20 *Dlic1*^{+/+} and *Dlic1*^{-/-} mice, showing the ER in the IS. Arrows in the left and right panels indicate normal or dilated ER, respectively. **(E)** Co-immunofluorescent staining of *Dlic1*^{+/+} and *Dlic1*^{-/-} MEFs with anti-GalT and anti-Giantin shows the colocalization of GalT and Giantin signals in both mutant and control MEFs. Images are representative retina sections or MEFs from at least three mice per group. Bar = 20 μ m **(A–C, E)**; 250 nm **(D)**.

tered in the IS and the ONL of the control retinas (Figure 5C). Thus, we verified the mislocalization of the ER in *Dlic1*^{-/-} photoreceptors. We suspect that it is probably due to *Dlic1* deficiency-mediated impairment of dynein transport function (see discussion).

As both the ER (Figure 5C) and rhodopsin (Figure 4A) were aberrantly accumulated in the OPL of *Dlic1*^{-/-} retinas, we suspected that mislocalization of rhodopsin could also be partially because of the aberrant accumulation of rhodopsin in the ER. If this were the case, rhodopsins should be colocalized with the ER marker, calnexin. Indeed, our co-immunofluorescent staining experiments with anti-rhodopsin and anti-calnexin showed that signals of colocalization of these two proteins were detected

not only in the OPL, but also in the ONL layer (perinuclei) of *Dlic1*^{-/-} retinas, while few to none were found in the controls (Supplementary Information, Figure S3). Ultrastructure analysis with electron microscopy also revealed that the ER in *Dlic1*^{-/-} photoreceptors was dilated (Figure 5D), a cellular phenomenon usually resulting from the aberrant accumulation of unfolded or misfolded proteins in the ER or treatment with an ER stressor [33]. The results presented above strongly indicate that *Dlic1* deficiency partially blocks vesicle export from ER. Therefore, we suggest that the aberrant accumulation of rhodopsin in the ER of *Dlic1*^{-/-} photoreceptors provides another mechanism underlying mislocalization of rhodopsin in *Dlic1*^{-/-} retinas.

DLIC1 is essential for the stability and proper distribution of dynein subunits in mouse photoreceptor cells

The mutant phenotypes of defective protein transport from the Golgi to cilium and mislocalization of the ER in the OPL of *Dlic1*^{-/-} retinas strongly suggest the dysfunction of dynein transport. Loss of the DLIC protein in *Drosophila* and *Aspergillus* results in destabilization of both the DHC and DIC and the dynein complex [10, 15], although depleting mammalian DLIC1 was reported to have no effects on the stability of dynein subunits or integrity of dynein complex [11, 14]. To understand the molecular mechanisms by which the *Dlic1* mutation impairs transport function of dynein in photoreceptor cells, we examined the effect of *Dlic1* deficiency on dynein in *Dlic1*^{-/-} retinas. Western blot analysis showed that the protein levels of DHC, DIC and DLC (Tctex-1) were significantly reduced in *Dlic1*^{-/-} retinas, while the protein levels of DLIC2 was not significantly altered, and dynactin p150 Glued were dramatically increased (Figure 6A). The decrease of DIC protein level was also observed in other tissues/cells of the *Dlic1*^{-/-} mice, such as brain, liver, and MEFs (Figure 6B and Supplementary information, Figure S4A). To investigate whether *Dlic1* deficiency regulates the expression of dynein subunits at transcriptional level, we evaluated the expression of both *Dhc* and *Dic1* at mRNA level by RT-PCR analyses, and found that mRNA expression of *Dhc* and *Dic1* was not significantly altered compared with the control (Supplementary information, Figure S4B). These results suggest that the DLIC1 protein is essential for the stability of dynein DHC, DIC and DLC subunits, and *Dlic1* deficiency may impair the transport ability of dynein by destabilizing dynein subunits.

Dynein subunits are apically distributed in the inner segment of photoreceptor cells [2, 34]. The apical distribution of Tctex-1 was demonstrated to be correlated to the apical transport of proteins in polarized MDCK cells [35]. If the apically distributed dynein is self-driven, then we expect that *Dlic1* deficiency should result in the loss or disturbance of the apical distribution of dynein. To address this question, we analyzed the distributions of DHC and DIC in *Dlic1*^{-/-} retinas by immunofluorescent staining. The results showed that the distributions of DHC and DIC in *Dlic1*^{-/-} retinas were dramatically disturbed (Figure 6C and 6D). For DHC, the control retina displayed very strong signals in the IS and the OPL; however, the *Dlic1*^{-/-} retina showed much weaker signals in the IS, although the DHC signal in OPL was similar to that of the control (Figure 6C). While there were strong DIC signals in the IS of the control retina, there were only diminished DIC signals in the IS of the *Dlic1*^{-/-} retina (Figure 6D). These results strongly demonstrate that

retrograde transport of dynein complex without DLIC1 subunit in *Dlic1*^{-/-} retina is impaired.

Defective ERG responses of the Dlic1^{-/-} photoreceptors

Finally, we determined whether *Dlic1* deletion affects retinal function. We used electroretinography (ERG) to analyze the rod- and cone-driven circuits in both scotopic and photopic conditions. Under scotopic and photopic conditions, the ERG detects responses from rod- and cone-driven circuits, respectively. We compared a-wave and b-wave of *Dlic1*^{-/-} mice under both scotopic and photopic conditions with those of control mice (Figure 7A-7D). Under scotopic conditions, while the average saturated a-wave and b-wave represent 373 μ V and 875 μ V, respectively, in control mice, the average saturated a-wave and b-wave were reduced to 104 μ V and 379 μ V correspondingly in *Dlic1*^{-/-} mice (Figure 7A and 7C). These data illustrate that the rod-driven circuit is significantly affected in the *Dlic1*^{-/-} mouse retina. Similar to the scotopic ERG responses, photopic ERG responses were weakened both at a-wave and b-wave in *Dlic1*^{-/-} mice. Under the photopic conditions, the a-wave and b-wave represent 10 μ V and 117 μ V in control mice and 0.9 μ V and 16 μ V in *Dlic1*^{-/-} mice, respectively (Figure 7B and 7D). These results demonstrate that both the scotopic and photopic responses of the *Dlic1*^{-/-} retina are impaired.

Discussion

Accumulating evidence demonstrates that mutations in genes encoding motor proteins, such as Kinesin-2 and dynein 2 responsible for anterograde and retrograde intraflagellar transport, respectively, result in ciliopathy [36]. Tctex-1, a light chain subunit of cytoplasmic dynein 1, has been recently proposed as a component of dynein 2 to negatively regulate cilium length [37] and/or participate in cilium disassembly, but not ciliogenesis in a dynein-independent manner [38]. We have now shown that primary ciliogenesis is impaired in both *Dlic1*^{-/-} photoreceptor and MEF cells, thus for the first time demonstrating that DLIC1 is a positive regulator of ciliogenesis, and that cytoplasmic dynein 1 also participates in the assembly of primary cilia. We have found that the apical localization of centrioles is not affected before the development of the OS, but Rab11 vesicles were ectopically localized in the ONL of *Dlic1*^{-/-} retinas. As Rab11-mediated vesicle transport from Golgi to the basal body is required for ciliogenesis, we conclude that *Dlic1* deficiency impairs ciliogenesis and OS development by blocking polarized vesicle transport, but not by interfering with the apical localization of basal bodies/centrioles.

Our study shows that *Dlic1* deficiency results in ecto-

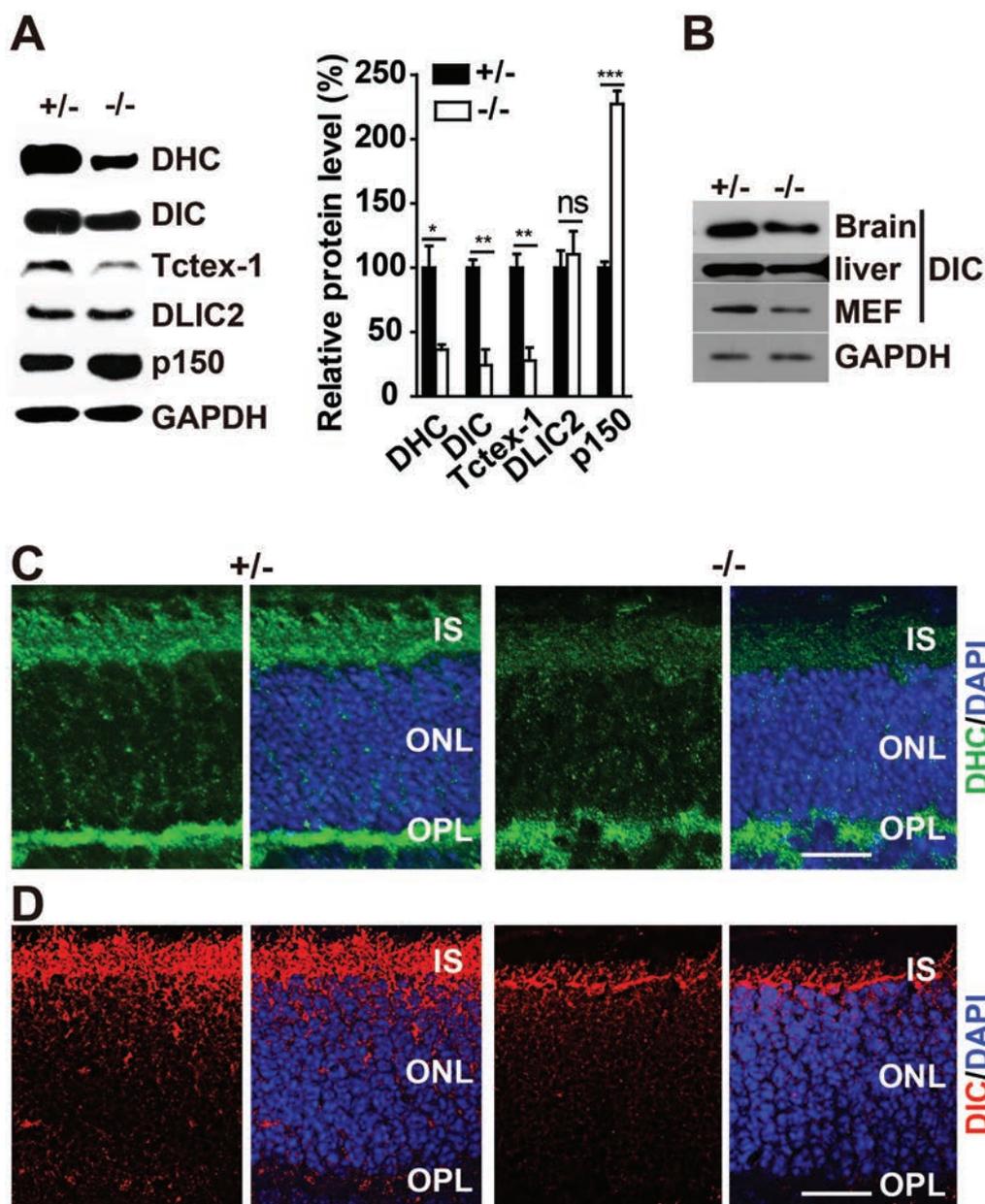


Figure 6 Deletion of *Dlic1* disrupts the distribution of dynein in mouse retinas and destabilizes other subunits of dynein. **(A)** Ablation of *Dlic1* leads to the decreased protein levels of dynein subunits in retinal cells. Western blot (left panel) and quantitative (right panel) analyses of the protein levels of dynein subunits in retinal cells. **(B)** The reduction of DIC level in *Dlic1*^{-/-} brain, liver and MEFs. Cell lysates from *Dlic1*^{+/-} and *Dlic1*^{-/-} retinas **(A)** or other tissues/cells as shown **(B)** were immunoblotted with the indicated antibodies. GAPDH was used as a loading control. **(C, D)** Loss of *Dlic1* results in the disruption of distribution of DHC and DIC in mouse retinas. Cryosections of P12 *Dlic1*^{+/-} and *Dlic1*^{-/-} retinas were immunofluorescently stained with anti-DHC **(C)** and anti-DIC **(D)**. Cell nuclei were stained with DAPI. Images and blots are representative from at least three mice per group. The values in **A** represent the means ± SEM of three mice. ns, not significant; **P* < 0.05, ***P* < 0.01, ****P* < 0.001. Bar = 20 μm.

pic accumulation of Rab11 vesicles in the ONL of retinas. Rab11 vesicle transport from the Golgi to the basal body is dynein dependent [13], and Golgi and the basal body are IS features in photoreceptor cells. However,

how does *Dlic1* deficiency-mediated loss of function for dynein lead to accumulation of Rab11 vesicles in the ONL rather than at an origin or terminus within the IS? In highly polarized photoreceptor cells, microtubules are

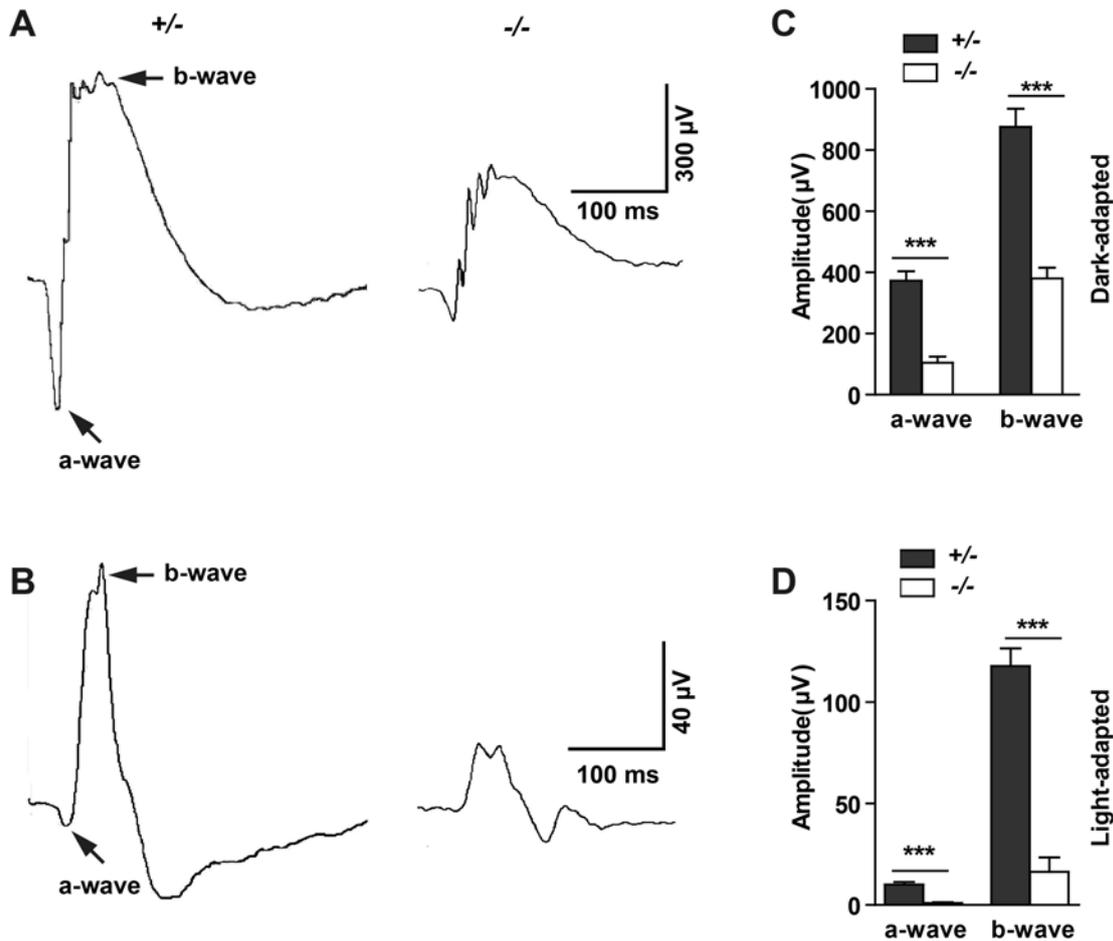


Figure 7 Functional characteristics of *Dlic1*^{+/-} and *Dlic1*^{-/-} retinas. **(A, B)** Saturating ERG responses of the retinas of the mice as indicated to 510 nm flashes at an intensity of -35 log scot. cd. s/m² under scotopic **(A)** and photopic **(B)** conditions. Each trace is the average of individual records from five mice. **(C, D)** Statistical analyses of the saturating amplitude of a- and b-waves of the mice as indicated under scotopic **(C)** and photopic **(D)** conditions. The *Dlic1*^{+/-} and *Dlic1*^{-/-} mice were 10-month-old littermates. Values in **C** and **D** represent the means ± SEM of five mice. ****P* < 0.001.

arranged with their minus ends clustered around a basal body near connecting cilium and their plus ends pointing towards OPL. Many of cargoes or vesicles including endosomes and ER are transported along polarized microtubules bidirectionally with plus- and minus-end motors present on the surface of cargo simultaneously. The direction of the net transport depends on the balance between plus- and minus-end directed motions. Thus, a possible explanation for ectopic accumulation of Rab11 vesicles could be that *Dlic1* deficiency results in an imbalance of cargo transportation between plus- and minus-end directions in cells. Impairing or enhancing minus-end motors should improve or reduce motion in the direction of plus-end motors and vice versa [39, 40]. We show here that deletion of *Dlic1* results in decreased protein levels of multiple dynein subunits, including DHC,

DIC and DLC (Tetx-1), but not DLC2. Therefore, *Dlic1* deficiency reduces the total number of minus-end motors in *Dlic1*^{-/-} photoreceptor cells, and the plus-end trafficking is then enhanced. This will facilitate the travel of Rab11 vesicles in the plus-end direction and retard their motion in minus-end direction. The final result is that Rab11 vesicles delocalize from their normal IS location and accumulate in the ONL.

Dlic1 deficiency results in ER mislocalization in the OPL of mouse retinas. The ER is a highly dynamic organelle that undergoes constant movement and reorganization. Dynein drives the ER mobility towards the minus end of microtubules, while kinesin drives ER movement towards the plus end of microtubules [41]. Therefore, *Dlic1* deficiency-mediated imbalance of cargo transportation between plus- and minus-end directions in cells

could also provide an explanation for ER mislocalization in the OPL of *Dlic1*^{-/-} retinas.

The role of DLIC proteins in the assembly of the core dynein complex has been controversial. The DLIC protein is required for the stability of both the DHC and DIC in *Drosophila* and *Aspergillus*, and is able to stabilize the interaction between DHC and DIC in *Aspergillus* [10, 15]. However, depleting DLIC1 or DLIC2 [11, 14] by RNAi in HeLa cells did not affect the protein levels of DHC, DIC and DLC (Tctex-1). We have now found that the deletion of *Dlic1* in mouse leads to the destabilization of DHC, DIC and DLC (Tctex-1) in many mouse tissues or cells, such as retinas and MEFs, but does not affect DLIC2. Our findings support the notion that the assembly of core dynein subunits, including the DHC, DIC and DLIC, is an interdependent process [10]. The discrepancy about the effect of DLIC1 on the stability of the dynein complex or subcomplex between HeLa cells and our *Dlic1*-deficient mouse cells may result from the different methods used to deplete DLIC1: knockdown by RNAi may not completely eliminate DLIC1 but our *Dlic1*^{-/-} mouse is a null mutant.

In mammalian cells the formation and budding of COPII-coated vesicles at ER export sites (ERES) is the first step of membrane trafficking in the secretory pathway [42]. The budded vesicles are then transported from ERES to the ER-Golgi intermediate compartment (ER-GIC) in a COPII-dependent step, and at ERGIC they are fused to form vesicular-tubular transport carriers (VTC) and finally transported to the *cis*-Golgi in a second COPII- and dynein-dependent step [43, 44]. Disruption of interaction between COPII complex and dynactin, a dynein adaptor linking dynein to cargoes, inhibits cargo export from the ER, but does not affect the motility of VTCs [44]. This study suggests that ER export may be dynein dependent. Here we show mis-accumulation of rhodopsins in dilated ER of *Dlic1*^{-/-} photoreceptors, suggesting partial blockage of vesicle export from the ER, and providing direct evidence to support the involvement of dynein or dynein subunit in ER export. However, the effect of DLIC on protein transport of dynein from the ER to Golgi is controversial, too. Palmer *et al.* [14] reported that knocking down *Dlic1* in HeLa cells resulted in blockage of vesicle export from ER and VTC transport from the ER to Golgi, and the failed VTC transport leads to fragmentation of the Golgi structure; but Sivaram *et al.* [11] showed that depleting DLIC1 in the same cell line had no effect on the Golgi structure at all. Here we demonstrate that the Golgi structure in both *Dlic1*^{-/-} photoreceptors and MEFs is intact and GalT is steady-stately localized in *trans*-Golgi cisternae in *Dlic1*^{-/-} MEFs, strongly indicating that VTC transport is not affected in

Dlic1^{-/-} cells. Therefore, we infer that *Dlic1* deficiency impairs vesicle export from ER, but not VTC transport from the ER to Golgi. The possible explanations for our results could be (1) the DLIC1 protein may play a specific role in ER export; (2) DLIC1 and DLIC2 proteins are functionally redundant for VTC transport from the ER to Golgi. However, these are our speculations and more experiments are needed to further verify the above possibilities.

Materials and Methods

Generation of *Dlic1*-deficient mice

To construct the *Dlic1* gene-targeting vector, we screened the 129/Sv mouse genomic phage library (Stratagene) and isolated a genomic DNA fragment containing *Dlic1* exon 3 to exon 9. In the gene targeting vector, we inserted a FRT-flanked β -gal reporter and neo expression cassette at the *NheI* site before exon 5, and a *Loxp*-flanked *NheI*-*SmaI* fragment containing exon 5. A diphtheria toxin A (DTA) expression cassette was used as a negative selection marker in the targeting vector. The linearized *Dlic1* targeting vector was electroporated into mouse embryo stem (ES) cells (W4/129S6 ES, Taconic Transgenic). After double selection, the recombinant ES cell clones were screened by PCR and further confirmed by Southern blot analysis with ³²P-labeled *PvuII*-*BglIII* fragment containing exon 3. Two recombinant ES clones were injected into C57BL/6J blastocysts to produce chimera mice. The chimera mice were crossed with PGK-*Cre* transgenic mice to delete exon 5 and generate *Dlic1*^{+/-galco} mice. Progenies were genotyped by PCR using primers: *Dlic1*SA: 5'-GGCAGAACTCCTAACCAGCAAG-3', *Dlic1*KO: 5'-GCTACTTATCCACAACCAACGCAC-3' and *Dlic1*WT: 5'-CATACCTGAGACCCACAAAA CAGC-3'. Then *Dlic1*^{+/-galco} mice were crossed with *Act-Flp* transgenic mice (003800, The Jackson Laboratory) to delete the β -gal and neo cassette to generate *Dlic1*^{-/-} mice. Progenies were genotyped by PCR using primers: *Dlic1*LR: 5'-TGGCTCAGTGGTAAAGGTCC-3', *Dlic1*LoxpR: 5'-TCAGGAAAAGCACTGG CTG-3', *Dlic1*FRTF: 5'-GGAAGATGTGACAAGACAGACACG-3'. All of the *Dlic1* mutant mice used for experiments were maintained on a C57BL/6 and 129/sv mixed genetic background. Mouse breeding and experimental manipulations were carried out following the general guidelines published by The Association for Assessment and Accreditation of Laboratory Animal Care. Animal-related procedures were reviewed and approved by the Animal Care and Use Committee of the Institute of Developmental Biology and Molecular Medicine at Fudan University.

Histological analysis and immunofluorescent staining

Frozen sections: after the animals were exterminated by cervical dislocation, eyes were dissected out and embedded in OCT. Samples were then quickly frozen in liquid nitrogen-cooled isopentane, and 10 μ m cryosections were collected. For histological analysis, frozen sections were stained with hematoxylin and eosin for light microscopy. OS length and ONL thickness in retinas were measured in sections (on a Leica DMRXA2 microscope at 400 \times magnification). OS length was measured along the long axis of the OS between the inner segments and the retinal pigment epithelium. ONL thickness was analyzed by counting rows of photoreceptor

cells in the outer nuclear layer. Five different areas in the central and midperipheral retina were measured and three mice for each genotype were used for statistics. Only representative images were used for statistical analysis. For immunofluorescent staining, the sections were stained with various antibodies as described previously [45] except that the sections were fixed in 0.2% PFA for 10 min for immunofluorescent staining with anti-PCNT antibody. Fluorescence micrographs were acquired using a Leica DMRXA2 fluorescence microscope equipped with a Leica DFC350FX camera. Histological micrographs were acquired using a Leica DMRXA2 fluorescence microscope equipped with a Leica DFC300FX camera. Images were processed using Adobe Photoshop.

Statistical analysis

Statistical analyses were performed using Microsoft Excel 2007 and GraphPad Prism 4. The results are presented as mean \pm SEM, and significant differences are indicated by a single asterisk * when $P < 0.05$ and a double asterisk ** when $P < 0.01$.

X-gal staining

Dlic1^{galco/galco} mice were used for X-gal staining to determine the expression pattern of *Dlic1*. Eyes were fixed in 0.2% glutaraldehyde for 10 h followed by cryoprotection in 30% sucrose for overnight at 4 °C and then 20- μ m frozen sections were collected. The X-gal staining was carried out following standard protocol [46]. The slices were subsequently counterstained with eosin.

TUNEL assays

TUNEL assays were performed as described previously [47]. For quantification, digital images were taken under fluorescence microscopy at 400 \times magnification, and the TUNEL-positive cells in the ONL of five randomly chosen fields in the central and mid-peripheral retina were counted for each retina. Three animals for each time point and each genotype were used for statistics.

Antibodies

A synthetic peptide consisting of the C-terminal 15 amino acids of the DLIC1 protein, VFPTTSPTEGEAS [12], was used to raise polyclonal antibodies. Antibodies were affinity-purified with Hitrap NHS-activated HP columns (Amersham Biosciences). Antibodies against the following proteins were used for immunofluorescence: GFAP (MAB3402, Chemicon); arrestin (PA1-731, Affinity Bioreagent); fluorescein peanut agglutinin (L-1020, Vector); calnexin (SPA-860, Stressgen); DIC (74 kDa) (MAB1618, Millipore); DHC (SC-9115, Santa Cruz); giantin (A488-114L, Covance); GalT (generously provided by Dr Jack Rohrer); tctex-1 (11954-1-AP, Proteintech). Antibodies against Rhodopsin (R5403) and Ac-tubulin (T7451) were from Sigma; Rab11 (610656), p150 Glued (610474), and PCNT (611814) were from BD Bioscience. Secondary antibodies: goat anti-rabbit IgG-FITC (F9887) and goat anti-rabbit IgG-CY3 (C2306) were from Sigma; goat anti-mouse IgG-FITC (FI-2020) was from Vector Labs. All the secondary antibodies were diluted at 1:300 when used.

Immunoprecipitation and western blot analyses

Freshly dissected eyes were enucleated and two eye cups were homogenized in 200 μ l TNP buffer [48] to make retina protein extracts. Preparation of cell lysates from other tissues, co-immunoprecipitation and western blot analyses with antibodies against

DIC, DHC, and DLIC1 were performed according to the procedures described previously [48]. For quantification of protein levels, appropriate film exposures were scanned, the density of bands were determined with image J and normalized to band intensity for GAPDH. Each experiment was repeated three times.

Quantitative PCR

Total RNA was extracted from mouse retinas using Trizol reagent (Invitrogen). cDNA was synthesized using RNA PCR kit (Takara) following the manufacturer's instructions. Quantitative PCR was carried out with 2 \times QuantiTect SYBR Green PCR master mix (Stratagen). The assay were repeated with five pairs of mice in two independent experiments. Values obtained for the target genes were calibrated with the housekeeping gene *Gapdh*. The mRNA levels of DHC and DIC were assessed as an increase or decrease in the mutant mice relative to control. Primers used were as follows: DIC-F1, 5'-AACCCCTATGTCTCCCTCTTCG-3', DIC-R1, 5'-CGTCTCCCAAGTTCTGAGTCTGAC-3', DHC-F1, 5'-GACAGGGATGGTGATAAGATGGC-3', DHC-R1, 5'-TA-ACCTTGGGCTTTTC TCCGC-3'.

Electron microscopy

Eyes were enucleated and eye cups were fixed in the fixative buffer (0.05% glutaraldehyde plus paraformaldehyde in PBS, pH 7.4) at 4 °C for at least 24 h, and then embedded in resin. Ultrathin sections were prepared using a Reichert ultramicrotome, contrasted with uranyl acetate and lead citrate and examined under a Philips CM120 electron microscope.

Mouse electroretinograms

Mouse ERG was performed according to the procedure described previously with some minor modifications [49]. Mice were dark adapted overnight and anesthetized with a mixture of ketamine (15 mg/g body weight) and xylazine (5 mg/g body weight) under dim red light. The pupils were dilated with a single drop of 1% atropine sulfate. A drop of 0.5% proparacaine hydrochloride was applied for corneal anesthesia. A small amount of 2.5% methylcellulose gel was applied to the eye. Mice were placed on a heating pad keeping at 37 °C inside a Ganzfeld dome (Roland Q400, Wiesbaden, Germany). A silver loop electrode was placed over the cornea to record the ERGs. Needle reference and ground electrodes were placed in the cheek and tail, respectively. The responses were differentially amplified (1-500 Hz). All stimuli were presented in the Ganzfeld dome. Light was spectrally filtered with a 500 nm interference filter. The intensities of flashes were -5.0, -15, -25 and -35 log scotopic candela-sec/m² (cd. s/m²). For photopic ERG, mice were recovered at intensities of 30 cd. s/m² for 10 min, and then tested at the intensity of 3.0 cd. s/m². Data were collected and analyzed with Port32.exe. The mice tested were at the age of 10 months.

Cilium induction and length measurements

We prepared MEFs from the control and *Dlic1*^{-/-} E13.5 embryos. Primary MEFs were grown in Dulbecco's modified Eagle's medium (Invitrogen) supplemented with 10% fetal bovine serum, 1% L-glutamine, and 1% penicillin-streptomycin (Invitrogen). To induce cilium growth, 3 \times 10⁵ cells were plated on a cover slide in a six-well plate, followed by serum starvation for 48 h to promote cilium growth. Then the cells were stained with anti-Ac-tubulin

antibody to label the cilium.

To quantify the cilium lengths of *Dlic1^{-/-}* photoreceptors and MEFs, fluorescent images of retinal sections or MEFs immunofluorescent-stained with anti-Ac-tubulin antibody from three independent experiments were acquired using a Leica DMRXA2 fluorescence microscope at 400× (for MEFs) or 1 000× (for photoreceptors) magnification. Cilium in images were traced and measured with software (Image J version 1.46, NIH).

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