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Infuence of Host Plants and Tending Ants on the Cuticular Hydrocarbon Profle of a Generalist Myrmecophilous Caterpillar

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Abstract

In myrmecophilous organisms, which live in symbiosis with ants, cuticular hydrocarbons (CHCs) play a pivotal role in interspecifc communication and defense against chemical-oriented predators. Although these interactions form complex information webs, little is known about the infuence of biotic environmental factors on the CHC profles of myrmecophiles. Here, we analyzed the efect of diferent host plants and tending ants on the larval CHC profle of *Synargis calyce* (Lepidoptera: Riodinidae), a polyphagous species with facultative myrmecophily. Groups of caterpillars were fed individually with three host plant species (without tending ants), and with two tending ant species. Through gas chromatography analysis, we compared the cuticular profles of treatments and found a high similarity between plants and caterpillars (65–82%), but a low similarity between caterpillars and their tending ants (30−25%). Cluster analysis showed that caterpillars, ants, and plants form distinct groups, indicating that *S. calyce* caterpillars have their own chemical profle. These results are similar to those observed for Lycaenidae caterpillars indicating that there is functional convergence in the chemical strategies used by myrmecophilous caterpillar species with similar ecology. Also, the results suggest that the cuticular compounds of *S. calyce* are primarily infuenced by their host plants rather than their tending ants. Thus, we propose that these caterpillars present a trade-of between camoufage and directly informing their presence to ants, maintaining their unique chemical profle, though slightly afected by biotic environmental factors.

Keywords Chemical compounds · Mutualistic interactions · Myrmecophilous butterfies · Riodinidae · *Synargis calyce*

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Introduction

Ants are predominant in many terrestrial ecosystems in terms of abundance and biomass (von Beeren et al. [2012](#page-14-0)). These eusocial insects live in complex societies where communication plays a crucial role in their functioning (Hölldobler and Wilson [1990](#page-12-0); Yamaoka [1990](#page-14-1); Lenoir et al. [1999;](#page-13-0) Akino [2008](#page-12-1)). In ants and other social insects, communication and particularly the recognition of nestmates is primarily based on chemical cues and signals (Yamaoka [1990](#page-14-1); Lenoir et al. [1999;](#page-13-0) Akino [2008](#page-12-1); Blomquist and Bagnères [2010;](#page-12-2) Nunes et al. [2014](#page-13-1)). Cuticular hydrocarbons (CHCs), the main class of cuticular compounds in ants, are colony-specifc and actively participate in nestmate recognition (Howard and Blomquist [2005;](#page-12-3) Hefetz [2007](#page-12-4); Ferguson et al. [2021\)](#page-12-5). The ecological success, wide distribution, and social organization of ants have contributed to the evolution of diverse associations with organisms from various kingdoms, including plants, fungi, and animals, particularly other insects (Casacci et al. [2019\)](#page-12-6). Organisms that interact with ants during at least part of their lifecycle are called myrmecophiles (Hölldobler and Wilson [1990;](#page-12-0) Kronauer and Pierce [2011;](#page-13-2) Hölldobler and Kwapich [2022\)](#page-12-7). There are about 10,000 species of myrmecophiles across various insect orders (Thomas et al. [2005](#page-14-2); Parker [2016](#page-13-3); Hölldobler and Kwapich [2022\)](#page-12-7).

Myrmecophily in Lepidoptera is primarily observed in the families Lycaenidae and Riodinidae, with 75% of the species in these families having immatures stages that interact with ants (Pierce et al. [2002;](#page-13-4) Casacci et al. [2019](#page-12-6)). These families commonly exhibit facultative and unspecifc relationships, involving interactions with various ant taxa. However, there are butterfy species that establish obligatory and specifc relationships with specifc ant taxa (Fiedler [1994](#page-12-8), [2021](#page-12-9); Kaminski [2008;](#page-13-5) Pierce and Dankowicz [2022](#page-13-6)). Interactions between butterfies and ants can range from mutually benefcial outcomes, such as mutualism, to interactions where butterfies beneft without harming ants in commensalism, and to antagonist interactions where butterfies may be preyed by ants or where ants may be harmed by butterfies, as seen in social parasitism and competition for resources (Fiedler [1995](#page-12-10), [1996](#page-12-11); Pierce and Dankowicz [2022\)](#page-13-6). Both lycaenid and riodinid species exhibit various adaptations resulting from the pressures exerted by their association with ants (Pierce et al. [2002\)](#page-13-4). These adaptations include highly specialized ant-organs involved in chemical and acoustic deception (Cottrell [1984](#page-12-12); Fiedler et al. [1996](#page-12-13); Pierce et al. [2002](#page-13-4); Elmes et al. [2002](#page-12-14); Barbero et al. [2012](#page-12-15); Hill et al. [2022\)](#page-12-16). For instance, nectary organs such as the dorsal nectary organ (DNO) in Lycaenidae and tentacular nectary organs (TNOs) in Riodinidae liquid secretions rich in sugar and amino acids (Newcomer [1912;](#page-13-7) Malicky [1970](#page-13-8); DeVries [1988](#page-12-17)). Caterpillars of these two families are also equipped with a pair tentacular organs (TOs) in Lycaenidae and anterior tentacular organs (ATOs) in Riodinidae, which emit chemical signals or vibroacoustic signals that modify the ant behavior (Henning [1983;](#page-12-18) DeVries et al. [1986](#page-12-19); DeVries [1988](#page-12-17); Gnatzy et al. [2017;](#page-12-20) Schönrogge et al. [2017](#page-13-9)).

Chemical strategies mediated by cuticular compounds enable myrmecophiles to overcome the barrier of chemical recognition employed by ants (von Beeren et al. [2012](#page-14-0)). One such strategy is chemical camoufage, where organisms resemble their background and avoid detection by chemically oriented predators (Silveira et al. [2010](#page-13-10)). In herbivorous organisms, achieving chemical camoufage with their host plants is possible through diet (Espelie et al. [1991](#page-12-21); Barbero [2016;](#page-12-22) Lima et al. [2024](#page-13-11)). In Lepidoptera, this strategy has already been demonstrated in both non-myrmecophilous and myrmecophilous caterpillars (Akino et al. [2004](#page-12-23); Portugal and Trigo [2005;](#page-13-12) Lima et al. [2021](#page-13-13)). One of the most extensively studied strategies is chemical mimicry, where organisms possess chemical compounds that close resemble those of other organisms such as ants. This strategy has been observed in social parasitic species (Akino et al. [1999](#page-12-24); Elmes et al. [2002](#page-12-14); Schlick-Steiner et al. [2004;](#page-13-14) Schönrogge et al. [2004;](#page-13-15) Akino [2008\)](#page-12-1). Additionally, some myrmecophilous caterpillars present low concentration of compounds on their surface, making their detection challenging - a strategy known as chemical insignifcance (Inui et al. [2015;](#page-13-16) Barbero [2016](#page-12-22)). Recently, a new strategy has been proposed in myrmecophilous caterpillars called chemical conspicuousness. In this strategy, caterpillars that do not provide caloric rewards for ants exhibit a distinct cuticular profle compared to ants or host plants. However, their profle is similar to that of other caterpillars that offer caloric rewards to ants (Lima et al. [2021](#page-13-13)).

Although ant-plant-herbivore systems embrace complex communication networks involving multiple species (e.g., Lima et al. [2021](#page-13-13), [2023\)](#page-13-17), there is still a lack of information regarding the infuence of phenotypic plasticity and biotic environmental factors on the cuticle compounds of generalist myrmecophilous species (Otte et al. [2018;](#page-13-18) Sprenger and Menzel [2020\)](#page-13-19). Therefore, our aim was to investigate whether the CHC profle of a polyphagous caterpillar changes (1) according to their diet on diferent host plant species and (2) according to their interaction with diferent tending ant species. Considering that the caterpillar species used here were fed on their host plants, we hypothesized that their CHCs would be infuenced primarily by their food source rather than the contact established with tending ants. Furthermore, due to the production of caloric rewards by caterpillars (trophobiosis) we predicted that caterpillars would exhibit a specifc chemical profle distinct of both host plant and tending ants.

Methods and Materials

Study System *Synargis calyce* C. Felder and R. Felder, 1862 (Lepidoptera: Riodinidae) (Fig. S1) is a Neotropical herbivorous and myrmecophilous butterfy whose caterpillars feed on several host plants in diferent families, including non-native species (Callaghan [1986;](#page-12-25) Beccaloni et al. [2008](#page-12-26); Alves-Silva et al. [2018](#page-12-27); Kaminski [2021\)](#page-13-20). Female butterfies tend to lay their eggs on plants that are frequented by attendant ants and caterpillars are tended by ants during all instars (Callaghan [1986](#page-12-25)). Although it presents facultative myrmecophily, the caterpillars are almost always found with tending ants of several genera, but mainly *Camponotus* species (Callaghan [1986](#page-12-25); Alves-Silva et al. [2018;](#page-12-27) Kaminski [2021](#page-13-20)). It is common to observe temporal turnover, with attendance by diferent species of ants during the day and night (LAK, personal observation). Due to the high degree of ecological plasticity of this butterfy in terms of both host plants and tending ants, it is an excellent model of a generalist myrmecophile.

Collection and Rearing of Study Species Insects were collected at the Universidade de São Paulo (21.1637° S, 47.8592° W), Ribeirão Preto Campus, SP, Brazil, between January 2021 and April 2022. To conduct our chemical analysis, we collected ~ 90 eggs of *S. calyce* for rearing in the laboratory. Initially, \sim 20 field observations were conducted to identify plant species used by butterfies for oviposition and seven species were identifed serving as host plants. Subsequently, eggs were collected from three of these host plants (*Senegalia polyphylla* (DC.) Britton and Rose (Fabaceae), *Inga laurina* (Sw.) Willd. (Fabaceae), and *Terminalia catappa* Linnaeus (Combretaceae)) at various study locations by harvesting branches where oviposition had been observed. In the feld, three ant nests of *Camponotus crassus* Mayr, 1862 (Formicidae: Formicinae) were collected and transferred to the laboratory. Each nest was placed in two connected boxes measuring $9 \times 26.6 \times 26.6$ cm. One box served as the nesting area and contained test tubes (15 cm long) flled with water, plugged with hydrophilic cotton. The other box served as the foraging area. The ant colonies were provided with a diet of *Tenebrio molitor* Linnaeus, 1758 larvae (Coleoptera: Tenebrionidae), diluted sugar solution (10%), and water *ad libitum*. Each colony consisted of approximately 150 workers, a queen, and some brood (eggs, larvae and pupae). Additionally, approximately, 600 workers of *Paratrechina longicornis* Latreille 1802 (Formicidae: Formicinae) were collected from three established colonies near the laboratory buildings. The insects were kept under controlled conditions at a temperature of 25 °C and a photoperiod of 12 h of light and 12 h of darkness.

Does the Chemical Composition of Caterpillars Change According to Their Food Sources? As the caterpillars studied here are polyphagous, we selected three host plant species on which the caterpillars feed in the study area: two native, *S. polyphylla*, and *I. laurina*, and one non-native, *T. catappa*. Our aim was to investigate whether the CHCs of the caterpillar change according to its diet in the absence of tending ants. To conduct the experiment, we placed individually each egg in a plastic container (250 ml). Once the eggs hatched, we provided the caterpillars with shoots containing young leaves and extraforal nectaries from the host plants. The shoots were replaced daily and kept in contact with moistened cotton to prevent them from drying. The *S. calyce* caterpillars were reared separately in plastic containers on three host plant species: caterpillar-*S. polyphylla* (*n*=10), caterpillar*-I. laurina* (*n*=10), caterpillar-*T. catappa* (*n*=6). After reaching the ffth instar, we killed the caterpillars by freezing and kept them at -20 °C until CHC extractions were performed. Additionally, we collected leaves of *S. polyphylla*

 $(n=10)$, *I. laurina* $(n=10)$, and *T. catappa* $(n=10)$ for CHC extractions (See Table S1).

Does the Chemical Composition of Caterpillars Change According to Their Tending Ants? In our feld site, the caterpillars are attended by four diferent ant species: *C. crassus, P. longicornis, Camponotus renggeri* Emery, 1894, and *Wasmannia auropunctata* Roger, 1863 (AVCG, personal observation). Thus, to assess whether the caterpillar CHCs change according to their tending ants, we individually reared *S. calyce* caterpillars in a plastic container as previously described, along with a group of associated ants. Specifcally, we reared the caterpillars with two experimental groups: (caterpillar-*C. crassus*) (*n*=10), and (caterpillar-*P. longicornis*) $(n = 10)$. These caterpillars were fed with the host plant *S. polyphylla*. Each caterpillar was placed together with 10 workers of *C. crassus* or 30 workers of *P. longicornis*. The number of ants was based on the average amount observed in the feld. The ant workers were replaced every day until the caterpillars were frozen for chemical extraction, as previously mentioned. For chemical analysis, we also collected worker ants from colonies reared in the laboratory for *C. crassus* ($n=8$ colonies; $n=20$ ants for chemical analyses) and from colonies established near the laboratory for *P. longicornis* ($n = 10$ colonies; $n = 300$ ants for chemical analyses) (See Table S1).

Chemical Analyses To perform the chemical analyses, we placed insects or plant shoots in glass vials (1.5 ml) and covered them with *n*-hexane (Macron Fine Chemicals, 95% *n*-hexane, USA) for 1 min (Lima et al. [2023\)](#page-13-17). For each sample, a ffth-instar larva of *S. calyce*, two workers of *C. crassus*, 30 workers of *P. longicornis*, and one young shoot with two leaves from each plant species were used individually. External standards were exclusively employed and the samples were not weighed. Subsequently, we left each vial at room temperature in a flow chamber to allow for drying. Once completely dried, we resuspended the contents in 5 µl of hexane, of which 2 µl were manually injected. The samples were analyzed with gas chromatography coupled to a mass spectrometer (GC/MS; Shimadzu, model QP2010 Plus), using a 30 m Rxi-1ms column, with helium gas flow rate set at 1 ml/min. The oven temperature was initially set to 40 °C and then increased by 3 °C min- 1 until reaching 310 °C (held for 15min), following da Silva et al. ([2021](#page-12-28)). The injector temperature was set to 250 °C. Data were analyzed by GC/MS Solutions for Windows (Shimadzu Corporation), and compounds were identifed based on their mass spectra, including diagnostic and molecular ions (Carlson et al. [1998](#page-12-29)). Additionally, a retention index was calculated for each identifed peak using a standard solution of diferent synthetic linear hydrocarbons $(n-C_{21}$ to $n-C_{40})$. We also consulted the Registry of Mass Spectral Data (Wiley) and National Institute of Standards and Technology (NIST) mass spectra search program (version 2.2) Libraries database for identifcation (Lima et al. [2023](#page-13-17)).

Statistical Analyses We used Morisita's Similarity Index (SI) which ranges from 0% (indicating no similarity) to 100% (representing complete similarity) (Krebs [1999\)](#page-13-21) to compare CHC profles of diferent groups, following the methodology of Lima et al. ([2021](#page-13-13)). This analysis was carried out using PAST software (Version 4.13) (Hammer et al. [2001\)](#page-12-30). Furthermore, to assess the overall chemical similarity or dissimilarity between groups, we performed a permutation analysis (PERMANOVA). This analysis was performed using the *adonis* function from the *vegan* package (Oksanen et al. [2013\)](#page-13-22) with 9999 permutations. In order to represent the multivariate chemical dataset and check for the cluster formation, we next performed a Principal Component Analysis (PCA). For this, we used the *prcomp* function of the *stats* package (R Core Team [2019](#page-13-23)). We also ran a multivariate similarity analysis (SIMPER) using the Bray-Curtis distance and adopting 999 permutations. The SIMPER analysis allowed us to determine the contribution of each chemical variable to the existing variation among samples. For this analysis, we used the *simper* function from the *vegan* package (Oksanen et al. [2013](#page-13-22)). For all tests, we determined the relative abundance percentages of each compound present in the cuticular extracts, treating the compounds as 100% and then analyzed the data. All these analyses were conducted using R version 4.0.2 (R Core Team [2019\)](#page-13-23).

Results

Overall Chemical Information A total of 78 peaks were identifed in the cuticular extracts from the diferent groups studied (Table [1](#page-4-0)). *Senegalia polyphylla* exhibited 22 peaks, *I. laurina* had 29 peaks, *T. catappa*, *C. crassus* had 28 peaks each, *P. longicornis* had 22 peaks, and *S. calyce* caterpillars had 25–30 peaks. These peaks corresponded to various chemical compounds, including branched hydrocarbons (mono-, di-, and trimethylated), linear alkanes, alkenes, alcohols, and aldehydes. The carbon lengths of the identifed compounds ranged from 18 to 36.

The cuticular profile of the three host plant species revealed a shared class of compounds, including linear alkanes, alcohols, and aldehydes. However, variations existed among them, particularly in the proportion and presence or absence of specifc alcohols and aldehydes. For instance, 1-hexadecanol, 1-hexacosanol, and hexacosanal were exclusively present in *I. laurina*, while 1-triacontanol acetate was found only in *T. catappa* (Table [1\)](#page-4-0). The SIs of the cuticular compounds of the plants varied according to the pairs of compared species. *Terminalia catappa* with *S. polyphylla* showed the highest SI of up to 77%, followed by *T. catappa* with *I. laurina*, which exhibited a SI of up to 65%, and *I. laurina* with *S. polyphyll*a, which showed a SI of up to 62%. *Senegalia polyphylla* and *I. laurina* shared 20 compounds, representing 69% and 91% of their respective cuticles. *Senegalia polyphylla* shared 15 compounds with *T. catappa*, representing 52% and 68.2% of their respective cuticles. *Terminalia catappa* shared 18 compounds with *I. laurina*, representing 60% and 62.06% of their respective cuticles. Although post hoc pairwise comparisons did not reveal a signifcant diference based on relative abundance or chemical composition (Table [2\)](#page-6-0), the PCA revealed that the three species form separate groups, in which the frst and the second principal component explained 30% and 13.77%, respectively. (Fig. [1](#page-7-0)). In terms of major compounds, *S. polyphylla* had the *n*-C29, 1-triacontanol and triacontanal; *I. laurina* had the *n*-C29, Hexacosanol and *n-*C31, and *T catapp*a had the *n*-C29 and n-C31.

Does the Chemical Composition of Caterpillars Change According to Their Food Sources? When comparing the compounds found in caterpillars reared on three diferent host plants, we found the following similarity indices: The cuticular profles of caterpillars reared on *T. catappa* showed a SI of up to 82% with *T. catappa*, up to 74% with *I. laurina*, and up to 61% with *S. polyphylla*. The cuticular profles of caterpillars reared on *I. laurina* showed a SI of up to 76% with *I. laurina* and *T. catappa*, and up to 60% with *S. polyphylla*. Finally, caterpillars reared on *S. polyphylla* exhibited a SI up to 65% with *S. polyphylla*, and up to 64% with *T. catappa*, and *I. laurina.* Moreover, certain compounds were exclusively found in groups of caterpillars reared on specifc plant species. For example, 1-hexadecanol was found only in the cuticular profle of caterpillars reared on *I. laurina*, and this particular compound was also identifed in the chemical profle of this plant. Similarly, 1-docosanol, identifed in *T. catappa*, was also detected in caterpillars reared on this plant but was absent in those reared on *S. polyphylla* (Table [1](#page-4-0)). The qualitative similarity varied according to the host plant. Specifcally, caterpillars shared 19 compounds with *S. polyphylla*, representing 70.37% and 86.36% of their respective cuticles. Caterpillars and *I. laurina* shared 23 compounds, representing 76.66% and 79.31% of their respective cuticles, while caterpillars and *T. catappa* shared 17 compounds, representing 58.6% and 60.7% of their respective cuticles. However, there was a low degree of similarity in the relative abundance of compounds between caterpillars and their host plants. The post hoc pairwise comparisons revealed signifcant diferences in relative abundance and in the chemical composition among caterpillars reared on diferent plants (Table [2\)](#page-6-0). However, most of the compounds were shared in the three groups and we found an overlap among caterpillars

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Fig. 1 Principal component analysis (PCA) of chemical compounds in *S. calyce* caterpillars (three groups reared on three plants species without ants and two groups reared with two ants species), the atten-

in the PCA (Fig. [1](#page-7-0)). Specifcally, caterpillar–*T. catappa* and caterpillar–*I. laurina* shared 27 compounds, representing 93.1% and 90% of their respective cuticles. Caterpillar*-T. catappa* and caterpillar*–S. polyphylla* shared 26 compounds, representing 89.7% and 92% of their respective cuticles. Caterpillar*–I. laurina* and caterpillar*–S. polyphylla* shared 25 compounds, representing 83.3% and 92.6% of their respective cuticles. In general, all caterpillar groups had *n*-alkanes (C29, C31, and C33) as their major compounds and they also showed a few methylated compounds, alcohols, and aldehydes, these last two also present in all plants.

Does the Chemical Composition of Caterpillars Change According to Their Tending Ants? The cuticular profles of caterpillars reared with *C. crassus*, or *P. longicornis* ants, as well as caterpillars reared without attendant ants, had a high SI (>90%). Moreover, the cuticular profiles of all caterpillar groups were qualitatively similar, with 24 shared compounds, representing 96% of their respective cuticles. Hence, we found an overlap among samples of caterpillars in the PCA (Fig. [1](#page-7-0)). The post hoc pairwise comparisons did not show diferentiation based on chemical composition among caterpillars reared with diferent species of attendant ants (Table [2](#page-6-0)).

In contrast, the cuticular profles of caterpillars and attendant ants showed low similarity indexes and varied according to the attendant ant (SI<30% for *C. crassus* and SI<25% for

dant ants (*C. crassus* and *P. longicornis*) and the host plants (*S. polyphylla, I. laurina*, and *T. catappa*)

P. longicornis). Caterpillars and *C. crassus* shared 14 compounds, representing 54% and 50% of their respective cuticles. Caterpillars and *P. longicornis* shared four compounds, representing 16% and 18.2% of their respective cuticles. The post hoc pairwise comparisons showed a signifcant diferentiation based on chemical composition between all groups of caterpillars and their respective attendant ants (Table [2](#page-6-0)). Moreover, the PCA revealed that ants and caterpillars form separate groups. In terms of major compounds, *C. crassus* had the *n*-C21, *n*-C30 and C33:1, and *P. longicornis* had the *n*-C29, 15-;13-;11-;9-MeC29, 15-;13-;11-MeC29 and 9.17-; 9.19-diMeC29 (Table [1](#page-4-0)).

In general, caterpillars, host plants, and attendant ants shared the *n*-C27, *n*-C29, and *n*-C31 compounds (Fig. [3](#page-9-0)). Only caterpillars and plants showed alcohols and aldehydes in their chemical profles (hexadecanol, octadecanol, eicosanol, octacosanal, docosanol, and triacontanal). The most signifcant diferentiating compounds among the groups, according to SIMPER analysis, were the n-C33, n-C31 and, *n*-C29, which were major compounds in caterpillars (Fig. [2;](#page-8-0) Table S2). Overall, we observed that all groups of caterpillars presented high similarity in composition and proportion of their cuticular compounds. Host plants and caterpillars had a higher number of compounds in common, with *n*-alkanes (*n*-C29 and *n-*C31) as their major compounds. Host plants also had various alcohols and aldehydes, which are also present in caterpillars, but in smaller proportions. The three plants exhibited qualitative

Fig. 2 Box plots of the relative abundance of the most important chemical compounds (*n*-C29, *n*-C31 and *n*-C33) that contributed to diferentiating the groups according to SIMPER. Groups 1 and 2

include attendant ants; 3–5 include host plants and 6–10 include caterpillars exposed to diferent conditions. Diferent letters represent *p*<0.05

similarities in their linear alkanes, with some variations in proportion. Additionally, there were qualitative differences in the alcohols and aldehydes among the three species. Thus, the cuticular profle of caterpillars was not infuenced by attendant ants. Ant species showed a greater diversity of compounds compared to caterpillars, including various branched alkanes, alkanes, and alkenes (Fig. [3\)](#page-9-0).

Discussion

We found that the cuticular profle of caterpillars are more similar to those of the host plants rather than to their tending ants. Moreover, caterpillars reared on diferent species of plants, without ants or in close contact with ants also had similar cuticular composition and proportion, meaning that in an overall perspective caterpillar cuticular composition is weakly afected by exogenous factors. However, some of

Fig. 3 Comparison of CHC profles among host plant, myrmecophilous caterpillar, and attendant ants. All compound identities can be found in Table [1](#page-4-0)

their compounds varied depending on the host plants where they were reared, which suggests that they acquired part of these compounds through their diet. Thus, *S. calyce* caterpillars have a chemical profle that it is slightly altered by their food. Consequently, our hypothesis that the caterpillar cuticular composition is infuenced by their food source was partially corroborated. Even though the caterpillars are not chemically identical to their host plants, there is a higher similarity between caterpillar-plant when compared to caterpillar-ant species. Given that some compounds, such as 1-hectacontanol, are present in one of the host plants and in caterpillars reared on that plant, but not in the other groups of caterpillars, this may suggest that at least part of the caterpillar chemical composition should derive from their food source. In this way, we suggest that the chemical composition of caterpillars is mainly genetically derived and slightly infuenced by the environment. Acquiring compounds from host plants through diet and the usage of them when interacting with ants stands out as a promising strategy and it has also been observed in other plant-herbivorous insect systems (e.g., Silveira et al. [2010;](#page-13-10) Lima et al. [2021,](#page-13-13) [2024\)](#page-13-11).

The three plant species used to feed the caterpillars exhibit qualitative similarities in their linear alkanes, as well as in certain alcohols and aldehydes. This suggests that *S. calyce* females likely tend to lay eggs on plants with similar chemical profles, a similar pattern was observed in lycaenids (Lima et al. [2021](#page-13-13)). There is evidence that alkanes and alcohols can serve as signals for host plant selection (Li and Ishikawa [2006;](#page-13-24) Barbero [2016;](#page-12-22) Bertea et al. [2020](#page-12-31)). In a recent study involving *S. calyce*, it was observed that butterfies sometimes mistakenly lay eggs directly on treehoppers because the treehoppers have a cuticular profle similar to that of the host plant (Lima et al. [2023](#page-13-17)). The compounds found on the surface of leaves have been described to play a role in the chemical defense of plants (aliphatic hydrocarbons, fatty and phenolic acids derivatives) (Martemyanov et al. [2015](#page-13-25): Bertea et al. [2020](#page-12-31)). Hence, there is a possibility that an evolutionary arms race between caterpillars and plants has driven caterpillars to develop mechanisms countering the chemical defenses of plants. This may involve detoxifcation of compounds through enzymes or the sequestration of such compounds. Consequently, caterpillars may exhibit a preference for specifc chemical compounds present in various host plants (Zu et al. [2020\)](#page-14-3).

Through chemical analyses, we found that the chemical profle of caterpillars were not infuenced by their attendant ants thus confrming our second hypothesis that *S. calyce* CHCs are not afected by the interactions that they establish with diferent ant species. The two ant species studied showed distinct chemical profles, while the groups of *S. calyce* caterpillars reared with diferent ant species or without ants had similar cuticular profles composition and proportion. This indicates that *S. calyce* caterpillars have cuticular profles that are independent of their attendant ants, indicating that they do not use chemical mimicry as a strategy when interacting with them. This fnding is supported by the dissimilarity in chemical composition between caterpillars and ants, compared to the similarity between caterpillars and plants. Additionally, *S. calyce* caterpillars have facultative association with several species of ants, and they do not exploit ant nests (Callaghan [1986;](#page-12-25) Alves-Silva et al. [2018](#page-12-27); Kaminski [2021\)](#page-13-20). Previous studies demonstrating chemical mimicry between caterpillars and ants have typically involved obligate interactions with a few specifc ant species, where the CHCs of caterpillars closely resemble those of the ants (Henning [1983](#page-12-18); Elmes et al. [1991](#page-12-32), [2002](#page-12-14); Dettner and Liepert [1994;](#page-12-33) Akino et al. [1999](#page-12-24); Schönrogge et al. [2004](#page-13-15); Hojo et al. [2009,](#page-12-34) [2014](#page-12-35); Thomas et al. [2013;](#page-14-4) Witek et al. [2013;](#page-14-5) Barbero [2016](#page-12-22); Casacci et al. [2019\)](#page-12-6). Chemical mimicry with attendant ants is an efective strategy for social parasitic caterpillars, as they typically inhabit ant nests and beneft from being perceived as members of the colony, allowing them to exploit valuable resources within the nest such as ant larvae or trophallaxis (Fiedler [1991;](#page-12-36) Barbero [2016;](#page-12-22) Casacci et al. [2019](#page-12-6)). We also ruled out the possibility of a chemical insignifcance strategy in *S. calyce* caterpillars as in previous studies where caterpillars and pupae used chemical insignifcance, their cuticular profle consisted of only a few hydrocarbons in very small proportions (Lohman et al. [2006;](#page-13-26) Inui et al. [2015](#page-13-16)). In contrast, *S. calyce* caterpillars showed 27–30 cuticular compounds, with some of them in high proportions.

Thus, it seems that this species maintains its own chemical profle in a chemical conspicuousness strategy (*sensu* Lima et al. [2021\)](#page-13-13). This strategy is the most likely since the chemical profles of caterpillars exposed to various conditions remained mainly unchanged, with all groups exhibiting a high degree of similarity. Chemical conspicuousness becomes advantageous when interacting with attendant ants, increasing the likelihood that ants will associate the reward with specific cuticular profiles (Hojo et al. [2014](#page-12-35)). Studies conducted with Neotropical Lycaenidae species have demonstrated that these species possess conspicuous chemical profles (Lima et al. [2021](#page-13-13)). Additionally, there are studies showing that caterpillars or pupae of facultative lycaenid butterfies from other regions have unique chemical profles recognized by ants, which helps maintain their attending behavior (Ômura et al. [2009](#page-13-27), [2012](#page-13-28); Hojo et al. [2014](#page-12-35); Mizuno et al. [2018\)](#page-13-29). However, to our knowledge, this is the frst study to explore cuticular hydrocarbons (CHCs) and investigate chemical strategies in a facultative species of the Riodinidae family.

In this study, caterpillars and plants shared several compounds when compared to ants. Notably, linear alkanes such as C29 and C31 are present in significant proportions across all groups, along with some alcohols and aldehydes.

Therefore, we cannot rule out the possibility that in certain instances, caterpillars may employ chemical camoufage. Moreover, caterpillars reared on one host plant species exhibited a similarity exceeding 80%, which has been previously demonstrated as sufficient to serve as chemical camoufage strategy in other insect groups (Silveira et al. [2010](#page-13-10)). Host plants play a crucial role in providing an efective background for herbivorous organisms, allowing them to avoid detection by visually or chemically resembling their surroundings. This strategy is observed in various organisms that have close relationships with ants, enabling them to interact with ants without being attacked (von Beeren et al. [2012;](#page-14-0) Barbero [2016](#page-12-22); Lima et al. [2024\)](#page-13-11). Consequently, we propose that these caterpillars may employ a trade-of strategy between camoufage and informing their presence to ants, which could vary depending on the presence of predators or mutualist ants. Given the presence of numerous non-attendant ant species that visit the host plants, many of which have extraforal nectaries, it is highly likely that chemical camouflage has been selected as an efficient strategy for *S. calyce* caterpillars to avoid attacks from diferent ants (Akino et al. [2004\)](#page-12-23). On the other hand, informing their presence to attendant ants also is efficient for the caterpillars. The complexity of CHCs profles is well-known, and it is likely that each chemical trait serves a distinct function (Sprenger and Menzel [2020](#page-13-19)). There may even be conficts or trade-ofs among the various functions of the chemical profle (Steiger and Stökl [2014](#page-14-6); Ingleby [2015](#page-13-30)). Camoufage might be the protagonist during encounters with non-attendant ants, while directly informing their presence becomes more prominent with attendant ants, as it aims to induce ants to associate caterpillars' chemical profle with the chemical reward, consequently securing ant protection (Hojo et al. [2014](#page-12-35)).

We suggest that specifc compounds such as some aldehydes present in both plants and caterpillars or methylated alkanes present in caterpillars, play a role in making the caterpillars blend with the background or informing their presence to attendant ants respectively. For instance, in studies conducted on *Lycaeides argyrognomon* (Bergsträsser, 1779) (Lepidoptera: Lycaenidae), pupal cuticular lipids were found to contain various long-chain aliphatics aldehydes, including 1-octacosanal and 1-triacontanal, which were found to suppress ant aggression (Mizuno et al. [2018\)](#page-13-29). Interestingly, these two compounds were found in all groups of *S. calyce* caterpillars and host plants, suggesting their potential importance in the interaction between *S. calyce* caterpillars and ants.

Studying tri-trophic relationships can present challenges when analyzing the results. For instance, in the PCA results, we observed a low percentage for PC1 and PC2. We suspect that these low percentages values may be linked to the number of groups included in the analysis, comprising five caterpillars, three plants, and two ant groups. The low existent variation within each main group (e.g. ant, caterpillar, and plant) may have contributed to an overall lower dissimilarity percentage when comparing all the groups at once. Thus, to elucidate our fndings, we employed more than one type of analysis. Using multiple approaches to analyze the data stands out as a useful strategy when working with complex systems.

Ants exert strong selection pressure on myrmecophilous caterpillars, leading to the development of multimodal adaptations (Pierce and Dankowicz [2022](#page-13-6); Marquis and Koptur [2022](#page-13-31)). These adaptations include morphophysiological, behavioral, chemical, and acoustic traits that caterpillars utilize to deceive, attract, alarm, or appease attending ants (Fiedler et al. [1996;](#page-12-13) Casacci et al. [2019](#page-12-6)). *Synargis calyce* caterpillars possess functional TNOs, which have been shown to contribute to the association with ants in other riodinids (DeVries [1988](#page-12-17); Kaminski and Carvalho-Filho [2012](#page-13-32); Kaminski et al. [2013;](#page-13-33) Mota et al. [2020](#page-13-34); Kaminski et al. [2021](#page-13-35)). To further enhance our understanding of the multimodal signaling in myrmecophile systems, future studies should conduct behavioral assays to experimentally confrm the chemical strategy employed and investigate the products and effects of caterpillars' ant-organs on attending ants, as well as compare them with the products of extraftoral nectaries from plants. This research will contribute to unraveling the specifc role of these chemical strategies and organs in the complex interactions between myrmecophilous caterpillars and ants.

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Author Contributions AVCG and LAK conceived, designed and conducted feld experiments. AVCG, RCS, ICCT, FSN and NPL conducted chemical analysis. AVCG, RCS and LDL conducted data analysis. AVCG and RCS wrote the frst version of the manuscript. All the authors read, wrote the fnal version and approved the fnal version of the manuscript.

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Data Availability Data was deposited in the supplementary section.

Declarations

Competing Interests The authors declare no competing interests.

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