

The role of eco-evolutionary experience in invasion success

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Abstract

Invasion ecology has made considerable progress in identifying specific mechanisms that potentially determine success and failure of biological invasions. Increasingly, efforts are being made to interrelate or even synthesize the growing number of hypotheses in order to gain a more comprehensive and integrative understanding of invasions. We argue that adopting an eco-evolutionary perspective on invasions is a promising approach to achieve such integration. It emphasizes the evolutionary antecedents of invasions, i.e. the species' evolutionary legacy and its role in shaping novel biotic interactions that arise due to invasions. We present a conceptual framework consisting of five hypothetical scenarios about the influence of so-called 'eco-evolutionary experience' in resident native and invading non-native species on invasion success, depending on the type of ecological interaction (predation, competition, mutualism, and commensalism). We show that several major ecological invasion hypotheses, including 'enemy release', 'EICA', 'novel weapons', 'naïve prey', 'new associations', 'missed mutualisms' and 'Darwin's naturalization hypothesis' can be integrated into this framework by uncovering their shared implicit reference to the concept of eco-evolutionary experience. We draft a routine for the assessment of eco-evolutionary experience in native and non-native species using a food web-based example and propose two indices (xp_{Focal} index and $xp_{Residents}$ index) for the actual quantification of eco-evolutionary experience. Our study emphasizes the explanatory potential of an eco-evolutionary perspective on biological invasions.

Keywords

Alien species, ecological novelty, ecological similarity, introduced species, invasibility, invasiveness, naïveté, non-indigenous species

Making the case for an eco-evolutionary perspective on biological invasions

A large number of hypotheses about the mechanisms that determine the success or failure of biological invasions have been proposed (reviews in Inderjit et al. 2005, Hufbauer and Torchin 2007, Catford et al. 2009, Jeschke et al. 2012). However, most of these hypotheses are restricted to specific processes (e.g. enemy release hypothesis, Keane and Crawley 2002, or novel weapons hypothesis, Callaway and Aschehoug 2000) and do not explain variation in invasion success on a more inclusive level. Thus, despite considerable progress in invasion ecology, the search for a more comprehensive and integrative understanding of biological invasions is still on-going (Davis 2009, Richardson 2011, Heger et al. in press). Accordingly, increasing efforts are being made to interconnect or even synthesize the growing number of hypotheses and concepts (e.g. Catford et al. 2009, Blackburn et al. 2011, Gurevitch et al. 2011).

With this conceptual paper we aim at contributing to this important development. We suggest that adopting an eco-evolutionary perspective on invasions is a promising approach to achieve a broader conceptual synthesis in invasion ecology (cf. Heger et al. 2013). Scientific awareness of evolutionary aspects in biological invasions has increased in the last decade (Sax and Brown 2000, Sakai et al. 2001, Hänfling and Kollmann 2002, Lee 2002, Sax et al. 2005, 2007, Facon et al. 2006, Kondoh 2006, Hufbauer and Torchin 2007). But the focus of most studies in this field lies particularly on the evolutionary *consequences* of invasions, i.e. the evolutionary response of species to invasions (see e.g. Cox 2004, Strayer et al. 2006, Carlsson et al. 2009, Orians and Ward 2010). Our focus, on the contrary, lies on elucidating the role that evolutionary *antecedents* may play for invasion success (see e.g. Cox and Lima 2006, Kondoh 2006, Mitchell et al. 2006, Sih et al. 2010, Thuiller et al. 2010). It is a general presumption in ecology that biotic interactions are influenced by the evolutionary legacy of the interacting species (Pianka 2000). During invasions, species reach areas where they are not native and interact with species that they have not evolved with (Heger and Treppl 2003, Cox 2004). Such settings lead to ‘novelty’ in biotic interactions in invaded areas, which may likely be decisive for the success or failure of invasions. In the following, we show on theoretical grounds that adopting an eco-evolutionary perspective on invasions (i) offers the possibility to consider the roles that both native *and* non-native species play in invasion success or failure, i.e. species invasiveness *and* community invasibility; (ii) allows an integrative and at the same time differentiated treatment of invasions that affect different types of ecological interaction (competition, predation, mutualism, commensalism); and (iii) has the potential to link so far apparently disconnected major invasion hypotheses in one common framework.

A framework for explaining variation in invasion success based on the concept of eco-evolutionary experience

During evolution, species adapt to biotic interactions in their native environment. They thereby accumulate what we propose to term ‘eco-evolutionary experience’ in

dealing with these interactions. We hypothesize that this inherited experience – possibly complemented by experience acquired during an individual's lifetime (e.g. predators getting better at capturing prey during successive encounters) – ultimately determines the species' proficiency to survive and prosper within new ecological contexts, as for example when invasions take place. For an introduced species, the biotic community in its exotic range may differ fundamentally from the one in its native environment. Biotic interactions that evolutionarily shaped the introduced species in its native environment may become interrupted (Mitchell et al. 2006). At the same time, the resident organisms in the exotic range are confronted with a species they have never met before. For instance, native prey species may not be familiar with the hunting strategy of a non-native predator, and at the same time the latter may be unprepared for having to compete for prey with other (resident) predators. Thus, as a consequence of biological invasions, biotic interactions arise that may be novel to both introduced and native species. Both sides then depend on their inherited eco-evolutionary experience to react appropriately to the new situation. Plasticity (e.g. in behaviour or morphology) resulting from adaptation to unstable environmental conditions in previous times may play an important role here (Nussey et al. 2005, Richards et al. 2006, Sol et al. 2008, Engel et al. 2011). The degree of eco-evolutionary experience available on either side may thus also be interpreted in terms of the introduced species' invasiveness and the native community's invasibility, respectively.

Figure 1 illustrates a conceptual framework to explain variation in invasion success based on the concept of eco-evolutionary experience. The framework consists of five hypothetical scenarios, corresponding to five major types of ecological interaction: the introduced species acting as prey (Fig. 1A), predator (including herbivores, parasites, and parasitoids; Fig. 1B), competitor (Fig. 1C), mutualist (Fig. 1D) or commensal (Fig. 1E). The graphs presented in each scenario are speculative, their exact shape still to be substantiated with empirical data in future studies. However, the scenarios formulate our generalized hypotheses about the relationship between the eco-evolutionary experience in the interacting introduced and native species on the one hand and the relative probability of the respective invasion to succeed on the other: For predator-prey and competitive interactions, the probability of a successful invasion is likely to be higher with a low degree of applicable eco-evolutionary experience in the native species and a high degree in the non-native species (Fig. 1A, B, C). Widely known examples where these circumstances likely apply include the invasion of purple loosestrife (*Lythrum salicaria*) in North America where it lacks herbivorous enemies that feed on it in its native range (Blossey and Nötzold 1995, Fig. 1A), mammalian invasions on oceanic islands causing the extinction of naïve local avifauna (Blackburn et al. 2004, Fig. 1B), and introduced diffuse knapweed (*Centaurea diffusa*) having allelopathic effects on competing resident native grass species in North America (Callaway and Aschehoug 2000, Fig. 1C). In mutualistic interactions, a high degree of experience in both the non-native and the native mutualist is likely to be advantageous for invasion success (Fig. 1D). This may be the case e.g. for yellow crazy ants (*Anoplolepis gracilipes*) associating with honeydew-producing hemipteran insects on Christmas Island

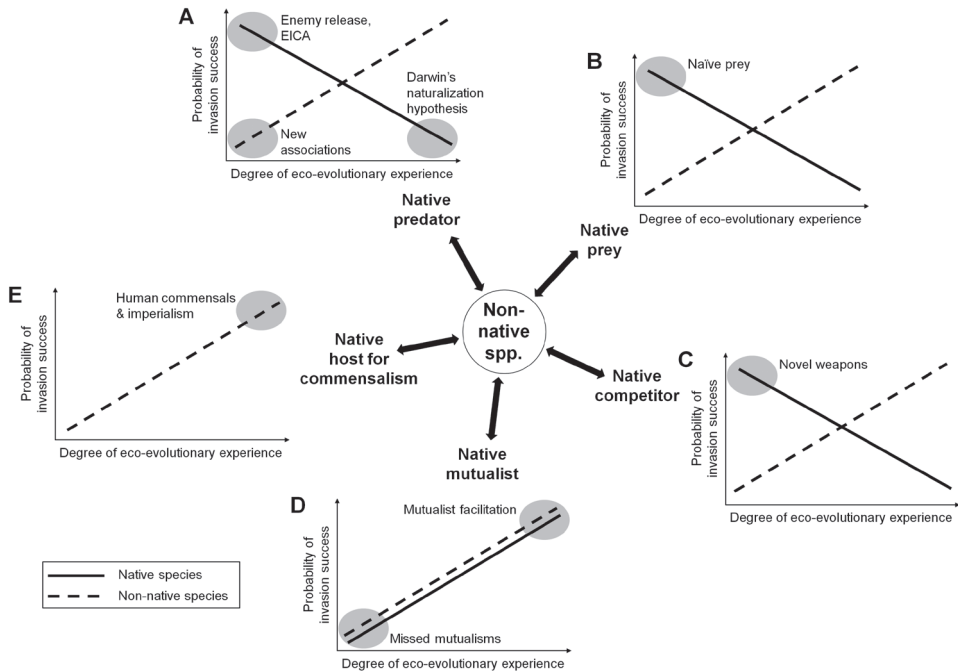


Figure 1. Framework of five hypothetical scenarios about the influence of eco-evolutionary experience in the non-native (dashed line) and native species (solid line) on the relative probability of invasion success, according to the type of ecological interaction (A/B: predator-prey, C: competition, D: mutualism, E: commensalism). In general, lower native experience (except in mutualistic interactions) and higher non-native experience is likely to be advantageous for invasion success. Shaded ovals exemplarily indicate parts of the framework covered by major hypotheses in invasion ecology that implicitly share a reference to the importance of evolutionary legacy for invasion success (see main text and Appendix I for details and references).

(O’Dowd et al. 2003, see also Styrsky and Eubanks 2007). In commensal interactions, eco-evolutionary experience may only have an influence on invasion success if the non-native species is the benefiting commensal (Fig. 1E). In such cases, a higher non-native experience for taking advantage of the native host should be favourable, while the experience level of native hosts is irrelevant, since *per definitionem* the host is not affected by the commensal. This may be the case e.g. for human affiliates like house sparrows (*Passer domesticus*) that successfully invade new areas by being able to reach high population densities in human settlements (cf. Jeschke and Strayer 2006). In the case of a native commensal (not shown in Fig. 1), experience on neither side should have an effect on invasion success, because the non-native host remains unaffected, and the facilitation of the native commensal does not necessarily bear on invasion success.

Notably, several major invasion hypotheses can be integrated into this framework. From an eco-evolutionary viewpoint, it becomes apparent that they actually share an implicit reference to the role of evolutionary legacy in invasion success. This includes

such often-cited hypotheses as ‘enemy release’ (Keane and Crawley 2002), ‘evolution of increased competitive ability’ (EICA; Blossey and Nötzold 1995), ‘novel weapons’ (Callaway and Aschehoug 2000) and ‘Darwin’s naturalization hypothesis’ (Daehler 2001, Procheş et al. 2008, Thuiller et al. 2010). Further examples are the hypotheses of ‘new associations’ (Hokkanen and Pimentel 1989, Mitchell et al. 2006), ‘naïve prey’ (Cox and Lima 2006), ‘missed mutualisms’ (Alpert 2006), ‘mutualist facilitation’ (Richardson et al. 2000), and ‘human commensals and imperialism’ (Jeschke and Strayer 2006). Most of the invasion examples given in the previous paragraph directly apply to one of these hypotheses. Appendix I provides examples of how the central reasoning of the hypotheses can be related to the concept of eco-evolutionary experience, which is visualized correspondingly in Fig. 1 by shaded ovals. Despite this implicit relatedness, they are usually considered separately, sometimes even as mutually exclusive. Only few studies consider potential interrelations between the hypotheses (but see Inderjit et al. 2005, Hufbauer and Torchin 2007, Sih et al. 2010, Gurevitch et al. 2011).

By adopting an explicit eco-evolutionary perspective, the framework provides a basis for interrelating the hypotheses (as defined in Appendix I) and conclusions based on them, but it also highlights their shortcomings: the hypotheses of enemy release, EICA, Darwin’s naturalization hypothesis, naïve prey and novel weapons consider the degree of experience only on the native species’ side (Appendix I: *a, b, d, e, f*), while new associations, missed mutualisms, and the human commensals and imperialism hypothesis focus on the non-natives’ experience (Appendix I: *c, g, i*). Only the mutualist facilitation hypothesis at least implicitly considers both sides (Appendix I: *h*). Thus, these invasion hypotheses emphasize either the invasibility of native communities or the invasiveness of non-native species and neglect that the outcome of an invasion is probably influenced by the degree of applicable eco-evolutionary experience on both interacting sides (cf. Sih et al. 2010). The framework presented here provides a basis for considering both sides simultaneously in order to achieve a more comprehensive understanding of variation in invasion success.

Quantifying eco-evolutionary experience: a food web-based example

Clearly, in connection with the framework presented here, practicable approaches to actually quantify eco-evolutionary experience are needed. Such approaches can build on the general assumption that more of the eco-evolutionary experience in species (native or introduced) will be applicable to a new interaction setting if that setting is ecologically similar to previous interactions. In other words, the degree of ecological similarity between new and previous interaction settings may be taken as a proxy for the degree of applicable eco-evolutionary experience in native and non-native species.

Ecological similarity of species is often assumed to be positively correlated with the taxonomic or phylogenetic relatedness between them (e.g. Agrawal and Kotanen 2003, Ricciardi and Atkinson 2004, Cavender-Bares et al. 2004, Strauss et al. 2006, Diez

et al. 2008, Procheş et al. 2008). Although convenient, this approach has important limitations. In particular, similarity – be it in respect to morphological, behavioural, or ecological traits – does not necessarily correlate with relatedness (Losos 2008, Thuiller et al. 2010). This becomes most evident in cases of convergent evolution where relatively unrelated species show a high degree of similarity (see e.g. Futuyma 2005). Thus, taxonomic classification and phylogenetic relatedness of species are unreliable indicators for their ecological similarity and therefore also for the similarity of biotic interactions of these species before and after an invasion event.

Our approach for quantifying eco-evolutionary experience of introduced and native species assesses the ecological similarity of the ecological *interaction settings* these species are part of before and after the invasion. Such comparisons can be done for any ecological network, e.g. plant-pollinator networks, seed-dispersal interactions, host-parasite systems or food webs. We here present an example for a quantification routine based on food webs (summarized in Appendix II), which covers predator-prey, competitive, and indirect mutualistic interactions (e.g. a predator and a primary producer indirectly benefitting from each other as the predator feeds on the herbivore that consumes the primary producer). We compare the food webs of the original ‘source’ area and a new ‘target’ area of the introduced species (hereafter called the ‘focal species’) regarding the occurrence and occupancy (in terms of number of species) of ecological guilds. Note that the term ‘guild’ as we use it here is not restricted to referring exclusively to “a group of species that exploit the same class of environmental resources in a similar way” (Root 1967). We use a broader definition, where guilds can also be, for instance, groups of species that share the same predators or anti-predator strategies. The exact definition should be chosen based on the particular context of a study. Other ecological groupings (e.g. functional groups or types) can be used instead of guilds as well (for more details on ecological groupings, see e.g. Hawkins and MacMahon 1989, Wilson 1999, Blondel 2003, Blaum et al. 2011).

Eco-evolutionary experience of the introduced focal species

In order to assess the experience of the focal species after its introduction to a target area, we compare the interactions in the food webs of these two areas from the perspective of the focal species (steps 1 to 4 in Appendix II). Both food webs will be composed of different trophic levels, each of which may contain species of different ecological guilds. For simplicity, we restrict our analysis to direct interactions and single-step indirect interactions (i.e. including one intermediate species as for example in exploitative competition) of the focal species with resident species (step 1 in Appendix II). These interactions can be assumed to have the most immediate consequences for the invasion success of the focal species. Separately for each type of interaction (i.e. the focal species acting as prey, predator, competitor or indirect mutualist), and for both the source and target area, the respective interaction partners are classified into their ecological guilds and the members of each guild are counted (steps 2 and 3 in Appendix

II). In this way, we obtain datasets for each type of interaction, with species numbers per guild in both the source and target area (see exemplary Table A in Appendix II).

To actually calculate the eco-evolutionary experience of the focal species (step 4 in Appendix II), we need an index of similarity. The Bray-Curtis similarity index (*sbc*) is often used in ecological studies when comparing the species composition of different samples, e.g. community samples:

$$sbc_{jk} = 1 - \frac{\sum_{i=1}^n |N_{ij} - N_{ik}|}{\sum_{i=1}^n (N_{ij} + N_{ik})}, \text{ (Eq. 1)}$$

where n is the total number of species considered, and N_{ij} and N_{ik} represent the number of individuals of species i in the samples j and k , respectively. Absolute abundance differences in all species are summed up in the numerator and standardized by the total number of individuals in all species from both samples in the denominator. However, while this index provides some grasp on the absolute difference between the samples, it does not consider the direction of change in numbers. But this is important from an eco-evolutionary perspective in the invasion context: for the focal species, it is decisive whether it encounters more or fewer interaction partners from particular guilds in the target area than in the source area. We thus adapted the Bray-Curtis index to account for this specific need. The new index is an index of experience rather than just similarity. We thus call it ‘ xp_{Focal} index’:

$$xp_{Focal} = 1 - \frac{\sum_{i=1}^n \max(0; N_{iT} - N_{iS})}{\sum_{i=1}^n (N_{iT} + N_{iS})}, \text{ (Eq. 2)}$$

where n is the total number of guilds considered, and N_{iS} and N_{iT} represent the number of species in guild i in the source (S) and target (T) area, respectively, that interact with the focal species. Values of xp_{Focal} range between 0 (no applicable experience in the target area) to 1 (maximum applicable experience). By considering not only the presence or absence of guilds but also how numbers of species occupying these guilds differ between source and target area, the xp_{Focal} index accounts for trait differences on the guild level as well as species level. In contrast to the Bray-Curtis index, however, the xp_{Focal} index only considers those differences in the number of guild members where $N_{iS} < N_{iT}$ by introducing the ‘max’ term in the numerator. From the perspective of the focal species, these are the relevant differences between the source and target area, because a larger number of interaction partners of a guild in the target area compared to the source area implies a reduced (or even absent) eco-evolutionary experience of the focal species in the new interaction setting.

This is obvious in cases where the focal species meets interaction partners of a guild in the target area that was entirely absent in the source area (i.e. when $N_{iS}=0$ and $N_{iT}>0$), being then unable to count on applicable eco-evolutionary experience

for these new interactions. But reduced experience is also expected when the focal species interacts with species even of a familiar guild if they occur in larger numbers in the target area as compared to the source area ($N_{is} < N_{it}$). This is reasonable to assume because also species of the same guild differ from each other. Although these differences are relatively small (otherwise the species would be classified into different guilds), they can still be relevant for the focal species. Thus, the more interacting species exist in the target area in comparison to the source area (i.e. the larger N_{it} is in relation to N_{is}), the higher is the probability that the focal species will have to respond to unknown ecological traits, and the lower is its experience in the target area. By contrast, the probability of having to respond to unfamiliar ecological traits of species of a particular guild is low when the focal species has already interacted with a larger number of species from that guild in the source area than in the target area. Our model makes the simplifying assumption of a threshold where the focal species has the maximum eco-evolutionary experience with the new interaction setting ($xp_{Focal} = 1$) when it has interacted with at least as many species in each guild in the source area as it encounters in the target area (i.e. if $N_{is} \geq N_{it}$). In future studies, alternative formulations without such a threshold may be explored.

To a certain degree, the xp_{Focal} index allows reduced experience with members of a particular guild to be compensated by experience in the same type of interaction with species of other guilds. For instance, in predator-prey interactions the focal species may not be familiar with predators of a particular guild in the target area, but may also not be entirely naïve because of having evolved in its source area in the presence of predators at least from other guilds. However, under the assumptions of the xp_{Focal} index, such ‘unspecific’ experience with a type of interaction (in this example ‘predation’) will not completely offset missing experience with a particular guild.

Eco-evolutionary experience of the resident species community

In order to assess the experience of the resident species community facing a new introduced species, we first determine the focal species’ guilds for each type of interaction, i.e. when it may act either as a predator, prey, competitor or indirect mutualist. We then count the number of resident species that are already present in these specific guilds in the target area (see step 5 and exemplary Table B in Appendix II). Finally, by calculating the following ‘ $xp_{Residents}$ ’ index’ separately for each type of interaction (step 6 in Appendix II), we can assess, in a first approximation, how much experience native species have with the focal species:

$$xp_{Residents} = 1 - \frac{1}{N_{i^*T} + 1}, \text{ (Eq. 3)}$$

where N_{i^*T} is the number of resident species in the same guild (i^*) as the focal species in the respective type of interaction. The fraction in this index provides an

estimate how ecologically ‘novel’ the focal species is for the resident community. The maximum novelty of the focal species (i.e. the least experience in resident species) can be expected if no resident species are present in the focal species’ guild before the invasion event. The novelty of the focal species gradually decreases with an increasing number of resident species that are in the same guild as the focal species. Subtracting the fraction from 1, we obtain the eco-evolutionary experience of the resident species community ($xp_{Residents}$), with values ranging again between 0 (no applicable experience of resident species with the focal species) to 1 (maximum applicable experience).

Having thus calculated both the eco-evolutionary experience of the focal species (xp_{Focal}) and the experience of the resident species community ($xp_{Residents}$) for different types of interaction, we can return to the framework in Fig. 1 and estimate the probability of the invasion to succeed.

Discussion

In the previous chapters, we introduced a framework that – by adopting an eco-evolutionary perspective – integrates so far unrelated approaches for explaining biological invasions, and we drafted a routine to quantify eco-evolutionary experience, which is the key variable in this framework. It has to be emphasized again that the framework is of conceptual nature. For instance, the assumed relationship between eco-evolutionary experience and invasion success has to be substantiated with empirical data beyond the hypothetical graphs presented in Fig. 1. Furthermore, the quantification routine makes several simplifying assumptions that have to be kept in mind for an appropriate interpretation:

- Species are adapted to virtually all of their biotic interactions in the source area, which constitutes the inherited eco-evolutionary experience that may matter in ecologically similar communities in the target area. In reality, species are not necessarily adapted to all interactions, e.g. due to weak selection pressure, evolutionary trade-offs, or gene flow. Furthermore, we assume there is no significant intraspecific variation in species traits, e.g. among different populations of the same species.
- Adaptation has no costs. Consider, for example, two focal species that face a single predator species of guild R3 from the example in Appendix II in their respective target areas. For both of them, we would calculate $xp_{Focal}=1$ if during their evolution in the source area they adapted to at least one predator species of the guild R3. The same xp value would be computed even if one of the focal species had adapted to additional predator species. In reality, such ‘over-adaptation’ would probably have generated costs, which could imply disadvantages when compared to the other focal species, but in our model it does not translate into a lower probability of invasion success.

- All interactions are assumed to be equal in strength and frequency. For instance, no distinction is made between generalists and specialists, or whether the focal species interacts in the target area with exactly the same species as in the source area or just with a member of the same guild.
- There is no amplifying effect within interaction types: an interaction partner is counted only once in each type of interaction, even if it maintains more than one ‘connection’ with the focal species within that interaction type (e.g. when competing with the focal species for several prey species).
- As mentioned above, only a subset of all interactions in the studied food webs is included in the analysis, i.e. direct and single-step indirect interactions, and the number of interacting partners in each guild depends on the particular guild definition chosen.

On a side note, we focused in this paper on novel *biotic* interactions that may influence invasion success in order to demonstrate the usefulness of an eco-evolutionary perspective in invasion research. This is not to argue, of course, against the substantial effect that other factors may have on invasion success as well. The significant influence of abiotic conditions has been indicated, for instance, by studies on the effect of climate change (Hellmann et al. 2008, Walther et al. 2009, Engel et al. 2011). Also, Mitchell et al. (2006, p. 734) correctly pointed out that biotic interactions may be influenced “not only directly through the gain and loss of enemies, mutualists and competitors, but also indirectly by putting interactions with the same species in a different environmental context“. Furthermore, among many other factors, the roles of propagule pressure or of intrinsic factors such as (lack of) genetic variability and reproductive systems have to be considered in this context.

We believe that the indices proposed here (xp_{Focal} and $xp_{Residents}$) constitute an important first step towards an efficient quantitative estimate of the influence of species’ evolutionary legacy on the success of biological invasions. A particular strength of this approach lies in its high flexibility: it allows considering not only food webs but also other ecological networks; different kinds of ecological groupings (ecological guilds, functional groups etc.) can be used; and it is applicable to all living organisms across taxonomic boundaries (e.g. plants and animals alike).

From an applied perspective, the further development of the framework and quantification routine to include less simplifying assumptions is certainly highly desirable and a stimulating research perspective. An important next step is to actually test the usefulness of our framework and the quantification routine for empirical case studies. Also, it should be investigated how the various xp values computed for the different types of interaction can best be integrated to provide an *overall* estimate of invasion probability. This could, for instance, be done by reducing complexity (and potential inconsistencies) considering only the most important type(s) of interaction in the respective case study, or it could comprise the development of a single, combined xp value.

Conclusion

An integrative and comprehensive conceptual treatment of conclusions derived from findings in both ecological *and* evolutionary research is still hard to find in invasion ecology. However, as we have outlined above, such an eco-evolutionary perspective would not merely add parenthetical historical information but would increase our potential to uncover invasion patterns. Our framework provides the means for interrelating seemingly isolated ecological invasion hypotheses by identifying implicit eco-evolutionary assumptions (Fig. 1, Appendix I). The framework thus helps to synthesize the conclusions drawn from these hypotheses, providing a stronger basis for a more general understanding of invasion mechanisms and reasons for variation in invasion success. It ties in with the idea of a ‘hierarchy of hypotheses’ (Jeschke et al. 2012, Heger et al., in press), where overarching conceptual ideas in invasion ecology (e.g. the concept of eco-evolutionary experience) branch into more precise and testable hypotheses at lower levels (e.g. enemy release, EICA, novel weapons etc.). Such a hierarchy helps to systematically organize the specific predictions of the large number of individual hypotheses and the evidence accumulated for or against them (Jeschke et al. 2012). This in turn allows evaluating the more general predictions represented by the complete branch of an overarching idea and to identify more fundamental patterns in biological invasions.

The framework generates new, although still very general conceptions on how invasion success depends on eco-evolutionary experience and emphasizes the importance of considering both interacting sides simultaneously: native and non-native species. It also takes into account that non-native species may take up different ecological roles in the exotic range and allows differentiated conclusions for the major types of ecological interactions that may be affected by the invasion.

We believe that the conceptual insights that can be derived from our framework and the quantification routine can be of significant help to guide future research. Ultimately, this research may lead to effective management measures to prevent the introduction of species that seem particularly ‘risky’ for a specific target area, or to adopt appropriate mitigation or restoration measures.

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Appendix I

The shared eco-evolutionary basis of major hypotheses in invasion ecology

The concept of ‘eco-evolutionary experience’ posits that biotic interactions maintained during the evolutionary history of species influence the outcome of interactions between native and introduced species in present times, i.e. (a) the invasion success of the introduced species and (b) the responses of natives. Several major hypotheses for explaining invasion success can be directly related to this concept based on their implicit reference to the logical consequence of a species being introduced into an area where it has not evolved (for references see main text):

- a) Specialized, i.e. eco-evolutionarily highly experienced native enemies of the introduced species may be missing (‘enemy release hypothesis’).
- b) Reduced predation due to inexperienced native predators (herbivores) may allow the introduced species to allocate more resources to traits that increase its competitive abilities (‘evolution of increased competitive ability hypothesis’).
- c) The introduced species may be inexperienced with native enemies and may therefore lack appropriate defence mechanisms (‘new associations hypothesis’).
- d) Introduced species with close relatives in the target area may be less successful because native predators may already be experienced with native congeneric prey species (‘Darwin’s naturalization hypothesis’).
- e) Native prey species may be unprepared, i.e. inexperienced for effectively countering novel predatory behaviour of an introduced species (‘naïve prey hypothesis’).
- f) Native species may not be adapted to, i.e. may be inexperienced with specialized competitive strategies of the introduced species (‘novel weapons hypothesis’).
- g) Mutualistic interactions may fail to develop because of missing experience between native and non-native species (‘missed mutualisms hypothesis’).
- h) Mutualistic interactions between a native and non-native species may be possible, provided that the degree of experience is high enough in both interaction partners (‘mutualist facilitation hypothesis’).
- i) Species that have evolved a strong commensal affiliation to humans may benefit from this eco-evolutionary experience when introduced to areas dominated by humans. This may be especially true for Eurasian species: they coevolved with Europeans and their plants, pathogens and livestock, which were dispersed all over the world during the European Imperialism period (‘human commensals and imperialism hypothesis’).

Appendix II

Routine for the quantification of eco-evolutionary experience: a food web-based example

1. Identify direct interactions and single-step indirect interactions (i.e. including one intermediate species) of the focal species in the food web of its source area and in the food web of the (potential) target area (see example in Fig. A).
2. Define ecological guilds (or other appropriate ecological groupings) for each type of ecological interaction (focal species acting as prey, predator, competitor or indirect mutualist). Assign the focal species and its interaction partners in the source and target area to the ecological guilds.

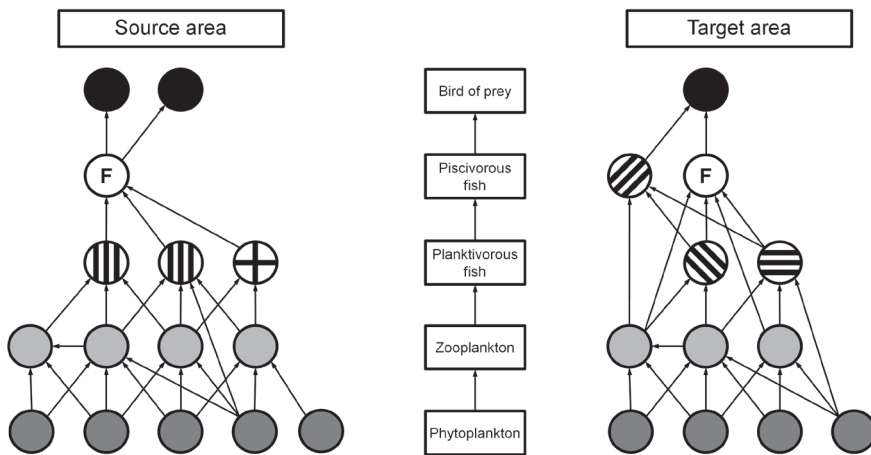


Figure A. Hypothetical food webs in freshwater lakes in source and target area. Circles represent species (F = focal species), different shading and patterning indicate different guilds (see steps 2 and 3).

Quantification of the focal species' eco-evolutionary experience (xp_{Focal}):

3. Determine the number of species that interact with the focal species per ecological guild in the source and target area, separately for each type of interaction (Table A).
4. Calculate the xp_{Focal} index (Eq. 2) for each type of interaction, obtaining the eco-evolutionary experience of the focal species regarding its interaction with resident species in the food web of the target area.

Quantification of the resident species' eco-evolutionary experience ($xp_{Residents}$):

5. Determine the number of resident species in the target area that are members of the same ecological guild as the focal species (regardless if they interact with the focal species or not), separately for each type of interaction (Table B).
6. Calculate the $xp_{Residents}$ index (Eq. 3) for each type of interaction, obtaining the eco-evolutionary experience of the resident species community regarding its interaction with the introduced focal species.

Table A. Numbers of species per guild that interact with the focal species in the food webs of the source and target area (taken from Fig. A), and the respective eco-evolutionary experience of the focal species (xP_{Focal}) in the target area (R1-R5: predator guilds, P1-P5: prey guilds, C1-C5: competitor guilds, M1-M5: mutualist guilds).

Type of interaction Interaction partners of the focal species	No. of species in guild i in source area S					No. of species in guild i in target area T					xP_{Focal}	
	$i =$	$R1$	$R2$	$R3$	$R4$	$R5$	$R1$	$R2$	$R3$	$R4$		$R5$
Predators	$i =$	2	-	-	-	-	1	-	-	-	-	1.00
Prey	$i =$	2	1	-	-	-	-	-	1	1	2	0.43
Competitors	$i =$	-	-	-	-	-	1	1	1	-	-	0.00
Indirect mutualists	$i =$	4	1	-	-	-	3	4	-	-	-	0.75

Table B. Number of resident species in the target area that are members of the same guild as the focal species (note that species numbers are exemplary and not directly deducible from Fig. A), and the respective eco-evolutionary experience of the native community ($xP_{Residents}$) with the focal species.

	No. of resident species in same guild as the focal species	$xP_{Residents}$
Predators	1	0.50
Prey	0	0.00
Competitors	3	0.75
Indirect mutualists	2	0.67