2009-07-27

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https://hdl.handle.net/2144/3397
The 9-methyl group of retinal is essential for rapid Meta II decay and phototransduction quenching in red cones

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Cone photoreceptors of the vertebrate retina terminate their response to light much faster than rod photoreceptors. However, the molecular mechanisms underlying this rapid response termination in cones are poorly understood. The experiments presented here tested two related hypotheses: first, that the rapid decay rate of metarhodopsin (Meta) II in red-sensitive cones depends on interactions between the 9-methyl group of retinal and the opsin part of the pigment molecule, and second, that rapid Meta II decay is critical for rapid recovery from saturation of red-sensitive cones after exposure to bright light. Microspectrophotometric measurements of pigment photolysis, microfluorometric measurements of retinol production, and single-cell electrophysiological recordings of flash responses of salamander cones were performed to test these hypotheses. In all cases, cones were bleached and their visual pigment was regenerated with either 11-cis retinal or with 11-cis 9-demethyl retinal, an analogue of retinal lacking the 9-methyl group. Meta II decay was four to five times slower and subsequent retinol production was three to four times slower in red-sensitive cones lacking the 9-methyl group of retinal. This was accompanied by a significant slowing of the recovery from saturation in cones lacking the 9-methyl group after exposure to bright (>0.1% visual pigment photoactivated) but not dim light. A mathematical model of the turn-off process of phototransduction revealed that the slower recovery of photoreponse can be explained by slower Meta decay of 9-demethyl visual pigment. These results demonstrate that the 9-methyl group of retinal is required for steric chromophore–opsin interactions that favor both the rapid decay of Meta II and the rapid response recovery after exposure to bright light in red-sensitive cones.

INTRODUCTION

A remarkable feature of our visual system is the ability to resolve changes in light intensity that span more than 12 orders of magnitude. The upper end of this range is mediated by cone photoreceptors, which can report fast changes in ambient illumination and come in several types that are sensitive to different parts of the wavelength spectrum. A key requirement for cones to encode this wide range of light intensities is their ability to adapt sensitivity based both on the background light level (light adaptation) and on the extent of bleaching of their visual pigment (bleaching adaptation).

Both light adaptation and bleaching adaptation allow the cones to reduce their sensitivity as the background light intensity increases, the goal of which is to prevent response saturation and extend the operating range of the cell (for review see Fain et al., 2001 and Burns and Arshavsky, 2005). Light adaptation is subserved by mechanisms that are mediated by intracellular Ca2+ and counteract the effect of steady light by reducing the rate of cGMP hydrolysis and accelerating the rate of its synthesis. Brighter background lights that cause a significant bleaching of the visual pigment can also induce bleaching adaptation, which like background adaptation results in reduced sensitivity and speeded response time course, and occurs by residual activation of phototransduction by bleached pigment (Cornwall and Fain, 1994; Leibrock et al., 1998). Under these conditions, the recovery of sensitivity ultimately requires the regeneration of the visual pigment, a process called dark adaptation (for review see Lamb and Pugh, 2004).

Pigment regeneration takes place via a series of reactions collectively referred to as the visual cycle, which in cones has been recently shown to occur through reciprocal interactions with the Müller cells (Mata et al.,...
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2002; Wang et al., 2009) as well as the retinal pigment epithelium (Wang et al., 2009). In this process, the all-trans retinol (ROL) generated during the bleaching of the cone visual pigment is transported to both cell types. In Müller cells, ROL is reconverted to 11-cis retinol and sent back to the cones, which first oxidize it and then use it to regenerate pigment. In the retinal pigment epithelium, 11-cis retinal is produced directly, which after translocation of receptors, is used to regenerate visual pigment. Key to the recovery of sensitivity is ultimately the regeneration of the pigment, a process that can take minutes (Lamb and Pugh, 2004); however, cones are known to recover fully their circulating current and thus responsiveness after exposure to bleaching light on a much faster time scale (20–100 ms) (Kenkre et al., 2005). It should be noted that this time is substantially faster than the 20-min recovery time for rod photoreceptors measured under similar light conditions (Thomas and Lamb, 1999). We sought to understand mechanistically how the fast recovery of the cone-circulating current occurs before the pigment is regenerated.

It should be noticed that the speed of recovery, as one indication of dark adaptation, can be characterized by various parameters. Thus, faster recovery may mean shorter time in saturation measured between the moment of strong bleaching flash and first sign of recovery of circulating current. A closely related but not identical measure is the steepness of recovery of preflash “dark” current after the saturation ceased. Finally, full recovery demands complete restoration of dark-adapted sensitivity, the process that can take a substantially longer time than the restoration of the dark current (Kruepel et al., 2003). In the present study, we use the term “accelerated recovery” to designate both shortening time in saturation and steeper restoration of the circulating current (recovery of response amplitude to saturating flashes).

Recently, we examined the role of cone visual pigment structure in recovery of response amplitude of red-sensitive cones (hereafter referred to as red cones) exposed to bright light (Estevez et al., 2006). In this study, we observed that red cones regenerated with 11-cis 9-demethyl (9-DM) retinal, an analogue of retinal lacking the 9-methyl group, were up to 10 times slower to recover their circulating current after intense light exposure compared with cones containing native visual pigment. This correlated with slower recovery of steady-state guananyl cyclase rates in these cells, suggesting that slow response recovery was due to sluggish termination of phototransduction activation. We concluded that visual pigment structure, particularly the steric interaction of red cone opsin with the 9-methyl group of retinal, plays an important role in ensuring the rapid response recovery after bright light exposure. However, the specific mechanisms of that role remained unresolved.

Early biochemical studies that examined the effect of the 9-methyl group of retinal on opsin had suggested that this group may play a steric role in the efficient hydrolysis of the Schiff base linkage between retinal and red cone opsin (metarhodopsin [Meta] II decay), followed by the efficient release of all-trans retinal from the opsin binding pocket (Das et al., 2004). Our results were consistent with this hypothesis; however, no direct measurements of Meta II decay in intact cone photoreceptors containing 9-demethyl–based visual pigment had yet been made. The experiments presented here were designed to directly test this hypothesis. Accordingly, we have made microspectrophotometric measurements of the rates of visual pigment photolysis as well as the time course of ROL formation in isolated salamander red cones containing normal visual pigment or pigment lacking the 9-methyl group of retinal. Direct measurements of the reduction of all-trans retinal to ROL by retinol dehydrogenase were also made by single-cell microfluorometry. All of these results were correlated with single-cell electrophysiological measurements of flash response kinetics and the time course of recovery of circulating current in darkness after bleaching exposure. These results have been fitted with a kinetic model of quenching of the phototransduction cascade. Accordingly, we conclude that the 9-methyl group of retinal is critical for rapid Meta II decay as well as rapid flash response recovery in salamander red cone photoreceptors.

MATERIALS AND METHODS

Animals and preparation

All experiments were performed on red cone photoreceptors isolated from the retinae of larval salamanders. Two different species of salamanders were used in this study: Ambystoma tigrinum and Ambystoma mexicanum. Physiological, microspectrophotometric, and microfluorometric experiments on cones from both species were performed in the laboratory at Boston University School of Medicine. Microspectrophotometric experiments conducted in the laboratory at The Russian Academy of Sciences in St. Petersburg, Russia, were performed only on Ambystoma mexicanum cones, as only that species was available. When data from a single species are illustrated or discussed, appropriate notation is provided in the text or figure legends. Data derived from experiments on both species were compared to ensure that species differences did not introduce confounding variables. Ambystoma tigrinum were purchased from Charles D. Sullivan, Inc. Ambystoma mexicanum used for experiments in the St. Petersburg laboratory were purchased from a local breeder; those used for experiments in Boston were purchased from the Ambystoma Genetic Stock Center. All experiments were performed according to procedures approved by the Institutional Animal Care and Use Committee at Boston University School of Medicine and in accordance with the Guide for the Care and Use of Laboratory Animals and the Animal Welfare Act.

Salamanders were dark-adapted overnight, killed by decapitation, and double-pithed under dim red illumination. Eyes were enucleated and hemisected in physiological solution. These as
well as all subsequent manipulations were viewed under infrared illumination. The physiological solution used for all dissections and subsequent measurements contained (in mM): 110 NaCl, 2.5 KCl, 1.6 MgCl₂, 1.0 CaCl₂, 10 dextrose, 10 HEPES, and 100 mg/l bovine serum albumin. The pH was adjusted to 7.8. Retinas were separated from the retinal pigment epithelium by dissection with fine forceps, and photoreceptors were isolated by chopping the retina with a small piece of razor blade. The resulting suspension containing intact cells was transferred to a chamber mounted on a microscope stage designed for microspectrophotometric, electrophysiological, or microfluorometric experiments. Individual cells were viewed using infrared-sensitive CCD cameras connected to the microscopes. All experiments were performed at 20°C. The methods are similar to those described previously (Cornwall et al., 1983, 1990, 2000; Ala-Laurila et al., 2006; Estevez et al., 2006).

**Pigment regeneration**

The native visual pigment chromophore contained in dark-adapted cones of both species is a mixture of 11-cis retinal (A1) and 11-cis,3,4-dehydroretinal (A2). This native pigment was removed by exposure to bright light (bleaching) and replaced with an exogenous chromophore, either 11-cis retinal or 9-DM retinal. 11-cis retinal and the 9-DM analogue used in these studies were exclusively in the A1 form. These were synthesized and purified in the Department of Ophthalmology at the Medical University of South Carolina (Crouch et al., 2002). Retinoid solutions were prepared by dissolving 50 µg retinoid in 3 µl ethanol and diluting in 3 ml of physiological solution to create a working concentration of 30–50 µM retinoid in solution containing <0.3% ethanol as a retinoid carrier. This retinoid solution was added directly to the bath containing bleach-adapted cells by application with a pipetter. The final concentration of retinoid in the bath was 18–30 µM, and cells were exposed to this solution for ~5–8 min before washing with normal physiological solution.

The preparation was monitored for ~20 min before subsequent microspectrophotometric, electrophysiological, or microfluorometric experiments were performed. Finally, after this 20-min period, the cell was tested to ensure a fully recovered steady state of regeneration by measuring flash response sensitivity or absorbance spectra. The rates of chromophore binding and pigment regeneration by 11-cis retinal and 9-DM retinal have been shown previously to be roughly the same (Kono, M., personal communication). Therefore, the absorbances at three wavelengths chosen close to absorbance maxima of PSB and Meta I (450 nm), Meta II and all-trans retinal (380 nm) and retinol (340 nm), and two polarizations can be analyzed by series of linear equations whose solutions at any moment of time yield the contribution of each photoproduct to the recorded spectra (Kolesnikov et al., 2003; Ala-Laurila et al., 2006). The absorbances can further be converted to concentrations normalized to bleached pigment by taking into account relative molar extinctions of the products. Meta I and II are in rapid equilibrium, and for kinetic computations can be considered as a single product, hereafter referred to as Meta (Ala-Laurila et al., 2006).

**Micorspectrophotometry**

A high-speed dichroic microspectrophotometer was used to measure the time course of photolysis of visual pigment and ROL.
RESULTS

Microspectrophotometric measurements of pigment photolysis

Fig. 1 compares average absorbance spectra of the native visual pigment of dark-adapted *Ambystoma mexicanum* red cone outer segments and spectra measured after exhaustive bleaching and regeneration with either 11-cis retinal or 9-DM retinal. The native dark-adapted spectrum had an absorbance maximum at 592 nm (Fig. 1, gray line). This represents a mixture of pigments, each composed of the same opsin, one containing the A1 form of 11-cis retinal and the other with the A2 form. The pure A1 form has a spectral maximum at 567 nm, and the A2 form has its maximum at 615 nm. Native dark-adapted spectra were fitted by a procedure that convolved the two pure pigment spectra into one with the measured spectral shape as described in Materials and methods. This procedure allowed an estimate of the proportion of each form; the average native spectrum shown represents a mixture of 0.33 A1/0.67 A2.

Bleaching and subsequent regeneration with 11-cis retinal yields the visual pigment whose absorbance spectrum (Fig. 1, black line) can well be fitted with the A1-based template with $\lambda_{\text{max}} = 567$ nm. On the other hand, the 9-DM-based pigment (Fig. 1, red line) peaks at 520 nm, and the spectrum can be described with the A2-based 520-nm template plus a small ($\approx 7\%$) admixture of the native A2 chromophore. This value is significantly blue-shifted with respect to 11-cis–based pigment, a finding consistent with previous studies (Das et al., 2004; Estevez et al., 2006). Variation in A1/A2 ratios among animals, and possible admixture of native chromophore(s) in regenerated pigment, may account for a small difference in $\lambda_{\text{max}}$ values of salamander red cone pigment reported here and elsewhere (Makino et al., 1999; Estevez et al., 2006).

![Figure 1](image)

Normalized absorbance spectra of *Ambystoma mexicanum* red cones containing different visual pigment chromophores. Averaged T-spectra of red cones in their native dark-adapted state (containing a mixture of A1 and A2 retinal [gray curve]; $\lambda_{\text{max}} = 592$ nm; $n = 11$), red cones regenerated with 11-cis retinal (11-cis [black curve]; $\lambda_{\text{max}} = 567$ nm; $n = 11$), or red cones regenerated with 9-DM retinal (red curve; $\lambda_{\text{max}} = 520$ nm; $n = 9$). Smooth lines show corresponding template fittings. Native pigment is best approximated with a 30/70% mixture of A1/A2 chromophore ($\lambda_{\text{max}} = 567$ and 615 nm, respectively). 11-cis–regulated pigment is pure 567 nm A1. 9-DM spectrum can be represented by the sum of 93% 520-nm 11-cis A2 pigment and 7% of native 11-cis A2, 615 nm chromophore.
value is represented in curves 2 of all panels and can be visualized by the reduced amplitude of the initial 380-nm peak. The all-trans retinal released from meta-products is then converted to ROL, as indicated by the progressive short-wave shift of the absorbance maximum, finally to a peak below 350 nm (curves 3 in all panels; peak not shown). Both retinal and retinol can be followed more clearly with the spectra recorded at L-polarization (not depicted).

A remarkable feature of the photolysis of the 9-DM pigment is its substantially slower rate compared with both the A1 and A2 pigments containing methylated chromophores. Although the Meta peak in the native and the 11-cis retinal–regenerated pigments decays to a half of its original height at ~5 s post-bleach, the half-decay time of the 9-DM metapigment is near 30 s (Fig. 2, curves 2 in all panels).

As explained in Materials and methods, recording spectra at two polarizations and reading absorbances at three wavelengths allow computation of the concentrations of all bleaching products. The time courses of formation and decay of Meta, retinal plus its PSB, and retinol are shown in Fig. 3. The decay of metapigments in native cones and those with the 11-cis chromophore can satisfactorily be described with a single exponent having the time constant of 5.1 and 4.4 s, respectively (Fig. 3, top, gray and black curves, respectively). A two-exponential approximation was necessary for the decay of 9-DM metapigment, with time constants of 18 and 830 s, and amplitudes of 0.91 and 0.09 (Fig. 3, top, red curve). The half-decay times (τ₁/₂) were 3.5, 3.0, and 14 s for native, 11-cis–, and 9-DM–regenerated pigment, respectively. Collectively, these data show that the decay of metapigments (and hence retinal release from opsin) containing the 9-DM chromophore is roughly fivefold slower than with its methylated counterparts.

Microfluorometric measurements of ROL
A primary goal of the present study was to determine if the sluggish response recovery of red cones containing 9-DM visual pigment after their exposure to bright light is due to slowed hydrolysis of activated visual pigment. However, the decay of metapigments may not be the sole factor that controls the recovery from saturation after extensive bleach. All-trans retinal released from the chromophore pocket may secondarily bind to opsin forming complexes that retain some residual activity (Buczylko et al., 1996; Jäger et al., 1996). Hence, the reduction of retinal to retinol may be important for further quenching of the light-activated visual pigment. Our microspectrophotometric data allow determining the time course of retinol production, yet the calculations need an analysis of a complex mixture of photoproducts based on absorbance measurements in the spectral range where the performance of the instrument is rather poor and effects of possible impurities
can interfere with the measurement. To address this issue, we also used an alternative approach to measure the rate of retinol formation: microfluorometry. This method monitors the increase in intrinsic retinol fluorescence within bleached outer segments of isolated cones as all-trans retinal is reduced to ROL by retinol dehydrogenase, and then cleared from the outer segment (see Materials and methods). This method has the advantage that it does not rely on absorbance measurements of a mixture of bleached photoproducts, but rather on the production of ROL alone.

Microfluorometric experiments were performed on cones of both species. Fig. 4 (inset) shows an example of fluorescence production and clearance measured in a single red cone from _Ambystoma tigrinum_. First, the dark-adapted cone was drawn inner segment first into a pipette filled with the same physiological solution as the extracellular bath. The pipette was attached to a current amplifier to permit electrophysiological recording. The pipette served to stabilize mechanically the cell and permitted extracellular membrane current recordings of flash responses. Responses were recorded from a cell at two different wavelengths of light (440 vs. 620 nm) to establish that the cone was red sensitive (not depicted).

**Figure 3.** The time course of bleaching products in _Ambystoma mexicanum_ red cones containing different visual pigments as assessed by microspectrophotometry. The time course of metapigment decay (top; Meta I + II), the appearance of all-trans retinal and its protonated Schiff-bases (middle; RAL + PSB), and the formation of ROL (bottom) are plotted. Concentrations of bleaching products are expressed as a fraction of bleached visual pigment. Measurements were made from native red cones (gray; n = 11), or red cones regenerated either with 11-cis retinal (black; n = 11) or 9-DM retinal (red; n = 9). All curves were generated from the spectra shown in Fig. 2. The data in the top panel were fitted with a single-exponential decay function in the case of native and 11-cis pigments; however, a two-exponential approximation was required to fit the decay of 9-DM pigment. The data points in the middle and bottom panels were connected by straight lines. Note in the top panel, the time course of Meta decay in red cones containing 9-DM pigment (half-decay time $\tau_{1/2} = 14.1$ s) is four to five times slower than in red cones containing native or 11-cis-regenerated pigment ($\tau_{1/2} = 3.5$ and $\tau_{1/2} = 3$ s, respectively). Error bars show ± SEM.

**Figure 4.** The time course of retinol formation in salamander red cones containing different visual pigments as assessed by microfluorometry. The normalized relative fluorescence intensity is plotted as a function of time after a >95% bleach in red cones from _Ambystoma mexicanum_ under different visual pigment conditions. Data were normalized relative to the peak value and fitted with a single-exponential function. Red cones contained their native A1/A2 mixture (gray; n = 6; $\tau = 17.5 \pm 1.2$ s), or were regenerated with either 11-cis retinal (black; n = 5; $\tau = 20.8 \pm 0.9$ s) or 9-DM retinal (red; n = 6; $\tau = 70.9 \pm 6.7$ s). Error bars show ± SEM. Note that between the first and second measurements in all three cases, there is a rapid initial increase in fluorescence that is present in all fluorescence recordings; this has been reported previously (Tsina et al., 2004; Ala-Laurila et al., 2006) and is of unknown origin. The inset illustrates an experiment in which the rise and fall of retinol fluorescence were measured in a single red cone from _Ambystoma tigrinum_ under all three visual pigment conditions. First, the red cone was measured with its native pigment (gray squares), then containing 9-DM retinal visual pigment (red circles), and finally containing 11-cis retinal visual pigment (black triangles). 100 µM IRBP was present to facilitate the exchange of chromophore. Data are normalized relative to the peak value.
A fluorescence measurement was recorded (150 ms exposure to 360 nm excitation light) to establish the dark-adapted level of ROL in the cell. Then, the cone was bleached (>95%) for 10 s with 620 nm light in the presence of 100 µM IRBP. Immediately after the bleach, the rise and fall of outer segment fluorescence, corresponding to the formation and clearance of ROL, were recorded (Fig. 4, inset, gray squares). The dark current and sensitivity of the cone were monitored after the bleach until the cell had reached a new steady state. Then, the bleached cone was regenerated with exogenous 9-DM retinal, and again the dark current and sensitivity were monitored until the cell reached a fully dark-adapted state. The cell was then exposed to second identical bleach, again in the presence of 100 µM IRBP, and the rise and fall of outer segment fluorescence were again recorded (Fig. 4, inset, red circles). It is apparent from comparison of the rising phases of these recordings that fluorescence production was considerably slowed when the pigment was in its 9-demethyl form relative to that when it contained native chromophore. IRBP was again added to the bath during the bleach to facilitate removal of excess exogenous 9-DM retinal from the preparation. After washing the bleached cone with IRBP, the dark current and sensitivity were monitored until the cell again reached a new steady state. Then, the bleached cone was regenerated with exogenous 11-cis retinal, and the dark current and sensitivity were monitored. Finally, the cell was exposed to a third bleach in the presence of IRBP, and the rise and fall of outer segment fluorescence were recorded (Fig. 4, inset, black triangles). Here, the fluorescence rose rapidly, similar to the original condition in which the cell contained native visual pigment; however, it decayed somewhat more slowly than was the case after the first bleach. The reason for this is not known.

Thus, the production and clearance of fluorescence (ROL) was recorded from the outer segment of a single red cone under three different conditions: first when the cell contained its native complement of visual pigment, then when the cell contained regenerated 9-demethyl visual pigment, and finally when the cell contained regenerated 11-cis visual pigment. In the cases in which the cone contained either native pigment or pigment regenerated with exogenous 11-cis retinal, the rate of rise of retinol fluorescence was very similar, reaching a maximum value with time constants $\tau = 15.0$ s or $\tau = 19.8$ s (fitted curves not shown). This is consistent with the mean time constant for the post-bleach rise of retinol fluorescence in tiger salamander red cones ($\tau = 23.9 \pm 4.8$ s) reported previously (Ala-Laurila et al., 2006). In contrast, when the cone contained pigment regenerated with 9-DM retinal, the rise of retinol fluorescence was significantly slowed, reaching a peak with a time constant $\tau = 95.9$ s.

Fluorescence experiments performed on red cones from Ambystoma mexicanum produced essentially identical results. These average data are plotted in the main panel of Fig. 4, which has been shown on an expanded time scale to highlight the rising phase of the fluorescence. Here, the time constant of post-bleach retinol formation in red cones containing 9-DM retinal ($\tau = 70.9 \pm 6.7$ s; $n = 6$) was ~3.4 times slower than in cones regenerated with 11-cis retinal ($\tau = 20.8 \pm 0.9$ s; $n = 5$), and 4 times slower than in native cones ($\tau = 17.5 \pm 1.2$ s; $n = 6$).

Fig. 5 compares measurements of retinol formation obtained by the fluorescence technique (Fig. 4, main panel) with those obtained by microspectrophotometry (Fig. 3, bottom). Notice that in Fig. 4, the fluorescence data are scaled to unity at their maximum to facilitate comparison of their time courses. However, the fluorescence signal has no intrinsic standard. To convert the

Figure 5. A comparison of time courses of ROL production derived from microspectrophotometry versus microfluorometry. Data from Figs. 3 (bottom) and 4 have been replotted on the same graph for comparison. Data obtained by microspectrophotometry (MSP) are shown in black: $n = 11$ cells (native), 9 cells (9-DM), and 11 cells (11-cis). Fluorescence data are shown in gray: $n = 6$ cells (native), 6 cells (9-DM), and 5 cells (11-cis). Concentration of ROL in microspectrophotometry data are expressed as a fraction of bleached cone pigment. Fluorescence data are scaled for best visual match to microspectrophotometry. Error bars show ± SEM.
levels of fluorescence into relative concentrations of retinol normalized with respect to bleached visual pigment, fluorescence curves were scaled for best visual fit to the concentration time courses deduced from microspectrophotometry (Fig. 5). This provides the concentration of retinol as a fraction of bleached visual pigment, that is, expressed as mol retinol per mol rhodopsin. The average time course of retinol concentration estimated from microfluorimetry is illustrated by closed gray circles; retinol concentrations measured from microspectrophotometry are shown by open black circles. Data obtained with the two methods in native and 9-DM-regenerated cells are in a good agreement (Fig. 5, top and middle). The concentration of retinol in cells regenerated with the 11-cis chromophore increased more rapidly as judged from microspectrophotometry than that measured by microfluorimetry (Fig. 5, bottom). The reason for this discrepancy is unclear.

Because a saturating exponential function provides a good description of retinol time course (Fig. 4), we can estimate the initial rate of retinol production (in mol ROL/mol Rh per second) as $k_{ROL} = \frac{ROL_{\text{max}}}{\tau}$. Here, $ROL_{\text{max}}$ is maximum retinol concentration with respect to bleached visual pigment, as judged from microspectrophotometry (Figs. 3 and 5), and $\tau$ is the time constant of the front of the ROL curve (as assessed with microfluorimetry; see above). This yields the values of $0.022 \text{ s}^{-1}$, $0.007 \text{ s}^{-1}$, and $0.029 \text{ s}^{-1}$ for native, 9-DM–, and 11-cis–regenerated visual pigment, respectively. Thus, the retinol production in 9-DM cones proceeds three- to fourfold slower than in cones containing methylated chromophore.

**Electrophysiological recordings of Ambystoma mexicanum red cones under different visual pigment conditions**

The rationale for performing the microspectrophotometric measurements of visual pigment photolysis (Figs. 2 and 3) and the microfluorimetric measurements of ROL formation (Fig. 4) was motivated by our previous work (Estevez et al., 2006) in which salamander red cones regenerated to contain 9-DM visual pigment demonstrated sluggish flash response recovery after exposure to light that bleached >0.2% visual pigment. These experiments were performed on red cones isolated from *Ambystoma tigrinum* (Estevez et al., 2006). To provide a direct comparison between photoproduct decay rate and recovery of physiological responses in the same species, analogous electrophysiological measurements were made on red cones of *Ambystoma mexicanum*. These results are shown in Fig. 6. Cells were pre-bleached and regenerated with either 11-cis retinal (black traces; $n = 6$)

![Figure 6](image.png)

**Figure 6.** Flash response properties of *Ambystoma mexicanum* red cones containing different visual pigments. Shown are comparisons of average normalized flash responses to 500-ms test flashes that bleached various pigment fractions (pigment fraction shown in top right corner of each panel) in red cones. Cones were regenerated with 11-cis retinal (black traces; $n = 6$) or 9-DM retinal (red traces; $n = 4$). All traces have been smoothed by the Savitsky-Golay method (5 points to left and right). A significant difference in the photoresponse recovery between red and black traces is first observed at 0.1% bleach.
or 9-DM retinal (red traces; n = 4). The average photoreponses to 500-ms test flashes calculated to have bleached from 0.001 to 45% of the cone visual pigment are illustrated. Light intensities were adjusted with neutral density filters so that test flashes produced equivalent levels of photoactivation of the two chromophores, based on calculations described in Materials and methods. After each flash exposure, visual pigment was regenerated by excess retinoid partitioned in outer segment membranes (see Estevez et al., 2006). Flash responses elicited in cells containing 11-cis retinal and 9-DM retinal had identical time courses at low flash strengths (bleaching <0.1% of the pigment). However, as can be seen from comparisons of responses, exposure to flashes of increased light intensity that bleached ≥0.1% of the pigment resulted in significant differences in flash response shape; flash responses from cells containing 9-DM visual pigment were progressively slower to recover from saturation and re-store dark current as flash strength increased.

**Kinetic scheme of quenching of cone PDE activity**

Firsov et al. (2005) have shown that the progressive retardation of the recovery from saturation with increasing stimulus strength, similar to that seen in Fig. 6, can be explained by accumulation of metaproduts that stimulate, with a low efficiency, the phototransduction cascade. According to this idea, responses to low-intensity flashes are turned off by fast rhodopsin phosphorylation and transducin/phosphodiesterase (PDE) quenching. At supersaturating flash intensities, this fast phase is completed while the cell response still remains in saturation, and the recovery is now controlled by long-lived metaproduts. This idea is in general agreement with the findings presented here that show that slower photolysis of 9-demethyl metaproduts is accompanied by slower recovery of photoresponses to bright flashes.

To quantitatively test the idea, we applied the model of cascade turn-off suggested by Firsov et al. (2007) as illustrated in Fig. 7.

The model assumes that the photoexcited visual pigment, \( MII^* \), is quenched in a series of reactions, each step resulting in a progressive decline of \( MII^* \)’s ability to activate transducin \( (T) \). These steps include the phosphorylation of \( MII^* \) (formation of \( MII-P \)), subsequent arrestin binding (formation of \( M-P-Arr \)), decay of phosphorylated and arrestin-bound metaproduts to all-trans retinal and opsin \( (Ops) \), and eventual regeneration of the dark pigment \( (Rh) \). Each reaction is assumed to proceed accordingly to first-order kinetics with rate constants \( k_P \), \( k_A \), \( k_D \), and \( k_R \), respectively. GTPase (operating with rate constant \( k_E \)) quenches the active transducin \( T^*-PDE^* \) complex. Fractional (with respect to fully active \( MII^* \)) activities of the intermediates toward transducin are \( a_{M*} > a_{MP} > a_{MA} > a_{Ops} > a_{Rh} \). This is adapted from Firsov et al. (2005), with permission. (Insets) Cone responses to a dim 20-ms flash (noisy gray curves) with superimposed phototransduction model response (smooth black line) for regenerated 11-cis (top) and regenerated 9-DM (bottom) red cones. \( k_P \) and \( k_E \) were obtained from modeling. Light stimulus marks are shown below the curves.
proceed accordingly to first-order kinetics and is characterized by a corresponding rate constant \((k_p, k_c, k_b, \text{ and } k_0)\). (Assumed first-order kinetics of phosphorylation of the visual pigment does not necessary imply that it is a random one-step process. Sequential multiple phosphorylation may also result in approximately single-exponential quenching, as seen from detailed modeling by Hamer et al., 2005 [their Fig. 5]. This is irrelevant for our model. A second issue is that we do not know the level of phosphorylation of the 9-DM pigment relative to that of pigment regenerated with 11-cis retinal. The rod 9-DM pigment only has \(\sim 25\%\) the phosphorylation level as the native pigment [Morrison et al., 1995]. Phosphorylation levels for the 9-DM cone pigments have not yet been measured.) In turn, the active transducin \(T^*\)-PDE* complex is quenched by GTPase operating with the rate constant \(k_R\). Fractional (with respect to fully active MII*) activities of the intermediates toward transducin are:

\[
a_{M} > a_{MP} > a_{MA} > a_{mP} > a_{HR}.
\]

The kinetic scheme of Fig. 7 is described by a system of five linear differential equations whose solution provides the time course of the light-induced activity \(\text{PDE}^*(t)\). On the other hand, the \(\text{PDE}^*(t)\) curve can also be deduced from electrical photoresponses by analyzing the saturation time versus intensity function. It has been shown by Firsov et al. (2005) that

\[
p(t) = \frac{1}{I}.
\]

where \(p(t)\) is the wave of PDE activity elicited by a flash of unit intensity, \(c\) is a constant (the criterion level to which \(\text{PDE}^*(t)\) should decay to result in a fixed recovery of the circulating current), and \(I\) is light intensity (Firsov et al., 2005). In other words, fractional activity of PDE at a certain time \(t\) is inversely proportional to the intensity of the flash that results in saturation time \(t_{sat}\) = \(t\). Hence, plotting the inverse of the stimulus intensity versus time of response saturation provides the time course of PDE turn-off scaled by the factor \(c\).

Data from six cones regenerated with 11-cis retinal and four cones regenerated with 9-DM retinal processed this way are shown in Fig. 8. It is seen that up to \(\sim 2\) s, the PDE quenching with both types of the chromophore follow a common curve. At longer times, however, PDE activity decays substantially more slowly in cells containing 9-DM visual pigment (red symbols) compared with cells containing normal visual pigment (black symbols).

To see whether slower decay of Meta can explain slower turn-off of the cascade in 9-DM cones, we fitted the experimental data with the model depicted in Fig. 7. The fitting procedure was based on the fact that the solution of the system of the equations comprising the model is a sum of five exponentials, with five amplitudes and five rate constants. Rate constants are equal to corresponding rate constants of the kinetic scheme, whereas amplitudes can be expressed via \(a\)- and \(k\)-values. The amplitudes are bound to the condition that their sum is zero to ensure that \(\text{PDE}^*(t)\) starts from \(0\). Therefore, formal five-exponential approximation of the experimental data ideally yields entire information necessary to calculate all rate constants and catalytic activities of the bleaching products. Resulting formulas relating kinetic parameters of the scheme to the parameters of fitting are cumbersome and are not given here.

Of course, a straightforward nine-parametric fitting of this sort would mostly yield meaningless results. To reduce the number of free parameters, some of them should be estimated from independent experimental data. The rate of Meta decay \(k_0\) was obtained by microspectrophotometric measurements (Fig. 3, top). The
value of 0.02 s\(^{-1}\) was accepted as a crude estimate of the rate of recombination of opsin with 11-cis retinal (\(k_b\)), based on the fact that complete recovery of the circulating current after strong bleaches takes \(\sim\)3 min. We found that uncertainty of this value does not markedly affect values of other significant parameters derived from modeling. \(a_{th}\) was set to 0 because the activity of regenerated pigment is probably well below the levels detectable in our experiments. To further reduce the freedom of fitting, we estimated \(k_p\) and \(k_b\) by fitting nonsaturated flash responses with a complete model of phototransduction by Kazmin et al. (2004). The model is similar to that by Hamer et al. (2005) and includes all firmly established mechanisms of activation, quenching, and Ca\(^{2+}\) feedback regulation of the phototransduction cascade. It appeared that nonsaturated flash responses in 11-cis- and 9-DM–regenerated cones can be fitted with the same values of \(k_p\) and \(k_b\) after just a small adjustment of the kinetics of Ca\(^{2+}\) feedback (Fig. 7, inset). The values shown in Fig. 7 were therefore fixed in further analysis.

Thus, only four parameters, namely the rate of arrestin binding to phosphorylated metaproducts (\(k_i\)), fractional activity of phosphorylated but arrestin-unbound metaproducts (\(a_{MII}\)), fractional activity of metaproducts phosphorylated and capped with arrestin (\(a_{MP}\)), and fractional activity of “naked” opsin (\(a_{Ops}\)), have to be determined from fitting. An additional consideration during fitting is that the values of PDE\(^*\)(\(t\)) encompass a range of \(\sim\)5 log units. Therefore, points at the low-amplitude slow tail should be assigned higher weights to be properly treated with the fitting algorithm. We assigned to each experimental point the weight inverse to the point’s value.

Weighted least-square fits of experimental data with a sum of five exponentials made under constraints discussed above are shown in Fig. 8 by smooth bold lines. The parameters of the kinetic scheme derived from fitting are provided in Table I. The model fits are consistent with the assumption that fivefold slower Meta decay in 9-DM–regenerated cones is the main cause of the differences observed in the recovery from saturation after exposure to bright light. In addition, the model suggests that the residual activity of partially quenched 9-DM Meta intermediates is several-fold higher than in methylated 11-cis pigment (Table I).

### DISCUSSION

The experiments performed here were designed to test two related hypotheses. The first is that removal of the 9-methyl group from the retinal moiety that is covalently attached to opsin in red cones results in a visual pigment whose activated state (Meta II) decays unusually slowly compared with that of native visual pigment. The second related hypothesis is that this slowed Meta decay results in sluggish recovery of flash responses elicited by bright light. To test these hypotheses, we correlated microspectrophotometric measurements of post-bleach visual pigment photolysis and microfluorometric measurements of ROL formation with electrophysiological measurements of flash response kinetics in intact red cone photoreceptor cells isolated from salamander retina. Measurements were made under dark-adapted conditions when cells contained a full complement of native visual pigment, or when the visual pigment had been fully bleached and regenerated to contain either 11-cis retinal or 9-DM retinal.

We summarize our results as follows. First, visual pigment was successfully regenerated in the outer segments of bleached red cones of *Ambystoma mexicanum* to contain 9-DM retinal. The absorbance spectrum of the pigment was significantly blue-shifted (\(\lambda_{\text{max}} = 520\) nm) compared with visual pigment either containing the native 11-cis retinal/dehydroretinal mixture (\(\lambda_{\text{max}} = 592\) nm) or 11-cis retinal (\(\lambda_{\text{max}} = 567\) nm) (Fig. 1). Second, the time course of Meta decay measured in situ in red cones containing 9-DM retinal (\(T_{1/2} = 14\) s) was approximately five times slower than that regenerated with 11-cis retinal (\(T_{1/2} = 3\) s), and four times slower than in native pigment (\(T_{1/2} = 3.5\) s) (Fig. 3). Almost no Meta III was observed in red cones, consistent with earlier reports (Ala-Laurila et al., 2006; Golobokova and Govardovskii, 2006). Third, microspectrophotometric and microfluorometric measurements revealed that the initial rate of retinol production was several-fold slower

### Table I

| \(k_{f}\) \(s^{-1}\) | \(a_{II}^\ast\) | \(k_{i}\) \(s^{-1}\) | \(a_{MII}\) | \(k_{II}\) \(s^{-1}\) | \(a_{MP}\) | \(k_{III}\) \(s^{-1}\) | \(a_{Ops}\) | \(k_{MP}\) \(s^{-1}\) | \(a_{MP}\) |
|----------------|-------------|-----------------|---------|-----------------|--------|-----------------|--------|----------------|--------|
| 11-cis         | 5           | 1               | 1       | 0.022           | 0.21   | 2.7 \times 10^{-5}| 2 \times 10^{-6}| 4 \times 10^{-6}| 3.5    | nd             |
| 9-DM           | 5           | 1               | 1.05    | 0.004           | 0.042  | 1.1 \times 10^{-4}| 2 \times 10^{-6}| 4 \times 10^{-6}| 3.5    | nd             |

\(k_{f}\) and \(k_{b}\) are determined from fitting the full model of phototransduction to nonsaturated flash responses; \(k_{b}\) values are obtained by single-exponential fitting to Meta decay in Fig. 3; \(k_{II}, a_{MII}, a_{MP}\), and \(a_{Ops}\) from fitting the kinetic scheme above to \(T_{II}\) versus bleach data, using previously found \(k_{b}, k_{p}\), and \(k_{II}\). These results, together with Fig. 7, suggest that, in addition to slower decay, the residual activity of 9-demethyl metaproducts (\(a_{MP}\) and \(a_{MII}\)) is substantially higher than that of 11-cis pigment. nd, not determined.
in cone outer segments containing pigment lacking the 9-methyl group of retinal (Figs. 4 and 5).

We have compared these microspectrophotometric and microfluorometric results to electrophysiological results obtained under identical conditions of pigment bleaching and regeneration. Consistent with earlier results obtained on red cones from *Ambystoma tigrinum* (Estevez et al., 2006), red cones regenerated with 9-DM retinal had photoresponses to dim flashes that appeared to be identical in time course to those of red cones regenerated with 11-cis retinal. However, at brighter flashes that bleached from ~0.1 to 45% of the visual pigment, red cones lacking the 9-methyl group had progressively slower response termination. For example, after a 33% bleach, red cones containing visual pigment lacking the 9-methyl group exhibited a flash response plateau that was approximately seven times longer than cones regenerated with 11-cis retinal; further restoration of dark current was also greatly retarded (Fig. 6). Slower recovery from saturation paralleled slower decay of Meta in 9-DM cones.

The role of decay of metaproducts in regulation of quenching of responses in rods and cones

The importance of Meta decay in regulating the time course of dark adaptation of rods is well established. Exposure of rods to bright bleaching light leads to the persistence of long-lived photolysis intermediates that are able to activate, with low gain, the phototransduction cascade for a time beyond the end of light stimulation. These persistent effects have been observed in physiological (Donner and Hemilä, 1975; Lamb, 1980; Leibrock and Lamb, 1997; Leibrock et al., 1998; Kolesnikov et al., 2003) as well as biochemical (Okada et al., 1989; Jäger et al., 1996) experiments in rods. They are understood to provide at least a partial basis for the slowness of photoresponse recovery that occurs after exposure to bright light.

Alternative mechanisms have been suggested to explain the slowness of the cascade quenching at high bleach intensities. Lyubarsky and Pugh (1996) proposed that in this condition the cascade turn-off is retarded ($k_e$ decreased) due to local saturation of the PDE pool that serves as a GTP-ase activating factor for T* quenching. Alternatively, biochemical experiments on carp cone membranes provide evidence that the phosphorylation power (hence rhodopsin turn-off) of cones may be limited at high light intensities (Tachibanački et al., 2005). However, these mechanisms do not account for parallel effect of 9-DM chromophore on the rate of Meta decay and recovery of saturation observed by us in salamander red cones.

An example illustrating this critical relationship between Meta II decay and photoresponse recovery was shown in a recent study by Imai et al. (2007). They accelerated Meta II decay 10-fold and Meta III twofold in mammalian mouse rods by replacing rhodopsin with its E122Q mutant, and observed no effect on the dim-flash response kinetics unless arrestin was concurrently knocked out (indicating that phosphorylation and subsequent binding of arrestin is fast enough to precede Meta II decay). However, after exposing to brighter flashes that bleached 20% of the pigment, flash responses elicited in mutant rods exhibited a faster rate of recovery. This included both reduced time in saturation and steeper return to the dark current level. Thus, these results on rod photoreceptors support the idea that different mechanisms control the response recovery depending on the amount of light present, and that it is Meta decay that controls the photoresponse recovery after bright illumination. The sluggish photoresponse recovery of red cones lacking the 9-methyl group of retinal observed in response to bright but not dim light suggests that this is also true for cones.

This idea is also consistent with conclusions from electrophysiological studies by Firsov and coworkers on goldfish cones (Firsov et al., 2007) and frog rods (Firsov et al., 2005). According to their model, at low intensities the cells recover to the prestimulus state by using the fast mechanisms that quench dim flash responses (phosphorylation, arrestin binding, and Ca*2* feedback). Apparently, these reactions in cones have no dependency on the presence of the 9-methyl group. However, after flashes that bleach larger fractions of the pigment (>3% in goldfish cones and >0.1% in rods), these fast steps have run their normal courses while the cell’s response still remains in saturation, and thus, residual activity of decaying Meta and subsequent pigment regeneration control the time course of photoresponse recovery.

Mathematical modeling based on the Firsov et al. (2005) kinetic scheme of cascade quenching further supports this idea. Application of this model to our electrophysiological data (Fig. 6) shows that the difference in the time courses of PDE turn-off in cones containing normal and 9-DM chromophores can be predicted based on the kinetics of photolysis of corresponding visual pigments obtained microspectrophotometrically (Fig. 8). This idea is also consistent with the results from our previous work on *Ambystoma tigrinum* in which we demonstrated that slowness of response recovery in 9-DM cones was due to persistent activations of the transduction cascade (Estevez et al., 2006).

Thus, targeted modification of the visual pigment chromophore results in concomitant changes in the photolysis kinetics and the speed of photoresponse recovery. Our results indicate that the 9-methyl group of retinal is required for steric chromophore–opsin interactions that favor both the rapid decay of Meta and the rapid termination of the photoresponse after exposure to bright but not dim light in red cones. These findings provide a partial mechanistic explanation for the rapid time course of dark adaptation observed in red cone photoreceptors.
Steric interactions of opsin and the 9-methyl group

Previous biochemical studies of chicken rod and cone visual pigments have also investigated the opsin–chromophore interactions that influence the lifetime of Meta II in rod and cone visual pigments (Imai et al., 1997, 2007; Kuwayama et al., 2002, 2005). For instance, Kuwayama and coworkers replaced the proline at position 189 in chicken green cone pigment with the corresponding rhodopsin residue (isoleucine) (Kuwayama et al., 2002). They observed a 27-fold slower Meta II decay rate in these mutated cone pigments. Similarly, they demonstrated an acceleration of the Meta II decay rate by the reverse, that is, by replacing the isoleucine residue as position 189 in rhodopsin with the corresponding residue (proline) in chicken green cone pigment (Kuwayama et al., 2002). In other studies, an accelerated Meta II decay rate in rhodopsin was induced by mutation at residue 122 with the corresponding residue in cone pigments (Imai et al., 1997). This group concluded that the 50-fold difference in Meta II decay rate and the 700-fold difference in Meta III decay rate between chicken rhodopsin and chicken green cone pigment is regulated by the residues at positions 122 and 189 (Kuwayama et al., 2002, 2005).

Could these mutations at positions 122 and 189 that modulate Meta II decay rate in chicken green cone pigment also have their effects through the 9-methyl group of retinal? No definitive experiments have been performed to investigate this possibility. However, Kuwayama and coworkers note that the CG1 carbon atom of I189 in bovine rhodopsin is 3.8 Å from the 9-methyl group of retinal, whereas the peptide amine of I189 is 4.6 Å from the 9-methyl group (Kuwayama et al., 2002). They speculate that I189 may have steric interactions with the 9-methyl group in chicken rhodopsin that may regulate Meta II decay. However, the role of the 189 residue (proline) in salamander red cone pigment is not known, nor is it known if this residue interacts with the 9-methyl group of retinal in salamander rod or cone pigment. The chicken green cone pigment has higher homology in amino acid sequence with vertebrate rhodopsins (70%) than with other cone visual pigments (40%) (Wang et al., 1992). Based on these differences, the speculations of chromophore–opsin interactions between the 9-methyl group of retinal and residue 189 in chicken green cone pigment cannot be directly translated to salamander red cone pigment, but they provide impetus for further investigation.

Han et al. (1997) in a separate study have indicated that the 9-methyl group of retinal interacts with Gly121 of rhodopsin. In the salamander, the Gly121 in rhodopsin is conserved in red cone pigment. However, this residue may be in a different structural location in red cone pigment, and there is no evidence that it interacts with the 9-methyl group of retinal. How the 9-methyl group of retinal may interact with various amino acid residues in salamander red cone pigment is not known. The data presented here do not address this point. However, it is clear that the absence of this methyl group on retinal causes significant changes in the biochemical and physiological properties of the red cone opsins that strongly affect Meta II decay and flash response recovery kinetics under bright light exposure.

This work was supported by National Institutes of Health grants EY-001157 (to M.C. Cornwall) and EY-004939 (R.K. Crouch); unrestricted grant to MUSC from Research to Prevent Blindness, Inc., New York, NY; The Russian Foundation for Basic Research grant 06-04-48914 and the Biological Department of The Russian Academy of Sciences (to V.I. Govardovskii); and a grant from the Academy of Finland (grant 123231 to P. Ala-Laurila). R.K. Crouch is a Research to Prevent Blindness Senior Scientific Investigator.

Edward N. Pugh Jr. served as editor.

Submitted: 20 March 2009
Accepted: 13 July 2009

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