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1 **Title: Gender-biased nectar targets different behavioural traits of flower visitors**

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18

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20 **Author contributions**

21 MB and GB share the first authorship. MG, MN, LC and LB conceived and designed the
22 experiments. GB, MA and MG performed the experiments. HPLC analyses were executed by
23 MN. MB and GB analysed data and wrote the paper. All authors read, provided editorial
24 advice and approved the final manuscript.

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26

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35

36 **Abstract**

37 Floral nectar is a chemically complex aqueous solution within which several secondary
38 metabolites have been identified and that affect attractiveness for pollinators. Understanding
39 preferences and aversions to nectar quality in flower visitors is crucial since this may
40 influence the patterns of insect floral visitation with consequences on the plant fitness. We
41 hypothesise that nectar chemical variation through different floral sexual phases may affect
42 the number of insect visits that each phase receives. The study was realized on a population of
43 *Echium vulgare* L. growing in a natural area close to Bologna. Nectar was collected from
44 functionally male and female flowers to investigate its chemical composition through the
45 HPLC technique. A total of 200 mins of behavioural observations on foraging insects were
46 also carried out. Variation in nectar traits has been detected for the amino acid spectrum. The
47 proportion of protein amino acids appeared to be significantly higher in male-phase flowers.
48 This may explain the significantly higher number of visits on male flowers than expected
49 observed for all bee taxa (except *Hoplitis adunca* females). Functionally male flowers
50 presented higher concentrations of phenylalanine, whilst proline was highly represented in

51 functionally female flowers. Since a recent study demonstrated that hymenopterans can
52 oxidize proline at a high rate for ATP production, we can hypothesise that the quality of
53 nectar offered by the two sexually distinct floral phases targets different insect behavioural
54 traits and likely ensures an optimal pattern of visit among flower sexes, which are unequally
55 distributed within and among individuals in the population.

56

57 **Keywords:** *Echium vulgare*, flower visitors, inbreeding avoidance, nectar chemistry, plant-
58 pollinator interactions

59

60 **Introduction**

61 Floral nectar is a chemically complex aqueous solution in which the main components
62 comprise sugars, followed by amino acids (Nicolson and Thornburg 2007). In recent decades
63 considerable progress has been made in providing evidence that points to the involvement of
64 nectar chemistry in the interactions between plants and a variety of organisms (Nepi 2014;
65 Stevenson et al. 2017). Although there is wide variability in nectar traits (Pacini et al. 2003;
66 Nocentini et al. 2013; Irwin et al. 2014), a general paradigm shared by plants is balancing
67 nectar chemical composition in order to not deter specific pollinators exceeding their
68 tolerance thresholds (Baker and Baker 1975; Adler 2000; Nicolson and Thornburg 2007;
69 Wright et al. 2013; Stevenson et al. 2017). For example, a small increase in nectar sugar
70 concentration can increase its viscosity (Harder 1986; Nicolson and Thornburg 2007), which
71 is strongly related to the energy required by nectar consumers to visit flowers (Corbet 1978;
72 Josens and Farina 2001; Borrell and Krenn 2006; Nepi and Stpicyńska 2006; Kim et al.
73 2011).

74 After sugars the most abundant nectar solutes are the amino acids (Baker and Baker 1982;
75 Nepi et al. 2012; Bogo et al. 2019). A study conducted by Inouye and Waller (1984) showed a
76 general decline in nectar consumption in honeybees as amino acid concentrations increased,
77 despite evidence supporting the preference for amino acid enriched sugar solutions in insects
78 (Alm et al. 1990; Bertazzini et al. 2010; Bogo et al. 2019). Amino acids also contribute to the
79 taste of nectar, stimulating specific insects' labellar chemoreceptors (Gardener and Gillman
80 2002). Among protein amino acids, Inouye and Waller (1984) found that phenylalanine and
81 leucine were phagostimulant for honeybees at all concentrations tested, even at those that in
82 the case of other amino acids resulted in deterrence. In the same way, a preference in
83 honeybees for proline enriched artificial nectar was reported (Carter et al. 2006; Bertazzini et

84 al. 2010), as well as a strong phagostimulatory activity (Nicolson and Thornburg 2007;
85 Petanidou 2007).

86 Beside primary metabolites (such as sugars and amino acids) an array of secondary
87 metabolites with different chemical natures have been identified in nectar and all of them
88 positively or negatively affect attractiveness to pollinators, showing effects which depend on
89 metabolite concentration and pollinators' sensitivity (Baker and Baker 1977; Faegri and van
90 der Pijl 1979; Baker and Baker 1982; Adler 2000; Stevenson et al. 2017). Among them non-
91 protein amino acids (NPAAs) have been detected in nectar (Nicolson and Thornburg 2007;
92 Petanidou 2007; Nepi et al. 2012). Despite that they can constitute a large portion of the
93 amino acidic content of floral nectar, little is known about their role in determining
94 pollinators' preferences and feeding behaviour. For some of those, such as γ -aminobutyric
95 acid, a phagostimulant function has been reported in some caterpillars and adult beetles
96 (Mitchell and Harrison 1984; Shoonhoven et al. 2005), whilst Bogo et al. (2019) found that
97 both bumblebees and honeybees showed higher consumption of sucrose solution enriched
98 with β -alanine, but exhibited the effect at different concentrations.

99 Understanding preferences and aversions to nectar traits is crucial since they likely influence
100 the patterns of floral visitation by nectar consumers and thus the plant inbreeding and
101 outbreeding rate within a population. Minimal inbreeding is predicted when pollinators visit a
102 small fraction of the open flowers on a plant (Iwasa et al. 1995; Ohashi and Yahara 2001):
103 this behaviour may be enhanced by within-plant variation in nectar, as occurs in plants
104 showing gender-biased nectar production (Feinsinger 1978; Pike 1978; Rathcke 1992).

105 Despite many studies having already addressed the subject of gender-biased nectar
106 composition, most of them investigated the existence of bias in relation to nectar volume or
107 sugar content only (Langenberger and Davis 2002; Canto et al. 2011; Fisogni et al. 2011;
108 Stpiczyńska et al. 2015; Antoń et al. 2017; Jacquemart et al. 2019; Konarska and

109 Masierowska 2020) and few reported the observation of insect visit bias (Carlson and Harms
110 2006 and references therein).

111 In this study we focused on the many-flowered hermaphrodite species *Echium vulgare* L., a
112 self-compatible plant which shows both herkogamy and incomplete protandry, that avoids
113 self-pollination within the same flower, but within which geitonogamy can still occur
114 (Rademaker et al. 1999). Melsner et al. (1999) reported evidences of inbreeding depression in
115 *E. vulgare*, finding a significant decline in siring success when selfing occurs. A study on
116 geitonogamy conducted by Rademaker et al. (1999), though, found a consistently lower
117 percentage of selfing rate than expected. Also, they reported that bumblebees visited only a
118 small fraction of the flowers on *E. vulgare* as a result of the presence of different flower
119 stages simultaneously occurring on a single individual plant.

120 *E. vulgare* represents an important food resource for many insect visitors, despite containing
121 toxic pyrrolizidine alkaloids in both nectar and pollen (Lucchetti 2017). The pollen contains
122 high concentrations of pyrrolizidines, whilst more than 500 times lower concentrations are
123 found in nectar (Lucchetti et al. 2016). For this reason, only a few taxa show oligolecty or
124 floral constancy on *E. vulgare* by actively collecting pollen for larval nourishment (Cane and
125 Sipes 2006; Burger et al. 2010; Filella et al. 2011), even if its flowers are visited by a wide
126 spectrum of insect taxa among which bumblebees have often been reported as main
127 pollinators (Corbet 1978; Klinkhamer and de Jong 1990; Pappers et al. 1999; Rademaker et
128 al. 1999).

129 Here, we examined if floral visitation pattern may be influenced by variations in the chemical
130 composition of nectar through different floral stages, and thus we investigated (i) whether *E.*
131 *vulgare* produces a gender-biased nectar for volume, sugar and amino acid composition and
132 (ii) if flower visitation rates of insects looking for nectar varied among different floral stages.

133

134 **Material and Methods**

135 **Study site**

136 The activity in the field was carried out in June 2018 and took place in the Parco Belpoggio, a
137 public park managed since 2010 by the WWF, in San Lazzaro di Savena (Bologna, Italy). The
138 area is situated close to the protected area Parco dei Gessi Bolognesi e Calanchi
139 dell'Abbadessa (44°27'14.5"N 11°22'58.3"E). The studied population was located on an open
140 prairie along the public pathway.

141

142 **Study species**

143 *Echium vulgare* L. is a perennial hemicryptophyte belonging to the family Boraginaceae. It is
144 distributed in Europe, Asia and North America and it shows a long flowering period, ranging
145 between June and October. Flower anthesis lasts 3-4 days and flowers show an incomplete
146 protandry (Melser et al. 1997): the anthers are often dehiscent already at the bud stage, while
147 the stigma becomes receptive only hours after the flower opening.

148 In this study we considered three phases of floral development: closed flower (Bud),
149 functionally male (M) and functionally female (F) flowers. The male phase was represented
150 by an open flower presenting pollen with non receptive stigma, whilst the female phase was
151 recognised as soon as the stigma became bifid and receptive.

152

153 *Plant phenology*

154 On the first day of the study we counted all plants and inflorescences per plant constituting
155 the population (approximately 600 m² of extension) and we observed all open flowers to
156 assess whether the phenomenon of gynodioecy, firstly described in *E. vulgare* populations by
157 Darwin (1877), occurred in our study population. Each day, prior to visitor observations, on
158 the same patch we recorded the number of flowers per developmental stage. Two fixed

159 patches were alternatively considered: the first one was a single plant carrying 6
160 inflorescences while the second one was made up of 6 plants carrying one or two
161 inflorescences each.

162

163 **Nectar quality**

164 *Sampling*

165 We collected nectar samples by means of Drummond Microcaps (3-5 μL ; Drummond
166 Scientific Co., Broomall, PA), we transferred samples to Eppendorf tubes filled with 100 μL
167 of pure ethanol, and then we took them to the laboratory in thermal bags where they were kept
168 at 5°C until analyses. We collected each sample from multiple flowers at the same floral stage
169 in order to reach a minimum volume of 2 μL needed for the sugar and amino acid analyses. In
170 order to let the nectar accumulate, flowers were bagged in the morning for 2 hours prior to
171 sampling; all nectar present in the selected flowers was collected.

172 We collected a total of 8 nectar samples each one from 3-13 male flowers belonging to 1-7
173 plants, and a total of 8 samples from 2-9 female flowers belonging to 1-3 plants. Both sugar
174 and amino acid compositions were investigated on these samples. We then collected 14
175 additional samples from 1-22 buds belonging to 1-10 plants. Since the amount of nectar
176 presents in the buds was very low, the minimum volume of 2 μL needed for amino acid
177 analysis could not be reached and thus these samples were tested for sugar composition only.

178

179 *Sugar analysis*

180 Sugar content was analysed by HPLC technique through a Waters LC1 with refractive index
181 detector (Waters 2410) connected to the output of a REZEX RCM Monosaccharide column
182 (Phenomenex, 300 mmx7.8 mm, grain 8 μm) maintained at 85°C. Water (MilliQ, pH 7) was

183 used as mobile phase at a flow rate of 0.6 mL min⁻¹; 20 µL of sample and standard solutions
184 of sucrose, glucose and fructose were also injected (Nocentini et al. 2012).

185

186 *Amino acids analysis*

187 Amino acid analysis was performed by gradient HPLC with an ion exchange Novapack C18
188 (15 mm x 4.6 mm) cartridge with guard column maintained at 37°C and a Waters 470
189 scanning fluorescence detector (excitation at 295 nm, detection at 350 nm). A solvent
190 composed of TEA-phosphate buffer (pH 5.0) mixed with a 6:4 acetonitrile-water solution was
191 used as mobile phase at a flow rate of 1.0 mL min⁻¹. According to AccQtag protocol (Waters
192 Corp.), the selected volume of each reconstituted sample was amino acid derivatized (Cohen
193 and Micheaud 1993) with AQC fluorescent reagent and 0.02 M borate buffer (pH 8.6). In
194 addition to all the protein amino acids, standard solutions of β-alanine, citrulline, L-
195 homoserine, α-aminobutyric acid (AABA), γ-aminobutyric acid (GABA), hydroxyproline,
196 ornithine and taurine were also used (Nocentini et al. 2012).

197

198 **Flower visitors' observations**

199 We carried out observations on flower visitors on the two fixed patches described previously,
200 on 7 non-sequential days. Every survey consisted of two 15-mins periods separated by 10
201 mins of rest, adapting the protocol of Fisogni et al. (2016). Every day we performed 1 to 3
202 surveys, between 10:30 am and 3:00 pm and under favourable weather conditions, for a total
203 of 200 mins of observation. Once a visitor left the patch, we counted the following
204 approaching insect belonging to the same taxon as a different individual. Recorded data
205 concerned the food resource collected (nectar or pollen, observing if the insect inserted its
206 mouth-parts deeply inside the corolla or if it manipulated the anthers) and the number of male

207 and female flowers approached per visit. We also recorded the visitor's taxon, indicating the
208 taxonomic level in as much detailed as possible, and its sex.

209 After each observation period, we performed a 15-mins period of net sampling throughout the
210 area, collecting insects that alighted on flowers of *E. vulgare*. Captured individuals were put
211 in separate vials with ethyl acetate and brought to the laboratory where they were pinned in
212 entomological boxes and inspected under a dissecting microscope for taxonomic
213 identification.

214

215 **Data analysis**

216 Sugar and amino acid quantities and the mean nectar volume were calculated per single
217 flower. Total sugar concentration was calculated as the sum of sucrose, fructose and glucose
218 concentrations.

219 Data on nectar composition were grouped by floral stage and tested to assess homogeneity of
220 variances and normality of distribution (Bartlett test and Shapiro Wilk test).

221 Data on sugars per flower, total sugar concentration and sucrose per flower were square root
222 transformed to achieve normality. When the transformed data failed to match normality, we
223 applied the corresponding non-parametric analyses.

224 To investigate whether the floral stage affected sugar content and volume a one-way ANOVA
225 followed by Tukey's HSD post hoc test with Benjamini-Hochberg correction for 'false
226 discovery rate' (Verhoeven et al. 2005) were performed. When distribution was not normal a
227 Kruskal Wallis H-test followed by a Mann Whitney pairwise comparison with Benjamini-
228 Hochberg correction were carried out instead.

229 Data on single amino acid concentrations were ln transformed to achieve normality when
230 needed and a Student t-test was applied in all analyses.

231 For both phenological stages (functionally male and functionally female flowers), three
232 diversity indices were calculated on the nectar amino acid composition. The first index was
233 the reciprocal Simpson's diversity index $1-D$ of the nectar amino acidic spectrum. D was
234 calculated as $D = \sum_{i=1}^n \left(\frac{n_i}{n}\right)^2$, where n_i is the abundance of the i th amino acid and n is the
235 total mean concentration (Ranjbar et al. 2017). This index ranges from 0 (one amino acid
236 dominates the spectrum) to 1 (all amino acids equally represented) (Harper 1999).
237 The second was the Shannon's H - index, by taking into account mean amino acid
238 concentrations as well as the total mean concentration of amino acids. The index is calculated
239 as $H = -\sum_i \frac{n_i}{N} \ln \frac{n_i}{N}$, where n_i is the mean concentration for the i th amino acid and N is the
240 total number of amino acids (Magurran 2004). This index varies from 0 for a spectrum with
241 only a single amino acid to high values for a spectrum with many amino acids, each
242 represented by relatively low concentrations (Harper 1999; Hubálek 2000; Fattorini et al.
243 2016).
244 The third one was the Buzas and Gibson's evenness index, a measure of the relative
245 abundance of the different amino acids within the floral stage. The index is calculated as the
246 proportion of equally dominant amino acid in the phenological stage $E = e^H/S$, where H is
247 Shannon's H index and S is the number of amino acids within the floral stage. This index
248 ranges from 0 (highest dominance by a single amino acidic species) to 1 (all amino acids have
249 the same abundance) (Buzas and Hayek 2010; Fattorini et al. 2016).
250 Insect visit data were first analysed by comparing the observed number of male and female
251 flowers visited to the expected ones by χ^2 test. The expected number of visits was calculated
252 on the basis of the ratio between the functionally male and the functionally female flowers
253 occurring in the population.
254 Frequencies of male flowers visited by each taxon were compared by a Kruskal Wallis H-test
255 followed by a Mann-Whitney pairwise comparison with Benjamini-Hochberg correction.

256 All data are presented as mean \pm SE and all statistics were performed using R software
257 (version 3.6.1) with the significance level set at 0.05.

258

259 **Results**

260 **Plant phenology**

261 In June 2018, the studied population contained 47 flowering individuals, all hermaphrodites.
262 The mean number of inflorescences per plant was 3.17 ± 0.44 , while the mean number of
263 cymes per inflorescence was 14.30 ± 0.81 . Moreover, the mean number of male flowers per
264 inflorescence was 2.69 ± 0.171 , while the mean number of female flowers per inflorescence
265 was 21.07 ± 0.858 . On the basis of the data collected on the population structure the ratio of
266 male and female floral stages in the observation patches was determined at 1:9.

267

268 **Nectar analyses**

269 *Sugars and volume*

270 Mean nectar volume per flower showed a clear trend of increasing in relation to floral age,
271 with volume in buds statistically lower than in both male- and female-phase flowers ($U = 15$,
272 $p = 0.009$ and $U = 2$, $p = 0.001$, respectively). A significant difference for mean sugar
273 quantity per flower was also reported between buds and female-phase flowers (Tukey's HSD:
274 $p = 0.028$), whilst sugar concentration did not differ significantly among floral stages (Table
275 1).

276 A more in depth analysis on sugars reported that hexose sugar quantity per flower in the bud
277 stage differed significantly from both male- and female-phase flowers ($U = 12$, $p = 0.008$ and
278 $U = 19$, $p = 0.018$, respectively), whilst sucrose quantity per flower found in bud differed
279 statistically only from the average amount found in the female stage (Tukey's HSD: $p =$

280 0.021; Table 1). Mean percentage of sucrose per flower did not appear to be significantly
281 different among floral stages (Table 1).

282

283 *Amino acids*

284 There was no significant difference for total, protein, and non-protein amino acid quantity per
285 flower between male and female flowers, while the ratio between protein and non protein
286 amino acid concentrations was significantly higher for male-phase flowers (Table 1).

287 The only amino acid with a statistically significant difference was phenylalanine ($t_{14} = 2.94$, p
288 $= 0.011$), showing a higher concentration in male floral phase ($M = 352.7 \pm 63.2 \text{ nmol mL}^{-1}$
289 and $F = 143.6 \pm 32.6 \text{ nmol mL}^{-1}$; Fig. 1).

290 Among all protein amino acids, proline and phenylalanine showed the highest concentrations:
291 the former appeared to reach higher concentrations in the functionally female stage ($674.8 \pm$
292 $243.5 \text{ nmol mL}^{-1}$), whilst the latter in the functionally male stage ($352.7 \pm 63.2 \text{ nmol mL}^{-1}$).

293 Among non protein amino acids, in both male and female stages GABA showed the highest
294 concentration ($51.4 \pm 12.2 \text{ nmol mL}^{-1}$ and $202.0 \pm 73.4 \text{ nmol mL}^{-1}$, respectively).

295 The number of different amino acids (richness) detectable in the male stage was significantly
296 lower than number of amino acids in the female stage ($t_{15} = 3.54$, $p = 0.003$; 16.5 ± 0.6 and
297 19.0 ± 0.3 , respectively), while no differences were found in Simpson, Shannon and Evenness
298 indices between male and female stages (Table 2).

299

300 **Insect visit analyses**

301 *Flower visitors' abundance*

302 A total of 215 insect visits were recorded on *Echium vulgare* during 200 minutes of field
303 surveys (Table 3).

304 Visitors belonged to three order: Hymenoptera (87.4%), Lepidoptera (9.8%) and Diptera
305 (2.8%). The order Hymenoptera was mainly represented by individuals belonging to the
306 family Megachilidae (59%), followed by the family Halictidae (26.5%) and Apidae (14%).
307 The order Lepidoptera was represented mainly by individuals belonging to the species
308 *Macroglossum stellatarum* (43%) and the family Pieridae (43%). The order Diptera was
309 represented only by 6 individuals belonging to the families Bombyliidae and Syrphidae. The
310 most frequent visitors were solitary bees of the species *Hoplitis adunca* (42%).

311

312 *Flower visitor observations*

313 Among the 215 insects visiting the plant, we fully recorded data for 189 individuals.
314 Statistical analyses were carried out only on the 112 individuals which were looking for
315 nectar and for which the number of total visits exceeded 5 (*Macroglossum stellatarum*,
316 Pieridae, *Anthidium florentinum*, *Apis mellifera* and *Hoplitis adunca*). The family Pieridae
317 was analysed as a single taxon in order to reach a total number of visits above 5. Since
318 *Hoplitis adunca* was the most abundant taxon and the only species strongly oligolectic on
319 *Echium*, we therefore decided to analyse the sexes separately.
320 Although nectar is produced before flower opening and insects can force the bud searching
321 for nectar (personal observation), this event occurred very rarely. Consequently, we did not
322 consider the phenological stage bud in these analyses.
323 For each insect taxon, we compared the number of visits to male and female flowers with the
324 expected ones, calculated according to the ratio 1:9 between male and female flowers
325 registered in the studied population.
326 Regarding the number of male flowers visited, no significant difference was reported for
327 lepidopterans (Pieridae spp., *Macroglossum stellatarum*) and for females *Hoplitis adunca*,
328 while *Anthidium florentinum*, *Apis mellifera* and *Hoplitis adunca* males visited more male

329 flowers than expected (Table 4). The number of female flowers visited was never statistically
330 different from that expected.

331 The frequency of male flowers visited in relation to the total number of flowers visited among
332 taxa was statistically different ($H_4 = 14.01$, $p = 0.016$). Statistical analyses confirmed that the
333 female *Hoplitis adunca* visited fewer male flowers than did *Anthidium florentinum* ($U = 65$, p
334 $= 0.002$), *Apis mellifera* ($U = 48$, $p = 0.002$) and *Macroglossum stellatarum* ($U = 28.5$, $p =$
335 0.043 ; Fig. 2).

336

337 **Discussion**

338 Our studied population did not show the phenomenon of gynodioecism, as all flowers were
339 hermaphrodite, and our data confirmed the ratio of 1:9 found by Rademaker et al. (1999)
340 between functionally male and functionally female flowers.

341 Our analyses confirmed that nectar is secreted in the bud, as reported by Chwil and
342 Weryszko-Chmielewska (2011). Contrary to Klinkhamer and de Jong (1990), we found that
343 nectar volume, as well as sugar quantity per flower, increased with the age of the flower (from
344 bud to female phase), although the positive trend between male and female phases was not
345 statistically significant. Both quantity of hexose sugars and sucrose per flower increased with
346 the age of the flower, the latter reaching a mean almost 7 fold higher in functionally female
347 flowers than the mean amount found in the bud stage and almost twice the amount found in
348 functionally male flowers. At the same time, the mean percentage of sucrose per flower
349 appeared to be lower in male-phase flowers, even though not significantly, meaning that the
350 total sugar increase in relation to floral age is due to the rise of nectar volume, since total
351 sugar concentration and composition remained constant during the entire flower phenology.

352 The existence of nectar homeostasis mechanisms which actively maintain a constant nectar

353 sugar concentration to ensure pollinator visits has been previously reported in other species
354 (Nepi and Stpiczyńska 2008; Nepi et al. 2011).

355 When we compared the number of insect visits on male and female flowers observed to the
356 expected ones, all bee taxa except female *Hoplitis adunca* showed a higher number of visits to
357 male flowers than expected. This result could be explained by the higher proportion of protein
358 amino acids found in the male stage: preferences have often been reported in bees for protein
359 amino acid enriched solutions (Inouye and Waller 1984; Bertazzini et al. 2010; Hendriksma et
360 al. 2014), suggesting that flower visitors may actively choose to visit functionally male
361 flowers. Comparable results have been reported by Klinkhamer and de Jong (1990) and by
362 Rademaker et al. (1999) on bumblebees: when calculating the probabilities of visits on
363 different floral stages, the oldest female stage was less likely to be visited than a male-phase
364 flower. Females of *Hoplitis adunca* are the only bees collecting both pollen and nectar on *E.*
365 *vulgare*: this different foraging behaviour might explain the difference from the other bee
366 species.

367 Individuals of *Lasioglossum* sp. were observed visiting the flower and collecting pollen only.
368 A tendency for afternoon trips for nectar only have been reported for the subfamily Halictinae
369 by Michener (2003) so we cannot conclude that *Lasioglossum* sp. does not exploit *E. vulgare*
370 nectar since the species may simply collect the resource at different time of the day.

371 Despite Lepidoptera having been reported to prefer nectar rich in PAAs (Baker and Baker
372 1986; Erhardt and Rusterholz 1998), our study reports that Pieridae butterflies visited as many
373 male flowers as expected, indicating that these insects did not actively look for functionally
374 male flowers (containing a higher proportion of protein amino acids). A study conducted by
375 Alm et al. (1990) showed that male individuals of the species *Pieris rapae* do not discriminate
376 between artificial nectars containing sugar only or sugar solution enriched with protein amino
377 acids, and Romeis and Wäckers (2000) reported that feeding and source-selection in *Pieris*

378 *brassicae* is elicited by sucrose more than protein amino acids. We report a similar result for
379 the species *Macroglossum stellatarum*, but to date no study has been done in order to assess
380 amino acid preferences in the species and whether taste receptors on the proboscis can sense
381 their presence in nectar remains unsubstantiated (Stöckl and Kelber 2019).

382 Nectar of male-phase flowers in *E. vulgare* presented, among all the amino acids, the highest
383 concentration of phenylalanine, representing an average of 35% of total amino acid content.
384 Phenylalanine is an essential protein amino acid (de Groot 1953) and several studies proved
385 that it exerts a phagostimulatory effect on several insects, especially on honey bees, and it is
386 strongly correlated with pollinator preferences (Inouye and Waller 1984; Hendriksma et al.
387 2014; Tiedge and Lohaus 2017; Seo et al. 2019). Consequently, this could explain the higher
388 frequency of visit on male flowers than expected. A correlation between phenylalanine
389 concentration and nectar feeding by Megachilids, that were the more numerous pollinators in
390 our study, was demonstrated in a phragmic community, a plant association typical of the East
391 Mediterranean (Petanidou et al. 2006).

392 Proline, instead, represented the most concentrated amino acid in functionally female flowers,
393 and the second in the early-stage functionally male flowers (representing more than 30% and
394 almost 20% of the total amino acid content, respectively). This non-essential amino acid,
395 commonly found in nectar (Nicolson and Thornburg 2007), can stimulate the insect salt cell
396 increasing intensity of feeding behaviour (Hansen et al. 1998; Wacht et al. 2000). Proline also
397 represents an energy substrate to fuel the earliest or most expensive stages of insect flight
398 (Micheu et al. 2000; Gade and Auerswald 2002), resulting in short-term bursts of energy
399 production (Teulier et al. 2016).

400 Finally, in both male- and female-phase flower nectar GABA showed the highest
401 concentration among the non-protein amino acids representing more than 5% and 9% of total
402 amino acid content, respectively. Recent studies indicated that GABA could affect both

403 insects' physiology and behaviour, feeding rate and flight muscles performances (Shelp et al.
404 2017; Felicioli et al. 2018; Bogo et al. 2019). Besides GABA, or possibly the combination of
405 GABA and NaCl, can constitute an important nectar phagostimulant and its presence
406 correlates with visits by an array of pollinators such as long tongued bees, ex-anthophorid and
407 andrenid bees, as well as anthomyiid and syrphid flies (Petanidou 2007 and reference
408 therein).

409 The spectrum of visitors recorded through our observations confirm that reported by previous
410 studies stating that flowers of *E. vulgare* are visited by hummingbird hawkmoths (Aguado
411 Martin et al. 2017), bees, bee flies (Proctor et al. 1996) and syrphids (Willmer and Finlayson
412 2014). Also, even though the species has often been reported as mainly pollinated by
413 bumblebees (Corbet 1978; Klinkhamer and de Jong 1990; Pappers et al. 1999; Rademaker et
414 al. 1999), we observed only one individual of *Bombus pascuorum* visiting the
415 flowers. Pollinators of wide spread plant species can vary in relation to their geographical
416 distribution (Armbruster 1985; Thompson 2006; Pérez-Barrales et al. 2007) and, moreover, as
417 reported by Lázaro et al. (2010), the plant and pollinators assemblages of an entire community
418 may also influence the composition of visitors of a particular species by determining, for
419 instance, the strength of competition or the intensity of attraction to that species rather than
420 another. Thus, the scarcity of bumblebees observed on *Echium vulgare* in 2018 may either
421 depend on several factors and/or reflect a temporal fluctuation in the species composition of
422 the pollinator community, as previously reported by many studies (Cane et al. 2005;
423 Petanidou et al. 2008; Dupont et al. 2009).

424

425 **Conclusions**

426 The inbreeding avoidance hypothesis states that some mechanisms develop within a species in
427 order to prevent breeding among related individuals and its damaging effects on fitness

428 (Darwin 1876, 1877; Charlesworth and Charlesworth 1987). In dichogamous species, gender-
429 biased nectar often occurs (Carlson and Harms 2006; Stpiczyńska et al. 2015; Konarska and
430 Masierowska 2020), and this, according to the mentioned above hypothesis, may contribute to
431 decrease geitonogamous selfing through its effects on a pollinator's behaviour (Carlson and
432 Harms 2006). Our results suggest that the quality of nectar offered by the two sexually
433 distinct floral phases may target different insect needs, thus affecting simultaneously different
434 behavioural traits and ensuring an optimal pattern of visit among functionally different floral
435 stages, unequally present in the population throughout the anthesis period. The more
436 nutritional nectar found in the less frequent sexual phase occurring in the population (male
437 flowers) may enhance movements among plants by encouraging "better-resource hunt",
438 whilst the flight efforts accomplished for doing so may be sustained by a rapidly oxidable fuel
439 such proline offered in female-phase flowers. In the light of this hypothesis, it appears clear
440 that gender-biased nectar studies in dichogamous, many-flowered species should be
441 undertaken in relation to the occurrence of floral sexual phases in the population (when a bias
442 in the frequency of sex occurrence exists).

443 Despite no study yet providing strong scientific evidence that gender-biased nectar in fact
444 reduces inbreeding (Carlson and Harms 2006), it is reasonable to assume that by offering
445 variable quality nectar through sexually different floral phases the plant may produce a
446 mosaic of food targeting different pollinator behavioural traits aiming to promote cross-
447 pollination.

448

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453

454 **Conflict of interest**

455 The authors declare that they have no conflict of interest.

456

457 **Data availability**

458 Data available from the Zenodo Digital Repository: <http://doi.org/xxxxxxxxx> (Barberis,

459 Bogo et al. 2020)

460

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Codice campo modificato

714 **Figure captions**

715 **Figure 1.** Amino acid concentrations (nmol mL⁻¹) detected in functionally male (dark bars)
716 and in functionally female (light bars) flowers (mean ± SE). Amino acids hydroxyproline,
717 homoserine, citrulline, cysteine, histidine, glutamine, asparagine and L-thyronine were not
718 detected in either floral stages and thus not shown in the graph. The asterisk denotes a
719 statistically significant difference according to Student t-test. NPAA = non-protein amino
720 acids; PAA = protein amino acids.

721

722 **Figure 2.** Frequency of male flowers visited by each taxon. Different letters denote statistical
723 differences according to Kruskal Wallis H-test followed by Mann-Withney pairwise
724 comparison with Benjamini-Hochberg correction ($p < 0.05$).

725

726 **Table 1.** Comparison of nectar volume, sugar and amino acid (AA: amino acids; PAA: protein
 727 amino acids; NPAA: non-protein amino acids) compositions among the three phenological
 728 stages (bud, male and female flowers). Values (expressed by mean \pm SE) marked with different
 729 letters were significantly different according to one-way ANOVA or Kruskal-Wallis test
 730 followed by the respective post hoc test with Benjamini-Hochberg correction.

Nectar parameters	Bud	Male flower	Female flower	Test value	p-value
Volume ($\mu\text{L flower}^{-1}$)	0.159 \pm 0.019 a	0.427 \pm 0.080 b	0.669 \pm 0.135 b	$H_2 = 16.83$	< 0.001
Total sugar ($\mu\text{g flower}^{-1}$)	0.013 \pm 0.006 a	0.040 \pm 0.013 ab	0.070 \pm 0.026 b	$F_{2,27} = 5.78$	< 0.001
Total sugar concentration ($\mu\text{g } \mu\text{L}^{-1}$)	0.089 \pm 0.033	0.094 \pm 0.022	0.090 \pm 0.020	$F_{2,27} = 0.45$	0.642
Hexose sugars ($\mu\text{g flower}^{-1}$)	0.005 \pm 0.004 a	0.007 \pm 0.001 b	0.008 \pm 0.002 b	$H_2 = 11.43$	0.003
Sucrose ($\mu\text{g flower}^{-1}$)	0.009 \pm 0.003 a	0.033 \pm 0.012 ab	0.061 \pm 0.024 b	$F_{2,27} = 5.63$	0.007
Sucrose (% per flower)	82.278 \pm 7.824	72.896 \pm 5.776	81.900 \pm 3.817	$H_2 = 4.10$	0.129
Total AA (nmol flower^{-1})	-	0.367 \pm 0.061	1.349 \pm 0.611	$U = 21$	0.270
PAA (nmol flower^{-1})	-	0.321 \pm 0.054	1.058 \pm 0.467	$U = 23$	0.372
NPAA (nmol flower^{-1})	-	0.045 \pm 0.007	0.290 \pm 0.145	$U = 15$	0.083
PAA:NPAA ratio	-	7.31 \pm 0.670	4.65 \pm 0.437	$t_{14} = -3.34$	0.005

731

732

733 **Table 2.** Comparison of diversity indices calculated on nectar amino acid concentration
734 between male and female phases (8 samples for both floral phases).

Diversity indices	Male flower	Female flower	t	p-value
Amino acids richness	16.50 ± 0.627	19.00 ± 0.327	3.54	0.003
Simpson	0.793 ± 0.035	0.822 ± 0.024	0.68	0.506
Shannon <i>H</i>	2.109 ± 0.103	2.233 ± 0.111	0.82	0.428
Evenness	0.527 ± 0.059	0.511 ± 0.050	-0.20	0.842

735

736

737 **Table 3.** *Echium vulgare* visitors recorded in June 2018 (215 visits in total), their abundance
 738 and the percentage of them looking for nectar as reward.

Order	Family	Species	Relative frequency	Looking for nectar (%)
Hymenoptera	Apidae	<i>Apis mellifera</i> Linnaeus, 1758	0.079	100
Hymenoptera	Apidae	<i>Bombus pascuorum</i> (Scopoli, 1763)	0.005	100
Hymenoptera	Apidae	<i>Ceratina</i> (Latreille, 1802) sp.	0.023	100
Hymenoptera	Apidae	<i>Eucera</i> (Scopoli, 1770) sp.	0.018	100
		<i>Lasioglossum interruptum</i> (Panzer, 1798)		
Hymenoptera	Halictidae	<i>Lasioglossum laticeps</i> (Schenck, 1869)	0.233	0
		<i>Lasioglossum corvinum</i> (Morawitz, 1878)		
Hymenoptera	Halictidae	<i>Halictus subauratus</i> (Rossi, 1792)	0.005	100
Hymenoptera	Colletidae	<i>Hylaeus</i> cfr. <i>angustatus</i> (Schenck, 1859)	0.005	100
Hymenoptera	Megachilidae	<i>Anthidium florentinum</i> (Fabricius, 1775)	0.102	100
Hymenoptera	Megachilidae	<i>Hoplitis adunca</i> (Panzer, 1798)	Male: 0.191 Female: 0.219	Male: 100 Female: 66.6 ^a
Diptera	Bombyliidae	<i>Bombylius</i> (Linnaeus, 1758) sp.	0.009	100
Diptera	Syrphidae	Syrphidae (Latreille, 1802) sp.	0.019	0
		<i>Hesperia comma</i> (Linnaeus, 1758)		
Lepidoptera	Hesperiidae	<i>Thymelicus acteon</i> (Rottemburg, 1775)	0.019	100
Lepidoptera	Papilionidae	<i>Iphioides podalirius</i> (Linnaeus, 1758)	0.005	100
		<i>Pieris brassicae</i> (Linnaeus, 1758)		
Lepidoptera	Pieridae	<i>Pieris mannii</i> Mayer, 1851 <i>Colias croceus</i> (Fourcroy, 1785) <i>Pontia edusa</i> (Fabricius, 1777)	0.042	100
Lepidoptera	Sphingidae	<i>Macroglossum stellatarum</i> (Linnaeus, 1758)	0.042	100

739 ^avalue calculated only on individuals with fully recorded data (n = 21)

740

741 **Table 4.** Male (a) and female (b) flowers visited by each taxon (mean \pm SE). Chi-square test is
 742 calculated on the basis of the ratio 1:9 between male and female flowers occurred in the studied
 743 population.

a)				
Taxon	Male flowers visited	χ^2	d.f.	p-value
<i>Anthidium florentinum</i>	0.96 \pm 0.192	37.80	21	0.014
<i>Apis mellifera</i>	1.59 \pm 0.384	39.39	16	<0.001
<i>Hoplitis adunca</i> male	0.51 \pm 0.100	70.51	40	0.002
<i>Hoplitis adunca</i> female	0.14 \pm 0.143	8.50	13	0.810
<i>Macroglossum stellatarum</i>	2.33 \pm 0.799	4.54	8	0.806
Pieridae	0.33 \pm 0.236	5.21	8	0.735
b)				
Taxon	Female flowers visited	χ^2	d.f.	p-value
<i>Anthidium florentinum</i>	3.95 \pm 0.826	4.20	21	1.000
<i>Apis mellifera</i>	7.47 \pm 1.652	4.38	16	0.998
<i>Hoplitis adunca</i> male	2.37 \pm 0.312	7.84	40	1.000
<i>Hoplitis adunca</i> female	1.64 \pm 0.199	0.94	13	1.000
<i>Macroglossum stellatarum</i>	15.67 \pm 14.696	0.50	8	1.000
Pieridae	4.22 \pm 1.656	0.58	8	1.000

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