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The Influence of Biological Rhythms on Host-Parasite Interactions

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Abstract

Biological rhythms—from circadian control of cellular processes to annual cycles in life history—are a main structural element of biology. Biological rhythms are considered adaptive because they allow organisms to partition activities to cope with, and take advantage of, predictable fluctuations in environmental conditions. A flourishing area of immunology is uncovering rhythms in the immune system of animals, including humans. Given the temporal structure of immunity, and rhythms in parasite activity and disease incidence, we propose that the intersection of Chronobiology, Disease Ecology and Evolutionary Biology holds the key to understanding host-parasite interactions. We review host-parasite interactions while explicitly considering biological rhythms, and propose that (1) rhythms influence within-host infection dynamics and transmission between hosts, (2) rhythms might account for diel and annual periodicity in host-parasite systems, and (3) rhythms can lead to a host-parasite arms race in the temporal domain.

Keywords: biological rhythm, circadian, circannual, infection, parasite, season, temporal niche

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Biological Timekeeping

Environmental rhythms are a ubiquitous feature of our planet. Many rhythms are caused by geophysical cycles, including diel, tidal, lunar, and annual rhythms. These rhythms are highly predictable and have resulted in the evolution of biological clocks throughout the tree of life [1]. Most biological activities have rhythmic time structure, which scales from gene expression to life history events such as breeding and hibernation. Rhythmic time structure allows organisms to partition and prioritize life history activities—whether they are molecular or behavioral—relative to predictable fluctuations in environmental conditions. For example, for cyanobacteria, which are an ancient lineage, sunlight provides both energy and risk. Cyanobacteria have adapted to this challenge by temporally partitioning photosynthesis from UV-sensitive DNA replication [2]. Likewise, throughout the year, organisms must meet survival needs, while seasonally requiring further resources for reproduction and other activities. This leads to annual cycles of life history, when animals alternate between reproductively active states and inactive states such as dormancy, hibernation, or migration, a retreat to wintering grounds that buffer against resource scarcity [1].

Over evolutionary time, organisms have adapted to environmental fluctuations by an internal representation of time—endogenous biological clocks—that perpetuate biological rhythms even when environmental conditions are kept constant. These rhythms are characteristically innate, evidenced by

38 the observation that individuals who have never experienced environmental fluctuations display rhythmicity [1]. Endogenous biological rhythms oscillate with period lengths that approximate those of
39 geophysical cycles, and are accordingly called circadian, circatidal, circalunar, and circannual. Circadian
40 rhythms, which are most heavily studied, are driven by cell-specific transcription-translation feedback
41 loops that are integrated across the organism. The evolutionary origin of internal clocks is ancient,
42 with circadian clocks being a unifying feature of eukaryotes and cyanobacteria [3], and a new area of
43 research in other bacterial lineages [4]. The endogenous circannual clock underlying seasonal rhythms
44 is also thought to be evolutionarily conserved, since circannual clocks are found in organisms ranging
45 from dinoflagellates [5] to mammals and birds [6, 7].
46

47 An internal representation of time enables the anticipation of favorable environmental conditions,
48 ensuring that activities are initiated in advance to match the opportune time. For example, to rear
49 offspring at the time of maximal food abundance, many species activate their reproductive system and
50 copulate far in advance, potentially under harsh conditions. If individuals initiated breeding activities
51 when food abundance was maximal, offspring would be reared outside the optimal environmental win-
52 dow [6]. Endogenous biological clocks function in concert with the geophysical cycles to which they
53 synchronize [8, 9]. Synchronizing cues (also called *zeitgebers*) include diel and annual changes in light,
54 temperature, and other factors. Jetlag, the overturning of rhythms resulting from changing time zones,
55 and the subsequent re-synchronization of the circadian clock, is familiar to many of us. Species and even
56 populations vary greatly in the way their clocks interact with the environment. They assume different
57 phases, e.g., of activity or reproduction, relative to the environmental cycle, and also differ in the use of
58 synchronizing cues. To varying degrees, organisms retain the ability to adjust their rhythms to respond
59 to current, less predictable, conditions. While some species' rhythms show considerable phenotypic plas-
60 ticity (e.g., the reproductive rhythm of Great tits, *Parus major*), other species have rigid rhythms that
61 impose fitness costs under rapid environmental change (e.g., the seasonal phenology of Snowshoe hare
62 coat color, *Lepus americanus*) [8, 10, 11]. In addition to phenotypic plasticity, evolutionary malleability
63 of biological rhythms is supported by directional evolution of time adjustments in multiple species, which
64 include heritable shifts in the seasonal timing of life history events such as reproduction, dormancy, and
65 migration [12, 13].

66 Biological rhythms are observed across biological processes. In addition to substantial diel and annual
67 fluctuations in activity, reproduction, and metabolism, there is also overwhelming evidence for temporal
68 structuring of immunity. Importantly, such fluctuations cannot be comprehensively characterized as
69 changes in overall immunity; rather, they are a selective re-organization of structural and functional
70 aspects of the immune system [14–16]. Differentiated temporal structuring of immune defenses can
71 arise from heterogeneous requirements and costs of specific defenses, investment in self-maintenance
72 versus immunity, or the integration of immunity with other aspects of physiology [17, 18]. In light of
73 this, we review biological rhythms pertinent to host-parasite interactions, and propose that rhythms of
74 hosts, parasites, and the environment impose temporal structure on epidemiological and evolutionary
75 dynamics.

76 **Timekeeping in the Host-Parasite Context**

77 Interactions between hosts and parasites (i.e., microparasites and macroparasites) are embedded within
78 environmental rhythms (Figure 1A). In addition to the environment, host immunity imposes selective
79 pressure on parasites, whilst parasite-driven morbidity and mortality reduces host fitness. These multiple
80 selective forces make optimal timing of allocation of limited resources to survival and reproduction

81 particularly tricky. For hosts, massive investment into parasite resistance, for instance, might only be
82 energetically feasible during a resource pulse (i.e., opportunity) also favorable for reproduction, resulting
83 in an optimization problem for resource allocation to survival versus reproduction [19]. Yet hosts also
84 undoubtedly face the challenge of mitigating the deleterious effects of parasites when resources are
85 scarce, a situation that might favor investment into parasite tolerance versus resistance. For parasites,
86 not only does the host immune response impose risk, additional risks can be introduced by environmental
87 regimes during transmission [20] or environmental life stages [21]; which has led to parasite risk avoidance
88 strategies such as climate-driven arrested development [22]. For both hosts and parasites, therefore,
89 external environmental conditions impose selective pressure by providing fluctuating opportunity for
90 reproduction and risk of mortality. These exogenous factors need not be identical for hosts and parasites,
91 although they co-occur in the same physical environment. For example, we need not expect that
92 rhythms in parasite reproduction, host reproduction, and host immune investment be synchronized. An
93 empirical case of this is the seasonal influence of temperature and humidity on development of the free-
94 living nematode parasite (*Trichostrongylus*) of rabbits, which results in an autumn peak in the force of
95 infection; whereas, the rhythm in host immunocompetence has a peak in the springtime [23].

96 The temporal structure of host immunity and parasite success suggests that constraints (1) preclude
97 hosts from maintaining high levels of parasite resistance, and (2) prevent parasites from sustaining
98 high reproductive output (fitness). Such constraints would result in trade-offs between investments in
99 opportunity versus risk avoidance [19, 24]. Consequently, we hypothesize that both hosts and parasites
100 time their biological processes with reference to both the external environment and each other, and that
101 therefore in many cases periodic incidence of infectious disease is a consequence of biological rhythms,
102 as has been suggested elsewhere (e.g., [25]). Below, we first lay out empirical evidence for the role
103 of rhythms in host-parasite interactions. In order to inspire quantitative study of biological rhythms in
104 host-parasite systems, we utilize a transmission model to illustrate the epidemiological consequences of
105 rhythms. We then formulate a conceptual evolutionary model for understanding host-parasite dynamics
106 embedded within the rhythmic context in which they are evolving.

107 **Biological Rhythms in Host and Parasite Traits**

108 The incidence of many infectious diseases displays substantial seasonality [26–28]. Seasonally struc-
109 tured disease incidence can be discussed from the viewpoint of hosts or parasites. From a host's perspec-
110 tive, parasite exposure can be influenced by host behavior, such as seasonal aggregation; a contemporary
111 example being epidemic seasonality of mycoplasmal conjunctivitis in house finches [29]. However, phys-
112 iological factors influencing host susceptibility to infection and symptomatic disease, such as seasonal
113 changes in immunity, can also drive disease seasonality [23, 28, 30]. Table 1 summarizes some known
114 diel and annual rhythms in host immunity and parasite traits (see [16, 31–34] for extensive reviews).
115 Although rhythms in immunity are observed across a broad array of taxa, including plants [35, 36] and
116 animals, we focus on mammalian and avian hosts to enhance the link to human health. A very active
117 area of biomedical research is characterizing temporal structure in both innate and adaptive immunity,
118 and correspondingly, in disease susceptibility [32, 33]. For now, biomedical studies use model organisms;
119 but in the future should include non-model organisms [37], which are either directly relevant for un-
120 derstanding natural host-parasite systems, or enable a broader understanding of within-host dynamics
121 of infection. Longitudinal studies of wild or captive animals compare immune parameters during active
122 versus resting phases and are typically combined with experimental approaches. In the wild these may
123 include repeated immune challenges [38–40], and in captivity may involve constant conditions, shifted
124 environmental rhythmicity, or biological clock disruption. Studies of rhythms in immunity under natural

125 conditions—wild immunology—are important for understanding how non-model organisms deal with
126 exposure to multiple co-occurring parasites [41]. Studies of wild systems allow us to test, for instance,
127 how seasonal allocation into defense against one parasite can result in enhanced susceptibility to an-
128 other [42], and whether temporal variation in immune status covaries with other physiological traits and
129 is influenced by nutritional status and parasite exposure [18, 43, 44]. Laboratory studies, in turn, are
130 necessary for distinguishing between endogenous rhythms in immunity versus variation that occurs as a
131 result of patterns of infection or other biotic factors.

132 By profiling the response to infection across time, much progress has been made in characterizing
133 biological rhythms in immunity. For example, in mice, the circadian rhythmicity of a receptor that rec-
134 ognizes pathogens substantially influences the inflammatory response and survival prospects (for details,
135 see Box 1). The health implications of circadian rhythms in immunity have also been demonstrated using
136 hosts entrained to different light-dark cycles, and mice with genetically modified circadian clocks. Such
137 recent studies have revealed that the immune system is fundamentally circadian in nature [33, 45–49],
138 which is highlighted by the local circadian clock of macrophages [46], and the feedback between immu-
139 nity and molecular, cellular, and behavioral rhythms. The emerging picture is that the immune system
140 is an active component of integrated whole-body circadian rhythms in animals [50] and plants [35, 36],
141 lending support to the idea that sophisticated mechanisms of immune defense were also present in their
142 common ancestor [51].

143 Annual cycles in immunity are not as well characterized as circadian cycles because of the time
144 scale of experimentation [16, 31] but are epidemiologically relevant [28]. Longitudinal studies under
145 controlled captive conditions have revealed substantial annual changes in immune parameters (Table 1).
146 These included a down-regulation of key aspects of immunity during the time of reproductive activation,
147 induced solely by photoperiodic simulation [14, 16, 34, 52]. Such rhythms might have evolved from a
148 trade-off between immune defense and demanding life-cycle stages, and can underlie annual patterns of
149 disease incidence, as suggested, for example, by rhythms of bactericidal capacity of whole blood (Box 1;
150 Figure 1D [53]). It is important to note, however, based on the existing evidence for both circadian and
151 annual immunomodulation, that temporal patterns can differ between innate and adaptive immunity
152 and among traits even within the same immune cell subset [16, 33].

153 In addition to immunomodulation, several other aspects of host rhythmicity can have population-scale
154 consequences for host-parasite dynamics. Relevant host annual cycles include aggregation [29, 54, 55],
155 sexual contacts (with regard to STDs), habitat use, migration [56–58], and birth pulses that (1) act
156 to replenish the pool of susceptible individuals, (2) can influence the critical community size required
157 for parasite persistence, and (3) can determine the geographic synchrony of outbreaks [59–63]. The
158 sweeping effects of annual cycles in host physiology on disease incidence are exemplified by White-
159 Nose Syndrome (WNS), which is drawing many North American bat species near extinction. A new
160 longitudinal study indicated that neither birth pulses nor social behavior affected transmission and
161 intensity of WNS. Instead, WNS is associated with hibernation [30], which in mammals that have been
162 studied in captivity, is a programmed circannual rhythm [6]. We speculate the link between WNS and
163 hibernation is mediated by hibernation-associated changes in immunity. Evidence thus far suggests that
164 there are large adjustments in immunity in hibernating mammals, including a 90% decrease in circulating
165 white blood cells [64], down-regulation of the acute-phase response to LPS [65], and modifications of
166 intestinal immunity [64, 66].

167 Migration is another host rhythmicity receiving attention in infectious disease ecology. In monarch
168 butterflies, the protozoan parasite *Ophryocystis elektroscirrha* displays a seasonal pattern of prevalence
169 and a spatial gradient along the monarchs' migratory flyway. Parasite prevalence declines as monarchs
170 migrate, which is likely due to migratory culling [57]. In migratory culling, the coupled energetic
171 demands of migration and fighting infection result in increased mortality of infected individuals during
172 fall and spring migrations. The uninfected are most likely to survive the journey to the breeding or
173 wintering grounds, allowing the destination to be relatively parasite-free. Thus, migratory culling is a
174 direct intersection of host seasonal rhythms and disease prevalence [56], and anthropogenically-driven
175 disruption of this rhythm results in elevated disease burden [58].

176 Taking the parasite's perspective, rhythmic patterns in parasite dissemination can be influenced by
177 fluctuating abiotic and biotic conditions that affect parasite survival and transmission. Clear examples
178 of abiotic influences are (1) the role of temperature and humidity in transmission of influenza [67],
179 which might be responsible for latitudinal clines observed in influenza incidence [68, 69], and (2) the
180 UV sensitivity of sporulation in *Isospora*, which might have driven the remarkably robust diel pattern of
181 oocyst outputs [21]. In addition to abiotic effects, biotic influences can stem from rhythms of vectors [70]
182 and other parasites [71]. Effects of vector circadian rhythmicity have been studied in the malaria vector
183 *Anopheles gambiae*, whose rhythmic gene expression persists under constant conditions. Rhythmically
184 expressed genes include those implicated in the melanization immune response, which encapsulates
185 the Plasmodium parasites, and can thereby affect mosquito to human transmission. Vectors can also
186 temporally structure parasite transmission via their diel patterns of feeding [72] and their phenology [73].
187 Interspecific influence of parasites on one another's rhythm, to our knowledge, has only been described
188 for *Drosophila* parasitoids, which gain a fitness advantage by temporally segregating circadian rhythms
189 in egg oviposition [71], hypothesized to alleviate competition.

190 Perhaps then unsurprisingly, parasites—faced with rhythms in their abiotic environment, hosts, and
191 vectors—display what seem to be biological rhythms. Documentation of parasite rhythms dates back
192 over 100 years, long before the discovery of biological clocks. In fact, the early observation that
193 both malaria parasites and microfilariae are abundant in the blood of hosts at night was instrumental
194 to the discovery of mosquitoes as the malaria vector [74]. Experimental studies of parasites report
195 diel and annual rhythms, as measured by fluctuations in parasite burden and infectivity (Table 1),
196 but disentangling the contributions of host and parasite to these rhythms is difficult [75]. To our
197 knowledge, the only described example of a parasite life history event that depends on a host rhythm is
198 reproduction in the ectoparasitic rabbit flea, a vector of myxoma virus. To reproduce, rabbit fleas must
199 undergo maturation on a pregnant or newborn nestling host, and flea maturation is controlled by host
200 hormone cues associated with pregnancy and parturition, thereby synchronizing parasite and host life
201 cycles [76–78]. There is solid evidence for adjustment of diel parasite rhythms to those of the host, for
202 example from trematodes like *Schistosoma mansoni*—an agent of schistosomiasis in humans—whose
203 emergence from snail hosts is initiated by light [79, 80]. These parasites display diel cycles that shift to
204 match perturbations in their hosts' circadian rhythm [81]. Similar results were found in rodent-infecting
205 *Trypanosomes*. Rats infected with *T. lewisi* and housed under a normal light-dark cycle (LD 12:12
206 h) experienced a peak in circulating *T. lewisi* during the early part of night and a trough in the early
207 morning. However, when the host photoperiod was inverted, the parasite rhythm was also reversed [82].
208 In *Isospora*, the characteristic diel pattern of oocyst output persisted under continuous light in the high
209 Arctic, although feeding and activity rhythmicity of their avian hosts was greatly diminished, suggesting
210 synchronization to subtle host rhythms or possibly self-sustained parasite rhythms [21].

211 Experimental studies give clear evidence that synchronization to host rhythms impacts parasite fit-
212 ness. For example, murine host rhythms were experimentally mismatched to that of their malaria parasite
213 (*Plasmodium chabaudi*). This mismatch resulted in a 50 per cent reduction in both parasite replication
214 and production of transmissible life-stages [83]. Follow up experiments have now revealed additional
215 complexities, with the effect of mismatch manifesting differently between parasite life stages, and down-
216 stream effects on host disease severity. Mismatch can confer a substantial cost to the parasite, and this
217 cost is experienced at the onset of infection, rather than acquired throughout infection [84, 85]. This
218 suggests there can be intense selective pressure on parasites to maintain a specific phase position relative
219 to their host rhythms, or to vector rhythms, since parasite ability to infect vectors is also time-of-day
220 dependent [72]. Fortunately, the amassing knowledge of biological clocks might help identify host cues
221 used for entrainment of parasite rhythms. For example, the nocturnally-peaking hormone melatonin
222 is a core circadian feature of many vertebrates, and applying this knowledge produced indication that
223 parasites might be using melatonin to synchronize their circadian cell cycle [21, 86].

224 While we still lack unambiguous evidence for endogenous circadian or circannual rhythms of parasites,
225 recent research suggests the intriguing possibility that parasites can actively manipulate host and vector
226 rhythms to their advantage. For example, various parasites interrupt host diel activity at specific times
227 of day to enhance transmission [87, 88]. The diel timing and synchrony of host behavioral manipulation,
228 along with candidate molecular mechanisms of manipulation, strongly implicate circadian clock pathways
229 [87, 89]. By integrating chronobiology with infectious disease ecology, we might be able to identify,
230 for example, the mechanism by which the trematode *Dicrocoelium* manipulates diel host behavior,
231 inducing suicide, and facilitating trophic transmission [90], and how the notoriously manipulative fungus
232 (*Ophiocordyceps unilateralis s.l.*) seemingly breaks the host circadian clock to perpetuate transmission
233 [87]. Transkingdom cross-regulation between prokaryotic and eukaryotic rhythms is plausible because
234 it has already been documented for other systems (e.g., in bioluminescent squid light organ symbionts
235 and in mammalian gut microbiota) [4, 91].

236 Despite our knowledge of (1) rhythmic host immunity and physiology, (2) rhythms in parasite repro-
237 duction and transmission, and (3) enticing evidence that host rhythms can impact parasite fitness and
238 be exploited by parasites, the effects of biological rhythms on host-parasite dynamical processes remain
239 poorly understood. We surmise that careful consideration of biological rhythms in infectious disease
240 ecology and evolution will provide a better understanding of (1) daily and annual patterns of diseases,
241 (2) within-host parasite dynamics, and (3) parasite transmission.

242 **Models for Investigating Host-Parasite Contributions to Rhythms in In-** 243 **fectious Disease**

244 In order to determine how biological rhythms impose temporal structure on host-parasite dynamical
245 processes, we can integrate empirical data on host and/or parasite rhythms into epidemiological and
246 evolutionary models. Biological rhythms research has great potential for feedback between laboratory
247 studies, field ecology, and dynamical systems modeling. First, rhythms in immunity characterized under
248 laboratory or field conditions can be used in transmission models to make predictions about the epi-
249 demiological consequences of those rhythms in nature. Second, observations of diel and annual cycles in
250 infection—characterized via disease incidence, parasite abundance, or host serological markers of infec-
251 tion history—can be used to make predictions regarding rhythms generating such patterns. A new study
252 of the first type [92] explores the effect of annual and biannual rhythms in births (in bats) on the per-
253 sistence of filoviruses (i.e., Marburgvirus and Ebolavirus). Transmission models predict that filoviruses

254 can persist in species with biannual birth pulses—making them potential reservoirs of infection—and
255 this prediction is supported by serology data showing that species with biannual birth pulses are more
256 likely to be seropositive for filoviruses; demonstrating that explicit consideration of host rhythms can
257 inform targeted surveillance and control of emerging zoonotic diseases. A study of the second type is
258 that of [23], in which long-term field data on nematode infections in European rabbits were used to
259 discriminate among multiple potential seasonal rhythms in the host-parasite system. This led to identi-
260 fication of epidemiologically relevant seasonality in host immunity; the endogenous nature of which can
261 be tested in the lab.

262 Building upon the examples above, as well as other transmission models that incorporate reproduc-
263 tive rhythms [59–61, 93, 94], here we provide a Susceptible-Infected-Recovered (SIR) model of a directly
264 transmitted hypothetical bacterial infection in the Siberian stonechat to illustrate the numerous entry
265 points for biological rhythms into epidemiological processes (Figure 1B). We narrate our model with
266 seasonal rhythms in mind; however, this can be extended to circadian rhythms. The model incorporates
267 ambient temperature as a covariate influencing parasite transmission as well as empirical data on host
268 circannual cycles in reproduction, immunity and migration (Figure 1CD; cf. Box 1) [53, 95]. For migra-
269 tion, the timing is defined empirically, while the model assumes migratory culling of infected individuals
270 only during the autumn migration, when host bacterial killing activity is lowest [53]. Importantly, we
271 propose that circannual cycles in host immunity can influence (1) the transmission rate, (2) the recovery
272 rate, and (3) the pathological consequences of infection, which manifests as symptomatology and en-
273 ters the model as the report rate. The multiple rhythms: temperature, births, bacterial killing activity,
274 and migratory culling act collectively to shape the observed incidence of disease, which is the model
275 output shown in Figure 1E. We define the resulting seasonal window of elevated disease incidence as
276 the parasite’s temporal niche. The seasonal incidence that arises from this model matches the expec-
277 tations from the underlying data. However, in contrast to most models of seasonal infectious diseases,
278 which only place sinusoidal seasonality in the transmission rate, it contains multiple axes of seasonal
279 forcing. Thus, we provide this model to encourage the inclusion of empirically characterized rhythms
280 into models as covariates. Such models can be used to explore the epidemiological consequences of host
281 and parasite rhythms, although for simplicity the parasite is not explicitly modeled here. Host-parasite
282 systems where modeling parasite rhythms is particularly compelling include nematodes with seasonal
283 arrested development [22, 96], microfilariae which display both circadian and seasonal cycles [97], and
284 *Plasmodium* within-host circadian cycles [75]. Major challenges of incorporating biological rhythms
285 into epidemiological or within-host models, will be (1) recognizing which host and/or parasite rhythms
286 are epidemiologically relevant, and (2) identifying the functional relationships between rhythms and
287 epidemiological parameters, including: transmission, recovery, and symptomatology.

288 In addition to the epidemiological consequences of rhythms, we can benefit from understanding the
289 feedback between host and parasite rhythms and the multiple axes that shape their temporal structure.
290 Thus, we provide a conceptual evolutionary model for understanding how hosts and parasites time their
291 biological processes with reference to each other while being embedded in environments with temporally
292 structured risk and opportunity. We pose this model in evolutionary terms, but depending on the varying
293 degrees of plasticity of biological rhythms, individuals may also adjust their rhythms during their life.

294 Our evolutionary model illustrates three idealized scenarios of how host immune defense varies sea-
295 sonally, relative to fluctuating environmental conditions (Figure 2A–C). These scenarios are motivated
296 by life history theory and by empirical observations of seasonal immunity in mammals and birds (Table
297 1). The scenarios assume that immune defense, specifically, parasite resistance, either parallels the

298 availability of resources (A, “resource-driven”), or is reduced when resources are used for reproduction
299 (B, “traded-off”). The third is an extension of the “resource-driven” scenario with modulation related
300 to life history events that can lead to complicated, but potentially important, annual patterns. In our
301 example of migration (C) we assume down-regulation of immune defense during migration (see Figure
302 1; this could also occur during other vulnerable times such as molting or hibernation [65]), and a shall-
303 low trough under favorable conditions in the wintering grounds. We then (D) switch to the parasite’s
304 perspective and illustrate how parasites are subject to two axes of seasonal fluctuations: (i) seasonal
305 environmental conditions outside the host and, (ii) seasonal immune defense of the host. We propose
306 that together the two seasonal axes shape parasite transmission (i.e., parasite fitness; which is captured
307 by the basic reproductive number).

308 The last and crucial component of our model is the evolutionary feedback between host and parasite
309 rhythms. We propose that due to parasite-induced host morbidity and mortality, selection can drive
310 changes in host seasonal immune defense. Subsequently, since host immune defense is one of the
311 seasonal axes influencing parasites, selection will favor changes in the parasite rhythm. This interplay
312 can continue, driving hosts and parasites to sequentially alter their seasonal rhythms while working
313 within the constraints of environmental conditions. Figure 2E shows these steps.

314 We suggest that under certain conditions this can escalate into an evolutionary arms race. In this
315 framework, the prerequisite for an arms race is that parasite fitness is sufficiently impacted by the
316 temporal structure of the host immune response, and that the host immune response is predictably
317 rhythmic. To be clear, when considering the temporal structure of immune defense, reference to “low”
318 host immune defense pertains to the parasite in question. However, it must be appreciated that a
319 time of diminished resistance to one parasite (e.g., a helminth) can be a time of high investment into
320 fighting another (e.g., a virus). Furthermore, infection with one parasite can seasonally elevate host
321 susceptibility to another, as is exemplified by concomitant infections of myxoma virus and nematodes [98]
322 and increased susceptibility to bovine TB resulting from helminth coinfection [18]. The arms race itself
323 has two requirements. First, hosts must be able to shift their immune defense to counter exploitation
324 by parasites (host changes in Figure 2E). Changes in host rhythms then translate into a new landscape
325 of time-structured risk and opportunity for parasites. Upon experiencing a new temporal landscape, a
326 dynamic host-parasite arms race can arise only if the second requirement is met: parasites shift their
327 rhythm by changing reproduction within hosts, or release from hosts (parasite changes in Figure 2E).
328 As with other host-parasite arms races, an arms race in the temporal domain is subject to tradeoffs for
329 both the host and the parasite that might constrain the extent to which their rhythms can be altered.
330 Host tradeoffs can include an immunity-reproduction tradeoff [99]; whereas, tradeoffs for the parasite
331 can include a transmission-virulence or transmission-recovery tradeoff [99–101]. Also, due to the rapid
332 generation time of parasites, relative to hosts, evolution of host rhythm shifts might be slow relative to
333 the evolution of parasite rhythms, but this would not preclude an arms race from occurring.

334 Conclusion

335 There is enticing evidence that biological rhythms are structuring elements of host-parasite interac-
336 tions, both in within-host processes and in epidemiological dynamics. Host circadian rhythms in the
337 immune system influence the progression of infection and parasite burden, and annual rhythms might
338 have similar effects [23, 30]. The existing evidence leads us to conclude that the effects of biolog-
339 ical rhythms on the perpetuation of parasites, and on host reactions during infection, can generate

340 population-level rhythms in infectious disease incidence [25], which we here define as the parasite's tem-
341 poral niche. To formalize temporal niches across parasite taxa and life history strategies, we will need
342 novel integration of epidemiological, immunological, and life history data of both hosts and parasites.

343 Importantly, circadian rhythms in immunity have direct implications for transmission, and practical
344 application for (i) timing of antibiotic, antiviral, and anthelmintic treatment, and (ii) managing im-
345 munopathology, such as cytokine storms. The rhythms of parasites themselves can drive patterns of
346 exposure and illness, as is evident in malaria and filarial infections. Similarly, rhythms in parasitemia
347 and parasite release from hosts can impose temporal structure on transmission, which can be leveraged
348 for interventions such as deworming campaigns.

349 We believe that a multi-disciplinary approach at the intersection of Chronobiology, Disease Ecology
350 and Evolutionary Biology holds the key to understanding how biological rhythms influence host-parasite
351 interactions. We have outlined open questions that will bring us closer to understanding the underlying
352 biological interactions in the temporal domain (refer to Outstanding Questions Box). We hope that
353 this Opinion will generate discussion on how to leverage rhythms for translational medicine, for instance
354 to counter the evolution of resistance; and we hope the insights provided here inspire new avenues for
355 interrogating transmission models with host-parasite data from the laboratory and the field, ultimately, to
356 better understand the forces structuring disease incidence and the immunology of non-model organisms.

357 Finally, although it is beyond the scope of this Opinion, the importance of accounting for biolog-
358 ical rhythms is accentuated by the accumulating data on anthropogenically-driven disruptions and
359 mismatches of biological rhythms that are occurring across taxa. Circadian disruption due to light-
360 at-night [102] and altered environmental seasonality due to climate change [13] are challenging the
361 plasticity of rhythms and modifying the fitness advantages of their endogenous basis. For example, the
362 adverse effects of circadian disruption have already been seen in human health and gut microbiota [91].
363 Given the pervasiveness of rhythms in host immunity, vectors and parasites, we might soon be faced
364 with palpable effects of rhythm disruptions on infectious diseases [73, 103, 104].

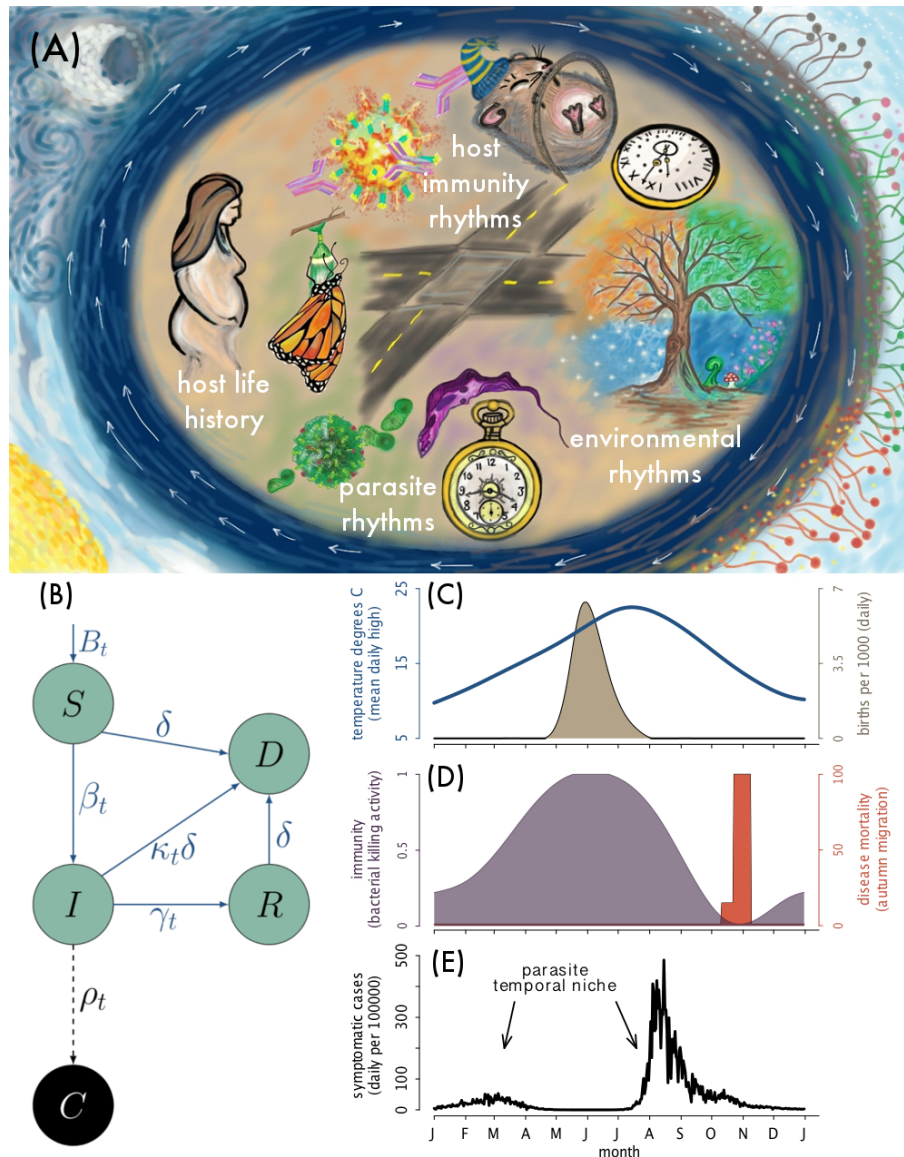


Figure 1: Rhythms and Temporal Niche. (A) The timing of host and parasite activities falls in the intersection of environmental rhythms, host life history, host immunity rhythms, and parasite rhythms. This intersection is embedded within geophysical rhythms, diel and annual cycles. (B) Biological and environmental rhythms can enter into epidemiological models in multiple ways. The schematic shows a *Susceptible-Infected-Recovered* model, *SIR*, with natural and disease-induced deaths, *D*. The model distinguishes between infections, *I*, and the subset of infections that are observed as symptomatic cases, *C*. The model is parameterized using the life history of Siberian stonechats. Four seasonal rhythms enter into the model (births, temperature, immunity, and migration). Host births, B_t , are seasonal. The transmission rate, β_t , is a function of (1) an environmental rhythm (i.e., temperature) that influences parasite transmissibility, and (2) the seasonal immune status of hosts. We assume seasonal immunity also influences the recovery rate, γ_t , and the probability of symptoms, ρ_t . We also assume infected individuals suffer disease-induced mortality, κ_t , associated with the autumn migration (i.e., migratory culling), which multiplies the (here constant) rate of natural mortality δ . (C) Annual fluctuations in temperature and birth seasonality in Siberian stonechats [95]. (D) Annual host immunity is based on bacterial killing activity [53], elevated mortality during autumn migration is inferred from natural migratory timing. (E) Incidence of symptomatic cases assuming: temperature has a positive correlation with transmission, bacterial killing activity reduces transmission, reduces the probability of symptoms, and increases the recovery rate. The four seasonal rhythms act collectively to determine the parasite's temporal niche, the time of year when the parasite is abundant and disease outbreaks occur.

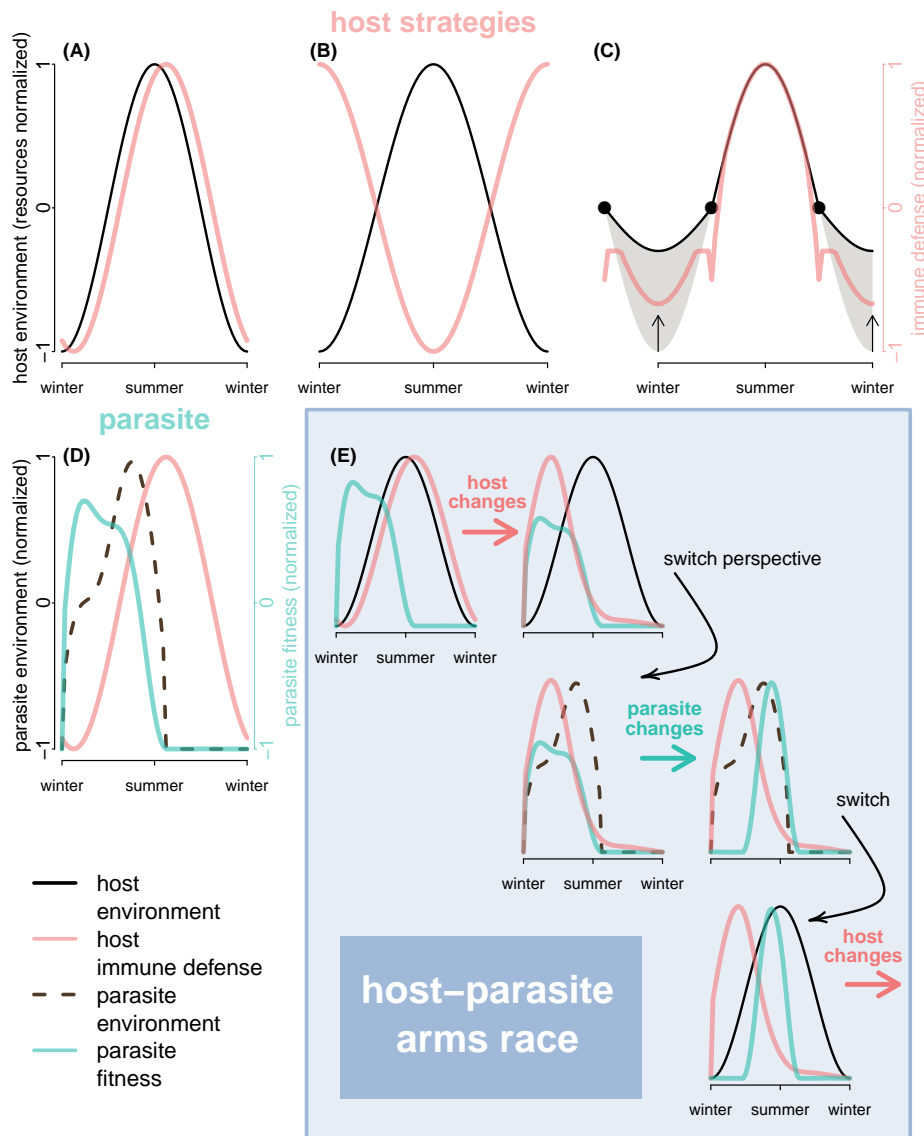


Figure 2: Conceptual model for investigating host-parasite contributions to rhythms in infectious disease.

(A) Host immune defense is resource-driven and tracks the host's environmental conditions (i.e., host resource availability). (B) Host immune defense has an inverse relationship with environmental conditions; this could occur due to a trade-off against investment into reproduction during high resource availability. (C) Resource-driven immune defense in a migrating species that has reduced immune defense during migration. Migrations (indicated by black points) result in shallower environmental troughs since individuals migrate to regions with higher resource availability. For all scenarios, we consider immune defense to be resistance to the parasite in question, although we acknowledge that this simplifies the complexity of the immune system (e.g., independent immunomodulation of innate or adaptive immune parameters). For the resource-driven host immune defense strategy, in (D) we show seasonal parasite fitness shaped by both environmental conditions and seasonal host immune defense. Although host and parasite co-occur in the same physical environment, the environmental rhythms pertinent to the parasite need not be identical to the environmental rhythms pertinent to the host, which is why we distinguish host versus parasite environmental rhythms. (E) Arms race between host and parasite. For illustrative purposes, the arms race is initiated with resource-driven host immune defense and parasite seasonality. The host changes the seasonal timing of peak immune defense to coincide with peak parasite fitness. We then switch to the parasite perspective to consider the parasite's environment. In response to the new host seasonality, the parasite changes its timing of peak reproductive output. These cycles can continue, with both host and parasite seasonality shifting within the bounds of their respective environmental constraints.

Table 1: Rhythms in Hosts and Parasites. Diel (circadian) and annual (circannual/seasonal) rhythms in host immunity, parasite reproduction, and parasite release.

| Type | Period | Host Trait or Parasite Description | Organism/Species | Rhythm | Citation |
|----------|--------|---|---|--|--------------|
| Immunity | Diel | Macrophages (detection and restriction of parasite invasion) | Mice | 8% of transcripts are circadian; autonomous macrophage circadian clock controls rhythm | [46] |
| Immunity | Diel | Natural Killer Cells (early defense against viruses and intracellular bacteria) | Humans | Circadian trafficking between the blood and organ compartments; inverse trafficking compared to T-lymphocytes | [105] |
| Immunity | Diel | Toll-like receptor 9 (evolutionarily conserved receptor that recognizes bacteria and viruses) | Mice | Expression and function controlled by circadian clock | [47] |
| Immunity | Diel | T-lymphocytes (surveillance for infected cells) | Humans | Cytokine production | [106,107] |
| Immunity | Diel | Leukocytes | Humans | Abundance in the blood follows a circadian rhythm for neutrophils, lymphocytes, monocytes, and eosinophils | [108] |
| Immunity | Diel | Whole blood response to LPS stimulation | Humans | Cytokine and chemokine production is circadian in an environment free of time cues | [109] |
| Immunity | Diel | <i>Salmonella</i> colonization and host cytokine response to infection | Mice | Mice have a different immunological response to infection depending on whether infection challenge occurs at day or night; infection during the day results in more inflammation; this effect is due to clock-controlled gene expression | [49] |
| Immunity | Diel | Clock genes and pro-inflammatory cytokines in spleen; inflammatory response | Birds (captive) | mRNA of cytokines and clock genes are rhythmic under LD cycles and constant conditions; inflammation is rhythmic under LD cycles | [110] |
| Immunity | Diel | Cellular (PHA) and humoral immune response | Birds (captive) | PHA response and antibody production depend on time of challenge, but their peaks are phase-inversed | [17] |
| Immunity | Annual | Bacterial killing activity | Birds (captive), turtles (wild), humans | Lower bactericidal activity during migration, especially in autumn (birds); Higher bactericidal activity during breeding season (turtles); Higher bacterial killing by neutrophils in summer (humans) | [53,111–113] |
| Immunity | Annual | Leukocytes | Birds (captive) | Annual cycles in several immune traits, no effect of <i>Coccidia</i> on annual cycle of immune measures | [114] |
| Immunity | Annual | Lysis | Birds (wild) | Lower ability of plasma to lyse foreign cells during migration and winter | [38] |
| Immunity | Annual | Sickness behavior in response to LPS | Birds (captive and wild), hamsters | Repression of sickness behavior during reproduction in summer (birds); repression of sickness behavior during winter (hamsters) | [39,115] |
| Immunity | Annual | Acquired Immunity | Rabbits (wild) | Resistance against nematodes | [23] |
| Immunity | Annual | Spleen size (spleen is important for both innate and adaptive immunity) | Birds (wild) | Regression of spleen during migration | [116] |

Continued on next page

| Type | Period | Host Trait or Parasite Description | Organism/Species | Rhythm | Citation |
|-----------------------|--------|--|---|---|------------------------|
| Immunity | Annual | Cytokine production stimulated by bacterial endotoxin | Humans, rats, hamsters | Seasonal differences in pro- and anti-inflammatory cytokine production (humans); Summertime photoperiod increases production of pro-inflammatory cytokine TNF- α and extends (rats) or elevates (hamsters) disease symptoms | [117–119] |
| Immunity | Annual | Vaccine response | Humans | Seasonal variation in symptoms following live influenza vaccine | [120] |
| Immunity | Annual | Intestinal immunity | Ground squirrels (captive) | Increase in intestinal leukocytes, pro- and anti-inflammatory cytokines during hibernation | [66] |
| Susceptibility | Diel | Bacterial burden, pathogenesis, and/or virulence of infection | Mice | Timing of infection can affect: (a) bacterial burden due to circadian variation in monocyte trafficking and/or gene expression at site of infection, (b) disease severity from sepsis due to circadian TLR9 expression, and (c) virulence | [45, 47, 49, 121, 122] |
| Susceptibility | Annual | Susceptibility to fungal growth | Bats (wild) | Hibernating bats have temperatures that match that of hibernacula, allowing explosive growth of WNS fungal pathogen <i>Pseudogymnoascus destructans</i> | [30] |
| Parasite reproduction | Diel | <i>Plasmodium</i> species (Malaria parasite) asexual reproduction | Various mammalian hosts | Parasite cohorts of millions of individuals synchronously burst from red blood cells at a particular time in LD cycle | [75] |
| Parasite reproduction | Annual | Microfilariae (heartworm) | Dogs | Strong seasonal rhythm in microfilaria abundance in dogs infected in the lab | [97] |
| Parasite development | Annual | Nematodes (parasitic roundworms) | Domestic mammals | Parasites engage in seasonal hypobiosis, arrested development within hosts that allows for persistence when environmental conditions are unfavorable for transmission between hosts | [96] |
| Parasite discharge | Diel | Coccidia | Birds (wild and captive) | Oocyte release is strictly circadian; although parasites are vulnerable to sun exposure the rhythmic pattern persists under continuous light during the Arctic summer | [21, 123] |
| Parasite discharge | Diel | <i>Echinostoma</i> (parasitic flatworms), Pinworms, <i>Schistosoma</i> | <i>Echinostoma</i> in mice, Pinworms and Schistosomes in humans | The release of <i>Echinostoma</i> eggs by hosts occurs during night when mice are active. Human pinworms migrate out of anus during the night to lay eggs. In contrast, <i>Schistosoma</i> eggs are discharged in urine during the day. | [74, 124] |
| Parasite discharge | Annual | Nematodes (roundworms) | Dall's Sheep (wild) | Seasonal variation in intensity of parasite larvae shed in feces | [125] |

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| Type | Period | Host Trait or Parasite Description | Organism/Species | Rhythm | Citation |
|--|--------|---|--------------------------|--|----------|
| Parasite manipulation of host behavior | Diel | <i>Dicrocoelium</i> trematode | Ants (intermediate host) | In the evening infected ants affix themselves to the top of blades of grass and enter torpor until the next morning, allowing them to be eaten by grazing mammals (definitive hosts) | [90] |
| Parasite manipulation of host behavior | Diel | Manipulating fungus <i>Ophiocordyceps</i> | Ants | <i>Ophiocordyceps</i> manipulates host behavior and causes host death at characteristic times of day | [87] |

365

366 **Box 1. Circadian and Seasonal (circannual) Modulation of Host Immune** 367 **Defense.**

368 **Circadian Immune Cycles.** In order for hosts to mount an immune response against an infecting
369 pathogen, the immune system must first detect the presence of the pathogen. One way that animals
370 detect pathogens is by immune surveillance for pathogen-associated molecular patterns (PAMPs), which
371 are shared across groups of pathogens. Host cells express pattern recognition receptors (PRRs) that
372 recognize PAMPs. Toll-Like Receptor 9 (TLR9) is an important PRR that can recognize both viruses
373 and bacteria, and is highly evolutionarily conserved. In 2012 Silver et al. discovered that the expression
374 of TLR9 by macrophages and B cells follows a circadian rhythm. Importantly, the circadian rhythm of
375 TLR9 has a significant effect on the immune response and disease severity because the rhythm of TLR9
376 also produces a rhythm in inflammatory cytokines. The implications were experimentally demonstrated
377 by inducing sepsis. Sepsis can occur during bacterial infections when a severe inflammatory immune
378 response causes damage to the host. Bacterial infection was induced in laboratory mice using a puncture
379 that allowed commensal bacteria to enter the body cavity from the intestine. Mice were entrained to a
380 light-dark cycle (LD 12:12 h), and infection was induced either at the midpoint of the light period or
381 of the dark period. Mice that were infected during the night, when the TLR9 inflammatory response
382 was elevated, had higher bacterial burdens, earlier mortality, and worse disease scores, hypothermia, and
383 tissue damage than mice that were infected during the day. This study demonstrated that the functional
384 response of the immune system varies according to a circadian rhythm, and this variation is biologically
385 relevant because it can have a significant effect on the dynamics of infection [47].

386 **Annual Immune Cycles.** Faced with stark annual fluctuations in environmental conditions and
387 resources, many avian and mammalian species partition life history events such as reproduction, growth,
388 and hibernation into distinct times of year, and their immune system also undergoes seasonal changes.
389 Versteegh et al. 2014 set out to investigate whether annual variation in immunity is due to seasonal
390 adjustments directly driven by environmental or physiological conditions, or originates from a genetically-
391 based circannual rhythm that allows organisms to prepare for changes in the environment. They looked at
392 5 different immune measures, including bactericidal competence of whole blood as a proxy for functional
393 implications.

394 To determine whether seasonal immunity is a genetically encoded circannual rhythm, genetically
395 distinct subgroups of a widespread songbird, the stonechat (*Saxicola torquata*), were bred and raised
396 in a common garden experiment. The subgroups chosen for this experiment differ in their seasonal life

397 history and traits. They included a (i) long-distance migrant, (ii) short-distance migrant, and (iii) a
398 non-migrant, along with hybrids. The prediction was that if seasonal immunity is a direct response to
399 seasonal environmental conditions and energy demands, then by raising birds in an environment where
400 (a) they have ample food, (b) they are not allowed to migrate or breed, and (c) the only fluctuation to
401 which they are exposed is changing day length, their annual rhythms in immunity would be lost.

402 The authors found that not only did the annual rhythm in immunity persist under these controlled
403 conditions but also that the subgroups and hybrids of the birds showed specific patterns. The long-
404 distance migrants displayed seasonality in 4 immunity parameters, which included bacterial killing ability
405 (Figure 1D). The short-distance migrants displayed seasonality in only 3 immunity parameters, and the
406 non-migrants displayed seasonality in only 2 parameters. Furthermore, the amplitude of the annual
407 fluctuation was greatest in the long-distance migrants. The inheritance of the rhythm in hemolysis (the
408 ability of antibodies and their complement system to lyse foreign cells) was also quite striking. Both
409 the long- and short-distance migrants showed reduced hemolysis during the time of the natural autumn
410 migration. The reduction in hemolysis in the long-distance migrants was much more extreme than that
411 of the short-distance migrant, and intermediate in F1- hybrids. Together, this work demonstrates that
412 an inherited, biological clock controls seasonal immunity in stonechats, and these rhythms vary across
413 groups that differ in their seasonal life history [53]. Related avian studies generally confirm annual cycles
414 in immune parameters, and although species differ, there is a common tendency for greater seasonal
415 immunomodulation with increasing migratory lifestyle.

Outstanding Questions.

1. Are rhythms in the immune system adaptive for fighting infection?
2. Are parasite rhythms adaptive for dealing with host rhythms or environmental conditions?
3. Are observed parasite rhythms truly endogenous?
4. If parasite rhythms are endogenous, are they entrained by host rhythms?
5. Are host circannual rhythms in immunity an adaptive response to (a) seasonal parasite exposure or (b) resource limitations and life history trade-offs?

Glossary.

Adaptive immune system - cells of the adaptive immune system include B cells and T cells. The initiation of the adaptive immune response occurs after the initiation of the innate immune response. Receptors of adaptive immune cells require genetic recombination and alteration to generate, resulting in antigen specificity and immunological memory [126,127].

Annual cycle - cycle with a length of approximately 1 year, with phases occurring consistently at a particular time every year, on an annual/seasonal basis

Circadian - cycle that occurs with an approximate 24-hour period, used in reference to endogenous rhythms

Circannual - cycle that occurs with a period of approximately 1 year, used in reference to endogenous rhythms

Diel cycle - cycle with a length of approximately 1 day, with phases occurring consistently at a particular time during the day-night cycle

Immune defense - immune defense includes (1) resistance against the establishment of infection and the reproduction of parasites, and (2) parasite tolerance, in which the host mitigates the pathological consequences of infection, but tolerates infection

Innate immune system - cells of the innate immune system include macrophages and neutrophils. Innate immune cells are immediate responders to infection. A fundamental distinction between innate and adaptive immune cells is that innate immune cell receptors responsible for immune recognition are encoded in the germline; whereas, receptors of adaptive immune cells require genetic recombination and alteration to be generated. It was previously thought that the innate immune response is not parasite-specific and lacks memory, but that characterization is now considered incorrect [127].

LD-cycle - cycle in which light and darkness alternate, and each last for a given duration, for example in LD 12:12 h, light and darkness both last for 12 hours

LPS - Lipopolysaccharide is a component of the outer membrane of Gram-negative bacteria that is used in experiments to elicit an anti-bacterial immune response

Macroparasites - parasites that are large and typically metazoans (e.g., helminths)

Microparasites - parasites that are small and often unicellular (e.g., pathogenic viruses, bacteria, and fungi)

Macrophage - phagocytes, often referred to as big eaters because they engulf invading bacteria and are responsible for clearance of dead (apoptotic) cells. Macrophages are one of the cells responsible for detection and restriction of parasite invasion.

Parasite Resistance - The ability of the host's immune response to prevent infection from establishing or limit parasite replication. Parasite resistance has a negative impact on parasite fitness [128].

Parasite Tolerance - The ability of the host to mitigate the pathological consequences of infection, rather than mitigate infection itself. Parasite tolerance does not necessarily have a negative impact on parasite fitness [128].

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