

1 **Season-specific carry-over of early-life associations in a monogamous bird**
2 **species**

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25

26 **Abstract**

27 Social relationships can have important fitness consequences. Although there is increasing
28 evidence that social relationships carry over across contexts, few studies have investigated
29 whether relationships formed early in life are carried over to adulthood. For example,
30 juveniles of monogamous species go through a major life-history stage transition—pair
31 formation—during which the pair bond becomes a central unit of the social organization. At
32 present, it remains unclear if pair members retain their early-life relationships after pair
33 formation. We investigated whether same-sex associations formed early in life carry over into
34 adulthood and whether carry-over was dependent on season, in a monogamous species.
35 Moreover, we investigated the role of familiarity, genetic relatedness and aggression on the
36 perseverance of social associations. We studied the social structure before and after pair
37 formation in captive barnacle geese (*Branta leucopsis*), a highly social, long-lived,
38 monogamous species. We constructed association networks of groups of geese before pair
39 formation, during the subsequent breeding season, and in the following wintering season.
40 Next, we studied how these associations carried over during seasonal changes. We found that
41 early-life associations in females were lost during the breeding season, but resurfaced during
42 the subsequent wintering season. In males, the early-life associations persisted across both
43 seasons. Association persistence was not mediated by genetic relatedness or familiarity. The
44 high level of aggressiveness of males, but not females, in the breeding season suggests that
45 males may have played a key role in shaping both their own social environment and that of
46 their partners. We show that early-life social relationships can be maintained well into later
47 life. Such relationships can be sustained even if they are temporarily disrupted, for example
48 due to reproductive behaviour. Our findings therefore highlight that the early-life social
49 environment can have life-long consequences on individuals' social environment.

50

51 **Key-words:** aggression, early life, familiarity, genetic relatedness, monogamous, pair
52 formation, social relationships, social associations

53

54 **Introduction**

55 The adaptive nature of sociality is a long-standing topic in ecology and evolution (Alexander,
56 1974; Krause & Ruxton, 2002), particularly the potential costs and benefits of maintaining
57 stable social relationships. Repeated social interactions between the same individuals have,
58 for example, been linked to faster predator-evasion responses, increased foraging success
59 (Carter et al., 2009; Griffiths et al., 2004), and can, ultimately, affect fitness (Beletsky &
60 Orians, 1989; Cameron et al., 2009; Kohn, 2017; Silk, 2007; Silk et al., 2009, 2010).

61 There is increasing evidence showing that social relationships carry over across time,
62 place, and context. Shizuka et al. (2014) showed that golden-crowned sparrows (*Zonotrichia*
63 *atricapilla*) that flocked together in one winter flocked together in the subsequent winter more
64 often than expected based on the degree of home range overlap. Roosting associations in
65 Bechstein's bats (*Myotis bechsteinii*) and Natterer's bats (*Myotis nattereri*) were found to
66 remain stable across several years despite high fission-fusion dynamics (Kerth et al., 2011;
67 Zeus et al., 2018). Stanley et al. (2018) revealed that, despite seasonal fluctuations in
68 gregariousness and overall weak social associations, semi-feral ponies (*Equus caballus*)
69 maintained stable association preferences over three years. Finally, Firth and Sheldon (2016)
70 showed that great tits' (*Parus major*) winter social associations carried over into their
71 subsequent breeding season, as individuals bred nearer to those they were most associated
72 with during winter. Despite evidence that social associations between individuals can persist
73 across time and context, it is still largely unknown when these relationships are formed in the
74 first place.

75 Experiences in early life are known to regularly carry over to adulthood, influencing
76 survival and reproductive performance (Lindström, 1999). Early-life social conditions are
77 generally known to have long-lasting effects (Langenhof & Komdeur, 2018; Leris & Reader,
78 2016; Stanton & Mann, 2012; Szipl et al., 2019). Despite the likely significance of the early-

79 life period for the formation of social relationships, only a few studies have tracked the social
80 associations of the same individuals from their early life into adulthood (e.g., Carter et al.,
81 2013; Linklater & Cameron, 2009; Mitani, 2009), and almost exclusively in mammals (but
82 see Frigerio et al., 2001 for long-term associations between female siblings in geese). As they
83 mature, individuals of many taxa go through major life-history transitions, such as dispersal
84 (Blumstein et al., 2009; Linklater & Cameron, 2009) and pair formation (Kurvers et al.,
85 2013). These transitions will strongly impact the structure of an individual's social
86 environment, but perturbations in the social environment do not necessarily impact the
87 stability of already-established relationships (e.g., Kerth et al., 2011). It remains an open
88 question if a life-history transition such as pair formation will cause preferred associations
89 formed in early life to disappear or whether these may persist into adulthood.

90 Here, we examined the social structure of barnacle geese, a highly social species,
91 before and after pair formation, in order to study the persistence (or disappearance) of early-
92 life same-sex associations after a major life-history transition: pair formation. Barnacle geese
93 are long-lived monogamous birds, which generally find a partner at 2–3 years of age
94 (Choudhury & Black, 1994). Barnacle geese are very selective in choosing a mate; they
95 sample one to six potential mates in so-called “trial liaisons” before settling with a permanent
96 partner (Choudhury & Black, 1993; van der Jeugd & Blaakmeer, 2001). The pair bond is
97 extremely strong and pair members generally stay in close proximity to each other and remain
98 together until one of them dies (Black, 2001; Owen et al., 1988). Pairs with a longer pair bond
99 show higher reproductive success than pairs with a shorter pair bond (Black, 2001). Pair
100 formation is thus a crucial step in the life history of geese. Here we took advantage of
101 controlled experiments to study how pair formation impacted same-sex foraging associations
102 formed early in life, allowing us to circumvent problems with missing individuals in
103 naturalistic data. We followed a captive population of barnacle geese over several years,
104 quantifying the dyadic association strength between individual geese before and after pair

105 formation. To study mediating factors underlying the potential persistence of these dyadic
106 associations, we furthermore quantified the degree of familiarity and genetic relatedness in
107 the population, as well as individuals' aggressive interactions. We focused on the persistence
108 of same-sex associations because, in comparison to inter-sex associations, these relationships
109 are less well studied in monogamous social species, but may have important adaptive
110 benefits, for example stress-reduction (Sachser et al., 1998), especially in competitive
111 contexts such as foraging.

112 In the wild, barnacle geese, like many migratory waterfowl, show a high site fidelity
113 to their breeding, wintering and staging areas (Percival, 1991; Robertson & Cooke, 1999),
114 creating opportunities for long-term maintenance of social relationships established early in
115 life. Both sexes show natal philopatry, though females have substantially higher levels of
116 natal philopatry than males (van der Jeugd, 2001; van der Jeugd et al., 2002). The extent to
117 which individuals maintain non-sexual relationships across seasons is not well understood,
118 partly because of the challenges of following individuals across space and time. van der Jeugd
119 et al. (2002) showed that during breeding, female, but not male, barnacle geese nested closer
120 to siblings than expected by chance. This occurred not only when the female siblings bred in
121 the same island as their parents, but also when they nested on a different island. Moreover,
122 this relationship was only observed for siblings hatched in the same year. Similar sex-specific
123 patterns have been observed in terms of resting proximity in semi-feral greylag geese, *Anser*
124 *anser*, in winter (Frigerio et al., 2001). Given these observations and females being the more
125 philopatric sex, we predicted that females would show a higher likelihood of maintaining their
126 early-life relationships than males after mating.

127 We constructed association matrices from foraging observations collected during the
128 breeding and wintering seasons. Geese are generally more gregarious during the wintering
129 season (Black et al., 2014; Szípl et al., 2019), while being more aggressive and territorial
130 during the breeding season (Owen & Wells, 1979). We, therefore, predicted that associations

131 with early-life companions after pair formation would be stronger in the wintering than the
132 breeding season. Finally, based on the expected direct and indirect adaptive benefits of
133 associating with kin, for example by providing and receiving social support (Black & Owen,
134 1989; Raveling et al., 2000; Scheiber et al., 2009; Scheiber et al., 2005), we predicted that
135 genetic relatedness would positively impact the strength of the associations. Similarly, based
136 on the expected benefits of reduced aggression between familiar individuals (i.e. “dear enemy
137 effect”; van der Jeugd, 2001; Ydenberg et al., 1988), we predicted that long-term familiarity
138 would positively influence dyadic foraging association strength.

139

140

141 **Methods**

142

143 **Study subjects**

144 In late 2007 we obtained two mixed-sex groups of barnacle geese (see Fig. 1 for study
145 timeline, and Kurvers et al., 2013 for more details). The two groups consisted of 21 (13
146 female, 8 male) and 23 individuals (8 female, 15 male) respectively. The groups were housed
147 in separate outdoor aviaries (12 x 15 m) at the Netherlands Institute of Ecology (NIOO) in
148 Heteren, the Netherlands. Groups were visually, but not acoustically, isolated from each
149 other. The aviaries consisted of bare soil with a large pond (6 x 1 m) with running water for
150 drinking and bathing. Birds received ad libitum food consisting of a mixture of grains and
151 pellets occasionally supplemented with grass. Most individuals (40 of 44) were hatched in
152 2007 and were thus approximately 5 months old upon arrival. All birds were captive-hatched,
153 wing-clipped and (upon arrival) fitted with uniquely coded white leg rings for identification.
154 Birds from the two groups had different origins, implying that birds within (but not between)
155 groups could have a high genetic relatedness (see below). Birds within a group will

156 henceforth be referred to as “familiar” individuals, and birds between groups as “unfamiliar”.
157 Geese lived for approximately 1.5 years in these familiarity groups before the start of the
158 social network observations (see below).

159

160 **Genetic relatedness**

161 We determined the genetic relatedness between each pair of individuals using a high-
162 resolution 374 single nucleotide polymorphism (SNP) marker set, developed for the barnacle
163 goose (Jonker et al., 2012). We took a small blood sample (± 1 ml) from each individual from
164 the brachial vein and preserved it in ethanol. Entire genomic DNA was isolated using the
165 Gentra Systems Puregene DNA purification kit. Genotyping was conducted with Vera Code
166 assays on an Illumina BeadXpress (described in Kraus et al., 2011). We calculated the
167 pairwise genetic relatedness (r) using the program Coancestry (version 1, Wang, 2011). To
168 determine which relatedness estimator best fitted our data, we used the empirical SNP allele
169 frequencies of our population and simulated 500 dyads of geese of varying relatedness
170 coefficients. Meeting real conditions as closely as possible, we simulated mainly unrelated
171 dyads ($N = 430$) but also dyads of close familial relationships, namely 30 full sibs dyads ($r =$
172 0.5), 10 half sibs dyads ($r = 0.25$), and 30 first cousins dyads ($r = 0.125$). Based on this
173 simulation, we found that the maximum-likelihood estimator of Milligan (2003) performed
174 best and used it for the final estimation of r . This produced a strong correlation with expected
175 values of r ($r^2 = 0.9$; analysis of the simulated data set carried out with default settings).
176 Subsequently, all pairwise relatedness values of the experimental geese were obtained from
177 Coancestry with standard settings (see Kurvers et al., 2013 for more details).

178

179 **Observations prior to pair formation: early life**

180 After living for approximately 1.5 years in their respective familiarity groups—and prior to
181 any pair formation—we separated all geese into two single-sex groups (June 2009; Fig. 1) to

182 study the factors shaping intra-sexual social association preferences while avoiding additional
183 ‘noise’ caused by inter-sexual trial-liaisons (van der Jeugd & Blaakmeer, 2001). Moreover, by
184 separating individuals in single-sex groups, we could control the timing of pair formation and
185 keep this comparable and tractable for all subjects. Geese were sexed by visual inspection of
186 sexual organs in the cloaca. The sex of one individual could not be reliably determined at this
187 time. This individual was not included in the early-life observations, but joined the flock after
188 these observations finished. This individual was excluded from all analyses. Association
189 observations were conducted in the home aviaries during foraging. These foraging
190 associations were studied on five grass patches (40 x 20 cm, 1.5m apart) which were replaced
191 twice a day to avoid depletion. Other food sources were removed during the observations.

192 Each single-sex group was observed (9 a.m.–1 p.m.) for 15 days (females: 22 June–12
193 July 2009; males: 13–30 July 2009). The presence of all individuals on the patches was
194 recorded every 4 minutes. This interval was longer than the mixing time among individuals
195 (i.e., the time individuals need to exchange who they are association with), in order to ensure
196 independence of observations (Croft et al., 2008). Associations at feeding patches was rarely
197 the same in consecutive records (females: 5.9%; males: 7.9%). Observations were
198 occasionally interrupted for 10 minutes in the event of an external disturbance. Since patch
199 size (40 x 20 cm) and group size (mean females = 1.9, range 1–5; mean males = 2.0, range 1–
200 5; Fig. A1a, b) were small, we assumed that animals grazing on the same patch during a
201 sampling period were associating (a.k.a. gambit of the group, Franks et al., 2010; Whitehead,
202 2008).

203 The results of the foraging associations before pair formation are reported in Kurvers
204 et al. (2013). In brief, in both sexes, familiarity and genetic relatedness predicted association
205 strength, whereas boldness and dominance did not. Therefore, we focus here on the role of
206 familiarity and genetic relatedness in governing the stability of long-term associations.

207

208 **Observations after pair formation: breeding and wintering**

209 Geese remained in the single-sex groups for six months and were then placed together in one
210 group (January 2010; Fig. 1). By then, most geese were 2.5 years of age, approximating the
211 average age of final pair formation in barnacle geese (Choudhury & Black, 1994; van der
212 Jeugd & Blaakmeer, 2001). Pair formation started rapidly, and most geese quickly formed a
213 stable pair bond. Before starting the first post-pair formation association observations, 37 (of
214 44) geese had formed stable pair bonds. In total, geese formed 12 pairs (11 male-female pairs,
215 and one male-male pair), four triplets (three geese continuously moving together as one unit
216 without any aggression; two triplets consisted of two females and one male, and two triplets
217 of one female and two males) and seven geese (3 males, 4 females) remained unpaired. The
218 occurrence of triplets has also been observed in the wild, with a third party joining a pair
219 between 10 months and 4 years, however incidence in the field was lower (Black et al., 1996).
220 As reported in Kurvers et al. (2013) genetic relatedness did not play a role in mate choice,
221 whereas geese actively selected against familiarity in selecting a mate. After the geese formed
222 pair relationships, and during their first breeding season, we again conducted observations of
223 foraging associations. We placed 10 grass patches (40 x 40 cm) in the aviary. Observations
224 were conducted for five weeks (31 May–2 July 2010, 25 observation days) in two 2-hour
225 blocks per day (8 a.m.–1 p.m.) following the same observation protocol described above.
226 During the breeding season, many of the paired individuals started building nests and laying
227 eggs. We regularly checked all nests and removed any eggs to avoid the hatching of goslings.

228 To study the stability of associations across seasons, we repeated the observations of
229 foraging associations six months later, during the wintering season, following the same
230 protocol (17 December 2010–4 February 2011, 13 observation days). Comparing the pair
231 status of all geese between the breeding and wintering seasons, we observed that all pair
232 relationships but one remained the same—the one pair relationship that changed was the
233 male-male pair. This reflects the strong and long lasting pair bonds in barnacle geese, which

234 generally stay together until one of the pair members dies (Black, 2001). One individual died
235 in between the breeding and wintering observations. To facilitate comparison across seasons,
236 we removed this individual from all analyses. The final sample size was thus 42 individuals
237 (20 females, 22 males).

238

239 **Agonistic interactions**

240 During all three observation periods (early life, breeding, wintering), we collected data on
241 agonistic interactions to study the role of aggression on the stability of early-life associations.
242 In between scoring the presence of individuals on the patches, we scored the winner and loser
243 of agonistic interactions, defined as a direct confrontation between two geese and ranging
244 from threats with lowered head and neck to active chases with flapping wings. In the early-
245 life observations, we identified a total of 1,429 interactions in the female group and 2,619
246 interactions in the male group. We then identified 3,411 and 786 interactions during the
247 subsequent breeding and wintering periods, respectively.

248

249 **Statistical analyses**

250

251 *Stability of early-life connections*

252 We used social network analysis to investigate carry-over effects of dyadic associations
253 between seasons. Within the networks, nodes represent individuals and are connected by
254 edges that represent associations. The edge weight varied with dyadic association strength.
255 From the observation data, we thus generated undirected weighted networks (i.e., networks
256 based on associations without initiators or receivers) for each of the three periods (early life,
257 breeding season, and wintering season). Edge weights in the networks were calculated using
258 the simple ratio index (SRI) as an association measure using the ‘asnipe’ package (Farine,
259 2013) in R (v. 3.4.4). The SRI indicates the probability of observing two individuals in

260 association with each other given that one was observed. Values range from 0 (two
261 individuals were never observed together) to 1 (two individuals were always observed
262 together). The SRI is considered an effective measure of dyadic association strength provided
263 there are no large sampling biases (Farine & Whitehead, 2015; Ginsberg & Young, 1992;
264 Hoppitt & Farine, 2018). Since all observations were performed within aviaries in which all
265 individuals feeding on all patches could be easily observed, we did not expect a strong
266 sampling bias.

267 We first analysed whether early-life associations (within the same sex) were
268 maintained after pair formation, i.e. during the subsequent breeding and wintering season.
269 Within each sex, we performed Mantel tests using the built-in node permutation test from the
270 ‘vegan’ library Mantel test function—running 50,000 permutations and using the spearman
271 correlation (Oksanen et al., 2019)—to compare dyads’ association strength between (i) early
272 life and breeding season, and (ii) early life and wintering season. We started with a separate
273 matrix for each sex for the early-life associations, one matrix (containing both sexes) for the
274 breeding season, and one matrix for the wintering season. To compare the matrix correlation
275 for females (/males), we therefore first subset all females (/males) from the breeding and
276 wintering matrix, constructing single-sex matrices. To confirm that our results were not
277 mainly driven by the associations of the unpaired individuals (as we were primarily interested
278 in the possible continuation of associations after pair formation), we repeated these analyses
279 after further excluding associations between the unpaired individuals. To investigate if our
280 results were driven by a few strong associations, we also repeated this analysis while
281 excluding all associations with an SRI value higher than 0.1.

282

283 *Genetic relatedness and familiarity*

284 To investigate which factors might underlie a possible continuation of associations after pair
285 formation, we studied the effect of genetic relatedness and familiarity on dyadic association

286 strength during the breeding and wintering season. We tested the effect of familiarity and
287 relatedness using multiple regression quadratic assignment procedures (MRQAP; Krackhardt,
288 1988) using the `mrqap.dsp` function in the R package `asnipe` (Farine, 2013). With this
289 procedure, the association matrix was first regressed against two matrices, one with data on
290 relatedness and one on familiarity. We used a version of the function allowing us to combine
291 the `mrqap.dsp` model with pre-network permutations (Farine, 2017). Specifically, we
292 generated 1,000 random networks by re-organising the observations of individuals in the
293 original group-by-individual matrices following the method first described by Bejder et al.
294 (1998). We then compared the resulting distribution of coefficient values from these permuted
295 networks to the coefficient value generated from the original observation data to obtain P -
296 values. Further, because our random distributions were not centred on 0 (see Farine, 2017),
297 we rescaled measures to an effect size by taking the difference between the observed
298 coefficient values and the mean of the corresponding distribution of coefficient values based
299 on the permuted networks. We performed the MRQAP separately for each season and sex.

300

301 *Patch visits and aggression*

302 As we found sex differences in the continuation of early-life associations (see below), we
303 investigated the role of patch visitation rates and aggression as potential underlying
304 mechanisms. For each individual, for each period, we calculated the mean number of patch
305 visits per hour. We then used a Mann-Whitney U test to determine whether the sexes differed
306 in their patch visitation rates in each of the three seasons. Similarly, for each individual and
307 for each season, we determined the mean number of aggressive interactions initiated per hour,
308 and tested whether the sexes differed in their likelihood to display aggression in each of the
309 three seasons.

310

311 **Ethical permission**

312 The animal ethical committee of both the Royal Netherlands Academy of Arts and Sciences
313 (KNAW) and the Wageningen University approved all experiments [protocol numbers:
314 2010008.b (blood sampling)]. Geese were obtained from a waterfowl breeding farm (Kooy
315 and Sons, 't Zand, the Netherlands) and returned there after the experiments had finished.

316

317

318 **Results**

319

320 **Sex- and season-dependent effects of persistence of early-life associations**

321 We quantified the foraging association networks for each of the three observation periods
322 (i.e., early life, breeding season, and wintering season, see Fig. 2), and compared the strength
323 of the dyadic associations that were formed early in life to those in subsequent seasons. We
324 found that a female's dyadic association strengths from early life (i.e. prior to pair formation)
325 were not significantly correlated with its dyadic association strengths to the same pool of
326 individuals (i.e. other females) in the subsequent breeding season ($r = 0.04$; $P = 0.32$; Fig.
327 3a). A female's early-life dyadic association strengths were, however, significantly correlated
328 with its dyadic association strengths in the subsequent wintering season ($r = 0.18$; $P = 0.019$;
329 Fig. 3b). A male's early-life dyadic association strengths were significantly correlated with its
330 dyadic association strengths to other males in both the subsequent breeding ($r = 0.26$; $P =$
331 0.002 ; Fig. 3c) and the following wintering season ($r = 0.31$; $P = 0.002$; Fig. 3d).

332 When excluding the associations among individuals that remained unpaired (7 of 42;
333 but maintaining the associations between paired and unpaired individuals), we obtained
334 similar results (females: early life vs. breeding: $r = 0.00$, $P = 0.49$; early life vs. wintering: r
335 $= 0.19$, $P = 0.018$; males: early life vs. breeding: $r = 0.26$, $P = 0.003$; early life vs. wintering:
336 $r = 0.31$, $P = 0.003$).

337 When excluding all dyadic associations with an SRI value above 0.1 (which excluded
338 38 associations for females (7% of all associations) and 26 for males (5%)), we, again,
339 obtained similar results (females: early life vs. breeding: $r = 0.00$, $P = 0.52$; early life vs.
340 wintering: $r = 0.18$, $P = 0.022$; males: early life vs. breeding: $r = 0.23$, $P = 0.004$; early life
341 vs. wintering: $r = 0.34$, $P = 0.002$).

342

343 **Genetic relatedness and familiarity do not drive associations in females or males**

344 In females, there was no effect of genetic relatedness or familiarity on dyadic association
345 strength in the breeding season (effect sizes: genetic relatedness: -0.00006; familiarity:
346 0.0001) or in the wintering season (genetic relatedness: 0.0016; familiarity: 0.0001).
347 Likewise, for males, we found no effect of genetic relatedness or familiarity on dyadic
348 association strength in the breeding season (genetic relatedness: -0.00035; familiarity: -
349 0.00006) or in the wintering season (genetic relatedness: 0.0002; familiarity: -0.00026).
350 Thus, although familiarity and genetic relatedness positively impacted single-sex associations
351 in females and males in early life (see Kurvers et al., 2013 and see Fig. A2 and Fig. A3), these
352 factors did not drive same-sex associations after pair formation (all $P > 0.1$).

353

354 **Elevated aggression in males and during the breeding season**

355 We investigated two mechanisms that might drive the disappearance of social associations in
356 females—but not males—during the breeding season. First, females may simply visit the food
357 patches less than males during the breeding season, which would lower their opportunities for
358 maintaining relationships. Though females visited patches less than males in the single-sex
359 groups before pair-formation ($W = 50$, $P < 0.001$; Fig. 4a), females visited the patches equally
360 often as males in the breeding and wintering season (both $P > 0.35$; Fig. 4b, c), ruling out this
361 explanation.

362 A second mechanism might be that males, being generally the more dominant member
363 of a pair, play a stronger role in determining the association members of a pair than do
364 females, especially in the breeding season. Before pair formation, males showed a higher
365 level of aggression than females in the single-sex groups ($W = 142$, $P = 0.049$; Fig. 4d).
366 Likewise, males showed a substantially higher level of aggression than females during the
367 breeding ($W = 8$, $P < 0.001$; Fig. 4e) and wintering seasons ($W = 39$, $P < 0.001$; Fig. 4f).
368 Males displayed equal levels of aggression towards males and females in the breeding season
369 (Wilcoxon signed-rank test: $V = 76$, $P = 0.10$, Fig. 4e) but slightly higher levels of aggression
370 towards males than females in the wintering season ($V = 50$, $P = 0.02$; Fig. 4f). As expected,
371 male geese displayed higher levels of aggression in the breeding than in the wintering season
372 (Wilcoxon signed-rank test, males only: $V = 240$, $P < 0.001$; Fig. 4e, f). In accordance, the
373 mean group size at patches was higher during the wintering than during the breeding season
374 (mean group size wintering: 2.4; breeding: 2.0; Fig. A1c, d). Moreover, paired individuals
375 were more tolerant to the presence of other individuals (i.e., not belonging to the pair) at a
376 patch during wintering season. In the breeding season, in 80% of cases when pair members
377 were observed together on a patch, there were no other individuals present. In the wintering
378 season, this percentage dropped to 60% (mean number of non-pair individuals at a patch with
379 a pair during breeding: 0.26, during wintering: 0.76; Fig. A4c, d). This is also apparent in the
380 network graphs showing more edges during the wintering season (Fig. 2c, d).

381

382

383 **Discussion**

384 Maintaining stable social associations across time and contexts can have adaptive benefits
385 (Kohn, 2017). Yet the importance of the early-life period for the formation of such long-term
386 stable relationships has so far received little attention. Here we show that early-life same-sex

387 foraging associations can persist after a major life-history transition—pair formation—in a
388 monogamous and long-lived bird species. However, which associations were maintained
389 depended on sex and season. Early-life associations in females were lost during the breeding
390 season, but resurfaced again during the subsequent wintering season. In males, the early-life
391 associations persisted across both seasons. We found no evidence of genetic relatedness or
392 familiarity on association persistence. Elevated male aggression likely influenced the limited
393 number of contacts outside of the pair bond during the breeding season—and thereby the
394 extent to which early-life associations could be maintained, especially by females during the
395 breeding season. Our findings extend the understanding of how social relationships develop
396 and are maintained over different life-history phases and how their importance to individuals
397 may vary with season.

398 Across taxa, females are well-known for maintaining long-term social relationships
399 (Cameron et al., 2009; Carter et al., 2013; Ilany et al., 2015; Silk et al., 2003), which is
400 commonly attributed to their reproductive strategies. In our study, males, not females,
401 maintained their associations throughout both the breeding and wintering season. Benefits of
402 social relationships in males have been observed in other species: male red-winged blackbirds
403 (*Agelaius phoeniceus*) that bred close to familiar neighbours fledged more offspring (Beletsky
404 & Orians, 1989) and male Assamese macaques (*Macaca assamensis*) with stronger social
405 associations sired more offspring (Schulke et al., 2010). Males can thus clearly form, and
406 benefit from, long-lasting social relationships.

407 Social relationships in males often take the form of coalitions or alliances (Connor et
408 al., 2017; Gilby et al., 2013; Schulke et al., 2010), benefiting individuals by providing
409 cooperation partners in agonistic interactions with conspecifics. But reduced aggression, for
410 instance, via dear-enemy effects (Temeles, 1994), can likewise be an important benefit of
411 maintaining long-term social relationships, especially in territorial animals (Chuang et al.,
412 2017; Jaeger, 1981; Siracusa et al., 2019). Additionally, repeated association with certain

413 individuals can influence vigilance behaviour, with individuals showing reduced vigilance in
414 the proximity of well-known conspecifics (Carter et al., 2009; Kutsukake, 2006). In a
415 monogamous prey species such as barnacle geese, in which the male spends much of its time
416 during the breeding season on vigilance at the expense of foraging (Forslund, 1993), such
417 benefits can be especially substantial. The expected benefits of maintaining long-term social
418 relationships vary with ecological and social conditions (Connor et al., 2017; Kappeler et al.,
419 2013; Maher & Burger, 2011), and studying the social structure of both sexes in taxa with
420 distinct space use (e.g., natal philopatry), life history (e.g., long life-span), social organization
421 (e.g., fission-fusion), and mating system (e.g., monogamy) characteristics, offers us greater
422 insight into the drivers and constraints of maintaining long-term stable associations in animal
423 societies.

424 The apparent lack of social association persistence for females in the breeding season
425 is surprising, but adds to our understanding of social flexibility by showing that the
426 persistence of social relationships not only varies between species and individuals (Kappeler
427 et al., 2013) but can also have sex-specific effects across seasons. Female chacma baboons
428 (*Papio hamadryas ursinus*) were similarly found to vary in their social preferences depending
429 on ecological context (Henzi et al., 2009): social preferences were more pronounced when
430 food resources were scarce, but females acted more “gregariously” (i.e., without social
431 preferences) when resources were plentiful. For geese, winter is a time when resources
432 become scarce; barnacle geese, however, are highly gregarious in winter (Black et al., 2014),
433 possibly to access social information on foraging opportunities (Drent & Swierstra, 1977;
434 Kurvers et al., 2009; Kurvers et al., 2010) (but see Kurvers et al., 2014). Alternatively, the
435 fluctuations in female social preferences may have been the result of the male partner
436 unselectively excluding his partner’s—but not his own—early-life companions during the
437 breeding season. The high level of aggression displayed by the males during the breeding
438 season compared to the wintering season supports this hypothesis.

439 The absence of continued association with early-life companions in females in the
440 breeding season appears to be in contrast to previous findings. An earlier study on wild
441 barnacle geese found that female, but not male, geese exhibited social preferences in terms of
442 nesting proximity, with females nesting closer to familiar females (van der Jeugd et al., 2002).
443 It is possible that nest choice offers female geese an alternative way to maintain early-life
444 social associations, circumventing the potentially controlling influence of their partner during
445 foraging. Possibly, breeding in proximity to familiar and/or related conspecifics provides
446 context-specific benefits that foraging together does not. Intraspecific brood parasitism and
447 adoption is common in waterfowl (Anderholm et al., 2009a; Andersson et al., 2019;
448 Choudhury et al., 1993; Forslund & Larsson, 1995) and breeding close to related individuals
449 may decrease the costs, through inclusive fitness benefits, of having to care for additional
450 offspring. Moreover, breeding in proximity could enable siblings to defend each other's nests
451 against unrelated brood parasites. Female siblings may also actively or passively support each
452 other in the acquisition of high-quality nest locations. Indeed, in geese, social support from
453 family members is known to give individuals an advantage in competitive interactions,
454 starting from an early age (Black & Owen, 1989; Raveling et al., 2000; Scheiber et al., 2009;
455 Scheiber et al., 2005). Lastly, neighbouring barnacle geese pairs are known to defend their
456 nests together (Black & Owen, 1995) and nesting closely to familiars may facilitate
457 cooperative nest defence (Grabowska-Zhang et al., 2012; Olendorf et al., 2004). Nest
458 predation is a major threat for geese (Drent & Prop, 2008) that exposes the females in
459 particular to considerable predation risk (Samelius & Alisauskas, 2006).

460 Our captive study design had several important limitations compared to natural
461 settings. First, the group size under study was relatively small compared to natural groups. In
462 natural groups, the number of genetically related and/or familiar individuals may be
463 substantially higher, allowing geese more opportunities to associate with these types of
464 individuals. Second, the space available to our subjects was reduced as compared to natural

465 conditions. This may have led to higher levels of aggression, especially among males, and/or,
466 more and stronger associations as compared to more natural settings. Third, we removed the
467 eggs of breeding females to avoid the undesired hatching of more experimental animals. This
468 may have caused heightened aggression, and may have had repercussions for social
469 relationships. The pair bonds did, however, almost all remain intact till the next wintering
470 season, suggesting that egg removal did not cause major disruption of pair bonds.

471 Taken together, our findings suggest that different types of social associations may be
472 beneficial in different contexts and that the early-life period can be a crucial time for the
473 formation of these associations. The next step is to disentangle whether individuals actively
474 choose to (re)associate with earlier companions depending on season- and context-dependent
475 benefits, or whether the observed fluctuations in social association persistence are an
476 emergent property following relatively simple season-dependent social processes, such as
477 heightened aggression. The first scenario may have important implications for our
478 understanding of the cognitive abilities of animals. Notably, Scheiber et al. (2011) found that
479 six-week-old juvenile greylag geese can already discriminate between two of their siblings,
480 showing that individual-level recognition is already present from an early age. Our findings
481 here suggest that geese may be able to keep track of multiple types of relationships in a large
482 fission–fusion society, despite extended breaks, supporting similar observations in wild
483 barnacle geese (Black & Owen, 1995), and that they can re-evaluate the benefits of these
484 relationships depending on context. Given the strong evidence for birthplace-independent
485 long-term kin discrimination in both migrating and captive barnacle geese (Anderholm et al.,
486 2009b; Kurvers et al., 2013; van der Jeugd et al., 2002), this level of cognitive ability is
487 certainly feasible and makes the long-lived barnacle goose an interesting study system to
488 further examine such mechanisms. Complex social patterns can be driven by cognitive ability,
489 but also emerge from relatively simple processes (Kappeler, 2019), such as site fidelity
490 combined with season- and sex-dependent aggression. Unravelling how these mechanisms

491 underlie social complexity in a diversity of social systems will be central to our understanding
492 of the evolution of animal societies.

493

494

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508

509 **Data Availability Statement**

510 The raw social association observations, the matrices containing the SRI values per sex per

511 season, the matrices containing the familiarity and genetic relatedness data per sex, and the

512 raw aggression data are uploaded on the Open Science Framework for referee inspection

513 ([link](#)). These files will be made public in case of acceptance.

514

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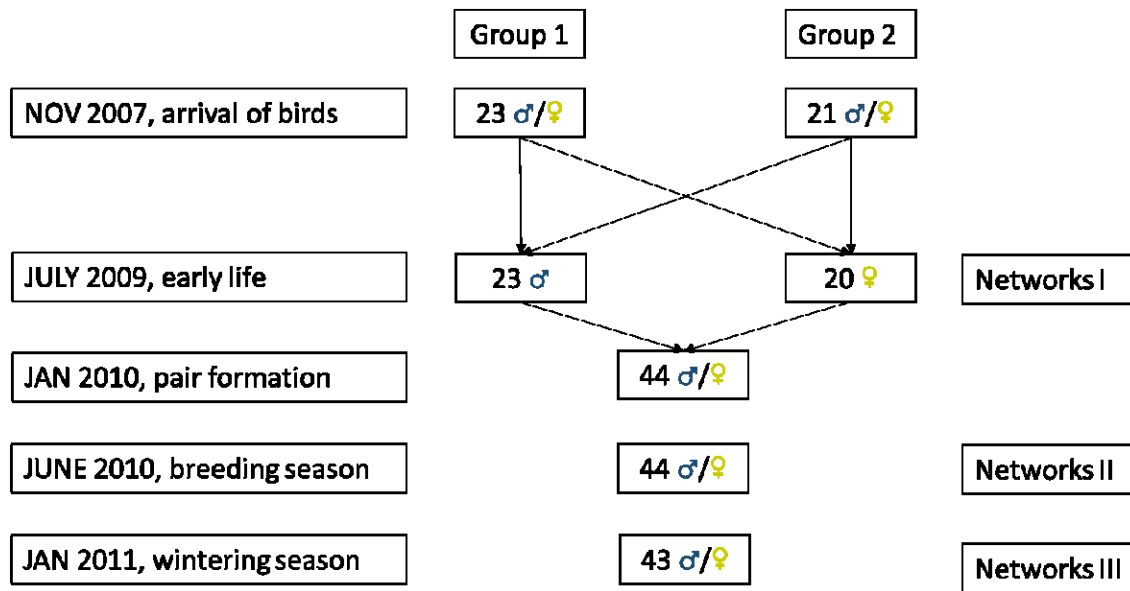
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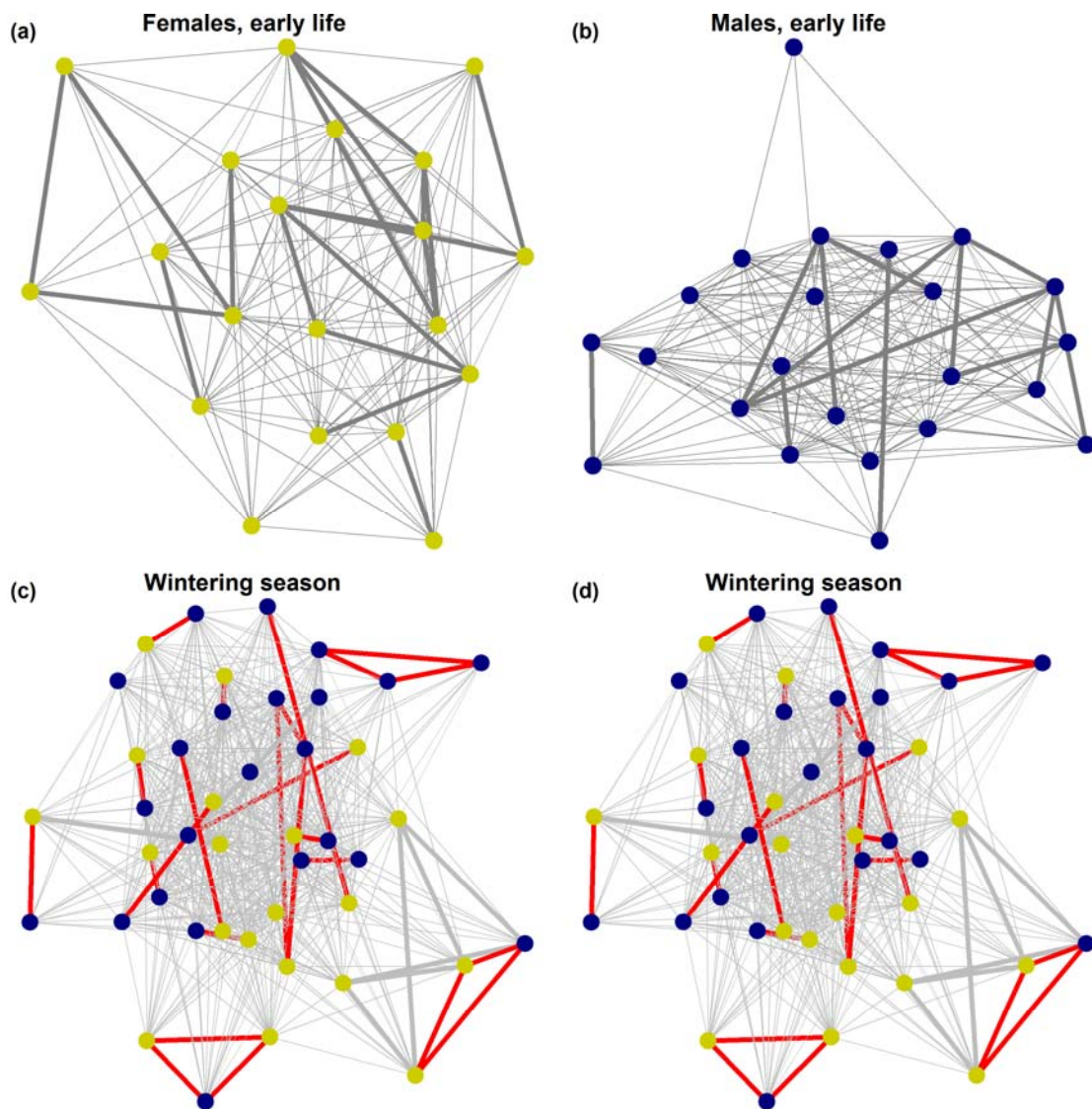


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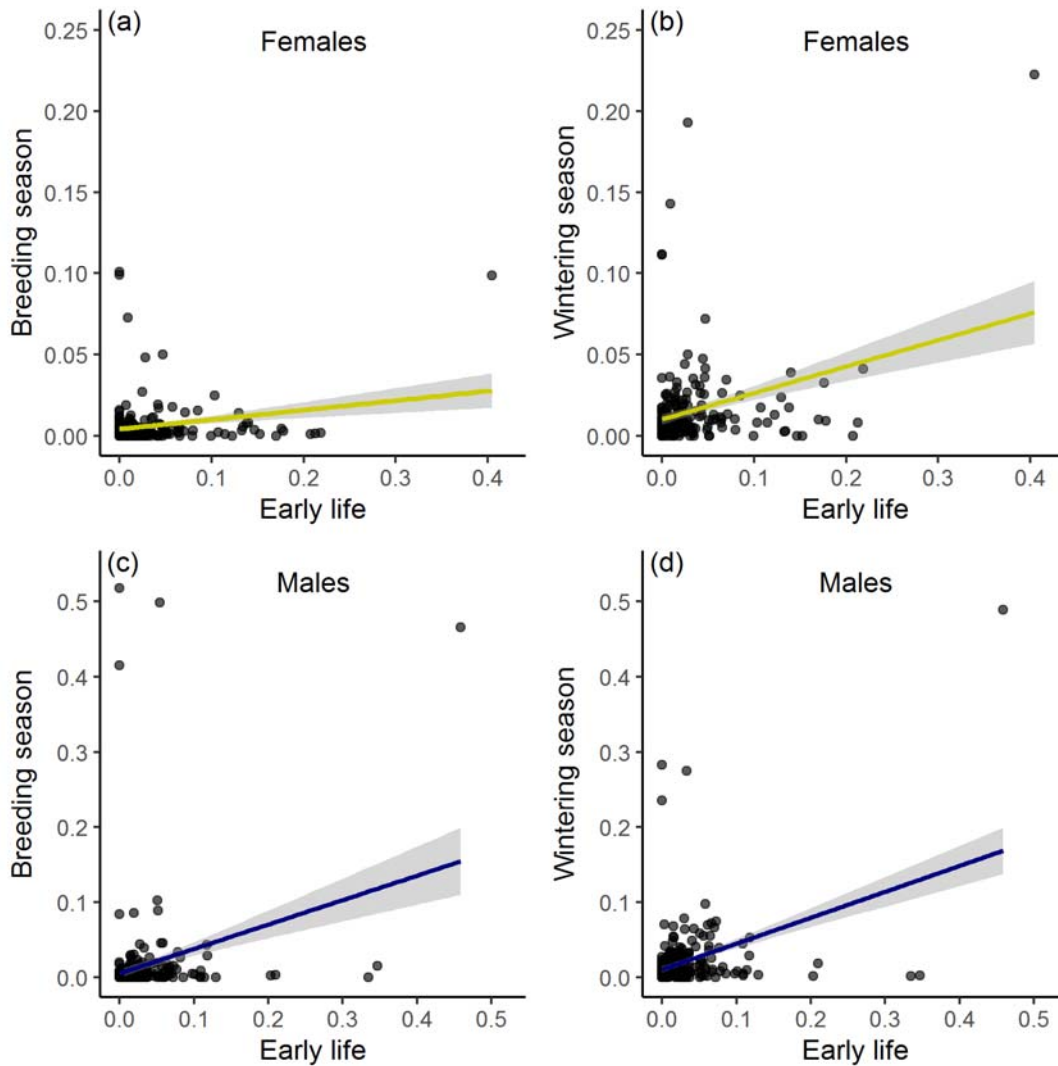
758 **Figure 1.** Schematic overview of the experimental procedures. Geese were procured in
759 November 2007—at which point almost all were juveniles (hatched in 2007)—and placed in
760 one of two familiarity groups. In July 2009, males and females of both groups were placed
761 together and we quantified the social network structure of both groups (to examine the role of
762 familiarity and genetic relatedness on social structure, Kurvers et al., 2013). In January 2010,
763 all geese were placed together in one group and individuals rapidly started to form pairs. In
764 the subsequent breeding season (June 2010) and wintering season (January 2011), we
765 quantified the social network structure of the entire group. Note that the sex of one individual
766 could not be reliably determined at the time of the first network observation and hence was
767 not included in the early-life observations. Another individual died in between the breeding
768 and wintering season. Both individuals were excluded from all analyses, resulting in a final
769 sample size of 42 (20 females, 22 males).

770



771
772 **Figure 2.** The (a) female and (b) male network in single-sex groups prior to pair formation
773 and the network of all individuals after pair formation during the (c) breeding, and (d)
774 wintering season. Yellow/blue circles represent females/males respectively. For visualization,
775 we removed associations below SRI values of 0.005. Thin/thick lines represent SRI values
776 below/above 0.1. Coloured lines indicate a pair bond. Networks were created with the `ggnet2`
777 function in R using the Fruchterman-Reingold algorithm for node placement.

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780 **Figure 3.** The relationship between the association strength of female dyads comparing (a)

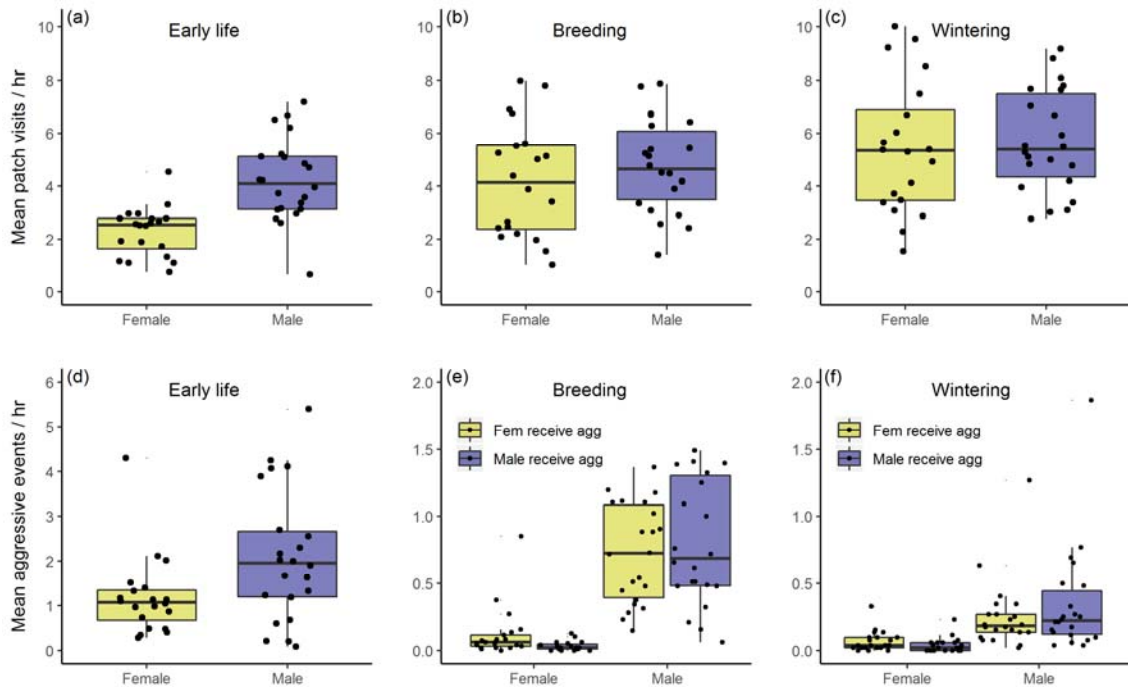
781 early life and breeding season, and (b) early life and wintering season, and the relationship

782 between the association strength of male dyads comparing (c) early life and breeding season,

783 and (d) early life and wintering season. Lines are linear regression lines, including 95%

784 confidence bands.

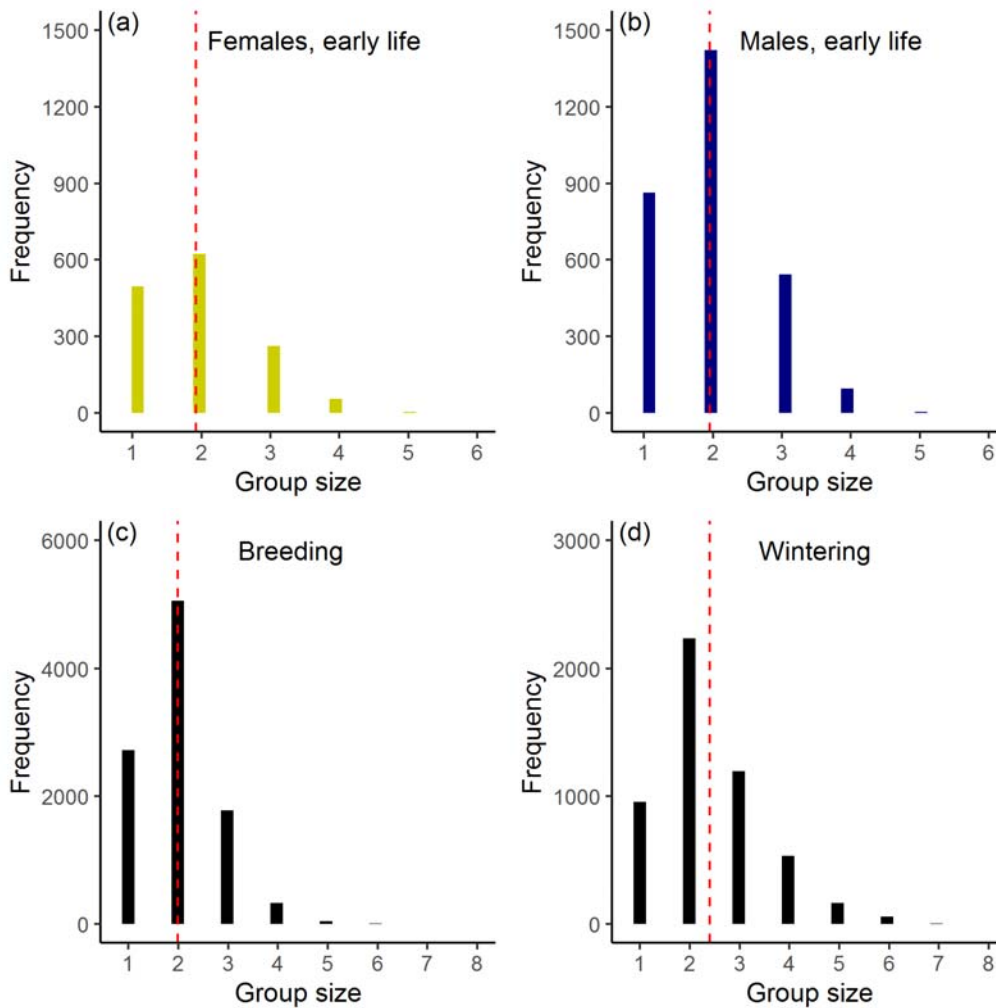
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787 **Figure 4.** The mean number of patches visited per hour for (a) females and males in the
788 single-sex groups prior to pair formation, and for both sexes during the (b) breeding and (c)
789 wintering seasons after pair formation. The mean number of aggressive events initiated per
790 hour for (d) females and males in the single-sex groups prior to pair formation, and for both
791 sexes during the (e) breeding and (f) wintering season after pair formation. (e, f) Colours
792 indicate the sex of the individual receiving the aggression. Dots represent individuals.
793 Boxplots show median, interquartile ranges and whiskers show the lowest/highest values
794 within the 1.5 interquartile range.

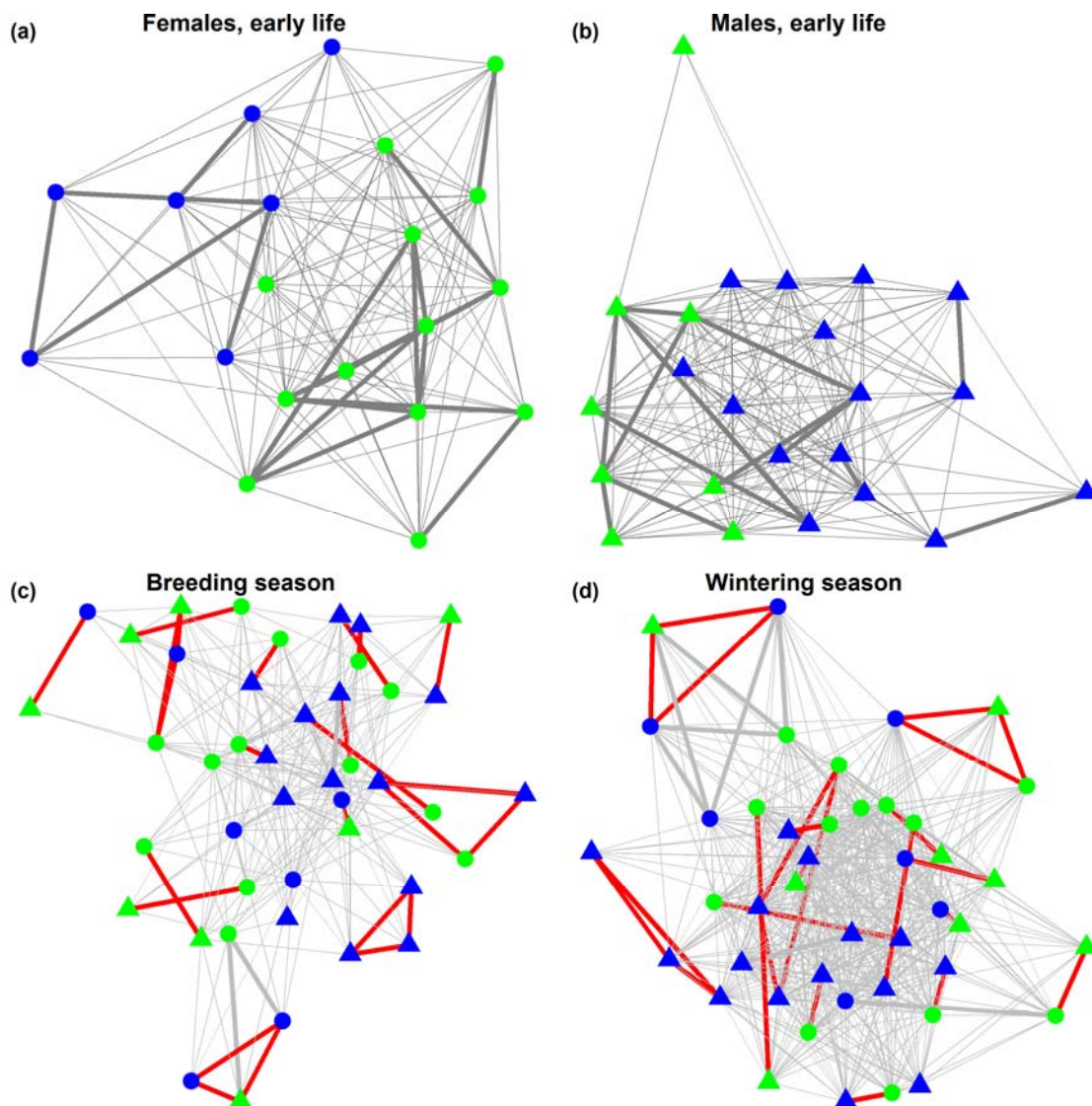
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797 **Figure A1.** (a, b) The frequency of observed group sizes during the early-life social network
798 observations for (a) females and (b) males. Red dashed lines represent mean group size (mean
799 group size females: 1.92; males: 1.96). Males frequented patches at a higher rate (see also Fig.
800 4a). (c, d) The frequency of observed group sizes during the (c) breeding and (d) wintering
801 season. Red dashed lines represent mean group size (mean group size breeding: 2.0;
802 wintering: 2.4). Note that the overall higher frequency of group sizes in the breeding season is
803 due to more observation days during breeding ($N = 25$ days) than during wintering season (N
804 $= 13$ days). The mean patch visit rate was in fact slightly higher in wintering than in breeding
805 season (see Fig. 4b, c).

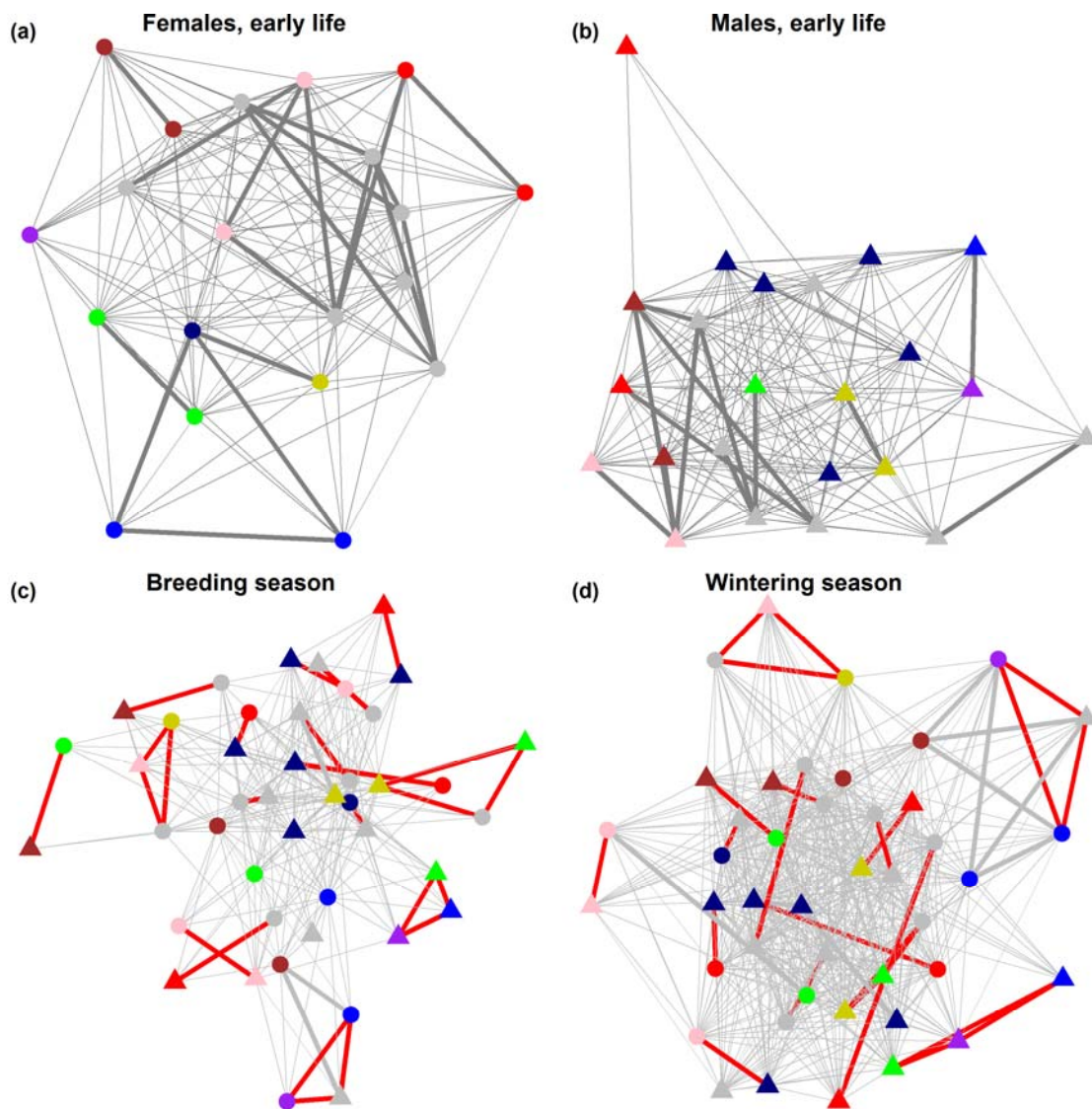
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808 **Figure A2.** Nodes with the same colour indicate individuals coming from the same familiarity
809 group. In the single-sex groups prior to pair formation, geese associated more with individuals
810 from their own familiarity group as shown by the strong segregation of colours (and as
811 reported in Kurvers et al., 2013). This pattern is, however, not present anymore in the
812 breeding and wintering season. Circles represent females, and triangles males.

813



814

815 **Figure A3.** Nodes with the same colour indicate individuals with high genetic relatedness.

816 Each cluster of colours has a mean genetic relatedness above 0.3. Grey-coloured nodes

817 represent individuals not sharing a high genetic relatedness with any individual in the

818 population. In the single-sex groups before pair formation, geese associated more with

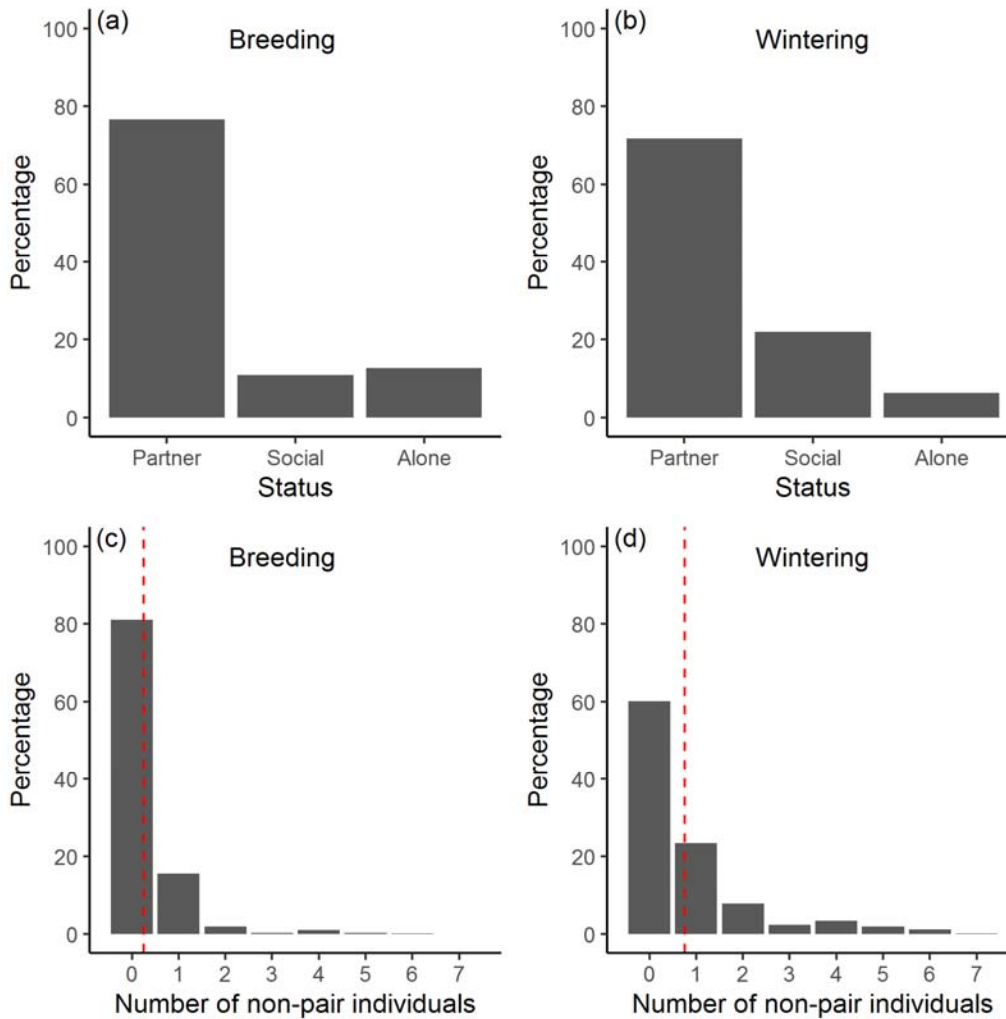
819 genetically related individuals colours (as reported in Kurvers et al., 2013). This pattern is,

820 however, not present anymore in the breeding and wintering season. Circles represent

821 females, and triangles males.

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825 **Figure A4.** (a, b) The observed likelihood that when a paired individual was present on a
826 patch that its partner was also present ('partner'), that it was together with another individual
827 but not its partner ('social'), or that it was alone ('alone'), for the (a) breeding, and (b)
828 wintering season. (c, d) The observed likelihood of the number of individuals not belonging to
829 the pair, which were present at a patch in the presence of a pair during the (c) breeding, and
830 (d) wintering season. To illustrate, in the breeding season, in 80% of cases when pair
831 members were observed together on a patch, there would be no other individuals (i.e., not
832 belonging to that pair) present. In the wintering season, this percentage was around 60%. Red
833 dashed lines represent the average number of non-paired geese present with a pair (mean
834 during breeding: 0.26, during wintering: 0.76).