Social network centrality predicts dietary decisions in a wild bird population

Keith McMahon[1]*, Nicola M. Marples[2], Lewis G. Spurgin[3], Hannah M. Rowland[4,5], Ben C. Sheldon[1] &
Josh A. Firth[1]*

[1]Department of Biology, Oxford University, Oxford, UK

[2]Department of Zoology, School of Natural Sciences, Trinity College Dublin, Eire

[3]School of Biological Sciences, University of East Anglia, Norwich, UK

[4]Max Planck Institute for Chemical Ecology, Jena, Germany

[5]Department of Zoology, University of Cambridge, Cambridge, UK

*Correspondence: Keith.McMahon@biology.ox.ac.uk & Josh.Firth@biology.ox.ac.uk

SUMMARY (150 words)

How individuals balance costs and benefits of group living remains central to understanding sociality.
In relation to diet, social foraging provides many advantages but also increases competition.
Nevertheless, social individuals may offset increased competition by broadening their diet and
consuming novel foods. Despite the expected relationships between social behaviour and dietary
decisions, how sociality shapes individuals' novel food consumption remains largely untested in
natural populations. Here, we use wild, RFID-tracked, great tits to experimentally test how sociality
predicts dietary decisions. We show that individuals with more social connections have higher
propensity to use novel foods compared to socially-peripheral individuals, and this is unrelated to
neophobia, observations, and demographic factors. These findings indicate sociable individuals may
offset potential costs of competition by foraging more broadly. We discuss how social environments
may drive behavioural change in natural populations, and the implications for the causes and
consequences of social strategies and dietary decisions.

KEYWORDS

Social Networks, Diet, Social Behaviour, Foraging, Social Structure, Competition.

INTRODUCTION

Groups of foraging animals benefit from sociality in many ways [1], ranging from social foraging
allowing complex cooperative hunting [2] to other benefits such as learning about how food is
distributed or avoiding predators [1-5]. It is also likely that social learning may help individuals
decide which new foods to explore and consume [6-8]. Indeed, novel foods pose a challenge to
foragers because they can differ from familiar foods in their nutritional quality, and may also contain
unfavorable chemicals or defensive toxins [10]. How individual foragers in the wild differ in their
propensity to explore and use novel foods and how individual sociality affects foraging decisions
about novel foods remains a central topic in understanding dietary decision making [10].

As well as many benefits, foraging in groups also imposes costs through intraspecific competition
[11]. When resources are limited, there is greater competition between group members [12] and

this can result in carry-over costs through 'interference competition' e.g. fighting for resources, with
attendant costs in time and risk of injury [13]. This interference competition can reduce the time
available to an individual to make decisions about which food items to consume, and can reduce the
number of profitable food items encountered while foraging [14]. Exploiting new food sources, in
this situation, can alleviate within-group competition through niche partitioning [15-17].

When faced with decisions about which new foods to explore and consume, many species exhibit
dietary wariness [18, 19]. Dietary wariness is comprised of two behavioural processes: neophobia
and dietary conservatism [20]. Neophobia describes the initial fear/apprehension of novel objects or
foods, and is observed in many animal groups, including fish, mammals and birds [19, 21, 22]. This
aversion is usually brief, and is followed by investigation of the novel food or object [23]. Once
neophobia has waned, some individuals within a population continue to avoid novel food long after
the initial exposure, and this is termed dietary conservatism [20]. Dietary conservatism (DC) is a
spectrum: individuals differ in their willingness to consume novel food. Adventurous consumers (AC)
are those individuals which show little or no hesitation in consuming novel food once neophobia has
passed, while conservatively foraging individuals continue to avoid the novel food for extended
periods [24]. Following this general classification, dietary conservatism has been observed in a wide
variety of species, particularly in various studies in birds [22, 25] and fish [26-28]. It is important to
consider individual differences in the propensity to eat novel foods when discussing the strategies
that animals use to mitigate resource competition during social foraging, as this is of direct relevance
for the study of the factors that shape sociality and resource acquisition [29] .

The advent of animal tracking technologies has revolutionised our ability to observe individuals'
social foraging associations in the wild [30], and animal social networks have now been quantified
across a range of animal systems [31]. Social network analysis provides a framework for quantifying
variation in intraspecific sociality [32] and allows the estimation of various metrics of individuals'
social behaviour [33]. This provides fine-scale information about an individual's own social
associations, as well as the wider social environment they inhabit [34], who they associate with,
when they associate with them and where the associations happen. The ability to quantify social
networks within the wild, while simultaneously tracking individuals' foraging behaviour, presents the
opportunity to determine empirically how intraspecific differences in sociality relate to the various
aspects of dietary wariness in natural settings.

In this study, we use novel food experiments to test individual-level dietary wariness in a RFID-
tracked social system of wild great tits (*Parus major*). Using this approach, we are able to examine
dietary wariness and novel food usage independently, and use social network analysis to determine
how individual sociality predicts individuals' foraging decisions. A priori we predicted that there
would be variation in the use of novels foods among foraging great tits. This variation would be
driven in large part by underlying propensities to consume novel food, dietary wariness, but also
that social network position would have a major role in influencing decisions made by foraging birds.
Thus, we were able to directly test the prediction that more social individuals have a greater
propensity to eat novel foods whether that be due to the need to mitigate the potential costs of
interference competition or because they have more access to information about the profitability of
novel food sources. These potential explanations need not be mutually exclusive. As a consequence
these more social, more informed individuals should show lower levels of dietary wariness
compared to less social individuals. We are able to separate out other elements (observation-related
factors and demographic traits) when assessing the relationship between foraging decisions and a
suite of social measures. We discuss the implications of these experimental results for
understanding how competition shapes social foraging, and the wider insights this may offer into the
interplay between individual-foraging decisions and social behaviour.

METHODS

Study System

Wytham Woods, Oxford, United Kingdom (51° 46' N, 1° 20' W) is home to a long-term study population of great tits [35]. These birds are captured and tagged with British Trust for Ornithology rings (as adults and as nestlings) during the spring as they breed in the intensively-monitored nest boxes [35], and immigrant birds are captured during the winter during regular mist-netting sessions throughout the woodland. As well as recording standard morphological information during capture, since 2007, all captured great tits have also been fitted with radio-frequency identification (RFID) tags. Each RFID tag possesses a unique ID code which allows automated recording of the times and locations of individuals' occurrence at feeding stations over the winter. Each feeding station is equipped with an RFID antenna which successfully records>99% of RFID tagged individuals visits to feeders[36, 37]. These RFID feeding stations allow the recording of individual feeder usage (see Methods: Experiment data) and also the inference of flock structures and arising social networks (see Methods: Social network data). The antennae scan for RFID-tagged individuals 16 times per second from pre-dawn until post-dusk (i.e. over the entirety of the great tits' foraging hours).

The study was conducted at two separate sites within Wytham Woods, approximately 1km apart. Within the timeframe of the study, this 1km distance between sites effectively ensures two separate local populations; of the 105 birds recorded as part of this study, only one individual was observed at both sites (see supplementary information for details). Previous work has estimated that >80% of locally-occurring great tit individuals are RFID tagged [9].

Social Network Data

Prior to beginning the experimental trials, we gathered detailed baseline information regarding individuals' usage of a familiar food, and their social connections to one another. From 11/01/2018 to 22/01/2018 an RFID feeding station containing non-coloured granulated peanut was placed at each site. Granulated peanut is a familiar food source which is commonly used by great tits in Wytham Woods and the surrounding area, as well as throughout the UK [38]

Each RFID station automatically recorded the unique identity of each individual detected along with the associated time-stamp. Because these birds forage in loose fission-fusion flocks [36], this produced a temporal data stream made up of bursts (as flocks arrive and feed) interspersed with intermittent quiet periods [39, 40]. These bursts of activity (the flocking events) were detected automatically (without the need for subjective specifications) using a Gaussian Mixture Model (GMM – an unsupervised learning algorithm) [40] which returns a group-by-individual matrix [32] specifying which individuals were detected within each of these flocking events. Following this, social networks can be derived for any desired time period by applying the widely used 'Simple Ratio Index' (SRI) [41] to the 'groups' (i.e. flocking events) observed within that time period, derived as a proportion of flocking events in which the focal dyad (A and B) were seen together as $Flocks_{A,B}/(Flocks_A+Flocks_B-Flocks_{A,B})$, where $Flocks_A$ is the number of flocks that individual *A* was seen in, irrespective of the observation of *B*. In this way, a weighted, symmetrical, social network was produced for all three periods of the study (baseline, experiment 1, and experiment 2).

In these social networks, the individuals are represented as the network 'nodes', and the social connections between them as the network 'edges', and the weight of these edges are the dyadic

association scores (as specified in the dyadic association matrix). These weights denote the strength
of the social affiliation between each of the dyads [40].

This approach to calculating social networks has been extensively used for this population and
methodological examination of this system has found that the GMM approach outperforms other
potential methods of identifying associations [39, 40]. Large-scale observational studies have shown
that the derived social networks are consistent across time [42] and contexts [37], and linked to
other processes such as mating [43, 44], territory acquisition [37, 45], and information flow [46].
Furthermore, detailed experimental tests have confirmed the social network's consistency [47, 48],
and its relation to biologically meaningful outcomes [49-51].

We quantified individuals' social network centrality from the weighted social networks. The most
common and intuitive metric of social network centrality is weighted strength, which is the sum of
the focal individual's social connections to all other individuals. We also calculated two other
measures of social centrality, namely *(i)* 'average edge weight' which measures the typical strength
of an individual's social bonds by taking the mean weight of their non-zero dyadic social association
scores, and *(ii)* 'eigenvector centrality' which measures their position within the wider network by
summing the social connections of their associates, and thus represents the sociability of their social
associates.

As well as computing these social metrics, we also calculated for each individual the mean size of the
flocking events (i.e. the grouping events automatically identified from the feeder co-occurrence
records) they occurred in (i.e. their average group size), and the number of unique individuals they
were seen with, across all observations. In this way, we were able to separate the influence of
individuals' social network metrics from simpler social measures (see Methods; Statistical Analysis).

Experiment Data

Each of the experimental trials were carried out after 12 days of baseline data collection. The same
general protocol was used at both sites. The first novel food experimental trial took place
immediately after the baseline data collection. The single RFID feeder (containing familiar food) was
swapped for two RFID feeders at either side of the original feeder position, within 1m of one
another. One of these RFID feeders contained the familiar peanut granules, while the other feeder
contained peanut granules which were made novel by dying them either green or red, under
standardised methods, see [52] for details. Both feeders were made of transparent plastic to allow
the birds to see the colour of the food This experimental trial ran for four days, recording all visits by
RFID tagged birds to each of the feeders.

Following this first experimental trial, a second novel food experimental trial was then carried out, in
which the feeder containing the novel food was swapped to contain different coloured novel food.
In the first experimental trial, the novel food RFID feeder at Site 1 was filled with red-dyed granules
while the novel food RFID feeder at Site 2 contained green-dyed granules. In the second
experimental trial. this was switched so that the novel food RFID feeder at Site 1 was filled with
green-dyed granules while the novel food RFID feeder at Site 2 contained red-dyed granules. In both
trials, familiar coloured food was provided in the other feeder at each site. The second experimental
trial was carried out for four days (the same length as the first trial), and all visits by RFID tagged
birds to the feeders were recorded.

During the experimental trials, we also aimed to reduce any additional influences on the birds'
feeding behaviour that may be caused by either human presence causing disturbance, or through
positional effects of feeder placement. We ensured that all required activity at the feeders (i.e.
placement changes and associated device checks) were carried out when the great tits were not
using the feeders (i.e. after dusk). Even though the familiar-food feeder and the novel food feeder
were next to one another (>1m apart), we also aimed to reduce any remaining fine-scale positioning
effects by swapping the feeders' positions every other day during the experiment (see Table S1).

Statistical Analysis

Novel Food Usage

For each of the experimental trials, we examined how prior social centrality (i.e. their network
centrality before the experiments began) was related to subsequent usage of novel food during the
trials. As we aimed to consider individuals' relative use of the novel food, rather than just their total
feeder use in general, we treated the proportion of their total activity which took place on the
feeder containing novel food as a measure of individual propensity to use novel food. Therefore, we
carried out logistic regressions for each of the trials separately, whereby the response variable in the
generalised linear model (GLMs) was set as a binomial variable with the number of detections on the
novel food feeder as 'successes' and the number of detections on the familiar food feeder during the
trial as 'fails'. In this way, the total feeder usage, and also confidence in their propensity (i.e.
strength of their bias/preference) to use novel vs familiar food, was considered directly within the
response variable. Because GLMs with binomial error-distributions are vulnerable to over-dispersion,
we used a quasi-binomial error distribution, which removed this issue of over-dispersion. The
models were set to include fixed effects of the factors that could potentially be related to individual
novel food usage propensity. We specified the primary explanatory variable of interest as social
network centrality (weighted strength) prior to the experimental trial. For each trial, the social
centrality used for the analysis were derived from the period immediately before the trial. As such,
for the model assessing the novel food usage during the first trial, individual social centrality
calculated from the network directly before the trial began (i.e. during the baseline data collection
period) was used. For the model assessing the second trial, weighted strength during the period
directly before the second trial began (i.e. during the first trial data collection period) was used. We
also aimed to account for other variables that may affect novel food usage, and included site (i.e.
which of the two areas the individual was detected in), sex (whether they were male or female), age
(specified as either adult, or juvenile), and immigratory status (whether they had arrived in the
Wytham Woods study area that year or not) as explanatory variables in the model. In order to
directly consider individual differences in feeder usage, we also included the number of detections
on the feeders in the period prior to the experimental trial.

Although social centrality was set as weighted network strength in the main models, we also
quantified it using other commonly used network metrics (see Methods *Social Network Data*).
Therefore, we ran supplementary models using other common measures of social network centrality
(average edge weight, eigenvector centrality) calculated from the period prior to the trial, while the
rest of the model structure remained the same. Furthermore, it is also possible that other more
basic measures of sociality (i.e. non-network based metrics) might act as potential explanatory
variables (see Methods *Social Network Data*). To test this, we ran the same models again using each
of the simple individual-level social metrics (i.e. not based on networks) obtained from the period

prior to the experimental trial (average size of the flocking events they were observed in, and
number of unique individuals they were seen with).

Neophobia

Individual variation in the observed usage of novel food in the experimental trials could potentially
be due to differences in the propensity to first approach the novel food (i.e. avoidance/neophobia)
rather than variation in propensity to use the food once any potential neophobia is overcome. We
considered this directly by employing the same models as described above, but instead of setting
proportional novel food usage as the response variable, we used a binary variable of whether or not
they were detected on the feeding perch of the novel food feeding station when they first arrived at
the feeding site during the experimental trial. We used a GLM with a quasi-binomial error
distribution, and fitted the same fixed effects of the main models (prior social centrality, individual
sex, age, immigrant status, experimental site, and prior number of feeding detections).

Another measure of neophobia is the latency to first approach the novel food (as opposed to the
'likelihood of using the novel food upon the first visit' as described above). Therefore, we also
calculated two related temporal measures of individual novel food neophobia; 'time to use the novel
food since the experiment began', and 'time to use the novel food since first detected during the
experimental trial'. We set each of these in turn as response variables in the same model structure
as described above, but using a gaussian-error distribution instead of binomial due to the
distribution of these response variables.

After modelling how the explanatory variables were related to measures of novel food neophobia,
we also re-assessed the models examining novel food usage propensity but only considering
individuals' behaviour once any neophobia had been overcome. We re-calculated each individual's
proportional usage of novel and familiar foods but this time only within the time-period following
their first detection on the perch of the novel food feeder. Following the primary model structure,
we fitted this as the response variable in a GLM with binomial error-structure (with the novel food
usage as 'successes' and familiar food usage as 'fails'), along with the explanatory variables (as
stated previously) to examine how this predicted novel food usage once individuals had already used
the novel food feeder. We additionally evaluated how model structure related to the observed
results using randomisations (see below).

Network Randomisations

Individuals' positions within social networks are dependent on one another [32]. Social network
data, by definition, violates the assumption of the independence of data points made under the
standard maximum likelihood statistical tests. Therefore, network randomisations are commonly
used when estimating the statistical significance of observed parameters computed from standard
tests [32]. Such randomisation techniques allow the creation of null models using a given
permutation procedure, and from these null models the same parameters can be re-calculated using
the permuted data (instead of the observed data) to provide the distribution of this parameter that
is expected given the underlying network structure, and the non-independence of data. More
broadly, null models based on permutations of the observed data can also act as an additional, and
intuitive, test of significance of observed statistics across various contexts. We employed a
hierarchical node attribute permutation procedure controlling for space and time [37] whereby
individuals were randomly reassigned the attributes (response variable of consideration) of another
node individual in the same area during the same period of consideration as themselves. Following
this, we re-ran the models and stored the estimated effect size (Coefficient) of each of the predictor

variables on the permuted response variable, while keeping everything else in the model the same
(i.e. maintaining the exact distributions of all the variables, and the covariance between the
predictor variables). By running 10,000 of these permutations, we generated the null distribution of
the effect size parameter for each model's predictor variables and calculated the significance of the
observed data test statistics by comparing it to these null distributions. In this way, the p-value (p_{rand})
represents each observed statistic's position within the corresponding null distribution, whereby
$p_{rand}<0.05$ indicates that the observed statistic lays outside of the 95% range of the null distribution
for this predictor variable (i.e. below the bottom 2.5% or above the top 97.5%, i.e. it detects a
significant effect).

RESULTS

During the study, 105 unique RFID tagged great tits were detected: 85 during the baseline data
collection period, and 75 and 61 in the first and second experimental trial, respectively. The average
number of detections of each RFID tagged individual over the 19-day experiment was 3234±409
(mean±SE), with a total of 210,579 detections of all individuals during the baseline period and
60,727 and 68,311 during the first trial and second trial, respectively. We detected 2393 flocking
events for the baseline period, and 767 and 764 for the first and second trail, respectively. The
typical group size (i.e. group size encountered by the average individual [53] was 6.8±0.03. The social
networks inferred from these flocking events (see Methods) were relatively dense networks within
sites (Figure 1), with a total number of unweighted social network connections of 1266 in the
baseline period and 892 and 697 for the first and second experimental trials, respectively.

Social Centrality and Novel Food Usage

An Individual's propensity to use novel food during each of the experimental trials was significantly
predicted by their prior social centrality (Figure 2): the GLMs showed a strong relationship between
proportion of novel food usage and the individuals' prior weighted strength for both trials (Trial 1 -
Table S2a: Coefficient= 0.529±0.235, t=2.25, p=0.028, p_{rand}=0.01. Trial 2 - Table S2b: Coef=
0.467±0.150, t=3.11, p=0.003, p_{rand}=0.012). None of the other individual characteristics in the models
(age, sex, immigrant status, previous feeder usage) were significant predictors of novel food usage
(Table 2). The first experimental site had a strong colour preference for red over green when each
colour was novel (Figure 2), the first trial had a reduced novel food usage for site 1 (initially using
green novel food) over site 2 (using red novel food) and the reverse effect for the second trial when
the novel food colours were swapped (Trial 1 - Table S2a: Coef= 3.40±0.71, t=4.8, p<0.001. Trial 2 -
Table S2b: Coef= -1.59±0.27, t= -5.96, p<0.001). This apparent effect of colour preference persisted
through all of the models (See Supplementary Tables).

Supplementary analysis which considered two alternative measures of centrality ('average edge
weight' and 'eigenvector centrality') confirmed the findings that prior social network position
significantly predicted novel food usage. (Figure S1; Table S3-S4). The average edge weight was
significantly related to the proportion of novel food usage across both trials (Trial 1 - Table S3a:
Coef= 16.5±7.2, t=2.28, p=0.027, p_{rand}=0.022. Trial 2 - Table S3b: Coef=15.9±4.5, t=3.5, p=0.001,
p_{rand}=0.004) as was eigenvector centrality (Trial 1 - Table S4a: Coef=1.74±0.76, t=2.29, p=0.026,
p_{rand}=0.05. Trial 2- Table S4b: Coef=1.71±0.58, t=2.93, p=0.005, p_{rand}=0.012). A further line of

supplementary analysis confirmed the importance of using network centrality as a robust measure
of sociality, as novel food usage was not significantly related to more basic social measures (Figure
S2; Table S5-S6) that simply quantified an individual's average flock size (Trial 1 - Table S5a; Coef=
0.049±0.17, t=0.29, p=0.77, p_{rand}=0.68. Trial 2 - Table S5b; Coef=-0.14±0.15, t=-0.92, p=0.36,
p_{rand}=0.23) or their total number of flock mates (Trial 1 -Table S6a; Coef=0.036±0.048, t=0.75,
p=0.45, p_{rand}=0.18. Trial 2 - Table S6b; Coef=0.0565±0.0385, t=1.47, p=0.15, p_{rand}=0.10.)

Novel Food Neophobia and Social Centrality

The majority of individuals (92%) recorded during the experimental trials were detected on the novel
food feeder during the trial, indicating that complete neophobia (unwillingness to try the novel food
at all) was extremely rare. Furthermore, 95% of those that were detected using the novel food
feeder during the trial were recorded using it on the first day of the trial, again indicating that
neophobia generally was not a persistent barrier to novel food usage.

However, we also aimed to examine whether any individual variation in initial avoidance of the novel
food (i.e. neophobia) was related to individuals' network position. By using the very first record of
each bird during the experimental trial, we found that whether or not individuals landed on the
novel food feeder when they first arrived at the experimental trial was not significantly related to
social network centrality in either the first trial (Figure 3a – Coef=0.29±0.41, t=0.7, p=0.48,
p_{rand}=0.29, Table S7a) or the second trial (Figure 3b - Coef=-0.55±0.57, t=-0.96, p=0.34, p_{rand}=0.28,
Table S7b). Although only 30% of individuals immediately tried the novel food when first arriving at
the experimental trials, none of the individual characteristics included in the GLM were predictive of
which individuals landed on the novel food feeder as they first arrived during the experiment (Table
S7).

In line with this result, supplementary analysis also showed that network strength was not related to
the amount of time taken for each individual to first land on the feeding perch of the novel food in
each trial (Table S8;S9). This was true when time was quantified as the time of day they were first
recorded on the novel food (Table S8), or when quantified as the total elapsed foraging time since
they were first detected at the site during the trial (Table S9).

As a direct assessment of whether the relationship between sociality and proportional usage of
novel food exists regardless of any neophobia, we also found that prior network strength
significantly predicted the proportion of novel food (over familiar food) that individuals used after
they had first tried the novel food feeder (Figure 4; Table S10) i.e. after any neophobia was
overcome and only dietary conservatism was active. Again, this was true for both the first trial
(Figure 4a - Coef=0.55±0.25, t=2.15, p=0.037,p_{rand}=0.006, Table S10a) and second trial (Figure 4b -
Coef=0.53±0.168, t=3.17, p=0.003,p_{rand}=0.010, Table S10b), and the site/colour preference effect
was again evident (Figure 4;Table S10).

DISCUSSION

By quantifying wild great tit social networks, and tracking their behaviour in experimental trials
aimed at testing dietary decisions, we found that individuals' prior social network centrality
predicted their subsequent propensity to use novel food, and that this was unrelated to various
measures of neophobia. This link between individual sociality (as measured as social network
position) and dietary decision making has important implications for understanding how different

behavioural strategies influence resource acquisition, and for understanding the emerging
consequences of these strategies and decisions [16].

Through measuring the sociality of individuals using social network analysis, we were able to
quantify the individual level of sociality with this generalisable and powerful approach [54], and
measure individuals' own social propensity and experienced social environment using metrics that
take into account individuals' own general sociality ('strength' Figure 2), as well as their average
bond strength to others ('edge weight' Figure S1a;b), and the social centrality of their associates
('eigenvector centrality' Figure S1c;d). This approach outperforms simpler methods of attempting to
measure sociality [55] when making wider inferences e.g. using estimates of group size (Figure
S2a;b) or number of group members (Figure S2c;d). Here, the positive relationship between network
centrality and novel food usage in this free-living system of socially foraging individuals supports the
prediction that competition in social environments can predict foraging strategies in natural
populations. Specifically, individuals that are very social may be able to offset the competitive costs
of reduced resources by using other food sources. Furthermore, our approach allowed us to
demonstrate that this effect of prior social network centrality on subsequent novel food usage was
unlikely to be due to more social birds just generally being more exploratory or less averse to
approach novel-looking food, as there was no significant relationship between individuals' social
centrality and their delay in approaching the novel food. Thus, it would appear that highly social
great tits which experience a more competitive social environment (i.e., by having more social ties)
may be alleviating the costs of competition by foraging more broadly. These findings could be
explained by optimal foraging theory [56-59] which states that individuals' foraging decisions should
maximise their net rate of energy intake given their environment, as the more socially central great
tits (i.e., those experiencing a more competitive social environment) are more likely to expand their
diets by consuming novel food .

Our findings that birds showing the highest degree of dietary conservatism (i.e. those eating least
novel food) held the least central network positions, may suggest that these individuals generally
experience a reduced level of competition compared to those willing to eat the novel food. In a
previous study investigating dietary conservatism and competition in wild-caught captive blue tits
(*Cyanistes caeruleus)* [52], a high proportion of individuals displayed a strong aversion to novel food
when foraging alone, i.e. dietary conservatism, preferring to forage only on familiar food. However,
with the introduction of a second individual this aversion was quickly overcome, resulting in
consumption of novel food. These findings support our own finding which suggest that less social
individuals may not experience the same level of competition felt by their more social conspecifics,
and thus they may not have to resort to exploiting a novel food source in order to forage efficiently.
Furthermore, it could also be argued that these conservative foragers were demonstrating resource
partitioning behaviour i.e. reducing dietary overlap with their more social counterparts by excluding
this novel food source from their diet [60]. This kind of behaviour has been demonstrated in other
social species, where 'specialised' individuals exist. For example, Sheppard, Inger et al. [17] found
that individual banded mongooses (*Mungos mungo*) with substantially narrower resource niches
compared to other members of their social group experienced reduced intraspecific competition
through niche partitioning. There are many other instances where individual resource-level
specialisation has been examined [15, 16, 61] and where competition has been implicated in driving
this resource use variation among individuals of the same population [62]. Here our results suggest
that a potential mechanism regulating these kinds of processes might be linked to social network
position. Those more conservatively foraging individuals with fewer foraging social ties, and

therefore experiencing a less competitive social environment, do not have to risk expanding their
diet to include foods of unknown profitability. In contrast, those with more social ties and therefore
more potential competitors can expand their diets and include novel food to mitigate the potential
effects of any intraspecific competition for themselves.

As well as competition-driven effects, social information may play a role in foraging decisions.
Indeed, information about the profitability and nutritional value of the food may be transmitted to
group members through social facilitation [63] or local enhancement [1]. For example, prior work
within our great tit population has demonstrated that individuals use social information to locate
new foraging locations, and that more central individuals are most likely to learn the location of new
resources faster [64, 65]. The relationship between social centrality and information also appears in
other species, such as the acquisition of information in social groups of wild baboons (*Papio ursinus*;
[66]. In this study, it is unlikely that social network position shaped the propensity for individual
great tits to find the novel food (as it remained in a set location), and also unlikely that it shaped
their propensity to try the novel food (as there is no significant relationship between social centrality
and timing of using the novel feeder). However, it may be the case that social influence potentially
played a role in the extent to which individuals exploited the novel food following discovery of it. For
instance, McMahon, Conboy et al. [67] showed that conservatively foraging domestic chicks (*Gallus
gallus domesticus*) were more willing to consume novel food when they were able to see
conspecifics consuming novel food, essentially treating conspecifics as sources of social influence. As
such, more socially central great tits may be more likely to be socially associated with others using
the novel food resource (simply due to having more social ties) and thus more likely to increase
usage of the novel food themselves. On the other hand, less social individuals with fewer social ties
may experience less social influence for using the novel food (due to having fewer links to others in
general). Indeed, individuals displaying this dietary conservatism may be simply more efficient at
exploiting foods with which they are familiar and therefore remain more rigid in their foraging
decisions. [68]. Others have also shown that individuals with higher network centrality may tend to
have a more proactive personality [9] and that these individuals could also be important in the
spread of information because they move more between groups [69].

The positive effect of social centrality on novel food exploitation may also have consequences for
considering the evolution of conspicuous prey defences. Novel conspicuous prey are expected to
suffer high initial attack risk from naïve predators [21, 70] and our results support this idea:
Adventurous consumers foragers `attacked` novel food. This makes the initial evolution of
conspicuous prey types paradoxical. However, just as birds use social information to find novel food
(as specified above) it is also known that individuals transmit social information about prey defences,
which can aid the evolution of aposematism [71, 72] . If more social individuals attack novel prey (as
suggested here), they may play a larger role in providing information about prey defences for others,
and this could influence how social information spreads in the predator population. Further research
into the fine-scale interactions between dietary wariness (or specifically lack of wariness), sociality,
social avoidance, and social learning, may be valuable for understanding the evolution of
conspicuous prey types.

Finally, an additional finding of our experiments was a strong preference for red novel food over
green novel food. Colour preferences have been reported previously in relation to dietary decision
making, such as captive blue tits and great tits preferring red food over green food [73], and captive
domestic chicks (Gallus gallus domesticus) generally preferring red coloured food over green
coloured food [74] but with other colour preferences varying depending on the types of foods
offered [75] or experiences prior to being given a colour preference test [76].

Conclusion

By monitoring wild great tit activity at experimental feeders, we showed that socially central
individuals are more likely to use novel food than less social individuals. This finding suggests that
individuals experiencing a more social associations may be more likely to incorporate novel food
resources. Our results also suggest the relationship between social centrality and novel food usage is
unlikely to be due to individuals' initial aversion to first using a new food resource, meaning that
variation in exploratory behaviour, or differences in access to new social information, appear to be
improbable drivers of link between sociality and novel food usage. Future research into how social
influence over longer-time periods than initial discovery/aversion may govern novel food usage over
time will be useful for revealing how social mechanics may shape these relationships. Further
expansions of the investigation into social behaviour and individual dietary decisions may also
provide insights into topics such as the competition, foraging, sociality, and even the evolution
conspicuous prey defences.

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Main Text Figures and Legends: Social network centrality predicts dietary decisions in a wild

bird population

Figure 1. Graphics of the social networks at experimental site 1 (Figure 1a) and experimental site 2
(Figure 1b) in the baseline period. The nodes (points) represent the individuals and the edges (lines)
show the social connections between them. The size of the nodes and their shading indicates an
individual's network centrality (large dark nodes = high strength, small light nodes = low strength)
and are positioned using spring layout forced into a best-fit filled circle. The thickness of the lines
shows the weighted social bond between dyads where thick lines indicate strongly connected
individuals and thin lines show weak connections (the edge thickness is standardised by total sum of
social connections with the network). Although site 1 (Figure 1a) appears to be denser than site 2
(Figure 1b), the actual network densities (percentage of potential realised links) are very similar (70%
and 63% respectively), and the main visual difference comes from higher number of individuals in
site 1 (nodes=52) resulting in more connections (connections=931) than site 2 (nodes=33 ,
connections=335).

Figure 2. Prior social centrality (network strength – x axis) and subsequent novel food usage
(proportion of novel food usage – y axis) for the (a) first trial, and (b) second trial. The point positions
show the individual data points, point colour shows the colour of the novel food (red or green dyed
peanut), point shape shows which experimental site the individual was at (site 1 round, site 2
square), and point size indicates weight of the data point i.e. the total number of detections (at both
the novel, and familiar food feeder). The lines show the GLM fit, and the surrounding polygons show
the associated standard error around this estimate, with the red lines showing the fit for the red
novel food site, the green line showing the fit for the green novel food site, and the black line
denoting the overall fit. See Table S2 for full model details.

Figure 3. Prior social centrality (network strength – x axis) and probability of individuals trying the novel food upon arrival at the (a) first trial, and (b) second trial. The point positions show the individual network strength and whether they immediately tried the novel food (top) or not (bottom), point colour shows the colour of the novel food (red or green dyed peanut), and point shape shows which experimental site the individual was at (site 1 circles or site 2 squares). The lines show the GLM fit, and the surrounding polygons show the associated standard error around this estimate, with the red lines showing the fit for the red novel food site, the green line showing the fit for the green novel food site, and the black line denoting the overall fit. See Table S7 for full model details.

Figure 4. Prior social centrality (network strength – x axis) and subsequent novel food exploitation (proportion of novel food usage – y axis) after they had overcome any neophobia for the (a) first trial, and (b) second trial. The point positions show the individual data points, point colour shows the colour of the novel food (red or green dyed peanut), point shape shows which experimental site the individual was at (site 1 or site 2), and point size indicates weight of the data point i.e. the total number of detections (at both the novel, and familiar food feeder). The lines show the GLM fit, and the surrounding polygons show the associated standard error around this estimate, with the red lines showing the fit for the red novel food site, the green line showing the fit for the green novel food site, and the black line denoting the overall fit. See Table S10 for full model details.