

28 altruistic; explaining this behavior is conceptually challenging because one animal provides a
29 benefit to another at a cost to itself. The seminal contribution of W.D. Hamilton (2, 3) provided a
30 simple yet powerful framework for understanding altruism. This has since become known as
31 Hamilton's Rule and posits that a behavior or trait will be favored by selection, when $rb-c>0$,
32 where c is the fitness cost to the actor, b is the fitness benefit to the recipient, and r is their
33 genetic relatedness. Since then numerous studies across a very diverse range of organisms from
34 bacteria and yeasts to ants and mammals have demonstrated that the necessary conditions for
35 altruism are upheld (4, 1).

36 Crucial to understanding the evolution of altruism has been determining how animals
37 distinguish kin from non-kin because r must be >0 to satisfy Hamilton's rule. An unlikely tool
38 for studying altruism, it turns out, are parasites, the very antithesis of altruistic behavior.
39 Parasites have evolved ways to break the code within kin groups, benefiting from their altruism,
40 despite being completely unrelated to the donor ($r=0$). The classical example is the Common
41 Cuckoo (*Cuculus canorus*) that exploits the parental behavior of other bird species through egg
42 mimicry and selfish behaviors by the chick. In societies such as ants, where altruism is expressed
43 from sibling to sibling, diverse parasites ranging from other ants to beetles, flies, caterpillars and
44 even mollusks have evolved ways to break the code to act like cuckoos (5,6). For example,
45 caterpillars (i.e. *Maculinea rebeli*) perfectly mimic the chemical profile of larval ants and are
46 carried into the nest by foraging workers, where they are then fed colony resources and consume
47 the larval and egg stage ants (5,7,8,9). The general term for such organisms is social parasite and
48 studying their chemical ecology and behavior has provided many insights into the mechanisms
49 by which altruism works.

50 Although the color pattern of a cuckoo egg or the chemical cues of caterpillars entering ant nests
51 are complex, it is conceptually easy to imagine how they evolved from 'so simple a beginning'
52 (10). Indeed Darwin considered both cuckoo and socially parasitic ants, conjecturing that each
53 emerged from non-parasitic progenitors (Chapter 7). More difficult to conceptualize is whether a
54 parasite entering the body of an altruist can be recognized as a parasite since it is within the body
55 of a colony member who presents kin recognition cues to the rest of the colony. Here test
56 whether uninfected members of an ant colony can recognize siblings infected by a specialized
57 endoparasite. We will use as our model the entomopathogenic fungus, *Ophiocordyceps*

58 *unilateralis*, a highly specialized parasite of worker ants that manipulates host behavior to
59 achieve transmission.

60 In recent years a number of studies have demonstrated that species within the complex *O.*
61 *unilateralis s.l.* infect worker ants and adaptively manipulate host behavior by causing infected
62 individuals to leave the colony and bite into vegetation before dying (11). The function of such
63 manipulation is to provide a platform for spore release as post-mortem the fungus transitions
64 from growing with the body to growing externally, forming a large stalk from which spores are
65 produced and released onto the forest floor (12). Because the time from exposure and infection
66 to behavioral manipulation is between 9 days and 3 weeks, during which time the infected ant is
67 within the colony, this model offers the potential to examine how the colony responds to infected
68 individuals. In this study we develop a system for within colony observation of infected and
69 healthy individuals. We show that despite their status, infected ants are neither evicted from the
70 colony nor prevented from leaving the nest to die and spread disease to their siblings.

71 **Results**

72 **Infected ants receive food from siblings**

73 We first set out to determine if infected ants received food from their siblings inside the nest
74 area. We infected ten ants per colony in four different colonies of *Camponotus castaneus* species
75 with a strain of *Ophiocordyceps unilateralis* fungus, which naturally infects this species in the
76 wild (13). Ten ants were removed from the stock colony and injected with 1 μ l *O. unilateralis* in
77 solution with Grace's insect media supplement with 10% fetal bovine serum (FBS). A further 10
78 ants were sham treated, and inject with 1 μ l of Graces +FBS media. Both infected and sham
79 treated ants were maintained together with 15 additional untreated individuals in a wooden
80 chamber of volume 14.93 \pm 0.53 cm³ placed within a cage of 451.61cm² that served as a foraging
81 area and contained sand. These ants were given water and 10% glucose *ad libitum*. We began
82 continuous data recording from the third day post injection until day 18, moment at which there
83 was the least amount of infected individuals alive. Ant behavior was recorded inside the nest
84 with GoPro Heron 2 cameras for 24 hours/day. We then scored behavior from playback on
85 screens. We first focused on food exchange between individuals since our hypothesis was that
86 infected ants receive food at a different rate from non-infected ants. Worker ants cannot eat solid

87 food but instead exchange liquids in a process called trophallaxis. We followed 17 focal
88 individuals from one colony for a total observation 976.24 hours. Using a mixed effect linear
89 model, programmed in R, we used trophallaxis duration as a function of day post infection and
90 using ant identification as a random effect we found no significance $p\text{-value}=0.5156$. We found
91 no significant patterns in differences in either duration or count of trophallaxis (Figure 1). Our
92 quantification of observations includes only within nest exchange of liquids. We did however
93 also observe that ants infected by *O. unilateralis* would receive trophallaxis from nest mates
94 when outside the nest. Individuals were even fed in the minutes and hours before they were
95 behaviorally manipulated to ascend vegetation before biting bite into the twigs we provided and
96 dying. We therefore found no evidence that infected ants were refused food from other colony
97 members.

98 **Infected ants are not attacked by siblings**

99 It might be expected that infected ants are attacked more often. Although an infected ant can
100 only infect other ants following its own death and the subsequent growth of the stalk from its
101 head, it is possible that the increase of fungal cells within the worker ants changes some aspect of
102 its phenotype, such as smell, causing other ants to attack it. Because aggression might be rare and
103 fleeting we observed the behavior of 2 colonies (30 ± 5 ants/colony) 24 hours/day for 18 days
104 (still in progress). We saw no aggression between untreated control ants and either infected or
105 sham treated ants. We conclude based on continuous observations over the entire course of
106 infection that infected ants are not attacked by their siblings.

107 **Infected ants are initially distanced from colony members but this declines over time**

108 Although we found no aggression there were subtle indications of infected ant segregation. To
109 measure this we applied spatial point process approaches to within-nest ant locations. To
110 examine spatial interaction behavior between healthy and infected ants, we examine the K-cross
111 function (14) between healthy and infected ants, which is the expected number of infected ants
112 that would be found within a distance d of each healthy ant.

113 Our null hypothesis is that ant infection status does not affect the tendency to group together or
114 avoid other ants. We used a nonparametric permutation test (15) to test for significant deviation
115 from this null behavior. In Figure 2 we show the observed K-cross function (in red) for all the

116 data, as well as for days 3, 6, and 9, together with 1000 K-cross functions (in black) simulated
117 from the null model by randomly permuting the labels (e.g., healthy, infected, or sham) of the
118 ants 1000 times, and calculating the K-cross function between healthy and infected ants under
119 each of these permutations. Significant deviation from the null model is indicated by an
120 observed K-function that lies on the edges (tails) of the envelope of K-functions simulated under
121 the null model, and empirical p -values can be computed by considering the rank of the observed
122 K-function within the envelope. Significant deviation from null behavior only occurs day 3 at
123 short spatial distances (less than 8 millimeters; p -value <0.05), where the observed K-function
124 lies on the lower tail of the permuted K-functions, suggesting spatial segregation at small spatial
125 scales.

126 To examine this potential small-scale ant interaction behavior between healthy and infected ants,
127 we found the nearest neighbor to each ant at each time point. We then tested for deviation from
128 the null assumption that ants are equally likely to have any other ant as nearest neighbor by again
129 permuting the ant labels at each time point and recording the permuted label of each ant's nearest
130 neighbor. Table 1 shows the proportion of healthy ants with an infected nearest neighbor over
131 all observed days, and for day 3, day 6, and day 9, together with the empirical p -value under the
132 null, obtained using 1000 permutations of ant labels at each time point of observation. We see
133 significant differences on day 3 and when we pool all the data together.

134 **Discussion**

135 Our data suggests no aggression towards infected individuals. We also see no distinct differences
136 in the mean duration or counts of trophallaxis between infected and uninfected individuals (p -
137 value=0.5156). Our linear mixed effect model shows trophallaxis had no relation with treatment
138 and trophallaxis is stochastic. Other studies have used trophallaxis as a tool to study social
139 immunity, making similar observations to the ones we have made here, yet their results are an
140 increase the amount of trophallaxis that occurs 24 hours after infection with a fungal pathogen
141 (16, 17). An important factor these papers did not take into account is time, they only made
142 observation 24 hours after the infection, this experiment on the other hand observed the ontogeny
143 of behavior within the nest. By using a one chamber scenario in a cage where ants were able to
144 freely move and interact with one another enables us to observe more naturalistic interactions
145 that has been lacking in the ant-pathogen research.

146 Our data suggest there are no shifts in behavior towards infected individuals, suggesting healthy
147 individuals are unable to detect *Ophiocordyceps* infection. The spatial point process analysis
148 revealed that by and large there is no evidence for spatial segregation of infected ants. The only
149 exception was the slight differences in spatial segregation between healthy individuals and those
150 infected at small spatial scales on day 3 of the infection, but not on days 6 and 9. These minute
151 changes in spatial arrangement could be caused by changes in individual infected ant behavior
152 and are not likely to be indicators of social exclusion, which we would expect to increase in
153 strength with time from infection. We did not test for any relationship between spatial
154 segregation and the identity of the focal individuals in relation to who perform the most
155 trophallaxis. Data collection for both trophallaxis and distance data were collected on Colony 2
156 the sample sizes we have may be masking the effect of *Ophiocordyceps* on the infected
157 individuals.

158 Within nest distance observations has been done before by using images to determine spatial
159 fidelity and time budgets of *Leptothorax acervorum* (18, 19), their observations did not take into
160 account how pathogens may change social dynamics within a colony nor did they do continuous
161 behavioral observations. We were able to observe rare interactions and behaviors that have
162 previously not been described. Being able to follow individuals through time and space lends
163 itself to be a powerful tool for further understanding the ontogeny of behavior within infected
164 individuals. Although our trophallaxis and distance results were not significant we can still
165 progress our understanding between uninfected individuals and those being parasitized.

166 In order to establish if these results are caused by the evolutionary history between
167 *Ophiocordyceps* and host we should observe non-coevolved pathogen species. These behavioral
168 assays enable us to further explore the role of parasites in not only the behavioral of the single
169 host, but also in the colony host. The ability to combine behavioral observation and spatial
170 dynamics as a tool to make very fine detailed observations enables us to further tease out the
171 dynamics of the colony and those infected. Another powerful tool that we could add to this type
172 of behavioral assay is chemical cues, such as cuticular hydrocarbons.

173 Chemical communication is the method of communication within an ant colony (20,21).
174 Therefore individual odor changes could signify caste allocation (22) and colony members could
175 also use it as a methods to determine health. Using continuous, detailed observations and

176 cuticular hydrocarbons would give insight into how these infected individuals are perceived by
177 their nest mates.

178

179 **Methods and Materials**

180 **Ant collection and stock colony maintenance-** Ants were collected in South Carolina during
181 October 2012. Colonies were collected by following foragers to nest sites that were then dug up.
182 Colony 1 consisted of sexuals, brood and about 120 workers; collected 10/4/2012. Colony 2
183 formed by 100 workers and brood; collected 10/5/2012. Colony 3 formed by 100 workers and
184 brood; collected 10/3/2012. Colony 4 has queen, brood and 120 workers. Colonies were
185 maintained by providing them sugar water and water ad libitum and changed once a week. From
186 these colonies we collected individuals to run our experiment on.

187 **Infection techniques-** *O. unilateralis* Infections were done as described in de Bekker *et al.*,
188 2014/submitted. Single fungal colonies were placed in a sterile 2 mL tube with two 8/32 inch
189 metal balls (Wheels Manufacturing Inc.) and 200 μ L Grace's medium (Sigma) freshly
190 supplemented with 10% FBS (PAA laboratories Inc.). The colony tissue was lysed using
191 TissueLyser II (Qiagen) at room temperature for 60 sec. at 30 freq/sec. This process enabled
192 us to obtain single hyphae used at a mean concentration of $3.9 \times 10^7 \pm 1.1 \times 10^7$ hyphae/ml for
193 infection. Infections were done by injecting 1 μ L hyphal solution with a laser pulled 10 μ L
194 micropipette (Drummond) and aspirator tube (Drummond) into the thorax underneath the front
195 legs. Sham treatments were done in similar fashion using 1 μ L medium without hyphae (23).

196 **Treatments and individual identification-** Subcolonies were made of fifteen healthy, ten
197 injected with Grace's+FBS media used for *Ophiocordyceps* growth in the laboratory and another
198 ten were injected with *Ophiocordyceps* plus media. These individuals were collected by their
199 colonies by agitating the housing tubes within each colony and collect the individuals that
200 from the population. In order to follow individuals through time we used a dot system, each
201 individual had a different dot pattern painted on its body. We used an edding® 751 paint marker
202 to label the ants we used for the experiment.

203 **Behavioral observation set up-** We created sub-colonies containing 35 worker ants within a
204 wooden cage with a volume of 14.93 ± 0.53 cm³. In order to make 24 hour observations we used

205 a Go Pro camera (Hero 2 with IR lens) and an IR lamp was used for nocturnal observations. The
206 camera was located on top of the colony chamber and removed to change size video card three
207 times a day.

208 **Trophallaxis-** There was only one observer who made the observations of trophallaxis to reduce
209 observer bias. Trophallaxis was classified as starting when labrum was exposed and distended
210 between the two individuals. The event was as over when the mouth parts separated and the
211 individual parted ways. We observed a total of 976.24 hours of video for Colony 2 in order to
212 determine the amount of trophallaxis focal individuals were receiving on days 3-9,12,15 and 18
213 in trial one. A total of seven Infected individuals and five sham treated and five healthy ants
214 were followed over the course of the daylight session (7.68 ± 0.32 hours per day) on days
215 3,4,5,6,7,8,9,12,15 and 18 post injection. We analyzed days 3-9 since these are the days we have
216 most infected individuals inside the nest (Figure 1). The chambers in which ants were placed did
217 not restrict individuals to stay within the nest, we were only able to record behaviors for those
218 present within the nest at the time of observation.

219 **Aggression-** There was only one observer who took note of aggressive behavior to reduce
220 observational bias. The videos were observed in fast forward and stopped if any abnormal
221 behavior occurred. Colony 2 has a total of 76.77 hours observed and no aggression has been
222 seen. Further observations will be made in other colonies to see if non-aggression holds.

223 **Distance data collection-** Screen shots were made for every ten minutes of observation during
224 the day period (8.24 ± 0.34 hours per day). Individuals were identified using paint marks. We
225 then used an R program (version 2.15.1; created by Kezia Manlove) that calculated both pair-
226 wise distances and x-y coordinates for the individuals within the chamber. On average there were
227 24 ± 2 individuals inside the chamber, we recorded point distances on all the individuals visible to
228 us on days 3, 6, 9 and 12 for a total of 4,758 x-y coordinates and 61,000 pair-wise data points.

229 **Distance data analysis-** We focused on days 3, 6 and 9 (8.09 ± 0.43 hours per day) when
230 analyzing the data. We used point process models, which take a set of point locations in window
231 of space. Each of these points can be labeled with a mark to indicate a certain type or class. In
232 this case, the ant locations are the points of interest, the window of observation is the nest, and
233 the mark of each ant is their infection status; either untreated, infected, or sham treated. In point

234 process statistics, the interaction between points can be measured using a summary statistic
235 called the K-function (14, 15). For a given distance d , the K-function gives the expected number
236 of additional ants to be found within a radius of d of a focal ant.

237 Functions to compute the K-cross function from healthy to infected ants, and the nearest
238 neighbor to each ant at each time point, were created in R. The K-cross function finds the
239 average number of infected ants within a specified distance of a healthy ant, with the average
240 being over all healthy ants in the chamber at each time point, and over all time points within the
241 specified day. The permutation tests for the nearest-neighbor analysis were carried out by
242 permuting the labels (healthy, infected, or control) of ants in the chamber at each time point and
243 re-computing the nearest neighbor of each ant.

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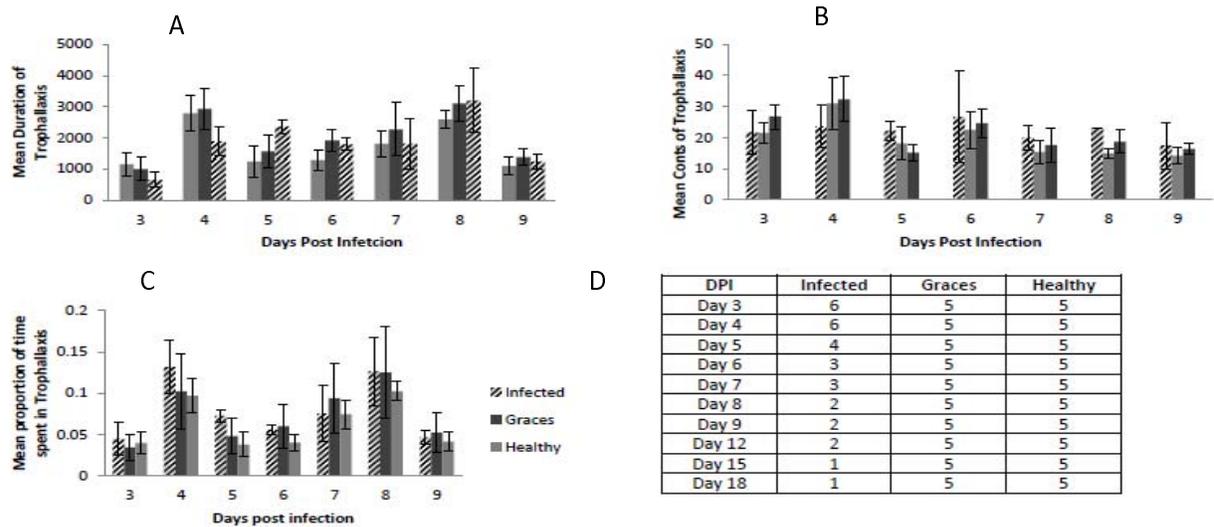
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301 **GRAPHS AND TABLES**

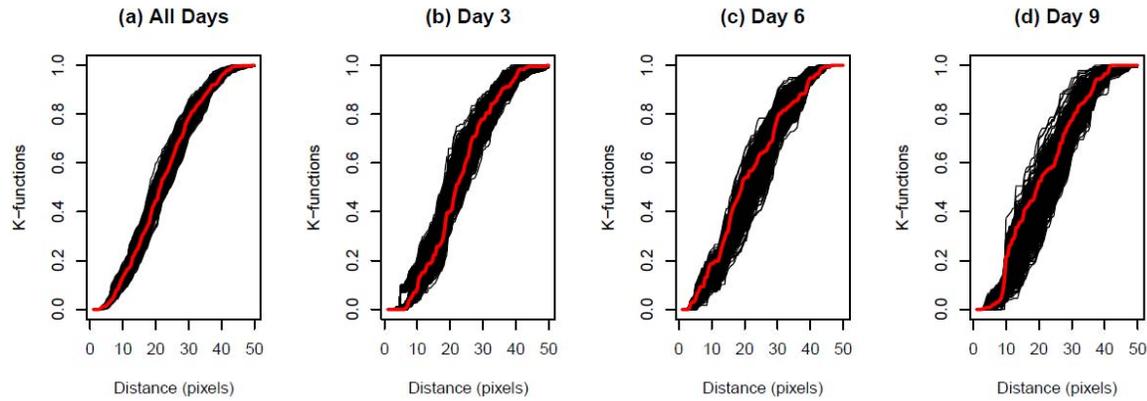
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303 **Figure 1-** Trophallaxis data collected from videos on days 3-9, error bard represent the standard
 304 error of the data. We were unable to see any significance difference between treated individuals
 305 and healthy. (A) Shows no differences between infected, graces and healthy ants although we do
 306 see an interesting pattern of duration increase on day 4 and 8. (B) Mean count of trophallaxis
 307 changes slightly throughout the days we have observed. (C) The proportion of time spent in
 308 trophallaxis.



309

310 **Figure 2- K function analysis**



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312

313

314 **Table 1-** Nearest neighbor analysis we can see there is a significant difference between healthy
315 and infected on when looking at all three days combined and only on day 3.

Time	Proportion	Permutation Test p-value
All Days	0.110	0.002
Day 3	0.108	0.004
Day 6	0.208	0.48
Day 9	0.104	0.24

316