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The Mysterious Food-Entrainable Oscillator: Insights from Mutant and Engineered Mouse Models

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Abstract

The food-entrainable oscillator (FEO) is a mysterious circadian clock because its anatomical location(s) and molecular timekeeping mechanism are unknown. Food anticipatory activity (FAA), which is defined as the output of the FEO, emerges during restricted feeding. FAA disappears immediately during *ad libitum* feeding and re-appears during subsequent fasting. A free-running FAA rhythm has been observed only in rare circumstances when food was provided with a period outside the range of entrainment. Therefore, it is difficult to study the circadian properties of the FEO. Numerous studies have attempted to identify the critical molecular components of the FEO using mutant and genetically engineered mouse models. Herein we critically review the experimental protocols and findings of these studies in mouse models. Several themes emerge from these studies. First, there is little consistency in restricted feeding protocols between studies. Moreover, the protocols were sometimes not optimal, resulting in erroneous conclusions that FAA was absent in some mouse models. Second, circadian genes are not necessary for FEO timekeeping. Thus, another non-canonical timekeeping mechanism must exist in the FEO. Third, studies of mouse models have shown that signaling pathways involved in circadian timekeeping, reward (dopaminergic), and feeding and energy homeostasis can modulate, but are not necessary for, the expression of FAA. In sum, the approaches to date have been largely unsuccessful in discovering the timekeeping mechanism of the FEO. Moving forward, we propose the use of standardized and optimized experimental protocols that focus on identifying genes that alter the period of FAA in mutant and engineered mouse models. This approach is likely to permit discovery of molecular components of the FEO timekeeping mechanism.

Keywords

circadian; food anticipatory activity; FEO; restricted feeding; food entrainment

Introduction

In 1922, Curt Richter described that the “spontaneous activity of the rat is very intimately related to the food habits of the animal.” When Richter fed rats every day at noon (for 25 min), he found that “activity increased rapidly... right up to the time of the next feeding

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period (Richter, 1922).” Thus was born the study of food anticipatory activity (FAA). Beginning in the 1970’s, Frederick Stephan intensively studied this FAA and showed that it was controlled by an autonomous circadian oscillator, the food-entrainable oscillator (FEO). It was so-named because FAA persisted in constant conditions (i.e. during fasting) and entrained to a circadian range of feeding intervals [for review see (Boulos and Terman, 1980; Mistlberger, 1994)].

After Stephan established that anticipation of daily restricted meals (called FAA) was an output of the circadian FEO, Mistlberger and others spent the next 2 decades searching for the anatomical locus of the FEO. The FEO is not in the suprachiasmatic nucleus (SCN), nor in dozens of other brain regions that Mistlberger and others, and later, Davidson and Stephan, explored [for review see (Davidson, 2009)]. Thus, the elusive anatomical location(s) of the FEO remains one of the unsolved mysteries in the field of circadian rhythms.

In 2003, another mystery surrounding the FEO began to emerge, namely that canonical circadian clock genes may not be required by the FEO for circadian timekeeping. In the SCN, the canonical circadian timekeeping mechanism is well-characterized. Heterodimers of Brain and Muscle ARNT-Like 1 (BMAL1) and Circadian Locomotor Output Cycles Kaput (CLOCK) activate the transcription of *Period* (*Per1, 2*) and *Cryptochrome* (*Cry 1, 2*) genes. As CRY and PER proteins accumulate, they form complexes and inhibit the transcription factor activity of BMAL1/CLOCK, thereby inhibiting their own transcription. However, Rae Silver and colleagues were the first to report that mice lacking the functional canonical circadian timekeeping mechanism had normal FAA, when they showed that Clock^{-/-} mutant mice had robust FAA that persisted during fasting in the light-dark cycle (Pitts et al., 2003). Subsequently, studies performed in *Cry1^{-/-}/Cry2^{-/-}* double mutant and *Per1^{-/-}/Per2^{-/-}/Per3^{-/-}* triple mutant mice, definitively showed that canonical timekeeping genes were not required for FAA, because FAA persisted during restricted feeding and subsequent fasting in constant darkness (when the SCN was disabled) (Iijima et al., 2005; Pendergast et al., 2012). Several other studies showed that the FEO was intact in most circadian gene mutant mice, but compromised in a few mutants. Herein we will review the studies of mutant and engineered mouse models (circadian gene mutants as well as other molecular pathways) that have been examined for their effects on FAA [for earlier reviews of FAA neurogenetics and methodological issues, see (Challet et al., 2009; Mistlberger, 2009)].

Measuring FAA that is controlled by an autonomous circadian clock

The system for food entrainment is organized similarly to the light entrainment pathway (Fig. 1A, Fig. 2). Each system is comprised of inputs, a pacemaker, and outputs. Light is the most salient input to the light-entrainable oscillator in the SCN. The SCN pacemaker then controls outputs, such as the nocturnal activity rhythm. Changes in the nocturnal activity rhythm could result from alterations in the input, pacemaker, or output pathways. Since we know the anatomical locus of the light-entrainable oscillator, experiments can be designed to determine which component of the pathway contributes to a phenotype. Food is the most salient (known) input to the FEO, which then controls FAA. Studying the autonomous FEO

is challenging. Since its anatomical location(s) is unknown, the FEO cannot be cultured in a dish, or lesioned, or manipulated with targeted injections of siRNA. Additionally, omics methods cannot be used to measure gene, protein, or metabolic oscillations in the FEO. Thus, when the magnitude of FAA is altered, it is difficult, or impossible, to conclude whether the input pathway, pacemaker, or output pathway is responsible. On the other hand, when the period of FAA is altered in a mutant mouse, it is more likely that the FEO pacemaker has been altered. However, measuring the period of FAA is challenging and has not been widely studied.

Interpreting restricted feeding studies of mutant mice is also difficult because the SCN and FEO are coupled (Fig. 1A) (Rosenwasser et al., 1984; Stephan, 1986; Ottenweller et al., 1990; Pendergast and Yamazaki, 2014). If the SCN is functional in a mutant mouse, then the SCN could be used as phase reference and FAA would appear normal. In addition, the SCN could moderate the period of the mutant FEO, resulting in apparently normal FAA. It is only when the SCN is disabled, ideally by lesioning the SCN, but also in circadian mutants that are arrhythmic in constant darkness (e.g. *Bmal1*^{-/-}; *Per1*^{-/-}/*Per1*^{-/-}), that we can study the output of the FEO alone (and not in a complex with the SCN). Therefore, to study the genes that are required for, or participate in, FEO timekeeping, the SCN must be disabled (Fig. 2B). Most of the studies of the FEO/FAA to date have not eliminated the SCN rhythm. Thus, our discussion will focus on how mutation of a specific gene affects FAA, noting that this could be a defect in input, timekeeping, or output of the FEO. We will also highlight the studies that have disabled the SCN, since these provide information about the genes that may regulate the timekeeping mechanism of the FEO.

Some studies combine temporal restricted feeding with caloric restriction. For example, a mouse will be given ~60% of its typically daily food intake at mid-day. The mice typically eat all of the food provided in several hours. Thus, this protocol is easier technically because food does not have to be removed from cages daily. However, this protocol could be problematic because caloric restriction changes the phase of the SCN circadian rhythm (Challet, 2010). Since the FEO is coupled to the SCN, then changes in the phase of FAA could result from the effects of caloric restriction on the SCN, rather than the FEO. We have reviewed the studies of FAA that used a caloric restriction protocol herein, but they should be interpreted with caution.

It is also critical to assess the endogenous, self-sustained nature of the FEO. This means that FAA must be assessed both during restricted feeding and in the absence of a feeding time cue. Since FAA disappears during *ad libitum* feeding, the endogenous nature of the FEO is assessed during fasting (Fig. 1C, D). Notably, 2 days of fasting are required. The appearance of FAA on the second day of fasting shows that it is the output of a self-sustained oscillator (rather than an hourglass timer which would still drive FAA on the first day of fasting). In this review, we will highlight studies that assessed FAA during both restricted feeding and fasting.

Circadian gene mutant mice and FAA

The SCN and peripheral tissue clocks use the canonical circadian timekeeping mechanism (Takahashi, 2016). However, mounting evidence strongly suggests that the FEO oscillation

does not require the canonical circadian transcription translation feedback loop(s) (Table 1). Two independent studies showed that *Clock*¹⁹ mice had FAA during restricted feeding and subsequent fasting in the light-dark cycle (Pitts et al., 2003; Horikawa et al., 2005). However, these experiments alone were not sufficient to demonstrate that the FEO used a non-canonical timekeeping mechanism. This is because the locomotor activity of *Clock*¹⁹ mice entrained to the light-dark cycle, thus the light-driven SCN oscillation could have controlled food anticipation. SCN-controlled locomotor activity becomes arrhythmic in constant darkness in homozygous *Clock*¹⁹ mutant mice (i.e. the SCN is disabled in constant darkness) (Vitaterna et al., 1994). Pitts et al. also showed that FAA was present during restricted feeding in *Clock*¹⁹ mice in constant darkness (2003). However, since they did not fast the *Clock*¹⁹ mutant mice after restricted feeding in constant darkness, the possibility remained that the SCN participated in food entrainment (Pitts et al., 2003). Two later studies of other circadian mutant mice did demonstrate that the canonical circadian loop is not required for timekeeping by the FEO. First, *Cry1*^{-/-}/*Cry2*^{-/-} double mutants had FAA during restricted feeding and fasting in the light-dark cycle (Iijima et al., 2005). Importantly, when the SCN was disabled in *Cry1*^{-/-}/*Cry2*^{-/-} mice (in either SCN-lesioned mice or in mice housed in constant darkness), FAA was still present during restricted feeding and fasting (Iijima et al., 2005). Similarly, we showed that FAA was present in *Per1*^{-/-}/*Per2*^{-/-}/*Per3*^{-/-} triple mutants during restricted feeding and fasting in constant darkness when the SCN was disabled (Pendergast et al., 2012). Storch et al. also showed that *Per1*^{-/-}/*Per2*^{-/-} double mutants in constant darkness had FAA during restricted feeding, but the mice were not subsequently fasted (2009). Together these studies show that, under ideal conditions to assess the timekeeping mechanism, the FEO keeps time in the absence of canonical clock genes.

Anticipation of food availability has also been examined in single *Period* mutant mice (Table 1). Several studies have shown that *Per1*^{-/-} and *Per3*^{-/-} mice have normal FAA (Feillet et al., 2006; Mendoza et al., 2010a; Li et al., 2015; Pendergast et al., 2017). However, the results from *Per2*^{-/-} mice were discrepant. *Per2* was implicated as essential for FEO timekeeping when Feillet et al. reported that *Per2*^{-/-} (Brdm1 line) mice lacked FAA during restricted feeding and fasting (Feillet et al., 2006). This finding was subsequently repeated by the same lab (in the same line and in a *Per2*^{fllox} line) and by another lab (after backcrossing with the C57BL/6J strain), although FAA during fasting was not examined in these studies (Mendoza et al., 2010a; Li et al., 2015; Chavan et al., 2016). In contrast, Storch and Weitz (2009) found that *Per2*^{-/-} (Brdm1 line) mice had FAA during restricted feeding. We also recently showed that both the Idc and Brdm1 lines of *Period2*^{-/-} mice had robust FAA during restricted feeding and fasting (Pendergast et al., 2017). It is likely that differences in experimental protocols were responsible for the discrepant results. Nonetheless, several studies have now observed robust FAA in *Per2*^{-/-} mice, demonstrating the *Per2* is not essential for FAA.

Bmal1^{-/-} mice are widely studied because *Bmal1* has been targeted to generate tissue-specific knockout or rescue of the circadian clock (Bunger et al., 2000). However, FAA has been difficult to study in *Bmal1*^{-/-} mice, in part because they are unhealthy (Table 1) (Rudic et al., 2004; Bunger et al., 2005; Kondratov et al., 2006). The first study of FAA in *Bmal1*^{-/-} mice showed that these mice had no FAA, but that FAA was rescued by injecting virus expressing *Bmal1* into the dorsomedial nucleus of the hypothalamus (Fuller et al., 2008).

However, several independent laboratories have since demonstrated that *Bmal1*^{-/-} mice had FAA (Mistlberger et al., 2008; Pendergast et al., 2009; Storch and Weitz, 2009; Mieda and Sakurai, 2011; Takasu et al., 2012; Izumo et al., 2014). The discrepancy in results could be attributed, in part, to the fact that *Bmal1*^{-/-} mice may have been ill during restricted feeding in the Fuller et al. study because food availability was not gradually reduced (for further discussion of FAA in *Bmal1*^{-/-} mice, see (Fuller et al., 2009; Mistlberger et al., 2009b; Mistlberger et al., 2009a). In sum, numerous studies, performed in different labs have demonstrated that BMAL1 is not necessary for FAA.

Although circadian genes are not necessary for FEO timekeeping, several studies have shown that they modulate FAA. Dudley et al. showed that mice lacking functional Neuronal PAS domain protein 2 (*Npas2*), the paralog of *Clock*, had delayed expression of FAA, but after several days of restricted feeding displayed FAA that was indistinguishable from wild-types (2003). Our lab also showed that the period of the FEO rhythm was shortened in *Per1*^{-/-}/*Per2*^{-/-}/*Per3*^{-/-} triple mutant mice (Pendergast et al., 2012). Likewise, Takasu et al. showed that the FEO period was shortened in SCN-lesioned *Cry1*^{-/-} mice and lengthened in SCN-lesioned *Cry2*^{-/-} mice (2012). Global and nervous system-specific *Nr1d1*^{-/-} (*Rev-erba* mutant mice) mice also had attenuated FAA (Delezie et al., 2016).

In sum, *Cry1*, *Cry2*, and *Per1/2/3* alter the period of FAA and therefore likely participate in the FEO timekeeping mechanism (although they are not necessary for FEO timekeeping; Fig. 2A). On the other hand, *Npas2* and *Rev-erba* (*Nr1d1*) modulate FAA via the input or output pathways, or FEO timekeeping itself (Fig. 2B).

FAA in neurotransmitter/neuromodulator mutant mice

Most neurotransmitter mutant mice examined to date expressed normal FAA, except mice with disrupted dopamine signaling (Table 2). Dopamine signaling is involved in reward-seeking behaviors (Baik, 2013). Rodents anticipate not only timed chow diet, but also rewarding timed palatable meals and stimulant drugs (Mistlberger and Rusak, 1987; Jansen et al., 2012; Keith et al., 2013; Flores et al., 2016b). In addition, dopamine-deficient mice do not forage for food (Szczyepka et al., 1999). Thus, it has been postulated that the FEO may require (or be located in) the dopaminergic circuitry. Several studies have examined the effects of pharmacological activation or inhibition of dopamine signaling on FAA. Mice co-treated with D1 and D2 receptor antagonists had attenuated FAA (Liu et al., 2012). In addition, mice developed anticipatory activity to timed treatment with a D1 receptor agonist even without restricted feeding (Gallardo et al., 2014). Consistent with these pharmacological studies, mice lacking functional dopamine D1 receptor (*Drd1*^{-/-}) had attenuated FAA during restricted feeding (Gallardo et al., 2014; Michalik et al., 2015). Restricted feeding with caloric restriction (which could alter the phase of the SCN rhythm) showed that FAA was normal in dopamine D2 receptor mutant (*Drd2*^{-/-}) mice (Gallardo et al., 2014). Moreover, virus-mediated rescue of dopamine expression in the dorsal striatum of dopamine-deficient mice (that do not express FAA because they do not forage at all) permitted the development of FAA during restricted feeding (Gallardo et al., 2014). Together these studies implicate the dopaminergic brain circuitry, and the dopamine D1 receptor, as a

regulator of FAA. There may be functional redundancy in the circuitry since FAA was not abolished in dopamine receptor mutant mice.

Studies have shown that FAA is normal in other neurotransmitter and neuromodulator mutant mice (Table 2). μ opioid receptor signaling modulates dopaminergic signaling before anticipation of reward in rodents, and therefore could regulate FAA via dopaminergic or other signaling pathways (Spruijt et al., 2001). However, mice with global knock out of the μ -opioid receptor had FAA during restricted feeding (Kas et al., 2004). Likewise, although serotonin signaling via the serotonin 2C receptor (5-HT_{2C}R) regulates food intake (Tecott et al., 1995), *5-HT_{2C}R*^{-/-} mice had FAA during both restricted feeding and subsequent fasting (Hsu et al., 2010).

In summary, disruption of reward-related pathways modulates the development and robustness of FAA. However, it is not clear if the components of the reward circuitry participate in the input pathway, the pacemaker, and/or the output pathway (Fig. 2B).

FAA in hormone and hormone signaling mutant mice

Several hormones in the peripheral circulation and brain regulate food intake and energy balance and therefore are poised to regulate FAA. However, contrary to their roles in homeostatic regulation of food intake, it appears that they modulate, but are not necessary for, FAA (Table 3).

Leptin is released from fat stores and suppresses food intake (Zhang et al., 1994). In rodents, the peak of the daily rhythm of plasma leptin coincides with meal timing (Ahima et al., 1998; Bodosi et al., 2004; Arble et al., 2011), suggesting that it could regulate FAA. On the other hand, the leptin rhythm, unlike FAA, does not persist during fasting, suggesting that leptin may not contribute to FAA (Martinez-Merlos et al., 2004). Indeed, *ob/ob* mice, which have mutated leptin, had enhanced or normal FAA during restricted feeding and normal FAA during subsequent fasting compared to control animals (Gunapala et al., 2011; Ribeiro et al., 2011). These studies demonstrate that leptin is not necessary for FAA, but can regulate its robustness.

Another hormone that could regulate FAA is ghrelin. Plasma ghrelin increases before habitual mealtime and it activates dopaminergic signaling in reward brain circuits in rodents (Bodosi et al., 2004; Abizaid et al., 2006). Several studies in transgenic mice have demonstrated that ghrelin modulates the expression of FAA. Mice with non-functional ghrelin receptors (*Ghsr*^{-/-} from Regeneron) had attenuated FAA during restricted feeding and fasting (Blum et al., 2009; LeSauter et al., 2009). *Ghsr1a*^{-/-} (from Astrogeneca) mice also had attenuated FAA during restricted feeding. Davis et al. reported that FAA was absent during restricted feeding in *Ghsr*^{-/-} mice (from Zigman & Elmquist) (Zigman et al., 2005; Davis et al., 2011). However, in this study, locomotor activity, measured as photobeam breaks, was measured only for 2h before scheduled feeding (Davis et al., 2011). We have shown that wheel-running activity anticipates food availability while general (non-wheel) locomotor activity occurs during scheduled feeding (Flores et al., 2016a). Therefore, because general (non-wheel) activity was measured during a brief time window prior to restricted feeding, it is unknown whether the *Ghsr*^{-/-} mice expressed FAA during food

availability. Mice lacking functional proghrelin, which is the precursor to ghrelin, or mice that lacked the ghrelin ligand, had normal FAA during restricted feeding (Szentirmai et al., 2010; Gunapala et al., 2011). Together, these studies show that ghrelin signaling modulates the expression of FAA, but is not necessary for FAA.

Leptin and ghrelin signaling pathways converge on the melanocortin system in the hypothalamus (Ellacott and Cone, 2006). Thus, while leptin and ghrelin are not necessary for FAA, it is possible that the central melanocortin mechanisms regulating food intake could mediate FAA. The hypothalamic melanocortin system regulates energy homeostasis by coordinating food intake and energy expenditure (Ellacott and Cone, 2006). Of the five melanocortin G-protein coupled receptors, melanocortin receptors 3 and 4 (*Mc3r* and *Mc4r*), regulate energy homeostasis. MCR agonists, such as the proopiomelanocortin (*POMC*)-derived α -melanocyte stimulating hormone (α -MSH), inhibit food intake, while MCR antagonists, such as agouti-related protein (AgRP), stimulate feeding (Fan et al., 1997; Anderson et al., 2016). *Mc4r* is widely expressed throughout the brain, while *Mc3r* expression is relatively limited and was detected at high levels in the hypothalamus and limbic/reward circuitry (e.g. ventral tegmental area) (Roselli-Reh fuss et al., 1993; Liu et al., 2003). Reward brain circuits have been implicated in regulation of FAA (see discussion above), suggesting that *Mc3r* could underlie FAA. *Mc4r*^{-/-} mice had normal FAA, while *Mc3r*^{-/-} mice had attenuated FAA, during restricted feeding, but FAA was not measured during subsequent fasting (Sutton et al., 2008). Since central melanocortin signaling is downstream of leptin, Ribeiro et al. determined whether mice lacking both leptin and *Mc3r* [*ob/obMc3r*^{-/-}] would have impaired FAA (2011). Instead, *ob/obMc3r*^{-/-} double mutant mice had enhanced FAA during restricted feeding compared to *ob/ob* mice (Ribeiro et al., 2011). Therefore, the melanocortin system is not necessary for FAA.

Melanin-concentrating hormone (MCH) regulates energy homeostasis and mice lacking functional MCH receptor (*Mch1r*^{-/-}) are hyperphagic (Chen et al., 2002; Marsh et al., 2002). Despite its role in regulating food intake, FAA in *Mch1r*^{-/-} mice was indistinguishable from wild-type controls (Zhou et al., 2005).

Together these data of mutant mice with non-functional hormone signaling pathways suggest that the well-characterized brain circuitries that regulate food intake and energy homeostasis are not necessary for FAA. Instead, ghrelin, leptin, and the melanocortin signaling pathways appear to regulate the magnitude of FAA during restricted feeding. This regulation could result from altering the input pathway, the amplitude of the FEO, and/or the output pathway (Fig. 2B).

FAA in neuropeptide mutant mice

Neuropeptide signaling pathways that are related to food intake and arousal could regulate FAA (Table 4). Orexin A and B are neuropeptides that signal via OX1R and OX2R and regulate arousal (Alexandre et al., 2013). Infusion of orexin A can stimulate food intake while OX1R antagonists reduce food intake, suggesting that the orexin signaling pathway could constitute the FEO (Lubkin and Stricker-Krongrad, 1998; Sakurai et al., 1998; Smart et al., 2002). Postnatal ablation of orexin neurons attenuated FAA in orexin/ataxia-3 mice (Akiyama et al., 2004; Mieda et al., 2004). However, the ablated orexin neurons in this study

also expressed dynorphin, an endogenous opioid, and galanin, a neuropeptide that has been implicated in feeding. A subsequent study of orexin mutant mice (*Hcrt*^{-/-}) in which only orexin was non-functional, showed that FAA was attenuated, but the anticipatory rise in body temperature prior to restricted feeding was normal (Kaur et al., 2008). FAA (activity and body temperature) persisted during fasting after restricted feeding in *Hcrt*^{-/-} mice (Kaur et al., 2008). Two other studies measured behavior prior to food availability using video cameras and found that *Hcrt*^{-/-} mice had either attenuated (Clark et al., 2009) or normal (Gunapala et al., 2011) FAA using a caloric restriction restricted feeding protocol (neither study measured FAA during subsequent fasting). Finally, disabling the high-frequency firing of orexin neurons, in mice lacking functional TWIK-related acid-sensitive potassium channels (TASK1/3), did not affect FAA during restricted feeding or subsequent fasting (Gonzalez et al., 2009).

Together these studies show that orexin is not necessary for FAA, but it modulates the expression of FAA (Fig. 2B).

Neuropeptide Y (NPY) released from neurons in the arcuate nucleus of the hypothalamus potently stimulates food intake (Loh et al., 2015). In addition, NPY expression increases in the hypothalamus prior to restricted feeding, suggesting that NPY could underlie or regulate FAA (Yoshihara et al., 1996). However, NPY mutant mice had normal FAA during calorically-restricted restricted feeding, demonstrating that NPY is not necessary for FAA (Gunapala et al., 2011).

FAA in other mutant and engineered mice

FAA has also been investigated in various other mutant and engineered mice (Table 5). Prokineticin2 (PK2) is a secreted protein that has been postulated to act as an SCN output factor (Cheng et al., 2002). This is, in part, because PK2 is highly expressed in the SCN, and PK2^{-/-} mice have low-amplitude activity and body temperature rhythms (but normal SCN circadian gene rhythms) (Li et al., 2006). If PK2 is an output of the SCN, then perhaps it is not surprising that PK2^{-/-} mice had normal FAA, since FAA persists in SCN-lesioned mice (Li et al., 2006).

Vasoactive intestinal peptide (VIP) signals through the VIP receptor2 to synchronize the rhythms of SCN neurons (Aton et al., 2005; Maywood et al., 2006). Mice lacking VIP receptor 2 (*Vipr2*^{-/-}) had impaired SCN and locomotor activity rhythms, but normal FAA during restricted feeding (Sheward et al., 2007).

Pituitary adenylate cyclase activating polypeptide (PACAP), together with glutamate, transmits light signals from the intrinsically photosensitive retinal ganglion cells in the retina to the SCN using the PACAP receptor 1 (PAC1). PAC1^{-/-} mice had normal FAA during restricted feeding (Hannibal et al., 2016).

The expression of FAA may rely on detection of the metabolic state of cells. Cellular NAD⁺/NADH levels are a readout of metabolism. Several studies have examined FAA in mice with mutations in the molecular machinery that participates in sensing and responding to cellular NAD⁺/NADH levels. Poly(ADP-Ribose) Polymerase 1 (PARP1) is an NAD⁺-

dependent ADP-ribosyltransferase that regulates the activity of transcriptional regulatory proteins (Schreiber et al., 2006). Thus, PARP1 is poised to sense changes in cellular metabolism and enact transcriptional responses. However, PARP1 is not necessary for FAA since PARP1^{-/-} mice had normal FAA during restricted feeding (Asher et al., 2010). Sirtuins are another family of proteins that require NAD, but not NADH, for their deacetylase activity (Imai and Guarente, 2010). Therefore, they also act as cellular metabolism sensors. Sirtuin 1 (SIRT1) has been shown to regulate physiological responses to diet restriction (Imai and Guarente, 2010). In addition, SIRT1 is required for increased locomotor activity during fasting in mice (Chen et al., 2005). Therefore, SIRT1 could regulate FAA since it both senses cellular metabolism and can regulate locomotor activity. One study showed that *Sirt1*^{-/-} mice had attenuated FAA, while brain-specific overexpression of SIRT1 increased FAA (Satoh et al., 2010). Therefore, while SIRT1 regulates FAA, it is not necessary for its expression (Fig. 2B).

One study investigated the role of the cerebellum in FAA (Mendoza et al., 2010b). The cerebellum coordinates locomotor activity and motivational processes (Caston et al., 1998). Moreover, the molecular circadian *Period1-Luciferase* rhythm advanced in cerebellar explants in response to daytime restricted feeding (Mendoza et al., 2010b), suggesting that the cerebellum could harbor the FEO. When the function of the cerebellar circuitry was impaired in Nancy hotfoot (*Grid^{ho/ho}*) mutant mice, which lack the coding sequence for the ionotropic glutamate receptor $\delta 2$ gene in Purkinje cells, FAA was attenuated, but still present, during restricted feeding (Mendoza et al., 2010b).

The neurodegenerative Huntington's disease is characterized by progressive motor symptoms and sleep and circadian disruption (Morton et al., 2005). The R6/2 mouse is a model of Huntington's disease that has the 5' end of the human Huntington gene with about 120 CAG repeat expansions (Mangiarini et al., 1996). R6/2 mutant mice have progressive disruption of circadian rhythms of locomotor activity and circadian gene expression in the SCN (Morton et al., 2005). Consistent with the numerous studies that have shown that the SCN and FEO are distinct, FAA during restricted feeding was intact in R6/2 mutant mice (Maywood et al., 2010) (Table 5). Importantly, restricted feeding restored daily rhythms of locomotor activity in late-stage R6/2 mice that previously had arrhythmic activity (Maywood et al., 2010). These data support the hypothesis that the FEO is an oscillator that can operate at the top of the circadian hierarchy when the SCN is disabled (Pendergast and Yamazaki, 2017). Restricted feeding could serve as a therapeutic to restore rhythms via the FEO in Huntington's disease patients.

The role of signaling molecules in regulating FAA have also been investigated (Table 5). RGS16 regulates G-protein coupled receptor signaling and is rhythmically expressed in the SCN and liver. Since *Rgs16* expression in the liver was altered by fasting and re-feeding, Hayasaka et al. (2011) examined FAA in mice with transgenic knockdown of *Rgs16* (Huang et al., 2006). *Rgs16* knockdown mice had attenuated FAA during restricted feeding (Huang et al., 2006). Protein kinase C γ (PKC γ) also regulates molecular pathways, is expressed almost exclusively in neurons, and in brain regions that entrain to restricted feeding (Saito and Shirai, 2002; Verwey and Amir, 2009). However, PKC γ ^{-/-} mice had normal FAA

during restricted feeding that was indistinguishable from wild-type mice (Zhang et al., 2012).

The Wolfram syndrome 1 (WSF1) gene is expressed in the DMH (and widely expressed in the brain), but only weakly expressed in the SCN (Luuk et al., 2008). The DMH was initially implicated as the locus of the FEO, although subsequent studies demonstrated that the DMH modulated FAA, but was not the FEO (Gooley et al., 2006; Landry et al., 2006; Mieda et al., 2006; Fuller et al., 2008; Moriya et al., 2009; Acosta-Galvan et al., 2011). Because of the DMH's putative role in food entrainment, and the expression of *Wsf1* in the DMH, one study examined FAA in *Wsf1*^{-/-} mice (Table 5). They found that FAA was normal in *Wsf1*^{-/-} mice (Luuk et al., 2012).

Brain-derived neurotrophic factor (BDNF) signals via TrkB receptors to suppress food intake and regulate energy homeostasis (Marosi and Mattson, 2014). The production of BDNF is, in part, regulated by tissue-type plasminogen activator (tPA) (Pang et al., 2004). Therefore, tPA could play a role in regulating FAA. However, FAA was normal in *tPA*^{-/-} mice, suggesting that mature BDNF signaling is not necessary for FAA (Krizo et al., 2018).

Conclusions

Despite decades of research, the molecular and anatomical substrates of circadian food entrainment remain a mystery. While mutant and engineered mouse models have been key to the discovery of many physiological processes and behaviors in the field of circadian rhythms, the use of these models for studying the FEO have been largely unsuccessful. One conclusion that is clear from studies of mouse models is that canonical circadian genes are not required for the expression of FAA or for FEO timekeeping. Although this conclusion may fall into the category of 'negative results,' the implications of these studies are exciting and far-reaching. That the FEO can keep circadian time in the absence of the canonical clock genes means that there is (at least) one other robust circadian timekeeping mechanism that can function independently from the SCN. This timekeeping mechanism remains to be discovered. The only other non-canonical circadian oscillators discovered to date are the reduction-oxidation cycles of peroxiredoxin proteins (Reddy, 2016). Thus, it is possible that the FEO uses a circadian timekeeping mechanism akin to the redox and metabolic oscillators.

Another overarching conclusion from the study of mouse models is that signaling pathways that regulate food intake and energy homeostasis, or respond to changes in cellular metabolism, regulate the magnitude of FAA, but are not necessary for FAA. Experiments have not been performed to examine the role of these hormones in FEO timekeeping (i.e. in SCN-lesioned mice during fasting subsequent to restricted feeding). Energy homeostasis signals converge on the dopaminergic brain circuitry, which has also been shown to be important for food entrainment (Gallardo et al., 2014; Michalik et al., 2015). However, it is impossible to determine if the FEO is functional in mice with abolished dopamine signaling. This is because dopamine-deficient mice do not forage (or have normal locomotion) and thus they die without intervention (Zhou and Palmiter, 1995).

Critical review of studies of FAA in mutant and engineered mouse models also reveals the technical challenges of studying the FEO. First, there is little consistency among protocols for studying FAA. Studies use different outcome measures (e.g. wheel vs. general activity), different phases and durations of restricted feeding (e.g. 3h or 4h window, or caloric restriction), and different quantifications of FAA (e.g. 3h vs. 1h before food presentation or normalized vs. not normalized to total activity). And, importantly, the majority of studies did not examine whether FAA persisted during fasting after restricted feeding to test the fundamental circadian property of persistence in constant conditions. These inconsistencies across studies make it difficult to systematically compare the factors that modulate FAA.

We have presented 2 protocols for studying FAA that we have optimized for detecting FAA in mutant and engineered mouse models. In the first protocol (Fig. 1A), coupling between the SCN and FEO is intact. In the second protocol (Fig. 1B), the SCN is disabled so FAA is measured without FEO-SCN coupling. Because genetically altered mice sometimes have known or unknown health problems, we have found that it is critical to gradually reduce food availability (from 8h- to 6h- and finally to 4h/day). In addition, it is sometimes necessary to provide food on the bottom of the cage, even if the mutant mice are healthy when given food in the hopper during *ad libitum* feeding (as with *Bmal1*^{-/-} and *Drd1*^{-/-} mice) (Pendergast et al., 2009; Gallardo et al., 2014). FAA is also more robust when wheel-running activity (compared to general activity) is measured. It is also important to note that the phase of FAA may be altered depending on the mode of activity. Wheel-running FAA typically occurs during the hours immediately before food availability and stops abruptly when food is presented. In contrast, general locomotor FAA (e.g. measured by a passive infrared sensor) typically occurs during food availability. Finally, the timing and duration of fasting should be carefully considered. We have found that 48 hours of fasting is ideal to reveal the expression of robust FAA. Fasting should be timed so that hours 40 to 48 of fasting occur during the predicted time of FAA. FAA may be weak or absent on the first day of fasting, but is typically robustly expressed on the second day of fasting. Moreover, fasting is necessary to determine if FAA is the output of an hourglass timer or a self-sustained oscillator. FAA on the first day of fasting could be the output of an hourglass timer, while FAA on the second day of fasting is the output of a self-sustained oscillator.

Finally, studies of FAA expression must be interpreted with caution. Changes in the robustness, or magnitude, of FAA could result from changes in the input or output, and not necessarily from changes in the FEO amplitude. The best evidence that a gene is important for FEO timekeeping is if mutating that gene causes a change in the period of FAA. However, measuring the period of the FEO is challenging (because it doesn't free-run in *ad libitum* feeding conditions). Only a few studies have reported free-running FAA under a feeding cycle that is outside the range of entrainment (Stephan, 1992; Pendergast et al., 2012; Takasu et al., 2012). The period of the FEO can be estimated in mutant and engineered mice by measuring the range of entrainment using different T cycles of food availability. However, this approach is costly, time-consuming, and difficult.

Acknowledgments

We believe we have reviewed all FAA studies in mutant and genetically engineered mice herein. If we have overlooked a study, we apologize and would like to be notified of said study. J.S.P is supported by National Institutes of Health (NIH) grants DK098321, P20 GM103527 (Junior Investigator), DK107851, and the University of Kentucky. S.Y. is supported by NIH grant R21 NS099809.

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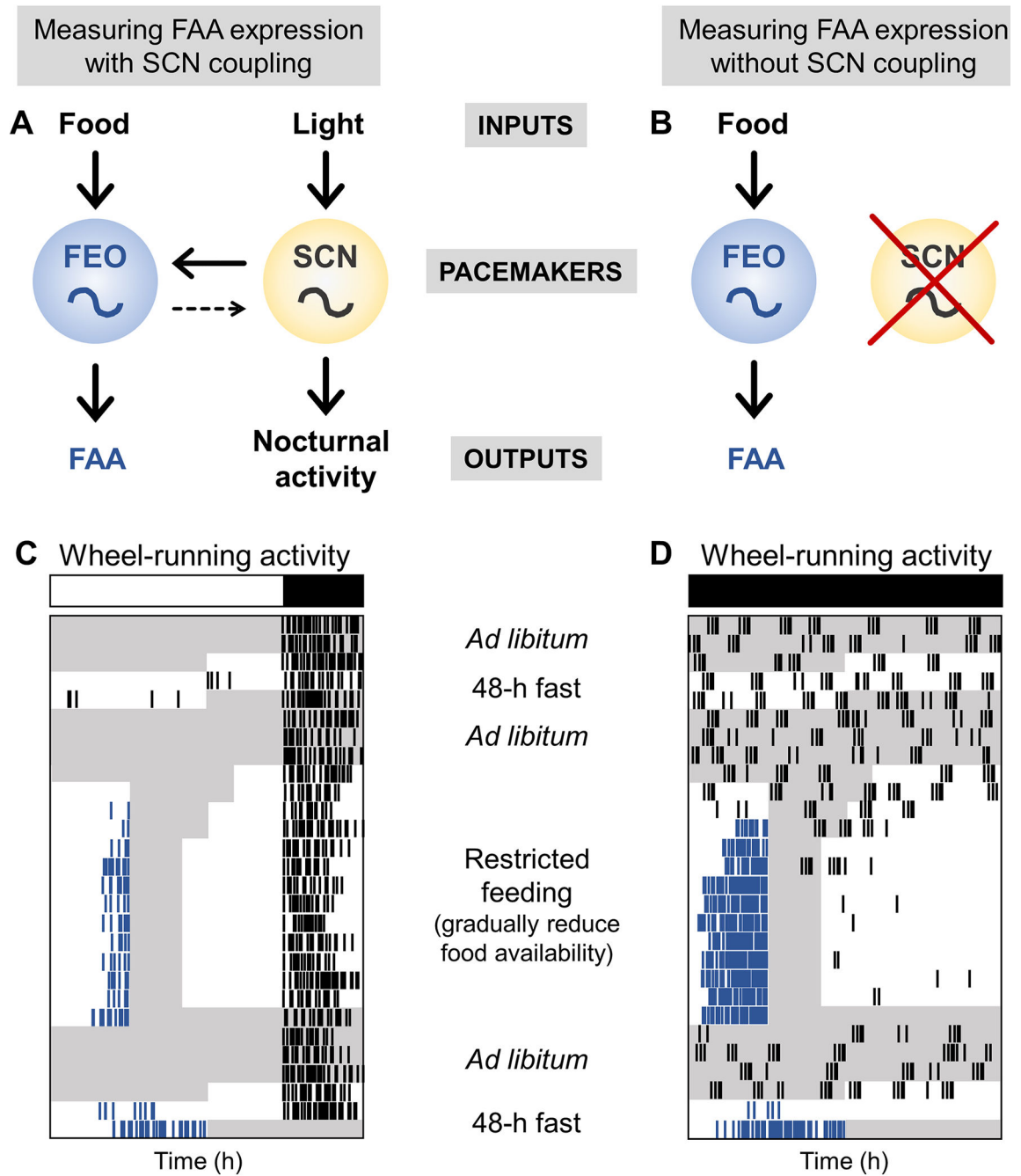


Figure 1. Experimental protocols to measure FAA expression.

A. The inputs, food and light, entrain the pacemakers, the FEO and SCN, which thereby regulate the outputs, FAA and nocturnal activity, respectively. The FEO is strongly coupled to the SCN, while the SCN is weakly coupled to the FEO. **B.** When the SCN is disabled, either by SCN lesion, or by housing arrhythmic mutant mice in constant darkness, FAA expression is measured without the influence of the SCN on the FEO. **C.** To measure robust FAA when the FEO is coupled to the SCN, mice are housed in 18L:6D and wheel-running activity is recorded. **D.** To measure FAA expression without the influence of the SCN, mice

(SCN-lesioned or arrhythmic mice) are housed in constant darkness. In both protocols, mice are fasted for 48-h before restricted feeding and then returned to ad libitum feeding. Food availability is then gradually reduced (8h/day to 6h/day to 4h/day). Food is manually added and removed 4h later. The mice are allowed to eat as much as possible during the restricted feeding window. Food may be placed on the bottom of the cage during restricted feeding. Mice are then fed for 4h/day for 10 days. Mice are then fed *ad libitum* for several days and then fasted for 48h. Fasting should be timed so the hours 40 to 48 of fasting occur during the predicted time of FAA expression. White and black bars represent light and dark, respectively. Gray shading shows when food is available. Blue ticks show wheel-running activity controlled by the FEO and black ticks show other wheel-running activity. The x-axis is time (in hours). Mice may be fed on non-24h cycles (in D) to determine the period of the FEO. The y-axis is days.

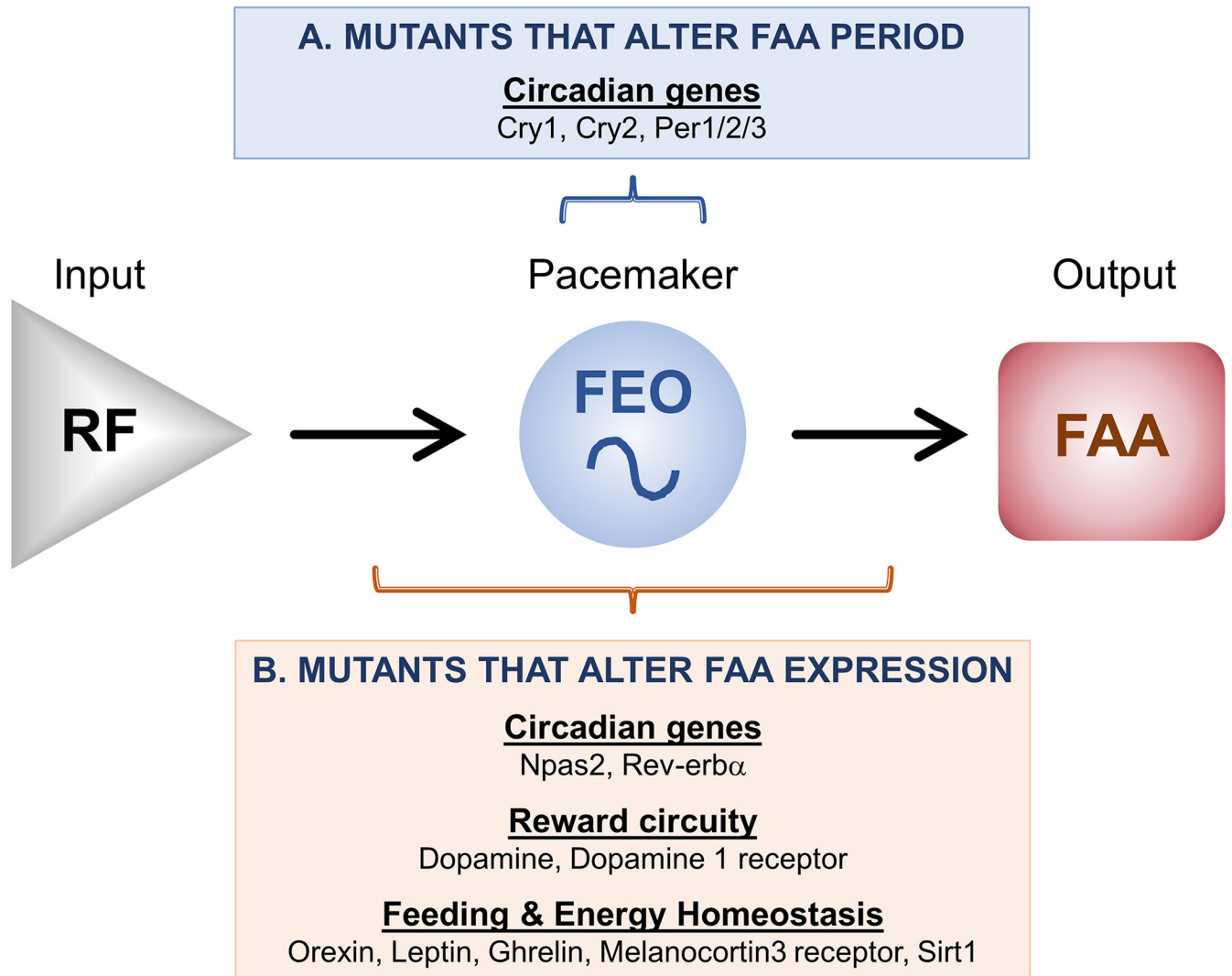


Figure 2. Genes that modulate FAA expression.

A. Several circadian mutant mice have altered periods of FAA expression, suggesting that these genes participate in the timekeeping mechanism of the FEO. **B.** Genes involved in circadian rhythmicity, reward circuitry, and feeding and energy homeostasis have altered FAA expression levels. These genes could modulate the input pathway, the FEO pacemaker, and/or the output pathway.

Table 1.

Food anticipatory activity in circadian gene mutant and engineered mice.

Gene disrupted	Line	Strain	RF phase (ZT)	Output	FAA during RF	FAA during fasting	References
<i>Clock</i>	19	C57BL/6J x BALB/cj	6-10	Wheel	Present ¹	Present	(Pitts et al., 2003)
	19	C57BL/6J	5-9	Thermal radiation detector	Present	Present	(Horikawa et al., 2005)
<i>Npas2</i>	Global	C57BL/6J (N9)	6-10	Wheel	Attenuated ²	ND	(Dudley et al., 2003)
<i>Cry1/Cry2</i>	Global	129 x C57BL/6 x ICR	5-9	Infrared sensor	Present ³	Present ³	(Iijima et al., 2005)
<i>Cry1</i>	Global	C57BL/6	SCN-X ⁴	Infrared sensor	Present ⁴	Present ⁴	(Takasu et al., 2012)
<i>Cry1</i>	Over-express C414A mutation	BDF1	5-9	Wheel	Present ¹	Present	(Okano et al., 2016)
<i>Cry2</i>	Global	C57BL/6	SCN-X ⁴	Infrared sensor	Present ⁴	Present ⁴	(Takasu et al., 2012)
<i>Bmal1</i>	Global	N5	4-8	Telemetry	Absent	ND	(Fuller et al., 2008)
	Global	C57BL/6J	6-10	Infrared sensor	Present	Present	(Mistlberger et al., 2008)
	Global	C57BL/6J	8-12 ⁵	Wheel	Present ¹	Present ⁶	(Pendergast et al., 2009)
	Global	N5	6-9 ⁷	Wheel	Present ¹	ND	(Storch and Weitz, 2009)
	Global	C57BL/6J	DD	Infrared sensor	Present	Absent	(Takasu et al., 2012)
	Nestin-Cre floxed	N5-6	5-8	Infrared sensor	Present	Present	(Mieda and Sakurai, 2011)
	Camk2a::iCre B floxed	Not reported	6-10	Wheel	Present	ND	(Izumo et al., 2014)
	Atoh1-Cre floxed	C57BL/6 x 129	6-12	Telemetry	Present	ND	(Bering et al., 2017)
<i>Period1</i>	Brdm1	C57BL/6x129S5/SvEvBrd	4-12 & 4:66% CR	Wheel	Present	Present ⁸	(Feillet et al., 2006)
	Brdm1	C57BL/6x129S5/SvEvBrd	6-12	Wheel	Present	ND	(Mendoza et al., 2010)
	Brdm1	C57BL/6J	4-8	Wheel	Present	ND	(Li et al., 2015)
	Idc	C57BL/6J	6-10	Wheel	Present	Present	(Pendergast et al., 2017)
<i>Period2</i>	Brdm1	C57BL/6x129S5/SvEvBrd	4-12	Wheel	Absent	Obscured by free-running SCN-controlled rhythm	(Feillet et al., 2006)
	Brdm1	C57BL/6x129S5/SvEvBrd	6-12	Wheel	Absent	ND	(Mendoza et al., 2010)

Gene disrupted	Line	Strain	RF phase (ZT)	Output	FAA during RF	FAA during fasting	References
	Brdm1	C57BL/6J	4-8	Wheel	Weak or absent	ND	(Li et al., 2015)
	Brdm1	129/C57BL/6-Tyrc-Brd	6-10	Wheel	Present	Present	(Pendergast et al., 2017)
	Idc	129xC57BL/6	6-9	Wheel	Present	ND	(Storch and Weitz, 2009)
	Idc	C57BL/6J	6-10	Wheel	Present	Present	(Pendergast et al., 2017)
	CMV-Cre floxed (global)	C57BL/6	4-12; 70%-80% CR	Wheel & telemetry	Attenuated	ND	(Chavan et al., 2016)
	Nestin-Cre floxed (brain)	C57BL/6	4-12; 70%-80% CR	Wheel & telemetry	Present ⁹	ND	(Chavan et al., 2016)
	Albumin-Cre floxed (liver)	C57BL/6	4-12; 70%-80% CR	Wheel & telemetry	Attenuated ⁹	ND	(Chavan et al., 2016)
<i>Period3</i>		C57BL/6J	6-10	Wheel	Present	Present	(Pendergast et al., 2017)
<i>Per1/Per2</i>	Idc	129	6-9	Wheel	Present ¹	ND	(Storch and Weitz, 2009)
	Brdm1	C57BL/6x129S5/SvEvBrd	DD	Wheel	Present	ND	(Mendoza et al., 2010)
<i>Per1/Per2/Per3</i>	Idc	C57BL/6J	8-14 ⁵	Wheel	Present	Present ¹⁰	(Pendergast et al., 2012)
<i>Nr1d1^{-/-}</i>	Global	C57BL/6J	6-12	Telemetry	Attenuated	ND	(Delezie et al., 2016)
	Nestin-Cre floxed	C57BL/6J	6-12	Telemetry	Absent or attenuated ⁹	Absent	(Delezie et al., 2016)

All experiments were performed in 12L:12D unless otherwise indicated. All mice had intact (not lesioned) SCN unless otherwise indicated. RF: restricted feeding; FAA: food anticipatory activity; ND: not determined; CR: caloric restriction

¹ FAA was also present in constant darkness

² FAA was attenuated and had delayed onset in constant darkness

³ FAA was also present during restricted feeding and fasting in constant darkness. Additionally, FAA was present in SCN-lesioned *Cry1/Cry2* double knockout mice during restricted feeding and fasting in LD

⁴ SCN-X: These mice were SCN-lesioned

⁵ These experiments were performed in 18L:6D and restricted feeding began 8h after lights on

⁶ In constant darkness, the mice are hyperactive and FAA could not be resolved

⁷ These experiments were performed in 16L:8D and feeding began 6h after lights on

⁸ Fasting was performed in constant darkness

⁹ Identical actograms of body temperature were incidentally presented for both Nestin-Cre-Per2flox and Albumin-Cre-Per2flox mice

¹⁰ In constant darkness, *Per1/Per2/Per3* knockout mice exhibited FAA during restricted feeding and fasting when food was provided on a 21-h cycle (but not 24-h cycle)

Table 2.

Food anticipatory activity in neurotransmitter/neuromodulator mutant mice

Gene disrupted	Line	Strain	RF phase (ZT)	Output	FAA during RF	FAA during fasting	References
<i>Drd1</i>	Global	C57BL/6J (N8)	6-10	Disc	Attenuated	ND	(Gallardo et al., 2014)
	Global	C57BL/6J (N8)	8; 60% CR ¹	Video	Present	ND	(Gallardo et al., 2014)
	Global	C57BL/6J (>N10)	6-10	Disc	Attenuated	ND	(Michalik et al., 2015)
<i>Drd2</i>	Global	C57BL/6J	8; 60% CR ¹	Video	Present	ND	(Gallardo et al., 2014)
<i>μ-opioid receptor</i>	Global	C57BL/6	12-15	Wheel	Present	ND	(Kas et al., 2004)
<i>5-HT2CR</i>	Global	C57BL/6J (N9)	4-8	Photobeam	Present ²	Present	(Hsu et al., 2010)

All experiments were performed in 12L:12D unless otherwise indicated. All mice had intact (not lesioned) SCN unless otherwise indicated. RF: restricted feeding; FAA: food anticipatory activity; ND: not determined; CR: caloric restriction

¹Experiments were performed in 13L:11D

²FAA was also present in constant dim red light

Table 3.

Food anticipatory activity in hormone mutant mice.

Gene disrupted	Line	Strain	RF phase (ZT)	Output	FAA during RF	FAA during fasting	References
<i>ob/ob</i>	Global	C57BL/6J	12-16, then 4-8	Wheel & home-cage activity	Enhanced	Present	(Ribeiro et al., 2011)
<i>ob/ob</i>	Global	C57BL/6J	10; 60% CR ¹	Behavior recorded by video	Present, but some behaviors reduced	ND	(Gunapala et al., 2011)
<i>SGhr</i>	Global (Regeneron)	C57BL/6	6-14	Wheel	Attenuated	Attenuated	(LeSauter et al., 2009)
		C57BL/6J x DBA	4-8	Wheel; infrared beam	Attenuated ²	Attenuated	(Blum et al., 2009)
	Global (AstraZeneca)	C57BL/6	12-14	Wheel	Attenuated	ND	(Verhagen et al., 2011)
	Global (Zigman & Elmquist)	C57BL/6J	12:00-16:00 (no ZT given)	Photobeam breaks	Absent	ND	(Davis et al., 2011)
<i>Preproghrelin</i>	Global	C57BL/6J	4-8	Telemetry	Present	ND	(Szentirmai et al., 2010)
<i>Ghrelin</i>	Global	C57BL/6J x SJL/J	10; 60% CR ¹	Behavior recorded by video	Present	ND	(Gunapala et al., 2011)
<i>Mc3r</i>	Global	C57BL/6	7-11; 50% CR	Wheel & spontaneous activity (CLAMS)	Attenuated	ND	(Sutton et al., 2008)
<i>Mc3r & ob/ob</i>	Global	C57BL/6J	4-8	Wheel	Enhanced	ND	(Ribeiro et al., 2011)
<i>Mc4r</i>	Global	C57BL/6	7-11; 50% CR	Wheel	Present	ND	(Sutton et al., 2008)
<i>Mchr1</i>	Global	F3 C57BL/6J x 129SvEv	5-9	Wheel	Present	ND	(Zhou et al., 2005)

All experiments were performed in 12L:12D unless otherwise indicated. All mice had intact (not lesioned) SCN unless otherwise indicated. RF: restricted feeding; FAA: food anticipatory activity; ND: not determined; CR: caloric restriction

Note that data collected under conditions of restricted feeding with purposeful caloric restriction are not presented herein

¹Experiments were performed in 13L:11D

²FAA was also present in constant darkness (Lamont et al., 2014)

Table 4.

Food anticipatory activity in neuropeptide mutant mice.

Gene disrupted	Line	Strain	RF phase (ZT)	Output	FAA during RF	FAA during fasting	References
<i>Hcrt (orexin)</i>	Orexin neurons ¹ ; post-natal	C57BL/6J	5-9	Area sensor & thermal radiation detector	Attenuated	Attenuated	(Akiyama et al., 2004)
	Orexin neurons ¹ ; post-natal	C57BL/6J (N4-N7) x 129Svev F1	4-8	EEG-EMG & open field activity	Attenuated	Attenuated	(Mieda et al., 2004)
	Global	C57BL/6J	4-8	Telemetry-activity	Attenuated	Attenuated	(Kaur et al., 2008)
	Global	C57BL/6J	4-8	Telemetry-core body temperature	Present	Present	(Kaur et al., 2008)
	Global	C57BL/6J (N8)	15; 60% CR	Video and EEG-EMG	Attenuated	ND	(Clark et al., 2009)
	Global	C57BL/6J	10; 60% CR ²	Video	Present	ND	(Gunapala et al., 2011)
<i>K_{2p}3.1 & K_{2p}9.1 (TASK1/3)</i>	Global; impairs high frequency firing in orexin neurons	C57BL/6J x R1	3-7	Wheel	Present	Present	(Gonzalez et al., 2009)
<i>Npy</i>	Global	129S1	10; 60% CR ²	Video	Present	ND	(Gunapala et al., 2011)

All experiments were performed in 12L:12D unless otherwise indicated. All mice had intact (not lesioned) SCN unless otherwise indicated. RF: restricted feeding; FAA: food anticipatory activity; CR: caloric restriction; ND: not determined

¹These neurons also co-express dynorphin and galanin

²Experiments were performed in 13L:11D

Table 5.

Food anticipatory activity in other mutant and engineered mice.

Gene disrupted	Line	Strain	RF phase (ZT)	Output	FAA during RF	FAA during fasting	References
<i>Prokineticin2</i>	Global	C57BL/6 × 129/Ola	3-7	Infrared beam breaks	Present	Present	(Li et al., 2006)
<i>Vipr2</i>	Global	C57BL/6J	4-8	Wheel	Present	ND	(Sheward et al., 2007)
<i>PAC1 receptor</i>	Global	129	4-8	Wheel	Present ¹	ND	(Hannibal et al., 2016)
<i>Poly(ADP-Ribose) Polymerase 1</i>	Global	C57BL/6J	3-9	Wheel	Present	ND	(Asher et al., 2010)
<i>Sirt1</i>	Global	FVB (N6)	6-10	Wheel	Attenuated	ND	(Satoh et al., 2010)
	Global	CD1	7; 60% CR	Video	Present	ND	(Assali et al., 2018)
	Y/Y mutant, Global, no catalytic activity	CD1	7; 60% CR	Video	Present	ND	(Assali et al., 2018)
	CAGG-Cre-ER floxed (global in adulthood)	C57BL/6	9; 60% CR	Video	Present	ND	(Assali et al., 2018)
	Tyrosine hydroxylase-Cre floxed	C57BL/6	9; 60% CR	Video	Present	ND	(Assali et al., 2018)
	Pomc-Cre floxed	FVB/N x C57BL/6J x 129	9; 60% CR	Video	Present	ND	(Assali et al., 2018)
	Nestin-Cre floxed	C57BL/6J	9; 60% CR	Video	Present	ND	(Assali et al., 2018)
	CamKII-alpha-Cre floxed	C57BL/6	9; 60% CR	Video	Present	ND	(Assali et al., 2018)
<i>Sirt1 over-expression</i>	Brain (BRASTO)	C57BL/7J	6-10	Wheel	Enhanced	ND	(Satoh et al., 2010)
<i>Grid2^{ho/ho}</i>	Hotfoot mutation (ho/ho)	C57BL/6J	6-12	Wheel and telemetry	Attenuated	ND	(Mendoza et al., 2010)
<i>Human Huntington Disease gene with CAG repeats</i>	R6/2	C57BL/6 x CBA F1	6-11	Infrared sensors	Present	ND	(Maywood et al., 2010)
<i>Rgs16</i>	Global	C57BL/6J	6-10	Area sensor	Attenuated	ND	(Hayasaka et al., 2011)
<i>PKCγ</i>	Global	C57BL/6 × 129SvJ	6-10	Wheel	Present	ND	(Zhang et al., 2012)
<i>Wolfram syndrome 1</i>	Global	129 Sv/EvTac	4-8	Wheel	Present	ND	(Luuk et al., 2012)
<i>Tissue-type plasminogen activator</i>	Global	C57BL/6J	6-10	Wheel	Present	ND	(Krizo et al., 2018)

All experiments were performed in 12L:12D unless otherwise indicated. All mice had intact (not lesioned) SCN unless otherwise indicated. RF: restricted feeding; FAA: food anticipatory activity; ND: not determined; CR: caloric restriction

¹ FAA was also present in constant dim red light