Social foraging in vampire bats is predicted by long-term cooperative relationships

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17 Abstract

18 Stable social bonds in group-living animals can provide greater access to food. A striking example is that female vampire bats often regurgitate blood to socially bonded kin and 19 nonkin that failed in their nightly hunt. Food-sharing relationships form via preferred 20 21 associations and social grooming within roosts. However, it remains unclear whether these 22 cooperative relationships extend beyond the roost. To evaluate if long-term cooperative 23 relationships in vampire bats play a role in foraging, we tested if foraging encounters 24 measured by proximity sensors could be explained by wild roosting proximity, kinship, or rates of co-feeding, social grooming, and food sharing during 22 months in captivity. We 25 26 assessed evidence for six hypothetical scenarios of social foraging, ranging from individual to collective hunting. We found that female vampire bats departed their roost individually, but 27 28 often re-united far outside the roost. Nonrandomly repeating foraging encounters were 29 predicted by within-roost association and histories of cooperation in captivity, even when controlling for kinship. Foraging bats demonstrated both affiliative and competitive 30 interactions and a previously undescribed call type. We suggest that social foraging could 31 have implications for social evolution if 'local' cooperation within the roost and 'global' 32 competition outside the roost enhances fitness interdependence between frequent 33 34 roostmates. 35

36 **Short title:** Social foraging in vampire bats

37 Introduction

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Socializing and foraging are two key determinants of reproduction and survival that can 39 influence each other in several interesting ways. Preferred social relationships can drive 40 41 foraging decisions (e.g. great tits: Firth et al. (2015)). Conversely, shared foraging behaviors 42 might shape how relationships form (e.g. bottlenose dolphins: Machado et al. (2019)). Social 43 relationships can determine access to food because closely affiliated individuals can 44 peacefully co-feed at a food patch, hunt together (Lang and Farine, 2017), cooperatively defend food patches (e.g. Emery et al., 2007; Robichaud et al., 1996; Seed et al., 2008), or 45 46 even give food to less successful foragers (e.g. chimpanzees: Samuni et al. (2018)). Access 47 to food is therefore one benefit of long-term cooperative relationships, i.e. stable preferred 48 associations that involve cooperative investments such as grooming and food sharing. For 49 example, grooming in chacma baboons promotes tolerance during foraging (King et al., 50 2011), and vervet monkeys strategically groom individuals that control access to food due to 51 social dominance (Borgeaud and Bshary, 2015) or an experimentally manipulated ability to 52 access food (Fruteau et al., 2009). A particularly clear non-primate example of cooperative 53 relationships providing food occurs in common vampire bats where females regurgitate 54 ingested blood to socially bonded kin and nonkin that failed to feed that night (Carter and Wilkinson, 2013; Wilkinson, 1984). 55

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57 Food-sharing relationships in vampire bats form as preferred associates escalate social grooming (Carter et al., 2020). These preferred associations and cooperative interactions 58 59 occur within the day roost. However, little is known about if or how cooperative relationships extend beyond the roost. For example, foraging with socially bonded roostmates might 60 increase efficiency in searching for prey or feeding from wounds, but it remains unclear if or 61 62 how vampire bats perform social hunting. Several authors provide anecdotal evidence for: groups of females apparently flying together, adult females departing roosts in groups of 2-6, 63 and groups arriving together at a pasture, or approaching and circling prey (Crespo et al., 64 65 1974; Greenhall et al., 1971; Wilkinson, 1985; Wilkinson, 1988). There are also observations of up to four individuals feeding simultaneously from different wounds on the same cow 66 67 (Greenhall et al., 1971), or pairs feeding on the same wound (Greenhall et al., 1971; Wilkinson, 1985). Wilkinson (1985) described evidence that mother-daughter pairs co-forage 68 69 and share wounds, but found no evidence that frequent roostmates forage together. 70

71 Social foraging can take many forms, from mere aggregations attracted to a common resource to coordinated foraging groups with differentiated roles. Socially-hunting species 72 73 can be placed on a spectrum of resource sharing from individual foragers competing to group-level sharing (Lang and Farine, 2017). The form of social foraging and the scale of 74 75 competition over resources outside the roost can have implications for the evolution of food-76 sharing relationships. Several evolutionary models of vampire bat food sharing as multi-level selection view them as foraging individually then sharing food at the group-level (Di Tosto et 77 78 al., 2007; Foster, 2004; Witkowski, 2007), but this view contrasts with evidence that food-79 sharing relationships within groups are reciprocal and highly differentiated (Carter and 80 Wilkinson, 2015; Wilkinson, 1984). An alternative possibility is that individualized relationships drive both within-roost resource sharing and social hunting. This hypothesis is 81 82 not mutually exclusive with group hunting, because even if individuals forage in groups, 83 specific pairs could be more likely to compete or share a wound or host (Delpietro et al., 84 2017; Greenhall et al., 1971; Wilkinson, 1985; Wilkinson, 1988).

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Here, we assessed the relative evidence for a range of hypothetical scenarios that vary in
degree of coordination of social foraging among socially bonded bats (Figure 1). Preferred
roostmates might not coordinate their behavior outside the roost. If instead bats optimize
individual foraging efficiency by preferentially depart, follow, or forage with their preferred
roostmates, then within-roost networks should predict co-departures or foraging encounters.
Alternatively, to maximize their collective search area, bats might prefer to forage with bats

92 outside their network of cooperative relationships and actually avoid foraging with their

frequent roostmates. If so, within-roost and outside-roost networks should be negatively correlated. Finally, if entire roosting groups also forage together, then we expect highly

- 95 correlated within-roost and outside-roost networks.
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97 To evaluate evidence for these scenarios, we tested whether nightly foraging departures and 98 encounters were predicted by: kinship, roosting associations based on two levels of proximity 99 (during the previous day or over the whole study), and rates of social grooming, food sharing, 100 and co-feeding in captivity. To document roosting associations and foraging encounters, we analyzed social encounter data from proximity sensors placed on 50 free-ranging vampire 101 102 bats. As additional predictors for 23 of these bats, we used unpublished data on captive cofeeding rates and published long-term rates of social grooming and food sharing (Ripperger 103 104 et al., 2019). Using simultaneous ultrasonic recording and infrared video, we also describe a 105 distinct new type of vampire bat call only observed during hunting interactions. Our findings 106 illustrate how within-roost cooperative relationships influence foraging in vampire bats and how social networks can vary across contexts. 107



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Figure 1. Hypothetical scenarios for how within-roost relationships predict foraging. 109 For the same roosting association networks, each scenario predicts different outcomes for 110 how preferred roosting relationships correlate with co-departures or encounters during 111 foraging. Preferred roostmates (shown as pair of light brown and dark brown bats) might 112 either: not coordinate their behavior outside the roost (A), coordinate only their departures 113 (B), depart independently and then reunite during foraging (C), coordinate departures and 114 foraging (D), or avoid foraging together (E). Alternatively, the bats could depart and forage as 115 116 a large group (F).

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118 Methods

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120 Subjects

Subjects were common vampire bats (*Desmodus rotundus*) including 27 wild-caught adult females that were tagged and released, and 23 previously captive females (17 adults and

their six subadult captive-born daughters) that had spent the past 22 months in captivity and

were then tagged and released back into their wild roost tree (see Carter et al., 2020;

125 Ripperger et al., 2019). See supplement for details.

126 127 *Kinship*

We assumed that known mother-daughter pairs had a kinship of 0.5. To estimate kinship for

all other pairs, we genotyped bats at 17 polymorphic microsatellite loci (DNA isolated via a

130 salt-chloroform procedure from 3-4 mm biopsy punch stored in 80 or 95% ethanol), then

used the Wang estimator in the R package 'related'. See supplement for details.

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133 **Past cooperative interaction rates in previously captive bats**

To measure cooperative relationships in the previously captive bats, we used previously

published rates of social grooming and food sharing from experimental fasting trials (Carter

et al., 2020). See supplement for details. To assess tolerance while feeding, we also

analysed previously unpublished data on co-feeding among the same captive vampire bats.

- Social interactions were observed at blood spout feeders while the bats were in captivity,
- including 1300 competitive interactions and 277 cases of co-feeding where two bats were
- observed feeding from the same blood spout at the same time (from 1050 h of observation
 from 70 nights). We used 201 co-feeding events with identified bats to construct a co-feeding
- 142 network of the number of dyadic co-feeding events (range = 0 to 6) for each pair.
- 143

To assess correlations between the captive co-feeding network and networks of food sharing 144 or social grooming, we used Mantel tests. To test the same correlation while controlling for 145 146 overlap in individual feeding times, we also used a custom double permutation test (Farine and Carter, 2020). This procedure calculates an adjusted co-feeding rate for each pair as the 147 difference between the observed co-feeding rate and the median expected co-feeding rate 148 from 5000 permutations of the co-feeding bat identities, permuted among the bats seen 149 150 within each hour. The results of this constrained permutation test and the unconstrained Mantel test were similar and gave the same conclusion, so we report only the results from 151 the double permutation test. To test for preferred captive co-feeding partners, we also used 152 the same within-hour permutations to test if social differentiation in co-feeding (the coefficient 153

of variation in co-feeding rates) was greater than expected from the null model.

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156 Association rates in the wild using proximity sensors

We placed custom-made proximity sensors on all 50 female common vampire bats (sensor mass: 1.8 g; 4.5-6.9 % of each bat's mass) that automatically documented dyadic
associations among all 50 tagged bats when those come within reception range (max. 5-10 m). To log encounters, each proximity sensor broadcasted a signal every two seconds to

update the duration of each encounter. We used 1 s as the duration of encounters that were

shorter than two successive signals (i.e., encounters shorter than two seconds). The

163 maximum signal strength of each encounter can be used as an estimate for a minimum 164 proximity between two tagged bats during the encounter by comparing the signal intensity to

165 a calibration curve (Ripperger et al., 2019; Ripperger et al., 2020b).

166

167 We collected association data on the free-ranging bats at Tolé, Panama (8°12'03"N

168 81°43'46"W), a rural area that is mainly composed of cattle pastures for meat production.

Around 200-250 common vampire bats roosted inside a hollow tree on a cattle pasture that

170 was about 15 ha in size. To create a stable food patch, we corralled ca. 100 heads of cattle

- at a distance of ca. 300 m from the roost from 6pm until 6am between the evening of
- September 21 until the morning of September 26, 2017 (days 1 to 5 in our study). Before and

after that time period, the cattle were ranging freely. A neighboring, much larger pasture westof the roost had about 1,500 heads of cattle within a distance of 1-2 km (Figure S1).

175

To construct networks of roosting association rates during each daytime period within the roost, we relied on roosting association data that had been used in a previous study (Ripperger et al., 2019). Based on the same two thresholds of signal strength as before, we defined two categories of proximity: "associations" (within a maximum of ca. 50 cm) and "close contacts" (within ca. 2 cm). Roosting network edges were rates of within-roost association or close contact, i.e. the total time two bats spent in association per unit of time. See supplement for details.

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To log presence and co-occurrence of foraging bats at points outside the roost during the 184 night, we placed base stations (which can detect tagged bats at distances of about 150 m) at 185 the roost and at 5 other locations in the surrounding cattle pastures. To identify departures 186 187 from the roost, we found the points in time where each bat lost connection from the roost 188 base station and almost all of the many tagged bats in the colony within communication range (i.e., a sudden drop in associations from many bats down to 0-3 bats; see figure 2 in 189 190 Ripperger et al. (2020b)). Departing bats may have also contacted base stations on the cattle pasture (Figure S1). We used the same kind of data to find the return times to the roost for 191

192 each bat and night.193

194 Of the 629 dyadic encounters that occurred one minute after leaving the roost and one

195 minute before arriving at the roost, we excluded 43 encounters from further analysis,

because a proximity sensor contacted the roost base station, suggesting that those

197 encounters occurred while bats were roosting at the entrance or on the outside of the roost 198 tree. The remaining 586 encounters occurred farther away, outside the communication range

of the roost base station, and we refer to these as "foraging encounters".

200

201 **Observing interactions of foraging vampire bats**

At Tolé, we only observed two occasions, where two bats stopped at the cattle pasture and 202 were associated (for 3.5 and 4.6 minutes). When releasing the corralled cattle in the 203 morning we observed bite marks but to avoid changing their behavior, we did not get close 204 205 enough to the cattle at night to record audio or video of bats interacting. To collect direct 206 observations on foraging behavior, we recorded simultaneous audio and video of bat foraging behavior at a different farm near La Chorrera, Panama (8°52'42"N 79°52'05"W) 207 using an infrared (IR) spotlight, IR-sensitive video camera (Sony AX53 4K camcorder) and a 208 209 Avisoft condenser microphone (CM16, frequency range 1 to 200 kHz) and digitizer (Avisoft USG 116Hbm, 1000 kHz sampling rate, 16-bit resolution) connected to a notebook 210 211 computer. One observer (SPR) moved with a herd of about 20 grazing cattle without visual light and observed the moving cattle through the viewfinder of the IR camera. To compare 212 213 social calls made during foraging with calls from inside a roost, we used the same recording 214 equipment to record social calls from a roost only a few hundred meters from the foraging 215 site.

215 216

217 Acoustic analysis of calls in foraging bats

We used Avisoft SASLab Pro (R. Specht, Avisoft Bioacoustics, Glienicke/Nordbahn, 218 219 Germany; version 5.2.13) to measure acoustic parameters of the social call types. Start and 220 end of calls were determined manually, based on the oscillogram. Subsequent, five acoustic parameters were measured automatically: one temporal (duration) and four spectral 221 222 parameters (peak frequency at maximum amplitude, minimum and maximum frequency, and 223 bandwidth). Acoustic parameter extraction was restricted to the fundamental frequency. 224 Spectrograms were created using a Hamming window with 1024-point fast Fourier transform and 93.75 % overlap (resulting in a 977 Hz frequency resolution and a time resolution of 225 226 0.064 ms). To estimate the frequency curvatures of the different call types, we measured the spectral parameters at 11 different locations distributed evenly over the fundamental 227 228 frequency of each call. To compare call structure from different contexts (roosting vs

foraging, antagonistic vs affiliative behavior) in multivariate space, we plotted the first two principal components after entering these measures into a principal component analyses

with varimax rotation (using the 'foreign' package in R).

232

233 Statistical analysis of foraging behavior

For every bat, we calculated the times over 9 days when it was clearly distant from the roost 234 tree (ESM File 2). To test whether the previously captive bats and never-captive control bats 235 236 differed in the departure time and duration of their foraging bouts, we fit linear mixed-effect models (LMMs) with type of bats and day as fixed effects and bat as a random effect (p-237 238 values estimated with Satterthwaite's degrees of freedom method using the R package ImerTest). To compare consistency of onsets and durations, we measure the unadjusted 239 240 repeatability (ICC or intra-class correlation) for each type of bat. To see how often tagged 241 bats departed together, we inspected cases where departure times were within one minute. 242

243 **Preferred associations during foraging**

To test if repeated foraging encounters occurred among the same bats more than expected 244 by chance, we used a custom data permutation test that compared observed and expected 245 246 social differentiation (the coefficient of variation in co-foraging rates, which increases when some pairs have more repeated encounters) while controlling for overlap in foraging times. 247 Since all bats were sampled evenly within each night and most foraging encounters were 248 brief (median = 1 second), we first used a simple and conservative measure of co-foraging 249 rates based on counting the presence or absence of an encounter during each hour outside 250 the roost over 9 days (counts varied from 0 to 15). For instance, if two bats met twice in the 251 same hour bin, this is still one encounter. These binary observations could be swapped in 252 our null model. To generate a null distribution of 5000 social differentiations expected by 253 254 chance, we permuted one bat in every dyad to a random possible partner that was also present outside the roost in that same day and hour (to control for overlaps in foraging 255 256 times).

257

258 **Predictors of social foraging**

To test predictors of social foraging, we constructed foraging encounter networks where 259 edges were based on either duration of total encounter time outside the roost (seconds) or 260 261 number of days with foraging encounters (0-9). The latter response variable is far more conservative because it only counts repeats across different days. We included the following 262 predictors: kinship, two proximity levels of within-roost association, social-grooming rate, 263 food-sharing rate. We also tested the effect of dyad type (i.e. both bats previously captive, 264 both bats never captive, one bat previously captive, and both bats captive-born juveniles). 265 We did not use number of nights with foraging encounters as a response for tests that only 266 267 included the previously captive bats, because 9 of them (including all captive-born bats) left 268 the roost during the study period (Ripperger et al., 2019). To measure how much longer 269 foraging encounters were between kin versus nonkin, we fit a linear mixed effects model with 270 log-transformed duration as the response variable, kinship greater than 0.1 as a binary fixed effect, and both bats' identities as random effects, then converted model coefficients into a 271 272 percentage difference.

273

To test the effect of predictor networks on a response network, we used regression quadratic 274 275 assignment procedure (QAP) for single predictors, or multiple regression quadratic 276 assignment procedure with double semi-partialling (MRQAP) for two predictors (using the 'asnipe' R package (Farine, 2013)). To create null models, we used constrained (within-day) 277 278 node-label permutations. This approach is necessary for preserving the daily and nightly network structure (e.g. distribution of edges and edge weights) and for controlling for the 279 280 presence or absence of bats in the roost each day. To control for foraging bout overlap in each pair, we included that measure as a covariate. We also used QAP to test whether the 281 282 within-roosting association on each day, predicted the subsequent foraging network that 283 night. We then bootstrapped the mean of the slopes across the eight days to test for an 284 overall paired day-night effect.

overall paired day-night effect

285 Consistency of individual social traits

286 To test whether bats that are more socially connected within the roost are also more connected in foraging networks, we tested if the nodes' degree centrality was correlated 287 between roosting and foraging networks. We measured degree centrality independently 288 289 within each day or night network when the bat was present and then took the mean for each 290 bat. Bats with no encounters in that day or night were considered missing for that day (i.e. not counted as zero degree). We fit general linear mixed effect models with foraging network 291 292 centrality as the response variable, roosting network centrality (either association and close contact) as fixed effect, and bat as random effect. P-values were calculated from 5000 293 294 permutations of the bat's foraging centralities within each night. These constrained nodelabel permutations (within night) are necessary to control for the fact that foraging and 295 296 roosting network centralities could be correlated simply by some bats being present at the 297 site longer.





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Figure 2: Network comparisons. Foraging encounter rates were predicted by roosting 300 associations, kinship, and previous long-term rates of social grooming and food sharing in 301 captivity. To facilitate visual comparisons, we fixed the spatial coordinates of each node 302 (except for the sparse night-by-night foraging networks), we scaled edge strength in each 303 network, and we removed nodes without edges. In the kinship network, only edges with 304 kinship estimates > 0.24 are shown and bats without kin in the group are not shown. In the 305 paired night-day networks of association in the wild, we only detected a clear correlation 306 between day and night on day 4 (Table S1). 307

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309 Results

310

311 Sampled bats did not depart together

The never-captive control bats departed from the roost 8.3 hours after sunset and returned 2.5 h later, on average (ESM 2). The previously captive bats foraged earlier and less predictably (see below). We observed only five cases where two bats departed within five seconds of each other and none of these cases was followed by a foraging encounter. For the cases where pairs did have a foraging encounter, the shortest differences in departure

- times were 8, 21, and 28 s.
- 318

319 **Previous captivity influenced departures and foraging**

320 Compared to the wild control bats, the previously captive bats departed the roost on average 321 1.6 hours earlier (t = -4.55, p<0.0001), but they did not forage consistently longer (LMM; t=1.29, p=0.2, ESM 2). The captive-born bats departed 2 hours earlier (t = -3.15, p = 0.002) 322 and also did not forage longer (t = -0.41, df = 47.8, p = 0.7) than control bats. All these 323 324 models control for departure times being on average 14 minutes later each day (t = 6.6, p < 325 0.0001, ESM 2), perhaps due to moonset times being 20-40 min later each day during the 326 study period. The total duration of foraging encounters did not clearly differ between types of pairs (Figure S3A), but pairs of control bats had significantly more nights with foraging 327 encounters (Figure S3B) compared to other types of pairs, possibly due to control bats 328 having more consistent foraging times. Departure times were more consistent across days 329 within each control bat (intraclass correlation coefficient (ICC) = 0.58) compared to within 330 each previously captive bat (ICC = 0.21) or captive-born bat (ICC = 0). The duration of the 331 332 longest foraging bout was also more consistent in wild control bats (ICC = 0.54) than the previously captive bats (ICC = 0.35) or captive-born bats (ICC = 0.15). 333

334

335 **Preferred associations in foraging encounter networks**

336 Foraging encounters were orders of magnitude shorter in duration than within-roost

- encounters, their median duration was 1 s, and they never exceeded 30 minutes (ESM 1,
- Figure S2). Of 151 pairs with a foraging encounter, 45 did this repeatedly across 9 nights.
- 339 The variation in number of hours in which two bats reunited was greater than expected from
- our null model that simulated random encounters among bats that were outside the roost in the same hour (observed coefficient of variation = 4.36; p < 0.001; 95% of expected values:
- the same hour (observed coefficient of variation = 4.36; p < 0.001; 95% of expected values: -
 2.2 to 2.4). Most of these foraging encounters occurred at locations outside our sampled
 areas, but 10 (among eight pairs) occurred near the other base stations on the surrounding
 cattle pastures (ESM 1, Figure S1), and only three foraging encounters (among three pairs)
 occurred at the corral that we created as a stable food patch about 300 m from the roost (two
 encounters on days one and three while the cattle were present and one encounter on day
- 346 encount347 seven).
- 347 348

349 Kinship predicts foraging encounters

Kinship predicted the number of nights with foraging encounters (QAP, β = 15.4, n = 46 bats, p < 0.0001) and foraging encounter time (β = 15.4, n = 47 bats, p = 0.022) even when controlling for bout overlap (MRQAP, β = 0.10, p = 0.002). The median duration of a foraging encounter for close kin (r > 0.1) was 9 s, which was 135% longer in duration relative to the duration of foraging encounters between nonkin (r < 0.1; median duration = 1s; β =0.85, df=175, p=0.001).

356

357 Within-roost association rates predicted foraging encounters

Bats that spent more time near each other within the tree during the day, also spent more time together outside the roost during the night (associations: QAP, β = 29.5, p < 0.001; close-contact: QAP, β = 24.7, p = 0.002) even when controlling for the foraging bout overlap

- 361 (associations: MRQAP, $\beta = 0.092$, p = 0.003; close-contact associations: MRQAP, $\beta = 0.078$,
- p = 0.015). Within-roost associations also predicted a greater number of nights with foraging
- encounters (associations: QAP, β = 0.07, p < 0.001; close-contact association: QAP, β =
- 364 0.04, p = 0.021). The relationship between the within-roost association network and the

corresponding night's foraging network was clear within only one of the 8 days (Table S1), but the paired relationships between day and night networks tended to be greater than zero overall (associations: mean $\beta = 0.026$, 95% CI = 0.004 to 0.051; close-contact associations: mean $\beta = 0.018$, 95% CI = 0.003 to 0.04).

370 Roosting degree centrality predicted foraging degree centrality

Bats that were connected to more associates in within-roost association networks also tended to have more associates in the nightly foraging networks (associations: $\beta = 0.034$, n = 48 bats, one-tailed p = 0.008; close-contact: $\beta = 0.055$, one-tailed p = 0.078; Figure S4).

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369

375 **Cooperative relationships in captivity predict foraging encounters in the field**

In the previously captive bats, foraging encounter time was predicted by social grooming (QAP, β = 26.5, n = 22, p = 0.032; MRQAP controlling for bout overlap: β = 0.12, p = 0.063) and food sharing (β = 38.7, n = 22, p = 0.015; MRQAP controlling for bout overlap: β = 0.20, p = 0.014), and food sharing when controlling for kinship (MRQAP, sharing: β = 0.16, p = 0.022; kinship: β = 0.14, p = 0.049). Kinship and cooperative relationship were therefore independent predictors of social foraging.

382

Co-feeding among familiar captive bats was not limited to cooperative relationships.

In contrast to the measures of social foraging in the field, we detected only weak evidence for preferred associations during co-feeding in captivity (social differentiation = 2.10, p = 0.047 when controlling for hour, p = 0.041 when not controlling for hour), and we found no correlation between captive co-feeding and social grooming (r=0.008, p=0.36), food sharing (r=0.015, p=0.28) or social foraging time in the wild (r=0.003, n=20, p=0.42).

389

390 Behavioral interactions during foraging

To record a sample of bat interactions during foraging encounters, we recorded infrared 391 video and ultrasonic audio of 14 interactions between foraging vampire bats (Tables S2 and 392 393 S3). Social calls during foraging had three general spectral shapes (Figures S5 and S6): "downward sweeping calls" have been recorded often in roosts and are produced by socially 394 isolated vampire bats in captivity (Carter et al., 2012; Carter and Wilkinson, 2016). "Buzz 395 calls" were noisy without clear tonal structure and occurred during antagonistic interactions. 396 We observed "n-shaped calls" produced by bats interacting while near cattle (Figure S6), and 397 398 to our knowledge this call type has never been seen in wild roosts, from confrontations at the 399 feeders in captivity (Sailler and Schmidt, 1978), or from individually isolated bats in captivity 400 (Carter et al., 2012, Carter, unpublished data).

401 402

403 **Discussion**

404

405 Long-term cooperative relationships predicted repeated foraging encounters

Tagged female vampire bats departed the roost individually, but often re-united far from the 406 roost during foraging bouts. The rates of these foraging encounters were consistently higher 407 408 than expected in specific pairs and predicted by roosting associations, kinship, and by the history of social grooming and food sharing in captivity, even when controlling kinship. 409 410 Previous experiments with female vampire bats suggest that these measures—roosting 411 proximity, social grooming, and food sharing-reflect an underlying cooperative relationship 412 (Carter et al., 2020; Carter and Wilkinson, 2013; Ripperger et al., 2019; Wilkinson, 1984; Wilkinson, 1985). In this study, we knew the cooperation histories among the previously 413 414 captive bats, and that these individuals had no interactions with the control bats for at least the previous 22 months. We could therefore infer that relationships that are typically defined 415 by associations and cooperative interactions within roosts, also extend beyond the roost and 416 may provide benefits during foraging. In addition to consistent social relationships across 417 418 context (from captivity to roosting to foraging), we found that bats that encountered more 419 associates in the roost during the days also encountered more associates while foraging

420 during the nights, suggesting consistent individual variation in social traits.

Although some foraging encounters may have occurred before or after foraging, most of 421 422 these encounters were likely to have occurred during foraging for several reasons. First, foraging encounters were brief, whereas associations among non-moving bats should be 423 much longer in duration (Figure S2). Second, foraging is likely to take up a substantial 424 425 amount of the limited time outside the roost (mean = 2.4 h). After commuting, searching, and 426 selecting a host, a vampire bat can take up to 30 minutes to select a wound site, 10-40 427 minutes to prepare the wound site, and 9-40 minutes to feed (Greenhall, 1972; Greenhall et 428 al., 1971). Third, we used infrared video to observe several interactions on or near cattle that were consistent with the short durations of foraging encounters in the proximity data (e.g., 429 430 Videos S1, S2, S4). Fourth, foraging encounters among close female kin had a median 431 duration of 9 s and were longer than among non-kin (median duration of 1s), which is 432 consistent with observations that affiliative interactions last longer.

433

434 No clear evidence for highly coordinated collective movements

For animals with fluid social structures (e.g. high fission-fusion dynamics), it is important to 435 436 clarify the ambiguous meaning of a "social group", and similarly, one must distinguish between different possible forms of "social foraging" (Lang and Farine, 2017). In bats, the 437 438 relative degree of social coordination during foraging can be difficult to assess and compare due to differing limitations in the observational methods and the lack of knowledge of 439 differentiated social relationships within the colony. In this study, we took advantage of well 440 described within-roost relationships to assess evidence for several alternative scenarios of 441 foraging behavior (Figure 1). Kinship and rates of association and cooperation led to longer 442 and more frequent foraging encounters, but we did not observe highly coordinated joint 443 444 departures or collective movements (Figure 1). This fluid pattern, of not moving in coordinated stable groups yet repeatedly encountering preferred associates during foraging, 445 446 is also reflected in co-roosting networks where individuals form roosting groups that frequently change composition, yet maintain preferred relationships over time (Wilkinson, 447 1985). Given the many unsampled bats inside the same tree (~200), it is possible that bats 448 449 departed with other unobserved roostmates, but we did not see departures of large groups (while catching bats outside the roost) nor did we see evidence for coordination between 450 451 roosting and departing in the tagged bats.

The ways that specific bats reunited with preferred associates therefore remains unknown,
but the downward sweeping calls that we recorded in foraging bats (Figure S6), are similar to
contact calls that captive vampire bats can use to find and recognize preferred partners

456 (Carter and Wilkinson, 2016). The role of calls, in particular a possibly foraging-specific call

type ("n-shaped call" Figure S6, Figure S7), warrants further investigation. In several other
 bat species, there is abundant evidence for socially influenced foraging based on

438 eavesdropping on echolocation calls (e.g. Cvikel et al., 2015; Dechmann et al., 2009;

460 Dechmann et al., 2010; Egert-Berg et al., 2018; Lewanzik et al., 2019). Greater spear-nosed

461 bats in Trinidad appeared to coordinate group foraging based on a group-specific contact call

462 (Wilkinson and Boughman, 1998). Observations of the fish-eating greater bulldog bat

- suggested that female roostmates depart individually, then assembled into small groups
- 464 outside the roost to forage together, possibly coordinating their movements with calls
- 465 (Brooke 1997). 466

467 Affiliative and competitive interactions

468 Given the difficulty of making a bite compared to the ease of drinking from an open wound, some individual vampire bats appear to exploit the bites already made by others, and fights 469 470 can occur over open wounds or hosts (Delpietro et al., 2017; Greenhall et al., 1971; Sazima, 1978; Schmidt, 1978; Wilkinson, 1985), but it remains unclear how often these competitive 471 472 interactions occur among familiar versus unfamiliar vampire bats. In our study, we observed foraging vampire bats engaging in both affiliative and competitive interactions (see Table S3, 473 474 Videos S1-5). The competitive interactions were far more aggressive than what we observed 475 among familiar captive bats feeding from an accessible and unlimited source of blood. This

- observation and our results above are consistent with the hypothesis that competitive
- 477 interactions are more likely between less familiar bats.
- 478

479 *Implications for social dominance*

480 The fluid nature of foraging encounters has potential implications for social dominance.

- 481 Dominance hierarchies should be common when animals move together in groups, because
- the same frequent groupmates will also be primary competitors for first access to food.
- 483 Dominance hierarchies among familiar female vampire bats, which do not always travel or
- 484 forage together, are indeed less clear and linear than among female mammals that do travel
- and forage in more stable groups (Crisp et al. unpublished data). Furthermore, blood from an
- 486 open wound is often not a limited resource, so competition over food might be relatively low
- among familiar vampire bats that tolerate each other (as observed in captivity) and even
 share food, compared to unfamiliar conspecifics that might "steal" a wound.
- 489

490 Implications for the evolution of cooperative relationships

Vampire bats might also benefit from foraging with socially tolerant partners by acquiring 491 information on where to feed or by gaining access to open wounds. A single open wound can 492 493 sequentially feed several bats, and leading a cooperation partner to an open wound on the same host would presumably be less costly to a successful forager than regurgitating blood 494 to that individual later at the roost. Put differently, socially bonded bats would benefit from 495 each other's foraging success, i.e. interdependence (Roberts, 2005). As seen in ravens, the 496 497 presence of a socially bonded partner might even allow for joint defense of food against thirdparties (Sierro et al., 2020; Szipl et al., 2015). 498

499

Such forms of social foraging in vampire bats may have implications for the spatial scale of 500 501 competition-a key factor shaping social evolution in humans (West et al., 2006) and other group-living animals (Radford et al., 2016). In female vampire bats, cooperation occurs 502 'locally' with specific frequent roostmates, and competition over food might occur more 503 504 'globally' with members of the much larger population. If so, a more 'global' scale of 505 competition could reduce conflict and increase interdependence among highly associated females. To test this idea, it would thus be useful to determine if sampled groups of vampire 506 507 bats consistently feed on the same or different prey individuals, and if vampire bats are more likely to approach or avoid the social calls of foraging bats that are frequent roostmates 508 509 versus unfamiliar conspecifics.

510

511 Implications for describing social structure

512 A major advantage of proximity sensors was the ability to continuously track associations among multiple individual bats both inside and outside their roost, which allows for the 513 514 construction of dynamic and multi-layer networks. Studies on social foraging and other social behaviors in bats and other small highly mobile vertebrates have historically been limited by 515 the available tracking technology (Ripperger et al., 2020b). Radiotelemetry has poor spatial 516 517 resolution and continuously tracking many individuals is difficult. Current GPS-tags for bats have rather short runtimes and the tags need to be recovered to download the data. On-518 519 board ultrasound recorders (e.g. Egert-Berg et al., 2018) do not reveal the identity of 520 encountered individuals. A major downside to proximity sensors was that many foraging 521 encounters occurred at unknown locations. However, placing proximity sensors or antennas 522 at more locations and on livestock would allow a better reconstruction of foraging behavior. A 523 combination of biologging approaches can also help to overcome existing challenges (e.g. Leoni et al., 2020: Ripperger et al., 2020a). Standardized high-throughput methods for 524 525 measuring social network structure across bats and other diverse groups allow for comparative studies that assess the relative ecological and evolutionary drivers of social 526 527 traits and social complexity across species.

528

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