

1 Gut transplants from bees fed an antipathogenic pollen diet do not confer pathogen resistance to  
2 recipients

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29 **Abstract**

30  
31 Pollinators are threatened by diverse stressors, including microbial pathogens such as *Crithidia*  
32 *bombi*. Consuming sunflower pollen dramatically reduces *C. bombi* infection in the bumble bee  
33 *Bombus impatiens*, but the mechanism behind this medicinal effect is unclear. We asked whether  
34 diet mediates resistance to *C. bombi* through changes in the gut microbiome. We hypothesized  
35 that sunflower pollen changes the gut microbiome, which in turn reduces *Crithidia* infection. To  
36 test this, we performed a gut transplant experiment. We fed donor bees either a sunflower pollen  
37 treatment or buckwheat pollen as a control treatment, and then inoculated recipient bees with  
38 homogenized guts from either sunflower-fed or buckwheat-fed donor bees. All recipient bees  
39 were then fed a wildflower pollen diet. Two days after the transplant, we infected recipients with  
40 *C. bombi*, and two days later, we provided another donor gut transplant. To quantify infection,  
41 we performed both fecal screens and dissections of the recipient bees. We found no significant  
42 differences in *C. bombi* infection intensity or presence between bees that received sunflower-fed  
43 microbiomes versus buckwheat-fed microbiomes. This suggests that sunflower pollen's effects  
44 on pathogen resistance are not mediated by gut microbiota.

45 **Main text**

46

47 Pollinators have widespread impacts on our environment and economy [1, 2], but they  
48 are susceptible to microbial pathogens, some of which are implicated in their declines [3]. One  
49 such pathogen is *Crithidia bombi* ('*Crithidia*' hereafter), which commonly infects bumble bees  
50 (*Bombus* spp.). *Crithidia* negatively impacts bumble bee fitness; it reduces learning and foraging  
51 ability [4], and under stressed conditions increases worker mortality [5] and reduces colony-  
52 founding by nearly 40% [6]. It is transmitted fecal-orally, on flowers or within colonies [7], [8].

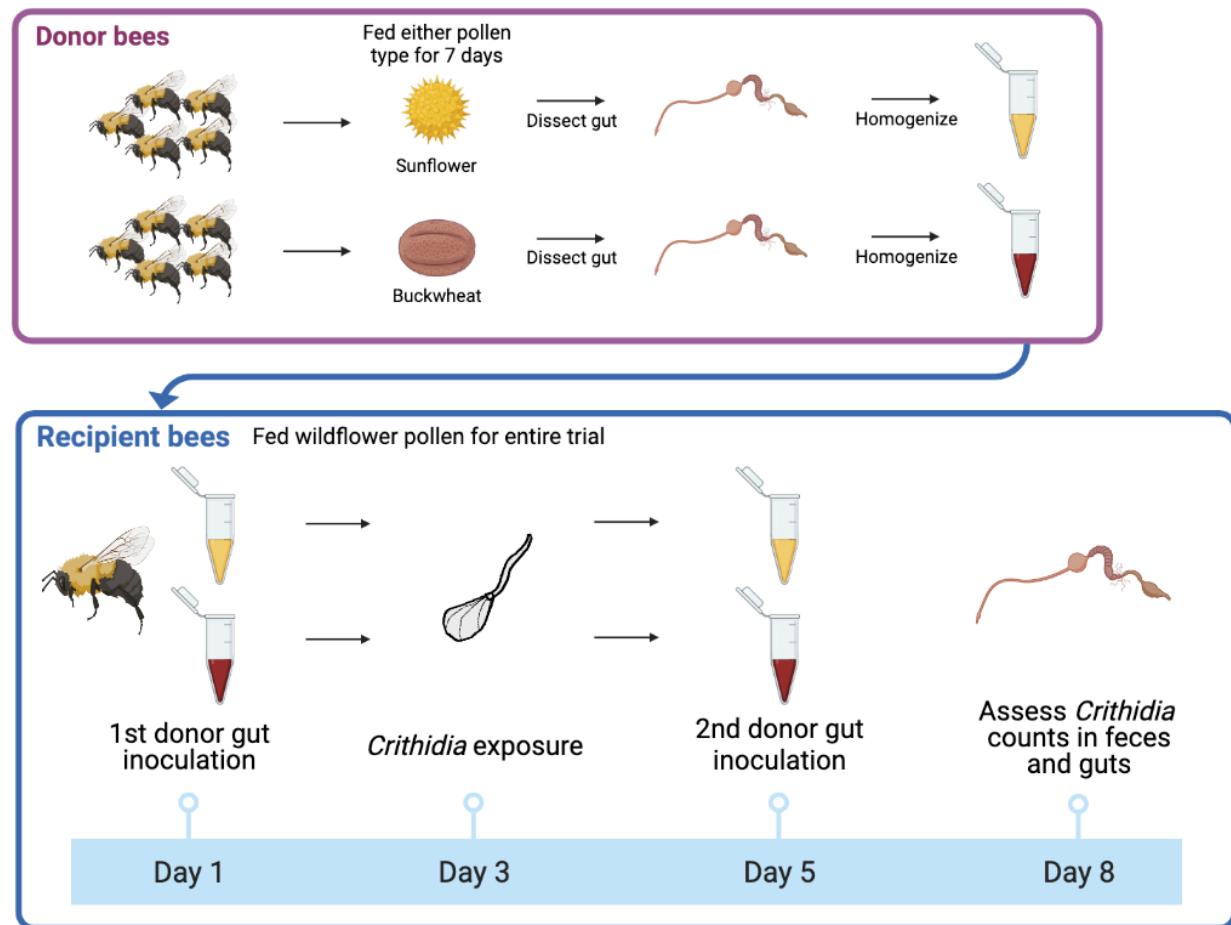
53 The gut microbiome can play a significant role in bee health; variation in the bumble bee  
54 gut microbiome predicts susceptibility to *Crithidia* infection [9]. Low *Crithidia* counts are  
55 associated with high microbial diversity, large gut bacterial load, and the presence of certain  
56 bacterial taxa such as *Apibacter*, *Lactobacillus* Firm-5, and *Gilliamella* spp. [10]. Furthermore,  
57 gut microbial communities are a stronger predictor of host susceptibility to *Crithidia* than host  
58 genetics [9].

59 In addition to the gut microbiome, diet can play an important role in bee resistance to  
60 pathogens. For example, sunflower pollen (*Helianthus annuus*) has consistently and dramatically  
61 reduced *Crithidia* infection in *Bombus impatiens* workers [11], and sunflower abundance on  
62 farms was associated with reduced worker infection and increased colony reproduction [12].  
63 Currently, two studies have linked diet to changes in bee gut microbiomes [13, 14], but none  
64 have connected these changes to pathogen resistance. We asked if the anti-pathogenic effects of  
65 a sunflower pollen diet are mediated by changes in the gut microbiome. Our study aims to shed  
66 light on the interactions between diet, pathogens, and the gut microbiome, which will improve  
67 our understanding of how floral resources affect bee health and may inform conservation  
68 strategies for these important pollinators.

69 Feeding bees different diets and observing effects on pathogen resistance does not  
70 distinguish whether resistance was due to direct effects of diet, or mediated by the diet-induced  
71 changes in the gut microbiome. To disentangle these effects we used a gut transplant experiment,  
72 inoculating a recipient host with the whole dissected gut contents of a donor host. This approach  
73 is a relatively inexpensive and straightforward way to assess effects of the gut microbiome on  
74 host phenotype, without requiring sequencing or culturing. Previous studies have used gut

75 transplants to determine the effects of gut microbial communities on *Crithidia* infections [9, 10, 15], but have not addressed the potential role of diet in these interactions.

77 Experiments took place at the University of Massachusetts Amherst campus from May to 78 September 2022. We fed “donor” *B. impatiens* workers either sunflower pollen or buckwheat 79 pollen and then dissected out their guts and fed them to experimental “recipient” bees (Fig. 1) 80 that were then inoculated with *Crithidia*. We hypothesized that recipients that received gut



81  
82 **Fig. 1** Diagram of the experimental procedure. We treated recipient bees with a gut solution from 83 the dissected donor bees. On day 3 we inoculated them with *Crithidia* and on day 5 we gave 84 them a second gut treatment. We performed fecal screens and gut dissections on day 8 to assess 85 infection

86  
87 microbes from sunflower-fed bees would exhibit lower *Crithidia* infections compared to those 88 that received gut microbes from buckwheat-fed bees. We conducted two rounds of the 89 experiment, using 36 recipient bees in the first round and 48 in the second, for a total of 84 bees

90 (42 per microbiome treatment; bees were maintained individually and are the unit of replication).  
91 We outline our basic methods and statistical analysis below; additional details can be found in  
92 the supplemental material.

93 We removed donor bees from their natal colony and assigned them to sunflower or  
94 buckwheat pollen diets. We placed groups of 5-10 bees (with the same diet and natal colony) in  
95 separate containers (15 cm x 15 cm x 9 cm; Biobest LTD., Ontario, Canada) in a dark incubator  
96 at 27 °C, and fed them their respective diet for 7 d. Then, we removed three bees per group,  
97 dissected their guts, pooled together the supernatant solution, and mixed it with equal parts 50%  
98 sucrose to create the microbiome solution.

99 Next, we removed recipient bees from the same natal colonies and randomly assigned  
100 them to a “buckwheat-fed microbiome” or “sunflower-fed microbiome” treatment. We used non-  
101 sterile adult workers who were presumably already inoculated with gut microbes from their  
102 nestmates, so that our treatments are an augmentation of the existing gut microbiome. We fed  
103 recipient bees 15  $\mu$ l of microbiome solution of their assigned treatment from donors originating  
104 from the same colony, observing until they consumed all the solution. We then isolated each bee  
105 in a 16 oz. deli cup and fed them a wildflower pollen ball and 15 ml of 30% sucrose solution,  
106 replaced every 2-3 days. Two days after the initial gut transplant treatment, we inoculated  
107 recipient bees with 15  $\mu$ l of *Crithidia* inoculum (600 cells/ $\mu$ l; 9000 cells total). Two days later,  
108 we administered a second dose of sunflower and buckwheat microbiome treatments to the  
109 recipients.

110 Five days after *Crithidia* inoculation, we quantified infection via both fecal screens and  
111 gut dissection. Since volumes of fecal samples were often too small to assess cell counts on a  
112 hemocytometer, we diluted feces with Ringer’s solution (Sigma-Aldrich). We counted moving  
113 *Crithidia* cells in 0.02  $\mu$ l aliquots of both diluted fecal samples and homogenized gut solution at  
114 400X using a compound light microscope. After counting cells in 0.02  $\mu$ l diluted feces, we  
115 calculated cells/ $\mu$ l feces after accounting for dilution, while in guts we retained counts per 0.02  
116  $\mu$ l as our response. We collected the right forewing of each bee and measured marginal cell  
117 length using ImageJ to estimate bee size [16].

118 We used R version 4.2.1 [17] for all analyses and the glmmTMB function [18] for all  
119 models. We analyzed four responses in separate models: *Crithidia* presence and cell counts in 1  
120  $\mu$ l of feces and *Crithidia* presence and cell counts in 0.02  $\mu$ l of gut solution. We used

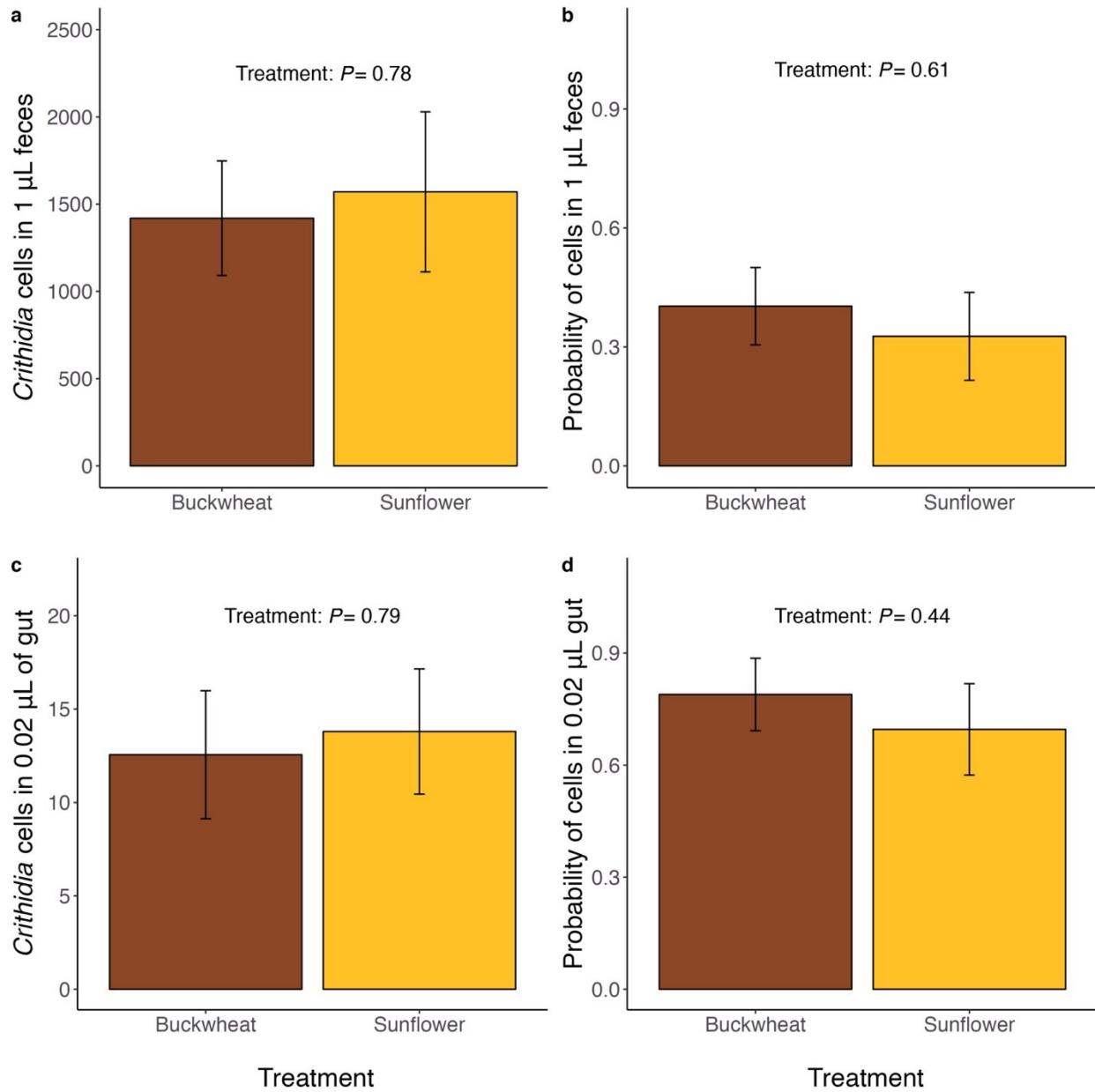
121 microbiome treatment and bee size (estimated as marginal cell length) as fixed effects, with  
 122 *Crithidia* inoculation date and colony as random effects in all initial models, including fecal  
 123 volume as an additional fixed effect in the fecal models. We used the DHARMA package [19] to  
 124 assess model fit, and AIC for model selection (AICcmodavg package [20]). Microbiome  
 125 treatment was always retained in the final models since that was our variable of interest. Colony  
 126 was retained as a random effect in the models for cells per  $\mu\text{l}$  feces and probability of cell  
 127 presence in 0.02  $\mu\text{l}$  of gut solution. Sample sizes for sunflower and buckwheat microbiome  
 128 treatments were 24 and 32 respectively for the fecal models, and 30 and 34 for the gut models.

129 We found no significant difference in *Crithidia* cell counts between buckwheat-  
 130 microbiome and sunflower-microbiome bees in feces or gut solutions (Fig. 2a and 2c, Table 1a  
 131 and 1c). There was also no significant difference in probability of *Crithidia* presence in feces or  
 132 gut solution (Fig. 2b and 2d, Table 1b and 1d). Surprisingly, higher fecal volume corresponded  
 133 with a higher probability of cells in feces (Table 1d), but lower cell counts (Table 1c). Larger  
 134

Terms	a. Cells/0.02 $\mu\text{l}$ gut		b. Probability of cells in gut		c. Cells/ $\mu\text{l}$ feces		d. Probability of cells in feces	
	$\chi^2$	p	$\chi^2$	p	$\chi^2$	p	$\chi^2$	p
Treatment	0.069	0.793	0.602	0.438	0.078	0.780	0.262	0.609
Bee size	2.962	0.085	5.024	<b>0.025</b>	0.098	0.754	7.182	<b>0.007</b>
Fecal Volume	n/a	n/a	n/a	n/a	5.536	<b>0.019</b>	6.482	<b>0.011</b>

135  
 136 **Table 1.**  $\chi^2$  and p values from final models predicting (a) *Crithidia* cells per 0.02  $\mu\text{l}$  of gut  
 137 solution, (b) the probability of *Crithidia* cells in 0.02  $\mu\text{l}$  of gut, (c) *Crithidia* cells per  $\mu\text{l}$  of feces,  
 138 and (d) the probability of *Crithidia* cells in 1  $\mu\text{l}$  of feces. Bold text indicates  $p < 0.05$

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 140



141

142 **Fig. 2** Effect of sunflower pollen on (a) *Crithidia* cells per  $\mu\text{l}$  of feces, (b) the probability of  
 143 *Crithidia* cells in 1  $\mu\text{l}$  feces, (c) *Crithidia* cells per 0.02  $\mu\text{l}$  of gut solution, and (d) the probability  
 144 of *Crithidia* cells in 0.02  $\mu\text{l}$  of gut solution in *Bombus impatiens* workers. Means estimated by a  
 145 generalized linear model; error bars indicate standard error back-transformed by emmeans

146

147 bees also had lower probability of *Crithidia* presence in feces (Table 1d) and a non-significant  
 148 trend for lower counts in gut solution (Table 1a).

149 Our results suggest that sunflower pollen's ability to reduce *Crithidia* infection in *B.*  
 150 *impatiens* is not due to changes in the gut microbiome. There may have been diet-induced

151 changes in the gut microbiome, but they did not significantly reduce *Crithidia* infection. We used  
152 workers with already established gut microbial communities, due to high mortality of newly  
153 emerged bees in prior versions of the experiment. This may have made it more difficult for the  
154 microbiome treatments to alter the existing microbial community. Additional experiments using  
155 newly emerged bees without an established microbiome may show stronger treatment effects.  
156 However, recent data show that *Bombus impatiens* workers fed sunflower pollen exhibited  
157 similar gut bacterial communities, in both composition and diversity, to those fed a wildflower  
158 pollen mix (Fowler et al., in prep). Compared to the control, bees fed sunflower pollen did not  
159 have higher prevalence of the bacterial taxa associated with lower *Crithidia* infections found in  
160 previous research [10]. This finding is consistent with our results that gut bacterial communities  
161 transplanted from sunflower-fed bees had no effect on *Crithidia* infection in recipients. Together,  
162 these results suggest that sunflower pollen does not alter the bee gut microbial community in  
163 ways that reduce *Crithidia* infection.

164 The mechanism underlying sunflower pollen's dramatic effect on *Crithidia* infection is  
165 still not established. However, new research suggests that the spiny sunflower pollen exine is  
166 responsible [21], which may mean the effect is mechanical rather than chemical. Uncovering the  
167 mechanism behind sunflower pollen's medicinal properties will help further our knowledge of  
168 how certain plants may benefit declining pollinator populations.

169

170 **References**

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230

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234

235 *Competing Interests*

236 The authors have no competing interests.

237

238 *Author contributions*

239 LSA and AEF conceived and designed the experiment. RTY collected the data and carried out  
240 the research. AEF analyzed the data. RTY wrote the first draft of the manuscript and all authors  
241 contributed to subsequent drafts. All authors read and approved the final manuscript.

242

243 *Data availability*

244 All data and code will be made publicly available at Scholarworks@UMass upon acceptance for  
245 publication.

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