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G PROTEIN-COUPLED INWARDLY-RECTIFYING POTASSIUM (GIRK) CHANNELS MEDIATE ENTRAINMENT OF CIRCADIAN RHYTHMS

by

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A DISSERTATION

Submitted to the graduate faculty of The University of Alabama at Birmingham, in partial fulfillment of the requirements for the degree of Doctor of Philosophy

BIRMINGHAM, ALABAMA

G PROTEIN-COUPLED INWARDLY-RECTIFYING POTASSIUM (GIRK) CHANNELS MEDIATE ENTRAINMENT OF CIRCADIAN RHYTHMS

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CELL, MOLECULAR, AND DEVELOPMENTAL BIOLOGY; GRADUATE BIOMEDICAL SCIENCES

ABSTRACT

Circadian rhythms are 24-hour cycles in biological and behavioral processes. These cycles enable an organism to predict changes in its environment, like changes in food availability and seasonality. Although endogenously driven, these rhythms can entrain or synchronize to daily changes in the environment, allowing the animal to adapt. One way entrainment occurs is shifts in circadian phase following the presentation of nonphotic, or non-light, stimuli, such as exercise, arousal, or stress at certain times of day. The molecular mechanisms underlying nonphotic entrainment are poorly understood - specifically, how nonphotic cues alter excitability within the suprachiasmatic nucleus (SCN) of the hypothalamus, the clock center of the mammalian brain, to change the timing of circadian rhythms. This dissertation tests the hypothesis that nonphotic stimuli activate G protein-coupled inwardly-rectifying potassium (GIRK) channels which decrease neuronal excitability, modulating the timing of circadian rhythms. We show that not only is GIRK channel protein and function regulated in a circadian manner within the SCN, but it is responsible for maintaining daytime resting membrane potential of these neurons even in the absence of a nonphotic signaling cue. Mice lacking the GIRK2 subunit have altered circadian entrainment, corresponding to decreased neuropeptide Y (NPY, a nonphotic neurotransmitter) signaling within SCN neurons. Loss of GIRK channel signaling also inhibits exogenous melatonin-induced phase-shifting of behavioral rhythms and decreased firing of SCN neurons, indicating that GIRK channels are necessary for two different nonphotic neurotransmitters within the SCN - NPY and melatonin. Finally, activation of GIRK channels is sufficient to mimic a nonphotic phase shift, indicating that GIRK channel activation may be a conserved response within the SCN to nonphotic stimuli. This dissertation is the first to demonstrate a direct link between nonphotic neurotransmitters, a specific ion channel at the membrane, and subsequent regulation of circadian rhythm timing both within the SCN and in animal locomotor behavior.

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INTRODUCTION

Circadian rhythms: an overview

We live in a complex environment. The sun drives critical dynamics such as day length, seasonality, temperature climates, resource competition, predator/prey dynamics, and ultimately, population growth. Most organisms on earth have developed 24-hour cycles in biological and behavioral processes that enable them to synchronize and predict daily changes in the environment. These cycles are called circadian rhythms. These rhythms include processes such as sleep/wake cycles, hormonal rhythms like cortisol and melatonin secretion, cognition, immune system function, and basic cellular metabolism (1, 2). Disruption of circadian rhythms has been linked to multiple diseases, such as depression (3, 4), epilepsy (5-7), cardiac arrhythmias (8-11), and many more.

At a cellular level, rhythms are controlled by a transcription/translation feedback loop commonly referred to as the "molecular clock." In brief, two basic helix-loop-helix transcription factors: circadian locomotor output cycles kaput (CLOCK) and brainderived muscle arnt-like factor (BMAL) bind together and activate the promotor region of period (PER) and cryptochrome (CRY). PER and CRY then are produced, posttranslationally modified, and are translocated back into the nucleus to inhibit the binding of the CLOCK/BMAL complex to their own promotors, creating a negative feedback loop and decreasing PER/CRY production (Figure 1). This feedback loop has multiple interconnecting circuits and influences rhythmic expression of clock-controlled genes governing a variety of cellular processes (12-14).

In mammals, circadian rhythms are orchestrated by the suprachiasmatic nucleus (SCN) of the hypothalamus within the brain. SCN lesions in several rodent models leads

to arrhythmic locomotor activity (15). Human patients with cancer-induced lesions of the SCN exhibit arrhythmic body temperature and disrupted sleep/wake rhythms (16, 17). Not only is the SCN necessary for rhythm maintenance, but several transplant studies have shown that implantation of a functioning SCN to the third ventricle in SCN-ablated rodents restores rhythmicity and is sufficient to determine free-running period (the intrinsic day-length of an organism in constant conditions without environmental resetting cues like light) (18, 19).

The SCN is unique in that each individual neuron regulates its excitability in a self-sustaining, oscillatory manner independently from other neurons in this area (20). During the day, action potential firing rates are higher within SCN neurons, corresponding to more depolarized resting membrane potentials. Conversely, at night these neurons are relatively silent and exhibit more hyperpolarized resting membrane potentials. These tightly regulated changes in electrical activity are due to highly coordinated expression and regulation of ion channels (21, 22). For example, slowinactivating sodium channels regulate the initiation of SCN action potentials (23). Voltage-gated calcium channels co-modulate the initiation of the action potential (24), are sufficient to drive oscillations at more depolarized membrane potential (24, 25), and may play a key role in linking membrane excitability to intracellular signaling and the molecular clock (26-28). A wide variety of potassium channels modulate the firing rate differentially between the day and night (29-36). For example, large conductance potassium (BK) channels modulate spike timing by increasing the amplitude of the action potential after-hyperpolarization at night (33, 34). Finally, potassium leak channels may play a role in governing resting membrane potential (22, 37, 38). It is important to note

that although these ion channels have been implicated in regulating SCN excitability, the SCN is a heterologous structure (39, 40). Indeed, studies of dissociated SCN neurons show that while most individual neurons exhibit rhythmic firing, the amplitude of these rhythms differs greatly among the population, indicating that ion channel regulation may differ throughout the SCN (41, 42).

Ultimately, these neurons are synchronized within the SCN by a variety of factors including vasoactive intestinal peptide (43-45) to coordinate and convey time-of-day information. From there, the SCN projects directly to multiple areas of the brain including the amygdala, arcuate nucleus, and paraventricular nucleus, regulating multiple processes throughout the body (46).

Although surprising, the SCN keeps the same rhythmicity (high firing during the day, low firing at night) regardless of the temporal niche of the animal, indicating that diurnality or nocturnality is downstream of the core circadian oscillator (47). Previous reports comparing laboratory rats to diurnal grass rat (Arvicanthis niloticus) have identified the lateral subparaventricular zone (SPZ) as a potential regulator of diurnality. These studies show day-active grass rats have synchronized expression levels for c-Fos protein that are high during the day and low at night in both SCN and SPZ, whereas night-active laboratory rats exhibit c-Fos expression that is out of phase between the SCN and SPZ (48-50). Indeed, data from our lab shows that exclusively diurnal tree shrews have higher firing rates in both the SCN and SPZ during the day, corresponding with significantly higher cFOS positive neurons in the day compared to the night (Figure 2).

A key feature of the circadian clock is their ability to adapt and predict what behavioral patterns are necessary for day-to-day survival in the organism. The way

circadian rhythms entrain (synchronize) to environmental stimuli is by shifting the phase of the rhythm, or altering the time of sleep-wake behavior the next day so that it starts earlier (phase advance) or later (phase delay) depending on the timing of the input. Traditionally, entrainment is broken into two categories: photic (light-driven) and nonphotic (exercise, stress, arousal, etc.,) (1, 47, 51). Each cue has a different efficacy over the course of the day. For example, light exposure in the early night phase delays rhythms, shifting the phase to a later time on subsequent cycles, whereas light exposure in the late night phase advances rhythms so that the cycle begins earlier on subsequent days. The magnitude of these shifts can be plotted over time, defining a phase response curve (PRC) (52).

Nonphotic entrainment

The first "nonphotic" phase response curve was developed in response to novel wheel running in Syrian hamsters (Figure 3). These animals were restricted to a closed wheel for 2 hours at different times throughout the subjective day in constant darkness. The results showed that animals advanced their activity rhythms after running in the wheel during the middle of the day and delayed activity rhythms after wheel running during the late night, with the magnitudes of phase shifts directly proportional to the amount of wheel running (53). In a separate experiment, subcutaneous injections of saline induce phase advances during the day with minimal effect at night (Figure 3) (54, 55). These "nonphotic" effects are very different from those induced by light, indicating there are alternative entrainment pathways within the SCN.

Neuropeptide Y as a nonphotic entrainment neurotransmitter

The SCN receives light information from the retina through two different pathways. The direct pathway is through the retinohypothalamic tract, which travels from the retina, and directly innervates the SCN. The indirect pathway leads from the retina to the intergeniculate leaflet (IGL) of the thalamus, which extends neuropeptide Y (NPY) positive projections to the SCN (56, 57). Initially, the IGL was thought to only convey light information, but later studies showed that ablation of the IGL lengthens the freerunning period of rodents and eliminates the effects of novel wheel on circadian phase, yet the animals have no difficulty entraining to light (58-61). These experiments demonstrated that the IGL projection to the SCN is not necessary for photic entrainment, though NPY does inhibit the effects of a light pulse in the early night (62-65), indicating that the IGL regulates some forms of photic signaling. Several experimental findings suggest that NPY release from the IGL is primarily involved in nonphotic neurotransmission to the SCN. First, wheel running activates the IGL, as measured by cFOS expression, yet not the SCN (55, 66). In addition, NPY injected into the third ventricle of hamsters produces a similar phase response curve as wheel running in rodents (67, 68). Also, when antisera to NPY is injected to the third ventricle, the effects of novel wheel are eliminated (69). Daytime microinjections to the SCN with NPY receptor Y_1 and Y_2 agonists induces phase advances of behavior in hamsters, with the Y_2 agonist-induced shifts greater than the Y_1 agonists (70). Finally, NPY knockout mice fail to alter free-running period in response to prolonged wheel-running activity (71).

Serotonin as a nonphotic entrainment neurotransmitter

The raphe nucleus is a cluster of serotonergic cells that projects directly to the SCN (72, 73). This area is mostly active when the animal is moving and awake. Initial studies depleting rats and hamsters of 5-HT positive neurons with 5,7dihydroxytryptamine showed that animals were still rhythmic, but had an advanced phase angle in a light/dark (LD) cycle (activity onset started earlier with respect to lights-off than control animals) (74) as well as lengthened free-running period in constant light conditions (75). These studies demonstrate that serotonergic projections regulate circadian activity, but are not required to maintain an endogenous rhythm. It is important to note that ablation of serotonergic neurons does not eliminate the phase advancing effects of a novel wheel in hamsters (76), nor do serotonin agonists influence the magnitude of activity-induced phase shifts (77), which is why serotonin is thought to play a secondary role in the nonphotic pathways, such as the development of circadian rhythms (78-80) and mediating arousal and mood cues (81, 82) to the SCN. However, serotonin injections into the third ventricle do induce phase advances similar to that of NPY (83). Also, application of 5-HT and 5-HT1 receptor agonists phase advance SCN firing rhythms during the day (84, 85), with delays to no effect at night, similar to NPY. Thus, serotonin has a traditional nonphotic phase response curve (Figure 3).

Melatonin as a nonphotic entrainment neurotransmitter

Unlike NPY and serotonin, the pineal gland, the primary source of melatonin in the mammalian brain, does not directly innervate the SCN. Instead, the pineal gland is regulated through a multi-synaptic pathway (from the SCN to the paraventricular nucleus, the intermediolateral cell column of the thoracic spinal cord, the superior cervical ganglion, which then innervates the pineal gland) (86), and regulates endogenous release of melatonin during the dark phase of the animal. As such, endogenous melatonin is not a typical nonphotic stimuli, but instead provides the organism with photoperiodic information encoding day length and acts as an anti-arousal cue (86-91). This being said, melatonin supplementation is an effective phase-shifting agent in humans and animal models. In rodents, melatonin application during the day phase advances both freerunning locomotor behavior and SCN firing rate (92-95). The phase response curve is similar to melatonin administration in humans, with an advancing effect in the late day, and a delay during the night (96-98), indicating that melatonin, when given at different circadian phases, acts as a nonphotic stimuli (Figure 3).

A conserved nonphotic pathway?

As indicated above, NPY, serotonin, and melatonin all have a similar nonphotic phase response curve, phase advancing rhythms during the day with a minimal delay zone at night (Figure 3). Another facet of serotonin and NPY signaling is that they can block the phase advancing effects of light (62, 63, 99-102). Not only are the PRCs and ability to modulate light similar, but when applied to the SCN, both NPY and melatonin hyperpolarize the membrane of individual SCN neurons and reduce firing rate within the SCN (103-106). Finally, based on several studies applying agonists to the G proteincoupled receptors for NPY, melatonin, and 5-HT, the main response from these neurotransmitters is conveyed by G_{i/o} heterotrimeric G protein signaling (76, 94, 107, 108). Indeed, the effects of melatonin within the SCN are pertussis toxin sensitive (109, 110). These attributes (PRCs, interaction with photic stimuli, influence on SCN neurophysiology, and reliance upon G protein signaling) suggest there is a conserved response to nonphotic stimuli within the SCN to entrain circadian rhythms in a similar manner.

GIRK channels as a nonphotic convergence point

In other areas of the brain, NPY and serotonin have been shown to decrease neuronal excitability by activating G protein-coupled inwardly-rectifying potassium (GIRK) channels (111-114). Also, NPY, melatonin, and serotonin receptors are coupled with pertussis toxin-sensitive G proteins (110, 115-117), which can activate GIRK channels. GIRK channels are inwardly-rectifying and can be identified as an inward potassium current at greatly hyperpolarized potentials (beyond physiological resting membrane potential) (118). At more depolarized potentials in the physiological range of most neurons, GIRK channels pass an outward potassium current, causing membrane hyperpolarization and decreased neuronal excitability (118) similar to the effects of NPY, serotonin, and melatonin. Thus, GIRK channel activation is an ideal candidate mediator of nonphotic cues within the SCN. Additionally, GIRK channels may play a role in modulating day-night differences in SCN excitability, influencing photic entrainment as well.

GIRK channel physiology

GIRK channels are tetrameric channels that can be composed of four different subunits, GIRK1-4 (119). GIRK1 and GIRK2 are found primarily postsynaptically, and GIRK3 is normally found presynaptically. GIRK1-3 are present at similar levels

throughout the brain, and increase with age (120). The GIRK1 subunit lacks a membrane targeting motif and thus requires another subunit as a binding partner to be expressed, but increases single channel conductance and opening probability of the channel (118, 121). Although most research has focused on the postsynaptic mechanisms of regulation by GIRK1 and GIRK2, mRNA for all four GIRK subunits have been found in the supraoptic nucleus (122), and GIRK4 has been found in the arcuate nucleus and in lower levels throughout the brain (123, 124), indicating that future research should include GIRK3 and GIRK4.

GIRK channels are coupled to and activated by the $\beta\gamma$ subunits of Gi/o heterotrimeric G proteins (118). These channels allow large inward potassium currents at negative potentials, with little outward current at more positive potentials. This inward rectification is caused by a cation block in the pore region of each GIRK subunit, where magnesium or large organic cations, like spermine or putrescine, bind and prevent outward current flow (125). The cation block in GIRK channels is not as efficient as other inward-rectifiers, and thus they are categorized as "moderate rectifiers (119)."

GIRK channels regulate intrinsic membrane properties and general firing characteristics of neurons throughout the brain. Within CA1, adenosine receptor A1coupled GIRK channels hyperpolarize resting membrane potential by ~8 pA in dorsal neurons, with little to no effect on membrane properties of ventral neurons (126). Serotonin-induced depolarization of neurons in the lateral amygdala requires both TRPC activation and GIRK channel inhibition (127). In addition, the endogenous opioid receptor ligand dynorphan inhibits proopiomelanocortin cells within the arcuate nucleus by GIRK channel activation (128). These modulatory effects of GIRK channels on firing

rate and resting membrane potential can be modulated by different firing patterns of neurons within the ventral-tegmental area (129). Thus, GIRK channels can have cell autonomous, neuron-to-neuron, and network regulatory roles (118). To date, little is known about whether GIRK channels are expressed in the SCN and how/if they regulate neuronal firing.

Hypothesis and major objectives

Based on what is known about nonphotic signaling within the SCN and GIRK channel physiology, the main hypothesis of this dissertation is: **nonphotic stimuli activate GIRK channels which decrease neuronal excitability, modulating the timing of circadian rhythms.** Three main questions will be discussed in the subsequent two chapters including: 1) are GIRK channels present within the SCN, and if so are they regulated in either a light-driven or circadian manner, 2) how does endogenous GIRK channel activation alter SCN excitability (such as spontaneous firing rate, and intrinsic membrane properties such as input resistance and resting membrane potential of individual SCN neurons), 3) can GIRK channels mediate multiple nonphotic cues, specifically NPY and melatonin (Figure 4), and finally, 4) is GIRK channel activation sufficient to cause a nonphotic phase shift. These questions will be answered with a variety of techniques including western blotting protein analysis, whole cell patch clamp electrophysiology, real time bioluminescence monitoring of the molecular clock, and circadian behavioral analysis.

This dissertation is broken into two chapters, the first investigating the role of GIRK channels in SCN physiology, and ultimately nonphotic entrainment. The second

chapter investigates how GIRK channels mediate the phase shifting and neuronal silencing effects of G protein-coupled melatonin signaling within the SCN. Ultimately, the goal of this dissertation is to have a better understanding of the fundamental mechanisms behind nonphotic entrainment, and how regulation of excitability within the SCN influences circadian phase of the SCN and the animal.



<u>Figure 1: A basic schematic of the molecular clock.</u> This diagram includes the negative feedback loop of Per and Cry inhibiting their own transcription, and the positive feedback loop of Rora and Rev-erba directly modulating BMAL expression.



Figure 2: The SCN and SPZ exhibit increased spontaneous firing rate and cFOS positive cells during the day in the diurnal tree shrew. Top: Spontaneous firing rates (mean \pm SEM) for SCN and SPZ neurons during the day and night (left). Representative loose-patch traces (right, 5s). *p < 0.05, n = 3 slices, >75 cells/group. Bottom: Number of cFOS positive cells (mean \pm SEM) for SCN and SPZ regions of interest during the day and night (left). Representative IHC during the day and night (right). *p < 0.05, n = at least 2 animals, \geq 3 slices/group.



Figure 3: Cumulative nonphotic phase response curves. Phase response curves to 2hr novel wheel (53), sub cutaneous saline injections (54), microinjections of NPY into the third ventricle (67), I.P. injections of melatonin (dotted line indicates predicted curve based on 2 time points) (130), and microinjections of serotonin agonists (83) were given to hamsters in constant darkness at different CTs. Notice most stimuli induce phase advances during the subjective day with minimal delays during subjective night.



Figure 4: A Model of GIRK mediated nonphotic entrainment. During the day both nonphotic stimuli-induced activation of the IGL releases NPY onto SCN neurons, and exogenous application of melatonin activates Gi/o coupled GPCRs, opening GIRK channels. Efflux of K+ ions hyperpolarizes the membrane and decreases firing rate of action potentials. This causes a phase advance of the molecular clock and wheel running activity.

SUPRACHIASMATIC NUCLEUS FUNCTION AND CIRCADIAN ENTRAINMENT ARE MODULATED BY G-PROTEIN COUPLED INWARDLY-RECTIFYING POTASSIUM (GIRK) CHANNELS

by

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CHAPTER 1

SUPRACHIASMATIC NUCLEUS FUNCTION AND CIRCADIAN ENTRAINMENT ARE MODULATED BY G-PROTEIN COUPLED INWARDLY-RECTIFYING POTASSIUM (GIRK) CHANNELS

Abstract

G protein signaling within the central circadian oscillator, the suprachiasmatic nucleus (SCN), is essential for conveying time-of-day information. We sought to determine whether G protein-coupled inwardly-rectifying potassium channels (GIRKs) modulate SCN physiology and circadian behavior. We show that GIRK current and GIRK2 protein expression are greater during the day. Pharmacological inhibition of GIRKs and genetic loss of GIRK2 depolarized the day-time resting membrane potential of SCN neurons compared to controls. Behaviorally, GIRK2 KO mice failed to shorten free running period in response to wheel access in constant darkness and entrained more rapidly to a 6-hr advance of a 12:12LD cycle than WT littermate controls. We next examined whether these effects were due to disrupted signaling of neuropeptide Y (NPY), which is known to mediate nonphotic phase shifts, attenuate photic phase shifts and activate GIRKs. Indeed, GIRK2 KO SCN slices had significantly fewer silent cells in response to NPY. This deficit likely contributed to the absence of NPY-induced phase-advances of *Per2^{Luc+/-}* rhythms in organotypic SCN cultures from GIRK2 KO mice. Finally, GIRK channel activation is sufficient to cause a nonphotic-like phase advance of PER2::LUC rhythms. These results suggest that rhythmic regulation of GIRK2 protein and channel

function in the SCN contributes to day-time resting membrane potential, providing a mechanism for the fine tuning responses to nonphotic and photic stimuli. Further investigation could provide insight into disorders with circadian disruption comorbidities such as epilepsy and addiction, in which GIRK channels have been implicated.

Introduction

The central circadian oscillator in the suprachiasmatic nucleus (SCN) of the hypothalamus contains ~20,000 coupled neurons (Welsh et al., 2010). A transcription/translation feedback loop called the "molecular clock" drives rhythmic gene expression in individual SCN neurons and regulates daily oscillations in action potential firing and excitability, with increased neuronal firing during the day and relative quiescence during the night (Kuhlman & McMahon, 2006). This day/night variation in electrical output from the SCN drives circadian rhythms in other brain areas and the body. The timing of these molecular and neurophysiological rhythms can be altered by environmental cues including photic (light) and nonphotic (stress, exercise, etc.) stimuli (Albrecht, 2012). These alterations or phase shifts occur daily, entraining the animal to its environment. Although much is known about the neurotransmitter systems that underlie these entraining pathways, the molecular mechanisms that couple receptor activation to the changes in SCN neuronal output that ultimately shift circadian phase and drive entrainment are not fully understood.

One mediator of intracellular phase shifting signals is G protein signaling, and many of these second messenger pathways are critical for maintaining SCN rhythmicity and enabling photic and nonphotic entrainment (Cheng et al., 2004; Aton et al., 2006;

Dahdal et al., 2010; Doi et al., 2011; Brancaccio et al., 2013). We hypothesize that this abundance of crucial G protein signaling in SCN neurons may activate G protein-coupled inwardly-rectifying potassium (GIRK) channels, which have been implicated in diseases such as epilepsy and addiction (Hibino et al., 2010; Luscher & Slesinger, 2010). Furthermore, GIRK channel activation within the SCN may alter neurophysiological function and the ability of environmental stimuli to reset circadian clock phase. We use a variety of electrophysiological, behavioral, and molecular assays to determine: when GIRK channel protein levels and activation are highest within the SCN, the effect of GIRK channel activation on SCN neuronal function, and the necessity of GIRK channels for circadian entrainment.

Methods

Ethical approval

All animal care, handling, and housing were in compliance with the University of Alabama at Birmingham's Institutional Animal Care and Use Committee guidelines.

Animals and housing

All mice in these experiments were 2-4 months of age to reduce developmental or aging phenotypes (Turek et al., 1995; Biello, 2009). Only male mice were used for western blotting and behavioral experiments (Ruiz de Elvira et al., 1992; Vyazovskiy et al., 2006). Three separate mouse lines were used: 1) C57/BL/6 mice (western blotting experiments); 2) GIRK2 knockout animals on a C57/BL6 background (Signorini et al., 1997) (electrophysiology and circadian behavior); and 3) Per2Luc+/- mice (Yoo et al.,

2004) on a C57/BL6 background crossed with the GIRK2 line for at least 2 generations before use in experiments (bioluminescence assays of molecular clock function). Separate cohorts of mice were used for each different type of experiment.

Western blotting

For light/dark (LD) experiments (Fig. 1AB), mice were group housed with food and water ad libitum in a 12:12 LD cycle. Mice were killed every 4 hours over a 24-hour period (Zeitgeber Time 0 or ZTO, defined as lights on). For constant darkness (DD) experiments (Fig. 1CD), mice were single-housed on running wheels in DD for two weeks. Animals were killed during either the subjective day at circadian time 4 (CT4; CT 12 defined activity onset) or subjective night (CT 16). Activity onset was predicted using linear regression analysis for the preceding seven activity onsets. For all experiments during the dark, mice were killed by cervical dislocation followed by enucleation with the aid of night vision goggles in order to prevent photic signaling during brain extraction. Hypothalamic slices (600 µm) were prepared using a vibroslicer (7000smz, Campden Instruments Ltd., Lafayette, IN), followed by dissection of the SCN under a Zeiss dissection microscope. Protein lysates were prepared, and immunoblotting for GIRK1 (1:500, Alamone Labs, Israel) and GIRK2 (1:750, Millipore, MA, USA) was performed by loading $10\mu g$ of protein per sample as analyzed by a BCA protein assay. Previous studies have shown that GIRK1 and GIRK2 both exhibit multiple bands in western blots within the brain. For GIRK1, the three bands indicate different glycosylation states (Koyrakh et al., 2005; Aguado et al., 2008). GIRK2 exhibits multiple splice variants (Lesage et al., 1995; Koyrakh et al., 2005; Aguado et al., 2008). For densitometry

analysis, all bands for each protein were quantified together for assessment of total protein levels, and the uppermost heavily glycosylated band of GIRK1 (75 kDa, Fig. 1) was analyzed for rhythmicity of GIRK1 glycosylation. For the LD analysis, each blot was normalized to the mean ZT5 time point as a positive control. β -Actin (1:40,000, Millipore, MA, USA) was used as a loading control.

Electrophysiology

Mice were killed either at ZT 2 or ZT10.5 (day and night recordings respectively) by cervical dislocation. For whole-cell electrophysiology, all recordings were made between projected ZT 3-9 (day) or ZT 12-17 (night). For loose-patch electrophysiology, all recordings were made between projected ZT 4-6 in the presence of 2.35 μ M NPY (American Peptide, Sunnyvale, CA) or vehicle (water). Brains were harvested, sectioned at 200 µm on a vibroslicer (7000smz, Campden Instruments Ltd., Lafayette, IN), and transferred to an open recording chamber (Warner Instruments, Hamden, CT) that was continuously perfused at a rate of 2.0 ml/min with extracellular solution consisting of (in mM) NaCl (124), NaHCO3 (20), Na2HPO4 (1), MgSO4 (1.3), glucose (10), KCl (3.5), CaCl2 (2.5; added the day of the experiment) with osmolality adjusted to 300-305 mOsm), bubbled with 5% CO2 / 95% O2, and heated to 34 ± 0.5 °C. Neurons were visualized with an Olympus BX51WI (Olympus America Inc., Center Valley, PA) using infrared-differential interference contrast optics. Electrodes with a pipette resistance of ~4-6 M Ω were filled with filtered, potassium gluconate solution consisting of (in mM): K-gluconate (135), KCl (10), HEPES (10), EGTA (0.5), then adjusted to pH 7.4 with KOH (as in Kuhlman & McMahon, 2004). Electrophysiological signals were processed

and controlled by a Multiclamp 700B amplifier, and pClamp 10.02 software (Axon Instruments, Union City, CA). Recordings were sampled at 20 kHz and filtered at 10 kHz. In order to block synaptic transmission (as in Fig. 2): bicuculline $(30\mu M)$ and CdCl2 (200µM) (Sigma-Aldritch, St. Louis, MO), D-AP5 (50µM) and CNQX (10µM) (Abcam, Cambridge, MA), and TTX (1μ M) (Tocris Biosciences, Minneanapolis, MN) were added to the bath solution. To isolate GIRK currents (Fig. 2) the concentration of KCl was increased from 3.5 mM to 30 mM in order to increase potassium conductance across the membrane (refer to (Fu et al., 2004)). The GIRK channel antagonist Tertiapin-Q (0.2 µM) (Alamone Labs, Israel) was used for experiments in Fig. 3. Input resistance and resting membrane potential were calculated as specified in (Kuhlman et al., 2003). Resting membrane potential, action potential amplitude, and firing rate were calculated from 30-s gap-free current clamp recordings. All cells included in these analyses had, in voltage-clamp mode, \leq 35 pA holding current to clamp membrane potential at -65 mV and an action potential amplitude of greater than 10pA during the current clamp step protocol. All data were collected within 6 min of membrane rupture to minimize any potential washout effects from the whole-cell recording (Schaap et al., 1999). Loose patch recordings were obtained in gap-free mode with an average seal resistance of 39 M Ω . Average spike rate was calculated from at least one minute of the two minute trace. Induced currents for ML297 and NPY application were determined by holding the cell at -80mV in gap-free voltage clamp mode in a high potassium (30mM) solution (as in (Hamasaki et al., 2013)). Either ML297 (10µM) or NPY (2.35µM) were bath applied for at least 1 minute. The change in current following drug application was compared to that of high potassium alone within the same cell. For all electrophysiological experiments, at least 3 biological replicates with at least 4 cells per animal were used, unless otherwise indicated. There was no specific regional bias when recording within the SCN.

Behavioral Analysis

All mice were housed in individual wheel cages. Wheel-running activity or general cage activity (via infrared motion-sensors from Spy Town, Melville, NY; as in Fig. 4AB) was recorded and analyzed using Clocklab software (Actimetrics, Wilmette, IL). For behavioral analysis, one WT animal was excluded due to low activity/equipment failure. Free-running period was measured by chi-square periodogram analysis of 7-10 days in constant dark conditions (Fig. 4). One KO mouse (out of 8) had a free running period of 23.3h, which was 6.5 SD's from the mean and was therefore excluded from behavioral analysis. Activity profiles from mice housed on wheels in constant darkness were acquired using Clocklab software. Each profile was aligned to CT 12, activity onset. Then, the average counts of WT and KO mice during the subjective day and subjective night were compared. Entrainment was defined as the first day when both the activity onset and alpha length (activity period) were within twenty minutes of those predicted 4 days prior to the light change. Clocklab software was used to determine activity onset. Offset was defined as the last activity bout where three out of six of the previous bouts were above 3 revolutions per minute. Alpha length was calculated by subtracting onset time from offset time.

Bioluminescence Assays

Organotypic SCN cultures from GIRK2 KO or WT Per2Luc+/- mice were prepared and treated within the first 7 days for one hour beginning at CT 3-5 (where CT 12 is defined as peak bioluminescence, drug timing specified for each experiment) with 2.35 μ M NPY (American Peptide, Sunnyvale, CA), 10 μ M ML297 (Days et al., 2010) or vehicle (either water or 0.02% DMSO respectively), using identical culture media and methods described in (Besing et al., 2012). Data were acquired and analyzed with Lumicycle Analysis software (Actimetrics, Inc, Wilmette, IL), and recordings with a goodness of fit greater than 85% were used for analysis. Phase shifts were calculated by comparing two predictions for the time of the first peak post-treatment: one prediction determined from at least three cycles before treatment and a second prediction determined from at least three cycles after treatment. The difference between these two predictions indicated the size of the phase shift.

Statistical Analysis

All statistical analysis was performed with PASW Statistics 18. GIRK channel protein expression was analyzed for rhythmicity using a cosinor analysis (as in (Bray et al., 2013)). For comparisons of means, an independent samples t-test or ANOVA was used for comparisons between two means or two or more means, respectively. Two factor designs were analyzed with a two-way ANOVA with repeated measures when appropriate. In cases of a non-normal distribution, a nonparametric Kruskal-Wallis test followed by median test post hoc analyses. Finally, to assess whether the number of silent cells changed in response to NPY treatment, a G likelihood-ratio test was used. Post hoc

analyses employed a Fisher's exact test, with a Bonferroni-corrected alpha of 0.025. For all other tests, significance was ascribed at P < 0.05.

Results

GIRK channel protein and function is regulated in a circadian manner within the SCN, influencing day-time neurophysiology.

To determine if GIRK1 and GIRK2 channel subunits were present within the SCN, and if levels of these proteins changed over the course of the day, we analyzed SCN samples from mice at 4-h time points across a 12:12 Light/Dark (LD) cycle. Using western blot analysis, GIRK1 and GIRK2 protein was present within the SCN, and GIRK2 (but not GIRK1) expression patterns significantly fit a 24-h rhythm (cosinor analysis; GIRK1: $R^2 = 0.114$, P > 0.05; GIRK2: R2 = 0.184, P < 0.05; n = 3-5/time point) with peak GIRK2 protein levels occurring at ZT 6.47 \pm 0.3h (amplitude, -0.25 \pm 0.1; mesor, 0.59 ± 0.1 ; Fig. 1AB). Because GIRK1 shows variation in the heavily glycosylated state, the glycosylated state was analyzed for rhythmicity and showed diurnal variation but failed to reach statistical significance ($R^2 = 0.195$, P = 0.06). Although GIRK2 expression exhibited a 24-h rhythm in LD, it was necessary to examine protein levels in the SCN from animals housed in constant darkness (DD) to assess whether this rhythm was endogenously generated rather than driven by the light cycle. GIRK2 protein expression from animals housed for at least 14 days in DD had significantly higher protein levels during the subjective day (CT4 based on activity onset) compared to night (CT 16; independent samples t-test, t(5) = 2.83, P< 0.05; n= 3-4/time

point). As expected, levels of both the heavily glycosylated form and total GIRK1 did not vary between time points (Fig. 1CD).

In order to determine whether higher levels of GIRK2 protein during the day could contribute to a change in basal GIRK activation over the course of the day, we used whole-cell, voltage-clamp electrophysiology and pharmacological inhibition of synaptic transmission (TTX (1µM), bicuculline (30µM), D-AP5 (50µM), CNQX (10µM), and CdCl2 (200µM)) along with increased KCl (30mM) for increased potassium conductance, in order to isolate currents from SCN neurons from wild-type and GIRK2 knockout animals in response to a slow ramp (2.5 s) from -140 mV to -20 mV. Peak inward current was significantly greater during the day in WT neurons (-88.1 ± 5.7 pA) compared to night (-61.3 ± 4.6 pA), and this difference was not seen in KO neurons (day: -64.4 ± 5.4pA; night:-67.9 ± 6.3pA) (two-way ANOVA, genotype by time interaction: F(1,122) = 6.53, P < 0.01; Tukey HSD post hoc comparison, P < 0.05 Fig. 2; n= ≥ 20 cells/group), showing that day-night differences in this current is specific to increased levels of GIRK2 (Fig. 2).

GIRK channels are known to decrease cellular excitability by hyperpolarizing the membrane (Luscher & Slesinger, 2010). To determine whether loss of GIRK2 plays a role in regulating SCN neuronal membrane properties, we measured resting membrane potential (RMP) of WT and KO SCN neurons during the day and night (n = \geq 25 cells/group). Consistent with previous results (Kuhlman & McMahon, 2004), there was an overall day/night difference in RMP (Kruskal Wallis test, H(3) = 37.403, P < 0.01; median post hoc test, P < 0.05 for both WT day compared to night, and KO day compared to night; Fig. 3AB). As predicted, KO neurons were significantly more
depolarized than WT neurons during the day only (WT day: -41.6 \pm 1.0 mV; KO day: -36.5 \pm 1.1 mV; median post hoc test, P < 0.05). To eliminate the possibility that the depolarized RMP in KO mice was driven by altered regulation of ion channel expression to compensate for loss of GIRK2, we applied a GIRK channel antagonist, Tertiapin-Q (TPQ) (0.2 μ M), to WT slices during the day (Fig. 3B). We found that TPQ caused a similar magnitude of depolarization (-36.9 \pm 1.0 mV, n = 38 cells), demonstrating that GIRK channels are necessary for maintaining normal day-time resting membrane potential. There were no significant differences in either input resistance (Kruskal Wallis test, H(3) = 2.662, P > 0.05), action potential firing rate (Kruskal Wallis test, H(3) = 4.879, P > 0.05), or peak current amplitude of action potential firing (Kruskal Wallis test, H(3) = 2.744, P > 0.05) among groups (Fig. 3CDE), indicating that GIRK channels primarily regulate resting membrane potential.

Loss of GIRK2 alters the behavioral response to nonphotic and photic cues.

After defining the temporal pattern of GIRK subunit expression and describing a role for GIRK channels in setting day-time resting membrane potential, we next evaluated the functional role of GIRK2 in the entrainment of behavioral locomotor rhythms. There were no behavioral differences between heterozygous and WT animals (data not shown), and therefore, the following results are from WT-KO comparisons only. First, we examined general cage activity of mice single-housed with the running wheel locked because several studies have shown that wheel running (a nonphotic entraining stimulus) shortens the period of behavioral locomotor rhythms in DD (Edgar et al., 1991a; Edgar et al., 1991b; Kuroda et al., 1997; Deboer & Tobler, 2000; Harrington

et al., 2007). Activity monitored with infrared motion sensors revealed no significant differences between genotypes in terms of average activity counts or circadian rhythmic amplitude in LD (Fig. 4AB; Table 1). In addition, WT and KO mice had very similar period lengths (or tau) when placed into DD (Table 1, n = 15-16/group). To test whether loss of GIRK2 alters the period shortening response to the nonphotic stimulus of wheel running (Harrington et al., 2007), we placed WT and KO animals on running wheels and assessed tau in DD (Fig. 4BC). In an LD cycle, there was no difference between KO and WT average activity counts (Table 1). In DD, KO animals had higher daily activity in general (Table 1); however, this effect was mostly driven by increased activity of KO mice during the subjective night (mean \pm SEM counts, WT night: 2,867.5 \pm 303.0, KO night: $3,996.6 \pm 252.7$) but not during the day (mean \pm SEM counts, WT day: $287.2 \pm$ 37.9, KO day: 296.0 \pm 60.5) consistent with reports of hyperactivity in these mice (Blednov et al., 2001). A repeated measures ANOVA revealed a significant time-of-day by genotype interaction (F(2,21) = 4.906, P < 0.05) with KO night activity significantly greater than WT night activity (Tukey HSD post hoc comparison, P < 0.05; Fig. 4BC). In contrast to general cage activity, KO mice with access to a running wheel did not exhibit the shorter free running period observed in WT mice (mean \pm SEM, WT: 23.80 \pm 0.04 h, KO: 23.95 ± 0.04 h; t(13) = 2.46, P < 0.05, n=7-8/group; $\eta^2 = 0.32$) indicating that GIRK2 is necessary for proper nonphotic signaling in response to wheel running.

Because the behavioral response to nonphotic cues was altered upon loss of GIRK2, we examined whether GIRK2 contributes to photic entrainment as well. Mice housed in a 12:12 LD cycle were subjected to a 6-h phase advance of the LD cycle (Fig. 5), and the number of days required for re-entrainment was determined. Entrainment was

defined as the first day in which the activity length returned to the same length observed before the light shift (see Materials and Methods). Mice lacking GIRK2 required approximately half the number of days to re-entrain to the new light cycle compared to WT (mean \pm SEM days to entrain, WT: 7.2 \pm 0.5, KO: 3.5 \pm 0.8; t(10) = 4.07, P < 0.01; n = 6/group). However, KO mice entrained to a 6-h phase delay at the same rate as WT (mean \pm SEM days to entrain, WT: 4.7 \pm 0.7, KO: 4.7 \pm 0.5; t(10)= -0.23, P > 0.05; n = 6/group). These results indicate that GIRK2 signaling slows the rate of re-entrainment to photic phase advances, but not delays.

Neuropeptide Y signaling in the SCN requires GIRK2.

In rodents, disruption of the nonphotic neurotransmitter NPY abolishes the period shortening effect of wheel access (Pickard et al., 1987; Pickard, 1994; Kuroda et al., 1997; Lewandowski & Usarek, 2002; Harrington et al., 2007). Because NPY signaling also antagonizes phase advances to light (Yannielli & Harrington, 2000; Lall & Biello, 2003; Yannielli et al., 2004), disruption of the NPY-dependent nonphotic signaling pathway may explain the loss of wheel running-induced period shortening and the enhanced photic entrainment of KO mice. Specifically, we hypothesized that NPY-induced phase shifts within the SCN requires GIRK2 activation because application of NPY to the SCN in vivo or in vitro during the day induces phase advances in behavior and neuronal activity rhythms (Yannielli & Harrington, 2000; Maywood et al., 2002; Lall & Biello, 2003; Yannielli et al., 2004; Besing et al., 2012) and NPY-induced effects in other brain regions require GIRK activation (Paredes et al., 2003; Fu et al., 2004).

First, we investigated whether the suppressive effects of NPY on SCN neuron firing rate (van den Pol et al., 1996; Besing et al., 2012) were lost in GIRK2 KO mice using loose patch electrophysiology with and without 2.35 μ M NPY in the bath (n = \geq 45 cells/group). Spontaneous firing rate was significantly different among the four groups (Kruskal Wallis test, H(3) = 79.271, P < 0.01). Surprisingly, firing rates of WT and KO neurons did not differ (median post hoc test, P > 0.05; Fig. 6A); however, KO neurons exhibited significantly higher firing rates than WT neurons in response to NPY (mean \pm SEM, WT: 0.8 ± 0.2 Hz, KO: 2.2 ± 0.3 Hz; median post hoc test, P < 0.01; Fig. 6AB). Contingency analysis revealed a significant effect of genotype and treatment on the percentage of silent cells ($\chi^2(3) = 53.193$, P < 0.01). Specifically, in vehicle-treated slices, the number of silent cells was not significantly different between KO and WT (Fisher's exact test, P > 0.05); however, WT slices treated with NPY had significantly more silent cells than did NPY-treated KO slices (Fisher's exact test, P < 0.01; Fig. 6BC), suggesting that NPY failed to silent many of the SCN neurons in absence of GIRK2. In order to determine whether GIRK channels directly mediate NPY-induced current, gap-free, whole-cell voltage clamp was used to hold the cells at -80mV in high potassium (30mM) in order to readily measure the GIRK-mediated current (as in (Hamasaki et al., 2013)). Upon application of NPY (2.35 μ M), 4 out of 5 cells responded to NPY with inward current (range, 15 - 119 pA; mean \pm SEM, 44.5 \pm 23.2 pA , n = 4 cells from two animals). Of these, 3 out of 4 cells had reduced inward current by $\sim 50\%$ in response to TPQ (0.2 μ M). TPQ alone had a net inward current of 3.1 ± 4.4 pA (n = 4 cells from two animals). These results indicate that GIRK channels mediate at least part of the NPYinduced current within SCN neurons, consistent with the partial effect of NPY in the

GIRK2 KO animals (Fig. 6). This failure to significantly reduce excitability in GIRK2 KO animals may impede NPY-dependent phase shifts to the circadian clock.

To test this hypothesis, we crossed GIRK2 WT and KO animals onto the Per2Luc+/- reporter line (see Materials and Methods) and measured the phase shifting effect of NPY on the molecular clock. Because these animals showed no difference in behavioral circadian rhythmicity (Fig. 4), it was not surprising that the PER2::LUC rhythms showed no difference in period between genotypes pre-treatment (mean period \pm SEM, WT: 24.42 \pm 0.12h, KO: 24.38 \pm 0.16h; t(28) = 0.99, P > 0.05). As has been previously published by Besing et al. (2012) and more recently by Belle et al. (2014), 1-h treatment with 2.35 μ M NPY during the early day (CT 4-5) produced ~3-h phase advances in PER2::LUC rhythms in WT mice; however, phase advances in KO animals were reduced to the level of controls (two-way ANOVA, genotype by treatment interaction: F(1,26) = 4.77, P < 0.05; Tukey HSD post hoc, P < 0.05, Fig. 7; n= 6-8 cultures/group). Thus, these results suggest that GIRK2 is necessary for NPY-induced phase advances within the SCN, and that loss of NPY signaling may be an underlying cause for the circadian entrainment alterations observed in GIRK2 KO mice.

GIRK channel activation is sufficient to cause a nonphotic-like phase shift.

We have shown that GIRK channels partially mediate NPY-induced silencing of SCN neurons as well as NPY-induced phase advances of the molecular clock. However, GIRK channels can be fully-opened by a variety of neurotransmitter signaling (Luscher & Slesinger, 2010), including other nonphotic signals. To test the hypothesis that GIRK channel activation was sufficient to mimic nonphotic signals, we applied 10µM ML297, a GIRK channel agonist (Days et al., 2010; Kaufmann et al., 2013), for one hour to PER2::LUC SCN cultures starting between CT3-4. This concentration was sufficient to induce GIRK currents in SCN neurons (-60.3 \pm 9.7, n= 3 cells), consistent with recent papers demonstrating ML297 specificity in cultured hippocampal neurons (Wydeven et al., 2014). Activation of GIRK channels with ML297 significantly phase advanced PER2::LUC rhythms compared to vehicle-treated controls (mean \pm SEM, ML297: 3.6 \pm 1.1, vehicle: 0.3 ± 0.2 ; t(5.23) = -3.10, P < 0.05; Fig. 8; n= 6 cultures/group), suggesting a broader role for GIRK channels in mediating nonphotic signals.

Discussion

Although G protein signaling is critical for circadian rhythmicity (Cheng et al., 2004; Aton et al., 2006; Zuberi et al., 2008; Doi et al., 2011; Brancaccio et al., 2013), the role of G protein-coupled potassium channels, and specifically GIRK channels, in modulating time-of-day cues has not been studied. This paper is the first to show that a single, circadian-regulated GIRK channel subunit can modulate SCN neurophysiology, mediate the effects of daytime NPY, and alter behavior in response to both photic and nonphotic cues. Specifically, we found that GIRK2 protein and function exhibited an endogenous rhythm within the SCN with a peak around midday. This increase during the day was necessary for proper maintenance of day-time resting membrane potential within SCN neurons since GIRK channel inhibition or GIRK2 knockout further depolarized resting membrane potential. Furthermore, animals with genetic loss of GIRK2 showed altered entrainment properties, such that the GIRK2 KO animals (compared to WT mice) re-entrained to a 6-h phase advance of the light-dark cycle more rapidly and did not

exhibit period-shortening with wheel access. These behavioral phenotypes may be due, in part, to a loss of NPY signaling within the SCN as indicated by the reduced effects of NPY on firing rate of SCN neurons in GIRK2 KO animals as well as impaired NPYinduced phase advances in GIRK2 KO organotypic cultures of the SCN. Finally, GIRK channel activation was sufficient to cause a daytime phase advance in PER2::LUC rhythms. Taken together, the results of our study indicate a role for circadian regulation of GIRK channels in modulation of neurophysiological rhythms and establishing proper responses to time-of-day stimuli.

The circadian regulation of neural activity in SCN neurons has been welldocumented with peak excitation occurring during the day (Kuhlman & McMahon, 2006). GIRK channels are inwardly-rectifying and can be identified as an inward potassium current at greatly hyperpolarized potentials (beyond physiological resting membrane potential). At more depolarized potentials in the physiological range of most neurons, GIRK channels pass an outward potassium current, causing membrane hyperpolarization and decreased neuronal excitability (Luscher & Slesinger, 2010). In the present study, GIRK2 channel protein and current was highest during the day (Fig. 1 and 2). Therefore, we hypothesized that GIRK channel activation during the day may act as a stop-gate on excitability. Indeed, we found that loss of GIRK2 channels or pharmacological GIRK channel inhibition resulted in depolarization of the resting membrane potential by ~5 mV (Fig. 3). This basal GIRK-mediated hyperpolarization is consistent with the effects of GIRK channel activation by neurotransmitters such as NPY, serotonin, melatonin, glutamate, acetylcholine, and GABA in a variety of central and peripheral areas (Krapivinsky et al., 1995; Nelson et al., 1996; Luscher et al., 1997; Fu et

al., 2004; Acuna-Goycolea et al., 2005; Luscher & Slesinger, 2010). For example, GIRK channels have been shown to couple directly with GABAB receptors (Luscher et al., 1997; Arora et al., 2011), and GABA signaling within the SCN is dependent upon Gi/o G proteins (Aton et al., 2006) which are necessary for GIRK channel opening.

The neurotransmitters listed above also play key roles in regulating the timing of circadian rhythms (Albrecht, 2012). Underlying mechanisms of nonphotic entrainment, or synchronization of rhythms to cues like exercise or stress, remain largely understudied even though they block the effects of light and reset clock phase both in rodents and humans (Hastings et al., 1998; Mistlberger & Skene, 2005). For example, both melatonin and NPY induce large phase advances of locomotor behavior and spike rate rhythms during the day and hyperpolarize the resting membrane potential in a potassium-sensitive manner (Jiang et al., 1995; Hall et al., 1999; Scott et al., 2010). Our results showed that GIRK2 activation was necessary and sufficient to induce these large nonphotic-like, phase advances of the molecular clock (Fig. 6 and 8). It is interesting to note that both NPY and melatonin hyperpolarize the membrane and reduce spontaneous firing rate of SCN neurons when treated during the subjective day (Jiang et al., 1995; Hall et al., 1999; Scott et al., 2010). Based on these observations, it can be speculated that these channels mediate multiple phase-resetting signals and may act as a convergence point for nonphotic signaling, resulting in similar phase response curves for serotonin, NPY, and melatonin (Yannielli & Harrington, 2004). An important function of increased GIRK current during the day could be to allow for daytime environmental cues to phase shift the SCN through hyperpolarization. This hypothesis is supported by the fact that activation of GIRK channels with ML297 during the early day is sufficient to cause a

phase-advance of molecular clock rhythms (Fig. 8). However, GIRK channel activation does not preclude possible co-activation of second messenger signaling cascades by nonphotic cues such as PKC activation in response to NPY (Biello et al., 1997). Indeed, NPY still has a partial effect on spike rate in GIRK2 KO animals (Fig. 6). Photic and nonphotic entraining stimuli can interact and in general, are mutually inhibitory. For example, NPY injection into the SCN region of light-exposed hamsters in the late night attenuates the characteristically ensuing phase advance and up-regulation of clock gene expression (Yannielli et al., 2004; Gamble et al., 2006). Conversely, NPY blockade of NPY Y5 receptors enhance light-induced phase advances (Yannielli et al., 2004). Similar experiments have been done with light mimicked by the glutamate receptor agonist NMDA, such that phase advances in SCN firing or wheel running behavior were diminished by NPY and NPY antagonists (Gamble et al., 2004; Soscia & Harrington, 2004; Yannielli & Harrington, 2004; Soscia & Harrington, 2005). In addition to acute phase shifts, NPY is also critical for other nonphotic behavioral effects, such as period shortening in response to wheel running. Elimination of the primary source of NPY to the SCN through lesions of the intergeniculate leaflet blocks period shortening induced by wheel access (Pickard et al., 1987; Pickard, 1994; Kuroda et al., 1997; Lewandowski & Usarek, 2002). This result is consistent with more recent evidence that NPY-deficient mice also fail to shorten free running period in response to wheel access (Harrington et al., 2007). The present data suggests that GIRK channel activation may be involved in these phenotypes. Specifically, loss of GIRK2 channels resulted in a lack of period shortening typically observed in response to nonphotic wheel access (Fig. 5) as well as enhanced phase advances in the light-dark cycle (Fig. 4). It appears that in

absence of GIRK signaling, entrainment to photic phase advances are functionally enhanced, while the nonphotic pathway is suppressed. Because there was no change in entrainment to photic delays, future studies should determine whether this photic phenotype is a broad strengthening of photic entrainment pathway, or simply a removal of nonphotic inhibition to photic phase advances in behavior. In addition, future studies should examine the role of GIRK in retinal ganglion cells and within IGL given its widespread expression in neurons (Signorini et al., 1997; Luscher & Slesinger, 2010). A better understanding of the nonphotic/photic interaction is an important area for future research given that an animal is unlikely to encounter a photic or nonphotic stimulus in isolation.

In the present study, we found that both GIRK2 protein expression and current amplitude were regulated over the day-night cycle. This result is consistent with the finding that GIRK2 mRNA expression in microarrays of SCN tissue is rhythmic (Pizarro et al., 2013), indicating that the gene transcribing GIRK2 mRNA (KCNJ6) may be under direct clock-control. In general, our understanding of the mechanisms underlying circadian regulation of ion channels is limited (Colwell, 2011), but some studies suggest epigenetic mechanisms or microRNA modifications may determine transcript stability (Wang, 2013), while other studies have shown that free radical homeostasis can also regulate ion channel gating (Wang et al., 2012). Determining clock control of ionic channel regulation in the SCN could provide insight into how ion channels are regulated in a circadian manner in other brain areas and tissues. GIRK channels have been implicated in regulating dopamine signaling within the VTA (Lomazzi et al., 2008; Arora et al., 2011). If GIRK channels are regulated in a time-of-day-sensitive manner in the

VTA (as in the SCN) then the present results may have broader implications for addiction and withdrawal and the circadian regulation of this disease.

GIRK channels have also been associated in diseases of hyper-excitability such as epilepsy (Loddenkemper et al., 2011; Zarowski et al., 2011), chronic atrial fibrillation (Capucci et al., 2012; Shusterman et al., 2012), and long QT syndrome (Jeyaraj et al., 2012; Takigawa et al., 2012). A key feature these disorders is that seizure or fibrillation onset is more likely to occur during the day in humans, suggesting that circadian control of GIRK channel function may underlie risk dependence on time-of-day. Indeed, microarray studies have shown that GIRK1 and GIRK4 transcripts are rhythmic in mouse whole-heart homogenate (Pizarro et al., 2013). Perhaps by understanding the interplay between GIRK channel dysfunction and circadian regulation of excitability throughout the body, novel time-of-day-sensitive therapeutics could be developed with fewer offtarget effects.

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		WT		KO		Independent samples t-test		
No Wheel		Avg	SE	Avg	SE	t	df	Sig. (2-tailed)
LD	Power	705.21	54.07	646.28	42.49	-0.86	24	0.400
	Avg Counts (counts/min)	3.02	0.26	2.56	0.18	-1.44	24	0.164
DD	Tau	23.92	0.04	23.85	0.04	-1.08	24	0.292
	Power	539.24	2.79	534.05	28.64	-0.15	24	0.885
	Avg Counts (counts/min)	2.91	0.20	2.87	0.20	-0.11	24	0.913
Wheel								
LD	Power	1099.63	59.83	1170.28	118.43	0.55	13	0.589
	Avg Counts (counts/min)	14.51	1.98	16.13	1.87	0.59	13	0.566
DD	Tau	23.80	0.04	23.95	0.04	2.46	13	0.029
	Power	1173.99	86.84	1430.68	86.55	2.08	13	0.058
	Avg Counts (counts/min)	11.85	1.65	18.72	1.95	0.76	13	0.018

Table 1. Circadian behavioral analysis of GIRK2 KO mice



Figure 1: GIRK2, but not GIRK1, protein levels are regulated in a circadian manner. A) Representative immunoblots for GIRK2 from mouse SCN out of an LD cycle (left). Relative optical density (mean \pm SEM) for GIRK2 throughout the day with lines indicating the predicted cosinor curve (*p < 0.05, right). n = 5-6 per time point. B) Representative immunoblot for GIRK1 in mouse SCN across an LD cycle (left). Relative optical density (mean \pm SEM) for GIRK1 at each time point throughout the day with predicted cosinor curve (right). n = 3-5 per time point. C) Representative GIRK1 (left) and GIRK2 (right) blots from mouse SCN out of DD at CT 4 (subjective day) and CT 16 (subjective night). D) Relative optical density (mean \pm SEM) between subjective day and night for GIRK1 and GIRK2. *p < 0.05, n = 3-4 per time point.



Figure 2: GIRK currents are increased during the day in SCN neurons. A) Peak inward current (mean \pm SEM) during the day vs. night (*p < 0.05) for GIRK2 knockout (KO) and wild-type (WT) mice. B) Representative voltage-clamp ramp traces. Recordings were done in the presence of TTX (1µM), bicuculline (30µM), D-AP5 (50µM), CNQX (10µM), and CdCl2 (200µM) in order to block synaptic transmission. All recordings were done in 30 mM KCl to increase potassium conductance. n = at least 3 animals and 20 cells per group.



Figure 3: GIRK2 knockout SCN neurons exhibit more depolarized resting membrane potential compared to wild-type controls. A) Resting membrane potential of WT and KO SCN neurons during the day and night. B) Representative gap-free current clamp of KO and WT SCN neurons during the day and at night. A representative trace from WT slices treated with the GIRK2 antagonist Tertiapin-Q (0.2μ M) during the day is also included. Dotted line indicates -40 mV. C, D, E) Input resistance, action potential frequency from gap-free current clamp recordings, and action potential amplitudes from gap-free current clamp recordings, respectively, from WT and KO neurons during the day and night. n = 3-5 animals and at least 25 cells per group, * p < 0.05.



Figure 4: GIRK2 knockout mice fail to shorten free-running period in response to wheelrunning activity. A) Representative, double-plotted actograms of WT (top) and KO (bottom) mice without wheel access (activity measured by infrared motion sensors), n =14-15 per group. B) Wheel-locked infrared free-running period (mean \pm SEM), and wheel-running activity-based free-running period (mean \pm SEM) for KO and WT mice. *p < 0.05. C) Representative actograms for WT and KO mice with wheel access (right, activity measured by wheel revolutions); n = 7-8 animals per each group. Time of lights off is indicated by dark gray; black tick marks indicate activity counts.



Figure 5: GIRK2 knockout mice entrain more rapidly to a 6-hour light cycle advance. A) Activity onset (mean \pm SEM) in days prior and following a 6-hour advance of an LD cycle (day 0) for WT and KO mice. Lights off represented by gray shading. B) Representative actograms of WT (top) and KO (bottom) mice housed on wheels in a 12:12 LD cycle. Lights off represented by gray shading; black tick marks indicate activity counts. n = 6 animals per group.



Figure 6: Loss of GIRK2 reduces effect of NPY on SCN neuron spontaneous firing rate. A) Frequency plot of individual WT and KO SCN neurons treated with 2.35 μ M NPY or vehicle from ZT 4 to 6. Mean value indicated by black solid line. Lowercase letters (a, b, c) indicate groups that are significantly different (*p<0.05). B) Representative loose patch traces (5 sec) of neurons in (A). C) Percentage of silent cells vs. non-silent cells with NPY treatment. n = 3 animals and ≥45 cells per group.



Figure 7: GIRK2 is necessary for NPY-induced phase shifts in the molecular clock. A) Phase shifts of PER2::LUC rhythms in response to a 1-hr treatment of 2.35 μ M NPY or vehicle at CT4 in WT and KO mice. *p < 0.05. B,C) Representative bioluminescence traces from two WT (B) or two KO (C) SCN cultures treated with vehicle (black) or NPY (gray) for three cycles before and after treatment (top) or day 1 after treatment (bottom). Time 0: onset of NPY application. Predicted peak time, determined from 3 cycles post treatment, is indicated by vertical bars for each group. n = 6-8 cultures per group.



Figure 8: Activation of GIRK channels induces nonphotic-like phase advances. A) Phase shifts of PER2::LUC rhythms in response to a 1-hr treatment of 10 μ M ML297 or vehicle starting between CT3-4.5. B) Representative bioluminescence traces from SCN cultures treated with vehicle (black) or ML297 (gray) for three cycles before and after treatment. n = 6 cultures per group.

GIRK CHANNELS MEDIATE THE NONPHOTIC EFFECTS OF EXOGENOUS MELATONIN

by

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CHAPTER 2

GIRK CHANNELS MEDIATE THE NONPHOTIC EFFECTS OF EXOGENOUS MELATONIN

Abstract

Melatonin supplementation has been used as a therapeutic agent for several diseases, yet little is known about the underlying molecular mechanisms by which melatonin acts within the brain to synchronize circadian rhythms. G protein signaling plays a large role in melatonin-induced phase shifts of locomotor behavior. Melatonin receptors have been shown to activate G protein-coupled inwardly-rectifying potassium (GIRK) channels in Xenopus oocytes. The present study tested the hypothesis that melatonin influences circadian phase and electrical activity within the clock center of the brain, the suprachiasmatic nucleus (SCN), through GIRK channel activation. The results showed that, unlike wild-type littermates, GIRK2 knockout mice failed to phase advance wheelrunning behavior in response to 3-day subcutaneous injections of melatonin in the late day. Loose patch electrophysiological recordings of SCN neurons revealed a significant reduction in the average intrinsic action potential rate in response to melatonin. This effect was lost in the presence of a GIRK antagonist, tertiapin-q (TPQ), and in SCN neurons from GIRK2 knockout mice. The melatonin-induced suppression of firing rate corresponded with an increased inward current that was blocked by TPQ. Finally, application of ramelteon, a potent melatonin receptor agonist, also significantly decreased

firing rate and increased inward current within SCN neurons in a GIRK-dependent manner. These results are the first to show that GIRK channels are necessary for the effects of melatonin and ramelteon within the SCN. This study suggests that GIRK channels may be an alternative therapeutic target for diseases with evidence of circadian disruption, including aberrant melatonin signaling.

Introduction

The hormone melatonin, produced by the pineal gland, is a potent regulator of circadian rhythms or 24-hour cycles in behavior and biological processes (Dubocovich, 2007). Exogenous melatonin has been used to treat a variety of diseases that exhibit circadian rhythm comorbidities such as epilepsy (Banach et al., 2011; Jain and Besag, 2013), delayed sleep phase syndrome (Mundey et al., 2005), cardiometabolic diseases (Paulis et al., 2012), and mood disorders such as depression (Racagni et al., 2007; Campos Costa et al., 2013; Comai and Gobbi, 2014; Laudon and Frydman-Marom, 2014). However, the molecular mechanisms linking melatonin-induced changes in neuronal activity to regulating the timing of circadian rhythms is poorly understood.

In both humans and rodents, exogenous melatonin administered during the late day advances the phase of circadian cycle, shifting activity onset to an earlier time (McArthur et al., 1991; Benloucif and Dubocovich, 1996; Hunt et al., 2001; Dubocovich et al., 2005; Mundey et al., 2005). Within the primary clock center of the brain, the suprachiasmatic nucleus (SCN) of the hypothalamus, melatonin application hyperpolarizes the resting membrane potential and suppresses spontaneous action potential rate in neurons within acute SCN slices from mice and rats (Jiang et al., 1995; van den Top et al., 2001; Scott et

al., 2010). These effects on the circadian system are thought to be mediated through G protein-coupled signaling. G protein-coupled melatonin receptors mediate melatonininduced phase shifts in behavior and changes in SCN firing rate (Hunt et al., 2001; Dubocovich and Markowska, 2005). In addition, melatonin-induced hyperpolarization of SCN neurons has been shown to be pertussis toxin-sensitive (van den Top et al., 2001), indicating Gi/o heterotrimeric G protein signaling is critical for the acute electrophysiological effects of melatonin. G protein-coupled inwardly-rectifying potassium (GIRK) channels are potential candidate mediators of this inhibitory effect of melatonin, given that GIRK currents are increased by melatonin receptor activation in a Xenopus oocyte expression system (Nelson et al., 1996). Recently, we have shown that GIRK channel activation varies over the day-night cycle and that day-time activation is sufficient to induce phase advances of the molecular clock within the SCN (Hablitz et al., 2014). We hypothesize that GIRK channels mediate the phase advancing effects of exogenous melatonin. Here, we use behavioral and electrophysiological techniques to ascertain: whether GIRK channels are necessary for the inhibitory and phase synchronizing effects of melatonin on SCN neurons and wheel-running behavior, and if ramelteon, a potent clinically relevant melatonin receptor agonist (Kato et al., 2005), requires GIRK channels to alter SCN electrophysiology.

Materials and Methods

Ethical approval

All animal care, handling, and housing were in compliance with the University of Alabama at Birmingham's Institutional Animal Care and Use Committee guidelines.

Animals and housing

All mice in these experiments were 2-4 months of age to reduce developmental or aging phenotypes (Turek et al., 1995; Biello, 2009). Only male mice were used for behavioral experiments (Ruiz de Elvira et al., 1992; Vyazovskiy et al., 2006). GIRK2 knockout (KO) animals on a C57/BL6 background (Signorini et al., 1997) and wild-type (WT) littermate controls were used for electrophysiology and circadian behavioral analysis. Although C57/BL6 mice are melatonin deficient, studies have confirmed that melatonin binding and phase shifting effects of melatonin are still intact and comparable to other mouse strains (Siuciak et al., 1990; Liu et al., 1997). Separate cohorts of mice were used for each different experiment. Unless otherwise stated, mice were group housed on a 12:12 light/dark (LD) cycle with food and water ad libitum.

Behavioral Analysis

All mice were housed in individual wheel cages. Wheel-running activity was recorded and analyzed using Clocklab software (Actimetrics, Wilmette, IL). After entrainment to a 12:12 LD cycle, mice were released into constant darkness (DD) for at least 10 days prior to treatment. Mice were then treated at approximately CT10 (circadian time 10; CT12 is defined as activity onset) with either subcutaneous injections of vehicle (3% ethanol/saline) or melatonin (90 μ g/mouse) for three consecutive days (as in (Dubocovich et al., 2005)). Phase shifts were measured as the difference between two predictions for the time of activity onset the day after treatment (using linear regression for each time period pre and post treatment for each animal). The first prediction was based on the 7 onsets prior to the first injection and the second prediction was based on

the 7 onsets after three stabilization days post injections. Mice that were injected before CT 9.5 were excluded from analysis. When possible, mice were treated with vehicle and melatonin using a crossover design. During the experiment, 9 of 13 WT and 6 of 7 KO mice received both treatments with at least 2 weeks between trials.

Electrophysiology

Mice were killed either at ZT8.5 (Zeitgeber time 8.5; ZT12 defined as lights off) by cervical dislocation. All recordings were made between projected ZT 10-12. Brains were harvested, sectioned at 200 µm on a vibroslicer (7000smz, Campden Instruments Ltd., Lafayette, IN), and transferred to an open recording chamber (Warner Instruments, Hamden, CT) that was continuously perfused at a rate of 2.0 ml/min with extracellular solution consisting of (in mM), NaCl 124, NaHCO3 20, Na2HPO4 1, MgSO4 1.3, glucose 10, KCl 3.5, CaCl2 2.5 (added the day of experiment). Osmolality was adjusted to 300-305 mOsm), bubbled with 5% CO2 / 95% O2 and heated to 34 ± 0.5 °C. Neurons were visualized with an Olympus BX51WI (Olympus America Inc., Center Valley, PA) using infrared-differential interference contrast optics. Electrodes with a pipette resistance of ~4-6 M Ω were filled with filtered, potassium gluconate solution consisting of (in mM): K-gluconate 135, KCl 10, HEPES 10, EGTA 0.5, then adjusted to pH 7.4 with KOH (Kuhlman and McMahon, 2004). Electrophysiological signals were processed and controlled by a Multiclamp 700B amplifier, and pClamp 10.02 software (Axon Instruments, Union City, CA). Recordings were sampled at 20 kHz and filtered at 10 kHz. In order to block synaptic transmission (as in Fig. 4-5): bicuculline (30µM) and CdCl2 (200µM) (Sigma-Aldritch, St. Louis, MO), D-AP5 (50µM) and CNQX (10µM)

(Abcam, Cambrige, MA), and TTX (1 μ M) (Tocris Biosciences, Minneanapolis, MN) were added to the bath solution. To isolate GIRK currents (Fig. 4-5) the concentration of KCl was increased from 3.5 mM to 30 mM in order to increase potassium conductance across the membrane (as in (Fu et al., 2004)). The GIRK channel antagonist Tertiapin-Q (0.2 μ M) (Alamone Labs, Israel) was used for experiments in Fig. 3-5. For experiments shown in Fig. 3-4, cells were treated with either vehicle (water) or melatonin (2 μ M; Sigma-Aldritch, St. Louis, MO). For experiments shown in Fig. 5 cells were treated with either vehicle (20 nM ethanol) or ramelteon (10 pM; Biotang Inc., Lexington, MA). All data were collected within 6 min of membrane rupture to minimize any potential washout effects from the whole-cell recording (Schaap et al., 1999). Average spike rate was calculated from at least 1 min of the 2-min trace. For all electrophysiological experiments, at least 3 biological replicates with at least 4 cells per animal were used. There was no specific regional bias when recording within the SCN.

Statistical Analysis

All statistical analysis was performed with SPSS 22. For comparisons of means, an independent samples t-test or ANOVA was used for comparisons between two means or two or more means, respectively. Two factor designs were analyzed with a two-way ANOVA with repeated measures (using a linear mixed model) when appropriate. In cases of a non-normal distribution, a nonparametric Kruskal-Wallis test was used, followed by median test for post hoc analyses. For all other tests, significance was ascribed at P < 0.05.

Results

GIRK2 is necessary for melatonin-induced phase advances of behavioral rhythms

It has been previously shown that three-day subcutaneous administration of melatonin in the late day (CT10) phase advances wheel-running behavioral rhythms of mice (Dubocovich et al., 2005). In order to test the hypothesis that these phase advances are mediated by GIRK channels, we performed a similar experiment using GIRK2 KO animals as a model of disrupted GIRK channel signaling, compared to WT littermate controls. Administration of melatonin to WT mice induced significantly larger phase advances in wheel running behavior compared to vehicle (linear mixed model ANOVA, main effect of treatment, F(1, 29.3) = 4.88, P = 0.035; WT: n = 12-14 per group). Moreover, KO animals responded similarly to melatonin and vehicle injections, and these shifts were overall reduced compared to WT animals (main effect of genotype, F(1, 29.3)) = 5.29, P = 0.029; genotype x treatment interaction, F(1,29.3) = 3.32, P = 0.079; KO: n = 7-8 per group), indicating that GIRK2 is necessary for the phase-advancing effects of melatonin on behavior (Fig. 1,2). Similar to our previous findings (Hablitz et al., 2014), GIRK2 KO animals exhibited a longer free-running period compared to WT controls (WT: $23.74 \pm 0.04h$, KO: $23.87 \pm 0.04h$; independent samples t-test, t(37) = 2.094, P = 0.043). Finally, melatonin increased free-running period equally in both WT and KO animals (post vehicle mean: $23.77 \pm 0.04h$, post melatonin mean: $23.85 \pm 0.05h$; repeated measures ANOVA, main effect of treatment, F(1, 34) = 4.324, P = 0.045), indicating that GIRK2 is not necessary for melatonin-induced effects on period length.

Melatonin-induced suppression of SCN neuronal activity is mediated by GIRK currents

Previous studies have shown that melatonin suppresses action potential rates of SCN neurons both in rat (Jiang et al., 1995; Zhou et al., 2000; van den Top et al., 2001) and mouse (Scott et al., 2010). Given that GIRK channels are necessary for the phaseadvancing effects of late-day melatonin on wheel-running behavior, we tested the hypothesis that GIRK channels are necessary for the inhibitory effects of melatonin in the SCN. First, we performed loose patch electrophysiology within the SCN. Melatonin application $(2 \mu M)$ in the late day (ZT10-12) significantly decreased the spontaneous action potential frequencies of SCN neurons (mean \pm SEM, Veh: 4.51 \pm 0.47 Hz, Mel: 3.07 ± 0.36 Hz; Kruskal–Wallis test, H(3) = 8.035, P = 0.045; median post hoc test, P = 0.048 for Veh vs. Mel). This effect was lost in the presence of tertiapin-Q (TPQ), a GIRK channel antagonist (TPQ + Mel: 5.04 ± 0.54 Hz, median post hoc test, P = 0.009 for TPQ + Mel vs. Mel). TPQ alone had no influence on spike rate of SCN neurons (TPQ: $4.84 \pm$ 0.53 Hz; Median post hoc test, P = 0.755, Veh vs. TPQ; n = >28 cells/group). Similar to TPQ, melatonin application to SCN slices from GIRK2 KO mice did not suppress firing rate when applied during the late day (KO Veh: 4.62 ± 0.45 Hz, KO Mel: 5.07 ± 0.48 Hz; independent samples t-test, t(88) = -0.692, P = 0.49, n = >44 cells per group), indicating that GIRK channels are necessary for the decreased firing rate in response to melatonin (Fig. 3).

Melatonin receptor activation has been shown to activate GIRK channels in a Xenopus oocyte expression system (Nelson et al., 1996), yet it remains unknown as to whether melatonin directly activates GIRK channels within SCN neurons. To measure a melatonin-sensitive GIRK current in SCN neurons, we used whole-cell, voltage-clamp electrophysiology and pharmacological inhibition of synaptic transmission (1µM TTX, 30µM bicuculline, 50µM D-AP5, 10µM CNQX, and 200µM CdCl2) along with increased KCl (30mM; in order to increase potassium conductance as in (Hablitz et al., 2014)), in response to a slow ramp (2.5 s) from -140 mV to -20 mV. We found that melatonin increased peak inward current during this protocol (mean \pm SEM, Veh: -56.64 \pm 5.79 pA, Mel: -113.59 \pm 10.21 pA; Kruskal–Wallis test, H(3) = 20.98, P = 0.0001; median post hoc test, P = 0.001 for Veh vs. Mel), and this effect was lost in the presence of TPQ (TPQ + Mel: -61.35 \pm 8.59 pA, median post hoc test, P = 0.001 for TPQ + Mel vs. Mel; n= >17 cells per group), demonstrating that melatonin directly activates GIRK channels in SCN neurons (Fig. 4).

GIRK channels mediate the effects of the melatonin agonist Ramelteon within the SCN

Ramelteon is a MT1/2 receptor agonist used in treating sleep disorders, depression, and delirium (Borja and Daniel, 2006; Hatta et al., 2014). This therapeutic agent can cause similar phase-shifts to the circadian cycle (Rawashdeh et al., 2011), and has a higher affinity for melatonin receptors than melatonin (Kato et al., 2005; Miyamoto, 2009). Here, we tested whether ramelteon acts upon similar mechanisms as melatonin within the SCN, thereby inducing GIRK currents and suppressing action potential rates. The results indicated that 2-hr ramelteon application (10 pM, from ZT10-12) significantly decreased the spike rates of SCN neurons (mean \pm SEM, Veh: 3.49 \pm 0.4 Hz, Ram: 1.99 \pm 0.3 Hz) and this effect was gone in the presence of TPQ (TPQ: 3.52 \pm 0.5, TPQ + Ram: 3.29 \pm 0.5; Kruskal–Wallis test, H(3) = 14.89, P = 0.002, Fig. 5). Moreover, this inhibition corresponded to an increased inward current that was sensitive to TPQ (two-
way ANOVA, ramelteon by TPQ interaction: F(1,119) = 8.122, P = 0.005; simple effects analysis: ramelteon vs. vehicle: P = 0.0001, Ramelteon vs. TPQ: P = 0.842; Fig. 5; n= >23 cells per group), similar to melatonin treatment (Fig. 5). Altogether, these results indicate that GIRK channels are a downstream target of melatonin receptor activation by ramelteon.

Discussion

The molecular mechanisms underlying decreased neuronal firing and phase shifts induced by late-day melatonin within the SCN are largely unknown. Here, we propose that GIRK channels mediate this suppression of firing rate, and are necessary for melatonin induced phase shifts. Indeed, we show that melatonin fails to phase advance circadian activity rhythms of mice in the absence of GIRK2 channels. Furthermore, melatonin did not suppress spontaneous action potential rates of SCN neurons when GIRK channels were blocked or genetically ablated. This melatonin-induced decrease in spike rate corresponded to activation of a TPQ¬-sensitive GIRK current. These results support the hypothesis that GIRK channel activation is necessary within the SCN to convey the phase shifting properties of melatonin on the circadian clock. Because melatonin is a key molecule to treat sleep disorders, depression, and several other diseases, we examined the effects of ramelteon, a clinical sleep aid and potent MT1/2receptor agonist, on SCN neurophysiological response. We found that ramelteon required GIRK channel activation to suppress SCN firing rate. To our knowledge, our study is the first to investigate this acute inhibitory effect of ramelteon on neuronal excitability.

Altogether, these results indicate that GIRK channels are a potential therapeutic target in diseases where melatonin signaling has been disrupted.

Similar to melatonin, previous work from our lab shows that neuropeptide Y (NPY) signaling within the SCN is also mediated through GIRK channels (Hablitz et al., 2014). Importantly, activation of GIRK channels during the day is sufficient to phase advance organotypic SCN cultures (as reported by PER2::luciferase), similar to in vivo phase response curves for both NPY (Huhman and Albers, 1994; Besing et al., 2012) and melatonin (McArthur et al., 1991; Hastings et al., 1992; Benloucif and Dubocovich, 1996; Hunt et al., 2001; Dubocovich, 2007). This type of phase response curve with maximal phase advances during the day is characteristic of several neurotransmitters that signal for the presence of 'nonphotic' stimuli (i.e., not driven by light-induced activation of the retinohypothalamic tract), such as arousal and exercise (Challet, 2007). Interestingly, these neurotransmitters (melatonin, NPY, serotonin and GABA) are mediated through Gi/o heterotrimeric G protein signaling (Muraki et al., 2004; Dubocovich, 2007; Fowler et al., 2007; van den Pol, 2012), indicating that there may be shared common mechanisms for resetting circadian clock phase. Here, we show that GIRK channels are necessary for melatonin-induced SCN neuronal response and behavioral phase advances. Taken together with our prior study, we propose that GIRK channel activation is a putative conserved mechanism for nonphotic signals to influence the circadian clock.

Although the present study found that melatonin-induced changes in SCN neurophysiology were mediated through GIRK channels, little is known about how exogenous melatonin may regulate the molecular clock, a transcription-translation

feedback loop (Roenneberg and Merrow, 2005; Partch et al., 2014) that drives circadian rhythms on a molecular level throughout different tissue types. A single injection of melatonin in the late day does not influence molecular clock levels on the first day, but does significantly change expression of Per1, Per3, Bmal1 and AVP on the second day (Poirel et al., 2003). Nuclear orphan receptors such as Reverb- α , which have been implicated in melatonin signaling (Agez et al., 2007), may mediate this effect. Also, mice with a CLOCK mutation show normalization of period length in the presence of melatonin or ramelteon (Shimomura et al., 2010). Future studies should investigate whether GIRK channel activation ultimately influences components of the molecular clock such as regulation of Reverb-α (Agez et al., 2007; Agez et al., 2009) influencing redox state of the cell (Bonnefont-Rousselot and Collin, 2010; Luchetti et al., 2010; Garcia et al., 2014), or second messenger pathways like PKC activation (Luchetti et al., 2010), all of which have been implicated in melatonin signaling. It is important to emphasize that although GIRK channels are necessary for melatonin-induced phase advances, this does not preclude the involvement of co-activation of PKC in response to melatonin. Indeed, studies measuring SCN ensemble firing in rats and rhythmic PKC expression in cell culture have demonstrated that PKC activation via MT1/2 receptors plays a key role in the phase shifting effects of melatonin (McArthur et al., 1997; Rivera-Bermudez et al., 2003; Rivera-Bermudez et al., 2004).

In addition to resetting circadian phase, endogenous melatonin also provides a seasonal cue, signaling photoperiodic day length via changes in hormonal circulation patterns released from the pineal gland (Coomans et al., 2014). Classic studies have shown that changes in the length of the photoperiod cause long-lasting effects on period

length, such that longer or shorter photoperiods lengthen or shorten period, respectively (Pittendrigh and Daan, 1976). In the present study, we found that administration of melatonin for three consecutive days at CT10 was sufficient to significantly increase period length by approximately 5 minutes, indicating that acute pulses of melatonin may provide day-length information. Although this result may seem biologically insignificant, it may provide insight into circadian timing disruption in affective disorders such as seasonal affective disorder and depression, which have been successfully treated with melatonin receptor agonists (Srinivasan et al., 2012) or have shown evidence of disrupted melatonin receptor disruption within the SCN in postmortem brain tissue (Wu et al., 2013).

In conclusion, melatonin signaling has been shown to influence neuronal excitability, metabolic state, time-of-day, inflammation, and much more (Dubocovich et al., 2003; Dubocovich, 2007; Uberos et al., 2010; Paulis et al., 2012; Srinivasan et al., 2012). Ramelteon is clinically available and has been used to treat depression, insomnia, delirium, and Alzheimer's disease symptoms (Borja and Daniel, 2006; Furuya et al., 2012; Hatta et al., 2014). Here, we show that the effects of both melatonin and ramelteon on the neurophysiological function of SCN neurons are mediated by GIRK channels. Diseases such as addiction, Down's syndrome, and epilepsy, which have strong circadian components (Loddenkemper et al., 2011; Zarowski et al., 2011; Cho, 2012; Churchill et al., 2012; Lott, 2012; Stores and Stores, 2012; Ramgopal et al., 2013; Parekh et al., 2015; Webb et al., 2015), are characterized by aberrant GIRK channel function (Kobayashi and Ikeda, 2006; Luscher and Slesinger, 2010; Arora et al., 2011; Kaufmann et al., 2013). Future studies could investigate whether pharmacological regulators of GIRK channels,

such as TPQ or ML297 (a GIRK channel agonist (Days et al., 2010; Wydeven et al.,

2014)), could be used as therapeutic agents in diseases presenting with circadian rhythm disruption. In support of this future research area, one study has shown that ML297 is effective in reducing signs of epilepsy in mice (Kaufmann et al., 2013).

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Figure 1: Melatonin-induced phase shifts in circadian activity of WT and GIRK2 KO mice. Representative, double-plotted actograms of individual WT (A,C) or KO (B,D) mice treated with vehicle (A,B) or melatonin (C,D). WT and KO mice were housed in constant darkness (indicated by gray shading) and treated 3 consecutive days at CT10 with either injections of vehicle (3% ethanol/saline, s.c.) or melatonin (90µg/mouse, s.c.). Treatment is indicated by white circles; best fit lines to activity onsets are shown in gray with the extended fit lines in black. White arrows indicate the day after injections (used in the phase shift predictions; see Methods). Black tick marks indicate activity counts.



Figure 2: GIRK2 is necessary for the phase-advancing effects of melatonin on wheel running activity. A) Average phase shifts (mean \pm SEM) of wheel-running activity rhythms in response to 3 days of subcutaneous injections at CT10 with either vehicle (3% ethanol/saline) or melatonin (90µg/mouse) in WT and KO mice. B) Free-running period (mean \pm SEM) for KO and WT mice before treatments. C) Cumulative free-running period (mean \pm SEM) post treatment with either vehicle or melatonin. *P < 0.05.







Figure 4: Melatonin induced an inward GIRK current in SCN neurons. A) Peak inward current (mean \pm SEM) for vehicle and melatonin (2 μ M) treated cells during ZT10-12 in the presence and absence of TPQ (0.2 μ M). *P < 0.05. B) Representative voltage-clamp ramp traces. Recordings were done in the presence of TTX (1 μ M), bicuculline (30 μ M), D-AP5 (50 μ M), CNQX (10 μ M), and CdCl2 (200 μ M) in order to block synaptic transmission and 30 mM KCl to increase potassium conductance.



Figure 5: GIRK channels mediated the inhibitory effects of ramelteon within the SCN. A) Action potential rates (mean \pm SEM) for vehicle and ramelteon (10 pM) treated cells during ZT10-12 in the presence and absence of TPQ (0.2 μ M). *P < 0.05. B) Representative loose patch traces (5 sec) of neurons in (A). C) Peak inward current (mean \pm SEM) for vehicle and ramelteon (10 pM) treated cells during ZT10-12 in the presence and absence of TPQ (0.2 μ M). *P < 0.05. B) Representative voltage-clamp ramp traces. Recordings for C-D were done in the presence of TTX (1 μ M), bicuculline (30 μ M), D-AP5 (50 μ M), CNQX (10 μ M), and CdCl2 (200 μ M) in order to block synaptic transmission. Recordings in C-D were done in 30 mM KCl to increase potassium conductance.

DISCUSSION

GIRK channels and nonphotic entrainment: a recap

In general, both photic and nonphotic information signals are transmitted to the SCN, altering cellular excitability and the timing of the molecular clock. Ultimately, this signaling changes the entrainment of the animal. This dissertation explores the role of GIRK channels in mediating these cues within the SCN (Figure 1). The main hypothesis of this dissertation is that **nonphotic stimuli activate GIRK channels which decrease neuronal excitability, modulating the timing of circadian rhythms.**

Chapter one of this dissertation focused on defining the role of GIRK channels in regulating the timing of circadian rhythms and regulation of SCN excitability. GIRK1 and GIRK2 protein was found in the SCN. GIRK2 protein levels were higher during the day, even under constant conditions, indicating direct regulation by the circadian clock. This high level of GIRK2 during the day corresponded with an increased inward current, as measured by whole-cell voltage clamp electrophysiology. Loss of GIRK2, utilizing a GIRK2 knockout mouse, eliminated this current and resulted in more depolarized daytime SCN neurons. GIRK2 knockout animals failed to shorten free-running period in response to running wheel access and shifted more rapidly to a 6-hour phase advance of the light cycle, suggesting a deficit of the nonphotic entrainment pathway. Indeed, the inhibitory effects of the nonphotic neurotransmitter, NPY, on SCN neuronal firing rate were diminished in GIRK2 KO brain slices. Finally, broad activation of GIRK channels via application of ML297 (a GIRK channel agonist) induced a nonphotic phase shift, indicating that activation of GIRK channels is sufficient to mimic a nonphotic cue.

Because GIRK channels mediate the effects of NPY within the SCN, and activation is sufficient to cause a nonphotic phase advance, Chapter 2 focuses on whether

GIRK channels mediate additional nonphotic cues; in this case, the circadian effects of the common sleep aid melatonin. Subcutaneous injections of melatonin in the subjective late day caused a phase advance of wheel running activity in mice. This effect was lost when GIRK2 was genetically ablated. In addition to the phase shifting effects of melatonin, melatonin application to SCN slices suppressed neuronal firing and elicited an inward current, both of which were dependent on GIRK channel function. Finally, we demonstrated that GIRK channels were necessary for the effects of ramelteon, a potent melatonin receptor agonist, on the electrical activity of SCN neurons. This study was the first to characterize the single-cell neurophysiological effects of ramelteon on any neuronal population.

Does GIRK regulation of excitability and phase extend to other brain areas?

Multiple areas of the brain exhibit rhythmicity in clock gene expression and/or excitability (46, 131), such as the hippocampus(132), and multiple regions of cortex (133). The medial habenula even has similar oscillatory waveforms and resting membrane potential regulation as the SCN (134). Yet, the regulation of these rhythms and underlying ion channels governing excitability has yet to be studied. GIRK channels are found throughout many of these brain regions (122), but it is unclear if and how they are regulated by the clock.

In addition to whether these neuronal oscillators regulate their excitability in a 24hour pattern in a manner similar to the SCN, another fundamental question is how the oscillators maintain their phase relationship to the SCN. Future studies should include: 1) clearly defining the phase relationships of circadian rhythmicity between multiple

oscillators in the brain, 2) determining the sensitivity of these regions to external phaseshifting stimuli (*e.g.*, developing phase response curves), and 3) defining the mechanisms of entrainment for different brain regions. Perhaps GIRK channels regulate entrainment of these regions in a similar manner as the in SCN, such that integration of multiple $G_{i/o}$ coupled cues determines the phase relationship to the SCN. Understanding the phase relationships throughout the brain and the mechanisms underlying them could have huge ramifications for diseases such as schizophrenia, depression, and bipolar disorder, all of which exhibit signs of circadian disruption and involve integration of signals from multiple brain regions (135, 136).

Epilepsy & heart arrhythmias

GIRK channels have been implicated in epilepsy (118, 137-139), a disease of hyperexcitability and seizure generation within the brain. Temporal lobe epilepsy has time-of-day dependent seizure generation, with most seizures occurring in the late evening (5, 6, 140). In addition to epilepsy, GIRK channels, expressed in both the atria and ventricles, have been implicated in both long QT syndrome (141) and chronic atrial fibrillation (142, 143) in the heart. These arrhythmias, caused by dysregulation of excitability in the heart, have time-of-day dependent onset (8, 9, 144).

GIRK channels regulate intrinsic membrane properties of excitable cells throughout multiple peripheral oscillators in the brain and the body. In addition, a GIRK channel agonist, ML297, was able to increase time to seizure onset, increase survival, and decrease then number of convulsions in mice subjected to either a maximal electrical shock model of epilepsy, or a pentylenetetrazol (a GABA_A antagonist) model of epilepsy

(139). Melatonin has been used both in rodent models and in humans to treat epilepsy, but with mixed results (145). Finally, NPY has anticonvulsant effects in animal models of both temporal lobe epilepsy and hippocampal kindling models (146), and application of NPY to hippocampal slices resected from epileptic patients results in decreased evoked action potentials from dentate granule cells (147). Future studies will determine if diurnal/circadian regulation of these channels extends to other oscillators in the brain and heart, and whether this could drive time-of-day dependencies of seizure and arrhythmia generation. Regulating time-of-day administration may increase the efficacy of antiepileptic and antiarrhythmic drugs.

Aging and development

GIRK channels are upregulated within the brain with age (120), and can be expressed differentially in cultured neurons in the presence of different neurotransmitters (113). In the cerebellar cortex, GIRK channel expression reaches maximum potential at P5 (148), and in cultured hippocampal neurons, GIRK channels mediate GABA_B-induced inward currents by day 14 in vitro (149). This tight control over GIRK channel expression during different developmental timelines could have interesting implications for circadian rhythms. For example, it is well known that sdolescents are typically phasedelayed compared to the general population (150). In addition, aging tends to dampen circadian rhythmicity and decreases the response to daily entrainment cues (151, 152). If GIRK channels are differentially expressed within the SCN across development, it could help explain these phenomena.

In addition to basic questions of whether GIRK channels are necessary in the development and maintenance of rhythms, they may also play a role in neurodevelopmental disorders such as Down syndrome (DS), which occurs when chromosome 21 is triplicated, or Rett syndrome (RTT), which is associated with loss-offunction mutations in the gene encoding the methyl CpG binding protein 2 (MeCP2). Both DS and RTT patients and mouse models (specifically the Ts65Dn model of DS where the mouse homolog of the "critical region" of human chromosome 21 is triplicated (153), and MeCP2^{-/y} mice as a model of RTT (154)) exhibit sleep/wake and circadian rhythm disruption (155-162). In the case of DS, the GIRK2 gene is located on chromosome 21 and is thus triplicated, leading to overall decreases in neuronal excitability throughout the brain (163, 164). In RTT, MeCP2^{-/y} mice show overall decreased excitability within the SCN (160). Also, some symptoms of RTT are alleviated with serotonin receptor agonists, which also activate GIRK channels (165, 166). This evidence suggests that GIRK channel regulation within the SCN may be disrupted in these developmental disorders and that further research is necessary to determine if recovery of proper GIRK channel regulation could help alleviate the sleep/wake and entrainment problems in these patients.

Implications for human entrainment

The results of this dissertation project have shown that NPY and melatonin are sufficient to modulate the timing of circadian rhythms via GIRK channel activation. Mice with a loss of GIRK2 fail to adjust their free running period in response to wheel running. We also demonstrate that loss of GIRK2 enhances entrainment to a 6-hour phase advance of the light-dark cycle. These results highlight the importance of nonphotic entrainment and GIRK channels in multiple aspects of circadian timing.

The efficacy of nonphotic entrainment in humans has been debated, mainly due to the smaller magnitude of shifts and the prevalence of highly irregular lighting cycles in day-to-day life that may decrease the efficacy of nonphotic stimuli (167, 168). However, it is possible to entrain blind individuals to scheduled wake up times and brief 10 minute bicycle exercise (169). Additionally, musically enhanced bird song can shift human sleep/wake patterns when given in the early morning (170, 171). Interestingly, scheduled daytime exercise decreases the time to re-entrainment altered light cycles, either in a controlled circadian experiment or due to flights to a different time zone (172, 173). Exercise in older individuals also alleviates sleep disruption phenotypes (174-177). These experiments suggest that nonphotic stimuli, including music and exercise, are effective at entraining the circadian clock in humans, and could help improve sleep and mood in the older population or individuals with circadian disruption.

Bright light therapy (LT) is a noninvasive, fast, and efficacious way to reset the circadian clock. In patients with major depressive disorder, LT has been shown to alleviate some symptoms within one week, compared to antidepressant drugs that have a minimal time of two to three weeks before the effects are noticeable (178). LT has also been successfully used to treat seasonal affective disorder, insomnia, bipolar disorder, and epilepsy (179-183). However, LT is most effective in the morning, which is a problem for those patients with delayed phase angles, and is most often prescribed for a half hour every day (184-186). This strict regimen hampers adherence. Understanding how nonphotic cues, their downstream targets, and photic signaling interact could have

major implication for LT, potentially decreasing the time investment needed to be successful. Specifically, future studies should focus on the phase advancing effects of nonphotic cues, and timing these stimuli in such a way that does not block the effects of light.

Conclusions

This dissertation demonstrates that not only does NPY and melatonin-signaling require GIRK channel activation within the SCN, but that GIRK channel activation alone is sufficient to mimic a nonphotic phase advance, suggesting that GIRK channel activation may be a conserved response to $G_{i/o}$ coupled cues within the SCN (see Figure 1 for overall model of entrainment). GIRK channels have been implicated in a wide variety of diseases that show evidence of circadian disruption, and with proper timing and dosage, GIRK channel activation could be a powerful chronotherapeutic target. Future studies should investigate how GIRK channels regulate excitability in peripheral oscillators throughout the brain and body, how GIRK channels determine phase of the circadian clocks in these regions, if GIRK channels mediate development of circadian rhythms within the SCN, and finally, how GIRK channels modulate the nonphotic interactions with the photic entrainment system.



<u>Figure 1: Model of entrainment.</u> Photic and nonphotic stimuli interact and converge upon the SCN, a cluster of highly synchronized individual neuronal oscillators. These signals are coded by altering neuronal excitability and changing the timing of the molecular clock. The SCN signals through downstream cues to synchronize peripheral oscillators, eventually leading to a fully entrained animal. We hypothesize that GIRK channels modulate the change in activity of SCN neurons in response to nonphotic cues.

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APPENDIX I

IACUC APPROVAL FORM



MEMORANDUM

DATE: 29-Jan-2015

FROM:

TO: Gamble, Karen Lynnette

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Robert A. Kesterson, Ph.D., Chair

Institutional Animal Care and Use Committee (IACUC)

SUBJECT: NOTICE OF APPROVAL

The following application was approved by the University of Alabama at Birmingham Institutional Animal Care and Use Committee (IACUC) on 29-Jan-2015.

Protocol PI: Gamble, Karen Lynnette

Title: GIRK Channel Modulation of SCN Excitability and Circadian Rhythms (Lauren Hablitz)

Sponsor: National Institute of Neurological Disorders and Stroke/NIH/DHHS

Animal Project Number (APN): IACUC-00453

This institution has an Animal Welfare Assurance on file with the Office of Laboratory Animal Welfare (OLAW), is registered as a Research Facility with the USDA, and is accredited by the Association for Assessment and Accreditation of Laboratory Animal Care International (AAALAC).

Institutional Animal Care and Use Committee (IACUC) | Mailing Address:

CH19 Suite 403 | CH19 Suite 403

FAX (205) 934-1188

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