Host-mediated, cross-generational intraspecific competition in a herbivore species

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Conspecific insect herbivores co-occurring on the same host plant interact both directly through interference competition and indirectly through exploitative competition, plant-mediated interactions and enemy-mediated interactions. However, the situation is less clear when the interactions between conspecific insect herbivores are separated in time within the same growing season, as it is the case for multivoltine species. We hypothesized that early season herbivory would result in reduced egg laying and reduced performance of the next generation of herbivores on previously attacked plants. **We tested this hypothesis in a choice experiment with box tree moth females (*Cydalima perspectalis* Walker, Lepidoptera: Crambidae). These females were exposed to box trees (*Buxus sempervirens* L., Buxaceae) that were either undamaged or attacked by conspecific larvae earlier in the season.** We then compared the performance of the next generation larvae on previously damaged *vs* undamaged plants. Previous herbivory had no effect on oviposition behaviour, but the weight of next generation larvae was significantly lower in previously damaged plants. There was a negative correlation between the number of egg clutches **laid on plants by the first generation** and **the performance of the next generation larvae**. Overall, our findings reveal that early season herbivory reduces the performance of conspecific individuals on the same host plant later in the growing season, and that this time-lagged intraspecific competition results from a mismatch between the oviposition preference of females and the performance of its offspring.

1 Main text

1.1 Introduction

Biotic interactions are strong factors affecting the fitness of interacting individuals, even when interactions are delayed in time or do not imply direct contact between individuals. Such interactions can be found in both plants through plant-soil feedbacks (Putten et al., 2016) and in animals (Fisher et al., 2019; Pfennig & Pfennig, 2020). For instance, insect herbivores exploiting the same plant can compete for food, even when interactions among individuals are separated in time (Kaplan & Denno, 2007). Insects may reduce the impact of interspecific competition by avoiding crowded plants, or plants that have been previously consumed by herbivores, which assumes that they can detect competitors or their effects on plants (Shiojiri & Takabayashi, 2003; De Moraes et al., 2001). For many species, the choice of the oviposition site by mated females is crucial in this respect. The preference-performance hypothesis — *aka* the '*mother knows best hypothesis*' — states that female insects evolved host searching behaviour that leads

them to oviposit on hosts where their offspring do best (Gripenberg et al., 2010). A good match between the preference of a mated female for a given plant and the performance of its offspring developing on the same plant implies that females can recognize cues that correlate with larval performance, for instance those related to plant defenses and nutritional quality. Yet, these cues can be largely modified by the simultaneous or sequential presence of other competing herbivores (Bultman & Faeth, 1986; Nykänen & Koricheva, 2004; Abdala-Roberts et al., 2019; Visakorpi et al., 2019). Therefore, initial herbivory **by a given species** may have time-lagged consequences on the preference and performance of herbivores **of another species** that subsequently attack the same plant in the same growing season (Poelman et al., 2008; Stam et al., 2014). However, while such time-lagged *interspecific* interactions between herbivores have long been documented (Faeth, 1986), surprisingly much less is known about delayed *intraspecific* interactions in multivoltine species having several generations per year.

Previous herbivory generally reduces the performance of later arriving herbivores on the same plant (Moreira et al., 2018), **although the opposite effect can also be observed (Sarmento et al., 2011; Godinho et al., 2016). Reduced performance of herbivores on previously damaged plants could occur through both exploitative competition — whereby the first attacking herbivore depletes the resource available to forthcoming herbivores (Kaplan & Denno, 2007) — or through changes in plant traits (Hilker & Fatouros, 2015; Abdala-Roberts et al., 2019; Marchand & McNeil, 2004; Blenn et al., 2012; Fatouros et al., 2012). By triggering an hormonal response and inducing the production of anti-herbivore defenses as well as resource reallocation in plant tissues**, herbivory may induce changes in plant quality that generally reduce the performance of late coming herbivores (Agrawal, 1999; Abdala-Roberts et al., 2019; Wratten et al., 1988). **Such an affect has long been documented in interspecific interactions(Kaplan & Denno, 2007; Moreira et al., 2018), but also in intraspecific interactions.** For instance, prior damage by the western tent caterpillar *Malacosoma californicum* Packard (Lepidoptera: Lasiocampidae) induces the regrowth of tougher leaves acting as physical defenses and reducing the fitness of the next tent caterpillars generation (Barnes & Murphy, 2018).

Previous herbivory can influence the oviposition preference of later herbivores. Several studies have demonstrated that mated females can discriminate host plants that have been previously attacked by insect herbivores (Wise & Weinberg, 2002; Stam et al., 2014; Moura et al., 2017; Barnes & Murphy, 2018; Moreira et al., 2018; Weeraddana & Evenden, 2019), thereby reducing competition between herbivores separated in time. Mated females can directly detect the present, past and possibly future presence of competitors themselves. For instance, Averill & Prokopy (1987) showed that female *Rhagoletis pomonella* Walsh (Diptera: Tephritidae) marks its oviposition site with an epideictic pheromone that deters conspecific females from laying eggs, thus reducing intraspecific competition at the larval stage. The frass of several Lepidoptera species was also found to act as an oviposition deterrent (Jones & Finch, 1987; Hashem et al., 2013; Molnár et al., 2017). **Mated females may also detect herbivory-induced changes in the physical and chemical characteristics of attacked plants, and consequently avoid laying eggs on less suitable plants**. However, several authors reported a mismatch between prior herbivory effects on female oviposition preference *vs* larval growth, consumption or survival of their offspring (Wise & Weinberg, 2002; Bergamini & Almeida-Neto, 2015; Martinez et al., 2017). For instance, Weeraddana and Evenden (2019) found that herbivory by the diamondback moth, *Plutella xylostella* (L.) (Lepidoptera: Plutellidae) on canola plants (*Brassica napus* L.) had no effect on subsequent oviposition by the bertha armyworm, *Mamestra configurata* Walker (Lepidoptera: Noctuidae) whereas its larvae had reduced growth on previously damaged plants. **Thus, in order to quantify the effect of prior herbivory on subsequent herbivore performance, we need to assess how it affects both female choice and progeny performance in attacked and non-attacked hosts.**

In the present study, we investigated the consequences of box tree (*Buxus* spp.) defoliation by the first generation of the box tree moth (BTM) *Cydalima perspectalis* Walker (Lepidoptera: Crambidae) larvae on (*i*) the oviposition behaviour of **the adults emerging from those larvae** and (*ii*) on the larval performance in the next generation. **Specifically, we hypothesized that plants that had previously been attacked by conspecific larvae would (i) receive fewer eggs (*i.e.* reduced preference) and (ii) host smaller larvae and chrysalis (*i.e.* reduced performance) of the next generation than previously undamaged plants. Our experimental design allowed us to separate the effects of**

previous herbivory on both preference and performance of conspecific herbivores attacking the same plant in sequence. By doing so, our study brings new insights into the understanding of cross-generational intraspecific competition in insect herbivores and further challenges the '*mother knows best hypothesis*'.

1.2 Materials and methods

1.2.1 Natural history

The BTM is a multivoltine moth species introduced to Europe in 2007 from Asia (Wan et al., 2014). In its native range, BTM larvae can feed on different host genera, whereas in Europe they feed exclusively on box trees (Wan et al., 2014). In the introduced area, **BTM larvae overwinter in cocoons tied between two adjacent leaves, mainly in the third instar. Therefore, defoliation restarts in early spring** at the beginning of the growing season. In Europe, damage is aggravated by the fact that the BTM has 3-4 generations a year (Kenis et al., 2013; Matošević et al., 2017). When several pest generations successively defoliate the same box tree, there are no leaves left to eat and the caterpillars then feed on the bark, which can lead to the death of the host tree (Kenis et al., 2013; Wan et al., 2014; Alkan Akıncı & Kurdoğlu, 2019).

1.2.2 Biological material

In spring 2019, we obtained box trees from a commercial nursery and kept them in a greenhouse at INRAE Bordeaux forest research station. Box trees were on average 25 cm high and 20 cm wide. We transferred them into 5 L pots with horticultural loam. For two months, we watered them every four days from the above (*i.e.* watering leaves too) to remove any potential pesticide remain.

We initiated BTM larvae rearing with caterpillars collected in the wild in early spring 2019, **corresponding to those that had overwintered**. We reared them at room temperature in 4320 cm^3 plastic boxes, and fed them *ad libitum*, with branches collected on box trees around the laboratory. **We used the next generation larvae to induce herbivory on box tree plants (experimental treatment, see below) and the subsequent adults for the oviposition experiment.** At 25°C, the larval phase lasts for about 30 days and the BTM achieves one generation in 45 days. Adults live 12-15 days. A single female lays on average 800 eggs.

1.2.3 Experimental design

On June 18th 2019, we haphazardly assigned box trees to *control* and *herbivory* experimental groups. The *herbivory* treatment consisted of $n = 60$ box trees that received five L3 larvae each. Larvae were allowed to feed freely for one week, after which we removed them all from plants. In order to confirm that the addition of BTM larvae caused herbivory, we visually estimated BTM herbivory as the percentage of leaves consumed by BTM larvae per branch, looking at every branch on every plant. We then averaged herbivory at the plant level. **In 8 plants, herbivory data were missing and was imputed as the average of herbivory measured in other plants.** In the herbivory treatment, the percentage of leaves consumed by BTM larvae ranged from 2.2 to 17.2% and was on average 9.1%. The *control* group ($n = 61$) did not receive any BTM larva. On July 8th, we randomly distributed plants of the *herbivory* and *control* treatments on a 11 × 11 grid in a greenhouse (*i.e.* total of 121 plants). We left 40 cm between adjacent pots, which was enough to avoid any physical contact between neighbouring plants **(Figure 1, Figure 2)**.

The same day, we released *ca* 100 BTM moths that had emerged from chrysalis less than two days before (*i.e.*, an uncontrolled mix of males and females). **We released moths at the four corners of the experiment to reduce the risk of spatial aggregation**. Moths were allowed to fly freely within the greenhouse. They could feed on small pieces of cotton imbibed with a sugar-water solution, disposed on the ground in the greenhouse.
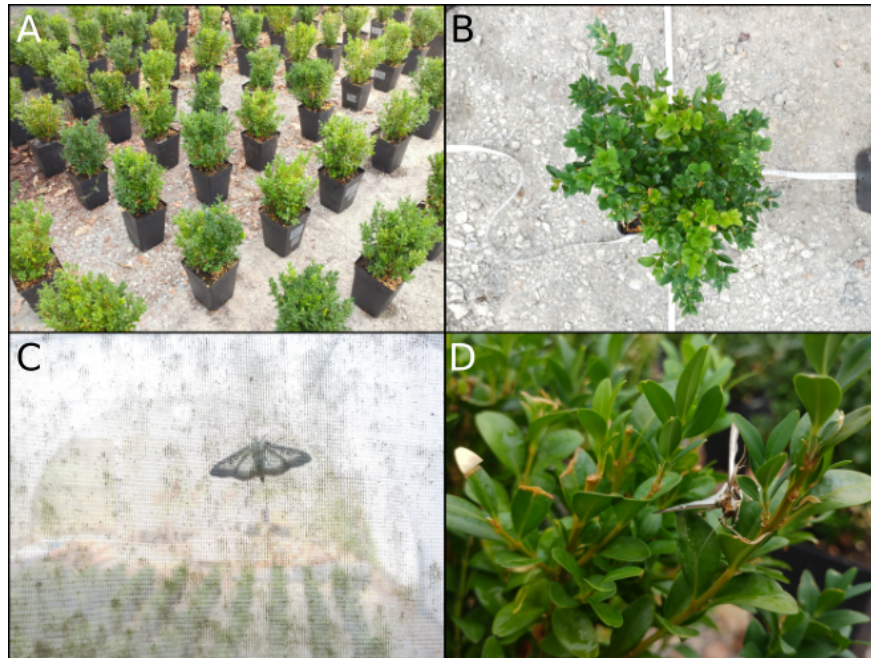
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Figure 1: Some photos, because it is nice to see what an experiment looked like. The two top photos (A, B) illustrate the experimental design and in particular distance among potted plants. Photo C is a view of the greenhouse from the outside, with an adult box tree moth in the foreground, and potted plants in the background. Photo D shows an adult box tree moth on a box tree branch, shortly after it was released.

It is important to note that at the time we released moths, there were no larvae feeding on experimental box trees anymore. In addition, at this time, plants in the herbivory treatment had been cleared of caterpillars for three weeks **(corresponding to the duration of the chrysalis stage)** during which they were watered every two to three days from above. Although larval frass may have been present in pots submitted to the herbivory treatment, it should have been washed out from leaves. Finally, we carried out our experiment in an enclosed greenhouse in which the potential effect of natural enemies on BTM behaviour can be neglected. The consequences are that any effect of prior herbivory on subsequent oviposition behaviour and larval performance should have been independent of cues emitted by BTM larvae themselves or by their frass (Sato et al., 1999; Molnár et al., 2017) and therefore were only plant-mediated.

1.2.4 BTM host choice

In order to test whether initial defoliation of focal plants influenced host choice for oviposition by BTM females, we counted egg clutches on every branch of every box tree on July 17th. Once eggs were counted, we moved box trees to another greenhouse. In order to prevent larvae from moving from one potted plant to another, we installed box trees in plastic saucers filled with a few centimeters of water (renewed regularly).

1.2.5 BTM growth rate

Fifteen days later (July 31st), we haphazardly collected up to five L3 BTM larvae per box tree **(only 6% of plants hosted less than five larvae)**. We kept them in Petri dishes without food for 24h **to make larvae empty their gut** and weighed them to the closest 10 µg. In some Petri dishes, we observed cases of cannibalism such that in some instances we could only weight two larvae (Schillé and Kadiri, *personal observation*). For each plant, we therefore calculated the average weight of a L3 larva, dividing the total mass
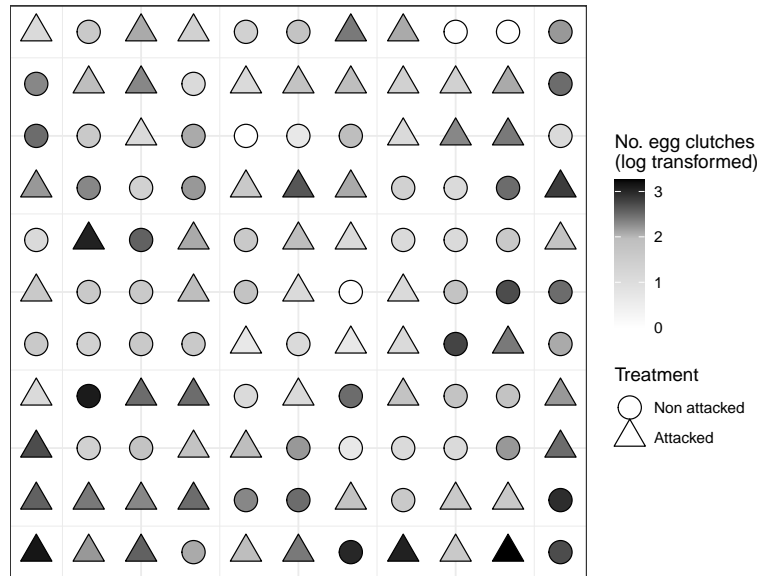
Figure 2: Experimental design. Pots were 40 cm apart. Circles and triangles represent non-attacked (control) and attacked trees. Scale colour represents the number of egg clutches per box tree (log-transformed).

156 by the number of larvae. Because we did not record the day every single egg hatched, we could not quantify
157 the number of days caterpillars could feed and therefore simply analysed the average weight of a L3 larva.

158 Larvae were allowed to complete their development on the potted box trees. After every larvae pupated, we
159 counted the number of chrysalis per box tree and weighted them to the closest 10 µg.

160 **1.2.6 Analyses**

161 All analyses were ran in R using libraries *nlme* and *car* (Team, 2018; Pinheiro et al., 2020; Fox et al., 2016).

162 We first looked for spatial patterns in female BTM oviposition. **We ran a generalized least square model**
163 **(GLS) testing the effect of potted tree location in the experimental design (through their x**
164 **and y coordinates, Figure 2))** on the number of clutches per plant (*log*-transformed) from which we
165 explored the associated variogram using the functions *gls* and *Variogram* in the *nlme* library. There was
166 evidence that oviposition was spatially structured, with strong spatial autocorrelation between 1 and 3m
167 **(Figure S1)**.

168 We tested the effect of prior herbivory on female BTM oviposition (***log*-transformed number of egg**
169 **clutches)** while controlling for spatial non-independence using two independent sets of GLS models. **In**
170 **the first one, we considered prior herbivory as a two-levels factor (attacked vs non-attacked)**
171 **and used the full data set, whereas in the second one, we treated herbivory as a continuous**
172 **variable, excluding data from the control treatment. In both cases,** we had no particular hypothesis
173 regarding the shape of the spatial correlation structure. We therefore ran separate models with different
174 spatial correlation structures (namely, exponential, Gaussian, spherical, linear and rational quadratic), and
175 compared them based on their AIC (Zuur, 2009). For each model, we computed the ΔAIC (*i.e.*, Δ_i) as
176 the difference between the AIC of each model i and that of the model with the lowest AIC (Burnham &
177 Anderson, 2002). We report and interpret the results of the model with the lowest AIC (see *Results*).

178 We then tested the effect of prior herbivory on BTM performance using a two-steps approach. We first used
179 two separate ordinary least square models, **with the mean weight of L3 larvae (*log*-transformed) or**
180 **the mean weight of chrysalis (untransformed) as a response variable, the herbivory treatment**
181 **(non-attacked *vs* attacked) as a two-levels factor and the number of egg clutches as a covariate.**
182 **Then, we restricted the analyses to plants from the herbivory treatment to test the effect of**

the percentage of prior herbivory, number of egg clutches and their interaction on the mean weight of L3 larvae (*log*-transformed) and chrysalis, separately. We deleted non-significant interactions prior to the estimation of model coefficient parameters. Finally, we tested the correlation between mean BTM larval weight and mean BTM chrysalis weight at the plant level using Pearson's correlation.

1.3 Results

We counted a total of 818 egg clutches and 593 larvae **on 117 out of 121 plants (*i.e.* 96.7%). The presence of egg clutches was comparable between control (plants with eggs, *i.e.* 47.1%) and herbivory treatments (49.6%).** However, at individual plant level, the number of egg clutches varied from 0 to 25 (mean $\pm$ SD: 6.76 $\pm$ 5.11, **Figure 2**).

When modelling the effect of prior herbivory on the number of egg clutches **using the full data set**, the best model (*i.e.*, model 5 with $\Delta_i = 0$, **Table 1**) was the model with a rational quadratic spatial correlation. It was competing with three other models with $\Delta_i < 2$ (**Table 1**). **When the analysis excluded data from control plants, the best model was that with a Gaussian spatial correlation (Table 1). It was competing with three other models, including that with a rational quadratic spatial correlation ($\Delta AIC = 0.2$). For sake of consistency, we therefore used this spatial correlation in further analyses, for it was common to the two analyses. The results were comparable with other spatial correlation structures.**

The herbivory treatment had no significant effect on the number of egg clutches per plant, **regardless of whether herbivory was treated as a categorical (model 5, full data set: $F_{1,119} = 2.91$, $P = 0.09$, Figure 3A) or continuous variable (model 5, herbivory treatment only: $F_{1,53} = 0.88$, $P = 0.353$)**.

The mean weight of BTM larvae varied from 6 to 54 mg (mean $\pm$ SD: 20 $\pm$ 9 mg). There was a significant, negative relationship between the number of egg clutches on a box tree and subsequent larval weight (**Table 2**, **Figure 3B**), suggesting intraspecific competition for food. BTM larval weight was lower on box trees that had been previously defoliated (**Table 2**, **Figure 3B**), **regardless of the amount of herbivory (Table 2)**. There was no significant interaction between the herbivory treatment and the number of egg clutches, indicating that intraspecific competition was independent of prior herbivory (**Table 2**). **The results were the same regardless of whether herbivory was treated as a categorical or continuous variable** (**Table 2**).

The mean weight of BTM chrysalis varied from 52 to 210 mg (mean $\pm$ SD: 145 $\pm$ 35 mg, n 104). There was a significant positive correlation between the mean weight of BTM larvae and the mean weight of chrysalis (Pearson's $r = 0.34$, t-value = 3.67, P-value = < 0.001). The effects of herbivory treatment and number of egg clutches on mean chrysalis weight were very comparable to those observed for BTM larvae: BTM chrysalis weight was lower on box trees that had been previously defoliated (Table 2, Figure 3C), regardless of the amount of

Table 1: Summary of AIC of GLS models testing the effect of prior herbivory on the number of egg clutches with different spatial correlation structures, for the full dataset and the data set excluding plants from the control treatment.

Model	Correlation structure	Full model		Herbivory treatment	
		AIC	\Delta	AIC	\Delta
Model 1	Exponential	249.8	0.4	116.6	0.3
Model 2	Gaussian	250.2	0.8	116.3	0.0
Model 3	Spherical	250.9	1.5	117.7	1.4
Model 4	Linear	255.1	5.7	121.2	4.9
Model 5	Rational quadratic	249.4	0.0	116.5	0.2

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²¹⁹ **herbivory. There was a significant, negative relationship between the number of egg clutches**
²²⁰ **on a box tree and subsequent chrysalis weight (Table 2, Figure 3C). There was no significant**
²²¹ **interaction between the herbivory treatment and the number of egg clutches (Table 2).**
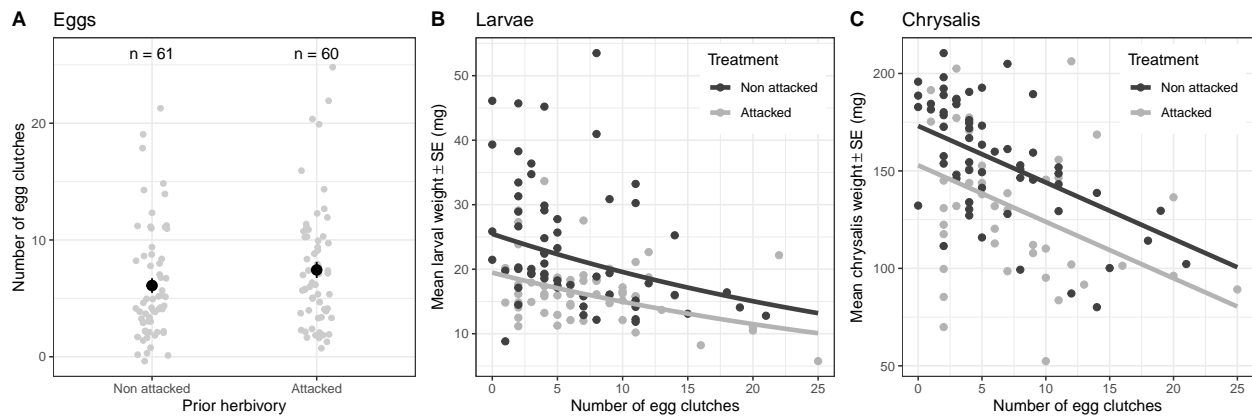

Figure 3: Effects of prior herbivory and conspecific density on (A) the number of egg clutches, (B) L3 larva weight and (C) chrysalis weight. In A, grey dots represent raw data. Black dots and vertical bars represent raw means (+/- SE). In B and C, dots represent raw data. Black and grey curves represent model predictions for control and herbivory treatments, respectively.

1.4   Discussion

²²³ Our findings reveal that early season herbivory reduces the performance of conspecific individuals that
²²⁴ subsequently attack the same host plant later in the plant growing season. This time-lagged intraspecific
²²⁵ competition results from a mismatch between female oviposition preference and the performance of its
²²⁶ offspring.

²²⁷ *Prior herbivory had no effect BTM oviposition choice.* One possibility for female BTM not choosing among
²²⁸ plants may be that that prior herbivory had no effect on box tree characteristics, or that female BTM were
²²⁹ indifferent to them at the time we conducted the experiment.

²³⁰ The first explanation seems unlikely as **we found clear evidence that prior herbivory reduced the**
²³¹ **performance of BTM larvae latter in the season**. This is fully in line with the numerous studies that
²³² have established that insect herbivory induces changes in plant physical and chemical traits, which have
²³³ profound consequences on herbivores or herbivory on the same host plant later in the season (Poelman et al.,
²³⁴ 2008; Abdala-Roberts et al., 2019; Wise & Weinberg, 2002; Stam et al., 2014; but see Visakorpi et al., 2019).
²³⁵ We cannot dismiss the second explanation that BTM females were indifferent to box tree cues related to
²³⁶ earlier herbivory. This may be particularly true in species whose females individually lay several hundred eggs,
²³⁷ **for which spreading eggs among several host plants may be an optimal strategy** (Root & Kareiva,
²³⁸ 1984; Hopper, 1999). Consistently, Leuthardt and Baur (2013) observed that BTM females evenly distributed
²³⁹ egg clutches among leaves and branches, and that oviposition preference was not dictated by the size of the
²⁴⁰ leaves. Assuming that this behavior is reproducible, the close distance between box-trees that we used in
²⁴¹ the present experiment (40 cm) could explain the lack of effect of initial defoliation on BTM oviposition
²⁴² behavior. In addition, Leuthard *et al.* (2013) showed that BTM larvae are able to store or metabolise highly
²⁴³ toxic alkaloid present in box tree leaves. Last, BTM larvae proved to be unable to distinguish between box
²⁴⁴ tree leaves infected or not by the box rust *Puccinia buxi*, while their growth is reduced in the presence of
²⁴⁵ the pathogenic fungus (Baur et al., 2019). Altogether, these results suggest that BTM female moths are not
²⁴⁶ influenced by the amount of intact leaves and probably not either by their chemical quality when choosing the
²⁴⁷ host plant, perhaps because of their strong ability to develop on toxic plants. **It remains however possible**
²⁴⁸ **that BTM adults use other cues to select their hosts such as the presence of conspecific eggs,**
²⁴⁹ **larvae or chrysalis.**

Table 2: Summary of models testing the effect of prior herbivory (with the full data set or the data set restricted to the herbivory treatment) and initial egg clutch density on mean BTM larvae and chrysalis weight

Data set	Response	Predictor	df	F-value	P-value	R^2	Estimate (SE)
Full	Larvae	Number of egg clutches	1, 117	26.31	< 0.001	0.27	-0.026 (0.006)
		Herbivory	1, 117	20.30	< 0.001		-0.269 (0.06)
		Eggs x Herbivory	1, 117	0.73	0.396		
	Chrysalis	Number of egg clutches	1, 100	33.74	< 0.001	0.31	-0.003 (0.001)
		Herbivory	1, 100	12.23	< 0.001		-0.02 (0.006)
		Eggs x Herbivory	1, 100	3.14	0.079		
Herbivory subset	Larvae	Number of egg clutches	1, 56	10.55	0.002	0.14	-0.022 (0.007)
		Herbivory	1, 56	0.16	0.691		-0.003 (0.009)
		Eggs x Herbivory	1, 56	1.74	0.193		
	Chrysalis	Number of egg clutches	1, 41	4.28	0.045	0.06	-0.002 (0.001)
		Herbivory	1, 41	1.08	0.306		-0.001 (0.001)
		Eggs x Herbivory	1, 41	0.39	0.535		

Prior box tree defoliation by the spring generation of BTM larvae reduced the performance of the next generation. Two alternative, non-mutually exclusive mechanisms can explain this phenomenon. First, the reduced performance of individuals of the second generation can have resulted from induced plant defenses. This explanation is in line with studies that have documented in several plant species reduced herbivore performance and changes in plant-associated herbivore communities linked to induced defenses after prior herbivory (Nykänen & Koricheva, 2004; Karban, 2011; Stam et al., 2014). In the case of multivoltine species, negative relationship between prior herbivory and subsequent larva growth rate could indicate intraspecific plant-mediated cross-generation competition between cohorts of herbivores separated in time (Barnes & Murphy, 2018), which could influence herbivore population dynamics and distribution across host individuals. However, **BTM is thought to have broad tolerance to variability in host traits, as suggested by previous observations that BTM larva growth rate did not differ significantly among box-tree varieties (Leuthardt et al., 2013). It is unknown whether herbivory induced changes in host traits are of the same order of magnitude as trait variability among varieties. However, assuming variability among varieties is greater, this result goes against the view that reduced performance of larvae of the summer generation resulted from box tree response to prior herbivory**. Secondly, reduced performance on previously defoliated plants may partly result from food shortage and increased exploitative competition among larvae of the same cohort. Although free living mandibulate herbivores were described to be less sensitive to competition (Denno et al., 1995), the effect of food shortage may have been exacerbated by the small size of box trees and interference competition (Kaygin & Taşdeler, 2019).

We detected a negative relationship between the number of egg clutches laid by BTM female moths and the subsequent growth of BTM larvae. This suggests the existence of intraspecific competition for food within the same cohort. Such competition has already been reported, particularly in leaf-miners (Bultman & Faeth, 1986; Faeth, 1992), which are endophagous insect herbivores whose inability to move across leaves makes them particularly sensitive to the choice of oviposition sites by gravid female. In our study, we prevented larvae from moving from one plant to another and noticed that some box trees were completely defoliated by the end of the experiment. Although we did not record this information, it is very likely that larvae first ran out of food in plants on which several egg clutches were laid. **We are however unable to determine whether the observed intraspecific competition in this cohort was determined by food shortage, or by herbivore-induced changes in resource quality, or both. In addition, we noticed that the number of chrysalis in 32 control plants (out of 61, *i.e.* 52%) was greater that the number of larvae, whereas this only happened in only one previously attacked plant (*i.e.* 2%). This indicates that in spite of our precautions some larvae could move from attacked to control plants (Table 3). Together with the fact that patterns of chrysalis weight were very similar to patterns of larval weight, these findings can be seen as another argument in favor**

of larvae escaping from intraspecific competition on previously attacked plants. However, this idea should be taken with caution as it is possible that such an experimental setup with small potted trees overestimated the effect of intraspecific competition.

Our findings may have profound implications on our understanding of BTM population dynamics. In many Lepidoptera species, all eggs are present in the ovarioles as the adult molt and larva body mass is proportional to fecundity (i.e., 'capital breeders', (Honěk, 1993; Awmack & Leather, 2002)). As a consequence, host plant quality during larval growth and development is a key determinant of individuals fitness (Awmack & Leather, 2002). Although the relationship between plant quality and herbivore fitness may vary among species (Moreau et al., 2006; Awmack & Leather, 2002; Colasurdo et al., 2009), we speculate that herbivory by the first BTM larva generation reduces the fitness of the second BTM generation, and that this effect may be further strengthened where high population density increases intra-specific cross-generational competition (Tammaru & Haukioja, 1996). We may therefore predict a strong density dependent feedback on population growth.

1.5   Conclusion

Our greenhouse experiment provides evidence for negative interaction across and within BTM generations, which are independent of BTM female choice for oviposition site. Such interactions may have consequences on BTM population dynamics and damage on box-trees. On the one hand, the slow-growth-high-mortality hypothesis states that any plant trait reducing the growth rate of herbivores can be seen as a resistance trait, because slow-growing herbivores are longer exposed to their enemies such as parasitoids, spiders or insectivorous birds (Benrey & Denno, 1997; Coley et al., 2006; Uesugi, 2015). It is therefore possible that a stronger top-down control can be exerted by generalist enemies on BTM larvae feeding on previously defoliated hosts which could reduce damage on box-trees. On the other hand, if herbivores take a longer time to complete development, they may cause more damage to plants, in particular to those with low nutritional quality as a result of compensatory feeding (Simpson & Simpson, 1990; Milanovic et al., 2014). The consequences of time-lagged intraspecific competition on the spread of and damage by BTM remain however to be investigated in the field. Particular efforts should be dedicated to the identification of host traits controlling the performance of BTM larvae and the interaction between these traits and the higher trophic level.

1.6   Acknowledgements

1.7   Data accessibility

Raw data as well as codes of statistic analysis are available in supplementary material and on the INRA dataverse: Castagneyrol, Bastien; van Halder, Inge; Kadiri, Yasmine; Schillé, Laura; Jactel, Hervé, 2020, "Raw data for the paper 'Host-mediated, cross-generational intraspecific competition in a herbivore species', https://doi.org/10.15454/KMUX39, Portail Data INRAE, V3.0.

1.8   Conflict of interest

The authors of this preprint declare that they have no financial conflict of interest with the content of this article. Bastien Castagneyrol is one of the *PCI Ecology* recommenders.

1.9 References

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2 Appendix

2.1 Supplementary material

Figure S1 - Semivariogram of the number of egg clutches as a function of distance among box trees.
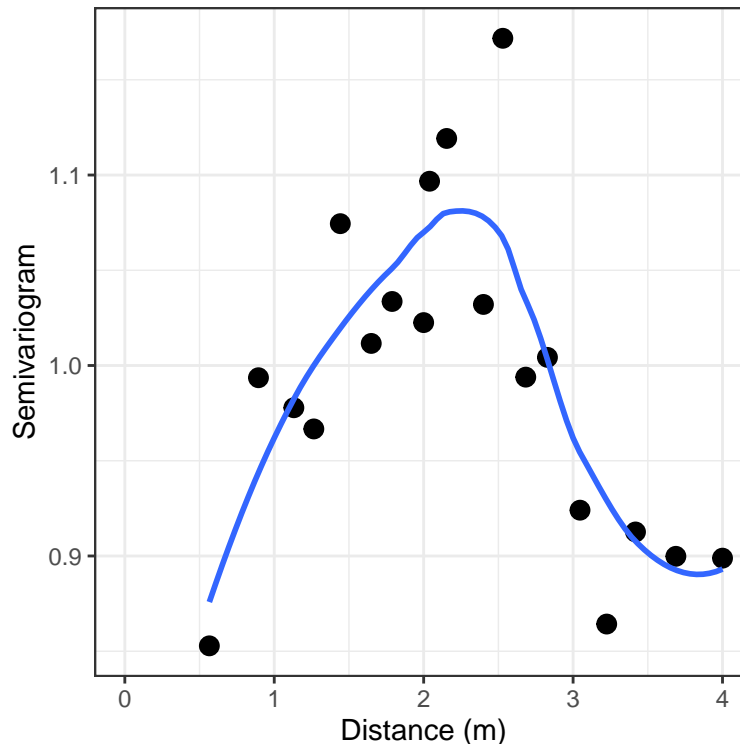
Table 3: Repartition of egg clutches, larvae and chrysalis across box trees with or without prior herbivory. Numbers correspond to mean ($\pm$ sd) and total number of egg clutches, larvae or chrysalis (n).

Response variable	Control	Herbivory treatment
Egg clutches	6.1 (4.87), n = 372	7.43 (5.3), n = 446
Larvae	4.84 (0.61), n = 295	4.97 (0.18), n = 298
Chrysalis	6.8 (5.78), n = 415	1.85 (1.79), n = 111

2.2   Raw data

Table S2 - Raw data used in the present manuscript: `x` and `y` are the position of each box tree in the green house; `Treatment` is the prior herbivory treatment; `Clutch.number` is the total number of egg clutches counted on a given box tree; `N.L3` is the number of retrieved L3 larvae, `L3.mean` is the mean weight of a L3 larvae (g); `N.chrysalids` is the number of retrieved chrysalis; `Chrysalid.mean` is the mean weight of a chrysalis. `Herbivory` is the % of leaves consumed by box tree moth larvae, which was either measured or estimated where raw data was missing (`Herbivory_source`).

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x	y	Treatment	Clutch.number	N.L3	L3.mean	N.chrysalids	Chrysalid.mean	Herbivory	Herbivory_s
1	1	Attacked	22	5	0.0221740	0	NaN	7.3	Estimated
2	1	Attacked	8	5	0.0183980	0	NaN	8.8	Estimated
3	1	Attacked	12	5	0.0187360	1	0.1020100	8.8	Imputed
5	1	Attacked	6	5	0.0146140	0	NaN	16.4	Estimated
6	1	Attacked	10	5	0.0165620	2	0.1101750	10.5	Estimated
8	1	Attacked	20	5	0.0110140	1	0.0961900	8.7	Estimated
9	1	Attacked	4	5	0.0132300	1	0.1438500	7.4	Estimated
10	1	Attacked	25	5	0.0057520	1	0.0891900	11.7	Estimated
1	2	Attacked	12	5	0.0226500	2	0.2062350	6.2	Estimated
2	2	Attacked	10	5	0.0162200	1	0.0524100	0.0	Imputed
3	2	Attacked	9	5	0.0200760	0	NaN	4.2	Estimated
4	2	Attacked	11	5	0.0211200	5	0.1465000	4.8	Estimated
7	2	Attacked	5	5	0.0112560	0	NaN	9.0	Estimated
9	2	Attacked	4	5	0.0161760	1	0.1338800	15.1	Estimated
10	2	Attacked	4	5	0.0173680	1	0.1706800	9.4	Estimated
1	3	Attacked	14	5	0.0159000	4	0.1686525	8.6	Imputed
4	3	Attacked	5	5	0.0159420	2	0.1380100	0.0	Imputed
5	3	Attacked	6	5	0.0121100	1	0.1319100	8.6	Imputed
11	3	Attacked	11	5	0.0101960	1	0.0836300	7.2	Estimated
1	4	Attacked	2	5	0.0111600	2	0.1224050	11.4	Estimated
3	4	Attacked	11	5	0.0157420	0	NaN	8.6	Imputed
4	4	Attacked	11	5	0.0158140	4	0.1557575	6.8	Estimated
6	4	Attacked	2	5	0.0238660	2	0.1728600	11.7	Estimated
8	4	Attacked	5	5	0.0187260	2	0.1527050	10.0	Estimated
11	4	Attacked	8	5	0.0181900	0	NaN	10.3	Estimated
5	5	Attacked	1	5	0.0201820	1	0.1914500	2.7	Estimated
7	5	Attacked	1	5	0.0148320	3	0.1752800	8.9	Estimated
8	5	Attacked	2	5	0.0176160	1	0.0853400	9.1	Estimated
10	5	Attacked	10	4	0.0171925	1	0.1453700	9.0	Estimated
1	6	Attacked	4	5	0.0158160	2	0.1774000	2.2	Estimated
4	6	Attacked	6	5	0.0161500	0	NaN	6.8	Estimated
6	6	Attacked	2	4	0.0148600	0	NaN	10.9	Estimated
8	6	Attacked	2	5	0.0273120	2	0.0698850	17.0	Estimated
2	7	Attacked	20	5	0.0104900	4	0.1364600	3.6	Estimated
4	7	Attacked	7	5	0.0275520	5	0.1299800	12.5	Estimated
6	7	Attacked	6	5	0.0143660	2	0.1127850	9.7	Estimated
7	7	Attacked	2	5	0.0145880	0	NaN	17.2	Estimated
11	7	Attacked	5	5	0.0129260	3	0.1438500	9.5	Estimated
1	8	Attacked	8	5	0.0161140	0	NaN	9.5	Estimated
5	8	Attacked	4	5	0.0336620	4	0.1512050	0.0	Imputed
6	8	Attacked	13	5	0.0136940	1	0.0916800	5.6	Imputed
7	8	Attacked	7	5	0.0119960	0	NaN	7.6	Estimated
11	8	Attacked	16	5	0.0082180	5	0.1013240	5.6	Imputed
3	9	Attacked	2	5	0.0124840	2	0.1309350	0.0	Imputed
8	9	Attacked	2	5	0.0140740	1	0.0996800	5.6	Imputed
9	9	Attacked	9	5	0.0147260	3	0.1120367	15.1	Estimated
10	9	Attacked	10	5	0.0121140	3	0.1454233	7.9	Estimated
2	10	Attacked	6	5	0.0183400	3	0.1203367	0.0	Imputed
3	10	Attacked	9	5	0.0159820	3	0.1078233	6.8	Estimated
5	10	Attacked	2	5	0.0291080	2	0.1450000	10.5	Estimated
6	10	Attacked	5	5	0.0185740	0	NaN	8.6	Estimated
7	10	Attacked	6	5	0.0177680	0	NaN	12.4	Estimated
8	10	Attacked	3	5	0.0149260	1	0.2025200	9.4	Estimated
9	10	Attacked	3	5	0.0195980	2	0.1319950	8.2	Estimated
10	10	Attacked	7	5	0.0157780	2	0.0985400	16.2	Estimated
1	11	Attacked	2	5	0.0161540	5	0.1175720	8.6	Imputed
3	11	Attacked	7	5	0.0190760	5	0.1385900	8.5	Estimated
4	11	Attacked	3	5	0.0178380	0	0.1771000	5.2	Estimated

2.3   R codes used to generate this report

```r
library(tidyverse)
library(knitr)
library(kableExtra)
library(nlme)
library(car)
library(here)

wd = here()

my.ggplot <- function(){
  theme_bw() +  theme(legend.key = element_blank())
}

knitr::opts_chunk$set(echo= F, warning= F, message = F, results = "hide",
                      fig.width=4, fig.height=4, dpi= 100, fig.pos = "H",
                      fig.path = paste0(wd, '/Figures/'),
                      output.dir = paste0(wd, '/Outputs/'))

d = read.csv(paste0(wd, '/Data/Castagneyrol_cydalima_data.csv'), header = T)

str(d)

d =
  d %>% mutate(L3.mean = Weight.L3 / N.L3) %>%
  mutate(Treatment = factor(Treatment, levels = c("Non attacked", "Attacked")))

knitr::include_graphics(paste0(wd, '/Figures/Figure 1.png'))

d %>%
  ggplot(aes(x, y, shape = Treatment, fill = log1p(Clutch.number))) +
  # geom_rect(aes(xmin = 1.5, ymin = 1.5, xmax = 10.5, ymax = 10.5), fill = 'grey90') +
  geom_point(size = 6) +
  scale_shape_manual(values = c(21, 24)) +
  my.ggplot() +
  labs(x = "", y = "") +
  scale_fill_gradientn(colours = c('white', 'grey', 'black'),
                       values = c(0,0.6,1), name = 'No. egg clutches\n(log transformed)') +
  theme(axis.ticks.x = element_blank(),
        axis.text.x = element_blank(),
        axis.ticks.y = element_blank(),
        axis.text.y = element_blank())
d.center = droplevels(d[d$x > 1 & d$x < 11 & d$y > 1 & d$y < 11,])
m1a = gls(log1p(Clutch.number) ~ Treatment,
          correlation = corExp(form = ~x + y, nugget = TRUE), data = d, na.action = "na.omit")
m1b = gls(log1p(Clutch.number) ~ Treatment,
          correlation = corGaus(form = ~x + y, nugget = TRUE), data = d, na.action = "na.omit")
m1c = gls(log1p(Clutch.number) ~ Treatment,
          correlation = corSpher(form = ~x + y, nugget = TRUE), data = d, na.action = "na.omit")
m1d = gls(log1p(Clutch.number) ~ Treatment,
          correlation = corLin(form = ~x + y, nugget = TRUE), data = d, na.action = "na.omit")
m1e = gls(log1p(Clutch.number) ~ Treatment,
```

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```r
          correlation = corRatio(form = ~x + y, nugget = TRUE), data = d, na.action = "na.omit")

l1 = list(m1a, m1b, m1c, m1d, m1e)

l2 = list(
m1a_2 = update(m1a, ~ Herbivory, data = d[d$Herbivory > 0,]),
m1b_2 = update(m1b, ~ Herbivory, data = d[d$Herbivory > 0,]),
m1c_2 = update(m1c, ~ Herbivory, data = d[d$Herbivory > 0,]),
m1d_2 = update(m1d, ~ Herbivory, data = d[d$Herbivory > 0,]),
m1e_2 = update(m1e, ~ Herbivory, data = d[d$Herbivory > 0,])
)

AIC_1 = round(unlist(lapply(l1, function(x){AIC(x)})),1)
delta_1 = AIC_1 - min(AIC_1)
AIC_2 = round(unlist(lapply(l2, function(x){AIC(x)})),1)
delta_2 = AIC_2 - min(AIC_2)

Table_AIC = data.frame(AIC_1, delta_1, AIC_2, delta_2)
plot.resid = function(m){
  df = data.frame(f = fitted(m), r = residuals(m))
  A = df %>%
    ggplot(aes(f, r)) + my.ggplot() + geom_point() +
    geom_smooth(method = "lm", se = F) + geom_hline(yintercept = 0)
  B = df %>%
    ggplot(aes(r)) + my.ggplot() + geom_histogram()
  cowplot::plot_grid(A,B)
}


m_larvae = lm(log(L3.mean) ~ Clutch.number * Treatment , d)
m_larvae2 = lm(log(L3.mean) ~ Clutch.number * Herbivory, d[d$Treatment == "Attacked",])
# plot.resid(m_larvae2)

d =
  d %>%
  mutate(Chrysalid.mean = Weight.chrysalids / N.weighted.chysalids)

m_chrys = lm(Chrysalid.mean ~ Clutch.number * Treatment , d)
m_chrys2 = lm(Chrysalid.mean ~ Clutch.number * Herbivory, d[d$Treatment == "Attacked",])
#plot(m_chrys2)
# lapply(l1, function(x) anova(x))
# lapply(l2, function(x) anova(x))
Table_AIC %>%
  mutate(Model = paste('Model', 1:5),
         'Correlation structure' = c('Exponential', 'Gaussian', 'Spherical',
                                      'Linear', 'Rational quadratic')) %>%
  select(Model, `Correlation structure`, everything()) %>%
  kable(col.names = c("Model", "Correlation structure", "AIC", "\\Delta", "AIC", "\\Delta"),
        caption = "Summary of AIC of GLS models testing the effect of prior herbivory on the number of
  kableExtra::kable_styling() %>%
  add_header_above(c(" " = 2, "Full model" = 2, "Herbivory treatment" = 2))
anova(m1e)
```

```r
Fig_3A =
  d %>%
  ggplot(aes(Treatment, Clutch.number)) +
  my.ggplot() +
  geom_point(position = position_jitter(0.1), colour = "grey80") +
  stat_summary(size = 0.6) +
  labs(x = "Prior herbivory", y = "Number of egg clutches") +
  annotate(geom = 'text', x = 1:2, y = 26, label = paste("n =", c(61, 60)))
res_larvae = anova(m_larvae, test = "m")
n = expand.grid(Treatment = levels(d$Treatment), Clutch.number = seq(0, 25))
p = predict(update(m_larvae, ~.- Clutch.number:Treatment), newdata = n)
n$Fit = 1000 * exp(p)

Fig_3B =
  d %>%
  ggplot(aes(Clutch.number, 1000 * L3.mean, colour = Treatment)) +
  my.ggplot() +
  geom_point(size = 2) +
  scale_colour_manual(values = c("grey25", "grey70")) +
  labs(x = "Number of egg clutches",
       y = expression("Mean larval weight" %+-% "SE (mg)")) +
  geom_line(data = n, aes(Clutch.number, Fit), size = 1.5) +
  theme(legend.position = c(0.8, 0.85))
r = with(d, cor.test(Chrysalid.mean, L3.mean))
r_corr = round(r$estimate,2)
r_tval = round(r$statistic,2)
r_pval = ifelse(round(r$p.value,3) < 0.001, "< 0.001", round(r$p.value,3))
res_chrys = anova(m_chrys)
n = expand.grid(Treatment = levels(d$Treatment), Clutch.number = seq(0, 25))
p = predict(update(m_chrys, ~.- Clutch.number:Treatment), newdata = n)
n$Fit = 1000 * p

Fig_3C =
  d %>%
  ggplot(aes(Clutch.number, 1000 * Chrysalid.mean, colour = Treatment)) +
  my.ggplot() +
  geom_point(size = 2) +
  scale_colour_manual(values = c("grey25", "grey70")) +
  labs(x = "Number of egg clutches",
       y = expression("Mean chrysalis weight" %+-% "SE (mg)")) +
  geom_line(data = n, aes(Clutch.number, Fit), size = 1.5) +
  theme(legend.position = c(0.8, 0.85))

cowplot::plot_grid(
  Fig_3A + labs(title = 'Eggs'),
  Fig_3B + labs(title = 'Larvae'),
  Fig_3C + labs(title = 'Chrysalis'),
  ncol = 3,
  labels = c('A', 'B', 'C')
)
f = function(model, response, data_set){
  ANOVA = anova(model, test = "m")
  Fval = function(ANOVA) {round(ANOVA[,4], 2)}
```

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```r
  Pval = function(ANOVA) {ifelse(ANOVA[,5] < 0.001, '< 0.001', round(ANOVA[,5], 3))}

  if(Pval(ANOVA)[3] < 0.05){
    b = round(summary(model)$coefficients[-1,1], 3)
    b_se = round(summary(model)$coefficients[-1,2], 3)
    Estimate = paste(b, ' (', b_se, ')', sep = '')

    }else{

    form = formula(paste("~",
                         paste(attr(model$terms, "variables")[[3]],
                             attr(model$terms, "variables")[[4]],
                             sep ="+")))
    b = round(summary(update(model, formula. = form))$coefficients[-1,1], 3)
    b_se = round(summary(update(model, formula. = form))$coefficients[-1,2], 3)
    Estimate = c(paste(b, ' (', b_se, ')', sep = ''), '')
  }

  return(df =
          data.frame(
            Data = c(data_set, "", ""),
            Response = c(response, '', ''),
            Predictor = c("Number of egg clutches", "Herbivory", "Eggs x Herbivory"),
            df = paste(ANOVA$Df[1:3], rep(ANOVA$Df[4], 3), sep = ", "),
            `F-value` = Fval(ANOVA)[-4],
            `P-value` = Pval(ANOVA)[-4],
            R2 = c(round(summary(model)$adj.r.squared, 2), "", ""),
            Estimate = Estimate))
}


rbind(
  f(m_larvae, response = "Larvae", data_set = "Full"),
  f(m_chrys, response = "Chrysalis", data_set = ""),
  f(m_larvae2, response = "Larvae", data_set = "Herbivory subset"),
  f(m_chrys2, response = "Chrysalis", data_set = "")) %>%
  kable(caption = "Summary of models testing the effect of prior herbivory (with the full data set or th
        col.names = c("Data set", "Response", "Predictor", "df", "F-value", "P-value", "R²", "Estimate
        escape = T, digit = 2) %>%
  kable_styling() %>%
  collapse_rows(columns = 1:2, valign = "top")


m0 = gls(log1p(Clutch.number) ~ x + y, data = d, na.action = "na.omit")
variog0 <- Variogram(m0, form = ~x + y, resType = "pearson", nugget = T)
variog0 %>%
  ggplot(aes(0.4*dist, variog)) +
  my.ggplot() +
  geom_point(size = 3) +
  geom_smooth(se = F) +
  labs(x = "Distance (m)", y = "Semivariogram") +
  xlim(0, 4)
d %>%
```

```
  select(Treatment, Clutch.number, N.L3, N.chysalids) %>%
  gather(Resp, Val, 2:4) %>%
  group_by(Treatment, Resp) %>%
  summarise(N = sum(Val), M = round(mean(Val),2), SD = round(sd(Val),2)) %>%
  mutate(Val = paste(M, ' (', SD, ')', ', n = ', N, sep = '')) %>%
  select(- N, - M, - SD) %>%
  mutate(Resp = factor(Resp, levels = c('Clutch.number', 'N.L3', 'N.chysalids'),
                       labels = c('Egg clutches', 'Larvae', 'Chrysalis'))) %>%
  spread(Treatment, Val) %>%
  kable(col.names = c('Response variable', 'Control', 'Herbivory treatment'),
        caption = "Repartition of egg clutches, larvae and chrysalis across box trees with or without p:
  kable_styling()
d %>%
  rename(N.chrysalids = N.chysalids) %>%
  select(x, y, Treatment, Clutch.number, N.L3, L3.mean, N.chrysalids, Chrysalid.mean, Herbivory, Herbivo
  kable() %>% kable_styling()
```

3 Responses to reviewers' comments

Dear Dr Magalhães,

We would like to thank you for your constructive and helpful comments. We revised the original manuscript accordingly. Significant changes in the manuscript are highlighted in bold characters. You may find our responses to your comments and to the two reviewers below, highlighted with bold characters. Wherever only minor changes were requested, we simply indicated "[R] — Done", otherwise, we justified what we changed, or not, in the manuscript.

We hope that the revised version of our manuscript has addressed every concerns and will be suitable for recommendation.

Best regards,

Bastien Castagneyrol, on behalf of co-authors.

Dear authors,

First of all, I deeply apologize for having taken so long to comment on this manuscript. I hope that the quality of the reviews compensates for this long wait... I found this article interesting and straightforward. I particularly appreciated the scale and nature of the experiment, being an intermediate between a lab and a field experiment. The thorough and insightful comments of the two reviewers also point in the same direction: they both enjoyed the manuscript very much. They do, however, suggest a number of changes that I think should be addressed in the revised version of the manuscript. In particular, they both suggest including more variables in your analysis, which they (and I) believe you can do based on the data you already have. If that is the case, I think it is a cost-effective means to make your article more complete.

[Response] — We followed recommendations made by the two reviewers and yourself and provide below detailed answers wherever necessary

Below I place my own comments, and I would be happy to look at a revised version of this paper soon. I only have two relatively major and a lot of minor comments.

1. I agree with one of the reviewers in that the introduction could be a bit further streamlined. If I understand correctly, the first paragraph is about preference-performance correlations, the second about preference, and the third a bit about performance and then another bit about preference-performance. You also go a bit back and forth concerning the effects of conspecifics and that of heterospecifics. I

don't want to impose my view on the introduction of a paper that is not mine, but I would like to feel that, whatever the structure chosen, it is apparent to the reader. In any case, I would exchange the order of second and third paragraph, first differences in performance and then oviposition choice, because the latter does not make sense without the former and the reverse is not true.

[R] — We changed the order of the two paragraphs as suggested and modified several sentences to improve the flow. Please note that we did not highlight every single change in the manuscript, only the most important additions.

2. I think the reader needs some information on how larval weight correlates (or may correlate) with fitness in this (or related) species. This would allow discriminating among the two main possible interpretations for this data set, namely (a) they don't discriminate because the consequences for fitness are not strong enough or (b) they don't discriminate because they don't have access to reliable cues.

[R] — We now discuss this question:

> *Our findings may have profound consequences on BTM population dynamics. In most of Lepidoptera species, all the eggs are present in the ovarioles as the adult molt and larva body mass is proportional to fecundity (i.e., 'capital breeders', (Honěk, 1993; Awmack & Leather, 2002)). As a consequence, host plant quality during larval growth and development is the key determinant of individuals fitness (Awmack & Leather, 2002). Although the relationship between plant quality and herbivore fitness may vary among species (Moreau et al., 2006; Awmack & Leather, 2002; Colasurdo et al., 2009), we speculate that herbivory by the first BTM larva generation reduces the fitness of the second BTM generation, and that this effect may be further strengthened where high population density increase intra-specific cross-generational competition (Tammaru & Haukioja, 1996).*

3. Minor comments:

4. I would remove "multivoltine" from the title. The cross-generational already gives the idea. . . .

[R] — Done

5. Line 19: replace "proposed" by "offered".

[R] — Done

6. Line 23: unclear if this number of eggs is from the previous or the current brood.

[R] — Changed

7. Line 37: replace "their" by "its".

[R] — Done

8. Line 54: "deters" instead of "deter".

[R] — Done

9. Line 60: incidentally, previous herbivory can also lead to increased performance in subsequent infestations (e.g., Sarmento et al 2011 Ecol Lett, Godinho et al 2016 Oecologia). This is just a side comment, you don't need to include this in the paragraph. . .

[R] — Thank you for these references.

10. Lines 94-100. This paragraph is a bit confusing. First, I suggest placing the sentences on the biological details of the system (lines 96-98) elsewhere, maybe in the very beginning of the Material and Methods section. Second, it is not very clear to me when were the moths placed on the experimental trees. Is "the overwintering generation" the same as the "caterpillars collected in the wild"? If so, please be clear about this. Also, I guess that by "their" adults you mean "the adults emerging from those larvae"? (also commented by one of the reviewers).

21

[R] — We restructured this paragraph, adding a new subsection ("*natural history*") and clarifying the description of first and second BTM generations (section "biological material").

11. Line 104: I would state "plants" instead of "plant individuals".

[R] — Done

12. Line 111: you seem to use "chrysalis" and "chrysalids" interchangeably. If these terms refer to the same thing (I guess they do. . .) please choose one.

[R] — Done

13. Results: Did you count the number of egg clutches per plant with at least one clutch or per experimental plant in general? That is, could there be a difference in the number of plants with no egg clutches among treatments?

[R] — We screened every single plant and counted the number of egg clutches on all plants. We added the information on the presence/absence of eggs to the '*results*' section.

14. I would not discard the data concerning chrysalids so easily. It is indeed a pity that you cannot discriminate whether more larvae died in one treatment vs the other or if there was active migration, as you state. But in any case, this means that more individuals of this second 'generation' are eventually found on the previously clean plants, and this is an interesting result per se. I would at least discuss this a bit further in the Discussion.

[R] — We do agree this is very unfortunate we have not been satisfyingly efficient in preventing larvae from moving among plants. Although we have been reluctant to present this data in first instance, we now reinjected them back into the manuscript, with words of caution in the discussion. Changes can be seen throughout the text in the "methods / analyses", "*results*" and "*discussion*" sections.

15. Lines 180-183: Maybe rephrase as to use a more fluid text style. Ex: One possibility for female BTM not choosing among plants may be that. . .

[R] — Done. Indeed, it reads smoother.

16. Lines 182-190: I think the main argument against this hypothesis comes from your own data: larval weight differs among treatments. Assuming this is correlated with fitness, there are consequences for the moths of their mothers' choice.

[R] — Yes! thank you for mentioning this. We have added this obvious argument.

17. Line 193: I don't understand why laying 200 eggs corresponds necessarily to a bet-hedging strategy. Maybe rephrase?

[R] —Done.

18. Lines 203-204: This paragraph is about the possible absence of cues, not about the possible absence of fitness consequences, so this sentence is best placed in the previous paragraph.

[R] — This paragraph, starting with "*Prior box tree defoliation by the spring generation of BTM larvae reduced the performance of the next generation*" is about herbivore growth. We dealt with possible absence of cues in the previous paragraph.

19. Line 206: I would remove "trait-mediated" from this sentence because I am not convinced that the dichotomy between the two explanations rests on this. Instead, I think that the two possible explanations are past vs current competition. Also is there a possibility to obtain the density of larvae in the two treatments? That is, the number of larvae per intact lead?

[R] — We removed "*trait-mediated*". Although we acknowledge that this would have been a powerful way to further address competition, we did not precisely count the number of larvae per plant or per shoot, mostly to avoid disturbance.

22

20. Lines 213-215: does this mean that larval weight has no effect on fitness / population growth? Please clarify.

[R] — Done:

> *the BTM is thought to have broad tolerance to variability in host traits, as suggested by previous observations that BTM larva growth rate did not differ significantly among box-tree varieties (Leuthardt et al., 2013). It is unknown whether herbivory induced changes in host traits are of the same order of magnitude as trait variability among varieties. However, assuming variability among varieties is greater, this result goes against the view that reduced performance of larvae of the summer generation resulted from box tree response to prior herbivory*

21. Line 219: although I agree with one of the reviewers that the possibility that food shortage may lead to cannibalism is fascinating, I would remove this sentence unless you have hard data on which to base this statement. In particular, if this were to be true, you would need to explain (a) why you still find the same overall number of larvae alive between the two treatments and (b) whether it is expected that this cannibalism does not compensate for food shortage in terms of larval weight. Overall, I think that this observation opens too many doors, so either it is solid or it better be left out of the Discussion.

[R] — We deleted the mention to cannibalism, because this is true we do not have hard data ta back it up.

22. Lines 223-225: These sentences fit best in the next paragraph.

[R] — We completely modified the corresponding paragraph to account for the several comments on the results and discussion.

23. Line 245: replace "in particular to plants" by "in particular to those".

[R] — Done.

24. Lines 359-361: please check formatting here.

[R] — Done.

Reviewed by Inês Fragata, 2020-09-08 23:55

In this manuscript the authors test whether female choice for oviposition impacts intraspecific competition across generations. In order to do this, the authors compare oviposition and larvae weight of box tree moths on box trees previously exposed to conspecific herbivores or un-attacked controls. They observe that previous herbivory does not affect where female choose to lay eggs, but it affects larvae weight. This suggests that there is a mismatch between female choice and larvae performance, which is against the preference-performance hypothesis. The question that the authors are trying to answer is very interesting and can help us to understand better how species avoid intra and interspecific competition, even across different generations. Unfortunately, the methodological problems with the chrysalids made it more difficult to fully explore the potential of the question and experimental design. I have some questions/suggestions that may allow to explore better the data set that you have here, and go a bit deeper into your questions.

[R] — Thank you for comments.

Questions/suggestions:

- If I understood correctly, you have the percentage of damage per plant that was done by the first infesting larvae. You could use this as a covariate in your choice experiment to see if damage was a better way of predicting female choice, in addition to your spatial correlation structure. I think it could also be interesting to use the initial percentage of herbivory on the analysis of the larvae weight.

[R] — We have now added this information and re-ran models accordingly. We therefore made appropriate changes in the *methods | Experimental design* and *Results* sections (but did

23

not list all changes here). The reason we did not consider herbivory data in first instance was that potted plants were initially used in a completely separate experiment, for another purpose. We could not match every pot tags between the two projects, leading to missing data. However, because we agree that testing BTM response to actual herbivory rather than to a qualitative attacked/non-attacked factor, we decided to bring original data back into the main text.

- Besides the effect on choice of the egg laying females, herbivory could also affect plant quality and manifest in other life stages. In addition to looking at the weight of the caterpillars, since you have a measure of egg to caterpillar mortality, it would be interesting to see if egg to larvae mortality was higher in attacked or control trees.

[R] — We agree that this would have been a great addition to the paper. However, we only counted and measured a subset of larvae and chrysalis (up to five), and therefore are not able to follow this advice

- Regarding the data on the chrysalids, why do you assume that it was the number of chrysalids that was wrong, and not the number of eggs or larvae (i.e. you could miss some larvae/eggs)? Does this excess occur more in non-damaged trees? Also, does the number of extra chrysalids match the number of missing caterpillars from nearby trees/attacked trees? Because it would be interesting to see if there was more dispersal for pre-attacked trees than for control trees. It is quite striking to have half your controls and 1 one attacked plant where this happens, so I wonder if there is something biologically interesting underlying this pattern. However, if you are planning to not analyse the data or speculate on it, I think it would be better to remove the chrysalid part, as the usefulness of the information is unclear.

[R] — Please see our response to Dr Magalhães, above*

- L148 - Did you release the moths in this region of the plots (between the 1 and 3)? because that could be a reason for the spatial structure to occur? alternatively did it had a source of light/heat or something alike? because it is rather strange that they clustered around that region.

[R] — We released moths at the four corners of the experiment to reduce the risk of spatial aggregation (information now added to the manuscript). We have no data to support any explanation regarding the aggregation of eggs in one particular part of the experiment. This could actually be because of light (the part of the greenhouse received more sunlight in the afternoon) or because of fresher air arriving from the doors.

- L152 – Why did you use these different spatial correlation structures? and what does it mean to have these different spatial correlation structures? This is important to explain what are you accounting for in the analysis.

[R] — We simply followed textbook recommendations when there is no *a priori* hypothesis on the shape of potential patterns.

> *we had no particular hypothesis regarding the shape of the spatial correlation structure. We therefore ran separate models with different spatial correlation structures (namely, exponential, Gaussian, spherical, linear and rational quadratic), and compared them based on their AIC (Zuur, 2009)*

- L163:166 – From table 2 you have 3 models (and not two) that have similar performance, and they are not significantly better or worse compared to the quadratic one, as you need at least a difference of 2 in the AIC, using the rule of thumb from Burnham & Anderson 2004

[R] — Thank you for noticing, we corrected the text.

- In the first section of the discussion, I think two hypotheses that you don't mention are that 1) the moths may need cues from other life stages, such as female conspecific oviposition or the chrysalids; 2) you let too much time pass and the cues related with the conspecific were not present anymore.

[R] — Thank you for these suggestions. We adapted the manuscript accordingly. As for (1):

> *or that female BTM were indifferent to them at the time we conducted the experiment.*
> *[. . .] It remains however possible that BTM adults use other cues to select their hosts*
> *such as the presence of conspecific eggs, larvae or chrysalis.*

as for (2):

> *it is also possible that induced defense reactions were delayed in box trees, or that*
> *they were already relaxed when we released BTM moths three weeks after the herbivory*
> *treatment (Karban, 2011), which remains to be evaluated.*

Text comments

- L17:20 – This sentence is a bit weird, suggestion: "We tested this hypothesis in a choice experiment with box tree moth females (*Cydalima perspectalis* Walker, Lepidoptera: Crambidae). These females were exposed to box trees (*Buxus sempervirens* L., Buxaceae) previously defoliated by conspecific larvae earlier in the season."

[R] — Changed.

- L30 – remove the thus from "Insects may thus reduce"

[R] — Done.

- L32 – I think you mean assumes instead of supposes

[R] — Changed.

- L38 – I don't understand why you say "in particular" here, are those the only traits that will be important for females to detect that correlate with larval performance? for example presence of predators and competitors could be other factors that females may detect before ovipositing.

[R] — We replaced *in particular* by *for instance.*

- L41 – "time-lagged consequences on the preference"

[R] — Done

- L49 – I would substitute a mix of, with both

[R] — Done

- L60 – I would replace the "later herbivores" by later arriving/appearing herbivores

[R] — Done

- L61 – I would replace the "late coming herbivores" by later arriving/appearing herbivores

[R] — We preferred keeping this sentence unchanged to avoid repetition with the previous one.

- L71:73 – This last sentence is not very clear. Maybe: "Thus, in order to quantify the effect of prior herbivory on subsequent herbivore performance, we need to assess how it affects both female choice and progeny performance in attacked and non-attacked hosts."

[R] — Done

- L99:100 – "Their adults. . . " whose adults? you mean that the larvae were used on the preference test and the adult stage on the performance test? maybe something like: "The adult stage of these larvae were used in. . . "

[R] — we clarified this point:

> _We initiated BTM larvae rearing with caterpillars collected in the wild in early
> spring 2019, corresponding to those that had overwintered**. We reared them at room

25

733   *temperature in 4320 cm^3 plastic boxes, and fed them_ ad libitum_, with branches*
734   *collected on box trees around the laboratory. We used the next generation larvae to induce*
735   *herbivory on box tree plants (experimental treatment, see below) and the subsequent*
736   *adults for the oviposition experiment.*

737   • L112 – feed on missing a space

738 **[R] — Done**

739   • L117 –Any specific reason for waiting the three weeks? is it the amount of time that they would take
740   to lay eggs again?

741 **[R] — We have added this information to the revised version:**

742   *In addition, at this time, plants in the herbivory treatment had been cleared of caterpillars*
743   *for three weeks **(corresponding to the duration of the chrysalis stage)** during*
744   *which they were watered every two to three days from above.*

745   • L132 – Why did you wait 24h for weighting them? and not measured them right away? Also was there
746   a high variance in larvae weight?

747 **[R] — We have added this information to the revised version:**

748   *We kept them in Petri dishes without food for 24h **to make larvae empty their gut***
749   *and weighted them to the closest 10 μg.*

750 **We present variability in larval weight in the *'Results'* section:**

751   *The mean weight of BTM larvae varied from 6 to 54 mg (mean ± SD: 20 ± 9 mg).*

752   • L140 – 61 instead of 60 (or otherwise you have the number wrong above)

753 **[R] — Changed**

754   • L140:141 – maybe "and only 1 previously attacked plant" instead of "(and only in 1 previously attacked
755   plant)"

756 **[R] — We rephrased this sentence.**

757   • L146 – x and y coordinates of what?

758 **[R] — Changed**

759   *We ran a generalized least square model (GLS) testing the effect of potted tree location*
760   *in the experimental design (through their x and y coordinates, Figure 2) on the number*
761   *of clutches per plant (log-transformed) from which we explored the associated variogram*
762   *using the functions gls and Variogram in the nlme library.*

763   • L166 – I would not call them competing models. Additionally, you should add whether they show
764   similar results, since you cannot say which one is best.

765 **[R] — *x* and *y* coordinates referred to the design of the experiment (Figure 2). We referred**
766 **to '*competing models*' after Burnham & Anderson textbook (2002). We now state that the**
767 **results would have been the same regardless of the spatial correlation structure. However,**
768 **because this information is not essential, we preferred not reporting the detailed model out-**
769 **puts. Interested or skeptical readers will have access to raw data and codes and will be able**
770 **to simple uncomment the corresponding lines of codes.**

771   • Fig1 – I would put this figure as supplementary material.

772 **[R] — We agree that this figure is not essential, but on the other hand we value this kind**
773 **of illustration showing what the experiment looked like, because the reality is sometimes**
774 **substantially different from what a '*Methods*' section give to imagine.**

775   • Fig 2- I would like to know what are the x and y axis? meters? random unities?

26

[R] — Pots were installed 40 cm apart. The numbers on axes were misleading. We removed them.

Reviewed by Raul Costa-Pereira, 2020-09-06 15:59

Castagneyrpol et al. present results from a well-designed experiment aiming to test the detrimental effects between conspecific individuals that have never meet. They studied a peculiar system where consumers (box tree moth larvae) have limited mobility, relying on their mother's oviposition decision to occupy good food patches (i.e., host plants). Interestingly, food patches are dynamic, and consumers can reduce resource quality to next-generation conspecifics by triggering defensive responses in host plants. Therefore, even if conspecific individuals never coexisted on the same individual host plant, the legacy of past "tenants" can reverberate negatively on current and future ones. Motivated by this interesting conceptual basis, the authors set up a greenhouse experiment to test how past herbivory affects preference (i.e., selection of oviposition sites by mothers, quantified as the number of egg clutches) and performance (i.e., individual consumer growth, quantified as average larvae body mass). Surprisingly, mothers did not avoid laying eggs on plants previously consumed by larvae, but larvae feeding on these plants with the legacy of past conspecifics were smaller. These results are exciting because they shed new light on the mechanisms shaping temporal dynamics of antagonistic interactions both between (plant-herbivore) and within (herbivore-herbivore) species.

[R] — Thank you for this very nice summary and positive appreciation!

I enjoyed reading the manuscript and think it is well-written and the figures are well-presented. The experimental design is creative and statistical analyses are solid (I particularly appreciated how the authors accounted for the underlying spatial structure of their experiment in the models). Below I describe a few major points that came up while reading the manuscript, as well as some minor points that I believe can be helpful. Please let me know if you have any questions, I am happy to clarify.

[R] — Thank you for your valuable and clear comments.

First, I feel that the conceptual framing of the manuscript is fascinating and could be contextualized and motivated more broadly in the introduction. Indeed, multivoltine insect herbivores are a great example of how individuals can affect conspecifics they have never met. Still, similar types of time-lagged interactions between conspecifics occur across diverse taxa (e.g., squirrels [Fisher et al. 2019 Ecology Letters], frogs [Pfennig & Pfennig 2020 Copeia]) and via different mechanisms (e.g., extend phenotypes, ecosystem engineers). Therefore, although the current structure of the introduction works well, I think that opening the manuscript with a more general view of ecological interactions among individuals separated in time would call the attention of a wider and more diverse readership. This approach could also help to reduce some overlap in ideas across the 1st and 2nd paragraphs of the introduction.

[R] — We really appreciated this suggestion. We added a couple of opening sentences to broaden the scope of the paper

> *Biotic interactions are strong factors affecting the fitness of interacting individuals,*
> *even interactions are delayed in time and do not imply direct contact among individuals.*
> *Examples of such interactions can be found in both plants through plant-soil feedbacks*
> *(Putten et al., 2016) and in animals (???; Fisher et al., 2019)*

Hypotheses and respective predictions could be more thoroughly presented to readers. The last paragraph of the introduction is concise and nicely describes the general hypotheses of the study (lines 84-85). However, readers will only find out how the authors investigated their hypotheses in the methods (e.g., lines 123-135), which creates a certain gap in the narrative flow. Thus, the authors could include their respective predictions as well at the end of the introduction (including 'operational variables' - e.g., We expect that plants that previously hosted larvae should [i] have fewer eggs and [ii] host smaller larvae). Moreover, as the experimental design allows inferring the contribution of purely spatial effects on oviposition patterns, the authors could at least mention this at the end of the introduction. By the way, I think that measuring and accounting for spatial structure in oviposition patterns is an exciting novelty of the manuscript. Thus, maybe the

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biological causes and implications of this spatial non-independence in mothers' oviposition choices could be more explored in the discussion.

[R] — We have modified the end of the introduction in order to introduce "*operational variables*" as suggested. As for the discussion on spatial analyses, we agree that our results could pave the way for further investigation. However, the experiment was not designed to explore such spatial effects. We only aimed at controlling possible bias in the design. Therefore, we preferred not putting to much emphasis on this issue as it would have been very speculative.

Finally, I have a quick suggestion about a potential additional dimension of performance that could be considered. To quantify the effects of past herbivory on offspring performance, the authors compared the average body mass of larvae across treatments. I fully agree this is a key aspect of *per capita* offspring performance, and results are exciting in this regard. In light of the natural history of the system, as the authors also measured the number of L3 larvae per plant (lines 130-131), I wonder whether the conversion rate egg $\rightarrow$ L3 larva (e.g., number of eggs/number of L3 larvae) could not be used as an additional metric of performance. Is there evidence in the literature that past herbivory can affect egg eclosion rates and/or early larval development (L1 $\rightarrow$ L2 $\rightarrow$ L3)? This alternative metric would capture a different facet of offspring performance not necessarily correlated with mean larvae weight (e.g., larvae mortality associated with lower foraging rates and/or increased toxins).

[R] — This would have been a great addition to the paper indeed. Unfortunately, we did not have such an information at hand for we counted *up to* five larvae per plant. The phrasing of the original version was ambiguous in this respect. We modified it accordingly.

Minor comments:

- Lines 1-2. The title is solid and general, but I am not sure if all readers will be familiar with the concept of 'multivoltine' (I guess it depends on the target journal).

[R] — We deleted the reference to *multivoltine* species.

- Lines 39-40. Given the idea of the last sentence in this paragraph, it would be good to emphasize here that "competing herbivores" refers to different species of herbivores (i.e., interspecific competition). I would say the same about Line 42.
- Lines 46-50. I feel that most of these ideas were already presented to readers in the previous paragraph. The argument presented in Line 50 sounds like an exciting way to begin this paragraph.

[R] — We deeply modified the introduction, please see our response to Dr Magalhães' comments.

- Line 50. I follow the meaning of 'passage of competitors' but it's possible that some readers may find it a bit confusing. One potential alternative (maybe not that accurate) would be something like 'the legacy' of past herbivores on host plants.

[R] — The corresponding sentence was deleted.

- Line 51. Maybe the authors could provide a brief view of what 'direct' detection means in this context, e.g. "...mated females can directly detect (e.g., via visual or olfactory cues) the present...".

[R] — We added "themselves" to make it clear that the female can detect herbivores, or herbivory-induced changes in plant traits.

- Line 61. Adding a 'triggering' or 'stimulating' before "defenses that generally" could make this idea clearer to readers.

[R] — We rephrased this sentence.

- Line 60. I am not an expert in plant-herbivore systems, but a first intuitive, simple mechanism seems to be the reduction of food biomass by previous consumers. Does it make sense?

[R] — Yes it does! We now mention interference competition and resource depletion.

- Lines 65-66. This is very interesting, and the following example illustrates well this mismatch between selection by mothers vs. impacts on offspring. However, I feel a follow-up conceptual sentence would help readers to crystallize this idea by clarifying that not necessarily the effects on preference and performance are congruent.

[R] — **We restructured the introduction to improve the reading**.

- Lines 74-75. Considering the broad readership of ecologists and evolutionary biologists this manuscript has the potential to reach, I recommend the authors to define the concept of 'multivoltine'. Not all readers may be familiar with it.

[R] — **Done**.

- Line 75. Tiny detail: remove the italic from spp.

[R] — **Done**.

- Lines 84-86. This is a matter of writing style, but one possibility here is to 'change the pace' of this key sentence to emphasize the potential effects on both preference and performance encapsulated by this hypothesis. A simple way to do this would be: "...early herbivory would (i) reduce oviposition..., and also (ii) reduce the performance ...

[R] — **Done**.

- Lines 85-86. As the last sentence of the introduction is often one of the most 'visited' by readers, I feel this one could deliver a 'self-standing', stronger message. For instance, instead of 'By addressing the above', one alternative could be 'By addressing the effects of previous herbivory by conspecifics on both preference and performance of subsequent..."

[R] — **Done**.

- Line 94. I think this initial sentence could be moved down in this paragraph.

[R] — **The sentence was moved to the "natural history" new paragraph.**

- Line 107. It is clear from the previous sentence, but it would be helpful to clarify that this plant-level herbivory metric represents the mean frequency of attacked leaves/branch.

[R] — **Done**:

> _ In order to confirm that the addition of BTM larvae caused herbivory, we visually estimated BTM herbivory as the percentage of leaves consumed by BTM larvae, looking at every branch on every plant. We then averaged herbivory at the plant level. In 8 plants, herbivory data was missing and was imputed as the average of herbivory measured in other plants. In the herbivory treatment, the percentage of leaf area consumed by BTM larvae ranged from 2.2 to 17.2% and was on average 9.1%._

- Figure 1. These photos are great for illustrating the experimental design and study system! I would just suggest adding more details in the legend.

[R] — **Done**.

- Line 113. feed on.

[R] — **Done**.

- Lines 115-116. I wonder if this difference of three weeks between caterpillars being removed (from the herbivory treatment) and moths oviposition reflect the phenology of this species in natural ecosystems. In other words, in light of the biology of BTM, a given box tree in nature could experience two separate groups/generations of caterpillars within three weeks?

[R] — **Yes, the duration of the pupal stage in the wild is long enough to have two separate generations**.

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- Line 118. Just to make it more straightforward: "...washed out from leaves".

[R] — We deleted this sentence.

- Line 120-121. This is an important point that could be briefly mentioned in the last paragraph of the introduction, e.g., "our experimental design allows us to quantify... of plant-mediated".

[R] — Done.

- Lines 126-128. I think that first presenting 'why' (i.e., "to prevent larvae from moving from one potted plant to another") and then 'how' (i.e., "we installed box trees in plastic saucers and interspaced plants and filled saucers with a few centimeters of water").

[R] — Done.

- Line 131. All plants had at least five larvae?

[R] — Done: "only 6% of plants hosted less than five larvae".

- Line 133. It would be helpful to emphasize that this value represents the average weight of larvae from one plant individual.

[R] — Done.

- Line 142. Thanks for such a careful explanation; I appreciate your transparency!

[R] — Thanks.

- Line 150. Could you please describe in more detail the structure of these models? (i.e., define response variable and predictors).

[R] — Done.

- Line 165. With other three models ($\Delta i < 2$ - Models 2, 3, and 4), right?

[R] — Yes, corresponding information is now available in Table 1.

- Line 177. How about finishing this sentence after 'season' and then start a new sentence with 'This time-lagged'?

[R] — Done.

- Line 182-183. Maybe the fact that larval frass was washed out could be a potential explanation?

[R] — We added the following sentence:

  However, we cannot exclude that some cues were mediated by larva frass, which was watched out from leaves when we watered plants.

- Lines 201-203. This an interesting explanation! I wonder if host plants in the native range of BTM have even higher toxic alkaloids than box trees.

[R] — This is an interesting question that would be worth digging further. We are not aware of dedicated studies.

- Lines 206-207. I do not follow this idea, could you please clarify? I feel that 'reduced performance of individuals... have been trait-mediated' requires some further explanation.

[R] — "Trait mediated" was misleading. We deleted these two words and believe the sentence reads better now.

- Line 219-220. The fact that food limitation can trigger cannibalism in this system is fascinating!

[R] — yes, we have been quite surprised to observe this, but did not try to investigate it further. However,in the absence of back-up data, we eventually deleted reference to this possible phenomenon.

30

- Line 221. To avoid repetition, I suggest replacing the first 'Herbivore' by 'Consumer'.

[R] — **This sentence was deleted**.

- Line 228. Would it be 'with' or 'within'?

[R] — **Changed to "within"**.

- Line 237. Perhaps 'negative interactions... generations' could communicate more clearly the results.

[R] — **Done**.

- Line 241. Because this idea expands to the next sentence, it would be good to mention their main enemies (e.g., parasitoids, predators).

[R] — **Done**.

- Line 244. It seems that a verb is missing in this sentence, 'causing more damage' is one option.

[R] — **Done**.

- Line 247. be investigated

[R] — **Done**.

- Line 248. dedicated

[R] — **Done**.

I hope the authors find these comments helpful. Best wishes, Raul.

[R] — **We did, thanks!**