Interspecific plant competition mediates the metabolic and ecological signature of a plant-herbivore interaction under warming and elevated CO2

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Abstract

1. Biotic interactions shape community evolution, but we lack mechanistic insights on how metabolic and ecological processes under climate change are altered by biotic interactions.

2. We used a two-trophic model community consisting of the aphid *Dysaphis plantaginea* feeding on the forb *Plantago lanceolata*, and a grass competitor *Lolium perenne* that does not experience herbivory by the aphid. Monocultures and mixtures were exposed to the herbivory treatment and to three relevant simulated environmental changes as prevalent under current climate change (increased temperature, CO_2, and increased temperature and CO_2)

3. Elevated CO_2 reduced the nitrogen content of *P. lanceolata*, while simultaneous increases of CO_2 and temperature modified the plant metabolic component and the magnitude of these responses in different directions. Elevated CO_2 enhanced defence systems in *P. lanceolata*, but these effects were not altered by warming. Interspecific plant competition did, however, neutralise these responses. There were no indications for indirect effects of climate on aphid population growth as mediated by changes in plant defence, nutritional quality or biomass.

4. We thus demonstrate interactions between abiotic and biotic processes on plant metabolite profiles, but more importantly, that climate change effect on a selection of the metabolic pathways are altered by herbivory and competition. Our experiment under semi-natural conditions thus demonstrates the non-additive and often neutralizing effects of biotic interactions on plant metabolism and species performance under climate-associated environmental change.

48 # Introduction

49 Interactions between insect herbivores and host plants are influenced by primary and
50 secondary metabolites. The concentrations and types of these molecules in plants is likely
51 affected by climate change (Bidart-Bouzat & Imeh-Nathaniel, 2008; Zvereva & Kozlov,
52 2006). For example, plants exposed to elevated CO_2 show higher concentrations of many
53 organic molecules, including carbohydrates (fertilization effect), but generally lower or
54 unaltered nitrogen levels, and consequently higher C:N ratios (Bezemer & Jones, 1998;
55 Lincoln *et al.*, 1993; Robinson *et al.*, 2012; Stiling & Cornelissen, 2007). Lower nitrogen
56 concentrations imply lower levels of leaf protein and amino acids and reduced nutritive value
57 for herbivores (Lincoln *et al.*, 1986). Consequently, the performance of insects generally
58 reduces under high CO_2 (Bezemer & Jones, 1998; Hunter, 2001; Lincoln *et al.*, 1993).
59 Contrary to elevated CO_2, moderate warming has positive effects on insect herbivore
60 performance by reducing development time (Bale *et al.*, 2002; Van De Velde *et al.*, 2016) and
61 increasing fecundity (Meisner *et al.*, 2014). Climate change may thus paradoxically constrain
62 rather than facilitate insect herbivore population growth by lowering plant quality (Bauerfeind
63 & Fischer, 2013; Jamieson *et al.*, 2015). These resource-quality mediated effects are,
64 however, anticipated to not fully counterbalance the direct effects of warming on
65 development, rendering the overall herbivore response to warming positive (Bauerfeind &
66 Fischer, 2013; Zvereva & Kozlov, 2006).

67

68 Secondary metabolites (e.g. lignin, tannins, phenolics and terpenoids), greatly affect tissue
69 quality by determining the nutritive value, palatability, digestibility and/or toxicity of foliage
70 (see Table 1 for a synthesis). Compared to primary metabolites, responses of secondary
71 metabolites to elevated CO_2 and warming are more variable and less understood (Bidart-
72 Bouzat & Imeh-Nathaniel, 2008; Robinson *et al.*, 2012). According to the carbon-nutrient

73 balance hypothesis (Bryant *et al.*, 1983), elevated CO_2 is expected to increase secondary
74 metabolite levels as a result of the 'excess' carbon (fertilizing). Yet, because of the relatively
75 lower N-availability, the increase of nitrogen-containing metabolites will be lower than
76 expected (Robinson *et al.*, 2012). On the other hand, trade-offs between growth and
77 secondary metabolisms are anticipated to promote primary metabolic activity under
78 warming and therefore favour photosynthesis and growth relative to defence. This
79 expectation corresponds to the growth-differentiation balance hypothesis (Herms & Mattson,
80 1992). The different components associated with climate change thus induce direct, and often
81 orthogonal responses in chemical defences (Bidart-Bouzat & Imeh-Nathaniel, 2008; Zavala *et*
82 *al.*, 2013; Zvereva & Kozlov, 2006).

83

84 Biotic interactions have a strong impact on life histories and physiology of species, and these
85 interactions may shift responses of plants towards climate change (Van de Velde et al. 2016).
86 Competition for limited nutrients for example, enhances the availability of carbon in a focal
87 plant relative to its demand. This, in turn, can increase carbon-based defences and thus reduce
88 herbivory compared with plants that are less constrained by competitors (Bryant *et al.*, 1983).
89 Also the identity of a neighbouring plant may affect the resistance of a focal plant by a
90 different outcome of the plant-plant competition (Barton & Bowers, 2006; Broz *et al.*, 2010).
91 Conspecific competition, in particular, may result in stronger decline of plant growth and
92 increased defence compared to heterospecific competition. Indeed, Broz *et al.* (2010) for
93 instance showed that focal plants with conspecific neighbours allocated more resources
94 towards production of carbon-based defence molecules, whereas those grown with
95 heterospecific neighbours allocated more resources towards growth. Hence, elevated CO_2 as
96 well as warming is anticipated to indirectly affect the defence of a focal plant, through species
97 specific stimulation of plant growth and thus plant-plant competition.

98 Although atmospheric CO_2 and temperature increase concurrently, empirical studies
99 investigating their combined effect on multi-trophic communities are surprisingly scarce and
100 insufficient to effectively guide theory or synthesis (Cornelissen, 2011; Dyer er al. 2013;
101 Zvereva & Kozlov, 2006). The current study therefore investigates effects of elevated CO_2 as
102 well as combined effects of elevated CO_2 and warming on a simple model community
103 consisting of three species: rosy apple aphid *Dysaphis plantaginea* Passerini (*Hemiptera:*
104 *Aphididae*) feeding on plantain, *Plantago lanceolata* L., and a heterospecific neighboring
105 plant species, perennial ryegrass, *Lolium perenne* L. The aphid does not feed on *L. perenne.*
106 The experimental design consisted of monocultures of *P. lanceolata* and mixtures of *L.*
107 *perenne* and *P. lanceolata* exposed to the two aforementioned simulated climate scenarios as
108 well as to a control scenario with the current climate. As the experiments were conducted
109 outdoors, we were also able to *post-hoc* study how the imposed climatic and competition
110 treatments affected mildew infections. Biotrophic parasites such as mildew extract nutrients
111 from living cells and have extended periods of physiological interaction with their hosts
112 (Agrios, 2005). Elevated CO_2 and warming have been shown to affect the severity of
113 pathogen infections (Mikkelsen *et al.*, 2015, Thomas & Blanford, 2003). We took advantage
114 of this unplanned infestation to study the impact of elevated CO_2 and warming on mildew
115 infestation and the putative additive effects of mildew infestation on plant-insect herbivore
116 interaction.

117

118 Based on the above, we anticipate that elevated CO_2 and warming will alter foliar nutrients
119 and defence molecules, that altered host quality and plant resistance will affect insect
120 herbivore performance, and that elevated CO_2 and warming should indirectly influence host
121 quality and plant resistance via effects on neighbouring plants. We predict investments in
122 defence to be higher under conspecific competition relative to heterospecific competition

because of increased resource competition. In addition, increased CO_2 is anticipated to increase chemical defence by promoting the production of secondary metabolites, while simultaneously increased temperature would in contrast promote plant growth. Depending on the relative rates of these physiological changes, net impact on herbivore populations may either remain constant, decline or increase if defence is respectively proportionally promoted or constrained relative to metabolism.

We do, however, aim to move beyond this generic perspective and to acquire a deep mechanistic understanding of the changing ecological interactions under elevated CO_2 and warming. To this end, we selected metabolites and groups of metabolites as molecular markers to evaluate defences and stress levels (see Table 1). In particular, we quantified an indicator of oxidative cellular damage (lipid peroxidation). Oxidative stress in plants is commonly induced under abiotic stress, but also by herbivory (Wu & Baldwin, 2010). Elevated CO_2 on the other hand, often reduces abiotic stress impact (Zinta et al. 2018). Herbivory damage also induces the production of secondary metabolites associated with the salicylic acid and jasmonic acid pathway (Kuśnierczyk *et al.*, 2011; Morkunas & Gabryś, 2011). Therefore, as an indicator of plant-insect interaction, jasmonic acid levels were determined. Simultaneously with the stress status, we evaluated the plant capacity to defend itself by quantifying protective metabolites. Here we determined molecular antioxidants (tocopherols, antioxidant capacity, carotenoids, proline) providing protection against oxidative stress (Gómez-Ariza *et al.*, 2007, Peshev *et al.*, 2013). We also quantified several groups of molecules that contribute to defence against insects (glycosides, tannins, lignin). Finally, elevated CO_2 as well as elevated temperature are known to affect stomatal aperture and therefore water transport. To evaluate the plant response to its water status we quantified osmolytes (proline, soluble sugars). Increase in these molecules reduces the cellular water

148 potential, improving water uptake. Given the anticipated complex interplay between elevated
149 CO_2, temperature and competition, we performed a multivariate grouping analysis to
150 document concerted responses in the quantified metabolites.

151

152 Table 1: Overview of molecular markers determined in this study and their principal function.

METABOLITE	PRINCIPAL FUNCTION	FUNCTION DESCRIPTION	REF.
TOCOPHEROL (VITAMINE E)	Lipophilic antioxidant	Components of biological membranes; act as scavengers of oxygen radicals generated by stress, in particular singlet oxygen. Tocopherols also stabilize membrane structures.	Blokhina et al (2003)
CAROTENOIDS	Lipophilic antioxidant	Lipophilic antioxidants localized in plastids of photosynthetic and non-photosynthetic plant tissues. Carotenoids react with lipid peroxidation products to end the chain reactions and dissipate the excess excitation energy.	Bartley and Scolnlk (1995)
PHENOLICS/ FLAVONOIDS	Antioxidant	Polyphenols have free radical scavenging activity and are involved in hydrogen peroxide neutralization. Flavonoids are also secondary ROS scavenging molecules in plants under excess excitation.	Blokhina et al (2003)
TOTAL ANTIOXIDANT CAPACITY (TAC)		Total antioxidant capacity (TAC) is the measure of the gross capacity of a plant extract to scavenge reactive oxygen species.	Das and Roychoudhury (2014)
MALONDIALDEHYDE, (MDA)	Lipid peroxidation marker	Oxidative stress induces lipid peroxidation disrupting membrane functions. Lipid peroxidation products (MDA) indicate membrane oxidation levels.	Das and Roychoudhury (2014)
SOLUBLE SUGARS (GLUCOSE, SUCROSE, FRUCTOSE)	Osmolytes	Soluble sugars protect plant cells through osmoregulation, improving water uptake and retention, and may impact cellular membranes through interaction with the lipid bilayer.	Shao et al., (2006)
PROLINE	Osmolytes/ antioxidant	Proline is an osmolyte, contributing to water uptake, as well as an antioxidant.	Szabados and Savoure (2010).
GLYCOSIDES: CATAPOL, AUCUBIN	Biotic stress defence	Secondary metabolites that help repelling herbivores, protects plant from UV and contributes to control of competitive interactions between plants.	Bowers (1991).
TANNINS	Biotic stress defence	Tannins are astringent bitter polyphenols and act as feeding deterrents to many insect pests.	Barbehenn et al., (2011)
LIGNIN	Biotic stress defence	A complex phenolic polymer that plant cell wall rigidity and hydrophobic properties, and is an important barrier against pests and pathogens.	Liu et al., (2018)
SALICYLIC ACID	Biotic stress	Salicylic acid is a phytohormone involved in regulation of	War et al.,

	defence signal	plant defence, and resistance to insect pests.	(2012)

Materials and methods

Experimental set-up (Fig S1.1 in supplementary material)

The study was performed at the Drie Eiken Campus, University of Antwerp, Wilrijk, Belgium (51° 09' N, 04° 24'E) in 12 sunlit, south-facing, climate-controlled chambers (details in Naudts *et al.* 2014). Three climate scenarios (four chambers per scenario) were simulated in an additive design: (1) current atmospheric CO_2 concentration and temperature (C); (2) future atmospheric CO_2 and current temperature (CO_2); and (3) future atmospheric CO_2 and temperature (TCO_2). Climate scenarios with elevated CO_2 had a target CO_2 concentration of 620 µmol mol^{-1} and future temperature chambers simulated a continuous 3 °C warming above fluctuating ambient temperatures. Climate manipulations were based on the IPCC-SRES B2-scenario prediction of moderate change for the year 2100 (IPCC, 2001).

The CO_2 concentration in each chamber was continuously measured and maintained at the target concentration with a CO_2 control group with an infrared analyser (WMA-4, PPSystems, Hitchin, UK). Air temperature and relative humidity were monitored every 0.5 h with a combined humidity–temperature sensor (Siemens QFA66, Erlangen, Germany), by averaging instantaneous readings in half-hour mean values. During the experiment, the CO_2 concentration was 382 ± 55 µmol mol^{-1} (SD) in the current climate, while it was 615 ± 70 µmol mol^{-1} (SD) in the climate scenarios with future CO_2 concentration (CO_2 and TCO_2). The monthly average air temperature in the C and CO_2 chambers was 16.2, 17.2 and 18.7 °C in June, July and August, respectively. TCO_2 chambers were 2.9 ± 1.0 °C (SD) warmer than current temperature chambers. Average vapour pressure deficit was 0.60 ± 0.34 and 0.64 ± 0.52 kPa (SD) in the climate treatments with ambient and warmed air, respectively. Irrigation

177 mimicked the rainfall outside (Naudts et al. 2014), with monthly totals equalling 64.4, 85.1
178 and 80.2 mm in June, July and August, respectively. Water freely drained while capillary rise
179 was prevented by a drainage system placed below the chambers. Future climate chambers
180 (TCO_2) received the same amount of water as current climate chambers, so that any increase
181 in water consumption would result in (aggravated) soil drought.

182

183 *Plant and insect communities*

184 We used two common co-occurring grassland species, *L. perenne* and *P. lanceolata*. *P.*
185 *lanceolata* is characterized by the presence of the iridoid glycosides aucubin and catalpol.
186 These compounds stimulate feeding and oviposition by specialist insects and can act as
187 defence against generalist herbivores (Bowers & Puttick, 1988; Puttick & Bowers, 1988).
188 Both plant species were sown at the end of March in a non-climate controlled greenhouse
189 with a time lag of one week to prevent size differences at the start of the experiment (Cotrufo
190 & Gorissen, 1997), and were watered twice a week. Four or five-week-old seedlings were
191 transplanted into PVC containers (24 cm inner diameter and 40 cm height), filled with sandy
192 soil (93.9% sand, 4.1% silt, 2.0% clay; pH 7.5; Kjeldahl-N 0.125 g kg^{-1}; 2.1% C in humus).
193 Each of the 12 chambers received 20 containers with two different compositions: (1) 10
194 monocultures of *P. lanceolata*, and (2) 10 mixtures of both plant species in a 50:50 ratio.
195 Each community contained 18 individuals planted in a hexagonal grid with 4.5 cm interspace.
196 Interspecific interactions were maximized by avoiding clumping. All communities were
197 fertilized with 10 g m^{-2} NH_4NO_3, 5 g m^{-2} P_2O_5, 10 g m^{-2} K_2O and micro-elements (Fe, Mn,
198 Zn, Cu, B, Mo), given dissolved in water in two equal amounts.

199 When the plants were three months old, *P. lanceolata* was involuntarily infested with
200 powdery mildew *Podosphaera plantaginis*. *P. plantaginis* is a biotrophic fungal pathogen
201 which means that it feeds on living plant tissue, but does not kill the infected host. The

202 powdery mildew was found in all climate scenarios and plant compositions but not all the
203 containers were equally affected. We took advantage of this unplanned infestation to study the
204 impact of elevated CO_2 and warming on mildew infestation and the putative additive effects
205 of mildew infestation on plant-insect herbivore interaction. Because these infections occurred
206 post-hoc, we do not have proper controls to quantify mildew effects on metabolite
207 composition.

208

209 The rosy apple aphid *D. plantaginea* was used as an insect herbivore. It overwinters as eggs
210 on apple trees, the primary host plant, and migrates in spring to the obligate alternate hosts, *P.*
211 *major* L. and *P. lanceolata* (Alford, 2014). On *Plantago* spp., they give birth to apterous
212 (wingless) morphs that reproduce by parthenogenesis (Blommers *et al.*, 2004). *L. perenne* is
213 not a host plant for *D. plantaginea*. The aphids were reared in small cages on *P. lanceolata*
214 under laboratory conditions of 22 ± 1 °C. They were introduced on *P. lanceolata* when the
215 plants were 20 weeks old. At this time, two adult, apterous aphids were placed with a dry
216 paintbrush on the apex of each *P. lanceolata* plant in monocultures and mixtures.
217 Consequently, at the start of the infestation each container was inoculated with 36
218 (monocultures) or 18 (mixtures) aphids. In each chamber, four monocultures of *P. lanceolata*
219 and four mixtures were randomly chosen for aphid infestation All containers were meshed,
220 and those not receiving aphids acted as controls. The meshes consisted of an 85-cm-tall
221 cylinder of lightweight netting to ensure aphids did not migrate between pots. The
222 infrastructure did not physically limit plant growth and did not cause photosynthetic stress
223 effects (F_v/F_m, the intrinsic efficiency of PSII in controls equalled 0.84 which is an optimal
224 value (Johnson *et al.*, 1993).

225

226 *Data collection*

In the fourth week after the aphid introduction we determined the mildew infestation. The degree of powdery mildew on *P. lanceolata* was categorized by a rating system: 1) healthy (no visible lesions); 2) 1% - 25% of the leaves damaged; 3) 26% - 50% of the leaves damaged; 4) 51% - 75% of the leaves damaged; 5) more than 75% of the leaves damaged. At the same time, aphid populations of four replicate communities per plant composition in each chamber were collected (totalling 4 replicate communities × 2 plant compositions × 12 chambers). Aphids were brushed directly into 70% ethanol with a dry paintbrush. It was not possible to collect all the aphids because their number per container was either too high or too low (hard to find), therefore a subsample of the population per container was collected: we searched for and collected aphids for 30 minutes. The total number of aphids per community was divided by the number of *P. lanceolata* individuals in that community. At the same time, we also harvested the plants of two replicate communities per treatment in each chamber (totalling 2 replicate communities × 4 treatments × 12 chambers). Alive aboveground plant biomass in each community was separated from dead biomass by species, dried at 70 °C for 48 h and weighed. These biomass values per species were likewise divided by the number of the species' individuals in the community, providing primary data for the statistical analysis. The alive biomass samples of *P. lanceolata* plants of the two harvested communities per treatment in each chamber (2 replicate communities × 4 treatments × 12 chambers) were ground in a mill. Three subsamples per community were analysed for nitrogen and carbon content using a NC element analyser (NC-2100 element analyser, Carlo Erba Instruments, Milano, Italy), which were averaged prior to data analysis.

A separate subsample of the milled live aboveground biomass of *P. lanceolata* of the two harvested communities per treatment in each chamber (2 replicate communities × 4 treatments × 12 chambers) was taken to quantify biochemical parameters. These parameters are indicators of stress in the organism (membrane damage as malondialdehyde levels, MDA),

252 indicators of antioxidant defences i.e. total antioxidant capacity (TAC) and antioxidant
253 molecules (tocopherols, carotenes, proline, phenols); indicators of biotic stress and defence,
254 i.e. glycosides (catapol, aucubin), tannin, lignin and salicylic acid; and metabolites involved in
255 plant water-deficit defence (proline, soluble sugars). We refer to Supplementary material 2 for
256 a detailed description of the methods used to quantify these metabolites.

257

258 *Data analysis*

259 We focused on *P. lanceolata* because this is the host plant for the aphid and powdery mildew.
260 In a first step we examined covariation in the metabolites responses to the three treatments
261 (aphid infestation, climate scenario and plant composition) by using a hierarchical clustering
262 analysis. Metabolites were normalized by Z-transformation and subjected to a hierarchical
263 clustering analysis with an Euclidean distance metric, and visualized as a heat map using
264 Multi Experiment Viewer (Mev) version 4.8 (Saeed *et al.*, 2003). A distance cut off of 0.3
265 was applied. The clusters were used in the structural equation models (see below).

266 In a second step, we fitted two piecewise Structural Equation Models (SEM), which combine
267 information from multiple separate linear models into a single causal network (Shipley,
268 2009). The first SEM investigated the effect of aphid infestation, climate scenario, plant
269 composition and mildew infestation on the metabolites and alive aboveground biomass of *P.*
270 *lanceolata*. In a second SEM, we separated the metabolites of *P. lanceolata* in control pots
271 from those with aphids. This allowed us to test the effects of the metabolites on the aphid
272 population, as well as to quantify effects of the aphid population on these metabolites. The
273 response of the aphid population was measured as their number at the end of the infestation.
274 The four clusters obtained by the clustering analysis were used to divide the metabolites in
275 separate groups and hence to avoid redundancy in the analyses. We standardized the
276 metabolites, the alive aboveground biomass of *P. lanceolata* and the number of aphids by

converting to Z-scores to equalize variances. To reduce the number of mildew categories, we rearranged degree of mildew in two categories: (1) no or mild mildew infection (category 1 - 2) (2) severe mildew infection (category 3 - 5).

Traditional SEM estimation methods assume that all variables follow a normal distribution and all observations are independent (Grace, 2006). In our analyses, we used piecewise SEM that allows fitting general linear mixed effect models that can incorporate random effects. Each mixed effects model was fitted using the "lme" function in the "nlme" package (version 3.1-128) in R. For each model, we fitted a random effect of chamber. The overall path model (the SEM) was fitted using the "piecewiseSEM" package (version 1.2.1) in R (Lefcheck, 2016). Goodness of fit was estimated using Shipley's test of d-separation, which yields a Fisher's C statistic that is Chi-squared distributed (Shipley, 2009). If the resulting P-value > 0.05, then the SEM can be said to adequately reproduce the hypothesized causal network.

As a final confirmatory step, all data were analysed with General Linear Mixed models (GLM) in SAS (version 9.2, SAS Institute Inc., Cary, NC) (Littell *et al.*, 1996) with chamber as a random factor nested within climate scenario. Climate scenario, plant composition, aphid infestation and two-way and three-way interactions between these predictors were included as fixed factors. Because mildew infestation did not have a significant effect on biochemical plant responses, this treatment was excluded from the GLM analysis. Non-significant factors were backwards-excluded from the model. In case of significant effects, *a posteriori* means comparisons using the Tukey test, corrected for multiple comparisons, were made. Effects were considered significant at $P \leq 0.05$. Repeated univariate analyses are subject to error proliferations. We deliberately decided not to correct results from these analyses for multiple testing, but instead interpret them in a conservative way in congruence with the higher described grouping analyses. Results and discussion of the series of univariate analyses are therefore provided in Supplementary Material 3.

302

303

Results

Hierarchical clustering analysis of metabolites

The variation in primary and secondary metabolites of *P. lanceolata* was subjected to a hierarchical clustering analysis in order to understand their grouping according to their covariation in response to the imposed herbivory, climate and competition treatments. We obtained four major clusters of metabolites (clusters 1 - 4 in Fig. 1).

The first cluster containing metabolites consistently upregulated by the presence of aphids contains the phenol antioxidants, the lipophilic antioxidant tocopherol, the C:N ratio, tannins, and the non-structural carbohydrates soluble sugars (including glucose, fructose) and starch (Fig. 1). The second cluster contains another group of a.o., antioxidant and defence molecules that are downregulated by elevated CO_2 and more so by aphids in mono-cultures, i.e. the carotenoids, as well as the growth regulator, jasmonic acid, total proteins, phosphorous and lipids (Fig. 1).

The nutrient N which contributes solely to the third cluster, is downregulated under elevated CO_2 . Carbon concentration, the C:P and N:P ratio, and lignin:N were classified in the fourth cluster that is specifically upregulated by aphids in monocultures under elevated CO_2 (Fig. 1). Cluster 4 also contains metabolites that play a role in plant defences against herbivores, such as salicylic acid and the iridoid glycosides (catalpol and aucubin), and also contains TAC, lignin, cellulose, proline and MDA. Notably in this cluster, metabolites are strongly induced in CO_2 and TCO_2.

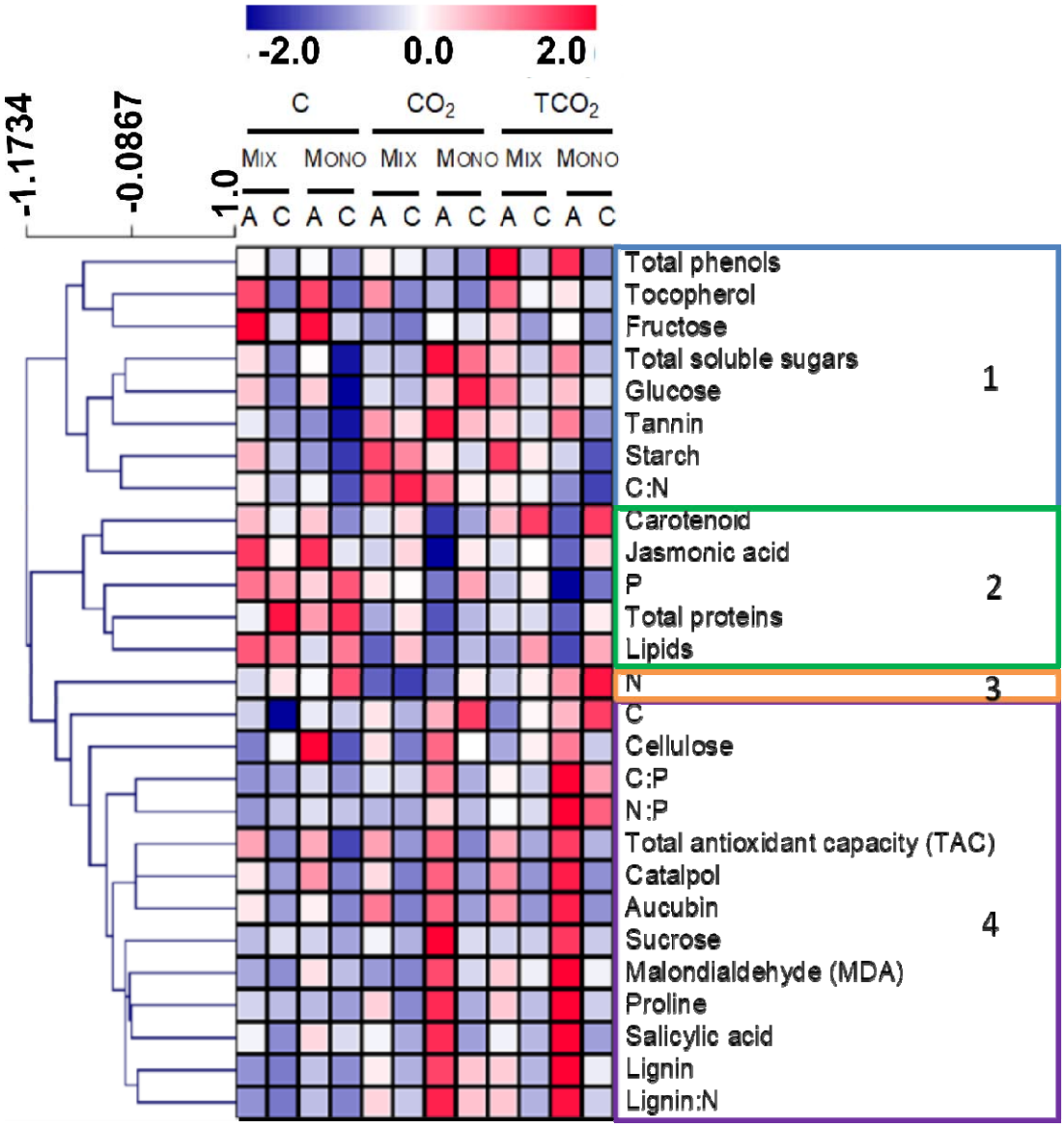


Fig. 1 *Heat map showing the metabolite levels in the leaves of P. lanceolata, normalized to Z-score for each metabolite (blue-white-red heat map). Blue and red colours indicate a low and high metabolite level, respectively. Clustering was based on the Euclidean distance for metabolites. Labels 1-4 and colours indicate the four prominent clusters. Labels C, CO_2 and TCO_2 indicate current climate, elevated CO_2 and combined warming and elevated CO_2, respectively. Plant communities consist of monocultures of P. lanceolata (mono) and mixtures of Lolium perenne and P. lanceolata (mix) with (A) and without aphids (C).*

333 *Causal changes as inferred by structural equation models and univariate analyses*

334 SEMs quantify the strength of both the direct and indirect interactions between the
335 environmental scenarios related to climate change, herbivory and plant phenotype. The first
336 SEM presents the effect of aphid infestation (qualitative effect), climate scenario, plant
337 composition and mildew infestation on the chemical composition (the metabolites of *P.*
338 *lanceolata* were subdivided into the four groups obtained from the hierarchical clustering; see
339 Fig. 1) and the live aboveground biomass of *P. lanceolata* (Fig. 2A). The hypothesized
340 structural relationship with climate scenario affecting mildew infection severity, and climate
341 scenario, plant composition and aphid infestation interactively affecting metabolites and
342 biomass adequately fits the data ($\chi^2 = 42.53$, df = 32, p = 0.101). Fig. 2A and SEM statistics in
343 Table S2.1 (in supplementary material 2) provide us with three insights. Firstly, TCO_2
344 increased the mildew infestation compared to C, but the mildew infestation, in turn, did not
345 affect the metabolites and the alive aboveground biomass of *P. lanceolata*. Secondly, neither
346 climate scenario, nor aphid infestation or plant composition had a significant impact on the
347 live aboveground biomass of *P. lanceolata*. Univariate analyses showed that live biomass
348 tended to be higher in CO_2 than C and was significantly lower in TCO_2 than CO_2, which
349 combined lead to similar values in TCO_2 and C (Fig. 3; see also supplementary material 3.1).

350

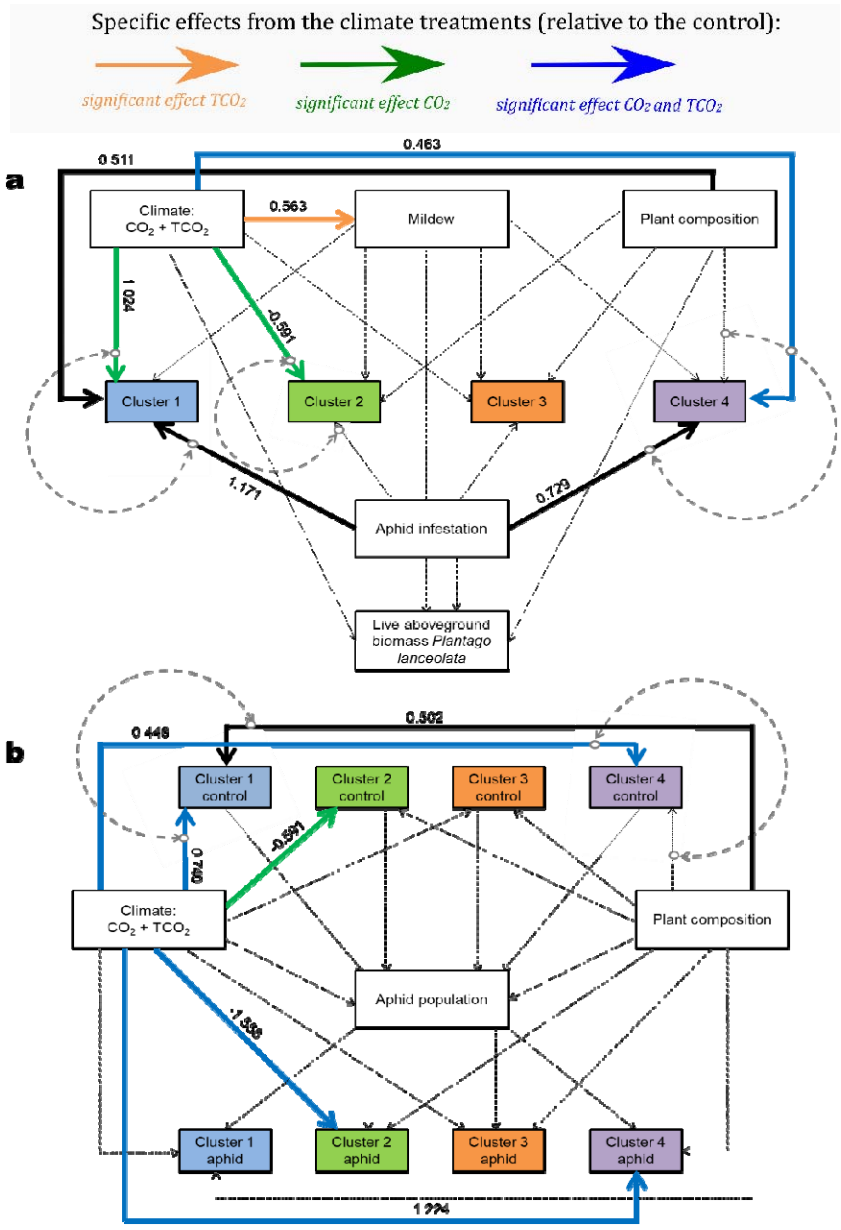
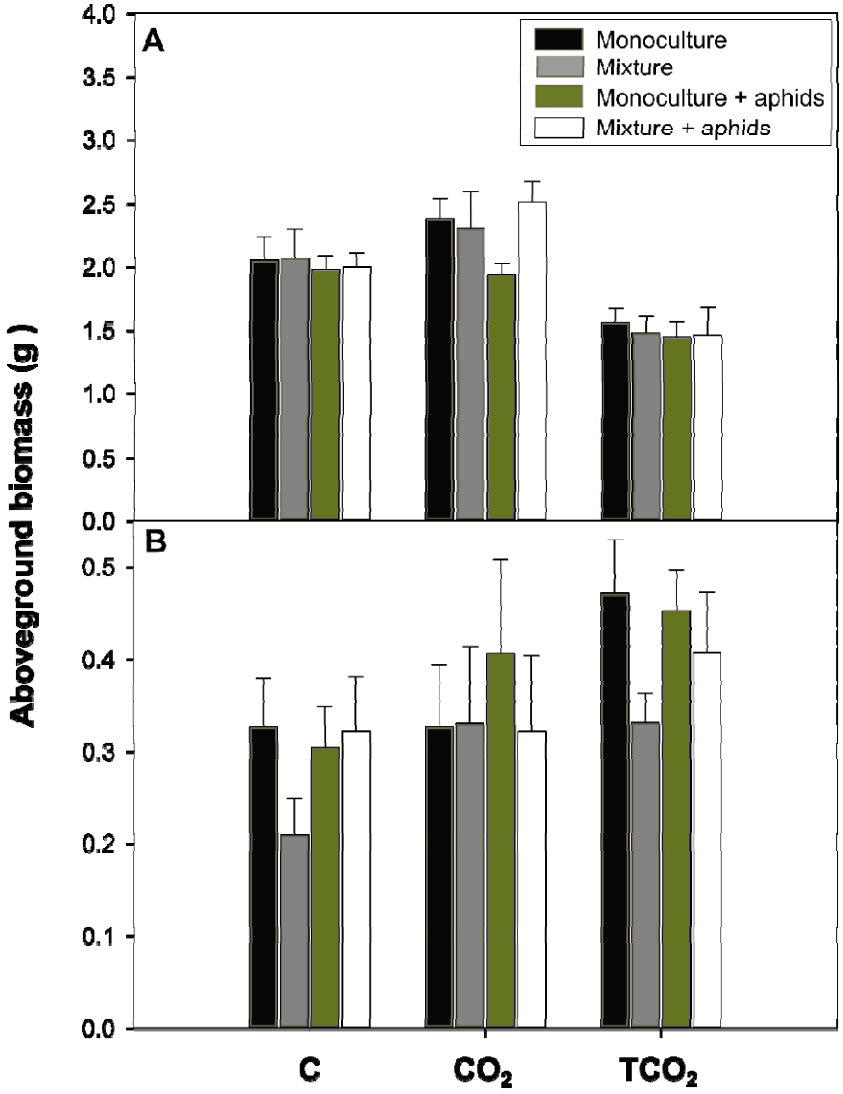


Fig. 2 (a) Structural equation model showing how climate scenario (CO_2 and TCO_2), mildew infestation, plant composition and aphid infestation affect the chemical composition and the live aboveground biomass of P. lanceolata. (b) Structural equation model showing how climate scenario, plant composition and chemical composition of P. lanceolata affect aphid population and how aphid population, in turn, affects the chemical composition of P. lanceolata. The four clusters refer to those obtained by the hierarchical clustering analysis (see Fig. 1). Solid black, green and orange arrows represent significant relationships ($P \leq 0.05$) and dashed grey lines significant interactions. Blue lines stand for significant effects of both CO_2 and TCO_2, green lines for significant effects of CO_2 and orange lines for significant effects of TCO_2. Light grey arrows represent non-significant relationships. Standardized path coefficients are shown next to pathways. For the effect of CO_2 and TCO_2, the average path coefficients are shown. The individual path coefficients of CO_2 and TCO_2 can be seen in Table S2.1 and Table S2.2 (see supplementary material section 2). Metabolites levels, live aboveground biomass of P. lanceolata and number of aphids were scaled before analysis.



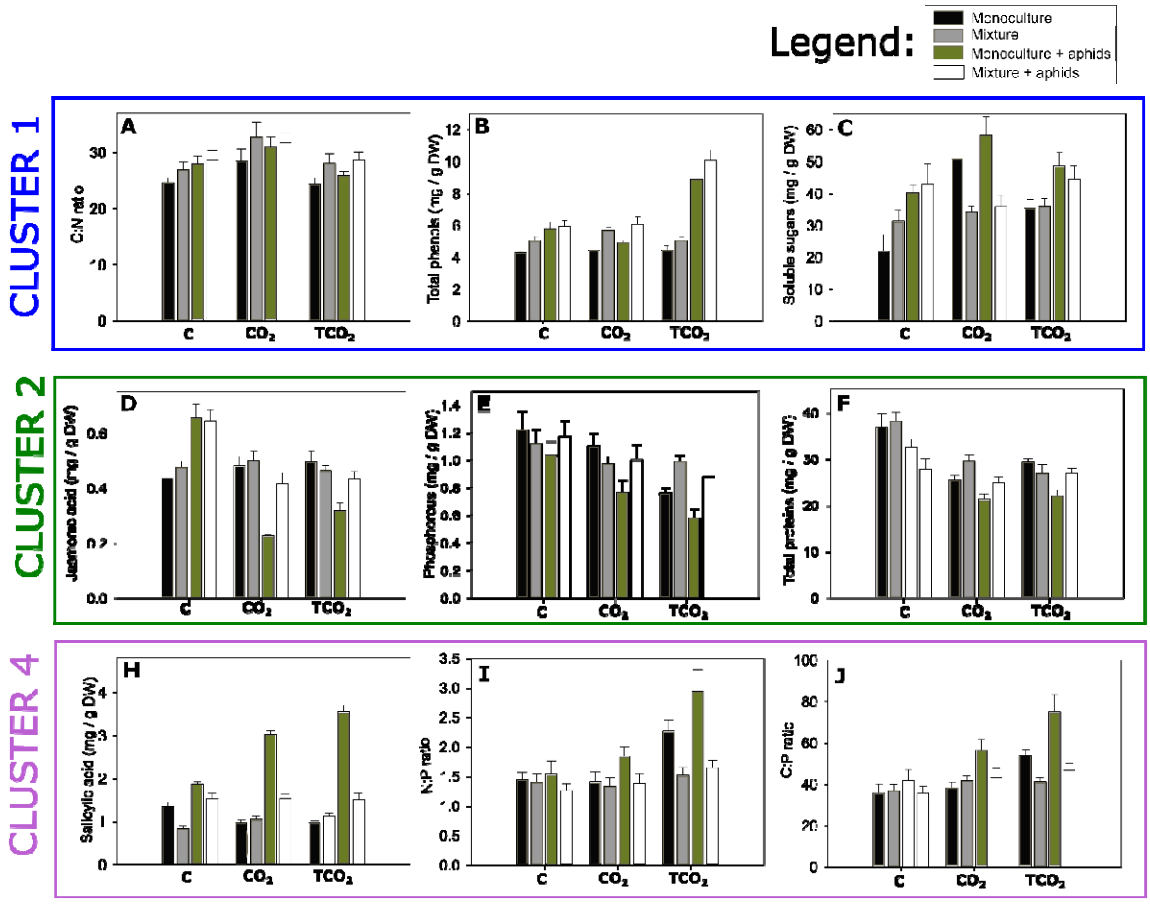
365

366 ***Fig. 3*** *Effect of climate scenario (C, CO_2 and TCO_2), aphid infestation and plant composition on the live (A)*
367 *and dead (B) aboveground biomass of individual P. lanceolata plants. Bars represent means ± SE. Plant*
368 *communities consist of monocultures of P. lanceolata and mixtures of Lolium perenne and P. lanceolata.*

369

370 Thirdly, climate scenario, aphid infestation and plant composition impacted the distinguished
371 clusters in a different manner. We noticed a significant interaction between climate scenario
372 and aphid infestation for metabolites associated with cluster 1 (see supplementary material
373 3.2; Table S3.1 and Fig. S3.1). Aphid infestation increased the concentrations of the
374 metabolites on average and pending on the environmental change scenario between 30 and
375 100% in cluster 1 and plant composition (except for C:N ratio; Fig. 4A-B). The

376 concentrations of glucose and soluble sugars increased at elevated CO_2 but only in
377 monocultures, while TCO_2 did not alter the concentrations of glucose relative to the CO_2
378 treatment but decreased the concentrations of soluble sugars in monocultures (Fig. 4C). In
379 general CO_2, compared to C, increased the concentration of tannin while TCO_2 did not impose
380 further differences. Elevated temperature and CO_2 reduced the concentration of starch (except
381 for mixtures with aphids) and the C:N ratio compared to CO_2.



382

383 ***Fig. 4*** *Effect of climate scenario (C, CO_2 and TCO_2), aphid infestation and plant composition on a selection of*
384 *relevant metabolites from the four distinguished clusters. Bars represent means ± SE. Plant communities*
385 *consist of monocultures of P. lanceolata and mixtures of Lolium perenne and P. lanceolata. As cluster 3 only*
386 *contained one metabolite, it is not depicted here.*

387

388 Also the metabolites in cluster 2 (see supplementary material 3.2 – Table S3.2, Fig. S3.2)
389 differed according to climate scenario and aphid infestation. More specifically, univariate

390 analyses of the metabolites indicates that the concentration of jasmonic acid in C increased
391 with aphid infestation in monocultures and mixtures while aphid infestation in CO_2 and TCO_2
392 decreased it in monocultures (Fig. 4D). Interspecific interactions between *P. lanceolata* and *L.*
393 *perenne* mitigated the negative effect of aphid infestation in these climate scenarios, and
394 hence have a neutralizing effect. Aphid infestation reduced the concentration of P in
395 monocultures, but did not alter it in mixtures (Fig. 4E). Furthermore, also CO_2, compared to
396 C, reduced the concentration of P and the concentration was further reduced in TCO_2.
397 Furthermore, aphid infestation decreased the concentration of proteins in mixtures in C but
398 did not alter the concentrations in monocultures and mixtures in CO_2 and TCO_2 (Fig. 4F). In
399 general, CO_2 compared to C decreased the concentration of proteins irrespective of the plant
400 composition while aphid infestation and TCO_2, compared to CO_2, did not alter it. The
401 concentration of lipids decreased with aphid infestation in monocultures and mixtures in all
402 climate scenarios (except for mixtures in C). Also, CO_2, compared to C, decreased the
403 concentration of lipids in monocultures without aphids and mixtures with aphids, but in
404 TCO_2, compared to CO_2, this concentration increased again to similar levels as C in
405 monocultures without aphids. Carotenoid concentration increased in TCO_2, compared to C
406 and CO_2. Leaf nitrogen was not impacted by any of the treatments (cluster 3; see
407 supplementary material 3.2; Table S3.3).

408 Aphid infestation, CO_2 and TCO_2 increased the metabolites in cluster 4 (see supplementary
409 material 3.2; Table S3.4, Fig. S3.3 & Fig. S3.4) and specifically inhibited the expression of
410 salicylic acid under the two environmental change scenarios (Fig 4G). Besides these single
411 factor treatments effects, climate scenario, aphid infestation and plant composition also
412 interacted with each other in affecting the metabolites in cluster 4. CO_2 and TCO_2
413 strengthened (salicylic acid and lignin) or induced (proline) an effect of aphid infestation but
414 only in monocultures (Fig. 4G). TCO_2, compared to CO_2 and C increased the N:P and the C:P

ratio in monocultures (Fig. 4H,I). Cellulose levels were always higher in monocultures and aphid infestation but the magnitude of increase was lowers under the two climate-related environmental change scenarios relative to the current conditions (control).

The second SEM presents the effect of climate scenario, plant composition and chemical composition of *P. lanceolata* on the aphid population and whether the aphid population, in turn, altered the chemical composition of *P. lanceolata* (quantitative effect). The mildew infestation was left out of this analysis since it is a parasite of the plants, not of the aphids. The results of this SEM model show the following goodness of fit statistics: $\chi^2 = 77.52$, df = 80 and $p = 0.558$ (Table S2.2 and Fig. 2b), which provides us with two insights. Firstly, climate scenario, plant composition and the chemical composition of *P. lanceolata* did not affect the number of aphids. Secondly, the aphid population size in turn, did not change the chemical composition of *P. lanceolata*. Bringing together the findings from both SEM models, we conclude that the treatment aphid infestation altered the chemical composition of *P. lanceolata* but the number of aphids had no effect on it.

Discussion

A future climate clearly altered leaf nutritional quality and the expression of defence molecules, while aphid infestation overall impaired nutritional requirements for insect herbivores. Despite the strong metabolic changes as induced by climate-related environmental change, no feedbacks on aphid population performance and herbivory were detected. More importantly, we show that interspecific plant competition neutralizes the positive effects of elevated CO_2 on the production of defence molecules in *P. lanceolata*. Induced effects of climate-related environmental change on plant performance can thus be altered and even nullified by competitive and food web interactions. We first discuss the general herbivore-induced responses followed by an in-depth discussion of their dependency on the imposed climate and competition treatments.

Herbivore-induced responses

Our experiment, as conducted under unique semi-natural conditions confirmed in first instance findings that aphid infestation reduces N, P, total proteins and lipids in *P. lanceolata* leaves, thereby impairing nutritional requirements of insect herbivores (Schoonhoven *et al.*, 2005). In contrast to previous research (Walling, 2000), however, aphid infestation promoted the salicylic acid and jasmonic acid pathway (Kuśnierczyk *et al.*, 2011; Morkunas & Gabryś, 2011) as parallelized by the increased production of catalpol, aucubin, tannin, lignin and cellulose.

Aphid herbivory increased oxidative stress in plants as well (Wu & Baldwin, 2010). We assessed the induction of oxidative stress by measuring MDA, which aphid herbivory enhanced considerably. A common defence response of plants against oxidative stress

454 includes the increase of antioxidant metabolites. Indeed, TAC and the well-known antioxidant
455 molecules phenols and tocopherols increased due to aphid herbivory. Moreover, *P. lanceolata*
456 responded to herbivory by increasing soluble sugars. At higher concentrations, also sugars can
457 act as antioxidants and may play a signalling role in regulating stress and defence responses
458 (Gómez-Ariza *et al.*, 2007, Peshev *et al.*, 2013). The induced resistance against oxidative
459 stress associated with aphid feeding appeared sufficiently effective to constrain biomass loss
460 from herbivory.

461

462 ***Climate dependency of the plant phenotype***

463 Elevated CO_2 and temperature did not change the impact of herbivory on *P. lanceolata*
464 biomass. This implies that warming and CO_2 did not affect the net interaction strength
465 between plants and herbivores under semi-natural conditions. In contrast to earlier studies
466 (Van De Velde *et al.*, 2016), our experiment allows a profound analysis of the underlying
467 biochemical processes.

468 First, we found elevated CO_2 to increase starch and soluble sugars (in monocultures) and to
469 lower leaf proteins, hereby increasing the C:N ratio. In addition, elevated CO_2 reduced P and
470 consequently increased the C:P ratio (in monocultures), thereby reducing the plant's nutritive
471 value to herbivores (Huberty & Denno, 2006; Lincoln *et al.*, 1986; Mattson, 1980;
472 Schoonhoven *et al.*, 2005). We demonstrate here that these effects can be countered when
473 elevated CO_2 is accompanied with a temperature rise. This contradicts findings based on a
474 limited set of primary metabolites (Murray *et al.* 2013), but urges to apply broader phenomic
475 approaches to fully understand plant responses to climate change.

476

Second, elevated CO_2 increased the defence molecules lignin and tannin, especially under aphid infestation. The induction of plant chemical defences (Bidart-Bouzat & Imeh-Nathaniel, 2008; Bidart-Bouzat *et al.*, 2005) is thus not consistent for all defence molecules, and are consequently anticipated to render prediction on future plant-insect herbivore interaction highly complex (Agrawal, 1999). In our study, the higher C:N ratio reflected higher lignin and tannin levels. While there is a consensus that the carbon-nutrient-balance hypothesis fails as predictive framework (Hamilton *et al.*, 2001; Lindroth, 2012), alternative hypotheses have been proposed stating that resource utilization for chemical defence is linked with photosynthesis, hormone regulation and the control of gene expression (Zavala *et al.*, 2017). We here provide evidence for this alternative mechanism as we found CO_2 to alter phytohormones (i.e., salicylic acid and jasmonic acid) that play an important role in promoting compounds responsible for herbivore defence (Wu & Baldwin, 2010). Since no further quantitative changes were observed under higher temperature, we can confidently conclude that climate-change-induced changes in metabolic profiles are more due to enhanced CO_2 than to temperature increases acting in parallel.

Competition: a biotic interaction mediating climate-induced effects

The presence and identity of neighbouring plants can influence the quality of the host plant by altering primary and secondary chemistry (Barton & Bowers, 2006; Broz *et al.*, 2010, Lankau, 2012; Thorpe *et al.*, 2011). We provide evidence that these biotic interactions are additionally able to alter climate-induced phenomic changes, and by that further impacting ecological interactions under climate-related environmental change. Energetic growth-defence-trade-offs have usually been attributed to these competition effects (Herms & Mattson, 1992), but recent studies demonstrated the role of light limitation for plant defence downregulation (Ballaré, 2014; Campos *et al.*, 2016). More specifically, a reduced red to far-red ratio downregulates

defences by a simultaneous inhibition of the jasmonic acid and the salicylic acid pathway (Wit *et al.*, 2013). The lower levels of defence molecules and salicylic acid in *P. lanceolata* support this hypothesis. Competition did, however, also counter the CO_2-induced changes in primary and secondary metabolites, as well as those induced by herbivores, rendering an understanding of plant defences far from trivial (de Vries *et al.*, 2017). It is well-understood that bottom-up effects from plant quality to herbivore abundance cannot be generalized across the feeding guilds (Bezemer & Jones, 1998; Robinson *et al.*, 2012; Stiling & Cornelissen, 2007), but phloem feeders such as aphids were at least expected to be facilitated. Despite the complex, high-dimensional changes in plant metabolomes under competition and climate change, the overall effects on aphid population sizes were small to non-existent.

We here thus demonstrate that the joint action of atmospheric factors associated with climate change and biotic interactions are able to induce pronounced changes in plant metabolomes and biomass, but that translation of these biochemical changes towards ecological responses appears non-trivial. Climate-change-imposed changes are shown to be completely neutralized when common biotic interactions are taken into consideration. Plant-plant interactions thus add a layer of complexity in mechanistic studies of climate change effects on plant-enemy ecological interaction.

520 **Supplementary data:**

521 **Supplementary material 1:** Schematic representation of the experimental setup

522 **Supplementary material 2:** Metabolite quantification methods and SEM statistics

523 **Supplementary material 3:** Results from the univariate GLM analyses (statistics and
524 figures)

525

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533

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705

706 **Figure legends**

707

708 **Fig. 1** Heat map showing the metabolite levels in the leaves of *P. lanceolata*, normalized to Z-
709 score for each metabolite (blue-white-red heat map). Blue and red colours indicate a low and
710 high metabolite level, respectively. Clustering was based on the Euclidean distance for
711 metabolites. Labels 1-4 and colours indicate the four prominent clusters. Labels C, CO_2 and
712 TCO_2 indicate current climate, elevated CO_2 and combined warming and elevated CO_2,
713 respectively. Plant communities consist of monocultures of *P. lanceolata* (mono) and
714 mixtures of *Lolium perenne* and *P. lanceolata* (mix) with (A) and without aphids (C).

715 **Fig. 2** (a) Structural equation model showing how climate scenario (CO_2 and TCO_2), mildew
716 infestation, plant composition and aphid infestation affect the chemical composition and the
717 live aboveground biomass of *P. lanceolata.* (b) Structural equation model showing how
718 climate scenario, plant composition and chemical composition of *P. lanceolata* affect aphid
719 population and how aphid population, in turn, affects the chemical composition of *P.*
720 *lanceolata.* The four clusters refer to those obtained by the hierarchical clustering analysis
721 (see Fig. 1).

722 Solid black, green and orange arrows represent significant relationships ($P \leq 0.05$) and dashed
723 grey lines significant interactions. Blue lines stand for significant effects of both CO_2 and
724 TCO_2, green lines for significant effects of CO_2 and orange lines for significant effects of
725 TCO_2. Light grey arrows represent non-significant relationships. Standardized path
726 coefficients are shown next to pathways. For the effect of CO_2 and TCO_2, the average path
727 coefficients are shown. The individual path coefficients of CO_2 and TCO_2 can be seen in
728 Table S1 and Table S2 (see supplementary material section 2). Metabolites levels, live
729 aboveground biomass of *P. lanceolata* and number of aphids were scaled before analysis.

730 **Fig. 3** Effect of climate scenario (C, CO_2 and TCO_2), aphid infestation and plant composition
731 on the live (A) and dead (B) aboveground biomass of *Plantago lanceolata*. Bars represent
732 means ± SE. Plant communities consist of monocultures of *P. lanceolata* and mixtures of
733 *Lolium perenne* and *P. lanceolata*.

734 **Fig. 4** Effect of climate scenario (C, CO_2 and TCO_2), aphid infestation and plant composition
735 on the on a selection of relevant metaboloites from the four distinguished clusters. Bars
736 represent means ± SE. Plant communities consist of monocultures of *P. lanceolata* and
737 mixtures of *Lolium perenne* and *P. lanceolata*.