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## **RESEARCH ARTICLE**



## Additive effects of plant chemotype, mutualistic ants and predators on aphid performance and survival

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## Abstract

- 1. Cascading effects in ecological systems acting across three or more trophic levels can be either of a resource-based (bottom-up) or natural enemy-based (top-down) nature. But, due to their complexity, these effects are often considered separately and their relative strength, acting simultaneously, remains unknown.
- 2. In a semi-natural field experiment using tansy (Tanacetum vulgare L.) and the specialised tansy-aphid Metopeurum fuscoviride Stroyan as a model system, we compared the effects of four distinct plant chemotypes (i.e., bottom-up), defined by the bouquet of their volatile terpenoids, on aphid population dynamics by manipulating the presence/absence of mutualistic ants and presence/absence of naturally occurring predators (i.e., top-down).
- 3. Predators reduced aphid abundance and colony survival but did not reduce initial growth rate due to a time-lag until predators arrived on the plants. Ants directly benefited initial aphid growth rates and abundance, even in the absence of predators, but not the number of days an aphid colony persisted on the plant.
- 4. Plant chemotype directly affected aphid growth rate and final abundances across the different plants and indirectly affected the abundances of tending ants and predators through effects on aphids. We found that tending ants were more abundant on one plant chemotype. Although ant abundance did not affect aphid population development, it became clear that ants had a preference towards aphids on certain chemotypes. However, a higher number of predators led to a lower number of aphids.
- 5. The results confirm the importance of plant chemical variation, acting through multiple effects on many species in arthropod communities, and support results from field studies. In a natural population, with a diverse selection of host-plant variants, aphid populations and their interacting species can therefore be structured at the level of an individual plant. Specialist aphids on patchily distributed host plants can exhibit metacommunity dynamics at very local scales. Plant within-species variation within a local population is often ignored in metacommunity ecology, yet our work shows that this can have strong effects on insectant-natural enemy dynamics, and therefore, future research should incorporate this into current theory and experimental studies.

#### KEYWORDS

bottom-up, exclusion experiment, Lasius niger, Metopeurum fuscoviride, mutualism, relative effects, top-down, tritrophic interaction

## 1 | INTRODUCTION

Herbivore populations and communities are considered to be regulated by bottom-up (resource-based) and top-down (e.g., predation) processes (Forkner & Hunter, 2000; Hanley & La Pierre, 2015; Hunter & Price, 1992; Kareiva & Sahakian, 1990; Ode, 2006; Walker & Jones, 2001; Wimp & Whitham, 2001). The relative importance of each process often depends on the developmental stage of the herbivore; for example, larvae and adults can experience a different set of selection pressures (Walker & Jones, 2001). Many research papers investigated such dynamics from a single perspective only, or emphasised only one of the two driving forces, hence limiting our understanding of population and community dynamics in terrestrial systems (Walker & Jones, 2001).

Top-down control by predators is assumed to play a crucial role in regulating herbivore populations, by negatively affecting their densities via direct predator-prey interactions (Schmitz, Hambäck, & Beckerman, 2000), or indirectly by affecting the prey's behaviour (e.g., predator avoidance behaviour; Clegg & Barlow, 1982). This can alter herbivore population growth, distribution (Roitberg, Myers, & Frazer, 1979) and stabilise associated communities (Halaj & Wise, 2001).

As herbivores directly depend on the availability and quality of their host plants (i.e., food source), variation in plant traits (e.g., nutritional quality, genotype or phenotype) can also directly influence herbivore population growth and distribution among host plants (bottom-up effects; e.g., Awmack & Leather, 2002; Dixon, Chambers, & Dharma, 1982; Johnson, 2008). In natural systems, it is unlikely that either bottom-up or top-down forces act in isolation, rather there is a combination of these forces (Hunter & Price, 1992). For instance, predators can be influenced by the direct influence of plants on herbivore densities that then affects predator density (density-mediated interactions; Bailey, Wooley, Lindroth, & Whitham, 2006), or plants can alter herbivore traits that change their susceptibility to predation (trait-mediated interactions; e.g., by sequestering plant secondary compounds; Brower, Zandt Brower, & Corvino, 1967; Mooney & Singer, 2012). Furthermore, prey can be indirectly affected by host-plant variation through direct effects of plants on predator density (Gassmann & Hare, 2005; Poelman et al., 2009). Plants have also been shown to directly influence predator traits, for example, the availability of plant extrafloral nectar can increase the longevity and searching efficiency of parasitoid wasps (Siekmann, Tenhumberg, & Keller, 2001). These mechanisms can indirectly influence herbivore abundances (density-mediated or traitmediated indirect effects on herbivores; Mooney & Singer, 2012). This means that bottom-up effects can affect herbivore populations not just directly but also indirectly, via effects of plant variation on

predator populations, thus changing the outcome of species interactions (Johnson, 2008; Johnson & Agrawal, 2005; Whitham et al., 2006, 2012 ; Williams & Avakian, 2015; Wimp & Whitham, 2001; Zytynska & Weisser, 2016).

Bottom-up effects mediated by intraspecific variation among plants can arise through various genetically based traits leading, for example, to differences in plant growth habit or metabolic diversity (Bálint et al., 2016; Johnson, 2008; Johnson & Agrawal, 2005; Kareiva & Sahakian, 1990; Mooney & Agrawal, 2008; Williams & Avakian, 2015; Züst & Agrawal, 2017; Zytynska & Weisser, 2016). Host-plant biochemistry is a key factor in affecting herbivore performance and often mediates herbivore preferences (Bernays & Chapman, 1994; Karban & Baldwin, 1997; Rosenthal & Berenbaum, 1992). One extensively studied type of variation in plant chemical diversity is the metabolic variation in secondary metabolites such as volatile organic compounds (VOCs; Azam et al., 2013; Eller, de Gouw, Graus, & Monson, 2012; Holopainen & Blande, 2012; Keskitalo, Pehu, & Simon, 2001; Lee, Sugawara, Yokoi, & Takahata, 2010). Plant VOCs can either be stored in specific morphological structures, constitutively synthesised and emitted (Clancy, Zytynska, Senft, Weisser, & Schnitzler, 2016) or emitted after induction by abiotic or biotic stress (Holopainen & Gershenzon, 2010). Plants use these volatile compounds for direct defence (Martin & Bohlmann, 2005) or for internal, intra- or interspecific communication (e.g., Riedlmeier et al., 2017) as well as for communicating with higher trophic levels (reviewed in de Vos & Jander, 2010; Holopainen & Blande, 2012; Paré & Tumlinson, 1999). One example is the recruitment of predators or parasitoids by herbivore-infested plants (plant-natural enemy-herbivores; e.g., Bálint et al., 2016; Linhart, Keefover-Ring, Mooney, Breland, & Thompson, 2005; Ninkovic, Al Abassi, & Pettersson, 2001). Some herbivore species (Erb & Robert, 2016; Goodey, Florance, Smirnoff, & Hodgson, 2015; Opitz & Müller, 2009; Prudic, Khera, Sólyom, & Timmermann, 2007) have also evolved to take advantage of host-plant-derived secondary metabolites (including nonvolatile defensive compounds, e.g., salicin derivatives or glucosinolates, and volatile defensive compounds, e.g., benzaldehyde) to use them in their own defence strategies against predation (Dyer, 1995; Gauld, Gaston, & Janzen, 1992). Thus, plant within-species variation in the abundance and composition of secondary metabolites, like VOCs, leading to so-called different plant chemotypes (i.e., a group of plants with similar chemical profiles; Clancy et al., 2016; Ghirardo, Heller, Fladung, Schnitzler, & Schroeder, 2012; Holopainen, Hiltunen, & von Schantz, 1987; Keefover-Ring, Thompson, & Linhart, 2009), can have multiple effects on herbivore populations and the associated arthropod community.

Plant-aphid-predator systems are ideal for simultaneously studying bottom-up and top-down effects. Aphids (Hemiptera;

Aphididae) are specialised insects feeding on the phloem sap of particular plants. Due to the highly specific nature of this interaction, plant within-species variation can have dramatic effects on the plant-aphid relationship. For instance, variation among plant genotypes or chemotypes (e.g., varying in camphor,  $\beta$ -pinene and linalool) can directly affect aphid performance (Kleine & Muller, 2011; Krauss et al., 2007; Linhart et al., 2005; Utsumi, Ando, Craig, & Ohgushi, 2011; Williams & Avakian, 2015). Furthermore, aphids are prey to a number of specialised and generalised predators, able to control aphid populations in a top-down manner (reviewed in Diehl, Sereda, Wolters, & Birkhofer, 2013).

Many aphid species are able to establish mutualistic relationships with ants. These plant-aphid-predator systems then gain in complexity (Stadler & Dixon, 2005). In exchange for aphid-produced honeydew, ants provide aphids with hygienic and protective services. Ants influence the predator-aphid relationship by attacking or carrying away (i.e., showing antagonistic behaviour) aphidophagous predators like syrphid larvae or ladybirds (reviewed in Way, 1963), and thereby increasing aphid fitness (Addicott, 1978; Buckley, 1987; Flatt & Weisser, 2000; Nixon, 1951; Stadler & Dixon, 2005). However, ants are not always beneficial for aphids but can also act as predators (Billick, Hammer, Reithel, & Abbot, 2007; Singh, Zytynska, Hanna, & Weisser, 2016). In many aphid systems, both mutualistic ants and predators can be influenced by intraspecific variation in the host plant. With this, both density-mediated and trait-mediated indirect effects on the third trophic level can come into effect. Moreira and Mooney (2013), for instance, could show that plant genetic diversity directly influenced aphid abundance which in turn affected the abundance of mutualistic ants and parasitoids (i.e., density-mediated indirect effects on the third trophic level). Host-plant-mediated changes in aphid traits (e.g., through plant architecture or biochemical diversity) are also known to affect the aphids' susceptibility to predatory attacks (e.g., increased hiding places) or alter ant preferences (and thus density) via variation in aphid honeydew composition across plants (i.e., trait-mediated indirect effect on the third trophic level; Cushman, 1991; Fischer & Shingleton, 2001; Johnson, 2008; Kareiva & Sahakian, 1990; Züst & Agrawal, 2017). The availability of ants can be crucial for some aphid species (e.g., obligate myrmecophilous species) and limit their realised distribution across different host plants (Senft, Weisser, & Zytynska, 2017; Wimp & Whitham, 2001).

In summary, aphids are subjected to a number of forces (e.g., plant chemotype, mutualistic ants and predators) defining their fitness. However, to which extent these different bottom-up and topdown forces contribute to observed patterns remains elusive.

One aphid-plant system that has been studied extensively is the tansy-aphid system. It offers ideal conditions to test relative effects of bottom-up and top-down effects (Stadler, 2004). It consists of (a) common tansy (*Tanacetum vulgare* L.; Asteraceae), an aromatic plant with a high chemical diversity regarding quantity and quality of stored and emitted VOCs (i.e., different plant chemotypes; Clancy et al., 2016; Forsén & Von Schantz, 1973; Rohloff, Mordal, & Dragland, 2004); (b) the highly specialised aphid *Metopeurum fuscoviride* 

Stroyan (Homoptera, Aphidoidea), an obligate myrmecophilous species, commonly tended by (c) ants such as *Lasius niger* L. (Formicidae); and (d) predated on by various common aphidophagous predators. In field studies on this system, the occurrence of aphids, tending ants and aphidophagous predators were associated with differences in the blend of volatile terpenoids across different plant chemotypes (Bálint et al., 2016; Clancy et al., 2016). This bottom-up effect of plant chemotype may therefore mediate effects of mutualists and predators on aphid populations (i.e., indirect effects of the chemotype), which could contribute to the distinct distribution of aphids observed in field surveys (Senft et al., 2017); for example, through higher predation pressure or reduced protection by ants on certain plant chemotypes.

So far, findings of potential plant chemotype, ant and predator effects on the aphid populations in the tansy-aphid system are mostly based on observational data, and thus remain correlational. Herein, we carried out a manipulation experiment that allowed us to explore how and to what extent direct and indirect bottom-up and top-down forces affect the aphid populations on tansy plants. While tansy terpenoid production (chemical volatiles), and thus also differences between chemotypes, has a genetic basis (i.e., different genotypes express different chemical phenotypes; Holopainen, Hiltunen, Lokki, Forsén, & Schantz, 1987), all our hypotheses are based on differences regarding variation in the plant's volatile chemical profile. Firstly, we hypothesised that aphid population growth rates (and subsequent abundance and colony survival) will vary across plant chemotypes. Secondly, we hypothesised that ants will benefit aphid populations by increasing growth and survival rates, whereas predators will have a negative effect on these parameters. Finally, the combined effect of ants and predators on aphid population dynamics (growth, abundance and colony survival) will depend on the plant chemotype, explained by effects of chemotype on the abundance of ants and predators on each plant.

## 2 | MATERIALS AND METHODS

## 2.1 | Plants propagation and GC-MS profiling of chemotypes

We used plants grown from seed collected from a field site of tansy (*Tanacetum vulgare* L.; Asteraceae) that was previously the focus of two studies about the spatio-temporal dynamics of tansy aphids (Senft et al., 2017), and the chemical diversity of tansy plants and how this affects early aphid colonisation (Clancy et al., 2016). Seeds were collected in late autumn 2013 from dried flower heads of plants growing at a distance between 3 and 21 m on a field site with 172 plant patches in Altenhausen, north of Freising in southern Germany (N 48°25'1.51"; E 11°46'1.19"). Tansy plants have a low rate of self-fertilisation (Lokki, Sorsa, Forsén, & Schantz, 1973). Therefore, the chemotypic profile of mother plants and their offspring can be different. We first grew 18 plants from seed collected from eight mother plants (1–3 seedlings per mother plant); these were chosen because they differed in their effects on aphids in the

field (Clancy et al., 2016). Plants were grown in individual pots (13 cm in diameter) containing commercial potting soil (Einheitserde<sup>®</sup>, Type SP ED63 T, Sinntal-Altengronau, Germany) and kept under greenhouse conditions (21.6°C mean ambient temperature, 67% mean relative humidity, 16:8 hr (light:dark) at Dürnast Experimental Station, Technical University of Munich, Freising, Germany) prior to the experiment. We analysed the chemical composition of all 18 plants using gas chromatography-mass spectrometry (GC-MS; see Supporting Information Table S1), based on 22 volatiles "putatively emitted from storage" (i.e., constitutively released from undisturbed glands on the plant's surface), which differentially defines tansy plants due to the variation in terpenoids (monoterpenes, oxygenated monoterpenes and sesquiterpenes) as in Clancy et al. (2016). From this, four plant individuals were chosen for the current experiment, based on chemotype effects in the field, including two plants low in 4-terpineol (higher aphid colonisation) and two high in this compound (lower aphid colonisation). Within these groups, the two plants were further chosen to be different in other aphid-related compounds from field data such as (E)-Dihydrocarvone. By comparing the chemical profiles (based on the relative concentrations

of the 22 volatile terpenoids "putatively emitted from storage"; Table S1) of the experimental plants to the profiles of field plants (data from Clancy et al., 2016) and by clustering them, we could confirm that the chemical profiles of the four experimental plants reflected the diversity of the chemical profiles of 172 plants from field sites (further information about the chemotype identification and the clustering analysis of experimental and field plants can be found in Supporting Information Appendix S1). The field and experimental plants clustered into the four main classes as previously described (Clancy et al., 2016). Two of four experimental plants belonged to class 2 chemotypes (experimental chemotypes 2.1 and 2.2) and two to class 4 chemotypes (experimental chemotypes 4.1 and 4.2; Figure 1). Class 2 chemotypes were dominated mostly by L-camphor (55.9% ± 2.4%; Figure 1) and supported early aphid colonisation in the field (Clancy et al., 2016). The terpenoid profiles of class 4 chemotypes were not clearly dominated by a single volatile; however, (Z)- $\beta$ -terpineol (from 0.0% to 55.0%) and eucalyptol (from 2.2% to 33.1%) were most abundant (Figure 1). Early aphid colonisation in the field was less likely on plants belonging to chemotype class 4 (Clancy et al., 2016).



**FIGURE 1** Chemotype clustering, chemotype profiles and experimental plants. Hierarchical cluster analysis of relative "likely emitted from storage" volatile concentrations from 172 plants from a small-scale field site and the four experimental mother plants (data from the field plants were used from Clancy et al., 2016). Four main classes were identified. Stacked bars show the mean relative concentrations of terpenoids in the different classes as well as the relative concentration of each experimental mother plant (2.1, 2.2, 4.1 and 4.2)

We then used vegetative propagation to obtain 20 clonal replicates of each of the four individual experimental chemotypes. This was achieved simply through splitting mother plants into multiple sections. Daughter plants were repotted in separate pots and regrowth occurred. We have previously shown that this method produces stable chemotype clones in tansy (Clancy et al., 2016). Three weeks after the last splitting event, and a week prior to the experiment, all but the longest stem were trimmed.

## 2.2 | Aphids and ants

*Metopeurum fuscoviride* aphids were collected from various tansy plants (mixed aphid genotypes) grown in the vicinity of the Weihenstephan campus of the Technical University of Munich, Freising, Germany. To avoid a bias due to variation among aphid genotypes, aphids were collected all in one glass jar and randomly selected for the experiment (note, crowding does not induce winged morph production in this species; Mehrparvar, Zytynska, & Weisser, 2013). Collected aphids were used the same day for the experiment (see experimental design section).

Lasius niger L. ant colonies (five colonies with each >2000 workers) were also collected around the University campus. All ant nests contained a high number of brood (>500). The colonies were housed in 10 L buckets and placed on the experimental field site near Dürnast Experimental Station, five days prior to the start of the experiment.

## 2.3 | Experimental design

To test the effects of plant chemotype, mutualistic ants and predators on aphid performance (aphid growth rate, abundance and survival), we used a fully factorial randomised block design with four chemotypes (chemotypes 2.1, 2.2, 4.1 and 4.2) from two different chemotype classes (class 2 chemotypes are known to support early aphid colonisation, while class 4 chemotypes did not support early aphid colonisation in the field), two ant treatments (presence and absence) and two predator treatments (presence and absence). Each of the 16 treatments was replicated five times (i.e., five blocks), resulting in 80 tansy plants. Each of the five blocks contained one repeat of each treatment in randomised order and a different *L. niger* ant colony (i.e., each of the five ant colonies was connected to 8 of the 16 plants in each block; Figure S1).

## 2.4 | Experimental setup

The experiment was performed near Dürnast Experimental Station (N 48°24'32". E 11°43'20") within a rectangular meadow with an approximate size of 90 × 30 m from mid-June until mid-July 2015. Only the part where the tansy plants were placed was mown  $(7 \times 7 \text{ m})$ . High grass and a variety of flowering plants surrounded the experimental area, ensuring sufficient habitat for natural enemies. The plants were transferred from the greenhouse to the field site and watered twice a day with tap water in case of no rain. Tube-like cages with a height of 30 cm and a diameter of 13 cm made out of PET transparencies (IP 2,912, H. Brunner GmbH, Achern, Germany) were placed on top of each pot and sealed (Figure 2). The cages contained a fine mesh on one side to allow ventilation and the top of each cage was closed with a removable mesh lid. PVC tubes (1 m length) were connected from the ant colonies to the cages allowing ants to enter the cages. Insect glue (Raupenleim grün, Schacht, Braunschweig, Germany) was used around the top of the cages and a fine mesh sealing the bottom of the pot to prevent ants from entering the cages opportunistically (Figure 2).

Ants were allowed to access the plants immediately after placing the plants outside (21 June). At the beginning, the mesh lids on top of the cages were closed to prevent predators from accessing the plants until start of the experiment. One day later (22 June), we added 10 × 1st instar larvae,  $10 \times 2nd/3rd$  instar larvae and  $5 \times 4th$  instar larvae/adult aphids to each plant, allowing them to settle overnight. On the next day (23 June), the aphids were counted and missing aphids (same age structure) were replaced



FIGURE 2 Experimental design: cages for predator and/or ant exclusion. Fine mesh was used to avoid aphid and ant movement between plants. Insect glue (Raupenleim grün) is a sticky substance, across which ants and other walking invertebrates cannot pass, and this was used to minimise access to plants by naturally occurring ants while allowing access to flying predator species (particularly important for the ant absence, predator present treatment combination) the next day. On 25 June (day zero of the experiment), the numbers of aphids per plant were recounted, and the experimental cages were opened in the predator treatments. Aphids, ants and predators in each cage were counted with a mechanical counter on days 2, 4, 6, 8, 11, 14, 17 and 20. The order of counting the cages (from 9 a.m. to 4 p.m.) was randomised every day to avoid diurnal effects. If no ant could be encountered on a plant, five ants were collected from the corresponding colony and transferred to the plant (these ants were not counted). At the end of the experiment on 15 July, the above-ground biomass of each tansy plant was measured by drying plants at 70°C for three days for dry weight determination.

### 2.5 | Statistical analysis

To quantify aphid population performance, we used three measures: (a) initial growth rate (up to day eight), which reflects the reproduction potential of a colony; (b) final abundance (day 20), which reflects the overall success; and, (c) survival, reflecting the persistence of a colony.

For each cage, a per capita initial growth rate (Agrawal, Underwood, & Stinchcombe, 2004) was calculated by subtracting the natural log of the aphid abundance on day eight (time when ant attendance dropped; Figure S2) by the log of the initial aphid abundance and then divided by the number of days. Linear models were used to analyse the aphid growth rate. Explanatory variables were either chemotype class (Class 2 and Class 4) or individual chemotypes (2.1, 2.2, 4.1 and 4.2), ant treatment (presence/absence) and predator treatment (presence/absence). In the full model, chemotype was allowed to interact with the ant and the predator treatment. As a fixed factor, we used experimental block (1-5) and plant biomass as a covariate. Nonsignificant interactions and explanatory variables were removed from the full model using the backwards stepwise method. In the result section, only minimal adequate models are presented. The strength of each effect was estimated by calculating the percentage of variance explained by each predictor (i.e., predictor sum of squares divided by the total sum of squares, multiplied by 100%).

To analyse treatment effects on the final aphid abundance, we calculated the log of the difference between the final and the initial aphid abundance. As described above, we used two linear models (log-transformed to achieve linearity) with the same variables as for the growth rate analysis.

The survival of aphid colonies was analysed using the survival functions "survfit" (package "survival"; Therneau, 2015) with standard settings in R, version 3.2.2 (R Development Core Team, 2014). Estimates of the different survival curves (censored data) were calculated using the Kaplan-Meier method by the "survfit" function, and a Cox proportional hazard model was used to analyse the effects of the different treatments (chemotypes, ants (presence/absence) and predators (presence/absence)) on aphid colony survival. A full model, containing all interactions, was fit

to the data and the backwards stepwise method was applied to remove nonsignificant terms.

We further analysed the abundance of ants and predators on each day by using a generalised linear mixed effect model fit by maximum likelihood (GLMER) in the R package "Ime4" (Bates, Maechler, Bolker, & Walker, 2015) with a poisson error distribution and log link function. In this analysis, each plant identity was included as the random factor due to repeated observations over the time course of the experiment. Continuous explanatory variables were centred and scaled using the scale function. As explanatory variables, we used predator or ant treatment, respectively, and chemotype or chemotype class. To distinguish between density- and trait-mediated effects of chemotype (mediated by aphids) on ant or predator abundances, we established two models: one with the number of aphids (during the day of observation) as covariate and one without (Mooney & Singer, 2012; Moreira & Mooney, 2013). Experimental block was used as fixed effect, and plant biomass and day of observation were used as covariates in both models. A full model, containing all explanatory variables and covariates, was fit to the data and the backwards stepwise method was applied to remove nonsignificant terms. Significance levels were calculated after model comparison through likelihood ratio tests.

We used linear models to determine whether (a) the mean number of ants was correlated with aphid colony growth rate (until day 8), that is, are more ants associated with a higher aphid colony growth rate? (b) a higher aphid colony growth rate led to higher aphid colony peak population sizes (log-transformed), and, (c) a higher mean predator abundance led to a decrease final aphid abundance. All statistical analyses were carried out in R, version 3.2.2 (R Development Core Team, 2014).

## 3 | RESULTS

The initial aphid population size was 40.0 ± 1.3 aphids (mean ± *SE*). The number of ants visiting aphid colonies decreased over time with more ants tending until day eight ( $\chi_1^2$  = 74.79, *p* < 0.001; Supporting information Figure S2 and Table S2). Final plant biomass varied across chemotypes ( $F_{3,76}$  = 5.98, *p* = 0.001) and was therefore included as a covariate in further models (i.e., plant biomass was highly correlated with plant growth rate (calculated as length growth per day): LM  $F_{1,78}$  = 147.45, *p* < 0.001).

Over the experimental period, five different kinds of aphidophagous predators were observed: syrphid larvae (Syrphidae;  $N_{cumulative} = 24$ ) on 14 plants (35% of plants where predators were allowed to enter); ladybirds and ladybird larvae (e.g., *Coccinella septempunctata* L., Coccinellidae;  $N_{cumulative} = 13$ ) on two plants (5%); lacewing larvae (Chrysopidae;  $N_{cumulative} = 6$ ) on five plants (12.5%); spiders (Araneae;  $N_{cumulative} = 6$ ) on three plants (7.5%); and parasitoids indirectly encountered through mummies (hardened shell of the host aphid after successful parasitism by a parasitoid wasp; Braconidae;  $N_{cumulative} = 43$ ) on eleven plants (27.5%). Despite the relatively high proportion of plants with mummies, the abundance per plant remained low, ranging between one and eight, and therefore parasitoids did not strongly contribute to any top-down regulation effect on aphid colony sizes, and thus were removed from further models.

### 3.1 | Aphid colony growth and abundance

To test the influence of plant chemotype, ants and predators on the performance of aphids, we analysed the initial growth rate (until day eight, after which ant tending was reduced; Figure S2) of aphid colonies on each plant. Plant chemotype explained 9.5% of the total variation in the model ( $F_{3,72}$  = 3.36, p = 0.023; Table 1). Here, the individual chemotype within the overall class was important since the aphids responded to the two chemotypes within class 2 differently. The difference was

mainly driven by chemotype 2.1 which had, by far, the highest growth rate both in the presence and absence of ants (Figure 3a).

The presence of ants on a plant had an overall positive effect on aphids by increasing colony growth rates, accounting for 13.6% of the total variation ( $F_{1,71}$  = 14.41, p < 0.001; Table 1, Figure 3a), but there was no evidence that higher mean ant abundances (within the ant presence treatment) on a plant resulted in higher aphid colony growth rates (LM  $F_{1,38}$  = 1.81; p = 0.187). We observed higher numbers of ants on plant chemotype 2.1 ( $\chi_3^2$  = 7.66, p = 0.053; Figure 3b; Supporting information Figure S3, Table S2). This chemotype effect on ant abundance is not only explained by the variable aphid density (i.e., actual aphid abundance;  $\chi_1^2$  = 9.60, p = 0.002) across the plants (i.e., density-mediated indirect chemotype effect), as the effect of chemotype on ants remains significant after including

TABLE 1 Effects of plant chemotype class, terpenoid chemotype, ants and predators on aphid colonies (growth rate and abundance)

	Growth rate <sup>a</sup>			Log (final-initial) <sup>b</sup>		
Response variable	df	F	р	df	F	р
Covariates						
Block	4,75	2.63	0.041	-	-	-
Biomass	-	-	-	1,78	15.04	<0.001
Variables						
Chemotype class (n = 2)	х	х	х	1,77	5.11	0.027
Individual chemotype (n = 4)	3,72	3.36	0.023	х	х	x
Ants (presence/absence)	1,71	14.41	<0.001	1,76	4.03	0.048
Predators (presence/absence)	-	-	-	1,75	25.58	<0.001

<sup>a</sup>Growth rate until day eight. <sup>b</sup>Models used were linear models. All full models contained interaction terms of main variables, but were not retained in the final models (not significant). All analyses were performed in R. "–" shows where a term was not retained in the most parsimonious model. "x" shows terms that were not included in the model.



**FIGURE 3** The effect of plant chemotype on aphid growth rate, ant and predator abundance. (a) Plant chemotype and ants affected aphid growth rates (N = 10). Aphids on chemotype 2.1 had a higher growth rate compared to aphids on other chemotypes and the presence of ants increased aphid growth rate across all chemotypes. (b) Plant chemotype affected ant abundance (N = 10, observations =9) with significant higher number of ants on chemotype 2.1 (independent of aphid abundance) than on chemotype 2.2 (c) Plant chemotype affected predator abundance (N = 10, observations = 9) with more predators found on chemotype 4.2 than on 4.2. Mean values ± *SE* are shown

aphid abundance in the model. This shows that there is a potential trait-mediated indirect effect of plant chemotype on ants ( $\chi_3^2$  = 7.93, p = 0.047; i.e., chemotype effects on aphid traits indirectly affects ants), leading to even higher ant abundances on chemotype 2.1, as expected with a simple increase in aphid numbers (i.e., higher antaphid ratio).

Predators on the other hand did not influence aphid colony growth, likely due to the low number of predators in the system until day eight (Table 1; Supporting information Figure S4). None of the interactive terms in our full model (i.e., all possible two- and threeway interactions between chemotypes, ants and predators) were significant and thus removed from the model.

A higher aphid growth rate inevitably led to a higher peak population size (LM  $F_{1,78}$  = 4.04, p = 0.048). The mean aphid peak population size during the experiment was 242.1 ± 12.0 (ranging between 80 and 541). The different aphid populations reached their peaks between experimental day 6 and day 20.

As a measure for the impact of the different treatments on aphid abundances across the whole experiment (i.e., beyond the peak population size), we calculated the log of the difference between the final and the initial aphid abundance. Ants increased the final abundance of aphids ( $F_{1.76}$  = 4.03, p = 0.048; Table 1), but only explained 3.2% of the total variation in the model. Predators had a strong negative impact, accounting for 20.5% of the total variation ( $F_{1.75}$  = 25.58, p < 0.001; Table 1); the final abundance of aphids was around four times higher when predators were absent. Aphids on chemotype class 2 plants had higher abundances at the end of the experiment, compared to those on plants of chemotype class 4  $(F_{1,77} = 5.11, p = 0.027;$  Table 1). In contrast to aphid growth rate, this effect was not driven by individual plant chemotypes. Chemotype class accounted for 4.1% of the variation in the model. Again, none of the interactive terms (i.e., possible two- and three-way interactions between chemotypes, ants and predators) were significant indicating that additive rather than interactive effects are present in our system.

#### 3.2 | Aphid colony survival and predation effects

The first plants without aphids (local extinction) were observed on day eleven (N = 2). At the end of the experiment (day 20), 18 plants were without aphids. Predators decreased the survival of aphid colonies (Cox proportional hazards model: LRT = 7.91, p = 0.005), but this did not vary across plant chemotypes (nonsig. term) and ants did not benefit colony survival (nonsig. term). None of the interactive terms in our full survival model (i.e., all possible two- and three-way interactions between chemotypes, ants and predators) were significant and thus removed from the model.

The abundance of predators increased over time ( $\chi_1^2$  = 12.57, *p* < 0.001; Supporting information Figure S4 and Table S3). Predators were more abundant on larger plants ( $\chi_1^2$  = 9.08, *p* = 0.003; Supporting information Table S3) and on chemotype 4.2 ( $\chi_3^2$  = 10.94, *p* = 0.012; Figure 3c; Supporting information Figure S3 and Table S3). Plant chemotype was significant when the covariate aphid abundance was both included in and excluded from the model showing that chemotype effects were rather trait-mediated indirect effects on predators than mediated through aphid density (Supporting information Table S3). Neither the ant presence treatment nor the abundance of ants reduced predator abundances on the plants (Supporting information Table S3 and Figure S5). In general, a higher mean predator abundance led to a lower final aphid abundance (LM  $F_{1.38}$  = 15.99, p < 0.001; Supporting information Figure S6).

## 4 | DISCUSSION

We found that plant chemical variation influenced aphid population dynamics across the host plants. There was a positive direct effect of certain plant chemotypes on aphid population growth and through this the final abundance of aphids, but not on aphid colony survival. Further, aphid population dynamics were indirectly affected via chains of direct interactions (Figure 4; Wootton, 1994). This means that plant chemical variation altered aphid densities, and higher aphid densities led to increased ant abundances (i.e., density-mediated indirect effect), but more ants did not lead to more aphids; yet, the presence of ants had a strong positive effect on aphid numbers. In addition, beyond effects via aphid densities, we also found trait-mediated indirect effects of plant chemotype on ants. Plant chemotype also affected predator abundances; however, higher aphid densities did not lead to higher predator abundances, suggesting trait-mediated indirect effects of plant chemotype on predators. Effects of predators on aphid population sizes increased as predator abundance increased. We did not detect any interaction modifications (i.e., higher-order interactions; Wootton, 1994), since plant chemotype did not alter the overall beneficial effect of ants on aphids or negative effect of predators on aphids, but simply enhanced these effects via chemotype effects on the ants and predators.

## 4.1 | Bottom-up effect of plant chemotype on aphid performance

Our results confirm previous field observations of tansy aphids that showed a beneficial effect of tansy plants in terpenoid chemotype class 2 (with camphor as dominating compound; Clancy et al., 2016); Clancy et al. (2016) could show that under natural conditions colonisation by winged aphids in the early part of the season was more likely on chemotype class 2 (43% probability of early aphid colonisation) than on plants from chemotype class 4 (17%) containing (Z)- $\beta$ -terpineol and/or eucalyptol (syn. 1,8 cineol) as dominating volatile terpenoids. As shown here, the mechanism underlying this could be the increased growth rate when feeding on these chemotypes that would inevitably lead to higher number of aphids within a short period of time, and thus enhance the chance of successfully establishing a colony in the field. However, whether this is driven by a reduced plant defence of chemotype class 2 against aphids or other characteristics being **FIGURE 4** The tansy chemotype acts directly on all species in the community. Aphid population development parameters are directly affected by the associated species community (mutualistic ants and predators; arrow strength indicates strength of effect, "+" and "-" signs indicate positive or negative effects). Plant chemotype indirectly affected higher trophic levels via aphids through trait-mediated indirect effects (i.e., ants and predators) and density-mediated indirect effects (i.e., ants of the provides of the provi



beneficial for aphid population development (e.g., higher defence capability of aphids against predators or diseases, like fungal infections; e.g., Züst & Agrawal, 2017) remains to be elucidated in future experiments.

# 4.2 | Bottom-up effect of plant chemotype on the associated community

The number of ants observed on plants varied with plant chemotype. Ants visited aphid colonies on plant chemotype 2.1 more frequently indicating a potential ant preference for aphids on this chemotype, especially as ants had the free choice between colonies on all four chemotypes. This plant-derived effect on ants is indirectly mediated by aphids. While the number of aphids had an effect on ant abundance in the statistical model (i.e., densitymediated indirect effect with more aphids leading to more ants), chemotype still explained residual variation also when accounting for aphid abundance. This suggests that the plant chemotype has a trait-mediated indirect effect on ant abundance (i.e., more ants per aphid; Mooney & Agrawal, 2008; Mooney & Singer, 2012, Moreira & Mooney, 2013). The latter implies that the plant chemotype affects aphid traits. Here, it is conceivable that differences in aphid growth rates among plant chemotypes mediate this effect (high growth/high reward) or that aphid honeydew production/ quality differs among chemotypes, for instance, by differently sequestering plant secondary compounds (beneficial or disadvantageous in terms of ant recruiting; Brower et al., 1967; Fischer & Shingleton, 2001; Pringle, Novo, Ableson, Barbehenn, & Vannette, 2014; Vantaux, van den Ende, Billen, & Wenseleers, 2011; Züst & Agrawal, 2017). Whether ants are indirectly more attracted to this particular chemotype or repelled by others remains unknown. It

also remains unknown whether direct effects of the plant chemotype on the third trophic level (i.e., directly affecting ant density or ant traits) come into play, as the experimental design did not allow to test for such direct effects. However, it is known that plants can repel ants with certain odour profiles making aphid colonies more susceptible to predatory attacks (Ghazoul, 2001; Junker, Gershenzon, & Unsicker, 2011). As described for another obligate myrmecophilous aphid species (*Chaitophorus populicola* Wimp & Whitham, 2001), aphids could be restricted to certain plant hosts via host-plant effects on ant preference.

Predators were also indirectly affected by plant chemotype effects on aphid traits (i.e., not a density-mediated indirect effect via variable aphid colony sizes; Mooney & Singer, 2012), resulting in higher abundances on chemotype 4.2. As described for ants, it remains unknown whether chemotype indirectly affects predator abundances via aphid traits (e.g., susceptibility to attacks) or whether the plant chemotype directly affects predator density. Nevertheless, due to a higher predation pressure on these chemotypes (i.e., more predators lead to a higher reduction of aphids), aphids probably have a reduced chance to establish a new colony on these chemotypes under natural conditions. This supports our field observations (Clancy et al., 2016).

# 4.3 | Top-down effect of ants and predators on aphid dynamics

In our system, ants increased aphid population growth and final abundance, confirming previous work showing the beneficial function of ants in this mutualistic system (Mehrparvar, Mansouri, & Weisser, 2014; Stadler, 2004). Despite the exclusion of ants, the obligate myrmecophilous aphid species *Metopeurum fuscoviride* was still able to maintain a relatively high colony growth rate. As consequence however, they often suffered from fungal infections (pers. observation), probably as these aphids are not able to remove honeydew efficiently and the nidus remained within the colony (Buckley, 1987; Nielsen, Agrawal, & Hajek, 2010).

Ant attendance decreased over time and made Metopeurum colonies prone to predatory attacks. Such a decrease (between June and July) is not unusual and reflects observations in the field under natural conditions where L. niger encounters started to decrease in the beginning of July (Senft et al., 2017). This may be associated with the mating flight of ants (mostly between July and August: see Hart. Hesselberg, Nesbit, & Goodenough, 2018). According to Edwards (1951), ants change their foraging behaviour from protein to sugar sources when their larvae start to pupate. This might change again, during/after mating flights, when a vast number of winged ant individuals leave the nest and thus the need of sugar-rich sources (i.e., honeydew) decreases abruptly; this might lead to the abandonment of aphid colonies. As shown elsewhere (Addicott, 1979; Rico-Gray & Oliveira, 2007), after promoting the growth of low-density aphid colonies, the positive impact of ants decreased with an increasing population of aphids. It is assumed that ants are not able to respond to the rapid increase of aphid densities or they already have sufficient resources from "medium-sized" colonies. In consequence, predators were able to reduce aphid abundance and lower colony survivorship across all treatments later in the season, even on plants where ants had access (as in Stadler, 2004).

Interestingly, our data suggest that there is a positive relationship between plant biomass and predator abundance as well as a negative relationship to aphid population development. Plants with higher biomass had more predator encounters and lower aphid abundances at the end of the experiment: For example, tansy chemotype 2.2 plants had a higher biomass compared to the other chemotypes, yet conferred lower aphid population growth rates. However, the causal relationship between these remains unknown, but we cannot rule out a potential positive fitness effect of lower aphid numbers on plant biomass production (Halaj & Wise, 2001).

### 4.4 | Intraspecific variation in chemotypes

Our results highlighted the extent of variation within the main chemotype classes (i.e., between the individual chemotypes). For example, we found higher growth rates of aphid colonies on chemotype 2.1 but not on chemotype 2.2. Therefore, it is not just the main compounds differentiating the plants, but also the minor compounds, or the terpenoid pattern, which affected the associated ecological community (Clancy et al., 2016). We also found inconsistencies in the preferences of ants and predators for aphid colonies on individual plant chemotypes within their respective chemotype class. Differences in aphid, ant and predator responses between individual tansy plants (beyond chemotype class) could also be caused by additional differences, for example, in phenotype (e.g., trichome density; Johnson, 2008) or metabolomic diversity of nonvolatile compounds (Clancy et al. 2018; Kleine & Muller, 2011). However, it is known that the tansy chemotype is genetically determined and associated with genetic variation and a number of morphological traits (e.g., shoot height, number of flower heads, corymb height or flowering time) (Keskitalo et al., 2001).

## 5 | CONCLUSION

By disentangling multiple factors under a controlled manipulated experimental design, we were able to quantify effects of plant chemotype (bottom-up), mutualistic ants and predators (top-down) on aphid population dynamics (Figure 4). We showed that, while direct effects between pairs of species had the strongest effect on structuring the community, effects of plant chemotype could also act indirectly on aphid populations through a chain of direct interactions via the higher trophic levels. This work confirms results from field studies, highlighting the ecological and evolutionary consequences of plant chemical variation for natural communities. Such variation can lead to structuring of communities at the plant level, with each plant variant (here, chemotype) having its own specific effect on the interacting herbivore, mutualist and predator populations, that is, individual plant-specific community dynamics. In a natural population of a patchily distributed host plant, individual variation such as chemical profile differences between plants can create a heterogeneous habitat for associated herbivores even within a single field site. Where this occurs, the community of herbivores and their associated mutualists and natural enemies may vary at the level of a single plant. This could lead to metacommunity dynamics at smaller scales than is often considered in current literature (often regional scales are used). We suggest that effects of within-species plant variation should be incorporated into studies of metacommunity dynamics, especially when the system is focused on sessile plantspecialist herbivores.

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#### AUTHORS' CONTRIBUTION

M.S., S.E.Z., J.-P.S. and W.W.W. designed the study. M.S. collected and analysed the data. M.V.C. and J-P.S. carried out the chemotype analysis. M.S. wrote the first draft, and all authors contributed critically to the drafts and gave final approval for publication.

#### DATA ACCESSIBILITY

Data deposited in the Dryad Digital Repository: https://doi. org/10.5061/dryad.g9n85sm, (Senft, Clancy, Weisser, Schnitzler, & Zytynska, 2018).

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## SUPPORTING INFORMATION

Additional supporting information may be found online in the Supporting Information section at the end of the article.

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