Behavioral/Systems/Cognitive

The Molecular Gatekeeper Dexras1 Sculpts the Photic Responsiveness of the Mammalian Circadian Clock

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The mammalian master clock, located in the suprachiasmatic nucleus (SCN), is exquisitely sensitive to photic timing cues, but the key molecular events that sculpt both the phasing and magnitude of responsiveness are not understood. Here, we show that the Ras-like G-protein Dexras1 is a critical factor in these processes. *Dexras1*-deficient mice (*dexras1*^{-/-}) exhibit a restructured nighttime phase response curve and a loss of gating to photic resetting during the day. Dexras1 affects the photic sensitivity by repressing or activating time-of-day-specific signaling pathways that regulate extracellular signal-regulated kinase (ERK)/mitogen-activated protein kinase (MAPK). During the late night, Dexras1 limits the capacity of pituitary adenylate cyclase (PAC) activating peptide (PACAP)/PAC1 to affect ERK/MAPK, and in the early night, light-induced phase delays, which are mediated predominantly by NMDA receptors, are reduced as reported previously. Daytime photic phase advances are mediated by a novel signaling pathway that does not affect the SCN core but rather stimulates ERK/MAPK in the SCN shell and triggers downregulation of clock protein expression.

Key words: Dexras1; MAPK; circadian; photic; entrainment; PACAP; suprachiasmatic

Introduction

Diverse life forms have evolved circadian timing mechanisms ensuring that biological processes are regulated at near 24 h cycles (Dunlap, 1999). The mammalian master pacemaker is located in the suprachiasmatic nucleus (SCN) (Ralph et al., 1990). The SCN receives external timing cues and provides signals that synchronize oscillations in brain and peripheral tissues. Timing cues (e.g., light) reset and entrain the SCN clock, ensuring that circadian rhythms of behavior and physiology maintain an appropriate temporal relationship with respect to each other and the environment.

A hallmark of circadian timing is that the clock responds to resetting signals in a phase-restricted manner. In nocturnal rodents, light is an effective entrainment cue at dusk and dawn but not during the midday (Decoursey, 1964; Pittendrigh and Daan, 1976). Conversely, behavioral arousal, a nonphotic stimulus, phase-shifts the clock only during the day (Bobrzynska and Mrosovsky, 1998). Temporal responses of the clock to entraining inputs are represented as a phase response curve (PRC) (Daan and Pittendrigh, 1976). The shape and amplitude of a PRC are established by the state of the molecular clock and the signal-

transducing capacity of the input pathways. Phase-restricted responsiveness and clock-controlled gating of clock inputs remain central tenets within the field.

The p42/p44 mitogen-activated protein kinase (MAPK) pathway has been implicated in circadian entrainment and represents a potential substrate for clock-controlled gating of clock inputs. This pathway is light-inducible in the SCN in a phase-restricted manner (Obrietan et al., 1998), couples light to transcriptional activation (Dziema et al., 2003), and mediates photic phase shifts (Butcher et al., 2002; Coogan and Piggins, 2003). MAPK activation is triggered by glutamate and pituitary adenylate cyclase activating peptide (PACAP), which are cotransmitters of the retinohypothalamic tract (RHT) and convey photic information to the SCN (Obrietan et al., 1998; Butcher et al., 2005).

Although it is recognized that the SCN controls its own sensitivity and therefore shapes the PRCs, the key signaling events that impart this sensitivity are unclear. Potentially, rhythmic availability of a key signaling molecule within an input pathway or, alternatively, a modulator of such a pathway confers both phase specificity and sensitivity to the stimulus. One such candidate is Dexras1, a Ras-like G-protein that exhibits clock-dependent rhythms of expression in the SCN (Cismowski et al., 1999; Panda et al., 2002; Ueda et al., 2002; Takahashi et al., 2003).

Biochemically, Dexras1 functions as a guanine nucleotide exchange factor (GEF) for $G_{i/o}$ α subunits (Cismowski et al., 2000). Dexras1 has been implicated as a receptor-independent activator of $G_{i/o}$ -protein signaling (Cismowski et al., 1999) as well as a context-dependent modulator of the MAPK cascade and other signal transduction pathways, including adenylyl cyclases (ACs)

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and NMDA receptor-nitric oxide (NO)-mediated signaling (Fang et al., 2000; Graham et al., 2004; Nguyen and Watts, 2005).

Here, we provide data revealing that Dexras1 plays a critical role in shaping the photic PRC and the signaling events through which it regulates clock entrainment. These findings provide new insight into the complex set of signaling events that underlie clock entrainment and PRC shape.

Materials and Methods

Generation of Dexras1(A178V) transgenic mice. The point mutation alanine 178 → valine was introduced into the I.M.A.G.E. clone encoding the cDNA of murine Dexras1 (American Type Culture Collection, Manassas, VA) using the QuikChange Site-Directed Mutagenesis kit (Stratagene, La Jolla, CA). Full-length Dexras1(A178V) was subcloned into the expression vector pIRES2-eGFP (Clontech, Mountain View, CA), and the Dexras1(A178V)-IRES2-eGFP cassette was subsequently moved into the tetracycline-inducible expression vector pTRE-TIGHT (Clontech). Using the construct TRE-TIGHT-Dexras1(A178V)-IRES-eGFP, three independent transgenic mouse lines were established, which showed similar patterns and levels of enhanced green fluorescent protein (eGFP) expression in founder screens. Data are presented from double transgenic mice generated from intercrosses between line #3901 and calcium calmodulin-dependent kinase II α (CaMKII α)-tetracycline transactivator (tTA) mice (Mayford et al., 1996). Experiments were performed on 8- to 16-week-old mice that had been maintained under standard, doxycycline-free conditions at the animal facility of The Ohio State University, in accordance with institutional guidelines.

Behavioral analyses. Mice were singly housed in cages equipped with a running wheel in light-tight, ventilated boxes with controlled lighting. Wheel-running activity was monitored and analyzed with VitalView and Actiview (MiniMitter, Sunriver, OR). Mice were entrained to a 12 h light/dark (LD) cycle (400 lux) for at least 1 week before any experimental manipulation. For the photic phase response curve, mice received a 15 min light pulse of 40 lux (at one of the following Zeitgeber (ZT) times: 14, 17, 20, or 23) and subsequently released into constant darkness (DD) for 7-10 d. For light pulses administered at ZT 2, 5, 8, and 11, on the day of the experiment, mice were put into DD at ZT 0 and received a 15 min light pulse of 40 lux at the appropriate ZT time. The animals remained in DD for at least 1 week after the pulse. For all other behavioral experiments, mice were entrained to a 12 h LD cycle (400 lux) and received a 15 min light pulse at a given light intensity (40 or 400 lux) exactly as described above or after 8-12 d in DD. Mice were maintained at the animal facility of The Ohio State University in accordance with institutional guidelines. All animal handling and experimental procedures were approved by the Animal Welfare Committee of The Ohio State University.

Cannulation and infusion. Mice were cannulated in the third or lateral ventricle as described previously (Butcher et al., 2002) and allowed to recover for a minimum of 10 d after cannulation. For the infusion, mice were restrained by hand under red light, and the infusate was delivered at a rate of 0.4 µl/min. Full-length PACAP was diluted in physiological saline to a final concentration of 40 μ M and injected at a volume of 3 μ l. PACAP 6–38 (500 μ M in physiological saline) was injected at a volume of 3 µl, 30 min before photic stimulation or PACAP infusion. 1,4-Diamino-2,3-dicyano-1,4-bis[2-amino-phenylthio]butadiene (U0126) was diluted to 10 mm in DMSO, and 3 µl was injected 30 min before light treatment. Control animals were infused with an equivalent volume of vehicle (saline or DMSO). We have consistently used a 3 μ l volume for lateral and third ventricle infusions (Butcher et al., 2002, 2005; Dziema et al., 2003) and have not detected any long-term effects on SCN-regulated physiology, signaling, or behavior. Likewise, SCN morphology is not affected. Thus, we are confident that the volume used and the slow rate of infusion do not cause complicating side effects. In addition, it should be noted that the volume change resulting from the infusion (+3 μ l) is partially offset by the loss of CSF (typically several microliters) that occurs after withdrawal of the cannula plug.

Cell culture and transfection. Embryonic rat cortical neuronal cultures were prepared from embryonic day 19–20 Sprague Dawley rat pups exactly as described previously (Lee et al., 2005). After 8–10 d *in vitro*, the

primary cortical neuron cultures were transfected in a 24-well plate format (1 µg of DNA per well) using Lipofectamine 2000 (Invitrogen, Carlsbad, CA) according to the instructions of the manufacturer. The following constructs were used: pcDNA3.1+Hygro, pCMV-SPORT6-Dexras1(wt), βARK-ct, pGL3-CRE-luciferase, E1B-luciferase, and Gal4-Elk1. Cells were treated 36 h posttransfection with the appropriate stimuli, and cell lysates were prepared 6 h after stimulation. Luciferase activity was measured using the Bright-Glo Luciferase Assay System (Promega, Madison, WI). To assess PACAP-mediated extracellular signal-regulated kinase (ERK) activation in culture, cortical neurons were treated as follows: tissue culture media were replaced with HEPES buffer (in mm: 137 NaCl, 25 glucose, 10 HEPES, 5 KCl, 1 MgCl₂, and 3 CaCl₂, pH 7.4) containing tetrodotoxin (1 μM) 30 min before experimental manipulation. Cells were pretreated (30 min) with PACAP 6-38 (200 nm) before a 15 min stimulation with PACAP (50 nm) and fixed (15 min) with paraformaldehyde (4% w/v). Phospho-ERK1/2 immunofluorescent labeling was performed as described previously (Butcher et al., 2005) using polyclonal antibodies against phospho-p44/42 MAP kinase (Thr202/Tyr204) (1:3000 dilution; Cell Signaling Technology, Beverly, MA). To identify neurons, cells were also labeled with a neuronal-specific nuclear protein (NeuN) monoclonal antibody (1:1000; Chemicon, Temecula, CA). Cells were processed with Alexa Fluor 488-conjugated goat anti-rabbit IgG antibody (1:1000; Invitrogen) and Alexa Fluor 594conjugated goat anti-mouse IgG antibody (1:1000; Invitrogen), and digital images were captured using a Zeiss (Oberkochen, Germany) 510 Meta confocal microscope.

Tissue collection and immunohistochemistry. Mice were maintained on a fixed 12 h LD schedule for a minimum of 2 weeks and dark adapted for 2 d before treatment. Unless otherwise stated, a single light pulse (15 min; 40 lux) was administered at circadian time (CT) 6 or CT 20, and the mice were returned to darkness before they were killed. p-ERK, c-Fos, and Per2 protein levels were assessed 30 min, 2 h, and 6 h after the start of the light pulse, respectively. For CT 6 light-induced p-ERK determination, a 5 min light pulse was administered. Control animals not exposed to light were killed at the same circadian time. Tissue processing, immunohistochemistry, and image analysis were performed exactly as described previously (Butcher et al., 2002). The following antibodies were used: phospho-p44/42 MAP kinase (Thr202/Tyr204) (1:3000 dilution; Cell Signaling Technology), c-Fos (1:40,000 dilution; Calbiochem, La Jolla, CA), Per2 (1:2500 dilution; Alpha Diagnostic International, San Antonio, TX), and biotinylated anti-rabbit IgG antibody (1:300; Vector Laboratories, Burlingame, CA). Immunodetection was accomplished using the horseradish peroxidase (HRP)-ABC technique (Vector Laboratories) according to the instructions of the manufacturer. Nickelintensified diaminobenzidine was used to visualize the signal. To quantify pERK expression, images of coronal hypothalamus were captured with a 10× objective and recorded using MetaMorph software (Universal Imaging, West Chester, PA). For analysis of pERK expression at CT 20, a digital oval 150 pixels (x-axis) \times 200 pixels (y-axis) was placed within the ventral SCN, and the mean pixel intensity was recorded. For analysis of pERK expression at CT 6, a crescent 150-pixel (x-axis)-wide was placed over the SCN shell, and the mean pixel intensity was recorded. Signal intensity for each SCN was normalized by subtracting the mean level recorded within a 200-pixel-wide circle placed within the hypothalamus lateral to the SCN. pERK data for each animal (two to three central SCN sections) were averaged, and then mean values were averaged for each time point. Absolute intensity measurements for pERK are expressed using a 0-4096 intensity scale.

Tissue used for fluorescent labeling was processed exactly as described previously (Butcher et al., 2005). The following primary antibodies were used: rabbit polyclonal anti-GFP (1:2000; University of Alberta, Edmonton, Alberta, Canada), and mouse monoclonal anti-phosphorylated ERK (1:2000; Sigma, St. Louis, MO). The following day, the sections were incubated with Alexa Fluor 488-conjugated goat anti-rabbit IgG antibody (1:1000; Invitrogen) and Alexa Fluor 594-conjugated goat anti-mouse IgG antibody (1:1000; Invitrogen). Fluorescent images were captured using a Zeiss 510 Meta confocal microscope (1- μ m-thick optical section).

Western blotting. SCN tissues were dissected and stored at -80° C until use. SCN tissues from four same-treated mice were pooled. Preparation

of protein lysate, PAGE, electroblotting, and immunodetection were performed as described previously (Lee et al., 2005). The following antibodies were used: affinity-purified rabbit polyclonal anti-cAMP response element (CRE)-binding protein (CREB) phosphorylated at Ser ¹³³ (1: 1000 dilution; Cell Signaling Technology); rabbit polyclonal anti-ERK1/2 (1:2000 dilution; Santa Cruz Biotechnology, Santa Cruz, CA); goat anti-rabbit IgG antibody conjugated to HRP (1:2000 dilution; PerkinElmer, Norwalk, CT); and goat anti-rabbit IgG antibody conjugated to alkaline phosphatase (AP) (1:2000 dilution; PerkinElmer). Bands were visualized using Renaissance chemiluminescent HRP or AP substrate (PerkinElmer).

cAMP assay. Mice were entrained to a fixed 12 h LD schedule for a minimum of 2 weeks and dark adapted for 2 d before treatment. At CT 20, mice received a 5 min light pulse of 40 lux and were decapitated immediately. The SCN was dissected, frozen on dry ice, and stored at -80°C until additional processing. Tissue from individual subjects was homogenized in 200 μl of 0.1 N HCl, and lysates were centrifuged for 2 min at 13,000 rpm. For each sample, a 100 μl aliquot of a 1:5 dilution was assayed using a direct method ELISA (Assay Designs, Ann Arbor, MI) according to the instructions of the manufacturer.

Results

The absence of Dexras1 restructures photic sensitivity as a function of the circadian phase

Given that Dexras1 is rhythmically expressed in the SCN (Panda et al., 2002; Ueda et al., 2002; Takahashi et al., 2003), and that clock entrainment is driven by phase-regulated signaling pathways, Dexras1 may be uniquely positioned to differentially contribute to photic entrainment as a function of circadian time. To begin to address this issue, we constructed a photic PRC of wildtype and dexras1^{-/-} mice. The PRC was constructed using light pulses (15 min; 40 lux) at 3 h intervals throughout the circadian cycle. As previously reported, a light pulse administered in the early subjective night (ZT 14) produced significant phase delays in wild-type but not in $dexras1^{-/-}$ mice (Fig. 1). In contrast, light presented in the late subjective night (ZT 20) induced robust phase advances in $dexras1^{-/-}$ mice but not in wild-type controls (Fig. 1), indicating that Dexras1 produces opposite effects on the phase-delaying and phase-advancing effects of light. Also surprising was that the absence of Dexras1 revealed a phase-advancing response to photic stimulation in the middle of the daytime (ZT 8) (Fig. 1). This is in marked contrast to wild-type mice (and most organisms) which, as expected, were not responsive to a light pulse administered in the mid-subjective day (ZT 8) (Fig. 1). The aggregate data indicate that Dexras1 plays a central role in regulating the phasic responsiveness of the clock to light.

The PAC1 receptor and the MAPK cascade couple light to clock entrainment during the late night in *dexras1*^{-/-} mice

Next, we identified the cellular signaling events that couple photic signals to phase advances during the late night in *dexras1*^{-/-} mice. A number of studies have shown that glutamate and PACAP are the principal neurotransmitters that convey photic information from the retina to the SCN (Ding et al., 1994; Chen et al., 1999; Kawaguchi et al., 2003). Interestingly, the role of PACAP as an effector of photic entrainment appears to be phase-biased: genetic ablation of PACAP or PAC1 receptors in mice strongly attenuates light-induced phase advances in the late night but has only a modest effect on phase delays in the early night (Hannibal et al., 2001; Kawaguchi et al., 2003). Given the role of PACAP in light-induced phase advances, combined with the finding that *dexras1*^{-/-} mice have potentiated light-induced phase advances raised the possibility that Dexras1 regulates PACAP-induced clock entrainment. To address this issue, we infused the PAC1 receptor-selective inhibitor PACAP 6–38 into

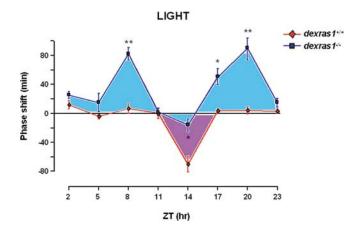


Figure 1. The photic response curve is altered in the absence of Dexras1. $dexras1^{+/+}$ (\spadesuit) and $dexras1^{-/-}$ (\blacksquare) mice were entrained to a fixed 12 h LD cycle of 400 lux. On their final LD cycle, the mice received a 15 min light pulse of 40 lux at designated ZT times (ZT 2, 5, 8, 11, 14, 17, 20, 23) and subsequently maintained in DD for at least 7 d. For ZT 2, 5, 8, and 11 experiments, lights remained off at ZT 0 on the day of the light pulse. Values are presented as mean \pm SEM phase shift (in minutes). Positive values on the *y*-axis denote phase advances, whereas negative values denote phase delays. n=6 per group. *p<0.05, **p<0.01 versus wild type (two-tailed Student's *t* test).

the third ventricle of $dexras1^{-/-}$ mice and assessed phase shifting in response to a single light pulse (15 min; 40 lux) administered at ZT 20. Vehicle-infused $dexras1^{-/-}$ mice showed significant light-induced phase advances (72 \pm 15 min; n = 12) (Fig. 2A). Importantly, infusion of PACAP 6–38 blocked light-induced phase advances in $dexras1^{-/-}$ mice (8.5 \pm 22 min; n = 13; p < 0.05 vs vehicle-infused control; two-tailed Student's t test) (Fig. 2A). These data reveal that Dexras1 antagonizes a late-night phase-advancing signaling pathway initiated by PACAP.

PACAP has been reported to stimulate activation of the MAPK signaling cascade (Barrie et al., 1997; Dziema and Obrietan, 2002; Butcher et al., 2005). Furthermore, in different experimental paradigms, Dexras1 has been shown either to promote or to inhibit MAPK activation (Cismowski et al., 1999; Graham et al., 2002). Given the central role of the MAPK pathway in the phase-delaying effects of light in mice (Butcher et al., 2002), we examined whether MAPK signaling was essential for the phaseadvancing effects of light during the late night in dexras1-/mice. To this end, dexras1^{-/-} mice were infused with the MAP kinase kinase 1/2 (MEK1/2) inhibitor U0126, and the phaseshifting effect of a single ZT 20 light pulse (15 min; 40 lux) was examined. Significant light-induced phase advances were observed in vehicle (DMSO)-infused $dexras1^{-/-}$ mice (72 \pm 7.1 min; n = 6) (Fig. 2B) but not in U0126-treated dexras1^{-/-} mice $(-12 \pm 14 \text{ min}; n = 6; p < 0.001 \text{ vs vehicle-infused control};$ two-tailed Student's t test) (Fig. 2C). Together, these data indicate that the ability of a signaling cassette formed by PACAP/ PAC1 and MAPK to convey photic timing cues to the clock is regulated by Dexras1.

Dexras1 represses late-night light-induced MAPK signaling in the SCN

The observation that late-night MAPK-dependent phase advancing is potentiated in $dexras1^{-/-}$ mice (Fig. 3B) compared with wild-type animals (Fig. 3A), combined with the recognized role of Dexras1 as a complex, context-specific regulator of the MAPK cascade (Cismowski et al., 1999; Graham et al., 2002), led us to examine the effects of Dexras1 on light-induced MAPK activation. In both wild-type and $dexras1^{-/-}$ mice, a single light pulse

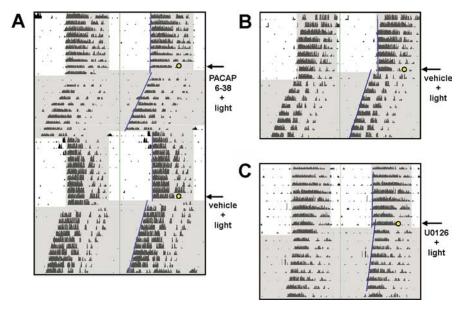


Figure 2. The enhanced late-night photic responsiveness of $dexras1^{-/-}$ mice is dependent on PAC1 receptor and p42/p44 MAPK activity. **A**, A representative actogram of wheel-running activity of $dexras1^{-/-}$ mice. Mice were entrained to a 12 h LD schedule (400 lux). Thirty minutes before a 15 min light pulse of 40 lux at ZT 20 (yellow circle), mice were infused with the PAC1 antagonist PACAP 6–38 (500 μ m; 3 μ l) via a guide cannula positioned in the third ventricle. Subsequent to the light pulse, animals were maintained in DD for at least 7 d. Animals were re-entrained, and the procedure was repeated for the counterbalanced treatment: vehicle infusion plus light pulse. Periods of darkness are shaded in gray. Activity onsets are indicated by blue lines. **B**, **C**, Representative actograms of wheel-running activity of $dexras1^{-/-}$ mice. Mice were entrained to a 12 h LD schedule (400 lux). Thirty minutes before a 15 min light pulse of 40 lux at ZT 20 (yellow circle), mice were infused with vehicle (DMSO) (**B**) or the MEK1/2 inhibitor UO126 (10 mm; 3 μ l) (**C**) via a guide cannula located in the third ventricle. Subsequent to the light pulse, animals were maintained in DD for at least 7 d.

(15 min; 40 lux) administered at CT 20 induced p-ERK expression in the SCN (Fig. 3C). However, the levels of light-induced p-ERK were significantly greater in the SCN of dexras1 -/- mice compared with those of wild-type controls. In addition, p-ERK immunoreactivity was observed throughout the SCN in dexras1^{-/-} mice, whereas in wild-type mice, p-ERK expression was confined to the ventral, retinoreceptive region of the SCN (Fig. 3C,F). The augmentation of light-induced p-ERK was paralleled by a greater increase in c-Fos-immunoreactive nuclei in the SCN of dexras1^{-/-} mice compared with that of wild types (Fig. 3D, G). Interestingly, under basal conditions, there was a significantly greater number of c-Fos-positive nuclei in dexras1 $^{-/-}$ SCN compared with wild-type controls (Fig. 3 D, G). As reported previously (Yan and Silver, 2002), light exposure in the late night did not induce Per2 expression in the SCN of wild-type mice (Fig. 3E). Consistent with the results from wild-type mice, light did not stimulate Per2 expression in dexras $1^{-/-}$ mice (Fig. 3E). The cumulative evidence indicates that Dexras1 suppresses photic responsiveness of the circadian system in the late night via a negative modulatory effect on light-induced MAPK activation.

Dexras1 negatively regulates PACAP-induced MAPK activation via both limbs of G-protein signaling

PAC1 receptors belong to the superfamily of G-protein-coupled receptors (GPCRs) that activate multiple intracellular signaling cascades (Spengler et al., 1993). The binding of PACAP to PAC1 induces the GTP-dependent dissociation of $G_s\alpha$ and $G\beta\gamma$ subunits, which independently engage multiple downstream effectors, including MAPK and AC (Spengler et al., 1993; Barrie et al., 1997). Given the finding that Dexras1 acts as a guanine nucleotide exchange factor for $G_{i/o}\alpha$ subunits and regulates $G\beta\gamma$ signaling, we assessed the potential negative regulatory effect of

Dexras1 on PACAP-mediated MAPK activation. To this end, PACAP-mediated activation of the MAPK pathway was monitored in cortical neuron cultures using a GAL4-Elk1/E1B-driven luciferase porter assay (Chung and Brautigan, 1999). Cotransfection of a Dexras1 expression vector attenuated PACAP-induced Elk1 transactivation relative to pcDNAtransfected controls by 55% (Fig. 4A). Interestingly, abrogation of $G\beta\gamma$ activity via cotransfection of a G $\beta\gamma$ scavenger, the C terminus of β -adrenergic receptor kinase (βARK-ct), reduced PACAP-induced Elk1 transactivation in pcDNA-transfected but not Dexras1-transfected cultures to 60% of control values (Fig. 4A). These data, which indicate that overexpression of a $G\beta\gamma$ scavenger phenocopies the effects of Dexras1, suggest that Dexras1 inhibits $G\beta\gamma$ activity and thereby decreases PACAP-induced MAPK activation.

The incomplete attenuation of PACAP-induced Elk1 transactivation by β ARK-ct indicates that an additional, $G\beta\gamma$ -independent pathway is engaged by PACAP and couples to the MAPK pathway. Along these lines, G_s -coupled receptors have been shown to stimulate the MAPK cascade via a cAMP-dependent mechanism (Norum et al., 2003). Interest-

ingly, as a GEF for G_{i/o}-protein subtypes, Dexras1 may be an effective modulator of AC-induced MAPK activation. To begin to examine the modulatory effects of Dexras1 on G_{i/o} activity and the consequence on PACAP-induced MAPK activation, we treated GAL4-Elk1/E1B-luciferase-expressing cortical cultures with the G_{i/o} inhibitor pertussis toxin (100 ng/ml) 2 h before stimulation with PACAP (100 nm). Pertussis toxin potentiated PACAP-induced Elk1 transactivation in pcDNA-transfected cultures by twofold (Fig. 4B). This is consistent with the removal of an inhibitory signal impinging after AC in the face of G_smediated stimulation. Pertussis toxin ADP-ribosylates $G_{i/o}\alpha$ and, thus, inhibits G_{i/o}-proteins only in the inactive, heterotrimeric form (Tsai et al., 1984). As noted above, Dexras1 acts as a guanine nucleotide exchange factor for $G_{i/o}\alpha$ subunits, thereby facilitating dissociation of $G_{i/o}\alpha$ from $G\beta\gamma$. Thus, Dexras1 overexpression should counter the effects of pertussis toxin if Dexras1 negatively modulates PACAP-induced MAPK via a $G_{i/o}\alpha$ -dependent mechanism. Indeed, we found that in neurons overexpressing Dexras1, pertussis toxin had no effect on PACAP-induced Elk1 transactivation (Fig. 4B). Given the mode of action of pertussis toxin, these data suggest that Dexras1 activates Gi/o-proteins, which have an inhibitory effect on PACAP-mediated, cAMPdependent, MAPK activation. To test this interpretation, the effects of Dexras1 on forskolin-induced Gal4-Elk1-mediated transcription were examined. Consistent with the effects on PACAP, Dexras1 attenuated forskolin-mediated activation of the MAPK cascade to 30% of control values (Fig. 4C). These data indicate that Dexras1 negatively regulates the capacity of G₂/cAMP to stimulate MAPK activation in neurons.

The CRE is found in a number of light-responsive genes and has been suggested to couple light to transcriptionally dependent entrainment of the SCN clock (Obrietan et al., 1999). Given the

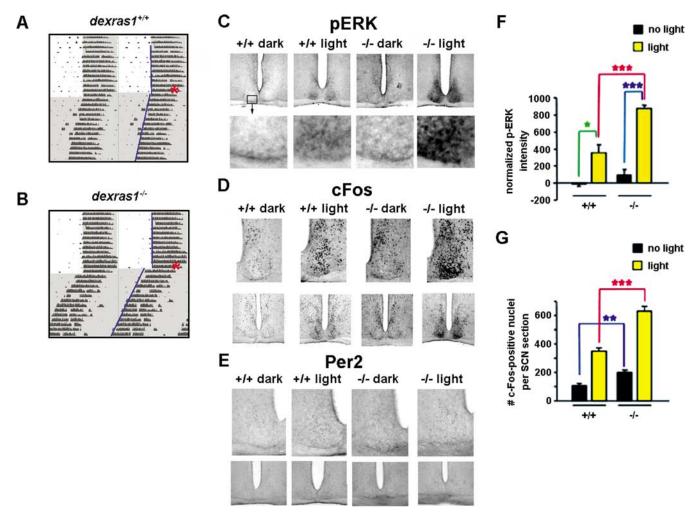


Figure 3. Disruption of Dexras1 potentiates the effects of late-night light exposure. *A*, *B*, Representative actograms of wheel-running activity of *dexras1*^{+/+} (*A*) and *dexras1*^{-/-} (*B*) mice. Mice were exposed to a single 40 lux light pulse for 15 min at ZT 20 (red asterisk), exactly as described in Figure 1. Activity onsets are indicated by blue lines. *C*–*E*, Immunohistochemical analysis of p-ERK (*C*), *c*-Fos (*D*), and Per2 (*E*) expression in response to a single 15 min light pulse at CT 20. Wild-type (+/+) and knock-out (-/-) mice were killed 0.5, 2, and 6 h after the light treatment, and brain sections were processed for phospho-ERK, c-Fos, and Per2, respectively. Dark control animals were not exposed to light but were killed at the same circadian time. *F*, Quantitation of p-ERK expression in the SCN. Data are presented as mean ± SEM densitometric intensity. Light-induced ERK activation was significantly enhanced in the SCN of *dexras1*^{-/-} mice relative to wild-type controls. *n* = 4 – 6 animals per group. **p < 0.05, ***p < 0.001 (two-tailed Student's *t* test). *G*, Quantitation of c-Fos expression in the SCN. Data are presented as mean ± SEM c-Fos-immunoreactive nuclei per SCN section. There was a significantly greater number of c-Fos-immunoreactive nuclei in the SCN of *dexras1*^{-/-} versus wild-type mice under basal conditions at CT 22 and 2 h after light treatment at CT 20. *n* = 6 –7 per group. **p < 0.01, ***p < 0.001 (two-tailed Student's *t* test).

capacity of Dexras1 to regulate G_s -mediated signaling, we tested its effects on CRE-mediated gene expression. Neurons were transfected with a CRE-regulated luciferase reporter plasmid and stimulated with forskolin. The coexpression of Dexras1 attenuated forskolin-induced CRE-mediated transcription by 60% relative to pcDNA-transfected controls (Fig. 4C). These data suggest that Dexras1 negatively modulates photic responses via an inhibitory effect on CREB/CRE-mediated transcription of clock genes per se or other target genes implicated in photic entrainment.

In a set of control experiments, we stimulated cortical neuronal cultures with PACAP (50 nm) for 15 min and assessed ERK activation. The levels of p-ERK were markedly elevated in PACAP-treated neurons compared with unstimulated cultures (Fig. 4D). Importantly, a 30 min pretreatment with the PAC1 antagonist PACAP 6–38 (200 nm) significantly suppressed p-ERK expression (Fig. 4D). These data indicate that PACAP-dependent signaling events actuated in cortical neurons are likely analogous to those observed in the SCN, although cortical neurons may not perfectly generalize to SCN neurons in terms of precise signaling mechanisms. Our cumulative observations sug-

gest that PACAP induces MAPK activation via both the $G_s\alpha$ and $G\beta\gamma$ limbs of G-protein signaling, and that Dexras1 not only inhibits $G\beta\gamma$, but it also provides additional inhibitory influences on G_s signaling via activation of $G_{i/o}$ -proteins.

Dexras1 inhibits cAMP production and cAMP-mediated signaling in the SCN

The results of the *in vitro* experiments described above suggested that cAMP-dependent signaling would be upregulated in *dexras1*^{-/-} mice. To examine this issue, we determined the levels of cAMP in SCN tissues of wild-type and *dexras1*^{-/-} mice during the late night (CT 20) under basal conditions and after light exposure. Basal levels of cAMP were demonstrably greater in *dexras1*^{-/-} SCN relative to wild type (Fig. 5A). Furthermore, a single light pulse at CT 20 evoked a significant increase in cAMP in the SCN of *dexras1*^{-/-} but not wild-type mice (Fig. 5A). In accordance with these data, basal and inducible levels of phospho-CREB were altered in the SCN of *dexras1*^{-/-} mice. In wild-type mice, basal expression of pCREB in the SCN was low at CT 20 (Fig. 5B) and light induced robust pCREB expression. In

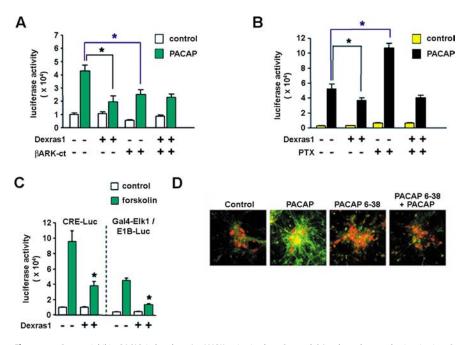


Figure 4. Dexras1 inhibits PACAP-induced p42/44 MAPK activation by a $G\alpha$ - and $G\beta\gamma$ -dependent mechanism *in vitro*. **A**, Primary cultures of rat embryonic cortical neurons were maintained for 8 d. Subsequently, they were transfected with an E1Bluciferase reporter gene construct and the Gal4-Elk1 expression vector, in combination with expression constructs for Dexras1 or a G $\beta\gamma$ scavenger, β ARK-ct. Thirty-six hours posttransfection, cultures were treated with 100 nm PACAP and assayed 6 h later. PACAP-induced Gal4-Elk1 activation was attenuated in Dexras1-overexpressing neurons compared with empty vector controls. BARK-ct expression reduced the magnitude of PACAP-induced Gal4-Elk1 activation in empty vector controls only but had no effect on Dexras1-overexpressing neurons. B, Cortical neurons were transfected as described in A, with or without cotransfection of the Dexras1 expression construct. Cultures received a 2 h pretreatment with 100 ng/ml pertussis toxin (PTX) followed by stimulation with 100 nm PACAP for 6 h. Pertussis toxin potentiated PACAP-induced Gal4-Elk1 activation in empty vector controls but had no effect on Dexras1-overexpressing neurons. C, Cortical neurons were transfected with a CRE-luciferase reporter gene construct or, alternatively, an E1B-luciferase reporter gene construct in combination with the Gal4-Elk1 expression vector. Neurons were cotransfected with empty vector or the Dexras1 expression construct. Thirty-six hours posttransfection, cultures were treated with 10 μ m forskolin and assayed 6 h later. In both CRE and E1B assays, the effects of forskolin were significantly reduced by cotransfection of Dexras1. Data are presented as the mean \pm SEM of quadruplicate determinations. *p < 0.05 versus empty vector control (two-way ANOVA). D, Rat cortical neuronal cultures were stimulated with PACAP (50 nm) for 15 min before a 30 min pretreatment with PACAP 6 – 38 (200 nm) (far-right panel). After stimulation, cells were fixed and the expression of p-ERK (green), and NeuN (a neuron-specific marker: red) was determined by immunocytochemistry. ERK activation was compared with cultures that had received no treatment (control, far-left panel), PACAP alone (center-left panel), and PACAP 6 – 38 alone (center-right panel).

dexras1^{-/-} mice, not only were basal pCREB levels in the SCN markedly greater compared with wild-type controls, photic stimulation elevated pCREB expression even further, beyond that observed in light-treated wild-type mice (Fig. 5*B*). Collectively, these data indicate that the effects of Dexras1 on cAMP levels are reflected in the transactivation potential of CREB, one of the central regulators of clock entrainment.

So far, our data indicate that Dexras1 functions as an inhibitor of signaling events that are initiated by PACAP/PAC1 receptors and impinge after the p42/p44 MAPK cascade. To test this possibility, we next examined the capacity of PACAP to elicit ERK activation in vivo in wild-type and dexras1^{-/-} mice. Infusion of PACAP (40 µm; 3 µl) into the lateral ventricle at CT 20 produced a significant increase in p-ERK levels in the SCN and periventricular region of both wild-type and dexras1^{-/-} mice (Fig. 5C,D). Quantitative analysis of p-ERK immunoreactivity in the SCN indicated that PACAP was more potent at stimulating ERK activation in the knock-outs compared with the wild types (Fig. 5D). Most notably, pretreatment with PACAP 6–38 (500 μ M; 3 μl) strongly inhibited the ability of PACAP to elevate p-ERK expression in $dexras1^{-/-}$ mice (Fig. 5C,D). Although these data demonstrate the efficacy of PACAP 6-38 as a potent antagonist of PACAP, we cannot rule out the possibility that PACAP 6–38

might have effects that are not attributed to PACAP. However, in support of its efficacy, PACAP 6–38 has been used in several *in vivo* studies (Chen et al., 1999; Bergstrom et al., 2003) as a PAC1-selective antagonist and was biochemically characterized in isolated membrane binding studies as a strong competitive antagonist of full-length PACAP (Robberecht et al., 1992).

In vivo abrogation of light-induced MAPK activation in the SCN of transgenic mice that express a constitutively active form of Dexras1

To provide in vivo data that Dexras1 expression in the SCN critically modulates the effects of nocturnal light, we generated an inducible transgenic mouse strain that overexpresses a constitutively active form of Dexras1. The alanine → valine mutation in codon 178 interrupts the guanine nucleotide binding pocket of Dexras1 and results in decreased affinity for GTP/GDP binding as well as increased exchange of GDP for GTP (Graham et al., 2001). Transgenic mice were generated, which expressed the A178V mutant form of Dexras1 bicistronically with eGFP under the control of a tetracycline-controlled promoter (Fig. 6A). Individual founder lines were bred with the CaMKII α -tTA mice, which drive transgene expression in the absence of doxycycline in a tissuemanner. Three of specific Dexras1(A178V)-IRES-eGFP founder lines, when intercrossed with CaMKII α tTA, showed transgene expression that was restricted to the SCN (Fig. 6A). The SCN specificity of one of these founder lines, Dexras1(A178V) #3901, becomes

apparent in low-magnification images where the SCN-containing coronal section can be viewed (Fig. 6*B*, left panels). This is in stark contrast to founder lines that exhibit broader expression encompassing the SCN, cortex, striatum, and hippocampus (Fig. 6*B*, right panels). All subsequent experiments were performed using the SCN-specific founder lines.

To address the role of Dexras1 in light-induced MAPK activation *in vivo*, SCN-specific Dexras1(A178V) double transgenic mice received a brief light pulse in the late subjective night. p-ERK expression was observed in the SCN in response to light but was notably absent in neurons that overexpressed Dexras1(A178V) (Fig. 6C, left panel). The absence of an overlap between the signals for p-ERK and the GFP marker was made more apparent under higher magnification to resolve individual neurons (Fig. 6C, right panel). Hence, using a tissue-specific transgenic approach in a whole animal model, we provide *in vivo* evidence that Dexras1 functions within the SCN to inhibit the coupling of light to the MAPK pathway.

Dexras1 gates the effects of daytime light exposure on MAPK activation and changes in clock gene expression

In nocturnal rodents, the circadian system is unresponsive to photic cues presented in the day, both at a behavioral as well as a cellular level (Pittendrigh and Daan, 1976; Obrietan et al., 1998). However, genetic inactivation of Dexras1 leads to a loss of daytime gating of light (Figs. 1, 7A, B). To examine where within the photic entrainment pathway Dexras1 was affecting the gate, we analyzed the expression of p-ERK, c-Fos, and Per2 after a 15 min light pulse administered during the subjective daytime. As reported previously (Obrietan et al., 1998), basal levels of p-ERK at CT 6 were strong and confined to the dorsal and mediolateral regions of the SCN of wild-type mice (Fig. 7C), and a single light pulse did not induce additional p-ERK immunoreactivity (Fig. 7C,F). In dexras $1^{-/-}$ mice, basal levels of p-ERK were moderately attenuated in the dorsomedial region of the SCN compared with wild-type mice (Fig. 7C,F). Furthermore, light induced a significant increase in p-ERK expression in the dorsomedial SCN (Fig. 7C,F). This region of the SCN encompasses a significant portion of the SCN "shell," the locus of the circadian pacemaker. It is important to note that this light-induced p-ERK expression pattern is distinct from the ventrolateral pattern of p-ERK expression that is normally observed after photic stimulation in the subjective night.

Phase-restricted gene expression was also abrogated in $dexras1^{-/-}$ mice. Consistent with light-induced p-ERK expression, we found that photic stimulation during the subjective day (CT 6) triggered a marked increase in expression of the immediate early gene c-Fos (Fig. 7D, G). Interestingly, under control conditions, $dexras1^{-/-}$ mice exhibited significantly less c-Fos expression in the dorsomedial SCN than wild-type mice (Fig. 7D, G). Consistent with previous reports (Rusak et al., 1992), light exposure during the day (CT 6) failed to increase the expression of c-Fos in wild-type mice (Fig. 7D, G).

Paralleling the c-Fos and p-ERK results, *dexras*1^{-/-} mice exhibited marked light-induced pCREB expression at CT 6, whereas in wild-type mice, CREB phosphorylation levels were not affected by light (Fig. 7*I*).

To assess whether daytime photic responses in $dexras1^{-/-}$ mice correlate with an alteration in the expression of circadian clock proteins, we processed tissue for the expression of Per2. In wild-type mice, a single mid-subjective day (CT 6) light pulse did not alter Per2 expression examined 6 h later (Fig. 7*E*,*H*). In striking contrast, $dexras1^{-/-}$ mice showed a significant light-induced reduction in Per2 immunoreactivity in the SCN shell (Fig. 7*E*,*H*). Interestingly, this light-induced decrease in Per2 expression mirrors the effects of nonphotic entrainment stimuli presented in the day (Maywood et al., 1999). Together, the data reveal that Dexras1 functions as an essential daytime gate that restricts access of photic information to the shell region of the SCN but is not necessary for gating of light-induced MAPK activation in the SCN core.

The p42/p44 MAPK cascade is required for daytime photic phase advances in *dexras1*^{-/-} mice

Nighttime photic entrainment of the circadian clock is driven by the MAPK cascade. To determine whether daytime light responses in $dexras1^{-/-}$ mice also require signaling via the MAPK pathway, we infused mice with U0126 and examined the behavioral effects of a single light pulse (40 lux; 15 min) administered at ZT 8. Vehicle-infused $dexras1^{-/-}$ mice showed robust light-induced phase advances (76 \pm 11 min; n = 6) (Fig. 7*J*). In contrast, U0126 abrogated the phase-advancing effect of light (4.2 \pm 8.2 min; n = 6; p < 0.001 vs vehicle-infused control; two-tailed Student's t test) (Fig. 7*K*). Together, the data indicate that Dexras1 functions as a molecular gate that suppresses MAPK-dependent entrainment to light in the day.

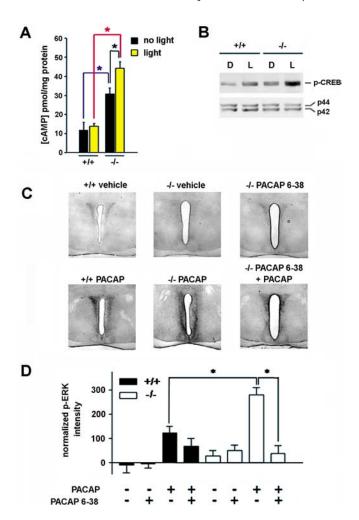


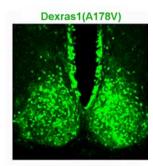
Figure 5. Dexras1 modulates cAMP-dependent signaling and PACAP/PAC1-mediated MAPK activation in the SCN. **A**, Wild-type and dexras $1^{-/-}$ mice received a single light pulse (5 min, 40 lux) at CT 20, and the SCN was dissected immediately and analyzed for cAMP content by ELISA. Control subjects did not receive a light pulse but were killed at the same circadian time. Light did not increase cAMP levels in wild-type mice (+/+). However, $dexras 1^{-/-}$ mice (-/-) exhibited a significant light-induced increase in cAMP. Additionally, under basal conditions, dexras 1^{-/-} mice had significantly elevated cAMP levels relative to wild-type mice. Values are presented as mean \pm SEM pmol cAMP per mg of protein. n=4-5 per group. *p<0.05(two-way ANOVA). **B**, Western blot analysis of light-induced p-CREB expression in the SCN. Wild-type and dexras $1^{-/-}$ mice received a single light pulse (15 min, 40 lux) at CT 20, and the SCN was dissected 30 min after the start of the light exposure. Pooled SCN extracts were probed for the expression of p-CREB. Expression of total ERK1/2 (p42, p44) was used as the loading control. L denotes light-treated mice, whereas D denotes dark controls. \boldsymbol{C} , \boldsymbol{D} . Wild-type (+/+)and dexras $1^{-/-}$ (-/-) mice were infused with the PAC1 antagonist PACAP 6 – 38 (500 μ M; 3 μ I) via a guide cannula positioned in the lateral ventricle 30 min before a second infusion of PACAP (40 μ M; 3 μ I). Thirty minutes after the second infusion, mice were killed, and brain sections were processed for p-ERK. ERK activation was compared with mice that received infusions of vehicle only, PACAP alone, or PACAP 6-38 alone. Representative micrographs are provided in **C**. Quantitation of p-ERK expression in the SCN is given in **D**. Data are presented as mean \pm SEM p-ERK-immunoreactive nuclei per SCN section. There was a significant effect of genotype on PACAP-mediated ERK activation as well as a significant effect of PACAP 6 – 38 on PACAP-induced p-ERK expression in the knock-out SCN. n = 6-7 per group. *p < 0.01 (twotailed Student's t test).

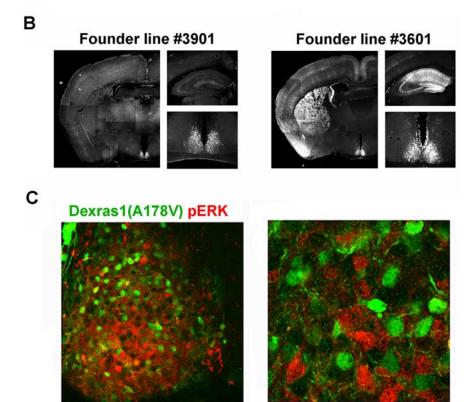
SCN-targeted expression of constitutively active Dexras1 leads to aberrant phase-shifting response to late-night light exposure

Thus far, we have shown that in the absence of Dexras1, late-night phase advances are potentiated and the gated unresponsiveness, which is characteristic of the subjective day, is lost. To provide another line of evidence validating the role of Dexras1 in photic

A Tet OFF: CaMKIIα polyA sv40 - Dox tTA Dox

Dexras1(A178V) IRES





eGFP

Figure 6. Tetracycline-controlled overexpression of constitutively active Dexras1 in the SCN abrogates MAPK activation in response to nocturnal light. A, Experimental schematic to drive SCN-specific expression of the constitutively active Dexras1 $(A178V)\ mutant\ using\ the\ tetracycline-inducible\ system.\ A\ single\ transgenic\ construct\ containing\ two\ genes,\ constitutively\ active$ Dexras1 harboring the A178V point mutation and eGFP, in a polycistronic unit under the control of the tetracycline-responsive (tet0) promoter was generated. The Dexras1 (A178V)-IRES-eGFP transgenic mice were bred with the CaMKII α -tTA mice. In the absence of the tetracycline analog doxycycline (Dox), double transgenic mice coexpress Dexras1(A178V) and eGFP in a tissuespecific manner. The Dexras1(A178V)-IRES-eGFP founder line #3901 shown here expresses specifically in the SCN. GFP immunoreactivity was found throughout the SCN, marking cells that express the Dexras1(A178V) transgene. B, Expression of eGFP in double transgenic mice generated from the breeding of the CaMKII α -tTA strain to Dexras1(A178V)-IRES-eGFP founder lines #3901 (left) and #3601 (right). Founder line #3901 exhibited SCN-specific expression, whereas transgene expression was more broadly expressed in founder line #3601 and observed in the SCN, cortex, hippocampus, and striatum. C, Light-induced MAPK activation in double transgenic CaMKII α -tTA::Dexras1(A178V)-IRES-eGFP mice. Double transgenic mice received a brief light pulse (15 min; 100 lux) at CT 20, and brain sections were processed 30 min later for expression of p-ERK (red) and GFP (green) using immunofluorescent labeling and confocal microscopy was used to visualize the signals. Cells that express the transgenes do not exhibit p-ERK immunoreactivity after light treatment (left panel). The absence of colocalization of GFP and p-ERK expression is more evident under higher magnification (right panel).

responsiveness, we tested the constitutively active Dexras1 transgenic mice in behavioral assays. Dexras1(A178V)-IRES-eGFP:: CamkII α -tTA double transgenic mice (which overexpress constitutively active Dexras1) and the Dexras1(A178V)-IRES-eGFP (which do not) single transgenic controls received a single light pulse (40 lux; 15 min) at ZT 8 or ZT 20 under entrained condi-

tions (Fig. 8). After 9-10 d of free running in DD, the mice received a second light pulse (40 lux; 15 min) at CT 8 or CT 20 (Fig. 8). Daytime light stimulation, whether administered under entrained or free-running conditions, had no phaseshifting effect in the single transgenic mice (Fig. 8A, E). A small phase advance was observed in double transgenic mice (Fig. 8 B, E) in response to a ZT 8 light pulse, but this was not significantly different from that of single transgenic controls. No phase advance was observed in double transgenics after a CT 8 light pulse given under free-running conditions (Fig. 8B,E). These results indicate that photic gating is not compromised in the presence of Dexras1. In contrast, the single transgenics did not phase-shift in response to a ZT 20 light pulse under entrained conditions but exhibited a modest phase advance when light was administered at CT 20 under free-running conditions (Fig. 8C,E). The response of Dexras1(A178V)overexpressing double transgenics to a ZT 20 light pulse was comparable with that of control mice (Fig. 8 *D*, *E*). However, rather than phase-advancing to light administration under free-running conditions, the double transgenics phase-delayed to CT 20 light (Fig. 8D, E). Interestingly, this phenocopies the behavior of PAC1^{-/-} mice, which show a similar phase delay in response to late-night light stimulation (Hannibal et al., 2001). Together, these data indicate Dexras1 is functioning to suppress PACAP-mediated signaling.

Discussion

Our results demonstrate the fundamental role of Dexras1 in shaping the phasedependent responsiveness of the mammalian circadian clock to photic entrainment cues. In the absence of Dexras1, the murine circadian system exhibits complex, phase-specific alterations in photic entrainment. Light-induced phase delays are attenuated in the early night, whereas photic phase advances in the late night are potentiated. Surprisingly, daytime gating of photic effects is lost in dexras1^{-/-} mice. The capacity to modulate MAPK signaling via discrete, time-of-day-specific signaling events appears to be the principal route by which Dexras1 regulates light sensitivity of the clock (Fig. 9).

Given that the MAPK modulator, Dexras1, is expressed in the SCN and is a

target of NMDA receptor/neuronal nitric oxide synthase-mediated activation (Fang et al., 2000), combined with evidence showing that photic resetting of the SCN clock requires activation of NMDA receptors, release of NO, and activation of the MAPK cascade (Colwell and Menaker, 1992; Ding et al., 1994; Shibata et

al., 1994; Mintz et al., 1999; Butcher et al., 2002; Hannibal, 2002), we hypothesized that Dexras1 may regulate aspects of the light entrainment process. As we demonstrated previously, the absence of Dexras1 was shown to attenuate light-induced phase shifts of behavioral rhythms in the early night as well as NMDA receptor signaling in the SCN (Cheng et al., 2004). This was coincident with a reduction in light-induced MAPK activation in the SCN of *dexras1*^{-/-} mice (Cheng et al., 2004).

In light of these results, it was entirely unexpected that the signaling events actuated by late-night light exposure are amplified in the absence of Dexras1, as demonstrated by a greater increase of light-induced p-ERK and c-Fos expression in the SCN of *dexras1*^{-/-} mice compared with wild types. Enhanced MAPK activation in *dexras1*^{-/-} mice is concomitant with a U0126-mediated blockade of phase advances, indicating a causal link between enhanced signaling through the MAPK cascade and a dramatic increase in the sensitivity of the clock to late-night light.

The initial signaling events that mediate the opposite effects of Dexras1 on light-induced phase delays (early night) and phase advances (late night) are clearly distinct. In contrast to the role of Dexras1 as a regulator of NMDA receptormediated early night phase delays, our data suggest that the exaggerated behavioral response of dexras1^{-/-} mice to latenight light exposure is mediated by enhanced PACAP/PAC1 signaling in the SCN. Several reports using genetically altered mice indicate that PACAP is a key regulator of late-night phase advances (Hannibal et al., 2001; Kawaguchi et al., 2003); however, there is some discrepancy with respect to the role of PACAP in earlynight phase delays (Hannibal et al., 2001; Kawaguchi et al., 2003; Colwell et al., 2004). Support for a connection between PACAP and Dexras1 came from data showing that pharmacological antagonism of PAC1 receptors blocked photic phase advances in dexras1^{-/-} mice, and that PACAP infusion triggered more robust MAPK pathway activation in dexras1^{-/-} mice. Moreover, whereas the absence of Dexras1 potentiated light/PACAP effects in vivo, overexpression of Dexras1 attenuated PACAP-mediated downstream signaling in vitro.

Our pharmacological data indicate that Dexras1 attenuates late-night light entrainment by reducing the capacity of PACAP to elicit MAPK activation via PAC1 receptors. However, we cannot exclude the possibility that the effects of PACAP 6–38 are restricted solely to PAC1. In consideration of this, another possible signaling cassette by which Dexras1 mod-

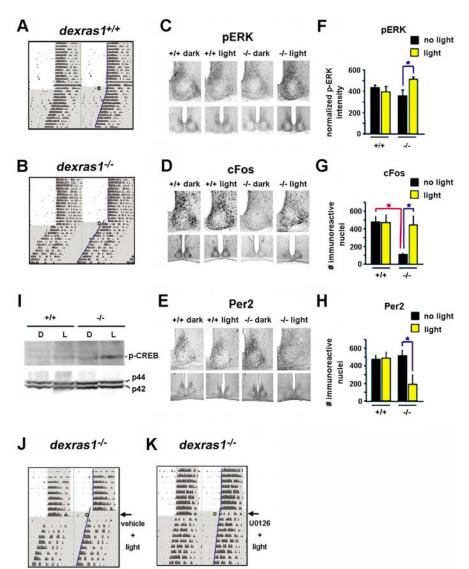


Figure 7. The absence of Dexras1 reveals a mid-day photic response in the SCN. A, B, Representative actograms of wheelrunning activity of dexras $1^{+/+}$ (A) and dexras $1^{-/-}$ (B) mice. Mice were exposed to a single 40 lux light pulse for 15 min at ZT 8 (yellow circle), exactly as described in Figure 1. Activity onsets are indicated by blue lines. C-E, Immunohistochemical analysis of p-ERK (C), c-Fos (D), and Per2 (E) expression in response to a single 5 min (for p-ERK) or 15 min (for c-Fos and Per2) light pulse at CT 6. Wild-type (+/+) and knock-out (-/-) mice were killed immediately (for p-ERK), 2 h (for c-Fos), or 6 h (for Per2) after the light treatment. Dark control animals were not exposed to light but were killed at the same circadian times. Light exposure in the subjective daytime had no effect on p-ERK, c-Fos, and Per2 levels in the SCN of wild-type mice. Baseline levels of c-Fos were reduced in dexras1^{-/-} mice relative to wild-type controls. A single light pulse in the mid-subjective day increased p-ERK and c-Fos immunoreactivity in the SCN of dexras1^{-/-} mice. Per2 immunoreactivity was decreased in dexras1^{-/-} SCN after light treatment. F-H, Quantitation of p-ERK (F), c-Fos (G), and Per2 (H) expression in the SCN. Data are presented as mean \pm SEM densitometric intensity (for p-ERK) or mean \pm SEM number of immunoreactive nuclei per SCN section (for c-Fos and Per2). n=4-6 per group. *p < 0.05 (two-tailed Student's t test). I, Western blot analysis of light-induced p-CREB expression in the SCN. Wild-type (+/+) and $dexras 1^{-/-}$ (-/-) mice received a single light pulse (15 min, 40 lux) at CT 6, and the SCN was dissected 30 min after the start of the light exposure. Pooled SCN extracts were probed for the expression of p-CREB. Expression of total ERK1/2 (p42, p44) was used as the loading control. L denotes light-treated mice, whereas D denotes dark controls. J, K, U0126 inhibits ZT 8 light-induced phase advances in dexras1^{-/-} mice. Representative actograms of wheel-running activity of $dexras 1^{-/-}$ mice are shown. Mice were entrained to a 12 h LD schedule (400 lux). On the day of the experiment, lights remained off at ZT 0. Thirty minutes before a 15 min light pulse of 40 lux at ZT 8 (yellow circle), mice were infused with vehicle (DMSO) (J) or U0126 (10 mm; 3 μ l) (K) via a guide cannula placed in the third ventricle. Subsequent to the light pulse, animals were maintained in DD for at least 7 d. Activity onsets are indicated by blue lines.

ulates MAPK activation within the SCN is the following: glutamate-vasoactive intestinal peptide (VIP)-VPAC2/PAC1-Dexras1-MAPK. VIP, which is expressed by retinorecipient SCN neurons, has been shown to phase-shift SCN firing rhythms *in vitro* in a manner similar to light or glutamate (Reed et al., 2001).

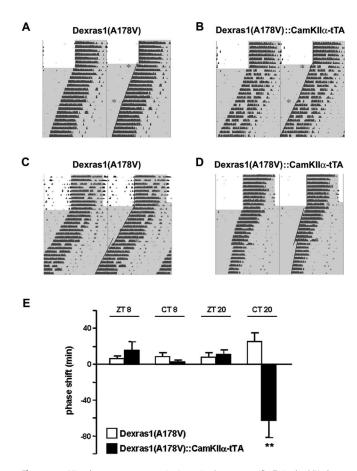


Figure 8. Mice that overexpress constitutive active Dexras1 specifically in the SCN phase-delay in response to late-night light exposure. A–D, Representative actograms of wheel-running activity of Dexras1(A178V) single transgenic (A, C) and Dexras1(A178V)-IRES-eGFP:: CaMKIl α -tTA double transgenic (B, D) mice. Mice were exposed to a single 40 lux light pulse (red asterisk) for 15 min at ZT 8 (A, B) or ZT 20 (C, D), exactly as described in Figure 1. Mice received a second light pulse (40 lux; 15 min) at CT 8 (A, B) or CT 20 (C, D) after 9 –10 d in DD. Activity onsets are indicated by blue lines. E, Quantitation of light-induced phase shifts. Values are presented as mean \pm SEM phase shift (in minutes). n = 6 - 8 per group. **p < 0.01 versus same-treated Dexras1(A178V) single transgenic mice (two-tailed Student's t test).

Moreover, VIP resetting effects are mimicked by a VPAC2 agonist and are abrogated by U0126 (Reed et al., 2001; Meyer-Spasche and Piggins, 2004). With regard to VPAC2 signaling, the receptor shows equal affinity for PACAP and VIP (Harmar et al., 1998); therefore, some cellular effects of PACAP within the murine SCN may be mediated in part by VPAC2 receptors (Reed et al., 2002).

To address the actions of Dexras1 in the late night, we focused on the role of PACAP/PAC1-G-protein-mediated signaling. In vitro, Dexras1-dependent attenuation of PACAP-induced MAPK activity was attributed to an inhibitory effect on both $G\beta\gamma$ and $G_s \alpha$ signaling. MAPKs are downstream effectors of $G\beta \gamma$ signaling and, as with other targets of $G\beta\gamma$ -mediated signaling, have been shown to be negatively modulated by Dexras1 (Takesono et al., 2002). At the same time, enhanced $G_i\alpha$ signaling in Dexras1expressing neurons likely accounts for the suppression of $G_s\alpha$ mediated MAPK activation. Although Dexras1 has been shown to inhibit GPCR-dependent activation of heterotrimeric G-proteins, the ability of Dexras1 to independently cause GTP/ GDP exchange on monomeric G_iα subunits has been demonstrated in biochemical studies (Cismowski et al., 2000). Pertussis toxin, which acts only on the inactive, heterotrimeric form of G_{i/o}-proteins, is without effect on PACAP-dependent MAPK activation in neurons that overexpress Dexras1. This supports our hypothesis that Dexras1 is an intracellular, receptor-independent promoter of the active, monomeric form of $G_i\alpha$, which provides an inhibitory tone to $G_s\alpha$ signaling.

In addition, we provide multiple lines of evidence that Dexras1 inhibits AC signaling (Graham et al., 2004; Nguyen and Watts, 2005). Forskolin-induced activation of CRE-dependent transcription is inhibited in Dexras1-expressing neurons *in vitro*. Moreover, basal and light-inducible levels of cAMP as well as active, phospho-CREB are greater in the SCN of *dexras1*^{-/-} mice versus wild types. These inhibitory effects of Dexras1 were realized at the level of the MAPK pathway: forskolin-induced MAPK activity was inhibited by Dexras1. The augmentation of PACAP/PAC1-mediated signaling in *dexras1*^{-/-} mice may in part be attributed to enhanced G_s/AC signaling, whereas the accumulation of cAMP in the SCN of *dexras1*^{-/-} mice may play a role in the late-night photic phase advances.

Most surprising was our finding that genetic ablation of Dexras1 revealed a response to daytime light. A lack of responsiveness to daytime light is a fundamental feature of all biological clocks. In light of a possible role of VPAC2 signaling in dexras1^{-/-} mice, it is interesting to note that photic gating is also compromised in VPAC2^{-/-} mice, which showed light-induced expression of p-ERK and c-Fos in both the subjective day and night (Hughes et al., 2004). Generally speaking, there are two possible explanations for how daytime light couples to the clock in dexras1^{-/-} mice. First, ablation of Dexras1 might reveal a capacity of photic stimulation to elicit a behavioral arousal-like entrainment cue, which in turn would result in a nonphotic phase shift. Second, ablation of Dexras1 might reveal a novel photic entrainment mechanism that is distinct from an arousal-like entrainment cue.

The possibility that the loss of Dexras1 reveals a photically activated arousal entrainment cue is supported by our data showing that a midday light pulse in *dexras1*^{-/-} mice suppressed Per2 protein expression in the SCN in a similar manner as exposure to a nonphotic stimulus (Maywood et al., 1999). In contrast, there are distinct differences between photic and nonphotic phase advances in the subjective day. For example, nonphotic stimulation leaves c-Fos levels in the SCN unchanged (Mead et al., 1992) or decreased (Mikkelsen et al., 1998), but we observed daytime photic induction of p-ERK and c-Fos in the SCN of *dexras1*^{-/-} mice.

If a direct photic entrainment mechanism underlies daytime clock entrainment in $dexras1^{-/-}$ mice, it would appear that the signaling events underlying daytime and nighttime lightentrainment are distinct. Although U0126 blocked the phase-shifting effect of light at ZT 8 and ZT 20, only a light pulse administered at ZT 20 evoked the expected p-ERK and c-Fos expression in the SCN core. Daytime light exposure induced the expression of p-ERK and c-Fos in the SCN shell without a similar response in the SCN core.

The mechanistic basis for light-induced MAPK activation exclusively in the SCN shell is unclear. Although the SCN core has been considered as the retinorecipient region, a recent study indicated that projections of melanopsin-expressing retinal ganglion cells were not restricted to this region but filled the entire murine SCN, including the core and shell subcompartments (Hattar et al., 2006). Hence, it is possible that light may trigger MAPK activation in the SCN shell without a requirement for concurrent activation in the SCN core. Alternatively, the SCN shell is densely innervated by galanin-positive afferents from the anterior hypothalamus/preoptic area (Abrahamson and Moore, 2001). The intergeniculate leaflet (IGL), which mediates both

photic and nonphotic effects, receives afferent inputs from the RHT and projects to a number of brain regions, including the SCN and the anterior hypothalamus (Morin and Blanchard, 1999). As such, the anterior hypothalamus may serve as a secondary route by which photic (and nonphotic) cues arising from the IGL reaches the SCN shell. Regardless of the mechanism, the results indicate that genetic ablation of Dexras1 eliminates a daytime gate that otherwise limits access of photic information to the SCN shell but does not compromise gating of the SCN core. Additional work will be required to determine both the synaptic circuitry and precise cellular signaling events that underlie light-induced phase advancing *dexras1*^{-/-} mice during the subjective day.

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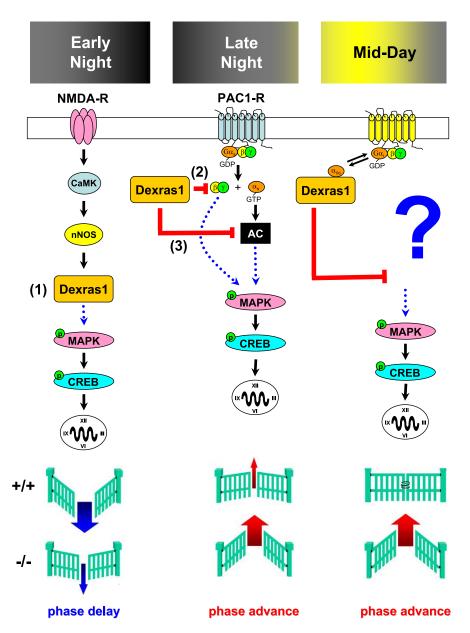


Figure 9. Proposed model of Dexras1 in modulating photic responsiveness of the circadian clock. Photic effects are mediated in part by NMDA and PAC1 receptors expressed in the SCN. In the early night, light-induced activation of NMDA receptors leads to a nitrosylation-dependent enhancement of the guanine nucleotide exchange activity of Dexras1. 1, As a result, Dexras1 activates the MAPK pathway and promotes photic resetting in the early night. Light exposure in the late night leads to activation of G_s -coupled PAC1 receptors, which signal via both the $G_s \alpha$ and $G\beta\gamma$ limbs to the MAPK cascade. Dexras1 inhibits PAC1-mediated MAPK pathway activation by suppressing $G\beta\gamma$ signaling events (2) as well as AC (3). Dexras1 may inhibit AC indirectly by a receptor-independent enhancement of tonic $G_{i/o}\alpha$ activity. Although we do not know the upstream signaling mechanisms that govern photic resetting in the midday in $dexras1^{-/-}$ mice, our data implicate the downstream activation of p42/p44 MAPK and CREB in the observed daytime photic responses. In summary, light induces smaller phase delays in the early night and larger phase advances in the late night in $dexras1^{-/-}$ mice. In the midday, a photic gate is lost in these animals, revealing an unusual phase advance response.

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