

Plant choice for oviposition in the phytophagous insect Bemisia tabaci: cytotype, including symbionts, knows best!

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1	Plant choice for oviposition in the phytophagous insect Bemisia
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20 Abstract

Most phytophagous insects harbour symbiotic microorganisms that may facilitate, at 21 22 multiple levels, the exploitation of plants. Actually, very little is known about the role of these symbionts in insects' host plant choice. The whitefly Bemisia tabaci is a good model to 23 explore these topics. It harbours a nutritional 'primary' endosymbiont allowing it to thrive on 24 plant sap, and up to seven 'secondary' endosymbionts (S-symbionts) whose roles in plant 25 utilization are currently under study. Here, we first investigated the influence of cytotype on 26 27 females' oviposition rate on hibiscus and lantana, respectively a favourable and an unfavourable host plant for *B. tabaci*. We then addressed whether cytotype affects whiteflies' 28 plant choice for oviposition. Using genetically homogeneous whiteflies harbouring different 29 cytotypes, we showed that oviposition rate was significantly lower on lantana than on 30 hibiscus, but that this reduction varied between cytotypes. Moreover, according to their 31 32 cytotype, whiteflies preferred hibiscus over lantana, or showed no preference. Thus, cytotype influenced plant choice for oviposition. Altogether, our results indicate that, without a 33 cytotype that facilitates the utilization of lantana, whiteflies may avoid this host plant. Since 34 35 cytotypes harbour different S-symbionts' combinations, we argue that the influence of these symbionts on whitefly oviposition behaviour is very likely. 36

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Keywords: Bemisia tabaci, cytotype, oviposition behaviour, plant choice, symbionts

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38 Highlights

- Cytotype (including symbionts) affects whiteflies oviposition rate
- Cytotype determines plant choice for oviposition
- Different symbionts' combinations affect whitefly oviposition behaviour

42 Introduction

Microbial symbiosis is widespread in eukaryotes. Some symbionts are parasites with 43 44 negative effects on their host. Other symbionts are mutualists with beneficial effects, or commensals with no apparent effects (Buchner, 1965; Ferrari & Vavre, 2011; Tipton et al., 45 2019). Infection by parasites often induces host behavioural changes. Some of these changes 46 improve transmission and/or survival of the parasite at the expense of host fitness and are 47 considered as 'parasite's manipulations' (Heil, 2016; Lewis & Lize, 2015; Poulin, 2010; Vale 48 49 et al., 2018). On the contrary, other behavioural changes benefit the host and participate to its defence strategies against parasites (Hart, 2011; Vale et al., 2018). While symbiont-driven 50 behavioural alterations have been frequently observed in conflicting interactions, host 51 52 behavioural modifications in mutualistic or commensal interactions have received less attention. However, recent discoveries indicate that these types of symbiotic interactions may 53 have various effects on insect behaviours related to locomotion (Schretter et al., 2018), 54 reproduction (Najarro et al., 2015; Sharon et al., 2010), foraging (Akami et al., 2019; 55 Angelella et al., 2018; Leitão-Goncalves et al., 2017; Leybourne et al., 2020; Qiao et al., 56 57 2019; Wong et al., 2017) or interactions with other organisms (Dion et al., 2011; Polin et al., 2014; Sochard, Bellec, et al., 2021) (for a review, see also Hosokawa & Fukatsu, 2020). 58

Most insect species are associated with microbial symbionts, which colonize their gut, body cavities or cells (Hansen & Moran, 2014). In phytophagous insects, mutualistic symbionts can facilitate the utilization of food sources that are unbalanced (*e.g.* phloem sap) or difficult to digest (*e.g.* wood) through the provisioning of essential and otherwise limiting nutrients, the detoxification of plant defence compounds or the breakdown of plant polymers, respectively (Frago et al., 2012; Hansen & Moran, 2014). Much less is known about the influence of symbionts in host plant selection. Recent works have shown that the gut

symbiotic bacteria can affect food choice and foraging decisions under laboratory conditions 66 67 in the fruit fly Drosophila melanogaster (Leitão-Gonçalves et al., 2017; Qiao et al., 2019; Wong et al., 2017) and the oriental fruit fly Bactrocera dorsalis (Akami et al., 2019). Other 68 studies on phloem sap-feeding hemipterans have demonstrated that infection by intracellular 69 bacteria can affect their interaction with the plants by modifying plant probing behaviour, like 70 in the cowpea aphid Aphis craccivora (Angelella et al., 2018) and the bird cherry oat aphid 71 Rhopalosiphum padi (Leybourne et al., 2020). On the other hand, intracellular symbionts have 72 no or only moderate influence on host plant selection in the pea aphid Acyrtosiphon pisum 73 (Sochard, Dupont, et al., 2021; Sochard, le Floch, et al., 2021). Studies on a broader spectrum 74 75 of host-symbiont associations are needed to better assess the role of symbionts in insect plant choice. This may also shed light on the potential impact of symbiosis on hosts' ecological 76 diversification and speciation (Brucker & Bordenstein, 2012; Vavre & Kremer, 2014). Indeed, 77 78 for phytophagous insects, being restricted to specific plants is likely to limit gene flow between populations (Bird et al., 2012; Schluter, 2001). 79

The tobacco whitefly Bemisia tabaci (Hemitpera: Aleyrodidae) constitutes an 80 interesting model to address this question. It represents a highly diversified cryptic species 81 complex and, to date, 42 putative species have been proposed in phylogenetic analyses based 82 83 on mitochondrial cytochrome oxidase 1 (*mtCOI*) sequences (de Barro et al., 2011; Dinsdale et al., 2010; Firdaus et al., 2013; Roopa et al., 2015). Previous studies reported significant 84 variation in plant preference and/or utilization capabilities between B. tabaci species. Some 85 are 'generalist' species with a broad host plant range, while others can be considered as 86 'specialist' species, with a restricted host plant range (Malka et al., 2018). Differences can 87 also be seen within species: indeed, in some species, genetic mitochondrial groups can be 88 identified, as in the case of the Mediterranean (MED) species, which can be divided into three 89 groups, Q1, Q2 and Q3 (Mouton et al., 2015; Vyskočilová et al., 2018). While the Q1 and Q2 90

91 groups are relatively polyphagous (Vyskočilová et al., 2019), field observations and 92 laboratory experiments indicate that the Q3 group, only identified in Burkina Faso to date, has 93 a more restricted host range and seems specialized on lantana (*Lantana camara*) (Romba et 94 al., 2018; Romba & Gnankiné, 2018), a host plant that is otherwise unfavourable to the MED 95 species (Benhamou et al., 2021). Other comparative experiments revealed host range 96 differences between Q1 and Q2, and among Q1 populations from different geographical areas 97 (Vyskočilová et al., 2019).

As other phloem sap-feeding insects, B. tabaci harbours an obligate 'primary' 98 intracellular symbiont (P-symbiont) that provide essential amino acids which are not present 99 in sufficient quantities in the phloem sap for whiteflies' development, survival and 100 reproduction (Baumann, 2005; Douglas, 2009). This y-proteobacterium, Portiera 101 aleyrodidarum (hereafter Portiera), is common to all members of the B. tabaci species 102 103 complex (Thao & Baumann, 2004). Portiera resides within the cytoplasm of specialized host cells, the bacteriocytes, associated to form an organ, the bacteriome (Gottlieb et al., 2008). 104 105 Portiera is vertically transmitted through a mechanism which is unique to whiteflies, with an 106 entire bacteriocyte being transferred to the next generation (Buchner, 1965; Luan et al., 2016,2018). In addition to this P-symbiont, B. tabaci harbour facultative 'secondary' 107 symbionts (S-symbionts). Seven bacterial genera of S-symbionts have been reported to date: 108 Arsenophonus, Cardinium, Fritschea, Hamiltonella, Hemipteriphilus, Rickettsia and 109 Wolbachia (Bing et al., 2013; Everett et al., 2005; Gottlieb et al., 2006; H. Wang et al., 2020; 110 111 Zchori-fein & Brown, 2002), with up to four present in the same insect body (Zchori-Fein et al., 2014). They all co-occur within the same bacteriocytes as Portiera, but some of them can 112 also colonize other tissues of the host (Gottlieb et al., 2008). B. tabaci S-symbionts are 113 predominantly vertically transmitted, but some of them can also be horizontally transmitted, 114 for instance through the phloem when whiteflies feed upon the same plant (Caspi-Fluger et 115

al., 2012; S. Li et al., 2016; Y. Li et al., 2017). Analyses of B. tabaci S-symbionts suggest that 116 some of them could contribute to their hosts' nutrition. Indeed, *Portiera*'s genome has 117 undergone drastic reduction (Rao et al., 2015) as a consequence of its prolonged intracellular 118 119 lifestyle (Mccutcheon & Moran, 2012; Moran et al., 2008). It has lost almost all genes involved in the synthesis of vitamins and cofactors, and several of its essential amino acid 120 pathways are incomplete (Rao et al., 2015), functions that S-symbionts may replace or 121 complement (Opatovsky et al., 2018; Rao et al., 2015; Ren et al., 2020; Y.-B. Wang et al., 122 2020). As an example, Hamiltonella can provide vitamins and cofactors (Rao et al., 2015; 123 Ren et al., 2020; Y.-B. Wang et al., 2020), and could also complete the missing steps of the 124 125 essential amino acid lysine biosynthesis pathway of *Portiera* (Rao et al., 2015). Interestingly, by producing metabolites able to interact with their hosts' neurons or sensory organs or that 126 are precursors of neurotransmitters, S-symbionts may directly influence their hosts' behaviour 127 (Engl & Kaltenpoth, 2018; Shropshire & Bordenstein, 2016). Importantly, S-symbiont 128 composition differs between mitochondrial groups, both across (Zchori-Fein et al., 2014) and 129 within (Gnankiné et al., 2013; Gueguen et al., 2010) species, raising the possibility that these 130 131 symbionts may contribute to *B. tabaci* species complex ecological diversification.

Recently, we demonstrated that S-symbionts might affect the range of the host plants 132 133 suitable for *B. tabaci*. Indeed, we have shown that cytotype (referring to the cytoplasmic characteristics of the insect line, including the mitochondria and intracellular symbionts) 134 affects whitefly capability to feed and oviposit on lantana (Lantana camara), an unfavourable 135 host plant to B. tabaci (Benhamou et al., 2021). The aims of the present study were to 136 investigate whether whiteflies exhibit plant choice for oviposition and whether the S-symbiont 137 community may influence their oviposition behaviour. To fulfil these objectives, we 138 performed choice experiments between a plant known to be favourable for *B. tabaci*, hibiscus 139

(*Hibiscus moscheutos*), and the unfavourable lantana, using genetically homogeneous
whiteflies harbouring different cytotypes (and, hence, different S-symbionts).

142 Material and Methods

143 Insects

Three B. tabaci parental lines (F0) from the Mediterranean (MED) species were used 144 145 in this study: AA(Q1-HW), BB(Q1-HR) and CC(Q2-ARW) that belong to the mitochondrial groups Q1 or Q2, and that are associated with different S-symbionts (Table 1). The first 146 uppercase double letters indicate females' diploid nuclear genotype (AA, BB or CC) (males 147 are haploid as *B. tabaci* reproduces by arrhenotoky). Characters within parenthesis refer to 148 their cytotype, including their mitochondrial group (O1 or O2) and S-symbionts (HW, HR or 149 150 ARW; A: Arsenophonus, H: Hamiltonella, R: Rickettsia, W: Wolbachia). Insect lines were reared on hibiscus plants in mesh cages in climate-controlled rooms at $26^{\circ}C \pm 1^{\circ}C$ and 70% 151 relative humidity with a 14L:10D photoperiod. The same climate-controlled conditions have 152 been used for the whole study. 153

154 Table 1. *Bemisia tabaci* laboratory lines used in this study.

	Nuclear . Genotype	Cytotype		Collection information			
Line		Mitochondrial group	S-Symbionts ^a	Year	Location (Country)	Plant	
AA(Q1-HW)	AA	Q1	HW	2012	Tympaki	Eggplant	
					(Greece)	Solanum sp.	
BB(Q1-HR)	BB	Q1	HR	2012	Les Ponts-de-Cé	Mandevilla	
					(France)	<i>Mandevilla</i> sp.	
CC(Q2-ARW)	CC	Q2	ARW	2018	Lyon	Lantana	
					(France)	Lantana camara	

^a A: Arsenophonus, H: Hamiltonella, R: Rickettsia, W: Wolbachia.

156 Collection of F1 females

To separate cytotype-related and genotype-related effects, experiments were 157 performed on F1 females obtained from the 9 possible crosses between the F0 parental lines 158 (Table 2, experimental design shown in Fig. S1), based on the procedure described in 159 Benhamou *et al.* (Benhamou et al., 2021). These crosses produced either F1 females with 160 identical nuclear genotypes (the two letters refer to the maternally (first) and paternally 161 (second) inherited nuclear genotype) but different cytotypes [e.g. AB(Q1-HW) and BA(Q1-162 HR), or the opposite, *i.e.* females with the same cytotype but different genotypes [*e.g.* 163 AA(Q1-HW), AB(Q1-HW) and AC(Q1-HW)]. The three lines were synchronized as follows: 164 adults from the hibiscus stock cages were allowed to mate and oviposit for 7 days on cotton 165 leaves. To keep them fresh, leaves were fixed on their adaxial side on a 1cm layer of 2% Agar 166 (Sigma-Aldrich®) in plastic Petri dishes (90mm in diameter). Petri dishes were placed upside-167 168 down to simulate the situation of leaves attached to a plant with the abaxial surface down. Cotton was used as it is not involved in the subsequent experiments, thus avoiding possible 169 effect of the F0 individuals' plant on the traits assessed on the following F1 generation (Fig. 170 **S1**). We chose cotton because it is a known favourable plant for the *B. tabaci* MED species 171 groups Q1 and Q2 (Romba & Gnankiné, 2018; Vyskočilová et al., 2018). F0 females were 172 collected within 5 hours after emergence. Individuals' sex was determined based on their 173 genitalia, by observation under a stereomicroscope (Leica EZ4, Leica Microsystems GmbH, 174 Wetzlar, Germany). F0 females (<5-h-old) were considered virgin as newly emerged adults 175 176 do not immediately mate (Li et al., 1989). As B. tabaci reproduces by arrhenotoky, which means that unmated females produce only haploid eggs that develop into males, this was 177 confirmed by the absence of females in their offspring. Each cross was made on cotton 178 179 (90mm in diameter Petri dishes) using groups of 15 F0 males and 15 F0 females (3 replicates for each). These 30 individuals were left for 1 week on the same cotton leaf to mate and 180

181 oviposit. Upon emergence, F1 virgin females (<5-h-old) were collected for further 182 experiments as described hereafter. The whole procedure (crosses and experiments) was 183 independently performed three times, treated as "experimental blocks" in the statistical 184 analyses.

			Mother (F0) ^a	
		AA(Q1-HW)	BB(Q1-HR)	CC(Q2-ARW)
Fathar	A(Q1-HW)	AA(Q1-HW)	BA(Q1-HR)	CA(Q2-ARW)
Father (F0) ^b	B(Q1-HR)	AB(Q1-HW)	BB(Q1-HR)	CB(Q2-ARW)
(FU)	C(Q2-ARW)	AC(Q1-HW)	BC(Q1-HR)	CC(Q2-ARW)

^a F0 and F1 Females identification: Nuclear genotype(Cytotype); Nuclear genotype: letters indicate the maternally (first) and the paternally (second) inherited nuclear genotype; Cytotype: Mitochondrial group (Q1 or Q2) + S-symbionts; S-symbionts: A: *Arsenophonus*; H: *Hamiltonella*; R: *Rickettsia*; W: *Wolbachia*.
 ^bF0 Fathers' genotypes (A, B or C) are indicated by only one letter as males are haploid.

185 Table 2. Nuclear genotypes and cytotypes of *B. tabaci* F1 females.

186 Cytotype determination

187 For each parental line (Table 1) and F1 cross (Table 2), mitochondrial group (Q1 or 188 Q2) and symbiont infection status (Portiera, Arsenophonus, Hamiltonella, Rickettsia, Wolbachia) were checked on 8 females to ensure that the expected cytotype was correct. Total 189 DNA was extracted from individual whiteflies using the NucleoSpin 95 Tissue kit (Macherey 190 Nagel GmBH, Düren, Germany) following the manufacturer's instructions. Samples were 191 crushed in a mix of 60µL of lysis buffer and 8µL of protein K (22mg/mL) with 1.4mm sterile 192 stainless-steel beads using a TissueLyser (Qiagen, Hilden, Germany). Elution was done in 193 100µL of buffer and DNA was then stored at -20°C. The mitochondrial haplotype was 194 checked by PCR-RFLP according to the method described in Henri et al., 2013), 195 allowing to differentiate between the Q1 and Q2 groups. The infection status was determined 196 using quantitative PCR (qPCR) using the following genes: 16S rRNA for Arsenophonus, 197

198 *dnaK* for *Hamiltonella*, 16S rRNA for *Portiera*, *gltA* for *Rickettsia* and *ftsZ* for *Wolbachia* 199 (**Table 3**). The β -actin gene from *B. tabaci* (**Table 3**) was used as a DNA extraction control. 200 The PCR amplifications were performed on a CFX-96 Real Time PCR machine (Bio-Rad, 201 Hercules, USA). Each reaction consisted of 5µL of SYBR® Green 2x Supermix (Bio-rad) in 202 addition to 2µL of DNA sample, 2µL of nuclease-free water, and 0.5µL of forward and 203 reverse primers (500nM).

0	Gene	Primers	D. t	Fragment	Hybridization	D
Organism			Primer sequence	size (bp)	T ° (° C)	Reference
Bemisia tabaci	β -actin	wf-Bactin-F	5'-TCTTCCAGCCATCCTTCTTG -3'	130	63	(Sinisterra et
		wf-Bactin-R	5'-CGGTGATTTCCTTCTGCATT-3'			al., 2005)
Portiera	16S	Port73-F	5'-GTGGGGAATAACGTACGG-3'	193	60	(Caspi-Fluger
		Port266-R	5'-CTCAGTCCCAGTGTGGCTG-3'			et al., 2011)
Arsenophonus	16S	ArsF3	5'-GTCGTGAGGAARGTGTTARGGTT-3'	765	63	(Duron et al.,
		ArsR3	5'- CCTYTATCTCTAAAGGMTTCGCTGGATG-3'			2008)
Hamiltonella	dnaK	dnaK-F	5'-GGTTCAGAAAAAGTGGCAG-3'	155	60	(Moran et al.,
		dnaK-R	5'-CGAGCGAAAGAGGAGTGAC-3'			2005)
Rickettsia	gltA	glt375-F	5'-TGGTATTGCATCGCTTTGGG-3'	199	60	(Caspi-Fluger
		glt574-R	5'-TTTCTTTAAGCACTGCAGCACG-3'			et al., 2011)
Wolbachia	ftsZ	F2	5'-TTGCAGAGCTTGGACTTGAA-3'	400	55	(Vavre et al.,
		R2	5'-CATATCTCCGCCACCAGTAA-3'			1999)

Table 3. qPCR primers used in this study.

205 Plants

Plants were grown in the same climate-controlled conditions as the ones used for rearing lines, in insect-free rooms. Cotton, *Gossypium hirsutum* L. (Malvaceae) cv. 'Elpida', and hibiscus, *Hibiscus moscheutos* L. (Malvaceae) cv. 'Honeymoon', were grown from seeds while lantana, *Lantana camara* L. (Verbenaceae) was vegetatively propagated from cuttings in pots using a 75:25 (v/v) mix of potting soil and vermiculite. For all experiments performed in this study, fully developed leaves, randomly collected from 2-month-old plants, have been used.

213 No-choice oviposition and egg hatching rate

The influence of whitefly cytotype on females' oviposition and egg hatching rate on 214 hibiscus and lantana was investigated using no-choice oviposition tests. F1 virgin females 215 216 (<5-h-old), which developed on cotton, were put individually on hibiscus or lantana leaf disks (20mm in diameter, to standardize the leaf area) fixed on an agar layer in Petri dishes (55mm 217 218 in diameter), and placed upside down. After 7 days, females were removed and the number of 219 eggs laid (oviposition) was recorded under a stereomicroscope. Ten days later, the number of eggs that hatched was estimated by counting the number of larvae (including both live and 220 221 dead larvae). This method was used because the larvae, even dead, are easier to identify than 222 the empty eggs. We also calculated the number of eggs that did not hatch (number of eggs laid - number of eggs hatched). Thirteen to 17 individual measures were obtained per 223 cytotype/genotype combination and per plant. 224

225 Plant choice for oviposition

Plant choice for oviposition was investigated in binary choice experiments between 226 227 hibiscus and lantana. The experimental setup used was inspired from the one developed by Shah and Liu (Shah & Liu, 2013). Choice arenas (produced 1 hour before use) consisted of 228 229 one lantana and one hibiscus leaf disk (20mm in diameter), fixed on their adaxial side on an agar layer in a Petri dish (90mm in diameter). Disks were arranged opposite and at equal 230 distance (20mm) from the center of the arena (Fig. S1). For this experiment, we used 2-d-old 231 females to ensure they were old enough to oviposit from the very beginning of the tests, at it 232 lasted less time than the previous no choice experiments. Newly emerged virgin females (<5-233 234 h-old), which developed on cotton, were transferred on fresh cotton leaves (55mm in diameter Petri dishes) for 2 days. Tests were initiated every day in the morning at the same hour with 235 the introduction of one F1 female (2-d-old) into the center of each arena, which were then 236

placed upside down. Thirty-two hours later, the female's location was recorded and the number of eggs laid on each leaf disk was counted under a stereomicroscope. As far as possible, experiments were run every day for all nine cytotype/genotype combination simultaneously. Measures were considered to be valid when females were found alive on a leaf disk and if they laid at least one egg at the end of the tests. Thereby, 31 to 37 valid measures were obtained per cytotype/genotype combinations.

243 Statistical analyses

For no-choice experiments, oviposition and egg average hatching rate were analysed 244 by a generalized linear model (GLM) (with a log link and a negative binomial or a quasi-245 246 binomial error structure, respectively) with the plant species, the insect cytotype, the insect nuclear genotype (as a nested effect within the cytotype) and the experimental block set as 247 explanatory variables. The average hatching rate was calculated according to the formula (a), 248 where i is the replicate number and N is the total number of replicates. Eggs that hatched were 249 considered as 'successes' ('No.successes'), and eggs that did not hatch as 'failures' 250 251 ('No.failures').

252 (a) average proprotion of successes =
$$\frac{\sum_{i=1}^{N} (No.successes)}{\sum_{i=1}^{N} (No.successes+No.failures)}$$

For the plant choice experiment, oviposition was analysed by a GLM (with a quasibinomial error structure) with the insect cytotype, the insect nuclear genotype (integrated into the model as a nested effect within the cytotype) and the experimental block set as explanatory variables. The response variable is expressed as the average proportion of eggs laid on lantana compared to the total number of eggs laid on lantana (accounted as 'successes') plus hibiscus (accounted as 'failures'), given by the formula (a). The number of eggs laid on hibiscus or lantana according to females cytotype and genotype in the plant

choice experiment was analysed using a GLM (with a negative binomial error structure). 260 From the complete models, we performed a model simplification through a backward 261 procedure (Crawley, 2013). Multiple comparison analyses were carried out by Tukey's test 262 (Tukey, 1949) to investigate statistical relationships between explanatory variables levels 263 using the emmeans R package (Russell, 2020). The emmeans R package was also used to 264 extract means, average proportions and 95% confidence intervals (95%CI) from the models. 265 All data analyses were performed using the R software (R Development Core Team 2011; 266 http://www.R-project.org). 267

268 Data accessibility

The datasets generated and analysed in this study can be obtained from the Zenodo
online repository at: https://zenodo.org/record/6414033.

271 **Ethical note**

This research adheres to the ASAB/ABS Guidelines for the Use of Animals in
Research, the legal requirements of France and the institutional guidelines of the University of
Lyon.

275 **Results**

276 *No-choice oviposition and hatching rate*

The influence of whitefly cytotype on females' ability to feed and produce offspring on favourable and non-favourable plants were investigated by measuring females' oviposition and egg hatching rate on hibiscus or lantana in no-choice experiments (**Fig. 1**). For the number of eggs laid, there was no difference between the three experimental blocks [$\chi^2(2)$ =

1.60, P = 0.45]. For a given cytotype, there was no difference between genotypes [$\gamma^2(6)$ = 281 3.57, P = 0.74]. There was also no tripartite interaction between plant, cytotype and genotype 282 $[\chi^2(6) = 1.43, P = 0.96]$. However, there was a significant interaction between the plant 283 species and the insect cytotype [$\chi^2(2) = 103.20$, P < 0.001]. This interaction results from the 284 fact that, while the number of eggs laid was homogeneous among cytotypes on hibiscus 285 (mean, 37.71; standard error [SE]: 1.31) (P > 0.99), it was significantly reduced on lantana for 286 all cytotype/genotype combinations (P < 0.001), but to different extents between cytotypes. 287 Indeed, females with the cytotype Q1-HW consistently laid a lower number of eggs on 288 lantana (mean, 5.98; SE: 0.74) than females with cytotypes Q1-HR (mean, 23.50; SE: 1.40) (P 289 290 < 0.001) or Q2-ARW (mean: 21.88; SE: 1.41) (P < 0.001) that laid a similar number of eggs on this plant (P = 0.98) (Fig. 1A). These results confirm our previous data (Benhamou et al., 291 2021) showing that lantana is globally a less favourable host plant than hibiscus, but also 292 293 highlight that some cytotypes mitigate this effect.

Regarding the average egg hatching rate (**Fig. 1B**), there was no difference between experimental blocks [F(2,202) = 1.91, P = 0.15]. We also did not detect any influence of the plant species [F(1,238) = 1.35, P = 0.25] or insect cytotype [F(2,238) = 0.65, P = 0.52], and no interaction between these two factors [F(2,238) = 1.26, P = 0.29]. There was no difference between genotypes for each cytotype [F(6,238) = 0.27, P = 0.95], and no interaction between plant, cytotype and genotype [F(6,238) = 0.72, P = 0.63].

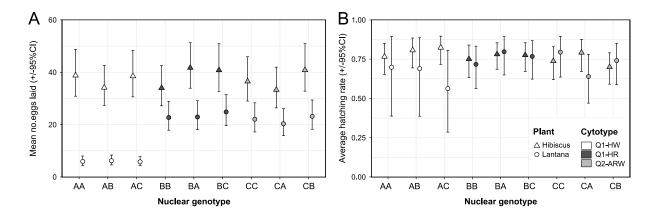


Figure 1. Effect of the insects' genotype and cytotype on *B. tabaci* females' oviposition 300 301 and egg hatching rate on hibiscus and lantana (no choice experiment). (A) Mean number of eggs laid per female [± 95% confidence interval (CI)] over 7 days. (B) Average egg 302 303 hatching rate (± 95% CI) after 10 days. All individuals are F1 females obtained from crosses of laboratory lines: AA(Q1-HW), BB(Q1-HR), or CC(Q2-ARW). F1 female legend key: 304 Nuclear genotype: letters indicate the maternally (first) and the paternally (second) inherited 305 nuclear genotype; Cytotype: mitochondrial group+S-symbionts (A: Arsenophonus, H: 306 307 Hamiltonella, R: Rickettsia, W: Wolbachia). Thirteen to 17 individual measures were obtained per type of F1 female and per plant. 308

309 Oviposition choice according to the insect cytotype

We then investigated the influence of the whiteflies' cytotype on female plant choice 310 for oviposition between hibiscus and lantana (Fig. 2). Oviposition choice is expressed here as 311 the mean proportion of eggs laid on lantana compared to the total number of eggs laid on the 312 two plants (Fig. 2A). There was no effect of the experimental block factor [F(2,278) = 0.13, P]313 314 = 0.88] and no difference between genotypes for a given cytotype [F(6,296) = 0.82, P = 0.56]. 315 In contrast, there was a significant effect of the insect cytotype on oviposition plant choice [F(2,296) = 17.46, P < 0.001]: females with the cytotype Q1-HW laid a greater proportion of 316 317 eggs on hibiscus (around 80%) than on lantana (around 20%), while females with the cytotype Q1-HR or Q2-ARW showed no oviposition preference, *i.e.* the average proportion of eggs 318 laid on hibiscus and lantana was merely 50% (Fig. 2A). Most females (95%) laid eggs on a 319 single leaf disk: the one they were on at the end of the experiment. Only a few of them (5%) 320 321 laid eggs on both leaf disks (Fig. S2). During this short-term time period (32h), females that have chosen a single leaf disk laid a similar amount of eggs, about 5, whatever their choice 322 (hibiscus or lantana), and whatever their cytotype or genotype, with one exception: females 323

with the cytotype Q1-HW not only chose lantana less often than hibiscus but, when they did, they laid a lower number of eggs (mean, 3.55; SE, 0.52) than the ones who chose hibiscus (mean, 5.16; SE, 0.30) [effect of the interaction between the plant species and the insect cytotype: $\chi^2(2) = 11.72$, P = 0.0028; effect of the insect's nuclear genotype : $\chi^2(6) = 3.62$, P = 0.73] (**Fig. 2B**). A similar number of eggs, about 5, was laid by females that oviposited on both leaf disks (**Fig. S3**).

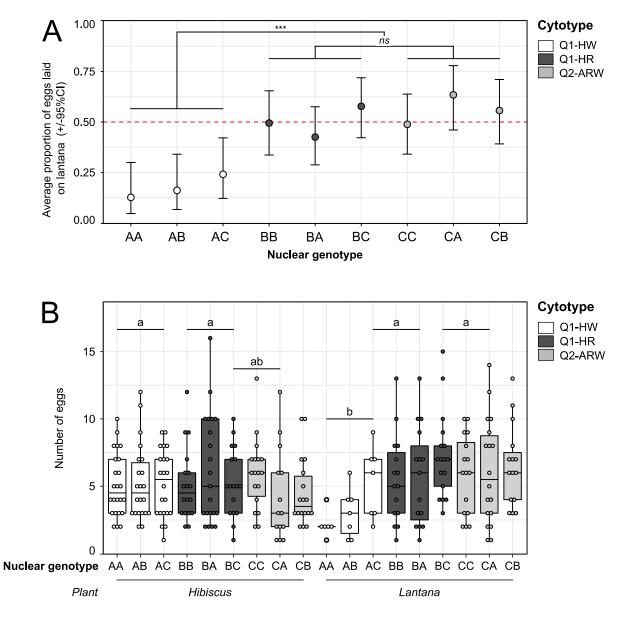


Figure 2. Effect of the insects' genotype and cytotype on *B. tabaci* females' oviposition
plant choice of hibiscus *vs* lantana. (A) Average proportion of eggs laid on lantana [± 95%
confidence interval (CI)] compared to the total number of eggs laid on the two plants over the

course of a 32h trial. The oviposition choice was investigated from 31 to 37 individuals per 333 cytotype/genotype combinations. Significance: p<0.001: ***, p>0.05: ns (Tukey's test). (B) 334 Number of eggs laid by females laying eggs exclusively on hibiscus or lantana (which 335 336 correspond to 95% of the females). Each dot represents one measure for one female; box length represents the interquartile range; whiskers indicate the lowest and largest data points, 337 excluding outliers. Black horizontal bars correspond to the median. To decipher the plant x 338 cytotype interaction, letters above the boxes indicate statistical groups determined by Tukey's 339 multiple comparison test (P < 0.05). Measures were performed on individual F1 females 340 obtained from crosses of laboratory lines: AA(Q1-HW), BB(Q1-HR), or CC(Q2-ARW). F1 341 females legend key: Nuclear genotype: letters indicate the maternally (first) and the paternally 342 (second) inherited nuclear genotype; Cytotype: mitochondrial group + S-symbionts (A: 343 Arsenophonus, H: Hamiltonella, R: Rickettsia, W: Wolbachia). 344

345 **Discussion**

In the present study, we demonstrate that whitefly cytotype is a major determinant of 346 347 both females' oviposition rate in no-choice experiments and oviposition behaviour in choice 348 experiments, while we did not find any influence of the insects' genotype on those traits. We acknowledge the possibility that these observations could result from genetic maternal effects. 349 such as the mother transmitting epigenetic states like DNA methylation or supplying eggs 350 351 with mRNA or proteins (English et al., 2015). The cytotype is composed of several components but differences due to potential genetic and/or gene expression variability of the 352 353 mitochondria or the P-symbiont between cytotypes seem unlikely due to (i) the extremely low mitochondrial DNA divergence between Q1 and Q2 populations (Vyskočilová et al., 2018), 354 and (ii) the similar genomic features and metabolic capabilities of *Portiera* across the *B*. 355 tabaci species complex (Santos-Garcia et al., 2020; Zhu et al., 2019, 2021). In contrast, 356

357 cytotypes differ in their S-symbiont composition, harbouring either *Hamiltonella* and
358 *Wolbachia* (Q1-HW), *Hamiltonella* and *Rickettsia* (Q1-HR) or *Arsenophonus*, *Rickettsia* and
359 *Wolbachia* (Q2-ARW). Therefore, it is likely that differences in traits and behaviours between
360 cytotypes result from this composition.

In this study, we also show that, in no-choice experiments, oviposition is lower on 361 lantana compared to hibiscus. Similarly to what we observed in a previous study, this could be 362 partially explained by the lower amount of the essential amino acids (EAAs) isoleucine, 363 leucine, phenylalanine and valine available in this plant compared to more favourable plants 364 (Benhamou et al., 2021). We also confirm that some cytotypes attenuate this effect: females 365 with cytotypes Q1-HR or Q2-ARW laid more eggs on lantana than females with cytotype Q1-366 HW, which could be due to the influence of the S-symbionts on their hosts' free amino acid 367 (AA) metabolism (Benhamou et al., 2021; Opatovsky et al., 2018; Rao et al., 2015). We 368 demonstrate here that this effect is dependent on the host plant species since oviposition is 369 homogeneous on hibiscus among cytotypes. 370

371 In choice experiments, females with cytotype Q1-HW oviposited more frequently on 372 hibiscus than on lantana, while other females did not show plant preference. Behavioural changes associated to symbionts may be attributable to the host's adaptive response to 373 symbiont infection or to the direct modification by the symbiont of the host behaviour. Here, 374 the choice looks adaptive: the oviposition preference of females with cytotype Q1-HW 375 towards hibiscus could be a response of the whitefly, driven by the lack of a cytotype 376 facilitating the utilization of lantana. Conversely, females with cytotypes Q1-HR and Q2-377 ARW, with no apparent plant preference, may be less selective considering their relatively 378 advantageous cytotype on lantana. Similar observations have been made in the olive fruit fly 379 B. oleae, in which experimental manipulation of the gut bacterial consortium resulted in 380 altered oviposition behaviour (Jose et al., 2019). Uninfected B. oleae showed less frequent 381

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oviposition on fruits, which may benefit the insect by avoiding oviposition on food sources that are difficult to use in the absence of the symbionts. It cannot be completely excluded that these observations are not an adaptive response of the host to S-symbiont infection. Indeed, the fact that half of the individuals of the cytotypes Q1-HR and Q2-ARW made non-optimal choices by choosing lantana could be due to behavioural manipulation (driven by the symbiont benefit), but the presumed benefit for the symbiont remains unknown.

388 The overall effect of the cytotype on whitefly oviposition behaviour may result as the net effect of the entire symbiotic community or may depend on the infection by a single S-389 symbiont species. Consistent with this second hypothesis, the avoidance/acceptance of lantana 390 391 for oviposition can be correlated to the absence/presence of the S-symbiont Rickettsia, as whiteflies with Rickettsia-associated cytotypes (Q1-HR and Q2-ARW) show the same 392 393 phenotype. Interestingly, *Rickettsia* has been associated to various phenotypic modifications 394 in B. tabaci, such as an increase of the fitness in some environments (Cass et al., 2016; Himler et al., 2011), heat tolerance (Brumin et al., 2011) or resistance to entomopathogenic 395 396 Pseudomonas syringae strains (Hendry et al., 2014) (for a review see also Milenovic et al., 397 2022). Moreover, Rickettsia was recently found to increase B. tabaci attraction towards tomato plants infected with the Tomato yellow leaf curl virus (Kliot et al., 2019), supporting 398 399 the hypothesis that this symbiont affects its host behaviour. Alternatively, it cannot be excluded that whiteflies do not have any preference but that the combination of *Hamiltonella* 400 and Wolbachia in females with cytotype Q1-HW decreases the preference for lantana. 401 However, less is known about the impact of this particular S-symbionts combination on their 402 hosts. 403

404 Mechanisms by which *Rickettsia* or any other S-symbiont may influence *B. tabaci* 405 oviposition behaviour are unknown to date. Studies on a broad spectrum of microorganisms 406 suggest that symbionts may influence their host behaviour directly, through the production of

a microbial signal (e.g. some EAAs) able to interact with their hosts' neurons or sensory 407 408 organs or that are precursors of neurotransmitters (Engl & Kaltenpoth, 2018; Shropshire & Bordenstein, 2016). Symbionts could also influence their host behaviour indirectly, through 409 410 their effects on host physiological functions (e.g. metabolism, immunity, endocrine system and pheromone production) that ultimately influence the host nervous system (Engl & 411 Kaltenpoth, 2018; Lynch & Hsiao, 2019; Shropshire & Bordenstein, 2016). In D. 412 melanogaster deprived of their gut bacteria, feeding on a diet depleted in EAAs elicits 413 preference for AA-rich food, while this behaviour can be rescued in flies inoculated with gut 414 bacteria (Leitão-Gonçalves et al., 2017). Therefore, it was proposed that flies are able to 415 detect changes in their inner free EAA concentrations, modulating their preference for AA-416 rich food. It was also proposed that D. melanogaster gut bacteria could provide their host with 417 EAAs, thereby modulating the nutritional status and subsequent food preference for this insect 418 419 (Leitão-Gonçalves et al., 2017). Similarly, S-symbionts may influence B. tabaci oviposition behaviour through their impact on their hosts' amino acid metabolism and nutritional status 420 421 (Benhamou et al., 2021; Opatovsky et al., 2018; Rao et al., 2015; Ren et al., 2020; Y.-B. 422 Wang et al., 2020). Regarding more specifically *Rickettsia*, genomic and experimental studies in B. tabaci MEAM1 (Middle-East Asia Minor 1) species suggest that this S-Symbiont could 423 424 complement the P-Symbiont and directly contribute to the biosynthesis of several EAAs, including lysine (Bao et al., 2021) and the branched-chain amino acids isoleucine, leucine and 425 valine (Opatovsky et al., 2018). According to this hypothesis, the apparent increase 426 427 production of EAAs previously reported in *Rickettsia*-associated cytotypes may explain why they show no preference between plants with high (i.e. hibiscus) and low (i.e. lantana) 428 amounts of these specific EAAs (Benhamou et al., 2021). Conversely, the lower level of 429 430 EAAs production in Q1-HW females may elicit preference for hibiscus. However, the

431 mechanisms by which insects could detect changes in internal free EAAs availability and432 subsequent changes in behaviour are not known.

In conclusion, this work shows that the host plant choice of *B. tabaci* for oviposition is 433 determined by the insect cytotype, most likely by the S-symbiont community. Whether these 434 effects depend on single or multiple S-symbiont species is unknown to date, and further 435 studies should investigate the respective contribution of each S-symbiont, including Rickettsia 436 as promising candidate, to host behaviour. This will require separation of S-symbionts from 437 other components of the cytotype, *i.e.*, selective elimination or transfer of symbionts to 438 whiteflies of identical genetic background. In this perspective, recent studies have made 439 encouraging advances in manipulating *B. tabaci* symbiotic communities by selectively 440 eliminating S-symbionts using heat (Shan et al., 2019) or antibiotic treatments (Ren et al., 441 2020; Shan et al., 2019; Y.-B. Wang et al., 2020). Mechanisms by which S-symbionts mediate 442 the insect behaviour also remain to be determined. Future investigations will help in 443 dissecting the molecular bases of whitefly behaviour to accurately determine the influence of 444 445 S-symbionts during each host plant selection step (e.g. olfaction or probing). Importantly, 446 acting on their hosts' niche choice, S-symbionts could contribute to build-up reproductive isolation between whitefly populations. However, in our framework, reproductive isolation is 447 not complete, as the choice of hibiscus is common to all cytotypes. Future work should 448 explore additional plant combinations to determine whether S-symbionts can lead to complete 449 reproductive isolation through differential host plant choice. Such directions would help to 450 better encompass the influence of symbionts on insects' ecological diversification and 451 speciation. 452

Declarations of interest

454 Declarations of interest: none

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466 **References**

- Akami, M., Awawing, A. A., Zhengzhong, C., Nan, J., Khaeso, K., Jurkevitch, E., Niu, C., & Yuval,
 B. (2019). Intestinal bacteria modulate the foraging behavior of the oriental fruit fly *Bactrocera dorsalis* (Diptera: Tephritidae). *PLoS* ONE, 14, e0210109.
 https://doi.org/10.1371/journal.pone.0210109
- Angelella, G., Nalam, V., Nachappa, P., White, J., & Kaplan, I. (2018). Endosymbionts differentially
 alter exploratory probing behavior of a nonpersistent plant virus vector. *Microbial Ecology*,
 76(2), 453–458. https://doi.org/10.1007/s00248-017-1133-5
- Bao, X., Yan, J., Yao, Y., Wang, Y., Id, P. V., Seal, S., & Id, J. L. (2021). Lysine provisioning by
 horizontally acquired genes promotes mutual dependence between whitefly and two intracellular
 symbionts. *PLoS Pathogens*, *17*(11), 1–23. https://doi.org/10.1371/journal.ppat.1010120
- Baumann, P. (2005). Biology of endosymbionts of plant sap-sucking insects. *Annual Review of Microbiology*, 59, 155–189. https://doi.org/10.1146/annurev.micro.59.030804.121041
- Benhamou, S., Rahioui, I., Henri, H., Charles, H., da Silva, P., Heddi, A., Vavre, F., Desouhant, E.,
 Calevro, F., & Mouton, L. (2021). Cytotype affects the capability of the whitefly *Bemisia tabaci*MED species to feed and oviposit on an unfavorable host plant. *MBio*, *12*(6), e00730-21.
 https://doi.org/10.1128/mBio .00730-21
- Bing, X., Yang, J., Zchori-fein, E., Wang, X., & Liu, S. (2013). Characterization of a newly
 discovered symbiont of the whitefly *Bemisia tabaci* (Hemiptera: Aleyrodidae). *Applied and Environmental Microbiology*, 79(2), 569–575. https://doi.org/10.1128/AEM.03030-12

- Bird, C. E., Fernandez-Silva, I., Skillings, D. J., & Toonen, R. J. (2012). Sympatric Speciation in the
 Post 'Modern Synthesis' Era of Evolutionary Biology. *Evolutionary Biology*, *39*(2), 158–180.
 https://doi.org/10.1007/s11692-012-9183-6
- Brucker, R. M., & Bordenstein, S. R. (2012). Speciation by symbiosis. *Trends in Ecology and Evolution*, 27(8), 443–451. https://doi.org/10.1016/j.tree.2012.03.011
- Brumin, M., Kontsedalov, S., & Ghanim, M. (2011). *Rickettsia* influences thermotolerance in the
 whitefly *Bemisia tabaci* B biotype. *Insect Science*, 18(1), 57–66. https://doi.org/10.1111/j.17447917.2010.01396.x
- Buchner, P. (1965). Endosymbioses of animals with plant microorganisms. In *John Wiley & Sons, Inc. New York.* John Wiley & Sons Inc.
- Caspi-Fluger, A., Inbar, M., Mozes-Daube, N., Katzir, N., Portnoy, V., Belausov, E., Hunter, M. S., &
 Zchori-Fein, E. (2012). Horizontal transmission of the insect symbiont *Rickettsia* is plantmediated. *Proceedings of the Royal Society B: Biological Sciences*, 279(1734), 1791–1796.
 https://doi.org/10.1098/rspb.2011.2095
- Caspi-Fluger, A., Inbar, M., Mozes-Daube, N., Mouton, L., Hunter, M. S., & Zchori-Fein, E. (2011).
 Rickettsia 'in' and 'out': Two different localization patterns of a bacterial symbiont in the same insect species. *PLoS ONE*, 6(6), 4–9. https://doi.org/10.1371/journal.pone.0021096
- Cass, B. N., Himler, A. G., Bondy, E. C., Bergen, J. E., Fung, S. K., Kelly, S. E., & Hunter, M. S.
 (2016). Conditional fitness benefits of the *Rickettsia* bacterial symbiont in an insect pest. *Oecologia*, 180(1), 169–179. https://doi.org/10.1007/s00442-015-3436-x
- 506 Crawley, M. J. (2013). The R book (Wiley-Blackwell, Ed.; 2nd ed.). John Wiley & Sons, Ltd.
- de Barro, P. J., Liu, S.-S., Boykin, L. M., & Dinsdale, A. B. (2011). *Bemisia tabaci*: a statement of
 species status. *Annual Review of Entomology*, 56(1), 1–19. https://doi.org/10.1146/annurev-ento 112408-085504
- 510 Dinsdale, A. B., Cook, L. G., Riginos, C., Buckley, Y. M., & de Barro, P. J. (2010). Refined global
 511 analysis of *Bemisia tabaci* (Hemiptera: Sternorrhyncha: Aleyrodoidea: Aleyrodidae)
 512 mitochondrial cytochrome oxidase 1 to identify species level genetic boundaries. *Annals of the*513 *Entomological Society of America*, 103(2), 196–208. https://doi.org/10.1603/AN09061
- 514 Dion, E., Polin, S. E., Simon, J., & Outreman, Y. (2011). Symbiont infection affects aphid defensive
 515 behaviours. *Biology Letters*, 7, 743–746. https://doi.org/10.1098/rsbl.2011.0249
- Douglas, A. E. (2009). The microbial dimension in insect nutritional ecology. *Functional Ecology*, 23(1), 38–47. https://doi.org/10.HH/j.1365-2435.2008.01442.x
- 518 Duron, O., Bouchon, D., Boutin, S., Bellamy, L., Zhou, L., Engelstädter, J., & Hurst, G. D. (2008).
 519 The diversity of reproductive parasites among arthropods: *Wolbachia* do not walk alone. *BMC*520 *Biology*, 6, 27. https://doi.org/10.1186/1741-7007-6-27
- English, S., Pen, I., Shea, N., & Uller, T. (2015). The information value of non-genetic inheritance in plants and animals. *PLoS ONE*, *10*(1), e0116996. https://doi.org/10.1371/journal.pone.0116996
- Engl, T., & Kaltenpoth, M. (2018). Influence of microbial symbionts on insect pheromones. *Natural Product Reports*, 35(5), 386–397. https://doi.org/10.1039/c7np00068e
- Everett, K. D. E., Thao, M., Horn, M., Dyszynski, G. E., & Baumann, P. (2005). Novel chlamydiae in
 whiteflies and scale insects: endosymbionts '*Candidatus* Fritschea bemisiae' strain Falk and

- 527 'Candidatus Fritschea eriococci ' strain Elm. International Journal of Systematic and
 528 Evolutionary Microbiology, 55, 1581–1587. https://doi.org/10.1099/ijs.0.63454-0
- Ferrari, J., & Vavre, F. (2011). Bacterial symbionts in insects or the story of communities affecting
 communities. *Philosophical Transactions of the Royal Society B: Biological Sciences*, *366*,
 1389–1400. https://doi.org/10.1098/rstb.2010.0226
- Firdaus, S., Vosman, B., Hidayati, N., Jaya Supena, E. D., Supena, J., Visser, R. G. F., & van Heusden, A. W. (2013). The *Bemisia tabaci* species complex: additions from different parts of the world. *Insect Science*, 20(0), 723–733. https://doi.org/10.1111/1744-7917.12001
- Frago, E., Dicke, M., & Godfray, H. C. J. (2012). Insect symbionts as hidden players in insect-plant
 interactions. *Trends in Ecology and Evolution*, 27(12), 705–711.
 https://doi.org/10.1016/j.tree.2012.08.013
- Gnankiné, O., Mouton, L., Henri, H., Terraz, G., Houndeté, T., Martin, T., Vavre, F., & Fleury, F.
 (2013). Distribution of *Bemisia tabaci* (Homoptera: Aleyrodidae) biotypes and their associated
 symbiotic bacteria on host plants in West Africa. *Insect Conservation and Diversity*, 6(3), 411–
 421. https://doi.org/10.1111/j.1752-4598.2012.00206.x
- Gottlieb, Y., Ghanim, M., Chiel, E., Gerling, D., Portnoy, V., Steinberg, S., Tzuri, G., Horowitz, A.
 R., Belausov, E., Mozes-Daube, N., Kontsedalov, S., Gershon, M., Gal, S., Katzir, N., & Zchorifein, E. (2006). Identification and localization of a *Rickettsia* sp. in *Bemisia tabaci* (Homoptera: Aleyrodidae). *Haematologica*, 72(5), 3646–3652. https://doi.org/10.1128/AEM.72.5.3646
- Gottlieb, Y., Ghanim, M., Gueguen, G., Kontsedalov, S., Vavre, F., Fleury, F., & Zchori-Fein, E.
 (2008). Inherited intracellular ecosystem: symbiotic bacteria share bacteriocytes in whiteflies. *The FASEB Journal*, 22(7), 2591–2599. https://doi.org/10.1096/fj.07-101162
- Gueguen, G., Vavre, F., Gnankine, O., Peterschmitt, M., Charif, D., Chiel, E., Gottlieb, Y., Ghanim,
 M., Zchori-Fein, E., & Fleury, F. (2010). Endosymbiont metacommunities, mtDNA diversity and
 the evolution of the *Bemisia tabaci* (Hemiptera: Aleyrodidae) species complex. *Molecular Ecology*, *19*, 4365–4378. https://doi.org/10.1111/j.1365-294X.2010.04775.x
- Hansen, A. K., & Moran, N. A. (2014). The impact of microbial symbionts on host plant utilization by
 herbivorous insects. *Molecular Ecology*, 23(6), 1473–1496. https://doi.org/10.1111/mec.12421
- Hart, B. L. (2011). Behavioural defences in animals against pathogens and parasites: parallels with the
 pillars of medicine in humans. *Philosophical Transactions of the Royal Society B: Biological Sciences*, *366*, 3406–3417. https://doi.org/10.1098/rstb.2011.0092
- Heil, M. (2016). Host manipulation by parasites: cases, patterns, and remaining doubts. *Frontiers in Ecology and Evolution*, *4*, 80. https://doi.org/10.3389/fevo.2016.00080
- Hendry, T. A., Hunter, M. S., & Baltrus, D. A. (2014). The facultative symbiont *Rickettsia* protects an
 invasive whitefly against entomopathogenic *Pseudomonas syringae* Strains. *Applied and Environmental Microbiology*, 80(23), 7161–7168. https://doi.org/10.1128/AEM.02447-14
- Henri, H., Terraz, G., Gnankiné, O., Fleury, F., & Mouton, L. (2013). Molecular characterization of
 genetic diversity within the Africa/Middle East/Asia Minor and Sub-Saharan African groups of
 the *Bemisia tabaci* species complex. *International Journal of Pest Management*, 59(4), 329–338.
 https://doi.org/10.1080/09670874.2013.869374
- Himler, A. G., Adachi-Hagimori, T., Bergen, J. E., Kozuch, A., Kelly, S. E., Tabashnik, B. E., Chiel,
 E., Duckworth, V. E., Dennehy, T. J., Zchori-Fein, E., & Hunter, M. S. (2011). Rapid spread of a

- bacterial symbiont in an invasive whitefly is driven by fitness benefits and female bias. *Science*, *332*, 254–256. https://doi.org/10.1126/science.1199410
- Hosokawa, T., & Fukatsu, T. (2020). Relevance of microbial symbiosis to insect behavior. *Current Opinion in Insect Science*, 39(March), 91–100. https://doi.org/10.1016/j.cois.2020.03.004
- Jose, A. P., Ben-yosef, M., Jurkevitch, E., & Yuval, B. (2019). Symbiotic bacteria affect oviposition
 behavior in the olive fruit fly *Bactrocera oleae*. *Journal of Insect Physiology*, *117*, 103917.
 https://doi.org/10.1016/j.jinsphys.2019.103917
- Kliot, A., Kontsedalov, S., Lebedev, G., Czosnek, H., & Ghanim, M. (2019). Combined infection with *Tomato yellow leaf curl virus* and *Rickettsia* influences fecundity, attraction to infected plants
 and expression of immunity-related genes in the whitefly *Bemisia tabaci. Journal of General Virology*, *100*(4), 721–731. https://doi.org/10.1099/jgv.0.001233
- Leitão-Gonçalves, R., Carvalho-santos, Z., Francisco, A. P., Tondolo Fioreze, G., Anjons, M., Elias,
 A. P., Itskov, P. M., Piper, M. D. W., & Ribeiro, C. (2017). Commensal bacteria and essential
 amino acids control food choice behavior and reproduction. *PLoS Biology*, *15*(4), e2000862.
 https://doi.org/10.1371/journal.pbio.2000862
- Lewis, Z., & Lize, A. (2015). Insect behaviour and the microbiome. *Current Opinion in Insect Science*, 9(3), 86–90. https://doi.org/10.1016/j.cois.2015.03.003
- Leybourne, D. J., Valentine, T. A., Bos, J. I. B., & Karley, A. J. (2020). A fitness cost resulting from *Hamiltonella defensa* infection is associated with altered probing and feeding behaviour in *Rhopalosiphum padi. Journal of Experimental Biology*, 223(1), jeb207936.
 https://doi.org/10.1242/jeb.207936
- Li, S., Ahmed, M. Z., Lv, N., Shi, P., Wang, X., Huang, J.-L., & Qiu, B.-L. (2016). Plant mediated
 horizontal transmission of *Wolbachia* between whiteflies. *The ISME Journal*, 11(4), 1–10.
 https://doi.org/10.1038/ismej.2016.164
- Li, T., Vinson, S. B., & Gerling, A. (1989). Courtship and mating behavior of *Bemisia tabaci*(Homoptera: Aleyrodidae). *Entomological Society of America*, 18(5), 800–806.
 https://doi.org/10.1093/ee/18.5.800
- Li, Y., Ahmed, M. Z., Li, S., Lv, N., Shi, P., Chen, X., & Qiu, B. (2017). Plant-mediated horizontal
 transmission of *Rickettsia* endosymbiont between different whitefly species. *FEMS Microbiology Ecology*, 93(6), 1–9. https://doi.org/10.1093/femsec/fix138
- Luan, J. B., Shan, H. W., Isermann, P., Huang, J. H., Lammerding, J., Liu, S. S., & Douglas, A. E.
 (2016). Cellular and molecular remodelling of a host cell for vertical transmission of bacterial symbionts. *Proceedings. Biological Sciences*, 283(1833). https://doi.org/10.1098/rspb.2016.0580
- Luan, J. B., Sun, X., Fei, Z., & Douglas, A. E. (2018). Maternal inheritance of a single somatic animal
 cell displayed by the bacteriocyte in the whitefly *Bemisia tabaci. Current Biology*, 28(3), 459–
 465. https://doi.org/10.1016/j.cub.2017.12.041
- Lynch, J. B., & Hsiao, E. Y. (2019). Microbiomes as sources of emergent host phenotypes. *Science*, 365(6460), 1405–1409. https://doi.org/10.1126/science.aay0240
- Malka, O., Santos-Garcia, D., Feldmesser, E., Sharon, E., Krause-Sakate, R., Delatte, H., van
 Brunschot, S., Patel, M., Visendi, P., Mugerwa, H., Seal, S., Colvin, J., & Morin, S. (2018).
 Species-complex diversification and host-plant associations in *Bemisia tabaci*: A plant-defence,
 detoxification perspective revealed by RNA-Seq analyses. *Molecular Ecology*, 27(21), 4241–
 4256. https://doi.org/10.1111/mec.14865

- Mccutcheon, J. P., & Moran, N. A. (2012). Extreme genome reduction in symbiotic bacteria. *Nature Reviews Microbiology*, *10*(1), 13–26. https://doi.org/10.1038/nrmicro2670
- Milenovic, M., Ghanim, M., Hoffmann, L., & Rapisarda, C. (2022). Whitefly endosymbionts: IPM opportunity or tilting at windmills? In *Journal of Pest Science* (Vol. 95, Issue 2, pp. 543–566).
 Springer Science and Business Media Deutschland GmbH. https://doi.org/10.1007/s10340-021-01451-7
- Moran, N. A., Degnan, P. H., Santos, S. R., Dunbar, H. E., & Ochman, H. (2005). The players in a
 mutualistic symbiosis: Insects, bacteria, viruses, and virulence genes. *Proceedings of the National Academy of Sciences of the United States of America*, 102(47), 16919–16926.
 https://doi.org/10.1073/pnas.0507029102
- Moran, N. A., McCutcheon, J. P., & Nakabachi, A. (2008). Genomics and evolution of heritable
 bacterial symbionts. *Annual Review of Genetics*, 42(1), 165–190.
 https://doi.org/10.1146/annurev.genet.41.110306.130119
- Mouton, L., Gnankiné, O., Henri, H., Terraz, G., Ketoh, G., Martin, T., Fleury, F., & Vavre, F. (2015).
 Detection of genetically isolated entities within the Mediterranean species of *Bemisia tabaci*: New insights into the systematics of this worldwide pest. *Pest Management Science*, 71(3), 452–458. https://doi.org/10.1002/ps.3834
- Najarro, M. A., Sumethasorn, M., Lamoureux, A., & Turner, T. L. (2015). Choosing mates based on
 the diet of your ancestors: Replication of non-genetic assortative mating in *Drosophila melanogaster. PeerJ*, 6(3), 1–11. https://doi.org/10.7717/peerj.1173
- Opatovsky, I., Santos-Garcia, D., Ruan, Z., Lahav, T., Ofaim, S., Mouton, L., Barbe, V., Jiang, J.,
 Zchori-Fein, E., & Freilich, S. (2018). Modeling trophic dependencies and exchanges among
 insects' bacterial symbionts in a host-simulated environment. *BMC Genomics*, *19*, 402.
 https://doi.org/10.1186/s12864-018-4786-7
- Polin, S., Simon, J., & Outreman, Y. (2014). An ecological cost associated with protective symbionts
 of aphids. *Ecology and Evolution*, 4(6), 836–840. https://doi.org/10.1002/ece3.991
- Poulin, R. (2010). Parasite manipulation of host behavior: an update and frequently asked questions. *Advances in the Study of Behavior*, 41(10), 151–186. https://doi.org/10.1016/S00653454(10)41005-0
- Qiao, H., Keesey, I. W., Hansson, B. S., & Knaden, M. (2019). Gut microbiota affects development
 and olfactory behavior in *Drosophila melanogaster*. *Journal of Experimental Biology*, 222(5),
 jeb192500. https://doi.org/10.1242/jeb.192500
- Rao, Q., Rollat-Farnier, P. A., Zhu, D., Santos-Garcia, D., Silva, F. J., Moya, A., Latorre, A., Klein, C.
 C., Vavre, F., Sagot, M., Liu, S., Mouton, L., & Wang, X. (2015). Genome reduction and
 potential metabolic complementation of the dual endosymbionts in the whitefly *Bemisia tabaci*. *BMC Genomics*, *16*(1), 226–238. https://doi.org/10.1186/s12864-015-1379-6
- Ren, F. R., Sun, X., Wang, T. Y., Yao, Y. L., Huang, Y. Z., Zhang, X., & Luan, J. B. (2020). Biotin provisioning by horizontally transferred genes from bacteria confers animal fitness benefits. *ISME Journal*, 14(10), 2542–2553. https://doi.org/10.1038/s41396-020-0704-5
- Romba, R., & Gnankiné, O. (2018). Comparative biology parameters of Q1 and Q3 genotypes of
 Bemisia tabaci MED (Hemiptera: Aleyrodidae) on two host plants in Burkina Faso, West Africa.
 African Entomology, 26(1), 116–123. https://doi.org/10.4001/003.026.0116

- Romba, R., Gnankine, O., Drabo, S. F., Tiendrebeogo, F., Henri, H., Mouton, L., & Vavre, F. (2018).
 Abundance of *Bemisia tabaci* Gennadius (Hemiptera: Aleyrodidae) and its parasitoids on vegetables and cassava plants in Burkina Faso (West Africa). *Ecology and Evolution*, 8(12), 6091–6103. https://doi.org/10.1002/ece3.4078
- Roopa, H. K., Asokan, R., Rebijith, K. B., Hande, R. H., Mahmood, R., & Kumar, N. K. K. (2015).
 Prevalence of a new genetic group, MEAM-K, of the whitefly *Bemisia tabaci* (Hemiptera: Aleyrodidae) in Karnataka, India, as evident from mtCOI sequences. *Florida Entomologist*, 98(4), 1062–1071. https://doi.org/10.1653/024.098.0409
- Russell, V. L. (2020). emmeans: Estimated Marginal Means, aka Least-Squares Means. R package
 version 1.4.8. https://cran.r-project.org/package=emmeans. https://cran.r project.org/package=emmeans
- Santos-garcia, D., Mestre-rincon, N., Ouvrard, D., Zchori-Fein, E., & Morin, S. (2020). *Portiera* gets
 wild: genome instability provides insights into the evolution of both whiteflies and their
 endosymbionts. *Genome Biology and Evolution*, 12(11), 2107–2124.
 https://doi.org/10.1093/gbe/evaa216
- Schluter, D. (2001). Ecology and the origin of species. *Trends in Ecology & Evolution*, 16(7), 372–
 380. https://doi.org/10.1016/s0169-5347(01)02198-x
- Schretter, C. E., Vielmetter, J., Bartos, I., Marka, Z., Marka, S., Argade, S., & Mazmanian, S. K.
 (2018). A gut microbial factor modulates locomotor behaviour in *Drosophila*. *Nature*,
 563(7731), 402–406. https://doi.org/10.1038/s41586-018-0634-9
- Shah, M. M. R., & Liu, T. X. (2013). Feeding experience of *Bemisia tabaci* (Hemiptera: Aleyrodidae)
 affects their performance on different host plants. *PLoS ONE*, 8(10), e77368.
 https://doi.org/10.1371/journal.pone.0077368
- Shan, H. W., Luan, J. B., Liu, Y. Q., Douglas, A. E., & Liu, S. S. (2019). The inherited bacterial
 symbiont *Hamiltonella* influences the sex ratio of an insect host. *Proceedings of the Royal Society B: Biological Sciences*, 286(1915). https://doi.org/10.1098/rspb.2019.1677
- Sharon, G., Segal, D., Ringo, J. M., Hefetz, A., Zilber-Rosenberg, I., & Rosenberg, E. (2010).
 Commensal bacteria play a role in mating preference of *Drosophila melanogaster*. *Proceedings*of the National Academy of Sciences of the United States of America, 107(46), 20051–20056.
 https://doi.org/10.1073/pnas.1009906107
- Shropshire, J. D., & Bordenstein, S. R. (2016). Speciation by symbiosis: The microbiome and
 behavior. *MBio*, 7(2), e01785-15. https://doi.org/10.1128/mBio.01785-15
- Sinisterra, X. H., McKenzie, C. L., Hunter, W. B., Powell, C. A., & Shatters, R. G. (2005).
 Differential transcriptional activity of plant-pathogenic begomoviruses in their whitefly vector
 (*Bemisia tabaci*, Gennadius: Hemiptera Aleyroadidae). Journal of General Virology, 86(5),
 1525–1532. https://doi.org/10.1099/vir.0.80665-0
- Sochard, C., Bellec, L., Simon, J.-C., & Outreman, Y. (2021). Influence of "protective" symbionts
 throughout the different steps of an aphid parasitoid interaction. *Current Zoology*, 67(4), 441–
 453. https://doi.org/10.1093/cz/zoaa053
- Sochard, C., Dupont, C., Simon, J.-C., & Outreman, Y. (2021). Secondary symbionts affect foraging
 capacities of plant-specialized genotypes of the pea aphid. *Microbial Ecology*, 82(4), 1009–1019.
 https://doi.org/10.1007/s00248-021-01726-7

- Sochard, C., le Floch, M., Anton, S., Outreman, Y., & Simon, J.-C. (2021). Limited influence of gain
 and loss of symbionts on host plant selection in specialized pea aphid genotypes. *Entomologia Generalis*, 41(1), 39–47. https://doi.org/10.1127/entomologia/2020/1076
- Thao, M. L., & Baumann, P. (2004). Evolutionary relationships of primary prokaryotic endosymbionts
 of whiteflies and their hosts. *Applied and Environmental Microbiology*, 70(6), 3401–3406.
 https://doi.org/10.1128/AEM.70.6.3401-3406.2004
- Tipton, L., Darcy, J. L., & Hynson, N. A. (2019). A developing symbiosis: enabling cross-talk
 between ecologists and microbiome scientists. *Frontiers in Microbiology*, 10, 292.
 https://doi.org/10.3389/fmicb.2019.00292
- Tukey, J. W. (1949). Comparing individual means in the analysis of variance. *International Biometric Society*, 5(2), 99–114. https://doi.org/10.2307/3001913
- Vale, P. F., Siva-Jothy, J. A., Morrill, A., & Forbes, M. R. (2018). The influence of parasites on insect
 behavior. In A. Córdoba-Aguilar, D. González-Tokman, & I. González-Santoyo (Eds.), *Insect behavior: from mechanisms to ecological and evolutionary consequences* (1st ed., Issue May, pp.
 274–291). Oxford University Press. https://doi.org/10.1093/oso/9780198797500.003.0018
- Vavre, F., Fleury, F., Lepetit, D., Fouillet, P., & Boulétreau, M. (1999). Phylogenetic evidence for
 horizontal transmission of *Wolbachia* in host-parasitoid associations. *Molecular Biology and Evolution*, 16(12), 1711–1723. https://doi.org/10.1093/oxfordjournals.molbev.a026084
- Vavre, F., & Kremer, N. (2014). Microbial impacts on insect evolutionary diversification: from
 patterns to mechanisms. *Current Opinion in Insect Science*, 4(1), 29–34.
 https://doi.org/10.1016/j.cois.2014.08.003
- Vyskočilová, S., Seal, S., & Colvin, J. (2019). Relative polyphagy of "Mediterranean" cryptic *Bemisia tabaci* whitefly species and global pest status implications. *Journal of Pest Science*, 92(3), 1071–
 1088. https://doi.org/10.1007/s10340-019-01113-9
- Vyskočilová, S., Tay, W. T., van Brunschot, S., Seal, S., & Colvin, J. (2018). An integrative approach
 to discovering cryptic species within the *Bemisia tabaci* whitefly species complex. *Scientific Reports*, 8(10886), 1–13. https://doi.org/10.1038/s41598-018-29305-w
- Wang, H., Lei, T., Wang, X., Maruthi, M. N., Zhu, D., Cameron, S. L., Rao, Q., Shan, H., Colvin, J.,
 Liu, Y. Q., & Liu, S. (2020). A newly recorded *Rickettsia* of the Torix group is a recent intruder
 and an endosymbiont in the whitefly *Bemisia tabaci. Environmental Microbiology*, 22(4), 1207–
 1221. https://doi.org/10.1111/1462-2920.14927
- Wang, Y.-B., Ren, F.-R., Yao, Y.-L., Sun, X., Walling, L. L., Li, N.-N., Bai, B., Bao, X.-Y., Xu, X.R., & Luan, J.-B. (2020). Intracellular symbionts drive sex ratio in the whitefly by facilitating
 fertilization and provisioning of B vitamins. *ISME Journal*, 14(12), 2923–2935.
 https://doi.org/10.1038/s41396-020-0717-0
- Wong, A. C., Wang, Q.-P., Morimoto, J., Senior, A. M., Lihoreau, M., Neely, G. G., Simpson, S. J., &
 Ponton, F. (2017). Gut microbiota modifies olfactory-guided microbial preferences and foraging
 decisions in *Drosophila*. *Current Biology*, 27(15), 2397–2404.
 https://doi.org/10.1016/j.cub.2017.07.022
- Zchori-fein, E., & Brown, J. K. (2002). Diversity of prokaryotes associated with *Bemisia tabaci*(Gennadius) (Hemiptera: Aleyrodidae). *Ecology and Population Biology*, 95(6), 711–718.
 https://doi.org/10.1603/0013-8746

- Zchori-Fein, E., Lahav, T., & Freilich, S. (2014). Variations in the identity and complexity of
 endosymbiont combinations in whitefly hosts. *Frontiers in Microbiology*, 5(310), 310.
 https://doi.org/10.3389/fmicb.2014.00310
- Zhu, D., Rao, Q., Zou, C., Ban, F.-X., Zhao, J.-J., & Liu, S. (2021). Genomic and transcriptomic analyses reveal metabolic complementarity between whiteflies and their symbionts. *Insect Science*. https://doi.org/10.1111/1744-7917.12943
- Zhu, D., Zou, C., Ban, F., Wang, H., Wang, X., & Liu, Y.-Q. (2019). Conservation of transcriptional
 elements in the obligate symbiont of the whitefly *Bemisia tabaci. PeerJ*, 7(9), e7477.
 https://doi.org/10.7717/peerj.7477

747

748 Supplementary material

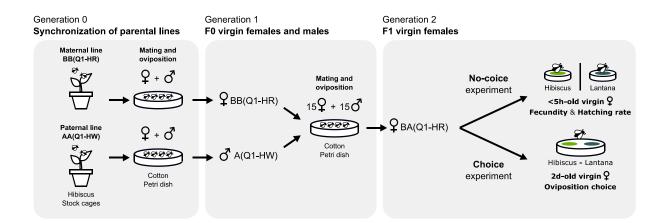


Figure S1. Experimental procedure used for no-choice and choice experiments on F1 749 750 females, using BA(Q1-HR) F1 females as an example. All experiments were performed on 751 individual F1 females. Oviposition choice was assessed by counting the number of eggs laid per female after 32 h on each plant (n=31-27). Fecundity (number of eggs laid) was assessed 752 over 7 days per F1 female after which the female was removed from the Petri dish. The 753 hatching rate was calculated as the ratio of larvae (observed 10 days after removing the 754 female), over the total number of eggs laid per F1 female (see formula (a) from main text) 755 756 (n=13-17).

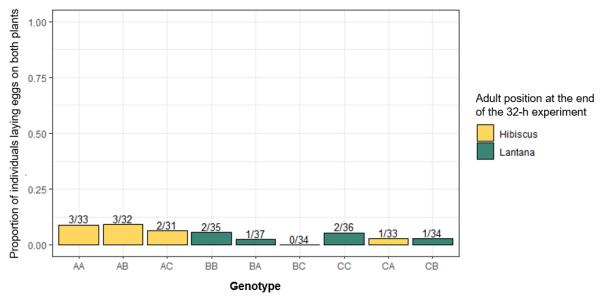


Figure S2. Proportion of *B. tabaci* females laying eggs on both hibiscus and lantana
according to their genotype in choice experiments. Most of females laid eggs on a single
plant species. Colors indicate the female's position at the end of the experiment. Numbers
above bars indicate absolute values (number of females laying eggs on both plants / total
number of females analyzed).

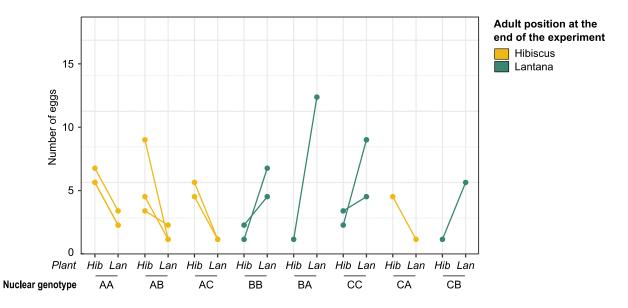


Figure S3. Number of eggs laid by whiteflies ovipositing on both plants in hibiscus *vs*lantana plant choice for oviposition experiments, according to their nuclear genotype.
There was no data for genotype BC, as no BC(Q1-HR) female oviposited on both plants. Each
dot pair corresponds to an individual females' oviposition on hibiscus (Hib) and lantana (Lan).
Colors indicate the female's position at the end of the experiment.