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- 7
- 8 Key words

9 Invasive species, Inbreeding, Cuticular hydrocarbons, Queen pheromone, Yellow-legged10 hornet.

11

12 Abstract

13 Invasive social insect populations that have been introduced to a new environment through a limited number of introduction events generally exhibit reduced variability in their chemical 14 15 signatures (cuticular hydrocarbons) compared to native populations of the same species. The reduced variability in these major recognition cues could be caused by a reduction of genetic 16 diversity due to a genetic bottleneck. This hypothesis was tested in an inbred European 17 population of the invasive hornet Vespa velutina nigrithorax. Our results show that, in spite of 18 the limited amount of genetic diversity present in the European population, the chemical 19 signatures of individuals were highly heterogeneous according to their caste, sex, and colony 20 21 origin. In queens, some specific saturated and unsaturated hydrocarbons were identified. These results suggest that epigenetic and/or environmental factors could play a role in 22 23 modifying cuticular hydrocarbon profiles in this introduced hornet population despite the observed reduction of genetic diversity. 24

25

26 Introduction

The insect cuticle is covered by a thin lipid layer, which largely consists of cuticular 27 28 hydrocarbons (CHCs). Because of their chemical properties, CHCs primarily act to limit insect water loss through the creation of a surface barrier. In addition, they also serve as 29 30 contact pheromones, i.e. recognition cues, for inter- and intraspecific recognition among insects. The CHC profile of an individual serves as a personalized chemical signature 31 32 (Blomquist and Bagnères 2010; Ginzel and Blomquist 2016). Several studies have 33 demonstrated a strong link between the social behavior of insects and their cuticular 34 signatures (Bagnères et al. 1991; Howard and Blomquist 2005; Leconte and Hefetz 2008).

35 CHCs are mainly involved in the recognition of nestmates (Howard and Blomquist 2005; van Zweden and D'Ettorre 2010), castes (Bagnères et al. 1998; Kaib et al. 2000; Smith and Taylor 36 1990), and worker's tasks (Rahman et al 2016). Therefore, chemical signatures can vary, and 37 the degree of variation depends on a wide range of endogenous factors, such as the endocrine 38 system or enzymes (Darrouzet et al. 2014b; Fan et al. 2004; Lengyel et al. 2007), and 39 exogenous factors, such as climate and season (Bagnères et al. 1990), diet (Liang and 40 Silverman 2000), parasitism (Bagnères et al. 1996; Lebreton et al. 2010; LeConte et al. 2015), 41 or time (Bagnères et al. 2011; Lebreton et al. 2009). The relative importance of these factors 42 43 varies among species (VanderMeer and Morel 1998). However, CHC profiles are modified 44 not only by the social or biotic environment but are also under genetic control in both social 45 and solitary insects (Bonavita-Cougourdan et al. 1996; Dronnet et al. 2006; Etges et al. 2009; Thomas and Simmons 2008; Vauchot et al. 1996). Consequently, variation in CHC profiles 46 47 has been used to identify species (Bagnères and Wicker-Thomas 2010) and to differentiate populations (Bonelli et al. 2015; Nowbahari et al. 1990). This type of chemotaxonomy, i.e. 48 49 the use of chemical compounds, has been productively used in numerous insect species. However, less attention has been paid to the extent or function of chemical variation within 50 51 invasive species, which in their introduced ranges often experience habitats and species interactions that are strikingly different from those of their native environments (Perdereau et 52 al. 2010). 53

Native to China, the invasive yellow-legged hornet, Vespa velutina nigrithorax, was 54 accidentally introduced to southwestern France around 2004, probably in imported ceramic 55 pottery (Haxaire et al. 2006). The species has successfully expanded its range to cover more 56 than 75% of France, and is currently colonizing neighboring countries (Spain, Portugal, Italy, 57 Belgium, Germany, and England) (Darrouzet et al. 2014a; Goldarazena et al. 2015; Monceau 58 et al. 2013; Robinet et al. 2017). Thus far, V. velutina invasion in France has two main 59 negative impacts. First, the species preys on several insect and arthropod taxa, thus potentially 60 affecting local biodiversity. In particular, V. velutina is a predator of the domestic honeybee, 61 62 Apis mellifera, and could induce economic losses via the destruction of bee colonies (Darrouzet et al. 2014a; Monceau et al. 2013). Second, the species presents a risk to human 63 64 health. Accidents have occurred, some fatal, when people have inadvertently approached the 65 hornet's nests (de Haro et al. 2010).

New colonies of *V. velutina* are established in the spring by mated queens, after the overwintering period. Colonies first pass through a period in which an increasingly large number of workers are produced in order to ensure colony growth, and then produce sexual

individuals (males and gynes) at the end of the summer (Monceau et al. 2013; Rome et al. 69 70 2015). These reproductive individuals emerge between late August and November. After reproductive flights, the gynes disperse and the males die. However, Darrouzet et al. (2015) 71 72 showed that several colonies produced diploid males all throughout the colony life. As authors explained, this result strongly suggests the occurrence of inbreeding in this species in 73 its invaded range (Europe). This hypothesis was supported by genetic analyses based on a 74 combination of mitochondrial and nuclear markers (Arca et al. 2015; Takeuchi et al. 2017). 75 The authors suggested that the yellow-legged hornet population in France was derived from a 76 77 single introduction event, and they showed evidence of a strong founder effect. Thus the French populations sampled in that study suffered a notable loss of genetic diversity 78 79 compared to native populations in China.

In the present study, we analyzed the chemical signature of the invasive hornet *V*. *velutina nigrithorax* in France, i.e. in an inbreeding context, to test whether this reduced genetic diversity translates into homogeneity in the insects' chemical signatures, as has been reported for other invasive social insect species (*Linepithema humile:* Brandt et al. 2009; *Wasmannia auropunctata:* Errard et al. 2005; *Reticulitermes flavipes*: Perdereau et al. 2010).

85

86 Materials and Methods

87 Insect collection

Hornet colonies were collected in 16 different locations in the Centre-Val-de-Loire region in
France (Table 1). In each colony, individuals from each caste and sex present (workers,
gynes, males, and queens when possible) were randomly selected, with a maximum of 10 per
caste. Insects were weighed and stored at -20°C before being further analyzed.

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93 *Cuticular hydrocarbon (CHC) analyses*

CHCs were collected by plunging each hornet in 1 ml pentane for 2 x 1 min (vortex). A 500-94 µl aliquot of pentane extract was transferred to a new tube. Immediately prior to analyses, 10 95 μ L of *n*-C20 (*n*-eicosane; concentration 10⁻³ g/ml) was added to each tube as an internal 96 standard. Then, 2-µL samples were analyzed using gas chromatography (GC) in a CPG 97 Agilent Technologies 7820A GC System equipped with a flame-ionization detector (FID) and 98 a capillary column (HP-5 Agilent Technology, Santa Clara, USA; 30 m x 0.32 mm x 0.25 99 μm) using helium as a carrier gas (1.7 ml/min). The following temperature program was used: 100 a first temperature ramp from 50 to 200°C at 8°C/min, a second temperature ramp from 200 101 to 315°C at 5°C/min, and a 5-min final hold at 315°C. 102

To identify the compounds (CHCs) in the chemical signatures according to caste and 103 104 sex (workers, gynes, and males), CHC extracts were pooled from nine random hornets, each from a different randomly selected colony (Table 1S). Workers and gynes were discriminated 105 according to their size and weight (Rome et al. 2015). CHCs from five random queens were 106 107 also extracted and pooled. The samples were dried, then stored at -20°C. Before being analyzed, 300 µl of pentane was added to each, with the internal standard as in the previous 108 analysis. To determine which compounds were present, we used an Agilent Technologies 109 7000C GC/MS Triple Quad with a 7890B GC System equipped with the same column as 110 111 above; we also used the same temperature program as above. Compounds were identified 112 following Blomquist (2010) and using standard *n*-alkanes.

To characterize the CHC profiles, the areas of the main peaks were determined using ChemStation software (Agilent Technologies), and the relative proportions of each peak were calculated as described in Bagnères et al. (1990).

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117 Calculations and statistical analyses

To determine whether we could use chemical (CHC) profiles to discriminate among colonies and/or among individuals within a colony on the basis of caste and sex, we performed multivariate principal component analyses (PCAs) using the R 2.14.1 software.

Differences in CHC composition among and within colonies were quantified using the Nei distance (Dronnet et al., 2006; Nowbahari et al 1990). The dissimilarity matrix, which included all possible types of pairs (intra-caste, inter-caste within a colony, and inter-colony within a caste), was constructed from the previously calculated mean relative areas of cuticular hydrocarbon peaks and were analyzed with a Kruskal-Wallis test using R 2.14.1.

For each individual, the quantity of each CHC was estimated with the following formula: $Qi = (Q_{C20} X xi) / (x_{C20}i)$. The quantity of each CHC was then divided by the mass of the individual to obtain the per-gram quantity of each compound.

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130 **Results**

131 Analyses of CHCs

Out of the peaks detected by GC, only the 75 most stable peaks (of which four were queenspecific) in individuals in each group were retained and used in our statistical analyses. Among the identified CHCs, which ranged in chain length from 21 to 35 carbon atoms, we found *n*-alkanes (13.48%, 12 compounds), mono- and dimethyl-alkanes (57.30%, 51 compounds), alkenes (17.98%, 16 compounds), and unknown compounds (11.23%, 10

- compounds). Queens possessed four specific alkenes (C27:1, C31:1, C33:1, and C35:1) which
 were never detected in other castes (Table 2, Figure 1).
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140 *Caste- and colony-related CHC differences*

In all colonies analyzed, PCAs showed that workers, males, gynes, and queens displayed
different CHC profiles, with queens always clearly differentiated on axis 2 (Figure 2, Table
3). A comparison of chemical signature among colonies also showed significant differences in
PCAs, with queens more or less following their offspring but always well differentiated from
other castes (Figure 3).

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147 *Chemical distances in CHC profiles*

With regard to the chemical distances associated with CHC profiles (Figure 4), we found 148 149 more difference between gynes and other castes within a given colony than among gynes from different colonies (Kruskal-Wallis, P<0.001). Instead, the chemical signatures of males 150 151 were not significantly different from those of other castes in the same colony or among different colonies (P>0.05; Figure 4). For workers, among-colony differences were larger than 152 153 the chemical distances within colonies and among castes (P<0.001; Figure 4). For gynes, males, and workers, intra-caste chemical distances (i.e. among individuals in the same caste, 154 whatever the colony) were always shorter than those among castes in the same colony or 155 between colonies (all P<0.001). Moreover, a comparison of intra-caste chemical distances 156 revealed that these distances were largest among workers (workers vs males P<0.001, workers 157 vs gynes P<0.01), and greater in gynes than in males (P<0.05). Among colonies, the chemical 158 distances among workers were larger than those among gynes and males (P<0.001 in both 159 160 cases), and were similar between gynes and males (P>0.05).

Figure 5 shows chemical distances between castes. The shortest distances were between
comparisons of males and gynes and males and workers. Comparisons with queens always
produced the highest distances.

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165 *Quantitative analysis of CHC profiles*

Quantitatively, queens possessed more CHCs per gram of body weight than did the other individuals within a colony (Figure 6) (Kruskal-Wallis, P<0.001). Gynes and males were similar to each other in terms of quantities of CHCs (P>0.05), but each had less than workers (P<0.001). The chromatograms of queens were characterized by the presence of two main

alkanes, *n*-C25 and *n*-C27, in greater quantities than in other castes (Figure 1) [(n-C25: 7.7x

more than in workers (Kruskal-Wallis P<0.001), 11.5x more than in gynes (P<0.001) and 10.9x more than in males (P<0.001); *n*-C27: 3.9x more than in workers (P<0.001), 10.1x more than in gynes (P<0.001), and 7.7x more than in males (P<0.001)], plus the queenspecific alkenes described above in the paragraph "*Analyses of CHCs*".

Figure 7 depicts differences between castes and genders in the chemical makeup of 175 176 CHCs. With respect to the amounts present in gynes, males, and workers, queens possessed higher quantities of *n*-alkanes (P<0.001 for each comparison), monomethyl-alkanes 177 (P<0.001), and alkenes (P<0.001), but the same quantities of dimethyl-alkanes (P>0.05). 178 179 Workers possessed more *n*-alkanes than gynes and males (P<0.05 with males; P<0.001 with gynes), more monomethyl-alkanes than gynes (P<0.05) but a similar quantity as males 180 181 (P>0.05), and more dimethyl-alkanes than males and gynes (P<0.001 for each comparison). Gynes possessed less *n*-alkanes than males (P<0.001), but the same quantities of 182 183 monomethyl-alkanes, dimethyl-alkanes, and alkenes (P>0.05 respectively).

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185 Discussion

Our results demonstrated that, in spite of inbreeding (Darrouzet et al. 2015) linked to a genetic bottleneck (Arca et al. 2015), the chemical signature of the invasive hornet *Vespa velutina* in France is clearly heterogeneous. Each colony, caste, and gender possesses a specific CHC profile. This study thus suggests that CHCs could serve as a useful marker for discriminating yellow-legged hornet populations in France, the main invaded area of this species in Europe.

The chemical heterogeneity that we observed within the introduced hornet population 191 192 in France was unexpected. Arca et al. (2015) demonstrated that native V. velutina populations near Shanghai (China) contained at least three different haplotypes. Little is currently known 193 about the chemical signatures of yellow-legged hornets in their native area, but it is possible, 194 195 indeed probable, that these populations have strongly heterogenenous CHC profiles. Instead, 196 the introduced population in France has just one haplotype (Arca et al. 2015), but has notable chemical heterogeneity in CHC profiles. This result is surprising, because to our knowledge 197 198 this type of hydrocarbon heterogeneity has not previously been detected in other introduced social insects. On the contrary, recent research has revealed homogeneity in CHC profiles in 199 200 introduced ants (Linepithema humile: Brandt et al. 2009; Wasmannia auropunctata: Errard et 201 al. 2005) and termites (*Reticulitermes flavipes*: Perdereau et al. 2010). These previous studies 202 suggested that the reduced chemical variability of introduced populations may arise from the limited number of introduction events into a new environment. Specifically, the reduced 203 204 variability of recognition cues, such as CHCs, in these introduced populations of ants and

termites has been attributed to a reduction of genetic diversity through a genetic bottleneck
(Tsutsui et al. 2000) or to 'genetic cleansing', i.e. selection against the less-common
recognition alleles (Giraud et al. 2002).

208 In the case of the invasive hornet V. velutina in France, which is an inbred population with a genetic bottleneck, the heterogeneity in CHC profiles may be affected by 209 environmental and/or epigenetic parameters. For example, diet or laboratory rearing 210 conditions are known to alter the chemical signature in termites (Shelton and Grace 1997; 211 Polizzi and Forschler 1998; Florane et al. 2004). A change in ants' chemical signature could 212 213 occurred over a short period of time, which is due to the ants walking over their prey (Liang and Silverman 2000). Because V. velutina is a generalist predator and can consume several 214 215 different types of prey, it is possible that a particular diet specific to the area around a given colony could explain the CHC heterogeneity we observed. However, as hornets have a 216 217 general diet, it is unlikely that any great overall differences could appear in such conditions. But this idea needs furthers studies to be verified. Seasonal variations could also alter V. 218 219 *velutina*'s chemical signatures, as has been observed in termite species (Bagnères et al. 1990); however, this cannot explain the differences reported here since we collected all hornet 220 221 colonies in the same period of time. Another possible contributing factor could be the 222 composition of the endosymbiont community in the digestive tract, as this has been reported to affect host chemical communication in termites (Matsuura 2001, 2003). Since these 223 symbionts are probably obtained by new individuals thanks to trophallaxis with adult workers, 224 it is possible that each V. velutina colony could possess a unique endosymbiont profile with 225 specific effects on the CHC profiles of their host. Further studies are clearly necessary to 226 analyze all these factors and their relative contributions to the hornet CHC profile. 227

These specific chemical signatures probably enable each individual to identify the 228 229 colony and caste origins of conspecifics. In social insect colonies, labor is divided between a reproductive queen, reproductive individuals (gynes and males), and sterile workers (Wilson 230 1971). The distribution of tasks among individuals requires interactions among colony 231 232 members which involve the recognition of caste membership and perhaps specific tasks inside the caste (more particularly in the worker caste) (Green and Gordon 2003). This process is 233 234 generally based on chemical cues such as CHCs. For example, in bees and termites, castes can be distinguished by differences in CHC profiles (honeybees: Kather et al. 2011; termites: 235 Bagnères et al. 1998, 2011; Darrouzet et al. 2014b; Haverty et al. 1996; Kaib et al. 2002; 236 LeConte and Hefetz 2008; Nunes et al. 2009). Such differences were also observed across 237 238 different age groups in Apis cerana workers and could serve as the basis for task allocation in

that species (Rahman et al. 2015). Wagner et al. (1998) supposed that these caste-related 239 differences in CHCs may be representative of the tasks performed by individuals, that is, the 240 type and quantity of CHCs present could be linked to the tasks and specific environment of a 241 given individual. For example, bee foragers are exposed more often than other workers to 242 unfavorable environmental conditions, such as hot temperatures, high humidity, and harmful 243 chemicals. Consequently, Rahman et al. (2015) suggested that the increased levels of n-244 alkanes observed in A. cerana foragers, compared with those of workers remaining inside the 245 hive, could play an important role in regulating water impermeability, as had been previously 246 247 proposed (Gibbs and Pomonis 1995). In the present study, we observed similar differences 248 between V. velutina workers and reproductive individuals collected in nests (gynes and 249 males). Such differences could be explained by the environment of individuals: most workers 250 must contend with variable environmental conditions during their foraging activities while the 251 reproductive individuals we analyzed (i.e. collected in their nests) stayed in the nest until their nuptial flight. However, eventually these gynes and males have to leave the nests and, like 252 253 workers, will be confronted with variable environmental conditions. It is possible that their chemical signature adapts prior to leaving the nest by increasing the amount of *n*-alkanes 254 255 present, though further studies are necessary to verify this hypothesis.

Interestingly, the CHC profiles of V. velutina queens possess two alkanes in great 256 quantities (n-C25 and n-C27) which seem to act as queen pheromones in other hymenopteran 257 species, such as Lasius ants (Holman et al 2010, 2013), the bumble bee Bombus terrestris 258 (Holman 2014), and the wasp Vespula vulgaris (Van Oystaeyen et al. 2014). These n-259 alkanes, which are present on queen cuticle, could induce sterility in workers. This 260 reproductive inhibition of the sterile caste seems to occur via two distinct processes: first, by 261 inhibiting the functioning of workers' ovaries, and second, by causing oocyte resorption (Van 262 263 Oystaeyen et al. 2014). These saturated linear alkanes seem to be the most conserved class of queen pheromones in several hymenopterans (Van Oystaeyen et al. 2014), and could be used 264 as a fertility signal in a common solitary ancestor of social hymenopteran species (Wilson et 265 266 al. 2013). Our result seems consistent with this idea and adds a second Vespa species to the group of species, spanning major clades of social hymenoptera, which has yielded clues about 267 268 the evolutionary history of queen and fertility signals. However, Amsalem et al (2015) tested three alkanes (n-C23, n-C25 and n-C27) in Bombus impatiens and found no evidence to 269 270 support this theory that a conserved class of hydrocarbons serve as queen pheromones or queen signals. Moreover, we observed four unsaturated hydrocarbons (C27:1, C31:1, C33:1, 271 272 and C35:1) specific to the chemical profile of V. velutina queens. As observed by Van

Oystaeyen et al. (2014), some other compounds, including unsaturated hydrocarbons, can also 273 be used as queen pheromones. In the Polistes, Apis, Bombus, and Lasius species in which they 274 275 have been studied, these alkenes derive from saturated hydrocarbons, and can be obtained 276 after biochemical modification during pheromone biosynthesis, from the up-regulation of 277 specific desaturases (Blomquist & Vogt 2003; Blomquist and Bagnères 2010; Qiu et al. 2012). Some alkene compounds are used as male attractants (sex pheromones), particularly in 278 solitary species (Blomquist and Vogt 2003; Blomquist and Bagnères 2010). As hypothesized 279 by Van Oystaeyen et al. (2014), different queen pheromone compounds have arisen as a result 280 of the multiple independent origins of eusociality, particularly by evolving from a fertility 281 signal directed to attract males in solitary ancestors. Further studies are necessary to test the 282 283 function of these alkenes in V. velutina nigrithorax.

284

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471	Legends
472	
473	Figure 1: Typical chromatograms of queens, gynes, males, and workers.
474	
475	Figure 2: Plot of the first two axes of the Principal Components Analysis (PCA) of the
476	proportion of CHCs in the different castes within V. velutina colonies.
477	
478	Figure 3: Plot of the first two axes of the Principal Components Analysis (PCA) of the
479	proportion of CHCs in the different castes (queens, males, workers, and gynes) among
480	different colonies.
481	
482	Figure 4: Calculation of chemical distances based on comparisons of CHC profiles of
483	workers, gynes, and males. "Intra-caste" represents the distance between individuals in the
484	same caste and in the same colony, "inter-caste" is between the analyzed caste and the others
485	castes in the same colony, and "inter-colony" is for the same caste but among colonies.
486	
487	Figure 5: Comparisons of inter-caste chemical distances. The different letters for each
488	distance were obtained with P<0.01 (Kruskal-Wallis test). G=gynes, m=males, q=queens,
489	w=workers.
490	
491	Figure 6: Quantitative analysis of total CHC profiles (gram of CHCs per gram of insect) in
492	workers, queens, gynes, and males.
493	
494	Figure 7: Quantitative analysis of alkanes, alkenes, monomethyl-alkanes, and dimethyl-
495	alkanes in CHC profiles (gram of CHCs per gram of insect) from workers, queens, gynes, and
496	males.

Table 1 Origins and dates of sample collection and sample sizes.

		Dates of				
		sample	Number of			
Colonies' origines	Lambert 93 coordinates	collections	Males	Workers	Gynes	Queens
Chinon	X:491725 Y:6677696	07/10/2011	10	0	0	0
Sepmes	X : 522947 Y : 6665711	17/10/2011	10	10	10	0
Barrou	X : 530076 Y : 6643026	24/10/2011	8	4	10	0
Ligueil (A)	X : 534307 Y : 6662641	25/10/2011	5	10	10	1
Azay sur Indre	X : 544621 Y : 6680754	21/11/2011	10	0	0	0
Ligueil (B)	X : 534307 Y : 6662641	26/11/2011	10	10	4	0
Civray de touraine	X : 552648 Y : 6694251	01/12/2011	10	10	10	0
Reignac sur Indre	X : 542394 Y : 6683401	06/12/2011	0	10	0	0
Lussault sur Loire	X : 542876 Y : 6701668	16/10/2012	10	10	0	0
Montlouis sur Loire	X : 536567 Y : 6700904	22/10/2012	10	10	10	1
Lerné	X : 482765 Y : 6674503	23/10/2012	10	10	10	0
Artannes	X : 518507 Y : 6688647	08/11/2012	10	10	10	0
Athée sur Cher	X : 542527 Y : 6693243	09/11/2012	10	10	0	0
St Avertin	X : 528458 Y : 6698865	27/08/2013	10	10	0	1
Tours	X : 525709 Y : 6701535	16/09/2013	10	10	0	1
Ligueil (C)	X : 534307 Y : 6662641	11/10/2013	10	10	0	1



Table 1S Origins and dates of sample collection for insect pools to identify CHC chemical
signatures (one individual was isolated from each colony).

		Dates of sample
pool of	Colonies from	collections
Males	Bourgueil	17/10/2011
	Sepmes	17/10/2011
	Azay sur Indre	21/11/2011
	Civray de touraine	01/12/2011
	Lussault	16/10/2012
	Montlouis sur Loire	22/10/2012
	Lerné	23/10/2012
	Athée	09/11/2012
	Tours	16/092013
Workers	Sepmes	17/10/2011
	Chinon	28/10/2011
	Azay sur Indre	21/11/2011
	Ligueil	26/11/2011
	Civray de touraine	01/12/2011
	Reignac sur Indre	06/12/2011
	Montlouis sur Loire	22/10/2012
	Lerné	23/10/2012
	Artannes	08/11/2012
Gynes	Sepmes	17/10/2011
	Barrou	24/10/2011
	Ligueil (b)	25/10/2011
	Ligueil(<i>a</i>)	26/11/2011
	Civray de touraine	01/12/2011
	Montlouis sur Loire	22/10/2012
	Lerné	23/10/2012
	Artannes	08/11/2012
	Orléans	12/12/2012
Queens	Ligueil (b)	25/10/2011
	Montlouis sur Loire	22/10/2012
	St Avertin	27/08/2013
	Tours	16/09/2013
	Liguei (c)	11/10/2013

Table 2 Identification and relative proportion of the cuticular hydrocarbon compounds
(CHCs) in the chemical signature of each caste of *Vespa velutina*. Compounds are the same
between males and females (workers and gynes). Peaks pQ1 to pQ4 are queen-specifics.

				Workers Males Gynes		Males		es	Queens		
Peaks	RT (min)	ECL	Identification	%	± SE	%	± SE	%	± SE	%	± SE
C20	23,12	20,00	n-eicosane								
p1	24,66	21,00	<i>n</i> -C21	0,23	0,31	0,48	0,63	0,56	1,22	0,20	0,06
p2	25,27	21,38	11-MeC21	0,59	1,17	0,50	0,46	0,26	0,35	0,19	0,12
р3	26,26	22,00	<i>n</i> -C22	0,03	0,05	0,10	0,17	0,08	0,13	0,04	0,01
p4	27,47	22,75	C23:1	0,09	0,13	0,14	0,16	0,24	0,24	0,03	0,02
p5	27,58	22,81	C23:1	0,03	0,09	0,03	0,05	0,03	0,05	0,01	0,00
p6	27,89	23,00	<i>n</i> -C23	2,03	2,41	4,14	3,83	5,01	6,65	2,99	0,58
p7	28,45	23,36	11-MeC23	0,11	0,14	0,21	0,20	0,29	0,20	0,05	0,02
p8	28,55	23,42	7-MeC23	0,01	0,02	0,02	0,05	0,03	0,04	0,01	0,00
p9	28,68	23,51	5-MeC23	0,00	0,00	0,00	0,01	0,01	0,02	0,01	0,00
p10	29,03	23,73	3-MeC23	0,11	0,07	0,16	0,11	0,24	0,11	0,09	0,03
p11	29,46	24,00	<i>n</i> -C24	0,36	0,18	0,59	0,51	0,48	0,21	0,83	0,13
p12	30,02	24,34	12-MeC24	0,08	0,06	0,10	0,06	0,13	0,03	0,02	0,01
p13	30,68	24,75	C25:1	1,21	0,63	1,27	0,74	2,10	1,46	1,53	0,73
p14	30,78	24,81	C25:1	0,17	0,10	0,23	0,10	0,31	0,09	0,09	0,03
p15	31,08	25,00	<i>n</i> -C25	14,93	4,56	17,94	5,38	14,96	4,98	39,84	4,60
p16	31,59	25,34	13-+11-MeC25	2,09	1,45	2,77	1,54	5,79	1,06	0,65	0,21
p17	31,70	25,41	7-MeC25	0,18	0,13	0,25	0,12	0,49	0,11	0,05	0,02
p18	31,83	25,50	5-MeC25	0,40	0,14	0,39	0,16	0,54	0,40	0,39	0,09
p19	32,05	25,65	C26:1	0,09	0,06	0,06	0,07	0,17	0,05	0,01	0,01
p20	32,19	25,73	3-MeC25	4,34	1,28	4,67	1,42	4,15	1,11	4,12	0,85
p21	32,33	25,83	5,11-+5,9-diMeC25	0,21	0,09	0,14	0,05	0,25	0,06	0,08	0,02
p22	32,58	26,00	<i>n</i> -C26	1,32	0,54	1,38	0,95	1,00	0,41	2,25	0,34
p23	32,68	26,07	diMe or triMeC25	0,15	0,07	0,08	0,04	0,19	0,11	0,14	0,05
p24	33,09	26,33	12-+10-MeC26	0,56	0,21	0,56	0,16	0,97	0,22	0,12	0,03
p25	33,21	26,41	?	0,00	0,00	0,01	0,02	0,03	0,04	0,02	0,00
p26	33,27	26,45	?	0,05	0,03	0,03	0,03	0,03	0,02	0,02	0,00
p27	33,35	26,50	?	0,02	0,04	0,01	0,02	0,03	0,03	0,03	0,01
p28	33,47	26,58	4-MeC26	0,37	0,16	0,25	0,09	0,17	0,05	0,29	0,04
pQ1	33,64	26,69	C27:1	0,00	0,00	0,00	0,00	0,00	0,00	1,59	0,86
p29	33,78	26,78	C27:1	7,98	5,83	5,88	1,66	6,58	2,23	3,39	1,31
p30	33,87	26,84	C27:1	3,57	5,99	0,78	0,22	0,86	0,15	0,19	0,13
p31	34,12	27,00	<i>n</i> -C27	10,73	6,18	12,23	3,76	8,21	2,06	19,95	2,62
p32	34,61	27,34	13-+11-+9-MeC27	6,99	2,58	8,79	2,67	13,37	3,34	1,19	0,60
p33	34,72	27,42	7-MeC27	0,96	0,38	1,04	0,38	1,79	0,59	0,39	0,13
p34	34,85	27,51	5-MeC27	0,80	0,43	0,63	0,39	0,65	0,21	0,69	0,12
p35	35,00	27,62	11,15-diMeC27	3,40	1,26	2,42	0,75	3,06	1,07	0,33	0,23
p36	35,06	27,66	9,15-diMeC27	0,33	0,60	0,04	0,18	0,37	0,53	0,00	0,00

p37	35,20	27,75	3-MeC27	7,47	2,56	7,22	3,24	4,68	1,46	5,66	0,83
p38	35,31	27,83	5,15-diMeC27	1,40	2,43	0,54	0,28	0,72	0,29	0,19	0,05
p39	35,38	27,88	?	0,39	0,64	0,41	0,39	0,10	0,11	0,07	0,04
p40	35,45	27,92	?	0,06	0,35	0,00	0,01	0,05	0,04	0,00	0,00
p41	35,56	28,00	<i>n</i> -C28	0,89	0,69	0,74	1,05	0,40	0,33	0,33	0,03
p42	35,64	28,06	diMe or triMeC27	0,89	0,76	0,72	0,49	0,55	0,28	0,50	0,12
p43	36,02	28,32	12-MeC28	0,50	0,47	0,44	0,23	0,67	0,21	0,15	0,06
p44	36,28	28,50	8-MeC28	0,54	1,32	0,18	0,76	0,02	0,02	0,05	0,02
p45	36,42	28,60	4-MeC28	0,67	1,01	0,83	1,89	0,28	0,14	0,15	0,04
p46	36,55	28,70	C29:2	0,58	1,00	0,60	1,47	0,10	0,05	1,42	0,62
p47	36,70	28,80	C29:1	3,94	2,68	2,81	1,17	2,29	0,81	0,62	0,20
p48	36,82	28,88	C29:1	2,45	1,72	0,98	0,30	0,71	0,17	0,28	0,15
p49	36,99	29,00	<i>n</i> -C29	1,49	0,76	1,61	1,13	0,84	0,34	1,32	0,20
p50	37,29	29,27	?	0,57	0,63	0,67	0,40	0,26	0,13	0,21	0,08
p51	37,44	29,40	13-+11-+9-MeC29	1,52	0,63	1,97	0,64	3,00	0,87	0,44	0,17
p52	37,57	29,51	7-MeC29	0,07	0,13	0,09	0,06	0,20	0,11	0,05	0,01
p53	37,82	29,73	11,15-+11,17-diMeC29	2,34	0,62	1,85	0,66	2,12	0,79	0,26	0,15
p54	37,90	29,81	3-MeC29	0,37	0,30	0,39	0,27	0,67	0,48	0,14	0,05
p55	38,02	29,92	5,17-diMeC29	0,56	0,31	0,44	0,20	0,24	0,15	0,48	0,07
p56	38,12	30,00	<i>n</i> -C30	0,46	0,22	0,26	0,15	0,27	0,11	0,18	0,05
p57	38,81	30,43	4-MeC30	1,30	1,30	0,94	0,90	0,78	0,59	0,20	0,09
pQ2	39,31	30,74	C31:1	0,00	0,00	0,00	0,00	0,00	0,00	0,94	0,28
p58	39,47	30,84	C31:1	0,49	0,28	0,40	0,15	0,28	0,10	0,12	0,03
p59	39,61	30,93	?	0,20	0,08	0,15	0,08	0,09	0,04	0,05	0,01
p60	39,73	31,00	<i>n</i> -C31	0,34	0,29	0,57	0,64	0,32	0,27	0,29	0,05
p61	40,14	31,31	15-+13-+11-MeC31	0,69	0,39	1,06	0,35	1,39	0,53	0,31	0,07
p62	40,36	31,48	?	0,06	0,18	0,21	0,41	0,04	0,09	0,03	0,01
p63	40,49	31,58	11,15-diMeC31	1,51	0,62	1,53	0,65	1,86	0,80	0,24	0,13
pQ3	41,94	32,69	C33:1	0,00	0,00	0,00	0,00	0,00	0,00	1,65	0,80
p64	42,07	32,79	C33:1	0,17	0,17	0,15	0,11	0,12	0,13	0,07	0,02
p65	42,34	33,00	<i>n</i> -C33	0,16	0,26	0,30	0,42	0,15	0,13	0,11	0,02
p66	42,75	33,26	15-+13-+11-MeC33	0,40	0,23	0,66	0,27	0,78	0,40	0,18	0,05
p67	43,15	33,53	11,19-+11,17-diMeC33	1,04	0,47	0,99	0,49	1,01	0,49	0,23	0,09
p68	44,46	34,38	?	0,36	0,52	0,41	0,44	0,22	0,30	0,42	0,23
pQ4	44,73	34,56	C35:1	0,00	0,00	0,00	0,00	0,00	0,00	0,41	0,27
p69	45,80	35,26	11-MeC35	0,24	0,14	0,38	0,21	0,37	0,23	0,12	0,04
p70	46,27	35,57	13,19-+11,17-diMeC35	1,19	0,63	1,10	0,65	0,95	0,54	0,25	0,09
p71	46,49	35,71	?	0,16	0,12	0,09	0,08	0,06	0,05	0,06	0,03

Table 3 Caste comparisons of each cuticular hydrocarbon compound (CHCs) in the chemical

513 signature (Kruskal-Wallis chi-squared test).

		Caste comparisons								
Peaks	global comparison	Male-Workers	Male-Queen	Males-Gynes	Workers-Queens	Workers-Gynes	Gynes-Queens			
p1	X ² = 23.7183, df = 3, p = 2.86e-05	p=0.001	N.S	N.S	N.S	p=0.01	N.S			
p2	X ² = 12.9491, df = 3, p = 0.004748	N.S	N.S	p=0.01	N.S	N.S	N.S			
р3	X ² = 21.2598, df = 3, p = 9.297e-05	p=0.01	N.S	N.S	N.S	p=0.001	N.S			
p4	X ² = 30.3086, df = 3, p = 1.188e-06	N.S	N.S	p=0.01	N.S	p=0.001	p=0.001			
р5	X ² = 14.4091, df = 3, p = 0.002398	N.S	N.S	N.S	N.S	p=0.05	N.S			
р6	X ² = 41.2662, df = 3, p = 5.742e-09	p=0.001	N.S	N.S	p=0.01	p=0.001	N.S			
р7	X ² = 68.7282, df = 3, p = 7.99e-15	p=0.001	p=0.01	p=0.01	N.S	p=0.001	p=0.001			
p8	X ² = 47.6831, df = 3, p = 2.487e-10	p=0.05	N.S	p=0.05	p=0.01	p=0.001	N.S			
р9	X ² = 99.9361, df = 3, p < 2.2e-16	N.S	p=0.001	N.S	p=0.001	N.S	p=0.001			
p10	X ² = 61.3123, df = 3, p = 3.082e-13	p=0.01	N.S	p=0.001	N.S	p=0.001	p=0.001			
p11	X ² = 55.6694, df = 3, p = 4.942e-12	p=0.001	p=0.001	N.S	p=0.001	p=0.01	p=0.001			
p12	X ² = 67.5535, df = 3, p = 1.426e-14	N.S	p=0.001	p=0.001	p=0.001	p=0.001	p=0.001			
p13	X ² = 14.5332, df = 3, p = 0.002262	N.S	N.S	p=0.01	N.S	p=0.01	N.S			
p14	X ² = 92.0405, df = 3, p < 2.2e-16	p=0.01	p=0.001	p=0.001	p=0.05	p=0.001	p=0.001			
p15	X ² = 55.2756, df = 3, p = 5.997e-12	p=0.05	p=0.001	p=0.01	p=0.001	N.S	p=0.001			
p16	X ² = 146.472, df = 3, p < 2.2e-16	N.S	p=0.001	p=0.001	p=0.01	p=0.001	p=0.001			
p17	X ² = 140.5225, df = 3, p < 2.2e-16	N.S	p=0.001	p=0.001	p=0.01	p=0.001	p=0.001			
p18	X ² = 21.9075, df = 3, p = 6.819e-05	N.S	N.S	p=0.001	N.S	p=0.01	N.S			
p19	X ² = 98.4494, df = 3, p < 2.2e-16	p=0.05	N.S	p=0.001	p=0.05	p=0.001	p=0.001			
p20	X ² = 7.2278, df = 3, p = 0.06498	N.S	N.S	N.S	N.S	N.S	N.S			
p21	X ² = 114.3282, df = 3, p < 2.2e-16	p=0.001	N.S	p=0.001	p=0.001	N.S	p=0.001			
p22	X ² = 63.2816, df = 3, p = 1.169e-13	N.S	p=0.001	p=0.001	p=0.001	p=0.001	p=0.001			
p23	X ² = 79.0834, df = 3, p < 2.2e-16	p=0.001	p=0.01	p=0.001	N.S	N.S	N.S			
p24	X ² = 134.2695, df = 3, p < 2.2e-16	N.S	p=0.001	p=0.001	p=0.001	p=0.001	p=0.001			
p25	X ² = 56.1558, df = 3, p = 3.891e-12	N.S	p=0.01	N.S	p=0.001	p=0.01	p=0.05			
p26	X ² = 35.8875, df = 3, p = 7.91e-08	p=0.001	N.S	N.S	p=0.01	p=0.001	N.S			
p27	X ² = 32.1362, df = 3, p = 4.899e-07	N.S	p=0.01	p=0.001	p=0.05	p=0.05	N.S			
p28	X ² = 86.5888, df = 3, p < 2.2e-16	p=0.001	N.S	p=0.001	N.S	p=0.001	p=0.001			
р29	X ² = 25.3108, df = 3, p = 1.329e-05	N.S	p=0.001	N.S	p=0.001	N.S	p=0.001			
p30	X ² = 94.0832, df = 3, p < 2.2e-16	p=0.001	p=0.001	N.S	p=0.001	p=0.001	p=0.001			
p31	X ² = 82.2795, df = 3, p < 2.2e-16	N.S	p=0.001	p=0.001	p=0.001	p=0.001	p=0.001			
р32	X ² = 132.0132, df = 3, p < 2.2e-16	p=0.05	p=0.001	p=0.001	p=0.001	p=0.001	p=0.001			
р33	X² = 107.1492, df = 3, p < 2.2e-16	N.S	p=0.001	p=0.001	p=0.001	p=0.001	p=0.001			
р34	X² = 12.6864, df = 3, p = 0.005366	p=0.01	N.S	N.S	N.S	N.S	N.S			
р35	X ² = 70.7854, df = 3, p = 2.898e-15	p=0.001	p=0.001	p=0.01	p=0.001	N.S	p=0.001			
р36	X ² = 27.8999, df = 3, p = 3.812e-06	N.S	N.S	p=0.01	N.S	N.S	N.S			
р37	X ² = 50.9824, df = 3, p = 4.935e-11	N.S	N.S	p=0.001	N.S	p=0.001	N.S			
p38	X² = 88.9472, df = 3, p < 2.2e-16	p=0.001	p=0.01	p=0.01	p=0.001	p=0.01	p=0.001			
p39	X² = 47.5966, df = 3, p = 2.595e-10	p=0.001	p=0.01	p=0.001	N.S	N.S	N.S			
p40	X² = 94.4655, df = 3, p < 2.2e-16	N.S	N.S	p=0.001	N.S	p=0.001	p=0.001			
p41	X² = 79.079, df = 3, p < 2.2e-16	p=0.001	N.S	p=0.01	p=0.001	p=0.001	N.S			

p42	X ² = 23.715, df = 3, p = 2.865e-05	p=0.01	N.S	N.S	p=0.01	p=0.001	N.S
p43	X ² = 86.0122, df = 3, p < 2.2e-16	N.S	p=0.001	p=0.001	p=0.001	p=0.001	p=0.001
p44	X ² = 27.807, df = 3, p = 3.987e-06	N.S	p=0.001	N.S	p=0.001	N.S	p=0.001
p45	X ² = 53.2645, df = 3, p = 1.61e-11	N.S	p=0.001	N.S	p=0.001	p=0.001	p=0.01
p46	X ² = 31.2824, df = 3, p = 7.413e-07	N.S	p=0.001	N.S	p=0.001	N.S	p=0.001
p47	X ² = 72.9925, df = 3, p = 9.757e-16	N.S	p=0.001	N.S	p=0.001	p=0.001	p=0.001
p48	X ² = 152.917, df = 3, p < 2.2e-16	p=0.001	p=0.001	p=0.001	p=0.001	p=0.001	p=0.01
p49	X ² = 63.6709, df = 3, p = 9.652e-14	N.S	N.S	p=0.001	N.S	p=0.001	p=0.001
p50	X ² = 72.4929, df = 3, p = 1.248e-15	p=0.001	p=0.001	p=0.001	p=0.01	p=0.01	N.S
p51	X ² = 121.4764, df = 3, p < 2.2e-16	p=0.01	p=0.001	p=0.001	p=0.01	p=0.001	p=0.001
p52	X ² = 73.8372, df = 3, p = 6.431e-16	p=0.01	p=0.05	p=0.001	N.S	p=0.001	p=0.001
p53	X ² = 58.9537, df = 3, p = 9.834e-13	p=0.001	p=0.001	N.S	p=0.001	N.S	p=0.001
p54	X ² = 28.7926, df = 3, p = 2.476e-06	N.S	p=0.05	p=0.01	N.S	p=0.001	p=0.001
p55	X ² = 75.3837, df = 3, p = 2.998e-16	N.S	N.S	p=0.001	N.S	p=0.001	p=0.001
p56	X² = 61.4547, df = 3, p = 2.873e-13	p=0.001	N.S	N.S	p=0.001	p=0.001	N.S
p57	X ² = 40.9767, df = 3, p = 6.614e-09	N.S	p=0.001	N.S	p=0.001	p=0.01	p=0.001
p58	X ² = 72.5886, df = 3, p = 1.191e-15	N.S	p=0.001	p=0.001	p=0.001	p=0.001	p=0.001
p59	X ² = 96.7315, df = 3, p < 2.2e-16	p=0.01	p=0.001	p=0.001	p=0.001	p=0.001	N.S
p60	X ² = 13.5855, df = 3, p = 0.003527	p=0.01	N.S	p=0.001	N.S	N.S	N.S
p61	X ² = 96.2835, df = 3, p < 2.2e-16	p=0.001	p=0.001	p=0.05	p=0.05	p=0.001	p=0.001
p62	X ² = 31.5711, df = 3, p = 6.444e-07	N.S	p=0.05	N.S	p=0.001	N.S	p=0.05
p63	X ² = 49.9463, df = 3, p = 8.202e-11	N.S	p=0.001	N.S	p=0.001	N.S	p=0.001
p64	X² = 21.9726, df = 3, p = 6.609e-05	N.S	p=0.001	p=0.05	p=0.01	N.S	N.S
p65	X ² = 17.8008, df = 3, p = 0.0004835	p=0.001	N.S	N.S	N.S	N.S	N.S
p66	X ² = 78.504, df = 3, p < 2.2e-16	p=0.001	p=0.001	N.S	p=0.01	p=0.001	p=0.001
p67	X ² = 40.1883, df = 3, p = 9.719e-09	N.S	p=0.001	N.S	p=0.001	N.S	p=0.001
p68	X ² = 15.8611, df = 3, p = 0.001211	N.S	N.S	p=0.01	N.S	N.S	p=0.01
p69	X ² = 45.4726, df = 3, p = 7.342e-10	p=0.001	p=0.001	N.S	p=0.01	p=0.01	p=0.001
p70	X ² = 36.0571, df = 3, p = 7.283e-08	N.S	p=0.001	N.S	p=0.001	N.S	p=0.001
p71	X ² = 34.6872, df = 3, p = 1.418e-07	p=0.001	N.S	N.S	p=0.01	p=0.001	N.S
pQ1	X ² = 243.6771, df = 3, p < 2.2e-16	N.S	p=0.001	N.S	p=0.001	N.S	p=0.001
pQ2	X ² = 243.6771, df = 3, p < 2.2e-16	N.S	p=0.001	N.S	p=0.001	N.S	p=0.001
pQ3	X ² = 243.6771, df = 3, p < 2.2e-16	N.S	p=0.001	N.S	p=0.001	N.S	p=0.001
pQ4	X ² = 243.6771, df = 3, p < 2.2e-16	N.S	p=0.001	N.S	p=0.001	N.S	p=0.001











Figure 5



Figure 6



