

1 **How do sunflower pollen mixtures affect survival of queenless microcolonies of bumblebees (*Bombus***
2 ***impatiens*)?**

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10 **Abstract**

11 The high pollen and nectar yield of Asteraceae flowers combined with the abundance and diversity of these
12 plants should make them an important pollen source for both specialist and generalist bees. However, studies have
13 found Asteraceae pollen to be a poor diet for some generalist bees, possibly owing to nutrient deficiencies or the
14 presence of secondary metabolites or digestive barriers that prevent nutrient assimilation. Mixing pollens could
15 allow bees to exploit these unfavourable pollens, if the different pollen sources help to complement nutrient
16 deficiencies or alleviate the effects of toxic secondary metabolites. In our study, we examined how the proportion of
17 sunflower (*Helianthus annuus*, Asteraceae) pollen in the diet of captive-reared bumblebees (*Bombus impatiens*)
18 affects the survival of bees in queenless microcolonies. Bees fed sunflower pollen had significantly shorter lifespans
19 than bees fed broad bean (*Vicia faba*, Fabaceae), rapeseed (*Brassica napus*, Brassicaceae), or Cucurbitaceae pollen.
20 However, survival on mixed pollen diets containing 50% sunflower pollen was as great as that on non-sunflower
21 diets, which suggests that the other pollens were able to compensate for the low nutritive quality of the sunflower
22 pollen. Due to agricultural intensification and a loss of wildflowers, farmland monocultures (e.g., sunflower crops)
23 can be important floral resources for bumblebees. Our study suggests that providing alternative floral resources of
24 high nutritive quality could help mitigate the potential harmful effects of a monofloral sunflower diet on
25 bumblebees.

26

27 **Keywords:** bumblebees, amino acids, protein, nutrition, pollen mixing, sunflower (*Helianthus annuus*)

28 **Electronic supplementary material**

29 The online version of this article contains supplementary material, which is available to authorized users.

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37 **Introduction**

38 Pollen can be a difficult resource for animals to exploit, yet bees (Hymenoptera: Anthophila) have
39 diversified on a pollen-based diet and most bee species rely on pollen as their sole source of protein (Willmer 2011).
40 However, all pollens are not equivalent as dietary substrates for bees. Several studies have shown that the pollens
41 used by certain specialist (oligolectic) bee species can be inadequate for development of non-specialists (Williams
42 2003; Praz et al. 2008; Sedivy et al. 2011; Haider et al. 2013; Haider et al. 2014), suggesting that physiological
43 adaptations may be required to tolerate “unfavourable” properties of various pollens. It is also well established that
44 pollen diets deficient in protein can negatively affect larval bee development (Levin and Haydak 1957; Regali and
45 Rasmont 1995; Roulston and Cane 2002) and shorten adult lifespan (in honeybees; Schmidt et al. 1987).
46 Furthermore, not all pollens collected by bees contain the 10 essential amino acids listed by De Groot (1953) for
47 honeybees; tryptophan and phenylalanine in particular are sometimes absent (reviewed by Roulston and Cane 2000),
48 as in dandelion pollen, *Taraxacum officinale* (Asteraceae; Auclair and Jamieson 1948). Some studies have found
49 that Asteraceae pollen possesses unfavourable properties that render it difficult to utilize for bees that are not
50 specialized on this pollen type (Levin and Haydak 1957; Guirguis and Brindley 1974; Sedivy et al. 2011). This
51 could be the reason that so many generalist bees appear to avoid Asteraceae pollen (Müller 1996; Williams 2003;
52 Müller and Kuhlmann 2008)—despite the fact that species in this family are often abundant, presenting copious
53 floral rewards and supporting numerous oligolectic species (*Helianthus* spp.: Hurd et al. 1980; Asteroideae: Müller
54 and Kuhlmann 2008).

55 Mixing pollens might be a strategy for generalist bees to exploit unfavourable pollens if the different pollen
56 sources help to complement nutrient deficiencies or alleviate the effects of toxic secondary metabolites (Eckhardt et
57 al. 2014). For example, *Megachile rotundata* (= *M. pacifica*; Megachilidae) larvae developed significantly more
58 quickly on a mixed pollen diet of *Melilotus* and *Medicago sativa* (both Fabaceae) than on a diet containing only
59 *Melilotus* pollen (Tasei and Masure 1978). Similarly, *Osmia cornuta* larvae developed well on mixed pollen
60 provisions containing their normal host plant *Sinapis arvensis* (Brassicaceae) and up to 50% *Ranunculus acris*
61 (Ranunculaceae) pollen, whereas a diet of solely *R. acris* pollen was toxic to most *O. cornuta* larvae (Eckhardt et al.
62 2014). Nevertheless, the nutritional quality of mixed pollen diets for bumblebee health and colony development
63 remains poorly understood (Baloglu and Gurel 2015; Moerman et al. 2016a), despite widespread interest in bee
64 conservation (e.g., Goulson et al. 2015) and pollinator management strategies (e.g., providing wildflower plantings
65 in agroecosystems; reviewed by Williams et al. 2015). Mixed pollen diets could benefit generalist (polylectic) bees

66 by minimizing the chances of vitamin, mineral, and/or protein deficiencies while simultaneously diluting the toxic
67 secondary metabolites found in certain pollen species (Eckhardt et al. 2014).

68 In this study, we examined how the proportion of sunflower (*Helianthus annuus*, Asteraceae) pollen in the
69 diet of captive-reared bumblebees (*Bombus impatiens*) affects the survival of worker bees. Previous studies have
70 demonstrated that sunflower pollen has deleterious effects on both honeybees (*Apis mellifera*: Schmidt et al. 1995)
71 and a European bumblebee species (*B. terrestris*: Regali and Rasmont 1995; Tasei and Aupinel 2008a), although the
72 precise mechanism of effect is unknown. We were interested in whether mixing sunflower pollen with other pollen
73 species could improve the quality of the pollen diet (e.g., through nutrient complementation or dilution of toxic
74 substances). Our mixed pollen diets contained either 0%, 25%, or 50% sunflower pollen, with an equal mixture of
75 Cucurbitaceae, rapeseed (*Brassica napus*, Brassicaceae) and broad bean (*Vicia faba*, Fabaceae) pollen making up the
76 rest of the diet. Each pollen type was also included as a monofloral pollen treatment in the experiment (i.e., broad
77 bean, Cucurbitaceae, rapeseed, and sunflower). We assessed the performance of *B. impatiens* microcolonies fed each
78 pollen diet. In queenless bumblebees microcolonies, one worker acts as “queen” by developing oocytes, suppressing
79 ovary development of subordinate workers, and laying male eggs (Cnaani et al. 2002; Cnaani et al. 2007) these
80 microcolonies make excellent systems for testing the nutritive value of different pollen diets (Génissel et al., 2002;
81 Tasei and Aupinel, 2008a; Vanderplanck et al., 2014; Moerman et al., 2016a; Moerman et al., 2017), and the results
82 can be extrapolated to queenright colonies (*B. terrestris*, Tasei and Aupinel, 2008b). We had three response
83 variables: i) worker mortality, ii) initiation of honeypot construction (a normal behaviour of growing colonies and
84 microcolonies), and iii) egg production. We also compared the amino acid composition of the four pollen types (i.e.
85 broad bean, Cucurbitaceae, rapeseed, and sunflower) to determine if any diet was deficient in essential amino acids
86 and, therefore, the potential for nutrient complementation among diets.

87 **Methods**

88 ***Study Species***

89 Five colonies of captive-reared *Bombus impatiens* Cresson were obtained from Biobest (Leamington,
90 Ontario, Canada) and fed a honeybee-collected, multifloral pollen mixture (Hawkins Honey Bee Pollen, Rockwood,
91 Ontario). The pollen was ground and mixed with deionized water to form pollen balls, and each colony received one
92 pollen ball (~0.5 g) per day. BIOGLUC® (Biobest Biological Systems), a nectar substitute, was included with each

93 colony and bees were fed ad libitum on this syrup solution. The colonies were kept at room temperature (21–22 °C)
94 at the University of Ottawa and were used as sources for the experimental microcolonies (described below).

95 ***Pollen Treatments***

96 Each bumblebee microcolony (described below) was assigned systematically to one of seven pollen
97 treatments (Table 1). The three SFM (“sunflower mixed”) diets contained either 0%, 25%, or 50% sunflower pollen
98 (97% *Helianthus annuus* and 3% other Asteraceae species [*Rudbeckia* sp., *Coreopsis* sp. and *Heliopsis* sp.]), with an
99 equal mixture of Cucurbitaceae (30% summer squash [*Cucurbita pepo*], 70% watermelon [*Citrullus lanatus*]),
100 rapeseed (*Brassica napus*), and broad bean (*Vicia faba*) pollen making up the rest of the diet. Each pollen type was
101 also included as a monofloral pollen treatment in the experiment (i.e. broad bean, Cucurbitaceae, rapeseed, and
102 sunflower).

103 Honeybee-collected pollen of four types (broad bean, rapeseed, sunflower, and watermelon) was obtained
104 from Changge Ruifenfang Bee Products Company (Changge City, Henan Province, China) and washed to remove
105 nectar residues. Three samples of each pollen type were stained with basic fuchsin and approximately 100 pollen
106 grains were examined under a light microscope to confirm pollen identity and purity (Kearns and Inouye 1993). The
107 samples were all found to be over 95% pure with only trace levels (i.e., < 5%) of other pollen species found in any
108 of the samples. Nectar-derived sugars can make up 40% of the dry weight of bee-collected pollen (Todd and
109 Bretherick 1942; reviewed by Roulston and Cane 2000), and analyses that ignore the nectar contribution are likely
110 to underestimate the concentration of protein in the pollen (Weiner et al. 2010). We therefore suspended 20 g of
111 pollen at a time in 800 mL of deionized water and filtered it through 6 µm cellulose filter paper (Little Chalfont,
112 Buckinghamshire, UK) using a vacuum pump to remove the nectar residues (see Kitaoka and Nieh 2009). Pollen
113 was then air-dried before being stored in a –20°C freezer. Pollen from *Rudbeckia* sp., *Coreopsis* sp., *Heliopsis* sp.,
114 and summer squash (*Cucurbita* sp.) was hand-collected from greenhouse-grown plants at the University of Ottawa
115 and added to the commercial pollen after the latter had been washed.

116 Pollen balls were made by combining either 1 g or 0.5 g of washed pollen with 10–15 drops of 35% w/w
117 sucrose solution; variation in the number of drops was due to the fact that different amounts of solution were
118 required to form pollen balls, depending on pollen type. We do not know if these minor differences in sucrose
119 content affected pollen consumption rates (we did not measure this response variable in our study), but the small
120 differences in the amount of solution added (maximum difference approximately 0.23 mL or 0.081g of sucrose, in

121 an environment with freely available sugar syrup) make an effect unlikely. The pollen balls were coated in honeybee
122 wax to prevent desiccation and to encourage egg production (cf. Gradish et al. 2013). Pollen balls were stored in the
123 freezer prior to use.

124 *Queenless Microcolony Set-Up and Experimental Design*

125 Pupal clumps of worker-caste bees were collected from source colonies and stored in plastic containers.
126 Bees were placed in microcolonies of 4–6 bees within 24 hours of their emergence (Table 1). Microcolonies were
127 housed in clear delicatesse containers (~650 mL, base radius = 4.6 cm, opening radius = 5.5 cm, height = 8.2 cm)
128 that were wrapped in duct tape to keep the bees in darkness and reduce stress (photos of microcolony set-up are
129 provided in Online Resource 1; design is adapted from Gradish et al. 2013). The base of each container was
130 removed and replaced with either plastic mesh (0.635 cm or 0.4 cm) or 18×16 nylon window-screen mesh. A plastic
131 painter pot (~15 mL) filled with 35% sucrose solution was placed underneath the mesh in a second deli container. A
132 cotton wick (1 cm diameter) was secured in the painter pot through a hole cut in the lid. The wick soaked up the
133 sucrose solution and allowed the bees to access the solution through the mesh base of their deli container. When
134 microcolonies were first established, honey—which is naturally scented—was placed on the tip of cotton wicks to
135 encourage the bees to feed. The sugar solution was changed every other day and the wicks were replaced every four
136 days. Within 24 hours of microcolony establishment, bees were given a 1 g pollen ball in a 6.35 cm diameter dish.
137 An additional 0.5 g pollen ball was provided one week later and replaced every four days over a four-week period.
138 The original 1 g pollen ball was removed if it remained uneaten at least seven days after microcolony establishment
139 to prevent the development of mold.

140 Microcolonies were assessed every day to note bee deaths, pollen consumption, honeypot initiation, and
141 egg production. No larvae were produced during the experiment. Microcolonies were kept for six weeks (42 days),
142 after which all remaining bees were culled. Over the course of the experiment, only 25 bees (10.4%) were culled;
143 thus, the majority died of other causes before the end of the 6-week period. In total, 43 microcolonies (3–13
144 microcolonies from each source colony) and 242 bees were included in the experiment, but only 237 were included
145 in the analysis. Two bees were omitted since they were accidentally killed during microcolony observations; another
146 bee died after she had escaped from her microcolony; and two bees were omitted since they were mistakenly added
147 after each of their microcolonies was established. Although there was variation in the number of microcolonies per

148 source colony, the only colony that significantly differed from others in terms of bee performance was the one with
149 only three microcolonies, and results were robust to the exclusion of the microcolonies from this source colony.

150 *Amino Acid Analysis*

151 The amino acid profiles of the four pollen types (broad bean, Cucurbitaceae, rapeseed, and sunflower mix;
152 n = 3 of each) were analyzed at the SPARC BioCentre of the Hospital for Sick Children (Toronto, Ontario, CA).
153 Each sample underwent three analyses: a standard amino acid analysis (AA analysis) and a free amino acid analysis
154 (FAA analysis), which both excluded cysteine and tryptophan, as well as a tryptophan analysis (Trp analysis), which
155 included both bound and free tryptophan. Cysteine was not quantified since this is not an essential amino acid for
156 bees (De Groot 1953). The Water Acquity UPLC system was used to analyze the amino acids, and the detection
157 limit for all three analyses was 25 pmol (Bidlingmeyer et al. 1984; Heinrikson and Meredith 1984; Cohen and
158 Strydom 1988). This system consists of a Binary Solvent Manager Module, a Sample Manager Module, a TUV
159 Detector Module and a Waters Acquity UPLC BEH C18 column (2.1 X 100 mm) (Milford, Massachusetts, USA).
160 Prior to analysis, each sample was pulverized and thoroughly mixed with a mortar and pestle. Approximately 0.0100
161 g of the pollen samples was used for each analysis. For the AA analyses, the samples were hydrolyzed in 6N HCl
162 with 1% phenol for 48 hours at 110°C. The Trp analyses required all samples to be hydrolyzed in 2.4N NaOH for 24
163 hrs at 110°C. For the FAA analyses, all samples were redissolved in 450 µL of 0.1M HCl. In addition, 50 µL of 25
164 µmoles/µL of norleucine were added to all samples of all analyses as an internal standard. Samples were dried under
165 vacuum using a centrifugal evaporator (Tomy CC-181 Centrifugal Concentrator with a Sargent-Welch Model 8821
166 Vacuum pump). The samples were then re-dried, derivatized and dissolved in sample diluent (pH 7.40). An aliquot
167 of the diluted sample was then injected into the column, running on a modified PICO-TAG gradient. Data were
168 collected, stored and processed using Waters Empower 3 Chromatography software.

169 *Statistical Analysis*

170 All statistical analyses were performed in R (version 3.3.3; R Core Team, 2017).

171 *Bee Survival and Microcolony Performance*

172 Cox proportional hazard (henceforth Cox PH) models were used to compare bee survival on the different
173 pollen treatments ('coxme' function, 'coxme' package; Therneau 2015). Random terms were included in the model
174 for both source colony and microcolony. The data were analyzed first with all data included (model 1) and second
175 (model 2) excluding microcolonies that did not consume visible amounts of pollen (as well as bees that died prior to
176 pollen consumption) and two outlier microcolonies. Specifically, two (of 6) microcolonies (12 bees) from the broad

177 bean treatment were strong outliers since they contained 3 and 4 bees respectively that survived until the end of the
178 experiment as well as several bees (3 and 2 respectively) that died earlier. Consequently, the proportional risk of
179 death for the bees in these two microcolonies, as well as the hazard risk relative to the other pollen treatments,
180 varied with time (Schoenfeld residuals; all bees: $\chi^2 = 5.65$, $P = 0.018$; only pollen-consuming bees: $\chi^2 = 16.0$, $P <$
181 0.0001)—inconsistent with the proportional-hazards assumption. We present here the results of model 1 (including
182 all data), but the results for both models are included in Online Resource 2, and we mention whenever results
183 differed qualitatively between models. A Cox PH model was also used to compare time to honeypot construction
184 across the different pollen treatments, with source colony as a random term. This model was consistent with the
185 assumption of proportional hazards (Schoenfeld residuals: $\chi^2 = 2.35$, $P = 0.89$), and all 43 microcolonies were
186 included in this analysis.

187 *Pollen Amino Acids*

188 The amino acid profiles (summed across AA and FAA for each amino acid except tryptophan) of the four
189 different pollen types (broad bean, Cucurbitaceae, rapeseed, and sunflower) were compared using non-metric
190 multidimensional scaling (NMDS) with the ‘metaMDS’ function in the ‘vegan’ package (Oksanen et al. 2017; cf.
191 Vanderplanck et al. 2014). Two NMDS ordinations were performed, both using Manhattan dissimilarity matrices, 2
192 dimensions, and 20 runs. These parameters were sufficient to reach a stress level less than 0.05 and model
193 convergence. Manhattan distances were used since this dissimilarity metric takes into account the sums of the
194 absolute differences between objects (e.g., pollen samples) across all variables (e.g., amino acids) and does not
195 ignore zero values (e.g., two samples containing zero amino acids of any kind would be considered identical in
196 terms of Manhattan distance) (Quinn and Keogh 2002). The first NMDS was based on absolute amounts of each
197 amino acid (i.e., μg of each amino acid per mg of dry sample); the second used proportions (i.e., each amino acid
198 expressed as a percent [by mass] of total amino acid content). Permutational multivariate analyses of variance
199 (perMANOVA; 1000 permutations) using the ‘adonis’ function were then performed to compare pollen types.
200 Pairwise permutation MANOVAs with a Bonferroni correction were also performed because the global test
201 indicated that pollen types differed significantly (‘pairwise.perm.manova’ function, ‘RVAideMemoire’ package;
202 Hervé 2017). Prior to these analyses, we verified homogeneity of group covariances using permutational
203 multivariate homogeneity of group dispersions tests (PERMDISP; 1000 permutations) with the ‘betadisper’
204 function. To determine if certain amino acids were representative of a particular pollen type, indicator values for

205 each amino acid (based on proportions of total amino acid content) were calculated ('indval' function, 'labdsv'
206 package; Roberts 2016).

207 **Results**

208 ***Bee Survival and Microcolony Performance***

209 There was no evidence of pollen consumption in 10 microcolonies (~23% of the total 43 microcolonies)
210 containing 53 bees. In addition, 15 bees died before any pollen was consumed in the remaining 33 microcolonies.
211 All 25 bees that survived until the end of the 6-week period (42 days) were from microcolonies with pollen
212 consumption. None of the bees in the 100% sunflower pollen treatment survived until the end of the 6-week period.
213 Both Cox PH models for bee survival revealed significantly shorter lifespans of bees provided 100% sunflower
214 pollen (median lifespan 19.5 d) than those in all other pollen treatments (median 20–27 d; $P < 0.05$) except 25%
215 sunflower mixed pollen (19.5 d; Fig. 1; Table 2); however, the difference between 100% sunflower and broad bean
216 was only marginally significant in model 2 ($P = 0.089$; Online Resource 2). Bees provided a mixed diet containing
217 50% sunflower pollen lived approximately 10 days longer than bees in the 100% sunflower treatment. In addition,
218 lifespans of bees provided the 25% sunflower mixed pollen were shorter than those in the 50% sunflower mixed
219 treatment (significantly so only in model 2; $P < 0.021$), the broad bean pollen treatment (model 1 only; $P < 0.028$),
220 and the rapeseed pollen treatment (model 2 only; $P < 0.037$). During the experiment, 19 of the 43 microcolonies
221 (44%) constructed honeypots, and microcolonies provided broad bean pollen were significantly more likely to
222 construct honeypots than those provided the 25% sunflower mixed pollen treatment ($P < 0.038$) and marginally
223 more likely to construct honeypots than the 50% sunflower mixed pollen treatment ($P = 0.058$) and the 100%
224 sunflower pollen treatment ($P = 0.095$) (Fig. 2; Table 3). Egg masses were visible in four microcolonies: one that
225 was provided 0% mix pollen, another that was given Cucurbitaceae pollen, and two that were given broad bean
226 pollen. However, all eggs that were laid were later destroyed or eaten, and no male bees emerged from any of the
227 microcolonies.

228 ***Pollen Amino Acids***

229 Free amino acids accounted for 3.49–6.43% of the total amino acid content for the four different pollen
230 types (Table 4, Online Resource 3). However, these results could underestimate the total FAA content of naturally
231 occurring pollen since some free amino acids could have been lost during the pollen washing process. Pollen types
232 differed in their absolute amounts of total amino acids (i.e., $\mu\text{g}/\text{mg}$ of dry sample; perMANOVA, $R^2 = 0.93$, $F_{3,8}$
233 = 36.41, $P = 0.0020$; Fig. 3, Online Resource 4). We found that broad bean pollen had both the highest total amino

234 acid and highest essential amino acid content, and sunflower pollen—followed closely by Cucurbitaceae pollen—
235 had the lowest total amino acid and lowest essential amino acid content (Table 4, Online Resource 3). However,
236 despite the overall difference among pollen types, no single pollen type differed significantly from any another in
237 pairwise comparisons once corrected for multiple comparisons ($P > 0.6$), likely due to low sample size. The
238 proportions of the different amino acids were similar across the four pollen types (perMANOVA, $R^2 = 0.36$, $F_{3,8}$
239 = 1.52, $P = 0.21$, Fig. 4), and none of the amino acids was indicative of a particular pollen type (Online Resource 5).

240 **Discussion**

241 ***Microcolony Success in Relation to Pollen Protein Content***

242 Bumblebees died sooner in the 100% sunflower treatment than in any other pollen treatment except 25%
243 sunflower mixed pollen. The difference in survival is correlated with both the lower total amino acid content and
244 essential amino acid contents of the sunflower pollen than the other pollen types. These results are consistent with
245 past studies that have found sunflower (*Helianthus annuus*) pollen to be low in protein (honeybee-collected pollen:
246 Kleinschmidt and Kondos 1976; Pernal and Currie 2000; Somerville and Nicol 2006; honeybee- and hand-collected
247 pollen: Nicolson and Human 2013). In addition, Tasei and Aupinel (2008a) observed the highest worker mortality
248 for *Bombus terrestris* fed sunflower pollen—higher than for bees fed five other pollen diets including *Castanea*
249 (Fagaceae), *Rubus* (Rosaceae), *Papaver* (Papaveraceae), *Actinidia* (Actinidiaceae), and *Cistus* (Cistaceae) pollen.

250 In contrast, broad bean pollen had both the highest total and essential amino acid content, which is also
251 consistent with the results of previous studies that have found legume (Fabaceae) pollen to be particularly protein-
252 rich (Hanley et al. 2008) and an important pollen source for bumblebees (Goulson et al. 2005). Although no larvae
253 were produced during our study, two of the four microcolonies that laid eggs were provided broad bean pollen, and
254 another microcolony that produced eggs was given 0% sunflower mixed pollen, which was one-third broad bean
255 pollen. The microcolonies provided broad bean pollen were also more likely to make honeypots, a normal colony
256 behaviour, than those in the mixed sunflower pollen treatments.

257 It is unclear why so few microcolonies in our experiment produced eggs and why the few microcolonies
258 that did lay eggs later destroyed them. Egg-eating (oophagy) can occur amongst workers that are competing for
259 reproduction (Michener 1969), and it might also occur if workers lay non-viable eggs (cf. Koedam et al. 1996).
260 Oophagy may also be a response to a lack of food or poor resource quality, as it allows bees to recycle nutrients for
261 future egg production (mentioned by Génissel et al. 2002). However, we do not believe the microcolonies in this

262 study were food-stressed, since they all had constant access to sugar solution and pollen. In addition, rapeseed pollen
263 is a suitable diet for queenless microcolony brood production by *B. terrestris* workers (Regali and Rasmont 1995),
264 and Fabaceae pollen (e.g., broad bean) is an important pollen source for many bumblebee species, as noted above.
265 *Bombus terrestris* queenless microcolonies can even rear brood on sunflower pollen, though fewer and smaller
266 males were produced on this diet than on other pollen diets (Regali and Rasmont 1995). This difference in
267 performance between *B. terrestris* and our study species, *B. impatiens*, could reflect species-specific requirements
268 for brood production. Gradish et al. (2013) also observed differences between these species, with few males
269 produced by *B. impatiens* queenless microcolonies under experimental conditions designed for *B. terrestris*.

270 Crude protein, estimated from nitrogen content, has traditionally been used as a measure of pollen
271 nutritional quality (reviewed by Roulston and Cane 2000). However, different pollen diets containing the same
272 overall protein content could have very different nutritional qualities for bees if some are deficient in essential amino
273 acids (Standifer 1967). For example, Tasei and Aupinel (2008a) found that *B. terrestris* queenless microcolonies fed
274 a mixed pollen diet that contained 9% sunflower pollen and had a lower nitrogen content than a pure sunflower diet
275 produced heavier larvae than microcolonies in the pure sunflower treatment. This result suggests that crude protein
276 content is not the only factor affecting pollen nutritive value and that the addition of other pollens may have
277 improved the nutritional properties of the mixed diet (Tasei and Aupinel 2008a). Similarly, in our study, the
278 Cucurbitaceae pollen treatment, which had an amino acid content that was only slightly higher than that of the 100%
279 sunflower treatment, did not reduce bee lifespan relative to the other pollen treatments, suggesting that another
280 nutritional factor besides protein content was responsible for the poor performance of bees in the latter treatment.

281 We found that the relative amounts of amino acids (i.e., percentages of total amino acid content and of total
282 essential amino acid content) were similar across the four pollen types. This finding is consistent with the results of
283 Vanderplanck et al. (2014) and could indicate that the amino acid profile of pollen is generally conserved across
284 plant species (but see Weiner et al. 2010). Nevertheless, the pollen of some Asteraceae taxa has also been found to
285 lack or be deficient in certain essential amino acids (Auclair and Jamieson 1948; Herbert et al. 1970; Wille et al.
286 1985 as cited by Müller and Kuhlmann 2008). In addition, Nicolson and Human (2013) found both hand- and bee-
287 collected sunflower pollen to be deficient in the essential amino acids methionine and tryptophan; however, other
288 studies on bee-collected sunflower pollen have not found essential amino acid deficiencies (Rayner and Langridge
289 1985; Somerville and Nicol 2006—although the latter excluded tryptophan). In our study, we found that all pollen

290 types either met or exceeded the minimum requirements for the 10 essential amino acids outlined for honeybees (De
291 Groot 1953). However, it is possible that the proportional amino acid requirements differ slightly between
292 honeybees and bumblebees. For example, Kriesell et al. (2017) found low proportions of isoleucine in the pollen
293 loads of seven European bumblebee species, which could indicate that bumblebees require proportionally less of this
294 amino acid. The amino acid requirements for bumblebees still need to be determined and could also differ among
295 *Bombus* species (Moerman et al. 2016b).

296 Moreover, pollens could be inferior diets if they lack other key nutrients, but few studies to date have
297 looked at the nutritional requirements of bees for other pollen constituents such as lipids, sterols, vitamins or
298 minerals (Vaudo et al. 2015). Pollen could also contain secondary metabolites, which could have toxic or harmful
299 effects on bees. For example, queenless *B. terrestris* microcolonies produced fewer and smaller males when
300 provided pollen treated with ecologically-relevant concentrations of the secondary metabolite D-lupanine, a
301 quinolizidine alkaloid found in the pollen of *Lupinus* spp. (Fabaceae) (Arnold et al. 2014).

302 Digestibility is another factor that could influence the nutritive quality of a pollen for bees (Human et al.
303 2007). For example, the thickness of the intine layer at the germinal apertures of pollen grains can affect pollen
304 digestion by *Osmia* bees (Suárez-Cervera et al. 1994). In addition, an experiment on honeybees (*A. mellifera*
305 *scutellata*) observed higher extraction efficiency in bees fed aloe pollen (*Aloe greatheadii* var. *davyana*,
306 Asphodelaceae) than those fed sunflower pollen, potentially due to differences in pollen morphology (Human et al.
307 2007). Aloe pollen grains are large and smooth, whereas sunflower pollen grains are smaller and covered in spines
308 (echinate), which could impede digestion (Human et al. 2007). In addition, the prominent oily coating (pollenkitt) of
309 sunflower pollen must first be digested before the cytoplasm can be extracted; this might further reduce nutrient
310 extraction efficiency (Human et al. 2007). Consequently, it is possible that we observed a shorter bee lifespan in the
311 sunflower pollen treatment due to nutritive factors other than the low amino acid content of this pollen treatment.
312 These factors could include the absence or deficiency of other essential nutrients, the presence of harmful secondary
313 metabolites, or pollen grain structure. These possibilities remain to be tested.

314 ***Advantages of a Mixed Pollen Diet***

315 In general, bumblebees are broadly polylectic (dietary generalist) species that can collect pollen from multiple
316 plant species while foraging (Leonhardt and Blüthgen 2012; Somme et al. 2015). Mixing pollen from different plant
317 species might allow bumblebees to exploit low-quality resources if the mixed diet compensates for nutrient
318 deficiencies and/or reduces the toxicity of the monofloral diet (Eckhardt et al. 2014). We observed that bees

319 provided sunflower pollen did not live as long as bees in the other pollen treatments. However, a mixed pollen diet
320 containing 50% sunflower pollen yielded worker survival as high as that of the non-sunflower diets, potentially due
321 to the presence of the other pollens in the diet which may have mitigated any harmful effects of the sunflower
322 pollen. For example, the 50% sunflower pollen treatment had a higher amino acid content than both the sunflower
323 and Cucurbitaceae pollen treatments (296 µg/mg compared to 250 µg/mg and 256 µg/mg, respectively), supporting
324 the idea that one benefit of the mixture may have been its increased nitrogen content. The other pollen mixtures also
325 had reasonably high amino acid content (342 µg/mg and 319 µg/mg for the 0% and 25% sunflower treatments,
326 respectively).

327 Other studies have observed better bee performance on a mixed pollen diet than a monofloral diet of
328 comparable protein content. For example, Baloglu and Gurel (2015) found that a mixed pollen diet containing equal
329 amounts of *Cistus* spp., *Papaver somniferum*, and *Sinapis* sp. was a superior pollen diet for queenright *B. terrestris*
330 colonies than *P. somniferum* pollen alone, even though the monofloral diet had a higher protein content than the
331 polyfloral diet (21.4% and 18.5%, respectively). In addition, Tasei and Aupinel (2008a) reported that protein
332 efficacy (weight of larvae/protein consumed) was ~4 times greater for queenless microcolonies of *B. terrestris* fed
333 mixed pollen diets than pure pollen diets. However, it is important to consider that monofloral diets can sometimes
334 be as good as, if not better than, mixed diets for colony performance. For example, Moerman et al. (2017) observed
335 that queenless microcolonies fed monofloral diets of *Cytisus scoparius* (Fabaceae) and *Sorbus aucuparia* (Rosaceae)
336 pollen produced larger larvae than microcolonies given difloral diets containing *Erica* sp. (Ericaceae) pollen,
337 perhaps because the latter contained the harmful sterol $\delta 7$ -avenasterol, which the former pollens did not (Moerman
338 et al. 2017). Similarly, the 25% sunflower pollen treatment in our study significantly shortened worker lifespan
339 compared to a monofloral diet of broad bean pollen. Finally, bumblebees do not mix pollen randomly but may prefer
340 pollens with a more favourable protein to lipid ratio (Vaudo et al. 2016a, b). In the future, it would be useful to
341 conduct a foraging study to determine whether bumblebees regulate their intake of sunflower pollen when given a
342 choice among pollen types, and if so, what proportion of sunflower pollen they choose to collect.

343 ***Conclusion and Implications***

344 Although we did observe improved survival of bees offered sunflower mix pollen diets (relative to the 100%
345 sunflower diet), additional trials with a greater variety of sunflower mixed diets (potentially with the addition of
346 other pollen species) will need to be conducted to determine what proportion of sunflower pollen is acceptable—
347 particularly considering the observed discrepancies in performance between the 25% and 50% sunflower mixed

348 pollen treatments, as well as the relatively low performance on the 0% mixed treatment. At present, we are unable to
349 explain these discrepancies in performance, but future studies using different pollen mixtures may shed light on our
350 results.

351 Host selection by bees is a multifaceted process that is influenced not only by reward quality but also by the
352 spatiotemporal availability of floral resources and the accessibility of floral rewards. As a consequence of
353 agricultural intensification and a loss of wildflowers, farmland monocultures can be important floral resources for
354 bumblebees (Westphal et al. 2003). However, a monotonous diet could have adverse effects on bumblebee colony
355 health, particularly if the floral rewards of the crop are of low nutritional value. Schmidt et al. (1995) noted that
356 commercial honeybee colonies established on or near sunflower crops, with little access to other floral resources
357 during sunflower bloom, might experience both colony stress and a shorter worker lifespan, which could in turn
358 reduce pollination services. Our results suggest that providing alternative floral resources of high nutritive quality
359 could help mitigate the potential harmful effects of a monofloral diet of sunflower pollen on bumblebees. These
360 alternative resources could be provided as hedgerows (Morandin and Kremen 2013) or wildflower plantings along
361 field margins (Williams et al. 2015) that ideally sustain flowering throughout the growing season. To support bee
362 populations, we need to improve our understanding of both the nutritional requirements of different bee species
363 (Vaudo et al. 2015) and how the quality of the available floral resources affects colony health (Leonhardt and
364 Blüthgen 2012). This knowledge could then be integrated into management strategies to support bee populations by
365 providing foraging habitats with high-quality resources.

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533 **Table 1** Pollen treatment allocation for the 43 microcolonies (MC) of *B. impatiens* used in the experiment. Each
 534 microcolony was exposed to one of seven pollen diets, which included 0%, 25%, and 50% sunflower mixed pollen
 535 (SFM) (97% *Helianthus annuus* and 3% other Asteraceae species [*Rudbeckia* sp., *Coreopsis* sp. and *Heliopsis* sp.])
 536 with an equal mixture of broad bean (*Vicia faba*), Cucurbitaceae (30% summer squash [*Cucurbita pepo*], 70%
 537 watermelon [*Citrullus lanatus*]), and rapeseed (*Brassica napus*) making up the rest of the diet. The remaining four
 538 diets contained only one kind of pollen (broad bean, Cucurbitaceae, rapeseed, or sunflower). There was no pollen
 539 consumption in ten of the microcolonies (~23%) included in the experiment. The numbers of bees in microcolonies
 540 that did and did not consume pollen are in brackets. Bees that died before pollen consumption are included with the
 541 microcolonies with no pollen consumption
 542

	Pollen Treatments							Total MC (bees)
	0% SFM	25% SFM	50% SFM	Broad Bean	Cucurbitaceae	Rapeseed	Sunflower	
MC (bees) with pollen consumption	4 (21)	6 (28)	4 (20)	5 (28)	5 (25)	5 (27)	4 (20)	33 (169)
MC (bees) with no pollen consumption	1 (8)	0 (4)	2 (11)	1 (6)	2 (14)	2 (14)	2 (11)	10 (68)
Total MC	5 (29)	6 (32)	6 (31)	6 (34)	7 (39)	7 (41)	6 (31)	43 (237)

543 **Table 2** Cox PH ratios for bee survival across the seven pollen treatments which included 0%, 25%, and 50%
544 sunflower mixed pollen (SFM) (97% *Helianthus annuus* and 3% other Asteraceae species [*Rudbeckia* sp., *Coreopsis*
545 sp. and *Heliopsis* sp.]) with an equal mixture of broad bean (*Vicia faba*), Cucurbitaceae (30% summer squash
546 [*Cucurbita pepo*], 70% watermelon [*Citrullus lanatus*]), and rapeseed (*Brassica napus*) making up the rest of the
547 diet. The remaining four diets contained only one kind of pollen (broad bean, Cucurbitaceae, rapeseed, or
548 sunflower). Sample size (number of bees) is included in brackets. Bees that did and did not consume pollen were
549 included in this analysis. Results of the analysis that omitted outliers and bees that did not consume pollen (model 2)
550 are provided in Online Resource 2. Hazard ratios less than one indicate higher chances of survival, and values
551 greater than one indicate reduced chances of survival, relative to the reference pollen treatment. Sunflower pollen
552 significantly reduced survival compared to all other pollen treatments except 25% SFM pollen

	Reference pollen treatment						
	0% SFM	25% SFM	50% SFM	Broad bean	Cucurbitaceae	Rapeseed	Sunflower
0% SFM (29)		0.66	1.11	1.63	1.03	1.13	0.35*
25% SFM (32)	1.51 ^a		1.67	2.47*	1.56	1.72	0.53
50% SFM (31)	0.90	0.60		1.47	0.93	1.02	0.32**
Broad bean (34)	0.61	0.40*	0.68		0.63	0.69	0.22**
Cucurbitaceae (39)	0.97	0.64	1.07	1.58		1.10	0.34**
Rapeseed (41)	0.88	0.58	0.98	1.44	0.91		0.31**
Sunflower (31)	2.85*	1.88	3.15**	4.64**	2.49**	3.23**	

553 * $P < 0.05$

554 ** $P < 0.01$

555 ^aHazard ratios above the diagonal are reciprocals of the corresponding values below the diagonal.

556 **Table 3** Cox PH ratios for microcolony honeypot construction across the seven pollen treatments which included
557 0%, 25%, and 50% sunflower mixed pollen (SFM) (97% *Helianthus annuus* and 3% other Asteraceae species
558 [*Rudbeckia* sp., *Coreopsis* sp. and *Heliopsis* sp.]) with an equal mixture of broad bean (*Vicia faba*), Cucurbitaceae
559 (30% summer squash [*Cucurbita pepo*], 70% watermelon [*Citrullus lanatus*]), and rapeseed (*Brassica napus*)
560 making up the rest of the diet. The remaining four diets contained only one kind of pollen (broad bean,
561 Cucurbitaceae, rapeseed, or sunflower). Sample size (number of microcolonies) is provided in brackets and all
562 microcolonies were included in this analysis. Hazard ratios less than one indicate reduced chances of honeypot
563 construction, and values greater than one indicate greater chances of honeypot construction, relative to the reference
564 pollen treatment. Microcolonies provided broad bean pollen were significantly more likely to construct honeypots
565 than the 25% SFM pollen treatments, and marginally significantly more likely to make honeypots than the 50%
566 SFM pollen treatment and 100% sunflower pollen treatment

	Reference pollen treatment						
	0% SFM	25% SFM	50% SFM	Broad bean	Cucurbitaceae	Rapeseed	Sunflower
0% SFM (5)		5.30	2.66	0.54	1.72	1.45	2.22
25% SFM (6)	0.19 ^a		0.50	0.10*	0.32	0.27	0.42
50% SFM (6)	0.38	1.99		0.20[§]	0.65	0.55	0.83
Broad bean (6)	1.87	9.89*	4.97[§]		3.21	2.71	4.14[§]
Cucurbitaceae (7)	0.58	3.08	1.55	0.31		0.85	1.29
Rapeseed (7)	0.69	3.64	1.83	0.37	1.18		1.52
Sunflower (6)	0.45	2.39	1.20	0.24[§]	0.78	0.66	

567 [§]0.05 < P < 0.075

568 *P < 0.05

569 **P < 0.01

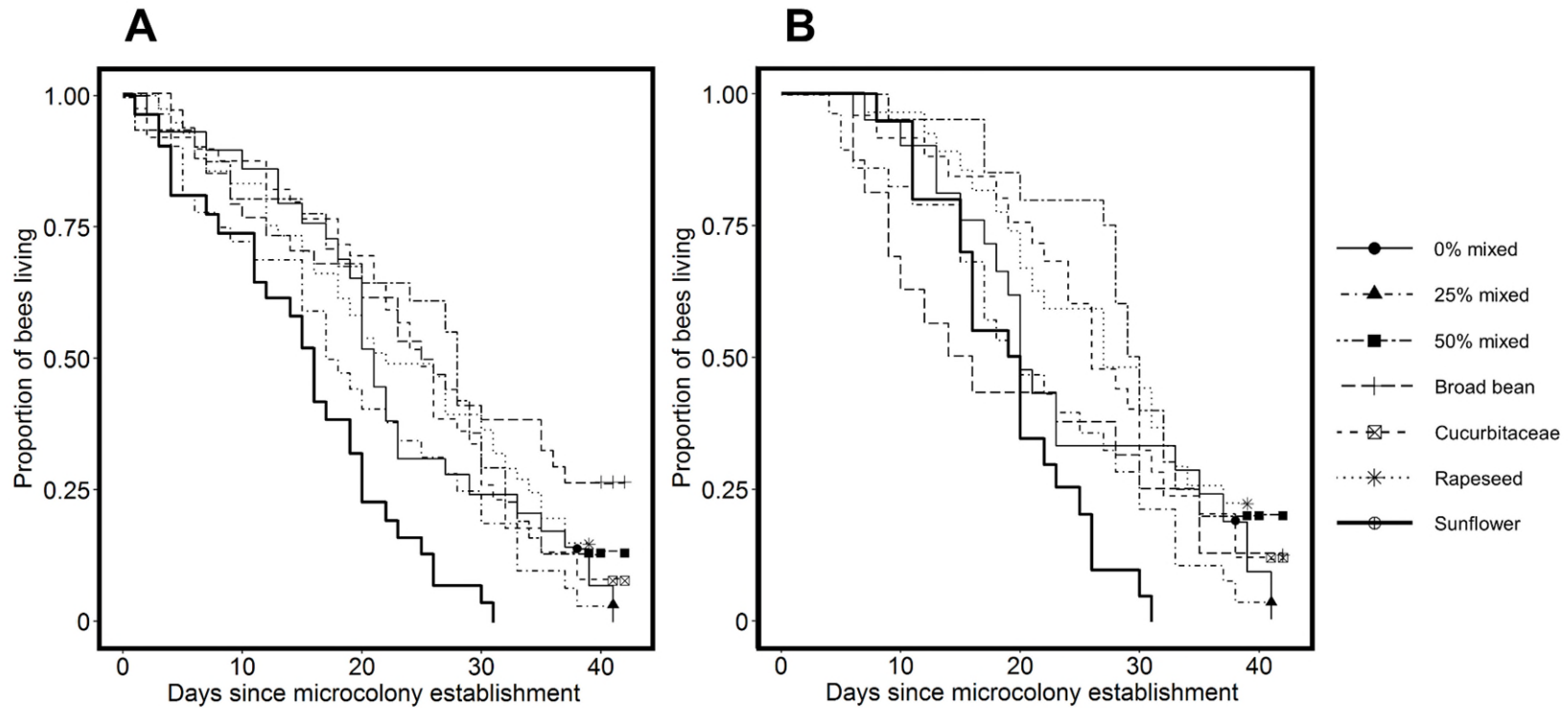
570 ^aHazard ratios above the diagonal are reciprocals of the corresponding values below the diagonal.

571 **Table 4** Mean \pm SEM for total amino acid (AA), essential amino acid (EAA) and percent free amino acid content
 572 (% FAA) for the four pollen types (broad bean (*Vicia faba*), Cucurbitaceae (30% summer squash [*Cucurbita pepo*],
 573 70% watermelon [*Citrullus lanatus*]), rapeseed (*Brassica napus*), and sunflower (97% *Helianthus annuus* and 3%
 574 other Asteraceae species [*Rudbeckia* sp., *Coreopsis* sp. and *Heliopsis* sp.]

Pollen Treatments	AA ($\mu\text{g}/\text{mg}$ of sample) ^a	EAA ($\mu\text{g}/\text{mg}$ of sample)	% FAA ^b
Broad Bean	420.89 \pm 16.28	216.29 \pm 4.20	6.06 \pm 0.41
Cucurbitaceae	255.99 \pm 12.51	129.47 \pm 2.25	6.43 \pm 0.27
Rapeseed	350.45 \pm 3.27	180.72 \pm 1.91	4.50 \pm 0.24
Sunflower	249.61 \pm 3.56	127.56 \pm 1.33	3.49 \pm 0.21

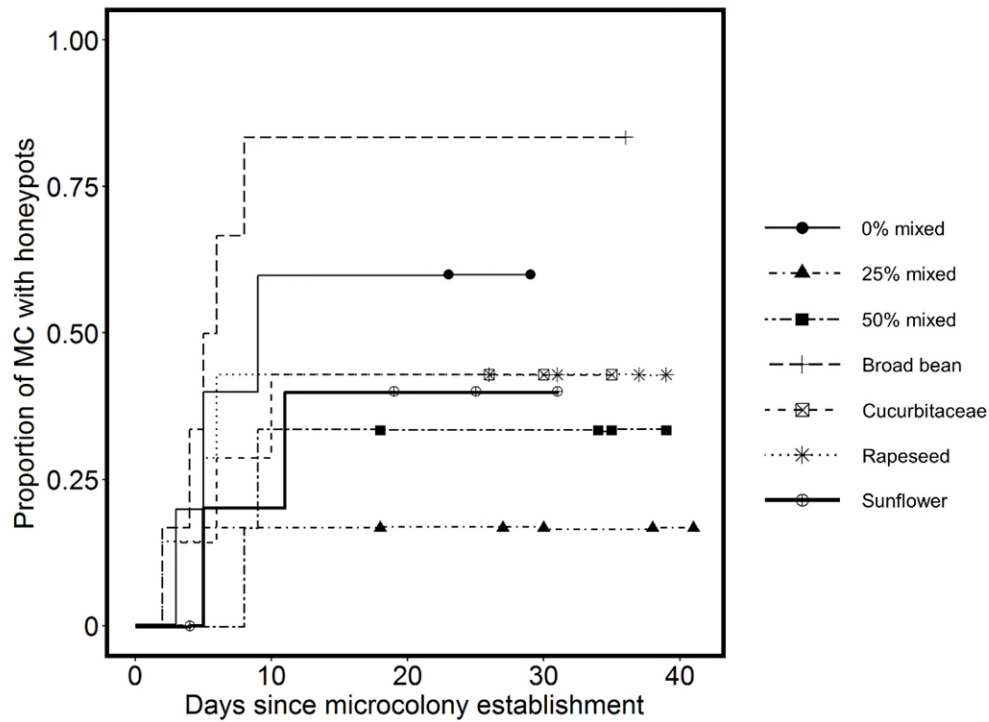
575 ^aSample weights are the dry weights of the pollen.

576 ^bThe proportion of free tryptophan could not be determined since this amino acid was analyzed in a separate analysis
 577 which did not distinguish between protein-bound and free amino acid. As a result, this amino acid was omitted for
 578 the % FAA calculation.



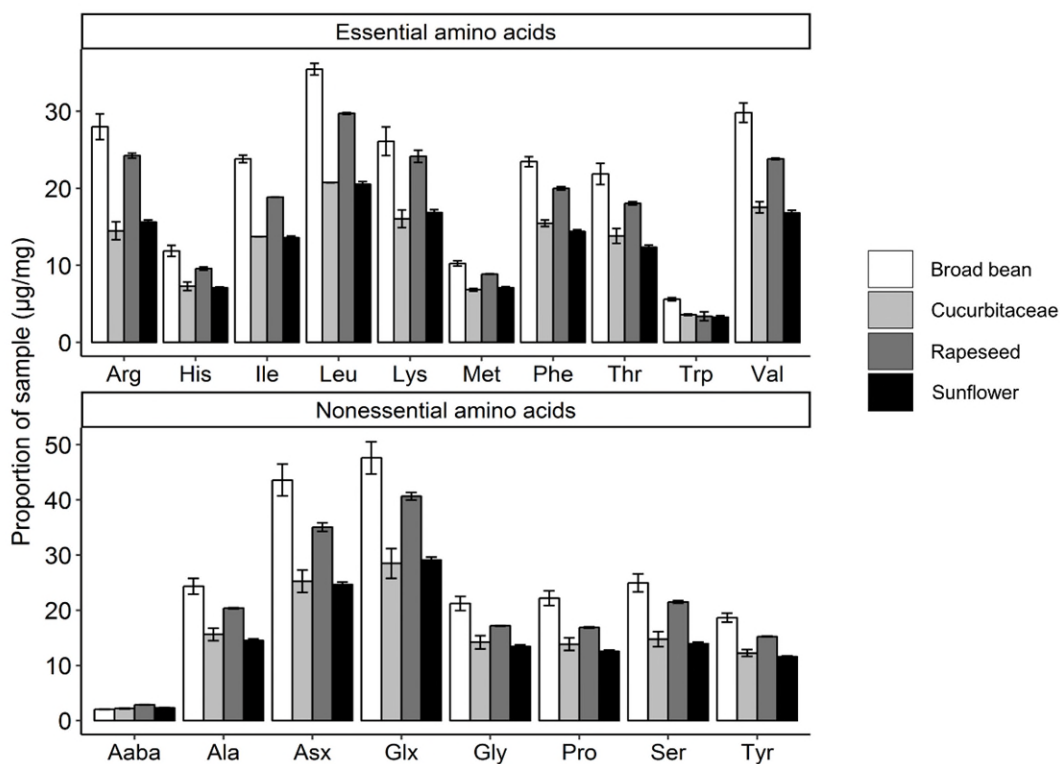
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581 **Fig. 1** Bee survival over time on the seven pollen treatments visualized using the ‘survfit’ function of the ‘survival’ package. **A** contains all bees (as in Table 2;
 582 model 1) and **B** contains only the bees from microcolonies with pollen consumption and omits two outlier microcolonies from the broad bean treatment (12 bees)
 583 (as in Online Resource 2, model 2). The seven pollen diets included 0% sunflower mixed (SFM) pollen diet (A: n = 29 bees; B: n = 21 bees), 25% SFM (33; 28),
 584 and 50% SFM (31; 20), broad bean (34; 16), Cucurbitaceae (39; 25), rapeseed (41; 27), and sunflower (31; 20). Survival was significantly reduced on sunflower
 585 pollen relative to all pollen treatments except 25% SFM pollen. Results were jittered both vertically and horizontally to reduce overlap between the different
 586 pollen treatments. Bees that were censored until the end of the 6-week period (42 days) are represented by the symbols on the curves



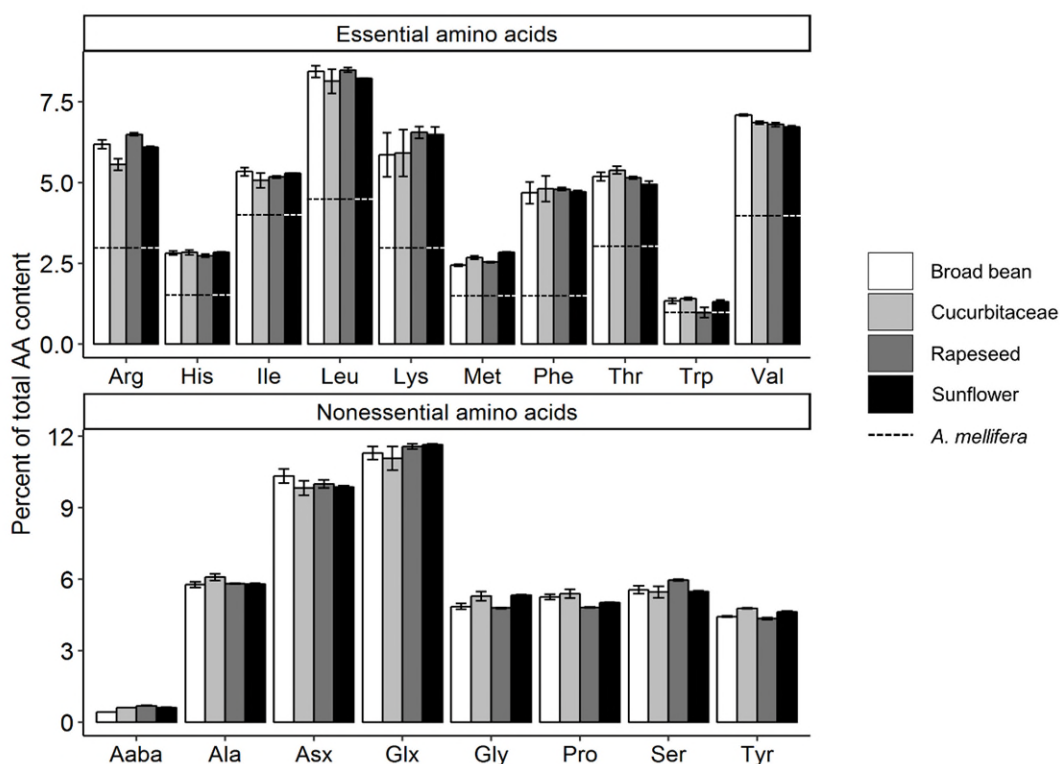
587

588 **Fig. 2** Honeypot construction over time on the seven pollen treatments visualized using the ‘survfit’ function of the
 589 ‘survival’ package. All microcolonies (“MC”) were included in this analysis. The seven pollen diets included 0%
 590 sunflower mixed (SFM) pollen diet (n = 5 microcolonies), 25% SFM (6), and 50% SFM (6), broad bean (6),
 591 Cucurbitaceae (7), rapeseed (7), and sunflower pollen (6). Microcolonies provided broad bean pollen were
 592 significantly more likely to construct honeypots than the 25% SFM pollen treatments, and marginally significantly
 593 more likely to make honeypots than the 50% SFM pollen treatment and 100% sunflower pollen treatment. Results
 594 were jittered both vertically and horizontally to reduce overlap between the different pollen treatments.
 595 Microcolonies without honeypot construction are represented by the symbols on the curves



596

597 **Fig. 3** The mean \pm SEM of essential amino acids (EAA) and nonessential amino acids (NAA) (including both
 598 protein-bound and free amino acids) in the four pollen types (broad bean (*Vicia faba*), Cucurbitaceae (30% summer
 599 squash [*Cucurbita pepo*], 70% watermelon [*Citrullus lanatus*]), rapeseed (*Brassica napus*), and sunflower (97%
 600 *Helianthus annuus* and 3% other Asteraceae species [*Rudbeckia* sp., *Coreopsis* sp. and *Heliopsis* sp.]); n=3) as a
 601 proportion of the dry mass of the sample ($\mu\text{g}/\text{mg}$ of sample). (EAA: Arg = arginine, His = histidine, Ile = isoleucine,
 602 Leu = leucine, Lys = lysine, Met = methionine, Phe = phenylalanine, Thr = threonine, Trp = tryptophan, Val =
 603 valine; NAA: Aaba = alpha-aminobutyric acid, Ala = alanine, Asx = asparagine + aspartic acid, Glx = glutamine +
 604 glutamic acid, Gly= glycine, Pro = proline, Ser = serine and Tyr = tyrosine)



605

606 **Fig. 4** The mean \pm SEM of essential amino acids (EAA) and nonessential amino acids (NAA) (including both
 607 protein-bound and free amino acids) in the four pollen types (broad bean (*Vicia faba*), Cucurbitaceae (30% summer
 608 squash [*Cucurbita pepo*], 70% watermelon [*Citrullus lanatus*]), rapeseed (*Brassica napus*), and sunflower (97%
 609 *Helianthus annuus* and 3% other Asteraceae species [*Rudbeckia* sp., *Coreopsis* sp. and *Heliopsis* sp.]); n=3) as a
 610 proportion of the total amino acid content. The dashed lines represent the EAA requirements for *Apis mellifera*
 611 described by De Groot (1953). (EAA: Arg = arginine, His = histidine, Ile = isoleucine, Leu = leucine, Lys = lysine,
 612 Met = methionine, Phe = phenylalanine, Thr = threonine, Trp = tryptophan, Val = valine; NAA: Aaba = alpha-
 613 aminobutyric acid, Ala = alanine, Asx = asparagine + aspartic acid, Glx = glutamine + glutamic acid, Gly= glycine,
 614 Pro = proline, Ser = serine and Tyr = tyrosine)