

Phytochromes and Cryptochromes in the Entrainment of the *Arabidopsis* Circadian Clock

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Circadian clocks are synchronized by environmental cues such as light. Photoreceptor-deficient *Arabidopsis thaliana* mutants were used to measure the effect of light fluence rate on circadian period in plants. Phytochrome B is the primary high-intensity red light photoreceptor for circadian control, and phytochrome A acts under low-intensity red light. Cryptochrome 1 and phytochrome A both act to transmit low-fluence blue light to the clock. Cryptochrome 1 mediates high-intensity blue light signals for period length control. The presence of cryptochromes in both plants and animals suggests that circadian input pathways have been conserved throughout evolution.

In many organisms, an endogenous, self-sustaining oscillator maintains rhythms of about 24 hours to control a wide variety of biological processes (1, 2). To function as a circadian clock, the endogenous oscillator must be entrained to the daily light and temperature cycles of the external environment. However, only a few components are known that connect the endogenous oscillator to the environment (2, 3), and no unequivocal connection between a specific photoreceptor and the entrainment of the circadian clock by light has been made in any organism to date (4).

Increasing light intensity often tends to lengthen period in nocturnal organisms and to shorten period in diurnal organisms, including plants (5). At least two types of unidentified photoreceptor systems mediate light input in the algae *Gonyaulax* and *Chlamydomonas* (6). To address this question in higher plants, we crossed *Arabidopsis* red light (RL) and blue light (BL) photoreceptor mutants with transgenic plants containing the firefly luciferase gene (*luc*) under the control of the clock-responsive *cab2* promoter (7).

Plant photoreceptors absorb primarily in RL [600 to 700 nm wavelength; the phytochromes (phy)] and BL [400 to 500 nm; the cryptochromes (cry) and *NPH1*] (8). Limiting RL input, either by reducing fluence rate or photoreceptor abundance [(9); supplementary figure 1, available at www.sciencemag.org/feature/data/985395.sh1] slows the pace of the oscillator, suggesting phytochromes are the likely mediators of RL signaling to the clock. To identify which of the five phyto-

chrome species (*PHYA-E*) (10) control RL input to the clock in *Arabidopsis*, we tested null mutant alleles of two phytochromes, *phyA* and *phyB*, for their effects on the free-running rhythm in constant RL.

A *phyA* deficiency affected period length of the clock in RL only in dim red light ($<1.0 \mu\text{mol m}^{-2} \text{s}^{-1}$) (Fig. 1A). These results support previous research showing that *phyA* mediates developmental responses to long-term, high-irradiance far-red (FR)-enriched light or to pulses of red and FR light (11). One exception is in the coupling of *phyA* to the photoperiodic control of flowering. Both in pea and *Arabidopsis*, white-light-grown *phyA* mutants have an altered flowering time,

suggesting an interaction between *phyA* and circadian timing (12). Our results indicate that this is not through a direct effect of *phyA*-mediated input to the clock, but more likely the result of clock-gated control of an independent *phyA* signaling pathway.

In contrast, the *phyB*-deficient mutant mediates high-fluence RL input to the circadian clock, showing a 1.5- to 2-hour lengthening of period, relative to the wild type, at fluence rates $>5.0 \mu\text{mol m}^{-2} \text{s}^{-1}$ (Fig. 1B). Overexpression (15-fold) of *phyB* (13) shortened period length by 1 to 3 hours, relative to the wild type, dependent on the fluence rate [(14); supplementary figure 2, available at www.sciencemag.org/feature/data/985395.sh1]. It is likely that other phytochromes (*phy C, D, or E*, or a combination of these) are also involved, because the free-running period in the *phyB* mutant was still shorter at the higher fluence range ($>5 \mu\text{mol m}^{-2} \text{s}^{-1}$) than in the phytochrome-deficient *hy1-6* line (14, 9), and it continues to shorten up to $200 \mu\text{mol m}^{-2} \text{s}^{-1}$ (Fig. 1B). Tests using single and multiple *phy* mutant combinations will be necessary to fully describe this signaling network (15).

Because phytochrome also absorbs BL, we examined the BL fluence rate-response curve of the *phyA* and *phyB* mutants. At fluence rates below 3 to $5 \mu\text{mol m}^{-2} \text{s}^{-1}$, the absence of *phyA* lengthened the free-running period by up to 3 hours, relative to the wild type (Fig. 1C), demonstrating that *phyA* is required for period control in low-fluence BL, and an additional photoreceptor is required to mediate the full range of response to BL. A deficiency in *phyB* had no period effect in BL (Fig. 1D).

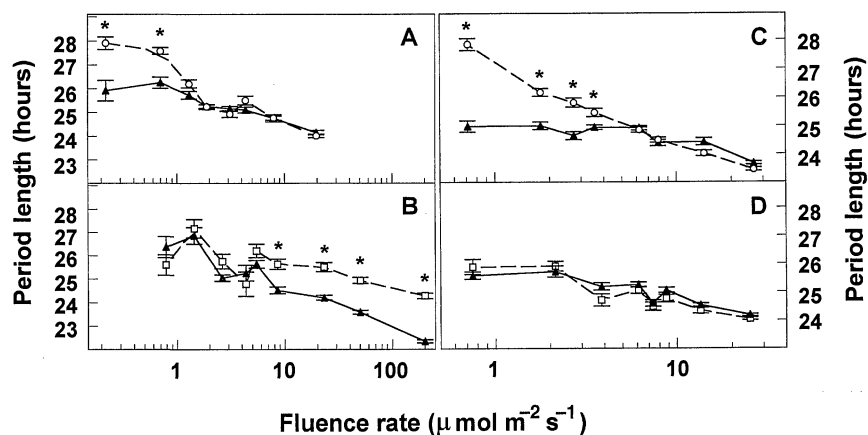


Fig. 1. Effect of RL fluence rate on free-running period length of *cab2::luciferase* (*cab2::luc*) expression in (A) *phyA-201* and (B) *phyB-1* mutant seedlings. Effect of BL fluence rate on free-running period length of *cab2::luc* expression in (C) *phyA-201* and (D) *phyB-1* mutant seedlings (23). ○, *phyA-201*; □, *phyB-1*; ▲, Laer wild type. Plants were germinated and grown in cycles of 12 hours white fluorescent light ($50 \text{ to } 60 \mu\text{mol m}^{-2} \text{s}^{-1}$) and 12 hours of dark for 6 days, then transferred for >110 hours to continuous red (600 to 700 nm) or blue (400 to 500 nm) light at the fluence rates indicated. Mean period length estimates were obtained by fitting a modified cosine wave function to the time series of each seedling (9, 24). Luminescence measurements of *cab2::luc* expression were described previously (9). Error bars are \pm SEM ($n = 7$ to 18). Asterisk, $P < 0.01$ (Student's two-tail heteroscedastic *t* test). Similar results were obtained in 2 to 11 independent experiments.

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To further establish the role of phyA in the BL entrainment pathway, a second type of assay was performed. Seedlings grown in white-light/dark (LD) cycles (12 hours/12 hours; $T = 24$ hours) were shifted to one of eight different BL/dark cycles (10 hours/10 hours; $T = 20$ hours), spanning a >30-fold range of BL fluence rates. We reasoned that at high fluence rates, the *phyA* mutant would entrain to the 20-hour T cycle, but at the lowest intensities, BL input to the clock would be impaired and the plants would show a 24-hour rhythm, in phase with the original white-light entrainment protocol.

At the highest fluence rate ($27 \mu\text{mol m}^{-2} \text{s}^{-1}$), both strains entrained rapidly (Fig. 2A). The wild type showed similar results at a much lower fluence rate ($2 \mu\text{mol m}^{-2} \text{s}^{-1}$), whereas the *phyA* mutant required at least three cycles to become stably entrained (Fig. 2B). In the mutant, peak luminescence after hour 20 came 4 hours later than the wild type (Fig. 2B; arrowhead), and there was no response to the lights coming on at hour 40. Poor entrainment also occurred at lower fluence rates ($0.8 \mu\text{mol m}^{-2} \text{s}^{-1}$), but at greater than $\sim 4 \mu\text{mol m}^{-2} \text{s}^{-1}$, the *phyA* mutant entrained as rapidly as the wild type [(14); supplementary figure 3, available at www.sciencemag.org/feature/data/985395.sh1]. These results match the range of fluence

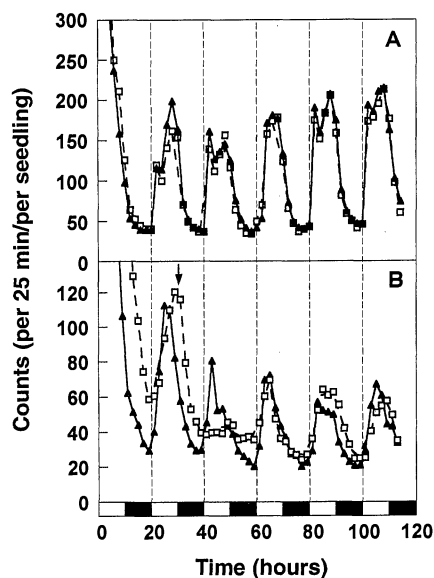


Fig. 2. Re-entrainment of *cab2::luc* expression rhythm to light/dark (10 hour/10 hour) cycles in wild-type and *phyA-201* mutant seedlings. Wild-type (Laer) (▲) and *phyA-201* mutant (□) seedlings were grown in cycles of 12 hours white fluorescent light (50 to $60 \mu\text{mol m}^{-2} \text{s}^{-1}$) and 12 hours dark for 6 days, then transferred (hour 0) to BL/dark (10 hour/10 hour) cycles at fluence rates of (A) $27 \mu\text{mol m}^{-2} \text{s}^{-1}$ or (B) $2 \mu\text{mol m}^{-2} \text{s}^{-1}$. For each trace, $n = 18$; error bars are omitted for clarity. Open boxes on the time axis indicate BL; filled boxes, darkness. Representative of two independent experiments.

rates over which a *phyA* deficiency causes a lengthening of period in the fluence rate-response curve (Fig. 1C). The similarity of the fluence rate range over which *phyA* affects entrainment under these two conditions supports the notion that light signaling to the clock in continuous illumination and light/dark cycles share some similar properties.

The cryptochromes (*cry1* and *cry2*) share similarity with photolyases and are present in both plants and animals (16, 17). Both *Arabidopsis* cryptochromes have been linked to blue-light-dependent processes in plant development, including flowering time (16). Overexpression of *cry1* shortened period by 1 to 1.5 hours in both high-fluence white ($50 \mu\text{mol m}^{-2} \text{s}^{-1}$) and blue ($18 \mu\text{mol m}^{-2} \text{s}^{-1}$) light (14). Conversely, loss of *cry1* resulted in period lengthening over two different ranges of BL fluence rates ($>10 \mu\text{mol m}^{-2} \text{s}^{-1}$; $<3 \mu\text{mol m}^{-2} \text{s}^{-1}$) (Fig. 3A). Between these two intensity ranges, the loss of *cry1* was inconsequential, suggesting that other BL receptors mediate signaling in this region. The absence of *cry2* caused minimal period lengthening in the high fluence rate range ($>10 \mu\text{mol m}^{-2} \text{s}^{-1}$) (Fig. 3B). Over a narrow intensity range (Fig. 3B; 3 to $4 \mu\text{mol m}^{-2} \text{s}^{-1}$), loss of *cry2* showed a slight but reproducible period shortening.

Cry1 and *cry2* may act additively and redundantly, and only the double mutant will reveal the combined effect of their overlapping actions. As well, at fluence rates <3 to $5 \mu\text{mol m}^{-2} \text{s}^{-1}$, *cry1* and *phyA* deficiencies each cause similar period lengthening, suggesting that the two together are required for normal BL signaling. This is supported by recent yeast two-hybrid interaction studies

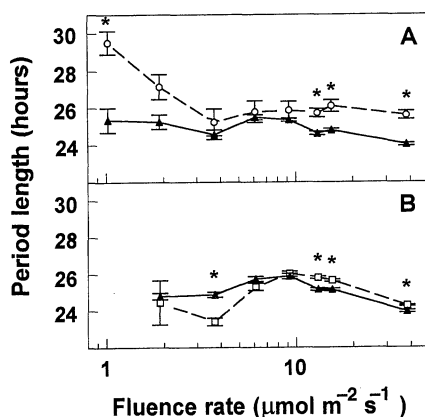


Fig. 3. Effect of BL fluence rate on free-running period length of *cab2::luc* expression in (A) *cry1* and (B) *cry2* mutant seedlings. ○, *cry1*; □, *cry2*; ▲, Laer or Col-1 wild type. Conditions of entrainment, luminescence measurements and data analysis are as in Fig. 1. Error bars are \pm SEM ($n = 5$ to 19). *, $P < 0.01$ (Student's two-tail heteroscedastic t test). Representative of four independent experiments (25).

showing a direct interaction between *cry1* and *phyA* polypeptides (18).

When grown in white light, *cry2-1* is day-length-insensitive and late-flowering (19). To determine whether this effect on photoperiodic timing correlates with a change in period length, we determined the free-running period of all individuals of an F_2 population segregating for *cry2-1* in white light. Although the population segregated 1:3 ($P > 0.10$) for long-period:wild-type-period, the 16 *cry2-1* homozygotes were distributed as 1/4 of the individuals ($P > 0.50$) within each of the two period classes, as expected for single gene segregation independent of period length (Fig. 4). Loss of *cry2*, therefore, does not correlate with a change in period length when grown in white light, and the slight period lengthening in BL (Fig. 3B) has no effect on flowering time (19). The effects of *cry2* on photoperiodic timing most likely arise through a circadian-clock gating of *cry2*-mediated signaling, and not through a direct signaling of the *cry2* to the oscillator.

Our evidence for multi-photoreceptor-mediated control of the pace of the circadian clock is in concert with recent physiological and molecular genetic studies of phototransduction in plants. Studies using photoreceptor-specific null mutants in *Arabidopsis* have indicated that the nature of the interactions among *phyA*, *phyB*, and *cry1* signaling pathways is strongly dependent on light quality and fluence rate (20). Similar conclusions can be derived from recent work showing direct molecular interactions between *phyA* and *cry1* (18).

For sedentary, light-dependent organisms such as plants, the development of compensatory mechanisms against the potentially disruptive effects of wide changes in the light environment on physiology and development may be essential. By recruiting a diversity of photoreceptors that can cover a wide range of fluence rates and spectral qualities, the plant can ensure that the pace of the circadian oscillator remains little affected. This in turn

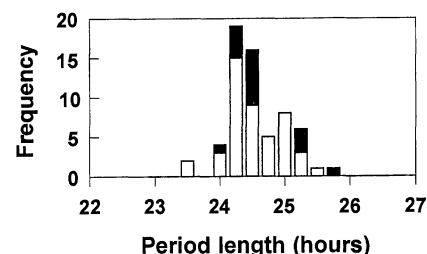


Fig. 4. Period length distribution in a seedling population segregating for *cry2-1*. Entrainment protocol, free-run conditions (50 to $60 \mu\text{mol m}^{-2} \text{s}^{-1}$ white light), luminescence measurements, and period analysis are as in Fig. 1. Period bins are labeled with the upper bound. Dark fill shows *cry2-1* homozygotes (25). Representative of two independent experiments.

ensures that circadian-controlled processes maintain the appropriate phase relationship to environmental cues. Photoreceptor diversity and redundancy, therefore, appear to be the key features in the photocontrol of the circadian clock in higher plants.

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23. Individual *cab2:luc*-expressing lines homozygous for *phyA-201* or *phyB-1* were selected from a cross of *phyA-201* (21) with 2CAC (C24 ecotype) (9), selfed, and harvested individually (*phyA* F₃ pools), or selected and selfed from a second backcross of 2CAC to *phyB-1* (21) (*phyB* F₃ pools). 2CAC/Laer(7x) F₂ seedlings [from the seventh backcross of 2CAC with ecotype *Landsberg erecta* (Laer)] were used as the wild-type control. Homozygous *phyA-201* and *phyB-1* mutants were selected on the basis of hypocotyl length in far-red and red light, respectively. Light sources in this work were described previously (9). Repeat trials used two to three independent F₃ pools.
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selected by morphological phenotype from a cross of *cry1* (*hy4-2.23N*) (22) with 2CAC and selfed. Eight F₃ families were pooled to generate one population; a second population was derived from a single individual. 2CAC/Laer(7x) F₂ seedlings were used as the wild-type control. F₂ seedlings homozygous for the *cry2* mutation were selected by polymerase chain reaction from a cross of *cry2-1* (Col-4 ecotype) (19) with 2CAC/Col-1(6x). The upstream (5'-CAGCTGCT-CACGAAGGATCT-3') and downstream primer (5'-GCAGTTATTGGCATCAACCG-3') amplified a fragment of 466 base pairs from the wild-type *CRY2* gene in a region deleted in the mutant *cry2-1* allele. Among the trials, two independently isolated F₃

pools were used. 2CAC/Col-1(6x) was generated by six introgressions of 2CAC into the Columbia-1 ecotype.

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Role of Mouse Cryptochrome Blue-Light Photoreceptor in Circadian Photoresponses

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Cryptochromes are photoactive pigments in the eye that have been proposed to function as circadian photopigments. Mice lacking the cryptochrome 2 blue-light photoreceptor gene (*mCry2*) were tested for circadian clock-related functions. The mutant mice had a lower sensitivity to acute light induction of *mPer1* in the suprachiasmatic nucleus (SCN) but exhibited normal circadian oscillations of *mPer1* and *mCry1* messenger RNA in the SCN. Behaviorally, the mutants had an intrinsic circadian period about 1 hour longer than normal and exhibited high-amplitude phase shifts in response to light pulses administered at circadian time 17. These data are consistent with the hypothesis that CRY2 protein modulates circadian responses in mice and suggest that cryptochromes have a role in circadian photoreception in mammals.

Circadian rhythms are oscillations in the biochemical, physiological, and behavioral functions of organisms with a ~24-hour periodicity (1). Circadian rhythms are synchronized with light-dark cycles, but the molecular basis of this "photoentrainment" is not known (2). Indeed, there is no consensus on the nature of the circadian photoreceptor. Three classes of pigments have been considered as candidates: opsin/retinal-based photopigments (3), tetrapyrrole-based heme pigments (4), and pterin/flavin-containing cryptochrome blue-light photoreceptors (5, 6). Cryptochromes were first identified in plants as structural ho-

mologs of the DNA repair enzyme DNA photolyase (7), but they lack DNA repair activity (8) and are involved in mediating growth (9), flowering time (10), and phototropism (11) in response to blue light. Recently, two human and mouse homologs of the plant cryptochromes were discovered (5, 12). Cryptochromes 1 and 2 (CRY1 and CRY2, respectively) lack DNA repair activity (5) and are expressed in the mouse retina (6), and *mCry1* exhibits circadian oscillations of expression in the SCN (6) wherein the central pacemaker of the body resides. These observations led to the proposal that cryptochromes were likely to be photopigments for circadian photoentrainment (6).

To test this hypothesis, we created a mouse strain that lacks the predominant form of cryptochrome found in the mouse retina, CRY2, and analyzed its circadian behavior using biochemical and behavioral tests. We generated *Cry2*^{+/-} heterozygous mutant mice by established methods (13) using the targeting construct shown in Fig. 1A. Interbreeding of heterozygotes yielded progeny of wild type:*Cry2*^{+/-}:*Cry2*^{-/-} at a ratio of 1:2:1 (Fig. 1B). The mutant mice were phys-

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