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# 1 A model of resource partitioning between foraging bees

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#### 27 Statement of authorship

- 28 CP, TD and ML designed the study. TD and JG developed the models. CP and TD performed the
- 29 analyses. CP, TD and ML wrote the first draft of the manuscript. All authors contributed substan-
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### 40 Abstract

41 Central place foraging pollinators tend to develop multi-destination routes (traplines) to exploit several 42 patchily distributed plant resources. While the formation of traplines by individual pollinators has been stud-43 ied in details, how populations of individuals exploit resources in a common area is an open question diffi-44 cult to address experimentally. Here we explored conditions for the emergence of resource partitioning 45 among traplining bees using agent-based models built from experimental data of bumblebees foraging on 46 artificial flowers. In the models, bees learn to develop routes as a consequence of feedback loops that change 47 their probabilities of moving between flowers. While a positive reinforcement of route segments leading to 48 rewarding flowers is sufficient for the emergence of resource partitioning when flowers are evenly distribut-49 ed, a negative reinforcement of route segments leading to unrewarding flowers is necessary when flowers are 50 patchily distributed. In these more complex environments, the negative experiences of individual bees favour 51 the spatial segregation of foragers and high levels of collective foraging efficiency.

- 52
- 53 Keywords: bumblebees; competition; resource partitioning; trapline foraging; vector navigation

# 54 Introduction

Animals competing for food resources are expected to self-distribute on feeding sites in order to maximize their individual energy gain (Fretwell and Lucas 1969; Giraldeau and Caraco 2000). Resource partitioning between individuals of different species is well documented, and often results from functional (Fründ *et al.* 2010; 2013) or behavioral (Nagamitsu and Inoue 1997; Valdovinos *et al.* 2016) differences. By contrast, how individuals of the same species optimally interact to exploit resources in a common foraging area is less understood (Johst *et al.* 2008; Tinker *et al.* 2012).

61 For pollinators, such as bees that individually exploit patchily distributed floral resources in envi-62 ronments with high competition pressure, efficient resource partitioning appears a prodigious problem in-63 volving the quality of food resources, their spatial distribution, their replenishment rate, and the activity of 64 other pollinators. As central place foragers, bees often visit familiar feeding sites in a stable sequence or 65 trapline (Janzen 1971; Thomson et al. 1997). Individual bees with exclusive access to an array of artificial 66 flowers tend to develop traplines minimizing travel distances to visit all the necessary flowers to fill their 67 nectar crop and return to the nest (e.g. bumblebees: Ohashi et al. 2008, Lihoreau et al. 2012a, Woodgate et al. 68 2017; honey bees: Buatois and Lihoreau 2016). This routing behavior involves spatial memories that can 69 persist several days or weeks (Thomson et al. 1996; Lihoreau et al. 2010).

70 How bees partition resources, when several conspecifics exploit a common foraging area, is however 71 still an open question. Experimentally the problem is challenging to address as it requires monitoring the 72 movements of several bees simultaneously over large spatial and temporal scales. In theory, bees should 73 develop individualistic traplines that minimize travel distances and spatial overlap with other foragers, there-74 by improving their own foraging efficiency and minimizing the costs of competition (Ohashi and Thomson 75 2005; Lihoreau et al. 2016). Best available data supporting this hypothesis come from observations of small 76 numbers of bumblebees foraging on potted plants (e.g. Makino and Sakai 2005, Makino 2013) or artificial 77 flowers (Lihoreau et al. 2016, Pasquaretta et al. 2019) in large flight tents. In these rather simple foraging 78 conditions, bees tended to avoid spatial overlaps as a consequence of competition by exploitation (when bees 79 visited empty flowers) and interference (when bees interacted on flowers) (Pasquaretta et al. 2019). 80 Computational modeling is a powerful approach to further explore how such partitioning might

81 emerge from spatial learning and competitive interactions. At the individual level, trapline formation has

82 been modeled using an iterative improvement algorithm by which a bee compares the net length of the route 83 it has just traveled (sum of the lengths of all movement vectors comprising the flower visitation sequence) to 84 the length of the shortest route experienced so far (Lihoreau et al. 2012b). If the new route is shorter (or 85 equivalent), the bee increases its probability of using all the movement vectors composing the new route in 86 its subsequent foraging bout. After several iterations, this route-based learning algorithm typically leads to 87 the discovery and selection of a short (if not the shortest possible) trapline. While this approach can accurate-88 ly replicate observations across a wide range of experimental conditions (Reynolds et al. 2013), it makes the 89 strong assumption that bees can compute, memorize and compare the lengths of multi-leg routes upon return 90 to their nest. Recently, it was proposed that such behavior could also emerge from vector-based learning (Le 91 Moël et al. 2019), which is more parsimonious and plausible considering the current understanding of spatial 92 computation in the insect brain (Stone et al. 2017). So far, however, none of these traplining models have 93 accounted for social interactions and current models of pollinator populations do not take into account indi-94 vidual specificities of movements based on learning and memory (Becher et al. 2014; 2016; 2018). Thus 95 presently, there has been no realistic exploration of how resource partitioning between interacting bees might 96 form. 97 Here, we investigated the behavioral mechanisms underpinning resource partitioning among 98 traplining bees by comparing predictions of agent-based models integrating route learning and social interac-99 tions. First, we developed models implementing biologically plausible vector navigation based on positive

100 and negative reinforcements of route segments leading to flowers. We used different models to test the indi-

101 vidual and combined influences of these feedback loops on route learning. Next, we explored how these

102 simple learning rules at the individual level can promote complex patterns of resource partitioning at the

103 collective level, using simulations with multiple foragers in environments with different resource distribu-

104 tions.

105

# 106 METHODS

#### 107 Model overview

108 We developed two agent-based models in which bees learn to develop routes in an array of flowers (Fig. 1).

109 In both models, all flowers contain the same quality and volume of nectar that is refilled between foraging

110 bouts (flower visitation sequence, beginning and ending at the colony nest entrance). In simulations with 111 multiple bees, all bees start their foraging bout synchronously and the flowers are filled after the last bee has 112 returned to the nest. At each foraging bout, each bee attempts to collect nectar from five different flowers in 113 order to fill its nectar crop (stomach) to capacity. For simplicity, this rule was set constant across all 114 simulations. For each bee, flower choice is described using movement vectors (orientated jump between two 115 flowers or between the nest and a flower). The initial probability of using each possible movement vector is 116 based on movement vector length. This probability is then modulated through learning when the bee uses a 117 vector for the first time. The learning rule varied depending on the model. If the ending flower of the vector 118 contains nectar, it is rewarding and the probability to reuse the vector later increased (positive reinforcement, 119 models 1 and 2). If the flower is empty, it is not rewarding and the probability to reuse the vector is either 120 left unchanged (model 1) or decreased (negative reinforcement, model 2). A flower is empty if it has 121 previously been visited in the same foraging bout by the same bee or another bee (exploitation competition). 122 If two bees visit a flower at the same time, a fight to secure the possible reward occurs (interference 123 competition). Only one bee can stay on the flower and access the potential nectar reward with a random 124 probability. The winner bee takes the reward if there is one. The loser reacts as for an empty flower. Both 125 bees update their probabilities to reuse the vector accordingly. Route learning thus depends on the experience 126 of the bee and its interactions with other foragers. For simplicity, we restricted our analysis to two bees, but 127 the same models can be used to simulate interactions among more bees (see examples with five bees in 128 Video S1, Appendix S1). The complete R code is available at 129 https://gitlab.com/jgautrais/resourcepartitioninginbees/-/releases.

130

#### 131 Environment

132 Simulations with one foragers

133 To calibrate our models, we ran simulations in environments replicating published experimental studies in

- 134 which individual bumblebees (*Bombus terrestris*) were observed developing traplines between five equally
- 135 rewarding artificial flowers in a large open field (Lihoreau et al. 2012b; Woodgate et al. 2017). To our
- 136 knowledge, these studies provide the most detailed available datasets on trapline formation by bees. Lihoreau
- 137 et al. (2012b) used a positive array of flowers (regular pentagon, Fig. S1A) in which the distance and
- 138 direction between flowers were positively linked, i.e. nearest neighbor flowers could be visited using lowest

139 angular deviation between consecutive vectors. The authors tracked seven bumblebees, which we judged

140 enough to run quantitative comparisons with model simulations (raw data are available in Table S1 of

141 Lihoreau et al. 2012b). Woodgate et al. (2017) used a negative array of flowers (narrow pentagon, Fig. S1B)

142 in which the distance and direction between flowers were negatively linked. Here, however, the small sample

143 size of the original dataset (three bumblebees, data shared by J. Woodgate) only enabled a qualitative

144 comparison with the model simulations (Appendix S2).

145

146 Simulations with two foragers

147 We explored conditions leading to resource partitioning by running model simulations with two foragers.

148 Here we simulated environments containing 10 flowers (each bee had to visit five rewarding flowers to fill

149 its crop to capacity). To test whether model predictions were robust to variations in resource spatial

150 distribution, we simulated three types of environments characterized by different levels of resource

151 patchiness: (i) a patch of 10 flowers, (ii) two patches of five flowers each, and (iii) three patches of five,

152 three and two flowers respectively (see examples in Fig. S2). We generated flower patches into a spatial

153 range comparable to the one used in both experimental setups (Lihoreau *et al.* 2012b; Woodgate *et al.* 2017):

154 about 500  $m^2$  with a minimum distance of 160 m between each patch center. Within a patch, flowers were

155 randomly distributed according to two constraints: (i) flowers were at least 20 m apart from each other, (ii)

156 the maximum distance of each flower from the center of the patch was 40 m. This ensured that each patch

157 had a maximum diameter of 80 m and inter-flower distances were smaller between all flowers of the same

158 patch than between all flowers of different patches.

159

#### 160 Movements

161 At each step, a bee chooses to visit a target location (flower or nest) based on a matrix of movement

162 probabilities (Fig. 1). This matrix is initially defined using the inverse of the square distance between the

163 current position of the bee and all possible target locations (Lihoreau *et al.* 2012b; Reynolds *et al.* 2013). The

164 probability of moving from location *i* to the location *j* among n possible targets, is initially set to:

166 (1) 
$$P(i \to j) = \frac{1}{d_{ij}^2} / \sum_{k \neq j} \frac{1}{d_{ik}^2}$$

1	67	
T	07	

168	Where $d_{ij}$ is the distance between locations <i>i</i> and <i>j</i> . Before choosing its destination, the bee lists all possible
169	target locations. For simplicity, the bee excludes its current location, thus preventing looping flights to and
170	from the same flower or nest, which are rare in experienced bumblebee foragers (Ohashi et al. 2007; Saleh
171	and Chittka 2007; Lihoreau et al. 2010; Lihoreau et al. 2012a; Lihoreau et al. 2012b; Woodgate et al. 2017).
172	The bee also excludes the location it had just come from. This simulates the tendency of bumblebees to avoid
173	recently visited (and thus depleted) flowers (Ohashi et al. 2007; Saleh and Chittka 2007; Lihoreau et al. 2010;
174	Lihoreau et al. 2012a; Lihoreau et al. 2012b; Woodgate et al. 2017). The foraging bout ends if: (i) the bee
175	fills its crop to capacity, (ii) the bee chooses the nest as a target destination, or (iii) the bee reaches a
176	maximum travelled distance of 3000 m. The latest was added to avoid endless foraging trips in the model. Its
177	value was chosen based on the observation that bumblebees typically forage within a distance of less than 1
178	km from their nest (Osborne et al. 1999; Wolf and Moritz 2008; Woodgate et al. 2016).
179	
180	Learning
181	Learning modulates the probability of using movement vectors as soon as the bee experiences the chosen
	Detaining includates the producting of using increment rectors as soon as the cee experiences are enosed
182	target (online learning) and only once within a foraging bout (the first time the movement vector is used; Fig.
182 183	<ul><li>target (online learning) and only once within a foraging bout (the first time the movement vector is used; Fig.</li><li>1). This approach has the advantage of implementing vector navigation (Stone <i>et al.</i> 2017; Le Moël <i>et al.</i></li></ul>
182 183 184	<ul> <li>target (online learning) and only once within a foraging bout (the first time the movement vector is used; Fig.</li> <li>1). This approach has the advantage of implementing vector navigation (Stone <i>et al.</i> 2017; Le Moël <i>et al.</i></li> <li>2019) and thus avoids unrealistic assumptions about computation and comparison of complete routes</li> </ul>
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182 183 184 185 186 187 188 189 190	target (online learning) and only once within a foraging bout (the first time the movement vector is used; Fig. 1). This approach has the advantage of implementing vector navigation (Stone <i>et al.</i> 2017; Le Moël <i>et al.</i> 2019) and thus avoids unrealistic assumptions about computation and comparison of complete routes (Lihoreau <i>et al.</i> 2012b; Reynolds <i>et al.</i> 2013). Positive reinforcement was implemented in models 1 and 2. It occurs when a bee uses a vector leading to a rewarding flower. The probability of using this vector is then multiplied by 1.5 (other vectors probabilities are scaled accordingly to ensure that all sum up to 1). This positive reinforcement is based on the well-known tendency of bumblebees to return to nectar rewarding places through appetitive learning (Goulson 2010). Negative reinforcement was implemented in model 2 only. It occurs when a bee uses a vector leading to a non-rewarding flower. The bee reduces the probability
182 183 184 185 186 187 188 189 190 191	target (online learning) and only once within a foraging bout (the first time the movement vector is used; Fig. 1). This approach has the advantage of implementing vector navigation (Stone <i>et al.</i> 2017; Le Moël <i>et al.</i> 2019) and thus avoids unrealistic assumptions about computation and comparison of complete routes (Lihoreau <i>et al.</i> 2012b; Reynolds <i>et al.</i> 2013). Positive reinforcement was implemented in models 1 and 2. It occurs when a bee uses a vector leading to a rewarding flower. The probability of using this vector is then multiplied by 1.5 (other vectors probabilities are scaled accordingly to ensure that all sum up to 1). This positive reinforcement is based on the well-known tendency of bumblebees to return to nectar rewarding places through appetitive learning (Goulson 2010). Negative reinforcement was implemented in model 2 only. It occurs when a bee uses a vector leading to a non-rewarding flower. The bee reduces the probability of using that vector by multiplying it by 0.75 (here also rescaling the other ones). This negative
182 183 184 185 186 187 188 189 190 191 192	target (online learning) and only once within a foraging bout (the first time the movement vector is used; Fig. 1). This approach has the advantage of implementing vector navigation (Stone <i>et al.</i> 2017; Le Moël <i>et al.</i> 2019) and thus avoids unrealistic assumptions about computation and comparison of complete routes (Lihoreau <i>et al.</i> 2012b; Reynolds <i>et al.</i> 2013). Positive reinforcement was implemented in models 1 and 2. It occurs when a bee uses a vector leading to a rewarding flower. The probability of using this vector is then multiplied by 1.5 (other vectors probabilities are scaled accordingly to ensure that all sum up to 1). This positive reinforcement is based on the well-known tendency of bumblebees to return to nectar rewarding places through appetitive learning (Goulson 2010). Negative reinforcement was implemented in model 2 only. It occurs when a bee uses a vector leading to a non-rewarding flower. The bee reduces the probability of using that vector by multiplying it by 0.75 (here also rescaling the other ones). This negative reinforcement rule is based on the tendency of bumblebees to reduce their frequency of revisits to
<ol> <li>182</li> <li>183</li> <li>184</li> <li>185</li> <li>186</li> <li>187</li> <li>188</li> <li>189</li> <li>190</li> <li>191</li> <li>192</li> <li>193</li> </ol>	target (online learning) and only once within a foraging bout (the first time the movement vector is used; Fig. 1). This approach has the advantage of implementing vector navigation (Stone <i>et al.</i> 2017; Le Moël <i>et al.</i> 2019) and thus avoids unrealistic assumptions about computation and comparison of complete routes (Lihoreau <i>et al.</i> 2012b; Reynolds <i>et al.</i> 2013). Positive reinforcement was implemented in models 1 and 2. It occurs when a bee uses a vector leading to a rewarding flower. The probability of using this vector is then multiplied by 1.5 (other vectors probabilities are scaled accordingly to ensure that all sum up to 1). This positive reinforcement is based on the well-known tendency of bumblebees to return to nectar rewarding places through appetitive learning (Goulson 2010). Negative reinforcement was implemented in model 2 only. It occurs when a bee uses a vector leading to a non-rewarding flower. The bee reduces the probability of using that vector by multiplying it by 0.75 (here also rescaling the other ones). This negative reinforcement rule is based on the tendency of bumblebees to reduce their frequency of revisits to unrewarded flowers with experience (Pasquaretta <i>et al.</i> 2019). We applied a lower value to negative

195	than negative stimuli (visits to non-rewarding flowers) (review in Menzel 1990). Sensitivity analyses of these
196	two parameters show that increasing positive and/or negative reinforcement increases the speed and level of
197	resource partitioning (see Appendix S1).
198	
199	Competitive interactions
200	We implemented competitive interactions between foragers in the form of exploitation and interference (Fig.
201	1). Exploitation competition occurs when a bee lands on a flower whose nectar reward has already been
202	collected by another bee. If the flower is empty, the probability to reuse the vector is either left unchanged
203	(model 1) or decreased (negative reinforcement, model 2). Interference competition occurs when two bees
204	encounter on a flower. Only one bee can stay on the flower and access the potential nectar reward with a
205	random probability (p=0.5). After the interaction, the winner bee takes the reward if there is one. The loser
206	bee reacts as for an empty flower.
207	
208	Statistical Analyses

209 All analyses were performed in R version 3.3 (R Development Core Team 2018).

210

211 Simulations with one forager

212 For each model, we compared the results of the simulations to the reference observational data, either

213 quantitatively (for Lihoreau et al. 2012b) or qualitatively (for Woodgate et al. 2017; see Appendix S2). We

stopped the simulations after 50 foraging bouts to match the experimental conditions of the published data

215 (22-37 foraging bouts in Lihoreau et al. 2012b; 47-61 foraging bouts in Woodgate et al. 2017). We run 500

216 simulations for each model (without and with negative reinforcement) and we estimated how models fitted

217 the experimental data using two main measures:

218

219 (*i*) the quality of each route QL, calculated as:

221 (2) 
$$QL = \frac{\frac{F^2}{d}*1}{QL_{opt}}$$

0	0	2
Z	Z	2

223 Where F is the number of rewarding flowers visited during a foraging bout and d is the net length of all 224 vectors traveled during the foraging bout. QL is standardized in [0; 1] by the quality of the optimal route in 225 each array  $QL_{opt}$  (shortest possible route to visit all flowers). 226 227 (ii) a similarity index  $SI_{ab}$  between flower visitation sequences experienced during two consecutive foraging 228 bouts *a* and *b* as follows: 229  $(3) \qquad SI_{ab} = \frac{s_{ab}}{2l_{ab}}$ 230 231 232 Where  $s_{ab}$  represents the number of vectors between two flowers found in both sequences, and  $l_{ab}$  the 233 length of the longest flower visitation sequence between i and j multiplied by 2 to make sure that  $SI_{ab} = 1$ 234 occurs only when two consecutive sequences sharing the same vectors also have the same length.

We applied generalized linear mixed effect models (GLMM) with binomial error, using the glmer
function in 'Ime4' package (Bates et al. 2014), to assess whether the estimated trends across foraging bouts
for *QL* and *SI<sub>ab</sub>* obtained from model simulations with one forager differed from trends obtained from
observational data. In each model, we used a random structure to account for the identity of bees. *Simulations with two foragers*We generated 10 arrays of flowers for each of the three types of environment, and ran 100 simulations for

each of the two models (6000 simulations in total). We compared the simulation outcomes of both modelsusing four measures:

244

*i*) the frequency at which each bee experienced exploitation competition (i.e. flower visits when the reward
has already been collected by another bee) and interference competition (i.e. flower visits when two bees
encounter);

249 *ii*) the similarity index  $SI_{ab}$  between successive foraging bouts by the same bee;

250

251	iii) the degree of resource partitioning among bees, based on network modularity $Q$ (Pasquaretta and Jeanson
252	2018; Pasquaretta et al. 2019). $Q$ is calculated using the computeModules function implemented in the R
253	package 'bipartite' (Dormann et al. 2008) using the DIRTLPAwb+ algorithm recently developed by Beckett
254	(2016). Although $Q$ ranges between 0 (the two bees visit the same flowers) and 1 (the two bees do not visit
255	any flower in common), the comparison of modularity between networks requires normalization because the
256	magnitude of modularity depends on network configuration (e.g., total number of flower visits) (Dormann
257	and Strauss 2014; Beckett 2016). For each network, we computed :
258	
259	(4) $Q_{norm} = \frac{Q}{Q_{max}}$
260	
261	where $Q_{max}$ is the modularity in a rearranged network that maximizes the number of modules (Pasquaretta
262	and Jeanson 2018).
263	
264	<i>iv</i> ) an index of collective foraging efficiency, $QL_{group}$ , computed for each foraging bout <i>i</i> , to estimate the
265	cumulated efficiency of all foraging bees, as:
266	
267	(5) $QL_{group,i} = \frac{\sum_{j=1}^{n} QL_{j,i}}{QL_{optimal}}$
268	
269	where $QL_{j,i}$ is the route quality of the individual j during bout i, n the number of bees and $QL_{optimal}$ is the
270	maximum value of all the possible sums of individual route qualities.
271	
272	To assess whether the trends across foraging bouts obtained from simulations with two bees
273	differed between models (Fig. 1) and type of environments (Fig. S2), we applied GLMMs for each of the

274 following response variables: (i) frequency of competition types (Poisson error distribution), (ii)  $SI_{ab}$ 

(Binomial error distribution), (*iii*)  $Q_{norm}$  (Binomial error distribution) and (*iv*)  $QL_{group}$  (Binomial error distribution). In each model, we used a random structure to account for bee identity nested in flower arrays (i.e. 100 simulations of each spatial array for each model). To statistically compare the trends across foraging bouts we estimated the marginal trends of the models, as well as their 95% confident intervals using the *emtrends* function in 'emmeans' package (Lenth 2018). When the 95% confidence intervals of the estimated trends included zero, the null hypothesis was not rejected. Statistical models were run using the g*lmer* function in 'lme4' package (Bates *et al.* 2014).

282

### 283 **RESULTS**

### 284 Simulations with one forager

285 We first tested the ability of our models to replicate trapline formation by real bees, by comparing

simulations with one forager to published experimental data. In the positive array of flowers (Fig. S1A), real

287 bees developed routes of increasing quality and similarity with experience (GLMM route quality: Estimate =

288  $0.128 \pm 0.021$ , P < 0.001; GLMM route similarity: *Estimate* = 0.148 \pm 0.027, P < 0.001). Simulated bees with

289 positive and negative reinforcements (model 2) or positive reinforcement only (model 1) developed routes of

similar qualities as real bees (Fig. 2A; GLMM  $_{model 1}$ : *Estimate* = 0.024 ± 0.021, *P* = 0.244; GLMM  $_{model 2}$ :

*Estimate* =  $0.017 \pm 0.021$ , *P* = 0.416). In both models, the simulated bees also increased route similarity as

292 much as real bees (Fig. 2B; GLMM  $_{model 1}$ : *Estimate* = 0.048 ± 0.027, *P* = 0.074; GLMM  $_{model 2}$ : *Estimate* =

293  $0.044 \pm 0.027$ , P = 0.105). In the negative array of flowers (Fig. S1B), a qualitative match between the

simulations of both models and the experimental data was also observed (Fig. S5; see Appendix S2). Thus,

295 overall, positive reinforcement was sufficient to replicate the behavioral observations. The addition of

296 negative reinforcement had no major effect on route quality and similarity.

297

#### 298 Simulations with two foragers

299 Exploitation and interference competition

300 Having calibrated our models with one forager, we next explored conditions for the emergence of resource

301 partitioning within pairs of foragers. Here experimental data are not available for comparison. We first

302 analyzed exploitation competition by quantifying the frequency of visits to non-rewarding flowers by each

303 bee during each foraging bout. Simulated bees with positive and negative reinforcements (model 2)

304 decreased their frequency of visits to non-rewarding flowers with time, irrespective of the type of

305 environment (Fig. 3A; GLMM <sub>one patch</sub>: *Estimate* = -8.94e-03, 95%CI = -9.16e-03 | -8.71e-03; GLMM <sub>two patches</sub>:

306 *Estimate* = -1.88e-02, 95%*CI* = -1.91e-02 | -1.86e-02; GLMM three patches: *Estimate* = -1.05e-02, 95%*CI* = -

307 1.07e-02 | -1.03e-02). However, with positive reinforcement only (model 1), bees behaved differently in the

308 different environments. In the one patch environment, bees decreased their visits to non-rewarding flowers

309 (Fig. 3A; GLMM <sub>one patch</sub>: *Estimate* = -4.26e-03, 95%CI = -4.47e-03 | -4.05e-03), whereas in the two and three

310 patch environments, bees tended to increase their visits to non-rewarding flowers (Fig. 3A; GLMM two patches:

311 *Estimate* = 6.27e-03, 95%*CI* = 6.09e-03 | 6.44e-03; GLMM three patches: *Estimate* = 6.65e-03, 95%*CI* = 6.46e-

312 03 | 6.84e-03).

We analyzed interference competition by quantifying the number of interactions on flowers at each foraging bout between the two bees. Bees with positive and negative reinforcements (model 2) decreased their frequency of encounters on flowers with time irrespective of the type of environment (Fig. 3B; GLMM  $_{one patch}$ : *Estimate* = -1.53e-02, *95%CI* = -1.61e-02 | -1.45e-02; GLMM <sub>two patches</sub>: *Estimate* = -1.66e-02, *95%CI* = -1.73e-02 | -1.59e-02; GLMM <sub>three patches</sub>: *Estimate* = -1.01e-02, *95%CI* = -1.07e-02 | -0.94e-02). Here again, bees with positive reinforcement only (model 1) behaved differently in the different environments. In the one

319 patch environment, bees decreased their frequency of encounters on flowers (Fig. 3B; GLMM one patch:

320 *Estimate* = -4.57e-03, 95%CI = -5.29e-03 | -3.85e-03), whereas in the two and three patch environments,

321 bees increased their frequency of interactions (Fig. 3B; GLMM two patches: Estimate = 1.05e-02, 95%CI =

322 1.01e-02 | 1.10e-02; GLMM three patches: *Estimate* = 9.16e-03, 95%*CI* = 8.64e-03 | 9.68e-03).

Thus overall, negative reinforcement was necessary for reducing exploitation and interference competition. By allowing bees to avoid empty flowers, negative reinforcement facilitated the discovery of new flowers and thus gradually relaxed competition. In the absence of negative reinforcement, both types of competition increased in environments with several flower patches.

327

328 *Route similarity* 

329 We analyzed the tendency of bees to develop repeated routes by comparing the similarity between flower

330 visitation sequences of consecutive foraging bouts for each individual (Fig. 3C). In the two models, bees

331 increased route similarity through time in all types of environments (Fig. 3C model 1; GLMM one patch:

 $\begin{aligned} & 332 \quad Estimate = 8.68e-02 \ , \ 95\%CI = 8.56e-02 \ | \ 8.80e-02 \ ; \ \text{GLMM}_{\text{two patches}}: \ Estimate = 8.07e-02 \ , \ 95\%CI = 7.96e-\\ & 333 \quad 02 \ | \ 8.19e-02 \ ; \ \text{GLMM}_{\text{three patches}}: \ Estimate = 6.77e-02 \ , \ 95\%CI = 6.66e-02 \ | \ 6.88e-02; \ \text{Fig. 3C model 2}; \\ & 334 \quad \text{GLMM}_{\text{one patch}}: \ Estimate = 6.80e-02 \ , \ 95\%CI = 6.69e-02 \ | \ 6.91e-02 \ ; \ \text{GLMM}_{\text{two patches}}: \ Estimate = 5.40e-02 \ , \\ & 335 \quad 95\%CI = 5.29e-02 \ | \ 5.50e-02 \ ; \ \text{GLMM}_{\text{three patches}}: \ Estimate = 3.56e-02 \ , \ 95\%CI = 3.46e-02 \ | \ -3.65e-02). \ \text{Note} \\ & \text{however that the presence of negative reinforcement (model 2) reduced the final level of route similarity. In \\ & \text{however that the presence of negative reinforcement (model 2) reduced the final level of route similarity. In \\ & \text{however that the presence of negative reinforcement (model 2) reduced the final level of route similarity. In \\ & \text{however that the presence of negative reinforcement (model 2) reduced the final level of route similarity. In \\ & \text{however that the presence of negative reinforcement (model 2) reduced the final level of route similarity. In \\ & \text{however that the presence of negative reinforcement (model 2) reduced the final level of route similarity. In \\ & \text{however that the presence of negative reinforcement (model 2) reduced the final level of route similarity. In \\ & \text{however that the presence of negative reinforcement (model 2) reduced the final level of route similarity. In \\ & \text{however that the presence of negative reinforcement (model 2) reduced the final level of route similarity. In \\ & \text{however that the presence of negative reinforcement (model 2) reduced the final level of route similarity. In \\ & \text{however that the presence of negative reinforcement (model 2) reduced the final level of route similarity. In \\ & \text{however that the presence of negative reinforcement (model 2) reduced the final level of route similarity. In \\ & \text{however that the presence of negative reinforcement (model$ 

337 these conditions bees learned to avoid revisits to empty flowers and showed greater variation in their

338 visitation sequences, as a result of searching for new flowers.

339

#### 340 *Resource partitioning*

341 We analyzed the level of resource partitioning by quantifying the tendency of the two bees to use different

342 flowers. With positive and negative reinforcements (model 2), pairs of bees showed a significant increase of

resource partitioning with time for all types of environments (Fig. 3D; GLMM one patch: Estimate = -3.82e-02,

344 95%*CI* = 3.69e-02 | 3.95e-02; GLMM two patches: *Estimate* = 2.91e-02, 95%*CI* = 2.78e-02 | 3.04e-02; GLMM

345 three patches: *Estimate* = 2.66e-02, 95%*CI* = 2.53e-02 | 2.78e-02). With positive reinforcement only (model 1),

346 bees showed an increase of resource partitioning in environments with one patch (Fig. 3D; GLMM one patch:

347 *Estimate* = 3.24e-02, 95%CI = 3.11e-02 | -3.36e-02), and a decrease in environments with two or three

348 patches (Fig. 3D; GLMM two patches: *Estimate* = -1.06e-02, 95%CI = -1.19e-02 | -0.93e-02; GLMM three patches:

Estimate = -8.08e-03, 95% CI = -9.40e-03 | -6.77e-03). In these patchy environments, negative reinforcement

350 facilitates the movements of bees between more distant flowers in different patches and thus the

351 establishment of spatially segregated routes.

352

#### 353 *Collective foraging efficiency*

354 To quantify the collective foraging efficiency of bees, we analyzed the capacity of the two foragers to reach

355 the most efficient combination of route qualities (i.e. minimum distance traveled by a pair of bees needed to

visit the 10 flowers ). With positive and negative reinforcements (model 2), pairs of bees increased their

357 cumulated foraging efficiency with time in all types of environments (Fig. 3E; GLMM <sub>one patch</sub>: *Estimate* =

358 3.05e-02, 95%*CI* = 2.91e-02 | 3.19e-02; GLMM two patches: *Estimate* = 3.05e-02, 95%*CI* = 2.91e-02 | 3.19e-02;

359 GLMM three patches: *Estimate* = 1.85e-02, 95%CI = 1.73e-02 | 1.98e-02). With positive reinforcement only

360 (model 1), bees increased their cumulated foraging efficiency in environments with one patch (Fig. 3E;

GLMM <sub>one patch</sub>: *Estimate* = 3.24e-02, 95%CI = 3.11e-02 | 3.37e-02), but decreased or maintained low levels in environments with two or three patches (Fig. 3E; GLMM <sub>two patches</sub>: *Estimate* = -5.05e-03, 95%CI = -6.33e-03 | -3.76e-03; GLMM <sub>three patches</sub>: *Estimate* = 3.07e-03, 95%CI = -1.42e-03 | 4.32e-03). Cumulated foraging efficiency thus follows the same pattern as resource partitioning. Note that bees can reach higher levels in environments with one patch than in environments with two or three patches because the probability to encounter new flowers within a patch is higher in environments with one patch.

367

### 368 **Discussion**

369 Central place foraging animals exploiting patchily distributed resources that replenish over time are expected 370 to develop foraging routes minimizing travel distances and interactions with competitors (Possingham 1989; 371 Ohashi and Thompson 2005; Lihoreau *et al.* 2016). Here we developed a cognitively plausible agent-based 372 model to explore the behavioral mechanisms leading to resource partitioning between traplining bees. In the 373 model, bees learn to develop routes as a consequence of feedback loops that modify the probabilities of 374 moving between flowers. While a positive reinforcement of route segments leading to rewarding flowers is 375 sufficient for resource partitioning to develop when resources are evenly distributed, a negative 376 reinforcement of route segments leading to unrewarding flowers is necessary in environments with patchily 377 distributed resources. 378 When considering bees foraging among uniformly distributed plant resources (one patch), the initial 379 probabilities to encounter each resource are similar (Fig. S6), and the likelihood of bees to discover new 380 rewarding resources after encountering unrewarding ones is high. Consequently, two bees are very likely, 381 over time, to learn non-overlapping foraging routes and show resource partitioning. However, in 382 environments with non-uniformly distributed resources (two or three patches), the added spatial complexity 383 can interfere with this process. The initial likelihood of moving between distant patches is relatively low. 384 Thus, the only implementation of positive reinforcement for route segments to rewarded flowers, often does 385 not enable bees to explore all possible patches, so that the paths of competing bees overlap and interfere 386 within a subset of the available patches. Adding a negative reinforcement for route segments to unrewarded 387 flowers increases aversion for these flowers, the spatial segregation of foraging paths between competing 388 bees and the collective exploitation of all available patches, even if the initial probabilities of moving to

distant patches are low. This interplay between the influences of positive and negative experiences at flowers
on the spatial and competitive decisions of bees is in accordance with behavioral observations (Lihoreau *et al.*

391 2016; Pasquaretta *et al.* 2019).

392 The need of negative reinforcement rules to enhance discrimination between different options or 393 stimuli is well known in both learning theory and behavioral studies (Beshers and Fewell 2001; Garrison et 394 al. 2018; Kazakova et al. 2020). This is especially notable in collective decisions making by groups of 395 animals and robots (Sumpter 2010), where negative feedbacks enable individuals to make fast and flexible 396 decisions in response to changing environments (Robinson et al. 2005; Seeley et al. 2012). At the individual 397 level, negative experiences also modulate learning. For both honey bees and bumblebees, adding negative 398 reinforcement to a learning paradigm (e.g. quinine or salt in sucrose solution) enhances fine scale color 399 discrimination (Avarguès-Weber et al. 2010) and performances in cognitive tasks requiring learning of rules 400 of non-elemental associations (Giurfa 2004). The insect brain contains multiple distinct neuromodulatory 401 systems that independently modulate negative and positive reinforcement (Schwaerzel et al. 2003) and the 402 ability of bees to learn negative consequences is well-established (Vergoz et al. 2007). Even so, the utility of 403 negative reinforcement to enhance efficient route learning and the consequences of this for the emergence of 404 effective resource partitioning has not been commented on previously. It may be that this is a general 405 phenomenon with applicability to other resource partitioning systems.

406 Our study implies that some very simple learning and interaction rules are sufficient for trapline 407 formation and resource partitioning to emerge in bee populations. Several improvements of the model could 408 be considered for future theoretical and experimental investigations of bee spatial foraging movements and 409 interactions, for instance by implementing documented inter-individual variability in cognitive abilities and 410 spatial strategies of bees (Chittka et al. 2003; Raine and Chittka 2012; Klein et al. 2017), the variability in 411 the nutritional quality of resources (Wright et al. 2018; Hendriksma et al. 2019) and the specific needs of 412 each colony (Kraus et al. 2019), or the well-known ability of bees to use chemical (Leadbeater and Chittka 413 2005) and visual (Dunlap *et al.* 2016) social information to decide whether to visit or avoid flowers. For 414 instance, it is well-known that foragers of many species of bees leave chemical cues as footprints on the 415 flowers they have visited (bumblebees and honeybees: Stout and Goulson 2001; several species of solitary 416 bees: Yokoi and Fujisaki 2009). Bees learn to associate the presence or absence of a reward to these 417 footprints and to revisit or avoid scented flowers (Leadbeater and Chittka 2011). Such a pheromone system is 418 an advantageous signal for all participants in the interaction (Stout and Goulson 2001). Our model suggests 419 that this additional information could significantly enhance the negative value experienced by a bee thus 420 increasing resource partitioning to the benefit of all bees coexisting in the patch (Appendix S3), even of 421 different species that learn to use these cues (Stout and Goulson 2001; Dawson and Chittka 2013). 422 Our agent-based model thus builds on previous attempts to simulate trapline foraging by individual 423 bees (Lihoreau et al. 2012b, Reynolds et al. 2013; LeMoël et al. 2019) and fills a major gap of current 424 pollinator population dynamic models (Becher et al. 2014; 2016; 2018). As such, it constitutes a unique 425 theoretical modeling framework to explore the spatial foraging movements and interactions of bees in 426 ecologically relevant conditions within and between plant patches. This holds considerable promising tool to 427 guide novel experiments in a wide range of pollinators exhibiting routing behavior. It can readily be used to 428 test predictions with more than two bees (see examples Video S1 and Appendix S3), several colonies, or 429 even different species of bees (e.g. honey bees) and may be used to complement current predictions about 430 pollinator population dynamics (Becher et al. 2014; 2016; 2018). Ultimately, the robust predictions of the 431 spatial movements and interactions of bees over large spatio-temporal scales, through experimental 432 validations of the model, has the potential to help illuminate the influence of bees on plant reproduction 433 patterns and pollination efficiency (Ohashi and Thomson 2009).

434

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# 602 Figure legends

603 Figure 1. Flowchart summarizing the agent-based models (1 and 2). Rectangles represent actions performed

604 by a bee. Diamonds indicate conditional statements. Arrows connect the different modules. Colored

605 rectangles describe the negative reinforcement rule specific to model 2. R code is available in GitLab:

606 https://gitlab.com/jgautrais/resourcepartitioninginbees/-/releases

607

608 Figure 2. Comparisons of route qualities (A) and similarities (B) between simulations and experimental data

609 for one forager (positive array of flowers as in Lihoreau *et al.* 2012b). Model 1: positive reinforcement only.

610 Model 2: positive and negative reinforcements. For each dataset, we show the estimated average trends

611 across foraging bouts (colored curves), along with their 95% CI (gray areas). For the sake of eye comparison,

612 in the simulation plot we represent estimated 95% confidence intervals for a random subsample of 7

613 simulated bees (N = 7 bees in Lihoreau *et al.* 2012b). Average trends were estimated over 100 simulation

fl4 runs, using GLMM Binomial model with bee identity as random effect (bee identity nested in simulation

615 identity for simulated data).

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617 Figure 3. Results of simulations with two foragers in environments with 10 flowers. Light blue: simulations 618 with positive reinforcement only (model 1). Dark blue: simulations with positive and negative reinforcement 619 (model 2). The x axis is the number of completed foraging bouts by the two foragers. The y axis represents 620 respectively: A) the estimated mean frequency of visits to empty flowers; B) the estimated mean frequency 621 of encounters on flowers; C) the similarity index  $SI_{ab}$  between two successive flower visitation sequences; 622 D) the index of resource partitioning  $Q_{norm}$  (0: both bees visit the same set of flowers; 1: bees do not visit any flower in common); E) the collective foraging efficiency index  $QL_{group}$ . Average trends for each model 623 624 are estimated across all types of environments (one patch, two patches and three patches; see Fig S2). 625 626

627

#### 629 Supplementary materials

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#### 631 Appendix S1. Sensitivity analysis of positive and negative reinforcements.

632 We ran a sensitivity analyses for the two main parameters: the positive and negative reinforcement. As we 633 had no a priori understanding of how the model behaved with different values of the two main parameters of 634 the model (positive and negative reinforcement), we explored the spectrum of potential outcomes with 635 different sets of parameters. We ran simulations on ranges of positive (1, 1.2, 1.4, 1.6, 1.8, 2) and negative (0, 636 0.1, 0.2, 0.3, 0.4, 0.5, 0.6, 0.7, 0.8, 0.9, 1) reinforcement factors for a total of 66 sets of parameters, for each 637 environment type explored (one, two and three patches, Fig. S2). We simulated 10 environments for each 638 environment type and computed 100 simulations of 50 foraging bouts per iteration (i.e. 1000 simulations per 639 environment type and set of parameters). 640 Since our observations focused on resource partitioning, we extracted the  $Q_{norm}$  index values for 641 these simulations and compared them. We plotted a heatmap showing the value of the mean  $Q_{norm}$  index at 642 the last foraging bout for each set of parameters and environment type (Fig. S3). In all types of environments, 643 the positive reinforcement has a strong effect on the final  $Q_{norm}$  index value. High values of resource parti-644 tioning are obtained for positive reinforcement values > 1.5. For the negative reinforcement factor, however, 645 it appears that high values of resource partitioning were obtained for negative reinforcement values larger 646 than 0.75. Negative reinforcement has little to no impact on the environment type with only one patch. 647 We also looked at how this same index evolved over successive foraging bouts, for each pair of 648 bees. Fig. S4 shows the dynamics of mean  $Q_{norm}$  index across 50 foraging bouts for each combination of 649 positive reinforcement (1.0, 1.2, 1.4, 1.6, 1.8, 2.0) and negative reinforcement (0, 0.1, 0.2, 0.3, 0.4, 0.5, 0.6, 650 0.7, 0.8, 0.9, 1) parameters, and each environment type (one patch, two patches, three patches). Higher val-651 ues of both positive and negative reinforcement factors most often lead to faster resource partitioning (with 652 some uncertainty due to the probabilistic nature of the model). Combinations of values in which the negative 653 reinforcement factor is missing (violet gradient curves) led to a decrease in partitioning. 654

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656 Appendix S2. Qualitative analysis for the negative array case.

657 We ran simulations with one forager to compare model outcomes to observational data using a second 658 reference field study (Woodgate et al. 2017). In this study, the authors used a 'negative array' (narrow 659 pentagon, Fig. S1B) in which the distance between flowers and the directionality of movement were 660 negatively linked, i.e. moving between nearest neighbor flowers required bees to make sharp deviations 661 between consecutive vectors. Four bumblebees were tested during 27 to 61 foraging bouts each (flower 662 visitation sequences were provided by Joe Woodgate). One of these bumblebees was tested over different 663 days and was therefore removed from the analyses. In these conditions, none of the bumblebees developed a 664 stable trapline, although all individuals significantly increased their foraging efficiency with time (e.g. 665 reduced travel distance and duration, increased similarity between two consecutive flower visitation 666 sequences).

667 The addition of a negative reinforcement had no major effect on route quality nor on route 668 similarity trends (Fig. S5). Model simulations showed good qualitative fit the traplining behavior observed in 669 real bees exploiting a negative array of flowers - i.e. there is a trend of increasing route similarities across 670 foraging bouts. Note however that the model tends to overestimate the bee ability to develop stable routes in 671 arrays of flowers where proximity and directionality are negatively linked (Woodgate et al. 2017). This 672 imperfect match could be due to the low amount of available experimental data (three individuals in 673 Woodgate et al. 2017). Alternately, the model does not integrate any kind of stochastic exploration so that at 674 each new step, bees do not provide the possibility to choose unknown spatial targets. Real bees, on the 675 contrary, show phases of stochastic explorations during and after trapline formation (Woodgate *et al.* 2017; 676 Kembro et al. 2019). Future experiments with more bees in a larger diversity of arrays of flowers will be 677 useful to further quantitatively calibrate the model.

678

### 679 Appendix S3. Predictions with more than two bees.

Here we explored the emergence of resource partitioning in groups of 5 bees, and how this varies as a consequence of number of available flowers and number of foraging bees in environments of 20, 25, 30, 40, 50, 70 and 100 flowers, thus encompassing a gradient of competition pressures from conditions where there is not enough flowers for all bees (20) to conditions where there is four times more flowers than necessary for all bees (100). For simplicity, in all these environments, flowers were evenly distributed (i.e. environment with one patch). The positive reinforcement factor was set at 1.5, and the negative reinforcement factor was

686	set at 0.75. For each flower density, we generated 10 environments, and ran 100 simulations of 100 foraging
687	bouts, for a total of 1000 simulations per density value. We extracted the resource partitioning index $(Q_{norm})$
688	at each foraging bout. Unsurprisingly, the mean final $Q_{norm}$ was higher in environments with most flowers
689	(Fig. S7). Plotting the mean final $Q_{norm}$ (final foraging bout) as a function of the number of available
690	flowers confirms that bees converge to a plateau where increasing the number of flowers only provokes
691	increase of partitioning (i.e. around 50 flowers) (Fig. S7).
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693	
694	Figure S1. Arrays of artificial flowers for the experimental dataset. A. 'Positive array' (regular pentagon) in
695	which the distance between flowers and the directionality of movement are positively linked, i.e. nearest
696	neighbor flowers can be linked using lowest angular deviation between consecutive vectors. Modified from
697	Lihoreau et al. (2012b). B. 'negative array' (narrow pentagon) in which the distance between flowers and the
698	directionality of movement are negatively linked, i.e. moving between nearest neighbor flowers requires bees
699	to make sharp deviations between consecutive vectors. Modified from Woodgate et al. (2017)
700	
701	Figure S2. Examples of simulated environments. Distribution of 10 flowers (gray circles) and a nest (black
702	pentagon) in three environments defined by different levels of resource patchiness. A flower patch was
703	characterized by: 1) a uniform distribution of flowers, 2) a lower distance between flowers within the patch
704	than between all flowers from different patches.
705	
706	Figure S3 Heatmap showing the mean $Q_{norm}$ Index value after 50 foraging bouts (mean over 1000 simula-
707	tions on 10 arrays of the same environment type), for each combination of positive reinforcement (1.0, 1.2,
708	1.4, 1.6, 1.8, 2.0) and negative reinforcement (0, 0.1, 0.2, 0.3, 0.4, 0.5, 0.6, 0.7, 0.8, 0.9, 1) parameters, and
709	for each environment type (one patch, two patches, three patches). For simplicity we inverted the values of
710	negative reinforcement here. 0 being models without negative reinforcement.
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Figure S4. Dynamic of the mean  $Q_{norm}$  Index across foraging bouts for each combination of positive (1.0, 1.2, 1.4, 1.6, 1.8, 2.0) and negative (0, 0.1, 0.2, 0.3, 0.4, 0.5, 0.6, 0.7, 0.8, 0.9, 1) reinforcement factors and

714	for each environment type (one patch, two patches, three patches). For simplicity we inverted the values of
715	negative reinforcement here. 0 being models without negative reinforcement.

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719	Figure S5. Qualitative comparisons of route qualities (A) and similarities (B) between simulations and
720	experimental data (negative array of flowers as in Woodgate et al. 2017) for one forager. Light blue:
721	simulations with positive reinforcement only (model 1). Dark blue: simulations with positive and negative
722	reinforcement (model 2). For each dataset, we show the estimated average trends across foraging bouts
723	(colored curves), along with their 95% CI (gray areas). For simplicity, in the simulation plot we only
724	represent a subsample of three random bees with their estimated 95% CI. Average trends were estimated
725	using GLMM Binomial model with bee identity as random effect (bee identity nested in simulation identity
726	for simulated data).
727	
728	Figure S6. Boxplot representing the distribution of the initial probabilities of visiting each flower in different
729	environments (one patch with uniformly distributed flowers; two patches with five uniformly distributed
730	flowers per patch; three patches of five, three and two uniformly distributed flowers; see examples Fig. S1).

731 Median (solid horizontal line), first and third quartiles (box) and maximum and minimum values are

732 represented.

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734	Figure S7. Evolution of the mean final $Q_{norm}$ index (after 100 foraging bouts) as a function of increase
735	resources availability. The model run has the positive reinforcement factor set at 1.5, and the negative rein-
736	forcement factor set at 0.75 with 5 bees in environments of one patch.
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738	Video S1. Example of simulation of five bees foraging in an environment with one patch of 50 flowers. Both
739	positive and negative reinforcement rules are implemented (model 2). Bees performed 100 foraging bouts.
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# 760 **Figure 1**



**Figure 2** 







798 Figure S2













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Foraging bout









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