

Ciliary neurotrophic factor (CNTF) protects retinal cone and rod photoreceptors by suppressing excessive formation of the visual pigments

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The retinal pigment epithelium (RPE)-dependent visual cycle provides 11-*cis*-retinal to opsins in the photoreceptor outer segments to generate functional visual pigments that initiate phototransduction in response to light stimuli. Both RPE65 isomerase of the visual cycle and the rhodopsin visual pigment have recently been identified as critical players in mediating light-induced retinal degeneration. These findings suggest that the expression and function of RPE65 and rhodopsin need to be coordinately controlled to sustain normal vision and to protect the retina from photodamage. However, the mechanism controlling the development of the retinal visual system remains poorly understood. Here, we show that deficiency in ciliary neurotrophic factor (CNTF) up-regulates the levels of rod and cone opsins accompanied by an increase in the thickness of the outer nuclear layers and the lengths of cone and rod outer segments in the mouse retina. Moreover, retinoid isomerase activity, expression levels of RPE65 and lecithin:retinol acyltransferase (LRAT), which synthesizes the RPE65 substrate, were also significantly increased in the *Cntf*^{-/-} RPE. Rod *a*-wave and cone *b*-wave amplitudes of electroretinograms were increased in *Cntf*^{-/-} mice, but rod *b*-wave amplitudes were unchanged compared with those in WT mice. Up-regulated RPE65 and LRAT levels accelerated both the visual cycle rate and recovery rate of rod light sensitivity in *Cntf*^{-/-} mice. Of note, rods and cones in *Cntf*^{-/-} mice exhibited hypersusceptibility to light-induced degeneration. These results indicate that CNTF is a common extracellular factor that prevents excessive production of opsins, the photoreceptor outer segments, and 11-*cis*-retinal to protect rods and cones from photodamage.

Phototransduction that converts light energy into electrical impulse in the retina begins with photoisomerization of 11-*cis*-

retinal (11cRAL)² to all-*trans*-retinal in the opsin visual pigments localized to the outer segments (OS), the light-sensing organelles, of rod and cone photoreceptors (1, 2). Because apopsins without the 11cRAL chromophore are not responsive to light, 11cRAL must be regenerated and recombined with apopsins to form light-sensitive visual pigments in the photoreceptor OS. RPE65 is a key retinoid isomerase in the RPE-dependent visual cycle responsible regenerating 11cRAL (3–5). It catalyzes synthesis of 11-*cis*-retinol, the reduced form of 11cRAL, from all-*trans*-retinyl esters synthesized by LRAT and other enzymes in the RPE (6–8).

Although rhodopsin is essential for initiating rod phototransduction it is also the mediator of rod degeneration induced by intense light (9). In addition, higher expression level and activity of RPE65 are associated with an increase in susceptibility of the photoreceptors to light-induced degeneration (10–12). These findings suggest that the expression levels and function of rhodopsin and RPE65 need to be controlled to maintain normal vision and to protect photoreceptors from light-induced degeneration. However, the molecular mechanisms that coordinately control expressions of both opsin and RPE65 during the development and maturation of the retinal light-sensing system remain largely unknown.

Expressed by Muller and RPE cells (13–15), CNTF is an extracellular signaling protein in the neuroretinal and the interphotoreceptor matrix, which is associated with the membranes of the RPE, Muller, and photoreceptor cells (16). CNTF initiates its signaling by interacting with its receptor- α , which then recruits the leukemia inhibitory factor (LIF) receptor- β and gp130 to form a functional receptor complex in the cell membrane (17). All of these receptor subunits are expressed in the RPE and photoreceptors (14, 18–20). Exogenous CNTF has

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² The abbreviations used are: 11cRAL, 11-*cis*-retinal; BC, bipolar cell; CAR, cone arrestin; CNTF, ciliary neurotrophic factor; DAPI, 4',6'-diamidino-2-phenylindole; ERG, electroretinography; INL, inner nuclear layer; IRBP, interphotoreceptor retinoid-binding protein; LIF, leukemia inhibitory factor; LRAT, lecithin:retinol acyltransferase; mGluR6, metabotropic glutamate receptor 6; ONL, outer nuclear layer; OS, outer segments; PKC α , protein kinase C α ; RPE, retinal pigment epithelium; TUNEL, terminal deoxynucleotidyl transferase dUTP nick end labeling; STAT, signal transducers and activators of transcription; PI3K, phosphatidylinositol 3-kinase; FBS, fetal bovine serum; PB, phosphate buffer; DMEM, Dulbecco's modified Eagle's medium; cd \times s/m², candelas \times s per m²; Rho, rhodopsin; OPL, outer plexiform layer; ROS, rod outer segments; MAPK, mitogen-activated protein kinase.

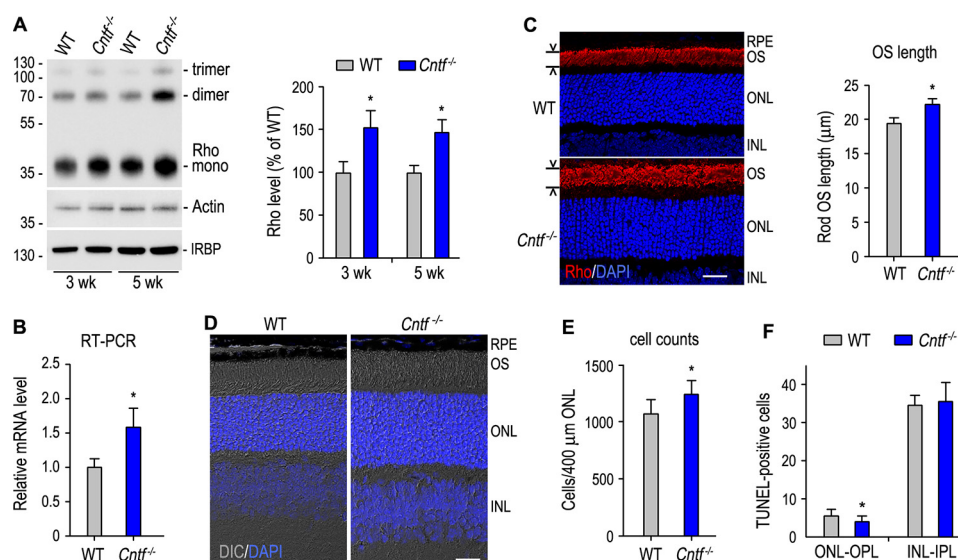


Figure 1. Increase in rhodopsin (*Rho*) expression, thickness of the ONL, and length of rod OS in *Cntf*^{-/-} mouse. *A*, immunoblot analysis of Rho and IRBP in 3- and 5-week-old WT and *Cntf*^{-/-} retinas. Actin was detected to normalize sample loading. The histogram shows percentages of the immunoblot intensities of Rho from *Cntf*^{-/-} retinas relative to the Rho immunoblot intensities of WT retinas. Asterisks in this figure indicate significant differences between WT and *Cntf*^{-/-} mice ($p \leq 0.05$); error bars indicate S.D. ($n \geq 3$). *B*, transcription level of Rho mRNA in 3-week-old *Cntf*^{-/-} retina was determined by quantitative RT-PCR and expressed as fold of Rho mRNA in age-matched WT retina. *C*, immunohistochemistry showing localization of Rho (red) to the rod OS in the superior retinas of 3-week-old WT and *Cntf*^{-/-} mice. Nuclei were stained with DAPI (blue). Histogram shows Rho-positive OS lengths in the sections. *D*, differential interference contrast (DIC) images of retinal sections from 3-week-old WT and *Cntf*^{-/-} mice. All scale bars denote 20 μm . *E*, rod nuclear numbers counted in a 400- μm wide superior ONL area that is 600 μm away from the optic nerve head. *F*, TUNEL-positive cell numbers in the ONL-OPL and inner nuclear layer (INL)-inner plexiform layer (IPL) of retinal sections from WT and *Cntf*^{-/-} mice at postnatal day 8.

been shown to inhibit rhodopsin expression in the rodent retina (21–23), but it induced expression of rod and cone opsins in the chick retina (24, 25). In addition, CNTF increased RPE cell survival and mitotic activity (26).

In animal models of retinal degeneration, CNTF treatment promoted photoreceptor survival (27–31) and regeneration of cone outer segments (32). Furthermore, recent clinical trials in patients with retinitis pigmentosa, Usher syndrome type 2, or geographic atrophy showed that CNTF treatment increased retinal thickness and slowed the progression of vision loss (33–35), although in some animal studies and human clinical trials CNTF treatment had no therapeutic benefit (36–38). Despite the neuroprotective effect in some clinical trials and animal studies, CNTF treatment significantly suppressed visual function, as determined by electroretinography (ERG) (29, 37, 39, 40). Intraocular CNTF administration also suppressed ERG responses in wildtype (WT) animals (23, 40–42).

A gene profiling study showed that intravitreal injection of CNTF induced numerous genes associated with inflammation and gliosis in the Muller cells (43). In addition, CNTF increased secretion of neurotrophin-3 and decreased secretion of vascular endothelial growth factor, interleukin-8, and transforming growth factor- β 2 in the RPE cells (14). These altered gene expression and protein secretion might cause many secondary effects in the CNTF-treated patients and animals, and therefore, increased difficulty in defining the primary role and mechanism of CNTF function in regulating development, function, and protection of the retinal photoreceptors and RPE. In this study, we analyzed neuroretinal and RPE phenotypes in *Cntf*^{-/-} mice, and identified CNTF as a common extracellular signal that down-regulates expression of opsins in rod and cone photoreceptors, as well as RPE65 and LRAT in the RPE, to avoid

excessive formation of both light-sensing organelles and light-sensitive visual pigments, which mediate phototransduction or photoreceptor degeneration, depending on the light intensity they have captured.

Results

Rod excessive development is associated with up-regulation of rhodopsin in the *Cntf*^{-/-} retina

CNTF added into retinal cultures displayed distinct effects on rod development depending on species: it promoted chick rod development and rod-opsin expression (24), but it inhibited rat rod differentiation and rod-opsin expression (21). To define the effect of CNTF deficiency on rod development and rod-opsin expression, we first compared expression levels of rhodopsin in WT and *Cntf*^{-/-} mice. As shown in Fig. 1*A*, expression levels of rhodopsin (Rho) in 3- and 5-week-old *Cntf*^{-/-} mice were 45–50% higher than those in age-matched WT mice. Rho mRNA expression levels in 3-week-old *Cntf*^{-/-} retina were increased by ~53% as compared with those in the same age WT retina (Fig. 1*B*).

To confirm these results, we analyzed morphology of WT and *Cntf*^{-/-} retinas. Immunohistochemistry showed that length of the rod outer segments (ROS) in a 3-week-old *Cntf*^{-/-} mouse was 15% longer than that of WT mouse (Fig. 1*C*). Differential interference contrast microscopy of retinal sections confirmed that both ROS length and thickness of the outer nuclear layer (ONL) in 3-week-old *Cntf*^{-/-} retina are clearly greater than those in age-matched WT retina (Fig. 1*D*). To know if numbers of rods are increased in the *Cntf*^{-/-} mouse, we counted nuclear numbers in a 400- μm wide region of the superior ONL that is 600 μm away from the optic nerve head. As

CNTF inhibits visual pigment generation to protect retina

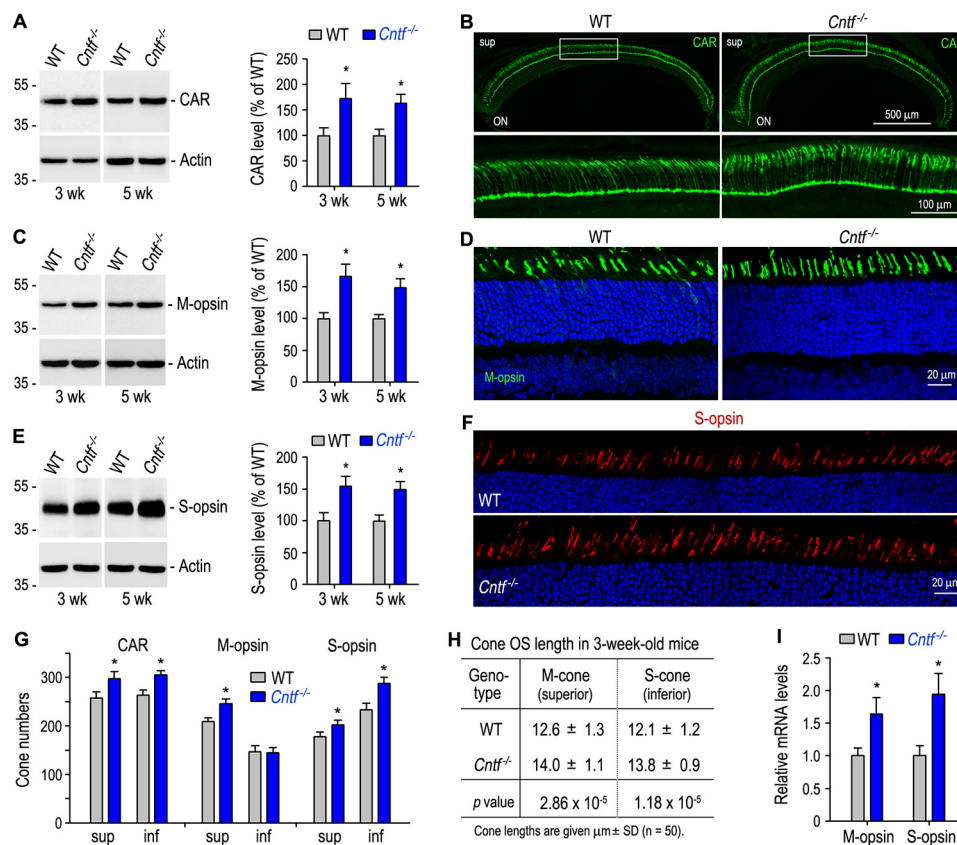


Figure 2. Increase in expression of cone-specific proteins and length of cone OS in *Cntf*^{-/-} mouse. *A*, immunoblot analysis of CAR in the retinas of 3- or 5-week-old WT and *Cntf*^{-/-} mice. The histogram shows percentages of immunoblot intensities of CAR from *Cntf*^{-/-} retinas relative to the CAR immunoblot intensities of WT retinas. *B*, immunohistochemistry showing CAR immunoreactivity in the superior (*sup*) retinas of 3-week-old WT and *Cntf*^{-/-} mice (*upper panels*). ON, optic nerve. The areas of the rectangles are shown in the higher magnification images (*bottom panels*). *C* and *E*, immunoblot analysis of M-opsin (*C*) and S-opsin (*E*) from 3- or 5-week-old WT and *Cntf*^{-/-} retinas. Histograms show percentages of immunoblot intensities of M-opsin (*C*) or S-opsin (*E*) in the *Cntf*^{-/-} retinas relative to M-opsin or S-opsin intensities in WT retinas. *D* and *F*, immunohistochemistry for M-opsin (*D*) and S-opsin (*F*) in the superior (*D*) or inferior (*F*) retinas of 3-week-old WT and *Cntf*^{-/-} mice. *G*, numbers of CAR- or cone opsin-positive cells in the superior (*sup*) or inferior (*inf*) retinas of 3-week-old WT and *Cntf*^{-/-} mouse retinal sections taken from the dorsal-ventral midline of the eye. *H*, lengths of M- and S-cone OS in 3-week-old WT and *Cntf*^{-/-} mouse retinal sections. *I*, relative mRNA levels of cone opsins in 3-week-old WT and *Cntf*^{-/-} retinas were determined by quantitative RT-PCR. All asterisks indicate significant differences between WT and *Cntf*^{-/-} mice ($p < 0.05$); error bars denote S.D. ($n = 3-4$).

shown in Fig. 1*E*, numbers of rods in the region of the *Cntf*^{-/-} retina were 11% greater than those in WT retina. Numbers of TUNEL-positive cells in the ONL and the outer plexiform layer (OPL) of the *Cntf*^{-/-} retina at postnatal day 8 were significantly smaller than those in age-matched WT retinal ONL-OPL (Fig. 1*F*).

Increase of cone-specific proteins in over-elongated OS of *Cntf*^{-/-} cones

Exogenous CNTF has been shown to promote cone differentiation in the developing chick retina (24, 25), suggesting that CNTF deficiency may reduce expression of cone-specific proteins. Unexpectedly, quantitative immunoblot analysis showed that cone arrestin (CAR), which expresses in both M- and S-cones (44), was increased ~50% in 3- and 5-week-old *Cntf*^{-/-} mouse retinas, as compared with age-matched WT mouse retinas (Fig. 2*A*). In agreement with this result, immunohistochemistry showed that length and numbers of CAR-positive outer segments in 3-week-old *Cntf*^{-/-} retina were clearly greater than those in WT retina (Fig. 2, *B* and *G*).

To know whether M- and S-opsins are also increased in the *Cntf*^{-/-} retina, we performed quantitative immunoblot analy-

sis for cone opsins. As shown in Fig. 2, *C* and *E*, expression levels of M- and S-opsins in 3- and 5-week-old *Cntf*^{-/-} retinas are significantly higher than those in age-matched WT retinas. This increase in expression of cone opsins is associated with an increase in M- and S-cone numbers as well as length of cone outer segments in the *Cntf*^{-/-} retina (Fig. 2, *D* and *F-H*). Quantitative RT-PCR revealed that transcripts for M- and S-opsins were increased by 60~70% in 3-week-old *Cntf*^{-/-} retina, as compared with those in age-matched WT retina (Fig. 2*I*).

Up-regulation of visual cycle enzymes in *Cntf*^{-/-} RPE

Increased rod and cone opsins may need an increase in 11cRAL synthesis to form functional visual pigments. We therefore tested whether the visual cycle enzymes are changed in the *Cntf*^{-/-} RPE. Immunoblot analysis showed that protein levels of RPE65 in 3- and 5-week-old *Cntf*^{-/-} RPE were 40–60% higher than those in age-matched WT RPE (Fig. 3*A*). Similarly, expression levels of LRAT are also significantly increased in the *Cntf*^{-/-} RPE, as compared with WT RPE (Fig. 3*B*). In agreement with these results, activities of the retinoid isomerase in the 3- and 5-week-old *Cntf*^{-/-} RPE were at least 45% higher than those in age-matched WT RPE (Fig. 3*C*).

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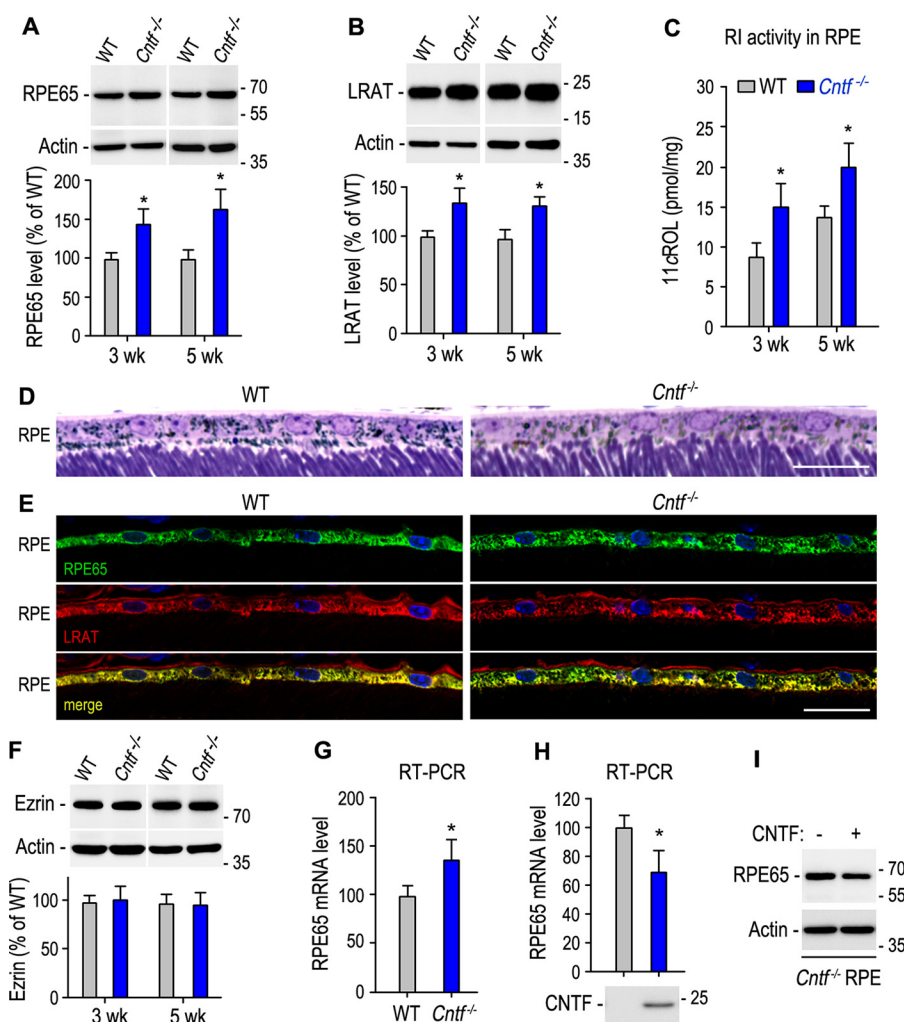


Figure 3. Up-regulation of the visual cycle enzymes in the *Cntf*^{-/-} RPE. A and B, immunoblot analysis of RPE65 (A) and LRAT (B) in the RPE of 3- or 5-week-old WT and *Cntf*^{-/-} mice. Histograms show immunoblot intensities of RPE65 or LRAT from the *Cntf*^{-/-} RPE relative to RPE65 or LRAT intensities in WT mouse RPE. C, retinoid isomerase (RI) activities determined by measuring synthesis of 11cROL from all-*trans*-retinol substrate incubated with RPE homogenates of 3- or 5-week-old WT and *Cntf*^{-/-} mice. D, light microscopic images of RPE layers in the superior retinal sections of 5-week-old WT and *Cntf*^{-/-} mice. E, immunohistochemistry showing distribution patterns of RPE65 and LRAT in the superior retinal sections from 3-week-old WT or *Cntf*^{-/-} mice. Scale bars in D and E denote 20 μ m. F, immunoblot analysis of Ezrin in the eyecups of 3- or 5-week-old WT and *Cntf*^{-/-} mice. The histogram shows relative immunoblot intensities of Ezrin in WT and *Cntf*^{-/-} eyecups. G, relative expression levels of RPE65 mRNA in WT and *Cntf*^{-/-} RPE were determined by quantitative RT-PCR. H, quantitative RT-PCR showing relative expression levels of RPE65 mRNA in *Cntf*^{-/-} mouse eyecup RPE incubated with media with or without CNTF. I, immunoblot analysis of RPE65 in *Cntf*^{-/-} mouse eyecup RPE in H. All asterisks indicate significant differences between WT and *Cntf*^{-/-} mice or between test and control groups ($p < 0.04$); error bars are S.D. ($n = 3$ or 4).

To know if the number of RPE cells is increased in *Cntf*^{-/-} mouse, we counted DAPI-positive RPE nuclear numbers in sections taken from the dorsal-ventral midline of the eye. We then calculated RPE cell density by dividing RPE nuclear numbers with arc length of retina in the sections. RPE cell densities in *Cntf*^{-/-} sections (28.6 ± 2.0 nuclei/mm, $n = 4$) were similar to those in WT sections (27.8 ± 1.8 nuclei/mm, $n = 4$). RPE morphology in the *Cntf*^{-/-} retinal section was similar to that in WT retinal section (Fig. 3D). Distribution patterns of RPE65 and LRAT in the *Cntf*^{-/-} RPE were also similar to those in WT RPE (Fig. 3E). Consistent with these results, expression levels of the RPE microvilli protein Ezrin in the *Cntf*^{-/-} RPE were similar to those in WT RPE (Fig. 3F).

To determine whether CNTF up-regulates RPE65 expression at the transcription level we performed quantitative RT-PCR. As shown in Fig. 3G, mRNA expression levels of RPE65 were increased ~30% in the *Cntf*^{-/-} mouse, as compared with

WT mouse. To confirm this result, we incubated *Cntf*^{-/-} eyecup RPE with media of 293T-LC cells transfected with pRK5 or pRK-CNTF. Quantitative RT-PCR and immunoblot analysis showed that both mRNA and protein levels of RPE65 were reduced in the eyecup RPE incubated with media containing CNTF (Fig. 3, H and I).

Rod hyperpolarization, but not rod ON bipolar depolarization, was increased in *Cntf*^{-/-} mouse

Because rod-opsin is up-regulated in the *Cntf*^{-/-} retina, we tested whether the *Cntf*^{-/-} rods display an increased photo-transduction. We recorded scotopic ERG responses of dark-adapted WT and *Cntf*^{-/-} mice to a series of increasing light flashes. Representative ERG responses elicited with 0~2 log candelas \times s/m² ($cd \times s/m^2$) flashes are shown in Fig. 4A. Amplitudes of *a*-waves elicited with 1~2 log $cd \times s/m^2$ flashes were significantly greater in *Cntf*^{-/-} mice, as compared with

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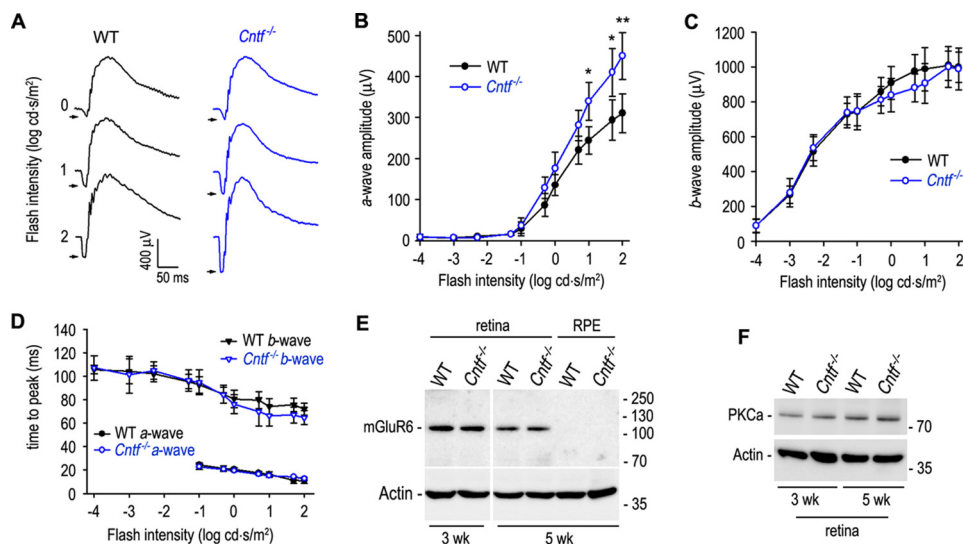


Figure 4. Enhanced rod phototransduction in *Cntf*^{-/-} mice. *A*, representative raw scotopic ERG responses of dark-adapted WT and *Cntf*^{-/-} mice to the indicated flashes. *B* and *C*, amplitudes of scotopic ERG *a*-wave (*B*) and *b*-wave (*C*) elicited with the indicated flashes in dark-adapted WT and *Cntf*^{-/-} mice. Asterisks indicate significant differences between WT and *Cntf*^{-/-} mice (*, $p < 0.04$; **, $p < 0.01$); error bars show S.D. ($n = 6$). *D*, times from stimulus onset to *a*- or *b*-wave peaks of WT and *Cntf*^{-/-} mouse rods. *E*, immunoblot analysis of mGluR6 in 3- or 5-week-old WT and *Cntf*^{-/-} mouse retinas or RPE. Actin was used as loading control. *F*, immunoblot analysis of PKC α in 3- or 5-week-old WT and *Cntf*^{-/-} retinas.

those in WT mice (Fig. 4, *A* and *B*). Unexpectedly, *b*-wave amplitudes in *Cntf*^{-/-} mice were similar to those of WT mice (Fig. 4, *A* and *C*).

To know if the synaptic function between rod and rod ON bipolar cell (ON-BC) was altered in the *Cntf*^{-/-} mouse, we measured *a*- and *b*-wave implicit times (times from stimulus onset to *a*- or *b*-wave peak). As shown in Fig. 4*D*, *a*-wave implicit times of *Cntf*^{-/-} rod ERG were similar to those of WT rod ERG. The *b*-wave implicit times of *Cntf*^{-/-} rod ERG were also similar to those of WT rod ERG elicited with $-4 \sim 0$ log cd \times s/m² flashes. When $0.7 \sim 1.7$ log cd \times s/m² flashes applied, the *b*-wave implicit times of *Cntf*^{-/-} rod ERG were 10.5–12.2% shorter than those of WT rods ($p = 0.056$, $n = 6$; Fig. 4*D*). Immunoblot analyses showed that expression levels of the ON-BC markers (mGluR6 and PKC α) in the *Cntf*^{-/-} retina were similar to those in WT retina (Fig. 4, *E* and *F*).

Faster recovery rate of rod light sensitivity is underpinned by accelerated visual cycle in *Cntf*^{-/-} mouse

Because rod opsin and the visual cycle enzymes are increased in the *Cntf*^{-/-} mouse, we tested whether the recovery rate of rod light sensitivity is accelerated in *Cntf*^{-/-} mouse. We exposed dark-adapted WT and *Cntf*^{-/-} mice to 800 lux light for 5 min, and then returned the mice to darkness. At different times, we recorded scotopic ERG responses to three different flash stimuli (10, 25, and 50 cd \times s/m²). WT and *Cntf*^{-/-} mice kept in darkness for 5 min displayed similar *a*-wave amplitudes (Fig. 5*A*). However, *a*-wave amplitudes of *Cntf*^{-/-} mice kept in darkness for 10–30 min were higher than those in WT mice under the same light conditions (Fig. 5*A*).

To know if this faster recovery of rod light sensitivity is supported by an accelerated visual cycle, we measured the regeneration rates of 11cRAL in WT and *Cntf*^{-/-} mice kept in darkness for different times after photobleaching the visual pigments. Because both the ONL thickness and the OS length are greater in *Cntf*^{-/-} retina than those of WT retina, we

expressed the levels of 11cRAL in photobleached WT and *Cntf*^{-/-} eyes as 100%. After keeping the mice in darkness for 15 or 30 min, the amounts of 11cRAL were increased 55 or 83%, respectively, in *Cntf*^{-/-} mice, but only 15 or 30% in WT mice (Fig. 5*B*). These data indicate that the regeneration rates of the 11cRAL in the *Cntf*^{-/-} eyes are faster than those in the WT eyes.

Hyper visual function of cones in *Cntf*^{-/-} mouse

The up-regulation of cone opsins in the *Cntf*^{-/-} retina (Fig. 2) prompted us to test if cone visual function was enhanced in the *Cntf*^{-/-} mouse. Under a rod-saturating background light, we recorded photopic ERG responses of WT and *Cntf*^{-/-} mice to a series of increasing achromatic flashes. As shown in Fig. 6, *A–C*, amplitudes of *a*- and *b*-waves elicited with a given intensity of flash were significantly higher in *Cntf*^{-/-} mice than WT mice. Implicit times to photopic *b*-wave peaks elicited with $0.5 \sim 2$ log cd \times s/m² flashes were markedly shorter in *Cntf*^{-/-} mice versus WT mice (Fig. 6*D*), suggesting that synaptic function between cone and cone ON-BC is normal or enhanced in *Cntf*^{-/-} mice. To confirm these results, we recorded photopic flicker ERG. Consistent with the above results, amplitudes of *b*-waves elicited with 10- and 20-Hz flashes at 10 cd \times s/m² intensity were significantly increased in *Cntf*^{-/-} mice, as compared with those in WT mice (Fig. 6, *E* and *F*). These results indicate that CNTF deficiency causes a “super vision” mediated by cones. These results also suggest that CNTF may have different roles in the development and/or function of cone ON-BC versus rod ON-BC systems.

Rods and cones in *Cntf*^{-/-} mouse exhibited hyper susceptibility to photodamage

Because the photoresponsiveness of rods and cones is increased in *Cntf*^{-/-} mice, we investigated the effects of intense light on the retinal function and structure of *Cntf*^{-/-} mice. We exposed mice to 12,000 lux light for different times (30–75

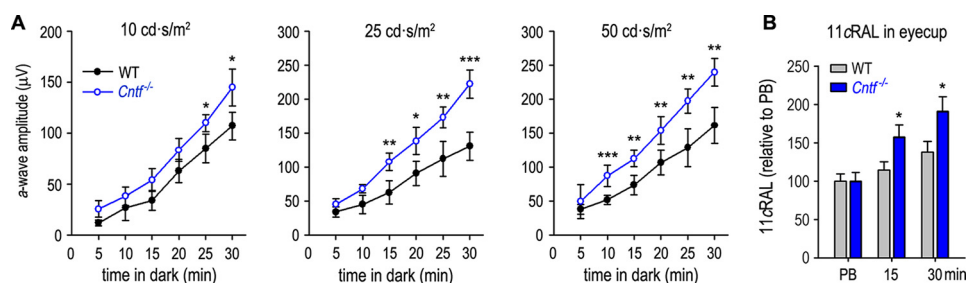


Figure 5. Faster recovery of rod light sensitivity in *Cntf*^{-/-} mice. A, amplitudes of scotopic ERG *a*-waves elicited with the indicated flashes (10~50 cd × s/m²) in WT and *Cntf*^{-/-} mice, which were kept in darkness for the indicated minutes after photobleaching the visual pigments. Asterisks indicate significant differences between WT and *Cntf*^{-/-} mice (*, *p* < 0.04; **, *p* ≤ 0.009; ***, *p* ≤ 0.003); error bars show S.D. (*n* = 6). B, relative contents of 11cRAL in eyecups of mice under the indicated light conditions: immediately after PB of the visual pigment or kept in darkness for the indicated minutes after PB. Error bars show S.D. (*n* = 4).

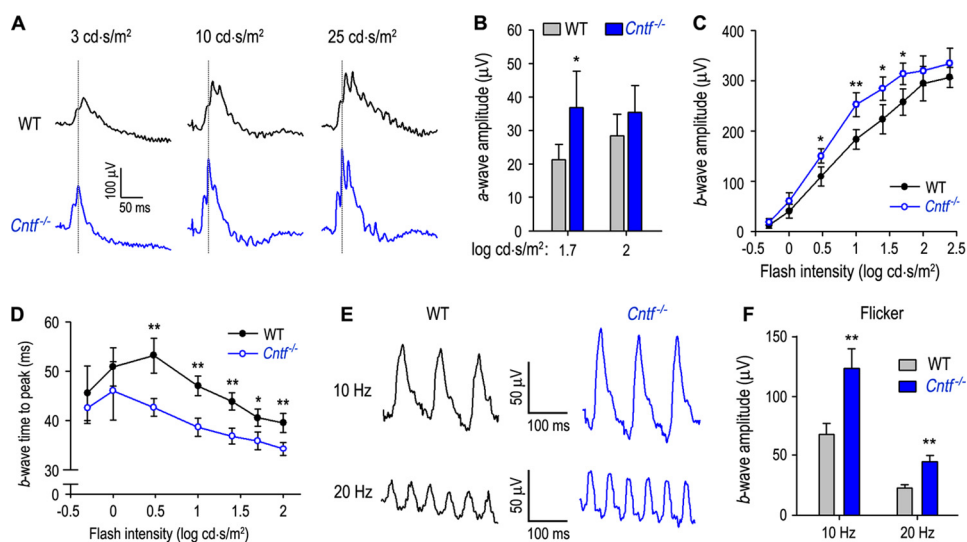


Figure 6. Hyper cone phototransduction in *Cntf*^{-/-} mice. A, representative raw photopic ERG responses of WT and *Cntf*^{-/-} mice to the indicated flashes under a rod saturating background light (32 cd/m²). B and C, amplitudes of *a*-wave (B) and *b*-wave (C) in photopic ERG responses elicited with the indicated flash intensities in light (32 cd/m²)-adapted WT and *Cntf*^{-/-} mice. Asterisks indicate significant differences between WT and *Cntf*^{-/-} mice (*, *p* < 0.03; **, *p* ≤ 0.001); error bars show S.D. (*n* = 7). D, times from stimulus onset to photopic *b*-wave peaks of WT and *Cntf*^{-/-} mouse cones. E, representative raw flicker ERG responses of WT and *Cntf*^{-/-} mice to 10 cd × s/m² flashes with frequencies of 10 or 20 Hz on the rod saturating background light. F, amplitudes of *b*-wave in the flicker ERG responses of WT and *Cntf*^{-/-} mice. Error bars show S.D. (*n* = 5, **, *p* ≤ 0.001).

min), and then kept them in darkness for 5 days. Retinoid analysis showed that the contents of 11cRAL and the total retinoids were significantly reduced in the *Cntf*^{-/-} eyes exposed to the intense light for 75 min, as compared with those in WT mice under that same light conditions (Fig. 7A). *Cntf*^{-/-}, but not WT, mice under that same light conditions displayed a significant reduction in rod visual function, as measured by scotopic ERG (Fig. 7B). Notably, both *a*- and *b*-wave amplitudes are markedly reduced in *Cntf*^{-/-} mice (Fig. 7B). This reduction of rod visual function is associated with a dramatic decrease in rod OS length and contents of rhodopsin in the *Cntf*^{-/-} retina (Fig. 7, C and D). Morphological analysis showed that 75 min exposure of 12,000 lux light caused a severe retinal degeneration in *Cntf*^{-/-}, but not WT, mice (Fig. 8, B and C). The superior ONL in *Cntf*^{-/-} retina was reduced in thickness to 2–8 nuclei versus 9–10 nuclei in WT retina whereas the inferior ONL in the *Cntf*^{-/-} retina was reduced to 4–8 nuclei versus 10–11 nuclei in WT retina (Fig. 8D).

Generally, cones are resistant to light-induced degeneration (45, 46). To know if CNTF deficiency increases susceptibility of cones to photodamage, we did immunohistochemistry for cones in WT and *Cntf*^{-/-} mice exposed to the intense light for

75 min. Lengths of both S-cone and M-cone OSs are significantly shortened in *Cntf*^{-/-} mice (Fig. 9, B and C). Consistent with these observations, immunoblot analysis showed that expression levels of S- and M-opsins were reduced at least 40% in the *Cntf*^{-/-} retinas, as compared with those in WT retinas (Fig. 9, D and E).

Discussion

Phototransduction and the visual cycle are the most important functions in sensing and converting light signal into biochemical and electrical signals in the retina. Light sensitivities of both rod and cone visual pigments rely on their 11cRAL chromophore provided by the visual cycle. In this study, we showed that CNTF deficiency resulted in up-regulation of opsins and the visual cycle enzymes, leading to an increase in thickness of ONL, length of rod and cone OSs, and the rates of the 11cRAL regeneration. These molecular and morphological changes led to 1) hyper phototransduction of rods and cones, 2) faster recovery of rod light sensitivity, and 3) increase in susceptibility of rods and cones to photodamage in *Cntf*^{-/-} mouse.

Up-regulation of opsins and the visual cycle enzymes (Figs. 1–3) are the most important primary phenotypes in the

CNTF inhibits visual pigment generation to protect retina

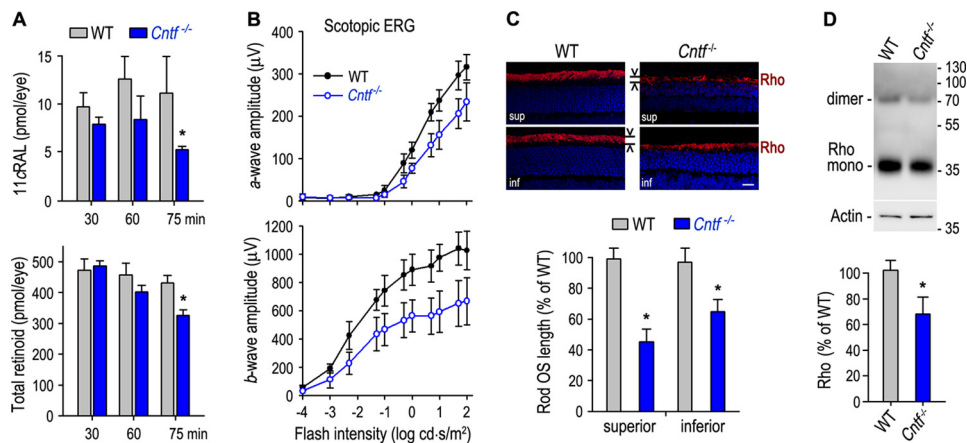


Figure 7. Reduction of retinoids, rod visual function, and rhodopsin contents in *Cntf*^{-/-} mice exposed to intense light. A, contents of 11cRAL and total retinoids in the eyecups of WT and *Cntf*^{-/-} mice kept in darkness for 5 days after exposing to 12,000 lux light for the indicated time. All asterisks indicate a significant difference between WT and *Cntf*^{-/-} mice ($p < 0.02$); error bars show S.D. ($n = 4$). B, amplitudes of a-wave (upper panel) and b-wave (bottom panel) in scotopic ERG responses of WT and *Cntf*^{-/-} mice to the indicated flashes. Error bars show S.D. ($n = 5$). All data shown in B–D are from mice kept in darkness for 5 days after exposing to 12,000 lux light for 75 min. C, immunohistochemistry for Rho in retinal sections of WT and *Cntf*^{-/-} mice. Histograms show relative lengths of Rho-positive OS in WT and *Cntf*^{-/-} retinal sections. Scale bar denotes 20 μ m. D, immunoblot analysis of Rho in WT and *Cntf*^{-/-} retinas. Histogram shows immunoblot intensities of Rho from *Cntf*^{-/-} retinas relative to Rho intensity in WT retinas.

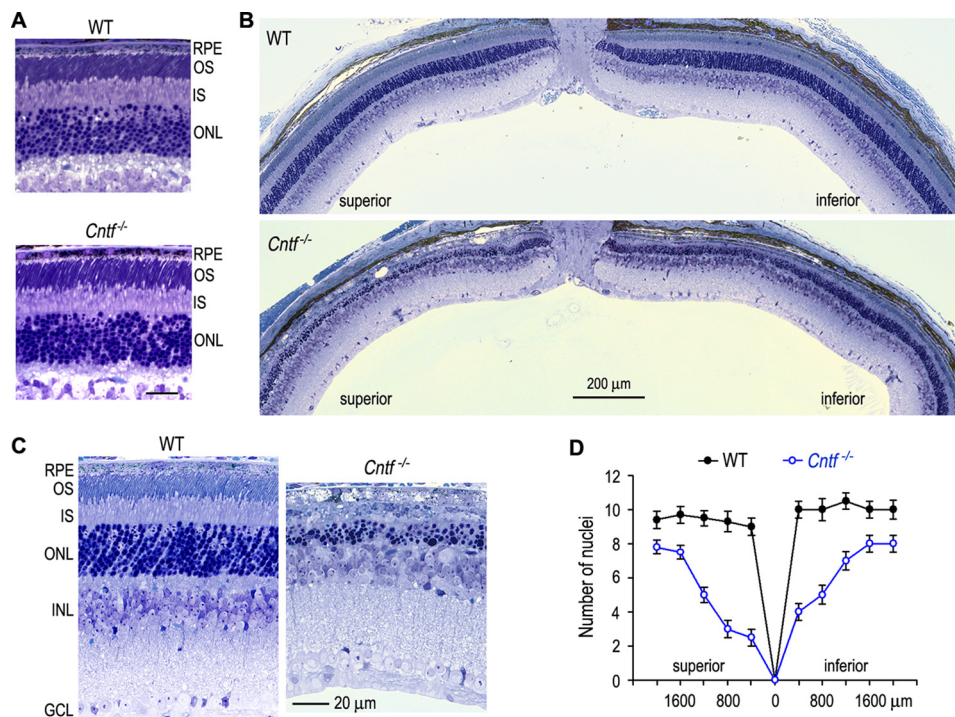


Figure 8. Retinal photodamage in *Cntf*^{-/-} mice. A, light microscopic images of WT and *Cntf*^{-/-} superior retinas before photodamage. Scale bar denotes 20 μ m. B, light microscopy showing severe retinal degeneration in *Cntf*^{-/-}, but not WT, mice kept in darkness for 5 days after exposing to 12,000 lux light for 75 min. C, higher magnification images of the superior retinas of WT and *Cntf*^{-/-} mice exposed to the intense light. D, thickness of the retinal outer nuclear layers in WT and *Cntf*^{-/-} mice in B. Numbers on the x-axis indicate distance from optic nerve head. Error bars show S.D. ($n = 4$).

Cntf^{-/-} visual system. CNTF might down-regulate these proteins through the heterotrimeric receptor complex (17), which can activate at least three distinct downstream pathways: Jak-STAT3, Ras-MAPK, and PI3K-AKT pathways (22, 47). The Jak-STAT3 pathway is the most studied pathway in CNTF/LIF signaling. Exogenous LIF activated STAT3 in mouse photoreceptors (48) and down-regulated rhodopsin (49), whereas CNTF did not activate STAT3 in any photoreceptors but down-regulated rhodopsin (23). CNTF might down-regulate rhodopsin by acting on Muller cells (23) or through the Ras-MAPK and/or PI3K-AKT pathways in rods.

Opsin up-regulation was accompanied by morphological changes in the *Cntf*^{-/-} retina. The ONL thickness and lengths of rod and cone OSs were increased in *Cntf*^{-/-} mice. The increase in the ONL thickness might be associated with the decrease in programmed cell death in the ONL of postnatal *Cntf*^{-/-} mice (Fig. 1). CNTF has been shown to promote programmed death of postmitotic rod precursor cells; and blocking the CNTF/LIF pathway reduced cell death during mouse retinal development, resulting a thicker ONL (50).

CNTF deficiency also resulted in an increase in lengths of both rod and cone OSs. This observation is consistent with

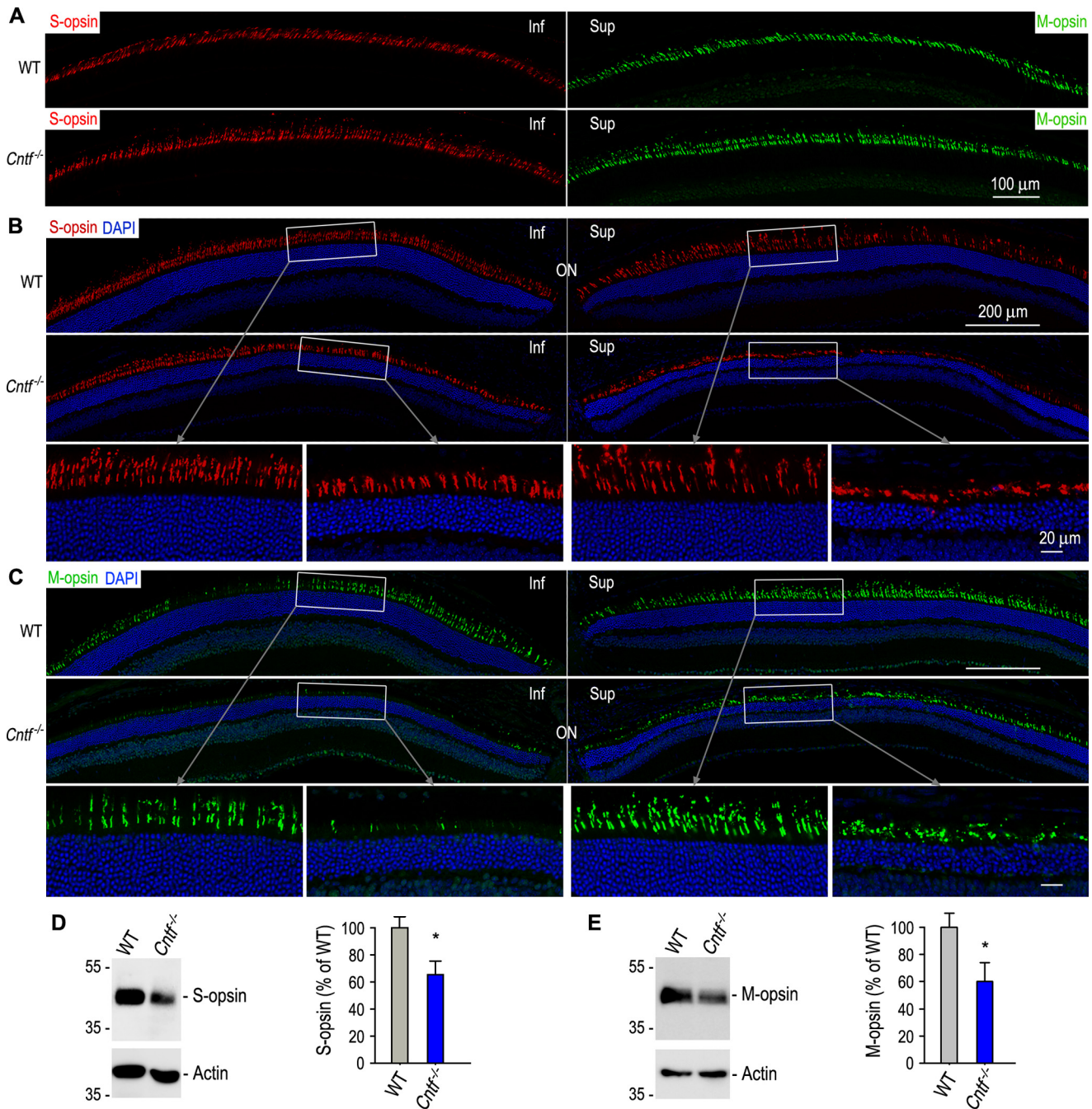


Figure 9. Light-induced degeneration of cones in *Cntf*^{-/-} mice. *A*, immunohistochemistry showing cone opsin-positive OS in WT and *Cntf*^{-/-} mice before photodamage. *B* and *C*, immunohistochemistry for S-opsin (*B*) or M-opsin (*C*) in retinal sections of WT and *Cntf*^{-/-} mice kept in darkness for 5 days after exposing to 12,000 lux for 75 min. Nuclei were counterstained with DAPI (blue). The areas of the rectangles are shown in the higher magnification images (bottom panels). *Inf*, inferior; *sup*, superior; *ON*, optic nerve. All data shown in this figure are from mice exposed to 12,000 lux light for 75 min. *D* and *E*, immunoblot analysis of S-opsin (*D*) or M-opsin (*E*) in WT and *Cntf*^{-/-} retinas. Actin was detected as sample loading control. Histograms show relative immunoblot intensities of S-opsin (*D*) or M-opsin (*E*) in WT versus *Cntf*^{-/-} retinas. Asterisks indicate significant differences between WT and *Cntf*^{-/-} mice ($p < 0.02$); error bars show S.D. ($n = 3$).

three previous studies: 1) CNTF induced a reversible rod OS shortening in WT rat (23), 2) CNTF reduced cone OS length in the *rds* mouse (51), and 3) transgenic LIF inhibited rod and cone OS maturation and/or elongation (52). In a rat line carrying a rhodopsin mutation, however, CNTF promoted regeneration of cone OS (32). The different effects of CNTF on cone OS morphology may depend on the presence or absence of strong inflammatory stimulation (53), which produce numerous signaling proteins in the retina (54).

11cRAL bound with opsins functions as a molecular switch for initiating phototransduction in response to light stimuli. Light-mediated isomerization of 11cRAL to all-*trans*-retinal induces opsin activation and phototransduction. To restore light sensitivity to opsins that have lost 11cRAL, 11cRAL must be regenerated and recombined with apo-opsins. RPE65 and LRAT are critical enzymes in the visual cycle that generates 11cRAL. We found that the synthesis rate of 11-*cis*-retinol in the *Cntf*^{-/-} RPE was increased due to up-regulation of RPE65

CNTF inhibits visual pigment generation to protect retina

and LRAT. RPE65 and LRAT might be up-regulated in part through the gp130-STAT3 pathway in the RPE (49). We further found that recovery of rod light sensitivity was accelerated due to an increase in 11cRAL synthesis in the *Cntf*^{-/-} RPE (Fig. 5).

Increase in opsin expression and 11cRAL synthesis suggest that the *Cntf*^{-/-} rods and cones contain more light-sensitive visual pigments. This might cause the enhanced responses of *Cntf*^{-/-} rods and cones to light stimuli (Figs. 4–6). The hyperphototransduction led to greater hyperpolarization of rods and cones, reflected in elevation of *a*-wave amplitudes in both scotopic and photopic ERGs of *Cntf*^{-/-} mice.

Although scotopic *a*-wave amplitudes were increased in *Cntf*^{-/-} mice *b*-wave amplitudes, implicit times were similar to those in WT mice. In addition, expression levels of mGluR6 and PKC α in the *Cntf*^{-/-} retina were also similar to those in WT retina. These results suggest that synaptic function between rod and rod ON-BC, which produce *b*-wave responses (55, 56), is not significantly changed or may be slightly reduced in *Cntf*^{-/-} mice. Overexpression of CNTF or LIF has been shown to cause BC disorganization (51, 57). Exogenous CNTF has also been shown to promote BC differentiation (21, 51) through increasing Ath3 expression and pleiotrophin secretion (58, 59). These studies and the similar *b*-wave amplitudes in *Cntf*^{-/-} and WT mice suggest that CNTF plays an important role in the development, differentiation, and function of rod ON-BCs.

Interestingly, cone *b*-wave amplitudes and implicit times in photopic ERG elicited with certain flash intensities were significantly higher or shorter in *Cntf*^{-/-} mice versus WT mice. These results are consistent with the previous studies that show significant reduction of photopic *b*-waves in WT animals overexpressing CNTF (41, 42). Because *b*-waves of photopic and flicker ERGs are mainly from depolarizing cone ON-BCs (60, 61), our results suggest that the synaptic function between cone and cone ON-BC, as well as cone-induced depolarization of cone ON-BCs are enhanced in *Cntf*^{-/-} mouse.

One of the pathological consequences of CNTF deficiency was retinal photodamage. *Cntf*^{-/-} rods exhibited hyper susceptibility to light-induced degeneration. This phenotype is consistent with the well-known phenomenon: the rodent superior retina has much higher susceptibility to photodamage, as compared with the inferior retina. Length of photoreceptor OS in the superior retina is ~30% longer than that in the inferior retina (62). The longer OSs in the *Cntf*^{-/-} and the rodent superior retinas may contain more rhodopsin, the primary mediator of retinal degeneration induced by light (9). Activated rhodopsin may increase the DNA-binding activity of the transcription factor AP-1 to promote light-induced apoptosis of photoreceptors (9, 63). In *Drosophila*, activated rhodopsin induced photoreceptor apoptosis by promoting clathrin-dependent endocytosis of rhodopsin-arrestin complexes (64).

It has been known that the higher expression level and activity of RPE65 cause an increase in susceptibility of retina to photodamage (10–12). Conversely, decrease in the visual cycle rate or RPE65 activity reduce retinal photodamage (65–67). RPE65 promotes retinal photodamage by facilitating the visual cycle that provides 11cRAL to generate light-sensitive visual pigments. Up-regulated RPE65 and LRAT have accelerated the synthesis rate of 11cRAL in the *Cntf*^{-/-} mouse. This acceler-

ated visual cycle plus increased expression of opsins resulted in an increase in the formation of light-sensitive visual pigment, therefore promoted retinal photodamage in *Cntf*^{-/-} mice.

In general, cones are resistant to photodamage (45, 46). In this study, however, we observed that intense light caused a severe degeneration of both M- and S-cones in the *Cntf*^{-/-} mouse. This result suggests that CNTF deficiency itself and increased formation of cone visual pigments contributed to cone photodamage in *Cntf*^{-/-} mouse. In addition, increased visual cycle and phototransduction rates may result in elevation of oxidative stress in cones due to an increase in contents of 11cRAL and all-*trans*-retinal in the *Cntf*^{-/-} cones. Retinaldehydes have been shown to play a critical role in retinal photooxidative damage (68–70). Cones are known to have high susceptibility to oxidative stress (71–74). Overexpression of NRF2, a master antioxidant transcription factor, effectively protected cones from degeneration in animal models of retinal degeneration (75). These studies and our results suggest that reduction of neuroprotective signals and increase in oxidative stress promoted light-induced cone degeneration in *Cntf*^{-/-} mice. In summary, we identified CNTF as a critical signal that down-regulates rod and cone opsins, as well as RPE65 and LRAT, to suppress visual pigment over formation and retinal photodamage.

Experimental procedures

Animals

We crossed *Cntf*^{-/-} mice (76) with WT 129S2/Sv mice (Charles River Laboratories), then intercrossed the heterozygous offspring to yield *Cntf*^{-/-} mice homozygous for the Leu-450 allele of the *Rpe65* gene. The homologous *Cntf* knockout mutation and the Leu-450 alleles were confirmed by PCR and DNA sequencing, as described previously (12, 76). Except where noted, mice were maintained in 12 h cyclic light at ~30 lux. All animal experiments were performed on both sexes of WT and *Cntf*^{-/-} mice at 3, 5, or 6 weeks of age in accordance with the Association for Research of Vision and Ophthalmology statement for the use of animals in ophthalmic and vision research and the protocols approved by the Institutional Animal Care and Use Committee for LSU Health Sciences Center.

Eyecup RPE, cell culture, and transfection

After removal of the anterior section and neural retina, the RPE in 3-week-old *Cntf*^{-/-} mouse eyecups were maintained in DMEM/F-12 medium (Thermo Fisher Scientific Inc.) supplemented with 10% heat-inactivated fetal bovine serum (FBS) and antibiotics (77). The 293T-LC cells maintained in DMEM (Invitrogen) with 10% FBS (78) were transfected with pRK5 or pRK5-CNTF plasmid DNA using the PolyJet transfection reagent (SignaGen Laboratories). After transfection, the cells were maintained overnight in DMEM/F-12 medium containing 5% FBS. These media with or without CNTF were incubated with the eyecup RPE for 6 h. All cultures were maintained in an incubator containing 5% CO₂.

Immunoblot analysis

Protein samples in the Laemmli buffer containing 50 mM DTT were incubated for 10 min at 70 °C (or room temperature

for opsins), separated by SDS-PAGE in a 10, 12, or 4–12% gradient polyacrylamide gel, and transferred to an Immobilon-P membrane (MilliporeSigma). The membrane was incubated in blocking buffer, primary antibody, and horseradish peroxidase-conjugated secondary antibody against rabbit, mouse, or goat IgG. Antibodies against RPE65 (79), LRAT (80, 81), CNTF (Santa Cruz Biotechnology), Ezrin (ProteinTech), interphotoreceptor retinoid-binding protein (IRBP) (82), mGluR6 (GeneTex), PKC α , rhodopsin, CAR, M-opsin (MilliporeSigma), S-opsin (MilliporeSigma, Santa Cruz Biotechnology), or β -actin (MilliporeSigma) were used as the primary antibodies. Immunoblots were visualized with the enhanced ECL-Prime and quantified (82).

Immunohistochemistry

Retinal cryosections were prepared from the dorsal-ventral midline of mouse eye as described previously (83). Briefly, enucleated mouse eyeballs were fixed overnight with 4% paraformaldehyde in 0.1 M phosphate buffer (PB). After removing cornea and lens, eyecups were immersed in 15% sucrose in 0.1 M PB for 2 h, in 30% sucrose in 0.1 M PB for 2 h, and then in a 1:1 mixture of 30% sucrose and optimal cutting temperature (OCT) medium (Sakura Finetechnical) overnight at 4 °C. After embedding eyecups in OCT, 15- μ m thick sections were cut on a Shandon Cryotome SME cryostat (Thermo Scientific). The sections were immunostained with the primary antibodies listed in the method of immunoblot analysis and secondary antibodies, as described previously (82). Nuclei were counterstained with DAPI (Sigma). Images were captured with a Zeiss LSM710 Meta confocal microscope with a $\times 20$ objective lens or $\times 40$ oil-immersion lens. Lengths of rod and cone OS as well as numbers of cells were measured using ImageJ software (National Institutes of Health).

TUNEL assay

This assay was performed using the *in situ* cell death detection kit (Roche Applied Science) following the manufacturer's protocol. Briefly, retinal cryosections washed with 0.1% sodium citrate in 0.1% Triton X-100, PBS were incubated with terminal deoxynucleotidyl transferase (TdT) and fluorescein-dUTP for 1 h at 37 °C. After rinsing three times in PBS containing 0.05% Tween 20, nuclei were counterstained with DAPI. Numbers of TUNEL-positive nuclear in the ONL-outer plexiform layer and in the inner nuclear layer-inner plexiform layer of whole retinal sections were counted separately using an Olympus BX61VS microscope equipped with a digital camera and VS-ASW FL software.

Quantitative RT-PCR

Total RNA was extracted from mouse retina or RPE using a PureLink RNA mini kit (Invitrogen), and was reverse-transcribed to cDNA using SuperScript III (Invitrogen). Quantitative PCR was performed on an iCycler iQTM Real-Time PCR Detection System (Bio-Rad) using a two-step qRT-PCR kit with SYBR Green (Invitrogen) and primer sets specific for mouse RPE65, opsin, and 18S rRNA. Three mice of each genotype were analyzed and all samples were run in duplicates. Starting templates were normalized after determining 18S rRNA C_t val-

ues for each sample (84). Relative mRNA levels of RPE65 were determined from the ΔC_t values.

Electroretinography (ERG)

Overnight dark-adapted 6-week-old mice were anesthetized with an intraperitoneal injection of 100 mg/kg of ketamine and 10 mg/kg of xylazine. The pupils were dilated with 1% tropicamide. ERG was recorded from the corneal surface using a silver-silver chloride wire electrode referenced to a subcutaneous electrode in the mouth. A needle electrode in the tail served as the ground. A drop of 2.5% methylcellulose was placed on the cornea to ensure good electrical contact and to prevent corneal desiccation during the entire procedure. Single-flash ERG recordings were performed in a Ganzfield dome (Espion e2, Diagnosys LLC) under dark-adapted (scotopic) and light-adapted (photopic) conditions. For scotopic ERG, single flash stimuli were presented with various intensities, reaching from $-4 \log \text{cd} \times \text{s/m}^2$ to $2.5 \log \text{cd} \times \text{s/m}^2$, in interstimulus intervals of 0.5–2 min (depending on the stimulus intensity). Three to four responses were averaged for each step. For photopic ERG, animals were light adapted for 10 min by exposing to a white 32 cd/m^2 light, and ERG responses were obtained with white flashes ($-0.3 \sim 2.4 \log \text{cd} \times \text{s/m}^2$) on the rod-saturating background light (32 cd/m^2). Five responses to 10-s interval flashes were averaged for each step. Intensity-response amplitude data were displayed on log-linear coordinates using SigmaPlot 11 software. Flicker ERG responses were recorded in light-adapted mice with flicker flashes on the rod-saturating background. The flicker stimuli had an intensity of $10 \text{ cd} \times \text{s/m}^2$ with frequencies of 10 or 20 Hz.

Retinoid isomerase assay

RPE homogenates in 20 mM HEPES buffer (pH 7.4) containing protease inhibitor mixture were prepared from mouse eyecups without the neural retina. The homogenates were irradiated for 10 min on ice with 365-nm light from a Spectroline Model EN-140L UV light source to destroy endogenous retinoids. Each assay mixture contained 200 μg of cell homogenate, 10 μM all-*trans*-retinol, and 6% BSA. After incubating for 2 h in darkness at 37 °C, retinoids were extracted with hexane and analyzed by HPLC, as described below.

Analysis of retinoids

Retinoids in mouse ocular tissues or *in vitro* enzyme assays were extracted with hexane and analyzed by normal-phase HPLC (84). In brief, retinoids in hexane extractions were evaporated, dissolved in 100 μl of hexane, and separated on a silica column (Zorbax-Sil 5 μm , $250 \times 4.6 \text{ mm}$, Agilent Technologies) by gradient (0.2–10% dioxane in hexane at 2.0 ml/min flow rate) or nongradient (10% dioxane in hexane at 1.0 ml/min flow rate) elution of mobile phase on an Agilent 1100 HPLC system equipped with a photodiode array detector (Agilent Technologies). Spectral data were acquired for all eluted peaks. Quantitation was performed by comparison of peak areas to calibration curves established with authentic retinoid standards.

Light microscopy

Mouse retinal sections were prepared as described previously (84). In brief, mice were fixed by intracardiac perfusion

CNTF inhibits visual pigment generation to protect retina

with a mixture of 2% paraformaldehyde and 2.5% glutaraldehyde in 0.1 M PB (pH 7.4). Light cautery was applied at the superior pole of the cornea to mark the orientation before enucleation of the eyeball. A window was cut in the cornea, and the eye was immersed in primary fixative and rotated at room temperature for 2 h. The anterior segment was removed, and the remaining eyecup was refrigerated overnight in primary fixative. The eyecup was trimmed into temporal and nasal hemispheres and immersed in 1% osmium tetroxide in 0.1 M PB (pH 7.2) for 1 h. Following dehydration in a graded series of alcohols, the hemispheres were embedded in an Epon/Araldite mixture (5:3, v/v). Care was taken to orient the eyecups so that sections were obtained through the vertical meridian to ensure sampling of both the superior, green-sensitive cones and the inferior, blue-sensitive cones. Sections were cut at 1 μ m and stained with 1% toluidine blue and 1% sodium borate, then photographed using a $\times 20$ objective lens or a $\times 60$ oil-immersion lens in the Olympus BX61VS microscope mentioned above. Except for whole retinal images, all images were obtained from retinal sections at a distance of 600 μ m superior or inferior to the optic nerve.

Light-induced retinal degeneration

WT and *Cntf*^{-/-} mice were dark-adapted for 3 days. After dilation of the pupils under dim red light (Kodak Wratten 1A), mice were exposed to 12,000 lux of white fluorescent light for 30, 60, and 75 min and then kept in darkness for 5 days. Retinoid contents, visual function, and retinal structures of these mice were analyzed as described above.

Statistical analysis

SigmaPlot Version 11 (Systat Software, Inc.) was used for statistical analyses. Data were expressed as the mean \pm S.D. of three or more independent experiments indicated in the figure legends. Differences between WT and *Cntf*^{-/-} mice were determined by single comparisons with an unpaired two-tailed Student's *t* test. The significance threshold was set at 0.05 for all statistical tests.

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CNTF inhibits visual pigment generation to protect retina

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