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Uncovering the Genetic Basis of Sleep: Use of Clock Mutant Mice

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13. ABSTRACT (Maximum 200 words)

Our overall goal is to use the mouse to identify genetic elements controlling sleep. We will use the Clock mutant mouse because of finding that Clock mutants, in addition to their circadian abnormalities, sleep less than wild-type mice.

Over the past year, we have examined how the Clock mutation alters sleep and body temperature rhythms under stressful conditions. Correspondingly, body temperature decreased during the recovery dark period in wild-types but did not change in Clock/Clock mice. Finally, a negative rebound in delta power was observed in wild-type mice, but not in Clock/Clock mice. These results indicate that the Clock gene disrupts the pattern and amount of sleep, as well as the amplitude of the body temperature rhythm under baseline and stressful conditions.

We have also examined sleep in neonatal pups to determine if pups could be used in mutagenesis screening. Our findings in mice, and in other studies in rats, that a simple behavior in the neonate can be used to detect genetic differences observed in adult sleep represent a "proof of principle" that it will be possible to use behavioral states in the neonate to uncover unknown genes involved in the regulation of the sleep-wake cycle of mammals.

Other studies carried out under this proposal are described in the next page.

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2. **Abstracts:**


3. **Scientific personnel**

   P.L.: Fred W. Turek was supported in part by this grant.

   Ph.D. Students: Supported by this grant was Mr. Erik Naylor, while other students worked on this project, but were supported by other grants, including Ms. Amy Easton and Ms. Sara Cronin.

   Research Technologists: Supported by this grant included Ms. Sally Battle, Ms. Susan Olsen, Mr. William Olsen and Ms. Julie Uherka.

   Postdoctoral Fellows/Research Associates: Supported by this grant included Dr. Martha Vitaterna and Dr. Muriel Koehl.

4. **Report of Inventions**

   None.

5. **Scientific Progress and Accomplishments.** The results of three separate sets of studies add to our previous progress reports. These include:

   1. **Clock Mutant Mice Show Altered Sleep and Body Temperature Rhythms Under Baseline and Stressful Conditions.**

   The circadian pacemaker is known to regulate the timing and distribution of sleep; however, the role of specific circadian genes in the regulation of sleep has just begun to be explored. The first mammalian circadian clock gene to be discovered, *Clock*, is highly expressed in the eye and suprachiasmatic nuclei and is an integral part of the pacemaker's molecular machinery. *Mice bred with a mutation of the Clock gene (Clock/Clock) exhibit significantly altered free-running rhythm in running wheel activity. In this study, we were specifically interested in determining whether recovery...
sleep following an acute stress was affected by the *Clock* gene mutation. Six wild-type and 8 *Clock/Clock* male mice bred on a C57/B1/6J background and maintained on a L:D 12:12 cycle. Mice were implanted with EEG and EMG electrodes and a Minimitter transmitter to collect data on body temperature and total locomotor activity. Baseline sleep was collected for 24 hours at least 1 week after recovery from surgery. On day 2, animals were restrained in a plastic tube for 1 hour between ZT3-5. Recovery sleep was recorded for the following 24 hour period immediately after the stress ended. *Clock/Clock* mice showed a decrease in NREM sleep (p<.001,t=4.04) during the light period and twice as much REM sleep during the dark period (p<.001,t=3.6) during baseline recording. In addition, *Clock/Clock* mice had less delta power during the dark period compared to wild-types and the amplitude of the body temperature rhythm was decreased in *Clock/Clock* mice, which corresponded with the changes in NREM and REM sleep across the light/dark cycle (p<.05,t=7.8). Following an acute stressor, *Clock/Clock* mice showed less recovery sleep than wild-type mice (p<.05). Correspondingly, body temperature decreased during the recovery dark period in wild-types but did not change in *Clock/Clock* mice. Finally, a negative rebound in delta power was observed in wild-type mice, but not in *Clock/Clock* mice (p<.01). These results indicate that the *Clock* gene disrupts the pattern and amount of sleep as well as the amplitude of the body temperature rhythm under baseline and stressful conditions.

2. In Mice Restraint Stress at Light Onset Produces an Increase in REM Sleep Whereas Restraint at Dark Onset Does Not.

**Introduction:** Restraint stress induces a specific increase in REM sleep in rodents. However, whereas most physiological, endocrine and behavioral consequences of restraint are strongly dependent on the time of day at which it takes place no study has directly addressed the question whether the REM sleep promoting effect of restraint stress is also time dependent.

**Methods:** Adult male C57BL/6J mice were implanted with permanent electrodes to record EEG and EMG. After at least 2 weeks of recovery, sleep-wakefulness patterns were recorded for 2 days. The first day was a baseline recording and on the second day the mice were subjected to 1h of restraint stress starting at either the onset of the light phase (n=8) or at the onset of the dark phase (n=8).

**Results:** Restraint stress at light onset resulted in a significant increase in REM sleep, especially in the subsequent dark phase. This increase in REM sleep was much larger than the amount of REM sleep that was lost during restraint and therefore could not be attributed to a rebound from sleep deprivation. In contrast, restraint stress at dark onset did not induce a significant increase in REM sleep, neither in the remainder of the dark phase nor in the subsequent light phase.

**Conclusions:** These results support earlier findings that restraint stress can induce an increase in REM sleep in mice. At the same time, however, the data show that this REM sleep inducing effect is strongly dependent on the time of day at which the restraint takes place.

3. Development of Sleep in Mice with Different Genetic Backgrounds and with Altered Sleep Phenotypes in Adults.

Understanding the relationship between sleep-wake states in neonates and adults is critical for any complete understanding of not only the ontogeny of sleep, but also for understanding the regulatory mechanisms underlying the transitions between states of vigilance, as well as for addressing critical questions concerning the function(s) or need for sleep. At the present time, there is considerable controversy as to the relationship between the active sleep (AS) and quiet sleep (QS) states in neonates and the EEG defined states of REM and NREM sleep in adults. Clear genetic differences in the characteristics of the behavioral sleep states in adult animals of different strains of rodents offer a potential window for examining, at least at the genetic level, how the neonatal and adult behavioral states are related to one another.

For this purpose, short-lasting recordings (3-h period during the light phase) of behavioral states of vigilance were performed in neonatal (8-day old) mice of four different strains (C57BL/6ByJ, BALB/cByJ, DBA/2J and C57BL/6J) which show different amounts of vigilance states as adults. Many of these differences were already evident when wake, QS and AS were monitored only via EMG recordings in 8-day old mice. Interestingly, despite the absence of circadian sleep-wake rhythms in neonates, it was also possible to detect sleep-wake differences between strains that appeared to be restricted later in life to the light or the dark phase in adult animals.

The present results indicate that behavioral sleep states in the neonate are under similar genetic control as the REMS and NREMS states in adult animals. While it is not known how neonatal AS and QS are physiologically related to adult REM and NREM sleep, respectively, the finding of similar differences in the amounts of sleep-wake states in neonatal and adult life argues strongly that at some level they are controlled by similar cellular/physiological mechanisms. Furthermore, the fact that these differences between strains of mice can be detected even before EEG defined sleep-wake states can be differentiated, indicates that the neonate can be used to screen for genetic abnormalities of sleep-wake states in a large number of animals. Such a screen is vital in order to use a forward genetics approach to elucidate the genetic
basis for sleep and wake states. Our findings in mice, and in other studies in rats, that a simple behavior in the neonate can be used to detect genetic differences observed in adult sleep represent a “proof of principle” that it will be possible to use behavioral states in the neonate to uncover unknown genes involved in the regulation of the sleep-wake cycle of mammals.

5. Technology transfer
   Not applicable.
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