1	Age-dependent and age-independent sexual selection on
2	multiple male traits in the lekking black grouse (Lyrurus
3	tetrix)
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6	Matti Kervinen <sup>* a</sup> , Christophe Lebigre <sup>b</sup> , Carl D. Soulsbury <sup>c</sup>
7	<sup>a</sup> Department of Biological and Environmental Science, P. O. Box 35, FI-40014
8	University of Jyväskylä, Finland
9	<sup>b</sup> Earth and Life Institute, Place de la Croix du Sud 4, Carnoy building, B-1348
10	Louvain-la-Neuve, Belgium
11	<sup>c</sup> School of Life Sciences, Joseph Banks Laboratories, University of Lincoln, Lincoln
12	LN6 7TS, UK
13	* Corresponding author: matti.m.kervinen@gmail.com
14	
15	Running headline: Age-(in)dependent sexual selection in L. tetrix
16	
17	Summary
18	1. Individuals' reproductive success is often strongly associated with their
19	age, with typical patterns of early life reproductive improvement and late
20	life senescence. These age-related patterns are due to the inherent trade-
21	offs between life history traits competing for a limited amount of resources
22	available to the organisms. In males, such trade-offs are exacerbated by the
23	resource requirements associated with the expression of costly sexual

traits, leading to dynamic changes in trait expression throughout theirlifespan.

- 26 2. Due to the age-dependency of male phenotypes, the relationship between 27 the expression of male traits and mating success can also vary with male 28 age. Hence, using longitudinal data in a lekking species with strong sexual 29 selection – the black grouse Lyrurus tetrix – we quantified the effects of 30 age, lifespan and age of first lek attendance (AFL) on male annual mating 31 success (AMS) to separate the effects of within-individual improvement 32 and senescence on AMS from selective (dis)appearance of certain 33 phenotypes. Then, we used male AMS to quantify univariate and 34 multivariate sexual selection gradients on male morphological and 35 behavioural traits with and without accounting for age and age-related effects of other traits. 36
- 37 3. Male AMS increased with age and there was no significant reproductive 38 senescence. Most males never copulated and of the ones that did, the 39 majority had only one successful year. Lifespan was unrelated to AMS, but 40 early AFL tended to lead to higher AMS at ages 1 to 3. AMS was related 41 to morphological and behavioural traits when male age was ignored. 42 Accounting for age and age-specific trait effects (i.e. the interaction 43 between a trait and age) reduced the magnitude of the selection gradients 44 and revealed that behavioural traits are under consistent sexual selection. 45 while sexual selection on morphological traits is stronger in old males.
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  4. Therefore, sexual selection in black grouse operates primarily on male
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  47 behaviour and morphological traits may act as additional cues to
  48 supplement female choice. These results demonstrate the multifaceted

49 influence of age on both fitness and sexual traits and highlight the
50 importance of accounting for such effects when quantifying sexual
51 selection.

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- 53 Key-words: ARS, mating success, lek, long-term data, LRS, selection differential,
  54 selection gradient, senescence, *Tetrao tetrix*
- 55

## 56 Introduction

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58 Sexual selection underpins the evolution of extravagant male ornaments either 59 through female preference for higher expression of these traits (e.g. colourful feathers) 60 or through the competitive advantage they offer during male-male competition (e.g. 61 male weaponry, Andersson 1994). Therefore, direct relationships between male traits 62 and mating success are generally interpreted as evidence for sexual selection on male 63 traits. However, mating success in a particular year of a male's life may not accurately 64 reflect the association between the trait and fitness, since individual mating success 65 typically increases in early life and decreases at later ages (Jones et al. 2008; Nussey 66 et al. 2013).

Age-dependent mating and reproductive success are typically explained by the inability to simultaneously maximise competing life history traits (e.g. growth, reproductive investment, survival), due to the limited amount of resources available to the organisms (Stearns 1992). Such trade-offs are responsible for the constraints on reproductive effort in early ages, when resources are allocated to growth, leading to shorter (reproductive) lifespan (Curio 1983; Forslund & Pärt 1995; Descamps *et al.* 2006; Reed *et al.* 2008). Therefore, individuals might delay the onset of breeding until 74 their condition allows them to display intensively and compete for mates and thus 75 reduce the fitness costs associated with an earlier onset of breeding (Curio 1983; 76 Beauchamp & Kacelnik 1990; Forslund & Pärt 1995). Apparent age-dependent 77 reproductive success may also arise due to selection acting against individuals in poor 78 condition leading to an increase in the proportion of 'high quality' individuals within a 79 cohort (selective appearance of good reproducers / selective disappearance of bad 80 reproducers; van de Pol & Verhulst 2006), or due to the negative effects of ageing on 81 individuals' physiological condition, which leads to a reduction in reproductive 82 success, i.e. senescence (Stearns 1992; Williams et al. 2006; Bouwhuis et al. 2010; 83 Nussey et al. 2013). All these mechanisms can explain the association between male 84 age and their ability to mate and hence produce offspring. Thus, fully understanding 85 the variation in individuals' fitness requires quantifying the temporal dynamics in 86 reproductive success that emerges from life history trade-offs.

87 In males of species with strong sexual selection, the allocation of resources to 88 reproduction embodies the expression of costly sexual traits that underlie males' 89 competitive ability and attractiveness (Andersson 1994; Höglund & Sheldon 1998). 90 Males' age has been shown to associate with the size and quality of sexual traits such 91 as antler size, tail feathers and plumage colouration (e.g. Nussey et al. 2009; Evans, 92 Gustafsson, & Sheldon 2011; Kervinen et al. 2015) most probably through changes in 93 body condition throughout individuals' lifespan (Höglund & Sheldon 1998; Emlen et 94 al. 2012). Therefore, quantifying the effect of the expression of a specific trait on 95 male mating success requires accounting for direct and indirect age-specific effects of 96 other traits.

97 Studies quantifying sexual selection on male traits while also accounting for age 98 are still relatively scarce (Coltman *et al.* 2002; Kruuk *et al.* 2002; Freeman-Gallant *et* 

99 al. 2010), with the majority examining single traits (though see Freeman-Gallant et al. 100 2010). One reason for the paucity of such studies is that accurate measures of male 101 mating success are difficult to collect in natural populations, because copulations can 102 be difficult to observe and animals move in and out from the study area (Coltman et 103 al. 1999; Kruuk et al. 2002). Molecular markers can be used to deduce female mating 104 behaviour and male mating success, but the accuracy of such methods will strongly 105 depend on our ability to quantify accurately male mating success and to identify 106 multiple matings, as well as on the occurrence of post-copulatory sexual selection 107 (Eberhard 1996; Birkhead 1998; Birkhead & Møller 1998). Therefore, better 108 understanding the magnitude of sexual selection acting on male traits requires using 109 study systems in which male mating success can reliably be measured and related to 110 the expression of the multiple traits across their entire lifespan (Clutton-Brock & 111 Sheldon 2010).

112 In the lekking black grouse (Lyrurus tetrix, formerly Tetrao tetrix), male mating 113 success can easily be recorded since copulations are very conspicuous and most 114 copulations take place on leks (Lebigre et al. 2007). Females usually mate only once 115 with one male and the parentage analyses demonstrated a very high concordance 116 between observed mating success and genetic reproductive success (in 94% of the 117 cases, the presumed father sired all the hatchlings of the brood; Lebigre *et al.* 2007). 118 Males are very faithful to the leks where they initially established their territories and 119 rarely switch lek sites (Rintamäki et al. 1995; Höglund et al. 1999; Lebigre et al. 120 2008). Hence, we can monitor the mating success of the same individuals over their 121 whole lifespan, as all males holding central territories and >90% of all territorial 122 males were ringed. Male mating skew is substantial due to the intense male-male 123 competition and clear female choice (Alatalo, Höglund & Lundberg 1991; Alatalo et 124 al. 1992). In theory, males with no mating success could gain indirect fitness benefits 125 by increasing the mating opportunities of their close relatives, but such kin selected 126 benefits amount only to ca. 3% of a copulation in this study system (Lebigre et al. 127 2014). Contrary to most previous studies that focused on one or two sexually-selected 128 traits, male black grouse express multiple morphological and behavioural traits that 129 are related to male mating success (these traits are partially correlated; see 130 Supplementary Table S1). Lyre length and quality (Höglund et al. 1994, Rintamäki et 131 al. 2001), the size of testosterone-dependent red eye combs (Rintamäki et al. 2000), 132 body mass (Rintamäki et al. 2001, Lebigre, Alatalo & Siitari 2013) and blue chroma 133 colouration of breast feathers (Siitari et al. 2007) are all positively correlated with 134 male mating success. Moreover, females prefer mating with males that have high lek 135 attendance (Alatalo et al. 1992), fight frequently and successfully against other males 136 (Hämäläinen et al. 2012), and occupy central territories on leks (Hovi et al. 1994). 137 However, a substantial amount of the variation in all these traits is explained by male age (Kervinen *et al.* 2015; unconditional  $R^2$  range in this study: 0.18–0.78, see 138 139 Nakagawa & Schielzeth 2013 for detailed methods).

140 We used longitudinal data describing male mating success and the expression of 141 multiple sexually-selected morphological and behavioural traits to quantify changes in 142 mating success with male age and determine the relative contribution of the age-143 specific trait expression to this pattern. We first tested the hypothesis that the variation 144 in male mating success is age-dependent by quantifying the effects of age, age of first 145 lek attendance (AFL), lifespan and terminal investment on males' annual mating 146 success (AMS). This enabled us to separate the effects of within-individual 147 improvement and senescence on AMS from selective appearance and disappearance 148 of certain phenotypes in the population. We then quantified the effect of each 149 morphological and behavioural trait on male AMS while accounting for other 150 unmeasured age-specific effects using univariate sexual selection differentials and 151 multivariate sexual selection gradients with male age as a covariate. We tested 152 whether these sexual selection gradients were age-specific by including an interaction 153 term between male age and trait expression. Finally, we re-quantified the univariate 154 sexual selection differentials and multivariate sexual selection gradients without male 155 age as a covariate to determine whether accounting for age effects (and hence the 156 expression of other age-related traits not directly used as covariates in the models) 157 changes the strength of sexual selection acting on the measured traits.

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#### 159 Materials and methods

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#### 161 STUDY POPULATION AND FIELD METHODS

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During 2002–2013, we collected longitudinal data describing male mating success and multiple sexual traits in five study sites in Central Finland (peat bogs with high visibility, *ca.* 62°15'N; 25°00'E). In addition, males were monitored during the mating season in spring 2014 to ensure the use of complete cohorts (no ringed males alive) in the analyses. Each study site is a local main lek with 5–40 territorial males, and as local hunting clubs have refrained from hunting on these areas, the age structure of the study population can reasonably be assumed to be natural.

Data on morphological traits were collected from the study sites annually in January–March by catching birds from the winter flocks with oat-baited walk-in traps (for methods, see Kervinen *et al.* 2012; Lebigre *et al.* 2012). Each captured individual was aged as yearling or older according to plumage characteristics and individually 174 ringed for future identification with an aluminium tarsus ring with a unique serial 175 number and three coloured tarsus rings. All captured birds were blood sampled (for a 176 parallel study) and measured for body mass, lyre (i.e. tail) length, eye comb size and 177 blue chroma colouration of breast feathers (Siitari *et al.* 2007; Lebigre *et al.* 2012).

Male mating success and lekking behaviour were recorded on the study sites 178 179 (leks) annually during the peak mating season (the period in late April – early May 180 when virtually all copulations take place, Ludwig et al. 2006). During this period, 181 typically lasting 1 to 2 weeks depending on the weather conditions, all study leks were 182 monitored daily from ca. 3:00 to 8:00 a.m. by observers located in hides near the lek 183 arenas (for detailed methods, see Kervinen et al. 2012; Lebigre et al. 2012). Male 184 behaviour was scan sampled at regular intervals and categorised as rookooing, hissing 185 (the two main vocal displays), fighting or inactive (methods and behaviours are 186 described in detail in Höglund, Johansson & Pelabon 1997). Lek attendance 187 (proportional to the most commonly present male on the same lek), the relative 188 proportion of each behaviour, and males' territory distances from the lek centre 189 (hereafter "territory centrality"; see Lebigre et al. 2012) were calculated for all 190 individually identifiable males from the records (see Rintamäki et al. 1995; Alatalo et 191 al. 1996). Finally, we documented the occurrence of copulations, which are easy to 192 observe, as males flap their wings conspicuously when mounting females, their 193 location and the individuals involved.

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195 DATA VALIDATION

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In these analyses, we only included the males that were first caught as yearlings andcould therefore be precisely aged. The data consisted of records of AMS, lekking

199 behaviour (lek attendance, fighting rate, territory centrality) and morphological traits 200 (body mass, lyre length, blue chroma, eye comb size) for 164 male black grouse with 201 known year of hatching (2001–2008), known AFL (at age 1: N = 89, at age 2: N = 66, 202 at age 3: N = 9) and known lifespan of 1 to 6 years (412 records in total); no males 203 with detailed life history survived >6 years old. Due to male philopatry for their initial 204 lekking site (Rintamäki et al. 1995; Höglund et al. 1999; Lebigre et al. 2008), lifespan 205 was defined as the age at which the male was last time seen on the lek. Complete 206 cohorts (i.e. no ringed males of the cohort alive in spring 2014) were available for 207 males hatched in 2001–2008. Cohorts hatched 2009 onwards were excluded because 208 of large proportions (>10%) of individuals being still alive in spring 2014.

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#### 210 STATISTICAL ANALYSES

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212 All analyses were run in R 3.0.3 (R Development Core Team 2014). We used 213 generalised linear mixed effects models (GLMM) to test the effects of age, AFL, 214 lifespan and terminal investment on AMS following the procedure detailed in van de 215 Pol & Verhulst (2006). We used varIdent variance error structure from Poisson 216 distribution ('gls' function from the R package 'nlme') to account for the overdispersion in the data. Age and age<sup>2</sup> represented the linear and quadratic within-217 individual change in AMS with age. A significant positive effect of AFL on AMS can 218 219 be interpreted as evidence for selective appearance of good reproducers. A significant 220 negative effect of lifespan (i.e. age at last lek) on AMS, in turn, can be interpreted as 221 evidence for selective disappearance of poor reproducers. The interactions of AFL 222 and lifespan with age were also introduced as fixed effects to account for potential life 223 history differences (e.g. reproduction at early ages and short lifespan vs. long lifespan

and reproduction at older ages). A binary fixed effect 'terminal event' (TE, indicates whether it was the male's last year alive or not) and its interaction with age was included to account for potential terminal investment in reproduction, as this can affect the interpretation of evidence of senescence (Rattiste *et al.* 2004; Hammers *et al.* 2012). Individual identity nested within study site was fitted as a random effect in all models to account for multiple measures from the same individuals and locations.

230 Model selection was carried out for the pre-selected set of biologically relevant 231 candidate models following the AIC-IT procedure (Table 1), and models within 95% cumulative model weight (ca.  $\Delta AIC_c < 6$ ) were averaged (Burnham, Anderson & 232 233 Huyvaert 2011; Grueber et al. 2011). Then, the relative importance and model-234 averaged coefficient estimates with unconditional SE and unconditional 95% CI were 235 calculated for each parameter in the averaged models (R package 'AICcmodavg' v. 236 2.0-1, Mazerolle 2014). As random slopes cannot be fitted in gls models, we repeated 237 the same analysis using 'lme' function from the R package 'nlme' with individual 238 identity nested within study site fitted as the random intercept and its interaction with 239 age fitted as the random slope to reduce the risk of potential type I and type II errors 240 (Schielzeth & Forstmeier 2009). However, as including random slopes did not 241 significantly change the interpretation of the results (Supplementary Table S2, S3, S4 242 and S5) and as the gls models better accounted for the overdispersion in the data 243 (based on the model residuals), we present the results of the initial gls approach.

Measures of senescence that do not account for within-individual age effects (random slopes) may mask within-individual patterns of ageing and mix together individuals with differing life history strategies (Bouwhuis *et al.* 2009). As including random slopes did not significantly alter the interpretation of our results, we could ignore variation in within-individual patterns of ageing. Thus, we carried out a

Poisson GLMM on AMS data after the age at which AMS showed its statistical peak with age and AFL fitted as linear effects only and individual identity nested within study site fitted as a random effect. In this additional analysis, a significant negative effect of age can be considered as evidence of senescence (Bouwhuis *et al.* 2009).

253 To quantify age-specific sexual selection, we first standardised male AMS by 254 dividing each male's AMS by the annual mean AMS of all males on the same lek: the 255 total number of copulations on the lek varied between sites and years (mean  $\pm$  SD = 256  $30.84 \pm 20.18$ , N = 49), but it was not correlated with individual AMS (Spearman's r = 0.05, p = 0.323, N = 412). We then standardised (i.e. zero-centred: mean = 0, SD = 1) 257 258 each morphological and behavioural trait using the yearly population mean as the 259 reference and related each standardised trait and their quadratic terms to male standardised AMS using age and age<sup>2</sup> as covariates to obtain linear and nonlinear 260 261 sexual selection differentials (s<sub>i</sub> and s<sub>i</sub>). In addition, the models included the 262 interaction of male age and trait expression to determine whether the magnitude and 263 direction of sexual selection on a specific trait changes with age. When the interaction 264 term or s<sub>i</sub> were nonsignificant, s<sub>i</sub> was quantified in the absence of these terms. 265 Univariate selection differentials measure both direct selection acting on each trait and 266 indirect selection resulting from selection acting on other traits with which the focal 267 trait is correlated. Therefore, we also carried out multivariate selection analyses where 268 multiple correlated (see Supplementary Table S1) traits and their quadratic terms were 269 used simultaneously to quantify linear ( $\beta$ ) and nonlinear ( $\gamma$ ) sexual selection gradients. 270 In the multivariate analysis, the parameter estimate for each trait accounts for the 271 selection acting on the other correlated traits, and hence allows direct comparison of 272 the relative contribution of each trait to the variance in male mating success. The 273 interactions of male age and trait expressions were included in the model, and when

274 the interaction term or  $\gamma$  were nonsignificant,  $\beta$  was quantified in the absence of these 275 terms. Due to our relatively limited sample size and the large number of parameters 276 (two parameters for age, seven parameters for directional selection, seven parameters 277 for nonlinear selection, seven interaction terms and the random effect term), the 278 multivariate analysis was conducted separately for male morphological and 279 behavioural traits which are partly independent (Rintamäki et al. 2001; Lebigre et al. 280 2012; Kervinen *et al.* 2015). Therefore, using age as a covariate in the models relating 281 for instance male mating success to the expression of the various morphological 282 ornaments enables us to account partially for the expression of the age-related 283 behavioural traits and other potentially unmeasured traits. Coefficients of the 284 quadratic terms ( $s_i$  and  $\gamma$ ) were doubled to measure balancing and disruptive selections 285 respectively (Stinchcombe et al. 2008). Finally, for comparison and to estimate the age-(in)dependency of sexual selection acting on male traits, we quantified  $s_i$ ,  $s_i$ ,  $\beta$  and 286 287  $\gamma$  as described above but without accounting for individual age.

288

289 **Results** 

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## 291 AGE EFFECTS ON MATING SUCCESS

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There was substantial variation in LMS as 52% (85/164) of the males did not mate, whilst the top 12 males accounted for 50% of all observed copulations (N = 480; maximum LMS = 32, Fig. 1). Most males reproduced only in a single year (68%; 54/79) and of these, 74% (40/54) died after this reproductive event. Of those 25 males that reproduced in multiple years, 16 males survived after the year of their maximum AMS and 20 males received >50% of their LMS in a single year (Supplementary Fig.
S1).

300 Nine of the pre-selected candidate models of AMS were within 95% 301 cumulative model weight which led to multimodel inference (Table 1). Age was 302 included in all and AFL in all but one of the averaged models and thus they had the 303 highest relative importance (Table 2). Moreover, age and AFL were the only variables 304 for which the 95% CI of the averaged coefficient estimate did not overlap 0. AMS 305 increased with age until the statistical maximum at age of 3.7 years after which it 306 levelled off but did not decrease significantly towards the end of life (GLMM: -0.709 307  $\pm$  0.878, t = -0.807, P = 0.423; Fig. 2 a). In addition to these direct age effects, males' 308 AFL was important; males lekking at age 1 tended to have higher AMS at age 2 than 309 males that began lekking at age 2, and males that began lekking at age 1 or 2 tended to 310 have higher AMS at age 3 than males that began lekking at age 3 (Fig. 2 b). Among 311 males known to be territorial in our study leks, these differences were absent at older 312 ages and all strategies led to similar LMS (mean  $\pm$  SE: 2.71  $\pm$  0.60, 3.20  $\pm$  0.69 and 313  $3.11 \pm 1.62$  for males with AFL = 1 (N = 89), AFL = 2 (N = 66) and AFL = 3 (N = 9), 314 respectively; ANOVA:  $F_{1,162} = 0.242$ , P = 0.623, N = 164). Lifespan was not 315 correlated with AMS and there was no clear evidence of terminal investment in 316 reproduction (Table 2). Lastly, there was a significant positive relationship between 317 AFL and lifespan ( $r_s = 0.40$ , N=164, P<0.001), indicating that males starting to lek 318 older had longer lifespans.

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### 320 SEXUAL SELECTION ON MALE TRAITS

322 Sexual selection operated on all behavioural traits but not on all morphological traits 323 when accounting for males' age. The significant interactions of lek attendance and 324 lyre length with age indicated that the magnitude of sexual selection operating on 325 these traits increased with age (Table 3 a, Fig. 3 a, b). Males' AMS increased non-326 linearly  $(s_i)$  with lek attendance and territory centrality (Fig. 4 a, b) and linearly  $(s_i)$ 327 with fighting rate (Table 3 a). Conversely, males' AMS increased linearly with body 328 mass but for other morphological traits s<sub>i</sub> and s<sub>i</sub> were nonsignificant. When accounting 329 for age and the expression of other morphological or behavioural traits (multivariate 330 sexual selection gradients), we found a significant interaction of age with body mass 331 on AMS, indicating stronger sexual selection on body mass in older males (Table 3 a; 332 Fig. 3 c). Moreover, males' AMS increased non-linearly ( $\gamma$ ) with lek attendance and 333 linearly ( $\beta$ ) with body mass (Table 3 a).

334 When not accounting for male age, the magnitude of univariate sexual selection 335 differentials increased and the sexual selection gradients became significant on 336 morphological traits. Specifically, AMS increased significantly linearly (s<sub>i</sub>) with lyre 337 length and eye comb size and non-linearly (s<sub>i</sub>) with blue chroma (Table 3 b). For body 338 mass, the age-corrected and age-independent sexual selection differentials changed 339 from significantly linear (s<sub>i</sub>) to significantly non-linear (s<sub>i</sub>), probably as a result of 340 within-age differences in AMS and body mass relationship. For behavioural traits 341 there were no substantial differences in the significance of s<sub>i</sub> and s<sub>i</sub> whether 342 accounting for age or not. Controlling for the expression of other morphological or behavioural traits but not for age revealed that males' AMS increased significantly 343 344 non-linearly ( $\gamma$ ) with body mass, lek attendance and territory centrality (Table 3 b).

345

### 346 **Discussion**

#### 348 AGE EFFECTS ON MATING SUCCESS

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350 Males' AMS was strongly associated with their age, as it first increased linearly until 351 reaching a peak or plateau at age 3 or 4 and levelling off until death. Terminal effects, 352 i.e. individuals investing differently in their last breeding attempt, could alter 353 conclusions about age-specific breeding success (Rattiste 2004). Reproductive effort 354 can increase when residual reproductive value decreases, i.e. towards the end of life 355 (sensu terminal investment: Williams 1966; Pianka & Parker 1975; Clutton-Brock 356 1984). Indeed, terminal investment in female reproduction has been documented in 357 many species (Creighton, Heflin & Belk 2009; Fisher & Blomberg 2011), but 358 evidence in birds and especially for males are scarce (but see e.g. Velando, Drummond & Torres 2006). In addition, several studies have found increased 359 360 reproductive effort and/or sexual signalling towards the end of life (e.g. moose Alces 361 alces: Mysterud, Solberg & Yoccoz 2005; mealworms Tenebrio molitor: Sadd et al. 2006), including in black grouse (Kervinen et al. 2015). However, we found here that 362 363 this recently documented late life increase in reproductive effort and sexual signalling 364 did not lead to changes in mating success towards the end of life in male black grouse. 365 Therefore, either these two processes are disconnected and females discriminate 366 against old and very active males in a similar way as they discriminate against very 367 active yearling males (Siitari et al. 2007; Kervinen et al. 2012) or, the previously 368 documented terminal investment in reproductive effort (Kervinen et al. 2015) 369 dampens a potential decline in mating success. The alternative to an increase in 370 reproductive effort is a decline in fitness, i.e. senescence. However, we did not detect 371 significant reproductive senescence in male black grouse in contrast to previous long372 term studies (e.g. Bouwhuis et al. 2009; reviewed in Nussey et al. 2013). This is 373 because most males that had any mating success reproduced only once and tended to 374 die directly after reproducing. This means that their highest AMS was in their last 375 year of life. Such high mortality possibly stems from the high energetic or 376 physiological costs of reproductive effort (Lebigre, Alatalo & Siitari 2013). However, 377 not all males had their highest AMS in their last year of life, but for those males that 378 bred more than once, most copulations still occurred in a single year and there were 379 sharp declines in AMS after this. This means that across the population, terminal 380 effects went in both directions depending on the number of reproductive episodes, 381 contributing to a lack of either terminal investment or reproductive senescence being 382 detected.

383 Male lifespan had low relative importance and its averaged coefficient estimate 384 was positive, but nonsignificant, indicating that short- and long-lived males did not 385 differ in their age-specific AMS. Previously we found significant positive effects of 386 lifespan on male morphological and behavioural traits (Kervinen et al. 2015). Our 387 results suggest that the trajectories of expression of male traits are partially 388 independent from male mating success, and that the outcome of sexual selection 389 cannot be directly linked to the expression of each trait separately. This is perhaps not 390 unexpected since age-specific trajectories typically vary across multiple traits (e.g. 391 Hayward et al. 2015), meaning that phenotypic, reproductive and survival traits may 392 be decoupled (Evans et al. 2011; Bouwhuis et al. 2012). However, relatively little is 393 known about age-specific trade-offs between sexually-selected traits and reproductive 394 traits (Cornwallis, Dean & Pizzari 2014). The fact that there are differences in trait 395 trajectories, strongly suggest that no single process underlies these patterns, but there

is critical need to quantify the physiological basis for such potential trade-offs(Cornwallis, Dean & Pizzari 2014).

398 Across the males that became territorial in the studied leks, there was a 399 significant negative effect of AFL on AMS, indicating that males that began lekking 400 at ages 1 or 2 tended to have higher age-specific AMS at ages 2 or 3 than the males 401 that were on their first lek at ages 2 or 3, respectively. This confirms a previous study 402 of the same black grouse population, but without accurate age estimates, which 403 showed that lekking experience is positively correlated with mating success (Kokko et 404 al. 1998). Thus, males with low AFL had higher age-specific AMS either because 405 they defended more central territories, or because they were better able to fight 406 against their neighbours due to their past experience in displaying with other males. 407 However, due to the high mortality among territorial yearling and 2-year-old males, 408 AFL was unrelated to LMS, which provides a mechanism to help maintaining the 409 coexistence of multiple male life history strategies in this population (Kervinen et al. 410 2015).

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## 412 SEXUAL SELECTION ON MALE TRAITS

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Given that both male mating success (this study) and the expression of a multitude of male morphological and behavioural traits (Kervinen *et al.* 2015) depend on age, our aim was to quantify whether accounting for such age effects (and hence indirectly for the effect of age-related traits not used in the selection gradients) affected our estimates of sexual selection operating on the morphological and behavioural traits. We found that sexual selection operated strongly on most morphological and behavioural traits when not accounting for male age but these effects strongly 421 declined when accounting for age. A similar effect has been found in common terns 422 (Sterna hirundo) in which accounting for age decreased the magnitude of viability 423 selection associated with individuals' trait expression (Zhang et al. 2015). Univariate 424 sexual selection differentials indicated sexual selection acting more generally on male 425 lekking behaviour than on male morphological traits when accounting for males' age. 426 This might primarily be due to age-specific sexual selection acting on body mass, lyre 427 length and lek attendance. Indeed, young males have low mating success regardless of 428 their body mass, lyre length, as they lack lekking experience and central territories. In 429 contrast, in prime age males and very old males that have the experience, high body 430 mass is selected for as it enables successful lekking performance (Lebigre, Alatalo & 431 Siitari 2013). To our knowledge, only two studies on ungulates have quantified age-432 specific sexual selection gradients; one showed that sexual selection in horn length of 433 big horn sheep increased with age (Coltman et al. 2002), and the other reported strong 434 selection for antler size in red deer during peak reproductive age (Kruuk et al. 2002). 435 In general, such age effects are not due to age *per se* but are most likely due to the 436 expression of other age-related traits that are not accounted for when quantifying the 437 selection gradient.

438 Multiple sexual signals can convey different messages about male quality over a 439 range of time periods (Doucet & Montgomerie 2003) leading to variation in female 440 choice or mate choice errors which may dampen or even mask sexual selection 441 (Chaine & Lyon 2008). This is very likely to apply to black grouse males' 442 morphological traits as lyre length and blue chroma reflect male condition during 443 moult which occurs shortly after the previous mating season (Siitari et al. 2007) while 444 body mass and eye comb size are more plastic and reflect shorter-term male condition 445 (Pérez-Rodríguez & Viñuela, 2008; Hämäläinen et al. 2012; Lebigre, Alatalo &

446 Siitari 2013). The stronger sexual selection differentials found on male behaviour 447 (such as lek attendance, fighting rate and territory centrality) might be explained by 448 their extreme variation and strong relationship with males' current body condition 449 (Hämäläinen et al. 2012; Lebigre, Alatalo & Siitari 2013), which is itself age-450 dependent in this species (Kervinen et al. 2015). Moreover, multiple traits together 451 might provide a better overall measure of quality than each cue alone in this system 452 (redundant signal hypothesis; see Candolin 2003). Indeed, it usually takes years for a 453 male to obtain a central territory on a lek (Kokko et al. 1998, 1999; DuVal 2012). 454 Therefore, female preference for males displaying on central territories (Höglund & 455 Lundberg 1987; Hovi et al. 1994; Kokko et al. 1998, 1999; Bro-Jørgensen & Durant 456 2003), indicates that both males' short- and long-term phenotypic quality matters 457 (viability, long-term lekking effort). Although it is also possible that there are 458 uninvestigated components of male phenotype that are important for female choice, 459 our analysis shows that behavioural traits are under age-independent sexual selection 460 but that these traits themselves may not be enough for females to discriminate the 461 quality of males at certain ages. As a consequence, morphological traits (body mass, 462 lyre length) may be used as additional cues to supplement female choice (Møller & 463 Pomiankowski 1993). Morphological traits may also have a dual utility, by serving as 464 honest signals of male fighting ability or dominance (Berglund, Bisazza & Pilastro 465 1996), but previous evidence suggests this is unlikely in black grouse (Rintamäki et 466 al. 2000).

We found evidence of non-linear sexual selection gradients ( $\gamma$ ) for several traits which are commonly interpreted as evidence of balancing (negative  $\gamma$ ) or disruptive (positive  $\gamma$ ) selection (Stinchcombe *et al.* 2008; Hunt *et al.* 2009). However, the significant positive  $\gamma$  and s<sub>j</sub> values found in this study more likely suggest "threshold"

471 values for male lek attendance and territory centrality as male mating success sharply 472 increases when males have a lek attendance >0.8, and defend territories within *ca*. 20 473 metres from the lek centre. Other species show similar nonlinear "threshold" 474 relationships between mating success and both physical (e.g. bighorn rams Ovis 475 canadensis and horn length: Coltman et al. 2002) and behavioural traits (e.g. topi 476 bulls Damaliscus korrigum and territory centrality: Bro-Jørgensen & Durant 2003). In 477 black grouse, territory centrality and lek attendance are strongly related and the 478 threshold values found in this study may reflect the difference between the dominant 479 males displaying very actively on central territories and the other, less successful 480 males that hold territories further away from the lek centre. It is important to note that 481 the relationship between lek attendance and mating success is not due to missed 482 mating opportunities of the males attending less to the lek, because copulations take 483 place on the leks when all territorial males are present (Kokko et al. 1999; Lebigre et al. 2007). 484

485

### 486 CONCLUSIONS

487

488 This study shows that the overall variation in male black grouse mating success is 489 strongly related to male age with substantial nonlinear increase over individuals' 490 lifespan without a significant decrease at the end of life. Hence, age-related changes in 491 male trait expression explains a substantial amount of the total variation in male 492 mating success. Age of first lek attendance was negatively related to age-specific 493 mating success, but unrelated to males' lifetime mating success. Young age of first lek 494 attendance was also associated with shorter lifespan, indicating that different male life 495 history strategies coexist in this population. Sexual selection seemed to operate on 496 both morphological and behavioural traits when male age was not used to measure the 497 sexual selection gradients. However, when accounting for the effect of other age-498 related traits, males' mating success was directly related only to behavioural traits, but 499 was related to male body mass, lyre length and lek attendance in an age-specific 500 manner (i.e. selection acted on these traits in old males only). Hence, sexual selection 501 in this species primarily operates on male behaviour and morphological traits are used 502 as additional cues to supplement female choice. This study shows that the complex 503 dynamics of the expression of males' sexual traits with age needs to be accounted for 504 and failing to do so might lead to an overestimation of the magnitude of sexual 505 selection acting on these traits and of their potential microevolutionary changes.

506

507 Data Accessibility

508

509 We will archive the data associated with this manuscript to Dryad Digital Repository510 (http://datadryad.org/) should the manuscript be accepted.

511

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513

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Figure 1. The distribution of lifetime mating success (LMS) in male black grouse (N
= 164). Mating skew was substantial: most males never mated at all, whereas the top
12 males accounted for roughly the half (238/480) of all copulations.

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Figure 2. Variation in mean annual mating success (AMS, mean  $\pm$  SE) with age for a) all males (N = 164) and b) separately for males that started lekking at age 1 (black circles, N = 89), at age 2 (grey squares, N = 66) or at age 3 (white triangles, N = 9). N denotes the age-specific sample sizes. Note the different scales on the vertical axes. For the figure, ages 5 and 6 were combined to age class 5+. The high mean AMS with large SE at age 5+ of the males that began lekking at age 1 is largely affected by one male that had 15 copulations at age 6.

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Figure 3. The interactions of age with a) lek attendance, b) lyre length and c) body mass on annual mating success (AMS), indicating that the strength of sexual selection on these traits increased with age. All parameters were standardised to account for the variation lek size and the total number of copulations on the lek (see Materials and methods).

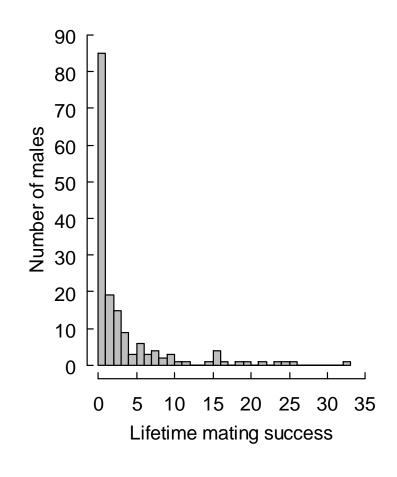
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Figure 4. Annual mating success (AMS) of yearling (filled triangles) and older (ages 2 to 6 years; open circles) male black grouse in relation to a) lek attendance and b) territory centrality (analysed as the male's territory distance from the lek centre). Significant  $s_j$  and  $\gamma$  values on these traits suggested thresholds after which AMS

- sharply increases. All parameters were standardised to account for the variation lek
- size and the total number of copulations on the lek (see Materials and methods).

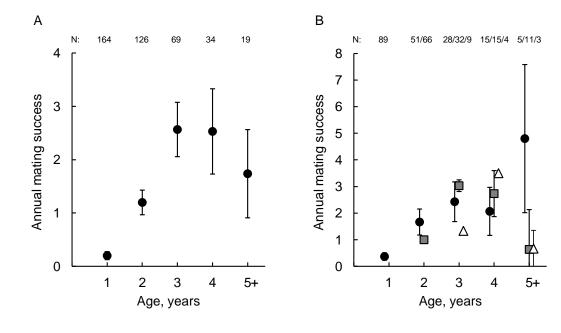
776 Figures

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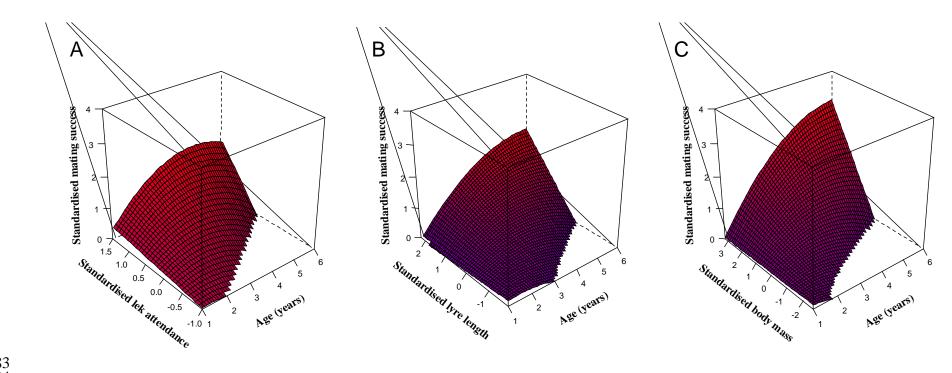


779 Figure 1.

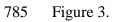


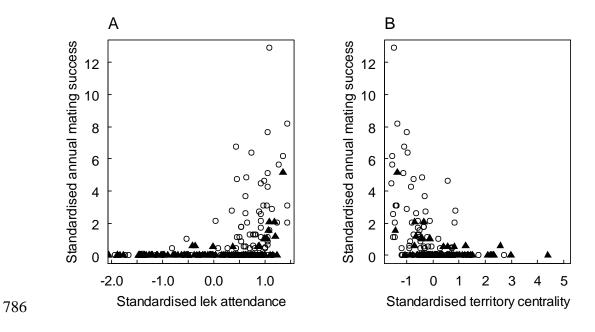


781 Figure 2.









787 Figure 4.

# 788 Tables

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Table 1. A suite of pre-selected biologically relevant candidate models predicting the effects of age, age<sup>2</sup>, age of first lek attendance (AFL),

791 lifespan (LS) and terminal event (TE; binary, indicates if it was the male's last year alive or not) on annual mating success in male black grouse,

with the number of estimated parameters for each model (k), Log-likelihood and AIC<sub>c</sub> values, model weights (*w<sub>i</sub>*), cumulative model weights

and evidence ratios (ER). Individual identity nested within study site was fitted as a random factor in all the models. Candidate models within

794 95% cumulative model weight were averaged (see Table 2).

Model	k	LogLik	AIC <sub>c</sub>	$\Delta AIC_{c}$	wi	Cum. w <sub>i</sub>	ER
$Age + Age^2 + AFL$	11	-858.52	1739.71	0.00	0.27	0.27	
Age + LS + TE + AFL + Age*LS + Age*TE	14	-855.56	1740.19	0.48	0.21	0.48	1.27
$Age + Age^2 + AFL + Age*AFL$	12	-858.03	1740.85	1.14	0.15	0.63	1.77
$Age + Age^2 + AFL + LS + TE + Age*LS + Age*TE$	15	-855.28	1741.77	2.06	0.10	0.72	2.81
Age + AFL + LS + TE + Age*AFL + Age*LS + Age*TE	15	-855.34	1741.90	2.20	0.09	0.81	3.00
$Age + Age^2 + AFL + LS + TE$	13	-858.01	1742.94	3.23	0.05	0.87	5.03
$Age + Age^2 + AFL + LS + TE + Age*AFL + Age*LS + Age*TE$	16	-855.08	1743.53	3.83	0.04	0.91	6.77
$Age + Age^2 + AFL + LS + TE + Age*AFL$	14	-857.54	1744.15	4.44	0.03	0.93	9.21
$Age + Age^2$	10	-862.01	1744.56	4.86	0.02	0.96	11.34
$Age + Age^2 + TE + Age*TE$	12	-860.60	1745.98	6.27	0.01	0.97	23.05
$Age + Age^2 + TE$	11	-861.98	1746.61	6.91	0.01	0.98	31.63
Age + LS + TE + Age*LS + Age*TE	13	-860.30	1747.52	7.82	0.01	0.98	49.82
Age + AFL	10	-863.69	1747.93	8.22	0.00	0.99	60.99
	$\begin{array}{l} Age + LS + TE + AFL + Age *LS + Age *TE \\ Age + Age^2 + AFL + Age *AFL \\ Age + Age^2 + AFL + LS + TE + Age *LS + Age *TE \\ Age + AFL + LS + TE + Age *AFL + Age *LS + Age *TE \\ Age + Age^2 + AFL + LS + TE \\ Age + Age^2 + AFL + LS + TE + Age *AFL + Age *LS + Age *TE \\ Age + Age^2 + AFL + LS + TE + Age *AFL \\ Age + Age^2 + AFL + LS + TE + Age *AFL \\ Age + Age^2 + TE + Age *TE \\ Age + Age^2 + TE \\ Age + LS + TE + Age *LS + Age *TE \\ \end{array}$	Age + Age² + AFL11Age + Age² + AFL + AFL + Age*LS + Age*TE14Age + Age² + AFL + Age*AFL12Age + Age² + AFL + LS + TE + Age*LS + Age*TE15Age + AFL + LS + TE + Age*AFL + Age*LS + Age*TE15Age + Age² + AFL + LS + TE + Age*AFL + Age*LS + Age*TE13Age + Age² + AFL + LS + TE13Age + Age² + AFL + LS + TE + Age*AFL + Age*LS + Age*TE16Age + Age² + AFL + LS + TE + Age*AFL14Age + Age² + AFL + LS + TE + Age*AFL14Age + Age² + AFL + LS + TE + Age*AFL14Age + Age² + AFL + LS + TE + Age*AFL11Age + Age² + TE + Age*TE12Age + Age² + TE + 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14	Age + AFL + Age*AFL	11 -862.82 1748.30	8.60 0.00 0.9	99 73.65
15	$Age + Age^2 + LS + TE$	12 -861.81 1748.41	8.70 0.00 1.0	00 77.52
16	$Age + Age^2 + LS + TE + Age*LS + Age*TE$	14 -860.00 1749.05	9.35 0.00 1.0	00 107.08
17	Age + AFL + LS + TE	12 -863.38 1751.55 1	1.85 0.00 1.0	00 373.39
18	Age + AFL + LS + TE + Age*AFL	13 -862.54 1752.00 12	2.29 0.00 1.0	00 467.29
19	Age	9 -866.88 1752.20 12	2.50 0.00 1.0	00 517.40
20	Age + TE	10 -866.80 1754.15 14	4.44 0.00 1.0	0 1366.80
21	Age + TE + Age*TE	11 -866.73 1756.13 10	5.42 0.00 1.0	0 3685.64
22	Age + LS + TE	11 -866.74 1756.13 10	5.43 0.00 1.0	0 3687.43
23	Null	8 -880.90 1778.15 38	8.45 0.00 1.0	$2.23 \times 10^8$

- 797 Table 2. The relative importance and averaged coefficient estimates with unconditional standard errors and 95% unconditional confidential
- <sup>798</sup> intervals of age, age<sup>2</sup>, age of first lek attendance (AFL), lifespan (LS) and terminal event (TE; binary, indicates if it was the male's last year alive
- 799 or not) and their relevant interactions on annual mating success in male black grouse.

Parameter	Relative importance	Estimate	Unconditional SE	95% unconditional CI
Age	1.00	2.15	0.49	1.19, 3.11
AFL	0.98	-0.37	0.16	-0.67, -0.06
Age <sup>2</sup>	0.66	-0.24	0.12	-0.48, 0.01
LS	0.52	0.22	0.20	-0.18, 0.61
TE	0.52	-0.46	0.38	-1.20, 0.28
Age:LS	0.44	-0.30	0.17	-0.63, 0.04
Age:TE	0.44	0.10	0.26	-0.41, 0.60
Age:AFL	0.31	-0.13	0.15	-0.43, 0.17

800 Table 3. Standardised linear  $(s_i)$  and nonlinear  $(s_j)$  univariate sexual selection differentials and linear  $(\beta)$  and nonlinear  $(\gamma)$  multivariate sexual

801 selection gradients in male black grouse with and without controlling for individual age. Estimates are provided with their standard errors. Bold

802 indicates significance at P < 0.05. Individual identity nested within study site was fitted as a random factor in all the models.

a) Controlled for individual age						b) Not controlled for individual age				
Trait	Si	Sj	age*trait	β	γ	age*trait	Si	Sj	β	γ
Morphological traits										
Body mass	$0.763 \pm 0.206$	$0.129\pm0.077$	$0.332\pm0.249$	$-0.366 \pm 0.136$	$0.034\pm0.113$	$0.728 \pm 0.301$	$0.209 \pm 0.050$	$0.222 \pm 0.074$	$0.133 \pm 0.069$	$0.208 \pm 0.088$
Lyre length	$-0.520 \pm 0.285$	$-0.161 \pm 0.163$	$0.743 \pm 0.376$	$-0.075 \pm 0.085$	$-0.053 \pm 0.115$	$0.783 \pm 0.438$	$\textbf{0.208} \pm \textbf{0.055}$	$0.198 \pm 0.104$	$0.064\pm0.074$	$0.027 \pm 0.114$
Blue chroma	$-0.028 \pm 0.040$	$0.110\pm0.069$	$0.076\pm0.119$	$-0.028 \pm 0.042$	$0.098\pm0.072$	$0.132\pm0.171$	$0.076\pm0.039$	$0.150\pm0.067$	$-0.010 \pm 0.042$	$0.098 \pm 0.072$
Eye comb size	$0.048 \pm 0.061$	$0.012\pm0.059$	$0.170\pm0.177$	$0.023\pm0.066$	$-0.033 \pm 0.064$	$0.336\pm0.227$	$\textbf{0.193} \pm \textbf{0.043}$	$0.040\pm0.058$	$0.092\pm0.060$	$-0.005 \pm 0.061$
Behavioural traits										
Lek attendance	$0.129\pm0.110$	$\textbf{0.455} \pm \textbf{0.078}$	$\textbf{0.471} \pm \textbf{0.172}$	$0.131\pm0.113$	$1.367 \pm 0.334$	$0.398 \pm 0.346$	$\textbf{0.450} \pm \textbf{0.045}$	$\textbf{0.508} \pm \textbf{0.077}$	$0.189 \pm 0.114$	$1.301 \pm 0.337$
Fighting rate	$0.186 \pm 0.066$	$-0.080 \pm 0.121$	$0.055\pm0.160$	$0.098 \pm 0.062$	$-0.026 \pm 0.108$	$-0.198\pm0.182$	$0.252\pm0.065$	$-0.121 \pm 0.122$	$0.103\pm0.063$	$-0.048 \pm 0.109$
Territory centrality*	$-0.552\pm0.080$	$0.342\pm0.066$	$-0.323 \pm 0.181$	$-0.049 \pm 0.064$	$0.134\pm0.075$	$-0.358 \pm 0.243$	$-0.620\pm0.078$	$0.370\pm0.066$	$-0.225\pm0.102$	$0.155 \pm 0.075$

803 \* Territory centrality was analysed as the male's territory's distance from the lek centre.