

# Volatile Organic Compounds Induced by Herbivory of the Soybean Looper *Chrysodeixis includens* in Transgenic Glyphosate-Resistant Soybean and the Behavioral Effect on the Parasitoid, *Meteorus rubens*

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**Abstract** Transgenic soybean plants (RR) engineered to express resistance to glyphosate harbor a variant of the enzyme EPSPS (5-enolpyruvylshikimate-3-phosphate synthase) involved in the shikimic acid pathway, the biosynthetic route of three aromatic amino acids: phenylalanine, tyrosine, and tryptophan. The insertion of the variant enzyme CP4 EPSPS confers resistance to glyphosate. During the process of genetic engineering, unintended secondary effects are likely to occur. In the present study, we quantified volatile organic compounds (VOCs) emitted constitutively or induced in response to herbivory by the soybean looper *Chrysodeixis includens* in transgenic soybean and its isogenic (untransformed) line. Since herbivore-induced plant volatiles (HIPVs) are known to play a role in the recruitment of natural enemies, we assessed whether changes in VOC profiles alter the foraging behavior of the generalist endoparasitic larval parasitoid, *Meteorus rubens* in the transgenic line. Additionally, we assessed whether there was a difference in plant quality by measuring the weight gain of the soybean looper. In response to herbivory, several VOCs were induced in both the conventional and the transgenic line; however, larger quantities of a few compounds were emitted by transgenic plants. *Meteorus rubens* females were able to discriminate between the odors of

undamaged and *C. includens*-damaged plants in both lines, but preferred the odors emitted by herbivore-damaged transgenic plants over those emitted by herbivore-damaged conventional soybean plants. No differences were observed in the weight gain of the soybean looper. Our results suggest that VOC-mediated tritrophic interactions in this model system are not negatively affected. However, as the preference of the wasps shifted towards damaged transgenic plants, the results also suggest that genetic modification affects that tritrophic interactions in multiple ways in this model system.

**Keywords** Herbivory; parasitoids; shikimic acid pathway; induced volatiles · Lepidoptera · Noctuidae · Hymenoptera · Braconidae

## Introduction

Co-evolution of plants with herbivores has resulted in plant recognition of foreign molecules and cellular damage that activates a number of induced responses to herbivory in plants (Hare 2011). These induced responses include changes in specialized morphological structures such as trichomes, as well as in chemical compounds that directly influence the colonization and performance of herbivores (Karban and Baldwin 1997; Mithöfer and Boland 2012). In addition, plants can defend themselves indirectly by recruiting natural enemies of the attacking herbivore (Arimura et al. 2009), by offering an increase in resources such as extrafloral nectar (Turlings and Wäckers 2004), or through the emission of herbivore-induced plant volatiles (HIPVs) that attract predators and parasitoids of the herbivore. Herbivore-induced plant volatiles comprise an array of compounds from different biosynthetic pathways, some of which can be synthesized *de novo* in

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response to herbivory (Dudareva et al. 2006). During the last two decades, HIPVs have been studied extensively from molecular to community levels because manipulation of this ecological mechanism offers the possibility to improve biological control and develop novel, environmentally-friendly, pest management strategies (Kaplan 2012; Turlings and Ton 2006).

The development of soybeans resistant to the herbicide glyphosate through genetic modification has changed soybean production globally, including in Brazil, where the use of transgenic plants has been authorized since 2005. Roundup Ready® (RR) soybean plants express a variant of the enzyme 5-enolpyruvylshikimate-3-phosphate synthase (EPSPS). This enzyme participates in the shikimic acid pathway, the biosynthetic route of three aromatic amino acids: phenylalanine, tyrosine, and tryptophan (Maeda and Dudareva 2012). In addition to their role in the synthesis of proteins for plant development (Hermann and Weaver 1999), these aromatic amino acids are precursors of secondary metabolites involved in plant defense (Buchanan et al. 2000). Tryptophan is the precursor of indole, a volatile compound induced in response to herbivory in various plant species, including maize and soybean (Rostás and Eggert 2008; Turlings et al. 2005). In maize, this compound is essential in priming maize plants (Erb et al. 2015), and recently it has been demonstrated that indole acts as a direct defense against herbivores (Veyrat et al. 2016). L-Phenylalanine is the precursor of volatile phenylpropanoids/benzenoids, such as methyl salicylate (MeSA), which also is induced by soybean plants in response to herbivory (Michereff et al. 2011; Rostás and Eggert 2008). Methyl salicylate has important ecological functions in arthropod-plant interactions by acting as an infochemical for predators (De Boer and Dicke 2004) and by preventing plant colonization by aphids (Mallinger et al. 2011). Moreover, it is a mobile signal for the development of systemic acquired resistance against pathogens (Park et al. 2007). Salicylic acid (SA), its precursor, is involved in the activation of direct and indirect defenses against pathogens and phloem-feeding insects (Loake and Grant 2007; Zarate et al. 2007 and references therein). Recent studies have shown that the insertion of a gene that expresses an insensitive form of the EPSPS to confer resistance to glyphosate affects soybean metabolism. Altered expression of seed protein (Brandão et al. 2010; Garcia-Villalba et al. 2008), as well as altered accumulation of lignin in roots compared to conventional lines already has been reported (Zonetti et al. 2012). Cheng et al. (2008) did not find any significant differences in gene expression in soybean leaves between conventional and transgenic lines. However, the authors reported down-regulation of genes involved in cysteine protease inhibitor activity and dihydroflavonol-4-reductase activity in transgenic lines, possibly as an unintended consequence of genetic transformation (Cheng et al. 2008). According to Arruda et al. (2013), genetic modification itself contributes to alterations in a variety of soybean plant traits in

a cascading manner. Based on this, not only primary but also secondary metabolism may be modified due to the insertion of the gene for CP4 EPSPS.

Because the behavior and performance of herbivores, and indirectly of natural enemies, is influenced by primary and secondary plant metabolism, any variation in the physical and chemical traits may affect plant-insect interactions. In transgenic maize plants engineered to express delta-endotoxins from *Bacillus thuringiensis*, emissions of HIPVs are altered, possibly as a result of a combination of altered insect feeding and changes in carbon allocation, but no negative effects on the recruitment of two parasitoids have been reported (Dean and De Moraes 2006; Moraes et al. 2011; Turlings et al. 2005). Recently, we have shown that herbivory by the velvet bean caterpillar, *Anticarsia gemmatalis* (Lepidoptera: Erebidae) induces much larger emissions of individual volatile compounds in transgenic plants expressing resistance to glyphosate compared with plants from the isogenic line (Strapasson et al. 2016). However, no negative effect on the foraging of the predatory bug, *Podisus nigrispinus* (Hemiptera: Pentatomidae) was observed.

In this study, we addressed whether transgenic soybean plants that harbor the gene for CP4 EPSPS resistance to glyphosate emit a VOC profile different from the isogenic line in response to feeding by the soybean looper, *Chrysodeixis includens* (Lepidoptera; Noctuidae). We also assessed whether any variation in HIPV emissions between lines affects the behavior of the parasitoid, *Meteorus rubens* (Hymenoptera: Braconidae). Finally, we tested the performance of the soybean looper on both transgenic and isogenic soybean lines. The soybean looper was regarded as a secondary pest of soybean crops in Brazil until 2003. Since then, this species has become a major soybean pest, likely due to elimination of natural enemies because of excessive pesticide sprayings, and due to the expansive planting of soybean in monoculture (Bueno et al. 2007). *Meteorus rubens* is a gregarious larval endoparasitoid of various species of noctuids. In Southern Brazil, it commonly is found in soybean fields parasitizing the soybean looper (Strapasson et al. unpublished data).

## Methods and Materials

**Living Material** Soybean seeds (*Glycine max*) from a transgenic cultivar (CD 202 RR) and its isogenic line (CD 202) were provided by COODETEC (Cascavel, Paraná, Brazil). Germinating seeds were placed in the dark for 4–5 d at 25 °C. After this period, seeds were planted in plastic pots (top 6 × 6 cm; bottom 4.5 × 4.5 cm; height 5.5 cm) filled with commercial substrate (Vida Verde, São Paulo, Brazil). Plants were kept under artificial lighting (14:10 h L:D regime), and watered as needed. They were used in the experiments when they reached the V3 phenological stage as described by Fehr

et al. (1971), approximately 5 wk. after planting. A colony of *C. includens* was started in the laboratory from eggs purchased commercially (BUG, Piracicaba, São Paulo, Brazil), and kept in plastic pots containing a bean-based artificial diet (Parra 2001). Larvae were kept under controlled conditions of  $25 \pm 1$  °C and a photoperiod of 14:10 L:D until pupation. Pupae were kept in vermiculite and, just prior to emergence, were transferred to PVC tubes (15 cm) lined with white paper as an oviposition substrate. As adults emerged, they were fed on a honey-based solution [honey, sorbic acid, methyl paraben, sugar, and distilled water, then mixed with beer (3:1 v/v)] according to Hoffman Campo et al. (1985). Every 2–3 d, eggs were collected to start a new cycle. *Meteorus rubens* used in the bioassays were recovered from parasitized larvae of the soybean looper collected in the experimental station “Gralha Azul” from the Pontifícia Universidade Católica do Paraná - PUCPR (Fazenda Rio Grande, Paraná, Brazil). Larvae were kept as previously described until the emergence of the parasitoids. Wasps were identified by Dr. Eduardo Shimbori (Universidade Federal de São Carlos, São Carlos, Brazil) and Dr. Helmuth Aguirre (University of Wyoming, Laramie, USA). Adults were maintained on a 10 % honey solution until they were used in the bioassays.

**VOC Collection and Analyses** Volatiles were collected from undamaged (control) and *C. includens*-damaged plants (treatment) from both, CD 202 and CD 202 RR. Herbivore induction was accomplished by infesting each plant with 10 3rd-instar larvae for a 24-h period. Prior to sampling, larvae were removed, and the unpotted plants were wrapped in aluminum foil in order to collect VOCs from the above ground parts only. Four plants (one per treatment) were sampled simultaneously. For this, each plant was placed in a sealed 1-L chamber. Humidified, charcoal-filtered air was pumped into each chamber at a rate of 1 L/min through Teflon tubes. VOCs were trapped onto ca. 20 mg HayeSep® Q 80–100 mesh in a glass tube for 24 h. Trapped VOCs were eluted with 300  $\mu$ l of twice-distilled HPLC-grade hexane, and 10  $\mu$ l of tetradecane (50-ppm solution) were added as an internal standard (IS). Samples were concentrated to 100  $\mu$ l, and 1  $\mu$ l of the extract was injected automatically in splitless mode (injector temperature 250 °C), and analyzed by GC-MS (Shimadzu QP 2010 Plus) with an RTX-5 column (30 m  $\times$  0.25 mm i.d., 0.25 mm film thickness; Restek, Bellefonte, PA, USA). The column oven temperature was held at 40 °C for 1 min, increased to 250 °C at 7 °C min<sup>-1</sup>, and held for 10 min. Helium was the carrier gas at a column head pressure of 170 kPa. The quantification of individual compounds was carried out based on the peak area of the IS. Emissions were expressed as ng/g dry weight/24 h. Quantitation of dry weight (DW) was obtained by cutting and drying the above ground plant parts for 48 h at 60 °C. Tentative identification of individual compounds was carried out by comparing the mass spectrum to those in

reference libraries (NIST 27 and NIST 147), and by comparing their calculated Kovats Indexes (KI) with those reported in the literature. GC co-injections with synthetic compounds also were conducted for (*Z*)-3-hexen-1-ol, (*Z*)-3-hexenyl acetate, octanal, 1-octen-3-ol, (*Z*)- $\beta$ -ocimene,  $\beta$ -linalool, decanal, methyl salicylate, indole, (*Z*)-jasnone,  $\beta$ -caryophyllene, and  $\alpha$ -farnesene. A total of 40 plants, 10 from each treatment ( $N = 10$ ) were sampled.

**Behavioral Response of *M. rubens*** The responses of *M. rubens* were tested in dual-choice bioassays in a Y-tube olfactometer ( $\emptyset = 2$  cm; main arm = 18 cm; smaller arms = 13 cm). A humidified, charcoal-filtered airflow of 1 L/min was equally split in two and channeled to each small arm. At the distal ends of the small arms, odor sources consisting of 20  $\mu$ l of the extract (trapped soybean plant VOCs eluted in hexane and concentrated to 100  $\mu$ l as previously described) or hexane were placed on a piece of filter paper (1  $\times$  1 cm). At the end of the main arm of the olfactometer, 3–4-d-old mated female parasitoids were introduced individually and observed for a period of 10 min. We considered that an insect had responded when it passed the halfway point of one of the small arms, and remained there for at least 5 s; otherwise, the insect was considered as non-responsive, and excluded from the data analysis. The odor source was replaced, and the olfactometer was rotated 180°, after each observation. Every 5 observations, the position of the odor sources was switched. The following tests were conducted: a) undamaged vs. herbivore-damaged plants (isogenic line), b) undamaged vs. herbivore-damaged plants (transgenic line), 3) herbivore-damaged plants from the isogenic line vs. the herbivore-damaged plants from the transgenic line. The bioassays were conducted between 0900 and 1700 h at 25 °C over a period of 6 d.

**Feeding Performance of *C. includens*** One-d-old larvae were weighed and kept singly in plastic pots under controlled conditions of  $25 \pm 1$  °C and a photoperiod of 14:10 L:D. For 7 d, they were fed on leaves excised from either CD 202 (isogenic line) or CD 202 RR (transgenic line) plants (at the V3 stage). After this period, they were weighed again. The initial and final weights were recorded ( $N = 19$  and  $N = 22$  for CD 202 and CD 202 RR, respectively). Larvae that died before completing the bioassay were not included in the data analysis.

**Statistical Analyses** Individual VOCs were log transformed ( $X + 1$ ) and subjected to a Principal Component Analysis (PCA). Analysis of Variance (ANOVA) followed by Contrast Analysis was performed for those PCs that explained the main percentage variance to visualize differences between treatments. For a more detailed analysis, individual compounds and total VOC emissions were analyzed using one-way ANOVA followed by a Tukey HSD post hoc test. The

preference of *M. rubens* was analyzed by performing a *Chi-squared* test. The performance of the soybean looper was checked for normality and analyzed by performing a *t*-test to compare the means between weight gain (final – initial weight) by the larvae fed on leaves of the transgenic or the isogenic line. Data were analyzed in the “R” Environment (R Development Core Team, 2006), except for the analysis of feeding preference that was performed on SPSS 21.0 for Windows (SPSS Inc., Chicago, IL, USA).

## Results

**VOCs Emitted by Transgenic and Isogenic Soybean Plants** Volatiles detected in the headspace of soybean plants from the various treatments and comparisons between treatments are presented in Table 1. Herbivory induced the emission of (*Z*)-3-hexenyl-2-methylbutyrate and (*Z*)-jasmone, plus two unidentified compounds (1 and 2) in the isogenic line; and (*Z*)-3-hexen-1-ol, 3-octanone,  $\beta$ -linalool, (*Z*)-3-hexenyl-2-methylbutyrate, (*Z*)-jasmone,  $\beta$ -caryophyllene,  $\alpha$ -caryophyllene, and TMTT, plus three

unidentified compounds (1–3) in the transgenic line. The PC1 explained 85.42 % of the variance. There were significant differences between treatments (ANOVA  $F = 4.76$ ;  $P = 0.007$ ), and the VOC emissions of transgenic *C. includens*-damaged plants differed from the rest of the treatments ( $P < 0.05$ ), mainly because of the variation of  $\alpha$ -farnesene, indole, (*Z*)- $\beta$ -ocimene, and (*Z*)-3-hexen-1-ol (Figs. 1, 2). There were differences in the total amount of VOCs between treatments (ANOVA  $F = 18.357$ ;  $P \leq 0.001$ ). However, the total induced VOC emissions in transgenic plants did not differ from those in conventional plants (Tukey HSD post hoc  $P = 0.138$ ). From those compounds that were significantly increased in response to herbivory (*Z*)- $\beta$ -ocimene (Tukey HSD post hoc  $P = 0.027$ ),  $\beta$ -linalool (Tukey HSD post hoc  $P = 0.007$ ), and indole (Tukey HSD post hoc  $P = 0.026$ ) were emitted in larger quantities by transgenic than by conventional plants.

**Behavioral Preference of *M. rubens*** *Meteorus rubens* discriminated between the odors from undamaged and *C. includens*-damaged plants from the isogenic line

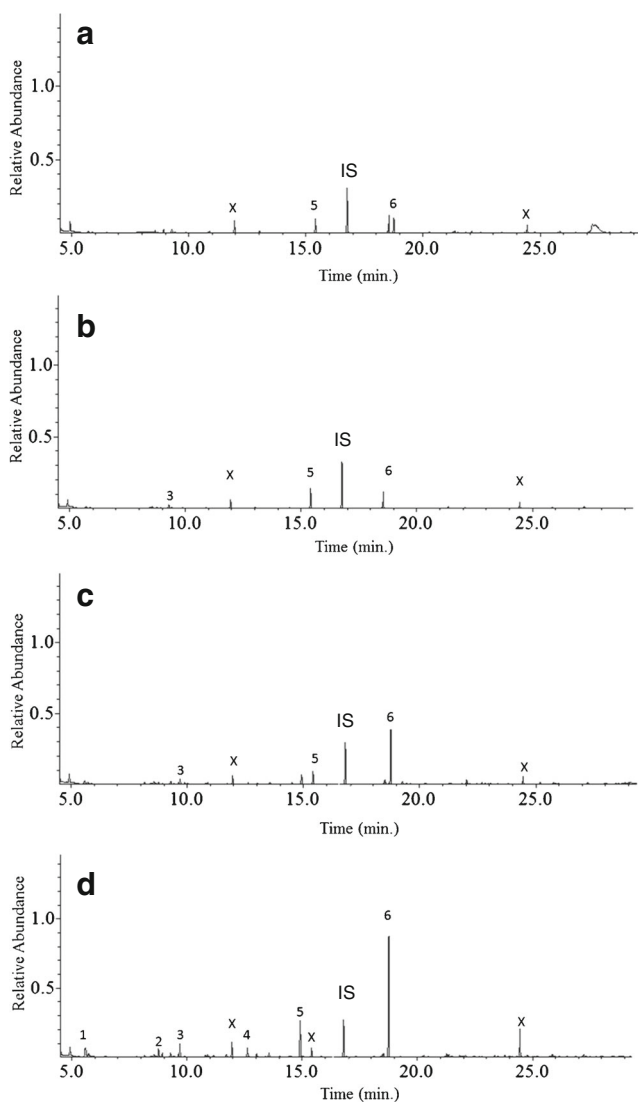
**Table 1** Emissions of individual volatile organic compounds (ng VOC/g dry weight/24 h) BY UNDAMAGED AND *Chrysodeixis includens*-damaged plants in a soybean transgenic line and its isogenic line (Mean  $\pm$  S.E. M)

| Compounds                               | R.T.   | K.I. (c) <sup>a</sup> | K.I. (r) <sup>b</sup> | Undamaged plants           |                    | <i>C.includens</i> -damaged plants |                    |
|-----------------------------------------|--------|-----------------------|-----------------------|----------------------------|--------------------|------------------------------------|--------------------|
|                                         |        |                       |                       | Isogenic (IU) <sup>c</sup> | Transgenic (TU)    | Isogenic (IH)                      | Transgenic (TH)    |
| ( <i>Z</i> )-3-Hexen-1-ol               | 5.581  | 851                   | 857                   | 0.04 $\pm$ 0.03 b          | 0.00 b             | 1.42 $\pm$ 0.53 a                  | 2.43 $\pm$ 0.77 a  |
| 1-Octen-3-ol                            | 8.155  | 978                   | 980                   | 0.15 $\pm$ 0.10 cb         | 0.01 bc            | 0.23 $\pm$ 0.06 ab                 | 0.55 $\pm$ 0.18 a  |
| 3-Octanone                              | 8.293  | 985                   | 985                   | 0.00 b                     | 0.00 b             | 0.00 b                             | 0.04 $\pm$ 0.02 a  |
| Octanal                                 | 8.668  | 1003                  | 1004                  | 0.12 $\pm$ 0.03 a          | 0.09 $\pm$ 0.02 a  | 0.11 $\pm$ 0.02 a                  | 0.14 $\pm$ 0.03 a  |
| ( <i>Z</i> )-3-Hexenyl acetate          | 8.757  | 1007                  | 1007                  | 0.14 $\pm$ 0.09 bc         | 0.05 $\pm$ 0.02 c  | 0.51 $\pm$ 0.16 ab                 | 0.76 $\pm$ 0.20 a  |
| ( <i>Z</i> )- $\beta$ -Ocimene          | 9.680  | 1051                  | 1043                  | 0.10 $\pm$ 0.04 c          | 0.03 $\pm$ 0.01 c  | 1.51 $\pm$ 0.47 b                  | 3.80 $\pm$ 1.24 a  |
| $\beta$ -Linalool                       | 10.796 | 1100                  | 1100                  | 0.01 c                     | 0.00 c             | 0.24 $\pm$ 0.07 b                  | 0.61 $\pm$ 0.14 a  |
| Nonanal                                 | 10.877 | 1106                  | 1104                  | 0.56 $\pm$ 0.12 a          | 0.38 $\pm$ 0.07 a  | 0.45 $\pm$ 0.06 a                  | 0.73 $\pm$ 0.12 a  |
| Compound 1                              | 11.711 | 1144                  | -                     | 0.00 bc                    | 0.00 bc            | 0.37 $\pm$ 0.13 ab                 | 0.90 $\pm$ 0.39 a  |
| Compound 2                              | 12.608 | 1235                  | -                     | 0.00 b                     | 0.00 b             | 0.57 $\pm$ 0.29 a                  | 0.88 $\pm$ 0.30 a  |
| Methyl salicylate                       | 12.878 | 1201                  | 1206                  | 0.14 $\pm$ 0.05 ab         | 0.04 $\pm$ 0.02 bc | 0.26 $\pm$ 0.11 ab                 | 0.31 $\pm$ 0.06 a  |
| Decanal                                 | 13.007 | 1206                  | 1204                  | 0.62 $\pm$ 0.15 a          | 0.45 $\pm$ 0.11 a  | 0.55 $\pm$ 0.09 a                  | 0.97 $\pm$ 0.17 a  |
| ( <i>Z</i> )-3-Hexenyl-2-methylbutyrate | 13.549 | 1234                  | 1231                  | 0.00 b                     | 0.00 b             | 0.55 $\pm$ 0.18 a                  | 1.11 $\pm$ 0.41 a  |
| Indole                                  | 14.905 | 1304                  | 1304                  | 0.12 $\pm$ 0.05 c          | 0.01 c             | 4.26 $\pm$ 1.32 b                  | 8.08 $\pm$ 2.01 a  |
| Compound 3                              | 16.344 | 1380                  | -                     | 0.03 $\pm$ 0.01 a          | 0.00 a             | 0.09 $\pm$ 0.07 a                  | 0.04 $\pm$ 0.01 a  |
| ( <i>Z</i> )-Jasmone                    | 16.846 | 1406                  | 1394                  | 0.00 cb                    | 0.00 cb            | 0.05 $\pm$ 0.02 ab                 | 0.17 $\pm$ 0.08 a  |
| $\beta$ -Caryophyllene                  | 17.306 | 1433                  | 1428                  | 0.06 $\pm$ 0.02 b          | 0.00 b             | 0.06 $\pm$ 0.02 b                  | 0.17 $\pm$ 0.04 a  |
| $\alpha$ -Caryophyllene                 | 17.921 | 1454                  | 1454                  | 0.01 $\pm$ 0.00 b          | 0.00 b             | 0.01 $\pm$ 0.01 b                  | 0.05 $\pm$ 0.01 a  |
| $\alpha$ -Farnesene                     | 18.731 | 1515                  | 1508                  | 4.39 $\pm$ 2.07 b          | 2.34 $\pm$ 1.02 b  | 15.44 $\pm$ 4.68 a                 | 30.97 $\pm$ 8.50 a |
| TMTT                                    | 19.928 |                       | 1581                  | 0.02 b                     | 0.00 b             | 0.07 $\pm$ 0.02 b                  | 0.18 $\pm$ 0.04 a  |

<sup>a</sup> K.I. (c) = Calculated kovats indexes

<sup>b</sup> K.I. (r) = Reference kovats indexes

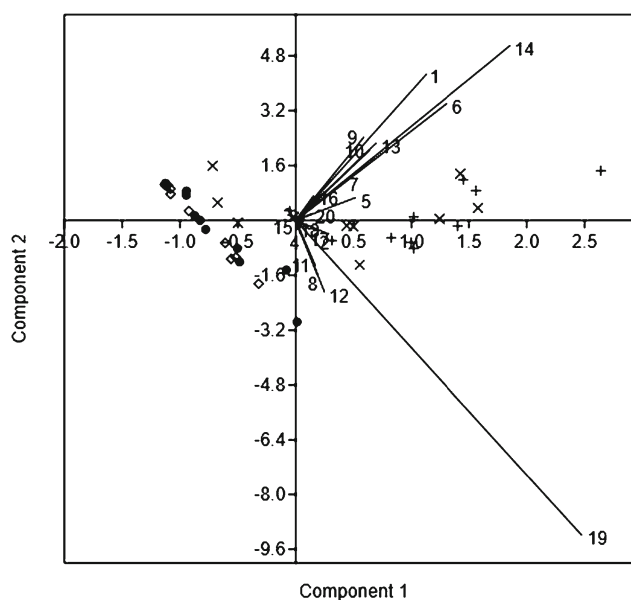
<sup>c</sup> Letters denote treatment comparisons after one-way ANOVA followed by Tukey HSD post hoc test



**Fig. 1** Typical total ion chromatograms (TIC) of volatile organic compounds (VOCs) emitted by an undamaged soybean plant of the isogenic **a** and transgenic line **b**, and a *Chrysodeixis includens*-damaged soybean plant from the isogenic **c** and transgenic line **d**. Major VOCs detected are: 1) (Z)-3-hexen-1-ol; 2) (Z)-3-hexenyl acetate; 3) (Z)- $\beta$ -ocimene; 4) unidentified compound 2; 5) indole; and 6)  $\alpha$ -farnesene. The X denotes contaminants in the sample. IS denotes the internal standard. Intensities are not normalized for dry weight of the plants

( $X^2 = 9.00$ ;  $P = 0.003$ ) and the transgenic line ( $X^2 = 12.10$ ;  $P < 0.001$ ). However, when we compared the odors from herbivore-induced plants of both lines, *M. rubens* preferred the odors induced in transgenic plants ( $X^2 = 10.53$ ;  $P = 0.001$ ) (Fig. 3).

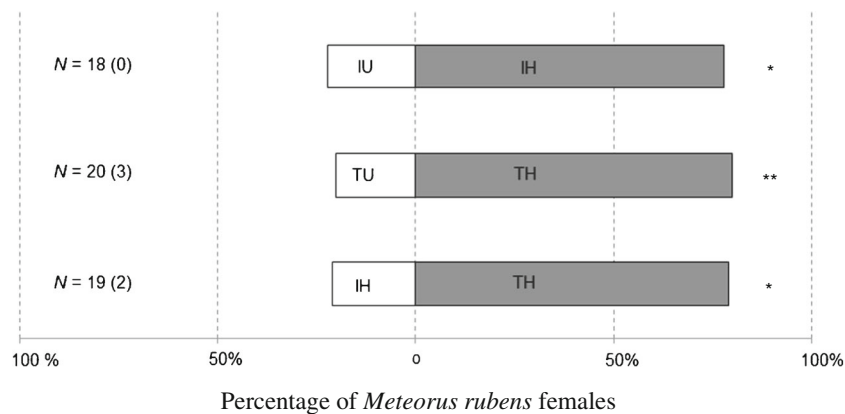
**Feeding Performance of the Soybean Looper** There was no difference between the weight increase of larvae fed on leaves from transgenic (Mean  $\pm$  SEM,  $11.36 \pm 1.80$ ) and leaves from conventional (Mean  $\pm$  SEM,  $8.59 \pm 1.20$ ) plants over one week ( $T = 1.135$ ;  $P > 0.05$ ).



**Fig. 2** Scores of individual plants from undamaged (●) and herbivore-damaged soybean plants (x) from the isogenic line, undamaged plants (◇) and herbivore-damaged plants (+) from the transgenic line, and volatile organic compounds (VOCs) projected on the first (PC1) and second principal components (PC2). Vectors show the loading of each (numbered) compound and the length represents the relative magnitude of its contribution to the different treatments. Analyzed VOCs: 1) (Z)-3-hexen-1-ol; 2) 1-octen-3-ol; 3) 3-octanone; 4) octanal; 5) (Z)-3-hexenyl acetate; 6) (Z)- $\beta$ -ocimene; 7)  $\beta$ -linalool; 8) nonanal; 9) unidentified compound 1; 10) unidentified compound 2; 11) methyl salicylate; 12) decanal; 13) (Z)-3-hexenyl 2-methylbutyrate; 14) indole; 15) unidentified compound 3; 16) (Z)-jasnone; 17)  $\beta$ -caryophyllene; 18)  $\alpha$ -caryophyllene; 19)  $\alpha$ -farnesene; and 20) TMTT

## Discussion

There is evidence that genetic transformation of plants to express the delta endotoxins of *Bacillus thuringiensis* (Bt) or resist the pathogen *Venturia inaequalis* can result in variation of HIPVs compared to conventional plants (Dean and De Moraes 2006; Turlings et al. 2005; Vogler et al. 2010). Studies using Bt-plants suggest that changes in VOC profiles may be the result of carbon allocation due to genetic transformation (Turlings et al. 2005), in addition to expected altered feeding patterns as a result of the Bt-toxin (Dean and De Moraes 2006; Turlings et al. 2005). Here, we demonstrated that, under controlled conditions of light and temperature, the emissions of VOCs induced in response to herbivory by the soybean looper are larger in transgenic plants expressing resistance to glyphosate than in non-transgenic plants. These results are in agreement with a previous study that showed that the emissions of VOCs in response to herbivory by the velvet bean moth are larger in transgenic RR soybean plants than in the isogenic line (Strapasson et al. 2016). Since the same soybean lines were used in both experiments, variation in the magnitude of individual compounds induced by the two species may be due to distinctive feeding behaviors and/or



**Fig. 3** Choice of mated *Meteorus rubens* females toward the odors from undamaged (U) and *Chrysodeixis includens*-damaged plants (H) in the transgenic (T) and isogenic line (I), and between the odors of herbivore-damaged plants emitted by both lines. The asterisks denote the level of

significance ( $P < 0.01$  \*,  $P < 0.001$  \*\*). The numbers on the left side of the graph show the total number of insects that responded to any of the odors ( $N$ ); the number of insects that did not respond is shown in brackets

different oral elicitors present in the velvet bean moth and the soybean looper (Erb et al. 2012 and references therein). However, the question of why transgenic plants that express resistance to glyphosate emit greater amounts of VOCs in response to herbivory by two lepidopteran larvae remains unanswered. Considering the fact that transgenic RR plants express a variant of the 5-enolpyruvylshikimate-3-phosphate synthase (EPSPS), an enzyme that participates in the biosynthesis of tryptophan, tyrosine, and L-phenylalanine via the shikimic acid pathway, a possible explanation would be that the insensitive EPSPS altered the synthesis of VOC precursors. This may be the case of indole, a tryptophan-derived volatile, whose emission is directly related to the activity of EPSPS (D'Alessandro et al. 2006). However, this may not explain why transgenic plants emitted larger amounts of terpenoids, which are synthesized from isopentenyl diphosphate and its allylic isomer via the mevalonic acid (MVA) pathway and the methyl erythritol phosphate (MEP) pathway (Dudareva et al. 2006).

A central event in the activation of plant responses to abiotic and biotic stress is the accumulation of phytohormones, mainly jasmonic acid (JA), salicylic acid (SA), and ethylene (ET) (Bari and Jones 2009). From these, JA is the main signaling molecule responsible for triggering induced responses (including HIPV emissions) in response to attack by mites and chewing insects such as lepidopteran caterpillars (Arimura et al. 2009; Menzel et al. 2014 for a review). Expression of JA-dependent genes relies, however, on a positive or negative interaction (cross-talk) between signaling pathways involved in plant defense (Pieterse et al. 2012). Cross-talk between SA and JA has been widely studied. Usually, accumulation of SA reduces the accumulation of JA and vice-versa, even though positive and neutral cross-talk between these two phytohormones also has been reported (Agrawal et al. 2014; Pieterse et al. 2012 and references therein). Considering the fact that the aromatic amino acid L-phenylalanine is precursor of SA

(Dudareva et al. 2006), a second explanation might be that genetic transformation somehow affected the cross-talk between signaling pathways and, in turn, JA-dependent induced responses. This effect on cross-talk may have directly affected induction of volatiles or another trait that directly affected herbivore feeding behavior, which in turn altered the emissions of volatile compounds.

Herbivore-induced plant volatiles are reliable cues for parasitoids (see Clavijo McCormick et al. 2012 for a review), which can cope with natural genetic variation in plant VOC bouquets (Kappers et al. 2011). Our results from the bioassays showed that *M. rubens* is able to discriminate between herbivore-induced and constitutively-emitted volatiles in both transgenic and conventional lines. This is not surprising, as herbivory in both lines resulted in clear qualitative and quantitative changes compared with the constitutive VOC blend. However, when *M. rubens* was challenged to discriminate between the HIPVs emitted by the conventional line and those emitted by the transgenic line, there was a clear preference of the wasps for the odors from the transgenic line. These results suggest a shift in the preference of the wasps, and that tritrophic interactions are affected in multiple ways, in this model system at least. VOC-mediated plant-insect interactions are complex, and the preference of natural enemies for particular genotypes within one plant species may be due to the abundance of individual compounds or changes in VOC ratios (Bruce et al. 2005). That is, natural enemies can respond differently to HIPVs depending on the amount of individual or total VOCs emitted by the plant (Girling et al. 2011; Horiuchi et al. 2003; Shiojiri et al. 2012). For example, the predatory mite, *Phytoseiulus persimilis* is more attracted to the odors emitted by cucumber plants infested with its prey, *Tetranychus urticae*, in varieties that emit larger quantities of (*E*)- $\beta$ -ocimene, TMTT, and two other unknown compounds (Kappers et al. 2011). In our study, when we compared different groups of plants, the volatile emissions in response to

herbivory from transgenic plants differed from the other three groups. Clearly, herbivore-damaged transgenic plants emitted larger quantities of  $\alpha$ -farnesene, indole, (*Z*)- $\beta$ -ocimene, and (*Z*)-3-hexen-1-ol. Therefore, our results suggest that changes in the concentrations of a few compounds may have modified the preference of the parasitic wasps. To our knowledge, this is the first study assessing the behavioral response of this Neotropical parasitoid to plant volatiles. Nothing is known about the innate response of *M. rubens* or its ability for associative learning. Associative learning is an important process in the foraging behavior of parasitoids, and their ability to learn allows them to distinguish between specific volatiles (De Jong and Kaiser 1991). Specialist parasitoids depend strongly on host-derived stimuli and particular plant VOC compounds that evoke invariable responses, but are likely influenced by associative learning (Clavijo McCormick et al. 2012). In contrast, generalist parasitoids such as *M. rubens* that must cope with variable VOC bouquets, benefit from associative learning (Ngumbi et al. 2012). Nevertheless, responses of parasitoids to volatiles may depend not only on the degree of specialization of the parasitoid, but also on the herbivore. Since we used naïve individuals in our experiments, further studies addressing associative learning would be interesting to understand better the behavior of *M. rubens*. Host traits can affect parasitoid performance. For example, host weight can influence parasitoid development (Hardy et al. 1992). However, we did not find any evidence that feeding on leaves of the transgenic line can affect weight gain of the soybean looper. These results suggest that the performance of wasps that develop in larvae fed on RR plants may not be negatively affected. However, as we did not follow the development of the parasitoid, these results are not conclusive and deserve further investigation.

Our results suggest that, in the studied model system, VOC-mediated tritrophic interactions are not negatively affected in plants engineered to express resistance to glyphosate. However, the shift in the preference of the wasps towards herbivore-induced odors from RR transgenic plants suggests that tritrophic interactions are altered. In addition to tritrophic interactions, altered volatile emissions may affect other ecological interactions mediated by airborne signal molecules. Indole, for instance, which was emitted in much larger quantities by damaged transgenic plants, is a signal molecule for priming maize plants (Erb et al. 2015) and can directly affect herbivore performance and behavior (Veyrat et al. 2016). Since genetic expression within and between conventional and transgenic lines are likely to exist (Cheng et al. 2008) further studies need to be conducted with other lines.

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