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Title: Photoperiodic regulation of avian physiology: from external coincidence to seasonal reproduction.

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Running Title: Photoperiodic regulation of avian seasonality

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Data availability statement: The data that support the findings of this study are available from the corresponding authors upon reasonable request

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Research highlights:

- Two pathways are involved in the external coincidence of avian photoperiodism, a light-detection and an interval timer pathway.
- D- and E-Box binding motifs in *EYA3* suggest a connection between the circadian clock and initial photoinduction.

26 **ABSTRACT**

27 Seasonal cycles of environmental cues generate variation in the timing of life history transition
28 events across taxa. It is through the entrainment of internal, endogenous rhythms of organisms to
29 these external, exogenous rhythms in environment, such as cycling temperature and daylight, by
30 which organisms can regulate and time life history transitions. Here, we review the current
31 understanding of how photoperiod both stimulates and terminates seasonal reproduction in birds. The
32 review describes the role of external coincidence timing, the process by which photoperiod is
33 proposed to stimulate reproductive development. Then, the molecular basis of light detection and the
34 photoperiodic regulation of neuroendocrine timing of seasonal reproduction in birds is presented.
35 Current data indicates that vertebrate ancient opsin is the predominant photoreceptor for light
36 detection by the hypothalamus, compared to neuropsin and rhodopsin. The review then connects light
37 detection to well characterized hypothalamic and pituitary gland molecules involved in the
38 photoperiodic regulation of reproduction. In birds, Gonadotropin-releasing hormone synthesis and
39 release is controlled by photoperiodic cues via thyrotropin-stimulating hormone- β (TSH β)
40 independent and dependent pathways, respectively. The review then highlights the role of D-box and
41 E-box binding motifs in the promoter regions of photoperiodic genes, in particular Eyes-absent 3, as
42 the key link between circadian clock function and photoperiodic time measurement. Based on the
43 available evidence, the review proposes that at least two molecular programs form the basis for
44 external coincidence timing in birds: photoperiodic responsiveness by TSH β pathways and
45 endogenous internal timing by gonadotropin synthesis.

46

47 **Keywords:** Birds, photoperiod, opsins, coincidence, GnRH

48

49 Introduction:

50 Annual seasonal changes encompass a wealth of environmental cues which may contribute to
51 timing seasonal physiology and include ambient temperature, nutritional availability, and the
52 changing length of daylight (Nichelmann et al 1999, Helm and Gwinner 1999, Helm and Gwinner
53 2005, Flinks et al 2008, Helm et al 2009). This is especially true for migrating birds in which the
54 timing of life history transitions is dependent on the monitoring and response to changing
55 environmental conditions over time (Gwinner and Dittami 1990, Raess and Gwinner 2005, Helm
56 2009). The annual change in daylength is the predominant predictive cue that birds used to time
57 seasonal rhythms (Dawson et al., 2001; Tolla & Stevenson, 2020). Of all the avian models currently
58 in use for the study of photoperiod and reproductive development, few are more studied than Japanese
59 quail (*Coturnix japonica*) and migratory buntings (Huss et al 2008, Kumar et al 2021). Japanese quail
60 show incredibly robust changes in reproductive physiology and in their neuroendocrine response to
61 increased photoperiod (Stevenson et al 2022). Buntings show similar photoperiod response dynamics,
62 especially within their reproductive phenotypes. Unlike Japanese quail, the effects of photoperiod on
63 migratory behaviour have been documented in buntings (Tewary and Kumar 1982a, Kumar and
64 Tewary 1983, Kumar et al 2002, Rani et al 2005, Majumdar et al 2014, Singh et al 2020). This review
65 will primarily use literature derived from quail and bunting species to describe the mechanistic basis
66 of photoperiodic time measurement through coincidence timing and seasonal reproduction.

67 Seasonal rhythmicity in physiology can also occur in the absence of proximate photoperiodic
68 and supplementary environmental cues. Birds possess an internal seasonal clock, referred to as a
69 circannual clock, that generates a nearly annual rhythm (Gwinner 1977, Berthold 1984). Endogenous
70 circannual clocks in birds maintain their rhythmicity under constant environmental conditions
71 (Gwinner 2003). Both the initial activation and synchronisation/entrainment of these circannual
72 rhythms are dependent on aspects of the environment, and most notably the experienced photoperiod.
73 Hence, the timings of avian life history events (migration, reproductive investment etc.) are
74 synchronised by seasonal photoperiodic changes (Follett 2015). Gwinner and colleagues

75 experimentally demonstrated the significance of photoperiod on the timing of migration in white-
76 crowned and golden-crowned sparrows (Gwinner et al 1971). Upon the reduction of photoreception
77 by application of a black hood or injection of black ink under the skin of the head, birds showed lower
78 levels of migratory restlessness (*zugunruhe*) than control groups. Similar work by Donald Farner
79 further highlights the entrainment of avian rhythms in neuroendocrinology and reproductive
80 physiology (Farner and Wingfield 1980, Hiatt et al 1987) in white-crowned sparrows and other
81 migratory passerines.

82 Internal timekeeping mechanisms are key for the successful timing of seasonal life history
83 transition events, but also for the monitoring and regulation of several biological processes. These
84 internal clocks may be based upon cycles of hormone secretion but may also be dependent on the
85 control of regulatory calendar clock gene expression (Lincoln et al 2003). The circadian clock is often
86 used to explain how daily changes in light duration are integrated at a cellular level (Green and
87 Besharse 1996, Millar 2004, Goto 2013). Thus, the interplay between circadian clocks and seasonal
88 interval timers entrained by photoperiod is of particular interest, given the ability of circadian clock
89 genes to code light information (e.g., duration) and induce physiological responses. The circadian
90 clock is therefore a critical component for timing the photoperiodic control of seasonal reproduction.
91 This review will first cover endogenous circannual timing and external coincidence timing of light
92 cues for the photoperiodic timing of seasonal reproduction in birds. Then, the review will describe
93 the molecular steps from light detection by brain photoreceptors to the seasonal variation in the
94 Gonadotropin-releasing hormone (GnRH) system. Lastly, we will discuss the role of circadian clock
95 genes and distribution of genome binding motifs, D- and E-boxes, as a link between coincidence
96 timing of light cues and circadian clock gene pathways.

97

98 **Circannual oscillations and photoperiodic timing in birds**

99 Endogenous circannual rhythms are particularly widespread among avian taxa, but are also
100 present within mammals, plants, and even plankton (Harris 1963, Zamm et al 2016). Entrainment of

101 these rhythms by photoperiod allows birds to respond with significant physiological change during
102 the transitions between life history events. While the entrainment of circannual rhythms has been well
103 documented within birds, and particularly within temperate birds, where photoperiodic variation is
104 relatively great, this entrainment is less pronounced in other geographical areas (Gwinner 1977).
105 Within equatorial latitudes, where temperature and photoperiod remain relatively constant, native
106 birds and migrants to these environments are unable to rely on the variability of these factors to entrain
107 their endogenous pacemakers. Here, avian breeders show asynchronous circannual rhythms, as
108 opposed to rhythms which synchronise with seasonal change (Lincoln 2019). The endogenous
109 circannual rhythms underpinning reproduction initiate at variable times throughout the year, and
110 sometimes asynchronously with respect to conspecifics. Given the lack of environmental variation
111 and consistently suitable conditions for breeding, tropical species such as the Sooty tern
112 (*Onychoprion fuscatus*) display variable timing in reproductive investment (Bollen et al 2018).
113 Therefore, such species are much more dependent on internal endogenous rhythms with annual
114 timekeeping mechanisms and are less influenced by environmental variation (Wikelski et al 2008,
115 Helm and Stevenson 2014). It stands to reason that these circannual endogenous rhythms provide an
116 adaptive benefit, and may be ancestral, conserved in species which have since geographically
117 diverged (Helm 2006). The characteristics of these circannual rhythms, and entrainment thereof by
118 photoperiod, can be analysed experimentally.

119 While it is known that circannual timekeeping rhythms rely less on environmental variation,
120 many species are known to show a synchronisation of these rhythms to environment. Helm and
121 colleagues (2009) examined that the interplay between endogenous circannual rhythms and
122 experienced photoperiod in European (*Saxicola rubicola*) and Siberian (*Saxicola maurus*) stonechats
123 under both constant and changing light conditions. It was found that both variants of stonechats
124 respond to photoperiod in accordance with what is expected based on their circannual timekeeping,
125 and hence their expected geographical location. The predominant migratory behaviour used to assess
126 circannual rhythmicity is the nocturnal zugunruhe or ‘restlessness’ displayed by many captive

127 songbirds. Patterns of zugunruhe appear to align in period of activity and in direction of restlessness
128 when compared to migratory conspecifics in the wild. Captive stonechats display variable timing of
129 zugunruhe with a lack of periodicity, demonstrating sensitivity to entrainment by photoperiod (Helm
130 2006, Helm and Gwinner 2006). However, the observed patterns of zugunruhe are similar when
131 comparing European migrants with African residents in captivity. Therefore, the differences in
132 zugunruhe observed between these groups outside of captivity are highly dependent on the
133 experienced photoperiod. Similarities between these endogenous free-running cycles in captivity
134 indicate similar underlying programs. Differences between migrants and non-migrants arise during
135 the expression of migration, rather than within endogenous cycles underpinning migration. This
136 research suggests that rhythmic zugunruhe is ancestral, conserved within present-day non-migratory
137 stonechats. Responses to environmental change are therefore characterised by the state of an
138 individual's circannual timekeeping resulting from entrainment. By considering the internal state of
139 avian circannual timekeeping mechanisms, and how these mechanisms may coincide with incident
140 light, we can begin to understand the mechanism by which birds accurately respond to stimulatory
141 photoperiod for seasonal adaptation.

142

143 **Photoperiodic Coincidence Models:**

144 The neural and cellular mechanisms by which vertebrates both detect and respond to a critical
145 length of photoperiod (critical photoperiod) to induce reproductive change has historically been a
146 topic of study. Critical photoperiod is generally defined as the 24-hour light-dark ratio at which half
147 the population of a species switches from non-stimulatory to stimulatory physiological states
148 (Hazlerigg and Wagner 2006). Thus, a photoperiod longer than the CD is considered a stimulatory
149 long photoperiod by the animal. The critical photoperiod length has been reported as ~11.5 hours in
150 Japanese quail (Simpson and Follett 1982), but the critical length of photoperiod is variable among
151 avian taxa (Dawson 1987, Rani et al 2005) and within different environmental conditions, including
152 latitude (Silverin et al 1992). Overall, this implies a mechanism by which animals measure

153 photoperiod length, termed photoperiodic time measurement. Two leading models have been
154 postulated to explain the role of photoperiodic time measurement: the internal coincidence model,
155 and the external coincidence model. The internal coincidence model describes a theoretical system in
156 which two or more internal timers (i.e. circadian clock genes) are entrained by photoperiod. Changing
157 daylengths may cause these timers to coincide in such a way that initiates induction of seasonal
158 responses. In birds, this model lacks experimental support. Whilst some parties welcome the internal
159 coincidence model as an explanation for aspects of photoperiodic timing (Underwood et al 1997,
160 Yadav and Chaturvedi 2014), others, like Farner and Gwinner, suggested that the model is too
161 complex or have provided experimental evidence against internal coincidence in photosensitive birds
162 (Farner et al 1977, Meier 1981). Instead, the external coincidence model posits that the oscillating
163 external light stimulus entrains one or more internal timers of sensitivity to annual photoperiod, and
164 its photoinducible phase (ϕ_i ; Figure 1). Coincidence between the photoinducible phase and a light
165 stimulus, are required for successful induction of reproductive development. Evidence for the external
166 coincidence model has been demonstrated in several birds (Table 1) and other taxa (Saunders 1979,
167 Prasad et al 1982, Simpson and Follett 1982, Tewary and Kumar 1982b, Tewary and Prasad 1983,
168 Ravikumar and Tewary 1990). The precise nature of the molecular substrates that provide interval
169 timing for the external coincidence are described in detail below.

170

171 **Photoperiodic Regulation of Seasonal Life History Transitions in birds:**

172 In most long day breeding birds, the annual change in day length results in three distinct life-
173 history photoperiodic states: photosensitive, photostimulated, and photorefractory (Nicholls et al.,
174 1983; Stevenson et al., 2012). European starlings held in short days (<11.5hrs) for longer than 2 weeks
175 results in birds that are physiologically responsive to stimulatory long days and are referred to as
176 photosensitive (Dawson and Goldsmith 1997; Stevenson et al., 2012). In this state birds are
177 physiologically responsive to light stimulation during the photoinducible phase (Follett et al 1992).
178 The importance of the specific timing of light exposure on the avian response to photoperiod has

179 often been a point of interest for researchers. Follett and Sharp showed experimentally in 1969 that
180 the gonadal response of Japanese quail to a stimulatory photoperiod is dependent on the state of their
181 endogenous circadian rhythm, hence why a single pulse of light during a photoinducible phase is
182 sufficient to initiate significant reproductive change (Follett and Sharp 1969). Follett and colleagues
183 (1974) showed that when transferring photosensitive white-crowned sparrows from a light schedule
184 consisting of 8 hours of light, and 16 hours of darkness, to complete darkness, 8-hour pulses of light
185 between 2 and 100 hours after lights off were sometimes able to stimulate plasma luteinising hormone
186 (LH) concentration, indicative of a reproductive response to photoperiod (Follett et al 1974). Further,
187 the periods of time in which these light pulses were stimulatory occurred regularly at approximately
188 daily intervals. Hence, Follett and colleagues provided experimental evidence to suggest that not only
189 was an underlying endogenous circadian rhythm involved in reproductive development, but also that
190 this rhythm could be activated through stimulation by photoperiod. Similar dynamics of an
191 endogenous circadian element in photoperiodic stimulation of reproduction were also uncovered in
192 the house sparrow (Farner et al 1977).

193 The significance of the dark period during a photoperiodic cycle was explored experimentally
194 in other species. Jenner and Engels first verified the stimulatory nature of long photoperiod (16h) in
195 male juncos and white-throated sparrows before placing birds on a schedule of 8.25h light followed
196 by darkness interrupted by 1.75h light in the middle of the dark (Jenner and Engels 1952). They found
197 that this short light pulse was sufficient to induce gonadal growth and mature sperm development in
198 males. This evidence provided empirical support for the presence of a photoinducible phase, first
199 coined by Bünning in 1936, in the avian photoperiodic response (Bünning 1936). Light exposure
200 during the photoinducible phase results in reproductive development during which birds become
201 photostimulated. In this state, birds are reproductively active and engage in high rates of nest building,
202 copulatory behaviors and egg laying (McDonald 1976, Sharp 1993). Tewary and Tripathi (1983)
203 identified that female migratory buntings (*Emberiza bruniceps*) have significant ovarian growth when
204 exposed to longer day lengths consistent with those observed in summer months (Tewary and Tripathi

205 1983). Long days also inhibit reproduction evidenced by spontaneous regression after prolonged
206 exposure (Bentley et al., 1998). Thus, long days have a dual role to stimulate and then inhibit
207 reproductive physiology. Reproductive involution in response to long days is referred to as absolute
208 photorefractoriness (MacDougall-Shackleton et al., 2009). Some birds, such as the Japanese quail,
209 maintain reproductive function on long days and don't exhibit spontaneous regression therein
210 (Robinson and Follett 1982). Rather, a reduced photoperiod is required to initiate a reversal of
211 reproductive investment. Therefore, photorefractoriness here is defined as relative (MacDougall-
212 Shackleton et al., 2009).

213 It is known that most bird species do not require functioning retina in order to perceive light
214 and respond appropriately to photostimulatory photoperiods, including Japanese quail (Homma et al
215 1972). Blinded quail exhibit the same patterns of gonadal growth upon stimulation of a long
216 photoperiod compared to untreated birds, suggesting that the presence of extraretinal photoreceptors
217 ~~are~~ is responsible for light detection for the seasonal response. Similar research has been performed
218 in house sparrows, with several papers demonstrating an unchanged photoperiodic response when
219 comparing blinded and non-blinded birds (Menaker et al 1970, Underwood and Menaker 1970). Such
220 research has been crucial in directing research towards the discovery of deep brain photoreceptors,
221 now known to be widespread across avian taxa (Pérez et al., 2019). The avian photoperiodic response
222 includes three main components, (i) hypothalamic light receptors, (ii) the reproductive hypothalamo-
223 pituitary gonadal axis, and (iii) the circadian clock. Here, we will describe the neuroendocrine signal
224 transduction cascade, involving photostimulation of deep brain photoreceptors (opsins) and
225 subsequent regulation on deiodinase type 2 (*DIO2*) and type 3 (*DIO3*) expression, as a photoinducible
226 molecular link between photostimulation and GnRH expression. In this way, we propose that two
227 pathways are involved in the external coincidence model. There is a light detection pathway that links
228 deep brain photoreceptors with *DIO2/3* expression in tanycytes and a second interval timer pathway
229 that resides in the GnRH-gonadotropes system. The coincidence timing of the two pathways is
230 responsible for triggering the avian photoperiodic response.

231

232 Opsins: molecular substrates for light detection in birds

233 Opsins are a family of light-sensitive, G-protein coupled receptor membrane proteins.
234 Structurally, almost all the opsins consist of a single polypeptide chain of 340-500 amino acids
235 forming seven alpha-helical transmembrane regions with the cytoplasmic or extracellular loops and
236 domains connecting the transmembrane regions (Palczewski *et al.*, 2000). Vitamin A chromophore
237 is an 11-cis retinaldehyde (A1) and binds to the seven-helix bundle which, when absorbing a photon
238 of light, converts to an all trans stage by photoisomerization. This conformational change of the
239 chromophore interacts with transducin (G-protein), triggering the photoinduction molecular cascade
240 and change in membrane potential (Palczewski *et al.*, 2000). All the photopigments have a
241 characteristic absorption spectrum with a specific maximal sensitivity (λ max). Three opsin
242 classes have received the most attention due to the anatomical localization, function, and action
243 spectrum in avian brain and include rhodopsin, neuropsin and Vertebrate Ancient opsin. Here, we
244 will focus on the evidence to suggest the role of each photoreceptor in the control of avian
245 photoperiodism.

246

247 Rhodopsin

248 Rhodopsins are classified into Type 1 molecules found in microbes like bacteria, archaea,
249 fungi and algae and Type 2, which are the visual pigments in vertebrates and invertebrates
250 (Palczewski *et al.*, 2000). The photoreceptors in the quail brain were found to fit to an action spectrum
251 of rhodopsin with the maximum absorption at 492nm (Foster & Follett, 1985). Rhodopsin monoclonal
252 antibody RET-P1 identified two cell populations of rhodopsin positive neurons: in the lateral septum
253 (SL), in the infundibular nucleus (IN), and in median eminence (ME) in ring doves (Wada *et al.*,
254 1998). The expression of rhodopsin mRNA has also been found to increase with increased duration
255 of light and vary in different physiological states of migratory buntings with peak expression in the
256 photostimulated state (Majumdar *et al.*, 2015b). Along with the photoperiod-dependent expression,

257 it has also been shown that rhodopsin expression has a circadian pattern, and light pulse in the night
258 can elevate rhodopsin expression (Korenbrod *et al.*, 1989). Thus, keeping in mind that light at the
259 photoinducible phase, which usually falls in the early evening of many bird species, induces gonadal
260 development, rhodopsin may be one of the brain photoreceptors implicated in light detection for
261 neuroendocrine regulation of gonadal development.

262

263 *Neuropsin (OPN5)*

264 Bioinformatic and molecular search for new opsins resulted in the isolation of neuropsin from
265 mice and human tissues (Tarttelin *et al.*, 2003). The gene shares only 20%-30% similarity with other
266 opsin families and so has been designated as a new family. It shows all the functionally important
267 characteristics of other opsins but along with a lysine retinaldehyde chromophore binding site in the
268 seventh transmembrane domain (Tarttelin *et al.*, 2003). Later it was shown by analyzing purified
269 OPN5 protein that it does not act as retinal photoisomerase but functions as a UV sensitive GPCR,
270 which couples with Gi type G protein (Yamashita *et al.*, 2010). OPN5-positive cells have been found
271 in the chicken retina, pineal gland, and paraventricular organ (PVO; (Yamashita *et al.*, 2010)).
272 Detailed localization of OPN5 in quail brain revealed its presence in CSF contacting neurons in the
273 PVO region and in the external zone of the ME, which connects to the pars tuberalis (PT) (Nakane *et*
274 *al.*, 2010). Experimentally, incident light with a lambda max of 415nm can stimulate OPN5 to induce
275 a change in voltage in *Xenopus* oocytes. It is therefore inferred that OPN5, present in CSF-contacting
276 neurons, acts as a light sensor and subsequently generate membrane currents in response (Nakane *et*
277 *al* 2010). In chickens, another OPN5-like gene, OPN5L2 has been shown to be present laterally in
278 the third ventricle colocalizing with GnRH-I cells, reinforcing its probable role as a deep brain
279 photoreceptor (Ohuchi *et al.*, 2012). However, a direct test for the functional significance of *OPN5* in
280 seasonal neuroendocrine events revealed an inhibitory role of this opsin on thyrotropin-stimulating
281 hormone- β (*TSH\beta*) which has a direct role in the photoperiodic gonadal induction. Injections of OPN
282 RNAi into the ventricular system of border canaries caused an upregulation of *TSH\beta* mRNA levels.

283 Further, the mRNA levels of *OPN5* did not change in response to light in the photoinducible phase
284 (Stevenson & Ball, 2012). Similar observations were made in photoperiodic red headed buntings
285 where neuropsin levels did not vary in different lengths of photoperiod, and a significantly negative
286 correlation was found between *TSH β* and *OPN5* expression (Majumdar *et al.*, 2015a). Thus, the
287 anatomical localization and functional considerations suggest a role of *OPN5* in the photoperiodic
288 regulated neuroendocrine system in birds, but the course of action on the mechanism is unclear.

289

290 *Vertebrate ancient opsin*

291 Vertebrate Ancient opsin (VA opsin) was first isolated from cDNA pool of salmon eye tissue
292 with the derived amino acid sequencing showing about 37% - 42% similarity with other opsin families
293 (Soni & Foster, 1997). Structurally, VA opsin has many similar features as of other opsins but on the
294 third cytoplasmic domain, four non-conserved substitutions with two amino acid deletions make it
295 unique as a photoreceptive molecule, making it a small-sized opsin with light-dependent
296 phosphorylation properties (Soni & Foster, 1997). In birds, VA opsin has two isoforms: 972 bp VA
297 and 1080 bp VAL, which encodes for a 323 and 359 amino acid long protein, respectively. Both
298 proteins have been shown to be capable of eliciting a retinal dependent light response with current
299 productions in patch clamp recordings (Halford *et al.*, 2009). VA opsin mRNA have been amplified
300 from chicken hypothalamic cDNA and antibodies against it have shown positive signal in anterior
301 hypothalamus adjacent to the third ventricle, supraoptic nucleus, PVN and BnSTm and ME adjacent
302 to the PT (Halford *et al.*, 2009). Both isoforms of VA opsin have an approximate lambda max of
303 490nm, which corresponds to the previously accepted avian photoperiodic response peak at 492 nm
304 (Davies et al., 2012).

305 The precise mechanistic link between light detection by opsins in the hypothalamus and the
306 synthesis and release of GnRH from the median eminence is not well characterized. Recent work has
307 identified a series of molecular steps during the photoinducible phase and photostimulation that
308 provide a framework to identify potential neural circuits.

309

310 Molecular substrates that link photodetection and seasonal reproductive physiology*311 Photoperiodic regulation of hypothalamic expression of GnRH*

312 Gonadotropin-releasing hormone-I (GnRH-I; (King et al., 1966) and GnRH-II (Miyamoto et
313 al., 1984)) were first isolated from domestic chicken while GnRH-III was isolated from lamprey and
314 later found in songbirds (Bentley et al., 2004). Anatomical localization of GnRH-I includes preoptic
315 area (POA) with the fibres extending up to ME along the third ventricle and in the lateral septal area
316 (Millam et al., 1995; Stevenson and Ball, 2009). GnRH cells expressed in the pre-optic area (POA)
317 of the hypothalamus forms one of the most important links between the brain and the downstream
318 reproductive physiology in birds (Gore, 2002). The annual change in photoperiod (or simulated
319 photoperiodic changes in laboratory) causes distinct changes in the GnRH expression in many avian
320 species (Figure 2) (Stevenson et al., 2013). This dramatic variation in GnRH expression is believed
321 to be very pronounced in avian species in comparison to other vertebrates (Stevenson et al., 2012).
322 Photorefractory birds exposed to short photoperiod increase GnRH after 10 days suggesting that
323 GnRH may be the neural substrate for gaining photosensitivity (Dawson and Goldsmith, 1997;
324 Stevenson et al., 2012). Then, photosensitive birds exposed to long stimulatory photoperiod display
325 a further increase in GnRH synthesis that is maintained during the breeding periods. Then prolonged
326 exposure to long days and the onset of photorefractoriness is caused by a decline in GnRH content.
327 Other opportunistic and flexible reproductive strategies in birds maintain relatively constant GnRH
328 content which likely functions to provide rapid changes in reproductive physiology in response to
329 environmental conditions for optimal for breeding (MacDougall-Shackleton et al., 2009).

330

331 GnRH release is triggered by thyrotropin-stimulating hormone driven changes in Deiodinase type-2

332 Increased hypothalamic triiodothyronine (T3) during the photoinducible phase and the
333 subsequent rise in the gonadotropin, luteinizing hormone mRNA expression, suggested a causal link
334 between thyroid hormone, photoperiod, and gonadal regulation (Yoshimura et al., 2003). In quail, it

335 was found that the common glycogen polypeptide (CGA) mRNA, which together with TSH β forms
336 the functional TSH hormone. In birds, induction of *TSH β* mRNA was subsequently shown in Border
337 canaries (Stevenson and Ball, 2012) and migratory red headed buntings (Majumdar *et al.*, 2014).
338 Hypothalamic thyroid hormone signalling is critical for the long-day induction of GnRH release
339 (Figure 2) (Yoshimura *et al.*, 2003). Search for molecular components in the mediobasal
340 hypothalamus (MBH) yielded *DIO2* and *DIO3*. *DIO2* was found to be upregulated by a long
341 photoperiod in the MBH (Yoshimura *et al.*, 2003). Further analysis found a high concentration of
342 active 3,5,3' -triiodothyronine (T3) and prohormone thyroxine (T4) in MBH tissue in response to
343 long days from short days, although no change was detected in plasma concentration of these
344 hormones (Yoshimura *et al.*, 2003). A causal role for TSH was established by ICV injections of a
345 bovine TSH, which induced expression of *DIO2* in a dose dependent manner in the ependymal layer
346 of the third ventricle (3rdV; Nakao *et al.*, 2008). *DIO3* expression decreases in response to long days,
347 the reciprocal expression of *DIO3* occurs in the MBH, but the primary activator of increased *DIO3*
348 transcription remains unresolved; Evidence in mammals suggests an epigenetic mechanism
349 (Stevenson and Prendergast, 2013). Along with the thyroid hormone related changes, it has also been
350 shown that the GnRH nerves undergo huge seasonal changes in their morphology (Yamamura *et al.*,
351 2004). The GnRH nerve terminals lie in very close proximity to the basal lamina in median eminence
352 when exposed to long photoperiod and the encasement of their terminals by glial endfeet is also
353 reduced (Yamamura *et al.*, 2004, Nakane and Yoshimura 2010). Later it was shown that relatively
354 higher levels of T3 reduces the amount of glial (tanycytic) processes contacting the basal lamina and
355 that these numbers were negatively correlated with the number of neurons contacting the basal lamina
356 (Yamamura *et al.*, 2006). These studies indirectly support they hypothesis that changes in thyroid
357 action due to change in photoperiod can cause morphological and anatomical changes to brain cells
358 along with associated molecular dynamics. Altogether, these data indicate that TSH β activation
359 during the photoinducible phase is the primary molecular substrate for photoperiodic regulation of
360 the external coincidence model (Figure 2). It is likely that other molecules, such as GnRH or FSH β

361 contribute to the endogenous internal timing for seasonal reproduction, and that regulation of these
362 molecules is initiated by upstream TSH β activation. Given the importance of the circadian clock in
363 timing these molecular changes which link photoperiod and investment into reproductive physiology,
364 it is pertinent to consider whether the clock itself plays a role in the regulation of such molecules,
365 outside of initial timing of stimulation.

366

367 *Link between circadian clock timing and photoperiodic response*

368 The rhythmic expression of circadian clock genes is essential for the regulation and
369 maintenance important biological functions, including immune response, hormone synthesis and
370 release, metabolism, and sleep-wake cycles (Ikegami et al 2019). The first instance of daylight in a
371 24-hour cycle synchronises the circadian clock to begin its daily rhythm (Cassone 2014). It is through
372 this synchronisation that individuals determine the beginning of their “subjective day”. The avian
373 circadian clock possesses both positive and negative elements, which upregulate and downregulate
374 aspects of the circadian clock itself, respectively (Dunlap 1999). The clock genes *BMAL1* (Brain and
375 Muscle ARNT-Like 1) and *CLOCK* (Circadian Locomotor Output Cycles Kaput) are initially
376 transcribed and translated into BMAL1 and CLOCK protein, which heterodimerize. The
377 BMAL1/CLOCK dimer promotes transcription of clock genes *PER2* and *3* (Period protein) and *CRY1*
378 and *2* (Cryptochrome protein, Yoshimura et al 2000, Yasuo et al 2004). Dimerization of PERs and
379 CRYs results in a complex which interferes with BMAL1/CLOCK dimerization, acting in a negative
380 feedback loop to reset the circadian clock. Expression patterns of the clock genes *BMAL1*, *CLOCK*,
381 *PER2,3*, *CRY1*, and *2* are known to be stable within the MBH of Japanese quail under variable light
382 conditions (Yasuo et al 2003). This allows for consistent timekeeping in the presence of
383 photoinducible light stimuli and maintains a regular photoinducible phase. It is known that the
384 BMAL1/CLOCK dimer regulates transcription of *CRY* and *PER* paralogues by binding to E-box
385 sequences (CANNTG, where N = any appropriate nucleotide) within the promoter regions of target
386 genes (Nakao et al 2007). Similarly, binding to D-box DNA cis-elements within gene promoter

387 regions promotes transcription. Therefore, by searching for E- and D-box promoter regions upstream
388 of a gene's start codon, we can determine whether they are likely to be regulated by the circadian
389 clock (Helfer et al 2006, Onoue et al 2019) (Table 2-5).

390 When investigating the photoperiodic response, the transcription genes involved in the
391 neuroendocrine signal transduction cascade (following exposure to a stimulatory photoperiod) is of
392 particular interest. It is expected that if a gene is clock-controlled, their expression pattern over a 24-
393 hour period should mirror that of their corresponding regulatory clock gene. The indication that a
394 gene is clock-controlled provides important insight into the function of the gene and may indicate a
395 role in coordinating a connection between photoperiod and circadian clock. D-box and E-box
396 positions (Table 2-3 and Table 4-5, respectively) in the promoter regions of various photoperiodic
397 genes of interest have been provided. Each relevant avian D-box and E-box motif has been cross-
398 checked within the promoter regions of these genes of interest, and presence highlighted. These
399 findings represent new data, suggestive of a potential link between the circadian clock and the
400 neuroendocrine response to photoperiod. Each highlighted gene is involved to some degree in the
401 avian neuroendocrine response to photoperiod, whether that role be in photoreception (e.g., Eyes-
402 absent-3 (*EYA3*)) or reproduction (e.g., GnRH). Data is provided for the best characterized bird
403 genomes: chicken (*Gallus gallus*) and Japanese quail (*Coturnix japonica*). The presence of E-Box
404 and D-Box in the promoter region of *EYA3* in chicken and quail indicate this genomic motif is critical
405 for linking the circadian clock and genes implicated in the initial stages of photoinduction. These data
406 suggest that light activation during the photoinducible phase drives circadian clock gene binding to
407 D-Box and E-Box motifs in *EYA3* and the key photoperiodic component for the external coincidence
408 model. Despite this suggestive evidence, at this stage a solid molecular link between the avian
409 circadian clock and neuroendocrine response to photoperiod has not been established experimentally.

410 *EYA3* is a member of EYA gene family first identified in *Drosophilla* as a transcription factor
411 in eye development and have two distinct domains: C terminal domain which interacts with other
412 proteins and a tyrosine rich N terminal domain which contributes to its transactivational activity

413 uniquely, *EYA3* has a dual function of transcription factor and has an intrinsic phosphatase activity
414 (Rebay *et al.*, 2005). Although the role of *EYA3* has been extensively studied in tissue development,
415 recent advances also suggest a possible link with the circadian clock in mammals (Wood *et al.*, 2020)
416 *EYA3* mRNA expression sites include muscles, retina, and in PT in MBH of both birds and mammals
417 (Nakao *et al.* 2008; Dardente *et al.* 2010). Recently it was also shown that EYA3 peptide is expressed
418 at elevated levels in long day (Majumdar *et al.*, 2014). With two E-box elements in promotor (making
419 it sensitive to CLOCK and BMAL1) and two D-box elements (which makes it sensitive to TEF1
420 (transcriptional elongation factor 1) /SIX (sine oculis homeobox) elements), it provides support as a
421 photoperiodic gene for the integration of external photic cues for coincidence timing (Dardente *et al.*,
422 2010) (Table 2-5). In birds, melatonin does not play any role in photoperiodic control of reproduction
423 (Juss *et al.*, 1993), *EYA3* has been suggested to be a part of the transduction of photoperiodic
424 information pathway rather than directly involved in induction (Majumdar *et al.*, 2014). Expression
425 of EYA3 peptide in corresponding areas where rhodopsin is expressed in the hypothalamus
426 (Majumdar *et al.* 2014) along with the evidence that *EYA3* mRNA expression correlates with the
427 rhodopsin mRNA expression in response to increasing photoperiods (Majumdar *et al.*, 2015b)
428 suggests that *EYA3* is directly linked with light detection for the avian photoperiodic response.

429

430 **Summary and concluding remarks**

431 Seasonal changes in photoperiod and the daily timing of light stimulation are crucial for the
432 control of seasonal breeding in birds. The external and internal coincidence models provide
433 explanations for the mechanism by which birds respond to photoperiod in a stimulatory manner.
434 Despite some debate as to the more appropriate model, experimental evidence generally favours the
435 external coincidence model, or Bünning hypothesis, as an explanation for avian photoperiodism.
436 Circannual endogenous rhythms show similar programming between species. Expression of such
437 endogenous cycles are altered through entrainment by photoperiod, a reliable indicator of seasonal
438 time. Birds show a heightened reception to stimulatory photoperiods during a photoinducible phase

439 occurring circa 11.5 hours after dawn in Japanese quail. Coincidence of a light pulse during this
440 photoinducible phase results in neuroendocrine and physiological changes linked to reproductive
441 investment and is particularly apparent in Japanese quail. Where species are similar in
442 photostimulation, differences are observed in photorefractoriness. Japanese quail require a period of
443 shortened days to reset their reproductive statuses (relative photorefractoriness) whereas red-headed
444 buntings show spontaneous regression of gonads when maintained on long days (absolute
445 refractoriness). Deep brain photoreception is widespread across avian taxa. Rhodopsin, neuropsin,
446 and vertebrate ancient opsin have all been implicated in the avian photoperiodic response. Although
447 the link between light reception by these opsins and the following GnRH synthesis and release from
448 the median eminence is not fully understood, GnRH is an essential messenger that connects the
449 neuroendocrine changes in the brain to downstream changes in reproductive physiology. The data
450 suggest that GnRH and/or gonadotropin synthesis may act as the endogenous interval timer that
451 connects hypothalamic responsiveness to stimulatory photoperiodic cues driven by TSH β signalling
452 (Figure 1).

453

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459

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