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26	Kinship Dynamics: Patterns and Consequences of Changes in Local Relatedness
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48 Abstract

Mounting evidence suggests that patterns of local relatedness can change over time in predictable ways, a process termed kinship dynamics. Kinship dynamics may occur at the level of the population or social group, where the mean relatedness across all members of the population or group changes over time, or at the level of the individual where an individual's relatedness to its local group changes with age. Kinship dynamics are likely to have fundamental consequences for the evolution of social behaviour and life history because they alter the inclusive fitness payoffs to actions taken at different points in time. For instance, growing evidence suggests that individual kinship dynamics have shaped the evolution of menopause and age-specific patterns of helping and harming. To date, however, the consequences of kinship dynamics for social evolution have not been widely explored. Here we review the patterns of kinship dynamics that can occur in natural populations and highlight how taking a kinship dynamics approach has yielded new insights into behaviour and life history evolution. We discuss areas where analysing kinship dynamics could provide new insight into social evolution and we outline some of the challenges in predicting and quantifying kinship dynamics in natural populations.

76 For over half a century kin selection has been a cornerstone of evolutionary biology, providing 77 a fundamental theoretical framework for understanding the evolution of social behaviour [1]. When developing the theory of kin selection, Hamilton [1] recognised that a key factor 78 79 influencing kin structure is the degree to which individuals disperse away from their natal 80 habitat or social group. In Hamilton's original formalisation of kin selection, however, the behavioural/demographic mechanisms by which the relatedness coefficient r was generated 81 were left undefined [1]. The effects of population structure and demography (group size and 82 dispersal rates) on patterns of local relatedness (mean pairwise relatedness to nearby 83 84 individuals) were explored by Taylor [2]. Rather than specifying the level of relatedness, Taylor [2] allowed relatedness to emerge from the model. Taylor demonstrated that although a 85 decrease in dispersal generated an increase in local relatedness in structured populations, the 86 benefits of helping kin were cancelled out by the costs of an increase in competition among kin 87 [2]. In this pivotal finding, Taylor highlighted the importance of considering how dispersal 88 generates patterns of relatedness in structured populations. Over the last 30 years, a significant 89 90 focus of theoretical work has been in identifying the factors that influence local relatedness [3]. As well as patterns of dispersal, mating patterns, overlapping generations and social-partner 91 92 choice can all influence local relatedness [4-10].

93 Although often considered a static property of a group, population, or species, local relatedness can be dynamic, changing predictably with time - referred to as kinship dynamics [11, 12]. 94 95 When kinship dynamics are predictable and repeatable, then they may have profound implications for social evolution [11, 12]. Currently, however, despite many species living in 96 kin-structured populations [13], studies of social evolution have tended not to consider the rich 97 implications of kinship dynamics. Here we review the different patterns of kinship dynamics 98 that can occur in natural populations and the potential consequences of kinship dynamics for 99 100 behaviour and life history evolution. We outline current gaps in our understanding of the patterns and consequences of kinship dynamics and highlight priorities for future research. 101

102

103 **1. Patterns of Kinship Dynamics**

Here we identify three levels at which kinship dynamics may occur in natural populations (Fig
1). First, average local relatedness may change over time at the level of the population
(population kinship dynamics; Fig 1A). Second, kinship dynamics may occur at the group level
(group kinship dynamics), where the average local relatedness of members of a social group

108 changes over time (Fig 1B). Finally, kinship dynamics may occur at the individual level (individual kinship dynamics), where the relatedness of a focal individual to its local group 109 changes systematically over time, often with age [11, 12] (Fig 1C). These levels can act either 110 independently or concurrently. For example, group level kinship dynamics may occur 111 independent of population kinship dynamics - the local mean relatedness of a social group may 112 change over time while the average local relatedness of the population remains constant over 113 time. Likewise, individual kinship dynamics do not imply population or group kinship 114 dynamics and the overall mean local relatedness (of a group or population) may remain 115 116 constant over time, even though the average relatedness of an individual female or male to the rest of its group may increase or decrease. 117

118

119 1.1. Population kinship dynamics

Population kinship dynamics can be driven by temporal shifts in population density, fecundity 120 121 (reproductive success), reproductive skew, dispersal and mortality. For example, some species 122 of voles, mice and grouse exhibit cyclical kinship dynamics, linked to changes in population density [14, 17, 18] (Fig 1A). For example, in male red grouse (Lagopus lagopus scoticus) 123 124 males form territories to gain access to females. In the autumn, young of the year males compete to gain a territory and recruitment of males to a population is dependent on them 125 126 successfully securing a territory. At low population density, males are less aggressive towards kin than non-kin and can even help young related males to gain a territory by relinquishing 127 territory or by helping in territorial contests with unrelated males [19]. This process forms 128 clusters of territories held by related males [20], which in turn generates a positive feedback 129 130 that leads to the proliferation of kin clusters. A point is reached, however, where space is limited due to the density of the population and the inclusive fitness benefits of helping kin no 131 longer outweigh the direct costs of increased competition. At this point, cooperation breaks 132 down and all individuals behave aggressively. This suppresses recruitment of new males as 133 134 territory holders and the population spirals into decline and kin clusters break up [14]. This process continues until the population reaches a point where competition is low and the males 135 once again start to help recruit related males to local territories setting the population off on 136 another cycle (Fig 1A). 137

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Changes in patterns of reproductive success and survival may also change patterns of localrelatedness over time [21]. In China, for example, demographic transitions have resulted from

141 a reduction in the number of children per family (due to the country's family planning program) while life expectancy has increased. The combined effects are a decline in the number of kin 142 relationships over time but the kin relationships that do occur are longer lasting [22]. 143 Ecological forces such as predation risk and disease may generate substantial variation in 144 fecundity between groups, which can have consequences for population relatedness [23-25]. 145 Under conditions where the variance in fecundity is high, for example, a predator wiping out 146 an entire family (clutch) of offspring, rather than removing individuals from multiple clutches, 147 then this can lead to high proportions of close kin being recruited to the population in 148 149 comparison to a scenario where mortality occurs at the level of the individual [25]. Cyclical changes in predation or disease risk could, therefore, in theory, drive cyclical kinship dynamics 150 at the level of the population. The potential for such processes to drive population kinship 151 dynamics, however, remain uninvestigated. 152

153

154 **1.2.** Group kinship dynamics

155 At the level of the social group, local relatedness may be dynamic and change over time. Such group level kinship dynamics can be driven by a range of demographic, social and ecological 156 157 factors. For example, in mammalian social groups, smaller groups, with higher reproductive skew tend to have higher within-group average relatedness in comparison to larger groups [26-158 159 28]. Changes in group size, for example, as a result of changes in the ecological environment, may thus drive changes in patterns of relatedness within groups. Exactly how relatedness 160 changes with a change in group size, however, will depend on the process of group formation. 161 In banded mongooses (Mungos mungo), for example, new groups are founded when single sex 162 groups disperse from their natal group and join with groups of dispersers from other packs to 163 found a new group [15]. This "budding dispersal" can lead to an increase in local relatedness 164 with group age [15] (Fig 1B). In recently founded groups, relatedness between the sexes is low. 165 However, both males and females are highly philopatric, remaining in their natal group to 166 breed, this drives an increase in local relatedness between males and females with the age of 167 the group (Fig 1B). Such group based kinship dynamics can occur across multiple generations, 168 with the longevity of the social group exceeding the longevity of individuals [15]. The process 169 of group formation may be tightly linked to the quality of the environment, population density 170 and mortality rate. For example, in white-winged choughs (Corcorax melanorhamphos) new 171 groups were only observed after a period of unusually high mortality as a result of a drought 172 [29]. The process of new group formation was associated with reduced intra-group relatedness 173

174 due to fusions of unrelated groups/individuals [29]. At high population density, dispersal opportunities may be limited, meaning that individuals are more likely to remain with their 175 local group [30] and thus changes in density at the local or population level may drive kinship 176 dynamics. In some social systems, relatedness may actually decline with the age of the social 177 group. For example, in several ant species, within-colony relatedness decreases with colony 178 age [e.g. 31, 32]. This is driven by new queens being more likely to remain in their natal colony 179 after mating, rather than dispersing to found new colonies as local colony density increases 180 [32, 33]. Thus increasing the number of queens within the colony (increasing polygyny), 181 182 diluting relatedness between workers and leading to long-term decreases in within-colony relatedness. 183

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185 Group fission may also be a key process leading to group level changes in kin structure. For example, when the size of a social group exceeds the optimal group size, groups often split. 186 187 During such fission events, individuals may segregate based on kinship resulting in daughter groups that are more assorted by kinship than the original group (e.g. [34, 35]). Group level 188 189 changes in kin structure may also be driven by group takeovers, whereby the dispersing sex (usually males in the case of mammals), forcibly move into a group [36-38]. Such takeover 190 191 events are often associated with the eviction of residents of the same sex and the infanticide of unrelated juvenile resident offspring [36-38]. If unrelated coalitions of males take over a group, 192 193 then reproduction may be shared among the males [38], which will reduce the relatedness of 194 resulting offspring in comparison to a group with high male reproductive skew. Group fission, fusion and takeover events clearly have the potential to drive group kinship dynamics. To date, 195 however, previous work has not explicitly quantified patterns of kinship dynamics under these 196 197 social processes.

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199 **1.3. Individual kinship dynamics**

Due to demographic processes – particularly the degree of philopatry and the extent of local mating - local relatedness may change across an individual's lifetime (individual kinship dynamics (Fig 1C)). Individual kinship dynamics were first formalised in theoretical models examining the evolution of menopause in humans and some species of toothed whales [11, 12] (see Fig 2). These models demonstrate that local relatedness can change as a function of age as some individuals disperse from their natal group, and others die and are replaced by relatives [11, 12]. Johnstone and Cant [12] investigated the consequences of different patterns of 207 dispersal and mating for changes in female local relatedness across the lifespan, highlighting how demographic processes can drive individual kinship dynamics (Fig 2). Under sex biased 208 dispersal and within-group mating, the relatedness of the dispersing sex to their group (after 209 dispersal) is predicted to increase with age [11, 12] (Fig 2Aii). When the dispersing sex takes 210 up residency with their new social group, they will initially have low local relatedness. Their 211 relatedness, however, will increase with age as they reproduce and subsequently their 212 philopatric offspring reproduce and produce philopatric grand offspring, which they are related 213 to. Thus the increase in local relatedness for the dispersing individual joining a new social 214 215 group is driven by an increase in relatedness to the philopatric sex (Fig 2 Aii). Even when there is no sex biased dispersal, kinship dynamics can still occur [12] (Fig 2 Aiii). For example, in 216 some species of toothed whales, neither sex disperses and mating occurs outside of the social 217 group. Under these conditions, females are born into a group without their father and have 218 comparatively low relatedness to males in the group. As females age and reproduce, their sons 219 will remain in the local group and their relatedness to local males is predicted to increase over 220 time [12]. Work to date on patterns of individual kinship dynamics has tended to consider the 221 general pattern of kinship dynamics at the level of the population for individuals belonging to 222 a particular class (e.g. males or females). However, within a given class of individuals, there is 223 224 likely considerable variation in the kinship dynamics that individuals experience, for example, due to individual differences in dispersal and fecundity. 225

226

Few empirical studies have quantified age-dependent kinship dynamics in natural populations. 227 228 In a comparative study across 19 human communities with diverse patterns of residence and dispersal, Koster et al. [39] examined the prediction that immigrants to a community will 229 230 initially have low relatedness to the local group with few kinship ties, but that their local relatedness will increase as they populate the local group with their descendants [11, 12]. The 231 232 general predicted pattern was observed, with patterns of local relatedness increasing with age in both immigrant males and females where they were the more commonly dispersing sex. 233 Some communities, however, displayed very different patterns, which were likely due to 234 unstable settlement histories, fluid patterns of residency and community size [39]. In human 235 societies, analysis of kinship dynamics could be extended to include affinal kin (in-laws) who 236 may be treated more like biological kin than unrelated friends [40], because they share genetic 237 238 interests in future generations [41].

240 Support for individual kinship dynamics also comes from work on non-human animals. For example, in work on wild populations of killer whales that show bisexual philopatry, we have 241 found a strong match with the theoretical prediction of increasing female relatedness with age 242 (Fig 1C, Fig 2Aiii) [16]. Analogous patterns have been found in several species of cooperative 243 breeders where relatedness of an individual to the local group changes as a function of age [42-244 245 44]. For example, in African wild dogs (*Lycaon pictus*), dwarf mongooses (*Helogale parvula*) and Lake Tanganyika cichlids (Neolamprologus pulcher) the relatedness of non-breeding 246 helpers to the dominant breeders decreases with helper age due to a combination of breeder 247 248 turnover, extra-pair paternity and helper immigration [42-44].

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251 2. The consequences of kinship dynamics for social evolution

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Individual kinship dynamics may directly impact patterns of helping and harming behaviour 253 254 across the lifespan due to a change in the immediate availability of kin with age, which will drive changes in patterns of social behaviour. However, when individual kinship dynamics 255 show predictable changes, they have the potential to influence the evolution of optimal life 256 257 history and behaviour [11, 12]. For example, models of life history evolution in humans and toothed whales predict that kinship dynamics shape the selective pressures on reproductive 258 259 lifespan and can help explain why females undergo menopause [11, 12]. Under patterns of kinship dynamics where female relatedness increases with age (which occurs in resident killer 260 261 whales (Fig 1C) and is thought to be the case in ancestral humans [11, 12]) younger females 262 are predicted to be under strong selection for harming behaviour, whereas older females are 263 under stronger selection for helping behaviour [11, 12 see Fig 2]. Cant and Johnstone formalised these asymmetries in selection for helping and harming behaviour as the 264 reproductive conflict hypothesis, which predicts that when old and young females in the same 265 local group compete for reproduction, younger females should outcompete older females 266 (because they are under stronger selection for harming behaviour). When taken together with 267 the benefits grandmothers can gain by helping their kin (the grandmother hypothesis [49]) the 268 reproductive conflict hypothesis provides a mechanism to explain why females undergo 269 menopause – they are under selection to switch from a harming strategy where they reproduce 270 271 to a helping strategy where they invest in helping their kin and no longer reproduce themselves [49]. There is strong support in both humans and killer whales that there is intergenerational 272 conflict between old and young females [16, 50, 51] and that this can contribute to selection 273

for the evolution of menopause. For example, in work on resident killer whales using over 40 years of individual-based demographic and social data we showed that when mothers and daughters co-breed, the offspring from the older generation had a significantly higher risk of mortality than offspring from the younger generation female [16].

278

Evolutionary models of individual kinship dynamics to date, have tended to focus on life 279 280 history trade-offs at a given point in time [11, 12]. Selection, however, may act on temporal trade-offs between current vs future fitness, which will be shaped by individual kinship 281 dynamics. Such trade-offs would not require an individual to project likely changes in kinship 282 283 dynamics itself - selection on social traits will be influenced by the expected future kin structure and the associated fitness costs and benefits. Although there has been little explicit analysis of 284 how kinship dynamics (i.e. changing levels of relatedness over time) influence inter-temporal 285 286 trade-offs, models have shown that kin selection can modulate the balance between current and future costs and benefits (even when relatedness remains constant) [52, 53]. For example, 287 288 Lehmann [52] demonstrated that when there is a tendency for future members of an individual's community to be kin, then selection can act on organisms in a way that produces 289 290 future benefits to its community. Sozou [53] extended this concept further to examine how an individual should value future benefits to itself (individual discounting) or its social community 291 292 (social discounting), finding that the social discounting rate was generally lower than the 293 individual (private) discounting rate. The extension of kinship dynamic models to consider 294 temporal trade-offs and social/individual discounting provides an exciting avenue for future 295 research.

296

To date, models examining the evolutionary consequences of individual kinship dynamics have 297 been restricted to studies investigating the evolution of menopause in humans and toothed 298 299 whales. There is no reason to expect, however, that the behaviour and life-history implications of kinship dynamics are restricted to the evolution of this unusual life-history trait. Kin 300 301 selection models have been integrated with life-history theory in considering the evolution of senescence, where an individual's longevity can influence the vital rates of local relatives [54, 302 55]. If individuals can provide benefits to kin in late life it may select for longevity [49, 56-303 58]. In support of this prediction, recent work comparing the longevity of females across non-304 human mammals found that females in species with grandparental care lived longer [59]. Thus 305 we hypothesise that in species where relatedness increases with individual age selection will 306

307 favour longevity if there are opportunities for late life helping. The costs of interacting with kin, which can reduce the fitness of relatives, may also drive life history evolution [54]. For 308 example, theoretical work has reported that low juvenile dispersal reduces the strength of 309 selection on adult survival [55], favouring the evolution of shorter lifespans [60, 61]. Thus we 310 predict that in species where interacting with kin carries significant costs, individual kinship 311 dynamics, where relatedness increases with age could result in selection for shorter lifespans. 312 Future research is needed to understand how kinship dynamics interact with both the 313 opportunities for helping kin and the costs of interacting with kin to shape the evolution of 314 315 senescence. Theoretical work has also shown that competition among kin can shape agespecific fecundity [55]. When dispersal is limited and competition among siblings is intense, 316 selection will favour individuals to reduce competition among kin during times of peak fertility 317 and spread reproduction throughout life [55]. Kinship dynamics thus have the potential to shape 318 reproductive decisions across the lifespan, well beyond the evolution of menopause and new 319 work is needed to examine how kinship dynamics influence reproductive decisions across the 320 lifespan. 321

322

At the level of the group - group kinship dynamics generate the possibility that patterns of 323 324 helping and harming within and between groups may change over time. For example, in the case of the banded mongoose, an increase in within-group relatedness will select for increased 325 326 helping within groups which could manifest as increased helping in rearing pups and/or dominant individuals being more tolerant of the breeding attempts of related subordinate 327 328 individuals in older vs younger groups [15]. Moreover, in social groups that have high levels of within-group relatedness selection can favour indiscriminate helping whereby individuals 329 330 do not adjust the level of help they provide other individuals within the group in response to their relatedness to them [45]. Group based kinship dynamics may have consequences for the 331 332 success of the group. For example, in matrilineal groups of red howler monkeys, the average group relatedness predicts female reproductive success with females in kin groups having 333 significantly higher reproductive success [46]. In addition, theoretical work has demonstrated 334 that changes in within-group or population relatedness can change the nature of intergroup 335 violence and warfare [47]. For example, an increase in within-group relatedness may increase 336 selection for belligerence and bravery behaviours, which can result in individuals paying a 337 personal cost of death so that relatives in the group can gain benefits such as new mating 338 opportunities or additional resources [47]. Predictable changes in patterns of local relatedness 339 340 at a group level may shape how social relationships form in groups. For example, one

possibility is that founding animals may invest more in forming stronger social relationships in groups where kinship increases over time. Future empirical and theoretical work is needed to examine the ontogeny of social relationships in social groups that are undergoing different patterns of group kinship dynamics.

345

Because kin selection is conditional on the kin structure in the social environment, kinship 346 347 dynamics may result in cyclical shifts in selection which could reduce the long term efficiency of kin selection [48]. Under population and group kinship dynamics, which can occur across 348 many generations, different social traits may be selected for under different levels of local 349 350 relatedness. Such condition-dependent selection may dilute the strength of selection when social conditions change across generations or differ between social groups [48]. In periods 351 when social traits are not under strong selection, genetic drift may be enhanced, which can lead 352 to accelerated evolution - referred to as the Red King process [48]. The efficiency of kin 353 selection thus may be weaker under greater conditionality (i.e. when periods of high relatedness 354 occur infrequently). Future work is needed to directly explore the consequences of population 355 and group kinship dynamics for the efficiency of kin selection. 356

357

358 **3. Predicting and quantifying individual kinship dynamics**

359

Current models predicting patterns of kinship dynamics have tended to focus on the 360 consequences of variation in dispersal and mating [11, 12]. It is likely however that other 361 factors also play a role in driving kinship dynamics in animal societies, including group size, 362 363 mortality patterns and variation in fecundity. How well model predictions match empirical patterns of kinship dynamics will depend on how well they capture the key processes driving 364 relatedness patterns [26]. Some insight into how population traits other than dispersal and 365 mating patterns may influence individual kinship dynamics comes from a study by Rodrigues 366 [62] who examined how variation in age-dependent changes in patterns of fecundity drive age-367 dependent kinship dynamics which in turn is expected to drive age-dependent patterns of 368 helping behaviour. Rodrigues found that in populations where fecundity increases with age, 369 the average local relatedness of an individual to their local group is predicted to increase with 370 371 age, which will select for an increase in helping with the actor's age. In predicting patterns of kinship dynamics, it is important to accurately evaluate demographic features of the population 372 that can drive kinship dynamics (e.g. dispersal patterns). In some instances, the predictability 373

of kinship dynamics may be limited due to stochasticity in demographic variables (including,

375 for example, mortality, reproductive success and dispersal).

376

New work is needed to fully examine how demographic and life history traits impact on 377 patterns of kinship dynamics. Kinship dynamics are likely to show considerable differences 378 between the sexes. For example, work in humans illustrates that kinship dynamics can show 379 380 surprisingly diverse patterns across communities and the sexes [39] with local relatedness both increasing and decreasing as a function of age. In the vast majority of natural populations, there 381 is a bias towards one sex dispersing. It is well documented that sex differences in dispersal 382 383 mean that kin selection can affect males and females very differently [63]. Currently, however, how the sexes differ in patterns of kinship dynamics and the resulting implications for sex 384 differences in life history evolution remains unexplored. Moreover, within a sex it is well 385 documented that both dispersal patterns [64] and reproductive success can vary among 386 individuals, which will generate inter-individual variation in kinship dynamics. The 387 consequences of this inter-individual variation in patterns of kinship dynamics for social 388 evolution has not been explored. 389

390

To quantify patterns of kinship dynamics, we need to decide how to quantify local relatedness. 391 392 In species where individuals form stable social groups, we can track patterns of kinship by examining patterns of relatedness between individuals within a local group over time. 393 394 However, in many animal societies, social groups are dynamic and individuals interact and 395 cooperate with animals from different social units. Here we need to capture social interactions 396 within and between groups. One approach is to use social network theory and community detection algorithms to define local kinship networks [65]. It would then be possible to quantify 397 local kin structure using weighted measures of association strength, using association indices 398 [66] to weight relatedness between individuals by the strength (weight) of their association. 399

400

401 **4.** Conclusions

402

Kinship dynamics mean that the strength and direction of kin selection can change
systematically over time [11, 12, 16, 62]. To date however, very few studies have incorporated
kinship dynamics into models of social evolution. Current theory, predicting the evolutionary
consequences of kinship dynamics has tended to focus on species that undergo menopause.

However, the evolutionary consequences of kinship dynamics are likely to extend well beyond
this unusual life history trait and new work is needed to develop a general understanding of the
evolutionary consequences of kinship dynamics at the level of the population, group and
individual.

411

A particularly exciting avenue for future research is the comparison of kinship dynamics 412 between the sexes. Life history differences between the sexes within a species are widespread 413 and significant attention has been given to examining the mechanisms driving sex differences 414 in life history evolution [67]. It is possible that in many species, the sexes experience very 415 different patterns of kinship dynamics that could have been a significant force driving sex 416 differences in life history evolution. Because kinship dynamics are driven by demographic 417 418 processes, kinship dynamics are likely to show profound differences between species. For 419 example, individual kinship dynamics are dependent on patterns of philopatry, the general 420 pattern of which differs between birds and mammals (e.g. in birds females normally disperse 421 more than males whereas in contrast in mammals males typically disperse more than females 422 [68]) leading to the prediction that general patterns of kinship dynamics may differ between taxa. Moreover, because group size, predation and fecundity likely play a key role in driving 423 424 patterns of kinship dynamics, it is likely that patterns of kinship dynamics differ between populations of the same species living under different ecological conditions. Future work 425 426 comparing patterns of kinship dynamics between species, populations and the sexes is eagerly 427 anticipated.

428

Patterns of kinship dynamics may help explain variation in social behaviour across the lifespan and contribute to patterns of social ageing. In humans and non-human primates there may be fundamental shifts in social interactions as individuals age [69]. This can include the propensity to engage in social interactions, the capability to influence others and the positive vs negative valence of social interactions [70]. How patterns of individual kinship dynamics influence the cost-benefit trade-off of social interactions across the lifespan and thus patterns of social ageing remains unexplored.

436

Future research should examine how active partner preferences shape kinship dynamics and
how individuals respond to demographic events in their social network and the consequences
of these events for kinship dynamics. For example, long term observations on female baboons

440 (*Papio cynocephalus*) has shown that following a mother's death, maternal sisters strengthen their social bond whereas the bonds with maternal aunts are weakened [71]. Feedbacks may 441 exist between changes in behaviour and changes in relatedness that generate further changes in 442 relatedness and behaviour. For example, excluding individuals from groups to minimise 443 competition could generate increases in local relatedness which further selects for increased 444 investment exclusion (since this is an altruistic act that benefits everyone in the group, it is 445 favoured by higher r). In principle, this kind of positive feedback could lead to a runaway 446 outcome that results in a very high level of local relatedness over time. Such feedbacks 447 448 however, have not been formalised in models of group formation and social evolution, providing an exciting opportunity for future research. 449

450

Here we have focused on changes in local relatedness over time, however, the other terms in 451 Hamilton's rule – the costs (c) and benefits (b) may also change over time [72] and interact 452 with individual kinship dynamics. For example, reproductive value (the expected contribution 453 of an individual to the future population) can significantly impact kin directed behaviours [73]. 454 The costs of a social act depend in part on the reproductive value of the actor and the benefits 455 on the reproductive value of the recipient, both of which can change with age [74]. In many 456 457 species, the ability of individuals to help and provide benefits to kin may also be age-dependent [75]. For example, as individuals age, they develop skills, knowledge and experience that 458 459 increase the benefits they can pass on to kin [76-79]. For example, in African elephants (Loxodonta africana) older females are better able to assess social and predatory threats, which 460 461 can increase the survival of their younger kin [78, 79]. Similar results have been reported in 462 resident killer whales where old post-reproductive females lead their group around foraging 463 grounds, especially in times of low food abundance [77]. Kinship dynamic models provide a conceptual framework to be able to examine how changes in patterns of local relatedness with 464 age interact with changes in the costs and benefits of social acts. 465

466

Given the ubiquity of kin-structured social groups in populations [13], kinship dynamics are likely to have widespread consequences for the evolution of social behaviour and life history in both males and females. Researchers working on kin structured social species often have the data needed to quantify kinship dynamics. We encourage researchers to consider patterns of kinship dynamics in their study systems and the potential consequences that they may have for social evolution.

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700 Figure 1. Examples of different patterns of kinship dynamics. (A) Population kinship dynamics 701 in red grouse (Lagopus lagopus scoticus) using data from an 8-year cycle in a natural population [figure redrawn from 14]. Here kinship dynamics show a cyclical pattern with local 702 703 relatedness increasing during population growth as established males help related males to 704 establish a neighbouring breeding territory. As population density increases the costs of competition with neighbouring kin outweighs the benefits leading to increasing aggression and 705 706 the break-up of kin clusters. (B) Group kinship dynamics of male – female relatedness in social groups of banded mongoose (Mungos mungo). New social groups form by the fission of 707 708 unrelated male and female groups. As social groups age, male-female relatedness increases due 709 to philopatric males and females inheriting the breeding positions [figure redrawn from 15]. (C) Individual kinship dynamics in Northern and Southern resident killer whales (Orcinus 710 orca) showing patterns of maternal relatedness with female age for a total of 200 whales over 711 846 whale-years [figure redrawn from 16]. Figure shows the relationship between female age 712 and mean relatedness to other females (red line), males (blue line) and average relatedness 713 across both sexes (black line). Standard error of mean relatedness across both sexes is shown 714 as dotted lines. As females age, their local relatedness to the group increases due to an increase 715 716 in local relatedness to males. Animal images from PhyloPic: red grouse image (uncredited) 717 published under Public Domain Dedication 1.0 license (https://creativecommons.org/publicdomain/zero/1.0/); banded mongoose image by Birgit 718 719 Lang published under Public Domain Dedication 1.0 license (https://creativecommons.org/publicdomain/zero/1.0/); killer whale image by Chris Huh 720 721 published under Creative Commons Licence 3.0 (https://creativecommons.org/licenses/by-722 sa/3.0/).

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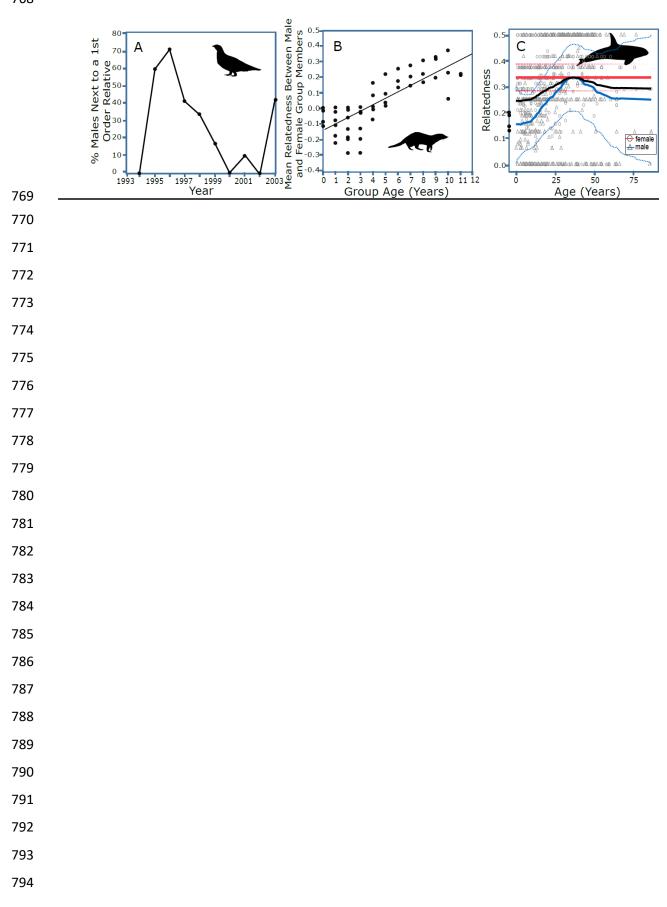
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Figure 2. Age-dependent kinship dynamics and their consequences for selection on social traits 725 726 under different demographic patterns (modified from [12]). A) Predicted age-dependent kinship dynamics, showing age-specific relatedness to a breeding female of other local males 727 728 (dotted lines) and females (solid lines) and the average relatedness across both sexes (dashed line). Age is scaled in mean generation lengths. Three different demographic patterns are 729 730 plotted: (i) local mating within the group with high male dispersal and low female dispersal; (ii) local mating within the group with low male dispersal and high female dispersal; (iii) non-731 local mating (i.e. mating occurs outside the group) with low dispersal by both sexes. B). The 732

733	consequences of age-dependent kinship dynamics for selection for helping (boosting the
734	fecundity of other local breeders) and harming (reducing the fecundity of other local breeders)
735	across the lifespan under different demographic patterns (i-iii as in A). A focal female can
736	perform social acts that result in an immediate gain of b offspring for other breeders at an
737	immediate $\cos t c$ to herself. Selection for helping and harming are indicated by shading on the
738	graph with blue areas indicating selection for helping and red areas indicate selection for
739	harming. For further details, see Johnstone & Cant [12].
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