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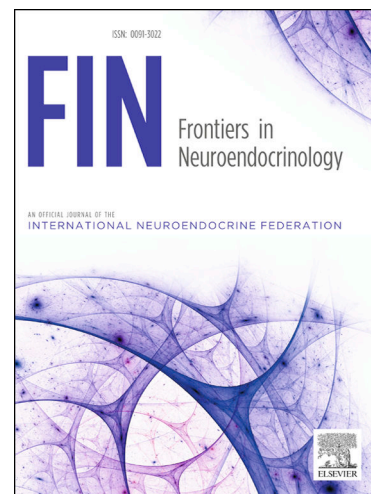
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Revision 3

**Timing is everything: circadian rhythms and their role
in the control of sleep**

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Abstract

Sleep and the circadian clock are intertwined and have persisted throughout history. The suprachiasmatic nucleus (SCN) orchestrates sleep by controlling circadian (Process C) and homeostatic (Process S) activities. As a "hand" on the endogenous circadian clock, melatonin is critical for sleep regulation. Light serves as a cue for sleep/wake control by activating retino-recipient cells in the SCN and subsequently suppressing melatonin. Clock genes are the molecular timekeepers that keep the 24 h cycle in place. Two main sleep and behavioural disorder diagnostic manuals have now officially recognised the importance of these processes for human health and well-being. The body's ability to respond to daily demands with the least amount of effort is maximised by carefully timing and integrating all components of sleep and waking. In the brain, the organization of timing is essential for optimal brain physiology.

Key words:

Circadian clock, Clock genes, Sleep, Sleep medicine, Sleep disorder

Introduction

The earth's daily rotational cycle has a pervasive influence on all living beings. It has guided biological events and all related processes of adaptation. Historically, the interactions of this daily cycle with organismal physiology has always been assumed but was not investigated scientifically until modern times. In the 1960s, Aschoff and his colleagues carried out a pioneering study of how humans respond to living in an underground bunker, i.e., artificially isolated from exposure to the earth's repetitive day and night cycles. The investigators found that, despite being restricted from experiencing normal environmental influences, the subjects' bodily rhythms of temperature variation and maintenance, urine excretion, and other behavioral activities persisted with a near 24 h periodicity. The finding that this periodicity diverged only slightly from the 24 h pattern observed in humans living under the influence of a regular light/dark (LD) cycle provided proof for the existence of an endogenous clock system (Aschoff, 1965). The term circadian (daily; L. *Circa*=about; *dies*=day; ~24h cycles) rhythm refers to any biological process which exhibits such an endogenous, entrainable oscillation of an approximately, but not exactly, 24 h period (Halberg et al., 1959). This endogenous circadian clock is remarkably widespread in all living organisms, from prokaryotes (e.g. cyanobacteria) to

eukaryotes (e.g. higher plants and animals). These circadian timing processes both regulate and coordinate a wide variety of rhythmic outputs which occur at molecular, physiological, and behavioral levels. These include processes involved in gene expression, regulation of the sleep/wake cycle, hormonal and metabolic regulation, thermoregulation, and a number of others (Cox and Takahashi, 2020; Kim and Lazer, 2020; Riji-Ferreira and Takahashi, 2019; Li et al., 2013; Sukumaran et al., 2010; Waterhouse et al., 2005; Richter, 1967, 1960).

The current review, on the other hand, will seek to provide a broad overview of the relationship between the circadian clock and sleep, which, according to some evaluations, continues to merit closer attention (Borbély et al., 2016; Dijk and Lockley, 2002; Foster, 2020). Some of the more recent findings of the implications of this relationship for physical and mental health will also be reviewed. In the presence of ambient time cues, circadian rhythms can be synchronised; yet, they can also persist even without them (Eban-Rothschild and Bloch, 2012; Honma et al., 2003; Mistlberger and Skene, 2004). The environmental LD cycle acts as the prominent *zeitgeber* (time giver) that resets the circadian clock daily to remain in synchrony with external time cues. However, the persistence of such near 24 h rhythms under prolonged periods of constant

environmental conditions demonstrates the existence of the endogenous circadian clock (Aschoff, 1965). This endogenously generated rhythm is sensitive to and can be changed by environmental geophysical information. This responsiveness helps to maintain a stable phase relationship between the organism and its environment, thereby adjusting those physiological variables as needed for survival (Pontes et al., 2010).

In humans, as in other mammals, these circadian processes are closely affected by a tiny group of melanopsin-expressing retinal ganglion cells (mRGCs) that are highly photosensitive to blue light. Among their other functions, these cells contribute to the mediation of circadian entrainment (Dacey et al., 2005; Esquivá et al., 2017; Hannibal et al., 2017; Nasir-Ahmad et al., 2019; Stachurska and Sarna, 2019; Zele et al., 2011). The retinohypothalamic tract (RHT) is sufficient to maintain the entrainment of circadian clock function (Dai et al., 1998; Klein and Moore, 1979; Sadun et al., 1984). Circadian self-sustaining oscillators are also found in most cells and are made up of a cell-autonomous transcriptional-translational feedback loop system that is decentralized throughout the body. It is, however, the basal hypothalamic suprachiasmatic nucleus (SCN), which is the master circadian clock in mammals, and it is their constituent nuclei which, on a daily basis, orchestrate the

overt rhythms which occur throughout the organism (Dai et al., 1998; Inouye and Kawamura, 1979; Moore and Lenn, 1972; Ralph et al., 1990; Stephan and Zucker, 1972). As a diurnal species, humans normally sleep at night (Lavie, 2001; Webb, 1994). However, there is considerable flexibility in sleep timing in a given individual. This flexibility, in turn, is determined by several factors and ultimately makes up what has been termed a two-process model. Herein, we discuss the current knowledge of sleep-wake timing, its known regulatory disorders, and its multiple implications for brain dysfunction.

Clock genes

At the molecular level, clock genes and their proteins control the rhythmic oscillations for 24 h periodicity. Initially, the transcription factors CLOCK and BMAL1 are encoded from *Clock* and *Bmal1*. These in turn produce heterodimers that permit the transcription of *Per* genes (*Per1*, *Per2* & *Per3*) and two *Cry1* & *Cry2* genes in mammals. Molecular complexes are then formed from the dimerization of PER and CRY proteins. These are then transferred into the nucleus from which they originated and then proceed to prevent the expression of their proteins. Additionally, REV-ERB α and ROR α interact with BMAL1 to repress and activate transcription, respectively. While it has not been consistently observed in the SCN, NPAS2 alternatively functions

as a dimerization partner for BMAL1 and can similarly regulate circadian processes in the forebrain. In addition, the products of clock genes i.e., their proteins are phosphorylated by *CKI ϵ* & *CKI δ* , as well as *glycogen synthase kinase3*, a homologue of *Drosophila shaggy*. Also, they are degraded by agents such as FBXL3 and β -TRCP1, which are components of ubiquitin ligase complexes. These work cooperatively to control the circadian oscillatory period by controlling the levels and transport of PER and CRY (Hastings et al., 2018; Wulff et al., 2010). Changes have been noted in the expression of clock genes of shift workers when compared to those who work normal shifts (Bracci et al., 2014; Taniyama et al., 2015). Moreover, subjects in sleep deprivation studies show similar alterations in clock genes, findings that underscore the importance of clock genes and sleep interactions (Franken and Dijk, 2009). The regulation of clock genes is associated with the usual steps involved in gene expression regulation, including, e.g., modifications in post-transcriptional and post-translational processes (Mure et al., 2018; Takahashi, 2017). A few significant proportions of the genome and transcriptome are expressed in a circadian oscillation. A new study on a primate diurnal transcriptome atlas, which contained a wide variety of major neural and peripheral tissues, found that 82 percent of genes were rhythmic in at least one region of the tissue (Mermet et al., 2017).

The circadian timing system interacts with the sleep/wake homeostatic process. Sleep advances or delays occur when lighting exposure is similarly advanced or delayed. The level of SCN electrical activity follows a normal pattern, with greater activity occurring in the daytime hours when compared to the biological night. Aside from the circadian regulation of firing rate, neurons in the SCN fire at higher rates during REM sleep and lower rates during NREM sleep (Deboer et al., 2003). It's been postulated that modifying the SCN neurons' firing rate during sleep can potentially alter the molecular feedback loops that are essential for sustaining the circadian timing system (Colwell and Michel, 2003). Inasmuch as there is close integration between the sleep homeostat and the circadian oscillator, the amount of slow wave and REM sleep that has taken place in the preceding sleep/wake cycle can be monitored by the circadian system. Furthermore, in mice, the SCN consolidates the sleep/wake cycle by generating an arousal signal during the subjective night (i.e., the active period), allowing it to change baseline sleep quantity (Easton et al., 2004). Although the SCN is not directly involved in sleep homeostasis, as demonstrated by an increase in electroencephalographic delta power following sleep deprivation, it does play a critical function in sleep regulatory control that extends beyond the timing of the states of vigilance.

Mutation of clock genes affects the homeostatic process which results in the alteration of sleep architecture and duration (Archer et al., 2018; Maire et al., 2014). Altered single nucleotide polymorphisms have been reported in four major clock genes (*Clock*, *Per*, *Cry*, and *Bmal1*), and have been found in autism spectrum disorder (ASD) (Nicholas et al. 2007; Yang et al. 2016), depression-related disorders (Serretti et al. 2005; Shi et al. 2016), and bipolar disorder (Benedetti et al. 2003; Karthikeyan et al. 2014). Similar dysfunctionalities in clock genes have been noted in attention deficit hyperactivity disorder (ADHD) (*Clock*) (Kissling et al. 2008; Xu et al. 2010), and schizophrenia (Takao et al. 2007; Zhang et al. 2011) (*Clock*). Other work has shown that, in addition to the genes noted above, clock gene involvement occurs in psychiatric disorders such as winter depression, ASD, and schizophrenia (*Npas2*), bipolar disorder (*ROR α* and *ROR β*), and ASD (*Timeless*, *DBP*, and *CK1 ϵ*) (Charrier et al., 2017). The overlap of identical clock gene occurrence in different psychiatric disorders implies that common pathways and etiopathology may underlie these disorders. Evidence indicates that the clock genes *Cry1*, *Cry2*, *Per2*, and *Npas2* are likely to have an association with major depressive disorder (MDD), while the gene *Timeless* is a promising candidate for association with sleep disorder in MDD (Melhuish Beaupre et al., 2018).

Studies have noted that the activity of the circadian timing system is affected significantly by the sex of the individuals. Certain disparities between men and women exist in the architecture, physiology, and pathology of the circadian endocrine rhythms (reviewed elsewhere, Nicolaides and Chrousos, 2019). Between males and females, differential regulation of daily rhythm of clock gene expression has also been seen at the molecular level; additional parameters include, but are not limited to, timing, duration, and phase of melatonin and body temperature rhythms, morningness-eveningness, and the entrained phase (Lim et al., 2013; Duffy et al., 2011). The SCN and its afferent and efferent circuits, and peripheral rhythms vary according to the sex of the individuals. The molecular dialogue between the circadian timing system and the hypothalamic-pituitary-adrenal (HPA) axis, as well as other behavioural rhythms, differs between men and women (Nicolaides and Chrousos, 2019). Factors that might affect the interpretation of these findings include uncontrolled experimental conditions, masking effects from environmental and behavioral factors, along with variation in timing and phase. Substantial circadian misalignment, as found in shift work, jet lag, sleep deprivation, or clock gene knockout animals, has been linked to major impairments in reproductive function. Alterations in hormone secretion patterns, reduced conception rates, greater

miscarriage rates, and an increased risk of breast cancer are just a few of these inadequacies. Because disrupted hormone cycles might further affect clock gene expression patterns, female health may be especially vulnerable to the impacts of desynchronizing work hours. Estrogen, for example, affects the expression of clock genes in the uterus, ovaries, and SCN (Mahoney, 2010).

Two-process model of sleep regulation

The two-process model of sleep regulation, proposed by Borbély and others, is based on the notion that the circadian clock and sleep are linked. (Borbély, 1982). The model includes a sleep-independent circadian mechanism (Process C) as well as a sleep-dependent homeostatic process (Process S) that regulates waking and sleeping times. The circadian component of sleep is closely related to circadian rhythms of metabolic and endocrine events and has an adaptive significance. In contrast, the impact of Process S is derived from sleep debt (i.e. build-up of homeostatic sleep drive or sleep pressure) accumulated during waking followed by an exponential decline during sleep. The level of Process S at sleep onset is responsible for the increase in sleep tendency, and is, therefore, heavily influenced by prior wakefulness. Process C is dictated by the recurrent variations in sleep tendency (SP) i.e. the ability to fall asleep) and thus presumably is controlled by a circadian

pacemaker. This model, therefore, accounts for how the combined action of the two processes is responsible for the onset and the duration of sleep.

This model was subsequently elaborated into a quantitative version of a model that included the ultradian (a period shorter than 24 h) dynamics of sleep as related to the NREM/REM sleep cycle (Borbély and Achermann, 1999). It was further noted that the time course of EEG slow-wave activity (SWA), which is a tightly regulated process, is the major marker of NREM sleep homeostasis. Individual differences in Process S were found, and particularly in its build-up and dissipation were found (Rusterholz et al., 2010). Subsequently, the same group observed that the time constants for dissipation seemed to follow two independent traits (Rusterholz et al., 2017). A modification of the two-process model proposed that Process S might increase during sleep and diminish during wakefulness (Putilov et al., 1999). Furthermore, circadian modulation affects displaced sleep processes (time constants and asymptotes), sleep deprivation, and homeostatic baseline sleep. The two-process model was subsequently developed to show that the reaction to sleep deprivation in both long and short sleepers could be predicted by using the same S time constants (Aeschbach et al., 2017). More S is found in short-sleepers than long-sleepers. It has

been found that, when compared to the S of long sleepers, higher levels occur in short sleepers. Nakao and coworkers developed a model of thermoregulatory feedback control modulated by two circadian oscillators, one a temperature rhythm, the other mediating sleepiness. The homeostatic pattern, sleep timing, and sleep deprivation aspects of sleep cycles were developed by integrating heat load during awake and heat loss during sleep (Nakao et al., 1995a, Nakao et al., 1995b). Kronauer conceived a model of how light affects human circadian rhythms. This model accounts for the influence of light pulses and their intensity on the pacemaker phase and amplitude. Additionally, it describes their entrainment effects (Klerman et al., 1996; Kronauer, 1990). The two-process model with homeostatic S and C processes was shown to be considerably comparable to a model based on mutual inhibition of sleep-promoting neurons in the ascending arousal system (Skeldon et al., 2014). This reciprocal inhibition model is commonly used as a tool for understanding a wide range of sleep-wake events, as well as for practical applications including the design of shift work schedules that are minimally disruptive to endogenous cycles. .

A study of the two-process model can thus provide insights into the functioning of mutual inhibition between the respective processes. Borbély recently modified the original two-process model. The SCN is now thought to orchestrate and integrate

rhythms rather than simply generate and drive them, and interactions between processes are now regarded as continuous (Borbély et al., 2016). Homeostasis of (as-yet-undefined components of) brain function, according to the model, can be achieved through the periodic transitions occurring between sleep and wakefulness, which themselves are produced by either an internal oscillator that is separate from the SCN or due to oscillations resulting from behavioral relaxation within Process S (Borbély et al., 2016). Definitive conclusions about this issue require additional research. An alternative hypothesis has suggested that a relaxation-type sleep-wake switch exists which is based on both orexinergic (wakefulness) and homeostatic (sleepiness) drives (Fulcher et al., 2014; Phillips et al., 2011). One of these models (Fulcher et al., 2014) proposes a combined action process, i.e., that wake-active processes are due to actions of monoaminergic neurons (in the brainstem and hypothalamus) that are mutually inhibitory, and that these processes operate in parallel with ventrolateral preoptic (VLPO) neurons in the hypothalamus.

According to this concept, orexinergic neurons stabilize prolonged waking episodes via excitatory input to monoaminergic neurons in the brainstem. The VLPO neurons inhibit both orexinergic and monoaminergic neurons during sleep, and the resulting diminution in orexinergic input indirectly promotes

sustained sleep. The model predicts that differences in orexin levels cause a change in sleep timing, with delays in the normal sleep episode being correlated with higher orexin levels (Fulcher et al., 2014). The orexinergic promotion of wakefulness is fully compatible with the two-process model because it is controlled by circadian processes and parallels the daily increase in Process C. These processes further support a model of cooperativity between sleep homeostatic mechanisms and the circadian clock.

Experiments carried out with clock gene mutant mice have revealed that clock genes may also be involved in sleep homeostasis, including changes in sleep length and depth. (see for example Kinoshita et al., 2020; Zaki et al., 2020). EEG investigations of slow wave activity (SWA) at various circadian phases, as well as additional modelling of sleep homeostatic processes, especially those happening outside of the regular circadian sleep phase, all support the conclusion that the circadian clock has a broad impact on sleep homeostasis. These effects could be attributed to the clock's influence on the duration and the quality of the sleep/wake process (Deboer, 2018; Lazar et al., 2015; Münch et al., 2010).

Neural basis of the sleep-wake cycle (Process S)

The neural model of sleep-wake regulation in common use until recently evolved from the initial findings of von Economo in patients with viral disease-causing encephalitis lethargica (Von Economo, 1930). This work suggested that the brainstem had wake-promoting effects on the forebrain while sleep promotion from the anterior hypothalamus opposed this waking effect. Subsequent refinement of this hypothesis proposed the existence of wake-promoting and sleep-promoting cell groups (Saper et al., 2005). In this revision cholinergic neurons in the pedunculopontine (PPT) and laterodorsal tegmental (LDT) nuclei of the upper brainstem, the serotonergic dorsal raphe nuclei (DRN), and noradrenergic locus coeruleus (LC) were thought to be responsible for wake promotion (Aston-Jones and Bloom, 1981; Jouvet, 1972). Sleep promotion was provided by GABAergic innervation from the median preoptic nuclei and ventrolateral preoptic nuclei to wake-promoting regions (Sherin et al., 1998; Suntsova et al., 2002; Suntsova and Dergacheva, 2003).

With the advent of new technologies, such as optogenetics, and CRISPR-Cas, there have been major advances in the understanding of this neuronal circuitry (Shiromani and Peever, 2017). The sleep-wake alterations in melanin-concentrating hormone (MCH) and hypocretin/ orexin (OX) activity, proposed earlier by Saper et al. (2005), have now been validated in

several studies (Fuller et al., 2006; Hassani et al., 2011; Jones and Hassani, 2013). In humans, for example, it has been found that OX levels in amygdala dialysates are higher during wake onset, while MCH levels are higher at sleep onset (Blouin et al., 2013). Also shown were correlations with emotional states. This work has supported the conclusion that activation of OX and MCH neurons in the lateral hypothalamus are respectively involved in regulating waking and REM sleep (Yamashita and Yamanaka, 2017). Moreover, fast glutamate and GABA neurotransmission have recently been proposed to be the backbone of the wake-sleep regulatory system, while the monoamine and cholinergic systems are relegated to a modulatory role (Jones, 2019; Saper and Fuller, 2017). In these fast neurotransmitter systems, there are vesicular transporters for GABA and for glutamate which are essential for synaptic neurotransmission (Tong et al., 2008, 2007). The processes of wake and sleep promotion are mutually inhibitory and therefore form the process S part of the Borbély theory (Boes et al., 2018). During waking the need for sleep gradually increases (a homeostatic process) and when sleep promotion outweighs wake promotion there is a rapid switch into sleep, which has been described as akin to an electric “flip-flop” switch (Saper, 2013). However, a higher degree of regulation is involved in the timing of this process. This occurs in the lateral hypothalamic

area outside of the SCN where it is influenced by neuronal input from the circadian system (Process C) inside the SCN. In addition, brain's adenosine system affects arousal and thus gating of slow wave sleep (SWS) expression is influenced by adenosine levels. This in turn can prolong or shorten the duration of the basic functions of sleep and its homeostasis (Lazarus et al., 2019).

Circadian rhythm regulation (Process C)

As previously stated, light is the most powerful synchronizer of the human circadian timing system. The following factors and/or consequences have been identified as key influencers of circadian responses: 1., the organism's circadian phase when exposure to light occurs; 2., properties of the light stimulus or organism-light interaction, such as wavelength, length of the exposure time, and intensity; and 3., the effect of artificial lighting or sunlight on circadian rhythm timing. (Gooley, 2017). Secondary, non-photoc factors in regulating circadian rhythms are food intake (Frank et al., 2021) and physical exercise (de Souza Teixeira et al., 2020). Other *zeitgeber* effects have been reported but full documentation of these effects is still ongoing (Atkinson et al., 2007; Cardinali et al., 2002; Youngstedt et al., 2019). The circadian timing system is more responsive to light during the

biological night, i.e., during normal sleeping hours. As a consequence, atypical sleeping schedules, such as delayed bedtime or early morning awakening, might result in a resetting of circadian clock when combined with nocturnal light exposure. This in turn will express as changes in SCN activity, as well as parallel changes in melatonin secretion. If an individual is exposed to light in the early part of the biological night, a phase delay in the circadian system may occur, i.e., melatonin secretion onset may take place later in the day after light exposure. This effect is similar to that experienced by transmeridian jet passengers when traveling westward (Czeisler et al., 1989). By contrast, individuals exposed to light late at night or in the early morning hours may experience a circadian phase advance.

The intensity of a light stimulus closely affects its circadian resetting potency. This effect is not linear however since the resetting effects occurring in the early part of continuous light exposure are stronger than those occurring later. This has led to the suggestion that a series of shorter light pulses could be more effective in resetting circadian rhythms than a single long exposure to continuous light (Najjar and Zeitzer, 2016). Amplitude differences in circadian responses to longer versus shorter wavelengths of light, as well as their

integration times, enable us to easily assess the relative contributions of melanopsin and cone photoreceptors to circadian photoreception (long for melanopsin, short for other opsins) (Van Der Meijden et al., 2016). In addition to the SCN, the RHT projects melanopsin-containing fibres to various parts of the brain, including those regions that are associated with the circadian and sleep/wake control, photic, and non-photic SCN regulation, and pupillary responses (Gooley et al., 2003; Morin, 2013) The extent to which these inputs may influence color vision is still under investigation (Stachurska and Sarna, 2019).

Until the beginning of the industrial revolution, humans lived in a mostly stable light exposure system, i.e., one that was characterized by predictable daily or seasonal oscillations in the occurrence of sunlight. Before the widespread adoption of artificial lighting, any oscillatory disruptions to the circadian systems of most living beings, particularly humans, were fairly uncommon. By contrast, people living in today's 24 h society are subject to various types of artificial illumination on a regular and continuing basis. These sources include, but are not limited to, ceiling lights, street lamps, television units, computers, and more recently, excessive use of tablets and smartphones. Many experimental studies have been conducted

to characterize the phase resetting effects of illumination, but few studies have been carried out on the adverse circadian effects of artificial lighting under conditions that prevail in the real world (Emens, 2017; Stothard et al., 2017). Artificial light created by electrical devices interacts with the human circadian system and this affects the health of individuals (Münch et al. 2020). The daily cycle of sleep and awakening is maintained by a combination of natural/artificial light during the day and darkness at night. An individual's history of light exposure, including its timing and duration, as well as other factors, such as its quantity, intensity, spatial distribution, and spectral power, are all important variables that affect the human circadian timing system (Chang et al. 2011 & Figueiro, 2013). When the internal time keeping system was exposed to artificial light in the evening, a phase delay occurred, whereas exposure to bright light triggered an advance in the circadian phase. Natural sunlight has a potential range of intensity of around 100,000 lx, this may be reduced to 25,000 lx in closed environments, while artificial lighting, such as that which occurs in offices, is around 400-500 lx (Blume et al., 2019). However, the human circadian system has a non-linear response to light, i.e., relatively low light levels can generate a relatively large response. It was also found in a small group of study that light intensities as low as 100 lx

were sufficient to entrain the human circadian system. The same study also found that entrainment occurred in one individual in response to a light intensity of only 25 lx (Gronfier et al. 2007). Confirmatory evidence has been provided by several dose-response studies showing that low light levels can be highly impactful for the circadian system, even those well below 100 lx (Boivin et al. 1996; Zeitzer et al. 2000; Phillips et al. 2019). In addition, a half-maximum response for phase shifting required only 80-160 lx (Zeitzer et al. 2000) while only 30 lx was needed to suppress evening melatonin levels by 50% (Phillips et al. (2019)).

In today's 24/7 society, night time exposure to a significant amount of artificial lighting is a pervasive and prominent phenomenon. It has been found that individuals who have only a reduced amount of natural sunlight exposure but who are also exposed to nocturnal electric lighting tend to exhibit a delayed onset of melatonin release. This finding of a late onset of subjective biological light has been associated with further disruptions in the in sleep/wake timing in such individuals (Wright et al. 2013). Light emitting devices appear to produce similar effects. One study found that individuals who read an electronic book (with a light emitting luminance of 30-32 lx) for 3-4 h prior to the bedtime exhibited delayed sleep

onset, decreased melatonin synthesis, and delayed timing of the circadian clock when compared to those who read with printed paper books (Chang et al. 2015). Further, it was found that individuals who were exposed in the evening to a range of light intensities (30-50 lx for 5 h) showed variations in melatonin suppression (Phillips et al. 2019) indicating the interindividual differences exist in how the human circadian timing system responds to day light and evening light exposure.

The negative impact caused by the exposure to light in the evening is significant. The effects of light-induced circadian disruption, which is directly the result of melatonin suppression and reduced melatonin synthesis, has further consequences which may result in sleep disruption, as well as impairments to thermoregulation, blood pressure responses, and glucose homeostasis (Gooley et al., 2011). It has thus been argued that night time exposure to artificial light, which differs in its spectral composition and intensity from natural sunlight, could be considered a potential physiological stressor (Grubisic et al., 2019).

Interactions between Process S and Process C

The circadian timing system and the sleep/wake cycle communicate in a variety of ways, most importantly in the timing of sleep onset (see Fig 1). However, the detailed neurobiology of these

interactions still needs to be elucidated. Defining the neural pathways, neurotransmitters, and neuromodulators that participate in these interactions will potentially lead to the development of novel treatment modalities for several circadian rhythm sleep dysfunctions.

<<<< Insert Figure 1 about here>>>>

Melatonin and circadian rhythms

Melatonin is synthesized in the pineal gland and secreted into the cerebrospinal fluid (CSF) of the posterior third ventricle and the blood. This activity occurs exclusively at night (Pandi-Perumal et al., 2006; Reiter et al., 2014). Melatonin levels in both the CSF and plasma remain low during the day but become elevated at night, all in accordance with a circadian rhythm that is under control of the master oscillator, the SCN; this cadence however is more apparent in the CSF, where its amplitude is greater than it is in the blood. The final control over melatonin's circadian rhythm is due to postganglionic sympathetic signaling, which itself is affected by inputs from several sources; in addition to the SCN, these include the hypothalamic paraventricular nucleus (PVN), polysynaptic projections to the intermediolateral column of the cervical spinal cord, and preganglionic (PG) projections to the superior cervical ganglion (SCG) (see Fig 2). The dim light

melatonin onset (DLMO) marks the beginning of nocturnal circadian activity. This prominent and reliable indicator corresponds to the first spike of melatonin synthesis and occurs in the early evening under low light conditions (Pandi-Perumal et al., 2007; Lewy et al., 1999).

Melatonin is essential for the normal regulation of sleep cycles in both diurnally and nocturnally active organisms, including humans. In humans, a significant urge in sleep tendency at night normally occurs two hours after the initiation of melatonin synthesis (Zisapel, 2018). The early evening rise in melatonin helps to trigger sleep onset while inhibition of melatonin secretion by bright evening light delays the rise in melatonin output. In addition to assisting with sleep onset, the duration of nocturnal melatonin activity serves to relay night length information to the brain, including the SCN, as well as to non-neural tissues. In both normally sighted and profoundly blind individuals, the circadian melatonin rhythm is closely linked to the sleep cycle. Other factors modify melatonin activity and these includes the effects of age, the presence of neurodegenerative disease, as well as certain pharmaceuticals (e.g., β -adrenoceptor blockers, clonidine, naloxone, and non-steroidal anti-inflammatory agents). These conditions or agents either reduce or abolish nocturnal melatonin synthesis and may

cause dysregulated sleep (Pandi-Perumal et al., 2008). In addition to promoting sleep onset and regulating circadian rhythms, melatonin, when it is administered exogenously, can benefit cardiovascular health and may slow cognitive decline in patients with Alzheimer's disease (Baltatu et al., 2019; Shukla et al., 2017). The observation that melatonin administration can reduce the normal suppression of the endogenous night signal has led to suggestions that this therapeutic approach could be of benefit for facilitating safe physical and mental aging (Srinivasan et al., 2009). It has been found that females tend to produce more melatonin than males, although the gender disparities in melatonin levels among elderly persons remain unclear (Obayashi et al., 2015). In a cross-sectional study (N=528; males=247; females=281), it was reported that the older females had significantly lower urinary melatonin excretion than older males (Obayashi et al., 2015). These findings thus underscore the importance of taking into consideration individual factors such as sex before initiating a program of melatonin supplement therapy (Olcese, 2020).

Disruption of these rhythms negatively impacts physical and mental health through its destabilizing effects on behavior, cognition, and mood (Albrecht, 2013; Bedrosian and Nelson, 2017; Benca et al., 2009; Richter, 1965; Schulz, 2007; Wirz-Justice,

2006). A large body of evidence has now established that chronic disruption of sleep and circadian rhythm can lead to the development of a number of physiological and psychiatric disorders (Karatsoreos, 2014; Melhuish Beaupre et al., 2018; Salgado-Delgado et al., 2011; Wirz-Justice, 2007, 2003). Zaki and coworkers reviewed evidence showing the importance of biological clock effects in mood disorders and depression (Zaki et al., 2018). Evidence from both animal model and epidemiological studies has shown that chronic circadian rhythm disturbance seems to raise the risk of metabolic diseases (Cardinali and Vigo, 2017; Karthikeyan et al., 2014; Reiter et al., 2012). It has been observed that time-restricted food intake activity without caloric restriction can promote and maintain greater metabolic homeostasis than such activity without restraints on time of food ingestion. This has led to the suggestion that such timing effects could be incorporated into a therapeutic strategy for managing metabolic disorders (Dashti et al., 2021).

Circadian rhythms play an integrative role in physiology and behavior, the clinical exploitation of which promises to have important utility for treatment (Panda, 2016). Evidence of circadian rhythm involvement in various physical and psychiatric disorders has been shown at multiple levels. Mice with a *Clock*

gene mutation exhibit behaviors that resemble the manic phase of human bipolar disorder. These have been shown to include hyperactivity, reduced sleep duration, and fewer behavioral symptoms of anxiety or depression (Roybal et al., 2007). Deletion of *Clock* genes in mice has been linked to oscillations of islet genes and subsequent disturbances in development, glucose metabolism, and insulin signaling. Again, these defects appear to be linked to genetic deficits in timing regulation. Mice with *Clock* and *Bmal1* mutations have compromised glucose resistance, decreased insulin secretion, and defects in pancreatic islet size. Further, these defects tend to proliferate and worsen with age (Marcheva et al., 2011). There is a strong association between sleep problems and ADHD, especially relative to sleep onset (Coogan et al., 2016). These are accompanied by a variety of rhythmic disruptions which can be detected physiologically, in, for instance, the endocrine system, as well as at the molecular and behavioral levels. Recent findings in these areas have suggested that phase advance techniques, currently used in sleep medicine to treat timing delays in circadian rhythms, may also help reduce symptoms of ADHD (Coogan et al., 2016). While having obvious applications in psychiatry, circadian realignment therapy could potentially be used for childhood developmental disorders which may be precursors to more serious problems in adulthood. Long-term

sleep restriction and circadian disruption in adolescent individuals have been shown to promote unhealthy behavioral activities and to increase the occurrence of substance use disorders (Logan et al., 2018).

Circadian Rhythm Sleep Disorders

Various sleep disturbances are strongly correlated with disruptions in circadian rhythms. The ICSD-3 (American Academy of Sleep Medicine, 2014) and DSM-5 (American Psychiatric Association, 2013) have classified circadian rhythm sleep disorders into several categories. The DSM-5 recognizes five disorders: delayed sleep-wake phase disorder (DSPS), advanced sleep-wake phase disorder (ASPS), non-24h sleep-wake disorder (N24SWD), irregular sleep-wake rhythm disorders (ISWRD), shift work disorder (SWD), and groups other disorders as parasomnias (Reynolds 3rd and O'Hara, 2013). The ICSD-3 also includes jet lag sleep disorders (JLD) (Sateia, 2014). Some of these disorders may be 'intrinsic', inasmuch as they are frequently chronic, familial, and related to genetic alterations. Extrinsic CRSDs, are triggered by environmental enforcement of an unfamiliar photoperiod (e.g. Jet Lag Disorder or JLD), work schedule (e.g. Shift-Work Disorder), or use of light emitting electronic devices (Duffy et al., 2021).

Insert Figure 2 about here

Diagnosis of circadian rhythm sleep-wake disorders

A recurrent or persistent pattern of sleep alteration, primarily due to changes in circadian regulation or a misalignment between the individual's internal body clock to geophysical LD cycle, and a sleep/wake pattern imposed separately by the schedule are required criteria for establishing the presence of a circadian rhythm sleep disorder. The disorder must also cause significant sleep disruption or distress in physical or social functioning. A further requirement is that the disruptions must be measurable. It is thus important to have quantified measures of the circadian phase (advance/delay; i.e. determining an alteration in the phase of the circadian rhythm), circadian peak, period length, and disruption of circadian oscillators is critical for delineating their function in sleep-wake disturbances (Dijk and Duffy, 2020). Traditionally, a precise assessment of the human circadian system has required extensive record keeping regarding, e.g., endogenous melatonin levels or core body temperatures, and it has been suggested that such protocols are often not clinically practical. As a consequence, alternative approaches for the assessment of circadian status are now being explored. These techniques, which are continuing to undergo development, utilize machine learning or use large datasets

obtained from the analysis of transcriptomes or metabolomes (Dijk and Duffy, 2020).

Delayed sleep phase syndrome (DSPS) is the most common occurring circadian rhythm sleep disorder (CRSD). An aberrant delay in the occurrence of a major sleep episode in proportion to the environmental zeitgeber is the most common symptom of this illness (Fahey and Zee, 2006; Nesbitt and Dijk, 2014). In some patients, a polymorphism in *Per3* has been discovered (Ebisawa et al., 2001). While in others, a variant in the *Cry1* gene has been discovered (Patke et al., 2017) however, most affected individuals have no known genetic abnormality. Many of the latter are adolescents and young adults who make extensive use of smartphones in the evening, a behavioral phenomenon so ubiquitous in modern-day life that it has been christened "cell phone addiction" (Gutiérrez et al., 2016). Recent evidence suggests that differences in endogenous period length (Micic et al., 2013) and/or light sensitivity (Abbott et al., 2020; Aoki et al., 2001; Watson et al., 2018) occur in this population.

The sleep phase in advanced sleep phase syndrome (ASPS) occurs unusually early in the evening compared to the night. Advanced sleep phase disorder (ASPD) patients have difficulties in remaining awake in the evenings and frequently fall asleep before completing personal and social tasks. Additionally, the

wake time of these patients is uncomfortably early and considered atypical in comparison to their peers. Several or all of these families share mutations that affect *Per2*, either in *CK1ε* or on *Per2*'s binding site (Toh et al., 2001; Zhang et al., 2016).

Non-24 h Sleep-Wake Phase Disorder (N24SWD) is a condition in which an individual's circadian clock does not synchronise to a 24 h day. As a result, patients' sleep-wake cycles typically display a gradual delay (usually minutes to hours), based on the circadian period of the individual. The timing of sleep gradually varies during a symptomatic phase, causing patients to suffer repeated bouts of daytime hypersomnolence and nighttime insomnia. The majority of individuals with N24SWD are entirely blind, although it can also affect people who have normal vision. Unlike the other CRSDs, a N24SWD diagnosis requires at least 14 days of consistent sleep/wake patterns, which are typically recorded using sleep diaries and/or actigraphy. Treatment of this disorder is very often successful with oral melatonin (Sack et al., 2000) or with the recently approved prescription medication Hetlioz® (tasimelteon, a melatonin analog) (Traynor, 2014).

Irregular sleep-wake disorder (ISWRD) is characterised by the absence of a discernible regular sleep-wake cycle and, as a result, the primary sleep episode lacks a temporal component.

Such irregular patterns have been reported in members of family groups with DSPD who also show mutations affecting CRY1 (Patke et al., 2017). It has been suggested that delay under restrictive environmental conditions could additionally cause symptoms of sleep fragmentation (Patke et al., 2017).

The American Academy of Sleep Medicine (AASM) guidelines for the management of circadian rhythm disorders have concluded that the most effective interventions can be broadly categorized into (1) Having a practice of regular sleep/wake schedule and/or physical activity/exercise, (2) Managing routine tasks with specific time and/or avoidance of sunlight & artificial light, (3) use of chronobiological agents altering or promoting sleep cycles; and (4) alternative therapies that influence sleep/wake habits by modifying bodily functions (i.e., somatic interventions) (Auger et al., 2015).

Circadian Draining of Brain Waste During Sleep

Recent theorizing suggested that a linkage may exist between sleep and the glymphatic system and that this relationship may serve an important rebalancing function. Homeostasis in the central nervous system, which requires an efficient elimination of metabolic waste products from the brain, exhibits circadian characteristics. This circadian control is manifested, for example, by the dependence of the

time of day on the permeability of the blood-brain barrier, demonstrated by the daily variation of the relative levels of metabolites between the CSF and the blood. The choroid plexus, which makes up an important part of the blood-CSF barrier, exhibits the rhythmicity of the circadian clock (Myung et al., 2018), regulating the exchange of fluids over time between blood and CSF. Glial cells of the hemolymph-brain barrier in fruit flies have a circadian clock that has been shown to regulate xenobiotic output (Zhang et al., 2018). **These findings support the idea that astrocytic glial cells regulate the blood-brain barrier (BBB) and influence the clearance of metabolic waste in mammals.**

Diffusion has long been used to explain solute changes in the extracellular space (ECS) of the central nervous system. It is now known that the process is somewhat more complex, with several local regulators such as adenosine, biogenic amines, NO, H⁺, and K⁺ working together to integrate blood flow and metabolism for the removal of waste products. More recently, and in parallel with conventional explanations for the process and function of removal of toxic products from the brain, a "glymphatic hypothesis", has been proposed. This model emphasizes the importance of active lymphatic-like movements in the cerebral ECS (Boespflug and Iliff, 2018; Iliff et al., 2013, 2012; Jessen et al., 2015). Although the CNS lacks lymphatic

vessels, it has been suggested that this process may nevertheless take place in the brain due to water exchange driven by aquaporin-4 (AQP4). The vascular lumen is altered by the passage of solute matter in perivascular astrocytes, which have high concentrations of Aquaporin-4 (AQP4). The AQP4 water channel is found mostly in the foot processes of astrocytes. The release of water through AQP4 may account for an actively guided exchange of fluid between the periarterial and perivenous spaces, which are linked in other areas of the ECS, such as the interstitial fluid, by the convective flow. Furthermore, it has been hypothesized that pulsation by arterioles, together with respiration-dependent venous collapse, may further increase this flow (Nedergaard, 2013).

Sleep, in particular NREM sleep, is strongly associated with the removal of brain waste (Boespflug and Iliff, 2018; Iliff et al., 2013, 2012; Jessen et al., 2015). An example is the elimination of amyloid peptide β ($A\beta$), which improves significantly during sleep. The discovery that clearance of injected radio-labelled $A\beta$ peptide is considerably decreased in AQP4 - / - mice lends support to the idea of $A\beta$ glymphatic clearance (Iliff et al., 2012). Inasmuch as during NREM sleep the cortical interstitial space expands by >60%, solute exchange between CSF and interstitial fluid (ISF) is at its maximum, and thus presents only limited resistance to CSF and ISF movement

during this sleep stage. The glymphatic system's contribution to the sleep repair mechanism is strongly reliant on age-related sleep profiles. Sleep among elderly subjects is associated with delayed and decreased fluid entry into the CSF along the perivascular tract and the pial wall (Zeppenfeld et al., 2017). Glymphatic system dysfunction is also visible in several neurological conditions, including Alzheimer's disease (AD), traumatic brain injury (TBI), and stroke (Boespflug and Iliff, 2018). Taken together, these findings show that the circadian clock and the REM/NREM cycle control many critical processes in the brain. Among other activities, these include the maintenance of regulated trafficking of substances into and out of the brain, brain metabolism, and sleep/wake control. Any tampering of the circadian clock and pathways may result in inefficient and dysregulated metabolic processes (Albrecht and Ripperger, 2018).

The impact of sex related variables on variations in sleep and circadian rhythms

The challenge of identifying how sleep is affected by circadian regulation, and more importantly about the consequences of disruptions to this regulation, is further complicated by how these processes are differentially affected in men and women. Men and women have differential responses in

hypothalamic-pituitary-gonadal (HPG) axis, hypothalamic-adrenal-pituitary (HPA) axis, and sleep-arousal systems. Understanding how these systems' circadian rhythms are disrupted differs by sex and is associated with malfunction and disease.

Understanding sex-specific circadian timing mechanisms may lead to more effective treatments for these illnesses (Bailey and Silver, 2014). Sexual differences in sleeping patterns have been underemphasized in experimental designs. Women in human studies and females in animal studies have always been historically underrepresented in research studies (Feldman et al., 2019), circadian and sleep-related studies are no exception.

In both basic and clinical research studies, sexual variations in normal and pathological sleep have been identified globally (Anderson and FitzGerald, 2020; Jonasdottir et al., 2021; Tang et al., 2017; Krishnan and Collop, 2006). Sleep characteristics such as sleep timing and quality are significantly influenced by age and sex (Li et al., 2021; Madrid-Valero et al., 2017). Females have a higher frequency of sleep difficulties than males, according to various surveys (Madrid-Valero et al., 2017; Stranges et al., 2012; Suzuki et al., 2015; Uhlig et al., 2014; Zhang and Wing, 2006; Ishigooka et al., 1999). The prevalence of insomnia increases with age, as reported in most population-based research undertaken in a

number of countries (Kerkhof et al., 2017; Leng et al., 2014; Ohayon et al., 2004; Middelkoop et al., 1996; Reyner et al., 1995). When compared to men, women of all ages report having sleep that is of poorer quality, more fragmented, and less satisfying (Tsai and Li, 2004). Some reviews have suggested that studies of sleep disorders among college students and other groups have suffered from a failure to properly account for how these problems manifest in males and females (Galambos et al., 2009; Carney et al., 2006). In a meta-analysis of the pooled prevalence study in the general population of China (N=115,988), no significant differences were found in the prevalence of insomnia between men and women (Cao et al., 2017). In another study, variations in the circadian amplitude of plasma melatonin and electroencephalographic (EEG) slow-wave activity (SWA) could not explain the sex differences in cognitive regulation (Santhi et al., 2016). Taken together, these research deficiencies point to the presence of a limited appreciation among many researchers for how sex differences can affect sleep and circadian regulation.

Another phenomenon which has been documented to affect circadian rhythms is shift work. There is growing evidence that suggests that the female shift employees are more likely than male shift workers to complain about sleep disorders, and

they have more trouble initiating and maintaining sleep after work. Hence, impairments in the quantity and quality of sleep are greater among women than men, and they report greater difficulties in adjusting to shift work as compared to their male counterparts (Chung et al., 2009; Admi et al., 2008). Night shift work contributes to a greater prolongation of pregnancy, reduces fetal growth in industrial shift workers, and increases the requirement for fertility therapy in young women (Fernandez et al., 2020; Zhu et al., 2004). Clinical and basic research findings point to the importance of sex steroids in sleep modulation, as well as to how neuroendocrine mediators and sex variables influence sleep (Mong and Cusmano, 2016). Sleep health providers should be aware of these discrepancies at all levels of disease prevention and health promotion to improve the quality of life for people with sleep-related health concerns. These include the diagnosis, treatment, management, and patient education (Boccabella and Malouf, 2017).

There are still major information gaps in research with respect to the sleep of men and women. More basically, there continues to be a general lack of awareness among the scientific community that many studies have frequently failed to include females in their experimental designs (Mallampalli and Carter, 2014). These deficits indicate that considerable work remains to

be carried out to explicate the impact of sex related differences on sleep disorders (Mallampalli and Carter, 2014), and more specifically suggest that future sleep studies should include more females. It is essential for the requirements of evidence based medicine, and especially for the creation of targeted therapy programs, that research on sleep disorders take sex related biological differences into account.

Conclusion

The discipline of sleep medicine has advanced significantly since the pioneering sleep investigations of Dement and Kleitman (1957), Jouvet and Courjon (1959), and Rechtschaffen et al (1966). It could be argued that, in terms of research and acknowledgment by the broader scientific community, the importance of circadian rhythms to overall health is now more fully appreciated. Despite the growing evidence for its efficacy, chronomedicine or chronotherapy (i.e. timed administration of a drug or other interventions according to the patient's biological clock) is still under-researched and under-utilized. For example, the use of light therapy, dark therapy, blue-blocking lenses, wake therapy, and effective utilization of chronobiotic substances such as melatonin and its derivatives. More research and clinical applications for chronotherapy are

hoped to result from the growing awareness of the significance of circadian rhythms in both sleep and overall health.

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Authors' Contribution

All authors met the ICMJE criteria for authorship. All authors contributed equally in the subsequent preparation of the manuscript. SRP, DPC, RJR, GMB conceived and designed the

outline of review; NFWZ, RK performed the literature search; SRP, DPC, RK wrote the paper; NFWZ, DWS, RJR, GMB performed critical review of the initial and/or copy edit the subsequent drafts; and all authors (SRP, DPC, NFWZ, RK, DWS, RJR, GMB) approved the final version of the manuscript prior to the submission.

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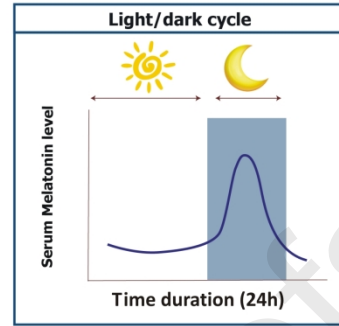
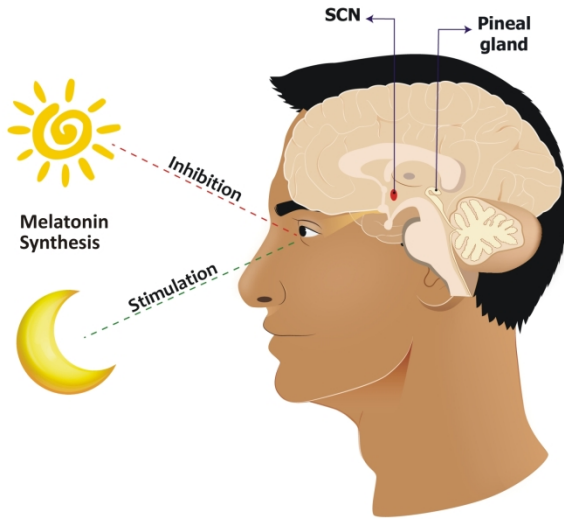
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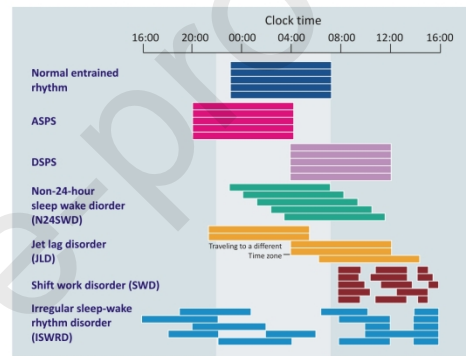
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Sleep regulation



AANAT & HIOMT - Light regulation of two key enzymes involved in the biosynthetic pathway of melatonin

Room Light (< 200 lux)
Constant light (< 3 lux)
Light at Night
Light pollution

Suppresses melatonin → Sleep disturbances

AANAT - Aralkylamine N-acetyltransferase
HIOMT - Hydroxyindole-O-methyl transferase

Figure 1.

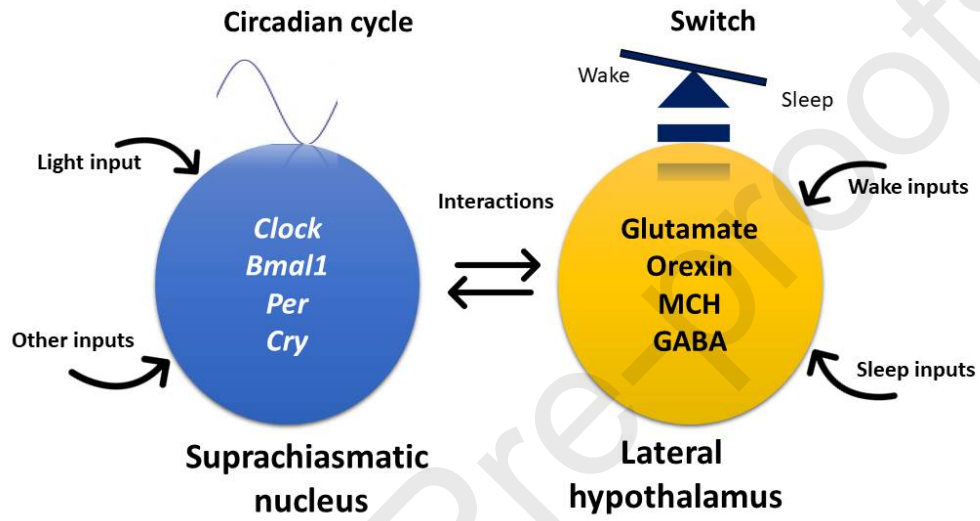


Figure 2.

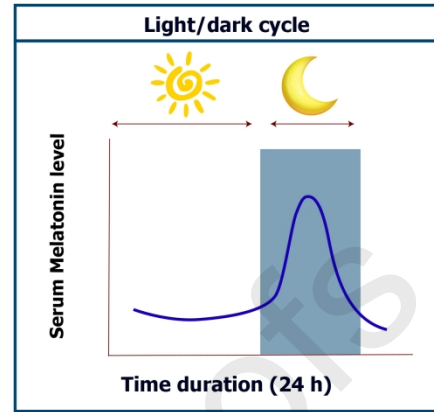
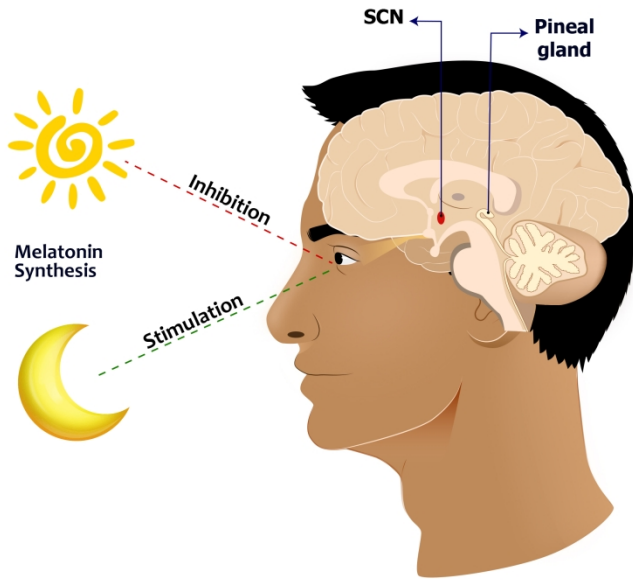
Legends for the figures

Figure 1. a) Depicts the circadian rhythm of melatonin secretion and its involvement in sleep regulation. Figure b & c addresses the role of light on pineal melatonin synthesis, suppression, and subsequent sleep disruption. Generally, indoor room light (<500 lx) has been proven in investigations to generate substantial melatonin suppression and phase shift responses. Again the effect varies with experimental conditions, the nature of light, the time, the intensity, and the duration of administration. d) Shows various types of circadian rhythm sleep disorders..

Figure 2. Putative pathways linking the SCN and LHA in regulating sleep and wake timing in humans. The cyclic process in the SCN is cued by light signals from the melanopsin containing retinal cells travel via the RHT to the SCN. The 24 h cycle in SCN is based on the binding of CLOCK and BMAL1, which initiates the transcription and translation of PER and CRY, then the translated PER and CRY proteins repress CLOCK-BMAL1 function. This cycle interacts with the switch process in the lateral hypothalamus by an undefined pathway. The LHA of the diencephalon receives multiple external wake and sleep promoting neurons that act on OX neurons to promote waking and MCH neurons to promote REM sleep. Together with glutamate and GABA neurons, these are thought to control the switch from waking to sleep. This system provides inputs to the SCN that are yet to be defined. **Abbreviations used.** LHA, lateral hypothalamic area; MCH, melanin-concentrating hormone; OX, orexin/hypocretin; RHT, retinohypothalamic tract; SCN, suprachiasmatic nucleus.

HIGHLIGHTS

- In the brain, the organization of timing is essential for optimal brain physiology and function.
- Sleep and waking are meticulously timed and integrated with bodily functions
- For optimal functioning, the circadian clock and external stimuli must be properly aligned.
- In humans, failure results in circadian desynchrony and sleep/wake disturbances.
- The adaptive value of these processes to human health in the periodic environment is now formally recognized.



Sleep regulation

AANAT & HIOMT - Light regulation of two key enzymes involved in the biosynthetic pathway of melatonin

Room Light (< 200 lux)
Constant light (< 3 lux)
Light at Night
Light pollution

Suppresses melatonin → Sleep disturbances

AANAT - Aralkylamine N-acetyltransferase
HIOMT - Hydroxyindole-O-methyl transferase

