

1 **REPRODUCTIVE CONSEQUENCES OF MATE RETENTION AND DIVORCE**  
2 **IN A SHORT-LIVED MIGRATORY PASSERINE**

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16

17 **Abstract**

18 In socially monogamous birds, pair bond duration varies widely across species, from  
19 single-breeding associations to long-lasting, multi-year bonds. Studies on mate retention  
20 and divorce have predominantly focused on long-lived species, while research in short-  
21 lived and migratory species is limited. Consequently, the fitness consequences of divorce  
22 or remating in these species remain unclear. Here, we used a long term (1987–2023) and  
23 individual-based dataset to investigate the reproductive consequences of mate retention  
24 and divorce in pied flycatchers (*Ficedula hypoleuca*), a short-lived, long-distance  
25 migratory passerine. First, we described the overall patterns of divorce and remating  
26 within the population. Then, we analyzed whether reproductive success (laying date,  
27 clutch size and number of fledglings) differed between individuals that remate and those  
28 that divorce, while controlling for confounding factors, such as age or breeding dispersal.  
29 Specifically, we compared the reproductive consequences in the season before remating  
30 or divorce (year  $t$ ) and the subsequent season (year  $t+1$ ). We found that remating rates  
31 were low (3.52%) compared to divorces (26.8%), and both remained stable over the study  
32 period. In terms of reproductive performance, pair bond duration (remating vs. divorce)

33 did not explain variation in laying date or clutch size. However, divorce was associated  
34 with an increase in the number of fledglings in the year following separation for both  
35 males and females, while remated individuals produced more fledglings in their first  
36 breeding season. The fitness consequences of divorce/remating persisted after accounting  
37 for confounding factors, supporting a direct influence of pairing status on reproductive  
38 success. These findings highlight the complex dynamics of mate relationships in short-  
39 lived migratory birds and emphasize the need for further research into the adaptive  
40 potential of pair-bond stability.

41 **Key words:** pair status, mate retention, breeding success, mate choice, pied flycatcher

## 42 **Introduction**

43 While social monogamy is the predominant mating system in birds, the duration of pair  
44 bond varies widely across species, from short associations limited to the parental care of  
45 a single breeding attempt to pair bonds that can last for multiple years (Black 1996,  
46 Kvarnemo 2018). The diversity in pair bond duration reflects the interplay between  
47 species-specific life-history strategies (e.g., investment in reproduction and survival) and  
48 ecological pressures (e.g., availability of breeding sites, food resources, predation risk),  
49 highlighting the importance of pair-bond strength in avian life histories. Since pair bond  
50 duration may strongly influence reproductive performance (Lack 1968, Bennett & Owens  
51 2002), investigating its specific consequences on reproduction is essential to  
52 understanding its adaptive significance and evolutionary implications.

53 Retaining the same mate between successive breeding attempts may provide several  
54 advantages. Established pairs save the time and energy in finding a new partner and  
55 building a bond (Pampus *et al.* 2005, Gabriel & Black 2013, Culina *et al.* 2020, D'amelio  
56 *et al.* 2024). In addition, familiarity between partners can enhance their coordination in  
57 critical activities, as territory defense or offspring provisioning (Griggio & Hoi 2011,  
58 Gabriel & Black 2013, Sánchez-Macouzet *et al.* 2014, Culina *et al.* 2020, Botero-  
59 Delgadillo *et al.* 2024). Thus, remating may translate into increased reproductive success  
60 relative to newly formed pairs due to earlier breeding or higher fledgling success,  
61 although such benefits are not consistently supported (Rowley 1983, Choudhury 1995,  
62 Culina *et al.* 2015, Naves *et al.* 2007, Lv *et al.* 2016, Pitera *et al.* 2020, Dumas *et al.*  
63 2024). Alternatively, individuals may switch mates between consecutive breeding seasons  
64 despite the former partner remaining alive, leading to "divorce". Divorce may arise from  
65 breeding dispersal or external constraints, such as differences in settlement time after  
66 migration, making it difficult for previous partners to overlap in space and time during  
67 the period of pair formation. Alternatively, divorce may result from adaptive decisions  
68 made by one or both pair members, as for example, when mate switching allows at least  
69 one member to improve their territory or mate quality (reviewed in Choudhury 1995).

70 Empirical studies on mate retention and divorce have primarily focused on long-lived  
71 species, while studies focused on short-lived and migratory species are much scarcer (e.g.,  
72 Botero-Delgadillo *et al.* 2024). Yet, short- and long-lived species face distinct ecological  
73 pressures that may profoundly influence pair bond dynamics (Covas & Griesser 2007,  
74 Tarwater & Arcese 2017). Further, differences in reproductive performance between

75 remated and divorced pairs may not be a consequence of the breeding strategy itself, but  
76 rather confounded by factors such as age or individual quality (Lv *et al.* 2016). For  
77 example, newly formed pairs may consist of young or low-quality individuals. However,  
78 longitudinal analyses controlling for those confounding factors are scarce in the literature  
79 (reviewed in Culina *et al.* 2015, Botero-Delgadillo *et al.* 2024, Dumas *et al.* 2024). Lastly,  
80 as explained above, divorce may be a consequence of mismatches in spatial or temporal  
81 arrival patterns from migration, rather than from active behavioral decisions (Choudhury  
82 1995). Nevertheless, previous work has rarely differentiated these scenarios by  
83 considering all instances of divorce together (i.e., individuals overlapping or not  
84 spatiotemporally), which may bias our understanding of the adaptive value of mating  
85 strategies.

86 Here, we take advantage of a 36-year (1987–2023), individual-based dataset to  
87 investigate the reproductive consequences of mate retention and divorce in the pied  
88 flycatcher (*Ficedula hypoleuca*), a short-lived, long-distance migratory passerine. This  
89 study first explores the patterns of remating and divorce across the population.  
90 Subsequently, using a longitudinal framework, we assessed whether breeding success -  
91 measured as laying date, clutch size and number of fledglings the season before (year  $t$ )  
92 and after (year  $t+1$ ) remating or divorce- differed between males and females based on  
93 their pairing status. Based on the literature (Choudhury 1995, Cézilly *et al.* 2000, Culina  
94 *et al.* 2015, Lv *et al.* 2016, Pitera *et al.* 2021, Botero-Delgadillo *et al.* 2024), we predict  
95 that divorce may be favored in the short term over remating due to the high mortality risk  
96 between breeding seasons, which could reduce the benefits of pair bond stability.

## 97 **Materials and Methods**

### 98 **Ethical Note and Animal Care**

99 Adults were captured during incubation or using a conventional spring trap, which is a  
100 harmless design (Friedman *et al.* 2008). Standardized ringing protocols and  
101 measurements were carried out for all captured individuals. We have a long experience  
102 in this procedure, the time required for it was restricted to a minimum (mostly less than  
103 15 min), and it was carried out as cautiously and efficiently as possible.

104 All applicable international, national and/or institutional guidelines for the capture  
105 and ringing of animals were followed and the study was approved by the Spanish  
106 institutional authorities. Doñana Biological Station-CSIC and Autonomous Communities

107 of Madrid and Castilla-La Mancha provided capture and ringing licences. Field  
108 procedures were approved by the CSIC Ethical Committee for the following projects  
109 (refs. PAC05-006-2, CGL2006-07481/BOS, CGL2009-10652, CGL2011-29694,  
110 CGL2014-55969-P and PID2022-141763NA-I00) and, more recently, by the  
111 Autonomous Community of Madrid (Ref.: PROEX 068.6/24 to DC) guaranteeing that  
112 they comply with Spanish and European legislation on the protection of animals used for  
113 scientific purposes.

114

### 115 **Study species and population**

116 The pied flycatcher is a small-sized (11-13 gr), hole-nesting, insectivorous passerine. It  
117 is a trans-Saharan migrant that breeds in the temperate forests of Eurasia (Lundberg &  
118 Alatalo 1992), occupying a wide variety of woodlands, both deciduous and coniferous.  
119 During the breeding season, pied flycatchers show sexual dimorphism in plumage, with  
120 females displaying a greyish-brown plumage in contrast to the black and white pattern of  
121 males (Lundberg & Alatalo 1992, Potti et al. 2014). Males arrive from the wintering areas  
122 before females, establish a territory around a nesting site, and try to attract a female. This  
123 species shows high natal and breeding philopatry (Camacho *et al.* 2013, 2016), is single-  
124 brooded and predominantly monogamous, although some males become socially  
125 polygamous (Lundberg & Alatalo 1992, Canal *et al.* 2020).

126 Data were collected between 1987 and 2023 in a population of pied flycatchers  
127 breeding in nest boxes near La Hiruela (41°04' N, 3°27' W, central Spain). The population  
128 occupies two distinct habitat plots that differ markedly in vegetation composition and  
129 structure: a deciduous oak forest covering 9.3 ha (*Quercus pyrenaica*), and a mixed  
130 coniferous plantation (predominantly *Pinus sylvestris*) of 4.8 ha, 1.1 km apart. There are  
131 239 nest boxes, 156 in the oak forest and 83 in the coniferous plot, spaced approximately  
132  $20 \pm 9.2$  meters apart from one another. The study area and field methodologies have been  
133 described in detail elsewhere (Camacho *et al.* 2018, Potti *et al.* 2018).

134

### 135 **General field procedures**

136 Breeding season begins approximately the second week of April (before the first males  
137 arrive from migration) and ends in early July. During this period, we routinely check to

138 monitor occupancy by pied flycatchers and other species (see Potti *et al.* 2021). Once a  
139 nest was occupied, we conducted daily inspections to record laying date (first laid egg),  
140 clutch size (typically, 5-7 eggs), and the number of fledglings (at 13 days old). Adult birds  
141 were captured during incubation (females) or while feeding the nestlings (both sexes)  
142 around 8 days post-hatching. The exact age of many individuals (ca. 53%) was known  
143 because all nestlings are ringed before fledging (Potti & Montalvo 1991) and local  
144 recruitment rate is among the highest reported for the species (on average 14% Potti &  
145 Montalvo 1991, Canal *et al.* 2014). Unringed individuals first caught as breeders were  
146 aged (yearling vs. older) and sexed based on plumage characteristics (Karlsson *et al.*  
147 1986, Potti & Montalvo 1991) and marked for future identification. Morphological  
148 measurements were taken from all individuals following standardized protocols.

149

## 150 **Statistical Analyses**

151 We used a longitudinal approach to investigate the influence of mate retention or divorce  
152 on breeding performance of males and females. We built a dataset by selecting individuals  
153 that experienced either a remating or a divorce throughout their lifespan, which requires  
154 that individuals bred at least twice. Remating cannot occur if one of the pair members  
155 dies, leading to a widowed individual, or, even if both are alive, they do not coincide in  
156 space and time. Consequently, we categorized an individual as divorced only if the  
157 following two conditions were met: 1) the mate of the previous year was recorded in the  
158 following year(s), and 2) both former mates co-occurred in space and time. To account  
159 for the spatial and temporal factors in the chances of encountering former mates, we  
160 applied a spatio-temporal filter based on the mating patterns and spatial use of the species  
161 during the breeding season (Canal *et al.* 2012, Canal *et al.* 2020). According to these  
162 patterns, we only considered individuals as divorced when their previous mate bred  
163 within 10 days before or after them and within 100 m. After applying the filters above,  
164 and because each divorced pair member mates with a new individual, we generated two  
165 separate datasets: one focused on females, including 414 breeding attempts from 174  
166 individuals, and another focused on males, including 414 breeding attempts from 167  
167 individuals. The consequences of divorce or remate on laying date and clutch size were  
168 analyzed using the female dataset, while the analyses on the number of fledglings were  
169 run with both the male and female datasets.

170 We explored the role of age and breeding dispersal to explain the patterns of  
171 divorce and remating within the population. We first used a generalized linear mixed  
172 model (GLMM) with binomial distribution to test whether the frequency of remating and  
173 divorce events varied over the study period using year as predictor. Next, we investigated  
174 whether breeding dispersal between year  $t$  and year  $t+1$  (the year when the event was  
175 recorded) differed between divorced and remated individuals, under the assumption that  
176 longer dispersal distances should result in decreased fitness due to loss of site familiarity  
177 (Greenwood & Harvey 1982, Stamps 1995, Piper 2011). Linear mixed models (LMMs,  
178 Gaussian distribution) on breeding dispersal were fitted separately for males and females  
179 using the pair status of individuals as predictor (remated or divorce) and the individual  
180 identity as random factor.

181 To investigate the influence of mate retention or divorce on three breeding  
182 performance parameters —laying date (only females), clutch size (only females) and  
183 number of fledglings (both males and females)—we used linear and generalized mixed  
184 models. The models on laying date (LMM, Gaussian distribution) and clutch size  
185 (GLMM, Conway-Maxwell-Poisson distribution for under-dispersed counts) had the  
186 same predictors and random-effects structure. Predictors included in these models are the  
187 following: habitat type (two-level factor: oak vs. pine forest), pair status (two-level factor:  
188 remate vs. divorce), time (two-level factor: year  $t$  vs. year  $t+1$  relative to the remating or  
189 divorcing event) and the interaction between pair status and time. Time was included in  
190 the models to test the impact of remating or divorcing on breeding performance in the  
191 year preceding/following each type of event. It is well established that age and individual  
192 experience may influence the reproductive performance of individuals (Forslund & Pärt  
193 1995, Fowler 1995). Although individuals in our study population show variation in the  
194 age of first reproduction (1-3 years; Potti & Montalvo 1991), the individuals included in  
195 the longitudinal analyses (selected based on the criteria outlined above) showed a very  
196 high correlation between age and age of first reproduction ( $r= 0.9$ ,  $p<0.001$ ), preventing  
197 to disentangle the effects of age and breeding experience on breeding performance.  
198 Consequently, we included only individual age (continuous variable) as predictor in the  
199 models. Female identity and year were included as random effects in both models, except  
200 for the clutch size model, where including both year and identity caused convergence  
201 issues. We therefore excluded year from the clutch size model, after verifying that  
202 excluding it yielded similar results to excluding identity (data not shown). The models on

203 the number of fledglings (GLMM, Conway-Maxwell-Poisson distribution for under-  
204 dispersed counts) were fitted separately for males and females, and included as predictors  
205 pair status, time and their interaction as well as habitat, laying date, and the age of the  
206 individuals (female or male in their respective models). As random effects, we included  
207 female identity, male identity, year, and nest-box identity.

208         Sampling intensity was limited in the years 2001 and 2003, and therefore, these  
209 years were excluded from the analyses. During the long-term monitoring of this  
210 population, we have performed several nest manipulations (e.g., clutch/nestlings cross-  
211 fostering; Potti & Canal 2011, Camacho *et al.* 2016) that might affect breeding success.  
212 Thus, data from manipulated nests were also excluded from the analyses. Similarly, data  
213 from socially polygynous matings and from replacement clutches were excluded from  
214 analyses. Lastly, it is worth noting that some individuals re-paired after a year breeding  
215 apart, and some divorced individuals were not recorded in the following year (e.g., due  
216 to early breeding failure or failure to capture). Analyses including both remating and  
217 divorce events, even when not occurring in consecutive years, yielded qualitatively  
218 similar results to those based only on consecutive years (see Table 1 in Supplementary  
219 material).

220         All analyses were performed in R version 4.4.1 (R Core Team, 2022). To fit the  
221 GLMMs, we used the glmmTMB package (Brooks *et al.* 2017), version 1.1.10, while the  
222 lme4 package (Bates 2014), version 1.1.35.5, was used to fit the LMMs. During model  
223 diagnostics, we checked for residual normality and homoscedasticity with the package  
224 DHARMA (Hartig 2018), as well as for multicollinearity issues using the VIF function of  
225 car package (Fox & Weisberg 2019). Model diagnostics showed no deviations from  
226 model assumptions.

227

## 228 **Results**

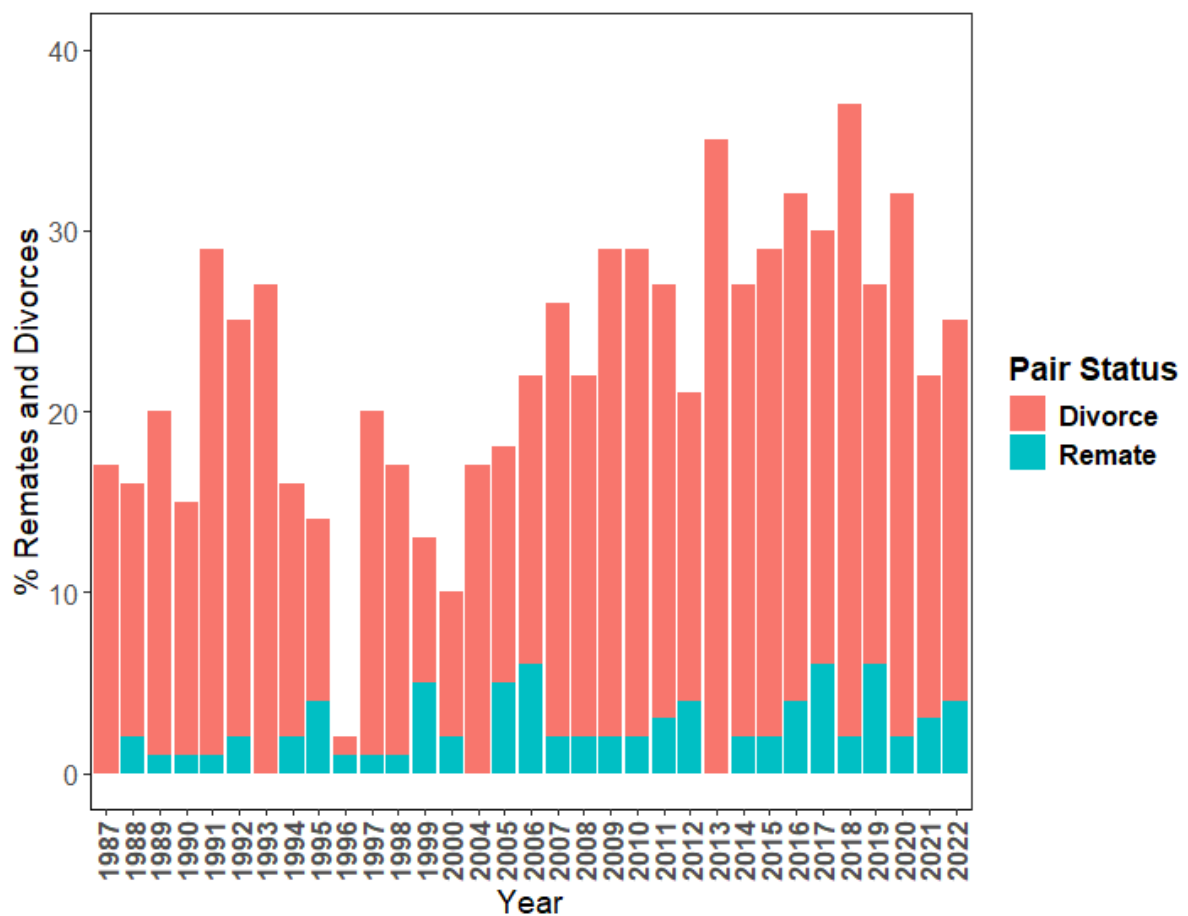
229 Overall, we observed 69 events of pair retention in consecutive years over the study  
230 period, representing 3.52 % of all breeding events (including divorces and widowing  
231 cases). Only 5 of the remating pairs continued breeding a third breeding season but not  
232 any longer. The number of divorces was 525, representing 26.8% of all breeding events.  
233 After applying the spatio-temporal filter to consider only those divorces occurring despite



234 previous mates co-occurring in space and time, the number of divorces was 92 cases  
235 (4.7% of all breeding events).

236 The frequency of remating events and divorces did not change over the study  
237 period (GLMM;  $p = 0.230$ ; Fig. 1). Breeding dispersal distance between year  $t$  and year  
238  $t+1$  was lower for remated than for divorced individuals (LMM;  $p < 0.001$  for females  
239 and males). Among divorced individuals, females showed greater breeding dispersal  
240 distances than males (Fig. 2).

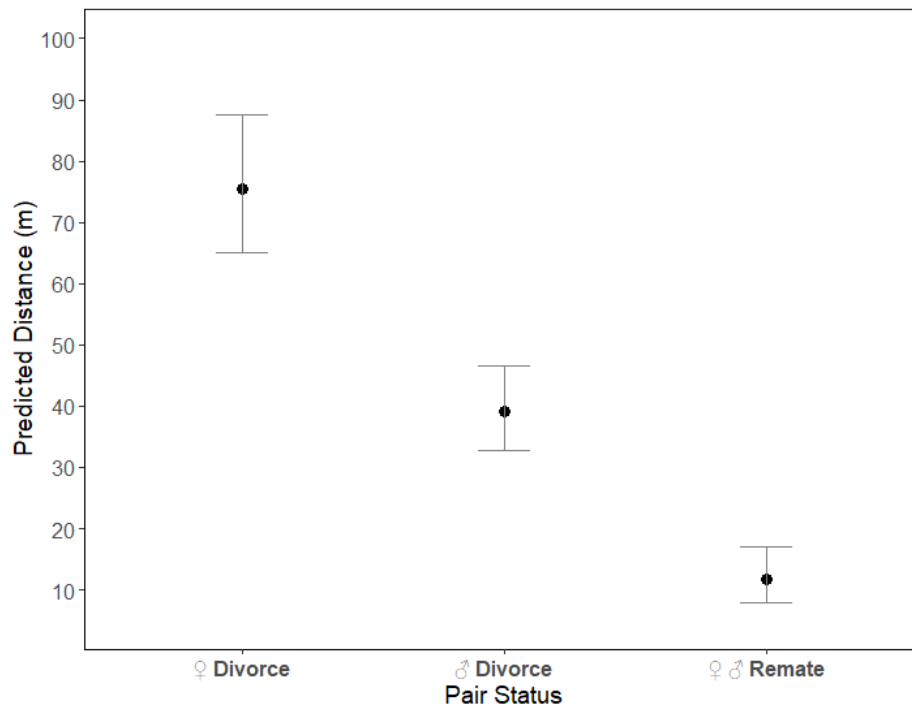
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242

243 **Figure 1.** Frequency of remates and divorces over the study period. Data from 2023 (last study year) are  
244 excluded, as assessing divorces and remating events required information from subsequent years. Data are  
245 unavailable for 1987 (the first study year), 2001-2003 (due to limited fieldwork), and 2004 (first study year  
246 after the 2001-2003 break). See main text for further details.

247



248

249 **Figure 2.** Breeding distance (mean  $\pm$  standard error) between year  $t$  and year  $t+1$  for divorced females  
 250 (left), divorced males (center) and remating pairs (right).

251 We found no significant effect of pair status (remating vs. divorcing) neither of its  
 252 interaction with time (year  $t$  vs. year  $t+1$ ) on laying date (Table 1a) nor on clutch size  
 253 (Table 1b). Instead, laying date was influenced by age, with individuals breeding earlier  
 254 as their age increased, while clutch size was negatively related to laying date and differed  
 255 between habitats (oak>pine). However, in the models on the number of fledglings, we  
 256 found an interactive effect between pair status and time (Table 1c-d) in both males and  
 257 females. This effect was driven by a positive impact of mate switching on the number of  
 258 fledglings in the year following divorce, whereas individuals retaining the same mate  
 259 tended to produce fewer fledglings (Fig. 1). The number of fledglings was also influenced  
 260 by laying date and habitat (oak >pine) in both sexes.

261 **Table 1.** Results of the models analyzing the influence of pair status (mate retention or divorce) on laying  
 262 date (LMM), clutch size and number of fledglings (both GLMM). The table presents estimated coefficients  
 263 ( $\beta$ ) and standard errors (SE) derived from the model. P-values were calculated via ANOVA type II due to  
 264 the presence of interaction terms.

a) Laying date			N = 322
Random effects	$\sigma^2$	SD	
Female identity	9.925	3.150	

Year	11.190	3.345		
<b>Fixed effects</b>	<b><math>\beta</math></b>	<b>SE</b>	<b>Wald <math>\chi^2</math></b>	<b>P value</b>
Intercept	21.479	1.142		
Pair status (remate)	-0.019	0.833	0.073	0.787
Habitat (pine)	0.976	0.710	1.894	0.169
Time ( $t+1$ )	-1.106	0.700	2.812	0.094
Age	-1.153	0.287	16.120	<b>0.001</b>
Pair status (remate) *Time ( $t+1$ )	0.370	0.988	0.140	0.708

**b) Clutch size** N = 322

<b>Random effects</b>	<b><math>\sigma^2</math></b>	<b>SD</b>		
Female identity	0.004	0.061		
<b>Fixed effects</b>	<b><math>\beta</math></b>	<b>SE</b>	<b>Wald <math>\chi^2</math></b>	<b>P value</b>
Intercept	1.810	0.031		
Laying date	-0.007	0.001	37.678	<b>0.001</b>
Pair status (remate)	0.032	0.018	1.806	0.179
Habitat (pine)	-0.041	0.014	8.253	<b>0.004</b>
Time ( $t+1$ )	0.027	0.015	1.732	0.188
Age	0.008	0.006	1.593	0.207
Pair status (remate) *Time ( $t+1$ )	-0.026	0.022	1.352	0.245

**c) Number of fledglings (females)** N = 317

<b>Random effects</b>	<b><math>\sigma^2</math></b>	<b>SD</b>		
Year	0.002	0.048		
Female identity	0.006	0.074		
Male identity	0.001	0.001		
Nestbox identity	0.004	0.066		
<b>Fixed effects</b>	<b><math>\beta</math></b>	<b>SE</b>	<b>Wald <math>\chi^2</math></b>	<b>P value</b>

Intercept	1.711	0.079		
Laying date	-0.008	0.003	8.992	<b>0.003</b>
Pair status (remate)	0.059	0.041	0.009	0.926
Habitat (pine)	-0.114	0.032	13.050	<b>0.001</b>
Time ( <i>t</i> +1)	0.074	0.038	0.619	0.431
Age female	-0.002	0.013	0.026	0.871
Age male	0.002	0.012	0.023	0.879
Pair status (remate) *Time ( <i>t</i> +1)	-0.119	0.054	4.812	<b>0.028</b>

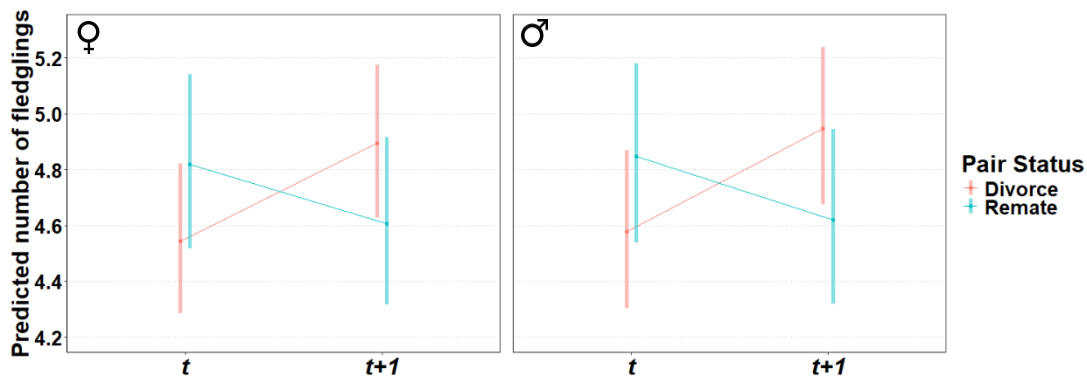
**d) Number of fledglings (males)**

N = 320

<b>Random effects</b>	$\sigma^2$	<b>SD</b>		
Year	0.005	0.069		
Female identity	0.001	0.001		
Male identity	0.001	0.023		
Nestbox identity	0.001	0.001		
<b>Fixed effects</b>	$\beta$	<b>SE</b>	<b>Wald <math>\chi^2</math></b>	<b>P value</b>
Intercept	1.684	0.078		
Laying date	-0.007	0.003	7.394	<b>0.007</b>
Pair status (remate)	0.057	0.042	0.063	0.802
Habitat (pine)	-0.120	0.029	16.711	<b>0.001</b>
Time ( <i>t</i> +1)	0.078	0.038	0.738	0.390
Age female	0.001	0.012	0.009	0.924
Age male	0.007	0.012	0.325	0.569
Pair status (remate) *Time ( <i>t</i> +1)	-0.126	0.057	4.832	<b>0.028</b>

265

266



267

268 **Figure 3.** Number of fledglings (mean ± standard error) produced by females (left panel) and males (right  
 269 panel) in relation to the pair status (remating or divorce) and time (year t vs t+1).

270

## 271 Discussion

272 Using a long-term and individual-based dataset spanning four decades, we analyzed the  
 273 reproductive consequences of remating and divorce in pied flycatchers (*Ficedula*  
 274 *hypoleuca*), a long-distance migratory and short-lived passerine, while controlling for  
 275 confounding factors such as habitat type, breeding dispersal or individual age. We showed  
 276 that pair status (remating vs. divorce) had no effect on laying date or clutch size but found  
 277 a positive impact of mate switching on fledgling numbers in the year following divorce  
 278 in both males and females.

279 Most research examining the consequences of pair retention and divorce has concentrated  
 280 on long-lived species, resulting in limited understanding of these processes in short-lived  
 281 and migratory species (Choudhury 1995, Culina *et al.* 2015, Botero-Delgadillo *et al.*  
 282 2024, but see Dumas *et al.* 2024). We found a positive influence of divorce on the number  
 283 of fledglings, which increased in the year following divorce compared to remated pairs  
 284 for both females and males. The observed improvement in reproductive success after  
 285 divorcing aligns with findings from a previous metaanalysis (Culina *et al.* 2015),  
 286 suggesting that the benefits of mate switching might outweigh the potential costs, such as  
 287 the energy and time needed to find a new mate or the risk of missing a breeding attempt.  
 288 Although comprehensively testing these costs are out of the scope of this study, some  
 289 characteristics of the study population, including high breeding synchrony and density  
 290 (i.e., several potential mates may be available at short distances; Canal *et al.* 2020,  
 291 Morales-Mata 2022), as well as a low probability of skipping breeding seasons (over 95%  
 292 individuals breed continuously until death once they enter the breeding pool), likely

293 minimize the cost associated with finding new mates. The positive effects of divorce on  
294 reproductive performance may stem from increased compatibility with the new mate or  
295 the acquisition of a breeding site or mate of higher quality (Montalvo & Potti 1992,  
296 Blondel *et al.* 2000, Cézilly *et al.* 2000, García-Navas & Sanz 2011). In relation to the  
297 latter idea, an interesting possibility is that the observed reproductive patterns related to  
298 breeding status may be (partially) driven by assortative mating. High-quality individuals,  
299 more likely to pair together, may have lower divorce rates due to limited opportunities to  
300 improve mate quality, whereas low-quality pairs may experience higher rates of mate  
301 switching, with one or both partners seeking improvement. This is supported by the fact  
302 that, in year  $t$ , reproductive success (number of fledglings) was higher for pairs that  
303 subsequently remained together compared to those that divorced in the following year.  
304 Overall, regardless of the underlying drivers, our results suggest that divorce may  
305 represent an adaptive strategy, potentially improving reproductive success after  
306 separation (Linden 1991, Dhondt & Adriaensen 1994, García-Navas & Sanz 2011, Culina  
307 *et al.* 2015). Future research evaluating the long-term fitness consequences of divorce or  
308 remating, such as survival prospects and offspring quality (e.g., offspring's lifetime  
309 reproductive success), are necessary steps to clarify the adaptive potential of these  
310 alternative strategies.

311 The potential reproductive benefits of pair-bond strength may be confounded by factors  
312 such as age, habitat quality or familiarity. Age is widely recognized as a crucial predictor  
313 of reproductive success, as long-term survival is itself an indicator of intrinsic individual  
314 quality, with older individuals typically possessing better territories and having superior  
315 foraging skills (Fowler 1995, Pärt 1995, Hatch & Westneat 2007). Further, early breeding,  
316 which is primarily age-dependent in migratory birds (e.g., Smith & Moore 2005, Stewart  
317 *et al.* 2002, Canal *et al.* 2020), is commonly associated with reproductive success in this  
318 and other temperate-region migratory birds (Newton 2008, Pitera *et al.* 2020). In line with  
319 these ideas, we found significant effects of age and/or laying date in most models,  
320 highlighting their importance in the reproductive success of individuals. Regarding  
321 habitat familiarity, divorced individuals dispersed farther between seasons ( $t$  vs  $t+1$ : on  
322 average of 76 m and 39 m in divorced females and males, respectively) compared to  
323 remating individuals (17 m), yet showed higher reproductive success in the year  
324 following divorce ( $t+1$ ). Although dispersal distances were relatively small and habitat  
325 conditions likely did not change markedly between consecutive years, breeding dispersal

326 itself entails additional energetic and time costs, such as locating and defending suitable  
327 nesting, foraging sites and repelling aggressions from prior residents. Thus, that fitness  
328 increased among divorced individuals despite the potential costs of dispersing longer  
329 distances point to an independent effect of pair status on the reproductive performance of  
330 individuals. It is also worth noting that divorce or, alternatively absence of remating, may  
331 simply result from asynchronous arrival at the breeding grounds or long-distance  
332 breeding dispersal, rather than being a reproductive strategy of at least one pair member.  
333 To address this confounding effect, and unlike most previous studies, we applied a  
334 spatiotemporal filter based on the species' behavior, to exclude cases where both pair  
335 members survived but did not coincide in space or time in the subsequent season, making  
336 remating virtually impossible (Gilsenan *et al.* 2017). Overall, the fact that fitness  
337 consequences of divorce/remating were detected after accounting for these confounding  
338 factors further support the direct influence of pairing status (divorced or remated) on  
339 breeding performance.

340 The apparent reproductive advantages associated with divorce compared to pairs  
341 remating raise the question of why remating persists, even at low frequency, in the  
342 population. In long-lived or non-migratory species, pair retention offers clear benefits,  
343 such as reduced mate-searching costs and increased cooperation (Bradley *et al.* 1990,  
344 Griggio & Hoi 2011, Gabriel & Black 2013, Sánchez- Macouzet *et al.* 2014, Culina *et al.*  
345 2020, D'amelio *et al.* 2024). However, in short-lived, long-distant migratory species such  
346 as the pied flycatcher, the probability that both pair members survive to the next breeding  
347 season may be low. Consequently, selection may not favor mate retention, as the time and  
348 energy costs of waiting for a mate that might not return are high (Choudhury 1995).  
349 However, the persistence of remating in the study population could be explained by  
350 benefits beyond the immediate reproductive parameters analyzed here (e.g., survival or  
351 offspring quality, rather than quantity), which could lead to weak selection pressure for  
352 divorcing over remating, thus allowing both strategies to coexist. Alternatively, given the  
353 low breeding dispersal distance observed in the population, particularly in remated pairs  
354 (17m with several pairs breeding in the same nest), remating could simply result from site  
355 fidelity or from stochastic processes, such as local encounter rates with the previous mate,  
356 rather than being an adaptive behavior (Montalvo & Potti 1992). Indeed, the relatively  
357 low and stable frequency of remating in the population over the study period, despite

358 these short breeding dispersal distances, could suggest some level of avoidance of former  
359 partners, though further research is needed to evaluate this possibility.

### 360 **Conclusions**

361 Using an individual-based dataset from a short-lived, long-distant migratory passerine,  
362 we found evidence that divorce improved reproductive success (number of fledglings) in  
363 the year following separation relative to individuals retaining the same pair between  
364 breeding seasons, which performed better in their initial breeding compared to divorced  
365 individuals. These findings underscore the complex dynamics of mate relationships in  
366 short-lived migratory birds and highlight key areas for investigation: i) Behavioural  
367 observations to investigate the mechanisms underlying divorce (e.g., whether it is  
368 initiated by a pair member or forced by a third party) and its reproductive consequences  
369 (e.g., acquisition of a better territory or mate); ii) experimental approaches to manipulate  
370 costs and benefits of each breeding strategy, and iii) studies of their long-term  
371 consequences (e.g., offspring performance) as well as iv) studies of the genetic bases of  
372 divorce and remating and its evolutionary potential are promising avenues of research.

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554

555 **Supplementary Material.**

556 **Table 1.** Results of the models analyzing the influence of pair status (mate retention or  
557 divorce) on laying date (LMM), clutch size and number of fledglings (both GLMM)  
558 including remating (n=81) and divorces (n=152) events that did not occur in consecutive  
559 years (e.g., because some individuals re-paired after one or more years breeding apart or  
560 because some divorced individuals were not recorded in the following year). The table  
561 presents estimated coefficients ( $\beta$ ) and standard errors (SE) derived from the model. P-  
562 values were calculated via ANOVA type II due to the presence of interaction terms.

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<b>a) Laying date</b>	N = 461
<b>Random effects</b>	<b><math>\sigma^2</math>      SD</b>

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Female identity	10.04	3.168		
Year	14.30	3.781		
<b>Fixed effects</b>	<b><math>\beta</math></b>	<b>SE</b>	<b>Wald <math>\chi^2</math></b>	<b>P value</b>
Intercept	20.510	1.029		
Pair status (remate)	-0.724	0.685	0.394	0.530
Habitat (pine)	1.969	0.528	11.169	<b>0.001</b>
Time ( <i>t</i> +1)	-1.398	0.589	6.486	<b>0.011</b>
Age	-0.864	0.230	14.126	<b>0.001</b>
Pair status (remate) *Time ( <i>t</i> +1)	0.7460	0.822	0.824	0.364

**b) Clutch size** N = 461

<b>Random effects</b>	<b><math>\sigma^2</math></b>	<b>SD</b>		
Female identity	0.002	0.047		
Year	0.001	0.014		
<b>Fixed effects</b>	<b><math>\beta</math></b>	<b>SE</b>	<b>Wald <math>\chi^2</math></b>	<b>P value</b>
Intercept	1.824	0.027		
Laying date	-0.007	0.001	47.768	<b>0.001</b>
Pair status (remate)	0.022	0.017	1.768	0.184
Habitat (pine)	-0.034	0.007	7.207	<b>0.007</b>
Time ( <i>t</i> +1)	0.015	0.014	0.981	0.322
Age	0.005	0.005	1.120	0.290
Pair status (remate) *Time ( <i>t</i> +1)	-0.010	0.022	0.209	0.647

**c) Number of fledglings (females)** N = 456

<b>Random effects</b>	<b><math>\sigma^2</math></b>	<b>SD</b>		
Year	0.002	0.049		
Female identity	0.013	0.113		
Male identity	0.001	0.001		
Nest-boxes	0.001	0.016		

<b>Fixed effects</b>	$\beta$	SE	Wald $\chi^2$	P value
Intercept	1.698	0.066		
Laying date	-0.008	0.002	14.578	<b>0.001</b>
Pair status (remate)	0.054	0.036	0.145	0.703
Habitat (pine)	-0.082	0.028	8.777	<b>0.003</b>
Time ( <i>t</i> +1)	0.061	0.029	1.716	0.190
Age female	-0.013	0.011	1.267	0.260
Age male	0.012	0.010	1.479	0.224
Pair status (remate) *Time ( <i>t</i> +1)	-0.085	0.046	3.382	0.066

**d) Number of fledglings (males)**

N = 461

<b>Random effects</b>	$\sigma^2$	SD		
Year	0.003	0.054		
Female identity	0.006	0.078		
Male identity	0.001	0.001		
Nest-boxes	0.001	0.001		
<b>Fixed effects</b>	$\beta$	SE	Wald $\chi^2$	P value
Intercept	1.606	0.068		
Laying date	-0.006	0.002	6.444	<b>0.011</b>
Pair status (remate)	0.056	0.037	0.207	0.649
Habitat (pine)	-0.066	0.026	6.385	<b>0.012</b>
Time ( <i>t</i> +1)	0.061	0.031	1.358	0.244
Age female	0.007	0.011	0.464	0.496
Age male	0.005	0.010	0.240	0.624
Pair status (remate) *Time ( <i>t</i> +1)	-0.089	0.051	2.991	0.083