

From immune homeostasis to inflammation, a question of rhythms

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Abstract

Numerous physiological processes vary according to the time of the day in mammals, including the immune defense mechanisms. These circadian rhythms are generated by circadian clocks located in most cell types including cells of the immune system. Recent research has shown that circadian clock function is key to maintaining the homeostasis of the immune system. Indeed, the development and differentiation of different immune cells, and their trafficking between tissues, rely on proper clock function. Disruption of immune system homeostasis can lead to the development of pathological processes such as inflammation or autoimmune diseases. Circadian clocks also control the response of the immune system to inflammatory challenges and infections, and clock dysfunction can provoke or enhance disease. This review focuses on the role of circadian clocks in the homeostasis of immune cells, such as cell recirculation in the peripheral blood and tissues, as well as in the context of inflammation in response to infectious challenges or autoimmune disorders.

Keywords: Circadian clock, clock genes, immune response, inflammation, infection, autoimmunity.

1. Introduction: circadian clocks in the immune system

Circadian clocks are composed of clock genes, such as *Clock*, *Bmal1*, *Per1-3*, *Cry1-2*, and *Rev-erb/Ror* genes, expressed in nearly all cells and tissues in mammals [1]. The clock genes and their protein products function within transcriptional-translational feedback loops [1]. As a consequence of these feedback loops, clock transcription factors such as CLOCK/BMAL1 and REV-ERB α display an activity that varies over the 24 h day, and thus, numerous target genes of these factors display circadian rhythms in their expression levels. Such mechanisms also occur within cells of the immune system. Therefore, a circadian variation of various aspects of the innate and the adaptive immune system has been observed in mammals, both at steady state and upon immune challenges, such as infection with pathogens or inflammatory conditions.

Clock genes are expressed in most cell types of the innate and adaptive arms of the immune system [2-4]. Within innate immunity (the first lines of defense against non-self-bodies), basophils [5], eosinophils [6], mast cells [6-8], monocytes/macrophages [9-13], dendritic cells [13,14], and natural killer cells [15] all express clock genes rhythmically. This is also the case for cells of the adaptive immunity (i.e. the antigen-specific immune response), T [16,17] and B lymphocytes [13]. Some consider intestinal epithelial cells and cells lining the lung bronchiole as “non-conventional” immune cells, which were also found to express clock genes [18,19]. Thus, over the past decade work in the emerging field of chronimmunology has highlighted the relevance of circadian rhythms in the regulation of the immune system to confer an evolutionary advantage to the organism in the context of environmental cycles of lighting, temperature, pathogen encounter, etc. This review will provide an overview of recent findings on the contributions of circadian clocks to the homeostasis and responses of the immune system.

2. Circadian clocks and immune cell development and differentiation

The expression of molecular clock components by most immune cells raises the question of whether the circadian clock in these cells is important for their development and function. A few studies have shown

that indeed, the clock is required for successful immune cell development and differentiation. This is the case of the T helper 17 (Th17) cells [20] and innate lymphoid cells [21]. In both cases the mechanism relies on a rhythmic transcriptional repressor called Nuclear Factor Interleukin 3-Regulated (NFIL3, also known as E4 Promoter-Binding Protein 4, E4BP4). NFIL3 is known to bind D-box promoter elements, where it competes with PAR transcription activators such as DBP [2]. As for all rhythmically-expressed transcription factors, NFIL3 can generate circadian rhythms of expression of its gene targets [2]. Within T cells, REV-ERB α (a transcription factor whose expression is activated by CLOCK/BMAL1) represses the expression of NFIL3. In turn, NFIL3 represses the expression of ROR γ t, the master transcriptional regulator of Th17 cell differentiation. Accordingly, Th17 counts (following CD4 T cell polarization) are higher when using cells of *Nfil3* KO mice and lower with cells from *Rev-erba* KO mice. Thus, NFIL3 can be viewed as a suppressor of Th17 cell differentiation [20]. A follow up study demonstrated a role for NFIL3 in the development of innate lymphoid cells from their precursors [20].

Whereas the differentiation of specific T cell subtypes may be relying on their endogenous clock, two studies highlighted that the development of T cells from progenitors is not affected by their own intrinsic clock. Indeed, in mice with a T cell-specific deletion of the essential clock gene *Bmal1* [17], as well as in irradiated wild-type mice grafted with *Bmal1* KO bone marrow [22], similar CD4 and CD8 T cell counts were found compared to the respective control mice.

In contrast, the development of B cells, which are involved in the humoral immune response, appears to be controlled by the circadian clock [22]. Indeed, in *Bmal1* KO mice, the number of fully developed B cells is lower than in WT mice [22]. Interestingly though, bone marrow chimera experiments mice showed that the need for clock function for cell development is not intrinsic to the B cells themselves, but is rather in the non-hematopoietic cells [22]. This is in line with data on B cell-specific *Bmal1* KO mice, showing intact B cells counts compared to wild-type littermates [17].

In sum, T cell and B cell development appears insensitive to disruption of the clock in these cells or their progenitors. However, the subsequent differentiation of these cells in specific subsets might be controlled by their endogenous clocks, as illustrated by the studies on Th17 cell differentiation.

3. Circadian clocks and immune cell trafficking

Immune cells recirculate to the different secondary lymphoid organs via the bloodstream and the lymph. A number of recent studies have uncovered a role for circadian clocks in regulating this immune cell trafficking. The blood and spleen counts of inflammatory monocytes present a daily rhythm in mice, with a maximum in the daytime and a minimum at night (Figure 1), and an antiphasic rhythm in the bone marrow [12]. This rhythm appears to be dependent on the clock within these monocytes, as it is lost in mice KO for the essential clock gene *Bmal1* specifically in myeloid cells (Figure 1). The data from this study, and from other reports [11,23], suggest that the clock in myeloid cells controls the circadian expression of C-C Motif Chemokine Ligand 2 (CCL2), a chemokine important for the trafficking of monocytes.

Clock control of adaptive immune cell trafficking has also been studied. T and B lymphocytes recirculate in a circadian manner in the blood and lymphoid organs [24-27]. Prior work has shown that the recruitment of white blood cells to tissues was regulated by the autonomic nervous system and adrenergic signaling, themselves under the control of the central circadian clock in the brain [28]. Several recent studies have started to uncover the mechanisms for the circadian control of T cell trafficking. Suzuki *et al.* found a rhythm of T cell counts in lymph nodes (LNs) and showed an involvement of the adrenergic pathways: the day-night difference of T and B cell counts in LNs, and the variation of lymphocyte egress from these organs, were lost after depletion of adrenergic nerves by treatment of mice with 6-OHDA, and in mice KO for the β_2 adrenergic receptor [26]. In parallel, Druzd *et al.* also studied the rhythm of T cell counts in the LNs, but they identified a distinct mechanism [27]. Using mice lacking clock function specifically in T cells, they showed that the T cell clock was essential for the LN T cell

count rhythm by regulating both the entrance into and exit from these organs (Figure 1). It does so by inducing rhythms of the expression of Sphingosine-1-phosphate receptor 1 (S1PR1) and C-C Motif Chemokine Receptor 7 (CCR7), two receptors important for T cell trafficking in the LN [27].

A more recent study shed new light on the T cell trafficking rhythm to lymphoid organs. Shimba *et al.* showed that in mice, glucocorticoids, via their receptor, act on the expression of IL-7 receptor, which varies as a function of time of day at the surface of T cells [29]. This in turns leads to a rhythm of the expression of C-X-C chemokine receptor 4 (CXCR4), which is involved in the recruitment of T cells to lymphoid organs. A parallel can be made with earlier studies in humans, which have also linked cortisol secretion with CXCR4 expression and blood T cell counts [30,31]. Indeed, these studies showed a negative correlation between T cell numbers and cortisol concentration in the blood. Both the endogenous morning cortisol rise and exogenous cortisol administration decreased T cell blood counts, and cortisol increased CXCR4 levels at the surface of T cells.

In humans, blood T cell counts exhibit a peak during the night while in mice, the peak is in the daytime [24,25,32]. Both the glucocorticoid rhythm in the blood and the rhythm of CXCR4 in T cells show an opposite phase in humans (diurnal) vs. mice (nocturnal). This opposite phase of glucocorticoids and CXCR4 might explain the opposite phases of T cell counts in humans vs. mice. However, Zhao *et al.* have recently proposed another mechanism. They used humanized mice, which were transplanted with human hematopoietic stem cells [33]. Thus, within the same animals, they were able to compare blood counts and protein expression for cells of both human and mouse origin. White blood cell count rhythms and CXCR4 expression rhythms had opposite phases in the mouse and human cells within these humanized mice, which is consistent with the previously described studies. However, Zhao *et al.* suggested a new possible mechanism for the circadian regulation of CXCR4 expression that involves reactive oxygen species (ROS) [33]. In this model, the opposite rhythms in mice and humans would be due to an opposite effect of MAP kinase activation on ROS generation. However, the lack of rhythms of

clock gene expression in the human white blood cells suggests that in this model, they might not be receiving the proper cues from the mouse environment, which may affect their trafficking.

4. Circadian clocks and the inflammatory response to microbe-derived molecules

The innate immune system constitutes the first line of defense against infections. As innate immune cells express the clock machinery, it is not a surprise that their responses are controlled in a circadian fashion. Innate immune cells recognize pathogens via receptors for molecules named pathogen-associated molecular patterns (PAMPs). These receptors, called pattern recognition receptors (PRR), were in some cases shown to have a daily rhythm of expression in mice. This is the case for Toll-like Receptor 5 (TLR5), which recognizes bacterial flagellin. This triggers the activation of the NF κ B pathway, involved in stimulating proinflammatory cytokine secretion [34]. In fact, CLOCK protein interacts with the p65 subunit of NF κ B and enhances its transcriptional activity on target genes including inflammatory cytokines [34]. TLR9 recognizes bacterial and viral DNA. The expression levels of this receptor fluctuate over the 24 h cycle in mouse peritoneal macrophages. These levels are correlated with the magnitude of the response to vaccination in the presence of a TLR9 ligand as adjuvant, with a maximum of the response around the middle of the night [35].

Macrophages can produce proinflammatory cytokines such as interleukin(IL)-1 β , IL-6, TNF α , in response to PAMPs. Several groups have demonstrated a circadian control of cytokine secretion by macrophages in response to the bacterial endotoxin lipopolysaccharide (LPS), both in mice and using cells *ex vivo*, with a higher secretion at the day/night transition [10-12,36,37]. This rhythmic regulation appears to be cell-autonomous (i.e. relying on the clock in macrophages), because a myeloid-specific KO of *Bmal1* abolished the morning/evening difference, with morning levels of cytokine response being increased to the evening levels [11,36]. The receptor for LPS is TLR4. A transcriptomic study on peritoneal macrophages showed a rhythm of transcripts for signaling molecules downstream of TLR4

[10]. Moreover, a recent proteomic study on Kupffer cells (the liver-resident macrophages) showed a 24 h rhythm of TLR4 protein (and downstream signaling molecules), with a peak during the early daytime [38]. Interestingly, the relationship between the macrophage clock and LPS response is bidirectional, as the LPS- and NF κ B-induced microRNA mir-155 was shown to act directly on BMAL1 expression. This pathway leads to a suppression of *Bmal1* expression in macrophages upon LPS stimulation, which in turn leads to an increased secretion of pro-inflammatory cytokines [36]. Other studies pointed to an impact of *Per2* and *Cry* gene mutation on the response to LPS [39,40]. Environmental circadian disruption can also affect the response to LPS. Indeed, simulated shift work in rats and repeated jet lag in mice led to an increase of the inflammatory responses after an LPS challenge [41,42]. In humans, it was shown that the circadian rhythm of response of monocytes to LPS became desynchronized from the rhythm of blood monocyte counts in subjects placed in a schedule simulating night shift work [32]. This showed that the regulation of these two rhythms depends on distinct mechanisms, and that abnormal sleep-wake timing can impact immune functions in humans.

Other TLRs were recently added to the list of PRRs showing rhythmic expression and/or function. The liver macrophage proteomic study already cited above also showed a daily rhythm of TLR3 and TLR8 in liver-resident macrophages [38]. A recent study provided evidence that a day-night difference in the severity of sepsis upon cecal ligation and puncture in mice relies on TLR2 (but not TLR9), and on the clock in leukocytes [43]. Silver and colleagues looked at the expression and function of many TLRs. All the PRRs they tested (TLR1-8) showed time of day-dependent mRNA expression in spleen cells, and some of their responses to their respective PAMPs showed time of day-dependence [44]. They also showed a diurnal rhythm of TLR3 response *in vivo*.

These studies confirmed a role of the circadian machinery in the cytokine response to bacterial endotoxin challenge. What is the molecular link between the clock and the cytokine response? Many studies from the past few years have pointed to REV-ERB α as a major intermediate. Mice KO for *Rev-*

erba and human cells with shRNA against this clock gene both show increased IL-6 response to LPS, and conversely, overexpressing REV-ERB α or stimulating its activity with a specific ligand blunted LPS-induced IL-6 expression and release [11,23]. REV-ERB α achieves a circadian expression of macrophage-specific genes by binding to enhancers selected by macrophage lineage-determining factors [45]. A similar role of REV-ERB α in modulating the inflammatory response was also found in a pulmonary infection model, in which LPS is aerosolized into the airway of mice. The inflammatory response to LPS inhalation follows a circadian rhythm, including a circadian rhythm of the production of the CXCL5 chemokine by the bronchiole epithelial cells, and consequently, of the recruitment of neutrophils to the lungs [18]. A recent study showed that in mice KO for *Rev-Erba* (either globally or specifically in the lung epithelial cells), there was an increased inflammatory response to LPS (with increased IL-6 and CXCL5 expression) and a loss of the rhythm of neutrophil infiltration in the lung [46]. Again, a REV-ERB α agonist led to a suppression of the IL-6 response [46].

Recently, a study from Pourcet and colleagues provided a mechanism for the action of REV-ERB α on the inflammasome, which is important for the response to non-infectious danger-associated molecular patterns [47]. The activation of the NLRP3 inflammasome is critical for the maturation and secretion of cytokines such as IL-1 β and IL-18. They showed that REV-ERB α directly regulates the expression of the *Nlrp3* gene, encoding the core component of the NLRP3 inflammasome. Thus, REV-ERB α , via the repression of *Nlrp3* but also by a direct action on the *Il-1 β* and *Il-18* genes, leads to a rhythm of NLRP3 inflammasome activity and cytokine expression. The study also shows the impact of this regulation in mouse models of peritonitis and fulminant hepatitis, in which REV-ERB α has a protective anti-inflammatory effect at certain times of the day via regulation of NLRP3 expression [47].

5. Circadian clocks and the control of infections

Over the past few years, studies have started to show the importance of circadian clocks for the control of different types of infection (Figure 2). This is the case for infection of mice with the bacteria *Diplococcus pneumoniae* [48] and *Salmonella enterica* serovar Typhimurium[37]: a day/night variation was found, with a higher mouse mortality or bacterial load after infection during the daytime compared to the night time. Moreover, in the case of *Salmonella*, a corresponding rhythm of mRNA expression of inflammation markers was observed in the colon [37]. These daily variations were abolished in *Clock* mutant mice, confirming a role of the clock in the control of the *Salmonella* infection. The clocks in immune cells also underlie a rhythm of the outcome of infection with the bacterium *Listeria monocytogenes*, with higher pathogenicity (and poorer mouse survival) in the second half of the light period vs. in the early day [12].

Two studies have shown an effect of the circadian clock on viral infections. Edgar and colleagues showed that the level of viral replication upon infection of mice with herpesvirus varies according to the time of time, a variation that is abolished (and overall levels increased) in *Bmal1* KO mice [49]. Another group showed that the inflammatory response, encephalitis and survival of mice following vesicular stomatitis virus infection was dependent on the time of infection, and that the clock protein REV-ERB α affects disease outcome [50]. One study has recently shown a time-dependent variation of the control of infection by the fungus *Aspergillus fumigatus* [51].

Finally, an implication of circadian clocks was studied in the context of parasitic infections. Indeed, infection of mice with the protozoan parasite *Leishmania major* leads to a higher parasite load when done in the early night compared to infection in the early day, both upon infection in the footpad or intra-peritoneally (i.p.) [52]. This can be put in parallel with a higher inflammatory response upon i.p. infection in the early night compared to the early day, both in terms of recruitment of neutrophils and macrophages (the cells that act as hosts for *Leishmania*) and chemokine/cytokine expression. All these rhythms were abolished in mice with clock dysfunction in hematopoietic cells [52]. Another group studied infection with the intestinal worm parasite *Trichuris muri*. The expulsion of the parasite and the cytokine response

depended on the time of infection [14]. Interestingly, mice with a dendritic cell (DC)-specific *Bmal1* deletion lost the morning/evening difference in worm burden and cytokine secretion, showing the importance of the DC clock in the development of the response to the infection [14]. The case of parasitic infections is interesting in that it raises the question of the contribution of circadian control in parasite-host interactions. The parasite has rhythms as well as the transmission vector (e.g. the sandfly in the case of *Leishmania*), and how this is coordinated with the host's immune system's rhythms has implications both for understanding the evolution of parasite-host interactions and to design new prophylactic strategies [53]. This is especially important in the context of the recent discovery of endogenous circadian rhythms of gene expression in the protozoan parasite *Trypanosoma brucei*, which suggests that it has its own circadian clock, although its molecular components have yet to be defined [54]. The synchrony between the parasite and the host was addressed for *Plasmodium* infection (*Plasmodium* is the causing agent of malaria). *Plasmodium* infection (in its red blood cell stage) displays a daily (24 h or a multiple of 24 h) rhythmicity, synchronous among red blood cells. Infection with parasite-bearing red blood cells of mice with a matched or reversed light:dark cycle was compared: parasites out of synchrony with the host had reduced proliferation and transmission potential [55].

6. Circadian clocks in animal models of autoimmune disorders

In autoimmune diseases, the immune system attacks self tissues. Given that circadian clocks regulate immunity, a deregulation of this rhythmic regulation could conceivably lead to excessive immune responses, and autoimmunity. Therefore, several recent studies have tested the impact of circadian clocks in animal models of autoimmune diseases. Rheumatoid arthritis (RA) is an autoimmune disease involving inflammation in the joints, and overgrowth of fibroblast-like synovial cells, and joint remodeling and destruction. Symptoms of RA display daily rhythms, being increased in the morning [56]. Experimental arthritis consists in injecting rodents with an antibody against type II collagen, or to immunize the animals with type II collagen and an immune adjuvant. Clock genes are expressed in

cultured human rheumatoid synovial cells [57] and in mouse synovial cells [58,59]. In the joints of rodents with experimental arthritis, clock gene expression is altered, and the extent of disease varies according to the time of arthritis-inducing treatment. Moreover, clock disruption by constant light or clock gene deletion or knockdown led to altered clock gene expression in the joints, exacerbated joint inflammation, and increased arthritis score [58,59].

Another autoimmune disorder that has been studied in the context of circadian clocks is multiple sclerosis (MS). In MS patients, the immune system attacks myelin in the central nervous system. Experimental autoimmune encephalomyelitis (EAE) is an experimental model of MS where mice are immunized with MOG₃₅₋₅₅ (a peptide antigen consisting in a fragment of a myelin protein) together with Freund's adjuvant. This leads to a demyelination of neurons similar to that occurring in MS patients. Two reports showed the importance of the circadian clocks of both T cells and myeloid cells for disease progression in EAE mice (Figure 3). Day/night differences were observed, with higher clinical score after MOG₃₅₋₅₅ immunization in the daytime. These day/night differences were abolished both in T cell-specific and myeloid cell-specific *Bmal1* KO mice, with the higher daytime disease score going down to the lower nighttime level, indicating a protective effect of the mutation [27,60]. It will be interesting to understand the mechanisms for the respective involvement of these two distinct circadian clocks in the control of the disease in MS. Interestingly, in humans, MS relapses showed a variation according to the time of year, with more cases during spring and summer, compared to the rest of the year. This was inversely correlated with levels of the circadian hormone melatonin. Using human cells and mice, it was found that melatonin affects the differentiation of cell types with opposite roles in MS: it inhibits the differentiation into Th17 (via NFIL3), which contribute to the pathology, whereas it promotes the production of regulatory T (Treg) cells, which are protective [61].

7. Conclusion

In this review, we mainly focused on the circadian control of innate arm of the immune system. The recent studies presented here highlight the importance of circadian clocks both in the control of the homeostasis and when the organism faces an infectious challenge. Recent work has started to point to a circadian control of the adaptive immune response too (beyond the T cell trafficking aspect described in the present review). Moreover, it will be necessary to understand the molecular mechanisms underlying the circadian control of immunity, to better define how to act on these pathways to design new therapies.

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Figure legends

Figure 1. Circadian rhythms of immune cell trafficking

Profiles of the rhythm of immune cell trafficking in the blood and lymphoid organs in mice. Inflammatory monocytes are shown for wild-type (thick lines) and myeloid-specific *Bmal1* knock-out mice (dotted lines). T lymphocytes are shown for wild-type (thick lines) and T lymphocyte-specific *Bmal1* knock-out mice (dotted line).

Figure 2. Circadian control of the infectious disease development

Circadian clocks regulate antiviral responses (in blue), anti-bacterial responses (in red), and responses to eukaryotic pathogens (in green) in mouse models. As a result, the pathogen load, the immune response and the host's survival are all susceptible to vary according to the time of day of initial infection. The right part of the figure displays the outcome of infections occurring in the daytime (or subjective day for experiments in constant darkness). The left part of the figure displays the outcome of infections occurring in the night time (or subjective night for experiments in constant darkness). The numbers are Zeitgeber times (hours after the onset of light in the morning, under a light:dark cycle) or Circadian times (corresponding times under constant darkness).

Figure 3. Control of the severity of experimental autoimmune encephalomyelitis (EAE) and of immune cell counts by the circadian clock

Day (left, yellow) and night (right, grey) levels of immune cells in wild-type mice (top) and cell type-specific *Bmal1* knock-out mice (bottom). Figure insets present a very simplified view of the clock feedback loop and graph insets present the time-dependence of the EAE clinical score in wild-type (WT) mice (top) and the lack of time-dependence in cell type-specific *Bmal1* knock-out KO mice (bottom). T lymphocyte counts are shown for experiments comparing WT vs. T lymphocyte-specific *Bmal1* KO

mice, whereas neutrophil and monocyte counts are shown for experiments comparing WT vs. mice for *Bmal1* KO in myeloid cells and granulocytes (LysM-Cre).

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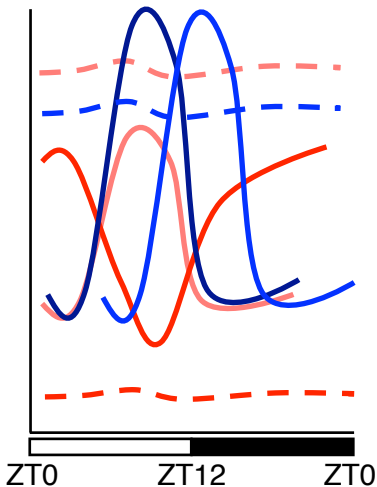
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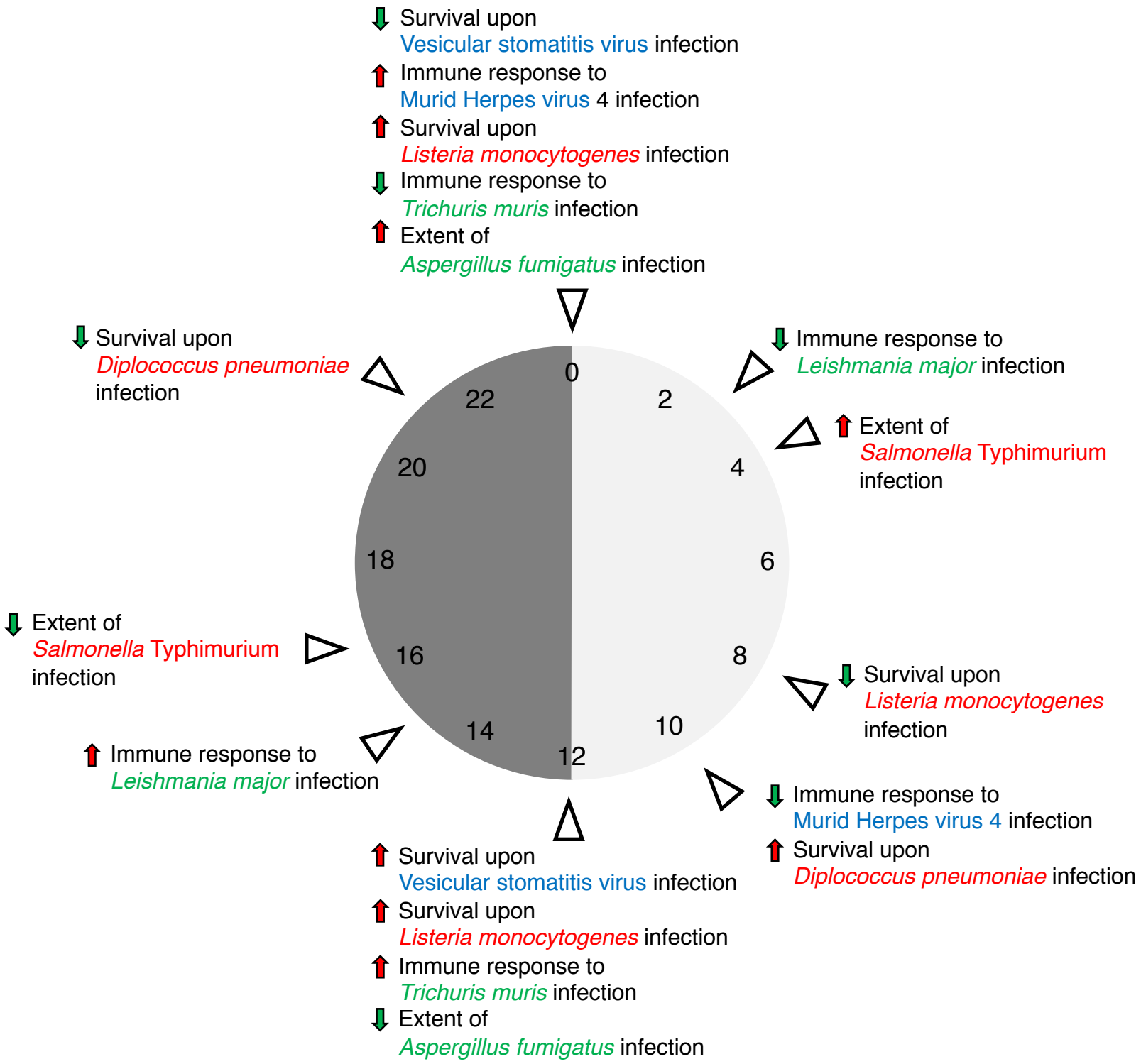
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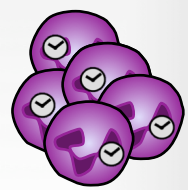
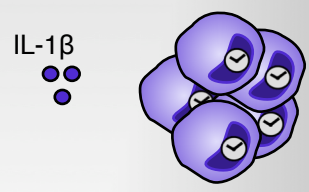
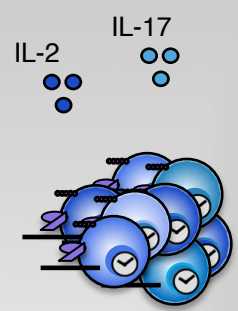
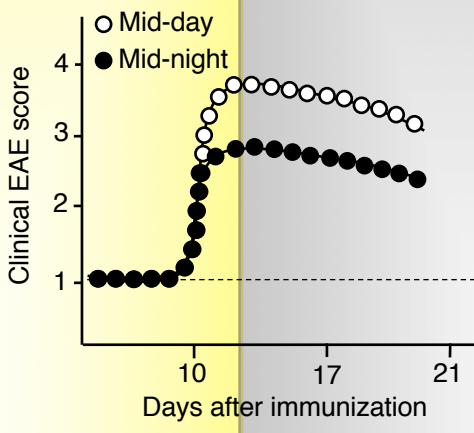
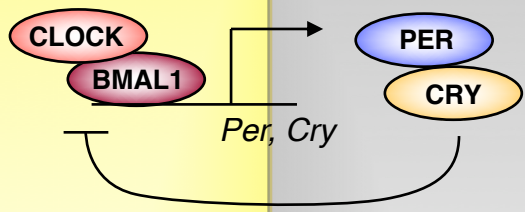
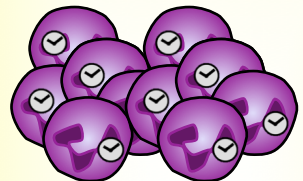
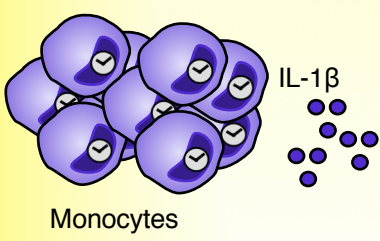
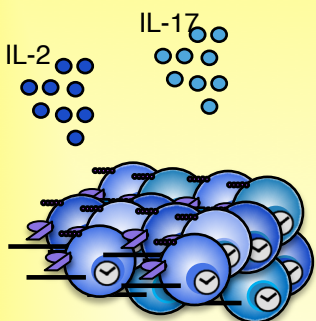
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Cell counts

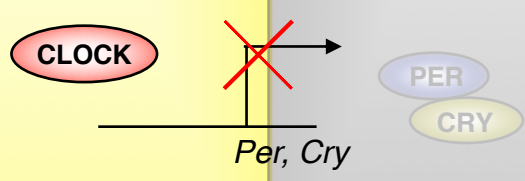
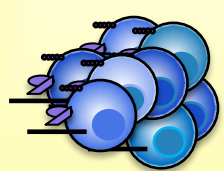
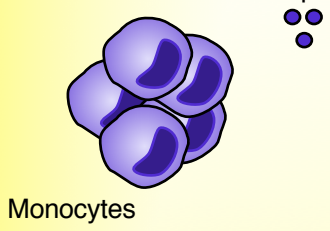
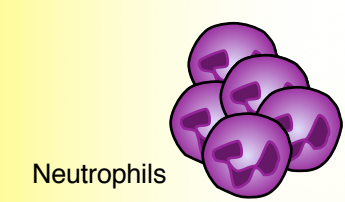
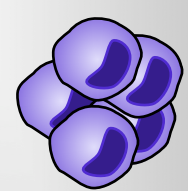
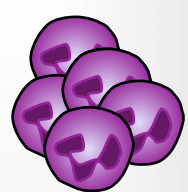
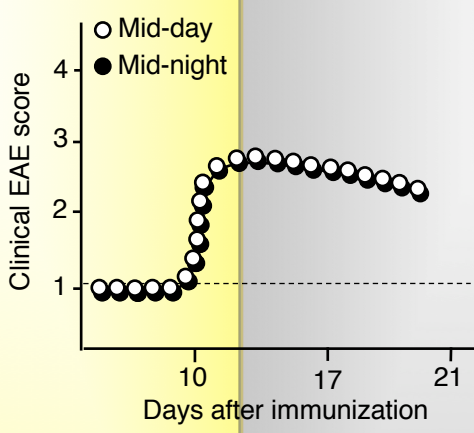






Experimental Autoimmune Encephalomyelitis

Myeloid-specific *Bmal1* KO mice



T lymphocyte-specific *Bmal1* KO mice