The quality of social relationships affects male health in a complex bird society

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Some animal societies, particularly in primate and corvid species, are based on

enduring social bonds, based on sophisticated cognitive skills. A key unresolved issue

for understanding the evolution of complex sociality and the associated advanced

cognition is to uncover the fitness advantages that social relationships convey to

individuals. Here, we investigated whether the social environment and the quality of

individuals' social relationships in captive groups of carrion crows affect excretion

patterns of coccidian oocysts, which have been shown to exert detrimental

consequences on bird health. In females, variables relating to the social environment

and the quality of individuals' social relationships did not have a strong influence on the

occurrence of coccidian oocysts. However, male crows with strong social bonds and

with a large number of relatives in their group were less likely to excrete feces

containing coccidian oocysts. Thus, our data present evidence that high quality social

relationships help male carrion crows to keep endo-parasite burden low, suggesting that

the social life might have potential fitness advantages by reducing parasite burden.

Ultimately, these advantages might be important in driving the evolution of complex

group living.

Significance statement

Uncovering the fitness benefits of close social bonds is a key factor in

understanding why close relationships between individuals have evolved in many

distantly related taxa, such as mammals and birds. Here, we investigated potential

health benefits of close social bonds in carrion crows. Carrion crows live in complex

male-biased social groups and possess high cognitive skills that allow managing social

relationships. Male crows with strong social bonds and with a large number of relatives

in their group were less likely to excrete feces containing coccidian oocysts. Our results

show that social bonds might have direct fitness implications in male crows, which

ultimately might drive the evolution of complex group living.

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Introduction

In many animals, the benefits of conspecific aggregations (e.g. protection against predators, increased access to food through shared defense or knowledge of resource locations) can arise in unstructured assemblages, where anonymous individuals show minimal interactions that they can manage with a relatively small brain and unsophisticated cognitive abilities (e.g. fish schools, ungulate grazing herds). However, some animal societies, notably those in some primate species, are based on enduring social bonds and according to the 'social brain hypothesis', the management of these complex relationships requires sophisticated cognition [1].

Currently, the key issue for understanding the emergence of this kind of complex sociality and the associated cognitive skills is to uncover the fitness advantages that bonding conveys to individuals [2]. In spite of much interest, this question is largely unresolved. Research in primates has shown that the quality of social relationships may affect individuals' longevity and reproductive success. The largest body of evidence supporting this hypothesis has been found in humans, where a rich social network and social support yields measurable positive consequences for health [3]. Other primate species show similar patterns. Social relationships help female chacma baboons (*Papio hamadryas ursinus*) cope with stressful events and increase longevity [4], social bonds in male Assamese macaques (*Macaca assamensis*) are directly linked to the number of offspring sired [5], and in Barbary macaques strong social bonds have a stress reducing effect [6]. These studies, together with more limited information on other mammalian species [1, 7], suggest that the human tendency to engage in enduring social bonds may have deep evolutionary roots [8]. To test this fascinating hypothesis, however, we

crucially need to investigate the fitness consequences of social bonds in a larger variety

of taxa and types of social organizations [2].

Recent research has shown that advanced sociality associated with sophisticated

cognition is not unique to humans and closely related primate species, but can be found

in a variety of taxa [9, 10]. Among birds, corvids show complex social relationships,

proving a suitable model to investigate the fitness consequences of social bonding.

Common ravens (*Corvus corax*), for example, are capable of forming valuable

relationships, not only within reproductive pairs but also between short/long-term

coalition and alliance partners in their social group [11, 12]. Such bonds are

characterized by low levels of aggressive and high levels of affiliative behaviors

exchanged in a reciprocal way over extended periods of time [13]. Within such valuable

relationships, individuals support each other in agonistic encounters [12, 14] and share

information and resources [12, 15, 16], whereas competition dominates in other social

contexts [17, 18].

Carrion crows (*Corvus corone*) are also capable of developing enduring non-

reproductive social bonds, especially among males, which in some populations can

share and defend a territory, and cooperate in rearing a communal brood [19, 20].

Within-groups, linear dominance hierarchies help reduce conflicts and keep the group

cohesive [21, 22]. As expected, cognitive skills mirror the social complexity in these

species. For example, crows can discriminate between reliable and unreliable

conspecifics [23] and show inequity aversion, being able to quantify other individuals'

efforts and payoff in comparison with their own in token exchange tasks [24]. Carrion

crows are therefore a suitable model to investigate the fitness correlates of social

bonding and, ultimately, the evolution of the social brain [25].

In this study, we investigated whether the social environment and the quality of

an individual's social relationships affects health in captive groups of carrion crows that

varied in size and composition. In particular, we focused on the pattern of excretion of

coccidian oocysts, as a proxy of health in carrion crows. Coccidian protozoans are

among the most common endoparasites of birds [26, 27] and are widespread among

corvids, including the carrion crows [28–30]; this study, see below. Infections with

large numbers of coccidia clinically manifest as 'coccidiosis', causing infections in the

intestinal tract [31]. This severely affects individual body condition, longevity and

fecundity, by inhibiting the uptake of essential dietary components [32, 33]. The

pathogenicity is well documented in poultry [34–36], and evidence for it is further

accumulating in wild bird species (e.g. [37, 38]). Coccidia are transmitted via the fecal-

oral route. Coccidian oocysts excreted in the feces of host individuals become infectious

by undergoing sporogony within 24 hours [35].

In this study, we address three non-mutually exclusive hypotheses: 1. Group size

hypothesis: parasite excretion patterns depend on the size of the group and/or the

number of related individuals. We expect infections to spread more easily in large

groups, although the presence of relatives in the group may counteract the effect of

crowding by reducing the occurrence of stressful conflicts and competition [39] that can

affect vulnerability to parasite infection. 2. Social bonds hypothesis: parasite excretion

patterns depend on the quality of the social relationships that an individual maintains,

with individuals with strong social bonds being less likely to be infected than

individuals with weak social bonds. 3. Intrinsic factors: parasite excretion rates are

explained by an individual's sex and age. We did expect sex differences, because males

are more competitive than females within their groups [21] and are probably more

exposed to the negative consequences of stress, which is expected to impair their ability

to fight against parasites [40]. Also, previous results in wild and captive carrion crows

suggest that males are more sensitive to the presence/absence of relatives [41] and

therefore are expected to be more likely to show a health response to the composition of

their social group compared to females. Age must also be controlled for, as it could

affect parasite infection rate, although no specific predictions can be formulated based

on the current knowledge of the host species.

Methods

Study subjects and ethics statement

We collected data over four different periods, from November 2008 to January 2009,

from January to February 2013, from May to June 2013 and from January to February

2014, from a total of 34 crows, which were housed in a large outdoor aviary (30 x 12 x

6 m) in Northern Spain (Navafria, León, 42°36'33 N 5°26'56 W). The aviary comprised

four major compartments (12 x 6 x 3 m), connected through eight smaller testing

compartments (3 x 3 x 3 m). The major compartments were equipped with wooden

perches, natural vegetation and rocks. An enriched diet consisting of fruit, vegetables,

bread, meat and milk products was provided on a daily basis. Water and dry food were

available *ad libitum*. Keeping crows in captivity was authorized by Junta de Castilla y

León (núcleo zoológico 005074).

Birds were kept in groups that mirrored the social aggregations that naturally occur

in the wild: (A) 'Flock': three or more juvenile individuals; in 2008 some juveniles

were initially kept in 'sibling pairs' and subsequently merged into a juvenile flock; (B)

'Pairs': adult male and adult female; in some cases adults were also kept in trios; (C)

'Cooperative Family': a reproductive pair with its own already independent offspring.

Group composition, however, varied due to the death of some individuals and

reproduction of breeding pairs, as summarized in Table S1.

Behavioral protocol data

A total of 1075 individual focal protocols have been recorded. Each focal protocol

lasted 5 minutes, and all occurring behaviors where recorded. However, for the present

study, we focused on frequencies of agonistic behavior (threat, chase flight, and fight)

and affiliative behaviors (approach, allopreening and sitting in contact). The identity of

the interacting individuals was noted, as well as their role (initiator/receiver). All

behavioral protocols (see sample size in Table S1) were recorded on video and analyzed

by CW.

Composite sociality index

For each period of data recording, we calculated a composite sociality index (CSI)

for each crow dyad within a group according to Silk, et al. 2010 [4]. A CSI was

constructed for five affiliative behaviors as follows: $(AI_{ij}/AI_{ave} + AR_{ij}/AR_{ave} +$

$CS_{ij}/CS_{ave} + API_{ij}/API_{ave} + APR_{ij}/APR_{ave})/5$. AI_{ij} represents the mean number of

approaches initiated by individual i towards individual j per focal protocol, and AI_{ave} is

the mean number of approaches initiated by all members of the group per social

protocol. Similarly, AR refers to approaches received, CS to contact sit, API to

allopreening initiated and APR to allopreening received. Note that we used mean values

instead of absolute numbers because the sample of focal protocols varied among

individuals (see Table S1). A second CSI for four agonistic behaviors (aggression

initiated won, aggression initiated lost, aggression received won, aggression received

lost) was calculated as follows: (AIWij/AIWave + AILij/AILave + ARWij/ARWave +

ARLij/ARLave)/4. For each period of data collection, we determined the average and

maximum affiliative CSI for each dyad. Further, over the entire study period and all

observed dyads, we calculated a mean affiliative and aggressive CSI. When an

individual fulfilled both conditions of having a higher than average affiliative CSI and

lower than average aggressive CSI with another individual, they were considered

closely bonded. In each period of data taking we determined the number of group

members to which each individual was closely bonded.

Parasitological examination

During the entire study period, 399 individual fecal samples were collected, directly

after defecation (for a detailed overview see Table S1). Further, from April 2013 to

October 2014, 178 fecal samples were collected from 30 different groups belonging to a

long term studied population of wild carrion crows [42] in the rural area 'La Sobarriba',

Northern Spain (42°37′ N, 5°26′ W). For the purpose of this study, after observing a

crow group for up to 30 min, we approached the area where the crows were foraging

and looked for fresh fecal samples on the ground. We could assign the samples to a

given group but not to individual crows. Sex and age of the wild crows were unknown.

Eggs and oocysts of intestinal parasites were searched in fecal samples. In 2008, we

used a modified version of the flotation method [43] to examine the

occurrence/abundance of parasite products in the fecal samples. Fresh feces (0.1g) were

suspended in a 2 ml collecting tube with 1 ml saturated saline. Collection tubes were

shaken for 10 seconds and afterwards centrifuged for 5 minutes at 3000 rpm. After

centrifugation, the collection tubes were filled with saline solution and a cover slip (18 x

18 mm) was positioned onto the tube. The high density of the saline solution causes the

parasite eggs and oocysts to float up and be caught on the cover slip [44]. After 10

minutes, the cover slip was moved onto an object slide and the parasite eggs and

oocysts were identified (by size and shape) and counted. From January 2013 onwards,

we used a McMaster counting chamber to examine the occurrence of parasite products

in the fecal samples. The entire fecal sample was weighed, then diluted with 3 ml

saturated NaCl solution per 0.1 g of feces and thoroughly mixed. Afterwards, the

solution was filled into both McMaster counting chambers. After 10 minutes of resting

period we counted parasite eggs and oocysts in both chambers and calculated the

number of parasite products per 1 ml of feces.

We used a binocular with 100-fold and 400-fold amplification to identify parasite

products. We found Coccidian oocysts, eggs of several nematode species (*Capillaria*

sp., *Ascarida sp.*, *Syngamus sp. and Heterakis sp., Trichostrongylus tenius*) and cestode

eggs to a varying degree. The proportion of positive samples was highest for coccidian

oocysts (24.7 %, n = 336), whereas the prevalence of eggs of nematode and cestode

species ranged between 1 and 17 %. We therefore restricted our analyses to coccidia,

the frequency of which allowed for robust statistical testing. Coccidia are a diverse

group of protozoan parasites which frequently infect the avian intestinal tract, causing

lesions [45]. Because the abundance of parasite cocciadian oocysts in the feces depends

on the stadium of the parasite cycle, which is difficult to control, we analyzed

presence/absence instead of absolute numbers.

Data analysis

Parasites with different life cycles and virulence may exert diverse and

synergetic effects on birds [43]. Exploring the causes and consequences of multiple

parasitism exceeded the aims of this study, which focused on the most common

infection found in crows at the study place, i.e. coccidia, and that represents a threat for

many species of wild birds [46, 32, 47]. Thus, to reduce the environmental noise

associated with the presence of multiple parasites, we restricted our analyses to samples

infected only by coccidia, i.e. with no indications of other parasitic taxa (nematode

eggs, cestodes). Likewise, we considered control ('healthy') samples those that were

found free from any of the parasitic taxa described above.

We used generalized linear mixed models (GLMMs) with binomial error

distributions and logit link functions in the package lme4 [48] to examine the variation

in the occurrence of coccidian oocysts (presence of oocyst in the feces = 1, absence =

0). We used an information-theoretic approach to assess which variables and models

best explained the variation in the data [49]. The information-theoretic approach

recognizes the uncertainty associated with biological data, and allows calculating

model-averaged parameter estimates and standard errors by considering non-exclusive

candidate models. Such analytical approach has been shown to be particularly suitable

for the comparison of alternative hypotheses – *i.e.* competing models – when multiple

variables may depict the data [50, 51], which is the case in our study. We built up and

compared models composed of explanatory variables associated with the three sets of

hypotheses that may influence the occurrence of coccidia infection as described in the

introduction section. We considered three age categories (first-year, second-year and

adult) and assessed the differences by Tukey post-hoc testing in the multcomp package

[52]. We conducted separate analyses for each sex because of different expectations

regarding the influence of social factors on the occurrence of coccidian oocysts between

males and females (see introduction and supplementary materials). Apart from sex, we

did not include highly correlated variables (see supplementary materials)

simultaneously in the same model to avoid potential spurious results due to

multicollinearity. For each model, we fitted individual identity as a random term to

control for the potential dependence associated with multiple samples from the same

individuals. Given the low number of individuals per group for each sex (19 males in 14

groups, 15 females in 12 groups), we did not fit group identity as a random term in the

models to avoid redundancy. Overall, we ran two groups of models (16 and 11 models

for males and females, respectively) that included combinations of low-correlated

explanatory variables as well as the intercept-only models, i.e. the null models. We

ranked the models for each candidate group of models according to Akaike's

Information Criterion corrected for small sample sizes (AICc) and Akaike model

weights [49, 53] by using the MuMIn package [54]. Models with lower AICc values

and higher Akaike model weights are better supported by the data, with the sum of all

model weights being one [49]. We calculated model-averaged parameter estimates and

standard errors for each parameter by considering all candidate models within each

group of models. Statistical analyses were performed in R 3.2.2 [55].

Results

Social bonds and parasitological examination

We observed 57 bonded dyads, 31 of which were male-female dyads (19 between related individuals and 12 between unrelated individuals), 18 were male-male (5 between related and 13 between unrelated individuals) and 8 female-female (6 related, 2 unrelated). On average, males and females had 1.7 ± 1.71 and 1.64 ± 1.60 bonded partners respectively. The mean CSI of related male-female bonds was 2.135 ± 0.846 and 3.034 ± 2.165 for unrelated male-female bonds. The mean CSI of male-male bonds was 2.345 ± 0.952 for related and 2.754 ± 1.798 for unrelated individuals. In female-female bonds, the mean CSI was 4.806 ± 1.549 and 1.556 ± 0.101 for related and unrelated partners respectively.

Overall, 83 out of 336 samples contained coccidian oocysts, whereas 253 samples were not infected (i.e., free of parasite products of any of the considered taxa). Coccidian oocysts occurred in 57 of 198 samples in males, and in 26 of 138 samples in females. In the droppings collected from wild carrion crows, 84 out of 178 samples contained coccidian oocysts, showing that coccidia are a common parasite in groups of wild carrion crows.

(a) Occurrence of coccidian oocysts in males

The selection of models that examined the occurrence of coccidian oocysts in males yielded a high-ranking model that accounted for a weight of 0.41 (Table 1). The maximum CSI value and the number of related individuals in the group were included

in the best model. Crows with a strong social partner and living with a larger number of

related individuals were less likely to contain coccidia (Table 2, Figures 1a and 1b).

The addition of the factor group size to the model with the maximum CSI value

and the number of related individuals did not improve model performance, as indicated

by the higher AICc value and the similar log-likelihood value of the second-ranking

model compared to the best model (Table 1). Further indication of the low influence of

group size on the occurrence of oocysts arises from the fact that the 95% confidence

interval of the model-averaged parameter estimate for group size included zero (- 0.187,

+ 0.433; see Table 2).

Model averaging results also suggested a tendency for males with fewer

affiliated individuals to be more likely to host coccidia compared to males with more

affiliated individuals (Table 2). However, the number of affiliated individuals was fitted

in models with rather low weights (0.01 - 0.06, Table 1), which suggests a low

influence of this variable. Likewise, the average CSI value was fitted in low-ranking

models (Table 1) and its model-averaged parameter estimate clearly included zero

(Table 2), suggesting low explanatory power.

The third-ranking model included age as a fixed term (Tables 1 and 2). Post-hoc

analyses by Tukey contrasts showed that first-year males were less likely to host

coccidia than second-year birds (second-year – first-year: estimate $\pm$ SE $= 2.44 \pm 0.89$, z

$= 2.733$, $p = 0.016$) and adults (first-year – adult: estimate $\pm$ SE $= - 2.62 \pm 0.90$, $z = -$

2.896, $p = 0.010$), whereas no difference was found between the latter two age

categories (second-year – adults: estimate $\pm$ SE $= -0.17 \pm 0.55$, $z = -0.312$, $p = 0.946$).

Although age was highly correlated with the maximum CSI value and the number of

related individuals (see Methods section), the addition of these two variables

significantly improved the performance of the model based on age (Anova test: AICc =

203.14 vs 208.08 respectively for the model with the three variables and the model with

the age only, Chi-squared = 9.172, df = 2, p = 0.010). Thus, both the age, the maximum

CSI value, and the number of related individuals had a strong influence on the

occurrence of coccidian oocysts in males.

The negative relationship between the maximum CSI value and the occurrence

of oocysts may be difficult to interpret. On the one hand, it may indicate that birds with

a highly affiliated group member are less likely to excrete fecal samples containing

coccidian oocysts, but, on the other hand, it may also suggest that only healthy crows

are able to maintain strong social bonds. To disentangle the two possibilities, we ran

two additional analyses. First, we fitted a binomial GLMM with the maximum CSI

value at a given day t as a predictor of the occurrence of coccidian oocysts later in time

($t + n$ days), including individual identity as a random term and two additional

explanatory terms, namely the occurrence of parasites at time t and the time elapsed

between measuring maximum CSI values and assessing the occurrence of oocysts (n

days). Neither of the two latter variables were significant (occurrence of parasites at

time t: estimate ± SE = 0.611 ± 0.520, z = 1.175, p = 0.240; number of days elapsed:

estimate ± SE = 0.001 ± 0.004, z = 0.320, p = 0.749), whereas crows with higher

maximum CSI values proved less likely to host coccidia later in life (estimate ± SE = -

0.191± 0.077, z = -2.477, p = 0.013). This significant relationship supports a causal

effect of maximum CSI value on the occurrence of the parasites. In a second analysis,

we addressed the possibility that infected birds were less able to maintain strong social

bonds. In this case, we expected that the presence of oocysts in the fecal samples of a

given individual at time t would predict a low maximum CSI value later in life ($t + n$

days). However, a linear mixed-effects model, where individual identity was fitted as

random term, showed that the presence of coccidia at time t did not significantly reduce

the maximum CSI value at time $t + n$ days (estimate ± SE = -0.104± 0.126, t = -0.834, p

= 0.406), after controlling for the maximum CSI value at t (estimate ± SE = 0.981 ±

0.018, t = 55.038, p < 0.001) and the number of days elapsed between assessing the

occurrence of oocysts and measuring maximum CSI values (estimate ± SE = -0.005 ±

0.001, t = -4.456, p < 0.001). The results of this second model therefore also supports

the conclusion that a strong social bond enhanced resistance against coccidian

endoparasites.

(b) Occurrence of coccidian oocysts in females

The selection of models that examined the occurrence of coccidian oocysts in

females yielded five high-ranking models that accounted for more model weight than

the intercept-only model (Table 1). The best model included age as a fixed term and

was approximately 10 times more likely to be the best model than the null model (i.e.,

model weights: 0.49 vs 0.05). Post-hoc analysis by Tukey contrasts showed that first-

year females were less likely to host coccidia than adult females (first-year – adult:

estimate ± SE = - 1.37 ± 0.50, z = -2.729, p = 0.017), whereas no difference was found

between second-year and adults (second-year – adult: estimate ± SE = - 1.27 ± 0.69, z =

-1.847, p = 0.151), nor between second-year and first-year (second-year – first-year:

estimate ± SE = 0.11 ± 0.73, z = 0.143, p = 0.989).

379    The second-ranking model included group size as an explanatory variable (Table

380 1). Despite model-averaging analyses suggesting that females in smaller groups tended

381 to be less likely to host coccidia (Table 2), this model had a ΔAICc value of only 1.68

382 compared to the null model. Moreover, the second-ranking model was only around two

383 times as likely to be the best model as compared to the null model (model weights: 0.11

384 vs 0.06), suggesting a limited influence of group size on coccidian infection in female

385 crows.

387    The average and maximum CSI values, as well as the number of affiliated

388 individuals and the number of relatives, were fitted in rather low-ranking models

389 (ΔAICc to the null model < 0.75, Table 1). Examinations of model-averaging parameter

390 estimates confirmed a low influence of these explanatory variables on the occurrence of

391 coccidian oocysts in females (Table 2).

Discussion

In the present study, we have shown that carrion crow males with closely

bonded social partners (high CSI scores) proved less likely to excrete coccidian oocysts.

The fact that high CSI values predicted low occurrence of coccidia at a later point in

time and that, conversely, the presence of oocysts in the feces did not predict the

strength of future social bonds, confirms the causal link between the quality of social

relationships and reduced occurrence of coccidian oocysts, which can be considered an

estimate of health [32, 33].

Health correlates of high quality social ties have been reported in humans (e.g.

[56]) and a large body of research has focused on the causal psychological and

biological processes that underlie this strong link [3, 57]. Similar evidence has been

recently found in non-human primates [4, 5], suggesting that the tendency to form

strong social ties may have deep phylogenetic roots [58]. Furthermore, as Silk [2]

pointed out, it is important to recognize that social complexity is not limited to primate

species. Indeed, coalition formation has been reported in several mammalian and bird

species, e.g. African elephants *Loxodonta africanus* [59], spotted hyenas *Crocutta*

crocutta [60], bottlenosed dolphins *Tursiops aduncus* [61], ravens [62], and carrion

crows [20]. Our study uncovered for the first time, potential health consequences of

high quality social relationships in a bird species, showing similarities with primates

and suggesting that the direct fitness benefits of strong social bonds may be a key factor

that triggers advanced sociality in a variety of taxa.

Cooperatively breeding groups of carrion crows form through two different

mechanisms, namely delayed dispersal of offspring, which remain in the natal territory

with their parents and siblings [63], and formation of long lasting social bonds among

distant relatives, most often males, that share all purpose territories and frequently mate

polyandrously [19, 20]. These bonds typically form when an immigrant male joins an

established family group and allies with the same sex resident breeder, to whom it is

related [20]. Cooperative breeding, with group members working together to raise the

brood, can only arise once stable groups have formed [64]. Both offspring delayed

dispersal and bonding between adults males are therefore necessary preceding steps that

eventually lead to cooperation, which has to be considered a consequence rather than

the cause for the formation of the social group [65–67]. In other words, to understand

why carrion crows form groups we need to understand the fitness advantages that

sociality conveys, independently of the payoff derived from cooperation at the nest. In

carrion crows, as well as in many other cooperatively breeding species, the benefits of

delayed dispersal for offspring are well studied [42], but little is known about the fitness

advantage of forming long lasting social bonds. We suggest that the health benefits of

high quality relationships may be an important factor for establishing stable alliances

between males.

Besides the positive effect of strong social bonds, we found that the number of

relatives in the group also affected parasite excretion patterns in carrion crows, with

individuals living in groups with more related individuals excreting less samples

containing coccidian oocysts. Sociality is based on kinship in wild cooperatively

breeding carrion crows. Offspring remain in the natal territory with their parents for

years and more distant relatives are actively recruited to form cooperative alliances with

the resident breeders [42]. As a result, social groups in cooperatively breeding carrion

crow are extended families, comprising members with different degree of relatedness.

Indirect fitness benefits are known to be a primary driver of kin-based sociality in many

taxa of animals [69]. However, our results now show that kin living can also accrue

immediate direct benefits to carrion crows in terms of improved health, as estimated via

parasite product excretion.

Male versus female sociality

In cooperatively breeding carrion crows, group sex ratio is male biased [68].

Within the social groups, conflicts are more likely among males, which form stable

linear hierarchies that are much stronger than those of females. This sex difference

mirrors the level of competition for resources, such as territory and reproduction, which

is higher in males than females [21, 68]. Females of a group also generally gain

unrestricted access to food sources, whereas males often interfere with each other and

fight [39]. In their groups, males are therefore likely to undergo higher levels of social

stress than females. This may explain why the observed benefits of social bonding in

carrion crows arise in males but not in females. First year males were also less likely to

excrete feces containing coccidian oocysts than second year and adult males. This may

also be due to the increasing level of competiveness that males experience when they

age and face the challenge of acquiring a territory and breeding position in the

population.

Conclusions

Social complexity, associated with high cognitive abilities, has deep

phylogenetic roots, being widespread not only among primate species but also in a

variety of other mammals and birds. Understanding its evolution, however, is difficult

because of our scant knowledge of the fitness consequences of sophisticated social

behavior across taxa. We are just starting to fill the gaps, and this study shows a

measurable positive health effect of the quality of social relationships in a bird species

that may trigger social bonding, setting the stage for a complex form of cooperation, i.e.

cooperative breeding, to arise. Importantly, our results parallel previous findings in

primate species, suggesting a general pattern in vertebrates for the evolution of

sociality.

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Figure 1: Occurrence of coccidan oocysts in carrion crow droppings. Occurrence
probabilities were calculated as the predicted values (solid lines) and 95% CI (dashed
lines) from the model-averaged parameter estimates for maximum CSI value (a) and
number of related individuals (b).

Table 1. Model selection of analyses that examined the occurrence of coccidian oocysts

in males and females. The occurrence of oocysts was related to variables associated

with three sets of hypotheses: the "group structure" hypothesis (group size, number of

related individuals), the "social bonds" hypothesis (maximum CSI value, average CSI

value, number of affiliated individuals) and the "age" hypothesis. Individual identity

was fitted as a random term in all models. LogLik: log-likelihood, df: degrees of

freedom, ΔAICc: difference in AICc to the best model. Models are ranked according to

their Akaike weight (Weight). N = 198 samples of 19 males and 138 samples of 15

females.

Sex	Hypotheses	Models	df	LogLik	AICc	ΔAICc	Weight
males	Group structure + Social bonds	No. related ind., Max. CSI	4	-98.16	204.52	0.00	0.41
	Group structure + Social bonds	No. related ind., Max. CSI, Group size	5	-97.64	205.59	1.08	0.24
	Age	Age	4	-99.94	208.08	3.56	0.07
	Group structure + Social bonds	No. related ind., No. affiliated ind.	4	-100.06	208.32	3.80	0.06
	Age + Social bonds	Age, Average CSI	5	-99.05	208.42	3.90	0.06
	Group structure	No. related ind.	3	-101.64	209.41	4.89	0.04
	Group structure + Social bonds	No. related ind., No. affiliated ind., Group size	5	-99.78	209.88	5.36	0.03
	Social bonds	Max. CSI	3	-102.10	210.33	5.81	0.02
	Social bonds	No. affiliated ind.	3	-102.39	210.90	6.38	0.02
	Group structure	No. related ind., Group size	4	-101.61	211.42	6.91	0.01
	Null model	Intercept-only model	2	-103.99	212.04	7.52	0.01
	Group structure + Social bonds	Group size, Max. CSI	4	-102.01	212.23	7.71	0.01

	Social bonds	Average CSI	3	-103.36	212.84	8.32	0.01
	Group structure + Social bonds	Group size, No. affiliated ind.	4	-102.37	212.94	8.42	0.01
	Group structure	Group size	3	-103.81	213.75	9.24	0.00
	Group structure + Social bonds	Group size, Average CSI	4	-103.08	214.38	9.86	0.00
females	Age	Age	4	-62.22	121.74	0.00	0.49
	Group structure	Group size	3	-64.76	135.69	2.95	0.11
	Group structure	No. related ind.	3	-65.22	136.62	3.88	0.07
	Group structure	Group size, No. related ind.	4	-64.32	136.94	4.20	0.06
	Group structure + Social bonds	Group size, No. affiliated ind.	4	-64.44	137.19	4.45	0.05
	Null model	Intercept-only model	2	-66.64	137.37	4.63	0.05
	Social bonds	No. affiliated ind.	3	-65.71	137.60	4.86	0.04
	Group structure + Social bonds	Group size, Max. CSI	4	-64.67	137.63	4.89	0.04
	Group structure + Social bonds	Average CSI, Group size	4	-64.74	137.79	5.05	0.04
	Social bonds	Max. CSI	3	-66.29	138.77	6.03	0.02
	Social bonds	Average CSI	3	-66.44	139.06	6.32	0.02

689 **Table 2.** Model-averaged parameter estimates and standard errors for analyses that

690 examined the occurrence of coccidian oocysts in males and females. Significant values

691 ($p \leq 0.05$) are highlighted in bold.

Sex	Parameters	Estimate ± SE	z	p
males	Intercept	0.230 ± 0.837	0.275	0.783
	Group size	0.123 ± 0.158	0.782	0.434
	Number of related individuals	-0.731 ± 0.289	2.529	**0.011**
	Number of affiliated individuals	-0.354 ± 0.205	1.729	0.084
	Maximum CSI value	-0.262 ± 0.095	2.746	**0.006**
	Average CSI value	-0.208 ± 0.170	1.226	0.220
	Age (first-year)	-2.686 ± 0.907	2.961	**0.003**
	Age (second-year)	-0.233 ± 0.562	0.415	0.678
females	Intercept	-0.816 ± 0.448	1.822	0.069
	Group size	-0.167 ± 0.101	1.650	0.099
	Number of related individuals	-0.273 ± 0.232	1.176	0.240
	Number of affiliated individuals	-0.311 ± 0.347	0.895	0.371
	Maximum CSI value	-0.076 ± 0.141	0.537	0.591
	Average CSI value	-0.145 ± 0.487	0.298	0.765
	Age (first-year)	-1.373 ± 0.508	2.704	**0.007**
	Age (second-year)	-1.269 ± 0.693	1.831	0.067