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Circadian Clock Control of Endocrine Factors

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Abstract

Organisms experience dramatic fluctuations in demands/stresses over the course of the day. In order to maintain biological processes within physiologic boundaries, it is imperative that mechanisms have evolved for anticipation of, and adaptation to, these daily fluctuations. Endocrine factors undoubtedly play an integral role in homeostasis. Not only do circulating levels of various endocrine factors oscillate over the 24 period, but so too does responsiveness of target tissues to these signals/stimuli. Emerging evidence suggests that these daily oscillations do not occur solely in response to behavioral fluctuations associated with sleep/wake and feeding/fasting cycles, but are orchestrated in part by an intrinsic timekeeping mechanism known as the circadian clock. Disruption of circadian clocks, through genetic and/or environmental means, appears to precipitate numerous common disorders, including cardiometabolic diseases and cancer. Collectively, these observations, which are reviewed within the current article, have led to suggestion that strategies designed to realign normal circadian rhythmicities hold a therapeutic potential for the treatment of various endocrine-related disorders.

INTRODUCTION

Consideration of the temporal relationship between processes/events is critical for our understanding of the molecular-basis of physiology, as well as the pathogenesis of disease. It is imperative that biological processes occur in an appropriate order, thereby preventing concurrent activation of potentially incompatible mechanisms. Should this level of control become impaired, the outcome can be catastrophic, often resulting in pathology. Temporal regulation is observed at multiple time scales, ranging from seconds (*e.g.*, ion homeostasis) to years (*e.g.*, development). One time scale that has received increasing attention over the

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past several decades is the circadian (*i.e.*, 24 hour) period. Rejuvenated enthusiasm in chronobiology has been fueled, in part, by identification of the molecular underpinnings of the mammalian circadian timekeeping mechanism, revealing that this mechanism orchestrates a plethora of critical biological functions, including those precipitating death.

Endocrine factors are undoubtedly at the heart of circadian biology. Not only is the generation, secretion, and abundance of various endocrine factors subjected to stringent and often predictable time-of-day-dependent control, but so too is the sensitivity of target organs to these signals. Emerging evidence strongly suggests that these rhythms are driven not only by behavior-associated factors, but also by an intrinsic timekeeping mechanism known as the circadian clock. The purpose of this article is to review the importance of circadian clock control of endocrinology. Following a general overview of known endocrine factor oscillation characteristics, we will discuss evidence regarding mediation of endocrine rhythms by circadian clocks, the pathologies associated with disruption of normal rhythmicity, and the tactics employed to restore orchestration of circadian processes.

CIRCADIAN RHYTHMS IN STIMULUS-RESPONSE COUPLING

Stimulus-response coupling overview

By definition, hormones and endocrine factors are substances produced in one organ or cell type that act at a site that is distinct from the origin. As such, these factors are critical in complex species, such as humans and other vertebrates, for inter-organ communication (*e.g.*, between the central nervous system and peripheral tissues). Such communication is essential for maintenance of homeostasis and adaptation to environmental changes or stresses. A substantial body of evidence suggests that various components of the endocrine system exhibit time-of-day-dependent rhythms, facilitating anticipation of and/or adaptation to environmental/behavioral fluctuations. In principle, cellular hormone action is governed by: 1) the concentration of the hormone relative to its affinity for its specific receptor; 2) the abundance and/or availability of the receptor; and 3) the activity status of post-receptor signaling components, including mechanisms of desensitization/downregulation of hormone signaling. Time-of-day has been shown to exert a modulatory effect on all of these governing mechanisms. Here, we will provide a brief summary of what is known regarding 24-hour rhythms in the level of the stimulus (*i.e.*, endocrine factors) and target tissue responsiveness.

Daily oscillations in endocrine factors

Among the best-studied examples of endocrine factors that fluctuate over the course of the day are those whose production is governed by hypothalamic-pituitary axes, such as cortisol, growth hormone (GH), prolactin (PRL), thyroid hormone, and gonadal steroids.^{1–5} Other factors, such as nutrient-sensitive hormones (such as insulin and adipokines) vary their circulating levels in part in response to the environment/behaviors, such as light/dark and feeding/fasting cycles, that typically occur in time-of-day-dependent patterns.^{6, 7} Figure 1 and Table 1 summarize what is known regarding time-of-day-dependent oscillations in these and other endocrine factors, while below we provide additional detail for several of these factors.

Melatonin—Circulating melatonin is produced in the pineal gland, under tight control by the central circadian pacemaker in the suprachiasmatic nucleus (SCN) region within the hypothalamus.⁸ A primary function of melatonin is to relay information regarding the environmental light-dark cycle, especially in response to changes in day length, triggering endocrine changes.⁹ Melatonin production is characteristically low in the presence of light and increases during the night, during which time it induces and supports sleep.¹⁰ It is important to note that additional functions of this hormone have been reported, including vasoconstrictor/vasodilator, anticonvulsant, and antioxidant properties.¹¹ Interestingly, melatonin deficiency has been associated with increased incidence of colorectal and breast cancer, while polymorphisms in melatonin receptors have been associated with diabetes risk.^{10, 12} Melatonin supplementation has been used in the clinical setting, for the treatment of winter depression, numerous sleep disorders, and as an adjuvant therapy for epilepsy.^{13, 14}

Cortisol and Hypothalamo-pituitary-adrenal axis—Release of cortisol from the adrenal cortex is under the regulation of the hypothalamo-pituitary-adrenal (HPA) axis.¹⁵ Corticotropin releasing hormone (CRH) is a component of this axis that is secreted by the hypothalamus and conveyed via the hypophyseal portal vasculature to the anterior pituitary gland. There, CRH triggers release of adrenocorticotropic hormone (ACTH) into the general circulation; upon binding to receptors in the adrenal gland, ACTH stimulates cortisol release. Cortisol levels exhibit a robust time-of-day-dependent rhythm in humans, normally peaking during the morning (0700–0800).¹⁶ In doing so, this catabolic hormone is believed to prepare the body for typical stresses associated with waking (e.g., energetic demand); the steady rise in cortisol levels during the sleep phase is consistent with its role in anticipation of wakefulness and increased activity.^{15, 16}

Growth Hormone—Growth hormone (GH) is produced in the anterior pituitary and secreted in response to the integrated stimulatory and inhibitory effects of GH releasing hormone (GHRH) and somatostatin (SMS), respectively, both of which emanate from the hypothalamus.^{17, 18} GH has specific and powerful effects in metabolism, typically in opposition to insulin (*e.g.*, decreasing glucose utilization, increasing lipolysis).¹⁹ Circulating GH levels exhibit both circadian and ultradian variation, as well as sexual dimorphism. In human females, GH is secreted in frequent non-discrete peaks of more uniform amplitude throughout the day with an increased mean secretion at night (during sleep). Males secrete GH in fewer more discrete pulses throughout the day, again with increased peak amplitude at night.^{1, 20} This dimorphism is also observed in children, with peaks in each case being greater than in adults.^{21, 22} Although the exact purpose of increased growth hormone availability at night is unknown, possibilities may include promotion of insulin resistance, mobilization of triglyceride, and/or protein synthesis/repair at this time.

Adiponectin—Adiponectin is one of several adipokines (*i.e.*, adipose-derived endocrine factors) that exhibits a time-of-day-dependent rhythm in humans, peaking in the circulation between 1200 and 1400.^{7, 23} Adiponectin is best known as an insulin sensitizer, whose circulating levels vary inversely with body mass index (BMI).²⁴ Interestingly, hypoadiponectinemia is associated with metabolic syndrome, but conversely, elevated levels

are seen in chronic heart failure and chronic renal failure.²⁴ The purpose of increased adiponectin secretion during the early afternoon is unclear, but this rhythm may provide anticipation of increased insulin levels later in the day (thereby promoting insulin action).

Insulin—Insulin is produced by beta cells of pancreatic islets, and is secreted in response to increased levels of circulating nutrients, particularly glucose.²⁵ Its primary function is at the level of metabolism, stimulating glucose utilization and protein synthesis by peripheral tissues, such as liver, skeletal muscle, and fat.²⁶ This anabolic hormone also inhibits fatty acid beta oxidation, and channels fatty acids into triglyceride.²⁶ Insulin secretion displays a diurnal rhythm in humans, peaking at roughly 1700h (with a nadir at 0400h).⁶ Such a rhythm is consistent with nutrient storage during the awake/fed state, and subsequent mobilization during the sleep/fasted period.

Daily Oscillations in Sensitivity to Endocrine Factors

In contrast to information concerning circulating levels of hormones and endocrine factors, our understanding of the impact of time-of-day on end-organ hormonal sensitivity is less well developed. Standard techniques of evaluation make this variable more difficult to study, especially in humans. However, a few examples exist in which time-of-daydependent rhythms in hormone sensitivity have been characterized, including responsiveness to ACTH and insulin. As mentioned in the previous sub-section, cortisol exhibits a robust diurnal variation in humans. However, this rhythm does not appear to be driven by fluctuations in ACTH levels; in contrast, the sensitivity of the adrenal cortex to ACTH exhibits a time-of-day-dependence, thereby mediating cortisol oscillations.¹⁵ Glucose homeostasis also exhibits interesting time-of-day-dependent changes, particularly in humans. The "dawn phenomenon", for example, occurs in normal individuals and is exaggerated in type 1 and type 2 diabetes.^{27–31} This phenomenon is characterized by a spontaneous increase in plasma glucose and/or insulin requirement without nutrient intake prior to awakening (typically roughly 5-8 AM). This is believed to arise from increased hepatic glucose production that prompts (or requires) increased insulin levels to ensure glucose disposal and restrain hepatic glucose production. While the rise in hepatic glucose production corresponds temporally to rising cortisol, epinephrine, and norepinephrine (all counter-regulatory hormones), experiments in humans indicate that the dawn phenomenon is not accounted for by increases in these hormones, but rather to antecedent surges of GH earlier in the nocturnal period.^{27, 32}

MEDIATORS OF CIRCADIAN RHYTHMS IN ENDOCRINE FACTORS

Role of intrinsic versus extrinsic influences

Daily rhythms in the abundance of many endocrine factors (as well as in the sensitivity of distinct tissues to these factors) are potentially mediated by a combination of influences. A classic view has often been that fluctuations in nutrients, hemodynamic stresses, body temperature, metabolism, sympathetic and autonomic tone, as well as various autocrine/ paracrine factors are mediated by behavioral changes across the sleep/wake and feeding/ fasting cycles.^{33–35} However, this 'extrinsic' concept has been challenged recently, as a wealth of evidence from both human and animal studies are consistent with a significant

contribution of an intrinsic timekeeping mechanism towards these oscillations. For example, elegant studies by Van Cauter & colleagues were among the first to assess the contribution of sleep on oscillations a number of endocrine factors in humans, by enforcing a state of arousal during the night, followed by sleep during the subsequent day. These studies revealed that some (*e.g.*, cortisol, TSH), but not all (*e.g.*, growth hormone, prolactin), oscillations in endocrine factors occur independently of sleep/wake cycles.^{36–38} This raised the possibility that fluctuations in distinct endocrine factors over the course of the day may be driven, at least in part, by an endogenous mechanism (*i.e.*, independent of behavioral rhythms). Similarly, Scheer & Shea have begun to dissociate the relative contribution of intrinsic versus extrinsic influences on biological processes in humans, through enforcement of multiple 20- or 28- hour contiguous days; these studies have successfully uncovered the contribution of the intrinsic circadian system on daily rhythms of multiple endocrine factors, including epinephrine and PAI-1.^{39–42}

The endogenous circadian system in mammals is the circadian clock.^{43, 44} Circadian clocks are cell autonomous molecular mechanisms that enable a cell to perceive the time of day. In doing so, molecular clocks are able to alter biological processes over the course of the day, independent of extracellular stimuli/stresses. Circadian clocks are classically described as being transcriptional in nature, composed of a collection of transcriptional modulators that interact to form positive and negative regulatory feedback loops.^{43, 44} At the heart of the mechanism are two transcription factors, CLOCK (circadian locomotor output cycles kaput) and BMAL1 (brain and muscle ARNT-like 1), that, upon heterodimerization, bind to Eboxes within promoters of multiple target genes.^{45, 46} The translation products of a number of these target genes are themselves core clock components, which feedback upon CLOCK/ BMAL1 activity, including the period (PER1/2/3) and cryptochrome (CRY1/2) proteins.^{47, 48} An additional well-established negative feedback loop involves REV-ERBa, which negatively influences BMAL1 at the transcriptional level.⁴⁹ Those genes that are regulated by the circadian clock, yet do not directly influence the activity of core clock components, are termed clock controlled/output genes. It has been estimated that 7% to 13% of a cell's transcriptome is under circadian control, including genes encoding for modulators of transcription, signal transduction, protein turnover, and metabolism; through fluctuations in clock controlled genes, the circadian clock influences cellular/organ function in a time-ofday-dependent manner.^{50–52} It should be noted that although the circadian clock is often described as a transcriptionally-based mechanism, a host of post-translational modifications (PTMs) are essential for its normal function, including phosphorylation, ubiquitination, sumovlation, acetylation, ADP-ribosylation, and O-GlcNAcylation.^{53–58} Many of these PTMs not only facilitate exquisite sensitivity of the clock mechanism to numerous extracellular factors (ensuring rapid entrainment to alterations in the environment), but also serve as an additional means by which the clock influences cellular processes. Interestingly, Reddy and colleagues have recently described a highly conserved circadian mechanism that is independent of transcription; redox biology is at the core of this ancestral clock mechanism, which can be monitored through oscillations in the oxidative state of peroxiredoxin.59

Influence of the SCN versus non-SCN clocks

Circadian clocks can be broadly divided into two major classes: the central clock within the SCN and peripheral (or non-SCN) clocks.⁴³ Through a classic series of lesion and transplant studies, the central circadian pacemaker (localized to the SCN of the hypothalamus) has been established as the orchestrator of the intrinsic circadian system in mammals. Specifically, lesions of the SCN result in severe arrhythmicity of multiple biological functions, including sleep/wake behavior, drinking/food intake, oxygen consumption, and stool output, as well as adrenal, pineal, and pituitary hormone release.^{60–62} Transplantation of fetal SCN tissue into the third ventricle of lesioned rodents re-instates locomotor rhythms with the rate of the donor clock tissue.^{63–65} Even when SCN grafts are encapsulated, behavioral rhythmicity is restored, suggesting that the SCN output signal involves, at least in part, a paracrine/endocrine signal.⁶⁴ One releasable factor thought to be involved in signaling rhythmic locomotor behavior is prokineticin 2.⁶⁶ Because transplantation does not restore rhythmic release of glucocorticoids, melatonin or luteinizing hormone, it is clear that many endocrine rhythms require a neural connection.

Advances over the last few decades have demonstrated that the circadian network is in reality a multiple oscillator system, in which different tissues/cells are capable of exhibiting self-sustained oscillations in isolation (although oscillations tend to dampen after several days for non-SCN tissue cultures).^{67–70} This realization has raised the possibility that circadian rhythms in various biological processes may be driven in part by local tissue oscillators. While melatonin rhythmicity is undoubtedly dependent upon central SCN clock control⁷¹, local clock control is evident in other peripheral endocrine tissues, such as the adrenal gland and pancreas (Figure 2). For example, transplantation of Per2/Cry1 mutant adrenal glands into wild-type animals results in a dampened rhythm of corticosterone release, despite normal ACTH levels.⁷² Similarly, adrenal-specific Bmal1 knockdown produces arrhythmic plasma corticosterone secretion.⁷³ Taken together, these data suggest that the adrenal molecular clock locally regulates ACTH sensitivity (although it should be noted that the latter is also dependent on SCN signals transmitted by sympathetic fibers).¹⁵ Evidence also supports the concept that the pancreatic β -cell clock influences insulin secretion. Allaman-Pillet *et al* have previously reported that synchronized β -cells in culture secrete insulin in a rhythmic fashion, with a periodicity of 24-hours.⁷⁴ More recently, β -cell specific Bmal1 null mice were shown to display impaired insulin secretion.^{90, 91} In summary, these results suggest that the local molecular clock plays a critical role in many endocrine tissues, and that disruption of this mechanism results in impaired function.

PATHOLOGIC CONSEQUENCES OF CIRCADIAN DISRUPTION

Lessons learnt from animal models of impaired circadian biology

As highlighted above, and in Table 1, it is clear that a large number of endocrine factors exhibit time-of-day-dependent oscillations in both humans and animal models. Furthermore, cell autonomous circadian clocks appear to contribute towards these rhythms to varying degrees. Observations such as these raise a host of critical questions, for which answers are not clear. For example, what is the physiologic purpose of daily oscillations in endocrine factors, and what is the pathologic consequence of disruption of these oscillations?

Impairment of normal circadian rhythmicity negatively impacts a host of biological processes in both humans and animal models.^{43, 75–77} Circadian disruption may occur secondarily to changes in the environment (*e.g.*, fluctuations in the light/dark, feeding/ fasting, sleep/wake cycles) and/or genetic abnormalities. The estrous cycle and conception success serves to illustrate this concept. Shift workers report decreased successful attempted conceptions during periods of night shift work, while in mouse models, modulation of the light/dark cycle, disruption of SCN and the pre-optic communication, or genetic manipulation of either CLOCK or BMAL1 all impede estrous cycling and successful pregnancies.^{78–82} The current sub-section will summarize key findings from experimental models with regards to the impact of disrupted circadian rhythms on endocrine factors and endocrine-regulated processes (particularly metabolism).

Manipulation of the light/dark cycle has proven to be a useful tool for the study of disrupted circadian rhythms on many biological processes, including endocrine- and metabolismrelated parameters. Following an abrupt alteration in the light/dark cycle, intrinsic circadian clocks reset at varying rates, in an attempt to resynchronize to the new environment. The SCN resets relatively rapidly to a large shift in the light-dark cycle (at rate of ~1 day per hour of shift in the LD cycle) compared to peripheral organs which invariably reset at a slower rate, lagging behind the SCN in a tissue-specific manner.^{83–86} Accordingly, dyssynchrony occurs at both the organism-to-environment and inter-organ levels, until all clocks (and clock-regulated processes) reset fully to the new light/dark cycle. Should the light/dark cycle be altered multiple times in succession, with an interval shorter than that required for complete resynchronization (e.g., jet lag protocol, wherein the light/dark cycle is phase advanced 6 hours either once or twice a week), chronic dyssynchrony ensues. Prolonged periods of dyssynchrony are associated with altered immune function, increased tumor growth, reduced body temperature, and increased adiposity.⁸⁷⁻⁹⁰ Additional models of light/dark cycle manipulation induced circadian dyssynchrony include housing rodents in constant light or shortened daily cycles (e.g., 20 hour days).⁹¹ Invariably these manipulations negatively impact endocrine-regulated processes, including metabolic homeostasis.92-97

The unveiling of core components of the mammalian circadian clock mechanism over the past two decades has been followed closely by their genetic manipulation and characterization of the phenotypic consequence in rodents. One of the first clock components to be genetically modified was CLOCK; chemical mutagenesis resulted in a truncated dominant negative mutant CLOCK protein lacking the transactivation domain.⁹⁸ Interestingly, although Clock mutant mice exhibit normal activity rhythms under light/dark cycles, feeding behavior appears to be altered, which is associated with increased adiposity and perturbations in endocrine factors (*e.g.*, insulin levels).⁹⁹ Both germline and tissue-restricted knockout mice have also been generated for distinct circadian clock components. Of these models, germline BMAL1 knockout mice exhibit one of the most striking phenotypes, being arrhythmic for most parameters under circadian conditions, developing markers of accelerated aging (*e.g.*, sarcopenia, cataracts, lipoatrophy), concomitant with reduced lifespan.¹⁰⁰ In terms of endocrine factors, Kennaway and colleagues have recently reported that circulating levels of insulin are lower, while adiponectin and leptin are higher,

in Bmal1 null mice (relative to wild-type mice).¹⁰¹ Indeed, it has been suggested that BMAL1 null mice should be considered a model of diabetes, due in large part to impaired insulin secretion.^{102, 103} Genetic manipulation of circadian clock components invariably alters insulin sensitivity, in a manner that is dependent upon whether a positive or negative clock component is disrupted (leads to increased and decreased insulin sensitivity, respectively).^{104, 105}

Endocrine and metabolic circadian abnormalities in humans

Misalignment of peripheral and central clocks with behavioral sleep-wake rhythms can be experimentally induced in the laboratory setting by use of a very short (20 h; 7 h sleep and 13 h wake) or long (28 h; 8 h sleep and 20 h wake) day that is outside the range of entrainment for a 24-h clock.¹⁰⁶ For example, three cycles of a 28-h enforced sleep-wake schedule results in sleep-wake behavior that is ~180° out of phase with the intrinsic circadian clock and behavior. After three additional cycles, behavior and the circadian clock are back in alignment. Thus, comparison of cycles with full alignment versus misalignment provide the opportunity to determine which endocrine rhythms follow sleep/wake/eating behavior, and which rhythms are more directly coupled to the intrinsic circadian clock. Using this protocol, Scheer et al found that glucose, epinephrine, and cortisol faithfully follow a ~24-h rhythm during behavioral misalignment, while leptin, insulin, and norephinephrine did not (although it is important to note that all of these endocrine rhythms were influenced to varying degrees by the behavioral cycle).¹⁰⁷ Overall leptin levels were lower during misaligned conditions, similar to when circadian misalignment is induced by a more subtle approach using a 24.6-h day. Volunteers in this condition for 25 days exhibited reduced leptin levels at physiologically significant levels,¹⁰⁸ which is likely to have a metabolic outcome. In fact, after just 3 cycles of laboratory-induced circadian misalignment (*i.e.*, a 28-h day), healthy volunteers show a dramatic loss of the number of rhythmic transcripts in the blood.¹⁰⁹ Importantly, this same protocol also causes glucose intolerance and insulin resistance, suggesting that circadian misalignment may cause insufficient β -cell compensation.¹⁰⁷ More recently, Leproult and colleagues have reported that then circadian misalignment is imposed on sleep deprived subjects, indices of insulin resistance worsen.¹¹⁰ Taken together, these studies are consistent with the concept that circadian misalignment impairs endocrine homeostasis, inducing a cardiometabolic state.

Aberrant metabolic homeostasis associated with experimental circadian misalignment is reminiscent of shift-work induced metabolic perturbations. It is clear that misalignment of melatonin and cortisol rhythms occur during shift work.^{111–113} In addition, shift workers are more likely to develop cardiometabolic syndrome, ¹¹⁴ potentially due to increased postprandial levels of insulin, glucose, and triacylglycerol during the night shift, ¹¹⁵ as well as increased energy intake and circulating triglycerides, as well as reduced insulin sensitivity and post-prandial ghrelin release.¹¹⁶ Shift workers also have an increased risk of developing cancer, gastrointestinal disorders, and cardiovascular diseases such as ischemic heart disease.^{107, 114, 117–120} Taken together, these observations underscore the importance of understanding synchronization of the endogenous circadian system with environmental/ behavioral and endocrine factors.

In addition to circadian misalignment induced by environmental/lifestyle influences, evidence suggests that desynchrony among intrinsic oscillators occur during pathologic states, particularly those associated with endocrine dysfunction. For example, clock gene rhythms in white blood cells are dampened in patients with diabetes.¹²¹ Interestingly, patients with type 2 diabetes do not show a circadian rhythm in insulin secretion, although rhythmicity in insulin sensitivity appears to remain intact.¹²² Even first-degree relatives of patients with Type 2 diabetes show dampened and shortened rhythms in insulin secretion rates.¹²³ These clinical observations are supported by findings in animal models of metabolic dysfunction (Zucker rats, diabetic *db/db* mice, and leptin knockout mice), revealing that circadian clock and metabolic perturbations occur in parallel.^{124–128} Recently, it has been reported that short term high fat diet disrupts alignment of circadian clock gene rhythms in multiple tissues prior to metabolic dysfunction, suggesting that circadian misalignment may precede (and potentially cause) metabolic and endocrine abnormalities.^{129, 130}

FEASIBILITY OF NORMALIZING CIRCADIAN RHYTHMICITY

Lessons learnt from clock gene rescue studies in animal models

As mentioned above, disruption of the molecular clock in animal models often results in endocrine disruption and cardiometabolic impairment. Genetic rescue of clock-controlled gene transcription rhythms and associated metabolic/endocrine phenotypes may not only serve as proof-of-principal, but may shed light on feasibility of targeting this system for future treatment of disease states. For example, arrhythmic activity and metabolism as well as body weight and adiposity abnormalities observed in Bmall knockout mice can be rescued by germline over-expression of *Bmal2*.¹³¹ In addition, the low body weight and early death phenotypes of Bmall knockout mice can be rescued by muscle-specific, but not brain-specific, over expression of Bmal1.132 Rhythmicity of a number of clock controlled genes is also restored by these genetic rescue strategies. It should be noted that a sub-set of transcriptional rhythms normally display 12-h periodicity in peripheral tissues such as the liver, heart, kidney, and lungs from wild type mice.¹³³ However, these 12-h rhythms become 24-h when CLOCK mutant mice are rescued only in the brain (using a Tet-OFF expression system).¹³⁴ This finding is interesting in light of the fact that time-of-day-restricted feeding (during the inactive period, between 1 to 9 hours after lights on) in wild-type mice also converts these shorter, ultradian rhythms to 24-h rhythms.¹³³ Taken together, these studies suggest that circadian desynchrony can be rescued, and that the central clock interacts with environmental/systemic signals to produce ultradian rhythms.

The utility of time-of-day-restricted feeding

Evidence to date undoubtedly suggests that the intrinsic circadian system plays an important role in daily rhythms of a host of endocrine factors. In several cases (*e.g.*, cortisol, insulin, *etc*), non-SCN clocks appear to contribute in a significant manner. This raises the question whether strategies designed to selectively entrain peripheral circadian clocks would be beneficial for resynchronizing oscillations in endocrine factors during disease states (such as cardiometabolic disease states). Damiola & colleagues were among the first to report that restricting food access in rodents to a distinct period during the sleep phase causes a phase

shift in circadian clocks within peripheral tissues (e.g., liver, kidney, pancreas, heart), while the SCN clock remains entrained to the light/dark cycle (*i.e.*, relatively unresponsive to feeding entrainment).¹³⁵ The strength of this intervention is exemplified by the demonstration that circadian clock gene oscillations are re-initiated in peripheral tissues of CLOCK mutant mice by time-of-day-restricted feeding.¹³⁶ These initial studies led to various investigations into the negative impact of restricted feeding-induced circadian dyssynchrony on energy homeostasis. A consensus of these studies appears to be that consumption of an inappropriately large number of calories during the end of the active period and/or during the sleep phase, is associated with metabolic dyssynchrony and cardiometabolic disease development.^{93, 137–141} Importantly, evidence exists that observations in rodent models translate to humans. For example, night eating syndrome is associated with obesity.¹⁴² In addition, Qin et al have reported consumption of a large meal at night leads to an uncoupling of the relationship between plasma glucose and insulin levels, indicative of metabolic dysfunction.¹⁴³ These findings have led several investigators to hypothesize that restriction of food intake to an appropriate time of the day may prove beneficial, via resetting peripheral clocks and reestablishment of circadian synchrony during cardiometabolic disease states. In support of such a concept, Kudo et al reported that restricting food access to the active phase normalized liver clock function in db/db mice, and concomitantly improved various cardiometabolic parameters (including lowering circulating insulin levels).¹⁴⁴ Similarly, active phase-restricted feeding is able to normalize peripheral circadian clock gene oscillations concomitant with metabolic and endocrine abnormalities in both a rodent model of shift work, as well as during chronic high fat feeding.^{137, 145, 146} Clearly additional studies are required to elucidate further the potential utility of time-ofday-restricted feeding on normalization of circadian dyssynchrony and metabolic/endocrine factors.

The therapeutic potential of drugs that can alter circadian clock function

Treatment strategies involving genetic manipulation or alterations in a patient's daily behavior are clearly not ideal, for a multitude of reasons. A more promising therapeutic approach would be to utilize a pharmaceutical agent that selectively targets the molecular clock.¹⁴⁷ For example, one recently described small molecule compound is longdaysin, which has been shown to lengthen the circadian period by targeting multiple kinases simultaneously (CKI\delta, CKIa, and ERK2), leading to PER1 degradation.¹⁴⁸ Stabilization of CRY proteins (through prevention of ubiquitination and degradation) by another small molecule (KL001) also lengthens the period of the clock. Interestingly, KL001 is an attractive therapeutic prospect for diabetes since it has been shown to suppress hepatic glucose production induced by glucagon.¹⁴⁹ Additional clock components are known to impact metabolism, including REV-ERBa. REV-ERBa agonists (SR9009 and SR9011) not only enhance Per2 rhythms, suppress Cry2 rhythms, and shift Bmal1 rhythms, but SR9009 also rescues increased fat, dyslipidaemia, and hyperglycaemia in obesity models.¹⁵⁰ Targets of kinases such as glycogen synthase kinase 3 (GSK3) and c-Jun N-terminal kinases inhibitors (CHIR99021 and SP600125, respectively) are also effective at changing the period of molecular clock.^{151, 152} It is predicted that when coupled with time-of-day-specific delivery, many of these small molecule inhibitors may have therapeutic potential, through

restoration of circadian synchrony of biologic processes, including secretion of, and sensitivity to, endocrine factors.

CONCLUSIONS

It has long been appreciated that circulating levels of various endocrine factors exhibit a time-of-day dependence. It is clear that these rhythms are not solely the result of environmentally/behaviorally derived influences, but are also largely mediated by the endogenous circadian system. With regards to the latter, it is becoming increasingly apparent that cell autonomous circadian clocks modulate endocrine factor levels, as well as tissue responsiveness to these stimuli, over the course of the day. Disrupted orchestration of these rhythms occurs readily through environmental and/or genetic means, resulting in a state of desynchrony known to promote a host of pathologies, including cardiometabolic diseases and cancer. Initial studies designed to reinstate synchrony through environmental (*e.g.*, time-restricted feeding) and/or pharmacological means show promise for attenuated pathogenesis in animal models. With the recent advent of small molecule modulators of the circadian clock mechanism, the prospect of therapeutically targeting this mechanism in patients with endocrine disorders holds great potential.

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DECLARATION OF INTEREST

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Biographies

Karen L. Gamble, PhD is an Assistant Professor at UAB, who has studied circadian rhythms in hamsters, transgenic mice, and humans over the last 15 years. Her research includes investigation of the molecular clock gene regulation and neurophysiology of the SCN as well as circadian rhythm and sleep studies in humans, including sleep behavior strategies of shift workers, circadian dysfunction in adult ADHD patients with delayed sleep phase, and daytime sleepiness in adolescents. She holds memberships with the Society for Research in Biological Rhythms, Society for Neuroscience, Working Time Society, and the Society for Light Treatment and Biological Rhythms.

Ryan D. Berry, BS is currently in his third year of the National Institutes of Health Medical Scientist Training Program at the University of Alabama at Birmingham. Presently he is pursuing a PhD studying Growth Hormone Receptor signaling in the lab of Dr. Stuart J. Frank. He received his BS degree in Biochemistry and Spanish from Utah State University in 2011, during which time he presented posters at the Utah Conference on Undergraduate Research (2009) on the effects of mutations on the kinetics of PTP1B and YopH, and at the National Conference on Undergraduate Research (2011) on the promiscuous catalytic activity of PTP1B.

Stuart J. Frank, MD is Director of the Division of Endocrinology, Diabetes, and Metabolism in the Department of Medicine at the University of Alabama at Birmingham and Chief of the Endocrinology Section of the Birmingham VAMC Medical Service. Dr. Frank is also Professor of Medicine and Cell, Developmental, and Integrative Biology at UAB. He earned the M.D. at Harvard Medical School and was an Internal Medicine resident at Washington University, prior to Endocrinology fellowship and research training at the National Institutes of Health. Dr. Frank's laboratory studies molecular mechanisms of growth hormone and prolactin action, determinants of GH and prolactin sensitivity, and the roles of these hormones in health and disease.

Martin E. Young, DPhil is an Associate Professor of Medicine in the Division of Cardiovascular Diseases at the University of Alabama at Birmingham. Dr. Young earned his PhD at the Oxford University (Biochemistry) before completing post-doctoral fellowships at Boston University (Diabetes/Metabolism) and the University of Texas at Houston (Cardiac Metabolism). Dr. Young's current research interests include understanding the role of cell autonomous circadian clocks in metabolic regulation, and how desynchrony of these molecular mechanisms contributes towards cardiometabolic diseases.

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Box 1. Definitions of commonly utilized terms in chronobiology

Tau (period)

The length of time for one cycle of an oscillation to repeat. For example, the length of time for a distinct phase point (such as the peak) to re-occur during the subsequent cycle.

Amplitude

The size of an oscillation as measured by half the distance from the peak to the trough.

Phase

The timing of a consistent point in the cycle, such as the peak or trough.

Phase shift

A change in phase such that it occurs earlier or later, due to a displacement of the entire cycle. A stable phase shift will only produce a period change for one or a few "transient" cycle(s), after which the oscillation will continue with the same period as before the shift.

Zeitgeber

Literally means "time giver," and refers to any resetting stimulus that serves as a time cue for the external environment. Light and food intake are classic examples of zeitgebers for the SCN and non-SCN clocks, respectively.

Circadian rhythm

"Circadian" literally means "about a day" and refers to a rhythm whose complete cycle repeats close to every 24 hours. In order to demonstrate that a circadian rhythm is endogenously generated, the rhythm must be measured under constant conditions (*i.e.*, in the absence of a zeitgeber).

Diurnal rhythm

A physiological measurement (*e.g.*, gene transcription, protein expression, body temperature, behavior, *etc*) that exhibits daily changes across the 24-h light dark cycle such that the peak(s) and nadir occur ~12-h apart or in a frequency that is a harmonic of 24 hours. In conditions in which a zeitgeber is present (*e.g.*, a light-dark cycle), it is impossible to tell if a diurnal rhythm is endogenously generated or an acute response to the zeitgeber.

Entrainment

A stable state in which an external zeitgeber (e.g., light) induces small shifts in the phase of an endogenous oscillator on a daily basis so that the period of the physiological output of the oscillator (e.g., locomotor activity) is equal to the period of the external zeitgeber cycle with a specific phase relationship to one another.

Ultradian

A repeating rhythm that has a cycle period that is much less than 24 hours (*e.g.*, 12-h gene expression rhythm, 1-h pulsatile GnRH release, *etc.*).

REVIEW CRITERIA

PubMed and Google Scholar were utilized to search for original research and review articles published up to 2014; no abstracts from meeting reports have been cited. Search terms used included "circadian clock", "diurnal", "time of day", "endocrine factors", "insulin", "growth hormone", "cortisol", "metabolism" and "desynchrony". The references of the retrieved articles were also reviewed.

KEY POINTS

1. Many endocrine factors oscillate in a time-of-day-dependent manner.

- 2. Endocrine factor rhythms are driven by in part by circadian clocks.
- 3. Circadian desynchrony is associated with pathologic states.
- 4. Re-instatement of circadian rhythms improves metabolic homeostasis.

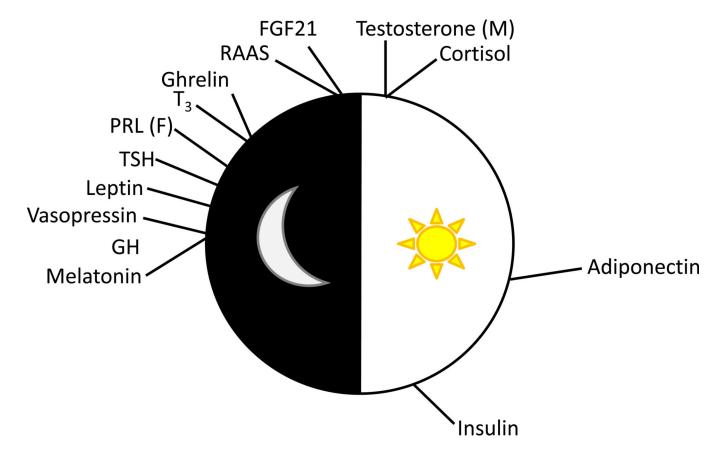
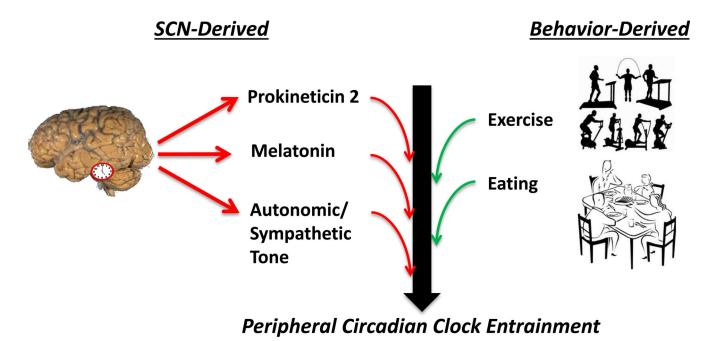


Figure 1.

Time-of-day at which circulating levels of key endocrine factors peak in humans. Abbreviations utilized include: GH, growth hormone; TSH, thyroid stimulating hormone; PRL, prolactin; T₃, triiodothyronine; RAAS, renin-angiotensin-aldosterone system; FGF21, fibroblast growth factor 21; (F), females only; (M), males only.

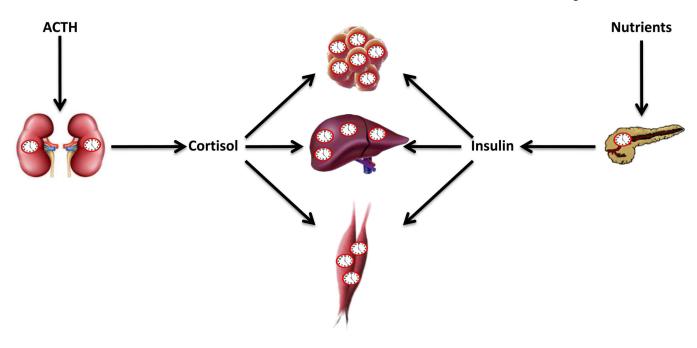
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Figure 2.

SCN and non-SCN derived entrainment of peripheral circadian clocks (A) and the impact of cell autonomous circadian clocks on endocrine factor release and sensitivity (B). As discussed in the text, peripheral circadian clocks are entrained (re-set) by both SCN- (*e.g.*, prokineticin 2, neural stimulation) and behavior- (*e.g.*, feeding, physical activity) dependent influences (A). Circadian clocks within the adrenal cortex contribute to diurnal variations in cortisol release, through modulating sensitivity to ACTH, while the β -cell circadian clock sithin target tissues (*e.g.*, adipose, liver, skeletal muscle) potentially modulate sensitivity to endocrine factors in a time-of-day-dependent manner (B).

Table 1

Endocrine factors known to oscillate with a periodicity of 24 hours in humans.

Hormone	Time-of-day- dependent Rhythm	Time of Peak	Citation
Cortisol	Yes	0700–0800h	Kalsbeek et al ¹⁶ ; Carroll et al ³
Growth Hormone	Yes	Pulsatile secretion, with Increased amplitude at night	Avram <i>et al</i> ¹ ; Jaffe <i>et al</i> ²⁰ ; Villadolid <i>et al</i> ²¹ ; Goji <i>et al</i> ²²
Follicle-stimulating hormone (females)	No	NA	Klingman <i>et al</i> ¹⁵³
Testosterone (males)	Yes	0700h	Walton <i>et al</i> ⁵
Prolactin	Yes	0200h (amplitude larger in females)	Freeman et al ⁴
Thyroid stimulating hormone	Yes	0100–0200h	Russell <i>et al</i> ²
Triiodothyronine	Yes	0230-0330h	Russell et al ²
Thyroxine	No	NA	Russell <i>et al</i> ²
Renin-angiotensin-aldosterone system	Yes	Early morning	White <i>et al</i> ¹⁵⁴
Fibroblast growth factor 21	Yes	0500–0800h	Lee <i>et al</i> ¹⁵⁵ ; Yu <i>et al</i> ¹⁵⁶
Ghrelin	Yes	0200–0430h (fed state) 1300h (fasted state)	Koutkia <i>et al</i> ¹⁵⁷ ; Natalucci <i>et al</i> ¹⁵⁸
Adiponectin	Yes	1200–1400h	Scheer <i>et al</i> ⁷ ; Gavrila <i>et al</i> ²³
Leptin	Yes	0100h	Gavrila <i>et al</i> ²³
Vasopressin	Yes	Middle of night	Forsling et al ¹⁵⁹
Insulin	Yes	1700h	Goel <i>et al</i> ⁶
Melatonin	Yes	Middle of night	Van Cauter et al ³⁶