

Rapid Eye Movement Sleep Is Reduced in Prolactin-Deficient Mice

Ferenc Obál Jr,^{1,2†} Fabio Garcia-Garcia,¹ Balint Kacsóh,³ Ping Taishi,¹ Stewart Bohnet,¹ Nelson D. Horseman,⁴ and James M. Krueger¹

¹Department of Veterinary and Comparative Anatomy, Pharmacology, and Physiology, College of Veterinary Medicine, Washington State University, Pullman, Washington 99164-6520, ²Department of Physiology, University of Szeged, A. Szent-Gyorgyi Medical Center, 6720 Szeged, Hungary, ³Division of Basic Medical Sciences and Department of Pediatrics, Mercer University School of Medicine, Macon, Georgia 31207, and ⁴Department of Molecular and Cellular Physiology, University of Cincinnati, Cincinnati, Ohio 45267-0576

Prolactin (PRL) is implicated in the modulation of spontaneous rapid eye movement sleep (REMS). Previous models of hypoprolactinemic animals were characterized by changes in REMS, although associated deficits made it difficult to ascribe changes in REMS to reduced PRL. In the current studies, male PRL knock-out (KO) mice were used; these mice lack functional PRL but have no known additional deficits. Spontaneous REMS was reduced in the PRL KO mice compared with wild-type or heterozygous littermates. Infusion of PRL for 11–12 d into PRL KO mice restored their REMS to that occurring in wild-type or heterozygous controls. Six hours of sleep deprivation induced a non-REMS and a REMS rebound in both PRL KO mice and heterozygous littermates, although the REMS rebound in the KOs was substantially less. Vasoactive intestinal peptide (VIP) induced REMS responses in heterozygous mice but not in KO mice. Similarly, an ether stressor failed to enhance REMS in the PRL KOs but did in heterozygous littermates. Finally, hypothalamic mRNA levels for PRL, VIP, neural nitric oxide synthase (NOS), inducible NOS, and the interferon type I receptor were similar in KO and heterozygous mice. In contrast, tyrosine hydroxylase mRNA was lower in the PRL KO mice than in heterozygous controls and was restored to control values by infusion of PRL, suggesting a functioning short-loop negative feedback regulation in PRL KO mice. Data support the notion that PRL is involved in REMS regulation.

Key words: stress; VIP; sleep deprivation; rapid eye movement; sleep; prolactin

Introduction

The first anecdotal account implicating prolactin (PRL) in rapid eye movement sleep (REMS) reported that systemic administration of exogenous PRL enhances REMS in hypophysectomized pontine cats (Jouvet et al., 1986). Since then, much evidence suggests that PRL modulates REMS (Roky et al., 1995; Obál and Krueger, 2000). Intracerebral or systemic administration of PRL enhances REMS but not non-REMS (NREMS) in rats, rabbits, and mice (Obál et al., 1989; Roky et al., 1993). Rats rendered chronically hyperprolactinemic, by means of grafting pituitaries under the kidney capsule, exhibit increases in REMS during daylight hours (Obál et al., 1997). Both REMS and NREMS are en-

hanced in hyperprolactinemic pseudopregnant rats (Zhang et al., 1995). In humans, plasma levels of PRL are REMS related and highest in the early morning hours (for review, see Roky et al., 1995; Obál and Krueger, 2000). In animal models of stress, either restraint stress or ether vapor exposure stress, elevated plasma PRL levels are associated with increases in REMS (Bodosi et al., 2000; Meerlo et al. 2001). Vasoactive intestinal polypeptide (VIP) releases pituitary PRL and enhances REMS (Obál et al., 1986, 1994). Similarly, other PRL-releasing peptides such as pituitary adenylate cyclase-activating peptide and peptide histidine methionine also promote REMS (Fang et al., 1995). Anti-PRL antibodies block VIP-enhanced REMS (Obál et al., 1994). Anti-PRL antibodies also reduce spontaneous REMS in rats (Obál et al., 1992) and block REMS induced by systemic VIP (Obál et al., 1994). Collectively, such data suggest PRL may modulate REMS.

Animal models with various PRL deficiencies also suggest a REMS modulating action of PRL. Thus, in mutant hypoprolactinemic rats, REMS is suppressed during the light period but enhanced during the dark period (Lobo et al., 1999). In a mutant mouse model, mice lacking an interferon type I receptor have very low levels of hypothalamic PRL mRNA and approximately a 30% reduction in REMS during both daylight and night hours (Bohnet et al., 2004). However, previously, REMS has not been evaluated in animals lacking PRL. In the current study, we report that REMS is reduced in mice that produce a biologically inactive

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[†]Deceased, June 29, 2004.

Correspondence should be addressed to Dr. James M. Krueger, Department of Veterinary and Comparative Anatomy, Pharmacology, and Physiology, Washington State University, P.O. Box 646520, Pullman, WA 99164-6520. E-mail: Krueger@vetmed.wsu.edu.

F. Garcia-Garcia's present address: Universidad Veracruzana, Instituto de Ciencias de la Salud, Col. Industrial Animas, C.P. 91190, Xalapa, Veracruz, Mexico.

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11 kDa PRL-immunoreactive polypeptide because of a targeted disruption of the PRL gene (Horseman et al., 1997). We also report that REMS is restored in these mice with PRL therapy and that their REMS responses to VIP and ether stress are attenuated.

Materials and Methods

Homozygous wild-type (+/+), heterozygous (-/+), and homozygous PRL knock-out (-/-) male mice, which had been made congenic on C57BL/6J by eight backcross generations, were used. Mice were developed initially by one of us (N.D.H.) (Horseman et al., 1997) and bred by mating (-/+) females with (-/-) males; heterozygous females lactate, thereby indicating that they have sufficient PRL for that purpose, and it allows them to be used for breeding without having to use surrogate mothers. Mice were genotyped at 3–4 weeks of age from tail-tip samples as described previously (Horseman et al., 1997). At 8–16 weeks of age, mice (25–30 g) were housed individually in Plexiglas cages placed in environmental chambers on a 12 h light/dark cycle at 30°C ambient temperature. Food and water were available continuously. Breeding and experimental protocols were approved by the Washington State University Animal Care and Use Committee.

Surgery

The mice were anesthetized using ketamine–xylazine (87 and 13 mg/kg, respectively). Stainless steel screw electrodes were placed on the top of the dura mater over the frontal and parietal lobes and the cerebellum. These electrodes were used to record the electroencephalogram (EEG). Two stainless steel wires were implanted into the dorsal neck muscles to record the electromyogram (EMG). The EEG and EMG electrodes were connected to a pedestal (0.4 g) implanted on the top of the skull. The surgeries were performed 7 d before recordings began.

Recording of sleep–wake activity

The mice were connected to lightweight recording cables and habituated to the experimental conditions during recovery (Bohnet et al., 2004). The cables were attached to commutators that were connected to amplifiers. The signals were digitized (128 Hz sampling rate) and collected by a computer and stored on compact discs. Power density spectra (0.25–15.0 Hz in 0.5 Hz bands) were calculated on-line for consecutive 10 s EEG epochs and stored with the EEG and EMG signals. For scoring, the EEG and EMG signals and the power density spectra were restored on the computer screen. The states of vigilance were determined for 10 s epochs (Morrow and Opp, 2005) by the usual criteria as NREMS (high-amplitude EEG slow waves and low muscle tone), REMS (highly regular theta EEG activity and loss of muscle tone with occasional twitches), and wakefulness (EEG activities similar to, but often less regular and with lower amplitude than, those in REMS and high EMG activity). The percentage of the time spent in each state of vigilance for 1 h periods was determined. The power values for the 0.5–4.0 Hz (Δ) frequency range were integrated. The mean of these integrated values was calculated for uninterrupted periods of artifact-free NREMS and used as an index of EEG slow-wave activity (SWA) during NREMS to characterize sleep intensity (depth of sleep) in each recording hour (EEG SWA data shown in Fig. 4). The absolute power density values (squared microvolts) were determined for 0.5 Hz frequency bins from 0–15 Hz during both NREMS and REMS. These values were then expressed as the percentage of total power (0–15 Hz) in each 0.5 Hz frequency bin; these values are shown in Figure 2.

Experimental schedule

Spontaneous sleep. Spontaneous sleep was determined in two separate experiments. First, wild-type mice [(+/+); $n = 10$] and heterozygous mice [(+/-); $n = 12$] were compared. In a second experiment, another group of (+/-) mice ($n = 10$) was used for comparison with PRL knock-out mice [(-/-); $n = 10$]. Most of the latter two groups were also used for PRL replacement (see below) and sleep deprivation (SD). Mice were recorded from for 2 consecutive days to obtain baseline values of spontaneous sleep–wake activity. On the third experimental day, the (+/-) and (-/-) mice were deprived of sleep. SD started at light onset (time 0 in the figures) and lasted for 6 h. The EEG and EMG were recorded

throughout SD, during the remaining 6 h of the light period, and for 11 h during the subsequent dark phase (collected data were backed up during the last hour of the dark period, and the records in this hour were discarded). SD was performed by gentle handling while the mice stayed in their home cage; whenever behavioral or EEG signs of sleep were observed, the mice were aroused by knocking on the cage or lightly touching them. It is assumed that stress is not a major factor determining the sleep response to SD, at least when SD is performed by gentle handling (Jouvet, 1994), although sleep loss itself may represent a stressor and may alter hypothalamo–pituitary–adrenal functions (Krueger et al., 1990).

PRL replacement. The mice used in the PRL replacement experiment [$n = 6$ (+/-) mice, $n = 10$ (-/-) mice] were implanted subcutaneously with osmotic mini-pumps (0.25 μ l/h for 14 d; Alzet 1002; Durect Corporation, Cupertino, CA) in the back (weight after loading, 0.5 g) (the same mice were used for spontaneous sleep determination; the ALZET pumps were implanted during a second surgery after spontaneous sleep was determined). The minipumps were filled with mouse PRL in physiological saline delivering 4.6 μ g of PRL per day in (-/-) mice. For (+/-) mice, saline was infused. Spontaneous sleep–wake activity of the mice implanted with the minipumps was recorded on day 11 and/or 12 of the infusion. The PRL preparation was for *in vivo* use and was a gift from Dr. A. F. Parlow (The National Hormone and Pituitary Program, National Institute of Diabetes and Digestive and Kidney Diseases, Bethesda, MD). After recordings were obtained, mice were killed and blood samples were taken for determination of PRL levels; only those mice showing circulating PRL were used for sleep and statistical analyses.

VIP treatment. Twelve (+/-) and 12 (-/-) mice received intraperitoneal injection of physiological saline (0.1 ml/10 g) at light onset as a control. On a separate day, VIP was administered at the same time of day at a dose of 0.3 μ g/g in a volume of 10 μ l/g body weight. An EEG and EMG were recorded for the next 23 h.

Ether stress. The mice [$n = 12$ (+/-), $n = 12$ (-/-)] were placed individually into a jar (10 L) containing ether vapor for 1 min (20 min before light onset) and then returned to their recording cage. A cotton cloth soaked with ether was in the bottom of the jar, and it was separated from the mouse via a plastic plate with holes. The lid of the jar was closed. The mice exhibited motor uncoordination near the end of the exposure, but they were still standing or rearing; anesthesia did not occur. As a control treatment, mice were placed in an identical jar that did not contain ether vapor. After ether exposure or control exposure, mice were recorded from for the next 23 h.

Determination of plasma PRL bioactivity

After (-/-) mice receiving PRL or (+/-) mice receiving saline were killed, trunk blood was taken to determine PRL bioactivity by means of the Nb2 lymphoma assay as described previously (Kacsóh et al., 1993; Kacsóh, 1997).

Determination of hypothalamic mRNA by reverse transcription-PCR

Mice were killed 2–4 h after light onset. Hypothalami were dissected from 12 heterozygous (+/-) and 12 knock-out (-/-) mice as described previously (Bohnet et al., 2004). Hypothalami from the (-/-) and (+/-) mice that received PRL replacement were also used. Total RNA was extracted by using Trizol reagent according to the manufacturer's protocol (Invitrogen, San Diego, CA). The RNAs were resuspended in RNase-free water and incubated with 2–4 U of DNase–RNase free and 1 μ l of SuperRNase In (Ambion, Austin, TX) at 37°C for 1 h, 30 min. RNA concentration was quantified by spectrophotometry at 260 nm. RNA integrity was verified by electrophoresis run on an agarose denaturing gel and subsequent visualization.

The synthesis of first-stranded cDNA was described in detail previously (Bohnet et al., 2004; Chen et al., 2004). Briefly, aliquots of RNA (2 μ g/4 μ l), 1 μ l of oligodT, and 1 μ l of 10 mM deoxynucleoside triphosphates were incubated at 65°C for 5 min. After chilling on ice, 200 U of Superscripts III RNase H reverse transcriptase (Invitrogen), 1 μ l of 0.1 M DTT, 1 μ l of RNase inhibitor, and 4 μ l of 5 \times cDNA synthesis buffer were added to a final volume of 20 μ l and incubated at 55°C for 60 min. Then the mixture was heated at 70°C for 15 min and digested with 1 μ l of

Table 1. Primers used for quantitative real-time, reversed-phase PCRs

Gene	GenBank	Primer	Sequence 5'–3'
Orexin	NM_010410	Sense	GCCGTCTCTACGAACTGTTC
		Antisense	CGCTTCCCAGAGTCAGGATA
Tyrosine hydroxylase	NM_009377	Sense	CTTCCAATACAAGCAGGGTGAG
		Antisense	TCCTCCAGGTAGCAATTTCC
NOS-1	NM_008712	Sense	GAAACTCTCGGAGGAGGATGC
		Antisense	CTCAGATCTAAGGCGGTGGTC
NOS-2	U43428	Sense	CAGGAAGAAATGCAGGAGATGG
		Antisense	GATGTCTGAACGTAGACCTTGG
VIP	NM_011702	Sense	CCTCACTACAGAAGGTGGTCCAAAGAGAGGCC
		Antisense	GAAGCCAGAAGCAAGCCTCAGTCTCTGGCA
PRL ^a	NM_011164	Sense	TGACTGCCAGACTTCTCTCCG
		Antisense	GACCATAAACTCAGGCTTTC
IFN type I receptor	NM_010508	Sense	GGCTAAATGTTGCAGGTTCC
		Antisense	ATTAAGCGTGGCACCAGAT
Cyclophilin A	NM_008907	Sense	AAATGCTGGACCAACACAAC
		Antisense	TTGATGCCTTCTTCCCTTC

^aThe primers for PRL amplify a product in the 5' region (161–290), and thus a product is amplified in the (–/–) mice (Horseman et al., 1997).

RNase H at 37°C for 20 min. Finally, cDNA was cooled to room temperature and stored at –20°C until additional analysis.

Real-time PCR was performed on an iCycler iQ Detection System (Bio-Rad, Hercules, CA) as described previously (Bohnet et al., 2004; Chen et al., 2004). The reaction mixture (25 μ l) contained 3 μ l of diluted cDNA (6 ng), 12.5 μ l of 2 \times Platinum Quantitative PCR SuperMix-UDG (Invitrogen), 0.25 μ l each of SYBR Green (1:1000 dilution) and fluorescein (1:1000 dilution), and 0.5 μ l of forward and reverse primers at 10 μ M. Primers were designed using Primer3 (www.genome.wi.mit.edu/cgi-bin/primer/primer3_www.cgi), and, if possible, an intron was spanned. A standard curve and efficiency analysis of the primers were performed, and the primer efficiencies were >90%. The primers used are listed in Table 1.

To activate uracil DNA glycosylase, the reactions were performed with one cycle at 50°C for 2 min and 95°C for 2 min. The amplification was followed by 40 cycles of denaturation at 94°C for 15 s, annealing at 58°C for 15 s, and extension at 72°C for 15 s. Finally, a melting curve was generated by stepwise increasing temperature (0.5°C increase every 10 s) for 80 cycles starting at 55°C. If multiple peaks were observed in melt-curve analysis, the data were not used. The threshold cycle (Ct) was determined using SYBR Green fluorescence.

The real-time PCRs were performed in triplicate, and average values were reported as Ct values. The Δ Ct values were determined by subtracting the reference cyclophilin A Ct values from the target gene Ct values. Relative gene expression was quantified with a comparative Ct method (User Bulletin #2 ABI Prism 7700 sequence detection system; Applied Biosystems, Foster City, CA). The fold change between the mRNA expression levels of the experimental samples (knock-out mice) and the control samples (heterozygous mice) was calculated (Table 2).

Statistical analyses

For sleep parameters, two-way ANOVA was used to compare hourly values of NREMS, REMS, and EEG SWA among the various groups and between baseline and experimental days, when appropriate. The two factors of the ANOVA were time effects and treatment effects (or group effects-independent samples). If significant differences were indicated, the Bonferroni's (all-pairwise) multiple comparison test was used. PCR data were analyzed by Student's *t* tests.

Results

Spontaneous sleep

Spontaneous NREMS was similar in all three groups [PRL (+/+), (+/–), and (–/–)] with no statistically significant differences (Fig. 1). In each group, NREMS was greater during daylight hours and lower during the night with a pronounced nadir of sleep at dark onset. The duration of REMS was similar between

the wild-type mice [(+/+)] and the heterozygous mice [(+/–)]. As with NREMS, REMS was greater during the daylight hours and less at night. The values of NREMS and REMS duration and their day–night distribution are similar to those reported previously for 129Sv mice (Bohnet et al., 2004), the background strain for the mice used. In contrast to these results, the duration of REMS in the PRL-deficient (–/–) mice was significantly less ($F_{(1,18)} = 28.47$; $p \ll 0.01$) during the day time compared with those amounts observed in (+/–) mice (Fig. 1, right). During the night, the duration of REMS was similar in the (+/–) and (–/–) mice. The EEG power spectra during REMS and NREMS in the (+/–) mice was similar to that observed in the (–/–) mice whether calculated during the sleep period (light) or for the entire 23 h recording period (Fig. 2).

PRL replacement

The duration of REMS was restored to 98.9% of (+/–) mice in those (–/–) mice that received PRL replacement for 10–11 d before recordings (Fig. 3). Thus, PRL replacement significantly increased REMS during the daylight hours in the (–/–) mice ($F_{(1,14)} = 25.34$; $p < 0.01$). These mice did not exhibit increases in REMS during dark hours, nor was their NREMS affected by this treatment. Heterozygous mice [(+/–)] receiving saline did not alter their sleep patterns from the baseline conditions (data not shown).

SD

In (+/–) mice, after 6 h of SD, there was an increase in the duration of REMS ($F_{(1,18)} = 14.39$; $p < 0.01$) and NREMS ($F_{(1,18)} = 12.70$; $p < 0.01$) during the remaining hours of the light period (Fig. 4). In the subsequent dark period, these (+/–) mice had NREMS values similar to those obtained from the same mice under control conditions ($F_{(1,18)} = 1.38$; $p < 0.25$). In contrast, in the (+/–) mice, REMS increased in the dark period after SD ($F_{(1,18)} = 6.86$; $p < 0.02$).

In (–/–) mice, 6 h of sleep loss also induced an increase in the duration of NREMS in the light period immediately after the deprivation period ($F_{(1,18)} = 12.35$; $p < 0.01$), but this increase did not extend into the dark period ($F_{(1,8)} = 0.40$; $p > 0.05$). SD of (–/–) mice also induced an increase in REMS in the light period ($F_{(1,18)} = 5.68$; $p < 0.05$) but not in the subsequent dark period ($F_{(1,18)} = 0.01$; $p > 0.9$). Averaged for the 17 h immediately after sleep loss, (+/–) mice increased REMS by 21 min ($F_{(1,18)} = 15.69$; $p < 0.01$), whereas the (–/–) mice only increased REMS by 8 min ($F_{(1,18)} = 1.34$; $p > 0.25$) above the amount of REMS that occurred during control conditions (57 vs 43 min, respectively). In a direct comparison of the amount of REMS occurring in (+/–) and (–/–) mice, there was significantly less REMS in the (–/–) mice after deprivation in hours 7–23 (Fig. 4) than in (+/–) mice ($F_{(1,16)} = 18.13$; $p < 0.003$).

EEG SWA during NREMS increased in both (+/–) and (–/–) mice in the first 4–6 h after sleep loss. After that, EEG SWA decreased compared with control values (Fig. 4). These responses were similar in both (+/–) and (–/–) mice.

VIP treatment

VIP induced significant increases in REMS in the (+/–) mice during the daylight hours (treatment: $F_{(1,11)} = 6.334$, $p < 0.05$; interaction: $F_{(11,121)} = 2.882$, $p < 0.05$). VIP had no significant effect on duration of REMS in the subsequent dark period in the (+/–) mice, nor did it alter the duration of NREMS in these mice. In contrast, in (–/–) mice, VIP failed to affect the duration

Table 2. Hypothalamic mRNA levels in heterozygous [(+/-)] and homozygous [(-/-)] mutant mice lacking PRL and in (-/-) mice receiving PRL and (+/-) mice receiving saline

Spontaneous difference (fold increase)	Orexin	Tyrosine hydroxylase	NOS-1	NOS-2	VIP	PRL	Interferon type I receptor
(-/-)	0.92 ± 0.042 ^a	0.77 ± 0.54	0.98 ± 0.297	0.85 ± 0.106	0.89 ± 0.106	4.45 ± 1.797	0.92 ± 0.042
(+/-)	1.01 ± 0.050	1.05 ± 0.109	1.03 ± 0.071	1.08 ± 0.145	1.04 ± 0.086	2.67 ± 1.288 ^a	1.01 ± 0.050
<i>p</i>	0.141	0.028*	0.622	0.153	0.262	0.411	0.141
Fold increase after PRL replacement	Orexin	Tyrosine hydroxylase	NOS-1	NOS-2	VIP	PRL	Interferon type I receptor
(-/-)	0.97 ± 0.094	1.29 ± 0.170	1.03 ± 0.051	1.04 ± 0.069	0.63 ± 0.103	5.22 ± 2.047	1.17 ± 0.63
(+/-)	1.05 ± 0.106	1.12 ± 0.176	1.05 ± 0.093	1.03 ± 0.083	1.05 ± 0.109	1.30 ± 0.486	1.05 ± 0.100
<i>p</i>	0.573	0.481	0.860	0.936	0.008*	0.096	0.289

^aValues are ΔΔCt values (see Materials and Methods) ± SEM. Variance in the PRL values was high, therefore values for (+/-) mice are not close to 1.0.

**p* < 0.05.

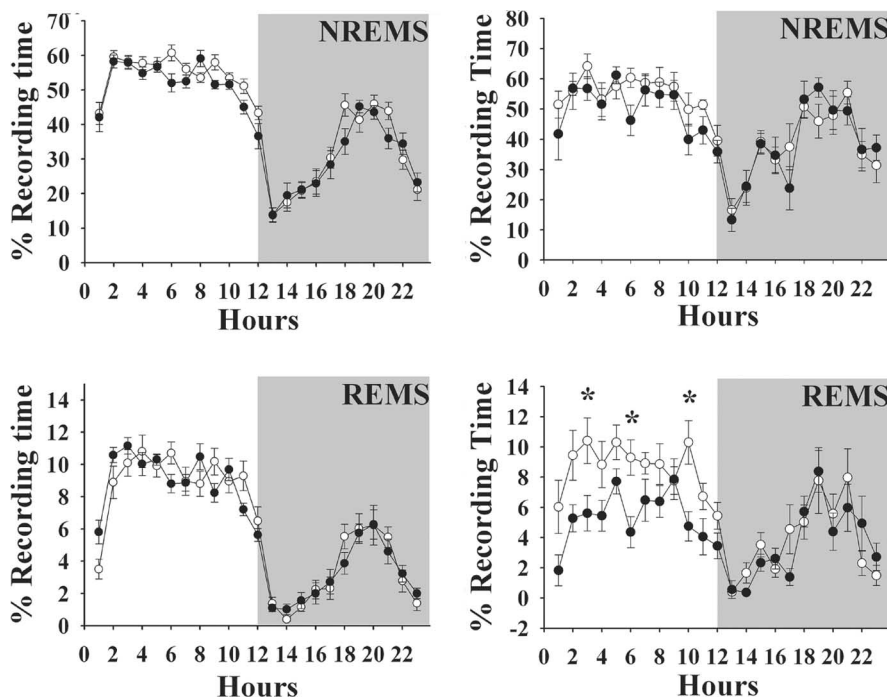


Figure 1. Comparison of sleep patterns between wild-type (*n* = 10; ○) and heterozygous (*n* = 12; ●) (left graphs) littermates and between heterozygous mice (*n* = 11; ○) and PRL knock-out mice (*n* = 10; ●) (right graphs). The top graphs show the percentage of time spent in NREMS, and the bottom graphs show the percentage of time spent in REMS. Sleep duration and its distribution across the day–night cycle were similar in the wild-type and heterozygous mice. In contrast, PRL knock-out mice had substantially less REMS, although day–night differences persisted. Means ± SEM are shown. The asterisks indicate significant difference (*p* < 0.05; *post hoc* test).

of REMS during the daylight or night hours, nor did it alter NREMS (Fig. 5).

Ether stress

Ether stress enhanced duration of REMS during the daylight hours immediately after exposure in the (+/-) mice (treatment: *F*_(1,11) = 11.479; *p* < 0.01). The increases in REMS became manifest ~4 h after exposure to ether and persisted for approximately another 8 h. In the subsequent dark period, REMS was near control values. Ether stress did not affect duration of NREMS. In the (-/-) mice, the ether stressor failed to affect any parameters of sleep. Thus, duration of REMS in the (-/-) mice was less than that observed in the (+/-) mice, but it did not increase after ether vapor exposure (Fig. 6).

Hypothalamic gene expression

There were no differences in mRNA levels between (+/-) and (-/-) mice for orexin, nitric oxide synthase 1 (NOS-1), NOS-2, VIP, PRL, and the interferon type I receptor; all of these substances are implicated in REMS regulation (Table 1). In contrast, tyrosine hydroxylase mRNA, the enzyme needed for dopamine synthesis, was significantly less in the (-/-) mice compared with the (+/-) mice. After PRL replacement in (-/-) mice, tyrosine hydroxylase mRNA was not significantly different from that observed in (+/-) mice, although after this treatment, VIP mRNA significantly decreased in the (-/-) mice.

Discussion

The major finding reported here is that PRL-deficient mice have less spontaneous REMS during the daylight hours than wild-type or heterozygous littermates. Furthermore, PRL replacement in the (-/-) mice results in REMS indistinguishable from that observed in heterozygous littermates. These data suggest a role for PRL in REMS regulation and are consistent with previous data that demonstrate the REMS-promoting activity of acute administration of PRL (Obál et al., 1989; Roky et al., 1993, 1994) and the excess REMS in rats rendered chronically hyperprolactinemic by means of pituitary grafts (Obál et al., 1997). In contrast, a hypoprolactinemic mutant rat strain exhibited increases in REMS with more REMS during the dark phase than in the light phase (Lobo et al., 1999). These rats have an altered sensitivity to photo-enhancement signals as well as other deficits such as being hairless, and thus it is not possible to directly link their changes in REMS to reduced PRL levels. However, in the male PRL knock-out mice used in the current study, there are no known deficits other than the lack of functional PRL and reduced REMS duration reported here.

Although previous observations suggested a REMS-promoting activity for PRL, it was not clear whether PRL was involved in the REMS rebound after sleep loss. Current results clearly indicate that PRL-deficient mice exhibit a REMS rebound, although it is attenuated compared with heterozygous mice. This

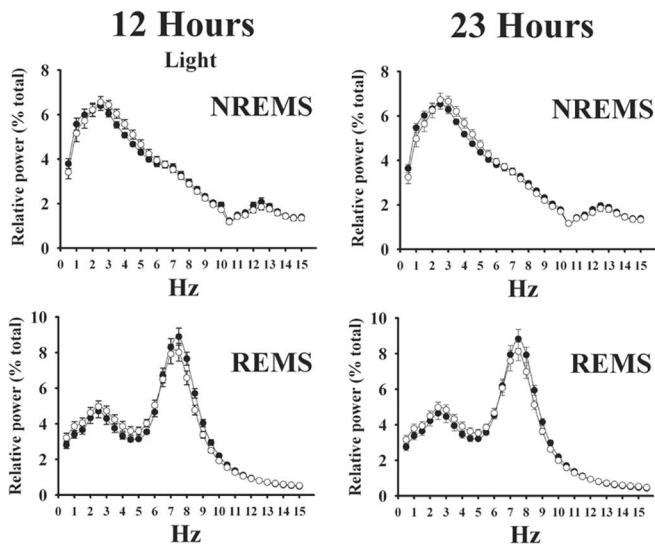


Figure 2. Relative EEG power (squared microvolts) expressed as a percentage of total power in heterozygous [(+/-); ○] and PRL knock-out [(-/-); ●] mice during the 12 h light period (left graphs) and the entire 23 h recording period (right graphs). The top graphs show relative EEG power during NREMS; characteristic patterns were observed in both (+/-) and (-/-) mice, with the greatest power in the low-frequency ranges. The bottom graphs show relative EEG power during REMS; both groups of mice exhibited the theta peak characteristic of REMS. There were no differences between (+/-) and (-/-) mice. Values are means \pm SEM.

suggests that PRL may enhance a REMS-promoting mechanism rather than be an initiator of REMS.

Jouvet (1994) suggested that stimulation of pyruvate dehydrogenase (PDH) in mesencephalic/pontine cholinergic neurons may serve as a final common pathway for various interventions that promote REMS. PRL stimulates PDH activity in the prostate (Costello et al., 1995) and the mammary gland (Waters and Rillemma, 1988). Enhanced PDH activity results in increased acetyl-coenzyme A and thereby stimulates the synthesis of acetylcholine (ACh). PDH is particularly enriched in cholinergic neurons (Milner et al., 1987; Rozanov and Parkhomenko, 1987). Furthermore, brain cholinergic neurons are vulnerable to the deficiency of PDH (Ruenwongsa and Pattenavibag, 1984). ACh levels are decreased by the PDH inhibitor 3-bromopyruvate in brain homogenates or in the basal forebrain after local injection of this compound (Arendt et al., 1990). Decreases in PDH activity elicited by experimental thiamine deficiency, are also associated with the suppression of REMS; thiamine is a coenzyme for PDH (Crespi and Jouvet, 1982). We propose that because PRL stimulates PDH in cholinergic neurons whenever spontaneous REMS occurs via cholinergic neurons, the quantity of ACh released increases, thereby facilitating spontaneous REMS but not initiating it. There is thus no need to assume that PRL acts via the suprachiasmatic nucleus on REMS as suggested previously by Roky et al. (1993, 1994). PRL is more effective in stimulating REMS during the light period than at night, simply because more spontaneous REMS occurs during the light than during the dark period.

Intracerebral administration of PRL or antibodies to PRL stimulates or inhibits REMS, respectively (Roky et al., 1993, 1994), suggesting a central action of PRL. PRL has several sources, including hypothalamic and brainstem neurons (Bole-Feysot et al., 1998; Freeman et al., 2000; Bohnet et al., 2004; Chen et al., 2004) and the anterior pituitary. Circulating PRL is transported into the brain via the choroid plexus (Walsh et al., 1987), and circulating PRL promotes REMS. Thus, systemically injected PRL (Jouvet et al., 1986; Obál et al., 1989), excess PRL released

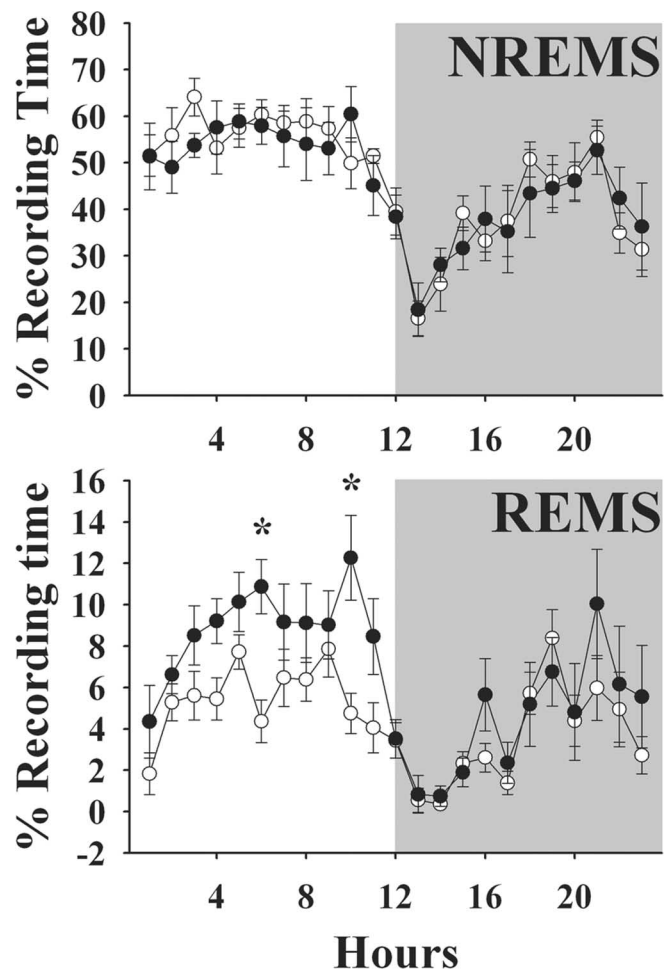


Figure 3. PRL replacement in PRL knock-out mice restores REMS. PRL was provided to PRL (-/-) mice via a mini-osmotic pump for 11–12 d before recordings were done (●). Values from (-/-) knock-out mice (○) are those from the mice shown in Figure 1 (right graphs). PRL replacement had no effect on the duration or diurnal distribution of NREMS; it did, however, enhance REMS during daylight hours in the (-/-) mice. Means \pm SEM are shown. The asterisks indicate significant difference ($p < 0.05$; *post hoc* test).

from pituitary grafts (Obál et al., 1997), or stimulation of endogenous PRL secretion from the pituitary (Obál et al., 1994) all promote REMS. Nevertheless, only slight decreases in REMS are observed when basal, nonstimulated circulating PRL is immunoneutralized (Obál et al., 1992). Although plasma PRL levels in humans are sleep related and highest in the early morning hours, patients with PRL-producing adenomas fail to display changes in REMS (Frieboes et al., 1998).

Hypothalamic PRL-containing neurons may innervate various structures in the brain including the brainstem (Paut-Pagano et al., 1993). However, the anti-ovine PRL antibody used in those rat studies also recognizes hypocretin (orexin), thereby suggesting that lateral hypothalamic immunoreactive prolactinergic neurons may, in fact, be hypocretinergic neurons (Risold et al., 1999). Regardless, the site of action where PRL modulates REMS is not clear. Indeed, when injected into the central amygdalar nucleus, PRL fails to affect REMS but inhibits NREMS (Sanford et al., 1998). Valatx et al. (1990) and Roky et al. (1994) suggest that PRL interferes with regulation of the diurnal rhythm of REMS, perhaps via PRL receptors in the suprachiasmatic nucleus. Their hypothesis, however, disregards the observation that PRL enhances REMS in pontine cats (Jouvet et al., 1986), and it

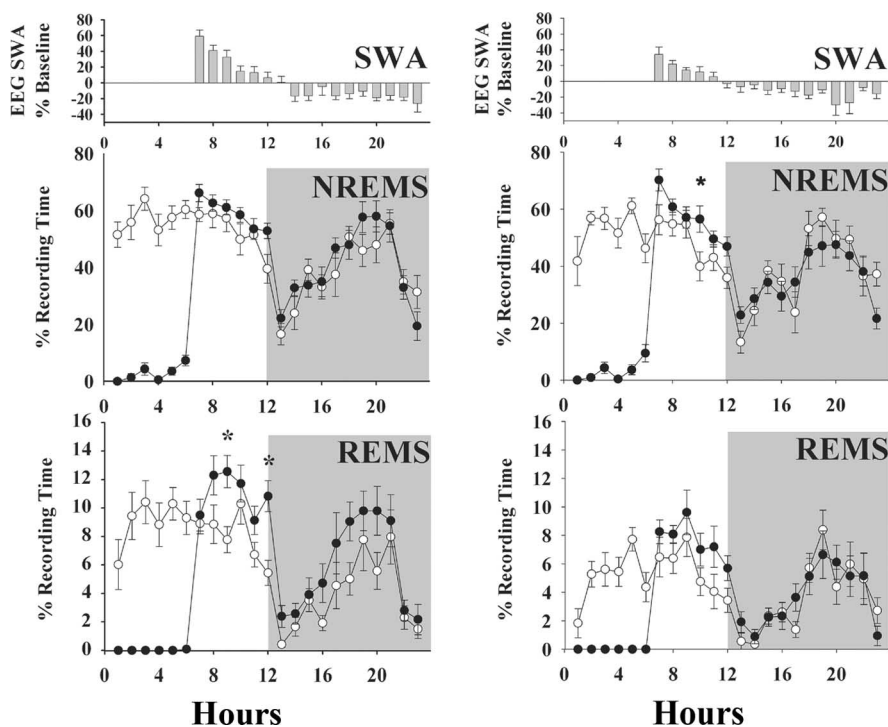


Figure 4. SD (6 h beginning at time 0) of heterozygous [(+/-); left graphs] and PRL knock-out [(-/-); right graphs] mice induces NREMS and REMS rebounds in both types of mice. The extent and timing of the NREMS rebound was similar in both (+/-) and (-/-) mice. However, the REMS rebound was limited to daylight hours, and its magnitude was less in the (-/-) mice compared with littermates [(+/-) mice]. Values are means ± SEM from a baseline control day (○) and the SD day (●). EEG SWA responses were similar in both sets of mice and were characteristic of that seen in other mice and rats initially characterized by an enhancement then followed by a decrease in EEG SWA. The asterisks indicate significant difference ($p < 0.05$; *post hoc* test).

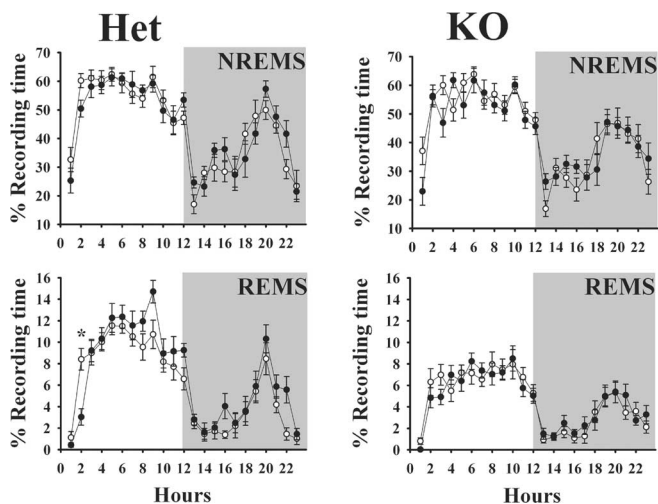


Figure 5. VIP enhances REMS in heterozygous mice [Het; (+/-); left graphs] but not in PRL knock-out mice [KO; (-/-); right graphs]. VIP (0.3 μg/g body weight) was injected intraperitoneally at light onset (time 0). The mice used for these experiments were different from those used in other experiments; the lowered spontaneous REMS in the (-/-) mice is evident by comparing the right and left graphs. ○, Control values; ●, values obtained after VIP. Values are means ± SEM. The asterisks indicate significant difference ($p < 0.05$; *post hoc* test).

does not consider subsequent evidence suggesting a brainstem PRL–nitric oxide (NO) interaction (Chen et al., 2004). PRL immunoreactivity, PRL mRNA, and PRL receptor mRNA are found in the brainstem (Harlan et al., 1989; Siaud et al., 1989; Freemark et al., 1995; Bole-Feysot et al., 1998). Therefore, we prefer a mech-

anism involving the brainstem cholinergic REMS-promoting mechanism, although the cellular origins of brainstem PRL and PRL receptors remain unknown.

Inbred mouse strains display significant variations in the EEG, particularly in the theta peak frequency, during both REMS and NREMS, suggesting the presence of genes influencing theta activity (Franken et al., 1998; Tafti et al., 2003). For instance, deficiency in the gene of the short-chain acyl-coenzyme A dehydrogenase (*Acads*) results in slowing of theta activity during REMS (Tafti et al., 2003). Nevertheless, the PRL-deficient mice used in this study exhibited a normal EEG power spectrum, suggesting that PRL is not directly involved in those mechanisms responsible for EEG power. This conclusion is consistent with the indirect role proposed above for PRL involvement in REMS regulation.

PRL-induced NO might be part of the REMS-promoting activity of PRL. There is involvement of NOergic neurons in the laterodorsal tegmental and pedunculo tegmental nuclei (LDT/PPT) in REMS regulation. PRL excites LDT neurons (Takahashi et al., 2000), and LDT neuron activation increases NO production and elevates extracellular NO (Leonard et al., 2001). PRL induces inducible NOS (iNOS) mRNA in glioma cells *in vitro* (Raso et al., 1999) and neuronal NOS (nNOS) protein expression in neurons in the paraventricular and supraoptic nuclei *in vivo* (Popeski et al., 2003). PRL receptor activation upregulates the expression of nNOS protein in GH₃ rat pituitary adenoma cells (Secondo et al., 2003). PRL also increases NO production in other cell types [e.g., immune cells (Kumar et al., 1997; Meli et al., 1997) and epithelial cells (Bolander 2002)]. NO reduces intracellular calcium through cGMP (Esplugues, 2002) and modulates cAMP signaling (Klein, 2002). The reduction in intracellular calcium or cAMP levels inhibits PRL gene expression in pituitary cells (Liu et al., 2002). Thus, NO may also form a feedback loop to inhibit PRL when the source of NO is close to PRL-containing cells.

We failed to observe differences in either iNOS or nNOS mRNA levels in the hypothalamus. We also failed to observe differences in mRNAs of other substances linked previously to REMS regulation, including hypothalamic VIP, interferon type I receptor, and PRL mRNAs. Such data are consistent with the notion that the actions of PRL on REMS are indirect, involving perhaps the brainstem PDH–ACh mechanism mentioned above. In contrast to these negative findings, the levels of tyrosine hydroxylase mRNA decreased in the (-/-) mice compared with their (+/-) littermates. Tyrosine hydroxylase is an enzyme involved in dopamine synthesis. Pituitary PRL release is under a continuous dopamine inhibition. Because the (-/-) mice are not producing functional PRL, there is a reduced need to inhibit pituitary release of PRL, and this is manifest as a reduced level of hypothalamus tyrosine hydroxylase. This observation is consistent with the established short-loop negative feedback regulation of PRL (Perkins et al., 1979; Reymond, 1990)

The lack of an effect of ether vapor exposure on REMS in the

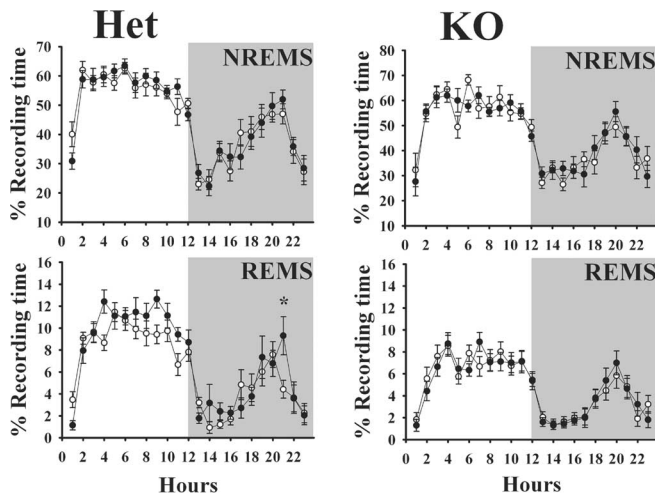


Figure 6. Ether stress enhances REMS in heterozygous mice [Het; (+/–); left graphs] but not in PRL knock-out mice [KO; (–/–); right graphs]. Mice were exposed to ether vapors for ~1 min during the last 20 min of the dark period (light onset; time 0). Ether stress did not affect NREMS. The mice used for these experiments were different from those used in other experiments. Spontaneous sleep values again illustrate that (–/–) mice have less REMS than littermate [(+/–)] mice. ○, Control values; ●, values obtained after ether stress. Values are means ± SEM. The asterisks indicate significant difference ($p < 0.05$; *post hoc* test).

(–/–) mice is consistent with the notion that stressor-enhanced REMS is mediated in part via PRL. Restraint stress and ether stress are associated with enhanced circulating PRL and enhanced REMS (Bodoss et al., 2000; Meerlo et al., 2001). Restraint stress also enhances hypothalamic release of PRL (Torner et al., 2004), although stressor-induced release of pituitary PRL is thought to be responsible for the increases in REMS after stress (Bodoss et al., 2000; Meerlo et al., 2001).

In conclusion, data reported here suggest that PRL plays a role in the regulation of spontaneous REMS and REMS responses to SD, systemic VIP, and an ether stressor. These and other data also suggest that these REMS-promoting actions of PRL are indirect, perhaps operating via a brainstem LPT/PPT–PDH cholinergic NO mechanism.

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