Conditional Deletion of Cytoplasmic Dynein Heavy Chain in Postnatal Photoreceptors

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PURPOSE. Cytoplasmic dynein-1 (henceforth dynein) moves cargo in conjunction with dynactin toward the minus end of microtubules. The dynein heavy chain, DYNCH1H1, comprises the backbone of dynein, a retrograde motor. Deletion of Dync1b1 abrogates dynein function. The purpose of this communication is to demonstrate effects of photoreceptor dynein inactivation during late postnatal development and in adult retina.

METHODS. We mated Dync1b1F/F mice with iCre75 and Prom1-CreER72 mice to generate conditional rod and tamoxifen-induced knockout in rods and cones, respectively. We documented retina degeneration with confocal microscopy at postnatal day (P) 10 to P30 for the iCre75 line and 1 to 4 weeks post tamoxifen induction (wPTI) for the Prom1-CreER72 line. We performed scotopic and photopic electroretinography (ERG) at P16 to P21 and were extinguished at P30. Cones were stable under rod-knockout conditions until P21 but inactive at P30. In iCre75 Dync1b1−/− photoreceptors, the IS/OS began shortening by 3wPTI and were nearly eliminated by 4wPTI. The ONL shrank significantly over this interval, indicating rapid photoreceptor degeneration following the loss of dynein.

CONCLUSIONS. Our results demonstrate dynein is essential for the secretory pathway, formation of outer segments, and photoreceptor maintenance.

Keywords: dynein-1, dynein heavy chain, conditional knockout

Transport along microtubules is accomplished by cytoplasmic dynein,1 which moves cargo toward the minus end of microtubules at the basal body.2–4 In photoreceptors, the minus end is located at the basal body, serving as a microtubule organizing center (MTOC) for review (see Baehr et al.5). During early photoreceptor development, dynein cargo may include nuclei,5 mitochondria,6 membrane vesicles,6,9 and other organelles (reviewed in Reck-Peterson et al.10). After ciliogenesis, dynein plays a key role in the secretory pathway transporting membrane vesicles and membrane proteins from the endoplasmic reticulum to the periciliary ridge, where cargo is unloaded.11

Dynein is composed of a pair of heavy chains (DYNCH1H1) (Fig. 1A) and a set of noncatalytic accessory components termed intermediate, light intermediate, and light chains.12–14 DYNCH1H1 serves as a scaffold organizing the distribution of dynein subunits and the three-dimensional structure of the dynein megastructure (Fig. 1B).15,16 The N-terminal DYNCH1H1 tail domain contains a dimerization domain17 and binding sites for intermediate chains (DIC1 or DIC2) and light intermediate chains (DLIC1 or DLIC2). The C-terminal half contains the motor domain and microtubule interaction site. The dynein motor domain is built around a ring of six AAA+ (ATPases Associated with Various Activities) of the heavy chain. The microtubule-binding domain sits at the tip of a coiled-coiled stalk emerging from AAA4. Dynactin and a cargo adaptor are essential for cytoplasmic dynein to move membrane vesicles along microtubules.18

Numerous mutations in the human DYNCH1 gene associate with spinal muscular atrophy and Charcot–Marie–Tooth disease.19–23 Germ-line deletion of mouse Dync1b1 (truncation after exon 1) is lethal as embryos do not survive beyond embryonic day 8.5 (E8.5).24 Four mouse mutants—legs at odd angles (Loa),25,26 Cramping 1 (Cra1),27,28 Sprawling (Stel),29,30 and a model of Charcot–Marie–Tooth disease31,32—display autosomal dominant behavioral phenotypes with no reported retina phenotype. A DYNCH1 nonsense mutation (Y3102X) underlies the cannonball (cnb) phenotype in zebrafish.4 Green fluorescent protein (GFP)—tagged rhodopsin has been mislocalized in both the cannonball mutant and dync1b1 morphant cells.3

Dynein is present prominently in photoreceptor inner segments (ISs), the outer plexiform layer (OPL), and the...
Deletion of Dynein-1 in Photoreceptors

**Figure 1.** Photoreceptor-specific knockout of Dyn1b1. (A) Schematic of mouse cytoplasmic dynein heavy chain (DYNC1H1, gene symbol Dync1h1). Locations of the truncation point (red arrow), six AAA motor domains (blue), and microtubule-binding domain (green) are indicated. Locations of previously identified mutations in mice and zebrafish are shown below the protein. Dyn1b1 knockout results in truncation of DYNC1H1 in its tail domain and eliminates the motor domain. (B) Cytoplasmic dynein drawn schematically with heavy chain (blue). Adapted from Hoang et al.22 (C) loxP sites of the floxed Dync1h1 allele are located in introns 23 and 25. (D) Following Cre-induced recombination, the null allele loses exons 24 and 25. (E) PCR amplification using P10 retinal DNA as template and primers X23F and loxP-R. The amplicon at 500 bp shows Cre-induced recombination has occurred in the Dync1h1 F/F genotype, while amplicon at 1600 bp represents Dync1h1 F/+ genotype in the inner retina. (F) Genotyping of Dyn1b1 F/F and Dyn1b1 F/+ alleles with primers loxP-F and loxP-R in intron 25. M, size markers; W, water control. (G) Genotyping of iCre75. +, tail DNA containing the iCre75 transgene. (H) Genotyping of Egfp-Cetn2. +, tail DNA containing the Egfp-Cetn2 transgene. (I) Genotyping of Prom1-ETCre mice.

inner plexiform layer, as well as, to a lesser extent, within the outer nuclear layer (ONL), inner nuclear layer (INL), and ganglion cell layer. Dynein is responsible for retina lamination, nuclear positioning, and vesicular trafficking of photoreceptor membrane proteins.13,33 We previously reported that conditional knockout of Dync1h1 in mouse retina (Dyn1b1 F/F;Six3cre or Dyn1b1 F/F; ret Dync1h1 −/−) resulted in rapid retinal degeneration within 2 postnatal weeks.13 Dync1h1 was undetectable in the Dyn1b1 F/F IS area as early as postnatal day (P) 6, and photoreceptors did not elaborate IS, outer segment (OS), or functional synaptic terminals. In the P8 Dyn1b1 −/− central retina, outer and inner nuclear layers were severely disorganized and lacked a recognizable OPL.13

In this report, we depleted DYNC1H1 in rod photoreceptors during postnatal development and in both rods and cones of adult mice by tamoxifen induction. These knockouts permit analysis of the consequences of DYNC1H1...
deletion during late postnatal development in rods and in photoreceptors of adult mice, respectively.

**RESULTS**

**Generation of DYNC1H1 Conditional Knockout Mice**

The dynein heavy chain DYNC1H1 (Fig. 1A) forms the backbone of cytoplasmic dynein, a mega complex of 1.5 Mda, composed of a pair of force-generating heavy chains and a set of accessory components termed intermediate, light intermediate, and light chains (Fig. 1B). The heavy chain serves as a scaffold organizing the distribution of dynein subunits. We previously generated a floxed Dync1b1 allele (Dync1h1^F/F), in which loxP sites are placed in introns 23 and 25 (Fig. 1C). Deletion of exons 24 and 25 truncates DYNClH1 (red arrow, Fig. 1A, D) removing the C-terminal motor domain and the microtubule binding site. We mated Dync1h1^F/F mice with iCre75 transgenic mice to generate rod knockout mice (Dync1h1^F/F;iCre75, abbreviated rod Dync1h1^-/-) and with CreERT2 mice (Dync1h1^F/F;Prom1-CreERT2) to generate rod/cone knockouts following tamoxifen induction (tam Dync1h1^-/-). In iCre75 mice, Cre is expressed under the control of a rhodopsin promoter, which drives Cre in the second postnatal week. In Prom1-CreERT2 mice, CreERT2 expression occurs under the control of the Prom1 promoter driving expression in rods and cones. Deletion of the Dync1b1 gene in rod Dync1h1^-/- mice was assessed by PCR with P10 retina DNA as template and primers X23-F and loxP-R to generate a 500-bp amplicon lacking exons 24/25 (Fig. 1E). Since dynein is expressed in all retina cells, a band at 1.6 kb corresponding to the Dync1b1^F allele is also observed. We did not perform an immunoblot as a blot using total retina tissue would be inconclusive due to the continued presence of DYNC1H1 in all nonrod retina cells. Genotyping with tail DNA confirmed the presence of loxP (indicating a Dync1h1^F allele) (Fig. 1F), iCre75 (Fig. 1G), EGFP-CETN2 (Fig. 1H), and Prom1-CreERT2 (Fig. 1I) in respective mice (see Methods for details).

**Electroretinography of Rod Knockouts**

Scotopic electroretinography as a function of light intensity showed diminished a-wave amplitudes at P16 (Figs. 2A, D), suggesting rod degeneration had begun. A-wave amplitudes were further attenuated at P21 (Fig. 2B) and essentially extinguished at P30 (Figs. 2C, D), indicating that

**FIGURE 2.** Scotopic and photopic electroretinography. (A–C) Average scotopic a-wave amplitudes as a function of light intensity (−4.5 to 2.4 log cd s·m−2) at P16 (A), P21 (B), and P30 (C). The scotopic rod Dync1b1^-/- a-wave amplitude decreases by P21 and is nearly extinguished at P30. (D) Summary of peak amplitudes at 2.4 log cd s·m−2; average peak scotopic a-wave amplitude and standard deviations are indicated. (E, F) Average photopic b-wave amplitudes as a function of light intensity (−0.1 to 1.9 log cd s·m−2) at P21 (E) and P30 (F). Decline of the P30 photopic b-wave is probably caused by bystander cone degeneration following loss of rods in the rod knockout.
rod photoreceptor degeneration progressed rapidly and was completed 2 weeks after onset. We assessed cone function in the DYNC1H1 rod knockout by assessing the photopic b-wave. Photopic b-wave amplitudes of control and rod knock-out mice were similar at P21 (Fig. 2E) but were extinguished at P30 (Fig. 2F). Assuming that cone bipolar cells are unaffected in the dynein rod knockout, the loss of photopic b-wave results from loss of cone function. This is consistent with previous findings in which deletion of rod-specific genes resulted in secondary loss of cones due to diminished rod-derived cone viability factor. Examples of cone loss following loss of rods have been documented in animal models of dominant or recessive retinitis pigmentosa (RP), the rd1 mouse, and bystander cone degeneration of human RP.

Rapid Rod Degeneration in the Absence of DYNC1H1

While DYNC1H1 was present in all retina cells, the highest concentrations were found in the photoreceptor IS, OPL, and, to a lesser extent, the ONL (Fig. 3, row A). Our iCre75 mice were maintained on the Egrp-Cetn2+ background, in which centrin 2 (CETN2) inherited the connecting cilium tethering the IS to OS (Fig. 3A, see P14 inset) and aided to identify photoreceptor compartments. Although the Dync1h1 gene was knocked out in rods by P10 (Fig. 1E), DYNC1H1 protein persisted in the rod ONL/IS (Fig. 3, row D), and the Dync1h1−/− morphology was indistinguishable from controls at P14. DYNCH1H1 levels were reduced in the IS beginning at P16, but some DYNC1H1 was still detectable at P30, presumably in cone inner segments (Fig. 3, rows B). DYNC1H1 in the OPL persisted as distal processes of rod bipolar cells were unaffected. Dync1h1−/− IS/OS regions and ONL began to shrink after P16 (Figs. 3E, F).

In the other half of the tamCh mice, the average OS thickness was reduced by 80% in mice with efficient tamoxifen injections, but DYNC1H1 was still detectable at P30, presumably in cone inner segments (Fig. 3, rows B). DYNC1H1 in the OPL persisted as distal processes of rod bipolar cells were unaffected. Dync1h1−/− IS/OS regions and ONL began to shrink after P16 (Figs. 3E, F).

OS Protein PDE6 and Synaptic Protein CtBP2/RIBEYE in Rod Knockouts

We determined the fates of rod PDE6 and RIBEYE, both of which are not directly dependent on dynein for transport to their destinations (OS and synapse, respectively). Rod PDE6 is a heterotrimeric protein consisting of isoforms PDE6α and PDE6β catalytic subunits and two PDE6γ inhibitory subunits (reviewed in Cote). Subunit delivery to the OS is dependent on diffusion by interaction with PDE6D, a solubilization factor. In the P16 Dync1h1−/− retina, PDE6 content was reduced significantly based on OS shortening (Fig. 4, row B). As observed with rhodopsin, PDE6 never accumulated in the IS during OS shrinking at P16 and P18.

RIBEYE, expressed in retina via a tissue-specific promoter within intron 1 of the CtBP2 (C-terminal binding protein 2) gene, is a major structural protein of rod/cone synaptic contacts and delineates the OPL of controls (Fig. 4, row C). RIBEYE is not a transmembrane protein, and its transport to the synaptic terminal (the microtubule plus end) is not expected to be dependent on dynein. Mutant photoreceptor synaptic termi have normal levels of RIBEYE at P14 and P16 but are nearly devoid of CtBP2/RIBEYE at P21 (Fig. 4, row D), consistent with attenuated scotopic electroretinography (ERG) measurements. RIBEYE-immunoreactive puncta of the P21 ONL likely belong to cone pedicles, as the photopic ERG is near normal at P21. RIBEYE is undetectable at P50, consistent with absence of functional photoreceptors and “flat” rod and cone ERG amplitudes.

Deletion of DYNC1H1 by Tamoxifen Induction

To explore consequences of DYNC1H1 deletion in the adult mouse, we deleted Dync1b1 in Dync1b1+/Prom1-CreERT2 1-month-old mice using tamoxifen induction. Prom1 expression in adult mice is seen in rods and cones along with cells within the brain, pancreas, intestine/colon, kidney, lung, and reproductive system (male and female). As observed for DYNC1H1 at 1wPTI and 2wPTI, Dync1h1−/− mice were abbreviated as tamDync1b1−/− mice (Figs. 5, 6). Intraperitoneal injection of tamoxifen for 5 consecutive days induced the nuclear translocation of Cre and the degeneration rate was assessed in retina cryosections of eyes harvested 1, 2, 3, and 4 weeks after the first injection. At 1 and 2 weeks post tamoxifen induction (wPTI), DYNC1H1 levels were indistinguishable among injected tamDync1b1−/−, un injected Dync1b1+/−;Prom1-CreERT2, and injected Dync1b1+/− or Dync1b1−/−;Prom1-CreERT2 mice (Fig. 5, rows A and B).

At 3wPTI, DYNC1H1 was nearly eliminated in half of the tamDync1b1−/− mice (see Fig. 5 legend). In tamDync1b1−/− mice in which elimination of DYNC1H1 was seen, the ONL thickness was reduced by 50% and the OS + IS length was reduced to ~30% that of control (Figs. 5C, E). In the other half of the tamDync1b1−/− mice, tamoxifen-induced knockdown of DYNC1H1 was inefficient and ONL and OS + IS measurements were normal or slightly reduced. DYNC1H1 antibody fluorescence in the OPL was undisturbed as DYNC1H1 levels in bipolar cells were unaffected in the tamoxifen-induced knockdown (Fig. 5C). At 4wPTI, ONL thickness was reduced by 80% in mice with efficient tamoxifen injections, but DYNC1H1 was still detectable in the deteriorating mutant IS (Fig. 5D, E), reflecting failure to clear tamoxifen from the system.

As observed for DYNC1H1 at 1wPTI and 2wPTI, rhodopsin and M-opsin levels were indistinguishable among injected tamDync1b1−/−, un injected Dync1b1+/−;Prom1-CreERT2, and injected Dync1b1+/− or Dync1b1−/−;Prom1-CreERT2 mice (Figs. 6, 7, rows A and B). At 3wPTI in mice with efficient tamoxifen injections, tamDync1b1−/− rod OS lengths were dramatically reduced and rhodopsin mislocalized in part to the IS and OPL (Fig. 6C, left column). Additionally, very few tamDync1b1−/− cone OSs were retained (Fig. 7C). Almost no rhodopsin or M-opsin accumulated at the perinuclear endoplasmic reticulum (ER) of tamDync1b1−/− photoreceptors at 3wPTI (Figs. 6C, 7C).

At 4wPTI, the mutant ONL was reduced to two to three rows of nuclei containing traces of rhodopsin and M-opsin (Figs. 6D, 7D). Average scotopic electroretinography traces at 1.4 log cd s·m−2 (log candela seconds per square meter) at 2wPTI were nearly identical (Fig. 6E). At 3wPTI, scotopic a- and b-wave amplitudes were reduced in the tamDync1b1−/−
FIGURE 3. Immunohistochemistry with DYNC1H1 and rhodopsin (Rho) antibodies. (A–D) Representative control (rows A, C) and rod Dync1h1−/− retinas (rows B, D) labeled with antibodies directed against DYNC1H1 (rows A, B) and rhodopsin (rows C, D) at P14, P16, P18, P21, and P30. DYNC1H1 is located in the photoreceptor IS and OPL (photoreceptor terminals and bipolar cell dendrites). The rod Dync1h1−/− ONL starts to shrink at P16, and only cone nuclei remain at P30. Rhodopsin-containing rod OSs are severely shortened by P21 and absent by P30. Note, almost no rhodopsin accumulates in the ONL. Scale bar: 20 μm. (E) Comparison of the average ONL thickness of control and rod Dync1h1−/− central retinas illustrates loss of rod photoreceptors between P16 and P30. Error bars show the standard deviation. *P < 0.05 (Student’s t-test). (F) Comparison of the average combined length of rod OS + IS for P10 to P30 control and rod Dync1h1−/− retinas shows loss of OS and IS in the absence of DYNC1H1. Error bars show the standard deviation. *P < 0.05 (Student’s t-test). For E and F, the numbers of mice for control and rodKO, respectively, were P10, n = 5; P12, n = 8; P14, n = 5; P16, n = 8; P18, n = 3; P21, n = 7; P30, n = 10.

retinas, compared to control injected or uninjected specimens (Fig. 6F). The reduction in scotopic a- and b-wave amplitudes of tam Dync1b1−/− mice was even more evident at 4wPTI (Fig. 6G). Similarly, average photopic b-wave amplitudes of tamoxifen-induced knockout mice were comparable at 2wPTI (Fig. 7E) but reduced at 3wPTI (Fig. 7F). The photopic a- and b-wave amplitude of tam Dync1b1−/− mice was nearly extinguished at 4wPTI (Fig. 7G).

Scotopic a-wave and photopic b-wave amplitudes as a function of light intensity did not display significant changes at 1wPTI for any of the light intensities (Fig. 8A). At 2wPTI, induced knockout (KO) did show as statistically different from control uninjected mice but not from the control injected mice. Control injected and control uninjected did not differ from each other at 2wPTI (Fig. 8B). At 3wPTI, scotopic a-wave and photopic b-wave amplitudes in the
induced KO were reduced, but this reduction was only statistically significant at the photopic 1.4 log cd s·m\(^{-2}\) intensity (Fig. 8C). At 4wPTI, the reduction in scotopic a-wave and photopic b-wave amplitudes for induced KO was significant at most intensities (Fig. 8D). These results indicate that in mice in which loss of DYNC1H1 was efficiently induced by tamoxifen injection, rod and cone degeneration began between 2 and 3wPTI, with further retinal degeneration occurring between 3 and 4wPTI.

**DISCUSSION**

In our previous study,\(^13\) we conditionally knocked out DYNC1H1 in retina using Six3Cre that initiates ablation of DYNC1H1 after embryonic day 9. We showed that segregation of the ONL/INL was absent at P6 and P8, and mutant photoreceptors did not elaborate inner and outer segments. Here, we show that deletion of DYNC1H1 in mouse photoreceptors after ciliogenesis led to loss of function (Fig. 2), shortening of the inner and outer segments, and shrinking of the ONL (Figs. 3, 4). In wild-type (WT) photoreceptors, dynein, in conjunction with dynactin, transports cargo toward the minus end of microtubules at the MTOC or basal body (for a recent review, see Dahl and Baehr\(^14\)). Once the basal body has docked to the IS cortex, inner segment biosynthetic activity is directed toward synthesis of outer segment proteins, transport of OS proteins to the periciliary ridge by dynein, movement through the connecting cilium either by diffusion or intraflagellar transport, and incorporation into nascent discs. As mouse photoreceptor outer segments are renewed every 10 days,\(^{42,43}\) sustained delivery of large amounts of transmembrane and peripheral membrane–attached and soluble proteins is essential.

When dynein is absent in photoreceptors before ciliogenesis (<P6), the ONL never forms correctly, and OS/synaptic structures are absent.\(^13\) In Dync1h1 rodents (this communication), the process was interrupted in the second postnatal week (>P16) (Figs. 3B, D; Figs. 4B, D). Interruption occurred in tamoxifen-induced knockouts between the second and third week postinduction (Figs. 5C, D; Figs. 6C, D). In the absence of dynein, delivery of OS proteins ceases and disc morphogenesis declined. In rod knockouts beginning at P16, Dync1h1 IS/OS shortened and were lost by P30, consistent with supply chain interruption of the secretory pathway. OS structures became unstable and essentially died by starvation. In tamoxifen-induced knockouts, rods and cones appeared to be stable for 2 weeks postinduction. As observed previously,\(^13\) impaired DYNC1H1 protein clearance delayed the onset of degeneration in rod (Fig. 3, row C) and tamoxifen-induced (Figs. 5C, D) knockouts of Dync1h1.

While Six3Cre-mediated excision of Dync1h1 exons 24 and 25 had occurred by P6, DYNCH1 clearance was slow and persisted past P16.\(^13\) In our rod knockout, excision of exons 24/25 was nearly complete at P10, but degeneration was not evident before P16, as judged by ONL thickness (Fig. 3E), presumably due to slow DYNCH1 clearance. Similarly, in our tamoxifen induction experiments, DYNCH1 persisted for 2 weeks after the initial intraperitoneal injection before onset of degeneration. In contrast to retina- and rod-specific deletion of DYNC1H1, the efficiency of tamoxifen induction was variable. For example, at 4wPTI, OS + IS...
FIGURE 5. Tamoxifen-induced DYNCH1 depletion in the adult retina. (A–D) tamDync1h1−/− (tamoxifen-induced KO, first column), uninjected Dync1h1F/F;Prom1-CreERT2 (second column), tamoxifen-injected Dync1h1F/F retinas (third column), and tamoxifen-injected Dync1h1F/F;Prom1-CreERT2 (fourth column) harvested at 1wPTI (A), 2wPTI (B), 3wPTI (C), or 4wPTI (D). Cryosections were labeled with anti-DYNC1H1 (red) and contrasted with DAPI (blue). Uninjected Dync1h1F/F;Prom1-CreERT2 and injected Dync1h1F/F or Dync1h1F/F;Prom1-CreERT2 retinas all maintain normal IS and ONL thickness throughout the experiment. In tamDync1h1−/− (induced KO) retina, the ONL and IS shrank between 2wPTI and 3wPTI, but this shrinking was not statistically significant until 4wPTI. In retinas with efficient knockout of Dync1h1, ONL thinning increased in severity by 4wPTI. (E) Plot of average central ONL thickness at 1, 2, 3, and 4wPTI showing ONL thickness decreases in the tamDync1h1−/− from 2wPTI to 4wPTI. *P < 0.05 (Welch’s ANOVA). (F) Plot of average combined OS + IS lengths shows that tamDync1h1−/− rod and cone OS shrink between 2wPTI and 4wPTI. *P < 0.05 (Welch’s ANOVA). For E and F, the numbers of mice for control injected, control uninjected, and tamDync1h1−/−, respectively, were 1wPTI, n = 5, 5, 5; 2wPTI, n = 5, 5, 3; 3wPTI, n = 5, 5, 4; 4wPTI, n = 11, 5, 6.
FIGURE 6. Effect of tamoxifen-induced DYNC1H1 deletion on rhodopsin expression. (A–D) Representative immunohistochemistry from tam^DYNC1H1^−/− (first column), uninjected Dyn^F/F;Prom1-CreERT2 (second column), tamoxifen-injected Dyn^F/F retinas (third column), and tamoxifen-injected Dyn^F/F;Prom1-CreERT2 (fourth column) harvested at 1wPTI (A), 2wPTI (B), 3wPTI (C), or 4wPTI (D) and probed with anti-rhodopsin (red). Nuclei were contrasted with DAPI (blue). tam^DYNC1H1^−/− retina reveals rod OS shortening and ONL thinning from 2 to 3wPTI. Rod OS shortening and ONL thinning increased in severity by 4wPTI, with minor rhodopsin expression in the ONL. (E–G) Average scotopic ERG traces at 1.4 log cd s·m−2 from control and tam^DYNC1H1^−/− at 2wPTI (E), 3wPTI (F), and 4wPTI (G). Scotopic responses decrease between 2 and 4 wPTI in the tam^DYNC1H1^−/−.

length was severely reduced to 12% of control when tamoxifen induction was efficient but remained 88% that of control when induction was inefficient (Fig. 5F). However, even in mice in which tamoxifen induction was inefficient, ONL thickness was reduced by 30%, likely reflecting a partial loss of DYNC1H1. These variances in the efficiency of tamoxifen induction were likely due to tamoxifen not being completely in solution.

The most abundant OS membrane protein, rhodopsin, is synthesized at the endoplasmic reticulum, posttranslation-
Deletion of Dynein-1 in Photoreceptors

**FIGURE 7.** Effect of tamoxifen-induced DYNC1H1 deletion on cone-opsin expression. (A–D) Representative immunohistochemistry from tamDync1h1−/− (first column), uninjected Dync1h1FF/Prom1-CreERT2 (second column), tamoxifen-injected Dync1h1FF retinas (third column), and tamoxifen-injected Dync1h1FF/Prom1-CreERT2 (fourth column) harvested at 1wPTI (A), 2wPTI (B), 3wPTI (C), or 4wPTI (D) probed with anti-M-opsin (red). ONL indicated by DAPI (blue). tamDync1h1−/− retina reveals loss/shortening of cone OS from 2 to 3wPTI. While some M-opsin is seen in the ONL at 4wPTI, cone OSs are absent. (E–G) Average photopic ERG traces at 1.4 log cd s·m−2 from control and tamDync1h1−/− at 2wPTI (E), 3wPTI (F), and 4wPTI (G). Photopic responses decrease between 2wPTI and 4wPTI in the tamDync1h1−/−.

ally modified in the Golgi, and transported to the OS base via the secretory pathway.11,44 Newly synthesized rhodopsin is predicted to accumulate in the ONL in the absence of dynein, as COPII vesicle transport from the ER to Golgi is impaired. Ununexpectedly, we observed little rhodopsin accumulation or mislocalization in the rodDync1h1−/− ONL (Fig. 3D), an observation that could mean that rhodopsin biosynthesis terminates. Alternatively, newly synthesized rhodopsin
Deletion of Dynein-1 in Photoreceptors

**FIGURE 8.** Summary of ERG responses as a function of intensity in \(1^{\text{lo}}\)Dync1h1\(^{-/-}\) retina. (A–D) Average scotopic a-wave amplitudes (left column) and photopic b-wave amplitudes (right column) as a function of light intensity shrink between 2wPTI and 4wPTI. Injected control (blue), un.injected control (green), and tamoxifen-induced knockout (red). At 1wPTI (A), average scotopic and photopic ERGs amplitudes did not differ significantly between induced knockout and control. At 2wPTI (B), the average scotopic a-wave amplitude was slightly reduced compared to controls. Photopic ERGs did not differ. At 3wPTI (C), induced KO shows reduced average ERG amplitudes for both scotopic a-wave and photopic b-wave, but this reduction was only statically significant for the photopic 1.4 log cd s·m\(^{-2}\) intensity. At 4wPTI (D), the reduction in average ERG amplitude for both scotopic a-wave and photopic b-wave for the induced KO was statically significant for most ERG flash intensities. *\(P < 0.05\) (unbalanced two-factor ANOVA, Tukey’s honestly significant difference criterion).
PDE6 is a key protein in the phototransduction cascade regulating the levels of cGMP, the internal transmitter of phototransduction. Rod PDE6 is peripherally membrane attached using prenyl side chains as anchors. PDE6D (PDE homolog of 2-hydroxyacid dehydrogenase, binding NAD) is thought to travel through the centriole identification without use of a specific antibody. It is unclear whether PDE6 transported in the absence of PDE6D depends on dynein. Mouse RIBEYE is a ribbon-specific protein, expressed under its own promoter within the CIBP2 gene. It has the capability to build the synaptic ribbon scaffold via multiple RIBEYE-RIBEYE interactions. The B domain of RIBEYE is a homolog of 2-hydroxyacid dehydrogenase, binding NAD and UNC119. RIBEYE is presumably not a membrane-associated protein, and while its transport mode to the synaptic region is unknown, it is not likely to be dynein dependent. Both PDE6 and RIBEYE disappear as rod photoreceptors degenerate rapidly.

Our study currently is limited to evaluation of photoreceptor degeneration by confocal microscopy and electron microscopy. We show that photoreceptor OS, IS, and ONL nearly simultaneously shrink in rod knockouts beginning at P16 and in tamoxifen-induced knockouts by the third week postinjection. OS likely shrink by failed delivery of newly synthesized proteins to the periciliary ridge near the basal body. It will be interesting to evaluate the status of the IS microtubule cytoskeleton in the absence of dynein by immunohistochemistry with anti-tubulin antibodies, the effect on disc morphogenesis as protein delivery ceases, and potential changes in synaptic structures by transmission electron microscopy.

Methods

Animals

All procedures were approved by the University of Utah Institutional Animal Care and Use Committee (Protocol 18-11005). Prom1-CreERT2/EGFP-Cetn2 mice (Stock No: 017743) and Egfp-Cetn2 mice Stock No. 008234-CB6-Tg(CAG-EGFP/CETN2)3-4Jgg/J were obtained from The Jackson Laboratory. iCre75 mice were generated in Utah. Dync1h1Jgg/J were obtained from The Jackson Laboratory. iCre75 (5′-CTCAAATCTTCTCTGGTTA-TCTCTGGAAAGGTTGGCAGA) and loxp-R (5′-GGATGCAGTTGCGGTTTGCTAGT). Amplicon sizes were Dynch1 WT allele (1.3 kb), Dynch1–/– allele (1.6 kb), and Cre-recombination allele (500 bp) (Fig. 1E). Primers flanking the loxP site of introns 25 to 26 were used to distinguish between wild-type and floxed Dynch1 alleles (loxP-F, 5′-TGATGGTCTTGCCATATTGGTG; loxP-R, 5′-GGATGCAGTTGCGGTTTGCTAGT) with amplicon sizes of 269 bp (floxed) and 202 bp (WT) (Fig. 1F). Presence of iCre75 transgene was verified with iCre75-F (5′-GGATGCCACCTCTGATGAAG) and iCre75-R (5′-CACACACCTTCTCTGACCG) primers (amplicon size 700 bp) (Fig. 1G). Presence of EGFP-CETN2 was determined by PCR using primers Cetn2-F (5′-CGATGCGGAGATCAGTTGCGGTTTGCTAGT) and loxP-R (5′-GGATGCAGTTGCGGTTTGCTAGT) with amplicon sizes of 269 bp (floxed) and 202 bp (WT) (Fig. 1H). Presence of iCre75 transgene was verified with iCre75-F (5′-GGATGCCACCTCTGATGAAG) and iCre75-R (5′-CACACACCTTCTCTGACCG) primers (amplicon size 700 bp) (Fig. 1I). Presence of EGFP-CETN2 was determined by PCR using primers Cetn2-F (5′-CGATGCGGAGATCAGTTGCGGTTTGCTAGT) and iCre75-R (5′-CACACACCTTCTCTGACCG) primers (amplicon size 700 bp) (Fig. 1I). Presence of EGFP-CETN2 was determined by PCR using primers Cetn2-F (5′-CGATGCGGAGATCAGTTGCGGTTTGCTAGT) and iCre75-R (5′-CACACACCTTCTCTGACCG) primers (amplicon size 700 bp) (Fig. 1I). Presence of iCre75 transgene was verified with iCre75-F (5′-GGATGCCACCTCTGATGAAG) and iCre75-R (5′-CACACACCTTCTCTGACCG) primers (amplicon size 700 bp) (Fig. 1I). Presence of EGFP-CETN2 was determined by PCR using primers Cetn2-F (5′-CGATGCGGAGATCAGTTGCGGTTTGCTAGT) and iCre75-R (5′-CACACACCTTCTCTGACCG) primers (amplicon size 700 bp) (Fig. 1I). Presence of EGFP-CETN2 was determined by PCR using primers Cetn2-F (5′-CGATGCGGAGATCAGTTGCGGTTTGCTAGT) and iCre75-R (5′-CACACACCTTCTCTGACCG) primers (amplicon size 700 bp) (Fig. 1I).

Generation of Rod Knockout Mice

Dynch1–/– mice were crossed with iCre75 transgenic mice kept on an Egfp-Cetn2 background to generate Dynch1–/–;iCre75,Egfp-Cetn2 mice. Mice were then backcrossed to Dynch1–/–;Egfp-Cetn2 mice to generate experimental animals. Expression of Egfp-Cetn2 allows connecting cilium and centrirole identification without use of a specific antibody.

Generation of Knockouts by Tamoxifen Induction

Dynch1–/– mice were bred to Prom1-CreERT2 mice to generate Dynch1–/–;Prom1-CreERT2 mice. In these mice, expression of CreERT2 is driven by the prominin 1 (Prom1) promoter/enhancer. Dynch1–/–;iCre75 mice were crossed to Dynch1–/– mice to generate mice for tamoxifen experiments. Tamoxifen was administered via intraperitoneal injection in adult mice (P21–P30 at time of first injection). Tamoxifen was dissolved in corn oil to a stock solution of 20 mg/mL. Mice were dosed with 150 mg/kg body weight for 5 consecutive days according to their weight on the first day of injections (i.e., 7.5 μL of 20 mg/mL tamoxifen solution per gram weight). ERG and collection of eyes for confocal immunolocalization were performed 1 to 4 weeks after the first injection.

Genotyping

Genomic DNA was extracted from fresh tissue by dissolving tail clips from P4 to P14 mice in 200 μL tail lysis buffer at 50°C to 60°C for 1 to 2 hours. Digests were then centrifuged at 15,000 rpm for 5 minutes. Supernatant was added to an equal volume of isopropanol and centrifuged at 15,000 rpm for 5 minutes. Pellet was rehydrated in 100 μL H2O. Genotyping was performed by PCR with EconoTaq DNA polymerase (Lucigen, Middleton, WI). To verify the deletion of exons 24 and 25 of Dynch1 in Dynch1–/– retina, P10 to P21 retinal genomic DNA was extracted and amplification was performed with primers exon 23-F (5′-TCTCTGGAAAGGTTGGCAGA) and loxp-R (5′-GGATGCAGTTGCGGTTTGCTAGT). Amplicon sizes were Dynch1 WT allele (1.3 kb), Dynch1–/– allele (1.6 kb), and Cre-recombination allele (500 bp) (Fig. 1E). Primers flanking the loxP site of introns 25 to 26 were used to distinguish between wild-type and floxed Dynch1 alleles (loxP-F, 5′-TGATGGTCTTGCCATATTGGTG; loxP-R, 5′-GGATGCAGTTGCGGTTTGCTAGT) with amplicon sizes of 269 bp (floxed) and 202 bp (WT) (Fig. 1F). Presence of iCre75 transgene was verified with iCre75-F (5′-GGATGCCACCTCTGATGAAG) and iCre75-R (5′-CACACACCTTCTCTGACCG) primers (amplicon size 700 bp) (Fig. 1G). Presence of EGFP-CETN2 was determined by PCR using primers Cetn2-F (5′-CGATGCGGAGATCAGTTGCGGTTTGCTAGT) and Cetn2-R (5′-ACTTGAAGACCTGCGACACAT) (amplicon size 600 bp) (Fig. 1H). Presence of Prom1-CreERT2 was verified with Prom1-F (5′-CAGGCGTGTAGCCTGGTTACT) and Prom1-CreERT2-R (5′-AGGCAAATTGGTACCG) primers (amplicon 320 bp) with Prom1-F and Prom1WT-R (5′-TGGTATGTTGCTCTGCTCTG) (amplicon 860 bp) serving as a control (Fig. 1I).
Immunohistochemistry

Sections were encircled using a PAP pen (Ted Pella, Redding, CA), warmed for 30 minutes at 37°C, and then rehydrated by washing 10 minutes three times in 1× PBS. All sections used were from the central retina near the optic nerve. Sections were blocked in 5% normal goat serum/0.1% TritonX-100 in 1× PBS for 1 hour. Antibodies, dilutions, and sources follow: DYNCH1 (dilution 1:250, 12345-1-AP; Proteintech, Proteintech Rosemont, IL), rhodopsin (1:1000, A15093; Abclonal, Abclonal Woburn, MA), PDE6 (1:1000; MOE Cytosign, Cytosignal, Cytosignal IRVINE, CA), OPN1MW/OPN1LW (1:500, AB5405; Millipore Sigma, Millipore Sigma St. Louis, MO), and CtBP2/RIBEYE (1:10,000, 612044; BD Biosciences, BD Biosciences Franklin Lakes, NJ). Primary antibodies were diluted in blocking buffer and applied to sections; sections were then incubated overnight at 4°C. Slides were washed for 10 minutes three times in PBS. Secondary antibodies and b-wave ERG response across multiple ages or weeks experimental and control animals for their quantified a-wave and b-wave ERG response were performed using Tukey’s honestly significant difference criterion. Statistical significance was determined using an a value of 0.05. All electroretinography statistics were performed using statistical toolbox “anovan” and “multcompare” functions in MATLAB (MathWorks, Natick, MA, USA).

We performed a Student's t-test assuming unequal variances to compare iCre75 control and KO ONL thickness or IS + OS length. Retina measurements used for these calculations were determined based on an average of three measurements per retina. Microsoft Excel (Microsoft, Redmond, WA, USA) function “T.TEST assuming unequal variances” was used to calculate the P value, with statistical significance determined using an a value of P < 0.05. Since we had unequal sample sizes and could not assume homogeneity of variances, we preformed Welch’s ANOVA test to compare control injected, control uninjected, and tamDync1h1−/− ONL thickness or IS + OS length using the xrcalstats data analysis add-on to Microsoft Excel. Statistical significance was determined using a P value of <0.05.

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References


Electroretinography

Scotopic and photopic ERG measurements were performed using P16, P21, or P30 mice for the iCre75 experiments and at 1, 2, 3, or 4 weeks after tamoxifen induction for the Prom1-CreERT2 experiments. Prior to ERG, the mice were light-adapted under a background light of 1.48 log cd s·m−2. Scotopic single-flash responses of control and knockout were recorded at stimulus intensities of 0.1 to 4.5 to 2.4 log cd s·m−2 for 5 to 10 minutes prior to measuring photopic responses. Photopic single-flash responses of control and knockout were recorded at stimulus intensities of −0.1 to 1.9 log cd s·m−2.

Statistical Analysis

We performed an unbalanced two-factor ANOVA to compare experimental and control animals for their quantified a- and b-wave ERG response across multiple ages or weeks after tamoxifen induction. Post hoc multiple comparison...


