Offspring social network structure predicts fitness in families

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Social structures such as families emerge as outcomes of behavioural interactions among individuals, and can evolve over time if families with particular types of social structures tend to leave more individuals in subsequent generations. The social behaviour of interacting individuals is typically analysed as a series of multiple dyadic (pair-wise) interactions, rather than a network of interactions among multiple individuals. However, in species where parents feed dependant young, interactions within families nearly always involve more than two individuals simultaneously. Such social networks of interactions at least partly reflect conflicts of interest over the provision of costly parental investment. Consequently, variation in family network structure reflects variation in how conflicts of interest are resolved among family members. Despite its importance in understanding the evolution of emergent properties of social organization such as family life and cooperation, nothing is currently known about how selection acts on the structure of social networks. Here, we show that the social network structure of broods of begging nestling great tits Parus major predicts fitness in families. Although selection at the level of the individual favours large nestlings, selection at the level of the kin-group primarily favours families that resolve conflicts most effectively.

Keywords: social networks; sexual conflict; parent–offspring conflict; begging; social evolution

1. INTRODUCTION

Social networks analyses have advanced our understanding of the evolution of animal societies [1,2], cooperation [3–5], the transmission of disease [6] and human social [7] and socio-economic [8] systems. Taking a networks approach to the study of social behaviour shifts emphasis away from variation in behaviour among individuals per se to how interactions among individuals shape variation [9]. This more realistically reflects the behaviour of individuals as being both the cause and the effect of their social environment [10,11]. However, all previous research has focused on the importance of social position within a network on the fitness prospects of interacting individuals [12,13] rather than the structure of the network of interactions themselves. As a result, very little is known about the relationship between the structure of social networks and fitness in natural populations, despite the importance of such information in understanding the evolutionary and ecological significance of social networks [9,14].

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of altricial species of birds provide an ideal study system to quantify between-group consequences of variation in SNS on fitness. There are no ‘gambit of the group’ issues (a common problem in social networks analyses where assumptions are made about social groupings of individuals based on their patterns of associations with one another; [1]) as all nestlings within each nest can be clearly and unambiguously assigned to a given group. In addition, because each individual within a brood interacts with all other individuals in the network and group membership is clearly defined, it is possible to make use of weighted networks metrics (i.e. incorporating the number of times or strength with which individuals interact with one another) as opposed to binary network metrics (where pairs of individuals in a network are simply classified as either associating with one another or not). This facilitates a measure of the strength of the behavioural interactions among network members (‘gregariousness’; [2]) and means no potentially important information is lost [2]. Finally, and perhaps most importantly, this form of study system allows the novel quantification of the fitness consequences of variation in social interactions among individuals at levels of selection higher than the individual through the use of replicated networks.

We used a nest-box population of great tits in the forests around Bern, Switzerland, to examine how the structure of social interactions among nestlings is related to the fitness of both offspring (recruitment into the population to breed) and parents (survival and breeding success in the year following the experiment). Sixty-three broods ranging in size from five to 10 nestlings were filmed during parental feeding events on day 10 post-hatch, when feeding rates are at their peak [25]. Variation in the hunger of nestlings (and, therefore their motivation to beg; [22]) in each brood was manipulated and the position and identity of the parent and the position, identity and begging intensity of all nestlings were recorded (see the electronic supplementary material for further details). Nestling positions were used to produce association matrices of begging nestlings. These matrices were then used to derive network metrics for each nestling within each network (social network position, SNP; which is an individual level trait) and summary metrics to describe the structure of each network (SNS, which is a group level trait). These were then used to relate SNP and SNS to measures of fitness and to quantify how SNS is related to interactions between parents and offspring (see §2).

Parental care is nearly always costly [26], and because individuals within families are not fully related to one another conflicts of interest are expected over the provision of parental investment [27–29]. The amount of parental investment provided depends on the outcome of such conflicts (the ‘resolution’), which are determined by how individuals interact with one another [18]. Great tit male and female parents do not differ to one another in feeding rate to nestlings at the population level but feed (predictably) from different positions in the nest and have different feeding rules to one another [22]. Mothers, but not fathers, respond to an increase in the begging calls of offspring by increasing their provisioning rate and preferentially feed hungry nestlings [22,23]. In contrast, fathers take longer to choose which offspring to feed and preferentially feed nestlings that jostle and compete for food most effectively [22,23]. These different feeding rules mean that the SNS of nestlings is predicted to be largely determined by whichever parent provides the most feeds and primarily controls the allocation of parental resources. Great tit nestlings respond more readily to greater female parental sensitivity to their state (i.e. they approach their mother when hungry and move or are displaced when fed; [22]). Broods where mothers, rather than fathers, primarily control the allocation of resources to nestlings are therefore expected to be composed of more strongly interacting (‘gregarious’) nestlings, as the higher responsiveness to variation in offspring state of mothers encourages nestlings to move around more in the nest in relation to the position of the feeding parent, reducing the variation in nestling state. In contrast, in broods where fathers provide most of the feeds, we expect a lower mean strength of interactions among nestlings because fathers preferentially feed the most competitive nestlings which can monopolize positions closest to where the male parent feeds from, so there will be less movement within the nest. The SNS of these broods is likely to be more clustered as more competitive nestlings can occupy the best positions near to the male parent [22] leading to increased variation in within-brood nestling state. If SNS is largely a consequence of variation in parental sensitivity to offspring state then the position of nestlings within the network (SNP) should be influenced by hunger and more strongly connected broods are expected to have a more uniform distribution of begging behaviour among offspring.

2. METHODS

Information on video analysis and preparation of data for social network analysis is provided in Kölliker et al. [22] and also in the electronic supplementary material.

(a) Data collection and experimental manipulation of nestling hunger

Data collection has been previously described in detail by Kölliker et al. [22]. We provide a brief synopsis here. A nest-box population of great tits nesting in the Bremgarten forest near Bern, Switzerland, was used for the study. Nestlings at experimental nests were ringed with numbered aluminium rings 9 days post-hatching and a dummy camera was installed in the nest-box to habituate the birds to the presence of a camera. On day 10 post-hatch individual nestlings at each nest-box were weighed (± 0.1 g) and uniquely marked on the head with paint. Two intermediate-sized nestlings were then temporarily removed and randomly assigned to one of two treatments: food-deprived or fed to satiation for 2 h with bee larvae. This allowed us to assess the effects of hunger on SNP. After 2 h, the two nestlings were then replaced in an arbitrary position in the nest and parental provisioning and nestling begging behaviour filmed from above using a camera with an infrared light source for 45 min at 63 nests. Brood size at the time of the experiment ranged from five to 10 nestlings (the mean brood size is 7.4 nestlings in this population).

(b) Social network analysis

For each nestling at each feed, we established the direct social associates (i.e. the identity of the individuals immediately next to, or touching, the focal individual). These positions
were calculated for each feed at each brood and the data used to produce association matrices for each brood. Association was measured as the proportion of feeding events in which individuals were immediate neighbours of each other member of the network; where a score of 1 between two given individuals means that they were always next to each other and a score of 0 means two individuals never associated. Data were extracted from data sheets involving feeding events at nests for calculation of matrices using MATLAB (MathWorks, Natick, MA, USA) and then UCINET [30] was used to derive our network metric, weighted degree (also known as node strength [1]), for individuals within broods. Weighted degree is defined as the total weight of the edges (social associations) connected to a node (individual), and provides a measure of SNP for each individual (i.e. the number and strengths of associations with other nestlings of a focal individual within the brood; [1]). Degree, whether unweighted or weighted, is a simple, robust metric that is widely used [1,2] and is a potentially important determinant of the evolution of cooperation [3,4]. We used weighted degree as our key network metric because individuals within broods all interacted with one another, so binary (unweighted) network metrics would be uninformative. In contrast, weighted degree allowed us to quantify the strength of interactions among individuals, which was our primary focus. Broods with higher mean weighted degree scores have greater overall strength of associations among interacting nestlings (i.e. nestlings moved around more so encountered other individuals more frequently; they were more gregarious, [2]) than broods with low mean weighted degree scores (SNS; figure 1).

(c) **Statistical data analysis**

Statistical analysis was performed using R version 2.14.2 (Copyright © 2007 The R Foundation for Statistical Computing) and SPSS version 16 (SPSS Inc., Chicago, IL, USA). We accounted for correlations among continuous predictors by including terms as covariates in models, rather than using residuals or other forms of variance partitioning, as this has been shown to be the most effective method for dealing with collinearity [31,32]. During model simplification, we removed non-significant interactions, followed by lower-order terms in turn from the maximal model until no further terms could be dropped without significantly reducing the model fit (minimum adequate model, MAM; [33]).

We used general and generalized linear mixed models (GLMMs) for analysis of individual level effects, with nest as a random term to account for non-independence of data from individuals in the same network. GLMM model simplification involved comparing maximum-likelihood (ML) models with and without each term. We took a term out of the model if its removal did not significantly increase the Akaike information criterion (AIC; [33]). For brood level effects we used general and generalized linear models (GLMs) and model simplification used analysis of deviance (a measure of the relative fit of the model compared to alternative models; [33]). Unless stated otherwise sample size for brood level analyses was 63 broods and for individual level analyses was 450 nestlings in 63 broods.

3. **RESULTS**

(a) **Recruitment probability of the brood**

The first question we addressed was whether there was any evidence that selection acts on the SNS of great tit broods. In order to answer this, we examined whether SNS explained a significant amount of variation in the proportion of nestlings in a brood that survived to recruit into the breeding population the following year. The relationship between the strength of associations among nestlings and recruitment success of broods was dependent upon brood size; being positive in small (five or six nestlings) and medium (seven or eight nestlings) sized broods, but negative in large broods (nine or 10 nestlings; table 1a and figure 2a). In contrast to smaller broods, large broods with weaker networks of interactions were more successful than those with stronger nestling associations (table 1a). Variation in the strength of associations among nestlings within broods (CV of SNS) did not explain variation in recruitment success.

(b) **Recruitment probability of individual nestlings**

We then examined whether recruitment probability at the level of the individual was best explained by individual
Sex and brood mean strength of associations (SNS) on the number of fledglings reared (table 1d). Males that reared broods with strongly associating offspring fledged a lower number of nestlings in the following year than females that reared strongly associating broods of nestlings (figure 2b). In addition to the significant sex × SNS interaction, the number of fledglings produced by surviving parents was significantly affected by whether they had a new partner in the following year or not (in virtually all cases new partners were present because the previous partner did not apparently survive); parents with new partners had lower reproductive success. However, neither brood size nor the mean mass of the nestlings significantly affected reproductive output in the following year (table 1d).

(c) Future reproductive success of parents

Does variation in SNS of broods also predict future parental success? Male parents had a lower probability of survival to the following year than females, but there was no significant effect of SNS on the survival of parents (table 1c). However, for parents that survived to breed there was a significant interaction between sex and SNS on the number of fledglings reared (table 1d). Males that reared broods with strongly associating offspring fledged a lower number of nestlings in the following year than females that reared strongly associating broods of nestlings (figure 2b). In addition to the significant sex × SNS interaction, the number of fledglings produced by surviving parents was significantly affected by whether they had a new partner in the following year or not (in virtually all cases new partners were present because the previous partner did not apparently survive); parents with new partners had lower reproductive success. However, neither brood size nor the mean mass of the nestlings significantly affected reproductive output in the following year (table 1d).

(d) Social network position

As predicted, the SNP of experimental nestlings was significantly related to manipulated levels of hunger; nestlings that were food-deprived had stronger associations with their siblings.
with other nestlings (i.e. were more gregarious) than satiated individuals (table 2a). Brood size was also positively related to SNP as expected (i.e. individuals had a greater number of associates in larger broods), and male nestlings were more gregarious than females, perhaps because males are larger than females, so became hungry more quickly. However, nestling mass at day 10 was dropped from the maximal model, as were all relevant interactions (table 2a).

(e) Social network structure and begging behaviour

If SNS reflects variation in the response of male parents compared with female parents to offspring behaviour, we predicted that SNS would be related to the evenness of begging within broods, with a positive relationship indicating that mothers primarily controlled feeding. As expected SNS was significantly related to the begging behaviour of nestlings. Broods with more strongly interacting nestlings had higher mean begging intensity and were more uniform in their pattern of begging behaviour than broods with less strongly associating nestlings (table 2b), with larger broods having a stronger relationship between SNS and evenness (1-CV) of begging than smaller broods (table 2b; figure 1c). The mean begging intensity of broods was negatively related to the mean number of feeds provided by the male, but was not related to brood size or the number of female parent feeds (table 2c): the harder the male worked the less the nestlings in the brood begged (i.e. the less hungry they were). Conversely, variation in begging intensity within broods was primarily driven by how hard the female parent worked: the evenness of begging behaviour within broods was positively related to the number of feeds provided by the female and not to brood size or the number of male feeds (table 2d).
Feeding behaviour

If SNS of broods is related to which parent primarily controls feeding we predicted that broods controlled by mothers would show a positive relationship between SNS and the number of feeds, and mothers would spend less time choosing which nestling to feed. As expected the number of feeds provided by parents to nestlings was negatively related to SNS (brood mean degree) for males, but positively related to SNS for females (figure 1d). Brood size dropped out of the model (table 3a). In contrast, females spent longer choosing which nestlings to feed when SNS was low, whereas males spent longer choosing when SNS was high (table 3b).

4. DISCUSSION

Despite widespread interest in animal social networks [1,2,14], and the recognition that understanding the evolutionary and ecological importance of the structure of social networks requires information on the relationship between SNS and fitness [9], there have been only two previous studies examining fitness in relation to social network metrics [12,13]. Moreover, both of these studies were concerned with the relationship between the position of individual adults within networks and correlates of fitness. To our knowledge, our study is therefore the first to relate fitness to variation in the structure of whole, replicated, networks, not just position within a single network. We show that SNS predicts fitness in
broods of great tits in the wild, but SNP does not. Selection acting at the level of the individual primarily favours large, well-nourished offspring, as might be expected, whereas selection acting at the level of the family depends upon how gregarious offspring are (SNS) in relation to the size of the network (brood) involved. Variation in social network attributes [35] or traits correlated with SNS, such as begging intensity or parental feeding behaviour, can be heritable [19,23,36,37]. If, as seems probable, SNS has a heritable basis, then family structure can evolve.

Selection acting at the level of the family emerges as a consequence of how interactions are distributed among nestlings during feeding by parents. The SNS represents the behavioural outcome of the resolution of within-family conflict over the provision of parental investment in terms of nestling positioning, which is a primary determinant of the probability of being fed in great tits [22]. Selection favours networks that are composed of strongly interacting (gregarious) individuals when broods are small and medium sized, with the evidence indicating that these patterns of associations are a result of offspring responding more readily to the feeding rules of mothers (who in turn are more responsive to variation in offspring state) than fathers [24]. Conversely, the finding that selection favours networks of interactions in large broods that are relatively weak indicates that nestling mobility can be constrained in large families owing to limited space, imposing costs on gregariousness when there are many mouths to feed. Thus, selection on network structure in great tit families depends on family size, a condition that may contribute to the maintenance of heritable variation in attributes of social networks or traits that correlate with SNS through genotype × family environment interactions.

As predicted, variation in the SNS of broods was related to differences in feeding behaviour by mothers and fathers. Broods with begging offspring that were more gregarious with one another were associated with mothers providing relatively more feeds than fathers, whereas broods with weakly interacting offspring were associated with fathers providing relatively more feeds. Consequently, the evidence indicates that resource allocation in broods of strongly associating nestlings is primarily controlled by mothers, not fathers. Furthermore, the mean time spent choosing before feeding nestlings showed the opposite pattern to the number of feeds. Female parents spent longer choosing which nestling to feed when the mean strength of interactions within broods was weak (i.e. when males provided more feeds), whereas male parents spent longer choosing nestlings when nestlings were highly gregarious (i.e. when females provided more feeds). Since it is expected that parents will take longer, on average, to decide how to allocate their resources when they have less information about brood need (i.e. when they provide a lower proportion of the number of feeds compared with their partner; [38]), this further supports the contention that high strength of interactions among nestlings (high gregariousness) in broods is associated with females primarily controlling feeding whereas weak interactions among nestlings are associated with greater male control.

Social network position of individual nestlings was primarily determined by hunger, and the consequent increased motivation to beg [22]. SNS was therefore strongly related to begging behaviour. Broods of highly gregarious nestlings had higher overall begging intensity and greater evenness of begging behaviour across the brood. Begging intensity was more evenly distributed across the brood when females provided more food, indicating that females are more responsive to nestling hunger than males [22,23]. However, the negative relationship between the number of feeds provided by males and mean brood begging intensity shows that attending to offspring demands is also dependent on providing sufficient food.

A probable explanatory scenario for the different relationships between SNS and the feeding behaviour of males and females is as follows: hunger drives offspring motivation to position themselves with respect to feeding parents [22]. Mothers are more responsive to variation in hunger than fathers, so nestlings move about more as they become hungry to gain access to feeds provided by mothers. Broods composed of strongly associating

### Table 3. Analyses of parental feeding behaviour. Only significant interaction terms are shown. Parameter estimates are given with s.e. in brackets.

<table>
<thead>
<tr>
<th>Model</th>
<th>Response variable</th>
<th>Terms in model</th>
<th>Terms dropped</th>
<th>Parameter estimates</th>
<th>Test statistic</th>
<th>d.f.</th>
<th>P-value</th>
</tr>
</thead>
<tbody>
<tr>
<td>(a) GLMM with normal errors (nest as random effect; n = 124 parents in 62 nests)</td>
<td>Mean number of parental feeds to brood</td>
<td>SNS × sex of parent</td>
<td>-0.46 (0.14)</td>
<td>9.73</td>
<td>1</td>
<td>0.0018</td>
<td></td>
</tr>
<tr>
<td></td>
<td></td>
<td>SNS</td>
<td>-0.17 (0.19)</td>
<td>0.80</td>
<td>1</td>
<td>0.372</td>
<td></td>
</tr>
<tr>
<td></td>
<td></td>
<td>Sex of parent</td>
<td>1.15 (0.36)</td>
<td>0.11</td>
<td>1</td>
<td>0.741</td>
<td></td>
</tr>
<tr>
<td></td>
<td></td>
<td>Intercept</td>
<td>2.43 (0.49)</td>
<td>0.10</td>
<td>1</td>
<td>0.749</td>
<td></td>
</tr>
<tr>
<td>(b) GLMM with normal errors (nest as random effect; n = 124 parents in 62 nests)</td>
<td>Mean time spent feeding brood</td>
<td>SNS × sex of parent</td>
<td>0.60 (0.24)</td>
<td>6.04</td>
<td>1</td>
<td>0.014</td>
<td></td>
</tr>
<tr>
<td></td>
<td></td>
<td>SNS</td>
<td>0.03 (0.27)</td>
<td>0.01</td>
<td>1</td>
<td>0.912</td>
<td></td>
</tr>
<tr>
<td></td>
<td></td>
<td>Sex of parent</td>
<td>-1.37 (0.60)</td>
<td>0.30</td>
<td>1</td>
<td>0.584</td>
<td></td>
</tr>
<tr>
<td></td>
<td></td>
<td>Intercept</td>
<td>3.37 (0.68)</td>
<td>0.53</td>
<td>1</td>
<td>0.466</td>
<td></td>
</tr>
</tbody>
</table>

*Test statistics for both models are likelihood ratios.*

(gregarious) nestlings are therefore characterized by female control of resources to offspring (i.e. higher feeding rate by the female compared with the male parent, a more even distribution of begging among nestlings and higher overall begging intensity). In contrast, broods characterized by relatively weak patterns of associations among nestlings had higher feeding rates by fathers relative to mothers, a more skewed distribution of begging behaviour and lower overall intensity of begging. Fathers are less responsive than mothers to offspring demands [22,23], so in broods where the feeds are primarily controlled by male parents it does not pay hungry nestlings to preferentially move towards fathers, who have feeding rules that primarily favour more competitive, not necessarily hungrier, offspring. Relatively high mean strength of interactions among nestlings is selectively advantageous in small- and medium-sized broods, whereas relatively weak interactions among nestlings are favoured when broods are large. The results indicate that fathers that put relatively more effort into provisioning appeared to gain more control over resource allocation, reflected in the SNS of the brood (lower gregariousness of nestlings), perhaps because they obtain more information about the need or quality of their brood [38]. However, the higher provisioning effort of these males may be offset by the increased efficiency of the allocation of their parental investment because males rearing broods with weakly interacting networks of offspring had higher future reproductive success than males with broods of strongly interacting nestlings.

Recent theoretical analyses have shown that the controllability of networks depends upon the distribution of behavioural interactions within networks [39]. Networks with dense, relatively homogeneous interactions among individuals are easier to control than sparse, heterogeneous networks [39]. In our population of great tits, dense, homogeneous networks (i.e. those with highly gregarious individuals) are characteristics of broods where mothers feed more than fathers. This provides further support that, in the most common sized broods, it is mothers that primarily control SNS not fathers. However, because it is more difficult for mothers to attend to offspring demands and/or the simpler feeding rules of fathers are more effective with many nestlings, in large broods selection favours weaker interactions among nestlings and a more skewed distribution of begging among nestlings had higher feeding rates by fathers relative to mother. This fits with theory showing that the benefits of male parental care to females are expected to co-evolve with clutch size; the larger the clutch the greater the benefit of increased male care [40]. Our results suggest a mechanism for how this might be maintained in great tits: the use of different feeding rules by mothers and fathers.

SNS can affect average group performance so it can shape the structure of social interactions within groups, and, therefore, social evolution [1,2,10,14,41], including the evolution of family life [19]. Further work is needed to determine causality in the relationship between parent–offspring interactions and brood SNS, and the mechanistic basis of the effect of nestling SNS on recruitment probability. However, the current study shows that selection can act at the level of the family on variation in parental–offspring behaviours that affect kin-group structure. How individuals interact with each other may be at least as important as the phenotypic characteristics of the interacting individuals in determining how selection acts on families. These results lend some support to a recent study on cooperation in humans by Fehl et al. [42] showing that coevolutionary relationships between behaviour and SNS can increase cooperation beyond direct reciprocity itself. In great tits, associations between feeding behaviours of parents and SNS affects how conflicts over investment are resolved, which may lead to selection on families that are most efficient at resolving conflicts (i.e. ‘cooperative families’). Our results are also applicable to any network of individuals whose behaviour is influenced by how they are ‘managed’ and whose success depends on group performance. In our study, selection acts on parent–offspring relationships in birds, but there are parallels in how individual humans interact in business and in team sports structures that would repay further investigation using our approach.

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REFERENCES
10 Moore, A. J., Brodie, E. D. & Wolf, J. B. 1997 Interacting phenotypes and the evolutionary process: I. Direct and


