Flower visitation by hoverflies (Diptera: Syrphidae) in a temperate plant-pollinator network

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14 ABSTRACT

Hoverflies (Diptera: Syrphidae) are among the most important pollinators, although they attract less 15 attention than bees. They are usually thought to be rather opportunistic flower visitors, although pre-16 vious studied demonstrated that they show colour preferences and their nectar feeding is affected by 17 morphological constraints related to flower morphology. Despite the growing appreciation of hoverflies 18 and other non-bee insects as pollinators, there is a lack of community-wide studies of flower visitation by 19 syprhids. The aim of this paper is to provide a detailed analysis of flower visitation patterns in a species 20 rich community of syrphids in a Central European grassland and to evaluate how species traits shape 21 the structure of the plant-hoverfly flower visitation network. We found that different species varied in the 22 level of specialisation, and while some species visited a similar spectre of flowers, others partitioned 23 resources more strongly. There was a consistent difference in both specialisation and flower preferences 24 between three syrphid subfamilies. Eristalinae and Pipizinae were more specialised than Syrphinae. 25 Trait-based analyses showed that relative flower visitation i) increased with plant height, but most strongly 26 in Eristalinae; ii) increased with inflorescence size in small species from all three subfamilies, but was 27 independent of inflorescence size in large species of Eristalinae and Syrphinae; and iii) depended on 28 flower colour, but in a subfamily-specific way. Eristalinae showed the strongest flower colour preferences 29 for white flowers, Pipizinae visited mostly white and yellow flowers, while Syrphinae were less affected 30 by flower colour. Exploration of the structure of the plant-hoverfly flower visitation network showed that 31 the network was both modular and nested. We also found that there were almost no differences in specialisation and relative visitation frequency between males and females. Overall, we showed that 33 flower visitation in syrphids was affected by phylogenetic relatedness, body size of syrphids and several 34 plant traits. 35

36 INTRODUCTION

Hoverflies (Diptera: Syrphidae) are one of the most abundant groups of flower visiting insects. Together 37 with other families of flies, their role in plant-pollinator interactions is often underappreciated (Inouye 38 et al., 2015). However, Diptera often make up a similar proportion of flower visitors as Hymenoptera and 39 are even the dominant group of pollinators in some habitats, e.g. in higher altitudes and latitudes (Kanstrup 40 and Olesen, 2000). Apart from being important pollinators of many wild plants (Orford et al., 2015; 41 Sakurai and Takahashi, 2017; Moquet et al., 2018), hoverflies play an important role also in pollination of 42 numerous crops (Ssymank et al., 2008; Inouye et al., 2015; Rader et al., 2016). For example, hoverflies 43 of a genus *Eristalis* have been successfully used for pollination of peppers in greenhouses (Jarlan et al., 44 1997), and are pollinators of rapeseed (Ohsawa and Namai, 1987, 1988; Jauker and Wolters, 2008; Rader 45 et al., 2009), apple trees (Solomon and Kendall, 1970), strawberries (Kendall et al., 1971), etc. (Inouye

et al., 2015). Hoverflies are thus an important group of pollinators not only from the perspective of 47 biodiversity conservation, but also for pollination of crops in agricultural settings. The interest in the role 48 of flies in general and Syrphidae in particular as pollinators has been increasing (Ssymank et al., 2008). 49

However, our knowledge of their preferences for different flowers and their partitioning of floral resources 50

51 is still limited.

Adults of all known syrphid species feed almost exclusively on pollen and nectar or honeydew

52 (Rotheray and Gilbert, 2011) and are usually considered as generalist flower visitors (Branquart and 53 Hemptinne, 2000). However, individual species cover a broad range from generalists to species with 54 strong preferences for a small number of plants (Branquart and Hemptinne, 2000; Colley and Luna, 2000). 55 Their flower preferences may, however, shift depending on local flower availability and plant phenology 56 (Cowgill et al., 1993; Colley and Luna, 2000). Several studies reported that selectivity of some hoverfly 57 species depends on certain plant traits. Overall, hoverflies seem to visit mostly open bowl-shaped flowers 58 (Branquart and Hemptinne, 2000), where they feed on both nectar and pollen (Gilbert, 1981), but some of 59 them have relatively long proboscises which allow them to reach nectar even in flowers with relatively 60 long spurs (Gilbert, 1981; Vlašánková et al., 2017). Additional plant traits affecting flower visitation by 61 at least some species are inflorescence height (Gervasi and Schiestl, 2017; Klecka et al., 2017). It has 62 been recently demonstrated that selective flower visitation by hoverflies can exert a selection pressure 63 strong enough to cause rapid evolutionary shifts in multiple plant traits (Gervasi and Schiestl, 2017; Zu 64

and Schiestl, 2017). However, there are still very few studies focusing on flower visitation patterns of 65 entire local assemblages of syrphids. 66

Ecological and evolutionary processes which are responsible for structuring of communities of 67 interacting insects and plants discussed above in the specific case of syrphids also lead to distinct 68 structural features of entire plant-flower visitor networks. Most of these networks are usually strongly 69 nested (Bascompte et al., 2003), modular (Olesen et al., 2007), or both nested and modular at the same 70 time (Fortuna et al., 2010). Modules pack species connected by numerous interactions and can be linked 71 to trait complementarity (Olesen et al., 2007). On the other hand, nested structure means that specialised 72 insects tend to interact with plants also visited by more generalised insects and vice versa. In this case, 73 mutual specialisation is rare, which has a stabilising effect on the structure of plant-flower visitor networks 74 (Burgos et al., 2007; Bastolla et al., 2009). 75 The aim of this paper is to advance our understanding of flower visitation by hoverflies by a thorough 76

analysis of flower visitation in a species-rich community of plants in a Central European grassland. We 77 focus on the level of specialisation of different species, their overlap in resource use, and the structure of 78 the entire plant-hoverfly flower visitation network. Our analyses show that hoverflies in our study area 79 varied in their level of specialisation, partitioned floral resources depending on their relatedness, body 80 size and several plant traits, and formed a plant-hoverfly flower visitation network which was nested and 81 modular at the same time. 82

METHODS 83

We conducted sampling of plant-flower visitor interactions in a small area in the southern part of the 84 Czech Republic between the northern edge of the city of Český Krumlov and nearby villages Vyšný and 85 Lazec in June-August 2015. We gathered observations from eight flower-rich grassland patches between 86 48°49'29.5"N, 14°18'59.5"E in the South, 48°49'42.6"N, 14°19'24.4"E in the East, and 48°50'7.0"N, 87 14°15'36.5" E in the North-West. Geographic coordinates of individual sites are given in Supplementary 88 Table 1. Sampling was conducted on public land and did not involve any protected species. For this 89 reason, we did not need to obtain any permits for this project. 90

We sampled flower-visiting insects by transect walks and collected all visitors to flowers of all 91 herbaceous flowering plants except grasses, although some syrphids are known to feed on pollen of 92 grasses and sedges (Ssymank and Gilbert, 1993). Sampling was carried out during sunny days with 93 no rain between 08:00 and 17:00 hours. In this paper, we focus on hoverflies (Diptera: Syrphidae), 94 which were among the most abundant groups of flower visiting insects in the study area. All collected 95 individuals were killed by ethyl-acetate, transported to the lab, pinned, and identified using keys of van 96 Veen (2010) and Speight and Sarthou (2014) to the species level with the exception of hoverflies of the 97 genus *Pipizella* and a few damaged specimens. We also identified the sex of all individuals. Several 98 individuals per species were photographed using a stereomicroscope Olympus SZX7 and a DSLR camera 99 Canon 700D controlled from a computer by digiCamControl software. We measured body length, thorax 100

Table 1. The list of species of Syrphidae collected during this study. The number of observations and the number of plant species visited is provided for each species and sex. Species-level identification was not possible in *Pipizella, Sphegina*, and in a few damaged individuals of *Platycheirus*.

	No.	of observa	tions	No. of plant specie		es visited
	Total	Females	Males	Total	Females	Males
Eristalinae						
Arctophila bombiformis	2	0	2	1	0	1
Cheilosia proxima	1	0	1	1	0	1
Cheilosia ruficollis	2	2	0	2	2	0
Chrysogaster basalis	1	1	0	1	1	0
Chrysogaster coemiteriorum	3	3	0	2	2	0
Chrysogaster solstitialis	32	20	12	4	2	3
Eristalis arbustorum	16	5	11	4	2	3
Eristalis horticola	2	2	0	2	2	0
Eristalis interruptus	56	32	24	12	10	8
Eristalis intricarius	1	0	1	1	0	1
Eristalis pertinax	2	1	1	2	1	1
Eristalis sepulchralis	1	0	1	1	0	1
Eristalis similis	1	0	1	1	0	1
Eristalis tenax	10	8	2	9	8	2
Helophilus pendulus	3	1	2	3	1	2
Myathropa florea	5	3	2	4	2	2
Neoascia podagrica	4	4	0	2	2	0
Orthonevra nobilis	2	1	1	1	1	1
Rhingia campestris	11	4	7	6	3	6
Rhingia rostrata	2	0	2	2	0	2
Sericomyia silentis	11	6	5	4	3	3
Sphegina sp.	1	1	0	1	1	0
Syritta pipiens	124	72	52	24	18	16
Vollucela bombylans	4	4	0	2	2	0
Vollucela pellucens	2	2	0	2	2	0
Pipizinae						
Heringia pubescens	1	1	0	1	1	0
Pipiza nocticula	5	5	0	4	4	0
<i>Pipizella</i> sp.	84	41	43	16	9	10
Syrphinae						
Chrysotoxum bicinctum	4	3	1	4	3	1
Chrysotoxum cautum	4	4	0	1	1	0
Chrysotoxum fasciatum	1	1	0	1	1	0
Chrysotoxum vernale	1	1	0	1	1	0
Chrysotoxum verralli	1	0	1	1	0	1
Didea alneti	4	1	3	2	1	2
Episyrphus balteatus	194	91	103	29	23	22
Eupeodes bucculatus	4	1	3	4	1	3
Lapposyrphus lapponicus	69	23	46	22	12	17
Melanostoma mellinum	18	8	10	10	6	7
Melanostoma scalare	1	0	1	1	0	1
Meliscaeva cinctella	1	0	1	1	0	1
Paragus haemorrhous	3	2	1	2	2	1
Parasyrphus lineolus	4	2	2	3	2	2
Platycheirus albimanus	2	2	0	2	2	0
Platycheirus peltatus	1	0	1	1	0	1
Platycheirus scambus	1	1	0	1	1	0
Platycheirus sp.	2	0	2	2	0	2
Scaeva pyrastri	18	6	12	10	6	7
Sphaerophoria scripta	237	143	94	47	38	34
Syrphus ribesii	58	35	23	16	14	10
Syrphus torvus	159	71	88	28	21	21
Syrphus vitripennis	18	12	6	13	9	6
Xanthogramma pedissequum	1	1	0	1	1	0

width, and head width of at least 8 individuals, or all individuals in species where less individuals were
 collected, using the Fiji distribution (Schindelin et al., 2012) of ImageJ (Rueden et al., 2017) and a plugin
 Microscope Measurement Tools. All specimens are deposited in Jan Klecka's collection at the Institute of
 Entomology, Biology Centre of the Czech Academy of Sciences.

We measured a set of three traits of all plant species visited by hoverflies to test their effect on flower visitation (Supplementary Table 4). Specifically, we measured plant height (the height of the top flower above ground), inflorescence size as the largest distance between any two open flowers within an inflorescence, and classified flower colour into four categories (blue, purple, white, and yellow), similar to previous studies (Haslett, 1989a). We conducted plant trait measurements in several patches for each species, measuring at lest 10 plant individuals per site, except in very rare species.

For data analysis, we pooled observations from the entire study area, because most sampled patches 111 were close to each other, often <1 km apart, well within dispersal range of most hoverflies (Rader et al. 112 2011; Moquet et al., 2018). We performed all analyses in R 3.2.3 (R Core Team, 2015). We visualised 113 the structure of plant-hoverfly flower visitation network using the package bipartite 2.08 (Dormann et al., 114 2009; Dormann, 2011). We used generalised linear models (GLM) with either a Poisson distribution with 115 overdispersion (quasipoisson), a Binomial distribution with overdispersion (quasibinomial), or Normal 116 distribution depending on the properties of the response variable. We used non-metric multidimensional 117 scaling implemented in vegan 2.4-4 package for R (Oksanen et al., 2017) to visualise diet overlap between 118 syrphid species. Diet overlap was calculated using Pianka's overlap index Pianka (1973) using the 119 plant-hoverfly flower visitation matrix (Supplementary Table 2) to estimate pairwise diet overlap values 120 between all pairs of hoverfly species. We performed the diet overlap analysis using EcoSimR 0.1.0 121 package for R (Gotelli et al., 2015). Finally, we performed analyses of modularity and nestedness of the 122 plant-hoverfly flower visitation network using the package bipartite 2.08 for R (Dormann et al., 2009; 123 Dormann, 2011). 124

125 RESULTS

We observed 1195 interactions between a total of 51 species of syrphids from three subfamilies (Eristalinae,
 Pipizinae, and Syrphinae) and 57 plant species from 20 families (Table 1). The network of plant-syrphid
 flower visiting interactions is shown in Fig. 1, raw data are available in Supplementary Tables 1 and 2.

The number of plant species visited by individual syrphid species increased with the number of 129 observations linearly on a log-log scale ($F_{1,49} = 873.41, P < 1 * 10^{-6}$) and differed significantly between 130 subfamilies, with Syrphinae visiting more plant species than Eristalinae and Pipizinae after accounting for 131 the number of observations ($F_{2,48} = 4.16$, P = 0.0216; Fig. 2A.). Species from the subfamily Syrphinae 132 were more generalised than the other two subfamilies also according to our calculation of a specialisation 133 index d' ($F_{2,15} = 4.81, P = 0.0243$; Fig. 2B.), which was restricted to species with at least five observations. 134 Body length had no effect on the number of plant species visited ($F_{1,47} = 0.47, P = 0.4986$), nor on the 135 value of the specialisation index d' ($F_{1,14} = 1.28, P = 0.2764$). The same results were obtained using 136 head width and thorax width as alternative measures of body size. Comparison of the specialisation index 137 d' revealed no consistent differences in specialisation between males and females (linear mixed effects 138 model with species as a random factor; $\chi_1^2 = 1.99, P = 0.1586$) (Table 2). 139

There was a clear differentiation between Syrphinae and Eristalinae in their flower preferences based 140 on nonmetric multidimensional scaling (NMDS) with values of pairwise diet overlap between all pairs of 141 species of syrhids (Fig. 3). Values of Pianka's overlap index of individual syrphid species pairs ranged 142 from 0 to 0.988 (mean = 0.363, Supplementary Table 3), i.e. from completely different to almost identical 143 pattern of visitation of flowers of individual plant species. We included only syrphid species with at least 144 five observations in this analysis. The two species of Pipizinae included in the analysis did not cluster 145 together, although they appeared distinct from the other two subfamilies (Fig. 3). Additional insight into 146 differences between the three subfamilies can be gained from a comparison of visitation frequency on 147 plants from different families shown in Fig. 4. 148

Males and females showed only minor differences in their preferences for flowers of different plant species. Comparison of the frequency of flower visits of males and females in individual plant species revealed a significant difference only in *Eristalis interruptus* (χ^2 test for contingency tables; $\chi^2 = 21.76, P = 0.0048; P$ estimated by 10000 Monte Carlo simulations). Males of *Eristalis interruptus* visited mostly *Daucus carota*, while females visited mostly *Centaurea scabiosa* and *Achillea millefolium* (Supplementary File 1). Other abundant species showed only minor differences in flower visitation, but



Figure 1. Flower visitation network of plants and hoverflies. Hoverflies are displayed in the upper row with blue boxes. The width of the boxes is proportional to the number of individuals observed. Plants are displayed in the lower row as green boxes whose width is proportional to the number of observations made at the individual plant species. The width of the connecting lines is proportional to the number of interactions observed between each plant-syrphid pair. The most abundant species are named, all species are identified by numbers - see Supplementary Table 2 for legend.



Figure 2. The level of resource specialisation of syrphids. (A.) The number of plant species visited depended on the number of observations. Variation around the regression line shows that species below the line were more specialised than expected and species above the line were more generalised. The green line is mostly hidden by the blue line. (B.) Syrphinae were more generalised than Eristalinae and Pipizinae.

the number of observations was low in many cases (Supplementary File 1); species with <10 observations
 per sex were not included in these analyses.

Relative visitation rate of plants by individual syrphid species significantly increased with plant height, 157 but with a different slope in syrphids from different subfamilies and different size classes (Figure 5A., 158 Table 3). The effect of inflorescence size also differed between small and large syrphids and between 159 species from different subfamilies (Figure 5B., Table 3). Specifically, relative visitation increased with 160 inflorescence size in small syrphids from all three subfamilies, but was independent of inflorescence 161 size in large syrphids (Figure 5B.). Flower colour also had a significant effect on visitation by syrphids, 162 but this effect varied between the three syrphid subfamilies (Figure 5C.-E., Table 3). Eristalinae clearly 163 visited mostly white flowers (Figure 5C.), Pipizinae had a similar visitation rate to yellow and white 164 flowers (Figure 5D.), while Syrphinae showed only minor differences in visitation of flowers of different 165 colours (Figure 5E.). When we look at the syrphid community as a whole, plants with yellow and white 166 flowers were overall most frequently visited, with 37.7% of visits to yellow flowers and 31.4% to white 167 flowers. Purple flowers received 26.1% visits, while blue flowers received only 4.9% of visits. In contrast 168 to subfamily, syrphid size class did not affect the dependence of visitation on flower colour (flower colour 169 x syrphid size class interaction, F = 1.25, P = 0.2921; Table 3). 170

Analysis of the structure of the plant-syrphid flower visitation network showed that the network 171 was both modular and nested at the same time. Modularity analysis detected four modules, with most 172 Eristalinae clustered in one module, while Syrphinae dominated two other modules, and the most 173 generalised species, Sphaerophoria scripta, was classified separately (Fig. 6). The network was not only 174 modular, but also nested (Fig. 7). Nestedness index was significantly different from random expectation 175 (NODF = 29.08, P < 0.001, based on 999 simulations). However, comparison of nestedness calculated 176 for syrphids and plants separately showed that only the syrphids had significantly nested pattern of 177 interactions (NODF = 18.02, P < 0.001), while the pattern for plants was not significantly different from 178 random expectation (NODF = 38.27, P = 0.221). 179

180 DISCUSSION

Flower visitation by Syrphidae was characterised by a variable degree of specialisation at the species
 level. Syrphids have been traditionally considered as generalised flower visitors. We showed that not only
 different species fell in different positions along a gradient from more specialised to truly generalised

Table 2. Values of the specialisation index d' of male and female syrphids. Larger values of the d' index correspond to more specialised flower visitation. Species where one or both sexes had <5 observations were excluded from the analysis.

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	Specialisation (d')		
Species	Females	Males	
Chrysogaster solstitialis	0.428	0.405	
Episyrphus balteatus	0.265	0.253	
Eristalis arbustorum	0.286	0.275	
Eristalis interruptus	0.273	0.262	
Lapposyrphus lapponicus	0.171	0.231	
Melanostoma mellinum	0.248	0.166	
<i>Pipizella</i> sp.	0.272	0.282	
Scaeva pyrastri	0.186	0.247	
Sericomyia silentis	0.341	0.287	
Sphaerophoria scripta	0.126	0.171	
Syritta pipiens	0.234	0.271	
Syrphus ribesii	0.163	0.214	
Syrphus torvus	0.158	0.258	
Syrphus vitripennis	0.067	0.249	



Figure 3. Results of Nonmetric multidimensional scaling (NMDS) show differences in flower preferences in Syrphinae and Eristalinae. NMDS analysis was run with a matrix of dissimilarities of the relative frequency of flower visitation on different plants by individual species of syrphids. The position of individual species in the plot corresponds to the center of the species label. *S. scripta* = *Sphaerophoria scripta*.



Figure 4. Comparison of the proportion of flower visits by the three subfamilies of Syrphidae to individual plant families. The bars show the proportion of observations of flower visits depending on plant family.



Figure 5. The effect of species traits on flower visitation by Syrphidae. A. Taller plants were visited more frequently by syrphids with a slope dependent on their body size and subfamily. B. Small syrphids visited more often plants with large inflorescences, while visitation by large syrphids was not affected by inflorescence size. C.-E. Flower colour affected visitation by the three subfamilies of syrphids differently.

Table 3. The effect of species traits on flower visitation by syrphids. Results of a GLM testing the dependence of relative visitation rate on species traits. Significance of all interaction terms in the model is shown.

Model term	df	F	Р
Log ₁₀ (plant height) x syrphid subfamily	2	4.09	0.0170
Log_{10} (plant height) x syrphid size class	1	4.39	0.0365
Log_{10} (inflorescence size) x syrphid subfamily	2	3.11	0.0449
Log_{10} (inflorescence size) x syrphid size class	1	7.31	0.0070
Flower colour x syrphid subfamily	6	4.07	0.0005
Flower colour x syrphid size class	3	1.25	0.2921

flower visitors, but that there were also significant differences in average specialisation between the three syrphid subfamilies. The pattern of higher specialisation of Eristalinae and Pipizinae compared to the more generalised Syrphinae was clear, although in the case of Pipizinae, we have to note that our observations included only three species of Pipizinae (counting individuals of *Pipizella* sp., which could be identified only to genus, as one species).

We also found pronounced differences in relative flower preferences both at a coarse level between 189 subfamilies, and at a finer level between species. Results of the NMDS showed that Eristalinae and 190 Syrphinae were nicely separated in the diagram, but also that species from the same genus clustered 191 together, e.g. the three species of each of the genera *Eristalis* and *Syrphus* (Fig. 3). Pairwise diet overlap 192 values were as high as 0.98 in *Eristalis arbustorum* and *Chrysogaster solstitialis* (the maximum possible 193 value is 1.0), and the three species of the genus Syrphus had diet overlap values between 0.84 and 0.88, 194 which suggests that they had almost identical diets. On the other hand, many species showed clear diet 195 partitioning evidenced by small values of diet overlap (Supplementary Table 3). This is noteworthy in 196 relation to ongoing debates about mechanisms of species coexistence. Conflicting theoretical explanations 197 of species coexistence showed that species can coexist only if they are sufficiently different according to 198 classic theory of limiting similarity (Hardin, 1960; MacArthur and Levins, 1967), or alternatively if they 199 are sufficiently similar as proposed by Hubbell's neutral theory (Hubbell, 2001). An emerging consensus 200 is that both explanations are correct, i.e. that species can coexist if they are either sufficiently different 201 or sufficiently similar (Scheffer and van Nes, 2006; Sakavara et al., 2018; Scheffer et al., 2018). Indeed, 202 there are examples of closely related coexisting species with different trophic niches, as well as examples 203 where they have a very similar niche (Goulson et al., 2008). In our case, we observed local coexistence 204 of some closely related species with very high diet overlaps, which is consistent with the argument that 205 similar species can coexist. Coexistence may be facilitated by differences in flower-visiting behaviour, 206 such as microhabitat choice (Janovský et al., 2013) or timing of activity during the day (Gilbert, 1980). 207

Trait-based analysis of relative visitation rate of flowers by individual syrphid species showed that 208 phylogenetic relatedness, i.e. belonging to the same or different subfamily, affected flower colour 209 preferences. We did not measure plant abundance in sufficient detail to test whether relative visitation 210 rates to different flower colour categories significantly deviated from a random pattern, but we can gain 211 insight into flower colour preferences from comparison of different species collected in the same study 212 area. This comparison revealed differences in the relative preference for white, yellow, purple, and blue 213 flowers between the three syrphid subfamilies. So, we can say that Eristalinae appeared to strongly prefer 214 white flowers, while Syrphinae were relatively indiscriminate in the colour of flowers they visited (Fig. 5). 215 Previous studies on the effect of flower colour in hoverflies suggested that they visit mostly white or yellow 216 flowers (Haslett, 1989a; Sutherland et al., 1999), with some exceptions, such as *Rhingia campestris* with a 217 preference for blue flowers (Haslett, 1989a), but previous studies were restricted to a small set of species. 218 We show that flower colour preferences varied between different syrphid subfamilies. Interestingly, 219 Haslett (1989a) observed that out of a group of six species, *Episyrphus balteatus* from the subfamily 220 Syrphinae was the least selective species towards flower colour, while several species from the subfamily 221 Eristalinae were more selective. Our data with a larger set of species provide compelling evidence of this 222 difference between subfamilies. There is not enough know about the visual system of different species, 223 but it is likely that interspecific differences in visitation of flowers of different colours represent foraging 224 preferences rather than differences in the visual system which seems to be quite uniform among flies 225 (Lunau, 2014). The dominant flower colour represents probably a relatively long-range visual signal, 226 227 while other cues may be used when the hoverfly approaches the flower. For example, small yellow spots are known to elicit an innate proboscis extension response and serve as cues visually guiding hoverflies 228 towards pollen, which is usually yellow (Lunau and Wacht, 1994). Not only anthers with exposed pollen, 229 but also other small yellow structures may thus guide the hoverfly towards pollen once it reached the 230 flower (Lunau and Wacht, 1994; Lunau, 2014). 231

Based on our results, Eristalinae and Pipizinae showed a stronger response to all plant traits, i.e. flower colour, plant height and inflorescence size, compared to Syrphinae. Taken together, these results highlight the differences in average specialisation level between the generalised Syrphinae on one side and more specialised Eristalinae and Pipizinae on the other side. Interestingly, Moquet et al. (2018) found that they could split hoverflies of Belgian heathlands according to an analysis of several life-history and ecological traits into two distinct groups roughly corresponding to the two dominant subfamilies, Eristalinae and Syrphinae. Our detailed analysis of flower visitation provides additional evidence of important ecological ²³⁹ differences between the syrphid subfamilies.

Apart from phylogenetic relatedness at the subfamily level, we found body size to be an important 240 trait modifying the responses of syrphid relative visitation rate to selected plant traits. Flower colour 241 was related mostly to phylogenetic relatedness, while plant height and inflorescence size were related 242 also to syrphid body size. Small syrphids preferred large inflorescences, which may be advantageous 243 because they could exploit resources clustered in one place (Akter et al., 2017). Another trait that has been 244 evaluated previously is the relationship between corolla depth and proboscis length. Some previous studies 245 showed a positive correlation between the average depths of flowers and proboscis length or length/width 246 ratio in bees (Stang et al., 2006, 2009) as well as hoverflies (Gilbert, 1981; Branquart and Hemptinne, 247 248 2000). We did not test this relationship mostly because we did not distinguish nectar and pollen feeding. Even species with a short proboscis are regularly visiting long-spurred flowers to feed on pollen and can 249 even lick nectar at the entrance to the spur without being able to reach deep inside (Vlašánková et al., 250 2017). Proper analysis of a morphological fit between the flowers and flower visitors would thus require a 251 more detailed data on mechanisms of feeding by individual species and on morphology of both the insects 252 and the flowers. 253



Figure 6. Modules detected in the plant-syrphid flower visitation network. Results of modularity analysis restricted to syrphid and plant species with at least 5 observations. Syprhidae are displayed in rows and plants in columns. The blue rectangles show observed interactions with more frequent interactions shown by darker colour The three syrphid subfamilies are distinguished by coloured circles next to the species names (see legend).

Network modularity (Olesen et al., 2007) partly reflected these patterns, because we found that the 254 plant-hoverfly flower visitation network could be partitioned into four modules; most Eristalinae were 255 clustered in one of them. Nestedness analysis showed that the syrphid flower visitation was significantly 256 nested, i.e. that more specialised species visited mostly flowers of plants which were a subset of those 257 visited by more generalised species, which is a typical pattern in plant-flower visitor networks (Bascompte 258 et al., 2003; Fortuna et al., 2010). However, nestedness of the plants did not differ from a random 259 expectation, so the nestedness pattern was asymmetric. This is likely because the plants were visited by a 260 range of other insects, not only hoverflies, so the network as we analysed it here was incomplete from the 261 plants' point of view. 262

Despite the clear patterns we found at the interspecific level, we detected very little differences in

flower visitation by males and females of species sufficiently abundant to allow such comparison. Both the 264 level of specialisation and the relative visitation rates to individual plant species were very similar in males 265 and females in most cases. Similarly, Sutherland et al. (1999) found that males and females of *Episyrphus* 266 balteatus showed very similar flower colour preferences. However, we did not distinguish nectar and 267 pollen consumption during our observations, so we cannot rule out a possible difference between sexes 268 in pollen vs. nectar feeding. Indeed, several previous studies reported that females of hoverflies feed 269 on pollen more frequently than males (Gilbert, 1981; Haslett, 1989b; Hickman et al., 1995), probably 270 because proteins from pollen are necessary for egg development. Males thus often feed less on pollen 271 and more on nectar which serves mostly as a source of energy for their active lifestyle, because they 272 are usually more active than females and spend a large amount of time by hovering (Haslett, 1989b). 273 However, no significant difference in pollen consumption between males and females was found in a few 274 other species, so the generality of this patterns is unclear (Irvin et al., 1999). 275



Figure 7. The plant-hoverfly flower visitation network was significantly nested. Syprhidae are displayed in rows and plants in columns. The blue rectangles show observed interactions.

276 Conclusions

Hoverflies and other dipterans are important pollinators, but they have been neglected compared to bees (Ssymank et al., 2008; Orford et al., 2015), and we need to learn more about they foraging biology, flower preferences, and pollination efficiency. We provided insights into some of these issues. However, there are questions we did not consider, such as feeding on pollen of wind-pollinated plants either by visiting flowers of grasses, sedges, trees, etc., or by eating pollen accumulated on the surface of leaves (Ssymank and Gilbert, 1993; Saunders, 2017). Apart from the need for more detailed understanding of foraging biology of hoverflies and other flower-visiting flies, there is a lot of unknowns about their pollination

- efficiency, although some studies demonstrated that hoverflies and other dipterans may provide pollination 284
- service comparable to bees (Kearns and Inouye, 1994; Inouye et al., 2015). Filling these knowledge gaps 285
- is urgently needed given the reported widespread declines of many native pollinators around the world 286
- (Potts et al., 2010). 287

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