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## Circadian rhythm as a therapeutic target

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### Abstract

The circadian clock evolved in diverse organisms to integrate external environmental changes and internal physiology. The clock endows the host with temporal precision and robust adaptation to the surrounding environment. When circadian rhythms are perturbed or misaligned, as a result of jet lag, shiftwork or other lifestyle factors, adverse health consequences arise, and the risks of diseases such as cancer, cardiovascular diseases or metabolic disorders increase. Although the negative impact of circadian rhythm disruption is now well established, it remains underappreciated how to take advantage of biological timing, or correct it, for health benefits. In this Review, we provide an updated account of the circadian system and highlight several key disease areas with altered circadian signalling. We discuss environmental and lifestyle modifications of circadian rhythm and clock-based therapeutic strategies, including chronotherapy, in which dosing time is deliberately optimized for maximum therapeutic index, and pharmacological agents that target core clock components and proximal regulators. Promising progress in research, disease models and clinical applications should encourage a concerted effort towards a new era of circadian medicine.

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In 1729, Jean-Jacques de Mairan documented that leaves of mimosa plants open and close with a daily rhythm, even in constant darkness. Such self-sustained rhythms suggested a so-called free-running, intrinsic biological timer as opposed to a passive response to environmental changes. Different *Drosophila* mutants with altered periodicity showed mutations that were mapped to the same locus, which was subsequently named *period*. The later cloning of the gene<sup>1,2</sup>, followed by the demonstration of negative-feedback regulation in *Drosophila*<sup>3</sup>, was ultimately proven to be a Nobel Prize-worthy discovery. We now have detailed knowledge of the genetic and molecular make-up of the biological timer in virtually all kingdoms of life, from cyanobacteria to humans<sup>4</sup>. It is generally believed that the biological timer, called circadian (from Latin *circa*, meaning ‘approximately’, and *dies*, meaning ‘day’) clock, has evolved independently in a convergent process to adapt to

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and use the daily geophysical cycle for maximum survival and competitive advantage. The endogenous period length is constrained by the natural environmental cycles dictated by Earth's rotation, and remarkably complex regulatory mechanisms have evolved to couple internal and external rhythms to ensure precise periodicity<sup>4</sup>. Clocks are readily responsive to a broad range of entraining signals, or so-called zeitgebers (German for 'time givers'), that determine or readjust the phase or period of the circadian rhythm. In addition to light, which is the dominant zeitgeber for photosensitive clocks, other external (such as feeding and exercise)<sup>5</sup> or intracellular (temperature)<sup>6</sup> cues can reset, or entrain, the endogenous rhythms.

### Zeitgebers

A German word (meaning 'time giver'), from *Zeit* (meaning 'time') and *Geber* (meaning 'giver'). It refers to any external or intracellular cues that can reset or entrain an organism's biological rhythms to the 24-hour day–night cycle of Earth.

Proper circadian rhythms confer a myriad of cell growth and survival advantages. Mirroring results from laboratory hamsters<sup>7</sup>, lesions of the suprachiasmatic nucleus (SCN) — the master pacemaker in the hypothalamus — in wild rodents shorten their lifespan<sup>8</sup>. Although circadian disruption does not lead to acute death in animal models or humans, short-term and long-term adverse effects on fitness and health can result from circadian disturbances<sup>9</sup>. The repertoire of clock-targeting therapeutic strategies now extends beyond the traditional chronotherapeutic applications<sup>10</sup>. In this Review, we will provide an up-to-date account of circadian timing systems in mammals, highlight several diseases closely linked with clock mechanisms and review interventional strategies and therapeutic pipelines that target the clock.

## The circadian rhythm in mammals

A circadian system consists of three components: inputs, the oscillator and outputs<sup>4</sup>. This basic design principle operates at two levels, systemic and cellular<sup>4</sup>. At the systemic scale (FIG. 1), light is a major input signal for the circadian system and resets the master pacemaker, the SCN<sup>11</sup>. A defining feature of the SCN is the robust networking among its oscillator cells compared with that in peripheral tissues<sup>12</sup>. For example, ex vivo SCN culture can sustain the rhythmicity of reporter expression for more than 1 year<sup>13</sup>, while brain regions outside the SCN show a loss or dampened rhythmicity in 120 hours<sup>14</sup>. The main function of the SCN is to coordinate other brain areas and peripheral tissues throughout the body by neural and hormonal signals<sup>15</sup>. Furthermore, the SCN also regulates the circadian secretion of diffusible endocrine signals, such as melatonin from the pineal gland, and glucocorticoids and catecholamines from the adrenal cortex, via the hypothalamic–pituitary–adrenal axis<sup>16,17</sup>.

At the cellular level, the ubiquitous, cell-autonomous molecular oscillator (FIG. 2) is composed of interconnected negative-feedback loops<sup>4,15,18</sup>. In mammals, the transcription factors circadian locomotor output cycles kaput (CLOCK)<sup>19</sup> and neuronal PAS domain-containing protein 2 (NPAS2) form heterodimers with aryl hydrocarbon receptor nuclear translocator-like protein 1 (ARNTL; also known as BMAL1)<sup>20</sup> to drive the expression of

the genes encoding period circadian protein homologues 1, 2 and 3 (*PER1*, *PER2* and *PER3*) and cryptochromes 1 and 2 (*CRY1* and *CRY2*) via direct binding to the E-box enhancer element in the daytime<sup>18</sup>. In the late afternoon or evening, PER and CRY proteins heterodimerize (and also interact with casein kinase 1 $\delta$  (CK1 $\delta$ ) and CK1 $\epsilon$ ), translocate into the nucleus and interact with CLOCK and BMAL1, thus suppressing their transcriptional activity<sup>18</sup>. In the meantime, the protein levels of PER1, PER2, CRY1 and CRY2 also decline by polyubiquitination and subsequent degradation via specific E3 ligase complexes ( $\beta$ -TrCP and FBXL3 (REFS<sup>21–23</sup>), respectively). With the gradual relief of this negative-feedback repression, CLOCK–BMAL1 transcription activity can be restored, and a new cycle will be resumed the next morning. This core feedback cycle takes approximately 24 hours. Two families of nuclear receptors function to stabilize the core loop and govern output transcription in a distinct phase. REV-ERB $\alpha$  and REV-ERB $\beta$ <sup>24</sup> (encoded by *NR1D1* and *NR1D2*, respectively) and RAR-related orphan receptors  $\alpha$ ,  $\beta$  and  $\gamma$  (ROR $\alpha$ , ROR $\beta$  and ROR $\gamma$ )<sup>25</sup>, which are also direct targets of CLOCK–BMAL1, antagonistically regulate *BMAL1* by competitively binding to ROR/REV-ERB-response elements (RORE) and cause an antiphase oscillation between *BMAL1* and *PER2*. A third transcriptional loop involves the proline and acidic amino acid-rich basic leucine zipper (PAR-bZIP) proteins (DBP, TEF and HLF) and the repressor E4 promoter-binding protein 4 (E4BP4; also known as NFIL3)<sup>15</sup>, which competitively bind to the D-boxes, and are driven by the CLOCK–BMAL1 loop or the REV-ERB–ROR loop, respectively.

Together, these three interlocking feedback loops can drive the expression of a large number of target genes by binding to the *cis* elements (E-box, RORE and D-box) in their promoters and enhancer elements<sup>15</sup> (FIG. 2). These genes, called ‘clock-controlled genes’, are highly tissue specific, and exhibit markedly different distributions of circadian phases in the two different tissues<sup>26,27</sup>. Moreover, in mice, an examination of 12 tissues revealed that 43% of all genes oscillate in at least one tissue<sup>28</sup>. Remarkably, increased profiling to 64 tissues and brain regions in male baboons showed that the proportion of protein-coding oscillatory genes in at least one tissue is more than 80%, establishing an overwhelming prevalence of circadian control of gene expression<sup>29</sup>. The tissue-specific rhythmic gene expression is jointly controlled by both central and local circadian clocks.

Circadian gene regulation is a complex, temporally orchestrated process that involves not only the circadian factors mentioned above but also a growing list of secondary or cell type-specific transcription factors, transcription co-regulators and epigenetic activities<sup>4</sup>. The mechanistic basis for tissue-specific circadian expression is emerging<sup>30,31</sup>. For example, hepatocyte nuclear factor 4 $\alpha$  (HNF4A) transcript is rhythmically regulated in the mouse liver and modulates circadian rhythms through repression of CLOCK–BMAL1 activity<sup>30</sup>. Furthermore, transcriptomic analysis of islet cells revealed the circadian expression of genes involved in insulin secretion<sup>31</sup>. Epigenetic regulation is also an active area of research, and histone modifications, chromatin remodelling and topological organization have significant impact on the regulation of both E-box and RORE elements<sup>4,32</sup>. Recent evidence has suggested that, besides transcriptional regulation, post-transcriptional and post-translational mechanisms also have major regulatory roles in rhythmic physiology in all organisms<sup>21,33,34</sup>.

## Melatonin and orexin signalling

The sleep–wake cycle is the most profound circadian rhythm and also a tightly orchestrated physiological process regulated by both sleep-promoting and arousal centres in the brain<sup>35</sup>. In all mammals, either nocturnal or diurnal, the rhythmic secretion of melatonin (5-methoxy-*N*-acetyltryptamine) is modulated solely by the input of light and the circadian clock in the SCN, not by physical activity<sup>36</sup>. During the day, when the SCN is activated by light input from the retina, melatonin production is inhibited. When light conditions turn low, the SCN triggers a neural output signal to induce the synthesis and release of melatonin<sup>36</sup>. Unlike its sleep-promoting effects in humans, melatonin administration failed to promote sleep in nocturnal animals<sup>37</sup> owing to the different mechanisms for circadian regulation of sleep. Therefore, melatonin levels are high at night in both diurnal (humans) and nocturnal species even though they have opposite activity periods. As an important endocrine signal that conveys the circadian information from the SCN to the peripheral tissues, melatonin functions mainly through the activation of two G-protein-coupled receptors, melatonin receptor 1 (MTNR1A; also known as MT<sub>1</sub> receptor) and melatonin receptor 2 (MTNR1B; also known as MT<sub>2</sub> receptor), which differ in terms of tissue distribution and the regulatory effects on sleep and circadian rhythms<sup>38,39</sup>. The MT<sub>1</sub> receptor is predominantly expressed in the SCN and is crucial for the inhibition of neuronal firing in the SCN, and thereby facilitates bedtime sleep onset<sup>38</sup>. Besides its direct effect on sleep, melatonin also exerts a phase-shifting function in the circadian rhythm, through MT<sub>2</sub> receptors in the SCN<sup>38,40,41</sup> (FIG. 3a). Notably, this phase-shifting effect of melatonin on circadian rhythms is thought to be independent of the modulation of circadian gene transcription in the SCN, given that injection of melatonin in rats did not result in immediate changes in the expression of circadian genes — such as *Per1*, *Per2*, *Per3*, *Cry1* and *Bmal1* — in the SCN<sup>42</sup>. However, some circadian rhythm oscillators are potential targets of melatonin. In an experimental mouse model of jet lag, both adaptation of PER1 and CRY1 to the new light–dark cycle in the SCN and the re-entrainment of locomotor activity rhythm were more profound in melatonin-proficient mice with intact MT<sub>2</sub> receptors<sup>43</sup>. Melatonin injections not only induced a phase advance in the *Nr1d1* mRNA expression rhythm in rats but also phase-shifted *Bmal1* mRNA expression, suggesting a link, although perhaps not direct, between melatonin and the core molecular circadian clock<sup>44</sup>. Collectively, melatonin is not only an important physiological sleep regulator but also serves as a time cue to synchronize the circadian physiological and behavioural rhythms, such as sleep–wake and locomotor activity, respectively.

### Phase advance

A phase shift in the circadian rhythm whereby the bedtime and wake-up time will move earlier in the day.

Orexins (orexin A and orexin B), also known as hypocretins, are a pair of excitatory neuropeptides that share 46% homology<sup>45</sup>. The levels of orexin A and orexin B, produced in orexinergic neurons located in the lateral hypothalamus, oscillate with the daily rhythm and peak in the awake phase under the control of the SCN circadian clock<sup>46</sup>. Orexins promote arousal and wakefulness by binding to the G-protein-coupled receptors orexin

receptor 1 (OX1R) and orexin receptor 2 (OX2R), which are widely and differentially distributed. They are expressed at high levels mainly throughout the central nervous system (FIG. 3b) and at relatively low levels in the peripheral organs (adrenal glands, testes and jejunum)<sup>47</sup>. Activation of orexinergic neurons gives rise to a rapid transition from sleep (non-rapid-eye-movement (NREM) sleep or rapid eye movement (REM) sleep) to wakefulness<sup>48</sup>. During this process, OX2R has a major role in stabilization of wakefulness by reducing the transition between the NREM sleep state and wakefulness, whereas OX1R and OX2R both contribute to suppression of REM sleep<sup>48</sup>. Furthermore, recent studies have started to show a reciprocal regulation between the SCN clock cells and the orexinergic neurons. For example, in mice, orexin was found to predominantly suppress expression of PER1-enhanced green fluorescent protein (GFP) in clock cells in the SCN via presynaptic modulation of GABAergic signalling in the daytime and direct activation of leak K<sup>+</sup> currents in the evening<sup>49</sup>. Thus, although orexin did not directly affect the clock phase, it may dampen the SCN control of the circadian rhythm by acutely and temporarily suppressing clock cells in SCN.

#### Non-rapid-eye-movement sleep

Also known as quiescent sleep, with little or no eye movement, rare dreaming and less muscle paralysation than in rapid eye movement sleep.

#### Rapid eye movement sleep

Also known as paradoxical sleep, a unique phase of sleep in mammals and birds characterized by rapid eye movement, low muscle tone throughout the body and vivid dreaming sometimes.

### Circadian rhythm signalling in diseases

There are several ways in which diseases are associated with circadian components or rhythms. First, various symptoms or disease onsets display clear circadian preference. Among the best-known examples are the early morning peak of myocardial infarction and stroke (due to rapid rise of blood pressure) and night-time exacerbation of asthma and other inflammatory diseases<sup>17,50,51</sup>. Some people experiences seasonal and circadian clusters of excruciating headache attacks, with remarkable circadian precision of headache episodes (similar time each day, within minutes)<sup>52</sup>. Furthermore, in mouse models, sundown syndrome, which is often observed in patients with mid-stage Alzheimer disease, is characterized by significant behavioural perturbations in late afternoon and evening<sup>53</sup>. Secondly, besides circadian onset of certain diseases, different disease states and natural ageing correlate with altered circadian rhythms, including the alternation in the expression of circadian genes, sleep–wake cycles and rhythms of key biomarkers such as melatonin and cortisol<sup>9,54</sup>. Although disease-related circadian dysfunctions are often complex, certain disorders display a primary circadian dysfunction. For example, sleep disorders and jet lag are associated mainly with phase dissociation (in this case, also known as dissociation of circadian rhythm and light), whereas ageing, neurodegenerative diseases and metabolic

disease all seem to be affected by attenuated or disrupted circadian rhythms<sup>5,54–56</sup>. Particularly, patients with Alzheimer disease initially exhibit out-of-sync sleep patterns and sleep fragmentations; as the disease progresses, the loss of sleep patterns becomes the primary driver for institutionalization<sup>56</sup>. Lastly, human genetic and genomic studies have uncovered direct links between gene mutations and circadian-related diseases. In addition to the mutations associated with familial sleep disorders, genome-wide association studies have also shown polymorphisms in circadian genes that correlate with diseases. For example, mutations in *MTNR1B* correlated strongly with elevated fasting glucose levels and type 2 diabetes risk in a cohort study<sup>57,58</sup>. In the following subsections, we describe several disease areas with significant research progress on circadian relevance.

### Phase dissociation

In the context of this Review, the dissociation of intrinsic circadian rhythm and the environmental light–dark cycle, as in jet lag and sleep disorders.

## Circadian genes and sleeping disorders

Circadian components have important roles in the regulation of sleep timing<sup>59,60</sup>. Several human mutations have been associated with circadian sleep disorders, most notably in genes encoding PERs and casein kinases that phosphorylate PERs before degradation<sup>60</sup>. For example, mutations in *PER2* (PER2-S662G) and *CSNK1D* (CK1δ-T44A) have been identified in familial advanced sleep phase syndrome, in which patients show significant circadian period shortening and early bedtime<sup>60</sup>. Studies of mouse models expressing these mutant forms validated the causality in the disease and revealed an important role of a PER phosphorylation cascade in gating sleep timing<sup>61</sup>. More recently, a missense mutation in *CRY2* was identified in a human family with familial advanced sleep phase syndrome with marked morning preference and early melatonin onset, and the same mutation led to shortened periods in a mouse model<sup>62</sup>. Genetically engineering the mutant human *CRY2* gene in mice led to elevated binding of CRY-targeting E3 ligase FBXL3, resulting in the degradation of CRY2 that is responsible for the period shortening<sup>62,63</sup>. Delayed sleep phase syndrome is the most common circadian sleep disorder in humans<sup>64</sup>, with an average bedtime onset in the early morning (around 3 a.m.). Previously, variants of human *PER3* gene were found to correlate with delayed sleep phase syndrome<sup>65</sup>. More recent studies in a family with delayed sleep phase syndrome revealed an A-to-C mutation in a splice site following exon 11 of the *CRY1* gene<sup>66</sup>. The mutant allele led to exon 11 skipping, producing a more potent CRY1 form (CRY1 11) that interacts with CLOCK and BMAL1 with greater affinity than the wild-type protein and consequently greater inhibitory activity.

### Delayed sleep phase syndrome

A circadian sleep disorder in which a person's sleep is delayed by 2 hours or more beyond what is considered a conventional bedtime, thus causing difficulty in waking up in the morning.

## Ischaemia and reperfusion injury

Ischaemia and reperfusion injury typically occurs as obstruction of blood flow to an organ in a wide range of diseases, including myocardial infarction, ischaemic stroke, liver injury and acute kidney injury<sup>67,68</sup>. Once the arterial obstruction is alleviated, reperfusion of the ischaemic tissue occurs, which can cause an even stronger inflammatory response and additional collateral tissue injury<sup>68</sup>. During ischaemia, the organ becomes profoundly hypoxic and hypoxia-inducible transcription factors are stabilized<sup>68,69</sup>. In many instances, this stabilization of hypoxia-inducible transcription factors enables the ischaemic tissues to adapt to limited oxygen availability — for example by transcriptionally enhancing the glycolytic capacity or by attenuating hypoxia-driven inflammation<sup>70,71</sup>. Daytime variations of the severity of ischaemia and reperfusion injury could be linked to an interaction between circadian rhythm and hypoxia signalling<sup>72,73</sup>. Initial evidence comes from studies of *Per2*- and *Hif1a*-mutant mice that show an abolished circadian rhythmicity of HIF1 $\alpha$  following *Per2* deletion<sup>74</sup>. *Per2*<sup>-/-</sup> mice have significantly larger myocardial infarct sizes and are not protected by ischaemic preconditioning<sup>74</sup> — an experimental strategy that alleviates myocardial injury and which is regulated by hypoxia signalling<sup>75</sup>. Different molecular strategies have been attempted to enhance circadian rhythm signalling for the treatment of ischaemia and reperfusion injury in animal models, including agonists of purinergic receptors — such as the A<sub>2B</sub> adenosine receptor<sup>74,76</sup> — intense light therapy and blue light therapy<sup>74,77,78</sup>. Intense light therapy is associated with higher cardiac PER2 levels, increased glycolytic capacity and significantly attenuated myocardial infarct sizes in a murine model of myocardial ischaemia and reperfusion injury<sup>74</sup>. In mice with endothelial-specific deletion of *Per2*, light therapy may function to promote endothelial barrier function via light-enhanced Hif1 $\alpha$ -mediated transcription of glycolytic genes<sup>78</sup>. Similarly, in mice, carbon monoxide can protect the kidneys from ischaemia and reperfusion injury through a mechanism involving purinergic receptors, including the A<sub>2B</sub> adenosine receptor, and PER2 (REF.<sup>76</sup>). Given these observations, clinical studies that investigate pharmacological targeting of circadian signalling as therapy for ischaemia and reperfusion injury seem justified. This could be particularly promising in the field of perioperative medicine, in which patients who are undergoing elective surgery can be pretreated with circadian therapeutics before an operation to attenuate subsequent ischaemia and reperfusion injury. For example, intense light therapy before cardiac surgery could be used as an intervention to provide cardioprotection from ischaemia and reperfusion injury.

### Ischaemic preconditioning

An experimental technique (usually via repeated short episodes of ischaemia via coronary artery occlusion) to increase tolerance to the loss of blood supply, and thus oxygen, if there is a subsequent prolonged insult.

The severity of ischaemia and reperfusion injury has been linked to the molecular clock<sup>79</sup>. Most studies — but not all — report larger infarct sizes or a higher incidence of heart failure following myocardial infarction in the early morning rather than at later times in the day<sup>80,81</sup>. However, daytime variations of perioperative myocardial injury may occur in patients undergoing cardiac surgery<sup>82</sup>. In a cohort study of 596 patients who underwent

surgical aortic valve replacement in either the morning or the afternoon, patients with afternoon surgery had lower incidences of major adverse cardiac events in a 500-day follow-up period<sup>82</sup>. In addition, in 88 patients randomly assigned to morning surgery ( $n = 44$ ) or afternoon surgery ( $n = 44$ ), patients assigned to afternoon aortic valve replacement surgery had a lower level of myocardial injury marker as assessed by perioperative cardiac troponin T release (with an estimated ratio for afternoon to morning of 0.79)<sup>82</sup>. Therefore, myocardial ischaemia and reperfusion injury likely has a circadian rhythm pattern, and optimization of surgery clock time based on the circadian rhythmicity of susceptibility to ischaemia and reperfusion injury will have to be further evaluated in clinical studies, as well as pharmacological approaches to enhance circadian rhythm signalling during ischaemia and reperfusion.

## Cancer

Since the early twentieth century, studies have indicated the potential connections between mammalian circadian rhythms and cell division cycles. In 1917, Droogleever Fortuyn-van Leijden described difficulty in observing mitosis in growing tissues from a cat particularly late at night<sup>83</sup>. As of today, epidemiological and genetic studies link disruption of circadian clock functions to increased risk of several types of cancer. In the past decade, it has become clear that circadian clock components influence cell growth and transformation in a cell-autonomous manner. Earlier studies showed circadian regulation of cell proliferation and DNA damage repair<sup>84</sup>. For example, partial hepatectomy in mice led to liver regeneration marked by a night-time control of mitosis, through a mechanism by which CLOCK and BMAL1 directly regulated expression of *Wee1* mRNA<sup>85</sup>. In addition, PER1 interacts with ataxia telangiectasia mutated (ATM) serine/threonine kinase proteins and regulates checkpoint kinase 2 (CHK2)-dependent DNA damage response, which may be an evolutionarily conserved mechanism<sup>86</sup>. Furthermore, mice with an in-frame deletion of *Per2* showed a predisposition to radiation-induced tumorigenesis, and dysregulated expression of multiple genes involved in cell cycle and DNA damage response, including *Myc*, which is directly controlled by BMAL1 (REF.<sup>87</sup>).

More recent studies have further delineated a causal role of circadian disruption in tumorigenesis. A chronic jet lag paradigm (8-hour phase shift) led to reduced lifespan in mice, dysregulated gene expression and dysfunctional metabolism in the liver, which eventually led to hepatocellular carcinoma (HCC)<sup>88</sup>. The bile acid pathway is closely involved in this process, as disruption of farnesoid X receptor increased jet lag-induced HCC, whereas loss of the tumour promoter constitutive androstane receptor (CAR) inhibited hepatocarcinogenesis<sup>88</sup>. Furthermore, gene expression analysis of HCC tumour specimens by use of the program CYCLOPS showed a largely normal oscillation of core clock genes, in contrast to significant alteration of genes involved in the Janus kinase (JAK)-signal transducer and activator of transcription (STAT) pathway, apoptotic pathways and metabolic output<sup>89</sup>. Another mouse study found that both experimental jet lag and disruption of *Per2* and *Bmal1* exacerbate lung tumorigenesis<sup>90</sup>. CRY2, as a component of the FBXL3–E3 ligase complex, also promotes ubiquitination and degradation of MYC in primary mouse embryonic fibroblast by interacting with MYC phosphorylated on Thr58 (REF.<sup>91</sup>). In human cancer cell lines, such as SW480 and A549, overexpression of CRY2 reduces MYC protein



levels and inhibits cell growth<sup>91</sup>. However, in some instances, circadian genes may also aggravate cancer progression. For example, *Cry1*- and *Cry2*-double-knockout mice showed inhibition of age-adjusted tumour development and extended lifespan in the absence of tumour suppressor p53 (REF.<sup>92</sup>). In addition, a recent functional small interfering RNA screen showed that *Clock* and *Bmal1* are necessary factors for murine acute myeloid leukaemia cell growth in vitro and in vivo<sup>93</sup>. Furthermore, genetic knockout or knockdown of BMAL1 and CLOCK in patient-derived glioblastoma stem cells significantly impedes cell growth in vitro and inhibits intracranial tumour formation in mice in vivo<sup>94</sup>. Finally, the impact of polymorphisms of the circadian rhythm genes on cancer development is highly dependent on the specific polymorphism and types of cancer<sup>95</sup>. From the findings taken together, clock genes have disparate roles in different types of cancer, and further studies are required to investigate how the distinct circadian phenotypes converge on tumorigenesis.

Cancer or cancer-causing processes can reciprocally impact circadian rhythms. Gamma irradiation, a common mutagen, was found to phase-advance mouse wheel-running behaviour when administered in vivo<sup>96</sup>. Moreover, although lung adenocarcinoma has little impact on the core clock, it strongly alters the circadian transcriptome and metabolome in the liver, resulting in disrupted metabolic homeostasis of tumour-bearing mice<sup>97</sup>. A recent study further illustrated a reciprocal control of MYC on circadian rhythms. Specifically, MYC induces REV-ERB gene expression via the consensus E-box on its promoter, which subsequently dampens the expression and oscillation of BMAL1 in neuroblastoma cell lines, and alters cell metabolism, including glucose metabolism and glutaminolysis in U2OS cells<sup>98</sup>. Although it is an attractive idea that a healthy circadian clock or robust circadian rhythms facilitate tumour suppression, the specific contribution of individual components may vary depending on the cancer type and genetic background, and additional mechanistic studies will be important to further define the molecular roles of circadian rhythm signalling in tumorigenesis.

### Metabolism and metabolic disorders

Energy metabolism is tightly linked with the natural daily rhythms of sleep–fasting and wake–feeding–activity cycles. Epidemiological evidence supports a greater risk of metabolic syndrome, including increased body mass index and insulin resistance, in shift workers with sleep deprivation<sup>99</sup>. Small-cohort human studies in laboratory circadian misalignment (recurring 28-hour ‘days’ or shiftwork) settings have shown acute effects of circadian desynchronization on metabolic fitness, such as dysregulated levels of glucose, insulin and leptin in circulation<sup>100</sup>, and diminished resting metabolic rate<sup>101</sup> (BOX 1).

Mirroring these human studies, circadian disruption in mice, via either genetic or environmental means, elicits metabolic disorders<sup>102</sup>. *Clock* 19/ 19-mutant mice with severely dampened circadian wheel-running rhythms, are hyperglycaemic, with exaggerated food intake during the rest period, and are susceptible to body weight gain with a high-fat diet<sup>103</sup>. Similarly, when challenged with a high-fat diet, *Bmal1*-null mice gained more weight than wild-type mice<sup>104</sup>. As the predominant zeitgeber, light, when administered at night, led to increased food intake during the rest period, accompanied by body weight gain and impaired glucose homeostasis in mice<sup>105</sup>.

### Circadian wheel-running rhythms

When rodents have free access to a running wheel, voluntary use of this wheel is active during the night and inactive during the day to generate a specific circadian rhythm, which serves as a particularly reliable and convenient measure of the output of the master circadian clock.

The strong link between clock function and metabolic disorders has prompted vigorous efforts to understand the underlying molecular and cellular mechanisms using animal models<sup>106</sup>. The levels of NAD<sup>+</sup>, as a crucial metabolite for redox balance and a key cofactor for NAD-dependent protein deacetylase sirtuin 1 (SIRT1) and poly(ADP-ribose) polymerase 1 (PARP1), are well known to fluctuate substantially during the circadian cycle<sup>107,108</sup>. CLOCK and BMAL1 directly target the promoter of nicotinamide phosphoribosyltransferase (NAMPT) — an essential enzyme during NAD<sup>+</sup> biosynthesis that catalyses nicotinamides and 5'-phosphoribosylpyrophosphate to nicotinamide mononucleotide — via its E-box element<sup>109</sup>. SIRT1, in turn, acts on clock protein substrates, including BMAL1 and PER2, completing a regulatory loop between the clock and metabolism<sup>109,110</sup>.

The clock controls metabolism in a tissue-specific manner<sup>28,29,111</sup>. CLOCK and BMAL1 are critically required for insulin secretion from pancreatic  $\beta$ -cells<sup>112</sup>, consistent with an over-representation of genes required for insulin secretion in the pancreatic circadian transcriptome in mice<sup>31</sup>. CLOCK and BMAL1 bind to distal promoter enhancers, colocalizing with the master transcription factor PDX1 for  $\beta$ -cells<sup>31</sup>, indicating a mechanism whereby the circadian machinery cooperates with local lineage factors to define tissue-specific metabolism. Circadian metabolic regulation has also been studied in other metabolic tissues, such as liver, muscle and adipose tissues<sup>106</sup>. Liver mass and cell size fluctuate as a function of the circadian cycle in mice, which closely follows cyclic ribosome assembly and protein accumulation<sup>113</sup>.

The hypothalamus, as an energy centre in the brain, plays an important part in the circadian regulation of energy homeostasis<sup>16</sup>. The SCN projects directly or indirectly to key regulatory centres, including the arcuate nucleus, the paraventricular nucleus, the lateral hypothalamic area and the dorsomedial hypothalamus<sup>35,114,115</sup>. Many of these regulatory centres express key neurochemicals and neuropeptides important for systemic metabolism. For example, arcuate nucleus neurons express orexigenic peptides such as neuropeptide Y (NPY) and agouti-related peptide (AGRP), as well as anorexigenic peptides such as pro-opiomelanocortin (POMC) and cocaine and amphetamine-regulated transcript (CART)<sup>116,117</sup>. Furthermore, the SCN, arcuate nucleus and ventromedial hypothalamus also form a functional axis with the dorsomedial hypothalamus to control thermogenesis, sleep, corticosteroid secretion, wakefulness and food-anticipatory behaviour<sup>35,115</sup>. SCN neurons also innervate the lateral hypothalamic area, which contains neurons that express orexin, which promotes wakefulness and energy expenditure<sup>118</sup>. Consequently, disruption of orexin or its receptor in mice leads to narcolepsy and an increased risk of diet-induced obesity, consistent with the elevated body mass index in patients

with narcolepsy<sup>119,120</sup>. Taken together, these are the many examples of how circadian rhythm signalling impacts metabolism and metabolic disorders. Targeting circadian rhythm signalling pharmacologically, by alteration in sleep–wake rhythmicity, or feeding timing and behaviour is a promising approach to alter metabolic outcomes in humans.

### Immune functions and sepsis

Inflammation serves as a natural defence mechanism against invading pathogens<sup>121</sup>. Given the high energy cost, the downside of collateral tissue damage, and the risk of sepsis and tissue injury, it is important to circumvent uncontrolled or unresolved inflammatory responses<sup>122–124</sup>. The circadian clock controls the timing and duration of expression of proinflammatory cytokines, such as interleukin 6 (IL-6), tumour necrosis factor (TNF), IL-17 and growth-regulated  $\alpha$ -protein (CXCL1), as well as other aspects of inflammation or innate immune responses<sup>125,126</sup>. Mice display rhythmic sensitivity to potentially lethal challenges of bacterially derived endotoxins<sup>127</sup> and are least able to respond to challenges just before (approximately 2 hours) activity onset. However, when mice go into the active phase, when inflammatory and injurious risks are greater, several aspects of inflammatory responses, including chemoattractant levels, leukocyte trafficking, phagocytosis and proinflammatory cytokines, are upregulated<sup>128,129</sup>. Moreover, the clock governs rhythmic haematopoietic cell homing to the bone marrow and leukocyte recruitment to skeletal muscle, partly through control of oscillatory expression of the intercellular adhesion molecule 1 gene (*Icam1*) and the chemokine gene *Ccl2* (REF.<sup>128</sup>).

BMAL1 is a key circadian regulator of innate immunity. Mice with myeloid-specific deletion of *Bmal1* showed greater susceptibility to sepsis, accompanied by elevated nuclear factor- $\kappa$ B (NF- $\kappa$ B) activity and the proinflammatory miR-155 (REF.<sup>130</sup>). In Ly6C<sup>hi</sup> monocytes, BAML1 rhythmically recruits a transcription co-repressor, polycomb repressive complex 2 (PRC2), to inhibit chemokine gene expression, which results in oscillatory trafficking to the inflammatory sites<sup>131</sup>. Moreover, BMAL1 regulates expression of the chemokine CXCL5 in epithelial club (Clara) cells, partly via regulation of binding of the glucocorticoid receptor to the CXCL5 promoter, which in turn controls neutrophil trafficking to the mouse lungs<sup>132</sup>. Other circadian proteins also have important roles. For example, REV-ERB $\alpha$  regulates transcription of inflammatory genes in macrophages, and mice deficient in REV-ERB $\alpha$  expression fail to control rhythmic IL-6 secretion following lipopolysaccharide challenge<sup>129</sup>. Therefore, the clock serves as a master gatekeeper of innate immunity, optimizing immune outcomes and the subsequent resolution and repair to avoid an exaggerated response and sepsis shock.

#### Lipopolysaccharide challenge

An experimental technique to elicit inflammation by administering lipopolysaccharide via intraperitoneal injection to mice.

Recent studies have established that adaptive immune cells are subjected to circadian control<sup>125</sup>. Several clock components play a critical part in lymphocyte development. Nuclear receptor ROR $\gamma$ t (also known as ROR $\gamma$ 2, an isoform of ROR $\gamma$ , which has a

different, shorter N terminus) is a master regulator for the development of IL-17-producing T helper cells (T<sub>H</sub>17 cells), an important immune cell type for autoimmunity<sup>133</sup>. NFIL3 serves as a repressor of ROR $\gamma$ t, thus functioning in a new circadian loop to regulate T<sub>H</sub>17 cell differentiation<sup>134</sup>. Similarly to innate immunity, trafficking and tissue recruitment of lymphocytes are controlled by the circadian clock. For example, both T cells and B cells oscillate in number in circulation, peaking during the rest phase<sup>135</sup>. Glucocorticoids, which are regulated by the central clock in the brain, serve as cell-extrinsic signals to govern the expression of the chemokine CXCR4, leading to the diurnal oscillation of circulating T cells in mice<sup>136</sup>. In addition, CLOCK and BMAL1 regulate the rhythmic expression of Toll-like receptor 9 (TLR9) via non-canonical E-box elements, and circadian rhythms of TLR9 strongly influence septic severity and immunization response<sup>137</sup>. For example, the septic severity in a mouse model of caecal ligation and puncture was elevated when the caecal ligation and puncture procedure was performed in entrained animals at zeitgeber time 19 (peak of TLR9 expression in the spleen) compared with zeitgeber time 7 (nadir TLR9 expression). Molecular pathways of circadian rhythm signalling could be targeted in patients with inflammatory or infectious diseases, for example in the context of critical illness such as sepsis<sup>138</sup>. Observational studies of windowed rooms and real-time use of ambient light in critical care units indicate that physiological light–dark patterns may support recovery from critical illness<sup>139</sup>.

#### Caecal ligation and puncture

A commonly used preclinical rodent model to study sepsis via ligation and perforation of the caecum, resulting in polymicrobial infection and systemic inflammation.

#### Zeitgeber time

A standardized 24-hour notation for the phase in an entrained circadian cycle with reference to environmental regularities or zeitgebers.

### Ageing and circadian rhythms

Besides its role in disease pathogenesis, the molecular clock output gained increasing attention from studies that focused on the physiological impact, such as in natural ageing. There has been growing evidence of the interaction between natural ageing and the alternations in circadian rhythmicity<sup>54</sup>. For example, aged mice maintain behaviour and cellular circadian rhythmicity in epidermal and muscle stem cells, but aged muscle stem cells show a reprogrammed oscillating transcriptome<sup>140</sup>. This reprogrammed oscillating transcriptome in aged mice is rewired to control expression of genes implicated in stress responses, rather than homeostasis. Similarly, ageing reprogrammes the circadian output in mouse liver, resulting in significantly reduced oscillation in protein acetylation<sup>141</sup>. In both scenarios, calorie restriction could reverse the ageing-related reprogramming, indicating a potential favourable impact of calorie restriction. Furthermore, from a therapeutic point of view, treatment with the NAD<sup>+</sup> precursor nicotinamide riboside deacetylates PER2 at Lys680 to improve genome-wide BMAL1 chromatin binding<sup>142</sup>. The modulation of

circadian rhythm by NAD<sup>+</sup> precursors restores the ageing-related reprogramming of circadian rhythmicity in the liver of old mice. Therefore, reprogramming of the circadian output associated with natural ageing can potentially be targeted by behaviour modification with calorie restriction and therapeutic interventions such as NAD<sup>+</sup> precursors.

Circadian rhythm genes also have a profound impact on the ageing process. A study using transgenic mice with modifications in critical circadian rhythm genes indicated that circadian rhythm genes affect the lifespan and contribute to ageing-related disease<sup>143</sup>. For instance, *Bmal1*-deficient mice show premature ageing indicated by reduced lifespan, with more than 90% of animals dying within 1 year, whereas wild-type mice have an average lifespan of 28 months<sup>143</sup>. *Bmal1* deficiency also results in ageing-related skin and eye disease accompanied by increased levels of reactive oxygen species in the kidney and spleen<sup>143</sup>. Moreover, SIRT1 and PER2 regulate each other reciprocally in circadian rhythm and ageing in mouse liver and human hepatocytes<sup>144</sup>. SIRT1 is a key NAD<sup>+</sup>-dependent deacetylase, and the absence of *Sirt1* leads to overexpression of *Per2* via acetylation of histone H4 on the promoter. On the other hand, *Per2* overexpression reduces *Sirt1* expression in mouse liver, causing increased levels of senescence markers, including the genes encoding phosphoenolpyruvate carboxykinase (*Pck1*) and glucose 6-phosphatase (*G6pc*)<sup>144</sup>. Finally, previous studies indicated that disruption of the sleep–wake cycle and disruption of circadian rhythm are risk factors for ageing-related diseases such as Alzheimer disease<sup>145</sup>. In summary, the relationship between ageing and the circadian rhythm is truly intricate, and direct targeting of the circadian rhythm genes could potentially reduce or slow down ageing-related disease.

## Targeting circadian rhythm in disease

### Circadian pharmacology

Circadian changes in drug pharmacokinetics, including absorption, distribution, metabolism and excretion, are well documented<sup>10</sup>. The term ‘chronotherapy’ usually refers to the timed dosing of drugs to maximize therapeutic index<sup>10</sup>. Various cardiovascular parameters display strong circadian rhythms<sup>17,50</sup>. In humans, blood pressure starts to rise before waking in the early morning, then peaks soon afterwards. However, 40% of patients with essential hypertension (without known causes) are non-dippers, which means that their blood pressure does not drop during night-time. Thus, several chronotherapies, such as bedtime intake of antihypertensive drugs — for example angiotensin-converting enzyme inhibitors or angiotensin receptor blocker — have shown increased cardiovascular efficacies in hypertensive patients by improving the nocturnal blood pressure profile<sup>50,146</sup>. In rodent models, 35 anticancer agents display a twofold to tenfold variation in tolerability in mice or rats when given at different times of the day<sup>10</sup>. These studies have motivated clinical trials in patients with cancer<sup>10,147</sup>, including assessing a combination therapy in which doxorubicin administered at 6 a.m., with cisplatin given 12 hours later, markedly increased survival of patients (a fourfold survival advantage) with advanced-stage ovarian cancer compared with patients who were given the same dose of the drugs at 6 p.m.<sup>147</sup>. The cytotoxic effects of cyclophosphamide correlated with the functional status of the CLOCK–BMAL1 complex, and the ablation of CRYs rendered tumours more resistant to the drug compared with

tumours in wild-type mice<sup>148</sup>. Chronotherapy with glucocorticoid receptor agonists has also been extensively applied to asthma<sup>51</sup>, and dosing of prednisone in the afternoon (3 p.m.) showed a much-increased efficacy in the management of night-time symptoms compared with prednisone administered in the morning or later in the evening<sup>149</sup>.

### Circadian rhythm sleep–wake disorders

**Insomnia.**—Approximately 25% of Americans develop insomnia each year, and up to 10% of them progress towards chronic insomnia, which is frequently associated with anxiety, depression, higher risk of cardiovascular disease, physical health problems and poor quality of life<sup>150</sup>. Owing to the close association between melatonin deficit in pathological conditions and natural ageing, and increased insomnia prevalence, a simple melatonin replacement therapy, as a chronobiotic, was first believed to reduce insomnia<sup>151,152</sup>. A randomized, double-blind placebo trial involving 791 adult outpatients (aged 18–80 years) with primary insomnia showed that prolonged-release melatonin (such as Circadin) reduced subjective sleep latency only in patients older than 65 years, demonstrating its specific efficacy in the older adult group but not in other age groups<sup>151</sup>. Particularly, Circadin, 2 mg per day orally, was recommended to increase sleep continuity and improve daytime wellness in insomnia disorder in adults aged 55 years or older in Europe<sup>152,153</sup> (FIG. 4a). As an indispensable component of the melatonergic signalling system, melatonin agonists have been studied extensively as pharmaceutical approaches for treatment of insomnia. Ramelteon, a high-affinity MT<sub>1</sub> and MT<sub>2</sub> receptor agonist (with an affinity approximately ten times higher than that of melatonin), has been approved for treating insomnia disorders characterized by difficulties in sleep onset in the United States and Japan<sup>153</sup> (FIG. 4a; TABLE 1). However, the effects of both exogenous melatonin and ramelteon on total sleep duration and maintenance are not so clear<sup>154</sup>. Notably, mice with deletion of *Mtnr1a* have a selective disruption of REM sleep and an increase of NREM sleep, whereas *Mtnr1b*-knockout mice have a selective disruption of NREM sleep<sup>155</sup>. These results highlight the opposing effects of the two receptors, which partially explain why melatonin or ramelteon only reduces sleep latency, without affecting total sleep time or sleep architecture. Thus, discovery of selective melatonin receptor agonists may promote the development of novel and more efficacious therapeutic agents.

#### Chronobiotic

An agent that is able to influence, directly or indirectly, the phase and/or the period of the body clock.

In turn, dual orexin receptor antagonists can enhance sleep maintenance and onset by attenuating cortical arousal and reducing the threshold of sleep state transitions associated with OX2R and OX1R inhibition, respectively<sup>156,157</sup>. Suvorexant, an orally active dual orexin receptor antagonist, was first approved by the FDA in 2014 for the treatment of insomnia marked by difficulty with sleep onset and/or sleep maintenance<sup>158</sup> (FIG. 4a; TABLE 1). Similarly to natural sleep, the preservation of night-time arousability to salient cues<sup>159</sup> and daytime cognitive performance<sup>160</sup> following administration of dual orexin receptor antagonists in experimental animals (monkey and rat, respectively), suggests further

advantages over conventional therapies such as benzodiazepines. From the findings taken together, suvorexant is more effective in treating sleep maintenance insomnia than sleep-onset difficulties.

**Jet lag and shiftwork.**—Mistimed circadian rhythms due to rapid travel across multiple time zones, together with the sleep deprivation from long flights, are the main contributing factors for jet lag symptoms<sup>161</sup>. The combination of hypnotic and phase-shifting properties of melatonin makes it an ideal drug to treat jet lag<sup>162</sup>. As discussed in the previous section, different melatonin receptors have different distributions and functions. Notably, animal studies revealed that melatonin phase-shifts the wheel-running activity rhythms, and this effect is diminished by the MT<sub>2</sub> receptor antagonist 4P-PDOT<sup>41</sup>. In a jet lag model, re-entrainment after the phase advance was significantly slower in mice lacking functional MT<sub>2</sub> receptors than in melatonin-proficient mice with intact MT<sub>2</sub> receptors<sup>43</sup>. Thus, the MT<sub>2</sub> receptor<sup>38,40,41</sup> is believed to be the main mediator of the re-entrainment effect of melatonin on the SCN<sup>163,164</sup> and the locomotor activity<sup>165</sup>. As the key to treating jet lag is to mimic the phase-shifting properties of melatonin, many studies have focused on phase-specific activation of melatonin receptors. Administration of tasimelteon, a melatonin receptor agonist with a higher affinity for the MT<sub>2</sub> receptor, resulted in a significant shift in endogenous circadian rhythms and an improvement in sleep initiation and maintenance in phase II ( $n = 39$ ) and phase III ( $n = 412$ ) trials involving healthy volunteers exposed to experimentally induced jet lag<sup>166</sup> (FIG. 4b; TABLE 2). The recent determination of the crystal structure of melatonin receptors<sup>167</sup> has led to the discovery of two MT<sub>1</sub> receptor-selective inverse agonists (UCSF7447 and UCSF3384)<sup>168</sup>, which, when administered at subjective dusk, advanced the onset of the running-wheel activity in mice in constant darkness. Furthermore, their effects on circadian phase advance were eliminated in *Mtnr1a*-knockout mice but not in *Mtnr1b*-knockout mice, indicating the impact of the MT<sub>1</sub> receptor in the phase-shifting function of melatonin<sup>168</sup>. Therefore, it is still unclear whether the MT<sub>1</sub> receptor is more important than the MT<sub>2</sub> receptor for the phase-shifting effects of melatonin under constant darkness, and clinical trials are needed to investigate their effectiveness of specific melatonin receptor agonists in real-life jet lag situations.

Most importantly, the timing of melatonin administration is critical for optimal therapeutic effects<sup>169</sup> (FIG. 4b). Melatonin phase-advances the physiological and behavioural circadian rhythm when administered in the late biological afternoon, whereas it delays the circadian clock when administered in the early biological morning<sup>170,171</sup>. Circadian status predicted by the timing of habitual sleep in the adapted departure time zone can be used to manage melatonin administration and begin shifting the circadian system in the desired direction even before a flight<sup>172</sup>: to delay the circadian system before westward travel, melatonin should be taken 1 hour after rising from bed (around 6 a.m.); to advance the circadian system before eastward travel, melatonin should be taken 6.5 hours before bed (around 6 p.m.). Similarly, melatonin could be used to shift the timing of the circadian system to the destination time zone during/after flights<sup>172</sup>. Although not approved for the treatment of jet lag, melatonin is still used to reduce jet lag symptoms following transmeridian flights<sup>161</sup>.

Similarly to jet lag, circadian misalignment also triggers shiftwork disorder<sup>173</sup>. Chronic circadian misalignment in shift workers can result in serious sleep disruption and a variety

of associated health issues, including increased risk of several cancers (such as breast cancer), metabolic syndrome (prone to weight gain) and cardiovascular disease<sup>100,174,175</sup>. There is low-quality evidence that melatonin is associated with increased sleep length after a night shift but not with reduced sleep latency time<sup>176</sup>. Limitations exist in that most trials were small with a duration of 1 week or less<sup>176</sup>. Besides melatonin, other phase-shifting chemicals might also be useful for promoting circadian adaptation to environmental changes. Administration of sildenafil, an inhibitor of cGMP-specific phosphodiesterase 5, significantly accelerates the entrainment of locomotor activity rhythms to the new light–dark cycle in hamsters<sup>177</sup>. In addition, a recent study showed that pharmacological perturbation of RuvB-like ATPase 2 (RUVBL2) by the adenosine derivative cordycepin shifts the circadian phase both in human cells and mouse tissue, and rapidly entrains the circadian phase in a mouse jet lag model. This phase-shifting effect is the result of disassembly of interaction between RUVBL2 and BMAL1, thereby leading to the release of repression of clock gene transcription<sup>178</sup>. Furthermore, the treatment of shiftwork focuses on combating the reduced alertness during night shifts. For instance, modafinil (200 mg) and armodafinil (150 mg), the dopamine reuptake inhibitors which indirectly activate the release of orexin and histamine, were associated with increased alertness and reduced sleepiness in shift workers. However, these drugs were also reported to lead to headache and nausea<sup>176</sup>. Therefore, the long-term benefits and adverse effects of melatonin and new-found phase-shifting compounds, in combination with wake-promoting medications, need to be further evaluated by clinical studies in patients with shiftwork disorder.

Therefore, for the fatigued flyer or shift worker in a state of internal and external desynchrony, the best strategy may still be the combination of timed medication, sleep and timed light treatment<sup>161</sup>.

### Targeting the circadian machinery

In addition to manipulating circadian output rhythms including feeding–fasting and sleep–wakefulness, direct manipulation of the core oscillator is an exciting new dimension to circadian medicine. Recent studies have aimed to develop candidate drug molecules targeting core components or the molecular oscillator (FIG. 2; TABLE 2).

**Drugs targeting REV-ERBs.**—REV-ERB $\alpha$  is a major modulator of the circadian expression of BMAL1, and the secondary feedback loop in the clock cycle. Ligands targeting REV-ERBs have been assessed in a broad range of disease models (TABLE 2). For example, GSK4112 — a small-molecule agonist of nuclear receptor co-repressor 1 (NCOR1) of REV-ERB $\alpha$ <sup>179</sup> — showed inhibitory effects against lipopolysaccharide induction of IL-6, CXCL11, CXCL6 and CCL2 in human monocyte-derived macrophages and the leukaemia-derived cell line THP1 (REFS<sup>129,180</sup>). Subsequently, two pyrrole compounds, SR9009 and SR9011 (agonists of REV-ERB $\alpha$ ), were derived, and showed strong *in vivo* activities, including metabolic regulation, anticancer and anxiolytic effects<sup>181–183</sup>. Mice treated with these compounds experienced acutely suppressed wheel-running behaviour and improved metabolic homeostasis in diet-induced obesity<sup>181</sup>. The compounds (SR9009 and/or SR9011) have also been assessed in other metabolic settings, including enhancement of mitochondrial content in skeletal muscle and exercise endurance



in mice<sup>184</sup>. In addition, SR9009 and SR9011 display selective cytotoxicity with regard to leukaemia and several solid tumours with different tumorigenesis drivers<sup>182</sup>. SR9009 down-regulated key autophagy-related mRNAs in primary hepatic stellate cells, suggesting an inhibitory role in autophagy<sup>185</sup>, a clock-regulated process that cancer cells use to meet heightened metabolic requirements. SR9011 also displayed anxiolytic effects<sup>183</sup>, consistent with genetic evidence showing enhanced anxiety in *Nr1d2*-knockout mice. In comparison, SR8278, a REV-ERB antagonist, attenuated anxiety when administered to mouse ventral midbrain<sup>186</sup> and ameliorated myocardial injury in an isolated heart system when administered at the activity onset<sup>82</sup>, indicating beneficial roles of REV-ERB inhibition. Together, these studies highlight a versatile yet complex role of targeting REV-ERBs in different diseases.

**Drugs targeting RORs.**—A few studies have also investigated compounds that enhance circadian amplitude, dubbed ‘clock amplitude-enhancing small molecules’<sup>187</sup>. One of the best-studied clock amplitude-enhancing small molecules is the natural flavonoid nobiletin, which was found to activate RORs directly in mice<sup>188</sup>, reaffirming a key role of the secondary stabilization loop in circadian amplitude regulation. Indeed, nobiletin potentiated circadian wheel-running activity and clock protein oscillation, reduced body weight and restored robust energy homeostasis in diet-induced obesity mice and *db/db* diabetic mice<sup>188</sup>. Other ROR modulators also have roles in metabolic diseases<sup>189</sup>. An inverse agonist of RORs, SR1555, was found to enhance energy expenditure and ameliorate obesity in a diet-induced obesity mouse model<sup>190</sup>. These results are reminiscent of the shared anxiolytic efficacies of SR9011 and SR8278, suggesting plasticity of circadian modulation against diseases. In addition to circadian amplitude-enhancing and metabolic regulatory effects, drugs targeting RORs can modulate numerous autoimmune disorders<sup>189,191</sup>. Furthermore, with canonical ligand-binding domains shared by nuclear receptors, RORs are bound by various endogenous and synthetic ligands<sup>189</sup>. For instance, digoxin and ursolic acid, as ROR $\gamma$ t inverse agonists, ameliorated autoimmune disorders including arthritis and encephalomyelitis via suppression of T<sub>H</sub>17 cell differentiation<sup>191,192</sup>. Consistently, several optimized ROR $\alpha$ /ROR $\gamma$  ligands attenuated expression of downstream cytokines and strongly alleviated autoimmune disease symptoms in mice (such as in animal models of multiple sclerosis and experimental autoimmune encephalomyelitis)<sup>192,193</sup>. Finally, the regulatory role of ROR $\gamma$ t in T<sub>H</sub>17 cell differentiation is governed by a circadian pathway that involves REV-ERBs and BMAL1 (REF.<sup>134</sup>), which supports the notion that these drugs modulate the circadian system against immune diseases. In conclusion, the data obtained from the in vivo ROR synthetic ligand studies are extremely encouraging, specifically for the regulation of circadian rhythms, metabolism and immunity. However, how these ligands affect the pathophysiology of the diseases has yet to be extensively studied.

**Drugs targeting PERs and CRYs.**—Several chemical screens have found that compounds that affect levels of PERs and CRYs can alter circadian periods. For example, a CRY1- and CRY2-stabilizing carbazole compound, KL001, was identified in a cell-based screen to lengthen the period and reduce the amplitude<sup>194</sup> (TABLE 2). Furthermore, evolutionarily conserved domains in both CRY1 and CRY2 can be targeted as binding surfaces for drug development. Mammalian CRY proteins belong to the evolutionarily

conserved photolyase protein family, and their photolyase homology region functions as a low-affinity FAD-binding pocket<sup>63</sup>. As illustrated by the compound KL001, which competes with the FBXL3 carboxy terminus at the FAD-binding pocket to stabilize CRYs<sup>195</sup>, even such a relatively shallow, suboptimal binding surface can mediate functional binding with chemical ligands. Besides circadian period-regulating function, KL001 derivative also has a role in metabolic regulation by normalizing glucose tolerance in diet-induced obesity mice<sup>196</sup>. Moreover, combination of KL001 and the REV-ERB agonist SR9011 inhibits cell cycle progression and reduces the self-renewal potential of patient-derived glioblastoma stem cells<sup>94</sup>. Notably, SHP656, one of the most promising KL001 derivatives, specifically inhibits the growth of patient-derived glioblastoma stem cells in vitro at a concentration of 2–30  $\mu\text{M}$ , while having an insignificant impact on differentiated glioblastoma cells or non-malignant epilepsy-derived neural cells<sup>94</sup>. SHP656 also shows improved in vivo safety and increased in vivo stability, and significant efficacy in the treatment of glioblastoma by reducing tumour growth and prolonging mouse survival<sup>94</sup>. Although the evolutionarily conserved features of CRY domains and the CRY FAD-binding pockets offer potential as targets for in silico screening or derivation of known ligands, this high sequence conservation between homologous proteins makes it particularly challenging to develop isoform-selective compounds. Recently, KL101 and TH301 were identified to stabilize CRY1 and CRY2, respectively, which is dependent on the disordered carboxy-terminal region outside the pockets, leading to enhancement of brown adipocyte differentiation in mice<sup>197</sup>. Finally, structural studies of clock proteins have pinpointed interaction hotspots with disproportionately high binding energy. For example, the PER–CRY crystal structure revealed several regulatory interaction surfaces on the CRY-binding domain of PER<sup>198</sup>, particularly in a region that competes with FBXL3 to bind to the extreme carboxy-terminal surface on CRY.

**Drugs targeting CK1.**—CK1 kinases are excellent pharmacological targets as they have a pivotal role in phosphorylating core clock proteins<sup>18</sup>. Although early inhibitors — including IC261 and CKI-7 — were only moderately active and selective, a specific inhibitor of CK1 $\epsilon$ , PF-4800567, was derived to confer an inhibitory activity greater than 20-fold over CK1 $\delta$ <sup>199</sup>. This ‘super’ strong inhibition has shown additional therapeutic promises in mice. For example, in mice rendered arrhythmic either by constant light exposure or by deletion of vasoactive intestinal polypeptide receptor 2 (*Vipr2*<sup>-/-</sup> mice), the CK1 $\delta$ -specific inhibitor PF-670462 was able to induce behavioural rhythms<sup>200</sup>, suggesting that a pharmacological agent that targets CK1 $\delta$  can be tested in patients with familial advanced sleep phase syndrome or animal models. Moreover, from high-throughput chemical screening, a novel compound called ‘longdaysin’, which is a purine derivative, was discovered to lengthen the circadian period in cell culture in vitro and in zebrafish in vivo by targeting CKI $\delta$ , CKI $\alpha$  and ERK2 (REF.<sup>201</sup>). Longdaysin could potentially serve as another promising novel compound for the treatment of circadian rhythm disorders.

### Environmental and lifestyle modifications

Environmental and lifestyle regimens are known to alter circadian timing and various clinical trials are ongoing to test their effects on health (TABLE 3). Blue light is a dominant zeitgeber, and the pervasive night-time light exposure in modern societies interferes with

neuroendocrine circuits to perturb sleep and dampen rhythms of melatonin and cortisol<sup>5</sup>. Conversely, bright light therapy, commonly conducted by exposure to broad-spectrum bright light (2,000–10,000 lux) for 1–3 hours in the morning, can improve sleep–wake cycles, mood and cognitive functions in patients with sleep or seasonal affective disorders<sup>202,203</sup>. In particular, in intensive care units — where patients suffer more from circadian arrhythmia due to both external (environmental) and internal (medical) factors — several randomized clinical trials showed that morning bright light can reduce delirium prevalence and shorten delirium episodes<sup>139</sup>. Therefore, safe, non-invasive chronotherapeutic interventions, such as light therapy, may be effective in maintaining and resynchronizing circadian rhythmicity and improve the short-term and long-term outcomes.

A number of lifestyle modifications, including timed meals, sleep and activity, have been investigated for their safety and their efficacy to achieve optimal circadian health<sup>5</sup>. Exciting advancements have been reported in recent years with regard to feeding regimens. For example, time-restricted feeding (TRF), characterized by food access for 8–9 hours during the active phase, has been found to exert strong preventive and therapeutic effects in metabolic disorders<sup>204,205</sup>. In mouse studies in which the overall caloric intake is largely constant between TRF and ad libitum feeding, considerable metabolic benefits, including energy balance, insulin sensitivity and reduced hepatic steatosis, were observed in the TRF group, independent of diet or initial body weight<sup>205</sup>. There was also a legacy effect in which benefits were maintained when TRF was applied for only 5 days a week and ad libitum feeding was applied for the remaining 2 days, mimicking the workdays and weekends in a human lifestyle. Furthermore, TRF showed the capacity to not only stabilize but also reverse the progress of metabolic diseases in mice with pre-existing obesity and type 2 diabetes<sup>205</sup>. Consistent with the notion that eating during the inactive phase diminishes energy expenditure and exacerbates weight gain, pilot human studies also showed that when total intake is maintained at a similar level, participants consuming a larger portion during breakfast showed a strong improvement in metabolic homeostasis in comparison with the calorie-rich dinner group<sup>206</sup>. In a randomized, crossover clinical study, an early TRF regimen, involving a 6-hour feeding period earlier in the day, was found to increase key insulin sensitivity and improve  $\beta$ -cell function in prediabetic men<sup>207</sup>. Conversely, patients with night-eating syndrome are prone to obesity and type 2 diabetes<sup>208</sup>. Such energy imbalance as a result of abnormally timed food consumption may explain metabolic dysregulation from nocturnal sleep deprivation or late sleep. In a human study in which a 5-day sleep deprivation was implemented to mimic a workweek, an increase in energy expenditure was observed in study participants, which led to more exaggerated energy intake, resulting in energy storage and bodyweight gain<sup>209</sup>.

Besides TRF, temporally controlled activity has also been studied in laboratory and clinical settings. Studies in hamsters showed that stimulated activity during the rest phase caused phase advance in activity onset the following day<sup>210</sup>, perhaps attributable to acute clock gene differential regulation in the SCN<sup>211</sup>. In aged mice, scheduled wheel activity also enhanced behavioural rhythmic amplitude and entrainment<sup>212</sup>. Likewise, in humans, exercise and activity can reset circadian melatonin rhythms<sup>213</sup>, and afternoon exercise has been found to improve sleep quality in older people<sup>214</sup>. Thus, it is conceivable that scheduled sleep routines can also improve circadian rhythms. Indeed, in a mouse model of

Huntington disease, induction of sleep by alprazolam (a benzodiazepine) during the inactive period improved cognition and mobility<sup>215</sup>.

## Human rhythm determination

Intrinsic circadian rhythms in humans are slightly over 24 hours, and manifest themselves as morning or evening chronotypes. Several questionnaires have been developed to assess human chronotypes<sup>216</sup>. For example, the self-assessment Morningness–Eveningness Questionnaire has shown usefulness in determining chronotypes and uncovering outliers by evaluating the circadian variation of oral temperature. People with morning types have a higher daytime temperature and lower post peak temperature than people with evening types<sup>217</sup>. The Munich Chronotype Questionnaire refined the method to distinguish workdays versus rest days, and collected detailed information on actual schedules of sleep and activity, as well as time spent outdoors<sup>218</sup>. Electronic devices can also monitor behavioural rhythms, including wrist actigraphy sensors and mobile phone apps<sup>219,220</sup>. Physiological markers that can be routinely assessed in the clinic include circulating hormones (melatonin and cortisol), core body temperature and midpoint sleep. As surrogates, human fibroblast primary cultures<sup>221</sup> and hair follicle cells<sup>222</sup> serve as minimally invasive in vitro systems to report behavioural rhythms. In clinical settings, common difficulties in assessing behavioural rhythms accurately include lack of information on the time of sample collection or obtaining a single time point — rather than a longitudinal — biopsy. To overcome these challenges, innovative statistical methods can assign circadian time or phase for human-derived samples, including blood metabolites<sup>223</sup>, gene expression in tumours<sup>89</sup> and post-mortem brain samples<sup>224</sup>.

### Chronotypes

Characterized by individual differences in the timing of sleep–wake schedules, biological parameters (such as core body temperature, melatonin and cortisol) and cognitive performance (such as attention).

### Circadian time

A notation for the phase in a circadian cycle that represents an organism's endogenous circadian clock, without reference to any environmental regularities or zeitgebers.

Most of the 662 currently registered clinical trials evaluating circadian-related compounds in different phases of the pipeline involve small-size cohorts and simply investigate circadian rhythms as an observational readout. Behavioural and physiological parameters, including sleep, activity, and melatonin and cortisol, are most commonly used to assess circadian rhythms. In some cases, expression of clock genes was also determined by use of skin biopsy samples. Among those that involve circadian intervention, the predominant regimens are bright light therapy, timed food intake and melatonin therapy (TABLES 1, 3). A meta-analysis showed that light therapy was effective in treating circadian rhythm sleep disorders, specifically insomnia<sup>225</sup>. Nevertheless, most studies are small to medium size. The timing

of food intake showed a more profound influence on the peripheral circadian rhythms than the timing of light<sup>226</sup>. In addition, melatonin therapy seems to be an encouraging treatment for sleep problems, including insomnia and other wake–sleep disorders<sup>176,227</sup>. However, the beneficial effects of melatonin supplements were often minor in humans, and there is a lack of high-quality, randomized controlled trials of melatonin therapy in circadian and sleep disorders published in the literature<sup>169</sup>. Other agents include orexin signalling modulators and vitamin B (TABLE 1). Although there are no registered clinical trials ongoing for agents that target the core clock factors, small human studies have been reported for some natural compounds. For example, a Japanese study of six cohorts of older patients revealed a preventive effect on the progression of the cognitive impairment associated with Alzheimer disease following prolonged intake of citrus extracts rich in the clock modulator nobiletin<sup>228</sup>. Given a causal role of perturbation in sleep and circadian rhythms in amyloid aggregation and cognitive abnormalities<sup>229</sup>, it will be interesting to investigate whether circadian enhancement can alleviate symptoms of Alzheimer disease in larger cohorts.

## Conclusions

The omnipresent circadian control permeates physiology and pathophysiology. This current growing interest in circadian rhythm signalling could propel discoveries of new circadian therapeutics and their functions and mechanisms in preclinical and clinical settings. Although existing studies in disease models illustrate the promising potential of targeting circadian rhythms as a therapeutic strategy, they also highlight the complex or even seemingly contradictory effects. These results are consistent with the notion that owing to integrated circadian feedback regulation, there may be an interventional range, in either direction, to manipulate circadian clocks and alter pathophysiology in a chronic setting. Importantly, although the levels of some circadian modulators are determined solely by light input, melatonin, as a typical representative, could show the opposite physiological impact: sleep-promoting effects in humans and awakesness-promoting effects in nocturnal animals. Thus, the difference in diurnal (humans) and nocturnal (rodents) species needs to be taken into careful consideration during circadian rhythm research.

Extending the preclinical efficacy of pharmacological manipulation of circadian rhythms to clinical trials with highly validated compounds — by themselves or in combination with existing drugs — will offer exciting insight to guide further mechanistic investigation and drug development efforts. With the remarkable elucidation of circadian expression of most protein targets for drugs<sup>29</sup>, a re-evaluation of circadian pharmacology for desired drugs, as well as discovering behaviour changes to modify the circadian output, will be a critical step for bringing circadian rhythm signalling into routine clinical practice. Besides serving as a treatment for circadian rhythm disorders, therapeutic targeting of circadian rhythm genes and modifying the circadian outputs could potentially be of benefit in other diseases. For instance, as demonstrated in a recent study of the relevance of daytime during cardiac surgery<sup>82</sup>, a straightforward chronotherapeutic application can markedly modify therapeutic outcomes and disease prognosis. Therapeutic targeting of the circadian rhythm pathway could also be beneficial in intensive care units, as patients in intensive care units often experience disruption in the circadian rhythm<sup>230</sup>, although more studies are required to dissect the mechanism and impact of circadian disruption. Overall, we believe that the

progress made in this area — which was recognized with the Nobel Prize in Physiology or Medicine in 2017 for discoveries on how internal clocks govern human biology — indicates that it is a timely and exciting challenge to harness the power of timing in disease intervention and drug discovery for the treatment of human diseases.

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**Box 1****Circadian parameters on disease progression**

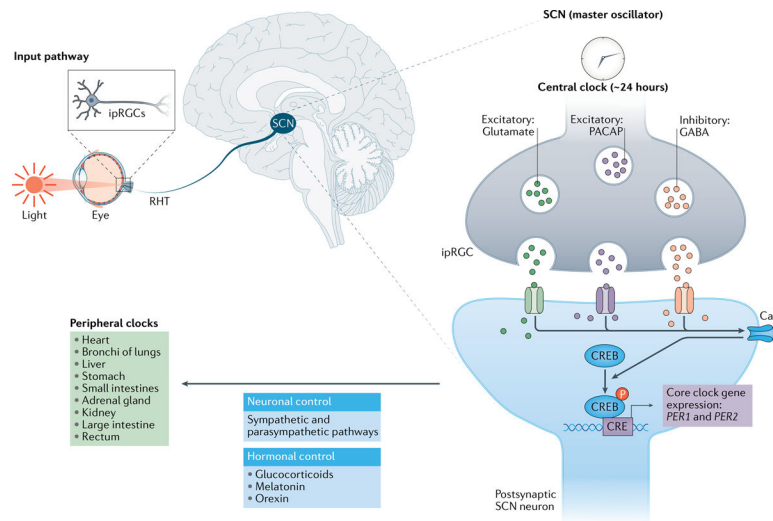
Numerous studies show that alterations of cardinal circadian parameters — period, phase and amplitude — are not only risk factors for diseases but are also closely related to disease progression. The difference between peak and trough values is the amplitude, and the time interval between two peaks or two troughs is called the ‘period’. The phase is defined as the timing of a reference event relative to a fixed activity (such as the beginning of daytime). Shiftwork, characterized by circadian misalignment (mainly phase misalignment), serves as a good example. Prevalent health problems can arise from this chronic circadian rhythm disruption, including sleep–wake and metabolic disorders, increased risk of several cancers (such as breast cancer) and cardiovascular disease<sup>100,175,243</sup>. In addition to phase shift, experimental animal models<sup>103</sup> and epidemiological data<sup>244</sup> also indicate a pivotal role of dampened circadian amplitude in metabolic disorders. *Clock 19/19* mice — which have a metabolic syndrome of hyperlipidemia, hepatic steatosis, hyperglycaemia and hypoinsulinaemia — show reduced levels of expression of circadian genes and diurnal feeding rhythm<sup>103</sup>. Furthermore, circadian dysregulation and cancer progression are also interdependent. Disturbance of circadian rhythm either locally or systemically is associated with reduced DNA repair, increased cancer cell proliferation and growth in vitro<sup>245</sup>, and higher cancer incidence and poor prognosis in vivo<sup>175,246</sup>. In mice, both physiological disruption (experimental jet lag) and genetic disruption (mutations in *Per2* and *Bmal1*) of the circadian machinery result in decreased survival and lung tumour growth and progression<sup>90</sup>. In addition, the tumour itself and its concomitant symptoms — such as acute or chronic pain, discomfort, or anxiety — may gradually disrupt a patient’s sleep, which in turn can result in disruption of endocrine and immunological rhythms formerly linked with cancer progression<sup>247,248</sup>. Circadian rhythm disturbances, in addition to being common symptoms of many neurodegenerative disorders, might also exacerbate the progression of disease. Recently, several longitudinal studies showed that attenuated amplitude and phase shifts of rest–activity rhythm even precede the clinical symptoms of Alzheimer disease and are associated with increased dementia risk, and that long daytime sleeping precedes the risk of Parkinson disease<sup>249</sup>.

Small-molecule modulators targeting clock components are emerging as powerful tools for understanding mechanisms of clock-related disorders as well as developing putative therapeutic strategies. For example, stabilization of period circadian protein homologues (PERs) by casein kinase 1 (CK1) inhibitors<sup>201</sup> or cryptochrome (CRY) by small-molecule circadian modulators<sup>194</sup>, respectively, could lengthen the circadian period. Moreover, the CRY stabilizer KL001 lengthens the period and reduces the amplitude<sup>192</sup>. In turn, several selective inhibitors of glycogen synthase kinase 3 $\beta$  (GSK3 $\beta$ ) can shorten the circadian period by reducing phosphorylation of both PER2 and CRY2 (REF.<sup>187</sup>). Early induction of *Per1/Per2* via the cAMP/cAMP-response element-binding protein (CREB) signalling pathway is involved in acute phase resetting in mammals<sup>187</sup>. Thus, both cAMP-inducing compounds and inhibitors of the cAMP-catabolizing enzyme phosphodiesterase 4 could result in phase delay by immediate *Per1/*

*Per2* induction<sup>187</sup>. In addition, with regard to the complexity of the factors affecting clock amplitude regulation, many small molecules show modifying capability in more than one parameter. For example, clock amplitude-enhancing small molecules could enhance reporter oscillatory amplitude and cause period shortening in vitro<sup>250</sup>. Nobiletin, a naturally occurring compound, and an RAR-related orphan receptor (ROR) agonist, could enhance clock amplitude and show robust *Clock*-dependent metabolic protection in diabetic mice<sup>188</sup>. In conclusion, with the increasing understanding of the circadian parameters in disease progression, restoring or enhancing clock function by small molecules might improve clock output physiology, thus alleviating many clock-related diseases.

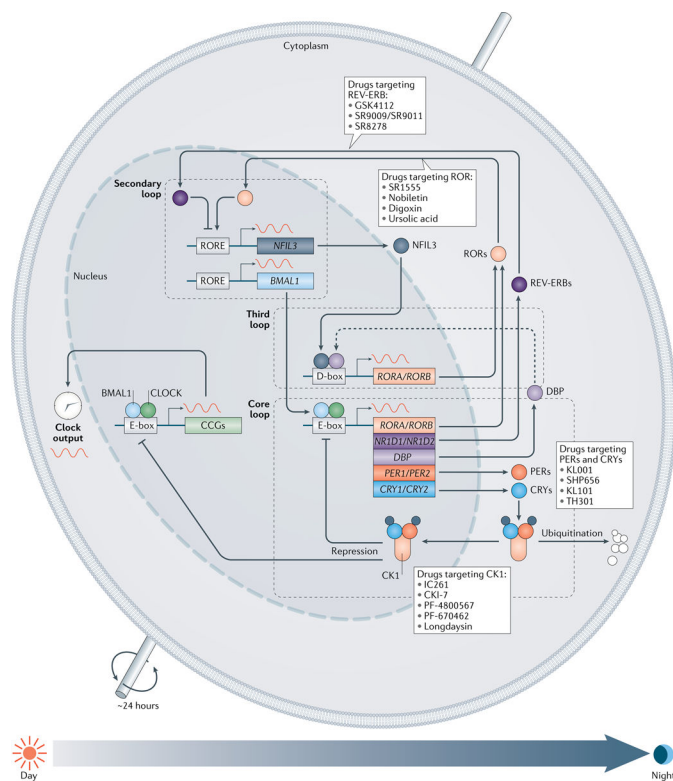
#### **Phase delay**

A phase shift in the circadian rhythm whereby the bedtime and wake-up time will move later in the day.



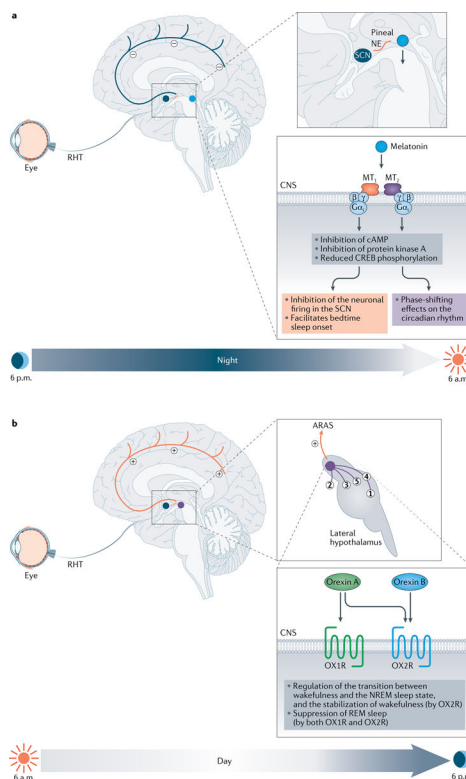
**Fig. 1 | Hierarchical organization of the mammalian clock system.**

The central pacemaker is located in the suprachiasmatic nucleus (SCN) of the anterior hypothalamus. In the input pathway to the SCN, light is received by the intrinsically photosensitive retinal ganglion cells (ipRGCs) expressing melanopsin, which sends electric signals to the SCN through the retinohypothalamic tract (RHT)<sup>203</sup>. Neurotransmitters released by ipRGCs, such as excitatory glutamate and pituitary adenylate cyclase-activating polypeptide (PACAP), cause membrane depolarization in postsynaptic SCN neurons. Changes in  $Ca^{2+}$  and cAMP levels, in turn, induce phosphorylation of cAMP-response element (CRE)-binding protein (CREB) and expression of immediate early genes including the core clock genes, *PER1* and *PER2*, thereby resetting SCN cellular oscillators<sup>240</sup>. Meanwhile, the inhibitory neurotransmitter  $\gamma$ -aminobutyric acid (GABA) decreases the sensitivity of non-image-forming behaviours at low light levels<sup>241</sup>. The SCN neurons are tightly coupled and control peripheral clocks in other brain regions and throughout the body via neuronal and hormonal signals.



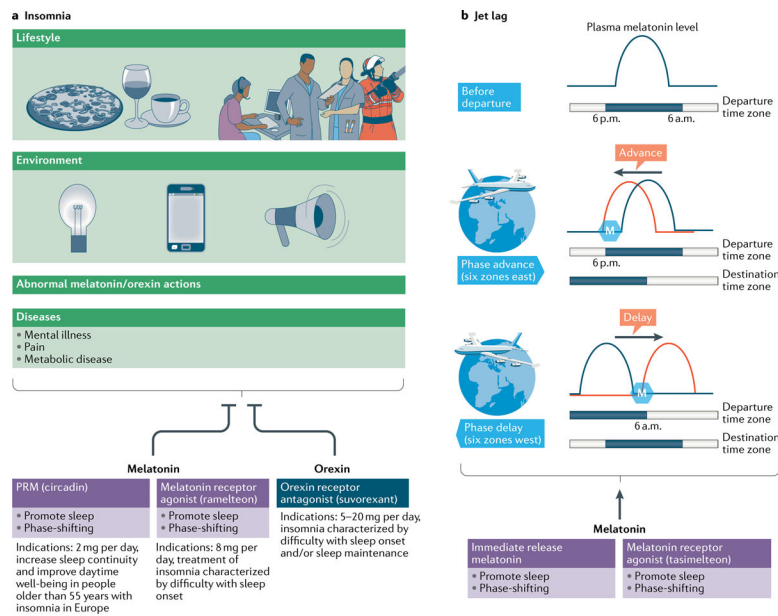
**Fig. 2 | The cell-autonomous core components of the circadian oscillator govern the ~24-hour cycle of gene expression.**

The oscillator consists of positive-feedback and negative-feedback loops that intersect via transcriptional and post-transcriptional regulatory mechanisms<sup>4</sup>. In the core loop, circadian locomotor output cycles kaput (CLOCK) and aryl hydrocarbon receptor nuclear translocator-like protein 1 (BMAL1, encoded by *ARNTL*) heterodimerize to drive expression of period circadian protein homologue (PER) and cryptochrome (CRY) genes via the E-box elements in the morning. PER and CRY proteins subsequently form a repressive complex to inhibit CLOCK and BMAL1 transactivation. The stability of the PER and CRY proteins is regulated by parallel E3 ubiquitin ligase pathways<sup>21–23</sup>. In the secondary, or stabilization, loop two subfamilies of nuclear receptors, REV-ERB $\alpha$  and REV-ERB $\beta$  (encoded by *NR1D1* and *NR1D2*, respectively) as repressors and RAR-related orphan receptors (RORs)  $\alpha$ ,  $\beta$  and  $\gamma$  as activators, antagonistically regulate *BMAL1* and other target genes at the ROR/REV-ERB-response element (RORE) promoter elements at night-time<sup>24</sup>. Another key circadian promoter element is the D-box, activated by the proline and acidic amino acid-rich basic leucine zipper (PAR-bZIP) proteins (such as D-box binding PAR bZIP transcription factor; DBP) and repressed by E4 promoter-binding protein 4 (E4BP4; also known as NFIL3)<sup>15</sup>. Together, clock-controlled genes (CCGs) are transcriptionally regulated by the three loops via activation of the E-box, RORE and D-box elements in their gene promoter regions. Various ligands (PER/CRY, ROR and REV-ERB ligands) and small molecules (casein kinase 1 (CK1) inhibitors) have been found to regulate core clock components and circadian functions.



**Fig. 3 | Melatonin and orexin signalling in circadian rhythm.**

**a** | Melatonin promotes sleep at night: the nightly release of noradrenaline (NE; in orange) stimulated by the suprachiasmatic nucleus (SCN) induces melatonin synthesis in the pineal gland (blue circle)<sup>36</sup>. Melatonin functions through activation of two G-protein-coupled receptors, melatonin receptor 1 (MT<sub>1</sub>) and melatonin receptor 2 (MT<sub>2</sub>) mainly in the SCN<sup>38,39</sup>. By interacting with pertussis toxin-sensitive Gα<sub>i</sub>, MT<sub>1</sub> receptor can inhibit cAMP-response element-binding protein (CREB) phosphorylation, thus is crucial for the suppression (minus symbols) of neuronal firing, whereas MT<sub>2</sub> receptor exerts a phase-shifting effect<sup>38,40,41</sup> and shortens the circadian period of bioluminescence in SCN explant cultures from *Clock*<sup>+/+</sup> mutant mice<sup>242</sup>. **b** | Orexin promotes wakefulness during daytime. The SCN receives signals from light through the retinohypothalamic tract (RHT) and activates orexinergic neurons (purple circle) in the lateral hypothalamic area<sup>46</sup>. Activated orexinergic neurons release orexin A and orexin B and project signals by binding to orexin receptor 1 (OX1R) and orexin receptor 2 (OX2R) in various neurons located throughout the central nervous system (CNS): (1) the locus coeruleus; (2) the tuberomammillary nucleus; (3) the dorsal raphe and median raphe nuclei; (4) the laterodorsal tegmental nucleus; and (5) the pedunculopontine tegmental nucleus<sup>47</sup>. Together, these activating neurons constitute the ascending reticular activating system (ARAS) and directly activate (plus symbols) the cortex, thus promoting wakefulness. Furthermore, OX2R has a major role in stabilization of wakefulness and suppression of sleep. NREM sleep, non-rapid-eye-movement sleep; REM sleep, rapid eye movement sleep.



**Fig. 4 | Pharmacological interventions for insomnia and jet lag.**

**a |** Changes in lifestyle, the environment and health conditions and abnormal action of orexin or/and melatonin could all trigger insomnia<sup>150</sup>. According to its sleep-promoting effect, melatonin and its agonists have therapeutic potential to treat insomnia. Circadin, a commercially available prolonged-release form of melatonin (PRM) has been licensed on the basis of improved nightly sleep quality for treating insomnia disorder in adults older than 55 years in Europe<sup>152,153</sup>. Circadin has a similar affinity for melatonin receptors. Conversely, ramelteon is a melatonin receptor agonist and its affinity for melatonin receptor 1 (MT<sub>1</sub> receptor) is eight times higher than that for melatonin receptor 2 (MT<sub>2</sub> receptor), thus exerting a stronger inhibition of the neuronal firing in the suprachiasmatic nucleus to facilitate bedtime sleep onset<sup>38,153</sup>. Finally, suvorexant, which is a dual orexin receptor antagonist, attenuates orexin receptor 2 (OX2R)-mediated cortical arousal and enhances sleep maintenance and onset associated with orexin receptor 1 (OX1R), respectively<sup>158</sup>. **b |** Melatonin and its receptor agonists can be used in the treatment of jet lag disorder. The key to treating jet lag is the phase response of melatonin. Therefore, tasimelteon, a melatonin agonist with higher affinity for the phase-shifting MT<sub>2</sub> receptor, has been shown to result not only in a significant shift in endogenous circadian rhythms but also an improvement in sleep initiation and maintenance<sup>166</sup>. The timing of melatonin administration is of the essence. Melatonin can phase-advance the physiological and behavioural circadian rhythm when administered in the late biological afternoon. Conversely, when administered in the early biological morning, melatonin delays the circadian clock<sup>41,168,170,171</sup>. Circadian status can be approximately predicted by habitual sleep timing in the adapted departure time zone, and the plasma melatonin level (blue curve) can be used to define circadian phase for the timing of melatonin administration. The adaptation to the destination time zone before transmeridian travel could be achieved as follows<sup>172</sup>: for eastward travel across six time zones, administration of melatonin (blue hexagon) in the afternoon at departure can advance circadian phase (left-shifted plasma melatonin level, red curve); for westward travel across

six time zones, administration of melatonin (blue hexagon) in the early morning can delay circadian phase (right-shifted plasma melatonin level, red curve).

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Table 1 |

Examples of current clinical trials involving drug molecules

Drug	Status	Diseases	Clinical trial identifier	Disease indications	Parameters
<b>Melatonin signalling</b>					
Melatonin	Phase III	Obesity; prediabetic state	NCT02681887	Night-shift work, blindness, myocardial syndrome,	Metabolic parameters, sleep, mood,
	Phase II	Migraine	NCT00849511	migraine, metabolic syndrome, ICU sleep disruption,	insulin and $\beta$ -cell function, blood
	Phase III	Sleep problems; haemodialysis	NCT00388661	dialysis, epilepsy, nocturnal enuresis, DSPS, non-24-hour sleep-wake disorder, jet lag, Smith-Magenis syndrome	coagulation, migraine attack frequency, seizure, incontinence
Tasimelteon	Phase II	Epilepsy	NCT00965575		
	Phase II	Smith-Magenis Syndrome	NCT02231008		
	Phase III	Circadian rhythm alterations	NCT02231008		
Ramelteon	Phase III	Non-24-hour sleep-wake disorder	NCT01430754		
	Phase II	Jet lag disorder	NCT03291041		
	Phase II	Sleep disorders; circadian rhythm	NCT00593736		
<b>Orexin signalling</b>					
Modafinil	Phase IV	Bipolar disorder	NCT01965925	Bipolar disorder, long-term shiftwork, sleep disorder,	Sleep, circadian rhythms
Armodafinil (CEP 10953)	Phase III	Excessive sleepiness; shiftwork sleep disorder	NCT00080288	Alzheimer disease	
Lemborexant	Phase II	Irregular sleep-wake rhythm disorder	NCT03001557		
Suvorexant	Phase IV	Shiftwork sleep disorder	NCT02491788		
<b>Combined or other signalling</b>					
Vitamin B <sub>12</sub>	Not applicable	Sleep disorders; circadian rhythm	NCT00120484	Night owl or DSPS	Sleep and circadian rhythm
Methylselenocysteine	Not applicable	Breast cancer; prostate cancer	NCT01611038	Night-shift work	Circadian rhythm and oestrogen receptor B levels
Melatonin and meal timing	Not applicable	Obesity; cardiometabolic diseases	NCT03490864	Obesity, chronic insomnia, DSPS	Sleep, circadian rhythm
Melatonin and light therapy	Phase IV	DSPS	NCT00834886		

Drug	Status	Diseases	Clinical trial identifier	Disease indications	Parameters
Ramelleon and multicomponent behavioural therapy	Phase IV	Insomnia	<a href="#">NCT01180855</a>		

DSPS, delayed sleep phase syndrome; ICU, intensive care unit.

Table 2 |

Preclinical studies of small-molecule modulators targeting clock components and key regulators

chemical modulators	Mechanism of action	Physiological effects	Refs
<b>CRY1/CRY2</b>			
KL001 and CRY-stabilizing derivative (SHP656, KL101 and TH301)	Stabilize CRY, lengthen period, reduce amplitude	Improvement of glucose tolerance in obese mice, inhibition of glioblastoma stem cell proliferation in vitro, inhibition of glioblastoma tumour growth in vivo, enhancers of brown adipocyte differentiation	94,194,196,197
2-Ethoxypropanoic acid (KS15)	Inhibits CRY, activates E-box transcription, reduces amplitude	Inhibition of breast cancer cell growth	231
<b>REV-ERBs</b>			
GSK4112	Enhances REV-ERB and NCOR peptide interaction	Inhibition of gluconeogenesis and inflammatory response in primary cells	129,179,189
SR9009, SR9011	Derived from GSK4112, they are selective agonists for REV-ERB that alter circadian behaviour, clock gene expression ( <i>BMAL1</i> , <i>PER1</i> , and <i>PER2</i> ) and expression of some glioblastoma stem cell markers (for example, <i>OLIG2</i> and <i>SOX2</i> )	Improvement of glucose homeostasis in obese mice, promotion of wakefulness, anxiolytic, inhibition of glioblastoma stem cell proliferation	94,181,183
SR8278	GSK4112-derived antagonist that increases expression of REV-ERB target genes (for example, <i>BMAL1</i> , <i>PCK1</i> and <i>G6PC7</i> ) in cells	Anxiety attenuation and amelioration of myocardial injury	82,189
<b>RORs</b>			
Nobiletin	Agonist, enhances amplitude and lengthens period	Metabolic homeostasis improvement in obese/diabetic mice; broad efficacies with regard to inflammation and atherosclerosis	188,232,233
Neuroscogenin	Agonist promoting ROR interaction with NCOA2/TIF2, activates BMAL1 expression	Activation of hepatic expression of ROR metabolic target genes	234
SR1001	Derived from T0901317 with high inverse agonist activity and selectivity for ROR $\alpha$ and ROR $\gamma$ t	Inhibition of T <sub>H</sub> 17 cell differentiation and autoimmunity	193
SR2211, SR1555, digoxin, ursolic acid, ML209	ROR $\gamma$ inverse agonist	Inhibition of T <sub>H</sub> 17 cell differentiation	189,191,192
SR3335	ROR $\alpha$ inverse agonist	Reduction of blood glucose level in obese mice	235
SR1078	ROR $\alpha$ agonist	Induction of apoptosis and inhibition of hepatoma cell growth	236
<b>CKI kinases</b>			
CKI-7, IC261, D4476, PF-670462, PF-4800567 longdayasin, LH846, compounds 1–3 among others	Inhibitors, lengthen period	Pharmacological inhibition of CK1, significantly lengthen circadian rhythms mainly by nuclear retention of PER2. CK1 is broadly involved in various pathophysiologicals, including familial sleep and mood disorders	199–201,237
<b>ADORA2B</b>			
BAY 60-6583	ADORA2B agonist, PER2 stabilization	Enhancement of adenosine signalling and cardioprotection against myocardial ischaemia	74,238

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chemical modulators	Mechanism of action	Physiological effects	Refs
<i>PER2</i> Lithium	Increased transcription of <i>PER2</i> , lengthens period and enhances amplitude of <i>PER2</i> rhythms	Treatment of bipolar disorder associated with circadian rhythm disruptions	239

For additional clock-modulating compounds, see REFS<sup>5,189,192,237</sup>. ADORA2B, A2B adenosine receptor; CK1, casein kinase 1; CRY, cryptochrome; NCOR, nuclear receptor co-repressor; PER2, period circadian protein homologue 2; ROR, RAR-related orphan receptor; TH17 cell, interleukin-17-producing T helper cell.

**Table 3 |**

## Clinical trials involving behavioural and environmental modifications

Modifications	Disease indications	Parameters	Examples
Light therapy	Stroke, ageing, Parkinson disease, Alzheimer disease, mood disorders, metabolic disease, night shift, PTSD, traumatic brain injury, sepsis, cancer-related fatigues	Circadian rhythms in sleep and blood parameters, cognition, activity, fatigue, depression and PTSD scores, inflammation markers	<a href="#">NCT02186392</a>
			<a href="#">NCT02502045</a>
			<a href="#">NCT02769858</a>
			<a href="#">NCT01855126</a>
			<a href="#">NCT01048294</a>
			<a href="#">NCT01747811</a>
Time-restricted feeding	Obesity, metabolic syndrome, shiftwork	Body weight, metabolic homeostasis, cardiometabolic parameters	<a href="#">NCT02204735</a>
			<a href="#">NCT03527290</a>
Sleep intervention	Night-shift work, traumatic brain injury	Sleep quality	<a href="#">NCT02838082</a>
			<a href="#">NCT02609373</a>
Scheduled activity	Diabetes, Alzheimer disease	Sleep, glucose homeostasis	<a href="#">NCT03553524</a>
			<a href="#">NCT01920672</a>
Combination: including sleep, light, exercise and lithium	Night-shift work, HIV-related fatigue, mood disorders, ICU stay	Circadian phase, sleep, metabolic parameters, mood rating, clock gene expression	<a href="#">NCT01284140</a>
			<a href="#">NCT01767181</a>
			<a href="#">NCT02126007</a>
			<a href="#">NCT01799733</a>
			<a href="#">NCT03405493</a>
			<a href="#">NCT01431573</a>

ICU, intensive care unit; PTSD, post-traumatic stress disorder.