The eco-evolution and biodiversity in antagonistic and mutualistic interactive populations

Abstract

Understanding environmental and ecological effects on ecosystem structure, function, and dynamics is crucial. However, our comprehension of such effects remains challenging due to the complex interplay of abiotic environment, ecological relationships among species, and coevolution within and across trophic levels. Here, we set out to investigate a wide range of combined eco-evolutionary responses in a plant-insect ecosystem. We study ecological population dynamics, coevolution and functional diversity in response to changes in (1) abiotic-biotic interactions, (2) within-trophic-level competitive interactions, and (3) between-trophic-level antagonistic and mutualistic interactions. We developed an eco-evolutionary and functional trait-based model to simulate environmental and ecologically mediated responses to these interactions. We show that mutualistic interactions render a positive feedback loop of increased population sizes followed by selection for even stronger biotic ecological interactions instead of abiotic interactions. Antagonistic interactions have the opposite effect and adaptation to abiotic environment becomes more important. Furthermore, our study reveals that benign conditions (high ecological opportunities, low within-trophic-level competition, and strong interactions between plants and insects) facilitate functional diversity, whereas harsh conditions impede diversity. These findings

21 provide valuable insights into mechanisms that underlie fundamental eco-evolutionary responses

22 to current environmental and ecological changes, offering potential contributions to ecosystem

23 conservation facing urgent challenges.

24 **Keywords:** Eco-evolution, biodiversity, antagonism, mutualism, herbivore, pollinator.

Introduction

The societal and economic benefits associated with ecosystems and several of the services within are indisputable (Alexandridis et al. 2021; Potts et al. 2016). Simultaneously, our actions are reshaping the environment, ecology, and biodiversity, which have evolved over extensive temporal scales (Miller, 2005). For example, agricultural intensification negatively impacts many insect species by reducing within-field habitat quality, among-field diversity, and the amount of semi-natural habitat (Díaz *et al.* 2019; Wagner *et al.* 2021). These anthropogenic endeavours, often driven by short-term objectives, are disrupting ecosystems and leading to biodiversity loss (Díaz et al., 2006; Hilderbrand et al., 2010). These actions pose a potential threat to the long-term sustainability of robust ecological dynamics and health (Cardinale, 2012; Loreau et al., 2022; Naeem, 2002).

Despite the paramount importance of understanding the maintenance of ecological systems, progress in understanding the processes that underlie ecosystem structure (e.g., biodiversity) and function has however been slow (Loeuille et al., 2013). One reason for the slow progress arguably lies in the complex interplay of environmental effects, ecological interactions and evolutionary processes that shape diverse ecosystems. Organisms engage not only with their abiotic surroundings but also with other biotic populations both within and between-trophic-levels. The ecological consequences of these interactions combined with plausible coevolution imply the importance of eco-evolutionary processes in dynamical ecosystems (Cortez & Ellner, 2010; Ellner et al., 2011; Urban & Skelly, 2006). Yet, understanding the eco-evolutionary processes that generate coevolution and functional diversity in ecosystems that include a complex mix of

environmental effects, as well as antagonistic and mutualistic interactions, remain elusive

(Pennekamp et al., 2018; Urban et al., 2016). Much remains to be learned regarding how these

different factors interact within ecosystems.

With the above being said, previous research provides some theoretical bases for understanding

the maintenance and diversification of complex ecosystems. Adaptation in functional traits and

adaptive radiations associated with some biotic and abiotic interactions are, considered to reflect

the reasons for coevolution and diversification processes driven by eco-evolutionary processes in

the ecosystems. For example, studies show that the coevolution and diversity of various organisms

can be driven by ecological opportunity, such as diversified habitats or climate conditions (Losos,

2010; Meyer & Kassen, 2007; Stroud & Losos, 2016; Yoder et al., 2010). These varied ecological

niche spaces allow the emergence of new species, and the gradual accumulation of genetic and

functional differences (Bolnick and Fitzpatrick., 2007; Wellborn & Langerhans, 2015). Another

important driver of the partitioning of niche spaces during diversification is the competition

pressure for limited resources (e.g., food, territory, or mates). The competition pressure can lead

to the emergence of new traits or ecological roles, which foster the coexistence of multiple spaces

in the same ecosystem (Dieckmann & Doebeli, 1999; Doebeli & Dieckmann, 2000; Mac Arthur

& Levins, 1967, Pontarp et al., 2012, Pontarp et al., 2015, Pontarp et al., 2017). Other studies

emphasize the antagonistic and mutualistic trophic interactions as the drivers of adaptation and

diversification. Theory shows the coevolution and adaptive radiations in an antagonistic

community context (Brännström et al., 2011; Braga, 2021; Ito et al., 2009; Loeuille & Loreau,

2005; Sauterey et al., 2017; Wang et al., 2021;), and in specific predation can induce evolution and

disruptive selection on prey populations and drive the evolutionary branching of prey (Brown &

Vincent, 1992; Ito et al., 2009; Pontarp & Petchey, 2018; Pontarp, 2021; Ripa et al., 2009; Wang

et al., 2019). Furthermore, mutualistic interaction can also affect population abundance (Aguilera

et al., 2020), trait evolution (Ramos & Schiestl, 2019) and biodiversity (Aguilera et al., 2020;

Häussler, J. et al., 2017). For instance, studies show that mycorrhizal fungi coevolve with different

host plants, which can cause the emergence of various fungal species, each specialized for specific

plant partners (Brundrett., 2002; Smith & Read., 2008).

Although the above theories focus on different specific aspects (i.e., ecological opportunity,

competition, antagonism and mutualism) that have their influences on the ecological function,

coevolution or diversification, these aspects likely impact the ecosystem in concert (Vellend 2016).

For example, the characteristics of an antagonistic herbivore population may have an indirect

impact on a mutualist population even though they do not interact directly with each other, but

indirectly through a third population (Ohgushi, 2005). Therefore, it is worth adopting a holistic

view of ecosystems as a food web of multiple types of interactions (e.g., Brodersen et al., 2018)

and investigating a wide range of combined eco-evolutionary responses. With this in mind, we

develop an eco-evolutionary and functional trait-based model to explore environmental and

ecologically mediated responses to these interactions. Our model focuses on three main types of

interactions that populations commonly face: abiotic-biotic interactions, within-trophic-level

competition, and between-trophic-level antagonistic and mutualistic interactions (Figure 1). It

follows that we focus on three different interacting trophic levels viewed here as herbivores,

pollinators, and plants. We explore the ecological responses (the change of population sizes) and

evolutionary responses (coevolution in functional traits and diversification in biological

populations with distinct functional traits) to the three types of interactions. We use the adaptive

dynamics approach (introduced by Geritz et al., 1998) for the analyses.

Methods

Overview

We focus on three main interactions that most biological populations will face in their ecosystems

(1) environmental selection pressures (i.e., environmental interactions), (2) within-trophic-level

competitive interactions, and (3) between-trophic-level antagonistic and mutualistic interactions

(Figure 1), and explore how these three interactions will affect ecological dynamics, trait

coevolution and biodiversity in an interactive ecological network. We view the populations in our

system as a set of antagonistic herbivores, mutualistic pollinators, and plants on which the

herbivores and pollinators rely (Figure 1). We assume that functional traits in the biological

populations drive all the interactions. Such traits may be any traits that are of relevance for

environmental adaptation and ecological interactions within or between populations. Body size is,

for example, a trait that is of particular relevance as size commonly correlates with many other

functional traits including physiology, phenology, feeding morphology, etc. (White et al., 2007).

Irrespective of what specific traits are considered we generalized them in a mathematical trait-

based model of ecological and eco-evolutionary dynamics (Brännström et al., 2013). More

specifically, we generalise traits by assuming that herbivores or pollinator populations possess the

traits of relevance both for their adaptation to the environment and their interaction with other

populations. We also assume that the optimal trait for the given environment and the trait that

maximises resource acquisition from the plants may be different, i.e., assuming a tradeoff between

abiotic environmental adaptation and biotic plant resource acquisition. Furthermore, by assuming

that traits can evolve, it follows that population sizes and functional traits change dynamically in

the system in response to environmental effects, resource availability and ecological interactions.

Herbivore and pollinator traits that affect antagonistic and mutualistic interactions in the system

evolve dynamically and can also branch and diversify, so the populations with the new traits

become new functional populations. The newly emerged distinct functional populations then

automatically join the interactions with the existing populations. We run the dynamics until the

whole eco-evolution and diversification process reaches the stable equilibrium. To this end, we

use the adaptive dynamics approach (Brännström et al., 2013; Dieckmann & Law., 1996; Doebeli

& Dieckmann., 2000; Geritz et al., 1998) (details in the section "Eco-evolutionary analysis").

Figure 1. Illustration of the three populations and interactions in the ecological network.
Interactions are denoted by numbers (1-3) and represented by black lines/arrows. (1) The strength of abiotic-biotic interactions is represented by the birth rates (r_A, r_B) and the death rates (d_A, d_B) while the antagonists (A) and the mutualists (B) live in abiotic environment. (2) The strength of within-trophic-level competitive interactions is represented by the carrying capacity of the antagonist and mutualist populations (K_A, K_B) to compete. (3) The strength of between-trophic-level antagonistic and mutualistic interactions is represented by the corresponding interactive convergent coefficients for trophic-level plant resources (c_A, c_{RA}, c_B, c_{RB}). The interactive ranges of the three interactions are measured by the niche widths in the three interactions: σ_{0a}, σ_{0b} in the interaction (1); the competition niche width: σ_{ca}, σ_{cb} in the interaction (2); and antagonistic and mutualistic interactive niche width: σ_a, σ_b in the interaction (3).

Trait-based ecological model

The focused three populations in the model are the plant resource population (R), the antagonistic herbivore population (A), and the mutualistic pollinator population (B). Population interactions are illustrated in Figure 1. In the interaction with abiotic environment (1), both the herbivore and the pollinator populations A and B are dependent on their adaptation to their environment based on how their functional traits a and b deviate from the optimal traits for the environment a_0, b_0 and corresponding interactive niche widths σ_{0a}, σ_{0b}. The strength of the interactions with the environment is represented by their birth rates r_A, r_B and death rates d_A, d_B (Figure 1). In specific, the degree of adaptation to the environment acts on the growth of the populations by the factor of

the normal kernel $\exp\left(-\frac{1}{2}\left(\frac{a_i-a_0}{\sigma_{0a}}\right)^2\right)$ for the herbivores and $\exp\left(-\frac{1}{2}\left(\frac{b_1-b_0}{\sigma_{0b}}\right)^2\right)$ for the pollinators

(eq. I) and their respective intrinsic growth rates r_A and r_B. It follows that adaptation to the

environment is strongest when the population trait equals an optimal adaptive environmental trait

value a_0 or b_0. Also, adaptation tapers off symmetrically as the trait value mismatches the optimal

value according to environmental adaptation niche width parameters σ_{0a} and σ_{0b}. We thus model

adaptation to the general environment using a concept of trait matching between the organismal

trait and the optimal trait value for a given environment. We combine such trait matching with the

concept of niche width where a wide niche width implies that the organism is a generalist that

suffers relatively less from having a mismatched environmental trait while a specialist suffers more

from such a mismatch. The growth rate and the death rates of the plant resource r and d are

assumed to be constant and so not affected by the environment (Lotka-Vertera Model, Lotka, 1920;

Lotka, 1925; Volterra, 1926).

Both the herbivore and pollinator populations engage in within-trophic-level competitive

interactions for their own limited mutual resource, denoted as carrying capacity K_A and K_B (Figure

1 interaction (2)). We assume that the competitive interaction among populations depends on the

same trait as the ones described above (in interaction (1)) and trait matching between populations

dictates competitive strength. Such a trait may again be body size which commonly correlate with

other more specific traits of relevance for competition for resource, nesting places, etc (White et

al., 2007). Similar to the niche concept presented above we assume that such interactions diminish

following the functional form of a normal distribution. More specifically, the strength of the

competition acts on population growth by the factor of $\exp\left(-\frac{1}{2}\left(\frac{a_j-a_i}{\sigma_{ca}}\right)^2\right)$ and $\exp\left(-\frac{1}{2}\left(\frac{b_k-b_l}{\sigma_{cb}}\right)^2\right)$

(eq. I), where σ_{ca} and σ_{cb} denote the competition niche width for herbivores and pollinators.

The herbivores and pollinators also have trophic-level interaction with their shared plant resource

R. The plant population thus have positive effects on both herbivores and pollinators (Figure 1

interaction (3)). The effect from the herbivores on the plants is negative since the herbivores feed

on the plants, whereas the effect from the pollinators on the plants is positive due to mutualistic

contributions (Figure 1 interaction (3)). Like the interaction strength mentioned above in

interactions (1) and (2), the strengths of the interaction between the plant and the herbivore

populations depend on the matching between herbivore population trait (a_i), their corresponding

optimal interactive trait with the plants (a_m), and also on the resource niche width σ_a. It follows

that the strength of the interaction is highest when the herbivore trait equals the optimal interactive

trait with plant trait (a_m), and tapers off when the herbivore and plant trait value mismatches

following a normal kernel $\exp\left(-\frac{1}{2}\left(\frac{a_i-a_m}{\sigma_a}\right)^2\right)$. We also introduce the conversion coefficients c_A,

c_{RA} which denote how much the strength of interaction between the herbivores and the plants are

converted into their own growth (eq. I). Similarly, for the positive mutualistic interactions between

pollinators and plants, the strength of their interaction follows a normal distribution

$\exp\left(-\frac{1}{2}\left(\frac{b_l-b_m}{\sigma_b}\right)^2\right)$, with the conversion coefficient c_B, c_{RB} for the two populations (eq. I). It is

worth noting that the pollinators and the herbivores also have indirect interaction with each other

through their interaction with their shared plant resources (dashed arrows in Figure 1).

191
$$\frac{dA_i}{dt}\frac{1}{A_i} = r_A \exp\left(-\frac{1}{2}\left(\frac{a_i - a_0}{\sigma_{0a}}\right)^2\right) - d_A - \frac{1}{K_A}\sum_j A_j \exp\left(-\frac{1}{2}\left(\frac{a_j - a_i}{\sigma_{ca}}\right)^2\right) + c_A \exp\left(-\frac{1}{2}\left(\frac{a_i - a_m}{\sigma_a}\right)^2\right)R$$

192
$$\frac{dB_l}{dt}\frac{1}{B_l} = r_B \exp\left(-\frac{1}{2}\left(\frac{b_l - b_0}{\sigma_{0b}}\right)^2\right) - d_B - \frac{1}{K_B}\sum_k B_k \exp\left(-\frac{1}{2}\left(\frac{b_k - b_l}{\sigma_{cb}}\right)^2\right) + c_B \exp\left(-\frac{1}{2}\left(\frac{b_l - b_m}{\sigma_b}\right)^2\right)R \quad (1)$$

193
$$\frac{dR}{dt}\frac{1}{R} = r - d - \frac{1}{K}rR - c_{RA}\sum_j \exp\left(-\frac{1}{2}\left(\frac{a_j - a_m}{\sigma_a}\right)^2\right)A_j + c_{RB}\sum_k \exp\left(-\frac{1}{2}\left(\frac{b_k - b_m}{\sigma_b}\right)^2\right)B_k$$

194   ($i, j = 1, 2, ..., N$ and $l, k = 1, 2, ..., M$, where N and M are the total number of populations of herbivores and pollinators)

196   **Table 1 The mathematical terms in the model**

	Resources (plants)	Antagonists (herbivores)	Mutualists (pollinators)
Population sizes	R	A_i	B_l
Trait values		a_i	b_l
Intrinsic growth rates	r	r_A	r_B
Intrinsic death rates	d	d_A	d_B
Carrying capacities	K	K_A	K_B
Optimal adaptive trait values		a_0	b_0
Optimal interactive trait values		a_m	b_m
Adaptive niche widths		σ_{0a}	σ_{0b}

Competitive niche width		σ_{ca}	σ_{cb}
Resource niche width		σ_a	σ_b
Interactive conversion coefficients	c_{RA}, c_{RB}	c_A	c_B

Eco-evolutionary model analyses

To investigate the frequency-dependent ecological population dynamics and evolutionary dynamics, we apply adaptive dynamics theory (Brännström et al., 2013; Dieckmann & Law., 1996; Doebeli & Dieckmann., 2000; Geritz et al., 1998). According to the theory, ecological dynamics are assumed to be faster than the evolutionary process, i.e., ecological equilibria are reached before any mutation-driven trait changes are introduced. For the ecological equilibrium, we used computational method (in R 4.2.0) to retrieve the ecological population size equilibrium, which happens when all the population sizes do not change (the equilibrium population sizes are also checked as the fitness of all populations being zero using the 'solve' function in R 4.2.0). At the ecological equilibrium, an evolutionary mutation will occur only when the mutational traits bring a positive fitness value to the focused mutating populations, which follow the idea of "invasion fitness" by Metz et al (1992), i.e. a population with a mutant trait will be able to invade the resident population given that the mutant population have a positive invasion fitness at ecological equilibrium. Evolutionary mutations are assumed to be small, i.e., the mutant trait values replace the nearby resident trait value gradually in the evolutionary process. For such gradual trait

evolutionary processes, we use the Canonical equation (Dieckmann & Law., 1996) to calculate the

trait changes as a function of time:

$$a_{i,t+1} = a_{i,t} + \frac{1}{2}\mu\sigma_\mu^2 s_\mu \partial f_{a_i} A_i \qquad (i = 1,2,\dots,n_1) \qquad (II)$$

$$b_{i,t+1} = b_{i,t} + \frac{1}{2}\mu\sigma_\mu^2 s_\mu \partial f_{\beta_i} B_i \qquad (i = 1,2,\dots,n_2). \qquad (III)$$

where $a_{i,t}$, ∂f_{a_i} and A_i in Eq. II represent the value of i-th antagonist's trait, selection gradient (the

derivative of the fitness value in respect to the trait value) and the population size at the time t

correspondingly (similarly for the mutualists in Eq. III), and μ, σ_μ^2, s_μ represent the mutation rate,

mutation variance, and mutation steps, which are fixed in our analysis.

Following the above procedure of ecological equilibrium and mutation invasions, novel traits will

continuously replace the resident trait until the resident fitness reaches the local maximum or

minimum in the fitness landscape (Brännström et al 2013). If a fitness maximum is reached,

evolution will stop as no more mutants can invade. However, if the resident trait evolves to the

local minimum in the fitness landscape, the fitnesses of mutating traits on both sides of the resident

trait are higher than the fitness of the resident trait, so disruptive selection emerges. The population

then diverge into two phenotypically distinct populations, and both populations join the

interactions with the ecosystem and the evolutionary dynamics continue (Geritz et al., 1998). For

example, the herbivore population or the pollinator population can diverge from the original

population A or B into two and later into several more population A_i ($i=1,\dots,n_1$) or B_l ($l=1,\dots,n2$)

with their corresponding traits a_i ($i=1,\dots,n_1$) or b_l ($l=1,\dots,n_2$). The competitive interaction then

emerges among all the populations A_i ($i=1,\dots,n_1$) and among all the populations B_l ($l=1,\dots,n_2$)

respectively. The newly generated populations will coevolve with the other existing populations

(eq. I). It is also possible that during this coevolutionary process, one or more population sizes

drop to zero and thus go extinct. All the populations continue to coevolve with the new diverged

populations as well as adapt to new extinctions until all the population trait values reach their local

maxima in the fitness landscapes. Under such conditions, when all populations are on peaks in the

fitness landscape, the evolutionary dynamics become stable and stop changing, which means that

the Evolutionary Stable Strategy has been reached (the *ESS*, Geritz et al., 1998). All results

presented below are analyzed at the *ESS*.

Model scenarios and analyses

Here we use the above eco-evolutionary model to delve into the mechanisms governing ecosystem

structure, function and dynamics, focusing on the impact of the three interactions: abiotic–biotic

interaction, within-trophic-level competitive interaction and between-trophic-level herbivory and

pollination interactions during the ecological and evolutionary process. To this end, we look into

how the interactive factors in the three main interactions (Figure 1) affect the population dynamics,

coevolution and diversity in an ecosystem context. For the three interactions, (1) we use abiotic–

biotic interaction related growth and death rates (r_A, r_B, d_A, d_B) to represent the strength of

interactions between abiotic environment and biotic populations (herbivore and pollinator

populations). A population with a higher growth rate or a lower death rate means that the

population has a higher positive interaction with the environment. (2) We use the competitive

interaction related carrying capacities (K_A, K_B) to represent the strength or intensity of competitive

interaction within the herbivore populations or the pollinator populations. A higher carrying

capacity inside the herbivore populations or the pollinator populations means that the negative

competitive interaction within focused populations is less intense. (3) Finally, we use the

herbivory and pollination interaction related conversion coefficients (c_A, c_{RA}, c_B, c_{RB}) to represent

the strength of the insect-plant interactions between the herbivores or the pollinators with their

communal plant recourses. A higher conversion coefficient between a population and the plant

resources means that positive plant-resource-based herbivory or pollination interaction is higher

for that population. In the research, we examine how these factors relating to the three interactions

can affect the change of ecological population sizes for both the herbivores and the pollinators.

We also study in what direction the functioning traits evolve across time, that is whether the

focused traits evolve to be more interactive with abiotic environment or with the biological

populations in the ecosystem.

Moreover, we investigate the impact of interactive niche widths in the three interactions on

population diversity. In specific, (1) we use the adaption niche widths (σ_{0a}, σ_{0b}) to represent the

interaction range between the biological populations and abiotic environment. A higher adaptive

niche width means that the trait-based population is adaptive to a wider range of environmental

conditions and less sensitive to environmental change. (2) We use the competition niche widths

(σ_{ca}, σ_{cb}) to represent the competition range within the herbivore populations or the pollinator

populations. Lower competitive niche widths mean that the competition interaction ranges of the

focused populations, i.e., herbivore populations (or pollinator populations), are narrow and so the

herbivore population is less likely to have the overlapping living niche with other herbivore

populations (or pollinator populations). In this case, the competition within the herbivore

populations (or the pollinator populations) is less intense. (3) We use the resource niche widths

(σ_a, σ_b) to represent the interactive range between the herbivore or the pollinator populations with

their mutual plant resources. A higher resource niche width means that the herbivore or the

pollinator populations can feed on a wide range of trait-based plant resources. Here we examine

how the niche widths in the three interactions affect the diversity in the herbivore and pollinator

populations.

Results

1. Ecological and evolutionary responses to the three interactions

(1) The effect of abiotic–biotic interaction on population sizes and trait evolutions.

The strength of abiotic-biotic interactions in the herbivore and pollinator populations are measured

by their growth (r_A, r_B) and death rates (d_A, d_B) while living in this abiotic environment, and we

explore how the antagonists and mutualists' ecological population sizes and their evolutionary trait

values can be affected the rates (r_A, r_B, d_A, d_B) (Table 1, Eq. I). As expected, our analyses show

that the population size of the focal populations increases with their own growth rate and decreases

with their own death rate for both the herbivores and pollinators (Table 2, Figure A1). The indirect

effects propagating through the ecological network to the other populations, however, are less

obvious. The increase in the growth rate (or the decrease in the death rate) of herbivores decreases

the population sizes of the pollinators, due to the herbivores' direct negative effects on the plants

as the herbivores consume the plants and so also brings subsequent negative indirect effects on

pollinator populations that live mutualistically with the plants. On the other hand, the increase in

the growth rate (or the decrease in the death rate) of the pollinators, increases the population sizes

of the herbivores, due to the pollinators' direct positive effects on the plants and so also brings

subsequent positive indirect contribution to the herbivore populations (Table 2, Figure A1 - note

that the corresponding descriptive results are shown in Table 2, and the detailed results for the

responses of population sizes and trait evolution to different interactive coefficient factors are

shown in Appendix Figures A1).

Such indirect effect also propagates across eco-evolutionary time scales (not only in the change of

ecological population sizes shown above, but also in the evolution of traits). Both the trait values

of herbivores and pollinators evolve towards the optimal environmental adaptive traits $a_0=0$, $b_0=0$,

and away from the optimal interactive traits $a_m=1$, $b_m=2$, when the birth rate of the antagonistic

herbivores (r_A) increases. This means that the whole system evolves to be less interactive with

other biotic populations in the system both for the pollinators and the herbivores as the antagonists

adapt better to the environment. Interestingly, when the birth rate of the mutualistic pollinators (r_B)

increases, the pollinator traits evolve towards the optimal environmental adaptive traits and away

from the optimal interactive traits. However, during such conditions, the herbivore traits evolve

towards the optimal interactive traits and away from the optimal environmental adaptive trait. This

result means that the pollinators evolve to rely more on abiotic environment to survive since they

already have a high growth rate from the environment, whereas the herbivores evolve to rely more

on the interactions with the biotic system to get more benefits while indirectly interacting with

high-growth-rate pollinator populations through the ecological net (Table 2, Figure A2). As for

the death rate, both the trait values of herbivores and pollinators evolve towards the optimal

interactive traits $a_m=1$, $b_m=2$, and away from the optimal environmental adaptive traits $a_0=0$, $b_0=0$,

when the death rate of the herbivores (d_A) increases. This means that the whole system evolves to

be more interactive with other biotic populations both in the pollinators and in the herbivores,

when the antagonistic herbivores suffer from higher death rates. On the other hand, when the death

rate of the pollinators (d_B) increases, both the trait values of the herbivores and pollinators evolve

away from the optimal interactive traits with the biotic system, and towards the optimal

environmental adaptive traits. Thus the whole system evolves to be less interactive with other

biotic populations when the mutualistic pollinators suffer from higher death rates (Table 2, Figure

A2).

Table 2 The ecological and evolutionary responses to the interactive coefficient factors

parameters		Population sizes		Trait values	
		A	B	a	b
Growth and death rates	r_A	+	-	o	o
	d_A	-	+	×	×
	r_b	+	+	×	o
	d_B	-	-	o	o
Carrying capacity	K_A	+	-	o	o
	K_B	+	+	×	×
Convergent coefficient	c_A, c_{RA}	-	-	o	o
	c_B, c_{RB}	+	+	×	×

+: population size moves in the same direction as the parameters.

-: population size moves in the opposite direction as the parameters.

×: traits move towards higher interaction with the biotic system when parameters increase.

o: traits move towards higher interaction with abiotic system when parameters increase.

(2) The effect of within-trophic-level competitive interaction on population sizes and trait

evolutions.

The strength of within-trophic-level competitive interactions in the herbivore or pollinator

populations is measured by their total carrying capacity (K_A, K_B) that is available for the herbivore

populations or the pollinator populations to live. Higher carrying capacity means that the

competitive interaction is less intense inside the herbivore populations or pollinator populations.

Here we explore how the antagonists and mutualists' ecological population sizes and their

evolutionary trait values can be affected by the carrying capacity (K_A, K_B) (Table 1, Eq. I). As

expected, the population size of either the herbivores or the pollinators increases as their own

carrying capacities increase. Furthermore, since the indirect effects propagate through the network,

the increase in the carrying capacity of herbivores indirectly decreases the population sizes of the

pollinators. And the increase in the carrying capacity of the pollinators, on the other hand,

indirectly increases the population sizes of the herbivores (Table 2, Figure A1). This result means

that a lower intensity of competitive interaction within the herbivores, amplifies the herbivores'

negative effects on the whole system, so decreases the population size of the pollinators; whereas

the lower intensity of competitive interaction within the pollinators, amplifies the pollinators'

positive effects on the whole system, so increases the population size of the herbivores.

Such indirect effect also propagates across eco-evolutionary time scales and shows in the evolution

of traits of both herbivore and pollinator populations. Both the trait values of herbivores and

pollinators evolve towards the optimal interactive traits $a_m=1$, $b_m=2$, and away from the optimal

environmental adaptive traits $a_0=0$, $b_0=0$, when the carrying capacity of the pollinators (K_B)

increases. This means that the whole system evolves to be more interactive with other populations

both in the pollinators and in the herbivores, when the mutualistic pollinators face less negative

competitive interactions in the system. On the other hand, when the carrying capacity of the

herbivores (K_A) increases, both traits evolve away from the optimal interactive traits, and towards

the optimal environmental adaptive traits. This means that the whole system evolves to rely more

on abiotic environment and is less interactive with other populations, when the antagonistic

herbivores face less negative competitive interactions. (Table 2, Figure A2).

(3) The effect of between-trophic-level herbivory and mutualistic interactions with plant

resources on population sizes and trait evolutions.

The strength of between-trophic-level antagonistic or mutualistic interactions is measured by the

conversion coefficients when the herbivore or pollinator populations interact with the shared plant

resources (c_A, c_{RA}, c_B, c_{RB}). High conversion coefficients mean that the antagonistic or mutualist

interactions are stronger between the shared plant resources and the corresponding antagonists or

the mutualists. Here we explore how the antagonists and mutualists' ecological population sizes

and their evolutionary trait values can be affected by the conversion coefficients (c_A, c_{RA}, c_B, c_{RB})

(Table 1, Eq. I). It is shown that when the conversion coefficients of the antagonistic herbivory

interaction (c_A, c_{RA}) increase, both the herbivore (direct effect from the antagonistic herbivore

interactions themselves) and the pollinator (indirect effect through the network system from the

herbivores to the pollinators) population sizes decrease, because the negative antagonistic

interaction in the system is higher. However, when the conversion coefficients of the mutualistic

pollination interaction (c_B, c_{RB}) increase, both the herbivores (indirect effect) and the pollinators

(direct effect) population sizes increase, because the positive mutualistic interaction in the system

is higher (Table 2, Figure A1).

Similarly, both the direct and indirect effects propagate across eco-evolutionary time scales and

also show in the evolution of traits of both herbivore and pollinator populations. When the

conversion coefficients of the herbivores (c_A, c_{RA}) increase, both the trait values in the herbivores

and pollinators evolve away from the optimal interactive traits with the biotic system but towards

the optimal environmental adaptive traits (which means less interaction with the other biotic

populations in the system, but more with abiotic environment), since the level of negative

antagonistic interaction in the system is higher. However, when the conversion coefficients of the

pollinators (c_B, c_{RB}) increase, both the trait values in the herbivores and pollinators evolve towards

the optimal interactive traits but away from the optimal environmental adaptive traits (which

means more interaction with the other biotic populations in the system, but less with abiotic

environment), since the level of positive mutualistic interaction in the system is higher (Table 2,

Figure A2).

2. The population diversity in response to the three interactions.

Here we study how the trait diversity in the evolutionary time scale is affected by the interactive

niche widths in the three interactions. The interactive niche widths of the herbivore and pollinator

populations in the three interactions (adaptive niche width σ_{0a}, σ_{0b}, competitive niche width σ_{0a},

σ_{0b}, resource niche width σ_{0a}, σ_{0b}, see Figure 1 and Table 1), determine how their interactive trait

mismatch can affect the growths of herbivore and pollinator populations in the dynamic ecosystem.

Higher interactive niche widths mean that the corresponding interactive niches are wider and so a

higher mismatch of trait values is allowed for the same growth rates. With different levels of

interactive niche widths in the three interactions, both the herbivore and pollinator populations

evolve and also diverge (a new population with a new distinct trait emerges) differently across

evolutionary time scales. We explore how the numbers of herbivore and pollinator populations

with distinct traits (as the indicator of diversity) respond to the changes of these interactive niche

widths when all the populations in the ecological system reach their highest fitness in the fitness

landscape in the eco-evolutionary dynamics, and so the populations no longer evolve or diverge

(the *ESS*, see the Method session).

Firstly, our results show that when the adaptive niche width of herbivores (σ_{0a}) or the pollinators

(σ_{0b}) is wider, the level of diversity is higher for the herbivore or the pollinator populations (Figure

2). This corresponds to circumstances when a broader range of abiotic environmental opportunities

(e.g., living habitats, geographic location, external food resource, etc) are allowed for the herbivore

or the pollinator populations to adapt to, in which case more diversified populations with distinct

traits adapting to specific environmental conditions emerge in the system. Secondly, our results

show that when within-trophic-level competitive niche width of herbivores (σ_{ca}) or the pollinators

(σ_{cb}) is narrower, the level of diversity is higher for the herbivore or the pollinator populations

(Figure 2). This means that the within-trophic-level competition is less intense with less

overlapping conflicting living space for the herbivore or pollinator populations, then diversity is

higher in the system. Thirdly, we find that when between-trophic-level resource niche width of the

herbivores (σ_a) or the pollinators (σ_b) is wider, the level of diversity is higher for the herbivore or

the pollinator populations (Figure 2). This is when the herbivore populations and the mutualist

populations can benefit from a broader range of plant resources in the system, and so more

diversified populations with distinct traits emerge and thrive in the system.

Notably, all these observations show a fundamental pattern: populations of herbivores or

pollinators have a higher level of diversify under mild living conditions, where adaptability to the

environment is high, within-trophic-level competition is lower, and herbivory or mutualistic

resource availability is abundant. In contrast, harsh living conditions, characterized by narrow

adaptive niches, intense within-trophic-level competition, and limited between-trophic-level

resource opportunities, stifle population diversification. Note that all these differences do not affect

the diversity in the other population (not shown in Figure 2).

Figure 2. The patterns that the interactive niche width affect the diversity (the number of populations with different traits) of antagonists and mutualists. The vertical axis of each panel shows the gradient of the competitive niche widths. All four panels show that the lower the competitive niche widths the higher the diversity in traits. The horizontal axis of each panel shows

the gradient of the adaptive niche width. All four panels show that the higher the adaptive niche

width, the higher the diversity in traits. The upper two panels show the narrow antagonistic

interactive resource niche widths (between the antagonists and plants) and the narrow mutualistic

interactive resource niche widths (between the mutualists and plants). The lower two panels show

the corresponding wide antagonistic and mutualistic interactive resource niche widths. Comparing

the upper two with the lower two panels, the wider interactive resource niche widths lead to higher

diversity for both antagonists and mutualists. The parameter values: $a_0=0$, $b_0=0$, $a_m=1$, $b_m=2$,

$r_A=0.5$, $r_B=0.5$, $r=1$, $d_A=0.1$, $d_B=0.1$, $d=0$, $K=10$, $K_A=3$, $K_B=3$, $c_{RA}=0.2$, $c_{RB}=0.2$, $c_A=0.1$, $c_B=0.1$,

σ_{0a}, σ_{ca}, σ_{0b}, σ_{cb} equal the values shown in the figures, and $\sigma_a=1$, $\sigma_b=1$ (upper left, upper right),

$\sigma_a=1.3$, $\sigma_b=1$ (lower left), $\sigma_a=1$, $\sigma_b=1.3$ (lower right). The parameter terms are explained in Table

1.

Discussion

In biological ecosystems, populations interact with a multitude of biotic entities, both horizontally

(e.g., competitors) and vertically (e.g., trophic-level predators, herbivores, mutualists, etc), as well

as interact with their abiotic surroundings, including habitats, geographic locations, and climate.

To unravel this intricate web of interactions and simplify our understanding of the ecological and

evolutionary processes, we identify three pivotal types of interactions that nearly all biological

populations face in ecosystems: abiotic-biotic interactions[1], within-trophic-level competition[2],

and between-trophic-level antagonistic or mutualistic interactions with the mutual plant resources[3]

(note that superscription (1)(2)(3) referred to the three interactions, illustrated in Figure 1, same as

below). We developed a trait-based eco-evolutionary model (Eq. I) to scrutinize the responses

elicited by the three interactions within an ecosystem, encompassing ecological dynamics,

coevolutionary patterns, and diversity phenomena. The general trait-based nature of the model

enables its use in various eco-evolutionary processes in diverse dynamic systems, including but

not limited to competition, cooperation, mutualism, herbivory, predation, and abiotic-biotic

interactions.

From our analyses, we find that the population sizes of both the mutualists and antagonists are

shaped by their three interactions (abiotic-biotic interaction[1], within-trophic-level competition[2],

between-trophic-level interaction with mutualists and antagonists' shared resources[3]) (Figure 1)

either directly from mutualists or antagonists themselves or indirectly from the other. The stronger

positive mutualists' interaction in all the three interactions, whether with abiotic environment[1]

(i.e., higher adaptivity to the environment) (Memmott., 2007), or among themselves[2] (reduced

competition), or with plant resources[3] (higher compatibility with the plant resources), lead to

increased population sizes for both the mutualists themselves (directly) and the antagonists

(indirectly). This rise is attributed to the positive influence of mutualists on the entire ecosystem.

For the antagonists, stronger interactions with abiotic environment[1] or among themselves[2] boost

their population sizes (directly) but decrease mutualists' population size (indirectly), due to the

amplified negative impact on the shared plant resources. Antagonists' intensified negative

interaction with communal plant resources[3], such as consuming plants, results in reduced

population sizes for both the antagonists' and mutualists, due to their higher detrimental impact on

the communal plant resources. Other studies such as the empirical evidence, demonstrated by

Fiegna & Velicer (2005), support this indirect effect in the ecological network, where cooperative

populations can experience negative effects on reproduction and even face extinction when

antagonistic interactions in the system are high.

The evolution of the trait values is also shaped by the strength of the three interactions by the

mutualists or antagonists both directly and indirectly. The traits of antagonists evolve to be more

interactive with biotic systems and less reliant on abiotic environment, when the mutualists exhibit

stronger interactions with the system (either being less competitive[2] or more compatible with

plant resources[3] or the environment[1]), or when the antagonists exhibit weaker corresponding

interactions with the system. This signifies the significantly positive contributions of mutualists to

the system, contrasting with the negative contributions brought by antagonists. On the other hand,

for the mutualists, strong positive interaction with the biotic populations (reduced competition[2]

or higher compatibility with the plant resources[3]), drives their traits to become more interactive

with the biotic system than with abiotic system. However, when their interaction with abiotic

environment[1] is stronger, their traits evolve to be more interactive with abiotic system. The

mutualists' traits also evolve to be less interactive with the biotic system and more interactive with

abiotic system when any of the three interactions with antagonists are stronger. To sum up, the

mutualists evolve to interact more with the biotic system when they gain more from their mutualist

relation with the biotic system, but evolve to interact less when they are more exploited by the

negative antagonistic or competitive relation in the biotic system. Meanwhile, antagonists evolve

to interact more with the biotic system when they benefit from mutualists' positive contribution to

the system, yet evolve to interact less with the biotic system when their own negative antagonistic

relation within the system intensifies.

Diversity among both antagonists and mutualists is also significantly influenced by all the three

interactions and mainly through their interactive niche width in each interaction (Leibold, M. A.,

1995). Our finding shows that a narrower competitive niche width[2], indicating a reduced intensity

of competition among populations, correlates with higher diversity. This finding aligns with the

theory suggesting that reduced competition intensity within populations, resulting from a narrower

overlap in functional traits, creates more opportunities for distinct adaptations, and therefore higher

diversified interactive populations emerge (Dieckmann & Doebeli, 1999; Doebeli & Dieckmann,

2000; Pontarp et al., 2015; Pontarp et al., 2017; Pontarp et al., 2012).

Our research also reveals that a wider adaptive niche width[1], indicating populations' greater

adaptability to a broader range of environments (reflecting higher ecological opportunity),

correlates with increased diversity. This aligns with other studies (Petanidou & Potts., 2006;

Wellborn & Langerhans, 2015; Wiens & Graham., 2005) that a wide range of ecological

opportunities provides varied distinct niches in habitat and other living resources, so that different

species can evolve to exploit their own specific niches, and thus adaptive radiation and coexistence

of species can happen. This phenomenon is often observed in isolated environments, such as

islands or newly formed habitats, where diverse ecological niches are available for colonization

(Grant & Grant., 2008; Losos., 2011; Thorpe & Malhotra., 1996; Yoder et al., 2010).   The

empirical review study by Bolnick and Fitzpatrick (2007) explains that sympatric diversification

often involves adaptation to different ecological niches, which can result in expended adaptive

niche widths and, consequently, lead to higher diversification rates within a single environment.

This suggests a further positive loop between the extension of adaptive niche width and the higher

diversified ecological populations, which is worth further investigation.

Finally, our study shows that a wider trophic-level biotic resource niche width[3] (e.g., here is the communal plant resource) is associated with higher diversity among herbivores or pollinators. Like the above adaptive niche width[1], provided with varied living-relied plant resources, herbivores or pollinators with different specific function-based traits (associated with plant consumption or pollination) can coexist (Valdovinos et al., 2013; Vazquez & Aizen., 2006). This result is well recorded in many studies that the biodiversity in the tropics is much more abundant than the poles (Dowle et al., 2013; Hillebrand., 2004; Mora & Robertson., 2005). Also herbivore insects are found to evolve specialized feeding behaviours or physiological adaptations to exploit specific plant defences and chemical compounds (Farrell., 1998; Wang et al., 2019); and pollinators, such as hummingbirds, are shown to evolve various bill shapes and lengths to match the specific floral traits of the plants they pollinate, showing the adaptation of pollinators to different plant opportunities (Curti & Ortega-Baes., 2011; Valdovinos et al., 2013).

To sum up, our study elucidates that biological populations tend to experience higher diversity under relatively milder conditions (wider abiotic environmental opportunity[1], reduced competition intensity[2], or broader benefits from other biotic resources[3]), but lower diversity under relatively harsher conditions. These findings underscore the important role of niche widths in shaping the biodiversity of biological populations. Milder conditions, with their wider adaptive or interacting possibilities, both abiotic and biotic, provide biological populations with ample spaces to thrive and diversify.

The study here identifies three very main interactions that are found to be important in most of the

ecological systems: between-trophic-level[3] and within-trophic-level biotic[2] and also abiotic-

biotic interactions[1]. These interactions are studied in many studies mentioned above, but most of

them are confined to one specific interaction. Here we have a comprehensive study to address all

these three interactions in concert with both antagonistic and mutualist contexts and try to build a

theoretical foundation to study the overall ecological network. We covered some common themes

such as mutualism, antagonism, competition, and adaptation to elucidate the underlying reasons

for population structure changes, trait evolution, and biodiversity in the interactive dynamic

ecosystem. We hope broad implications can be derived from these common themes and the related

mechanisms from this study. For example, several pressing and intriguing questions are often

discussed - why certain species face extinction or experience significant population declines due

to their currently changing abiotic or biotic interactions; why certain species alter their habitats or

traits, or coevolve dynamically with other species; why the biodiversity in certain areas is much

higher than others (e.g., areas with different latitude or elevation); how can we preserve

endangered species amidst challenging environmental or climate conditions; how can we explain

the loss of biodiversity and explore avenues to prevent further deterioration of these situations. We

hope that our theoretical analyses provide the preliminary foundation to address these critical

issues, and also aid in future predictions and interventions.

583 Appendix

Population Sizes

Figure A1 The responses of population sizes of antagonists (left) and mutualists (right) to the interaction factors (grow rates, death rates, carrying capacity, convergent coefficients in the 1st, $(2)^{nd}$, 3rd, 4th rows correspondingly). The parameter values: $a_0=0$, $b_0=0$, $a_m=1$, $b_m=2$, $r=1$, $d=0$, $K=10$, $\sigma_{0a}=\sigma_{ca}=\sigma_a=1$, $\sigma_{0b}=\sigma_{cb}=\sigma_b=1$, r_A, r_B, d_A, d_B, K_A, K_B, c_{RA}, c_{RB}, $c_A=0.5c_{RA}$, $c_B=0.5c_{RB}$ (here shows for c_A and c_B and omits c_{RA} and c_{RB}) equal the values shown in the figures. The parameter terms are explained in Table 1.

Trait Values

Figure A2 The responses of trait evolution of antagonists (left) and mutualists (right) to the interaction factors (growth rates, death rates, carrying capacity, and convergent coefficients in the 1st, 2nd, 3rd, 4th rows correspondingly). The parameter values: $a_0=0$, $b_0=0$, $a_m=1$, $b_m=2$, $r=1$,

598     $d=0$, $K=10$, $\sigma_{0a}=\sigma_{ca}=\sigma_a=1$, $\sigma_{0b}=\sigma_{cb}=\sigma_b=1$, r_A, r_B, d_A, d_B, K_A, K_B, c_{RA}, c_{RB}, $c_A=0.5c_{RA}$, $c_B=0.5c_{RB}$

599   equal the values shown in the figures. The parameter terms are explained in Table 1.

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781