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1 **Disentangling the effects of genetic and environmental factors on movement**
2 **behaviour**

3

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1 **ABSTRACT**

2 Individual variability in animal movement behaviour is well documented for many species.
3 However, it remains unclear whether this variability reflects genetic variation, environmental
4 variation, or a combination of the two. Here, we conduct a cross-fostering experiment with the
5 aim of investigating the role of these two components in movement patterns during the post-
6 fledging dependence period and early natal dispersal of 21 eagle owls *Bubo bubo*. Our
7 experiment showed that cross-fostering did not influence any of the movement parameters
8 considered. Movement parameters were, however, affected by the age and sex of the owlets. We
9 therefore suggest that individual variability and family resemblance in movement behaviour
10 during the post-fledging dependence period and early natal dispersal might not be due to the
11 common genetic origin of siblings, but rather that it originates from factors related to the rearing
12 environment.

13 *Key-words:* *Bubo bubo*, cross-fostering, eagle owl, natal dispersal, post-fledging dependence
14 period

1 INTRODUCTION

2 Individual variability is well documented for many behaviours and life-history traits (Vindenes
3 & Langanen, 2015), yet the causes and mechanisms behind it are still poorly understood.
4 Differences **between and within** individuals, as well as similarities among relatives, may be due
5 to genetic variation (G), environmental variation (E) or a combination of these two effects (G x
6 E) (**Boake et al., 2002**; Clark & Ehlinger, 1987; Kruuk & Hadfield, 2007). The role of these
7 factors has been investigated for several traits (reviewed in: **Dingemane, Kazem, Réale, &**
8 **Wright, 2010**; van Oers *et al.*, 2005). Some heritability studies have focused on individual
9 variation in movement behaviour (Hansson, Bensch, & Hasselquist, 2003; Massot & Clobert,
10 2000; Massot, Huey, Tsuji, & Van Berkum, 2003; Matthysen, Van De Castele, & Adriaensen,
11 2005; Noordwijk, 1984; Pasinelli, Schiegg, & Walters, 2004; Pasinelli & Walters, 2002). These
12 studies showed that, even if many movement behaviours have a significant heritable component,
13 it is the combination of genetic and environmental components (G x E) which mainly determines
14 individual behavioural variation. Understanding the factors and mechanisms driving movement
15 behaviour is indeed a key question because movement behaviour affects individuals' survival
16 probability and reproductive success and, at a broader level, population dynamics (Morales et al.,
17 2010).

18 In birds, the post-fledging dependence period (hereafter PFDP) is defined as the period
19 between fledging and independence from the parents (Delgado, Penteriani, & Nams, 2009),
20 whereas natal dispersal (hereafter dispersal) represents the movement of an individual from their
21 birthplace to their first breeding location and involves three successive phases: departure,
22 transfer and settlement (Bowler & Benton, 2005; Ronce, 2007). Together, the PFDP and
23 dispersal represent an intensive period of experience and learning (Delgado, Penteriani, Nams, &
24 Campioni, 2009) during which a juvenile attains the necessary physical condition and abilities to
25 survive and eventually find a suitable breeding area. During these phases, different interacting
26 factors, such as the internal state of the individual (e.g. sex, age, body condition) and
27 environmental features (e.g. landscape characteristics, interaction with conspecifics), influence
28 individual movement behaviours (Bowler & Benton, 2005; Delgado, Penteriani, Revilla, &
29 Nams, 2010; Muriel, Ferrer, Balbontín, Cabrera, & Calabuig, 2015; Van Overveld, Adriaensen,
30 & Matthysen, 2011). In addition, a recent study found that movements of eagle owl (*Bubo bubo*)

1 siblings during natal dispersal were not independent of each other, suggesting a potential ‘family
2 effect’ on dispersal behaviour (Penteriani & Delgado, 2011). However, it remains difficult to
3 unravel the role of genetic and common environment effects in non-experimental studies
4 (Matthysen et al., 2005).

5 Few studies have performed brood manipulation, such as cross-fostering, to
6 experimentally separate genetic from common environment effects in movement behaviour
7 (Boonstra & Hochachka, 1997; Massot et al., 2003; Roche, Brown, & Brown, 2011). The
8 technique of cross-fostering, which consists of pairing nests and switching an equal number of
9 chicks of the same age and sex between the two families, represents a widely used tool to
10 experimentally separate the effect of the two potential sources of similarity between relatives, i.e.
11 genetics and a shared environment (Matthysen et al., 2005). The main assumption is that if cross-
12 fostered offspring behave more similarly to their biological parents and/or full siblings, this
13 behaviour probably has an important genetic component. On the contrary, if cross-fostered
14 offspring do not behave like their biological siblings, and their behaviour is more similar to their
15 foster siblings, environmental factors might be the dominant component determining individual
16 variation (Clayton, 1990; Soler, Moreno, & Potti, 2003).

17 Here, we performed a cross-fostering experiment on eagle owls to disentangle the effect
18 of a common rearing environment (E) from the genetic component (G) on individual movement
19 behaviour during the PFDP and early dispersal. We hypothesized that, if environmental factors
20 primarily shape movement behaviour and determine similar behaviour within the same brood
21 during the PFDP and dispersal periods, we should not observe significant differences in
22 movement between cross-fostered and control individuals of the same nest (environmental
23 hypothesis). Alternatively, if owlets reared in the same nest show different movement patterns,
24 with individuals behaving more similarly to their biological siblings reared in a different nest, we
25 could then hypothesize that a genetic component is the main force driving movement behaviours
26 (genetic hypothesis).

27

28 **METHODS**

29 *Field methods*

1 This study was conducted over two years (2015 and 2016), from March to December, in the
2 Sierra Norte of Seville (Sierra Morena, SW Spain 37°30'N, 06°03' W). During these periods, the
3 field work included the following temporal sequence: (a) nest checking, (b) nestling age and sex
4 determination, (c) set up and running of the cross-fostering experiment, i.e. exchanging chicks
5 between nests, and (d) radiotracking owls during PFDP and dispersal. We determined the age of
6 the owlets based on plumage characters (Penteriani, Delgado, Maggio, Aradis, & Sergio, 2005)
7 and their sex using DNA extracted from blood (Griffiths, R., Double, M.C., Orr, K. & Dawson,
8 1998)

9

10 *The cross-fostering experiment*

11 For the experiment, we performed partial cross-fostering, which involves pairing nests and
12 swapping one or two chicks (depending on brood size) between the two families, while leaving
13 at least one chick in the original nest as a control (Hadfield, Nutall, Osorio, & Owens, 2007;
14 Mateo & Holmes, 2004; Morrison, Ardia, & Clotfelter, 2009).

15 To avoid initial variations in **brood structure** which may affect the outcome of the
16 experiment, hatch date, brood size and sex ratio were maintained constant in each nest during the
17 experiment. We switched an equal number of males and females to obtain a homogenous sample
18 (Matthysen et al., 2005; Nicolaus et al., 2012; Winney, Nakagawa, Hsu, Burke, & Schroeder,
19 2015). Following these criteria, we performed the cross-fostering experiment between 7 nests
20 (one nest was used in both years) for a total of 4 experimental blocks, each block consisting of a
21 pair of nests ($n_{2015} = 3$; $n_{2016} = 1$). **A total number of 24 owlets were included in the study.**
22 **With such a design, we obtained two treatment groups: non-fostered owls (namely**
23 **treatment group 0; n = 14 owlets; 9 males and 5 females; $n_{2015} = 10$; $n_{2016} = 4$), which**
24 **comprised those individuals that remained in their original nest as controls, and fostered**
25 **owls (namely treatment group 1; n = 10 owlets; 4 males and 6 females; $n_{2015} = 8$; $n_{2016} = 2$),**
26 **composed of those individuals that were switched between paired nests of each**
27 **experimental block.** Previous experience of natural and artificial adoptions performed in the
28 studied population have shown that the survival of juveniles is not affected by brood switching
29 (Penteriani & Delgado, 2008).

30 *Radiotracking procedure*

1 We radio tagged the birds and performed the cross-fostering experiment when owlets were 30-35
2 days old. Individuals were fitted with a Teflon ribbon backpack harness that carried a 30 g radio
3 transmitter (henceforth ‘tag’; see Delgado & Penteriani, 2008). The weight of the tag was < 3%
4 of the weight of the smallest adult male (1550 g, mean \pm SD = 1667 \pm 104.8) and 3.5% of the
5 smallest fledgling weight (850 g, mean \pm SD = 1267 \pm 226.4 g). Because the chicks were still
6 growing when they were radio tagged, backpacks were adjusted so that the Teflon ribbon could
7 expand and allow for the increase in body size. We manipulated and marked owls under: (1) the
8 Junta de Andalucía – Consejería de Medio Ambiente permits No. SCFFSAFR/GGG RS 260/02
9 and SCFFS AFR /CMM RS 1904 / 02; and (2) the approval of the Comité de Ética de
10 Experimentación Animal of the Estación Biológica de Doñana-CSIC (CEEA-EBD_12_41).

11 Locations of radio tagged owls were determined using a 3-element hand held Yagi
12 antenna connected to a BIOTRACK portable receiver via bi-angulation. The **accuracy of**
13 **localizations was 83.5 \pm 49.5 m, estimated when, after a radiolocation, we needed to locate**
14 **where exactly one individual was (e.g. if it died).** During the PFDP, owlets were followed by
15 radio tracking during night sessions, which took place with an interval of 10 days. During these
16 sessions, all the owlets were radio-located throughout the entire night, from one hour before
17 sunset to one hour after sunrise. The time interval between successive individual locations was
18 ca. 1 hour and 30 minutes. During dispersal, owls were located on a weekly basis at their
19 daytime roosting sites (mean time between consecutive locations \pm SE = 6.9 \pm 0.1 days). During
20 the experiment, six owls were predated ($n_{2015} = 5$; $n_{2016} = 1$) and we lost the signal of three other
21 individuals. Thus, our final sample size was 21 owls for the PFDP sample and 15 owls for the
22 dispersal sample.

23

24

25 *Determination of PFDP and dispersal phases*

26 PFDP started when the juveniles left the nest (mean age = 40-45 days), and lasted until the
27 juveniles started to disperse (Delgado, Penteriani, & Nams, 2009). To identify the start of
28 dispersal, we plotted both the beeline distance from the natal nest for each location and the
29 average of the beeline distance between the whole set of locations and the nest, with the latter
30 value representing the global mean distance covered by each individual during dispersal

1 (Delgado & Penteriani, 2008). Dispersal started when the distance between successive moves
2 became larger than the average distance travelled by each bird. This happens when the distance
3 of each location from the nest starts to increase rather than fluctuate around a low value (Delgado
4 & Penteriani, 2008).

5

6 *Movement parameters*

7 To compare the two treatment groups, we quantitatively described the movement behaviour of
8 individuals during PFDP and dispersal. For each juvenile ($n_{\text{PFDP}} = 21$; $n_{\text{dispersal}} = 15$) and for the
9 two phases, we estimated seven movement parameters. (1) Step distance: distance between
10 consecutive locations. (2) Total distance: total distance covered by the owl. (3) Net distance: the
11 distance between the first and the last location collected. (4) Distance between the nest and each
12 of the location points. (5) Distance between different owls, via individual locations recorded at
13 the same time. (6) Movement speed: obtained by dividing the step distance by the time interval
14 between consecutive locations. (7) Movement direction (i.e. turning angles) between successive
15 locations. In addition, we estimated the size of the area explored by each owl during the whole
16 PFDP using the 100% minimum convex polygon in QGIS 2.14.3 Geographic Information
17 System (QGIS Development Team, 2016).

18

19 *Statistical analyses*

20 **For the statistical analyses, we considered all parameters except the step distance, as this**
21 **variable was highly correlated with speed ($r = \text{XX}$, $P = \text{YY}$).** Correlations between the other
22 variables were low ($r < \text{ZZ}$ in each case). For each of the six movement parameters considered,
23 we built a set of competing models which included all possible combinations of explanatory
24 variables, starting from the simplest null model (intercept-only model) to a full model that
25 included all the explanatory variables (Tables S2 and S4). To check model assumptions
26 (normality, independence, the presence of outliers), we first conducted a graphical data
27 exploration. Response variables following a normal distribution were modelled using linear
28 mixed models. When residuals did not follow a normal distribution, data were log transformed

1 and then checked again for the assumptions. When log transformation was not sufficient, we
2 applied a generalized linear mixed model with gamma distributions.

3 In each set of competing models, we included treatment group, age (except for net and
4 total distance in the dispersal phase) and sex of individuals, as well as their interactions, as fixed
5 factors. Random effects were held constant. For modelling movement parameters during the
6 PFDP, we included 5 random hierarchical factors, organized as follows: (1) year, (2)
7 experimental block, (3) nest, (4) night of radio tracking, and (5) individual. The night of radio
8 tracking was not included in the models built for net and total distance, as we only had one
9 observation per night for these variables. For dispersal, we included the same random factors,
10 except the night of radio tracking, as we had weekly observations for this period. In the models
11 built for net and total distance of dispersal, we also excluded the individual as a random factor,
12 given that for these two parameters we only had one observation per owl.

13 We selected the best competing model based on the Akaike's Information Criterion
14 corrected for small sample sizes (AICc; Burnham & Anderson, 2002), and calculated two
15 additional statistics for each model: Δ AICc and Weighted AICc, indicating the probability that
16 the model selected was the best among the competing candidates (Gelman & Hill, 2006). We
17 considered models with Δ AICc values lower than 2 as competitive. For each set of models, we
18 then employed model averaging on the 95% confidence set to derive parameter coefficients and
19 the relative importance values (RIV) of each explanatory variable using the full-model averaging
20 approach (Burnham & Anderson, 2002). **When high model selection uncertainty exists, model**
21 **averaging allows formal inference based on the entire set of models (or, in our case, the**
22 **95% confidence set) (Burnham & Anderson, 2002; Grueber, Nakagawa, Laws, &**
23 **Jamieson, 2011; Symonds & Moussalli, 2011). Parameter estimates produced by model**
24 **averaging derive from weighted averages of these values across all models in the set**
25 **considered (Symonds & Moussalli, 2011). In particular, the relative importance value**
26 **(RIV) of each explanatory variable is calculated by summing Akaike weights across all**
27 **models which contain the variable (Burnham & Anderson, 2002).**

28 Significance was assessed using P-values derived from the model averaged parameter
29 estimates and was set at $P \leq 0.05$.

1 To assess the variability between/within nests and between/within individuals, we
2 calculated the intraclass correlation coefficient (ICC) (Stoffel, Nakagawa, & Schielzeth,
3 2017; Zuur et al., 2009) for two of the random factors included in our models: “nest” and
4 “owl”. This coefficient represents the fraction of the total variance attributable to variation
5 among groups of the same class (Stoffel et al., 2017) and, in our case, describes the
6 correlation between owls reared in the same nest (random factor “nest”) and the
7 correlation between observations of the same individual (random factor “owl”).

8 For all selected models, we calculated the conditional deviance following Nakagawa &
9 Schielzeth, (2013). All analyses were performed using R 3.2.5 statistical software (R
10 Development Core 2016). GLMMs were run using the “lme4” (Bates & Sarkar, 2007), and
11 “nlme” (Pinheiro, Bates, DebRoy, Sarkar, & R Core Team, 2017) packages. Multi-model
12 inference and model averaging were run using the “MuMIn” (Bartón, 2013) package.

16 **RESULTS**

17 *GENERAL PATTERNS OF OWL MOVEMENTS*

18 *Post-fledging dependence period*

19 **During a total of 21 nights of radio tracking ($n_{2015} = 12$; $n_{2016} = 9$) we registered 1213**
20 **locations ($n_{2015} = 887$ with $n_0 = 488$ and $n_1 = 399$; $n_{2016} = 326$ with $n_0 = 254$ and $n_1 = 72$). A**
21 **summary of post-fledging movement parameters is given in Table S1.**

22 **During this phase, the average of all parameters considered (except total distance)**
23 **increased with time, with females generally showing higher values than males (Figure 1a).**

24 *Dispersal*

25 **During a total of 62 days of radio tracking ($n_{2015} = 38$; $n_{2016} = 24$), we registered 258**
26 **locations ($n_{2015} = 183$ with $n_0 = 93$ and $n_1 = 90$; $n_{2016} = 75$ with $n_0 = 54$ and $n_1 = 21$).**

1 In 2015, dispersal started when owlets were 172 ± 5.9 days old (mean \pm SE) ($n = 11$; range =
2 148-220 days), and in 2016 when owlets were 158 ± 19.2 days old ($n = 4$; range = 123-192
3 days). For both years combined, dispersal started at the age of 168 ± 6.5 days ($n = 15$; range
4 = 123-220 days). A summary of dispersal movement parameters is given in Table S3.

5 As for the PFDP, female average values were generally higher than male values. Average
6 distances from the nest and between-owl distances increased with time, while average speed
7 decreased (Figure 2a).

8 To give an idea of how the owls were moving during the two phases considered, we plotted
9 the path followed by one selected owl at different ages (Figures 1b and 2b).

10

11 ***THE CROSS-FOSTERING EXPERIMENT***

12 *Post-fledging dependence period*

13 For all movement parameters, selected models always included age. Treatment group appeared
14 in selected models of all parameters except speed, while sex appeared in selected models of all
15 parameters (Table S2). However, P and RIV values only supported the effect of age (Table 1),
16 which was significant ($P \leq 0.05$) and had a relatively high importance ($RIV > 0.90$) in 5
17 parameters (net distance, total distance, speed, distance between each recorded location and the
18 nest, distance between owls). **The effect of treatment group was never significant**, nor did it
19 have an especially high RIV (always < 0.80 ; Table 1). **Values of ICC (Table 2) for the random**
20 **factor “owl” (i.e. individual) were always 0.00, which means there was no correlation**
21 **between observations of the same individual. ICCs for the random factor “nest” ranged**
22 **between 0.35 and 0.99, which means there was a correlation between individuals raised in**
23 **the same nest, indicating that within-nest variation was greater than between nest**
24 **variation.**

25 *Dispersal*

26 Out of 6 parameters considered, for two of them (net and total distance) the best model was the
27 null model (intercept-only model; Table S4). For all remaining parameters, the selected models
28 always included age and sex, while the treatment group appeared in 2 models. P and RIV values
29 suggested that age had the main effect, showing $P \leq 0.05$ and $RIV > 0.90$ in 3 parameters (speed,

1 distance between each recorded location and the nest, distance between owls; Table 3). **Sex had**
2 **a significant effect** and had a high RIV value in 2 parameters (distance between each location
3 and the nest, distance between owls). **The effect of treatment group was never significant.** In
4 the majority of cases, the RIV was low (< 0.50 ; Table 3), with the exception of distance between
5 owls, where $RIV = 0.86$. **Values of ICC (Table 2) for the random factor “owl” were 0.00 for**
6 **the speed and 0.36 for the distance between owls and the nest. ICCs for “nest” were higher**
7 **(always $ICC > 0.40$; except for the speed). This suggests that there was a relatively high**
8 **correlation between owls raised in the same nest.**

9

10

11 **DISCUSSION**

12 Our results showed that the treatment group of the cross-fostering experiment did not affect
13 movement parameters during the two successive phases of eagle owl development. **Moreover,**
14 **we found variability in movement characteristics of owls reared in different nests. This**
15 **outcome reveals a scenario where all owls reared in the same nest, regardless of whether**
16 **they were cross-fostered or not, behaved similarly between each other and differently from**
17 **the owls reared in the other nests.** Thus, our experiment does not support the hypothesis of a
18 genetic basis for movement behaviour during PFDP and early dispersal. Our findings are in line
19 with previous studies on other species (Greenwood, Harvey, & Perrins, 1979; Massot & Clobert,
20 2000; Matthysen et al., 2005; Pasinelli & Walters, 2002; Waser & Thomas Jones, 1989), which
21 found no evidence for the heritability of movement behaviour during the dispersal phase. These
22 authors suggested that individual variability and family resemblance in movement behaviour
23 may instead be explained by environmental and social factors, such as nest location, the
24 landscape surrounding the nest site, family bonds and other factors acting on the whole brood
25 during rearing. In contrast, few other studies on birds (Hansson et al., 2003; Pasinelli et al., 2004)
26 have found evidence of a genetic basis for movement behaviour. However, although these
27 studies support a heritable component, they acknowledge that this behaviour may also be partly
28 modulated by environmental and social factors.

1 The moderate sample size utilized in the current study might have decreased the statistical power
2 of our findings and, thus, the probability of detecting a small genetic effect. However, our results
3 clearly support the hypothesis that environmental and social factors have a stronger effect than
4 genetics on the development of individual behaviour, and hence in determining similarity in
5 movement behaviour between siblings. In particular, during the PFDP, offspring are reared
6 together in the same environment. This implies that young birds move in the same area during
7 this crucial stage of their life and, thus, they are exposed to the same external conditions.
8 Moreover, during the PFDP, young eagle owls spend most of their time together, often exploring
9 nest surroundings, and remain in constant contact with their parents (Delgado, Penteriani, &
10 Nams, 2009). All of the common factors that owlets share during rearing may lead them to adopt
11 similar behavioural movement responses during the PFDP.

12 Environmental characteristics (e.g. landscape structure and composition) of the habitat
13 explored by an individual during the dispersal phase have already been shown to play an
14 important role in shaping movement behaviour in this species (Delgado et al., 2010; Penteriani &
15 Delgado, 2011), as owls born in the same place displayed similar movement patterns and started
16 the dispersal process at a similar age (Penteriani & Delgado, 2011). Furthermore, different
17 individuals moving through the same area had similar movement patterns (Delgado et al., 2010).
18 These authors suggested that individuals born in the same nest and/or moving through the same
19 area may face similar constraints, and this may lead to similar individual movement decisions.
20 **Another study on natural brood-switching in fledglings of eagle owls (Penteriani &**
21 **Delgado, 2008), which somewhat anticipated our results, showed that switched owls not**
22 **only were adopted by foster parents, but moved similarly to the resident fledglings.**

23 Furthermore, our results also showed that age influences the movement behaviour of
24 individuals during both the PFDP and dispersal phases. During the PFDP, young owls are still
25 developing their flight and cognitive abilities and, thus, the way they move around the natal area
26 changes over time (Delgado, Penteriani, & Nams, 2009). These progressive changes have
27 already been recorded in this and other avian species (O'Toole L. T., Kennedy P. L., Knight R.
28 L., 1999; Wood, Collopy, & Sekerak, 1998). During dispersal, an age effect may be explained by
29 the fact that this phase represents a multistep process during which individuals may show
30 different behaviours depending on their physical condition, as well as on their abiotic and social
31 environments (Bowler & Benton, 2005; Delgado & Penteriani, 2008; Delgado et al., 2010). The

1 sex of individuals also showed some effect on some movement parameters during early
2 dispersal. Although sex-biased dispersal has not been detected in previous studies on this eagle
3 owl population (Delgado et al., 2010), sex differences in dispersal behaviour are relatively well
4 documented for several bird species (Paul J. Greenwood, 1980; Massot & Clobert, 2000;
5 Newton, 2002). Nonetheless, it is worth mentioning here that we only analysed movement
6 behaviour during the early phases of dispersal and that different patterns may emerge when
7 considering the whole dispersal phase.

8 Overall, our experimental approach, which aimed to disentangle the contribution of
9 environmental and genetic effects on the movement behaviour of a long-lived species, did not
10 support the hypothesis of a genetic predetermination of individual movement behaviour during
11 the early stages of its life. On the contrary, the experiment supported an effect of the local
12 environment in influencing movement behaviour. However, we acknowledge that our approach
13 represents a first step towards a better understanding of a very complex process in behavioural
14 ecology. Further studies including other populations, larger sample sizes and detailed dispersal
15 movements spanning the entire year are undoubtedly needed to confirm the potentially limited
16 effect of the genetic component on animal movement behaviour.

17

18

19

20

21 REFERENCES

22 Bartón, K. (2013). Model selection and model averaging based on information criteria (AICc and
23 alike). *The Comprehensive R Archive Network*, 1, 13.

24 Bates, D., & Sarkar, D. (2007). lme4: Linear mixed-effects models using S4 classes.

25 Boake, C. R. B. C. R. B., Arnold, S. J. S. J., Breden, F., Meffert, L. M. L. M., Ritchie, M. G. M.
26 G., Taylor, B. J. B. J., ... Moore, a. J. A. J. (2002). Genetic tools for studying adaptation
27 and the evolution of behavior. *The American Naturalist*, 160 Suppl(december), S143–S159.

- 1 <https://doi.org/10.1086/342902>
- 2 Boonstra, R., & Hochachka, M. (1997). Maternal effects and additive genetic inheritance in the
3 collared lemming *Dicrostonyx groenlandicus*. *Evolutionary Ecology*, (2), 169–182.
4 <https://doi.org/10.1023/A:1018447815825>
- 5 Bowler, D. E., & Benton, T. G. (2005). Causes and consequences of animal dispersal strategies:
6 relating individual behaviour to spatial dynamics. *Biological Reviews of the Cambridge*
7 *Philosophical Society*, 80(2), 205–225. <https://doi.org/10.1017/s1464793104006645>
- 8 Burnham, K. P., & Anderson, D. R. (2002). *Model Selection and Multimodel Inference: A*
9 *Practical Information-Theoretic Approach (2nd ed)*. *Ecological Modelling* (Vol. 172).
10 <https://doi.org/10.1016/j.ecolmodel.2003.11.004>
- 11 Clark, A. B., & Ehlinger, T. J. (1987). Pattern and adaptation in individual behavioral
12 differences. *Perspectives in Ethology*, 7, 1–47. <https://doi.org/10.1007/978-1-4613-1815-6>
- 13 Clayton, N. S. (1990). The effects of cross-fostering on assortative mating between zebra finch
14 subspecies. *Animal Behaviour*, 40(6), 1102–1110. [https://doi.org/10.1016/S0003-](https://doi.org/10.1016/S0003-3472(05)80176-9)
15 [3472\(05\)80176-9](https://doi.org/10.1016/S0003-3472(05)80176-9)
- 16 Delgado, M. M., & Penteriani, V. (2008). Behavioral states help translate dispersal movements
17 into spatial distribution patterns of floaters. *The American Naturalist*, 172(4), 475–485.
18 <https://doi.org/10.1086/590964>
- 19 Delgado, M. M., Penteriani, V., & Nams, V. O. (2009). How fledglings explore surroundings
20 from fledging to dispersal. A case study with Eagle Owls *Bubo bubo*. *Ardea*, 97(c), 7–15.
21 [https://doi.org/Doi 10.5253/078.097.0102](https://doi.org/Doi%2010.5253/078.097.0102)
- 22 Delgado, M. M., Penteriani, V., Nams, V. O., & Campioni, L. (2009). Changes of movement
23 patterns from early dispersal to settlement. *Behavioral Ecology and Sociobiology*, 64(1),
24 35–43. <https://doi.org/10.1007/s00265-009-0815-5>
- 25 Delgado, M. M., Penteriani, V., Revilla, E., & Nams, V. O. (2010). The effect of phenotypic
26 traits and external cues on natal dispersal movements. *Journal of Animal Ecology*, 79(3),
27 620–632. <https://doi.org/10.1111/j.1365-2656.2009.01655.x>
- 28 Dingemanse, N. J., Kazem, A. J. N., Réale, D., & Wright, J. (2010). Behavioural reaction norms:

- 1 animal personality meets individual plasticity. *Trends in Ecology and Evolution*, 25(2), 81–
2 89. <https://doi.org/10.1016/j.tree.2009.07.013>
- 3 Gelman, A., & Hill, J. (2006). Data Analysis Using Regression and Multilevel/Hierarchical
4 Models. *Cambridge University Press, Analytical*(July), 625. Retrieved from
5 <http://www.loc.gov/catdir/enhancements/fy0668/2006040566-d.html>
- 6 Greenwood, P. J. (1980). Mating systems, philopatry and dispersal in birds and mammals.
7 *Animal Behaviour*, 28(4), 1140–1162. [https://doi.org/10.1016/S0003-3472\(80\)80103-5](https://doi.org/10.1016/S0003-3472(80)80103-5)
- 8 Greenwood, P. J., Harvey, P. H., & Perrins, C. M. (1979). The Role of Dispersal in the Great Tit
9 (*Parus major*): The Causes , Consequences and Heritability of Natal Dispersal. *Journal of*
10 *Animal Ecology*, 48(1), 123–142.
- 11 Griffiths, R., Double, M.C., Orr, K. & Dawson, R. J. G. (1998). A DNA test to sex most birds.
12 *Molecular Ecology*, 7(8), 1071–1075. <https://doi.org/10.1046/j.1365-294x.1998.00389.x>
- 13 Grueber, C. E., Nakagawa, S., Laws, R. J., & Jamieson, I. G. (2011). Multimodel inference in
14 ecology and evolution: Challenges and solutions. *Journal of Evolutionary Biology*, 24(4),
15 699–711. <https://doi.org/10.1111/j.1420-9101.2010.02210.x>
- 16 Hadfield, J. D., Nutall, A., Osorio, D., & Owens, I. P. F. (2007). Testing the phenotypic gambit:
17 Phenotypic, genetic and environmental correlations of colour. *Journal of Evolutionary*
18 *Biology*, 20(2), 549–557. <https://doi.org/10.1111/j.1420-9101.2006.01262.x>
- 19 Hansson, B., Bensch, S., & Hasselquist, D. (2003). Heritability of dispersal in the great reed
20 warbler. *Ecology Letters*, 6(4), 290–294. <https://doi.org/10.1046/j.1461-0248.2003.00436.x>
- 21 Ingleby, F. C., Hunt, J., & Hosken, D. J. (2010). The role of genotype-by-environment
22 interactions in sexual selection. *Journal of Evolutionary Biology*, 23(10), 2031–2045.
23 <https://doi.org/10.1111/j.1420-9101.2010.02080.x>
- 24 Kruuk, L. E. B., & Hadfield, J. D. (2007). How to separate genetic and environmental causes of
25 similarity between relatives. *Journal of Evolutionary Biology*, 20(5), 1890–1903.
26 <https://doi.org/10.1111/j.1420-9101.2007.01377.x>
- 27 Massot, M., & Clobert, J. (2000). Processes at the origin of similarities in dispersal behaviour
28 among siblings. *Journal of Evolutionary Biology*, 13(4), 707–719.

1 <https://doi.org/10.1046/j.1420-9101.2000.00202.x>

2 Massot, M., Huey, R. B., Tsuji, J., & Van Berkum, F. H. (2003). Genetic, prenatal, and postnatal
3 correlates of dispersal in hatchling fence lizards (*Sceloporus occidentalis*). *Behavioral*
4 *Ecology*, *14*(5), 650–655. <https://doi.org/10.1093/beheco/arg056>

5 Mateo, J. M., & Holmes, W. G. (2004). Cross-fostering as a means to study kin recognition.
6 *Animal Behaviour*, *68*(6), 1451–1459. <https://doi.org/10.1016/j.anbehav.2004.01.017>

7 Matthysen, E., Van De Castele, T., & Adriaensen, F. (2005). Do sibling tits (*Parus major*, *P.*
8 *caeruleus*) disperse over similar distances and in similar directions? *Oecologia*, *143*(2),
9 301–307. <https://doi.org/10.1007/s00442-004-1760-7>

10 Morales, J. M., Moorcroft, P. R., Matthiopoulos, J., Frair, J. L., Kie, J. G., Powell, R. a, ...
11 Haydon, D. T. (2010). Building the bridge between animal movement and population
12 dynamics. *Philosophical Transactions of the Royal Society B Biological Sciences*,
13 *365*(1550), 2289–301. <https://doi.org/10.1098/rstb.2010.0082>

14 Morrison, E. S., Ardia, D. R., & Clotfelter, E. D. (2009). Cross-fostering reveals sources of
15 variation in innate immunity and hematocrit in nestling tree swallows *Tachycineta bicolor*.
16 *Journal of Avian Biology*, *40*(6), 573–578. <https://doi.org/10.1111/j.1600->
17 [048X.2009.04910.x](https://doi.org/10.1111/j.1600-048X.2009.04910.x)

18 Muriel, R., Ferrer, M., Balbontín, J., Cabrera, L., & Calabuig, C. P. (2015). Disentangling the
19 effect of parental care, food supply, and offspring decisions on the duration of the
20 postfledging period. *Behavioral Ecology*, *26*(6), 1587–1596.
21 <https://doi.org/10.1093/beheco/arv114>

22 Nakagawa, S., & Schielzeth, H. (2013). A general and simple method for obtaining R² from
23 generalized linear mixed-effects models. *Methods in Ecology and Evolution*, *4*(2), 133–142.
24 <https://doi.org/10.1111/j.2041-210x.2012.00261.x>

25 Newton, I. (2002). Ecology and conservation of owls. *Ecology and Conservation of Owls*.
26 <https://doi.org/10.2307/3849247>

27 Nicolaus, M., Michler, S. P. M., Ubels, R., van der Velde, M., Bouwman, K. M., Both, C., &
28 Tinbergen, J. M. (2012). Local sex ratio affects the cost of reproduction. *Journal of Animal*
29 *Ecology*, *81*(3), 564–572. <https://doi.org/10.1111/j.1365-2656.2011.01933.x>

- 1 Noordwijk, A. J. Van. (1984). Problems in the Analysis of Dispersal and a Critique on its
2 'Heritability' in the Great Tit. *Journal of Animal Ecology*, 53(2), 533–544. Retrieved from
3 <http://www.jstor.org/stable/4532>
- 4 O'Toole L. T., Kennedy P. L., Knight R. L., M. L. C. (1999). Postfledging of Golden. *Wilson*
5 *Bull.*, 111(4), 472–477.
- 6 Pasinelli, G., Schiegg, K., & Walters, J. R. (2004). Genetic and environmental influences on
7 natal dispersal distance in a resident bird species. *The American Naturalist*, 164(5), 660–
8 669. <https://doi.org/10.1086/424765>
- 9 Pasinelli, G., & Walters, J. R. (2002). Social and environmental factors affect natal dispersal and
10 philopatry of male Red-cockaded Woodpeckers. *Ecology*, 83(8), 2229–2239.
11 [https://doi.org/10.1890/0012-9658\(2002\)083\[2229:SAEFAN\]2.0.CO;2](https://doi.org/10.1890/0012-9658(2002)083[2229:SAEFAN]2.0.CO;2)
- 12 Penteriani, V., & Delgado, M. M. (2008). Brood-switching in Eagle Owl *Bubo bubo* fledglings.
13 *Ibis*, 150(4), 816–819. <https://doi.org/10.1111/j.1474-919X.2008.00831.x>
- 14 Penteriani, V., & Delgado, M. M. (2011). Birthplace-dependent dispersal: Are directions of natal
15 dispersal determined a priori? *Ecography*, 34(5), 729–737. <https://doi.org/10.1111/j.1600-0587.2010.06773.x>
- 17 Penteriani, V., Delgado, M. M., Maggio, C., Aradis, A., & Sergio, F. (2005). Development of
18 chicks and predispersal behaviour of young in the Eagle Owl *Bubo bubo*. *Ibis*, 147(1), 155–
19 168. <https://doi.org/10.1111/j.1474-919x.2004.00381>
- 20 Pinheiro, J., Bates, D., DebRoy, S., Sarkar, D., & R Core Team. (2017). {nlme}: Linear and
21 Nonlinear Mixed Effects Models. Retrieved from <https://cran.r-project.org/package=nlme>
- 22 QGIS Development Team. (2016). QGIS Geographic Information System. Open Source
23 Geospatial Foundation. Retrieved from <http://qgis.osgeo.org>
- 24 Roche, E. A., Brown, C. R., & Brown, M. B. (2011). Heritable choice of colony size in cliff
25 swallows: does experience trump genetics in older birds? *Animal Behaviour*, 82(6), 1275–
26 1285. <https://doi.org/10.1016/j.anbehav.2011.09.005>
- 27 Ronce, O. (2007). How does it feel to be like a rolling stone? Ten questions about dispersal
28 evolution. *Annual Review of Ecology, Evolution, and Systematics*, 38(2007), 231–253.

1 <https://doi.org/10.1146/annurev.ecolsys.38.091206.095611>

2 Soler, J. J., Moreno, J., & Potti, J. (2003). Environmental, genetic and maternal components of
3 immunocompetence of nestling pied flycatchers from a cross-fostering study. *Evolutionary*
4 *Ecology Research*, 5(2), 259–272.

5 Stoffel, M. A., Nakagawa, S., & Schielzeth, H. (2017). rptR: Repeatability estimation and
6 variance decomposition by generalized linear mixed-effects models. *Methods in Ecology*
7 *and Evolution*. <https://doi.org/10.1111/2041-210X.12797>

8 Symonds, M. R. E., & Moussalli, A. (2011). A brief guide to model selection, multimodel
9 inference and model averaging in behavioural ecology using Akaike’s information criterion.
10 *Behavioral Ecology and Sociobiology*, 65(1), 13–21. [https://doi.org/10.1007/s00265-010-](https://doi.org/10.1007/s00265-010-1037-6)
11 [1037-6](https://doi.org/10.1007/s00265-010-1037-6)

12 van Oers, K., de Jong, G., van Noordwijk, A., Kempenaers, & Drent, P. (2005). Contribution of
13 genetics to the study of animal personalities: a review of case studies. *Behaviour*, 142(9),
14 1185–1206. <https://doi.org/10.1163/156853905774539364>

15 Van Overveld, T., Adriaensen, F., & Matthysen, E. (2011). Postfledging family space use in
16 great tits in relation to environmental and parental characteristics. *Behavioral Ecology*,
17 22(4), 899–907. <https://doi.org/10.1093/beheco/arr063>

18 Vindenes, Y., & Langangen, O. (2015). Individual heterogeneity in life histories and eco-
19 evolutionary dynamics. *Ecology Letters*, 18(5), 417–432. <https://doi.org/10.1111/ele.12421>

20 Waser, P. M., & Thomas Jones, W. (1989). Heritability of dispersal in banner-tailed kangaroo
21 rats, *Dipodomys spectabilis*. *Animal Behaviour*, 37, Part 6, 987–991.
22 [https://doi.org/https://doi.org/10.1016/0003-3472\(89\)90142-5](https://doi.org/https://doi.org/10.1016/0003-3472(89)90142-5)

23 Winney, I., Nakagawa, S., Hsu, Y. H., Burke, T., & Schroeder, J. (2015). Troubleshooting the
24 potential pitfalls of cross-fostering. *Methods in Ecology and Evolution*, 6(5), 584–592.
25 <https://doi.org/10.1111/2041-210X.12341>

26 Wood, P. B., Collopy, M. W., & Sekerak, C. M. (1998). Postfledging Nest Dependence Period
27 for Bald Eagles in Florida. *The Journal of Wildlife Management*, 62(1), 333–339.
28 <https://doi.org/10.2307/3802296>

1 Zuur, A. F., Ieno, E. N., Walker, N. J., Saveliev, A. A., Smith, G. M., & Park, W. (2009). Mixed
 2 effects modelling and extensions in ecology with R.

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6 **Table 1.** Model-averaged coefficients and RIV values for the post-fledging dependence period.

<i>DEPENDENT VARIABLE</i>	<i>EXPLANATORY VARIABLE</i>	<i>MODEL-AVERAGED COEFFICIENTS AND RELATIVE IMPORTANCE VALUES</i>			
		<i>β</i>	<i>SE</i>	<i>P</i>	<i>RIV</i>
<i>NET DISTANCE</i>	<i>Intercept</i>	4.86	0.45	<2e-16	
	<i>Age</i>	0.01	0.00	0.0119	1.00
	<i>SexM²</i>	0.25	0.42	0.5549	0.55
	<i>Treatment1¹</i>	-0.20	0.41	0.6209	0.47
	<i>Age:SexM</i>	-0.00	0.00	0.6389	0.27
	<i>Age:Treatment1</i>	0.00	0.00	0.6359	0.25
	<i>Treatment1:SexM</i>	0.02	0.11	0.8637	0.08
<i>TOTAL DISTANCE</i>	<i>Intercept</i>	6.56	3.32	< 2e-16	
	<i>Age</i>	6.31	1.99	0.00162	1.00
	<i>SexM²</i>	1.77	1.96	0.36912	0.78
	<i>Treatment1¹</i>	-4.79	2.08	0.81896	0.45
	<i>Age:SexM</i>	-9.30	1.38	0.94661	0.19
	<i>Age:Treatment1</i>	7.60	2.09	0.71705	0.19
	<i>Treatment1:SexM</i>	-2.66	6.71	0.96864	0.09
<i>SPEED</i>	<i>Intercept</i>	9.19	1.77	2e-07	
	<i>Age</i>	3.13	1.14	0.00625	0.98
	<i>SexM²</i>	1.27	5.74	0.82442	0.38
	<i>Treatment1¹</i>	-5.45	5.36	0.91907	0.37
	<i>Age:SexM</i>	-5.18	4.68	0.91194	0.10
	<i>Age:Treatment1</i>	7.07	4.72	0.88092	0.10
	<i>Treatment1:SexM</i>	7.28	1.90	0.96942	0.04
<i>TURNING ANGLE</i>	<i>Intercept</i>	0.09	0.13	0.488	
	<i>Age</i>	0.00	0.07	0.990	0.41
	<i>SexM²</i>	-0.12	0.16	0.432	0.61
	<i>Treatment1¹</i>	0.21	0.19	0.266	0.74
	<i>Age:SexM</i>	-0.02	0.08	0.773	0.13
	<i>Treatment1:SexM</i>	-0.01	0.11	0.950	0.13
	<i>Age:Treatment1</i>	0.01	0.06	0.837	0.11
	<i>Intercept</i>	5.43	3.38	<2e-16	
	<i>Age</i>	3.90	2.10	0.0634	0.91
	<i>SexM²</i>	7.24	1.11	0.5152	0.64

<i>NEST</i>	<i>Treatment1</i> ¹	6.12	5.42	0.9103	0.42
<i>DISTANCE</i>	<i>Age:SexM</i>	-3.36	9.19	0.7150	0.23
	<i>Age:Treatment1</i>	3.02	4.54	0.9471	0.11
	<i>Treatment1:SexM</i>	1.86	2.58	0.9428	0.08
<i>AREA</i>	<i>Intercept</i>	4.08	0.64	<2e-16	
	<i>SexM</i> ²	-0.04	0.13	0.785	0.09
	<i>Treatment1</i> ¹	-0.17	0.12	0.151	0.77
	<i>Treatment1:SexM</i>	0.00	0.03	0.930	0.01
<i>DISTANCE BETWEEN</i>	<i>Intercept</i>	7.66	1.28	<2e-16	
	<i>Age</i>	1.57	7.38	0.0329	0.95
	<i>SexM</i> ²	1.16	2.05	0.5725	0.58
	<i>Treatment1</i> ¹	1.10	1.79	0.9109	0.53
	<i>Age:SexM</i>	-3.26	1.53	0.8311	0.17
<i>OWLS</i>	<i>Age:Treatment1</i>	4.09	1.56	0.7936	0.17
	<i>Treatment1:SexM</i>	-2.24	9.31	0.8102	0.12

¹ = Treatment group 1 (fostered owls)

² = Males

1 **Table 2. ICC values of the random factors “owl” and “nest” for the post-fledging**
 2 **dependence period and dispersal.**

<i>DEPENDENT VARIABLE</i>	<i>RANDOM FACTORS</i>	<i>ICC VALUES PFDP</i>	<i>ICC VALUES DISPERSAL</i>
<i>NET</i>	<i>Owl</i>	0.00	
<i>DISTANCE</i>	<i>Nest</i>	0.35	0.90
<i>TOTAL</i>	<i>Owl</i>	0.00	
<i>DISTANCE</i>	<i>Nest</i>	0.73	0.58
<i>SPEED</i>	<i>Owl</i>	0.00	0.01
	<i>Nest</i>	0.35	0.00
<i>NEST</i>	<i>Owl</i>	0.00	0.36
<i>DISTANCE</i>	<i>Nest</i>	0.74	0.44
<i>AREA</i>	<i>Nest</i>	0.99	

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1 **Table 3.** Model-averaged coefficients and RIV values for early dispersal.

<i>DEPENDENT VARIABLE</i>	<i>EXPLANATORY VARIABLE</i>	<i>MODEL-AVERAGED COEFFICIENTS AND RELATIVE IMPORTANCE VALUES</i>			
		<i>β</i>	<i>SE</i>	<i>P</i>	<i>RIV</i>
<i>NET DISTANCE</i>	<i>Intercept</i>	7.90	3.56	<2e-16	
	<i>SexM²</i>	-3.16	1.75	0.869	0.06
	<i>Treatment1¹</i>	2.15	1.55	0.900	0.04
	<i>Treatment1:SexM</i>	-5.61	3.49	0.999	<0.01
<i>TOTAL DISTANCE</i>	<i>Intercept</i>	9.99	2.13	<2e-16	
	<i>Treatment1¹</i>	1.15	8.92	0.909	0.04
	<i>SexM²</i>	4.80	6.87	0.952	0.04
	<i>Treatment1:SexM</i>	-6.47	8.53	1.000	<0.01
<i>SPEED</i>	<i>Intercept</i>	8.54	1.06	<2e-16	
	<i>Age</i>	-1.63	4.48	0.000298	1.00
	<i>SexM²</i>	-2.63	1.42	0.065174	0.90
	<i>Age:SexM</i>	1.11	6.04	0.66097	0.86
	<i>Treatment1¹</i>	9.14	3.32	0.978140	0.37
	<i>Treatment1:SexM</i>	-8.29	1.21	0.945717	0.09
	<i>Age:Treatment1</i>	2.98	1.31	0.981901	0.09
<i>TURNING ANGLE</i>	<i>Intercept</i>	0.14	0.23	0.540	
	<i>Age</i>	0.05	0.25	0.834	0.84
	<i>SexM²</i>	0.21	0.26	0.934	0.59
	<i>Age:SexM</i>	0.26	0.36	0.462	0.43
	<i>Treatment1¹</i>	0.04	0.24	0.854	0.41
	<i>Treatment1:SexM</i>	0.05	0.25	0.826	0.09
	<i>Age:Treatment1</i>	0.00	0.08	0.965	0.09
<i>NEST DISTANCE</i>	<i>Intercept</i>	6.92	0.36	<2e-16	
	<i>Age</i>	0.01	0.00	<2e-16	1.00
	<i>SexM²</i>	1.27	0.41	0.00193	1.00
	<i>Age:SexM</i>	-0.01	0.00	6e-06	1.00
	<i>Treatment1¹</i>	0.05	0.27	0.83999	0.45
	<i>Treatment1:SexM</i>	-0.13	0.33	0.70568	0.19
	<i>Age:Treatment1</i>	0.00	0.00	0.90387	0.12
<i>DISTANCE BETWEEN OWLS</i>	<i>Intercept</i>	6.81	0.51	<2e-16	
	<i>Age</i>	0.01	0.00	5.4e-06	1.00
	<i>Treatment1¹</i>	0.85	0.57	0.140769	0.86
	<i>SexM²</i>	1.35	0.46	0.003573	1.00
	<i>Age:Treatment1</i>	-0.01	0.00	0.172951	0.76
	<i>Age:SexM</i>	-0.01	0.00	0.000313	0.99
	<i>Treatment1:SexM</i>	-0.11	0.26	0.678530	0.31

¹ = Treatment group 1 (fostered owls)

² = Males

1 **Figure legends**

2 **Figure 1. a. Movement characteristics of owls in 20-day periods during the post-fledging**
3 **dependence period. Means of males (dotted lines) and females (full lines) are presented.**
4 **(1.) Net distance. (2.) Total distance. (3.) Speed. (4.) Distance between each location and the**
5 **nest. (5.) Distance between owls.**

6 **b. Example of real paths followed by one individual at different ages during the post-**
7 **fledging dependence period. The black points represent the starting point of the path.**
8 **(1.) Path at 66 days. (2.) Path at 104 days. (3.) Path at 118 days.**

9 **Figure 2. a. Movement characteristics of owls in 20-day periods during dispersal. Means of**
10 **males (dotted lines) and females (full lines) are presented.**

11 **(1.) Speed. (2.) Distance between each location and the nest. (3.) Distance between owls.**

12 **b. Example of a real path followed by one individual during the whole dispersal phase**
13 **considered. The black point represents the starting point of the path.**

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15 **Supplemental files**

16 **Table S1.** Movement parameter values for the post fledging dependence period.

17 **Table S2.** Comparison of the competing models built to study variation in eagle owl movements
18 during the post-fledging dependence period. A summary of model-averaged coefficients and RIV
19 values is shown for all candidate models. Competitive models are ranked from the lowest (best
20 model) to the highest AICc value.

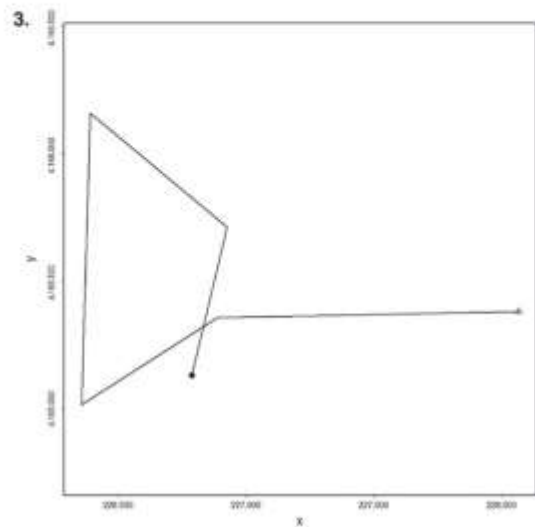
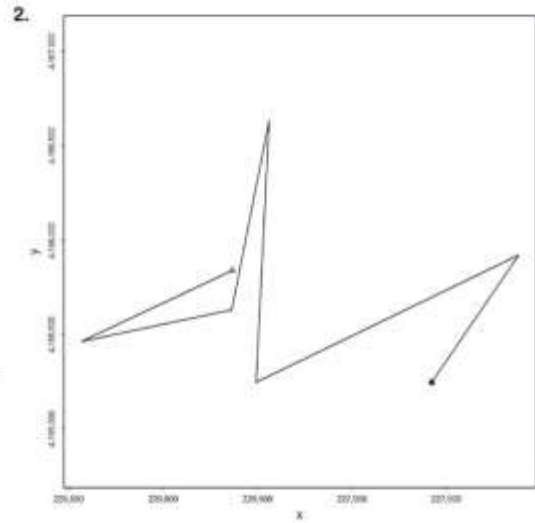
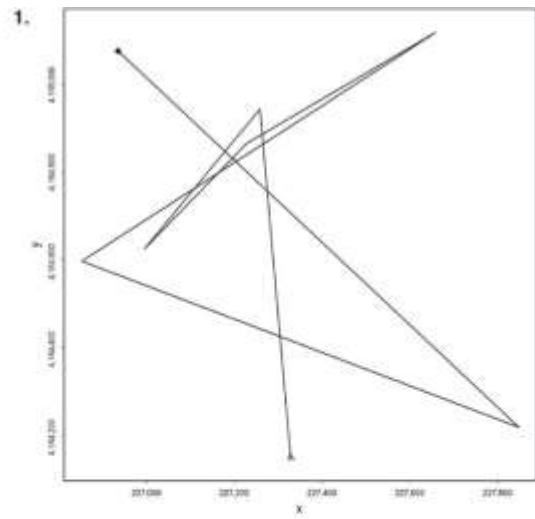
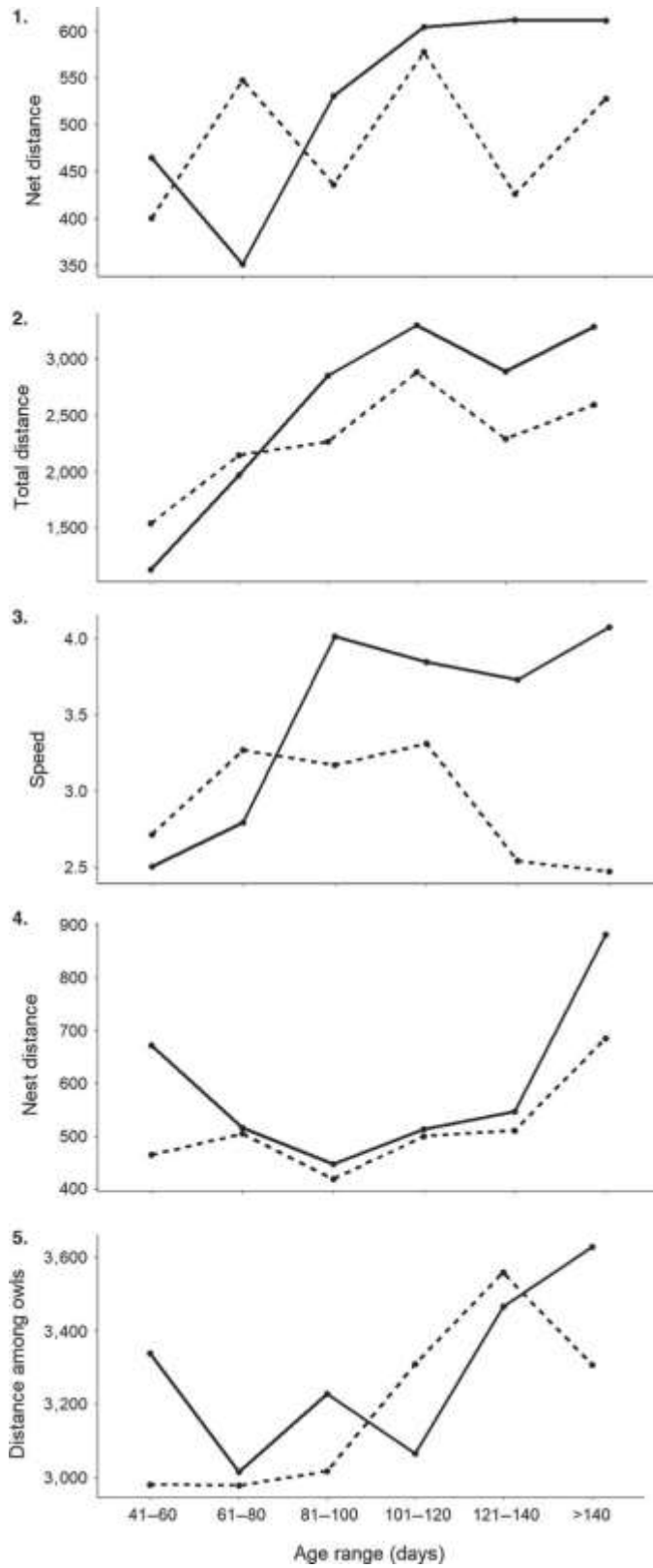
21 **Table S3.** Movement parameter values for early dispersal.

22 **Table S4.** Comparison of the competing models built to study variation in eagle owl movements
23 during early dispersal. A summary of model-averaged coefficients and RIV values is shown for
24 all candidate models. Competitive models are ranked from the lowest (best model) to the highest
25 AICc value.

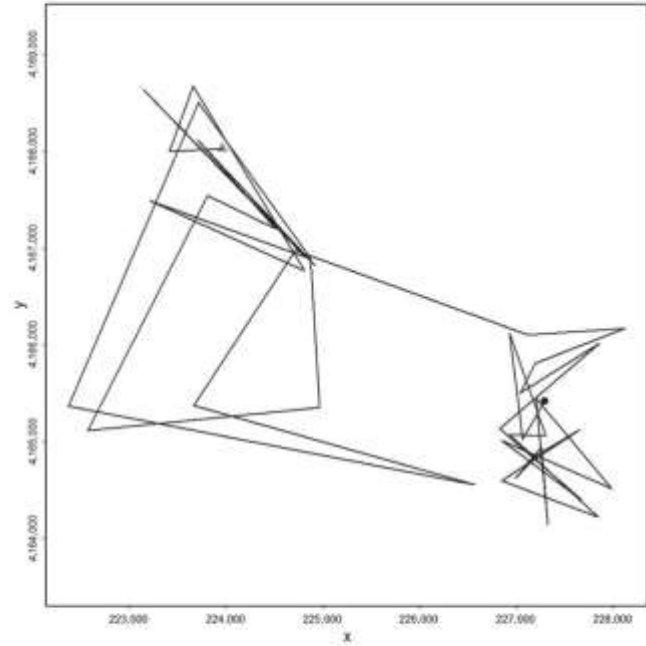
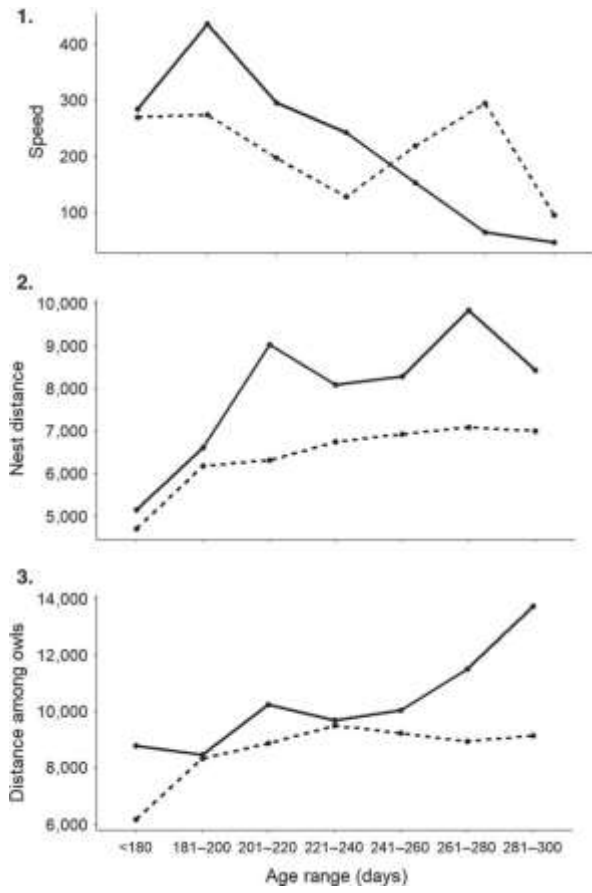
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1 Fig. 1.

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1 Fig. 2.



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