

1 **Discontinuity in the molecular neuroendocrine response to increasing daylengths in Ile-**
2 **de-France ewes : is transient *Dio2* induction a key feature of circannual timing ?**

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12

13 **Abstract**

14 In mammals, melatonin is responsible for synchronisation of seasonal cycles to the solar year.

15 Melatonin is secreted by the pineal gland with a profile reflecting the duration of the night and acts

16 via the pituitary *pars tuberalis* (PT), which in turn modulates hypothalamic thyroid hormone status

17 through seasonal changes in the production of locally-acting thyrotropin (TSH). Recently we

18 demonstrated that in the Soay sheep, photoperiodic induction of *Tshb* expression and consequent

19 downstream hypothalamic changes occur over a narrow range of photoperiods between 12- and

20 14-h in duration. In the present study, we sought to extend our molecular characterization of this

21 pathway, based on transcriptomic analysis of photoperiodic changes in the pituitary and

22 hypothalamus of ovariectomized, estradiol-implanted Ile-de-France ewes. We demonstrate that

23 photoperiodic treatments applied before the winter solstice elicit two distinctive modes of

24 accelerated reproductive switch off compared to ewes held on a simulated natural photoperiod,

25 with shut-down occurring markedly faster on photoperiods of 13-h or more than on photoperiods
26 of 12-h and less. This pattern of response was reflected in gene expression profiles of
27 photoperiodically sensitive markers, both in the PT (*Tshb*, *Fam150b*, *Vmo1*, *Ezh2* and *Suv39H2*)
28 and in tanycytes (*Tmem252* and *Dct*). Unexpectedly, the expression of *Dio2* in tanycytes did not
29 show any noticeable increase in expression with lengthening photoperiods. Finally, expression of
30 *Kiss1*, the key activator of GnRH release, was proportionately decreased by lengthening
31 photoperiods, in a pattern that correlated strongly with gonadotropin suppression. These data show
32 that stepwise increases in photoperiod lead to graded molecular responses at the level of the PT,
33 a progressive suppression of *Kiss1* in the hypothalamic arcuate nucleus and LH/FSH release by
34 the pituitary, in spite of seemingly unchanged *Dio2* expression in tanycytes. We hypothesize that
35 this apparent discontinuity in the seasonal neuroendocrine response illustrates the transient nature
36 of the thyroid hormone-mediated response to long days in the control of circannual timing.

37

38 **Abbreviations**

39 ISH: *in situ* hybridization; LP: long photoperiod; MBH: medio-basal hypothalamus; OVX:
40 ovariectomized; PD: *pars distalis* of the pituitary; PT: *pars tuberalis* of the pituitary; SP: short
41 photoperiod; TH: thyroid hormone; T3: triiodothyronine; TSH: thyrotropin; ZT: zeitgeber time.

42

43 **Data availability statement**

44 The data that support the findings of this study are available from the corresponding author upon
45 reasonable request.

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47

48

49 **Introduction**

50 Seasonal breeding is a common adaptive feature of most mammals living at temperate latitudes,
51 ensuring that birth takes place at the most favourable time of year in terms of environmental
52 resources. Photoperiod is the main synchronizer of seasonal functions with additional factors such
53 as food availability, stress and social interactions acting as modulators¹. In mammals, melatonin
54 is the endocrine messenger of photoperiod and coordinates seasonal switches in endocrinology
55 and metabolism. To achieve this, melatonin targets a peculiar population of thyrotrophs within the
56 *pars tuberalis* (PT) of the pituitary, leading to seasonal expression of *Tshb* and hundreds of genes
57 expressed in the PT and medio-basal hypothalamus (MBH)²⁻⁴. PT-specific TSH appears to be the
58 crux of photoperiod-dependent seasonal timing as it connects melatonin input with local
59 hypothalamic T3 output through induction of deiodinase 2 (*Dio2*) in tanycytes lining the infra-
60 lateral walls of the adjacent third ventricle⁵⁻⁸. While the basics of this TSH-DIO2-T3 axis have
61 been well characterized in birds and mammals, cellular and molecular mechanisms that link T3 to
62 seasonal GnRH output, and hence to control of the pituitary response and gonadal axis, remain
63 unclear. However, a role for the KISS1 neuronal population of the arcuate nucleus seems
64 inescapable⁹⁻¹¹.

65
66 Our understanding of how photoperiod controls this molecular neuroendocrine axis remains
67 limited. We lack a comprehensive view of the impact of increasing daylengths on the expression
68 of seasonal markers within the PT, tanycytes and hypothalamus. Indeed, the majority of studies in
69 quail, hamsters and sheep have focused on comparisons between a long and a short photoperiod
70 (typically 16h vs 8h). However, switches in physiology are triggered by intermediate photoperiods,
71 as exemplified by experiments that defined the length of the critical photoperiod (CP) in quail and
72 hamsters¹². The CP is the minimal duration of daylight exposure (as determined by stepwise
73 increases or decreases) that prompts the opposite reproductive state; i.e. activation in sexually

74 inactive animals and sexual arrest in reproductively active animals. In quail and hamsters the CP
75 lies somewhere between 12.5h and 13h¹³⁻¹⁶. We recently determined that the CP is comprised
76 between 11.75h and 12.5h in male Soay sheep¹⁷. While this study demonstrated a strong
77 correlation between the molecular response at the PT-tanycytes level and the endocrine output
78 (FSH and Testosterone), it also revealed that the endocrine response to TSH was logarithmic rather
79 than linear, such that a very small increase in PT-derived TSH could elicit a large increase in *Dio2*
80 expression. Interestingly, a recent study on maternal programming in hamsters also revealed non-
81 linearity in the photoperiodic response of TSH-*Dio2*¹⁸. We proposed that such non-linearity along
82 the molecular neuroendocrine axis of the MBH underpins the long-term effects of photoperiod on
83 intrinsic seasonal programs, i.e. circannual timers^{4,17,18}.

84
85 Here, we used an approach similar to that developed for Soay lambs (CP protocol) in
86 ovariectomized, estradiol-implanted (OVX+E2) ewes of the Ile-de-France breed. The OVX+E2
87 model normalizes the level of circulating E2, which uncovers the well-documented central
88 seasonal shift in the negative feedback action of E2 on gonadotropin secretion¹⁹. In this model,
89 serum levels of the gonadotropins LH and FSH provide a reliable index of the state of the GnRH
90 pulse generator^{3,19}. Our findings are consistent with the CP value defined in Soay rams as well as
91 with the existence of non-linearity in the gonadotropic response to long days, such that a small
92 increase in *Tshb* correlates with a large decrease in LH and FSH levels. Our findings also revealed
93 a strong linear correlation between *Kiss1* expression in the arcuate nucleus and plasma LH/FSH
94 levels, consistent with a major role for these neurons in the control of seasonal breeding. Finally,
95 we observed uncoupling between expression levels of *Tshb* and *Dio2*: levels of *Dio2* were similar
96 across the five groups at the end of the two months of photoperiodic treatments. We speculate that
97 this discontinuity in the molecular TSH/DIO2/KISS1 pathway reflects transient *Dio2* induction,

98 which would be consistent with the brief temporal requirements for long days and T3 in the
99 synchronization of the ovine circannual program.

100

101 **Material & Methods**

102 **Ethics statement**

103 All experimental procedures were performed in accordance with international (directive
104 2010/63/UE) and national legislation (décret n° 2013-118) governing the ethical use of animals in
105 research (authorization n° E37-175-2 and n°A38 801). All procedures used in this work were
106 evaluated by a local ethics committee (Comité d’Ethique en Expérimentation Animale Val de
107 Loire; n°2012-10-5).

108

109 **Experimental animals & procedures**

110 Experiments were conducted in 30 adult Ile-de-France ewes (3–5 years old; weight 60–80 kg) kept
111 under normal husbandry conditions at the research station of the Institut National de la Recherche
112 Agronomique (Nouzilly, Unité Expérimentale PAO n°1297 (EU0028)). The natural photoperiod
113 at the latitude of Nouzilly, France (47°N) ranges from ~16:8 (16h of light, 8h of night) at the
114 summer solstice to ~8:16 at the winter solstice. All ewes were ovariectomized (OVX) and
115 estradiol-implanted (E2; 1cm silastic implant) between Sept 2nd and Sept 6th (2012).

116

117 All surgeries were performed after sodium thiopental anesthesia (Nesdonal®, 1g/80kg), under
118 constant isoflurane administration (Vetflurane®) and all efforts were made to minimize suffering.
119 Following surgery, animals received an injection of antibiotics (oxytetracycline, Terramycine
120 LA®, 1ml/10kg) and an injection of a non-steroidal anti-inflammatory drug (Finadyne®, flumixin
121 megumine, 2ml/50kgs). Animals were followed daily throughout the experiment.

122

123 Following surgery, all ewes were initially kept outdoors (open barns) before being brought indoors
124 in a light-tight building on Nov 19th (**Figure 1A**). The natural photoperiod at this time of year in
125 Nouzilly is ~9h15min. Six ewes were randomly assigned to one of each photoperiodic treatment :
126 Simulated Natural Photoperiod (PP nat), 11:13 (i.e. 11h of light per day), 12:12, 13:11 and 14:10
127 (**Figure 1A**). For each treatment, ewes were housed by groups of 3 in two separate light-tight pens.
128 Ewes of the PP nat group were submitted to a photoperiodic treatment that mimics outdoor
129 conditions through weekly stepwise modifications of daily light exposure. All other ewes were
130 exposed to longer photoperiods by delaying the time of lights off (**Figure 1B**). Ewes were
131 submitted to these photoperiodic treatments for ~2 months (**Figure 1A**).

132
133 Blood samples were collected twice weekly by jugular venipuncture in heparinized tubes from
134 Sept 17th (2012) through to Jan 10th (2013). After centrifugation, serum was collected and frozen
135 until assayed for hormones. At the end of the photoperiodic treatments, all animals were
136 euthanized by decapitation under deep barbiturate anesthesia (Nesdonal®, 5mL). To minimize
137 issues linked to potential time-of-day effects on gene expression all ewes were killed in the early
138 morning between ZT2 and ZT4 (with ZT0 being the time of lights on; **Figure 1B**) on January 13th
139 (n=15) and January 14th (n=14). Hypothalamic blocks were sampled and immediately frozen in
140 isopentane kept on dry ice, before being stored at -80°C until sectioning, as described previously³.
141 One ewe of the 12:12 group died in December and, at the end of the experiment, no E2 implant
142 could be recovered from one ewe of the PP nat group and another ewe from the 13:11 group. Data
143 from these 3 ewes were removed from the analysis of hormonal profiles.

144
145 **Hormonal profiles**
146 Plasma levels of LH, FSH and PRL were assayed by RIA. All samples from one experiment were
147 included in a single assay and every sample was measured in duplicate. LH: the assay standard

148 was 1051-CY-LH (equivalent to 0.31 NIH-LH-S1). Intra- and inter- assay coefficients of variation
149 averaged 9% and 15%, respectively with an assay sensitivity of 0.1 ng/mL. FSH: levels were
150 measured using reagents supplied by Tucker Endocrine Research Institute (Atlanta, GA, USA).
151 Intra- and inter- assay coefficients of variation averaged 8% and 9%, respectively with an assay
152 sensitivity of 0.1 ng/mL relative to the standard (Tuener oFSHstd. 1 equiv to 1.0 NIH-FSH-S1).
153 The cross reactivity with ovine LH was 0.03%. PRL: Intra- and inter- assay coefficients of
154 variation averaged 7% and 12%, respectively with an assay sensitivity of 2.5 ng/ml.

155

156 ***In situ* hybridization (ISH)**

157 ISH was performed as described previously using validated ovine riboprobes^{3,20,21}. A complete list
158 of the riboprobes used in this study is provided in **Supplementary Table 1**.

159 Hypothalamic blocks for *in situ* hybridisation were cut into 20µm sections using a cryostat
160 (CryoStar NX70, ThermoScientific) and thaw-mounted onto SuperFrost Plus slides
161 (ThermoScientific). All radioactive cRNA riboprobes were prepared by plasmid linearisation and
162 *in vitro* transcription (Riboprobe System, Promega) including ³⁵S-UTP (Perkin-Elmer). The probe
163 was purified with Illustra Probe Quant G50 micro-columns (Fisher) and counted with a liquid
164 scintillation counter (Tri-Carb 2900TR, Packard). Slides were post-fixed at 4°C for 20 min in 4%
165 PFA, 0.1 M PB, rinsed with 0.1 M PB (2 X 5min), acetylated with 3.75% v/v of acetic anhydride
166 in 0.1 TEA, 0.05 N NaOH (10min) and finally rinsed with 0.1 M PB (2 X 5min). Slides were then
167 dehydrated through graded ethanol solutions (50%, 70%, 95% and 100%; 3min each) and dried
168 under vacuum for 60 min. Sections were hybridized overnight at 58°C with 10⁶ cpm of probe per
169 slide in hybridization buffer (50% deionized formamide, 10% dextran sulfate, 1 X Denhardt's
170 solution, 300 mM NaCl, 10 mM Tris, 10 mM DTT, 1 mM EDTA, 500 µg/ml tRNA). Sections
171 were then rinsed in 4 X SSC (3 X 5 min) and subjected to RNase-A digestion (20 µg/ml) in a
172 buffer containing 500 mM NaCl, 1 mM Tris, 1 mM EDTA for 30 min at 37°C. Stringency washes

173 in SSC (with 1mM DTT) were performed to remove non-specific probe hybridisation: 2 X SSC
174 (2 X 5 min), 1 X SSC (10 min), 0.5 X SSC (10 min), 0.1 X SSC (30 min at 60°C), 0.1 X SSC (5
175 min). Slides were then dehydrated through graded ethanol solutions (50%, 70%, 95% and 100%;
176 3min each), dried under vacuum for 60 min and exposed for 1 to 3 weeks (depending on the target
177 mRNA) to an autoradiographic film (BioMax MR, Kodak). Films were scanned on a transmittance
178 image scanner (Amersham, UK) along with a calibrated optical density (OD) transmission step
179 wedge (Stouffer, USA). Calibrated Integrated OD measurements of gene expression were
180 performed using ImageJ software.

181

182 **Data analysis**

183 Data were analysed using GraphPad Prism 6.0 and are reported as mean \pm sem. For hormonal
184 profiles, RIA data were analyzed by Repeated Measures (RM) 2-way ANOVA. One-way ANOVA
185 was also used to perform analyses limited to the two weeks before the photoperiodic transfer and
186 the last two weeks of the experiment. ISH data were analyzed by 1-way ANOVA using treatment
187 as a variable. The post-hoc Tukey test was used for multiple comparisons. $p < 0.05$ was considered
188 significant. Using individual values, linear regression analysis was performed to evaluate
189 correlations between *Tshb*, *Dio2* or *Kiss1* mRNA levels and mean LH/FSH levels over the last two
190 weeks of the experiment.

191

192 **Results**

193 **Hormonal profiles**

194 RIA was used to assess plasma levels for LH (**Figure 2A-C**), FSH (**Figure 2D-F**) and PRL (**Figure**
195 **2G-I**). For LH, RM two-way ANOVA revealed a strong time*group interaction ($P < 0.0001$), which
196 reflected the impact of the photoperiodic treatments, as no difference between groups were
197 observed prior to these. Indeed, further analysis revealed no differences between groups over the

198 2-wks period before the photoperiodic transfer (**Figure 2B** ; one-way ANOVA ; P=0.25), while
199 levels differed significantly over the last 2-wks period (**Figure 2C** ; one-way ANOVA ;
200 P<0.0001). For FSH, RM 2-way ANOVA revealed a strong time*group interaction (P<0.0001),
201 which mostly reflected the impact of the photoperiodic treatments. Indeed, further analysis
202 revealed differences between groups over the 2-wks period before the photoperiodic transfer
203 (**Figure 2E**) and the last 2-wks period of the experiment (**Figure 2F** ; one-way ANOVA ;
204 P<0.0001 for both comparisons). The difference before the photoperiodic transfer is accounted for
205 by slightly lower levels in the PP nat group compared to the 4 other groups. Linear regression
206 analysis showed that levels of LH and FSH during last 2-wks period of the experiment were
207 correlated ($R^2=0.4679$; P<0.0001).

208
209 For PRL, RM two-way ANOVA revealed a a strong time*group interaction (P=0.0004), which
210 may reflect the inter-individual and inter-group variabilities before the photoperiodic treatments
211 and slightly divergent trajectories after transfer. However, PRL levels did not differ between
212 groups before the photoperiodic transfer (**Figure 2H** ; one-way ANOVA ; P=0.53), while a trend
213 towards higher levels with increasing photoperiods was noticed for the last 2-wks period of the
214 experiment (**Figure 2I** ; one-way ANOVA ; P=0.053).

215

216 **Gene expression**

217 We used semi-quantitative ISH to assess the impact of the photoperiodic transfer on the mRNA
218 expression levels of multiple seasonal markers within the MBH (**Figure 3**). Data were analysed
219 by one-way ANOVA (**Figure 3A**). Photoperiod affected the expression of all PT markers except
220 *Chga* [$F_{4,24} = 0.32$; P=0.86] : *Tshb* [$F_{4,24} = 25.94$; P<0.0001], *Fam150b* [$F_{4,24} = 22.04$;
221 P<0.0001], *Vmol* [$F_{4,24} = 14.37$; P<0.0001], *Ezh2* [$F_{4,24} = 14.37$; P<0.0001], *Suv39H2* [$F_{4,24} =$
222 6.61 ; P=0.001] and *Eya3* [$F_{4,24} = 6.29$; P<0.005]. Post-hoc analysis revealed that expression

223 levels were not significantly different between the PP nat, 11:13 and 12:12 groups for any of these
224 PT-expressed genes. Compared to PP nat, expression of *Tshb*, *Fam150b* and *Eya3* were
225 significantly increased under 13:11. Expression of all markers was significantly increased by the
226 14:10 photoperiod compared to PP nat. Within markers of tanycytes, photoperiod significantly
227 affected mRNA expression of *Tmem252* [$F_{4,23} = 7.55$; $P=0.0005$] and *Dct* [$F_{4,24} = 6.11$; $P<0.005$]
228 but not *Dio2* [$F_{4,24} = 1.74$; $P=0.17$]. Similar to the PT markers, no significant differences were
229 found between PP nat, 11:13 and 12:12 groups, while longer photoperiods of 13:11 and 14:10
230 increased expression of *Tmem252* and *Dct*, compared to PP nat. No detectable expression of *Dio3*
231 was observed (not shown). Finally, the photoperiodic transfer affected expression of the
232 hypothalamic marker *Kiss1* [$F_{4,24} = 6.06$; $P<0.005$], with expression significantly reduced under
233 13:11 and 14:10 compared to PP nat. Representative images of autoradiograms for all these
234 markers at their peak of expression are provided in **Figure 3B**.

235

236 **Linear regression analyses**

237 Data for LH and FSH (using individual means for the last two weeks of treatment; data from Figure
238 2C and Figure 2F) are shown in **Figure 4A and 4B**, respectively. Levels of *Tshb* and *Kiss1* mRNA
239 were negatively and positively correlated with LH ($R^2=0.20$, $P=0.018$ and $R^2=0.66$, $P<0.0001$) and
240 FSH levels ($R^2=0.29$, $P=0.004$ and $R^2=0.53$, $P<0.0001$), respectively . Levels of *Dio2* mRNA were
241 not correlated with either LH ($R^2=0.04$, $P=0.34$) or FSH levels ($R^2=0.1$; $P=0.11$). For *Tshb* and
242 LH/FSH, data could be better fitted with a hyperbolic curve, which is shown instead of the linear
243 regression line.

244

245 **Discussion**

246 There is a strong intrinsic component to the seasonally timed changes in reproductive status in
247 sheep, which under constant long photoperiods manifests itself as a circannual rhythm of ~10

248 months^{22,23}. By employing OVX+E2 ewes of the Ile-de-France breed, this cycle of activation and
249 quiescence can be clearly seen in changes in gonadotropin secretion, and hence the 3 trajectories
250 of reproductive shutdown observed in the present study can be interpreted against the circannual
251 framework. These endocrine changes mirror changes in expression level of key molecular markers
252 in the MBH. However, our data also point to a discontinuity along the neuroendocrine
253 gonadotropic axis, since *Dio2* levels in tanycytes - unlike *Tshb* in the PT and *Kiss1* in the
254 hypothalamic parenchyma - were not correlated to the differential photoperiodic LH/FSH output
255 **(Figure 4)**.

256

257 Our molecular analysis supports the key role of the PT as a reliable decoder of the photoperiodic
258 message carried out by melatonin. Beyond *Tshb*, we and others recently identified hundreds of
259 PT-expressed genes displaying strong T3-independent seasonal changes in expression^{2,3}. The
260 expression profiles reported in this study are also consistent with the acute LP-responsiveness
261 of most of these markers (*Tshb*, *Fam150b*, *Vmo1*, *Ezh2* and *Eya3* ; see³). We note that
262 expression of *Chga*, a marker for SP in the PT^{2,3}, was not diminished upon exposure to increased
263 daylengths, consistent with its lack of responsiveness to acute LP exposure³. This finding is
264 compatible with a model in which most photoperiodic markers in the PT are induced/repressed
265 at different daylengths and/or located in pathways downstream of a core of "1st order LP-
266 responsive genes", in a parallel with the organization of the molecular circadian clock, with a
267 handful of core clock genes and a myriad of (tissue-specific) clock-controlled genes^{24,25}.

268

269 Our data are consistent with a role for PT-expressed *Eya3/Tshb* in the photoperiodic read-
270 out^{7,26}, and with the implication of *Kiss1*-expressing neurons of the arcuate nucleus in the
271 seasonal control of GnRH and gonadotropins^{10,11}. These data further indicate that a photoperiod
272 as short as 11:13 is sufficient to elicit a marked response in both LH and FSH, without

273 statistically significant changes in the expression of molecular seasonal markers. Current
274 evidence in mouse²⁷ and sheep^{2,26,28} points to a pivotal role for EYA3 in the photoperiodic
275 response of the melatonin-responsive PT. In sheep, we proposed that the response to LP is
276 triggered by the EYA3/SIX1 heterodimer, which acts as a co-activator of DNA-bound TEF at
277 the *Tshb* promoter^{1,7,26}. The LP increase in *Eya3* expression is crux to the system but the overall
278 *Tshb* transcriptional output rests on proper phase relationship between *Eya3* and *Tef*, which are
279 both circadian clock-controlled genes²⁶. *Eya3* expression is timed to occur ~12h after dark onset
280 (between 8h and 16h, tighter sampling schedule missing) and is potently inhibited by melatonin
281 through a mechanism that remains unknown (as is the mechanism for *Eya3* induction by LP).
282 This model predicts that longer photoperiods are more conducive to *Eya3* increase, due to
283 melatonin being absent in the morning, and also predicts that the rise in *Tshb* is necessarily
284 preceded by rising *Eya3*. Our data are consistent with this since *Eya3* has risen by 75% under
285 11:13 and by 250% under 12:12, when compared to PP nat (P<0.05 by t-test for both pairwise
286 comparisons). Comparatively, *Tshb* does not display any statistically significant changes in
287 expression. However, animals were killed in the early day (ZT2-4); a time which fits the
288 expected *Eya3* peak under longer photoperiods but also corresponds to a marked trough in *Tshb*
289 expression²⁶. This might account for the apparent lack of difference observed for *Tshb*
290 expression between PP nat, 11:13 and 12:12 photoperiods.

291

292 We recently used a similar methodology to investigate the impact of quantitative increases in
293 photoperiod on the neuroendocrine response in intact Soay ram lambs¹⁷. In that study, we also
294 found induction of *Eya3* at a shorter photoperiod (11.75h) compared to *Tshb* (12.5h), while the
295 pituitary and gonadal responses (FSH and Testosterone, respectively) were more clear-cut and
296 pointed to the existence of a CP between 11.75h and 12.5h. It is difficult to compare the two
297 studies with regards to the physiological output since we used intact ram lambs in the former

298 study and OVX+E2 ewes in the current study. However, in this OVX+E2 model, it has been
299 established that LH values above and below ~1ng/ml do correspond to active and anestrus states
300 of intact ewes of both Suffolk^{29,30} and Ile-de-France breeds^{3,31,32}. Considering this, ewes of the
301 13:11 and 14:10 groups, but not of the 11:13 and 12:12 groups, would have been in – or very
302 close to – a state of anestrus. Furthermore, LH levels are not significantly different between the
303 groups exposed to daylengths of 13h and 14h while *Tshb* levels are more than doubled under
304 14:10 compared to 13:11. This seems to indicate that *Tshb* levels attained with exposure to a
305 photoperiod equal to – and likely shorter than – 13h are sufficient to trigger maximal response
306 of the hypothalamic-pituitary axis. Therefore, we conclude that the critical photoperiod in Ile-
307 de-France ewes probably lies between 12h and 13h, similar to the situation in Soay ram lambs¹⁷.

308

309 In comparing the outcomes of these independent studies, it is also important to stress that the
310 duration of photoperiodic treatments differed between the two protocols: Soay rams were
311 exposed for 4 weeks, while ewes of the current study were exposed for ~8 weeks. This duration
312 has to be taken into account when interpreting the physiological status at the end of the
313 experiment. Considering the trajectory for LH (and FSH) in ewes of the 11:13 and 12:12 groups,
314 it seems likely that longer exposure to these photoperiods would eventually have led to LH
315 values <1ng/ml (signing an anestrus-like state ; see above). The impact of photoperiodic history
316 also has to be considered as the same photoperiod triggers opposite responses of the GnRH
317 pulse generator and LH output, according to the initial photoperiod to which ewes were
318 exposed³⁰.

319

320 In contrast to LH and FSH, plasma levels of PRL did not change significantly throughout
321 exposure to graded photoperiods, even though a trend towards higher levels was seen, most
322 obvious for the 14:10 group. We also note a much larger inter-individual variability, with

323 multiple peaks, before than after the transfer to the light-tight building. This most likely reflects
324 the well-characterized susceptibility of PRL secretion to various unpredictable stressors such
325 as noise or human activity and temperature fluctuations³³⁻³⁵, which are common in open barns
326 but very limited in our light-tight building. PRL displays a seasonal pattern of secretion, with
327 higher levels during spring and summer^{5,33,35-38} and rapid increase or decrease upon acute
328 exposure to longer or shorter daylengths, respectively^{26,28,39,40}. Photoperiodic history also
329 affects the long-day response of PRL secretion in ewes⁴¹. In sheep, gonadotropic (LH/FSH) and
330 lactotropic axes (PRL) display opposite responses to daylength, which are driven by two distinct
331 neuroendocrine axes^{12,42,43}: the gonadotropic axis uses the retrograde TSH/DIO2/T3 axis while
332 the lactotropic axis relies on anterograde signaling from the PT to the *pars distalis*,
333 independently of T3 (in both rams and ewes^{3,12,44,45}), through one or several endocrine factors
334 (known as tuberalin(s)) whose identity remains unclear⁴. Our findings that a 14h-daylength does
335 not significantly increase PRL is in line with early findings in Siberian hamster⁴⁶, which showed
336 that both axes have distinct CP, the CP for PRL being longer than that for LH/FSH.

337

338 Overall, the concept of CP is of theoretical value but must be used with caution as the CP value
339 differs according to the species and latitude of natural habitat, molecular or neuroendocrine
340 output considered, duration of exposure and moment in the seasonal (circannual) cycle at which
341 exposure occurs (i.e. photoperiod history ; see^{47,48}). CP might also differ slightly between sexes
342 since rams typically display an advance of their breeding season compared to ewes (~1 month),
343 which ensures that all ewes get pregnant - hence deliver lambs - within a very narrow time span
344 at the end of winter⁴⁹. However, our protocols with different breeds (Soay vs Ile-de-France),
345 intact ram lambs vs OVX+E2 adult ewes, different duration of photoperiodic treatments and a
346 temporal resolution of 1h in the current study are not adapted to rigorously test for small
347 differences in CP value between sexes.

348

349 Perhaps, the most striking difference with the Soay lamb study¹⁷ is the disconnection between
350 *Tshb* and *Dio2*: *Tshb* expression steadily increases with stepwise increases in photoperiod
351 while *Dio2* does not. This is very surprising considering our prior finding that *Dio2* (amongst
352 other genes, see below) is acutely induced – 4-fold increase as assessed by RT-qPCR – by
353 exposure to a LP of 15.5h³. However, here again, the length of the photoperiodic treatment, 3.5
354 weeks, was much shorter than the 8 weeks treatment of the current study. In addition, the ISH
355 methodology used here is less sensitive than RT-qPCR, which might be a problem considering
356 the modest (1.5-2-fold) seasonal amplitude of the *Dio2* rhythm in tanycytes of Ile-de-France
357 ewes³, compared to Soay rams⁵⁰. Nevertheless, considering the decrease in *Kiss1* expression
358 and the concurrent decreases in LH/FSH, we infer that photoperiodic treatments were efficient.
359 Since the DIO2-triggered hypothalamic local increase in T3 at the beginning of spring is crucial
360 to the progression of the ovine seasonal cycle towards reproductive arrest^{12,44,51-53}, we conclude
361 that *Dio2* induction occurred, but was not sustained throughout the 8 weeks. We therefore
362 hypothesize that the disconnection between *Tshb* and *Dio2* is due to a transient impact of TSH
363 upon *Dio2* and T3 production, which is nevertheless enough to entrain the circannual clock and
364 trigger the response of the hypothalamo-pituitary axis. We can exclude that this disconnection
365 reflects a general uncoupling between PT and tanycytes since two other tanycyte-specific
366 markers, *Tmem252*³ and *Dct*²¹, showed increased expression levels with increasing
367 photoperiods, in a pattern resembling that of PT-expressed genes. Interestingly, using
368 thyroidectomized OVX+E2 ewes we demonstrated that *Tmem252* induction by LP is genuinely
369 dependent upon T3³. The profile of *Tmem252* observed in this study would therefore indicate
370 that T3 levels remain high under LP in spite of constant *Dio2* expression. Overall, these
371 observations point to a *Dio2*-specific mechanism of negative feed-back, which rapidly follows
372 induction by LP.

373

374 Indeed, under natural conditions, Siberian hamster display sharp transient peaks of *Dio2* and
375 *Dio3* expression during the year⁵⁴. There is also strong evidence in sheep that a transient
376 exposure to LP – 45-60 long days – is sufficient to synchronize the circannual cycle^{12,55,56}. This
377 is in line with the idea that T3, even though it exerts a key organizing role, is required only
378 during a limited window of time in sheep and hamsters^{12,45,57,58}. Then, after being acutely
379 induced, *Dio2* expression would return to lower levels, in spite of increased PT-derived TSH.
380 The mechanisms for such an uncoupling are unknown but a few hypotheses can be made. The
381 induction of *Dio2* by TSH depends on the cAMP pathway ; *Dio2* is a CRE-dependent gene^{53,58}.
382 Considering that tanycytes are a hub for a host of signals^{4,59,60}, it seems plausible that other
383 GPCR-dependent signaling pathways, coupled to either G α s or G α i, also impinge on *Dio2*
384 expression. Furthermore, *Dio2* expression is enhanced in both hypothyroid rats⁶¹ and ewes³,
385 which indicates that T3 normally exerts a brake on *Dio2* expression in tanycytes. Such an
386 autocrine short feedback loop is predicted to keep *Dio2* levels in check during LP exposure,
387 and might be responsible for the apparent disconnection. Also, T3 strongly impacts local
388 metabolism⁶² and *Dio2* is itself sensitive to metabolic cues⁶³⁻⁶⁶, which provides yet another
389 potential short feedback loop. Other unrelated cues, such as LPS injection^{67,68} and hypoxia⁶⁹
390 also impact either *Dio2* expression or DIO2 stability. In conclusion, multiple signals – not
391 limited to PT-derived TSH – converge onto tanycytes and overall *Dio2* expression reflects
392 integration of all these inputs. We conclude that our experimental design, which examined
393 expression of genes after ~2 months of photoperiodic exposure, reveals the transient nature of
394 the LP impact upon the circannual clock.

395

396 Our prior work in Soay lambs¹⁷, Ile-de-France ewes³ and in Siberian hamsters¹⁸ has provided
397 hints that a very modest increase in PT-derived TSH at the end of winter might be enough to

398 yield a large increase in *Dio2* expression. We ascribed this logarithmic response to sensitization
399 of the TSHR pathway within tanycytes, a proposal which remains to be addressed rigorously.
400 Here, we report that very modest changes in *Tshb* expression trigger comparatively large
401 responses in LH/FSH (i.e. logarithmic response rather than linear), which are associated with a
402 rather linear and progressive decrease of *Kiss1* expression (**Figure 4**). Our data in OVX+E2
403 ewes are consistent with those in intact Soay ram lambs and point to a CP value comprised
404 between 12-13h. More importantly, our data provide strong evidence that non-linear responses
405 and discontinuity occur along the TSH/DIO2/T3/KISS1 axis. These features may be part of of
406 the circannual timing device and will have to be considered in the interpretation of future
407 studies.

408

409 **Acknowledgements**

410 We thank staff at the CIRE platform for assistance with surgical procedures, Olivier Lasserre,
411 Didier Dubreuil and Damien Capo from the Unité Expérimentale PAO no°1297 (EU0028) for
412 taking care of the animals and for blood sampling. We also thank members of the team
413 Molecular Neuroendocrinology of Reproduction for their input at various stages of this work.

414

415

416 **Figure legends**

417 **Figure 1:** Schematic of the experimental design. (A) Overview of the photoperiodic treatments.
418 OVX+E2 ewes were initially kept in open barns, exposed to the natural decrease in daylength,
419 before being brought indoors and exposed for ~2 months to either a simulated natural photoperiod
420 (PP nat) or to photoperiod of increasing duration: 11:13, 12:12 , 13:11 and 14:10. (B) Ewes were
421 exposed to constant photoperiods by delaying the time of lights off. All ewes were killed in the
422 early day (ZT2-4).

423
424 **Figure 2:** Hormonal profiling of the response to the five photoperiodic treatments. (A) Mean LH
425 levels in blood plasma of ewes sampled bi-weekly throughout the experiment. (B) Mean LH for
426 the two weeks (i.e. four time points) before the photoperiodic transfer. (C) Mean LH for the last
427 two weeks of the experiment. Different letters indicate statistically different groups ($P < 0.05$). (D-
428 F) FSH profiles – legends identical to those for LH. (G-I) PRL profiles – legends identical to those
429 for LH. (n=6 for groups 11:13 and 14:10 , n=5 for groups PPnat, 12:12 and 13:11; see M&M).

430
431 **Figure 3:** Impact of photoperiodic treatments on select photoperiod-responsive genes in the MBH
432 as assessed by ISH. (A) Bar charts showing normalized expression for PT markers (*Tshb* through
433 to *Chga*), tancytic markers (*Dio2*, *Tmem252* and *Dct*) and the hypothalamic marker *Kiss1*.
434 Different letters indicate statistically different groups ($P < 0.05$). (B) Images representative of high
435 ISH signal for each gene as revealed by autoradiography. (n=6 for all groups, except PPnat n=5;
436 see M&M).

437
438 **Figure 4:** Discontinuity in the molecular neuroendocrine pathway of seasonality revealed by linear
439 regression analysis. Individual means for LH and FSH correspond to the last two weeks of
440 treatment (i.e. same data as in Figure2) (A) Linear regression analysis reveals statistically
441 significant correlation between *Tshb* / *Kiss1* and LH, but not *Dio2* and LH. (B) Linear regression
442 analysis reveals statistically significant correlation between *Tshb* / *Kiss1* and FSH but not *Dio2*
443 and FSH. P values for linear regressions are provided on top of each panel. Note that the
444 relationships between *Tshb* and LH/FSH could be better fitted by a non-linear hyperbolic function,
445 which is shown, instead of the linear regression line. The dashed grey line arbitrarily set a $x=1$ is
446 meant to help visualization of the extent of the distribution. (n=6 for groups 11:13 and 14:10 , n=5
447 for groups PPnat, 12:12 and 13:11; see M&M)

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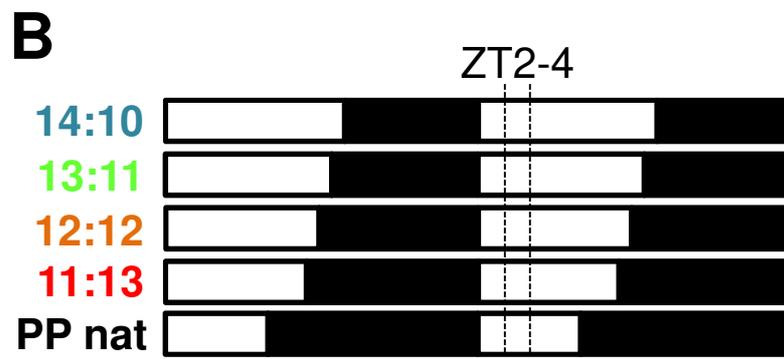
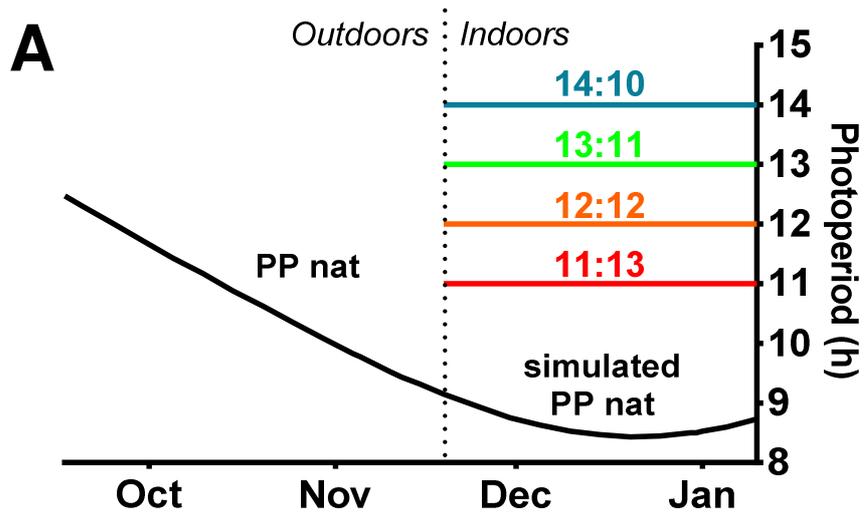


Figure 1

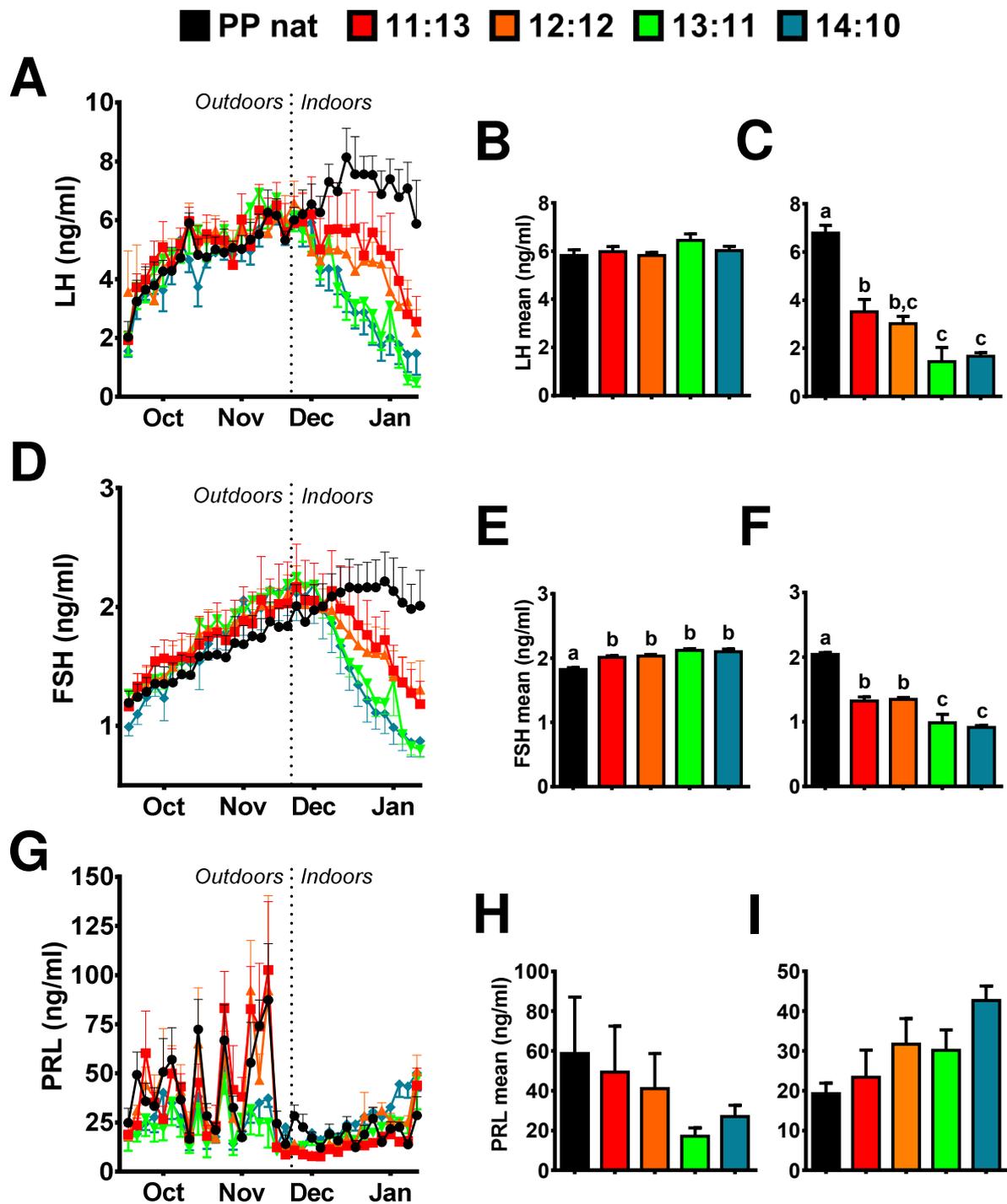


Figure 2

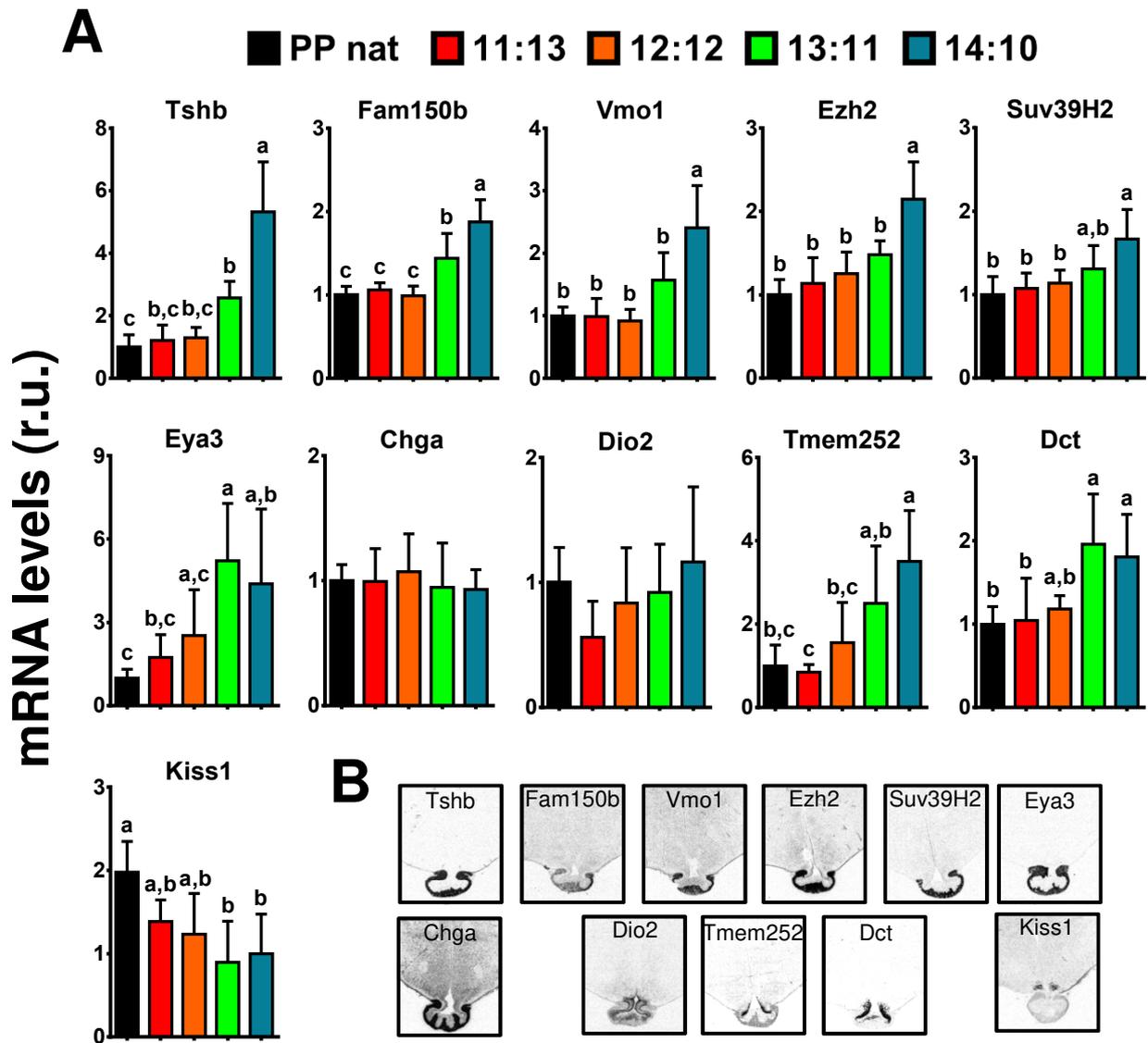


Figure 3

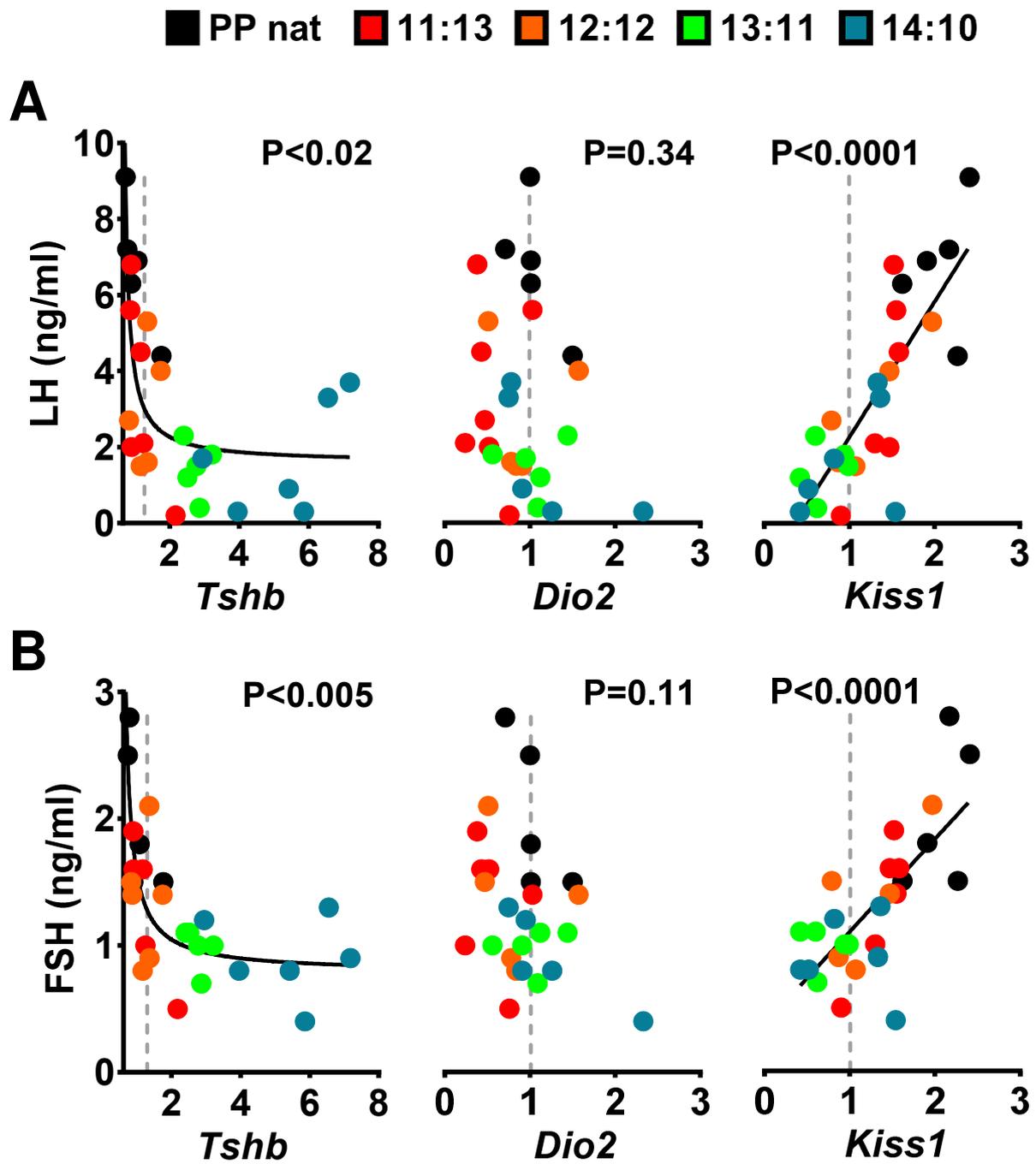


Figure 4