GRK1-Dependent Phosphorylation of S and M Opsins and Their Binding to Cone Arrestin during Cone Phototransduction in the Mouse Retina

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The shutoff mechanisms of the rod visual transduction cascade involve G-protein-coupled receptor (GPCR) kinase 1 (GRK1) phosphorylation of light-activated rhodopsin (R*) followed by rod arrestin binding. Deactivation of the cone phototransduction cascade in the mammalian retina is delineated poorly. In this study we sought to explore the potential mechanisms underlying the quenching of the phototransduction cascade in cone photoreceptors by using mouse models lacking rods and/or GRK1. Using the “pure-cone” retinas of the neural retina leucine zipper (Nrl) knock-out (KO, −/−) mice (Mears et al., 2001), we have demonstrated the light-dependent, multi-site phosphorylation of both S and M cone opsins by in situ phosphorylation and isoelectric focusing. Immunoprecipitation with affinity-purified polyclonal antibodies against either mouse cone arrestin (mCAR) or mouse S and M cone opsins revealed specific binding of mCAR to light-activated, phosphorylated cone opsins. To elucidate the potential role of GRK1 in cone opsin phosphorylation, we created Nrl and Grk1 double knock-out (Nrl−/−Grk1−/−) mice by crossing the Nrl−/− mice with Grk1−/− mice (Chen et al., 1999). We found that, in the retina of these mice, the light-activated cone opsins were neither phosphorylated nor bound with mCAR. Our results demonstrate, for the first time in a mammalian species, that cone opsins are phosphorylated and that CAR binds to phosphorylated cone opsins after light activation.

Keywords: cone opsin; phosphorylation; cone arrestin; phototransduction; coimmunoprecipitation; mouse retina

Introduction

Phototransduction mechanisms are well documented in rod photoreceptors and now are regarded as a classic model system of G-protein-coupled receptor (GPCR) signaling (Baylor, 1996; Leskov et al., 2000; Fain et al., 2001). The phototransduction cascade in cone photoreceptors is thought to be similar to that of rods because rod homologs of phototransduction components are expressed in cones. Nevertheless, the kinetics of photore- sponse in the two photoreceptor types are different. Cones are several hundred-fold less sensitive to light than are rods (Pugh and Lamb, 2000), and yet they recover sensitivity much faster than rods after light flashes that generate similar membrane currents (Baylor et al., 1979; Perry and McNaughton, 1991).

In rod photoreceptors the timely deactivation of photoactivated rhodopsin (R*) by GPCR kinase 1 (GRK1)-mediated R* phosphorylation followed by rod arrestin binding is critical for effective rod vision (Wilden et al., 1986a; Baylor and Burns, 1998). The low abundance of cones in most mammalian retinas has made it difficult to elucidate directly the biochemical and molecular mechanisms underlying cone phototransduction. It has been suggested that cones may rely primarily on regeneration for inactivation of photolyzed visual pigments, because patients with Grk1 null mutation have either normal or only slightly abnormal photopic vision (Cideciyan et al., 1998). However, mouse retinas lacking GRK1 expression display profoundly slowed recovery of cone photoresponses, suggesting that GRK1-dependent opsin phosphorylation may be involved in the shutoff of cone phototransduction in the mammalian retina (Lyubarsky et al., 2000). Identification of a cone-specific GRK7 (Hisatomi et al., 1998; Weiss et al., 1998, 2001; Chen et al., 2001) and cone arrestin (CAR) (Murakami et al., 1993; Craft et al., 1994; Craft and Whitmore, 1995; Hisatomi et al., 1997; Maeda et al., 2000; Smith et al., 2000; Zhu et al., 2002a,b) further supports the hypothesis that similar shutoff mechanisms exist in cone photoreceptors.

In this study we provide direct biochemical evidence of cone opsin phosphorylation and CAR binding to phosphorylated cone...
opsins during phototransduction, using the neural retina leucine zipper (Nrl) knock-out (KO, /−/−) mice (Mears et al., 2001). Nrl, a transcription factor of the basic motif leucine zipper family, is expressed preferentially in rod photoreceptors (Swaroop et al., 1992; Swain et al., 2001) and implicated in rod-specific gene regulation (Rehentulua et al., 1996; Bessant et al., 1999) and photoreceptor differentiation (Mears et al., 2001). The analysis of the Nrl /−/− retinas revealed a complete lack of rod function and rod-specific gene expression, with a concomitant increase in S-cone function and cone-specific gene expression, including S opsin, cone transducin, and CAR (Mears et al., 2001). Therefore, the photoreceptors of the Nrl /−/− mouse retina are functionally and biochemically cones, although they are proposed to be cone–rod intermediates because of their abnormal morphology.

Using the pure-cone retinas of the Nrl /−/− mice, we demonstrate that both S and M cone opsins are phosphorylated after light exposure and that CAR selectively binds to light-activated, phosphorylated cone opsins. We also created Nrl and Grk1 double KO (Nrl /−/− Grk1 /−/−) mice by crossing the Nrl /−/− with the Grk1 /−/− mice (Chen et al., 1999; Lyubarsky et al., 2000) and show that, in these double KO mice, neither S nor M opsin is phosphorylated either in light or in darkness, nor does CAR bind to the light-activated cone opsins, suggesting that GRK1 is responsible for cone opsin phosphorylation during phototransduction in the mouse retina.

Materials and Methods

Animals. C57BL/6J mice were purchased originally from the Jackson Laboratories (Bar Harbor, ME). The Nrl /−/− mice (Mears et al., 2001) and Grk1 /−/− mice (Chen et al., 1999; Lyubarsky et al., 2000) were described previously. To generate Nrl /−/− Grk1 /−/− double KO mice, we bred the Nrl /−/− mice with the Grk1 /−/− mice. After two rounds of breeding, mice homozygous null ( /−/−) for Nrl were identified by Southern blot analysis as previously described (Mears et al., 2001), and mice null for Grk1 were identified by genomic PCR, using primers specific for the Grk1 wild type (WT) or for the Grk1 KO construct. The WT and Nrl /−/− mice were reared under a 12 hr light/dark cycle, and the Grk1 /−/− and the Nrl /−/− Grk1 /−/− double KO mice were reared in total darkness.

Antiser generation. Rabbit antiser against the peptides of mouse S opsin (residues 1–11, MSGEDDFYLFQ) and M opsin (residues 3–16, QRLTGEQTLDHYED) were made for our research project by Zymed Laboratories (South San Francisco, CA) and affinity-purified against the peptides with the SulfoLink kit (Pierce, Rockford, IL) as previously described (Zhu and Craft, 2000).

Immunoblot analysis. Total retinal homogenates from normal C57 mice were used for immunoblot analysis with either anti-S or anti-M opsin antibody and HRP-conjugated anti-rabbit secondary antibody and were visualized by an enhanced chemiluminescence (ECL) kit (Amer sham Biosciences, Arlington Heights, IL) (Craft et al., 1998).

Immunohistochemistry. The protocol for immunohistochemistry on mouse retinal sections has been published previously (Zhu et al., 2002b). For cone opsin antibody characterization the sections were incubated with either the anti-M or anti-S opsin peptide polyclonal antibody, followed by incubation with a fluorescein anti-rabbit IgG. To visualize all cones, we incubated the sections with biotinylated peanut agglutinin (PNA; Vector Laboratories, Burlingame, CA) for 1 hr at room temperature (RT) and then with Texas Red-avidin D (Vector Laboratories) for 1 hr at RT. After washing, the slides were coverslipped and photographed.

For immunofluorescent triple labeling, the retinal sections were incubated with sequential primary antibodies, including a rabbit polyclonal [anti-M opsin, anti-S opsin, or anti-mCAR LUMII] (Zhu et al., 2002b) and a mouse monoclonal antibody [GRK1-specific D11 (Zha et al., 1998; Chen et al., 2001; Weiss et al., 2001), Affinity BioReagents, Golden, CO] at 1:1000 and 1:200 dilutions, respectively. After the washing steps the sections were reacted with a mixture of AMCA-anti-rabbit IgG (1: 100) and fluorescein anti-mouse IgG (1:100; both from Vector Labora-
(13,000 rpm, 4°C, 30 min); the supernatants were taken for isoelectric focusing (IEF) gels. Seven microfilters (one-tenth of a retina) of each sample were applied to a 1 mm IEF gel containing 5% Ready Mix IEF acrylamide (Amersham Biosciences), 6.3% pH 3–10 ampholytes (Amersham Biosciences), 13.3% glycerol, and 0.5% dodecyl maltoside. The samples were applied 4 cm from the cathode of a 13 cm gel with 1 M NaOH and 1 M phosphoric acid as the cathode and anode buffers, respectively. The gel was electrophoresed at 2500 V, 150 mA, 23 W for 2 hr at 10°C on a flat-bed IEF apparatus (LKB-Wallac, Gaithersburg, MD). The gel was preelectrophoresed for 30 min before the samples were applied. After electrophoresis the proteins were transferred to a PVDF membrane and probed with antibodies to S opsin, M opsin, or rhodopsin.

Immunoprecipitation (IP) was performed with the Protein A-Agarose IP kit (KPL, Gaithersburg, MD). Four hundred microfilters of the retinal supernatants (dark or light) were mixed with 400 µl of a 50% suspension of protein A-agarose in lysis buffer with no protease inhibitors or okadaic acid were added, and the samples were centrifuged at 14,000 rpm, 4°C, for 10 min to remove cell debris.

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**Results**

**Membrane association of mCAR in a light-dependent and GRK1-dependent manner**

To explore whether CAR contributes to quenching light-activated cone opsin in the mammalian retina, we analyzed the redistribution of mCAR to retinal membranes in response to light. The distribution of mCAR and mouse SAG (mSAG) was analyzed by immunoblot analysis of membrane and soluble proteins of retinal homogenates from either light- or dark-adapted mice. Figure 1 shows that 50% of the mCAR protein is in the membrane fraction (pellet) in the dark-adapted retina, whereas 81% is in the membrane fraction in the light-adapted retina (Fig. 1A), similar to the distribution of mSAG, which is mostly in the soluble fraction in the dark-adapted retina but redistributes to the membranes when exposed to light (Fig. 1C). These results are consistent with our previous observation that a portion of the mCAR immunoreactivity translocates to the cone outer segments in a light-dependent manner (Zhu et al., 2002b). Interestingly, in the Grk1−/− mouse retina (Fig. 1B,D) neither mCAR nor mSAG has a significant increase in membrane binding after light exposure as compared with the dark-adapted retinas, implying that the membrane binding of mCAR may be phosphorylation-dependent and that GRK1 may play a role in both rod and cone phototransduction in the mouse retina.

**Generation and characterization of anti-mouse S and M opsin antibodies**

To facilitate our cone opsin phosphorylation studies, we generated rabbit polyclonal antibodies against peptides derived from mouse S and M opsin peptide sequences and affinity-purified them against their respective peptide. Immunoblot analysis of retinal homogenates from normal C57BL/6J mice with the two affinity-purified antibodies identified a single band of the predicted molecular weight of the respective opsin, i.e., 37.5 kDa for S opsin and 39 kDa for M opsin (Fig. 2A,B). Minor bands were seen after a longer film exposure time (data not shown). To verify the specificity of the antibodies, we did a peptide-blocking experiment by incubating the primary antibody with 100 times excess...
throughout the whole cell body of the cone photoreceptors (Fig. 3) in contrast to mCAR, which has a diffuse staining pattern to rods, and is localized exclusively to the outer segments of the mouse retina. GRK1 is expressed in both S and M cones, in addition to the retina. Adult C57 mouse retinal frozen sections shows that mCAR also is localized throughout the whole photoreceptor outer segments (Fig. 4), consistent with the distribution of either GRK1 or mCAR with S and M opsins by immunohistochemistry. In the normal C57BL/6J mouse retina, both GRK1 and mCAR are expressed in both S and M cones (Fig. 4), consistent with the distribution of cone opsins by immunohistochemistry. Western blot analysis was performed with normal adult C57BL/6J mouse retinal homogenate, and immunohistochemistry was done on C57BL/6J mouse retinal frozen sections. Cone photoreceptor cells were labeled with biotinylated peanut agglutinin and visualized with Texas Red-avidin D in red (c, d), Dual immunofluorescence labeling verified both S and M opsin immuno-reactivities localized to cone cells (c, f), OS, Outer segments; IS, inner segments; ONL, outer nuclear layer. Scale bar, 20 μm.

Both GRK1 and mCAR are expressed in both S and M cone photoreceptors

Previous studies have shown the expression of both GRK1 (Lubarsky et al., 2000) and mCAR (Zhu et al., 2002b) in cone photoreceptors of normal mouse retinas. Both GRK1 and mCAR also are expressed in the retina of the Nrl<sup>−/−</sup> mouse by Western and Northern blot analyses, respectively (Mears et al., 2001). We examined the colocalization of either GRK1 or mCAR with S and M opsins by immunohistochemistry. In the normal C57BL/6J mouse retina, GRK1 is expressed in both S and M cones, in addition to rods, and is localized exclusively to the outer segments (Fig. 3) in contrast to mCAR, which has a diffuse staining pattern throughout the whole cell body of the cone photoreceptors (Fig. 3J,K). In the Nrl<sup>−/−</sup> mouse retina (Fig. 5) GRK1 is colocalized with either S or M opsin to the short outer segments of all photoreceptors, whereas mCAR is expressed throughout the whole photoreceptor layer, with the most intense staining in the outer segments and the synaptic terminals. mCAR also is colocalized with both S and M opsins in the outer segments of the Nrl<sup>−/−</sup> retinas.

Light- and GRK1-dependent phosphorylation of cone opsins

To determine cone opsin phosphorylation after light exposure, we examined the light-dependent incorporation of orthophosphate into the pure-cone retinas of the Nrl<sup>−/−</sup> mice. Exposure of isolated intact retinas from Nrl<sup>−/−</sup> mice to bright
sunlight (~8000 fc) resulted in phosphorylation of both S and M opsins, which was not observed in retinas from WT mice, presumably because of masking by the abundant phosphorylated rhodopsin (Fig. 6A). Retinas that were kept in darkness did not show appreciable levels of opsin phosphorylation in either of the two mouse lines (Fig. 6A). We chose sunlight as the stimulus because it is a naturally occurring light source, and it contains a strong UV component, which can activate the murine S opsin most efficiently (peak sensitivity at ~360 nm). However, 10 min of room light exposure also resulted in the phosphorylation of both S and M opsins (data not shown).

To investigate the role of GRK1 in cone opsin phosphorylation, as indicated by the analysis of the Grk1−/− mice (Lyubarsky et al., 2000), we bred Nrl−/− mice with Grk1−/− mice, and we analyzed the double knock-out mice lacking both Nrl and Grk1 (Nrl−/−Grk1−/−). Neither S nor M opsin was phosphorylated in the Nrl−/−Grk1−/− mouse retina after light exposure (Fig. 6A), although the double KO retinas had the same morphology as those of the Nrl−/− mice (data not shown), and both S and M opsins were expressed in the double KO mouse retina at equivalent levels to those in the Nrl−/− mouse retina (Fig. 6B).

Figure 6. In situ light-dependent phosphorylation of opsins ex vivo. A, In situ phosphorylation of opsins. WT, Nrl−/−, and Nrl−/−Grk1−/− mice were dark-adapted overnight and killed. The retinas were dissected under IR light and incubated in phosphate-free Krebs’ buffer containing 1.25 mCi/ml 32P orthophosphate for 30 min at RT in the dark. One retina of each strain was homogenized in SDS sample buffer in the dark (D) while the other retina was exposed to bright sunlight (L) for 10 min before homogenization. The proteins were resolved on an 11.5% SDS-PAGE gel and transferred to a PVDF membrane. O. Opin phosphorylation was detected by autoradiography on a PhosphorImager screen. B, Immunoblot analysis of the same membrane in A, using polyclonal antibodies to mouse S or M opsin or the rhodopsin monoclonal antibody 104, sequentially, to observe the quantity and location of the opsins relative to the radioactive phosphate (32P) identified in A. Note the slightly higher molecular weight of the phosphorylated as compared with the unphosphorylated species of S and M opsin and rhodopsin.

Phosphorylation of cone opsins at multiple sites
Recent evidence shows that multiple phosphorylation of rhodopsin at the C terminus is necessary for rapid and reproducible deactivation of rhodopsin (Mendez et al., 2000), which is consistent with earlier observations that up to nine phosphates could be incorporated into a single R* in vitro (Wilden and Kuhn, 1982) and that the addition of each phosphate reduces the ability of R* to activate transducin (Arshavsky et al., 1987; Wilden, 1995). Mass spectrometric analysis further confirmed multiple phosphorylation of rhodopsin in vivo (Kennedy et al., 2001). As many as 9 and 12 potential phosphorylation sites exist in the S and M opsin C-terminal sequences, respectively (Fig. 7A). To explore the possibility of multiple phosphorylation of cone opsins, we performed IEF with retinas from Nrl−/− mice. Immunoblot analysis of IEF gels of solubilized membrane fractions from Nrl−/− mouse retinas exposed to direct bright sunlight reveals potentially nine phosphorylated species of S opsin (Fig. 7B), implying that all serine and threonine residues near the C terminus of S opsin can be phosphorylated with exposure to naturally occurring bright light. Because of the much lower expression level of M opsin in these mouse retinas as compared with S opsin, we detected only five phosphorylated species of M opsin. However, other species with more than five phosphates may exist but may be beyond our sensitivity of detection. Under the same experimental conditions we detected only five phosphorylated species of rhodopsin in WT mouse retina, although up to six have been reported (Mendez et al., 2000). In the samples kept in darkness a monophosphorylated species of each opsin was detected in addition to the predominant unphosphorylated species (Fig. 7B).
Previous studies have shown that 2% of rhodopsin is monophosphorylated in the dark-adapted mouse retina either in the presence or absence of GRK1 (Chen et al., 1999). The significance of opsin monophosphorylation in the dark currently is unknown. Alternatively, the second band above the unphosphorylated species that is present in both dark and light conditions may be caused by other post-translational modifications of the cone pigments.

Coimmunoprecipitation of mCAR with light-activated, phosphorylated cone opsins
To examine whether mCAR interacts with light-activated, phosphorylated cone opsins, we performed IP on Nrl<sup>−/−</sup> mouse retinal proteins with either the mCAR antibody or cone opsin antibodies as well as the CRX antibody, which is also an affinity-purified rabbit polyclonal antibody (Zhu and Craft, 2000), as a negative control. Both the anti-mCAR antibody LUMIJ (α-mCAR) and the anti-cone opsin antibodies (α-S and α-M opsin) coimmunoprecipitated phosphorylated proteins (32P) only in the light-exposed retina, whereas the anti-CRX antibody (α-CRX) did not coimmunoprecipitate any phosphorylated proteins either in the light-exposed or dark-adapted retinas. The mCAR antibody precipitated an equal amount of mCAR protein from both the light-exposed and dark-adapted retinas. However, only in the light-exposed retinas, when cone opsins were activated and phosphorylated, did the mCAR antibody coprecipitate S and M opsins. Also, both the S and M opsin antibodies precipitated an equal amount of their respective opsin proteins from both the light-exposed and dark-adapted retinas, but the S opsin antibody coprecipitated mCAR only from the light-exposed retinal homogenate. The M opsin antibody, however, did not precipitate appreciable levels of mCAR protein either from the light-exposed or dark-adapted retinas (Fig. 8A), most likely because of the sensitivity limit of the assay.

In the Nrl<sup>−/−</sup> Grk1<sup>−/−</sup> mouse retina (Fig. 8B), where no cone opsin phosphorylation occurred after light activation (Fig. 6A), no 32P-labeled proteins were precipitated by any of the antibodies. The mCAR antibody did not pull down either cone opsin, nor
did the cone opsin antibodies pull down mCAR either in the light-exposed or dark-adapted retina, although each antibody pulled down equal amounts of its corresponding protein from both the light-exposed and dark-adapted retinas (Fig. 8). These results further confirm that the phosphorylated protein bands in the \( ^{32}P \) incorporation experiments (Fig. 6A) and in the coimmunoprecipitation experiments with the Nrl \(^{-/-}\) mice (Fig. 8A) are GRK1-phosphorylated cone opsins and that the binding of mCAR to cone opsins is phosphorylation-dependent, similar to rod arrestin binding to phosphorylated R*.

**Discussion**

In the vertebrate retina two distinct cell populations of photoreceptors, rods and cones, coexist. Rods are specialized for dim light vision, whereas cones provide high-acuity color vision. Although cones comprise only 5% of photoreceptors in the human retina, they are far more important than rods for vision in daylight. Even at night our world is flooded with enough artificial light so that our cones can function, and we can see clearly in color. Despite its importance in vision the cone phototransduction cascade and its underlying molecular mechanisms are poorly understood. In this study we aimed to explore the potential involvement of opsin phosphorylation and CAR binding in the cone phototransduction cascade, using mouse models lacking rods and/or GRK1.

**Cone opsin phosphorylation in the mammalian retina**

Two critical events, receptor phosphorylation and arrestin binding, participate in the deactivation of GPCRs (Freedman and Lefkowitz, 1996; Krupnick et al., 1997; Krupnick and Benovic, 1998; Pitcher et al., 1998). In rod photoreceptors the phosphorylation of R* by GRK1 initiates deactivation and limits the response amplitude (J. Chen et al., 1995; C. Chen et al., 1999). In the absence of arrestin, phosphorylation alone can decrease the catalytic activity of R* by 50% (Xu et al., 1997).

Cone opsins from the all-cone retina of lizard, from the cone-dominant retina of chicken, and from the rod-dominant retina of carp are phosphorylated in vitro either by the endogenous kinase or by bovine GRK1 (Walter et al., 1986; Fukada et al., 1990; Tachibanaki et al., 2001). In the present study, using the purecone retinas of the Nrl \(^{-/-}\) mice, we have provided the first biochemical evidence of cone opsin phosphorylation in a mammalian species. It is puzzling why, in the rhodopsin KO mouse retina, no light-dependent phosphorylation signals are observed, although these mice contain fully functional cones (Mendez et al., 2000). Because cones account for only 3% of the photoreceptors in the mouse retina (Carter-Dawson and LaVail, 1979; Jeon et al., 1998), the light-dependent cone opsin phosphorylation signals may have been masked by the nonlight-dependent phosphorylation signals from other phosphoproteins.

**GRK1 is responsible for light-dependent cone opsin phosphorylation in the mouse retina**

Seven distinct mammalian GRKs, each differing in tissue distribution and in regulatory properties, have been identified to date. Among these enzymes GRK1 is involved in phototransduction. Proof that GRK1 phosphorylation of R* is required for normal inactivation of R* in vivo has come from single-cell recordings of photoresponses in rods expressing mutant rhodopsin lacking the C-terminus (Chen et al., 1995) and, more recently, of responses of rods in Grk1 \(^{-/-}\) mice (Chen et al., 1999).

GRK1 is localized in both rod and cone photoreceptors in many species, including the rod-dominant human, monkey, and mouse (Zhao et al., 1998; Luibarsky et al., 2000; Weiss et al., 2001) and the cone-dominant chicken (Zhao et al., 1999). Although a cone-specific GRK7 has been reported recently in retinas of both cone- and rod-dominant species, the absence of GRK7 protein in the mouse photoreceptor outer segment layer rules out the possibility that GRK7 is responsible for cone opsin phosphorylation in the mouse retina (Chen et al., 2001). Functional analysis of mouse retinas lacking GRK1 expression shows defects in the recovery of cone-driven photoresponses (Luibarsky et al., 2000). Consistent with this observation, we found that mCAR lost its light-dependent binding to the membrane fraction in the Grk1 \(^{-/-}\) mouse retina, similar to the behavior of rod arrestin (Fig. 1B,D). The colocalization of GRK1 with both S and M opsins and the lack of light-dependent phosphorylation signals in the Nrl \(^{-/-}\) Grk1 \(^{-/-}\) double KO mouse retina suggest that GRK1 is the responsible kinase for cone opsin phosphorylation during phototransduction in the mouse retina. Electrophysiological analysis of the photoresponses of the photoreceptors of the Nrl \(^{-/-}\) Grk1 \(^{-/-}\) mouse retina will verify the functional significance of cone opsin phosphorylation.

In contrast to the Grk1 \(^{-/-}\) mouse phenotype, human patients with Grk1 null mutation have either normal or slightly abnormal photopic vision (Cideciyan et al., 1998). A detailed study of species-specific differences in GRK1 and GRK7 expression reveals that the cones of mice and rats express only GRK1, and the cones of pigs and dogs express only GRK7, whereas primate cones coexpress both GRK7 and GRK1 (Weiss et al., 2001). The expression of GRK7 in human cones may complicate partially for the loss of GRK1 in patients with Grk1 null mutation. Recent studies reveal that the deactivation kinetics of S cones in human patients with enhanced S-cone syndrome (ESCS) were much slower than long/middle wavelength-sensitive (L/M) cones in normal, ESCS, or GRK1-null Oguchi retinas (Cideciyan et al., 2003). Immunocytochemistry reveals no expression of either GRK1 or GRK7 in ESCS S cones, whereas ESCS L/M cones have no GRK7 and have only GRK1. These results suggest that the absence of both GRK1 and GRK7 causes a greater degree of abnormality in cone deactivation than that caused by the deficiency of either GRK alone (Cideciyan et al., 2003). Although the relative roles of GRK7 and GRK1 in primate cones still await elucidation, the above data clearly demonstrate that opsin phosphorylation is a critical component of the deactivation mechanisms in cone photoreceptors of the human retina.

The role of GRK1 in rods is to inactivate R* that, in concert with regeneration, leads to the recovery of sensitivity (Cideciyan et al., 1998). Because cones recover sensitivity much faster than rods and the maximal rate of visual pigment regeneration is 2000-fold higher in cones than in rods (Schnapf et al., 1990), it is expected that the rate of cone opsin phosphorylation is also much faster than that of rhodopsin. Indeed, Tachibanaki and coworkers have reported that phosphorylation is >20 times faster in cone cell membranes than in rod membranes in the carp retina after light activation (Tachibanaki et al., 2001). Recently, an alternate visual cycle that generates visual pigments at a rate 20-fold faster than the known visual cycle has been demonstrated, which is proposed to be responsible for sustained daylight vision mediated by cone photoreceptors (Mata et al., 2002). Studies of the kinetics of phosphorylation of mammalian rod and cone opsins after light activation are underway, using retinal membrane preparations from WT (rhodopsin) and Nrl \(^{-/-}\) mice (S opsin).
CAR is involved in the cone phototransduction cascade

Members of the arrestin family are involved in GPCR desensitization, internalization, and GPCR-mediated activation of MAPK pathways. In the rod photoreceptor the rod arrestin binding after receptor phosphorylation is necessary to complete the quench of the light-activated phototransduction cascade (Wilden et al., 1986b; Xu et al., 1997). CAR may be involved in the shutoff mechanisms of the cone phototransduction cascade by binding to light-activated arrestins and its cone photoreceptor localization suggest that CAR may play a key role in the modulation of phototransduction in cones as rod arrestin does in rods. Our previous studies showed that mCAR binds to purified embryonic chicken outer segment membranes in a light- and phosphorylation-dependent manner and that a portion of the mCAR protein translocates to the cone outer segments in the mouse retina after light exposure (Zhu et al., 2002b). The results presented here provide further evidence that CAR behaves like rod arrestin in response to light, suggesting that CAR may be involved in the shutoff mechanisms of the cone phototransduction cascade by binding to light-activated, phosphorylated cone opsins.

To elucidate further the function of CAR, we are creating mCAR KO mice. Characterization of the morphological, biochemical, and electrophysiological phenotypes of these mice will clarify the physiological role of CAR in the shutoff of cone phototransduction as well as verify its other functions in both the retina and the pigment gland.

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