

Circadian *Clock* Mutation Disrupts Estrous Cyclicity and Maintenance of Pregnancy

Brooke H. Miller,^{1,2} Susan Losee Olson,¹
Fred W. Turek,^{1,2} Jon E. Levine,^{1,2}
Teresa H. Horton,¹ and Joseph S. Takahashi^{1,2,3,*}

¹Department of Neurobiology and Physiology

²Northwestern University Institute for Neuroscience
Northwestern University
Evanston, Illinois 60208

³Howard Hughes Medical Institute
4000 Jones Bridge Road
Chevy Chase, Maryland 20815

Summary

Classic experiments have shown that ovulation and estrous cyclicity are under circadian control and that surgical ablation of the suprachiasmatic nuclei (SCN) results in estrous acyclicity in rats [1–3]. Here, we characterized reproductive function in the circadian *Clock* mutant mouse [4, 5] and found that the circadian *Clock* mutation both disrupts estrous cyclicity and interferes with the maintenance of pregnancy. *Clock* mutant females have extended, irregular estrous cycles, lack a coordinated luteinizing hormone (LH) surge on the day of proestrus, exhibit increased fetal reabsorption during pregnancy, and have a high rate of full-term pregnancy failure. *Clock* mutants also show an unexpected decline in progesterone levels at midpregnancy and a shortened duration of pseudopregnancy, suggesting that maternal prolactin release may be abnormal. In a second set of experiments, we interrogated the function of each level of the hypothalamic-pituitary-gonadal (HPG) axis in order to determine how the *Clock* mutation disrupts estrous cyclicity. We report that *Clock* mutants fail to show an LH surge following estradiol priming in spite of the fact that hypothalamic levels of gonadotropin-releasing hormone (GnRH), pituitary release of LH, and serum levels of estradiol and progesterone are all normal in *Clock/Clock* females. These data suggest that *Clock* mutants lack an appropriate circadian daily-timing signal required to coordinate hypothalamic hormone secretion. Defining the mechanisms by which the *Clock* mutation disrupts reproductive function offers a model for understanding how circadian genes affect complex physiological systems.

Results and Discussion

Clock Mutant Females Exhibit Defects in Estrous Cyclicity and Proestrus LH Release

The suprachiasmatic nuclei (SCN) of the hypothalamus coordinate circadian physiology and behavior by functioning as the master pacemaker in a hierarchical system of multiple circadian oscillators; the SCN receive photic input from retinal ganglion cells and subsequently phase-

coordinate the activity of tissue-specific oscillators via neuronal and humoral output [6]. Circadian output from the SCN plays a major role in the regulation of female reproduction. Previous studies in rats have shown that disruption of communication between the SCN and the gonadotropin-releasing hormone (GnRH) neurons responsible for regulating reproductive function—either by ablating the SCN or by severing the neuronal pathways between the SCN and the preoptic area—results in estrous acyclicity and infertility [2, 3]. Circadian output from the SCN can also be altered by disrupting core gene components of the molecular pacemaker, including *Clock* [4], *Bmal1* [7], *mPer1* and *mPer2* [8, 9], and *mCry1* and *mCry2* [10, 11]. Therefore, analysis of mice with “clock gene” mutations provides a way to dissect genetically the role of circadian rhythms in reproduction. Here, we characterized reproductive function in the female *Clock* mutant mouse, which carries a 51 amino acid deletion in the transcriptional-activation domain of the CLOCK protein [5, 12]. Wild-type, *Clock/+*, and *Clock/Clock* females were examined daily for the onset of vaginal opening, and upon reaching 10 weeks of age, they were monitored by vaginal lavage to record estrous cyclicity (see Supplemental Data for a detailed description of all Experimental Procedures). Although the timing of vaginal opening was normal (Table S1), there were significant differences in the characteristics of estrous cycles among genotypes (Figure 1A); whereas wild-type females and the majority of *Clock/+* females exhibited multiple consecutive 4–5 day-long cycles, *Clock/Clock* females had prolonged and irregular cycles characterized by significantly fewer days of proestrus and more days of estrus (Figures 1B and S1).

Abnormal estrous cycles could be due to either central or peripheral defects. Therefore, we began by examining ovarian function in intact cycling females. Wild-type, *Clock/+*, and *Clock/Clock* females were sacrificed on the afternoon of diestrus or proestrus, serum levels of estradiol and progesterone were measured by radioimmunoassay (RIA), and ovaries were fixed for histological analysis. In all three genotypes, estradiol (Figure S2A) and progesterone (Figure S2B) levels were low for diestrus and significantly elevated on the afternoon of proestrus, and histological analysis of ovarian tissue from proestrus *Clock/Clock* mice showed no gross morphological abnormalities (Figure S3, Table S2). Follicles at all stages of development were present, and both corpora lutea and Graafian follicles were present in *Clock* mutant ovaries in numbers comparable to those observed in wild-type tissue. These results suggest that the estrous cycle irregularities we observed in *Clock/Clock* females do not stem from an effect of the *Clock* mutation acting at the level of the ovary.

We then investigated the timing and amplitude of the luteinizing hormone (LH) surge by collecting serial blood samples from ovary-intact wild-type and *Clock/Clock* mice on the morning, afternoon, and evening of proestrus as defined by a nucleated vaginal smear. The proportions of wild-type (50%) and *Clock/Clock* (0%) mice

*Correspondence: j-takahashi@northwestern.edu

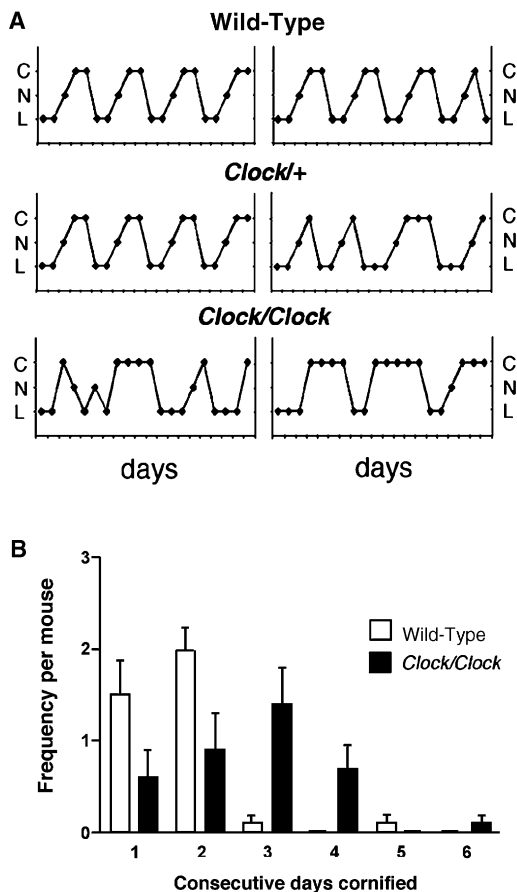


Figure 1. *Clock/Clock* Females Display Lengthened and Irregular Estrous Cycles

(A) Representative estrous cycles as measured by vaginal cytology in wild-type (top), *Clock/+* (middle), and *Clock/Clock* (bottom) females. C = cornified, N = nucleated, L = leukocytic.

(B) *Clock/Clock* females have significantly more days of cornified smears compared to wild-type females, as determined by an unpaired t test for the number of consecutive days cornified.

exhibiting an LH peak of 10 ng/ml or greater differed significantly (chi square, $p < 0.01$, Figure 2A). None of the *Clock/Clock* females exhibited LH concentrations exceeding 1.8 ng/ml, and concentrations in the majority of mutants never varied from baseline (0.2 ng/ml). Maximum serum LH concentrations were significantly lower in *Clock/Clock* than in wild-type females (Student's t test, $p < 0.05$, Figure 2B).

It is possible that *Clock* mutants have a normal LH surge that is not temporally associated with a nucleated smear, but the hormonal profiles we observed indicate that this possibility is unlikely. A minimum of 30 hr of exposure to estradiol is required for inducing GnRH release [13]; both vaginal cytology and serum sampling show that *Clock* mutants have low estradiol levels prior to the day of a nucleated smear. Therefore, a large LH surge is probably not occurring before the initial sampling time (ZT5). It is also unlikely that a surge is occurring long after the morning of a nucleated smear, because extended progesterone elevation results in inhibition of LH release [14], and *Clock* mutants exhibit

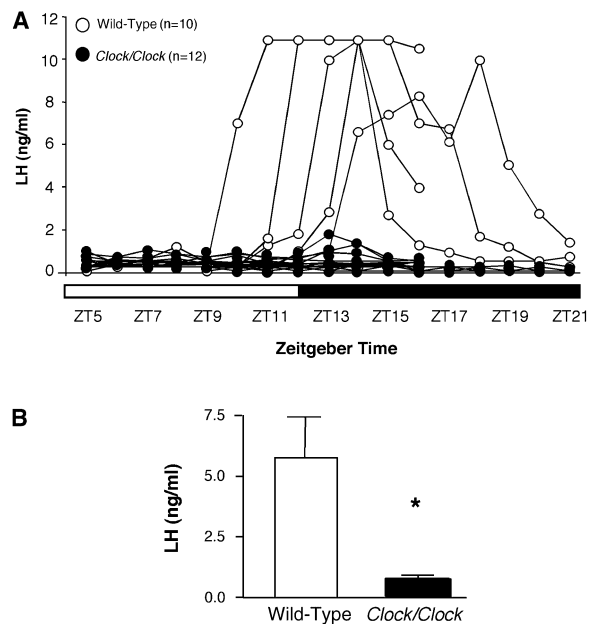


Figure 2. *Clock/Clock* Mutants Fail to Have a Coordinated LH Surge on the Day of Proestrus

(A) Individual LH traces from all wild-type (open circles) and *Clock/Clock* (closed circles) females sampled from either ZT5-16 or ZT9-21.

(B) Individual peak LH values in serum obtained from serially sampled mice. Peak *Clock/Clock* LH values were compared using an unpaired t test and found to be significantly lower than peak wild-type values (** $p < 0.01$). Due to the limited sample volume collected, samples with LH values exceeding the range of the RIA could not be reassayed at lower sample concentrations, preventing absolute measurements of serum concentrations above 10 ng/ml. Therefore, for analysis and presentation purposes, these values were set at 10 ng/ml, and we defined the minimum "surge" value as 14% of the maximum, or 1.4 ng/ml, as previous studies in rats have shown that 14 percent of the peak surge is the minimum value of LH required for ovulation [39]. Using these criteria, only one *Clock* mutant reached the minimum surge level at any time point.

elevated progesterone coincident with a nucleated smear. Thus, the most parsimonious explanation is that the small elevations of LH observed throughout the day of proestrus are sufficient to induce ovulation in individual *Clock* mutants.

Clock Mutant Females Exhibit Elevated Rates of Fetal Reabsorption and Pregnancy Failure

To our knowledge, circadian rhythm defects have never been implicated in the abnormal progression of pregnancy, although some researchers have suggested a connection [15], and anecdotal evidence from our laboratory suggests that *Clock* mutant females regularly failed to produce offspring. For verification and characterization of this defect, wild-type and *Clock/Clock* females were mated with wild-type males, and the initiation and progression of pregnancy was measured by observation of fetal reabsorption at 11, 14, and 17 days post copulation (dpc), and at full-term. Although the early stages of pregnancy were normal in *Clock* mutants, *Clock/Clock* females showed an increased rate of mid-gestation fetal reabsorption, and pregnancy failure at

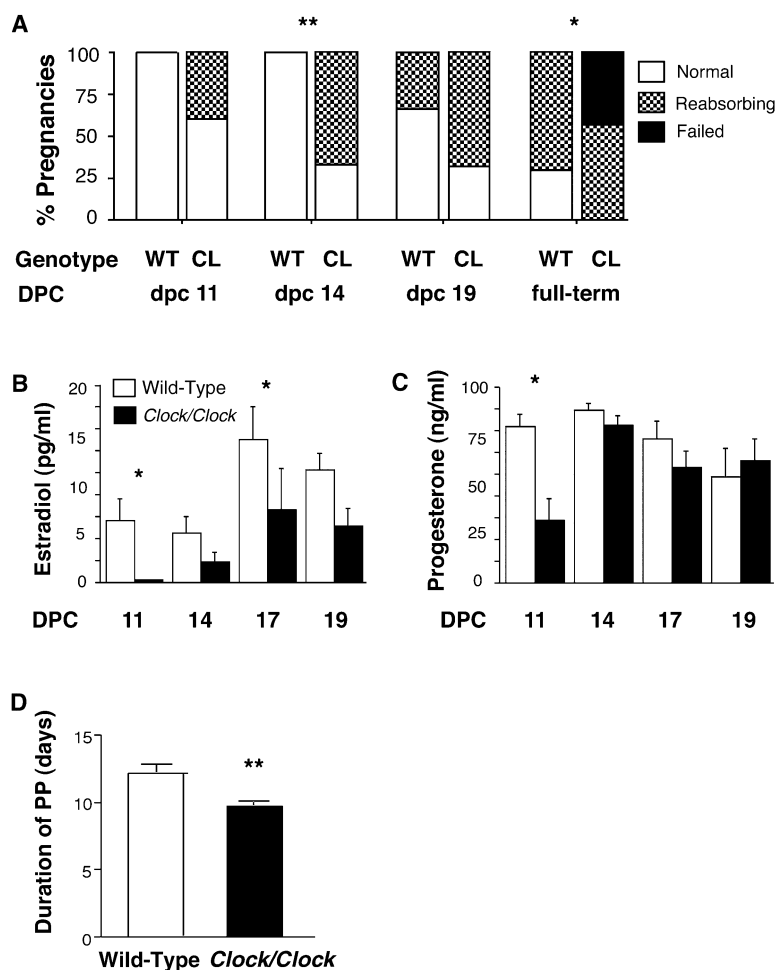


Figure 3. *Clock/Clock* Pregnancies Are Characterized by Increased Rates of Fetal Reabsorption, Reduced Serum Levels of Estradiol and Progesterone, and Possible Dysregulation of Prolactin Release

(A) The percentage of *Clock* mutant dam pregnancies showing any signs of fetal reabsorption is significantly greater than wild-type at dpc 14 (** $p < 0.01$) and full-term (* $p \leq 0.05$), as determined by chi square analysis. In many cases, both wild-type and *Clock* pregnancies showed some signs of reabsorption by full-term, but reabsorbing fetuses occurred more frequently in *Clock* pregnancies. While all wild-type pregnancies delivered normally, 40% of *Clock* mutant dams carried to full-term but failed to deliver. Importantly, *Clock* mutant females exhibited copulatory plugs as frequently as wild-type females, indicating that mating behavior was intact in the mutants. Additionally, the average number of fetuses per genotype at dpc 11 was almost identical between genotypes (7.6 fetuses per wild-type dam, 7.75 fetuses per *Clock/Clock* dam), indicating that the early stages of pregnancy, including ovulation, fertilization, implantation, and early fetal development, occur normally in *Clock* mutants. WT = wild-type, CL = *Clock/Clock*, DPC = days post copulation.

(B) Estradiol levels are significantly reduced in *Clock/Clock* mice compared to wild-types ($p < 0.001$). Estradiol in both genotypes is elevated in late pregnancy (dpc 17, 19) compared to mid-pregnancy (dpc 11, 14) ($p < 0.01$).

(C) Progesterone is significantly lower in *Clock/Clock* females compared to wild-types ($p < 0.05$) due to very low progesterone levels in *Clock/Clock* females at dpc 11. In both genotypes, progesterone levels at dpc 14 are elevated compared to dpc 11 and dpc 19

values ($p \leq 0.01$). For both (B) and (C), * $p < 0.05$. Prior to testing the hypotheses, data were checked to verify that they met the assumptions of normality and equality of variance required for analysis of variance. Because the data violated one or more of the assumptions, data were transformed using logarithms and statistical analysis (two-way ANOVA and Fisher's LSD) was performed on the log transformed data.

(D) The duration of pseudopregnancy is significantly shorter in *Clock/Clock* females compared to wild-type controls as determined by t test (** $p \leq 0.001$). Wild-type ($n = 10$) and *Clock/Clock* ($n = 26$) females were mated with vasectomized CD-1 males and the length of pseudopregnancy, as indicated by leukocytic vaginal cytology, was measured by vaginal lavage. The presence of a copulatory plug was designated as dpc 1, and the final day of leukocytic smears was designated as the last day of pseudopregnancy.

full-term, compared to wild-type females. The percentage of *Clock/Clock* pregnancies showing signs of fetal reabsorption was significantly different from wild-type pregnancies at two time points, dpc 14 and full-term, and the percentage of fetuses being reabsorbed by *Clock* mutant dams was elevated at dpc 14 and full-term (chi square, $p < 0.05$; Figures 3A and S4). All of the wild-type females and 57% (4 of 7) of the *Clock/Clock* females allowed to carry their pregnancies to full term delivered live litters on dpc 20. However, 43% (3 of 7) of the *Clock/Clock* females either went into an extended but non-productive labor or failed to enter labor and instead fully reabsorbed the full-term fetuses.

Estradiol and progesterone are vital for maintaining uterine receptivity to developing fetuses during pregnancy and promoting parturition [16]. Therefore, we measured estradiol (Figure 3B) and progesterone (Figure 3C) levels at dpc 11, 14, 17, and 19 (the day before

expected parturition) to determine whether altered ovarian hormone levels could explain the pregnancy phenotype in *Clock/Clock* females. Throughout pregnancy, *Clock* mutants exhibited significantly lower estradiol levels compared to wild-types, although both genotypes displayed an increase in estradiol levels from mid-pregnancy (dpc 11, dpc 14) to late pregnancy (dpc 17, dpc 19). At full-term, estradiol in *Clock* mutants was only one-third the level of estradiol in wild-type females. Because estradiol is important for enhancing uterine contractility [17], it is likely that low estradiol levels were at least partially responsible for the failure of some *Clock/Clock* females to initiate labor.

Progesterone levels were also reduced in *Clock* mutants, most notably at dpc 11. Mid-gestational levels of progesterone are particularly important for maintaining blood flow to developing fetuses [18], and previous studies have shown a quantitative relationship between pro-

gestosterone levels and maintenance of pregnancy [19]. Therefore, the reduced progesterone levels observed in *Clock* mutant dams at dpc 11 may explain the fetal reabsorption observed by dpc 14. By dpc 14, progesterone levels in *Clock* mutants had risen, and the rate of fetal reabsorption in *Clock/Clock* females remained similar until full-term to that in wild-types.

The Duration of Pseudopregnancy Is Shortened in *Clock* Mutant Mice

We hypothesized that the observed abnormalities in fetal reabsorption and ovarian hormones in pregnant *Clock/Clock* females could be explained by abnormal prolactin release during early and/or mid-pregnancy. In both rats and mice, prolactin release is induced by the copulatory stimulus, and thereafter occurs in two daily surges that are roughly coordinated to lights-on and lights-off, suggesting that release is under circadian control and is therefore vulnerable to gene mutations that alter the circadian pacemaker [20–22]. Prolactin is the only factor required to rescue and maintain the ovarian corpora lutea (CL), which produce the elevated serum progesterone levels characteristic of and necessary for pregnancy and pseudopregnancy [23]. At mid-gestation, around dpc 10, maternal prolactin release tapers off and the CL are supported for the remainder of pregnancy by the prolactin-like placental lactogen (PL-1) produced by fetuses [24]. If the mating was infertile, there is no PL-1 to rescue the CLs, and the female resumes estrous cyclicity. Thus, the termination of pseudopregnancy is a reliable indicator of the termination of mating-induced PRL release. Therefore, in order to determine whether a defect in maternal prolactin release might be responsible for low *Clock/Clock* progesterone levels at dpc 11 and subsequent elevated fetal reabsorption at dpc 14, we measured the duration of pseudopregnancy. This functional assessment was used rather than measurement of PRL levels because the expected variability in the timing of PRL secretion and prevalence of PRL release in response to stress would have made interpretation of the results difficult.

Pseudopregnancy was significantly shortened in *Clock/Clock* females compared to wild-type controls (Figure 3D), suggesting that PRL secretion may cease earlier in *Clock/Clock* mice than in wild-type females. The functional consequences of premature cessation of prolactin release are partial or complete CL regression, a drop in progesterone levels, and either abortion or an increase in fetal reabsorption [18]. These consequences are consistent with our observations. Although it is possible that there is a failure on the part of the *Clock/+* and *Clock/Clock* fetuses to begin to support the CLs at the appropriate developmental stage, the cessation of pseudopregnancy in mutant females often occurs several days prior to the onset of fetal PL-1 production. We have also observed that, when a litter of *Clock/Clock* embryos is transplanted into a pseudopregnant wild-type female, the pregnancy progresses normally, indicating that *Clock* mutant fetuses are capable of sustaining pregnancy in normal females (BHM, personal observation). Therefore, at least some of the *Clock* mutant pregnancy abnormalities are likely due to abnormal

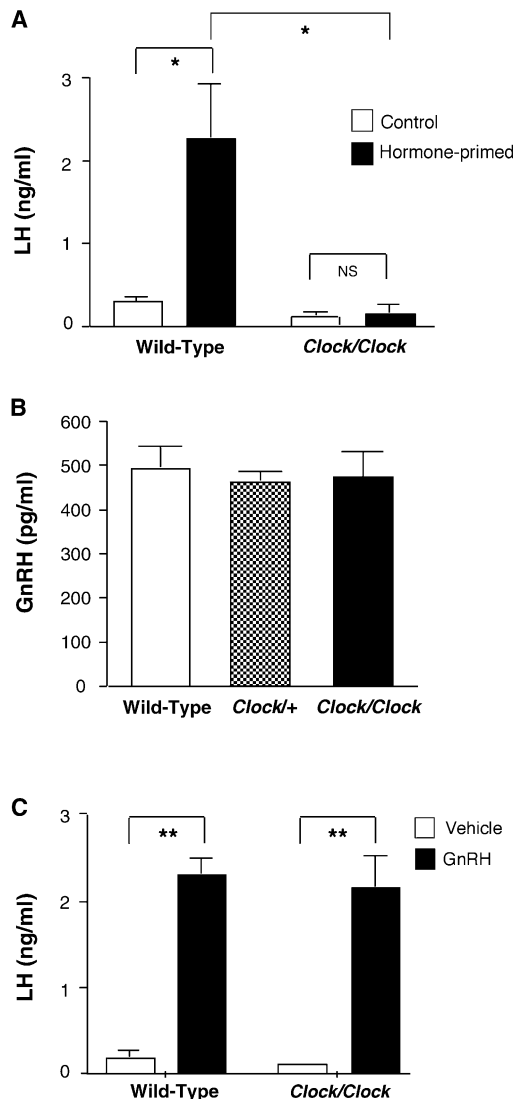


Figure 4. Hypothalamic-Pituitary Gonadal Axis Function in *Clock* Mutants

(A) Estradiol benzoate treatment resulted in a significant elevation in serum LH in wild-type ovariectomized females (* $p < 0.01$), but did not produce elevated LH levels in *Clock/Clock* females. NS = not significant. Two-way ANOVA. Treatment ($F = 8.9$, $df (1,24)$ $p < 0.01$); Genotype ($F = 11.8$, $df (1,24)$, $p < 0.01$). Mice were ovariectomized and implanted with estradiol capsules. Six days following implantation, mice received an injection of either estradiol benzoate or sesame oil at 0800h, and samples were collected the following evening at ZT13.

(B) There is no difference in hypothalamic GnRH peptide content among wild-type, *Clock/+*, and *Clock/Clock* females on the afternoon of proestrus. One-way ANOVA ($F = 0.1$, $df (2,11)$, $p = 0.9$).

(C) GnRH treatment (400 ng/kg, sc) resulted in significantly increased serum LH compared to control levels in both wild-type and *Clock/Clock* females that had been hormone primed as described above (** $p < 0.001$).

circadian control of maternal prolactin release. Our hypothesis is consistent with the findings of other groups that have described a link between the circadian system and prolactin secretion [25].

Abnormal LH Release in *Clock* Mutants Is Due to a Hypothalamic Defect

The role of circadian rhythms in LH release and in the estrous cycle has been more thoroughly described than it has been for pregnancy. Therefore, we focused on the estrous cycle to determine how the *Clock* mutation could result in altered reproductive physiology. We hypothesized that *Clock/Clock* females display irregular estrous cycles and fail to have a coordinated LH surge due to a disruption of the daily timing signal from the SCN to the GnRH neurons, rather than as a result of pituitary defects or inappropriate feedback from ovarian hormones.

In rodents, coordinated GnRH release on the afternoon of proestrus requires both a daily timing signal originating in the SCN and permissive levels of estrogen and progesterone [26]. To determine whether the observed defect in LH release in *Clock* mutants was due to inappropriate steroid feedback, ovariectomized mutant and wild-type mice received low-dose estradiol capsules followed by an injection of estradiol benzoate; a paradigm that induces an LH surge during the circadian-timed window. Estradiol capsules alone resulted in baseline levels of LH in both genotypes due to negative feedback on GnRH neurons (Figure 4A). Injection with estradiol benzoate resulted in positive feedback and a subsequent LH surge in wild-type mice, but failed to induce an LH surge in *Clock/Clock* mice. These data are consistent with the hypothesis that *Clock* mutants lack a coordinated daily timing signal triggering GnRH release, and rules out the possibility that improper levels of ovarian hormones prevent ovary-intact *Clock* mutants from mounting a coordinated LH surge.

Several studies show that GnRH transcription is under circadian control, and the *Clock* mutation results in the down-regulation of transcription of a number of genes in addition to the core clock genes, introducing the possibility that the *Clock* mutation alters GnRH synthesis [27, 28]. To test this hypothesis, we measured GnRH by RIA in hypothalami from wild-type, *Clock/+*, and *Clock/Clock* mice on the afternoon of proestrus. Hypothalamic GnRH content was normal in *Clock/+* and *Clock/Clock* females (Figure 4B); thus, the mutation does not interfere with the production of GnRH, and *Clock* mutants have an adequate supply of the peptide available for release on proestrus.

Finally, we evaluated pituitary function in vivo by measuring serum LH following treatment with GnRH in wild-type and mutant females. Expression of the core clock gene *Period2* is rhythmic in pituitaries in culture, raising the possibility that the *Clock* mutation might interfere directly with either LH production or release at the level of the pituitary [29]. However, pituitary release of LH following a GnRH challenge was identical in wild-type and *Clock/Clock* females (Figure 4C), indicating that pituitary responsiveness to GnRH is normal in *Clock* mutants.

Since all aspects of the hypothalamic-pituitary-gonadal (HPG) axis other than GnRH/LH release are normal in *Clock* mutants, estrous cycle defects in the *Clock* mutants appear to result from a disruption of the timing and/or coordination of GnRH release on proestrus. This disruption may be due to either altered output

from the SCN to GnRH neurons or an effect of the *Clock* mutation on GnRH neurons themselves. Several groups have shown that core clock genes, including *Bmal1* and *Per1*, are expressed and cycle in an immortalized GnRH cell line (GT1-7), suggesting that intracellular pacemakers within GnRH neurons may be important for the circadian regulation of GnRH release [30, 31]. It is possible that expression of the *Clock* mutation within GnRH neurons may disrupt the expression of other proteins necessary for GnRH release, such as hormone or ion channel receptors. In support of this, Chappell and colleagues found that transfection of GT1-7 cells with the CLOCK- Δ 19 dominant negative mutation decreases mean GnRH pulse frequency, whereas overexpression of *Cry1*, which inhibits CLOCK-induced transcription, increases GnRH pulse frequency [30]. However, the ultimate relationship between GnRH neuron pulsatility and the regulation of proestrus GnRH release remains undefined. It is therefore unclear whether the extent of the disruption of GnRH pulsatility induced by the *Clock* mutation in GT1-7 cells would be sufficient to cause the defects in LH release and estrous cyclicity that we observed in *Clock/Clock* mice.

Although a direct effect of the *Clock* mutation on GnRH neurons in vivo remains to be examined, we hypothesize that *Clock* mutants fail to have an LH surge because the SCN in mutant animals does not provide a coordinated time-of-day signal to GnRH neurons. Likely candidates for conveying the SCN-GnRH signal are the neuropeptides vasopressin and vasoactive intestinal peptide (VIP), both of which are expressed in neurons projecting from the SCN to the preoptic area [32]. VIP-containing fibers make direct projections to the population of GnRH neurons that expresses *c-fos* on the afternoon of proestrus [33, 34], and suppression of VIP synthesis in the SCN attenuates peak LH levels during estradiol-induced surges [35]. Vasopressin-containing neurons project to interneurons adjacent to GnRH fibers [36], and, in rats, inhibition of vasopressin signaling on the morning of a hormone-induced surge significantly attenuates LH release [37]. Furthermore, vasopressin expression is rhythmically controlled by CLOCK-induced transcription, and vasopressin content in the SCN is drastically reduced in *Clock* mutants, suggesting a molecular mechanism for the missing time-of-day signal in *Clock/Clock* females [38]. Further studies are necessary to refine our understanding of the roles these peptides play in controlling GnRH release. More generally, characterization of possible reproductive defects in other circadian mutants will help to clarify the role of the *Clock* gene specifically, as opposed to the role of the molecular pacemaker, in the circadian control of reproductive function.

Supplemental Data

Supplemental Data including Experimental Procedures, four additional figures, and two tables are available at <http://www.current-biology.com/cgi/content/full/14/15/1367/DC1/>.

Acknowledgments

We thank Brigitte Mann for performing the RIAs and Drs. Neena B. Schwartz and Teresa K. Woodruff for helpful advice. This work was supported by a predoctoral fellowship from the National Science

Foundation (B.H.M.), and NIH awards AG-18200 and HL-59598 (F.W.T). Joseph S. Takahashi is an Investigator in the Howard Hughes Medical Institute.

Received: April 14, 2004

Revised: May 28, 2004

Accepted: June 9, 2004

Published: August 10, 2004

References

- Hoffman, J.C. (1988). The influence of photoperiod on reproductive functions in female mammals. In *Physiology of Reproduction*, Knobil E., Neill J., eds. (New York: Raven Press), pp. 57–77.
- Brown-Grant, K., and Raisman, G. (1977). Abnormalities in reproductive function associated with the destruction of the suprachiasmatic nuclei in female rats. *Proc. R. Soc. Lond. B. Biol. Sci.* **198**, 279–296.
- Weigand, S.J., Terasawa, E., Bridson, W.E., and Goy, R.W. (1980). Effects of discrete lesions of preoptic and suprachiasmatic structures in the female rat. Alterations in the feedback regulation of gonadotropin secretion. *Neuroendocrinology* **31**, 147–157.
- Vitaterna, M.H., King, D.P., Chang, A., Kornhauser, J.M., Lowrey, P.L., McDonald, J.D., Dove, W.F., Pinto, L.H., Turek, F.W., and Takahashi, J.S. (1994). Mutagenesis and mapping of a mouse gene, *Clock*, essential for circadian behavior. *Science* **264**, 719–725.
- King, D.P., Zhao, Y., Sangoram, A.M., Wilsbacher, L.D., Tanaka, M., Antoch, M.P., Steeves, T.D.L., Vitaterna, M.H., Kornhauser, J.M., Lowrey, P.L., et al. (1997). Positional cloning of the mouse circadian *Clock* gene. *Cell* **89**, 641–653.
- Takahashi, J.S., Turek, F.W., and Moore, R.Y. eds. (2001). *Handbook of Behavioral Neurobiology: Circadian Clocks*, (Kluwer Academic/Plenum Publishers).
- Bunger, M.K., Wilsbacher, L.D., Moran, S.M., Clendenen, C., Radcliffe, L.A., Hogenesch, J.B., Simon, M.C., Takahashi, J.S., and Bradfield, C.A. (2000). *Mop3* is an essential component of the master circadian pacemaker in mammals. *Cell* **103**, 1009–1017.
- Zheng, B., Albrecht, U., Kaasik, K., Sage, M., Lu, W., Vaishnav, S., Li, Q., Sheng Sun, Z., Eichele, G., Bradley, A., et al. (2001). Nonredundant roles of the *mPer1* and *mPer2* genes in the mammalian circadian clock. *Cell* **105**, 683–694.
- Bae, K., Xiaowei, J., Maywood, E.S., Hastings, M.H., Reppert, S.M., and Weaver, D.R. (2001). Differential functions of *mPer1*, *mPer2*, and *mPer3* in the SCN circadian clock. *Neuron* **30**, 525–536.
- Vitaterna, M.H., Selby, C.P., Todo, T., Niwa, H., Thompson, C., Fruechte, E.M., Hitomi, K., Thresher, R.J., Ishikawa, T., Miyazaki, J., et al. (1999). Differential regulation of mammalian period genes and circadian rhythmicity by *Cryptochromes 1* and *2*. *Proc. Natl. Acad. Sci. USA* **96**, 12114–12119.
- van der Horst, G.T., Muijtjens, M., Kobayashi, K., Takano, R., Kanno, S., Takao, M., de Wit, J., Verkerk, A., Eker, A.P., van Leenan, D., et al. (1999). Mammalian *Cry1* and *Cry2* are essential for maintenance of circadian rhythms. *Nature* **398**, 627–630.
- Gekakis, N., Staknis, D., Nguyen, H.B., Davis, F.C., Wilsbacher, L.D., King, D.P., Takahashi, J.S., and Weitz, C.J. (1998). Role of the *CLOCK* protein in the mammalian circadian mechanism. *Science* **280**, 1564–1569.
- Bronson, F.H., and Vom Saal, F.S. (1979). Control of the preovulatory release of luteinizing hormone by steroids in the mouse. *Endocrinology* **104**, 1247–1255.
- Girmus, R.L., and Wise, M.E. (1992). Progesterone directly inhibits pituitary luteinizing hormone secretion in an estradiol-dependent manner. *Biol. Reprod.* **46**, 710–714.
- Endo, A., and Watanabe, T. (1989). Effects of non-24-hour days on reproductive efficacy and embryonic development in mice. *Gamete Res.* **22**, 435–441.
- Hodgen, G.D., and Itskovitz, J. (1988). Recognition and maintenance of pregnancy. In *The Physiology of Reproduction*, Knobil E., Neill J., eds. (New York, Raven Press, Ltd.).
- Thorburn, G.D., and Challis, J.R.G. (1979). Endocrine control of parturition. *Physiol. Rev.* **59**, 863–918.
- Milligan, S.R., and Finn, C.A. (1997). Minimal progesterone support required for the maintenance of pregnancy in mice. *Hum. Reprod.* **12**, 602–607.
- Csapo, I., and Wiest, W.G. (1969). An examination of the quantitative relationship between progesterone and the maintenance of pregnancy. *Endocrinology* **85**, 735–746.
- Freeman, M.E., Smith, M.S., Nazian, S.J., and Neill, J.D. (1973). Ovarian and hypothalamic control of the daily surges of prolactin secretion during pseudopregnancy in the rat. *Endocrinology* **94**, 875–882.
- Barkley, M.S., Bradford, G.E., and Geschwind, I.I. (1978). The pattern of plasma prolactin concentration during the first half of mouse gestation. *Biol. Reprod.* **19**, 291–296.
- Stirland, J.A., Hastings, M.H., Loudon, A.S., and Maywood, E.S. (1996). The tau mutation in the Syrian hamster alters the photoperiodic responsiveness of the gonadal axis to melatonin signal frequency. *Endocrinology* **137**, 2183–2186.
- Smith, M.S. (1980). Role of prolactin in mammalian reproduction. In *Reproductive Physiology III*, Greep, R.O., ed. (Baltimore: University Park Press), pp. 249–276.
- Linzer, D.I., and Fisher, S.J. (1999). The placenta and the prolactin family of hormones: regulation of the physiology of pregnancy. *Mol. Endocrinol.* **13**, 837–840.
- Palm, I.F., Van der Beek, E.M., Swarts, H.J.M., Van der Vliet, J., Wiegant, V.M., Buijs, R.M., and Kalsbeek, A. (2001). Control of the estradiol-induced prolactin surge by the suprachiasmatic nucleus. *Endocrinology* **142**, 2296–2302.
- Levine, J.E. (1997). New concepts of the neuroendocrine regulation of gonadotropin surges in rats. *Biol. Reprod.* **56**, 239–302.
- Gore, A.C. (1998). Diurnal rhythmicity of gonadotropin-releasing hormone gene expression in the rat. *Neuroendocrinology* **197**, 257–263.
- Panda, S., Antoch, M.P., Miller, B.H., Su, A.I., Schook, A.B., Straume, M., Schultz, P.G., Kay, S.A., Takahashi, J.S., and Hogenesch, J.B. (2002). Coordinated transcription of key pathways in the mouse by the circadian clock. *Cell* **109**, 307–320.
- Yoo, S.-H., Yamazaki, S., Lowrey, P.L., Shimomura, K., Ko, C.H., Buhr, E.D., Siepka, S.M., Hong, H.-K., Oh, W.J., Yoo, O.J., et al. (2004). *PERIOD2::LUCIFERASE* real-time reporting of circadian dynamics reveals persistent circadian oscillations in mouse peripheral tissues. *Proc. Natl. Acad. Sci. USA* **101**, 5339–5346.
- Chappell, P.E., White, R.S., and Mellon, P.L. (2003). Circadian gene expression regulates pulsatile gonadotropin-releasing hormone (GnRH) secretory patterns in the hypothalamic GnRH-secreting GT1-7 cell line. *J. Neurosci.* **23**, 11202–11213.
- Gillespie, J.M.A., Chan, B.P.K., Roy, D., Cai, F., and Belsham, D.D. (2003). Expression of circadian rhythm genes in gonadotropin-releasing hormone-secreting GT1-7 neurons. *Endocrinology* **144**, 5285–5292.
- Kalsbeek, A., and Buijs, R.M. (2002). Output pathways of the mammalian suprachiasmatic nucleus: coding circadian time by transmitter selection and specific targeting. *Cell Tissue Res.* **309**, 109–118.
- Smith, M.J., Jennes, L., and Wise, P.M. (2000). Localization of the VIP2 receptor protein on GnRH neurons in the female rat. *Endocrinology* **141**, 4317–4320.
- Krajnak, K., Rosewell, K.L., and Wise, P.M. (2001). Fos-induction in gonadotropin-releasing hormone neurons receiving vasoactive intestinal polypeptide innervation is reduced in middle-aged female rats. *Biol. Reprod.* **64**, 1160–1164.
- Harney, J.P., Scarbrough, K., Rosewell, K.L., and Wise, P.M. (1996). In vivo antisense antagonism of vasoactive intestinal peptide in the suprachiasmatic nuclei causes aging-like changes in the estradiol-induced luteinizing hormone and prolactin surges. *Endocrinology* **137**, 3696–3701.
- Watson, R.E., Jr., Langub, M.C., Jr., Engle, M.G., and Maley, B.E. (1995). Estrogen-receptive neurons in the anteroventral periventricular nucleus are synaptic targets of the suprachiasmatic nucleus and peri-suprachiasmatic region. *Brain Res.* **689**, 254–264.

37. Palm, I.F., Van der Beek, E.M., Wiegant, V.M., Buijs, R.M., and Kalsbeek, A. (1999). Vasopressin induces a luteinizing hormone surge in ovariectomized, estradiol-treated rats with lesions of the suprachiasmatic nucleus. *Neuroscience* 93, 659–666.
38. Jin, X., Shearman, L.P., Weaver, D.R., Zylka, M.J., De Vries, G.J., and Reppert, S.M. (1999). A molecular mechanism regulating rhythmic output from the suprachiasmatic circadian clock. *Cell* 96, 57–68.
39. Greig, F., and Weisz, J. (1973). Preovulatory levels of luteinizing hormone, the critical period and ovulation in rats. *J. Endocrinol.* 57, 235–245.