# Cockroach social network position does not influence dispersal tendency

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## Abstract

In group-living organisms, we might expect relationships between social position and dispersal, such as if certain patterns of social associations lead to dispersal or if certain personality types both link different groups and are more likely to disperse. However, directly testing these ideas is challenging as dispersal is hard to track in the field and is a difficult trait to measure in the laboratory. Furthermore, invertebrate dispersal is less commonly quantified as their small size makes them hard to track. Here we quantified social networks in multiple groups of the gregarious cockroach *Blaptica dubia* in the laboratory and related three measures of social network position (representing overall sociability, connectedness in local cliques, and connectedness to the wider network) to two different measures of dispersal. We found none of the social network measures related to either of the measures of dispersal, which were themselves not correlated. We also found that sociability and connectedness to the wider network were negatively correlated, with males falling towards the low-sociability/high global connectedness part of the continuum compared to females. We found no effects of mass, but there were differences between the two blocks our experiments were performed in, suggesting some effect of observer variability. Overall, our results suggest assays of dispersal may not effectively capture the underlying trait we hope to quantify, that cockroaches differ in the social behaviour along a syndrome integrating sociability and the tendency to move between groups, and that study replication should be encouraged so we can be confident identified trends are robust.

**Key words**: Blaptica, cockroach, dispersal, exploration, invertebrate, social network analysis

## Introduction

Organisms often disperse from their natal or breeding habitat to find resources elsewhere and avoid competition in their local environment (Matthysen, 2012). By facilitating the movement of individuals, their genetic material, and any parasites they bear (among other

things) to new locations, dispersal plays a vital role in various ecology and evolutionary

processes such as the spread of disease, population demographics, and the rate of adaptive

change (Clobert, 2012). Identifying drivers of dispersal and associations with other traits is

therefore key for understanding and predicting dispersal's importance and influence in the

natural world.

As organisms disperse to both avoid others due to competition and find others to mate with,

social interactions are likely to be key for predicting dispersal tendencies (Kurvers et al.,

2014; Louca et al., 2009; Wey et al., 2015). Animals engage in social interactions when they

mate and cooperate, but also when they fight or compete, and so essentially all animals

engage in social interactions of some sort (Székely et al., 2010). The great diversity of social

interactions in the natural world, both within and across species, makes it difficult to relate

individual social behaviour to other traits such as dispersal in a manner that facilitates the

synthesis of findings across taxa. A potential solution is to use a common quantitative

framework to analyse social interactions and then relate to traits such as dispersal. An

example of such a framework is social network analysis (SNA; Krause et al., 2007; Wey et

al., 2008). In SNA individuals are represented as "nodes" or "vertices" and are connected

with other individuals with "ties", "edges", or "links" (see: Croft et al., 2008; and: Krause et al.,

2014 for introductions). This approach is represented mathematically as a matrix, where

positive valued cells are present between individuals when they interact and 0 values are

used when they do not. We can then use the extensive suite of measures in SNA (Sosa et

al., 2021) to quantify different aspects of social behaviour, including those depending on both

direct and indirect interactions (Brent, 2015), and relate to other traits and compare across

species (Farine & Whitehead, 2015; e.g.: Sueur et al., 2011).

Although substantial information is available on how individuals dispersing into and out of

groups affects the group's social network structure (Borgeaud et al., 2017; Shizuka &

Johnson, 2020), it is less clear how the social position of individuals within the group

influences the tendency to disperse. It is conceivable that, social interactions are expected to

be related to dispersal in various ways. Negative interactions may stimulate an individual to

leave their current group or area (e.g. in the western bluebird Sialia mexicana; Aguillon &

Duckworth, 2015; Christian, 1970). In contrast, positive associations may encourage an

individual to stay (the "social cohesion hypothesis"; Bekoff, 1977; e.g. in yellow-bellied

marmots Marmota flaviventris; Blumstein et al., 2009). Meanwhile, an organism's own social

tendencies might be related to its dispersal tendency, for instance unsociable individuals

might be more willing to disperse from groups of high densities, while sociable individuals

might be more willing to disperse when at low densities (as seen in the common lizard

Lacerta vivipara; Cote & Clobert, 2007). In the males of the spotted hyena (*Crocuta crocuta*),

there is evidence that during ontogeny, network connections tend to weaken, which

eventually leads to individuals dispersing from their natal groups (Turner et al., 2018). These

relationships may differ between the sexes, for example if females compete for resources

they may disperse more at high density, while if males compete for access to females they

may disperse more at low densities (as seen in feral horses *Equus ferus caballus*; Marjamäki

et al., 2013). There may also be confounding factors for these relationships, for instance

higher body condition may effect position in a social group (for example in female reindeer

*Rangifer tarandus*; Holand et al., 2004) but also the probability of dispersal (such as in

greater flamingos *Phoenicopterus ruber*; Barbraud et al., 2003), creating the illusion of a

positive relationship between social phenotype and dispersal when in fact none exists.

Explicitly testing the relationship between dispersal and social behaviour, and how it

depends on context, is therefore essential. This is especially true in invertebrates, as their

dispersal is relatively understudied (Wey et al., 2015).

While using SNA allows us to quantitatively study social behaviour, relating social behaviours

to dispersal can be difficult as dispersal is often rare, hard to observe in the wild, and difficult

to re-create in the laboratory. Here we develop methods to quantify dispersal tendency using

two different approaches. We relate each measure of dispersal to three measures of social

behaviour based on social networks in a laboratory population of the cockroach *Blaptica*

*dubia*. We also compare the sexes and control for body mass. This allows us to measure

how different aspects of social behaviour relate to dispersal and how these traits are affected

by sex while controlling for key confounding variables.

We predict that individuals with a higher dispersal tendency will have a higher strength (a

measure of gregariousness), lower clustering coefficient (a measure of the density of only

local interactions), and higher closeness (a measure of connectedness across the entire

network). We also predict that males, which unlike females in this species have fully

developed wings (but with limited ability to fly; Kesel et al., 2009) and are likely the

dispersing sex, will show higher dispersal tendencies. Body condition could affect dispersal

in either direction; better condition individuals may have the stored resources to take on the

risk of dispersing (Barbraud et al., 2003), but alternatively poor condition individuals may

disperse to find more resources (Lowe, 2009). As such, we do not make a directional

prediction (see also the model of Bonte & De La Peña, 2009 which demonstrates how both

positive- and negative-condition dependent dispersal can evolve).

## Methods

Study population

A population of *B. dubia* was founded in the lab in Aberdeen in March 2021 by purchasing a mixed colony online. They were then maintained at 28°C, 50% humidity, and a 12:12 light:dark cycle. Stock was maintained in 48L plastic boxes (610 × 402 × 315 mm), with cardboard egg tray as shelter, carrot for hydration and Sainsbury's dog food for nutrition (approx. nutritional composition = 1527 kJ energy, 24 g protein, 12 g fat per 100 g). Boxes of immature nymphs were of mixed ages and sexes, with individuals that eclosed into adults moved to adult-only boxes within three-four days of eclosion. Some adult boxes were used for breeding and so contained both sexes, while others were for experimental animals and were single-sex. The colony showed no infection in this time and low mortality in all life stages, and so presumed to be healthy (for further details on the stock population see: Fisher, 2023). Data collection took place in two blocks with one researcher conducting the majority of the work in each, the first from 26/5/22 to 14/7/22 (led by SS), the second from 7/10/22 to 14/11/22 (led by NV). The methods were the same in each block unless we note otherwise.

Data collection

We collected data in each block in three stages, first measuring social associations to construct social networks, then assaying the cockroaches' dispersal in a group setting, before measuring dispersal tendency alone. To create groups to quantify social networks, we selected unmated adult males and females from single-sex stock boxes and placed them into single-sex groups of ten (selection of individuals was not random, as that would have required labelling each individual in the stock population, but we removed shelters from boxes before selecting individuals to avoid only selecting "bolder" individuals). We created 12 groups in the first block (six of each sex) and ten in the second block (five of each sex). The high level of replication of networks (22 in total) in a particular strength of our study; while most studies on social networks in animals only have a single or few networks, we have multiple independent networks. This replication prevents the behaviour of a few individuals influencing the social phenotypes of every individual in the study (as social network measures often depend on connections in across the whole network). We weighed each individual and, so that we could recognise individuals, for the first block marked each with a unique combination of two colours (red, green, blue, white and gold) on their wing cases using paint pens (Edding 780 Extra-fine paint markers). As we filmed the second block in darkness, here we instead marked individuals with a unique combination of dot locations (dots present/absent on head, left wing case and right wing case, with the head occasionally having 2 dots to reach a unique code for all 10 individuals) using only white paint pens. We

placed each group of ten into a plastic box (340 × 200 × 125 mm) with four cardboard egg-

tray shelters (approx. 100 x 120 mm), one in each corner taped along the long wall (see:

Fisher, 2023). We placed dog food (enough to ensure cockroaches have excess food,

around 3-5g) and carrot (for hydration, around 10g) in each box between the shelters and

replaced it each time we recorded associations. To record associations, we observed which

individuals were using the same shelter on six dates between two to four days apart (from

26/5/22 to 13/6/22 in the first block and from 7/10/22 to 19/10/22 in the second).

Observations were only roughly evenly spaced due to avoiding data collection at weekends

and other occasional scheduling constraints, but observations never took place on

consecutive days. After recording the associations of all individuals in a box they were

removed from the shelters and placed in the centre of the box, so any associations would be

re-formed each observation. In this manner we gain six snapshots of the social associations

in each group of cockroaches, much like observations of flocks of birds or herds of

ungulates. We then aggregated these data to build a single social network for each group by

assuming those in the same shelter are associated in some way (the "gambit of the group";

Whitehead & Dufault, 1999). While a single observation of two individuals sharing a shelter

does not imply they are close social associates, consistently observing them together would

allow us to infer some degree of social preference. Using the R package *asnipe* (Farine,

2013), we converted the data on co-occurrence in groups into an undirected weighted social

network, where edge weights equalled the "simple ratio index"; the number of times the two

individuals were using the same shelter divided by the total number of times they were seen

(Cairns & Schwager, 1987). This value is 1 if two individuals are always seen together and

near zero if they are very rarely seen together (it is exactly zero if they are never seen

together). We did this separately for each group, giving 22 independent social networks, 11

of each sex.

For each individual we calculated three social network measures, each representing a

different aspect of social behaviour. These measures all are weighted, and hence account

for the frequency of the associations. First was "strength", the sum of all an individual's

weighted interactions, representing overall gregariousness (function **rowSums** in the

package *Matrix*). Second was "clustering coefficient", the sum of the weighted edges

involved in *closed* triads centred on the focal node divided by the sum of all weighted edges

in triads centred on the focal node (Barrat et al., 2004; Opsahl & Panzarasa, 2009), which

represents whether individuals invest in a closed circle of associates by regularly interacting

with others who they share a mutual association with or not (function **clustering_local_w** in

*tnet*; Opsahl, 2009). Finally, there was "closeness", which is the inverse of the mean of the

weighted pathlengths between an individual and all others in the network (Freeman, 1978),

representing how well connected to all parts of the group the individual is (function **closeness** in *sna*; Butts, 2008).

After the sixth observation of associations, individuals in block 1 were immediately moved to the next stage of data collection. However, individuals in block 2 were observed a further four times (giving 10 in total). To better compare the blocks, we do not use the extra four observations in block 2 here (social network measures from networks based on six observations are either strongly or moderately positively correlated with measures from networks based on ten observations; Supplementary Materials Fig. S1). In the next stage we tested the individuals for dispersal tendency in group setting (hereafter "dispersal-group"). We removed the carrot from the groups for 24 hours in block 1 and 48 hours in block 2 to increase their need for hydration and so willingness to disperse; the change between groups was made to encourage more individuals to disperse (although this was unsuccessful as there was no difference in mean dispersal group time between the blocks, see Results). We then transferred all group members into a single large plastic box (610 × 402 × 315 mm) with the four shelters used in the social association stage placed at one end of the box, evenly spaced. At the other end of the box we placed 10g of carrot and three pellets of dry dog food (Sainsbury's Complete Nutrition Adult Small Dog Dry Dog Food, approx. nutritional composition = 1527 kJ energy, 24g protein, 12g fat per 100g) to act as a goal.  We placed each box in a room maintained at 22-26°C (although with no humidity control), with a pet heating pad placed beneath each box to raise the temperature closer to that of the stock room to encourage movement (22-24$^0$C). Above each box, at approximately 50cm, we placed an IP camera (ABUS IP video surveillance 8MPx mini tube camera) pointing directly downwards to record the box from above. We then left the room for 24 hours, with the camera recording continuously. For block 1, the lights were left on, while for block 2 the lights were turned off as, similarly to the above-mentioned difference in carrot deprivation between groups, this was to encourage more individuals to disperse, although as noted above this was unsuccessful.

We were able to record up to four groups simultaneously, and so we filmed them in batches of two to four (which means groups had different periods between the end of the recording of the social associations and the start of the dispersal-group trial). In block 1 we initially made recordings on 14-23/6/22, but due to computer error these were not useable, and so we repeated the trials on 1-6/7/22 (therefore starting at least 22 days after the final observation of social associations) and only used the data from the second run. In block 2 the recordings were made from 7-9/11/22 (therefore starting at least 18 days after the final observation of social associations). From the video recordings we extracted the number of minutes between the start of the trial and when each individual first crossed the half-way point of the

box (i.e., it left the shelter and travelled a distance of approx. 200 mm). This measure of dispersal is therefore low when individuals are quick to disperse, and high when they are slow/unwilling to disperse (with a theoretical maximum of 1440). We gave individuals who never dispersed values of NA to avoid a data distribution with a spike at high values (n = 11). We also repeated the analysis with those that did not disperse given a score of 1440; the clear results in the original analysis are maintained in the second analysis with the additional presence of a mass-dispersal relationship in males (see Supplementary Materials). After 24 hours we returned each group to the box it had recorded associations in with fresh carrot and cardboard shelters and returned them to the stock room.

For the third stage we measured a proxy for dispersal tendency while alone (hereafter "dispersal-alone") by placing each individual in a small plastic box (200 × 100 × 70 mm) with a grid of 18 rectangles (each 32 by 27 mm) drawn onto paper underneath the box's (clear) floor. In the same room as we recorded dispersal-group, again using pet heating pads to raise the temperature, we placed boxes with cockroaches in sets of four beneath the same IP cameras, switched off the lights (as *B. dubia* is more active in the dark; Bouchebti et al., 2022), and filmed continuously for 30 minutes. These cameras automatically use infrared LEDs to film in the dark, which we presume the cockroaches cannot see. We ignored the first 10 minutes of the recording as we did not want to record the individual's immediate response to being placed in a box, and instead counted the number of unique squares each individual visited in 20 minutes as a measure of dispersal tendency (counting unique areas visited as opposed to total amount of movement separates this from a measure of general activity and instead comes closer to "exploration"; Réale et al., 2007). In block 1 we made recordings on 12-14/7/22 (therefore starting at least 7 days after assaying dispersal-group), while in block 2 we made recordings on 10-14/11/22 (with trials staggered so that individuals were tested 3 or 4 days after being assayed for dispersal-group). After the dispersal-alone trials we returned individuals to the stock population. Individuals that died part way through the experiment had the values for the behavioural trials they completed retained in the dataset and NA values entered for any missing (n = 7).

## Data analysis

Our final dataset comprised of 215 individuals measured for each of the three social network traits, 199 of which recorded a non-NA value for dispersal-group, and 208 which recorded a measure of dispersal-alone. To determine how the different behaviours were associated, and if any were influenced by mass or sex, we entered the five behaviours into a multivariate mixed-effects model using function **MCMCglmm** in the R package *MCMCglmm* (Hadfield,

2010). We used each social network trait, dispersal-alone, and the log of dispersal-group as

response variables and estimated the covariances among each of them simultaneously (10

covariances in total). We included block number and sex as two-level categorical variables,

and cockroach mass as a continuous predictor, mean centred and scaled to a variance of 1

(Schielzeth, 2010). We also included the interaction between (scaled) mass and sex to test

whether the effect of mass differed between the sexes. We estimated an intercept and the

relationship with each fixed effect for each response variable, giving 25 coefficients in total.

We included the random effect of group identity, to estimate any among-group variation that

may have arisen due to emergent group dynamics and to account for differences in what the

groups experiences e.g., the slightly different periods of time between behavioural tests. We

did not estimate the among-group covariance in behavioural traits, leaving the covariance as

the phenotypic covariance, once the effect of the fixed effects and the variation in means

among groups was accounted for.

We ran the model for 550,000 iterations, discarding the first 50,000 (as parameter estimates

change most at the start of the chain), and using 1 in each 100 iterations to generate the

posterior distributions. We used a very weakly informative prior with a variance of 1 and a

degree of belief of 0.002 for the random effect and residual variance. For dispersal-alone we

used a Poison error distribution, while for the other four traits we used a Gaussian error

distribution as the raw data distributions were not too far from normality and we achieved

satisfactory convergence with these error distributions. We calculated the among-trait

correlations as the covariance divided by the square root of the product of the variances of

the two traits. We did this for each draw in our posterior distribution, allowing us to estimate

the mode and 95% credible intervals for each correlation; those with credible intervals that

do not overlap zero are considered clearly different from zero. For the fixed effects we used

the posterior distributions; those with 95% credible intervals that did not overlap zero have a

relationship with the trait in question. The author that ran the analysis (DNF) did not conduct

the behavioural trials or extract data from the video, and so was blind to any impressions

developed during those stages.

## Results

Closeness was negatively associated with strength (R = -0.643, CrIs = -0.715 to -0.540; we

show the raw correlations in Fig. 1 and the correlation posterior distribution modes and the

95% credible intervals estimated from the model in Fig. 2), and with clustering coefficient (R

= -0.270, CrIs = -0.388 to -0.117). Clustering coefficient and strength were not associated (R

= 0.010, CrIs = -0.130 to 0.150). Neither dispersal-group nor dispersal-alone were

associated with any of the social network traits (dispersal-alone – strength R = 0.046, CrIs =
-0.120 to 0.046; dispersal-alone – clustering coefficient R = -0.017, CrIs = -0.182 to 0.154;
dispersal-alone – closeness R = -0.043, CrIs = -0.227 to 0.092; dispersal-group – strength R
= -0.058, CrIs = -0.219 to 0.079; dispersal-group – clustering coefficient R = -0.067, CrIs = -
0.245 to 0.053; dispersal-group – closeness R = 0.032, CrIs = -0.137 to 0.160). The two
measures of dispersal were not associated with each other (R = -0.104, CrIs = -0.242 to
0.089). The residual variances and covariances are given in Table S1.

Figure 1. A pairs plot showing the relationship between each pair of traits. Females
are orange squares, males are blue triangles, with the statistically significant
correlations (closeness – strength and closeness – clustering coefficient) highlighted
in lighter colours.

Cockroach mass was not associated with any behavioural trait in either sex. Males had

higher closeness scores than females, and marginally lower strength scores, although this

latter effect overlapped with zero (Fig. 1). The sexes did not differ in any other trait.

Individuals in block 1 had lower strengths, higher closeness scores, and lower dispersal-

alone scores than individuals in block 2 (model results in Table 1).

Figure 2. Correlations between traits as estimated from the model., Points show the

posterior distribution mode, the horizontal lines show the 95% credible intervals. The

correlations that do not overlap with zero have been highlighted in red.

## Discussion

Our study investigated aspects of individual social network position within groups of *Blaptica*

*dubia* cockroaches and demonstrated that different measures of dispersal are not linked to

these positions. Two of the social network measures, strength and clustering coefficient,

were negatively correlated with a third, closeness, but not with each other. Meanwhile, our

two measures of dispersal were not correlated. Therefore, while we had predicted social

interactions may be associated with dispersal (with the direction of causality plausibly in

either direction, or both being affected by a third variable) we found no evidence for this

prediction. Instead, dispersal in *B. dubia* might depend strongly on other aspects of

behaviour, such as the motivation to mate and reproduce (which since all our individuals

were unmated did not vary in this study), such that the variables we did measure were

unimportant.

**Table 1**. Additional effect estimates from our statistical model. For each of the five traits

(dispersal alone, dispersal group, strength, closeness, and clustering coefficient) we give the

intercept, the effect of mass, the effect of sex (females are the default, so this effect represents

how males deviate from females), the mass:sex interaction (again, as females are the default

this effect indicates how the effect of mass on male traits differs from that for females, which

is indicated by the main effect of mass), the effect of block (block 1 is the default, and so this

effect indicates how block 2 differs), and the random effect of group. As group is a random

effect, we only report the posterior mode, mean, and lower and upper 95% credible intervals

(CrI), while as the remaining terms are fixed effects we also report the pMCMC value, which

represents two times the probability the effect is either greater or smaller than zero (whichever

is smaller), and can be interpreted in a similar way to traditional (frequentist) p-values. Terms

where the 95% CrIs do not overlap with zero are highlighted in bold.

| Variable | Posterior mode | Posterior mean | Lower 95% CrI | Upper 95% CrI | pMCMC | Effect |
|---|---|---|---|---|---|---|
| Intercept - Dispersal alone | 1.1293 | 1.0876 | 0.6622 | 1.5590 | 0.0002 | fixed |
| Intercept - Dispersal group | 4.9290 | 4.8186 | 4.0583 | 5.6829 | 0.0002 | fixed |
| Intercept - Strength | 2.3732 | 2.3380 | 1.8801 | 2.8451 | 0.0002 | fixed |
| Intercept - Closeness | 0.5342 | 0.5383 | 0.4646 | 0.6062 | 0.0002 | fixed |
| Intercept - Clustering Coefficient | 0.9498 | 0.9327 | 0.8895 | 0.9792 | 0.0002 | fixed |
| Mass – Dispersal alone | 0.0336 | 0.0876 | -0.2877 | 0.5124 | 0.6620 | fixed |
| Mass – Dispersal group | 0.0957 | 0.0411 | -0.4941 | 0.5529 | 0.8724 | fixed |
| Mass - Strength | 0.0754 | 0.0758 | -0.1602 | 0.3006 | 0.5204 | fixed |

| | | | | | | |
|---|---|---|---|---|---|---|
| Mass - Closeness | -0.0131 | -0.0142 | -0.0456 | 0.0179 | 0.3720 | fixed |
| Mass - Clustering Coefficient | -0.0010 | -0.0006 | -0.0121 | 0.0117 | 0.9256 | fixed |
| Sex – Dispersal alone | -0.0325 | -0.1244 | -1.0238 | 0.7681 | 0.7880 | fixed |
| Sex – Dispersal group | 0.2122 | 0.2178 | -1.2025 | 1.6551 | 0.7712 | fixed |
| Sex - Strength | -0.5928 | -0.6517 | -1.3758 | 0.0986 | 0.0876 | fixed |
| **Sex - Closeness** | **0.1430** | **0.1355** | **0.0324** | **0.2438** | **0.0148** | fixed |
| Sex - Clustering Coefficient | -0.0455 | -0.0450 | -0.1031 | 0.0117 | 0.1264 | fixed |
| Mass:Sex – Dispersal alone | -0.3949 | -0.1935 | -1.1410 | 0.7480 | 0.6692 | fixed |
| Mass:Sex – Dispersal group | 0.9968 | 1.0352 | -0.2887 | 2.2945 | 0.1184 | fixed |
| Mass:Sex - Strength | -0.1389 | -0.1580 | -0.7448 | 0.4142 | 0.5928 | fixed |
| Mass:Sex - Closeness | 0.0734 | 0.0586 | -0.0196 | 0.1377 | 0.1464 | fixed |
| Mass:Sex - Clustering Coefficient | -0.0157 | -0.0100 | -0.0403 | 0.0191 | 0.5028 | fixed |
| **Block – Dispersal alone** | **1.0619** | **1.0059** | **0.6806** | **1.3197** | **0.0002** | fixed |
| Block – Dispersal group | 0.4524 | 0.5098 | -0.2867 | 1.2855 | 0.2000 | fixed |
| **Block - Strength** | **0.6462** | **0.6529** | **0.1286** | **1.1858** | **0.0212** | fixed |
| **Block - Closeness** | **-0.1079** | **-0.0961** | **-0.1729** | **-0.0193** | **0.0208** | fixed |
| Block - Clustering Coefficient | 0.0217 | 0.0224 | -0.0253 | 0.0777 | 0.3840 | fixed |
| Group – Dispersal alone | 0.0013 | 0.0128 | 0.0002 | 0.0477 | N/A | random |
| Group – Dispersal group | 0.4688 | 0.6758 | 0.1722 | 1.3146 | N/A | random |

| | | | | | | |
|---|---|---|---|---|---|---|
| Group - Strength | 0.2555 | 0.3477 | 0.1277 | 0.6247 | N/A | random |
| Group - Closeness | 0.0069 | 0.0078 | 0.0032 | 0.0143 | N/A | random |
| Group - Clustering Coefficient | 0.0029 | 0.0038 | 0.0016 | 0.0066 | N/A | random |

Cockroach mass was not found to influence dispersal (or social network position) in either

sex, demonstrating either that animal condition is not a driver for dispersal or that mass is

not a good proxy for body condition in our system. There is evidence that in German

cockroaches (*Blattella germanica*) dispersal is linked to food availability and starvation (Ross

& Tignor, 1985; Tignor & Ross, 1987), which might indicate that in some cockroach

populations lack of food and hunger might initially be a driver for dispersal, but in a stable

laboratory population with ample access to resources such as ours, there is not sufficient

pressure on the lighter/poorer condition individuals to disperse. We could test this by

experimentally limiting food availability, beyond the 48 hours we used in block 2 of this study.

This manipulation would also be instructive for understanding the relationship between

mass/condition and social behaviour, as we found no relationship here. Studies on some

species, such as sleepy lizards (*Tiliqua rugosa*) and great tits (*Parus major*), have also

suggested limited relationship between body condition and social network structure (Godfrey

et al., 2013; Snijders et al., 2014), but other studies have suggested relationships between

connectedness and body mass (e.g., in prairie voles *Microtus ochrogaster*; Sabol et al.,

2020). As the number of studies using social network analysis grows, we may be able to

integrate across studies (either informally or formally as a meta-analysis) to determine the

typical relationship between mass/condition and centrality measures, and if this relationship

varies systematically across certain categories (such as females) or groups (such as

mammals) of animals.

Perhaps the most surprising result of our study is the lack of a clear relationship between our

two measures of dispersal. We had designed our assays to capture dispersal tendency in

two different contexts, but it is possible that instead we measured two different traits. One

possibility is that dispersal alone, which was quantified as searching a small arena, functions

to locate food or water, while dispersal in a group, which was assayed in a larger arena,

functions to location mating patterns. Along these lines, dispersal over longer distances in

the German cockroach (*Bl. germanica*) is often linked to searching for mates for breeding

(Ross & Tignor, 1985), which may assist with avoiding inbreeding (Lihoreau & Rivault, 2010).

In contrast, exploring the more immediate area (which is similar to our dispersal alone

assay) may serve to search for food, rather than dispersal (Durier & Rivault, 2001). Despite

the different functions, we might still have expected such similar behaviours to be positively

correlated, such as if individuals with a higher metabolic rate both moved around their

immediate environment more and travelled long distances more (Biro & Stamps, 2010). In

other species positive correlations are seen, such as in roe deer (*Capreolus capreolus*) and

a colonial spider (*Cyrtophora citricola*), where different measures of dispersal (exploration

beyond the home range and natal dispersal, and exploration of a new environment and

"tiptoeing" to release silk into air, respectively) are positively correlated (Debeffe et al., 2013;

Yip et al., 2021, p. 202). As Dochtermann (2023) highlights, we still have a limited

understanding of when and how different behavioural traits covary. Therefore, until we

generate better theoretical predictions, we will have to sift through published examples and

try to detect broader patterns.

A not-mutually-exclusive alternative to these measures of dispersal being different

behaviours is that dispersal shows such high within-individual variability that a correlation

across the two contexts (as well as across the time between the assays) was too weak to be

detected. Some personality traits in cockroaches are consistent (Arican et al., 2020; Planas-

Sitja & Deneubourg, 2018; Stanley et al., 2017), suggesting we might expect dispersal

behaviours to be consistent too. However, further data on cockroach dispersal in controlled

conditions (as opposed to in uncontrolled settings such as through apartment buildings;

Barcay et al., 1990; Crissman et al., 2010) is lacking for us to suggest why in our study we

might not have found this. In great tits (*P. major*) exploration of a novel environment is

repeatable and heritable (Dingemanse et al., 2002), and genetically correlated with dispersal

distance, yet the phenotypic correlation between exploration and dispersal is not different

from zero (Korsten et al., 2013), or is only positive for females in some seasons, and never

in males (van Overveld et al., 2014). In the case of the great tits, a negative environmental

covariance counters the positive genetic one to give the phenotypic null covariance (Korsten

et al., 2013); the same might be true in our system, but we would need further work on the

genetics underpinning cockroach behaviour to find out.

We found that males had higher closeness scores (and nearly lower strength scores) than

females. Work in another Blaberid, *Diploptera punctata*, found that females were more

gregarious and formed the core of a proximity network (Stanley et al., 2018). This result is

similar to ours, as in our study males were marginally less gregarious (lower strength) and

instead formed links with different parts of the network (higher closeness), implying they

were at the periphery of the network core (although note we did not assay male and females

together in a social network, whereas Stanley et al. did). These results suggest that males

and females may generally have different positions within cockroach aggregations, with

females possibly being more central as they are larger, and so more able to displace others
from preferred spots in an aggregation, or as they are preferred associates of both males
and other females, causing them to end up in the centre of an aggregation (Stanley et al.,
2018).

In a related result, we also found strength was strongly negatively correlated with closeness.
Therefore, within our study population there appears to be an axis of variation from high-
strength/low-closeness to low-strength/high-closeness, and females fall more towards the
first of these, and males the latter. However, the relationship is not entirely defined by sex, as
this covariance was present in our results despite modelling the effect of sex simultaneously.
We also found a negative relationship between clustering coefficient and closeness, which
likely represents a trade-off between making strong local connections with a few individuals
(giving a higher clustering coefficient) *or* making connections with a wide range of individuals
(giving a high closeness). The strong negative correlation between strength and closeness
suggests variation in social behaviour in *B. dubia*, with some individuals being less
gregarious and moving between groups, while others remaining in larger groups. Planas-
Sitja and Deneubourg (2018) found that collective aggregation formation in the American
cockroach (*Periplaneta americana*) is influenced by the social behaviour of individuals in the
aggregations, suggesting substantial differences among-individuals. Such differences in
social behaviour will likely expose an individual to a range of different selection pressures,
such as differential exposure to directly transmitted infections, variation in competition for
resources and mates, and different risk of desiccation. Understanding what drives this
variation in social behaviour and what consequences it has for the ecology and evolution of
cockroach groups are important next steps.

Our study has also demonstrated that three traits (dispersal alone, strength, and closeness)
differed between blocks, which indicates the presence of observer induced variation.
Observer-induced variation seems more plausible than an environmental change between
the boxes, as we used standardised environmental conditions. That the mean level of a
behaviour that is observed differs between blocks within a single study should be considered
in other studies and when making conclusions from single studies at single point in time by a
single laboratory, reinforcing the need for more extensive replication in such experiments
(Brecht et al., 2021; Farrar et al., 2021; Gaona-Gordillo et al., 2023).

In conclusion, our results suggest care should be taken when measuring dispersal in the
laboratory, as assays designed to capture dispersal may actually measure different traits or
not capture the consistent component of dispersal. We also found no support for the
suggestion that social behaviour and dispersal are associated. We found limited importance

of mass in our system, while there were some differences between males and females that reflected broader trade-offs in social behaviour. Finally, our results indicate that repeating experiments, even in the same lab, can give different results, and so add strength to the call that study replication should be an important component of animal behaviour research if we are to detect general pattens.

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No specific funding was received for this project.

## Conflict of interest
We have no conflicts of interest.

## Ethical approval
No specific approval was required for this work. We did follow the Association for the Study of Animal Behaviour's "Guidelines for the ethical treatment of nonhuman animals in behavioural research and teaching", notably using a non-invasive marking system (using non-toxic paint on the outer carapace) and performing no other procedures. Our level of replication, both number of groups and individuals per group, was high enough to detect clear trends but not unnecessarily large. Our study species is naturally aggregative and so is not stressed by being kept in groups, and periods kept alone were short (30 minutes). We used short periods (24 or 48 hours) without access to food to increase motivation to disperse, but all animals were given access to food in the subsequent behavioural trial and when returned to housing afterwards. At the end of the experiments individuals were returned to the stock population.

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