Archival Report

Circadian Disruptions in the *Myshkin* Mouse Model of Mania Are Independent of Deficits in Suprachiasmatic Molecular Clock Function


**ABSTRACT**

**BACKGROUND:** Alterations in environmental light and intrinsic circadian function have strong associations with mood disorders. The neural origins underpinning these changes remain unclear, although genetic deficits in the molecular clock regularly render mice with altered mood-associated phenotypes.

**METHODS:** A detailed circadian and light-associated behavioral characterization of the Na\(^+\)/K\(^-\)-ATPase \(\alpha_3\) *Myshkin* (*Myk*\(^/-\)) mouse model of mania was performed. Na\(^+\)/K\(^-\)-ATPase \(\alpha_3\) does not reside within the core circadian molecular clockwork, but *Myk*\(^/-\) mice exhibit concomitant disruption in circadian rhythms and mood. The neural basis of this phenotype was investigated through molecular and electrophysiological dissection of the master circadian pacemaker, the suprachiasmatic nuclei (SCN). Light input and glutamatergic signaling to the SCN were concomitantly assessed through behavioral assays and calcium imaging.

**RESULTS:** In vivo assays revealed several circadian abnormalities including lengthened period and instability of behavioral rhythms, and elevated metabolic rate. Grossly aberrant responses to light included accentuated resetting, accelerated re-entrainment, and an absence of locomotor suppression. Bioluminescent recording of circadian clock protein (PERIOD2) output from ex vivo SCN revealed no deficits in *Myk*\(^/-\) molecular clock function. Optic nerve crush rescued the circadian period of *Myk*\(^/-\) behavior, highlighting that afferent inputs are critical upstream mediators. Electrophysiological and calcium imaging SCN recordings demonstrated changes in the response to glutamatergic stimulation as well as the electrical output indicative of altered retinal input processing.

**CONCLUSIONS:** The *Myshkin* model demonstrates profound circadian and light-responsive behavioral alterations independent of molecular clock disruption. Afferent light signaling drives behavioral changes and raises new mechanistic implications for circadian disruption in affective disorders.

**Keywords:** Bipolar, Circadian, Light, Mania, Mood, Suprachiasmatic

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Bipolar disorder (BPD) is a debilitating mental health condition that affects approximately 0.7% to 0.9% of the population of Western societies (1). BPD is characterized by episodes of depression, euthymia, and mania, but its etiology and neural substrates remain poorly understood (2). Disruption of sleep and circadian rhythms is prevalent in many mental health diseases, including BPD, and as such, underlying circadian systems are implicated within BPD pathophysiology (3). Indeed, treatment of circadian rhythm abnormalities can alleviate symptoms of affective disorders (4,5), while circadian clock gene polymorphisms represent risk factors across neuropsychiatric conditions (6). Therefore, studying the circadian system may provide insight into the mechanisms and root causes of BPD.

Coordinated circadian rhythms in mammals, including humans, originate from the master circadian clock located in the hypothalamic suprachiasmatic nuclei (SCN) (7). SCN neurons contain an intracellular gene–protein transcription–translation feedback loop (TTFL) that is the molecular basis of circadian timekeeping, and the Period1/2 (Per1/2) genes and their protein products Period1/2 (Per1/2) are key components of this intracellular molecular clock (8). The TTFL drives the SCN neural network to exhibit electrically excited states during the day and relatively quiescent states at night (9). Such variation is key for individual SCN neurons to coordinate their internal clocks, as well as for the SCN to signal and exert temporal control on behavior and physiology (10). Consistent alignment of these central circadian rhythms to the external environmental light/dark (LD) cycle is important for health and well-being (11). Light information signaled directly from intrinsically photosensitive retinal ganglion cells to the SCN is critical in this process (12). This non–image-forming light input pathway, the retinohypothalamic tract, uses the excitatory neurotransmitter glutamate to activate SCN neurons, resetting
the phase of the TTFL and ultimately the timing of behavioral and brain states such as sleeping and waking (13,14). Further, both SCN-dependent and SCN-independent actions of light exert a powerful influence on mood pathology (15).

Interestingly, in mice, targeted disruption of core TTFL components alters circadian rhythms and consistently elevates the expression of aberrant behaviors resembling those of human affective disorders (16,17). However, because circadian clock genes are also expressed in mood-regulating brain centers (18,19), it is challenging to ascribe behavioral deficits directly to specific brain loci (20). Further, in human BPD, it is unclear whether sleep and circadian disruption arise as etiological drivers or as a consequence of wider pathophysiology. To gain insight into these problems, we used the Myshkin (Myk+/+) mouse, which possesses a heterozygous inactivating mutation in the neuron-specific Na+/K+-ATPase (NKA) α3 subunit, encoded by AtpTa3, and models the manic phase of BPD with face, construct, and predictive validity (20,21).

Importantly, we also found that the Myk+/+ SCN TTFL rhythms are intact, while the electrophysiological output of the Myk+/+ SCN neural network was damped. Critically, we show that period-lengthening effects on behavior of the Myshkin mutation are ameliorated through removal of the light input pathway. Circadian abnormalities in the Myk/+ mice arise through alterations in light signaling and processing by the SCN. This model provides new insights into the etiological mechanisms of circadian disruption in animal models of affective disorders that are independent of core circadian clock gene perturbation.

METHODS AND MATERIALS

Animal Housing and Breeding

Adult congenic Myk+/+ and wild-type (+/+) animals (2–6 months of age) used in this study were bred from pairs (male Myk+/+ × female +/+ ) of animals that had been backcrossed to the C57BL/6NCr strain for 20 generations (20). Pilot investigations revealed no obvious sex differences in behavioral measures in either genotype, so the data from male and female mice were combined (see also Supplemental Figure S1). To generate mice in which the dynamic activities of the molecular clock can be monitored in tissues ex vivo, Myk+/+ mice were crossed with mPer2LUC mice bearing a knock-in PER2-luciferase (LUC) construct (referred to here as PER2::LUC mice) (22). Congenic +/+ × PER2::LUC (+/+ × PER2) and Myk+/+ × PER2::LUC animals were generated through crosses of heterozygous male Myk+/+ and female PER2::LUC animals. All behavioral and in vitro studies of mice on the PER2::LUC background were performed on filial 1 generation animals. See Supplemental Methods for further details.

For assessment of daily rhythms in locomotor activity (with or without a running wheel), ingestive behavior (feeding and drinking), and metabolic activity, animals were housed singly as previously described (23,24). Most studies were conducted under 12-hour LD conditions; however, in some instances animals were assessed under a day-length (16-hour light/8-hour dark cycle) condition or in constant dark (DD) or constant light (LL). These in vivo investigations were conducted using previously established protocols (23,25); see the Supplement for study-specific details.

For in vitro assessments of electrophysiological activity, calcium transients, and bioluminescence rhythms in PER2::LUC, SCN-containing brain slices were made from adult mice using previously published protocols (26). Whole-cell current-clamp recordings and assessment of calcium transients were performed as previously described (9,26). Rhythms in whole SCN slice PER2::LUC expression were assessed using luminometry, while single cells in SCN slices were visualized and imaged using a Hamamatsu Image EM9100-13 electron-multipling-CCD (Hamamatsu, Welwyn Garden City, UK) (27,28).

Data Analysis and Statistics

Unless stated otherwise, genotype comparisons were made using two-tailed Student t test or by two-way analysis of variance with Sidak post hoc comparisons. For within-genotype comparisons, one-way analysis of variance was applied with Sidak corrections unless otherwise stated. Nonparametric equivalents and corrections for unequal variances were utilized where appropriate and are detailed in figure captions. The threshold for statistical significance was set at p < .05. See the Supplement for further details.

RESULTS

Circadian mechanisms as well as visual and non-image-forming light pathways influence normal and pathophysiological states including metabolism and mood behaviors (15). As such, we sought to determine if and how murine daily and circadian rhythms are influenced by the Myshkin mutation.

When singly housed under standard 12-hour LD conditions, the mania phenotype of Myk+/+ mice was characterized by an inability to restrict the normal nocturnally elevated wheel-running activity to the dark phase. Instead, Myk+/+ animals sustained vigorous activity into the first 3 to 4 hours of the lights-on phase (Figure 1A). Consequently, in comparison with +/+ animals, the duration of the daily active (alpha) phase was significantly lengthened by ~2.1 hours (Figure 1B), with the percentage of wheel running occurring during the day elevated by ~6.9% (Figure 1C) in Myk+/+ mice. Some Myk+/+ animals also showed unstable LD rhythms, with activity onsets variably preceding the initiation of the dark phase (Figure 1A). The effects did not rely on the provision of a running wheel, because monitoring of general locomotor activity via infrared detector indicated near-identical disruption in Myk+/+ animals (Supplemental Figure S4A–D). Similar effects of the Myshkin mutation on wheel-running activity were observed in a separate cohort of female mice (Supplemental Figure S1A–E).

In humans, extending daily exposure to light (~14 hours per 24 hours) associated with summer months can exacerbate the symptoms of mania (29,30). When transferred into longer day length (16 hours light/8 hours dark), Myk+/+ mice exhibited longer alpha, weaker rhythms, and ~40% higher wheel running in the light phase than +/+ mice (Supplemental Figure S2A–D). This reveals that the Myshkin mutation compromises behavioral consolidation and that increasing the
duration of the lights-on phase overtly disrupts rhythmic control of behavior.

Circadian disruption and mental illness can alter body weight regulation (31–33). Indeed, bipolar patients with mania can exhibit elevated basal metabolic rate (34), so we subsequently profiled metabolic activity in Myk/+ and +/+ mice. Using indirect calorimetry and monitoring of ingestion activity for 6.5 days under a 12-hour LD cycle, Myk/+ animals were found to exhibit elevated basal metabolic rate, heat production, and drinking activity (Figure 2A–C). The duration of their elevated metabolic activity was sustained into the lights-on phase (Figure 2A, E). Further, when released into an illuminated (\( \sim 1.5 \) \( \mu \text{W/cm}^2 \)) open-field test arena for 15 minutes during the early night (Zeitgeber time 15–18 [ZT15–18]), hyperlocomotor activity (as assessed by distance traversed) was marked in Myk/+ but not in +/+ mice (Supplemental Figure S3). Therefore, unlike other nocturnal rodents such as the Syrian hamster (35) as well as mice with TTFL mutations (36), Myk/+ mice do not exhibit pronounced negative masking, an effect overtly manifested in wheel running during the day.

Because Myk/+ mice exhibited disrupted rhythms under LD conditions, their intrinsic circadian rhythms in wheel-running behavior were initially assessed over 14 days in the absence of light (DD). Consistent with a previous report (20), behavioral rhythms of Myk/+ mice differed significantly from those of +/+ animals. Myk/+ mice displayed a lengthened period (Figure 3A–D; \( \sim 24.2 \) hours vs. \( \sim 23.5 \) hours) and an unusually elongated active phase (Figure 3E: alpha; \( \sim 19.0 \) hours vs. \( \sim 13.1 \) hours), while the amplitude, or strength (percent variance as measured by chi-square periodogram), of their behavioral rhythms was also markedly reduced (Figure 3F). In a separate all-female cohort, Myk/+ animals also showed similar changes in circadian rhythms of wheel running in DD (Supplemental Figure S1A, B, F–H). These effects of the Myshkin mutation were not dependent on the provision of a running wheel (Supplemental Figure S4A, B, E, F).

When assessed for an additional 14 to 21 days in DD, all +/+ animals maintained wheel-running activity, whereas that of some Myk/+ mice (\( n = 3 \) of 38; 8%) weakened and they became circadianly arrhythmic (Figure 3C). Additionally, some Myk/+ animals (but no +/+ mice) exhibited gradual unusual changes in free-running period (\( n = 8 \) of 38; 21%) either spontaneously or following transfer to a clean cage (Supplemental Figure S5A–C). Because rhythm amplitude and free-running stability are metrics for the output strength of the circadian system, this indicates that Myk/+ animals possess diminished and unstable central circadian regulation of behavior and physiology.
Altered Circadian Rhythms and Responses to Light in a Manic Mouse

For rodents in DD, light exposure during the night shifts subsequent onsets of activity; early night exposure (circadian time 14–18 [CT14–18]) delays circadian rhythms, while light given later in the night (CT20–24) advances rhythm onsets (39), so we next sought to determine if the Myshkin mutation affected photic resetting of the circadian system. Mice were released from 12-hour LD conditions into DD and, after 14 days, were exposed to a 1-hour light pulse either late (CT20–21; Figure 4A, B) or early (CT14–15; Figure 4C, D) in their active subjective night (an Aschoff type I protocol). The subsequent phase of the onset of their wheel-running rhythms was then measured (23). When exposed to light pulses at CT14, Myk/+ mice showed much larger phase delays than +/+ mice (~2.32 hours vs. ~1.6 hours, p = .008; Figure 4C–E). Light pulse treatment at CT20 evoked phase delays in Myk/+ activity that were unusual, as they were of the opposite direction to typical advances elicited at this time in +/+ mice (~1.3 hours vs. +0.5 hours, p ≤ .02;

Figure 2. Altered metabolic rhythms in Myk/+ mice. (A) Smoothed traces of wild-type (+/+ ) mice (blue; n = 10) and Myk/+ mice (red; n = 12) maximal oxygen consumption (Vol O2) over 6.5 days. Gray-shaded columns delineate lights off (dark). (B) Mean Vol O2 consumption is elevated in Myk/+ animals (+/+: 2.91 ± 0.14 L/kg/hour, Myk/+ : 3.86 ± 0.13 L/kg/hour; p < .0001). (C) Mean hourly heat production is elevated in the Myk/+ mice (+/+: 0.0148 ± 0.0007 kcal/g/hour, Myk/+ : 0.196 ± 0.0008 kcal/g/hour; p = .0003). (D) Altered ingestion behaviors in Myk/+ animals. Mean hourly food hopper visits were reduced by the Myshkin mutation (+/+: 92.6 ± 4.3 visits, Myk/+ : 64.6 ± 5.1 visits; p = .0005). (E) Mean hourly visits to drinking spout were increased by the Myshkin mutation (+/+: 90.0 ± 13.2 visits/hour, Myk/+ : 172.6 ± 31.7 visits/hour; p = .03). (F) Altered duration of daily peak (+/+: 10.37 ± 0.23 hours, Myk/+ : 13 ± 0.75 hours; p = .003) and nadir (+/+: 8.59 ± 0.30 hours, Myk/+ : 5.06 ± 0.75 hours; p = .0004) in Vol O2 activity. (G) The Myshkin mutation increases percentage of daily feeding occurring during the lights-on phase (+/+: 25.0 ± 1.9%, Myk/+ : 36.3 ± 1.3%; p < .0001). Data in panels (C–G) are plotted as mean ± SEM. *p < .05, **p < .01, ***p < .001. LD, light/dark.

Figure 3. Altered free-running rhythms in Myk/+ mice. (A–C) Example double-plotted actograms of wild-type (+/+ ) and Myk/+ animals released into constant dark (indicated by gray shading) following entrainment to 12-hour light/dark cycle. Most Myk/+ mice maintained rhythms in constant dark (B), but exceptionally, some (C) became arrhythmic. Red rectangles delineate timing of typical elevated wheel running in the Myk/+ mice. (D) The Myshkin mutation lengthened free-running period (+/+: 23.72 ± 0.03 hours, Myk/+ : 24.22 ± 0.02 hours; p < .0001) and (E) alpha duration (+/+: 13.14 ± 0.26 hours, Myk/+ : 18.99 ± 0.37 hours; p < .0001). (F) The amplitude of daily wheel-running rhythms is reduced by the Myshkin mutation (as measured by chi-square periodogram; +/+: 43.8 ± 3.1% variance, Myk/+ : 27.7 ± 2.4% variance; p < .001). Data in panels (D–F) are graphed as scatter plots with mean ± SEM. ***p < .001;
responses to 1-hour light pulses presented under Aschoff type II protocol (CT14, 0.08 hours, \( \text{Figure } 5\text{G, H} \)). This indicates that the phase shift in Myk/+ activity onset compared with the advance in activity onset of the +/+ animal. \( \text{Figure } 4\text{A, B, E} \). Example double-plotted actograms illustrating the shifting effects of a 1-hour light pulse (Aschoff type I protocol) given early in the subjective night (CT14) on the free-running rhythms of +/+ and Myk/+ animals. Myk/+ animals exhibited larger phase delays at CT14. \( \text{Figure } 4\text{E} \). Individual responses to 1-hour light pulses given under Aschoff type I protocol (CT14, +/+: 1.6 ± 0.1 hours, Myk/+: 2.3 ± 0.1 hours, \( p = .008 \); CT20, +/+: 0.5 ± 0.1 hours, Myk/+: 1.3 ± 0.4 hours, \( p = .002 \); CT23, +/+: 0.6 ± 0.1 hours, Myk/+: 0.2 ± 0.2 hours, \( p = .14 \)). \( \text{Figure } 5\text{G, H} \). Example double-plotted actograms showing the phase resetting responses of +/+ and Myk/+ mice to 1-hour light pulse given early in the subjective night (CT14) following transfer from light/dark to constant dark conditions (Aschoff type II protocol). Animals were exposed to a 1-hour light pulse within 48 hours following release into constant dark. \( \text{Figure } 4\text{H} \). Individual double-plotted actograms showing the phase resetting responses of +/+ and Myk/+ mice to simulated jetlag. Jetlag and other external disruptors to normal activity rhythms are associated with the presentation of episodes in BPD, and therefore sensitivity to external perturbation represents an important measure \( (40,41) \).

In response to the 8-hour advance (\( \text{Figure } 5\text{A-D} \)) or delay (Supplemental Figure S5D, E) of the LD cycle, Myk/+ mice rapidly altered their daily pattern of wheel running within 2 to 4 days, whereas +/+ animals took 6 to 8 days to resynchronize. Comparable rapid resynchronization to an 8-hour advance in the LD cycle was also observed in a separate cohort of female Myk/+ mice (Supplemental Figure S1A, B, I, J). To directly test if the Myshkin mutation enhances the photic resetting capabilities of the neural circadian system, animals were next subjected to a transient 7-hour advance of the LD cycle for 2 days then released into free-running DD conditions (\( \text{Figure } 5\text{E, F} \)). Again, Myk/+ mice demonstrated unusually large advances in the new phase of their activity onsets, and on the first day in DD these were of much greater magnitude than those observed by +/+ animals (\( \text{Figure } 5\text{G, H} \)). This finding indicates that mechanisms that normally brake the circadian system to prevent extraordinarily large resetting are dysfunctional in Myk/+ animals \( (42,43) \).

In rodents, exposure to LL suppresses wheel-running behavior and lengthens the period of circadian rhythms \( (24,44) \), so we next assessed how Myk/+ mice adapt to LL. Consistent with previous research, all +/+ animals showed a suppression of wheel running in LL (reduced by \( \sim 90\% \) from LD) and exhibited free-running rhythms with a period of \( \sim 25 \) hours (Supplemental Figure S1A, B, K). Myk/+ animals also showed longer period rhythms in LL (\( \sim 25 \) hours), but most \( (5 \text{ of } 7) \) exhibited markedly elevated wheel running in LL (\( \sim 403\% \) increase from LD). Therefore, while the period-lengthening effects of LL are observed in Myk/+ animals, some individuals sustain increased rather than decreased locomotor activity.

The NKA \( \alpha 3 \) pump is localized to several central nervous system structures, including retinal ganglion cells whose axons project along the optic nerve \( (45,46) \), so we next investigated if retinal input to the SCN contributed to the altered circadian rhythms of Myk/+ mice. To do so, mice free running in DD either underwent a sham surgical procedure or had their optic nerves crushed (see the Supplement for procedural details). Optic nerve crush in Myk/+ mice markedly shortened their circadian period by \( \sim 0.7 \) hours to \( \sim 23.6 \) hours, while +/+ animals showed no obvious change in circadian period (\( \text{Figure } 6\text{A–C} \)). Similarly, Myk/+ mice undergoing sham surgical procedure showed no change in free-running period (\( \text{Figure } 6\text{C} \)). This indicates that aberrant intrinsic activity of the light input pathway to the SCN contributes to the period-lengthening effect of the Myshkin mutation.

Glutamate is the main neurochemical of the retinal input pathway to the SCN, and stimulation of the alpha-amino-3-hydroxy-5-methyl-4-isoxazole propionic acid (AMPA)-type
Glutamate receptors excites SCN neurons, elevates intracellular Ca\(^{2+}\), and can shift the phase of behavioral rhythms (47,48). To determine if acute responses of SCN neurons to glutamate were altered by the Myshkin mutation, the actions of AMPA (5–20 \(\mu\)M) on intracellular Ca\(^{2+}\) were assessed in 1/1 and Myk/1 hypothalamic SCN brain slices. Neurons were loaded with the fluorescent calcium indicator dye Fura-2-acetoxymethyl ester to enable recording of somatic intracellular Ca\(^{2+}\) changes across populations of single SCN cells (Supplemental Figure S6A).

During the day (ZT4–10), AMPA treatments evoked changes in intracellular Ca\(^{2+}\) of similar magnitude and duration in both genotypes (Supplemental Figure S6B, C). However, with applications made during the night (ZT14–18), AMPA elicited increases in intracellular Ca\(^{2+}\) that were significantly larger in Myk/+ neurons compared with +/- SCN neurons (Figure 6D, E). At this time, response magnitude was dose dependent (5 and 10 \(\mu\)M), with baseline recovery from the 10-\(\mu\)M application taking significantly longer in the Myk/+ SCN neurons (Figure 6F). Such changes in AMPA responses indicate that, in addition to altering retinohypothalamic tract activity, the Myshkin mutation enhances the processing of a neurochemical mimic of this light input pathway to the SCN.

To gain further insight into the etiology of circadian disturbances arising from the Myshkin mutation, we next investigated the anatomy and TTFL molecular pacemaking of the Myk/+ SCN. The SCN exhibit characteristic, spatially distinct neuropeptide topography. Immunohistochemical staining for major SCN neuropeptides, vasoactive intestinal polypeptide and arginine vasopressin, showed no overt genotype differences in the pattern of expression (Supplemental Figure S7). Molecular clock activity was then tracked through expression...
of the circadian clock protein PER2 via PER2::LUC expression in SCN explants (Supplemental Figure S8). The characteristics of circadian rhythms in wheel running seen in Myk/+ mice (increased circadian period and alpha) were maintained in Myk/+PER2::LUC mice (Supplemental Figure S8A–E). Surprisingly, in SCN explants cultured ex vivo, no overt genotype differences were found in period or amplitude of whole-tissue PER2::LUC expression (Figure 7A–C). Similarly, in SCN explants imaged at single-cell resolution, neither the period nor the damping rate of single-cell rhythms nor the intercellular synchrony within SCN slices differed between the genotypes (Figure 7D–F). The persistence of +/+PER2-like rhythms ex vivo indicates that, in isolation, the Myk/+ SCN is a stable pacemaker. Together with the findings from the optic nerve crush experiment, this reinforces the idea that, in vivo, afferent signals such as those coming from the eye act to diminish the Myk/+ SCN’s control of physiology and behavior.

Neurons of the SCN control behavior and physiology in part by varying their spontaneous firing rate (SFR), with higher frequency discharge during the day than at night (49). Because neurophysiological studies suggest that the NKA pump influences SCN neurons (50), and because NKA a2b3 subunits affect membrane excitability (51,52), we next made whole-cell electrophysiological recordings and assessed the electrical activity of +/+ and Myk/+ SCN neurons maintained in brain slices. SCN neurons exhibit distinct electrophysiological states (9,53) and these were evident in both +/+ and Myk/+ SCN recordings, but no genotype differences were detected in most passive properties of these neurons (Supplemental Table S1). However, comparison of the SFR of Myk/+ and +/+ SCN neurons based on the time of recording indicated clear genotype-related differences. Unexpectedly, the mean SFR of Myk/+ SCN neurons did not differ from day to night recordings, whereas +/+ SCN neurons had significantly higher SFR during the day as compared with the night (Figure 7G).

Day/night variation in SFR is a key characteristic of the SCN network both in vitro and in vivo, and because the TTFL finding appears to be intact in the Myk/+ SCN, this damping in Myk/+ SCN neuronal activity most likely arises from exposure to altered photic afferent signals.

**DISCUSSION**

Here we have demonstrated that, in addition to increases in the period and active phase duration of circadian rhythms in behavior, the Myk/+ mouse exhibits instability in behavioral rhythms and unusually heightened circadian resetting/re-entrainment responses to light. Other mouse models expressing mania-like states, including Clock\(^{-/-}\), Reverb\(\alpha^{-/-}\), and DAT\(^{-/-}\) mice, also show heightened phase-shifting responses to light (54–56). Interestingly, in patients with mania, locomotor rhythms may also weaken with increasing severity of manic symptoms (57), and sensitivity to both white and blue light is heightened during manic episodes (58–60). Similar to bipolar patients (61,62), Myk/+ mice exhibit both altered sleep patterns (20) and circadian rhythm disturbance as well as elevated metabolic activity (34). Notably, the nocturnally elevated metabolic rate and locomotor activity of Myk/+ mice were not suppressed by light, indicating the absence of negative masking. Consistent with this, Myk/+ mice did not behaviorally adjust to a long day length, and in LL they increased wheel running, whereas +/+ mice markedly reduced it. Indeed, even in the absence of light, alterations in retinal afferents were sufficient to drive circadian behavioral disruption, because removal of afferent photic input to the SCN in vivo restored the circadian period of behavioral rhythms. Further, when cultured in vitro and assessed in isolation from retinal inputs, Myk/+ single-cell PER2 rhythms and whole-SCN explants behaved as stable pacemakers, comparable to +/+ SCN tissue. This indicates that the SCN molecular clock is largely intact in this mouse model of mania, a finding...
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**Figure 7.** The Myshkin mutation does not affect bioluminescent rhythms of PER2::LUC in whole suprachiasmatic nuclei (SCN) brain slices or single cells, but rather dampens electrophysiological activity. (A) Example luminometric recordings of rhythms of PER2::LUC output of whole SCN tissue explants from +/+ PER2::LUC (+/+ PER2) and Myk/+ PER2::LUC (Myk/+ PER2). (B) No genotype differences were detected in the period of PER2::LUC oscillations (+/+ PER2: 24.73 ± 0.05 hours, Myk/+ PER2: 24.82 ± 0.09 hours; p = .39). (C) Peak amplitude of PER2::LUC rhythms across the first 5 days (24-hour cycles) in culture did not differ between the genotypes (two-way analysis of variance: genotype p = .78; interaction p = .13). (D) Example of single-cell imaging from +/+ PER2 (n = 4) and Myk/+ PER2 (n = 4) explants at two time points over initial 24 hours ex vivo. (E) The period of single-cell rhythms did not differ between the genotypes (+/+ PER2: n = 140, 23.96 ± 0.06 hours; Myk/+ PER2: n = 132, 23.96 ± 0.07 hours; p = .99). (F) The synchrony (R) between single cells in the SCN slices did not differ between the genotypes at any of the 5 days (5 × 24-hour cycles) ex vivo (two-way analysis of variance: genotype p = .70). (G) Mean spontaneous firing rate (SFR) from whole-cell current-clamp recordings made over a 24-hour cycle. Day/night variation in SFR is seen across +/+ SCN neurons but not Myk/+ SCN neurons (two-way analysis of variance: time p = .0079; +/+ day: 2.0 ± 0.2 Hz, +/+ night: 1.2 ± 0.2 Hz, p = .0009; Myk/+ day: 1.6 ± 0.2 Hz, Myk/+ night: 1.4 ± 0.2 Hz; p = .31). Data in panel (E) graphed as scatter plot with mean ± SEM. Data in panels (B, C, F, G) plotted as mean ± SEM. ***p < .001. A.U., arbitrary units; CT, circadian time.

Concordant with the observation that the molecular clock is also intact in fibroblasts from patients with BPD (63), however, Myk/+ SCN slices lacked day/night variation in neuronal firing rate. Importantly, the low amplitude, long period behavioral rhythms, and damped SCN electrical activity of Myk/+ mice resemble similar measures made from +/+ mice exposed to long day lengths or LL (44,64–66). These findings reveal an important role for light inputs to the SCN in this mouse model of mania.

NKA a3 is localized to central and peripheral neurons, including retinal ganglion cells (46,67), and glutamate is the main neurotransmitter of image- and non-image-forming pathways from the eye (12,49). Both SCN and extra-SCN sites are implicated in masking effects of light (68,69) and the absence of negative masking in Myk/+ mice presumably arises as a consequence of altered glutamatergic signaling at these sites. Glutamatergic synapses are linked with mood disorders (70), and it is suggested that measures of glutamate in the brain vary in BPD, becoming elevated during mania and reduced in episodes of depression (71). Further, mood stabilizers used to treat BPD, such as lithium and valproate, act to restore glutamate levels (72,73) and can reduce sensitivity to light and directly alter SCN function (74,75). Interestingly, colocalization and functional coupling between NKA a3 and glutamate transporters has been demonstrated in rat brain, with a3 having similar neuronal localization to excitatory amino acid transporter 2/glutamate transporter 1, the most abundant subtype of glutamate transporter in the central nervous system (76). Previously, Kirshenbaum et al. (20) found that the duration of glutamate-evoked [Ca2+]i transients was prolonged in cultured cortical neurons from Myk/+ mice (20), raising the possibility that this mutation alters glutamatergic signaling in other brain pathways. Because metabolic rate is elevated in Myk/+ mice, it is plausible that the Myshkin mutation influences energy balance centers in the mediobasal hypothalamus (77). In high-firing hippocampal and cerebellar neurons, loss or reduction of NKA a3 is associated with neuronal hyperexcitability (20,51,78,79). Atp1a3 is expressed in the SCN (80), but Myk/+ SCN neurons that have a reduction in functional NKA a3 show damped daytime firing rate. To discharge action potentials, high-firing cells can require considerable adenosine triphosphate, and a likely consequence of a reduction in NKA pump activity is ionic imbalance and chronic depolarization (81). Because individual SCN neurons are comparatively low firing (typically <5 Hz), this suggests that their adenosine triphosphate requirements are low, such that a reduction in NKA a3 activity does not drive the cell into a chronic
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