INTERACTIONS OF INVASIVE PLANTS
WITH SOIL BIOTA

Inaugural-Dissertation
to obtain the academic degree
Doctor rerum naturalium (Dr. rer. nat.)
submitted to the Department of Biology, Chemistry and Pharmacy
of Freie Universität Berlin

by

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from Forst (Lausitz)

2012
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1st Reviewer:  Univ.-Prof. Dr. Matthias C. Rillig
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Date of defense:  February 19, 2013
FOREWORD

This dissertation is a cumulative work of manuscripts, either peer-reviewed, submitted or in preparation for submission. Therefore, this thesis is based on following articles, which are referred by their Roman numerals.


II  Bäucker C, Rillig MC (2012) Non-native *Ambrosia artemisiifolia* are more influenced by relative density and identity of neighboring plant species than arbuscular mycorrhiza. (In preparation for submission)

III  Bäucker C, Rillig MC (2012) Distinct seed morphs of *Galinsoga parviflora* (Asteraceae) give rise to different soil feedbacks. (Submitted to Acta Oecologica)
ACKNOWLEDGMENTS

First, I would like to thank Matthias Rillig for giving me the opportunity to do research into the exciting world of arbuscular mycorrhizal fungi, and for providing an inspiring working atmosphere. I am also very grateful to Kathryn Barto, Jeff Powell, Tancredi Caruso and Edith Hammer for discussions, statistical advice, or comments on drafts. Moreover, I thank the department of promotion of women’s of the FU Berlin for financial support of my conference participations, and the Dahlem Centre of Plant Sciences of the FU Berlin for technical support. Further, I would like to thank everyone in the workgroup ‘Plant Ecology’ for being such a very nice and friendly colleague.
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CHAPTER 1

General Introduction

Earth is a dynamic system. Episodes of enormous interchange of species have been repeatedly taken place, for example in consequence of tectonic activity or during Pleistocene ice ages (e.g. Vermeij 1991; CLIMAP Project 1976; Brown and Sax 2004). Besides geographical rearrangements and climate changes, many species also fluctuate in their range expansions as a result of biological interactions on the timescale of decades to years, e.g. empirically observable as succession. The interchange of species, therefore, is a constantly occurring biological process that should not be viewed as abnormal *per se* (Vitousek 1992; Lodge 1993).

Since the last century, however, earth’s biota is being homogenized rapidly as human activities increasingly introduce species outside their natural range (Elton 1958; Lodge 1993). After human-caused habitat destruction, biological invasions were exposed and afterwards heavily cited as the second largest component for current biodiversity loss (Vitousek 1992; Wilcove et al. 1998; Davis 2011). In fact, ecosystems worldwide face tremendous changes; the displacement of native species by non-native species has numerous direct and indirect effects on ecosystem functioning (D’Antonio and Vitousek 1992; Mack et al. 2000, Cassey et al. 2005; Wardle et al. 2011). Undoubtedly, the vast majority of habitat and community changes driven by the spread of non-native, invasive organisms must be regarded as irreversible (Sala et al. 2000). Furthermore, invasive species influence ecosystem services that are fundamental to human well-being resulting in substantial economic costs (see for USA: Pimentel et al. 2000, 2005; for South Africa, Hawaii, Great Lakes (USA): Pejchar and Mooney 2009; for Europe: Vilà et al. 2010; Keller et al. 2011; for China: Wan et al. 2010). Therefore, to understand why some species become the dominant component in communities, where they are not native, is of great scientific, economic and social interest as human activities such as international trade, transport and travel, which cause species dispersal into new ranges, continue to expand (Keller et al. 2011).

From the scientific perspective, biological invasions may also be regarded as ‘grand, but unplanned, biological experiments’ that give the opportunity to study ecological and evolutionary processes (Mooney and Cleland 2001; Brown and Sax 2004).
During the last 50 years, patterns and mechanisms of biological invasions have been increasingly investigated (Richardson and Pyšek 2008); the number of articles published on invasive topics per year has been exponentially growing since the last 30 years (Kühn et al. 2011).

Terminology in invasion biology lacks uniformity and agreements (e.g. Richardson et al. 2000a; Davis and Thompson 2000; Daehler 2001; Colautti and MacIsaac 2004; Pyšek et al. 2004a; Inderjit 2005; Valéry et al. 2008; Colautti and Richardson 2009; Young and Larson 2011; Webber and Scott 2012). The existing terminological difficulties result from the fact that invasion research is multifaceted: the perspectives range from population or community ecology to evolutionary and molecular biology, as well as restoration. Moreover, the usage of terms is related to research histories and traditions. In the German-speaking part, for example, plants introduced before the discovery of America are termed ‘archaeophytes’, and those introduced after 1492 are ‘neophytes’ (e.g. Schroeder 1969; Kowarik 2003; Pyšek et al. 2004b). Recently, European scientists also suggested the term ‘neobiota’ as a value-neutral approach (Kowarik and Starfinger 2009; Kühn et al. 2012). *Sensu* Kowarik (2002) neobiota are organisms, independent of their taxonomic rank, that occur in a region beyond their native range due to human agency or that evolved from such taxa. This ‘neobiota’ concept, however, is not generally accepted.

The English terminology, in contrast, terms species outside their native ranges ‘alien’ (e.g. Crawley et al. 1996), ‘imported’ (e.g. Williamson and Fitter 1996a), ‘non-indigenous’ (e.g. Mack et al 2000; Pimentel et al. 2000), ‘casual’ or ‘naturalized’ (e.g. Richardson et al. 2000a), but also ‘adventive’ (e.g. Mühlenbach 1979) or ‘non-native’ (e.g. Davis et al. 2000). Additionally, more ambiguous vocabulary like ‘exotic’ (e.g. Keane and Crawley 2002) or ‘invasive’ is very common. Above all, the term ‘invasive’ is the major focus of the terminological debate inclusive related terms like ‘invasion’, ‘invader’, ‘invasiveness’ or ‘invasibility’ (e.g. Richardson et al. 2000a; Colautti and MacIsaac 2004; Richardson and Pyšek 2006; van Kleunen et al. 2010; Catford et al. 2012). Defining invasive species, some authors put emphasis on a large negative ecological and/or economical impact (Davis and Thompson 2002; Inderjit 2005), while others see the disruption of the local target community as the most important feature (Keane and Crawley 2002).

In the present dissertation, I refer to the definition of an invasive plant species proposed by Richardson et al. (2000a). It states that invasive plants are ‘naturalized plants
that produce reproductive offspring, often in very large numbers, at considerable distances from parent plants and thus have the potential to spread over a considerable area’. Moreover, I see the invasion process as recently presented by Blackburn et al. (2011). Their concept can be regarded as a unifying framework; it incorporates classical concepts of Williamson (1996) and Richardson et al. (2000a), as well as other conceptual ideas (e.g. Heger and Trepl 2003; Colautti and MacIsaac 2004). The framework includes four stages, which are transport, introduction, establishment and spread. Moreover, it proposes that a species has to overcome a series of six barriers to become a fully invasive species. The barriers are: geography, captivity or cultivation, survival, reproduction, dispersal and environmental; hence, the framework makes no distinction between disturbed and undisturbed habitats (cf. Richardson et al. 2000a). Furthermore, the different possibilities of establishment (spontaneous and permanent) as proposed by Heger and Trepl (2003), are illustrated as a feedback loop between barriers to survival and reproduction in the unifying framework. Blackburn et al. (2011), however, do not rank first in conceptualizing a unifying approach for invasion ecology. Already 100 years ago, the Swiss botanist Albert Thellung suggested a universal framework (Kowarik and Pyšek 2012), and also other integrative concepts have been presented in the last few years (Barney and Whitlow 2008; Moles et al. 2008; Catford et al. 2009).

Aside from the passionate debate about terms and frameworks, invasion biology is a discipline that has been more characterized by theory accumulation than theory discrimination until recently (cf. Davis 2011); the number of theories explaining the exceptional success of invasive species is overwhelming. Some review articles, however, compose overviews on the existing leading theories (Sakai et al. 2001; Hierro et al. 2005; Mitchell et al. 2006; Catford et al. 2009). Catford et al. (2009) point out that many of the existing hypotheses are redundant as they ‘overlap, mirror, unite or share similarity with pre-existing hypotheses’. For example, resource availability in the new environment is an integral component of several hypotheses, e.g. fluctuating resource availability (Davis et al. 2000), disturbance (Sher and Hyatt 1999), opportunity window (Shea and Chesson 2002), dynamic equilibrium model (Huston 2004), or environmental heterogeneity (Melbourne et al. 2007). The excessive generating of new hypotheses in the past may be viewed as resulting from the intention to find the ‘holy grail’ of invasion, as well as related to the fact that most studies focused on one single aspect of invasions only (Richardson and Pyšek 2008; Catford et al. 2009).
Every invasion process, however, must be understood as highly context-dependent and linked to a combination of both abiotic and biotic factors, and multiple mechanisms (e.g. Daehler 2003; Richardson and Pyšek 2006; Barney and Whittlow 2008). Therefore, the success of invasive species cannot be explained with mono-causality. Moreover, all factors and mechanisms underlying the rapid range expansion of invasive species must be assumed to vary in time and space. Recently, two studies demonstrated that even enemy release, which is a crucial aspect of many theories in invasion ecology, does not persist forever (Mitchell et al. 2010; Diez et al. 2010).

To categorize the different approaches and hypotheses that explain the mechanisms of biological invasions, Heger and Trepl (2003) emphasize four different approaches: (1) to focus on the characteristics of the invading species, (2) on those of the ecosystems invaded, (3) on the relationship between these two factors (key–lock approach), and (4) the invasion process in time. Catford et al. (2009) also synthesized four categories/factors, into which hypotheses might be divided: (1) human interference, (2) propagule pressure, (3) abiotic and (4) biotic factors. Recently, Jeschke et al. (2012) evaluated six of the major leading theories, which were classified by their main focus into three groups: (1) invaders themselves, (2) ecosystems into which the invaders were introduced, and (3) invader-ecosystem interaction. According to Jeschke et al. (2012), I also group theories based on their main focus in the present thesis. Here, I differentiate between the following three foci/categories that might be derived from hypotheses in invasion biology:

i) features of the invasive species

ii) characteristics of the new environment/habitat

iii) interactions of invasive species with their new environment

The first category ‘features of the invasive species’ refers to hypotheses like ideal weed (Elton 1958; Baker 1965, 1974) or propagule pressure (Williamson and Fitter 1996b; Lonsdale 1999). The theory of propagule pressure, which implies that the chance for successful invasion is increased by a high supply and frequency of plant propagule introductions, had been found to be a significant predictor of invasion in a meta-analysis (Colautti et al. 2006). Other theories belonging into this group primarily focusing on invasive species traits might be lag-phase (Kowarik 1995) or evolution of increased competitive ability (Blossey and Nötzold 1995), although the latter also has a strong interaction focus.
The second category ‘characteristics of the new environment/habitat’ addresses hypotheses like fluctuating resource availability (Davis et al. 2000), disturbance (Sher and Hyatt 1999) or other theories explaining invasion success predominantly from the perspective of resource availability in the new environment (see above). This category is equivalent to the abiotic factor class suggested by Catford et al. (2009) and the aspect ‘ecosystems into which the invaders were introduced’ by Jeschke et al. (2012). Doubtless, abiotic characteristics play a major role for successful establishment and spread of invasive species. In a recent meta-analysis, globally widespread species were demonstrated to be better able to utilize increased resource amounts of nutrients, light and water compared to less widespread species (Dawson et al. 2012). Besides, Colautti et al. (2006) showed that disturbance and resource availability are significantly positively associated with invasibility. Furthermore, resource availability may synergistically interact with enemy release giving the advantage to non-native over native species (Blumenthal 2006; Blumenthal et al. 2009).

Among the proposed third category ‘interactions of invasive species with their new environment’ count hypotheses like enemy release (Keane and Crawley 2002), novel weapons (Callaway and Ridenour 2004), increased nitrogen cycling (Rout and Callaway 2009), enhanced mutualism (Reinhart and Callaway 2006), mycorrhizal degradation (Vogelsang et al. 2004), or invasional meltdown (Simberloff and Von Holle 1999). According to Jeschke et al. (2012), hypotheses considering invader–ecosystem interactions, such as enemy release, novel weapons, invasional meltdown are better supported than those, which exclusively focus on ecosystem properties or solely on invaders, like tens rule (Williamson and Brown 1986; Williamson and Fitter 1996a). Jeschke et al. (2012), moreover, report that the invasional meltdown theory has the highest level of support across both animals and plants in terrestrial, freshwater and marine habitats. The support, however, has considerably declined over time as it was found for all theories tested. For definition of all hypotheses mentioned in the thesis see Appendix A, Table A.1.

The aim of the present dissertation is to make a contribution to the field of biotic interactions of invasive plants with soil biota. Soil biota include a wide range of taxa, for example mites, collembola, nematodes, macro-arthropods as beetle larvae, earthworms, enchytraeid worms, fungi like Glomeromycota, Basidiomycota, Ascomycota, as well as bacteria, and archaea. This highly diverse belowground community is known to drive
aboveground community structure/functioning via direct and indirect pathways to plants (Wardle et al. 2004). Therefore, plant interactions with soil biota have been subject of numerous research projects in invasion ecology, and many studies found evidence for soil biota playing a crucial role in plant invasions, e.g. nematodes (van Ruijven et al. 2003; van der Putten et al. 2005), ectomycorrhizal fungi (Richardson et al. 1994; Wolfe et al. 2008; Nuñez et al. 2009; Trocha et al. 2012), arbuscular mycorrhizal fungi (e.g. Marler et al. 1999; Mummey and Rillig 2006; Stinson et al. 2006; Vogelsang and Bever 2009; Seifert et al. 2009), fungal pathogens (Mangla et al. 2008), fungal endophytes (Aschehoug et al. 2012), various N2-fixing bacteria (e.g. Vitousek et al. 1987; Parker et al. 2006; Rout and Chrzanowski 2009), or soil microbes < 20 μm (e.g. Klironomos 2002).

Recently, Inderjit and van der Putten (2010) synthesized an overview on how soil biota may directly and indirectly interact with invasive plants, and point out that many questions are open because soil is often used as ‘black box’ in experiments. Indeed, studies using whole soil as treatment have a low mechanistic resolution, but make that compromise to gain greater realism. For example, in soil feedback studies it had been found that the magnitude of soil biota net effects on invasive plants is considerably less negative or even positive in new ranges compared to the species’ native range (Reinhart et al. 2003; Callaway et al. 2004a; Reinhart and Callaway 2006; Kulmatiski et al. 2008; Callaway et al. 2011). Therefore, differences in soil feedback between ‘home’ and ‘away’ ranges may contribute to the successful spread of invasive plants, although these feedback differences must be assumed to become less important with increase in residence time. Diez et al. (2010) studied soil feedback responses of 12 non-native established plant species in New Zealand and found that those species that have been established longest (210 years) exhibited greater negative soil feedbacks than those with shorter residence time. The mechanisms behind these feedback changes in ‘away’ ranges over time is not fully understood, but most probably related to accumulating plant–pathogen interactions, alike novel plant–herbivore interactions aboveground (Verhoeven et al. 2009). However, pathogens may switch very slowly from native to introduced species. As Mitchell et al. (2010) demonstrated long established plants (since more than 400 years) still had 60% fewer pathogens in their new North American range compared to their native European range. Nonetheless, negative impacts of pathogens on non-native species must be assumed to accumulate over time. Aside from residence time, pathogen richness of non-native plants also may depend on geographic size of the introduced range, and if plants have a history of agricultural use (Mitchell et al. 2010).
This thesis primarily focuses on the impact of arbuscular mycorrhizal (AM) fungi on invasive plants. As non-native species, I studied *Ambrosia artemisiifolia* L. and *Galinsoga parviflora* Cav. in the new European range. Both plant species are annual and belong to the family of Asteraceae, whose members often form mycorrhizal symbioses with AM fungi, phylum Glomeromycota (Schüßler et al. 2001). The AM symbiosis is in most cases facultative for the plant partner, but always obligatory for the fungus (Helgason and Fitter 2009). The association most probably evolved in the Ordovician; fossil records date back to 460 million years ago (Redecker et al. 2000). AM fungi provide nutrients, predominantly phosphorous (P) to the plant side in exchange for carbohydrates (Allen 1991; Smith and Read 2008). In addition to the reciprocal nutrient fluxes, AM fungi mediate other functions to plants, such as pathogen protection (e.g. Newsham et al. 1995; Borowicz 2001; Wehner et al. 2010; Veresoglou and Rillig 2011) or improved resistance against drought stress (Augé 2001; Augé et al. 2004). Moreover, the mycorrhizal status of a plant alters their competitive ability, which maintains plant diversity (van der Heijden et al. 1998; Hart et al. 2003). Furthermore, multitrophic interactions as leaf-mining herbivores with parasitoids differ depending on mycorrhization of plants (Gange et al. 2003). However, under certain environmental conditions, such as high nutrient availability in fertilized systems or reduced photosynthesis, AM fungi seem to be less cooperative to the plant side resulting in reduced plant performance (Kiers et al. 2011). Therefore, the AM fungal symbiosis was viewed to act in a range of functions from mutualism to parasitism mediated by abiotic and biotic environmental conditions, including the plant and fungal genotype (Johnson et al. 1997).

Regarding the success of invasive plants, AM fungi have been found to be of particular importance in a number of cases. There are some hypotheses which highlight plant interactions with AM fungi as the critical factor for the plant’s invasive spread: enhanced mutualism (Reinhart and Callaway 2006) and mycorrhizal degradation (Vogelsang et al. 2004) (for details of the theories see Appendix A, Table A.1). A few review articles, moreover, cover the topic of plant interaction with AM fungi in the context of invasion (Richardson et al. 2000b; Wolfe and Klironomos 2005; Mitchell et al. 2006; Pringle et al. 2009; Shah et al. 2009). Recently, Moora et al. (2011) showed that the palm *Trachycarpus fortunei* associates with widely distributed AM fungal taxa when it was introduced to different new European ranges; hence, non-native mycorrhizal plants select for AM fungal generalists and, therefore, seem to be not limited by a lack of mutualistic fungi (Richardson et al. 2000b). Consequently, the role of AM fungi has to be taken into
account to properly explain invasive success of plant species (Mitchell et al. 2006). The present thesis, therefore, considers questions of invasive plants and natural arbuscular mycorrhizal (AM) communities.

I investigated three different issues, which correspond to the manuscripts I–III and following chapters two–four, respectively. Because AM fungal taxa and isolates have been shown to differ in their functions (e.g. van der Heijden et al. 1998; Bray et al. 2003; Munkvold et al. 2004; Scheublin et al. 2007), I aimed to conduct my experiments with comparatively high realism. In my studies, therefore, I always maintained the ecological context of soil and natural AM fungal communities.

**Manuscript I** (Chapter 2) reports about the relationship of the non-native plant *Ambrosia artemisiifolia* with natural AM fungal communities in the new European range at local scale. In a reciprocal inoculation experiment, I studied whether or not the mycorrhizal symbiosis between *A. artemisiifolia* and native AM fungal communities shows evidence of co-adaptation in a European roadside and cornfield population, respectively. I expected that plant performance and fitness of *A. artemisiifolia* is greater when the plants, soil and AM fungal community come from the same site. Further, I predicted that the AM fungal community from the roadside habitat would act cooperatively, while the AM fungal community from the agricultural field would show less cooperative behavior in its agricultural soil context.

**Manuscript II** (Chapter 3) reports about the influence of natural AM fungal communities on the competitive ability of *A. artemisiifolia*. The performance of *A. artemisiifolia* was studied in two different relative abundances (target and challenger arrangements), as well as in presence of different neighboring plant species of the European range. As neighboring plants, I tested *Conyza canadensis* L., *Artemisia vulgaris* L., *Daucus carota* L. and *Tanacetum vulgare* L., which I found co-existing with *A. artemisiifolia*. I expected that the mycorrhizal symbiosis would enhance the competitive ability of *A. artemisiifolia*; its invasive spread has been suggested to be promoted by AM fungi (Fumanal et al. 2006). Moreover, I investigated the influence of natural AM fungal communities on *A. artemisiifolia* and *D. carota* grown in pairwise arrangements of intraspecific and interspecific competition.
Manuscript III (Chapter 4) focuses on the aspect of heterocarpy of the non-native plant *Galinsoga parviflora* Cav. in a soil feedback study. *G. parviflora* produces two distinct seed morphs: seeds equipped with a pappus for long-distance dispersal and non-pappus seeds for maintaining the existing population. I asked if the different dispersal capacities of the two seed types might correlate to different soil feedback responses, which may contribute to the successful spread of *G. parviflora* in the new range. Therefore, I tested feedback responses of plants grown from the two seed types (non-pappus and pappus seeds) in soil trained over two plant generations by *G. parviflora*. Considering the different dispersal abilities of the two seed types, I hypothesized that plants arising from non-pappus seeds would exhibit better performance, i.e. less negative soil feedback, in soil trained by the mother plant than those grown from pappus seeds.
CHAPTER 2

Divergent responses of *Ambrosia artemisiifolia* to natural AM fungal communities in the new European range

Abstract

*Background and Aims* Recently, existence of coadapted plant-arbuscular mycorrhizal (AM) fungal interactions has been found, but knowledge about the extent to which such adaptations also occur during plant invasions is lacking. Here, we investigated whether or not the mycorrhizal symbiosis between *Ambrosia artemisiifolia* and natural AM fungal communities shows evidence of co-adaptation in the new European range.

*Methods* In a reciprocal inoculation experiment with ‘full soil strength’ inocula, we compared performance of genotypes from two different sites: a roadside and a cornfield habitat.

*Results* Natural AM fungal assemblages were mutualistic with *A. artemisiifolia* in roadside soil, but not in agricultural soil tested. Decreased plant growth in response to the less cooperative quality of the agricultural AM fungal community in the agricultural soil coincided with alterations of plant root systems towards greater fineness. We found no evidence for locally adapted plant-AM fungal interactions, but adaptation of roadside genotypes to a roadside soil environment.

*Conclusion* Our results highlight the importance of the soil context for mycorrhizal functions. Contrasting effects of natural AM fungal communities and processes of adaptation to novel soil conditions may play a crucial role in the early stages of the spread of non-native *A. artemisiifolia*. 
Introduction

Arbuscular mycorrhizal (AM) fungi, Phylum Glomeromycota, colonize plant roots gaining photosynthetically fixed carbon in exchange for mineral nutrients, predominantly phosphorus (e.g. Allen 1991; Pearson and Jakobsen 1993; Smith and Read 2008). In addition to the well-known reciprocal nutrient fluxes, AM fungi mediate multiple functions affecting plant traits (e.g. Newsham et al. 1995). For example, mycorrhizal plants are better defended against fungal pathogens (e.g. Borowicz 2001; Veresoglou and Rillig 2011), have greater reproductive output (Lu and Koide 1994; Koide and Dickie 2002; but see Allison 2002), and flower earlier (Sun et al. 2008). Furthermore, several studies report that plant species and genotypes with greater mycorrhizal responsiveness have root systems with a coarser root architecture than non-responsive species and genotypes (Hetrick 1991; Schultz et al. 2001; Berta et al. 2002; Seifert et al. 2009). The multidimensionality of plant responses to mycorrhization in part reflects the functional diversity of AM fungal taxa and isolates (e.g. van der Heijden et al. 1998; Klironomos 2003; Munkvold et al. 2004; Antunes et al. 2011).

In nature, plant species forming arbuscular mycorrhizas are typically colonized by a community of co-occurring AM fungal taxa, which form a complex underground mycelial network (Smith and Read 2008). There is accumulating evidence that plants interacting with this network are able to distinguish between cooperative and less cooperative AM fungi and promote more cooperative fungal partners with increased photosynthate allocation (Bever et al. 2009; Kiers et al. 2011). Therefore, selection pressure in arbuscular mycorrhizas would favor plant-fungi combinations which are most advantageous to both sides under the respective environmental factors (Helgason et al. 2002; Helgason and Fitter 2009; Johnson 2010). As a result of that coevolutionary selection process, both partners would specialize in their interactions and become locally adapted to each other and their abiotic environment (Thompson 2005; Hoeksema 2010).

Mycorrhizas might be understood as a dynamic ‘coadapted mycorrhiza–soil complex’ (Johnson et al. 1993), where both plant and fungal communities continuously adjust to the soil conditions and to one another. Therefore, if local circumstances allow a balanced trading partnership over time, co-adaptation in mycorrhizas and their local soil environment should be promoted (Johnson 2010). In this process, however, selection pressure may be stronger under extreme habitat conditions, for example in phosphorus and/or nitrogen limited soils, or if the interaction between native AM fungi and plant
communities experienced abrupt changes due to introduction and spread of an invasive plant (Pringle et al. 2009; Richardson et al. 2000) altering the mycorrhizal interactions of the resident plant species (e.g. Marler et al. 1999; Mummey and Rillig 2006; Vogelsang and Bever 2009; Zhang et al. 2010).

AM fungal associations are not always advantageous for the plant (Johnson et al. 1993). Under certain environmental conditions, such as high nutrient availability in fertilized systems or reduced photosynthesis, plant growth is decreased by AM fungi (Johnson et al. 1997; Verbruggen and Kiers 2010). Consequently, the plant–AM fungi interaction is viewed to act in a range of functions from mutualism to parasitism mediated by abiotic and biotic environmental conditions, including the plant and fungal genotype (Johnson et al. 1997). Hence, abiotic or biotic variables resulting in disadvantageous effects on one side of the plant–AM fungus relationship might counteract local adaptation in mycorrhizas.

Our understanding of the extent to which local adaptation might be important in arbuscular mycorrhizas is still limited. To date, only two studies focused explicitly on that question and used the approach of reciprocal inoculation of natural AM fungal communities (Johnson et al. 2010; Ji et al. 2010). Johnson et al. (2010) found evidence for co-adaptation because complete ‘home’ combinations of soil, whole soil inoculum and plant ecotype of Andropogon gerardii resulted in the highest fitness of the symbiotic partners in all populations tested. The experiment, moreover, showed that AM fungi and other soil organisms sharing a history in a nitrogen-limited soil were more effective in nitrogen supply to the plant; they were hence locally adapted to soil conditions. In contrast, Ji et al. (2010) demonstrated that adaptation of plants with AM fungal communities might depend on the plant species. Plant growth of Sorghastrum nutans was increased when soils were inoculated with the respective ‘local’ AM fungal spore community, while the origin of the inoculum had no effect on Schizachyrium scoparium. Further, the taxonomic composition of the AM fungal spore communities was also reported to change when the fungal spore inocula were introduced to novel soils. Other studies not employing reciprocal transplanting of AM fungi between study systems also indicated that plants and AM fungi may be adapted in their interactions (Schultz et al. 2001; Klironomos 2003; Pánková et al. 2008; Seifert et al. 2009). For example, Schultz et al. (2001) found a greater growth response by ecotypes of Andropogon gerardii to AM inoculation in phosphorus-limited soil when plants came from these nutrient poor conditions. Klironomos (2003), testing single AM fungal isolate–plant interactions, showed that plant performance was more
strongly affected, both positively and negatively, in ‘home’ combinations of plants and AM fungi compared to pairs where either the plant or the AM fungus were exotic. The equivocal results reported to date highlight the need for additional studies in different ecosystems.

Here we study the interaction between two natural AM fungal assemblages and *Ambrosia artemisiifolia* L. (Asteraceae) in the plant’s new European range. We ask if coevolutionary dynamics may have already led to a coadapted mycorrhiza–soil complex within the introduced range, since there is a potential for coevolution to drive rapid and far-reaching change (Thompson 1999). Thus, we were interested in testing rapid evolution of local adaptation, which has been suggested as an important mechanism and a fundamental issue in invasion ecology (e.g. Sakai et al. 2001, Colautti et al. 2009). Recently, Buswell et al. (2011) showed that rapid adaptive evolution of introduced species might be more common and of greater importance than previously thought.

Our study focus was the regional scale. Methodically, we used the approach of making comparisons between demes within habitats, which corresponds to the ‘local vs. foreign’ criterion (Kawecki and Ebert 2004). The ‘local vs. foreign’ contrast addresses the efficacy of divergent selection relative to other evolutionary processes, and has been proposed as diagnostic for the pattern of local adaptation (Kawecki and Ebert 2004). Therefore, we refer to the ‘local vs. foreign’ terminology; it has analogously already been used by Ji et al. (2010) in studying local adaptation in mycorrhizas in a two-site comparison.

We addressed our question using *A. artemisiifolia* because the plant’s invasive spread is thought to be facilitated by AM fungi (Fumanal et al. 2006). The species, moreover, is known to respond positively to mycorrhizal inoculation (Crowell and Boerner 1988). Further, the plant is ideal to quantify resource allocation to sexes because male and female functions are located in different types of flowers on each individual (e.g. Ackerly and Jasieński 1990; Friedman and Barrett 2011).

In a reciprocal inoculation experiment, we compared the performance of plant populations from two different sites: a roadside habitat and an agricultural field. Our hypothesis was that plant performance/fitness is greater when plants, soil and AM fungal community come from the same site (complete ‘local’ combinations of plant origin, soil and AM fungi) compared to combinations including plants from the other site (here defined as ‘foreign’ plant origin). Further, we ask how the respective AM fungal assemblages function in their own (defined as ‘local’) soil vs. when they are introduced to
new (defined as ‘foreign’) soils. For testing AM fungi in their ‘local’ soil context, we predicted that the AM fungal community from the roadside habitat would act cooperatively, while the AM fungal community from the agricultural field would show less cooperative behavior.

Materials and Methods

Site characteristics

In October 2008, seeds of Ambrosia artemisiifolia were collected from individual plants from two sites in southern Brandenburg, Germany. At each location, plants selected for collecting seeds were randomly chosen and distributed across a wide area (more than 3000 m²) of the population. The first site was a roadside habitat, where A. artemisiifolia plants form dense stands growing over a length of 0.5 km on either side of the road (51°44′02.20″N, 14°27′27.22″E). The second site was an agricultural field planted to Zea mays in the year of seed collection (51°45′15.20″N, 13°58′21.88″E). The distance between the sites was approximately 40 km. For both populations the year of introduction of A. artemisiifolia is unknown, but the region is found to be one large centre of the plant’s distribution in Germany (Brandes and Nitzsche 2006). In this area A. artemisiifolia occurs at the edges of cornfields, on fallows and stubble fields, in intercrop areas and on roadsides, typical of an uneven and disconnected distribution (Brandes and Nitzsche 2006).

Mycorrhizal fungal communities of the two sites exhibited obvious visual differences in terms of root colonization. Root colonization in the roadside habitat was dominated by a special group of AM fungi, which are described as ‘Glomus tenu’ or fine endophytes (Thippayarugs et al. 1999); hereafter referred to as FE. FE are typical for acid soils and characterized by hyphal diameters of less than 1.5 µm (Figure I.1a). In contrast, roots from the agricultural field harbored predominately the ‘normal’ AM fungi (Figure I.1b); hereafter termed coarse AM fungi. In October 2008, roots of A. artemisiifolia of 10 individuals from each site were stained and analyzed. For the roadside habitat we found colonization levels of 53 ± 6 % for coarse AM fungi and 20 ± 5 % for FE. In the cornfield soil, roots were less colonized and showed colonization levels of 17 ± 6 % for coarse AM fungi and 6 ± 3 % for FE.
Soil sampling and analyses

In March 2009, soil was collected from the two locations. We took soil samples only from areas that were occupied by *A. artemisiifolia* during the preceding autumn. At each habitat six soil samples of 10 L each were taken from the top 12 cm of soil. Soil samples per site were pooled, mixed and sieved (5 mm). Half of the soil from each habitat was pasteurized by steaming (Sterilo 1K, Harter Elektotechnik, Schenkenzell, Germany) for four hours at 90 °C. The other half of the soil was used for inoculum extraction.

To analyze soil we took soil samples from each soil after the steaming process. The samples were air-dried, sieved through a 2 mm sieve and analyzed for pH, water repellency, mineral nitrogen (N), mineral carbon (C) and plant available phosphorus (P). Soil pH was determined using both deionized water and in a 1:3 soil:0.01 M CaCl₂ suspension (van Lierop and MacKenzie 1977). Water repellency was measured as the water drop penetration time (Doerr 1998). Mineral N and C contents were determined using a CN analyzer (EuroEA3000-Single), and plant available P was analyzed as calcium-acetate-lactate soluble phosphorus content according to the German standard method DIN 3.4.1.30.2a (Blume et al. 2000).

Experiment

The experiment had a fully 2 x 2 x 3 factorial design and was replicated 12 times. It consisted of all combinations of seeds of *A. artemisiifolia* from the two habitats (seeds collected from the roadside: hereafter roadside seeds, seeds collected from the cornfield: hereafter cornfield seeds), soil from those two locations (roadside soil, cornfield soil), and three soil treatments with two types of inocula (mycorrhizal community from roadside soil, mycorrhizal community from cornfield soil, non-mycorrhizal control).

Mycorrhizal inocula are often prepared from a much smaller volume of soil than that used to fill pots in experiments, leading to potentially unrealistic inocula soil:experiment soil ratios of 1:10 or 1:100. We conducted the experiment with mycorrhizal fungal inocula at the more realistic ‘full soil strength’, meaning that we extracted inocula from the same volume of soil used to fill pots. Soils were wet sieved and inocula were prepared as filtrate (38–212 µm). Our approach of adding inocula as filtrate including AM fungal communities and other soil organisms was different from Ji et al. (2010) using AM fungal spores, but comparable to Johnson et al. (2010), who added
whole-soil inoculum. We chose this method to allow mycorrhizal colonization to establish starting from both AM fungal spores and hyphae (Klironomos and Hart 2002). Further, we checked our inocula for nematodes, which were not detected (binocular microscope). The control treatment received only a mixed microbial wash containing equal parts microbial wash (i.e. filtrate passing a 20 µm sieve) prepared from roadside and field soils (Koide and Li 1989). All treatment pots also received 15 mL of mixed microbial wash to correct for effects resulting from non-AM fungal microbial communities.

The seedlings were germinated in sterilized sand in the greenhouse (day temperature 25 °C, night temperature 16 °C, photoperiod 16 h). Before transplanting, 6 x 25 cm, 400 ml Conetainer pots (Stuewe and Sons, Oregon, USA) were filled with steam pasteurized roadside or cornfield soil. Then, 15 mL of a mixed microbial fraction was added to each pot, and those intended for inoculation received 15 mL of roadside or cornfield AM inoculum while non-inoculated controls received an equivalent amount of water. Plants grew in a fully randomized arrangement for ten weeks in a climate chamber (day temperature 20 °C, night temperature 18 °C, humidity 60 %). We chose a 14 hour photoperiod to encourage optimal development of A. artemisiifolia (Deen et al. 1998). Plants were watered as needed with the same amount of deionized water. Flowering status of the male inflorescences and number of seeds set were recorded weekly during the duration of the experiment.

At harvest, reproductive biomass was carefully removed from shoots. Roots were separated from soil and washed under a stream of water. Biomasses of roots, shoots and male inflorescences were determined after drying for four days at 40 °C. Both ripe and immature seeds were counted and weighed after air-drying to determine reproductive output.

To measure mycorrhizal root colonization, a root sample of each plant (ca. 100 mg of dry root material) was stained using the ink-vinegar method of Vierheilig et al. (1998). Mycorrhizal colonization levels were determined using the magnified intersect method of McGonigle et al. (1990) based on 100 intersections per root sample examined at 200X. Broad shifts in the composition of AM fungal communities in response to ‘local’ vs. ‘foreign’ soil were analyzed by assessing morphological structures of coarse AM fungi and FE (hyphae, arbuscules, vesicles) separately. To determine differences in root architecture, roots of 11 genotypes (six from the roadside, five from the field) were scanned (STD4800 Scanner, resolution 400 dpi) using WinRhizo image analysis system (version Pro 2007b,
Régent Instruments Canada Inc.). Roots with root diameters < 200 µm were classified as fine roots and those between 200–1000 µm as coarse roots.

Statistical analyses

In local adaptation studies plant performance is commonly based on more than one individual trait, leading to non-independent data per plant individual (Kawecki and Ebert 2004); we thus statistically analyzed plant responses as a plant trait syndrome. Therefore, influence of the main effects (seed origin, soil, soil treatment) and their interactions on plant performance were investigated using Principal Component Analysis (PCA) on the correlation matrix (Legendre and Legendre 1998; Gotelli and Ellison 2004). This ordination approach enabled us to reduce the dimensionality of the multivariate dataset to a few non-correlated new variables. We performed three PCAs focusing on three different plant trait aspects: biomass, root traits and mycorrhization. Each single PCA included a set of seven variables.

PCA on plant biomass was calculated on standardized data of shoot and root biomass, mass of male inflorescences and both ripe and immature seeds, number of seeds produced after five weeks of growth, and total seed number. To meet assumptions of normality and homogeneity data concerning seed number were log-transformed, whereas all other biomass variables were Box-Cox transformed. Time of flowering was not included in the analysis because of missing values. PCA on root traits was computed from standardized data of variables of root length per volume, root surface area, both fine and coarse root volume, both fine and coarse root length, and root diameter. Here, measures of root diameter were log-transformed to meet assumptions of normality. PCA on mycorrhization included standardized variables of percentage total AM fungal root colonization, and percentage root colonization by hyphae, arbuscules and vesicles of both coarse AM fungi and FE. The original data were log-transformed to achieve normality.

To test for differences among treatment groups, principal component scores from the first and second axis of the PCA were treated analogously to the univariate response variables, which were analyzed by Analysis of Variance (three-way ANOVA). Differences between treatment groups were compared with Tukey HSD post-hoc comparison tests ($P < 0.05$). To identify differences in soil properties we used a two sample $t$-test ($P < 0.05$). All statistical analyses were performed using R version 2.10.1 (R Development Core Team 2009).
Results

Soil properties

The soils of the locations differed significantly in soil pH, N, and water repellency (Table I.1). Extractable contents of C and P were similar.

Variance explained by first and second principal components of PCA

The first principal component (PC1) of the three PCAs on the different traits – plant biomass, root morphology, mycorrhization – of A. artemisiifolia accounted for more than 50% of the variance (Table I.2, Figure I.2). In the PCA on root traits, variance explained by PC1 was even higher (70%; see Table I.2). In all PCAs, a significant interaction term indicated that responses of PC1 scores to soil were mediated by mycorrhizal treatment (Table I.3). Moreover, for the PCAs on plant biomass and root traits we found that PC1 scores were also significantly influenced by the interaction between plant origin and soil. This effect, however, could not be shown for root traits with pairwise comparison tests (Tukey HSD; P > 0.05). Furthermore, PC1 scores were consistently strongly influenced by the factor soil (Table I.3).

Regarding the second principal component (PC2), patterns differed depending on the trait aspect analyzed. Details on analyses of variance (ANOVA) on PC2 are reported in the Supplemental Table A.I.1. PC2 of the PCA on biomass explained 20% of data variance (Table I.2), but PC2 scores were not significantly influenced by the factors tested (Supplemental Table A.I.1). PC2 of the PCA on root traits accounted for 25% of the variance (Table I.2) and was significantly influenced by plant origin only (Supplemental Table A.I.1). In contrast, PC2 of the PCA on mycorrhization explained 42% of the variance (Table I.2) and was significantly impacted by both the soil–mycorrhizal treatment and plant origin–mycorrhizal treatment interactions (Supplemental Table A.I.1). Effects of the plant origin–mycorrhizal treatment term could not be shown with pairwise comparison tests (Tukey HSD; P > 0.05).
Soil conditions had a strong effect on the plant traits tested. Plants growing in roadside soil had considerably less vegetative biomass (202.8 ± 11.5 mg) compared to those in cornfield soil (665.9 ± 25.9 mg). In addition, seed mass of ripe seeds was on average more than doubled in cornfield soil (94.9 ± 9.0 mg) compared to roadside soil (44.4 ± 3.9 mg). Male flower biomass was also higher in cornfield soil (77.5 ± 3.9 mg) compared to roadside soil (28.9 ± 2.0 mg). Flowering started significantly earlier in cornfield soil (roadside soil: 4.5 ± 0.1 weeks, cornfield soil: 3.7 ± 0.1 weeks, soil effect: F_{1,130} = 38.4; P < 0.001, ANOVA).

Concerning root traits, plants growing in cornfield soil had twice the coarse root length (2310.3 ± 91.9 cm) compared to those in roadside soil (1149.6 ± 90.1 cm). Fine root length, moreover, was also higher in cornfield soil (640.7 ± 41.3 cm) than in roadside soil (490.1 ± 47.9 cm), although the increase was proportionally smaller. Hence, plants in roadside soil had finer root systems with on average smaller root diameters (0.180 ± 0.004 mm) compared to those in cornfield soil (0.250 ± 0.006 mm).

In terms of mycorrhization, plants grown in soils with AM fungal assemblages both from the roadside and the cornfield soil were colonized with AM fungi (Supplemental Table A.I.2). Non-mycorrhizal controls, however, were rarely infected by fungi (0.30 ± 0.10 %), none of which could be classified as AM fungi. Percentage of total AM fungal root colonization was on average 31 % higher in roadside soil (78 ± 2 %) than in cornfield soil (47 ± 3 %).

Plant responses to soil and mycorrhizal treatment

The vast majority of response variables indicated that effects of mycorrhizal treatments (‘local’, ‘foreign’ soil inoculum, and non-mycorrhizal control) on plant traits differed depending on the soil type. Details on measured variables in response to soil and mycorrhizal treatment are reported in the Supplemental Table A.I.2.

Regarding the soil–mycorrhizal treatment effect for both the PCAs on plant biomass and root traits, PC1 scores were negative in roadside soil; in cornfield soil the values were positive (Figures I.3a and I.3b). For the PCA on biomass, higher or less negative PC1 scores should be interpreted as greater biomass, because both shoot and root biomass, and also number of seeds had the highest loading on PC1 (Table I.2). In terms of
root traits, higher PC1 scores reflect bigger and also coarser root systems, since variables of root surface area, root length per volume, and coarse root length had the highest influence on PC1 (Table I.2).

For plant biomass in roadside soil, non-mycorrhizal control plants had the lowest PC1 scores; hence, had significantly less biomass compared to other combinations tested. For ‘local’ and ‘foreign’ inoculum in roadside soil, PC1 scores were significantly higher compared to control, thus plant biomass was significantly increased in the presence of both AM fungal inocula (Figure I.3a). In cornfield soil, conversely, control plants had the highest PC1 scores, i.e. plants had greatest biomass in the absence of AM fungi. Moreover, ‘local’ soil inoculum caused a significant decrease in plant biomass compared to non-mycorrhizal controls in cornfield soil. Biomass of plants growing in cornfield soil with the ‘foreign’ inoculum was not different from those in control or ‘local’ inoculum treatments (Figure I.3a).

Regarding response patterns on root traits, PC1 scores did not significantly differ for the different mycorrhizal treatments in roadside soil (Figure I.3b). In cornfield soil, however, combination of cornfield soil with cornfield inoculum generated a significant decrease in PC1 scores compared to combinations with roadside inoculum (Figure I.3b). This means that in cornfield soil root systems were significantly smaller and less coarse when plants were treated with the ‘local’ compared to ‘foreign’ soil inoculum.

Concerning mycorrhization, we found strong proportional shifts between coarse AM fungi and FE colonizing roots among all treatment combinations of soil and mycorrhizal inoculum, which is indicated by both PC1 and PC2 scores revealing significant soil–mycorrhizal treatment interactions (Table I.3, PC2 soil x mycorrhizal treatment: F_{1,132}=16.41; P < 0.001, ANOVA). In spite of these shifts, cornfield inoculum always resulted in higher root colonization by coarse AM fungi than FE; for the roadside inoculum higher proportions of FE structures were typical (Supplemental Table A.I.2). Differences in root colonization resulting from inoculum application to ‘local’ or ‘foreign’ soils were mainly related to the dominant AM fungal component of the respective inoculum only. Treating roadside soil with cornfield inoculum increased root colonization by coarse AM fungal structures, while for cornfield soil with roadside inoculum a strong decrease in FE colonization was found (Supplemental Table A.I.2).

Focusing on PC1, scores were more negative the higher the root colonization with FE and total AM fungi; non-mycorrhizal controls had the highest scores (see Figure I.3c). Thus, PC1 scores were lowest for the combination of roadside soil with roadside inoculum,
which had highest colonization both with total AM fungi and FE (Figure I.3c, Supplemental Table A.I.2).

Plant origin–soil interaction and effects of plant origin

Regarding biomass patterns, PC1 scores significantly differed for ‘local’ and ‘foreign’ plant origin tested in roadside soil, with roadside origin plants being larger (Figure I.4). This indicates that in roadside soil plants with roadside origin produced significantly more biomass than plants originating from cornfield seeds (higher PC1 scores correlate with greater biomass, Figure I.2a).

In addition, function of the male gender was also significantly affected by the plant origin–soil interaction (plant origin x soil: $F_{1,132} = 11.82, P < 0.001$, ANOVA). Here, the difference was significant for cornfield soil only: plants with cornfield origin had significantly greater male flower biomass compared to plants with roadside origin (roadside origin: $66.0 \pm 4.1$ mg, cornfield origin: $89.3 \pm 6.0$ mg, $P = 0.01$ for pairwise comparison between roadside and cornfield plant origin in cornfield soil). Details on other variables in response to soil and plant origin are reported in the Supplemental Table A.I.3.

The factor plant origin significantly influenced PC2 scores of the PCA on root traits (plant origin: $F_{1,54} = 9.52; P = 0.003$, ANOVA). Because response variables describing fine root attributes had the highest loading on PC2 (Table I.2), we highlight fine root length and average root diameter. Plants originating from roadside seeds had on average smaller root diameters ($0.209 \pm 0.008$ mm) compared to those from cornfield seeds ($0.221 \pm 0.007$ mm). In addition, genotypes with roadside origin produced on average 43% more fine root length ($655.3 \pm 42.3$ cm) compared to those with cornfield origin ($457.6 \pm 44.4$ cm). Plant origin also significantly influenced time of flowering (plant origin: $F_{1,130} = 12.5, P < 0.001$, ANOVA). Plants originating from cornfield seeds flowered earlier ($3.9 \pm 0.1$ weeks) than those from roadside seeds ($4.3 \pm 0.1$ weeks).
Discussion

Mycorrhizal functions depending on soil and inoculum source

In this study, natural AM fungal assemblages were found to exhibit different qualities of cooperation with non-native *A. artemisiifolia* depending on mycorrhizal inoculum source and soil conditions. In less fertile roadside soil, presence of AM fungal communities significantly improved plant performance compared to the non-mycorrhizal control regardless of whether the inoculum came from the roadside or the cornfield habitat. The positive effect of mycorrhizal inoculation on *A. artemisiifolia* in roadside soil was reflected both in greater shoot and root biomass, and also higher number of seeds produced in comparison to the non-mycorrhizal treatment. As a result, enhanced vegetative plant growth was positively correlated with increased reproductive output indicating higher plant fitness: such a relationship is common in herbaceous plants (Allison 2002). In the comparatively adverse roadside habitat, which can also be expected to be corridors of invasion (Joly et al. 2011), mycorrhizal association may thus increase fitness of *A. artemisiifolia*, which may promote the plant’s spread as suggested by Fumanal et al. (2006).

In contrast, in cornfield soil, where plants on average had almost triple the biomass compared to the roadside soils, we found no evidence of a beneficial role of AM fungi on plant performance. Consistent with empirical and theoretical studies (e.g. Johnson et al. 1993; Johnson et al. 1997; Verbruggen and Kiers 2010), our results demonstrate that the AM fungal community from the managed agricultural system acted less cooperatively, i.e. decreased plant biomass, but only in its ‘local’ soil. Thus, less cooperative or ‘cheating’ behavior of the cornfield AM fungal community was only present when the ecological soil–AM fungi context was maintained. The importance of the ecological context for mycorrhizal functions, as recently comprehensively shown by Hoeksema et al. (2010), was also evident for the impact of the roadside AM fungal community. Treating agricultural soil with roadside inoculum mediated a neutral effect on plant growth compared to the non-mycorrhizal treatment.

Consequently, we found AM fungal assemblages having neutral to negative effects on *A. artemisiifolia* in more fertile agricultural soil, while the function in nutrient limited roadside soil was mutualistic as suggested by the trade balance model (Johnson 2010). We attribute these contrasting effects of mycorrhizal functions in the experiment to abiotic soil
conditions, and also the identity of AM fungi establishing the roots. The effect of soil was evident, since total root colonization was of similar magnitude lower in cornfield compared to roadside soil irrespective of inoculum origin, and although we always inoculated soil with ‘full soil strength’ inocula. Hence, ‘parasitic’ or ‘commensalistic’ functions of AM fungi in cornfield soil were not related to a different degree in mycorrhizal root colonization, but likely to different taxa of AM fungi establishing in the roots. Our assessment of fungal structures showed broad shifts in AM fungal taxa colonizing the plant roots depending on soil and inoculum identity, which may have also been influenced by cultivation bias (Sýkorová et al. 2007). Overall, the colonization levels of coarse AM fungi and FE differed significantly among all combinations of soil and mycorrhizal inoculum tested. The different inocula compositions, moreover, were reflected in a trade-off implying that when roots were highly colonized with coarse AM fungi, colonization with FE was low and vice versa. Therefore, divergent functions of AM fungal assemblages in cornfield soil were related to AM fungal taxa, which had colonized the roots. This finding, however, is not surprising given the fact that mycorrhizal function is known to vary among AM fungal genotypes (van der Heijden et al. 1998; Klironomos 2003; Munkvold et al. 2004; Antunes et al. 2011), as well as families (Powell et al. 2009).

Moreover, we found that the negative effect of the mycorrhizal symbiosis in agricultural soil inoculated with the ‘local’ AM fungal community coincided with increased root fineness compared to inoculation with ‘foreign’ AM fungi. This demonstrates that the presence of less cooperative AM fungal partners can result in greater branching enabling plants to acquire soil nutrients more directly via fine roots, which is advantageous in nutrient rich soils. For example, Schultz et al. (2001) reported that Andropogon gerardii had a more branched and finer root system, and was also less dependent upon mycorrhizal symbiosis, when genotypes came from high-nutrient soil. Similar relationships were also found for plant populations of Hypericum perforatum, which had greater root fineness and reduced mycorrhizal responsiveness in the non-native North American range (Seifert et al. 2009). Moreover, A. artemisiifolia may be less dependent upon mycorrhizal symbiosis in the new range than expected (Fumanal et al. 2006) because we found no evidence for root systems to be modified towards greater coarseness in the presence of mycorrhizal fungi compared to the control, which often is an indication of mycorrhizal dependency in other plant species (e.g. Hetrick 1991; Berta et al. 2002).
Adaptation to soil environment and plant origin effects

We found no evidence for co-adaptation between *A. artemisiifolia* and AM fungi in the new range because plants both originating from roadside and cornfield seeds grew equally with the respective AM fungal communities. However, plants with roadside origin performed significantly better in roadside soil, which demonstrates adaptation to the abiotic soil environment in the introduced range. In the native range, adaptation of *A. artemisiifolia* to roadside conditions, in particular to high salinity concentrations in roadside soils, has also been reported for the germination behavior of seeds (DiTommaso 2004). Further, results of a few studies indicated adaptation of *A. artemisiifolia* to climate. Recently, Hodgins and Rieseberg (2011) demonstrated adaptation to latitude and climate variables by comparing European with North American populations in common garden experiments. In addition, Chun et al. (2011) found effects of geographic location on reproductive allocation in introduced French populations. In a reciprocal transplant experiment, moreover, weather was found to have a huge effect on re-growth of *A. artemisiifolia* plants threatened with the herbicide imazethapyr, although the reproductive potential of the plants depended on seed origin, suggesting that genetic differences may be a result of evolution of different ecotypes (Leif et al. 2000).

Male function also indicated adaptation of plants with cornfield origin to ‘local’ soil. In cornfield soil, plants originating from cornfield seeds had significantly more male flower biomass (on average 35 %) than those from roadside seeds. Plant origin aside, more fertile soil conditions strongly increased resource allocation to male inflorescences, thus plants had 2.5 times greater male biomass, and flowered earlier. The correlation of greater plant biomass with a shift towards maleness in nutrient-rich soils was also observed by Ackerly and Jasieński (1990). In a recent study, Friedman and Barrett (2011) showed that sex allocation also varies with light conditions, where plants grown in the sun had higher male flower production.

Plant origin strongly affected root morphology. Plants with roadside origin had significantly finer root systems with smaller root diameters in all treatment situations tested compared to those with cornfield origin. This may result from genetic differences between the populations or maternal effects. Compared to plants in cornfield soil, conspecifics growing in roadside soil produced root systems with smaller root diameters and proportionally greater fine root length: apparently, these root alterations were more
efficient in nutrient acquisition in nutrient poor roadside soil because roots explored a larger soil volume per unit root surface area (Gahoonia and Nielsen 2003).

Conclusion

This study shows that performance of non-native *A. artemisiifolia* might be significantly influenced by mycorrhizal functions, leading to a positive effect in less fertile roadside soil and parasitism in more fertile cornfield soil *sensu* Johnson (2010). The effects of natural AM fungal assemblages are strongly soil context dependent (Hoeksema et al. 2010), thus mycorrhizal functions may be unpredictable when AM fungal communities are introduced to novel soils.

For the establishment phase, local adaptation between *A. artemisiifolia* and natural AM fungal assemblages may be of no relevance, highlighting the low host-specificity of AM associations (Richardson et al. 2000). In harsh environmental conditions such as roadside habitats, however, adaptation of *A. artemisiifolia* to soil may play a crucial role in the early stages of invasion, as well as the symbiosis with AM fungi enhancing plant growth and fitness, thus promoting invasive spread. Along the road, therefore, co-adaptation between AM fungi and *A. artemisiifolia* may not be utterly out of the question in later stages of the invasion process (Thompson 2005; Pringle et al. 2009; Hoeksema 2010).

Acknowledgements

We are deeply grateful to E. Kathryn Barto and Tancredi Caruso for discussion and statistical advice. We also thank Freie Universität Berlin for financial support.
Tables and figures

Table I.1 Characteristics of the soils used. Analyses of soil pH (as measured by deionized water and exchangeable in a 1:3 soil:0.01 M CaCl₂ suspension), water repellency, extractable contents of nitrogen (N), carbon (C), and plant available phosphorus (P) refer to soil properties after steaming. Values represent means ± SE (n = 3, except for water repellency n = 5). P-values relate to t-tests for independent samples. Values in bold indicate significance at P < 0.05.

<table>
<thead>
<tr>
<th>Soil</th>
<th>Roadside</th>
<th>Cornfield</th>
<th>P</th>
</tr>
</thead>
<tbody>
<tr>
<td>pH (H₂O)</td>
<td>5.54 ± 0.06</td>
<td>6.12 ± 0.03</td>
<td>&lt;0.001</td>
</tr>
<tr>
<td>pH (CaCl₂)</td>
<td>4.51 ± 0.03</td>
<td>5.31 ± 0.02</td>
<td>&lt;0.001</td>
</tr>
<tr>
<td>Water repellency (sec)</td>
<td>12.4 ± 4.2</td>
<td>2.2 ± 1.2</td>
<td>&lt;0.001</td>
</tr>
<tr>
<td>Total C in %</td>
<td>0.560 ± 0.153</td>
<td>0.581 ± 0.043</td>
<td>0.827</td>
</tr>
<tr>
<td>Total N in %</td>
<td>0.028 ± 0.007</td>
<td>0.047 ± 0.003</td>
<td>0.013</td>
</tr>
<tr>
<td>Plant available P (mg per 100 g soil)</td>
<td>6.25 ± 2.50</td>
<td>8.33 ± 2.20</td>
<td>0.339</td>
</tr>
</tbody>
</table>
Table I.2 Eigenvectors of the first five principal components (cumulative % variance explained > 97) of the Principal Component Analyses (PCA) performed on the correlation matrix of three different sets of plant traits of biological response variables listed in the first column.

<table>
<thead>
<tr>
<th>PCA</th>
<th>PC1</th>
<th>PC2</th>
<th>PC3</th>
<th>PC4</th>
<th>PC5</th>
</tr>
</thead>
<tbody>
<tr>
<td>Plant biomass</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Shoot biomass</td>
<td>0.475</td>
<td>-0.165</td>
<td>0.178</td>
<td>-0.054</td>
<td>0.365</td>
</tr>
<tr>
<td>Root biomass</td>
<td>0.434</td>
<td>-0.333</td>
<td>0.140</td>
<td>-0.031</td>
<td>0.529</td>
</tr>
<tr>
<td>Reproduction 5 weeks</td>
<td>0.220</td>
<td>0.634</td>
<td>-0.125</td>
<td>0.687</td>
<td>0.245</td>
</tr>
<tr>
<td>Total seed number</td>
<td>0.451</td>
<td>0.179</td>
<td>-0.294</td>
<td>-0.248</td>
<td>-0.297</td>
</tr>
<tr>
<td>Ripe seed wt.</td>
<td>0.166</td>
<td>0.571</td>
<td>0.629</td>
<td>-0.442</td>
<td>-0.093</td>
</tr>
<tr>
<td>Immat. seed wt.</td>
<td>0.415</td>
<td>0.056</td>
<td>-0.553</td>
<td>-0.233</td>
<td>-0.186</td>
</tr>
<tr>
<td>Male flower wt.</td>
<td>0.367</td>
<td>-0.315</td>
<td>0.382</td>
<td>0.461</td>
<td>-0.629</td>
</tr>
<tr>
<td>Proportion of total SS</td>
<td>53%</td>
<td>20%</td>
<td>12%</td>
<td>7%</td>
<td>5%</td>
</tr>
<tr>
<td>Root traits</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Root length per volume</td>
<td>0.443</td>
<td>0.059</td>
<td>-0.310</td>
<td>-0.221</td>
<td>0.017</td>
</tr>
<tr>
<td>Root surface area</td>
<td>0.443</td>
<td>-0.137</td>
<td>0.061</td>
<td>-0.184</td>
<td>0.426</td>
</tr>
<tr>
<td>Fine root volume</td>
<td>0.288</td>
<td>0.584</td>
<td>0.019</td>
<td>0.139</td>
<td>-0.661</td>
</tr>
<tr>
<td>Fine root length</td>
<td>0.264</td>
<td>0.613</td>
<td>0.114</td>
<td>0.293</td>
<td>0.570</td>
</tr>
<tr>
<td>Coarse root volume</td>
<td>0.384</td>
<td>-0.212</td>
<td>0.816</td>
<td>-0.182</td>
<td>-0.176</td>
</tr>
<tr>
<td>Coarse root length</td>
<td>0.428</td>
<td>-0.135</td>
<td>-0.451</td>
<td>-0.357</td>
<td>-0.132</td>
</tr>
<tr>
<td>Root diameter</td>
<td>0.351</td>
<td>-0.445</td>
<td>-0.131</td>
<td>0.808</td>
<td>-0.089</td>
</tr>
<tr>
<td>Proportion total SS</td>
<td>70%</td>
<td>25%</td>
<td>4%</td>
<td>0.01%</td>
<td>0.001%</td>
</tr>
<tr>
<td>Mycorrhization</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Total root colonization</td>
<td>-0.511</td>
<td>0.112</td>
<td>-0.170</td>
<td>0.367</td>
<td>-0.208</td>
</tr>
<tr>
<td>coarse AMF hyphae</td>
<td>-0.317</td>
<td>0.451</td>
<td>-0.233</td>
<td>-0.046</td>
<td>-0.611</td>
</tr>
<tr>
<td>coarse AMF arbuscules</td>
<td>-0.264</td>
<td>0.476</td>
<td>-0.415</td>
<td>-0.250</td>
<td>0.679</td>
</tr>
<tr>
<td>coarse AMF vesicles</td>
<td>-0.249</td>
<td>0.431</td>
<td>0.857</td>
<td>0.016</td>
<td>0.128</td>
</tr>
<tr>
<td>FE hyphae</td>
<td>-0.421</td>
<td>-0.348</td>
<td>0.022</td>
<td>0.200</td>
<td>0.062</td>
</tr>
<tr>
<td>FE arbuscules</td>
<td>-0.408</td>
<td>-0.362</td>
<td>0.021</td>
<td>0.325</td>
<td>0.289</td>
</tr>
<tr>
<td>FE vesicles</td>
<td>-0.405</td>
<td>-0.345</td>
<td>0.095</td>
<td>-0.809</td>
<td>-0.136</td>
</tr>
<tr>
<td>Proportion total SS</td>
<td>51%</td>
<td>42%</td>
<td>5%</td>
<td>1%</td>
<td>0.01%</td>
</tr>
</tbody>
</table>

SS, sum of squares; Ripe seed wt., weight of ripe seeds; Immat. seed wt., weight of immature seeds; FE, fine endophytes; coarse AMF, coarse AM fungi
Table I.3 Analyses of variance (ANOVA) on the first principal component score (PC1) of the Principal Component Analyses (PCA) on plant biomass, root traits and mycorrhization. Values in bold indicate significance at $P < 0.05$.

<table>
<thead>
<tr>
<th>Factors</th>
<th>d.f.</th>
<th>PCA</th>
<th>Plant Biomass</th>
<th>Root traits</th>
<th>Mycorrhization</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td></td>
<td>$F$</td>
<td>$P$</td>
<td>$F$</td>
<td>$P$</td>
</tr>
<tr>
<td>P.ori</td>
<td>1</td>
<td>1.98</td>
<td>0.162</td>
<td>1.93</td>
<td>0.170</td>
</tr>
<tr>
<td>Soil</td>
<td>1</td>
<td><strong>942.65</strong></td>
<td>&lt;0.001</td>
<td><strong>92.18</strong></td>
<td>&lt;0.001</td>
</tr>
<tr>
<td>Myc.treat</td>
<td>2</td>
<td>0.72</td>
<td>0.488</td>
<td>2.22</td>
<td>0.118</td>
</tr>
<tr>
<td>P.ori x Soil</td>
<td>1</td>
<td><strong>5.33</strong></td>
<td><strong>0.022</strong></td>
<td><strong>4.41</strong></td>
<td>0.040</td>
</tr>
<tr>
<td>P.ori x Myc.treat.</td>
<td>2</td>
<td>0.24</td>
<td>0.786</td>
<td>1.44</td>
<td>0.246</td>
</tr>
<tr>
<td>Soil x Myc.treat.</td>
<td>2</td>
<td><strong>17.00</strong></td>
<td>&lt;0.001</td>
<td><strong>3.75</strong></td>
<td><strong>0.030</strong></td>
</tr>
<tr>
<td>P.ori x Soil x Myc.treat.</td>
<td>2</td>
<td>1.10</td>
<td>0.337</td>
<td>0.11</td>
<td>0.897</td>
</tr>
<tr>
<td>Residuals</td>
<td>132</td>
<td>54</td>
<td>132</td>
<td></td>
<td></td>
</tr>
</tbody>
</table>

d.f., degree of freedom; P.ori., plant origin; Myc.treat., mycorrhizal treatment
**Figure I.1** Colonization with fine endophytes (a) and ‘coarse’ AM fungi (b) in roots of *Ambrosia artemisiifolia* in the experiment. Roots were stained with ink-vinegar method (Vierheilig et al. 1998). Photo credit: Cornelia Bäucker.
Figure I.2 Loading plots of the Principal Component Analyses (PCAs) performed on the correlation matrix of variables of different plant trait aspects: (a) biomass traits including weight of ripe seeds (Ripe seed wt.), number of seeds produced after five weeks (Reproduction after 5 weeks), total seed number (Seed number), weight of immature seeds (Immat. seed wt.), shoot biomass, root biomass, weight of male inflorescences (Male flower wt.); (b) root traits including fine root length, volume of fine roots (fine root vol.), root length per volume soil (Root length per vol.), coarse root length, root surface area (Surface area), volume of coarse roots (Coarse root vol.), root diameter; (c) mycorrhization including percentage total root colonization (Total root colo.), coarse AM fungal root colonization by hyphae (AMF hyph.), vesicles (AMF ves.), arbuscules (AMF arbus.), and fine endophyte root colonization by hyphae (FE hyph.), vesicles (FE ves.), and arbuscules (FE arbus.). The variable arrow coordinates are built from PC1 and PC2 eigenvector coefficients (Table I.3) and visualize how variables correlate with each other.
Figure I.3 Responses to soil and mycorrhizal treatments of *Ambrosia artemisiifolia* in the experiment: PC1 scores refer to the PCA on plant biomass traits (a); root traits (b); mycorrhization (c). Bar plots of mean (± SE) indicate PC1 score variation in response to treatments of soils inoculated with ‘local’ soil AM fungal inoculum (black), ‘foreign’ soil AM fungal inoculum (grey), and non-mycorrhizal controls (white). For (a) and (b) higher or less negative PC1 scores correspond to greater biomass and bigger root systems of greater coarseness, respectively. In (c) lower PC1 scores mean higher root colonization both with AM fungal hyphae in total and structures of fine endophytes (hyphae, arbuscules, vesicles). Different lower case letters on the bars indicate a significant difference ($P < 0.05$) among treatment groups according to Tukey’s HSD test.
Figure I.4 First principal component scores of the PCA on biomass traits of *Ambrosia artemisiifolia* indicating local adaptation to roadside soil in the experiment: higher PC1 scores (means ± SE) indicate greater plant biomass. Black bars correspond to the combination of soil with the respective ‘local’ plant origin and grey bars to ‘foreign’ origin. ‘Local soil plant origin’ means that plants grew in their respective own soil, i.e. plants from the roadside in roadside soil and plants from the cornfield in cornfield soil, respectively. The term ‘foreign plant soil origin’ refers to new combinations of plants and soil, i.e. when plants from the cornfield grew in roadside soil and plants from the roadside grew in cornfield soil, respectively. Different lower case letters on the bars indicate a significant difference ($P < 0.05$) among treatment groups according to Tukey’s HSD test.
References


CHAPTER 3

Non-native Ambrosia artemisiifolia are more influenced by relative density and identity of neighboring plant species than arbuscular mycorrhiza

Abstract

Arbuscular mycorrhizal (AM) fungi can play a crucial role in plant invasion processes by mediating a competitive advantage of invasive over native species. Since Ambrosia artemisiifolia, a plant native to North America, has been proposed to be facilitated by AM fungi in Europe, we investigated the effects of natural AM fungal communities on its competitive ability in two greenhouse experiments always maintaining the ecological context of soil and AM fungi. We studied A. artemisiifolia grown together with one of four co-existing mycorrhizal plant species in a 1:4 (target) or 4:1 (challenger) relative density. The neighbor species were Conyza canadensis, Daucus carota, Artemisia vulgaris, and Tanacetum vulgare. We found no clear evidence that the association with natural AM fungal assemblages increased the competitive ability of A. artemisiifolia. Regardless of presence/absence of AM fungi, A. artemisiifolia was highly dominant in all interspecific competitive arrangements under low soil fertility conditions. Further, we found that decreased shoot growth in the presence of conspecifics coincided with reduced number of male inflorescences in A. artemisiifolia, but also led to earlier development of female flowers. Likewise, in a pairwise competition experiment of A. artemisiifolia and D. carota, the invasive species was always dominant and we found no indication that AM fungi influenced the interspecific competitive outcome. However, AM fungi had a significant amplifying effect on intraspecific competition of A. artemisiifolia. Our findings suggest that the competitive ability of A. artemisiifolia, a species with a strongly developed ruderal life history, is very weakly influenced by natural AM fungal communities in presence of other mycorrhizal plant competitors under nutrient poor soil conditions. Hence, the invasive success of A. artemisiifolia in Middle Europe may be related to mechanisms other than facilitation by natural AM fungal communities.
Introduction

Invasive plant species are ‘naturalized plants that produce reproductive offspring, often in very large numbers, at considerable distances from parent plants and thus have the potential to spread over a considerable area’ (Richardson et al. 2000a). Worldwide, numerous ecosystems are affected by invasive plants in their functioning in a multitude of ways (Wardle et al. 2011). Invasive species, moreover, can damage ecosystem services that are fundamental to human well-being resulting in substantial economic costs (e.g. Pimentel et al. 2005; Pejchar and Mooney 2009; Vilà et al. 2010). Therefore, it is of increasing urgency to better understand the mechanisms involved in the invasion process as human activities, such as international trade, transport and travel, which cause species dispersal into new ranges, continue to expand (Keller et al. 2011).

Mechanisms by which invasive plants successfully spread into new areas may primarily be related to interactions of the invasive species with their new environment (Jeschke et al. 2012). Therefore, aboveground mutualistic interactions, such as pollination or seed dispersal, may play an important role for the success of introduced plants (Richardson et al. 2000b). In addition, belowground symbioses, such as mycorrhizal relationships, have also been shown to be a critical aspect (Richardson et al. 2000b; Pringle et al. 2009; Shah et al. 2009). For example, arbuscular mycorrhizal (AM) fungi, which form complex underground hyphal networks with roots of around two-thirds of plant species (Helgason and Fitter 2009), and provide multiple functions, especially increased phosphorus uptake to the plant in exchange for photoassimilates (Newsham et al. 1995), are known to be a crucial factor in some plant invasions. One way by which invasive plants can interact with natural AM fungal communities is by disrupting the formation of mycorrhizal associations upon which many native plant species depend (Stinson et al. 2006; Meinhardt and Gehring 2012). Another important possibility how invasive plants can interact with AM fungi has been described as facilitation (Reinhart and Callaway 2006; Shah et al. 2009), whereby invasive plants are positively influenced by the AM fungal association of the new range to the detriment of native species.

Several studies have documented that AM fungi can contribute to the dominance of invasive over native plants by altering competitive interactions (Marler et al. 1999; Callaway et al. 2003; Callaway et al. 2004; Callaway et al. 2005; Shah et al. 2008; Wilson et al. 2012; but see Bray et al. 2003; Emery and Rudgers 2012). However, the mechanism underlying this increased competitive ability of invasive plants has not been fully
elucidated; it might be caused by changes in AM fungal community composition in presence of the invasive species (Mummey and Rillig 2006; Hawkes et al. 2006; Zhang et al. 2007), or increased P-uptake of the invasive plant via AM fungal mycelia (Zabinski et al. 2002). Further, the outcome of competitive interactions between plants has been shown to be influenced by identity of AM fungi (Scheublin et al. 2007; Shah et al. 2008). Therefore, the ecological context of studies investigating the effect of AM fungi on competitive interactions between invasive and native plant species has to be carefully considered, as recently demonstrated by Hoeksema et al. (2010).

Here, we study the competitive ability of Ambrosia artemisiifolia L. (Astereceae), an annual plant native to North America. There, the species is known to be dominant during early stages of old-field succession in many parts of the eastern and midwestern United States (Bazzaz and Mezga 1973). Introduced to other continents, the species successfully spreads, and poses a risk for human health by producing pollen causing pollen allergy and allergic asthma (e.g. Török et al. 2003). A special trait of the monoecious genus Ambrosia is that sexes are located in different types of flowers on the individuals; hence, A. artemisiifolia allows quantification of resource allocation to functional genders (Friedman and Barrett 2011). The plant occurs in different types of disturbed habitats, for example, roadsides, construction sites, waste lands, and agricultural fields, where it is an important weed (Brandes and Nitzsche 2006). Moreover, A. artemisiifolia is a mycorrhizal plant and its spread has been suggested to be facilitated by the symbiosis with arbuscular mycorrhizal fungi in Europe (Fumanal et al. 2006). In a greenhouse experiment, Crowell and Boerner (1988) showed that A. artemisiifolia responds positively to inoculation with mycorrhizal fungi: plant shoot and total biomass were more than 20 times increased in the presence of AM fungi (the AM fungal taxon tested was Glomus etunicatum). The same study, further, reports that interspecific competition with the non-mycorrhizal Brassica nigra was stronger than intraspecific competition in the presence of AM fungi: here, AM fungal inoculation conferred no advantage to the mycorrhizal competitor, which is rather atypical for competitive situations of mycorrhizal with non-mycorrhizal plants (Moora and Zobel 2010). Other studies on competition, however, found that A. artemisiifolia had a large suppressive effect on other co-existing species; hence, A. artemisiifolia was competitively dominant (Miller and Werner 1987; Miller 1994; but see Leskovšek et al. 2012).

This study aimed to investigate whether the competitive ability of A. artemisiifolia is enhanced by the symbiosis with natural AM fungi in the presence of co-occurring plant
species in the new European range under conditions of a maintained ecological context of soil and AM fungal communities. Our expectations were: 1) AM fungi should confer a competitive advantage on *A. artemisiifolia* in presence of other mycorrhizal competitors; 2) the competitive outcome should be independent of the neighboring plant species; 3) the growth of neighbors should be more decreased in presence of a high than a low density of *A. artemisiifolia*.

**Materials and Methods**

We conducted two greenhouse experiments with comparatively high realism, i.e., we always maintained the ecological context of soil and natural AM fungal communities (Hoeksema et al. 2010). The first experiment focused on *A. artemisiifolia* in target and challenger arrangements in combination with four different neighboring plant species in presence/absence of AM fungi. As neighbor species we chose *Conyza canadensis* L. (Asteraceae), *Daucus carota* L. (Apiaceae), *Artemisia vulgaris* L. (Asteraceae), and *Tanacetum vulgare* L. (Asteraceae): all co-existing with *A. artemisiifolia* and mycorrhizal. Hereafter, we term this experiment target–challenger experiment. Because this target–challenger experiment indicated that the mycorrhizal association tended to have a growth reducing effect on *A. artemisiifolia* in presence of *D. carota*, we conducted a pairwise competition experiment with these two plant species: hereafter referred to as pairwise competition experiment.

**Target–challenger experiment**

The experiment had a fully 2 x 4 x 2 factorial design and was replicated seven times. It was comprised of two relative density arrangements of *A. artemisiifolia* (target and challenger), four neighboring plant species (*C. canadensis*, *D. carota*, *A. vulgaris*, *T. vulgare*), and two soil treatments (natural AM fungal community and non-mycorrhizal control). All arrangements were interspecific and had five plants per pot: one target plant in the centre surrounded by four challenger plants belonging to the same plant species (Figure II.1). Thus, the two different density levels of *A. artemisiifolia* were realized by either having *A. artemisiifolia* as target in a ratio of 1:4 (Figure II.1a) or as challenger in a ratio of 4:1 (Figure II.1b) in comparison to each of four selected neighboring plant species of the new
range. All neighboring plant species tested are mycorrhizal plants, but have some differences in life span and growth form: *C. canadensis* is annual and forms at first a leaf rosette in spring. *D. carota* is an upright biennial, and *A. vulgaris* and *T. vulgare* are upright perennials. We found these plant species co-occurring with the annual, upright growing *A. artemisiifolia*, which is in accordance with other observations for *A. artemisiifolia* in Middle Europe (Brandes and Nitzsche 2007).

In October 2007, we collected seeds of *A. artemisiifolia* and the four co-occurring plant species from plants from a ruderal site in Berlin, Germany (52°28’30.00”N, 13°21’46.45”E). The seeds were stored at room temperature or minus 20 °C (*A. artemisiifolia*). In February 2008, seeds of all plants species were pre-germinated in sterile playground sand in a greenhouse (day/night temperature 22/18 °C; ambient light conditions). Within one week all plant species started to germinate. Two weeks later we used the seedlings in the experiment.

The soil used in the experiment came from a field research station of the Humboldt Universität Berlin close to Berlin: Thyrow (52°15’N, 13°14’E). The Thyrow soil is characterized as sandy to silty sand soil (Ellmer et al. 2000), and has a low fertility (organic carbon = 0.52 %; total nitrogen = 0.04 %), as well as an acidic pH (soil pH = 5.2). We used that sandy soil, because competition has been shown to be greater for soil nutrients than for light under such nutrient poor soil conditions (Rebele 2000); hence, by using a low-fertility soil we expected an increased importance of AM fungi. In March 2008, we randomly took soil samples from the top 15 cm of soil. These soil samples were pooled and sieved through a 4 mm sieve. Except for 20 L of soil, which were used for mycorrhizal inoculum production, soil was pasteurized by steaming (Sterilo 7K, Harter Elektrotechnik, Schenkenzell, Germany) at 90 °C overnight. For mycorrhizal inoculum, non-steamed soil was thoroughly mixed with water and the supernatant was wet sieved through 500, 212, 53, 38, 20 µm sieves. Filtrate prepared from 53–38 µm soil fraction was used as mycorrhizal inoculum, and filtrate that passed through the 20 µm sieve was collected as microbial wash.

To set up the experiment, seedlings were planted in target and challenger arrangements in 3 L plastic pots filled with 3.5 kg of soil. To ensure that distances between plants were identical, we marked planting positions with a circle template in all 112 pots. After planting, pots intended for mycorrhizal treatment received 50 ml of the extracted mycorrhizal inoculum filtrate (inocula soil:experiment soil ratio 1:10), while an equivalent amount of water was added to the non-mycorrhizal controls. To correct for differences in
microbial background communities, all pots received 50 ml of the extracted microbial wash filtrate. Afterwards, pots were covered with a final 125 ml of steamed soil.

Plants grew in a fully randomized arrangement for seven weeks in a greenhouse (photoperiod 16 h; day/night temperature 22/18 °C), and were watered as needed with the same amount of water. At harvest, we found pots heavily penetrated by roots in all arrangements. Because it was impossible to separate roots from each other, we focused on shoot biomass of plant species, as well as male inflorescences and female flowers produced by *A. artemisiifolia*. Shoots were cut away from roots and number of male/female flowers was counted. Shoot weight was determined after drying for five days at 40 °C.

Pairwise competition experiment

Performance of *A. artemisiifolia* and *D. carota* was studied in situations of intra- and interspecific competition in a fully 3 x 2 factorial design. The experiment consisted of three competitive arrangements (intraspecific competition of *A. artemisiifolia*, hereafter AA; intraspecific competition of *D. carota*, hereafter DD; interspecific competition between *A. artemisiifolia* and *D. carota*, hereafter AD) and two AM fungal community treatments (presence, absence). For each treatment combination we set up 12 replicates.

In the experiment, the ecological context was strictly maintained (Hoeksema et al. 2010): plant seeds, soil and AM fungal community came from a same roadside site in Lower Lusatia, Germany (51°44′02.00″N, 14°27′27.10″E). The soil of the habitat was defined as sandy and low fertile (total carbon = 0.56 ± 0.15 %; total nitrogen = 0.03 ± 0.01 %; soil pH = 5.5 ± 0.1) (Bäucker and Rillig, unpublished manuscript). In October 2008, we collected seeds from 12 randomly chosen plants. In April 2009, ten soil samples (ca. 10 L each) were taken from the top 12 cm of soil. Soil samples were mixed, sieved (4 mm), and pasteurized by steaming for four hours at 90 °C (Sterilo 1K, Harter Elektotechnik, Schenkenzell, Germany). A portion of soil (15 L) was kept non-steamed and used for production of mycorrhizal inoculum and microbial wash filtrate (wet sieving method as described above). After extraction of AM fungal spores, the mycorrhizal inoculum was cleaned by using the sucrose centrifugation method (Johnson et al. 1999).

The experiment was set up in a greenhouse (photoperiod 16 h; day/night temperature 25/18 °C). To this, plastic pots (1 L) were filled with 700 ml of steamed soil. Corresponding to the three competitive situations (AA, DD, AD), we planted two seedlings
(two-week-old; germinated in sterile playground sand) per pot. Planting positions in the pots were marked with a circle template so that distances between seedlings were identical in all experimental units. After planting, 10 ml of mycorrhizal inoculum was added to roots in pots intended for mycorrhizal treatment (non-mycorrhizal controls received 10 ml water). Afterwards, all pots received 60 ml of the extracted microbial wash filtrate, and a final 110 ml of steamed soil. Plants grew in a fully randomized arrangement for six weeks, and were watered as needed (every day or every other day). At harvest, we removed shoot biomass, and carefully washed the root ball and separated roots of the two plants from each other. We determined shoot and root biomass per individuum after drying plant material for six days at 40 °C.

Mycorrhizal colonization in the roots

From both experiments we stained a representative root subsample from the total root biomass. As staining method we used the ink-vinegar procedure (Vierheilig et al. 1998). For this, roots were cut in pieces (1–2 cm) and treated in 10 % KOH for 35 min (water bath 90 °C). After decantation of KOH, roots were thoroughly rinsed with demineralized water, and cooked for another 15 min in ink-vinegar solution (1:1:8 Schaeffer ink, 10 % acetic acid and water). Afterwards, roots were de-stained in demineralized water and stored in lactoglycerol (1:1:1 lactic acid, glycerol and water). Presence of AM hyphae, arbuscules, and vesicles, as well as other root fungal structures were assessed by using the grid-line intersect method (McGonigle et al. 1990) based on 100 intersections per root sample examined at 200X (compound microscope, Leica Microsystems CMS GmbH, Wetzlar, Germany).

Statistical analyses

For the target–challenger experiment we performed standard three-way ANOVAs to test for significance of main effects (relative density arrangement of *A. artemisiifolia*, neighboring plant species, soil treatment) and their interactions. Data were transformed as necessary to meet assumptions of normality and homogeneity of variances, as indicated by Shapiro-Wilk test (on residuals) and Bartlett test, respectively. We used log-transformation for data of vegetative shoot biomass of *A. artemisiifolia* and BoxCox-transformation (exponent 0.4242424) for biomass data of neighboring plant species. Data of number of
male inflorescences and female flowers of *A. artemisiifolia* were square-root transformed. To make comparisons between plant species when grown as target (one plant in the centre per pot) and challenger (four plants belonging to the same species in a circle per pot), we used the average value of the four challenger plants per pot.

For the pairwise competition experiment of *A. artemisiifolia* and *D. carota* we performed two-way ANOVAs testing for significance of the main effects (competitive situation and soil treatment) and their interaction. The analyses on biomass and mycorrhization of the plant species were computed with two levels for competitive situation (*A. artemisiifolia* with level AA, AD; *D. carota* with level DD, AD). We based our analyses on the mean of the two plants in monoculture and the measure of the respective plant species in the mixed situation. To meet assumptions of normality (Shapiro-Wilk test) and homogeneity of variances (Bartlett test), we log-transformed data of shoot and root biomass of *A. artemisiifolia* and *D. carota*. To test for differences in plant performance depending on presence/absence of mycorrhiza in single situations of intra- and interspecific competition of *A. artemisiifolia* and *D. carota*, we used paired *t*-test’s for independent samples. Again, data were analyzed after log-transformation, except for shoot and root biomass of *A. artemisiifolia* in mixture.

Differences between treatment groups were always compared with Tukey HSD post-hoc comparison tests (*P* < 0.05). Computations were performed using R version 2.10.1 (R Development Core Team 2009).

**Results**

Target–challenger experiment

We found that the target or challenger arrangement of *A. artemisiifolia* had a substantial influence on all response variables measured (main effect of relative density always *P* < 0.001; Table II.1). When neighboring plant species were planted in the center surrounded by *A. artemisiifolia*, their pooled shoot biomass was on average smaller (mean ± SE; 0.185 ± 0.017 g) than if they grew in a circle (pooled shoot biomass of the mean of all neighbors per pot: 0.524 ± 0.040 g). In contrast, *A. artemisiifolia* produced on average considerably more vegetative shoot biomass per plant as target (3.184 ± 0.181 g) compared to challenger (1.158 ± 0.021 g) (Figure II.3a); hence, both *A. artemisiifolia* and
neighboring plant species had on average greater shoot biomass per plant in target-situations of *A. artemisiifolia*. Further, the number of male inflorescences produced by *A. artemisiifolia* was on average almost tripled when it grew as target (7.8 ± 0.9) compared to challenger (2.7 ± 0.2) (Figure II.3b). Conversely, the number of female flowers produced by *A. artemisiifolia* per plant was on average lower in target (0.3 ± 0.1) compared to challenger arrangements (1.7 ± 0.2).

Furthermore, identity of the neighboring plant species also strongly influenced shoot biomass produced by both *A. artemisiifolia* and the four neighboring plant species (significant main effect of neighboring plant species; Table II.1). Moreover, biomass of *A. artemisiifolia* and neighboring plant species depended on if neighboring plants where tested in target or challenger arrangements of *A. artemisiifolia* (significant two-way interaction term between relative density and neighboring plant species; Table II.1). Effects described hereafter are significant, unless otherwise stated. We found that *C. canadensis* performed poorly in all arrangements compared to the other neighboring plant species tested (Figure II.2). Moreover, when *A. artemisiifolia* was grown as challenger, *D. carota*, *A. vulgaris*, and *T. vulgare* showed no significant differences in aboveground biomass. In target arrangements of *A. artemisiifolia*, however, *A. vulgaris* had greater shoot biomass than *D. carota* (arrangement target *A. vulgaris–D. carota*: *P* = 0.009; after Tukey’s HSD pairwise comparison test) (Figure II.2). Overall, therefore, *C. canadensis* produced the smallest shoot biomass compared to all other neighboring species, and *D. carota* had less biomass compared *A. vulgaris* and *T. vulgare*, respectively (Figure 2; Supplemental Table A.II.2). Regarding *A. artemisiifolia*, we found that its shoot growth as target and overall was greater when grown together with *C. canadensis* compared to all other combinations of target/challenger arrangements and neighboring plant species tested (*P* < 0.05; Tukey HSD) (for data see Supplemental Tables A.II.2 and A.II.3; Figure II.3a).

Further, a significant three-way interaction term indicated that responses of shoot biomass and maleness of *A. artemisiifolia* when grown as target or challenger were influenced by neighboring plant species and mycorrhizal treatment (Table II.1). In presence of AM fungi, we found that shoot biomass of target *A. artemisiifolia* tended to be greater only when grown in combination to *C. canadensis* and *T. vulgare*, respectively (Figure II.3a). Conversely, when *A. artemisiifolia* was target to *D. carota*, shoot biomass tended to be decreased in presence of AM fungi compared to non-mycorrhizal control (Figure II.3a). For combination with *A. vulgaris* as the neighboring plant, we found that
shoot growth of target *A. artemisiifolia* was uninfluenced by presence/absence of AM fungi. Furthermore, the different AM fungal effects on *A. artemisiifolia* as a function of competing neighbor species were also found overall, as indicated by a significant interaction term between neighboring plant species and soil treatment: again, *A. artemisiifolia* tended to either increase in shoot biomass (with *C. canadensis* or *T. vulgare*), decrease (with *D. carota*) or was unaffected (with *A. vulgaris*) in the presence of mycorrhiza (Table II.1; Supplemental Table A.II.4). But, all shoot biomass effects mediated by presence/absence of AM fungi could not be shown with pairwise comparison tests (Tukey HSD; *P* > 0.05).

Considering number of male inflorescences of *A. artemisiifolia*, however, we found that more male flowers were produced when *A. artemisiifolia* grew as target surrounded by *C. canadensis* in presence of AM fungi compared to absence (arrangement target to *C. canadensis* mycorrhizal–non-mycorrhizal: *P* = 0.048; after Tukey’s HSD pairwise comparison test) (Figure II.3b). Overall, maleness of *A. artemisiifolia* also indicated a significant main effect of neighboring plant species, but this effect could not be shown with pairwise comparison tests (Tukey HSD; *P* > 0.05) (Supplemental Table A.II.2).

In terms of mycorrhization, plant roots in the pots treated with AM fungal inoculum were colonized by AM fungi: percentage colonization by AM hyphae 38.3 ± 5.7 %, arbuscules 12.7 ± 4.2 %, and vesicles 6.9 ± 1.0 %. Infection with non-AM fungi was very low (0.2 ± 0.2 %) in the mycorrhizal treatment. Non-mycorrhizal controls were also rarely infected by fungi (1.0 ± 0.3 %), none of which could be classified as AM fungi.

Pairwise competition experiment

We found that the roadside AM fungal community had divergent effects on performance of *A. artemisiifolia* and *D. carota* grown in pairwise competitive situations (significant main effect of mycorrhizal treatment; Table II.2). In the presence of AM fungi, *D. carota* produced always significantly more shoot and root biomass (Figure II.4a; Table II.3). Conversely, *A. artemisiifolia* had significantly greater shoot biomass without AM fungi; root biomass showed similar, but non-significant results (Figure II.4a; Table II.3). Furthermore, shoot biomass of both plant species was also divergently influenced by intraspecific and interspecific competition, respectively (significant main effect of competitive situation; Table II.2). While *D. carota* had greater biomass when grown with a
A. artemisiifolia produced significantly more biomass in the mixed situation (Figure II.4b).

Considering the influence of AM fungi on shoot and root biomass of A. artemisiifolia and D. carota in the different competitive situations, we found partially similar results as already indicated by the target–challenger experiment. When A. artemisiifolia grew with a conspecific, we found that its shoot and root biomass was significantly decreased with AM fungi compared to without (Table II.3). In mixture with D. carota and presence of AM fungi, A. artemisiifolia had also less biomass compared to the non-mycorrhizal control, although not significantly (Table II.3). A similar pattern, i.e., reduced shoot growth of A. artemisiifolia under condition of AM fungi and D. carota as competitor, was already found when A. artemisiifolia was target to D. carota. Regarding D. carota in mixture, shoot and root biomass was significantly increased in the presence of AM fungi compared to their absence (Table II.3). Such a positive mycorrhizal effect on D. carota was also already indicated by the target-challenger experiment, but results were non-significant (target arrangement with neighbor D. carota; shoot biomass with AM fungi: 0.546 ± 0.057 g; non-mycorrhizal control: 0.400 ± 0.042 g; Supplemental Table A.II.1). Furthermore, shoot and root biomass of D. carota in competition with a conspecific was also significantly increased in symbiosis with AM fungi than without (Table II.3).

Concerning mycorrhization, roots of both plant species were colonized by AM fungi when soil was treated with AM fungal inoculum. Overall, AM fungal colonization tended to be higher in roots of D. carota (mean ± SE; hyphae 44.7 ± 7.3 %; arbuscules 27.0 ± 5.1 %) than A. artemisiifolia (hyphae 36.2 ± 7.7 %; arbuscules 20.9 ± 5.1 %), but this difference was non-significant. Furthermore, percentage colonization by AM fungal hyphae, arbuscules and vesicles of both plant species showed no significant difference depending on competitive situation, although A. artemisiifolia tended to form more AM fungal structures in the roots when grown with D. carota (Supplemental Table A.II.5). Non-mycorrhizal controls were broadly similarly infected with non-AM fungi (0.3 ± 0.1 %) as plants growing in AM fungal treatment (0.2 ± 0.1 %).
Discussion

In contrast to our expectations, we found that the AM fungal symbiosis with natural AM fungal communities was of minor importance for the competitive ability of *A. artemisiifolia* in target and challenger arrangements with co-existing ruderal mycorrhizal competitors. Regardless of presence/absence of AM fungi, *A. artemisiifolia* grew considerably taller compared to neighboring plant species in all performed situations of interspecific competition; hence, *A. artemisiifolia* was highly dominant under the low fertile soil conditions tested here. This result is supported by findings of Miller (1994), who also reported *A. artemisiifolia* as an exceptionally good competitor that had strong direct suppressive effects on four co-existing plant species in North America. However, Leskovšek et al. (2012) showed that *A. artemisiifolia* was a poor competitor in competition with *Lolium multiflorum* L. under high resource availability (high levels of nitrogen and water). However, the same study also demonstrated that growth of *A. artemisiifolia* in competition was minimally affected by shortage of nutrients. Therefore, one explanation for the competitive dominance of *A. artemisiifolia* in our study may be its ability to compete for nitrogen in nutrient poor soils, such as those used by us. As reported by Tilman (1986), *A. artemisiifolia* had the greatest biomass as seedling when grown under low nitrogen levels in comparison to other eight co-occurring plant species.

The target–challenger experiment, further, showed that biomass of all species strongly differed in target compared to challenger arrangements of *A. artemisiifolia*. In accordance to our third hypothesis, growth of the neighboring plants was more decreased when they were surrounded by *A. artemisiifolia* (high relative density of *A. artemisiifolia*). Moreover, identity of neighbor species had a significant impact on shoot biomass of *A. artemisiifolia* and the neighboring species grown in target or challenger situations, which was contrary to our assumption and other observations. Miller (1994) found no evidence that growth of *A. artemisiifolia* was differently affected by presence of any other species. However, for other invasive plants such as *Centaurea melitensis* or *Centaurea stoebe* (formerly *C. maculosa*) it has been shown that plant neighbor identity matters (Callaway et al. 2003; 2004).

Considering the neighbors in our study, the annual *C. canadensis* performed poorly in all arrangements. The other species, i.e., *D. carota*, *A. vulgaris* or *T. vulgare*, grew equally poor in challenger arrangements, but performed differently when *A. artemisiifolia* was the target. In target situations of *A. artemisiifolia*, the upright perennials *A. vulgaris*
and *T. vulgare* had on average the greatest shoot biomass of the neighboring species selected, while the biennial *D. carota* grew less compared to *A. vulgaris*. Conversely, for target *A. artemisiifolia* we found that its shoot growth differed in presence of the most inferior competitor *C. canadensis* only. Here, target *A. artemisiifolia* could profit most, and had almost doubled its shoot biomass compared to arrangements with *D. carota*, *A. vulgaris* and *T. vulgare*. However, the substantial competitive advantage of *A. artemisiifolia* over *C. canadensis* may have been related to their different growth forms. During the whole period of the experiment, *C. canadensis* was in the leaf rosette-forming stage and, therefore, additionally shaded by leaves of the upright growing *A. artemisiifolia*.

In challenger arrangements, i.e., when *A. artemisiifolia* grew also with conspecifics, its shoot growth was strongly decreased compared to target arrangements irrespective of the neighboring plant and presence/absence of mycorrhiza, respectively. Since shoot performance of neighbors was also strongly suppressed when surrounded by *A. artemisiifolia*, we interpret our findings of reduced growth of challenger *A. artemisiifolia* related to a strong intraspecific competition within *A. artemisiifolia*. As shown by other studies, performance of *A. artemisiifolia* is reduced with increasing density of conspecifics (Miller and Werner 1987; MacDonald and Kotanen 2010). In our experiment, therefore, it seems that challenger plants of *A. artemisiifolia* were strongly competing with each other, which may have overridden other effects, such as the impact of mycorrhizal fungi.

Considering the mycorrhizal impact on *A. artemisiifolia* as target, we found that the AM fungal effect pointed towards different directions depending on the neighboring plant species, although not significantly. AM fungi tended to have a positive effect on target *A. artemisiifolia* when competing with *C. canadensis* or *T. vulgare*. In competition with *A. vulgaris*, however, the mycorrhizal symbiosis mediated a neutral effect on *A. artemisiifolia*. When *A. artemisiifolia* was surrounded by *D. carota*, shoot growth of *A. artemisiifolia* was reduced with AM fungi compared to without; here, the mycorrhizal symbiosis amplified effects of interspecific competition.

This growth reducing effect of AM fungi on *A. artemisiifolia* in the presence of *D. carota* was also indicated by the pairwise competition experiment solely focusing on *A. artemisiifolia* and *D. carota* in intra- and interspecific competitive situations. Again, we found that *A. artemisiifolia* tended to produce less biomass when growing in mixture with *D. carota* under mycorrhizal compared to non-mycorrhizal conditions. However, when *A. artemisiifolia* was tested in intraspecific competition we even found a stronger negative
effect of AM fungi on competing conspecifics; hence, the mycorrhizal symbiosis clearly amplified competition within *A. artemisiifolia*, which is in line with recent findings of a meta-analysis by Moora and Zobel (2010). They showed that AM fungi have an amplifying or neutral effect in intraspecific competition, while the effect is balancing in interspecific competition. Since we maintained the ecological context of soil, AM fungal community and plant species origin in our study, the AM fungal effect on *A. artemisiifolia* under natural conditions must be assumed to be negative in intraspecific competition. The effect in interspecific competition with *D. carota* might be rather neutral (because biomass was not significantly reduced). The findings of our pairwise competition experiment are contrary to a study by Shah et al. (2008) investigating the invasive plants *Anthemis cotula* in India. Similarly, they studied the effect of AM fungi on *A. cotula* and *D. carota* in intra- and interspecific competition. Here, invasive *A. cotula* was enhanced by the presence of local AM fungi when grown together with *D. carota*. Further, they found that *A. cotula* was even more promoted by AM fungi when grown in monoculture, which is the exact opposite of what we found for *A. artemisiifolia*. Shah et al. (2008) also showed that the degree of AM fungal root colonization of *A. cotula* was decreased in mixture with *D. carota* compared to monoculture. Our data, however, indicate that roots of *A. artemisiifolia* were more strongly colonized by AM fungi when grown in mixture with *D. carota*. In our experiment, furthermore, *D. carota* strongly profited from the mycorrhizal symbiosis both in intra- and interspecific competition; thus, the AM fungal symbiosis had a balancing effect on intraspecific competition of *D. carota*, which is in contrast to *A. artemisiifolia* and other plant species (Moora and Zobel 2010). Moreover, the competitive ability of *D. carota* was increased in interspecific competition with *A. artemisiifolia* in the presence of the roadside AM fungal assemblage tested here. Aside from AM fungi, *A. artemisiifolia* was most strongly affected by conspecific competition in both experiments. Conversely, *D. carota* performed better in intraspecific competition.

Considering reproductive traits of *A. artemisiifolia* in the target–challenger experiment, we found that the number of male inflorescences was significantly higher when it grew as target to the most inferior competitor *C. canadensis* under mycorrhizal compared to non-mycorrhizal conditions. Interestingly, increased production of male flowers coincided with greatest shoot biomass produced by *A. artemisiifolia*. Such positive effects of mycorrhizal symbiosis on shoot biomass and maleness were also found by Koide and Li (1991). They, however, studied *A. artemisiifolia* growing alone with the mycorrhizal fungus *Glomus etunicatum*. Therefore, our result is the first evidence that
inoculation with a natural AM fungal assemblage can also increase male gender function in *A. artemisiifolia* under certain circumstances of interspecific competition, as shown here with *C. canadensis*. In addition, *A. artemisiifolia* also showed protandry in specific combinations of interspecific competition: as target to *C. canadensis* (mycorrhizal and non-mycorrhizal treatment) and *T. vulgare* (only mycorrhizal). The phenomenon that sex allocation in *A. artemisiifolia* is adjusted to size and environmental conditions has been recently reported by Friedman and Barrett (2011). They showed that flowering can range from protandry in the sun to protogyny in the shade. Ackerly and Jasieński (1990), moreover, demonstrated that the variability in aboveground biomass and gender is higher under nutrient-rich soil conditions leading to a shift towards maleness in taller plants. Similarly, other studies also found that favorable conditions increase maleness in *A. artemisiifolia* (McKone and Tonkyn 1986; Traveset 1992; Bäucker and Rillig, unpublished manuscript). Therefore, our data once more demonstrate that greater shoot biomass in *A. artemisiifolia* results in higher number of male inflorescences.

Regarding female function of *A. artemisiifolia*, we found that number of female flowers was increased more than five-fold when *A. artemisiifolia* had a high relative density (grown as challenger) compared to low (target). Thus, our data suggest that reduced shoot growth and lower production of male flowers in competition with conspecifics led to earlier development of female flowers in *A. artemisiifolia*. A study by Lundholm and Aarssen (1994) also reported increased female function of *A. artemisiifolia* in the presence of neighboring plants. Besides, a shift towards greater femaleness under increased population density was also shown for the congeneric *Ambrosia trifida* (Abul-Fatih et al. 1979). However, an earlier flowering of female flowers in *A. artemisiifolia* does not necessarily imply greater seed output. There are some studies showing that plants, which produce less vegetative biomass also have lower seed mass (Bäucker and Rillig, present thesis, Chapter 2; Leskovšek et al. 2012). Moreover, Chikoye et al. (1995) found that increased population density of *A. artemisiifolia* correlates with decreased number of seeds per plant produced in the field.

To conclude, *A. artemisiifolia* was demonstrated to be an exceptionally good competitor at low and high relative density in comparison to co-occurring mycorrhizal plant species in the new European range under nutrient poor soil conditions. Moreover, *A. artemisiifolia* experienced strong competition by conspecifics, which caused decreases in shoot biomass and maleness, but earlier flowering of female flowers. Further, we found no evidence that growth performance or competitive ability of *A. artemisiifolia* was
enhanced in the presence of natural AM fungal communities under conditions of a maintained ecological context. In fact, our findings show that AM fungi had an amplifying effect on *A. artemisiifolia* in pairwise intraspecific competition, and a neutral effect in mixture with *D. carota*. Therefore, *A. artemisiifolia*, a successful pioneer plant and a species with a strongly ruderal life history, has a weak dependence on symbiosis with AM fungi. Therefore, the invasive success of *A. artemisiifolia* in Central Europe may not be related to facilitation by natural AM fungal communities, as previously proposed by Fumanal et al. (2006). However, since we studied the competitive ability of *A. artemisiifolia* and co-existing plant species at the seedlings stage, the outcome of interspecific competition may change as the species grow longer at a site under mycorrhizal conditions, which needs further research.

Acknowledgements

We thank Michael Baumecker for providing soil from Thyrow. Further, we thank Sabine Artelt and Sabine Buchert for help during the set up of the target–challenger experiment, and Ricarda Krüger, Susanna Duda, Stefan Zimmermann, as well as Jenny Schmidt for help during the set up and harvest of the pairwise competition experiment.
### Tables and figures

**Table II.1** Analyses of variance (ANOVA) on response variables of *Ambrosia artemisiifolia* and neighboring plant species in the target–challenger experiment with the factors relative density of *A. artemisiifolia* (dens), neighboring plant species (spec) and soil treatment (treat). Values in bold indicate significance at $P < 0.05$. Overview on analyzed data is given in Supplemental Table A.II.1.

<table>
<thead>
<tr>
<th>Response variable</th>
<th>Shoot biomass</th>
<th>Shoot biomass</th>
<th>Male flowers of</th>
<th>Female flowers of</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>A. artemisiifolia</td>
<td>neighboring species</td>
<td>A. artemisiifolia</td>
<td>A. artemisiifolia</td>
</tr>
<tr>
<td>Factors</td>
<td>d.f.</td>
<td>F</td>
<td>P</td>
<td>F</td>
</tr>
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<td>Dens</td>
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<td>490.10</td>
<td>&lt;0.001</td>
<td>152.51</td>
</tr>
<tr>
<td>Spec</td>
<td>3</td>
<td>15.76</td>
<td>&lt;0.001</td>
<td>43.26</td>
</tr>
<tr>
<td>Treat</td>
<td>1</td>
<td>0.61</td>
<td>0.600</td>
<td>1.58</td>
</tr>
<tr>
<td>Dens x spec</td>
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<td>16.53</td>
<td>&lt;0.001</td>
<td>6.16</td>
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<tr>
<td>Dens x treat</td>
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<td>0.359</td>
<td>0.56</td>
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<tr>
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<td>4.55</td>
<td>0.005</td>
<td>0.54</td>
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<tr>
<td>Dens x spec x treat</td>
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<td>3.37</td>
<td>0.022</td>
<td>1.61</td>
</tr>
<tr>
<td>Residuals</td>
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<td>95</td>
<td>95</td>
<td>95</td>
</tr>
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</table>
Table II.2 Analyses of variance (ANOVA) on biomass traits of *Ambrosia artemisiifolia* (A) and *Daucus carota* (D) in the pairwise competition experiment in intraspecific (AA, DD) and interspecific competition (AD). Values in bold indicate significance at $P < 0.05$; analyzed data are shown Table II.3.

<table>
<thead>
<tr>
<th>Factors</th>
<th>d.f.</th>
<th>$F$</th>
<th>$P$</th>
<th>$F$</th>
<th>$P$</th>
</tr>
</thead>
<tbody>
<tr>
<td>Shoot biomass</td>
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<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Comp</td>
<td>1</td>
<td>5.64</td>
<td>0.022</td>
<td>4.10</td>
<td>0.049</td>
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<tr>
<td>Treat</td>
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<td>5.26</td>
<td>0.027</td>
<td>49.08</td>
<td>&lt;0.001</td>
</tr>
<tr>
<td>Comp x treat</td>
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<td>0.04</td>
<td>0.851</td>
<td>0.30</td>
<td>0.584</td>
</tr>
<tr>
<td>Root biomass</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Comp</td>
<td>1</td>
<td>1.09</td>
<td>0.302</td>
<td>1.58</td>
<td>0.216</td>
</tr>
<tr>
<td>Treat</td>
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<td>1.33</td>
<td>0.255</td>
<td>24.42</td>
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</tr>
<tr>
<td>Comp x treat</td>
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<td>0.303</td>
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<td>0.456</td>
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<tr>
<td>Residuals</td>
<td>44</td>
<td>44</td>
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<td></td>
<td></td>
</tr>
</tbody>
</table>

comp, competitive situation; treat, soil treatment
Table II.3 Results of paired *t*-test on data of shoot and root biomass of *Ambrosia artemisiifolia* (A) and *Daucus carota* (D) in intraspecific and interspecific competition in presence or absence of AM fungi. In intraspecific competition, data (mean ± SE) represent the mean of two the two conspecifics growing together. *P*-values in bold indicate significance at *P* < 0.05.

<table>
<thead>
<tr>
<th>Response variable</th>
<th>Soil treatment</th>
<th></th>
<th>t-test</th>
<th></th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>AM fungi</td>
<td>non-mycorr.</td>
<td>d.f.</td>
<td><em>P</em>-value</td>
</tr>
<tr>
<td><strong>intraspecific competition (monoculture)</strong></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Shoot biomass A (g)</td>
<td>108.4 ± 9.5</td>
<td>133.9 ± 6.0</td>
<td>11</td>
<td>0.040</td>
</tr>
<tr>
<td>Root biomass A (g)</td>
<td>39.8 ± 7.1</td>
<td>53.2 ± 5.2</td>
<td>11</td>
<td>0.015</td>
</tr>
<tr>
<td>Shoot biomass D (g)</td>
<td>38.1 ± 5.4</td>
<td>8.1 ± 1.2</td>
<td>11</td>
<td>&lt;0.001</td>
</tr>
<tr>
<td>Root biomass D (g)</td>
<td>14.5 ± 2.8</td>
<td>4.5 ± 0.4</td>
<td>11</td>
<td>0.003</td>
</tr>
<tr>
<td><strong>interspecific competition (mixture)</strong></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Shoot biomass A (g)</td>
<td>140.8 ± 13.1</td>
<td>179.8 ± 19.4</td>
<td>11</td>
<td>0.191</td>
</tr>
<tr>
<td>Root biomass A (g)</td>
<td>36.6 ± 5.2</td>
<td>49.2 ± 10.8</td>
<td>11</td>
<td>0.394</td>
</tr>
<tr>
<td>Shoot biomass D (g)</td>
<td>26.5 ± 5.2</td>
<td>5.9 ± 0.8</td>
<td>11</td>
<td>0.002</td>
</tr>
<tr>
<td>Root biomass D (g)</td>
<td>10.5 ± 1.9</td>
<td>4.1 ± 0.3</td>
<td>11</td>
<td>0.010</td>
</tr>
</tbody>
</table>
**Figure II.1** Target versus challenger arrangement of *Ambrosia artemisiifolia* (A) in the target–challenger experiment: (a) shows *A. artemisiifolia* growing as target (ratio: 1/5), and (b) as challenger (ratio: 4/5) to the neighboring plant species (X).

**Figure II.2** Shoot biomass of the different neighboring plant species in response to target vs. challenger arrangements of *Ambrosia artemisiifolia* in the experiment (black = *Conyza canadensis*, light grey = *Daucus carota*; dark grey = *Artemisia vulgaris*, white = *Tanacetum vulgare*). Bars show means ± SE, and different lower case letters indicate significant differences (*P* < 0.05) among treatment groups according to Tukey’s HSD test.
Figure II.3 Average shoot biomass (a) and number of male inflorescences (b) of *Ambrosia artemisiifolia* in response to target vs. challenger arrangement, soil treatment (mycorrhizal vs. non-mycorrhizal) and different neighboring plant species (black = *Conyza canadensis*, light grey = *Daucus carota*; dark grey = *Artemisia vulgaris*, white = *Tanacetum vulgare*). Bar plots represent means ± SE. Different lower case letters indicate significant differences ($P < 0.05$) among treatment groups according to Tukey’s HSD test.
Figure II.4 Shoot biomass of *A. artemisiifolia* (black) and *D. carota* (grey) in response to the main effects soil treatment (a) and competitive situation (b). In (b) data for intraspecific arrangements were calculated from averaged values of both plants per pot. Bars represent means ± SE.
References


CHAPTER 4

Distinct seed morphs of Galinsoga parviflora (Asteraceae) give rise to different soil feedbacks

Abstract

Heterocarpy is the phenomenon that a single plant produces two or more distinct fruit types, which differ in dispersal mechanisms and ecological behavior. Galinsoga parviflora represents a heterocarpic plant that produces capitula with two seed morphs: seeds equipped with a pappus for long-distance dispersal and non-pappus seeds having a low dispersal potential. In laboratory conditions, we studied if the different seed types differ in soil feedback. Feedback is known to affect plant performance often negatively in ‘self-cultivated’ (trained) soil due to accumulation of disadvantageous soil organisms. We trained soil over two plant generations with plants arising from both seed types, and tested feedback responses of non-pappus and pappus progeny as ‘trained versus sterile’ soil contrasts. We found that plants from pappus seeds, which are produced for colonizing new sites, suffered greater negative feedback than plants from non-pappus seeds, which maintain the existing population. The feedback differences were most pronounced for reproductive traits, but also indicated by root biomass. Moreover, non-pappus progeny produced a higher portion of pappus seeds, i.e. showed proportionally greater investment in dispersal in disadvantageous conditions of trained soil compared to pappus progeny. The expected difference in fungal root colonization (greater infection with pathogenic fungi for pappus progeny) was not observed; hence, differences in magnitudes of negative feedback must have been related to other soil organisms than fungi. Interestingly, pappus progeny produced significantly more root biomass when they came from non-pappus mother/grandmother plants. Such maternal history effects need further evaluation and might be more common in heterocarpic plants.
Introduction

Many plant species have rather constant seed size (Harper et al. 1970) while others show intra-individual variation in seed form or behavior, termed seed heteromorphism (Venable 1985a). Such heteromorphism becomes most evident when two or more distinctly different fruit types with divergent ecologically functions are produced by one plant. This phenomenon was described as heterocarpy (Mandák 1997) and is often associated with differential dispersal mechanisms and differential germination responses of the respective fruit types (Tanowitz et al. 1987). Beside fruit, also the terms diaspora (i.e. dispersal unit) or seed in the broadest sense are used in the literature: in this study we refer to seed *sensu lato*.

Within the angiosperms, heterocarpy/heterospermy is frequently known in Asteraceae and Chenopodiaceae (Imbert 2002). Predominantly, heterocarpy occurs among annuals, pioneer species or plants faced with stochastic environments such as deserts or semi-deserts (Mandák 1997). It represents an adaptive trait evolving in situations of environmental unpredictability (Cruz-Mazo et al. 2009) and where adaptive tracking or the evolution of plasticity are partially impeded (Simons 2011). The most noticeable feature of heterocarpic plants is the diversification of offspring in terms of space or time, such as non-dispersive vs. dispersive or dormant vs. non-dormant seed types (Venable et al. 1995). In this context, dormancy has been found to be negatively correlated with dispersal (Venable and Brown 1988; Venable 1989): higher dormancy results in reduced selection for dispersal and vice versa (Venable et al. 1995).

The advantage of producing different kind of seeds per individuum under environmental unpredictability has often been considered from the perspective of bet-hedging, i.e. trading-off some potential short-term benefit for a long-term benefit (e.g. Venable 1985a; Philippi and Seger 1989; Venable 2007). Such a strategy ensures that at least some offspring are successful in a variety of environmental circumstances (Venable et al. 1995). As a consequence, bet-hedging traits do not maximize the expected fitness within a generation, but do maximize geometric-mean fitness across generations, which results in higher long-term success of bet hedgers compared to non-bet hedgers (Simons 2011). Other explanations for the evolution of heterocarpy are escape from negative density effects or sib competition (Venable and Brown 1988).

A whole body of literature exists on differences of seed morphs of heterocarpic plants, where patterns of seed size, dormancy and germination have been extensively
studied. For Asteraceae, a large number of studies indicate that achenes produced by central flowers of the capitula have lower dormancy and their germination is less restricted by specific temperature regimes (Baskin and Baskin 1976; Flint and Palmblad 1978; McEvoy 1984; Tanowitz et al. 1987; Venable and Levin 1985a; De Clavijo 2001; Brändel 2007; Sun et al. 2009; Aguado et. al 2011; but see for the opposite pattern Brändel 2004; Rai and Tripathi 1982). Moreover, the central achenes were frequently found to be lighter than the peripheral ones, although also some other patterns were found for seed mass and/or embryo weight (Rocha 1996; Brändel 2004; Venable and Levin 1985b). In a few species, however, heterocarpy is not associated with germination differences (Baker and O'Dowd 1982; Sorensen 1978; Imbert et al. 1996); hence, different seed morphology reflects different seed dispersal strategies and, possibly, other differences in ecological behavior.

Therefore, the influence of environmental conditions on heterocarpic plant traits, such as the ratio of non-dispersing to dispersing seeds produced depending on the habitat, was the focus of several studies (e.g. Ellner and Schmida 1984; Imbert and Ronce 2001; Kigel 1992; Cheptou et al. 2008). For example, Cheptou et al. (2008) demonstrated that *Crepis sancta* responds to urban habitat fragmentation with a shift towards a higher portion of seeds lacking dispersal structures, which reduces costs of dispersal. In a few other studies, the effects of density/competition and soil conditions, like nutrient or water availability, have been tested on plants grown from different seed types. For Asteraceae, some studies report about differences in competitive ability of plants originating from different seed morphs (Venable 1985b; Rai and Tripathi 1987; Imbert et al. 1997; De Clavijo and Jiménez 1998), while others found no such effects (Sorensen 1978; Baker and O'Dowd 1982; Brändel 2007). In Chenopodiaceae, plants arising from different seed types were also shown to differ in biomass (Ellison 1987), as well as resource allocation to the different seed types when grown both in mixture with each other and monoculture (Mandák and Pyšek 2005).

From the existing literature it can be seen that the phenomenon of heterocarpy is very complex. Since it was predominantly investigated from the perspective of the different seed types, our aim was to study another ecological aspect, which we hypothesize may vary between plants grown from distinct seed morphs of heterocarpic plants. We tested if plants arising from different seed types differ in their interaction with soil biota; hence, exhibit divergent responses of soil feedback. In short, negative feedback is caused by presence and accumulation of disadvantageous soil organisms, such as soil-borne...
bacteria, fungi and invertebrate fauna (Bever 1994; Klironomos 2002; Bever 2002; De Deyn et al. 2003). The negative soil effect becomes visible when species grow less well in ‘self-cultivated’ soil compared to soils trained by other plant types. Negative feedback is known to be an important local and large-scale mechanism influencing plant abundance and mediating plant coexistence (Bever 2003; Ehrenfeld et al. 2005; Kulmatiski et al. 2008; Petermann et al. 2008; Johnson et al. 2012).

To test for a relationship between dispersal ability and negative soil feedback we studied *Galinsoga parviflora* Cav. (Asteraceae). The species is known to produce different floret types within a flower head (capitulum), where a group of bisexual disc florets in the centre of the capitulum is surrounded by a ring of a few ray florets, which are female (Nielreich 1866). The different florets develop to distinctly different achene types: disc achenes bearing a pappus and ray achenes lacking a pappus as illustrated by Becker (1913). At maturity, each of the ray achenes (hereafter non-pappus seeds) remain enclosed in involucral bracts forming a winged structure (Espinosa-García and Sarukhán 1997), which might be dispersed or fall as a whole into the local habitat. Because non-pappus seeds lack visible adaptation to long-distance dispersal (Vibrans 1999), they are for short-distances dispersal. In contrast, disc achenes (hereafter pappus seeds) possess a crown of scales as appendages best-suited for long-distance dispersal predominately by human clothing (Holm et al. 1977; Vibrans 1999), but also by animal fur (Vibrans 1999) or wind (Terzioğlu and Anşin 2001).

Besides dispersal structures, the two seed types of *G. parviflora* differ in other aspects. Under low and high light regimes, non-pappus seeds germinate earlier and at higher percentages than pappus seeds (Rai and Tripathi 1982, 1987). The higher germination rate of non-pappus seeds is probably due to their greater seed weight and higher contents of reserves compared to pappus seeds (Rai and Tripathi 1982). The two seed types, moreover, differ in their dormant characters, and thus loss rates from the seed bank (Espinosa-García et al. 2003), and show differences in seedling survival, as well as competiveness (Rai and Tripathi 1987).

Based on morphological and ecological differences found for seeds of *G. parviflora* it can be stated that this species produces two functionally distinct seed types: the non-pappus seed for *in situ* persistence, and the pappus seed to reach new habitats; hence, founding of populations. Assuming that environments of newly colonized sites versus existing populations are highly dissimilar, especially with regard to soil biota composition and abundance, the different dispersal capacities in *G. parviflora* may correspond to
different soil feedback responses of pappus versus non-pappus seeds. By implication, we hypothesized that plants grown from non-pappus seeds would exhibit better performance in soil trained by the mother plant than those from pappus seeds; hence, progeny of non-pappus seeds experience less negative soil feedback compared to progeny of pappus seed.

Materials and methods

Study species

*Galinsoga parviflora* is an annual plant native to the mountainous region of Central America (Canne 1977). Several decades ago, the herb was already reported to have a worldwide distribution (Canne 1977; Holm et al. 1977), with human activity representing the most important vector (Warwick and Sweet 1983; Damalas 2008). The species occurs in disturbed habitats and agricultural areas, where it is an important weed (e.g. Holm et al. 1977; Warwick and Sweet 1983; Damalas 2008). *G. parviflora* reproduces via cross- and self-fertilization; hence, it needs one single seed only to start a new population (Warwick and Sweet 1983).

Seeds used in the experiment were collected in Warendorf Müssingen, Germany (51°58'06 N, 7°53'29 E) from plants growing on agricultural land. We ordered those seeds from the catalogue Index Seminum 2009 of the botanical garden of the Universität Münster (IPEN DE-0-MSTR-SA 8629).

Soils and soil preparation

The soils used for the experiment came from two different locations: Berlin-Dahlem (52°27' N, 13°18' E) and Thyrow (52°15' N, 13°14' E). The first site (Dahlem) is an experimental field of the Institute of Biology of Freie Universität Berlin, and the second belongs to a field research station of the Humboldt Universität Berlin. The urban Dahlem soil is classified as silty sand soil, while Thyrow soil has a higher content of sand; hence, is characterized as sandy to silty sand soil (Ellmer et al. 2000). Further, the Dahlem soil has a higher soil pH (pH = 6.1), higher contents of organic carbon (C = 1.01 %) and total nitrogen (N = 0.09 %) compared to the Thyrow soil (pH = 5.2, C = 0.52 %, N = 0.04 %) (Schweitzer 2010; Baumecker et al. 2009).
Soil samples were taken from the top 15 cm of soil and sieved through a 4 mm sieve. Half of each soil was directly used for the training phase. The other half was stored (4 °C), and later autoclaved two times at 121 °C (each time 30 min) to prepare both sterile background and control soil for the feedback experiment. Because the autoclaved soils had poor drainage, we mixed them both with sterilized playground sand (ratio 5:1). To prepare soil inoculated with trained soil inoculum, we thoroughly mixed trained soil into sterile background soil at a ratio of 1:10. In this process, the identity of replicates was always maintained so that seeds and soil inoculum used for testing soil-feedback responses had the same training phase context (Figure III.1).

Experiment

The experiment was based on the conceptual framework of plant-soil feedback (Bever 2003; Ehrenfeld et al. 2005). At first it included a soil training phase, which was performed over two plant generations, followed by the actual feedback experiment (Figure III.1). The final feedback experiment had a fully 2 x 2 x 2 x 2 factorial design. It consisted of all combinations of soils from two locations (Thyrow soil, Dahlem soil), two seed morph histories resulting from the training phases (non-pappus seed history, pappus seed history), two seed types tested (non-pappus seed, pappus seed), and two soil treatments (soil inoculated with trained soil inoculum, sterile soil as control). During the training phase, we had four replicates for the different combinations of soil and seed type history. At the feedback stage, we replicated three times at the genotype level, which allowed us to account for the genotype effect. In total we had 192 experimental units at the feedback step, where one individuum (plant originating from pappus seed type with pappus seed history in sterile Dahlem soil) died during the final experiment.

For the soil training, we let plants grow either from non-pappus or pappus seeds in two different soil types. After five weeks of growth, plants started to flower. Flowering stems with unopened capitula were separately enclosed in paper bags. Wrapped flowers produced seeds with no further manipulation, meaning that pollen transfer was intra-individual (self-pollination) only. That procedure excluded effects resulting from differences in mating system of the respective florets, which might contribute to ecological differences of the distinct seed types of heterocarpic plants (Olivieri et al. 1983; Cheptou et al. 2001). After an additional 11 weeks of growth, shoots and seeds were harvested while roots remained in the pots. Using seeds produced by plants, we started a second training
round consisting of 18 weeks of plant growth. Analogously to the seed morph scheme of the first training round, we planted either seedlings that originated from non-pappus or pappus seeds, i.e. the identity of genotypes in the pots was maintained (Figure III.1). Again, plants of the second training step reproduced geitonogamously, which minimized genetic variability. As a result of the first and second training round, we had non-pappus and pappus seeds produced by plants, which had either a non-pappus or pappus seed history for two plant generations and, further, corresponding soils trained by plants arising either from non-pappus or pappus seeds (Figure III.1). Trained soils and seeds produced by the last training were used for the feedback experiment.

Before using non-pappus and pappus seeds in the final feedback step, we determined average seed weight of the different seed types produced per plant (in groups of 10 seeds each). During the feedback experiment, we separately collected all seeds developing from a single flower head (up to four capitula per plant if possible). From a total number of 731 capitula, we quantified numbers of non-pappus and pappus seeds produced. At harvest, we counted the number of capitula per plant and removed shoots from roots. Roots were separated from soil and carefully washed under a stream of water. Biomasses of roots and shoots were weighed after drying for five days at 40 °C. To assess mycorrhizal and other fungal root colonization, we used the ink-vinegar method of Vierheilig et al. (1998) and stained a representative root sample of each plant (ca. 100 mg of dry root material). Arbuscular mycorrhizal (AM) root colonization levels (hyphae, arbuscules, vesicles), and analogously of non-AM fungal structures, were determined using the magnified intersect method of McGonigle et al. (1990) based on 100 intersections per root sample examined at 200X.

In all experimental phases, plants grew in 6 x 25 cm, 400 ml Conetainer pots (Stuewe and Sons., Oregon, USA) in fully randomized arrangements. The first training step was completed from May until September 2010 under greenhouse conditions (natural photoperiod, day temperature 21-34 °C, night temperature 18-19 °C). The second training round and also the feedback experiment were performed in a climate chamber (16 hour-photoperiod, day temperature 20 °C, night temperature 18 °C, humidity 60%). Plants were always watered as needed with the same amount of water.
Statistical analyses

To calculate soil feedback responses we used the approach of ‘home vs. away’ contrasts according to other feedback studies (e.g. Bever 1994; Klironomos 2002). In our experiment, soil inoculated with trained soil represented ‘home’, and sterile soil was defined as ‘away’. Negative feedback, therefore, was determined by subtracting the average measure of a given response variable in sterile soil from the average measure in soil inoculated with trained soil. Further, to calculate a value for individual investment in dispersal, we estimated the ratio between number of pappus seeds produced per individuum divided by the number of total seeds produced per individuum (in contrast to Cheptou (2008) using number of non-dispersing seeds as the numerator).

To answer if non-pappus and pappus seed of *G. parviflora* differ in soil feedback responses, we used linear mixed model analysis (Gotelli and Ellison 2004). The mixed model was specified with four factors as fixed effects (soil, seed history, seed type tested, soil treatment) and plant genotype as random effect. Treating genotype as random effect allows broader generalization of the results. Running the saturated model, we found no significant influences of the four-way and any of the three-way interactions on response variables tested (except for a significant interaction termed soil*seed type tested*soil treatment for total biomass). Therefore, we simplified the model to the hypothesized seed type-soil treatment interaction. Applying log-likelihood ratio tests, we tested if the model simplification was allowed, i.e. if the reduced model had a better fit than the saturated or respective higher-order interaction model.

To deal with heterogeneity within the data, we incorporated different variances per stratum in the model (Zuur et al. 2009) by using the ‘varIdent’ function in R (R Development Core Team 2009). Thus, the optimal model for number of total seeds and pappus seeds per plant was specified by a variance structure of varIdent(form=~1|soil). For root biomass and number of capitula per plant, models with varIdent(form=~1|soil*soil treatment) had the lowest AIC and, therefore, were selected. Total biomass data were analyzed with varIdent(form=~1|soil*soil history). Data estimated as pappus seed dispersal ratio showed homogeneity, but were Box-Cox transformed (exponent 2) to meet assumptions of normality. To achieve normality, exponentiation was also needed for data of seed weight of seeds produced by the second training round (exponent –0.3434343). To normalize percentages of AM fungal root colonization by hyphae and arbuscules we used
arcsin square root transformation. Data of percent root colonization by non-AM fungi were log-transformed.

To identify differences in root colonization (AM fungal structures of hyphae, arbuscules, vesicles and non-AM fungi) we used the linear mixed model approach, where data from the treatment ‘soil inoculated with trained soil’ were analyzed exclusively (fixed effects: soil, seed history, seed type tested; random effect: plant genotype). Differences between treatment groups indicated by mixed effect linear models were always compared with a posteriori pair-wise comparisons based on the resulting 95 % confidence limits (Zuur et al. 2009). For this purpose, standard errors (SE) of means and corresponding confidence limits were calculated from the mixed effect models that took into account the random effect of plant genotype. This procedure made these comparisons very conservative and facilitated interpretation in terms of which treatment groups differed significantly. Seed size of seeds produced during the second training round was analyzed by Analysis of Variance (three-way ANOVA; factors: soil, seed history and seed type; \( P < 0.05 \)). All statistical analyses were performed using R version 2.10.1 (R Development Core Team 2009).

Results

Always starting with the removal of the highest order interaction effect, we found that the models without the four and all three-way interactions fitted the data best. Overall, reproductive output of *Galinsoga parviflora* was strongly influenced by soil treatment, i.e. depended on if soil was sterile (hereafter sterile) or inoculated with trained soil (hereafter trained) (Table III.1). In sterile soil, plants produced significantly more flower heads (mean ± SE calculated from the intercept of the mixed model; sterile: 53.8 ± 1.7, trained: 42.0 ± 1.4), and had on average more seeds (sterile: 1668.1 ± 56.2, trained: 1469.6 ± 56.1). Total and pappus seed number per plant were highly positively correlated (\( r = 0.996 \), Pearson's product-moment correlation).

Feedback contrast between plants from non-pappus and pappus seeds

A significant interaction term indicated that most measured response variables to treatments were mediated by seed type tested (non-pappus vs. pappus) (Table III.1). For
plants grown from pappus seeds, root biomass and measurements of reproduction, as number of capitula, and average number of total seeds and pappus seeds per plant, were strongly increased in sterile compared to trained soil treatment (Table III.2). For plants grown from non-pappus seeds, we also found a significantly higher number of capitula, but differences in root biomass, pappus and total seeds per plant were less pronounced in sterile vs. trained soil treatment. However, dispersal ratio by plants arising from non-pappus seeds showed a significant shift towards a higher portion of pappus seeds in trained soil treatment compared to all other combinations of treatment and seed type tested (Table III.2). Negative feedback contrasts for root biomass, number of capitula, total and pappus seeds per plant were always larger for plants grown from pappus seeds than those from non-pappus seeds (Figures III.2a–c). In fact, feedback for dispersal ratio of plants grown from non-pappus seeds was even positive, while tending to be negative for plants grown from pappus seeds (Figure III.2d).

Seed-type treatment patterns had no correspondence with root colonization by putatively pathogenic fungi. In the trained soil treatment, roots were extremely rarely colonized by non-AM fungi (trained: 0.06 ± 0.04 %). We detected no significant effect of soil, seed history or seed type tested on non-AM fungal root colonization (Supplemental Table A.III.1). Furthermore, we found no significant effect of any of the factors tested on root colonization by AM hyphae and AM vesicles (Supplemental Table A.III.1). Merely colonization by AM arbuscules in trained soil treatment indicated a strong soil effect (soil: \( P < 0.001 \), mixed effect model). When plants grew in soil inoculated with trained soil from Dahlem, percentage colonization by arbuscules was significantly higher than for plants growing in soil treated with trained soil from Thyrow (trained Dahlem: 65.4 ± 3.9 %, trained Thyrow: 38.5 ± 3.9 %). AM hyphal root colonization was in both inoculated soils similar (trained Dahlem: 81.6 ± 3.8 %, trained Thyrow: 80.1 ± 3.8 %).

Effects of soil and soil–treatment interaction

Depending on soil type, biomass measures and dispersal ratio differed significantly (Table III.1). Overall, plants growing in Thyrow soil had greater root biomass (Thyrow: 0.444 ± 0.032 g, Dahlem: 0.323 ± 0.034 g), as well as more total biomass (Thyrow: 1.743 ± 0.132 g, Dahlem: 1.295 ± 0.141 g). The dispersal ratio was also strongly increased for plants in Thyrow soil (Thyrow: 0.775 ± 0.008, Dahlem: 0.732 ± 0.008). Further, a significant interaction between soil and soil treatment was found for biomass measures and
number of capitula per plant (Table III.1). In Thyrow soil, inoculation with trained soil decreased number of capitula per plant, while root and total biomass were only minimal influenced compared to the sterile treatment. In Dahlem soil, however, presence of trained soil had a strong growth reducing effect (Table III.3). Further, number of capitula per plant was significantly increased in sterile Dahlem soil compared to all other combinations of soil and treatment tested (Table III.3).

Plant growth depending on seed type and the seed history–seed type interaction

Plant root biomass was significantly influenced by the interaction between seed type and seed history over two plant generations (Table III.2). Plants grown from pappus seeds (papp), which also had a pappus history (papp hist), produced the smallest root biomass (papp and papp hist: 0.346 ± 0.039 g). But when plants grown from pappus seeds came from plants having a non-pappus history (non-papp hist), root biomass was found to be greatest compared to all other combination of seed history and seed type tested (papp and non-papp hist: 0.435 ± 0.039 g). For plants originating from non-pappus seeds (non-papp), the influence of the seed history tended in the same direction, but was less strong (non-papp and papp hist: 0.368 ± 0.039 g, non-papp and non-papp hist: 0.403 ± 0.039 g). Overall, root biomass for plants grown from the distinct seed types differed only marginally; the indicated significant seed type effect could not be shown with the 95% confidence limit (non-papp: 0.386 ± 0.028 g, papp: 0.391 ± 0.028 g).

Measurements of seed weight of seeds produced by plants of the second training round indicated a strong seed type effect (Supplemental Table A.III.2, seed type: F$_{1,23}$ = 38.27; P < 0.001, ANOVA). Non-pappus seeds were significantly heavier than pappus seeds (non-papp: 167.9 ± 2.7 µg, papp: 145.0 ± 2.9 µg).

Discussion

Under climate chamber conditions, we found that plants grown from distinct seed morphs of *G. parviflora* differed in the magnitude of soil feedback. The feedback differences between progeny of non-pappus and pappus seeds were most related to reproductive traits, such as number of capitula, total and pappus seeds per plant, but also indicated by root biomass. Consistent with our hypothesis, progeny of non-pappus seeds, which are
produced for *in situ* persistence, were less negatively affected by conditions of ‘self-cultivated’ (trained) soil than progeny of pappus seeds, which easily disperse from the capitula (Espinosa-García et al. 2003) and represent the long-distance dispersal type (Vibrans 1999). Therefore, the magnitude of negative feedback corresponds to the dispersal potential of the different seed types giving an advantage to non-pappus progeny in the existing population.

For dispersal ratio, we even found positive feedback for non-pappus progeny. They proportionally produced more seeds equipped with a pappus when soil was inoculated with trained soil. Thus, non-pappus progeny showed greater investment in dispersal under unfavorable conditions of ‘self-cultivated’ soil. This strategy would allow escape from sib competition and negative density effects (Venable and Brown 1988), although pappus seeds of *G. parviflora* have a higher risk of failure than non-pappus seeds (Venable 1985a, Simons 2011). Pappus progeny, in contrast, did not alter the ratio of the two seed types depending on soil treatment, and displayed neutral feedback for dispersal ratio. Less favorable environmental conditions, therefore, changed the proportion of seeds with dispersal structures in non-pappus progeny only. To our knowledge, this is the first evidence that shifts in the ratio of particular seed types produced by heterocarpic plants are associated with the seed morph from which the plant was arising. Besides, our result confirms other observations that dispersal ratio reflects a highly variable trait by which heterocarpic species adjust to environmental conditions (Mandák 1997). Interestingly, the direction of the adjustment for dispersal depends on the type of environmental stress. For example, nutrient depletion in *Crepis sancta* (Imbert and Ronce 2001) or high competition in *Hypochoeris glabra* (Baker and O’Dowd 1982) increases proportion of dispersing seeds, while habitat fragmentation in *C. sancta* (Cheptou et al. 2008) and increasing aridity in *Hedypnois rhagadioles* (Kigel 1992) or the genus *Picris* along a gradient from mesic to arid (Ellner and Schmida 1984) leads to a higher proportion of non-dispersing seeds.

Contrary to our hypothesis, divergent feedback of plants grown from the different seed types of *G. parviflora* did not coincide with differences in fungal root infection of non-pappus versus pappus progeny. Overall, percentage of root colonization by potentially pathogenic fungi in trained soil treatment was very low and not impacted by factors tested. Thus, we found no evidence for non-pappus progeny being less colonized and, therefore, less negatively affected by fungal pathogens. We also failed to find significant influences of the seed type tested on root colonization by AM fungi, which can also generate negative feedback (Bever 2002). Consequently, differences in negative feedback of non-pappus and
Pappus progeny must have been related to other microorganisms of the very complex soil community (Bever 2003): possibly nematode pathogens (De Deyn et al. 2003) or bacterial antagonists, which we did not evaluate in this study. However, influences of soil bacteria and fungi on different seed morphs of *G. parviflora* were investigated by Espinosa-García et al. (2003). Focusing on differential longevity of the two seed types within the soil, they found no correlation between different loss rates from the seed bank and their susceptibility to fungi or bacteria. Further, they report low incidence of fungal infection in the seeds. Given the fact that we also very rarely observed colonization by non-AM fungi in the roots, pathogenic fungi may be of minor importance for *G. parviflora*.

However, in contrast to Espinosa-García et al. (2003) studying *G. parviflora* in the native region (Damalas 2008), our experiment was conducted in the introduced range where most likely generalist pathogens rather than species-specific play a major role (Kulmatiski et al. 2008). As Klironomos (2002) demonstrated, introduced species often accumulate pathogens slowly and, therefore, experience only neutral feedback in the presence of a ‘self-cultivated’ pathogen/saprobe fraction. But, since we found negative feedback by inoculating with whole soil also including mutualists, our soil training over two plant generations was likely sufficient to generate an adequate pathogen load in the soil. Nevertheless, it would be interesting to test feedback responses of the two seed types to different soil fractions, and to disentangle what kind of soil biota might have caused the greater negative feedback of plants grown from pappus than non-pappus seeds.

Further, our data show that plants grown from pappus seeds had significantly more capitula and greater reproductive output under sterile conditions than those from non-pappus seeds. Under control conditions, therefore, the tendency of a colonizer to produce large numbers of viable seeds (Warwick and Sweet 1983) was more pronounced in progeny of pappus than non-pappus seeds. Thus, greater dispersal capacity of pappus seeds in *G. parviflora* might be associated with higher investment in reproductive output. Such a relationship would be adaptive because it permits pappus progeny a rapid population buildup after pappus seeds have reached new habitats.

Conversely, we did not find that non-pappus progeny, which maintain the existing population, invested more in biomass. This was surprising given the greater seed size of non-pappus compared to pappus seeds, and the observed positive relationship between seed weight and seedling growth after 9 days of emergence (Rai and Tripathi 1982). However, the positive correlation between seed size and plant growth/reproductive output was also shown to depend on environmental conditions, such as competition and nutrient
deficiency. Rai and Tripathi (1987) planting seedlings arising from the different seed types in monoculture and mixture demonstrated that greater seed weight of non-pappus seeds is only advantageous at low or medium fertilizer levels. At high fertilizer dose, plants grown from smaller pappus seeds performed better; hence, soil nutrient status altered competitive ability of non-pappus and pappus progeny. Therefore, it would be interesting to evaluate feedback of plants grown from the two morphs in pure and mixed competitive situations. Further, other studies of heterocarpic Asteraceae also report that greater seed size is not always converted into biomass growth, but becomes advantageous in the presence of competition (Imbert et al. 1997; De Clavijo and Jiménez 1998).

Our data, moreover, indicate that performance of plants grown from the two seed types was mediated by the seed type of the mother/grandmother plant. In the feedback step, pappus progeny produced significantly more root biomass when they had a non-pappus history, i.e. came from non-pappus mother/grandmother plants. For non-pappus progeny, we did not find such a history influence. Thus, by having a non-pappus mother/grandmother pappus progeny showed increased belowground growth, which might be advantageous in terms of improved efficiency of plant nutrient uptake (Imbert et al. 1997) and occupying space in a new colonized patch, respectively. Our result is the first evidence for a maternal history effect in heterocarpic species. This finding may be considered reliable because we used a very conservative approach for posteriori pair-wise comparisons. Therefore, further research into the phenomenon of heterocarpic species may not only consider differences of plants grown from the different seed types, but also investigate influences of the seed morph history of prior plant generations.

To conclude, progeny from distinct seed morphs of *G. parviflora* differ in their soil feedback: plants arising from non-pappus seeds are less affected by unfavorable conditions of ‘self-cultivated’ soil than plants from pappus seeds. Beside feedback, the differentiation between the two seed morphs in *G. parviflora* involves dispersal potential, dispersal pathway (Vibrans 1999), viability in the soil (Espinosa-García et al. 2003), as well as competitive ability (Rai and Tripathi 1987). Therefore, the different strategies of the two seed types are ecologically very complex. The greater dispersal capacity away from maternal plants of lighter pappus seeds is accompanied by reduced performance in home soil, shorter survival in the seed bank, and lower competitive ability in mixed populations under moderate soil fertility. Further, pappus progeny, which are produced for founding new populations, exhibit more the traits of a colonizer by having greater reproductive output under sterile conditions compared to non-pappus progeny. Conversely, plants
arising from heavier non-pappus seeds, which are produced to maintain the existing population, experience reduced negative feedback compared to plants from pappus seeds in the existing population, which might be adaptive. Therefore, in mixed populations and under moderate soil fertility the less negative feedback and greater competitive ability of non-pappus progeny can be converted into better performance in the local habitat compared to pappus progeny.

**Acknowledgements**

We are deeply grateful to Tancredi Caruso for statistical advice and E. Kathryn Barto for discussion. We also thank Anika Lehmann and Weishuang Zheng for help during the harvest of the feedback experiment. Further, we thank the Botanical garden in Münster for providing plant seeds, Michael Baumecker for the possibility to use soil from Thyrow, and Roland Buchhorn for providing greenhouse space of the Julius-Kühn Institute.
### Tables and figures

**Table III.1** Mixed effect model analyses on plant biomass and reproductive traits of *Galinsoga parviflora* in the feedback experiment. Plant genotype was treated as a random effect. Values in bold indicate significance at $P < 0.05$.

<table>
<thead>
<tr>
<th>Factors</th>
<th>Total biomass</th>
<th>Root biomass</th>
<th>Capitula</th>
<th>Total seeds</th>
<th>Papp. seeds</th>
<th>Dispersal ratio</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>$P$</td>
<td>$P$</td>
<td>$P$</td>
<td>$P$</td>
<td>$P$</td>
<td>$P$</td>
</tr>
<tr>
<td>Soil</td>
<td>0.0441</td>
<td>0.018</td>
<td>0.406</td>
<td>0.504</td>
<td>0.316</td>
<td>0.003</td>
</tr>
<tr>
<td>Seed.hist</td>
<td>0.3449</td>
<td>0.292</td>
<td>0.861</td>
<td>0.219</td>
<td>0.133</td>
<td>0.609</td>
</tr>
<tr>
<td>Seed.type</td>
<td>0.1132</td>
<td>0.050</td>
<td>0.447</td>
<td>0.114</td>
<td>0.079</td>
<td>0.213</td>
</tr>
<tr>
<td>Soil.treat</td>
<td>0.1527</td>
<td>0.191</td>
<td>$&lt;0.001$</td>
<td>$0.001$</td>
<td>$0.001$</td>
<td>0.120</td>
</tr>
<tr>
<td>Soil x Seed.hist</td>
<td>0.954</td>
<td>0.987</td>
<td>0.961</td>
<td>0.333</td>
<td>0.261</td>
<td>0.072</td>
</tr>
<tr>
<td>Soil x Seed.type</td>
<td>0.960</td>
<td>0.690</td>
<td>0.675</td>
<td>0.617</td>
<td>0.610</td>
<td>0.659</td>
</tr>
<tr>
<td>Seed.hist x Seed.type</td>
<td>0.090</td>
<td>0.020</td>
<td>0.653</td>
<td>0.873</td>
<td>0.996</td>
<td>0.345</td>
</tr>
<tr>
<td>Soil x Soil.treat</td>
<td><strong>0.001</strong></td>
<td><strong>0.001</strong></td>
<td><strong>0.018</strong></td>
<td><strong>0.077</strong></td>
<td><strong>0.102</strong></td>
<td>0.681</td>
</tr>
<tr>
<td>Seed.hist x Soil.treat</td>
<td>0.927</td>
<td>0.162</td>
<td>0.391</td>
<td>0.119</td>
<td>0.111</td>
<td>0.282</td>
</tr>
<tr>
<td>Seed.type x Soil.treat</td>
<td>0.581</td>
<td>0.049</td>
<td><strong>0.034</strong></td>
<td><strong>0.037</strong></td>
<td><strong>0.017</strong></td>
<td><strong>0.003</strong></td>
</tr>
</tbody>
</table>

Seed.hist, seed history; Seed.test, seed type tested; Soil.treat, soil treatment; Capitula, number of capitula per plant; Total seeds, average number of total seeds per plant; Papp. seeds, average number of pappus seeds per plant; Dispersal ratio, ratio of number of pappus seeds per individuum to number of total seeds number per individuum.
Table III.2 Responses of the distinct seed types of *Galinsoga parviflora* to soil treatment in the experiment. Values of mean ± SE were calculated from the intercept of the mixed effect model; hence, genotype as random effect was always incorporated in SE. Different lower case letters indicate differences according to 95% confidence limit.

<table>
<thead>
<tr>
<th>Variable</th>
<th>Non-pappus seed type</th>
<th>Pappus seed type</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>sterile soil</td>
<td>trained soil</td>
</tr>
<tr>
<td>Root biomass (g)</td>
<td>0.396 ± 0.031</td>
<td>0.370 ± 0.030</td>
</tr>
<tr>
<td>Capitula per plant</td>
<td>51.7 ± 2.2</td>
<td>43.1 ± 1.6</td>
</tr>
<tr>
<td>Pappus seeds per plant</td>
<td>1370.5 ± 65.9</td>
<td>1338.3 ± 65.9</td>
</tr>
<tr>
<td>Total seeds per plant</td>
<td>1581.5 ± 71.5</td>
<td>1514.1 ± 71.5</td>
</tr>
<tr>
<td>Dispersal ratio$^2$</td>
<td>0.742 ± 0.01</td>
<td>0.771 ± 0.01</td>
</tr>
</tbody>
</table>

$^1$ soil inoculated with trained soil inoculum; $^2$ ratio of number of pappus seeds per individuum to total seeds number per individuum.

Table III.3 Response variables of *Galinsoga parviflora* to soil and soil treatment in the experiment. Values of mean ± SE were calculated from the intercept of the mixed effect model with genotype as a random factor. Different lower case letters indicate differences according to 95% confidence limit.

<table>
<thead>
<tr>
<th>Variable</th>
<th>Dahlem soil</th>
<th>Thyrow soil</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>sterile</td>
<td>trained</td>
</tr>
<tr>
<td>Total biomass (g)</td>
<td>1.498 ± 0.148</td>
<td>1.098 ± 0.147</td>
</tr>
<tr>
<td>Root biomass (g)</td>
<td>0.369 ± 0.035</td>
<td>0.252 ± 0.036</td>
</tr>
<tr>
<td>Capitula per plant</td>
<td>58.0 ± 3.4</td>
<td>37.5 ± 2.9</td>
</tr>
</tbody>
</table>
Figure III.1 Design of the feedback experiment with a soil training phase performed over two plant generations. During the first training step, the initial soil was either trained by plants originating from non-pappus or pappus seeds of *Galinsoga parviflora*. For the second training round, roots of plants of the first training remained in the pots and plants germinating from non-pappus or pappus seeds (first training progeny) were planted congruently with the first training scheme (i.e. pots trained by non-pappus seeds were trained again by non-pappus seeds, and equivalently for pappus seeds). In all training stages, plants reproduced by self-fertilization (autogamy/geitonogamy), where flowering stems with unopened capitula were separately enclosed in paper bags allowing intra-individual pollen transfer (self-pollination) only. For the feedback experiment, progeny and soil of the second training round were used. To prepare ‘soil inoculated with trained soil’, trained soil was thoroughly mixed into sterile background soil (ratio 1:10). Identity of replicates was always maintained.
Figure III.2 ‘Trained vs. sterile’ soil contrasts of plants grown from non-pappus (grey) and pappus seeds (white) of *Galinsoga parviflora* in the experiment: (a) plant root biomass; (b) number of capitula per plant; (c) pappus seeds per plant; (d) dispersal ratio. Contrasts were determined by subtracting the average measure of a given response variable in sterile soil from the average measure in soil inoculated with trained soil, which follows the calculation of ‘home vs. away’ contrasts in plant–soil feedback studies (Bever 1994; Klironomos 2002). Error bars were calculated from the errors (SE) of the linear mixed effect model with genotype as a random factor.
References


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CHAPTER 5

Summary

Numerous ecosystems worldwide are influenced by invasive species in their functioning in a multitude of ways (Wardle et al. 2011). Invasive species, moreover, can damage ecosystem services that are fundamental to human well-being resulting in substantial economic costs (e.g. Pimentel et al. 2005; Pejchar and Mooney 2009; Vilà et al. 2010). Therefore, it is of increasing urgency to better understand the mechanisms involved in the invasion process as human activities such as international trade, transport and travel, which cause species dispersal into new ranges, continue to expand (Keller et al. 2011).

The success of invasive plant species must be regarded as highly context-dependent and linked to a combination of both abiotic and biotic factors, and multiple mechanisms (e.g. Daehler 2003; Richardson and Pyšek 2006; Barney and Whitlow 2008). In particular, biotic interactions of invasive species with their new environment may be the key driver for the successful spread into new areas (Jeschke et al. 2012). However, despite a large number of articles published both on experimental and theoretical topics in invasion biology per year (Kühn et al. 2011), it is still difficult to give precise statements on why a particular plant species becomes a dominant component in a plant community where it is not native.

In the present dissertation, I report about the interaction of invasive plant species with belowground organisms, especially arbuscular mycorrhizal (AM) fungi. I conducted a series of experiments to examine the importance of the AM fungal association for the successful spread of Ambrosia artemisiifolia in the new European range; A. artemisiifolia has been proposed to be facilitated by the symbiosis with AM fungi in Central Europe (Fumanal et al. 2006). Further, I focused on the phenomenon of heterocarpy of the non-native plant Galinsoga parviflora in a soil feedback study. I investigated if the different dispersal capacities of the two distinct seed types correlate to different soil feedback responses, which may contribute to the success of G. parviflora in the new range. In all experiments, I always maintained the ecological context of soil and natural AM fungal
community; hence, my findings have a comparatively high realism. Below I summarize the main results from the three manuscripts.

**Manuscript I** (Chapter 2) Divergent responses of *Ambrosia artemisiifolia* to natural AM fungal communities in the new European range.

**Background and Aims** AM fungi may act more or less cooperatively in association with plants exhibiting functions from mutualism to parasitism depending on soil and light environments (Johnson et al. 1997; Kiers et al. 2011). Therefore, natural selection pressure in arbuscular mycorrhizas may favor the AM fungi–plant combinations that are the most fit under their respective local environmental circumstances, promoting local adaptation and co-adaptation in AM associations (Helgason and Fitter 2009; Hoeksema 2010). Recently, existence of coadapted AM fungal–plant interactions has been found (Johnson et al. 2010; Ji et al. 2010), but knowledge about the extent to which such adaptations also occur during plant invasions is lacking. In this chapter, I investigated whether or not the mycorrhizal symbiosis between *A. artemisiifolia* and native AM fungal communities shows evidence of co-adaptation in the new European range. In a ‘local vs. foreign’ reciprocal inoculation experiment, I compared performance of plant genotypes from two different sites: a roadside and a cornfield habitat.

**Results** Natural AM fungal assemblages were found to be mutualistic with *A. artemisiifolia* in low fertility roadside soil, but not in agricultural soil (Figure I.3a). Decreased plant growth in response to the less cooperative quality of the agricultural AM fungal community in the agricultural system coincided with alterations of plant root systems towards greater fineness. I found no evidence for locally adapted plant-AM fungal interactions, but adaptation of roadside plants to a ‘local’ roadside soil environment (Figure I.4). Further, soil conditions had a strong effect: plants growing in more fertile cornfield soil produced more biomass, had greater total seed weight, and flowered earlier than plants in less fertile roadside soil.

**Conclusion** The results of this study indicate that performance of non-native *A. artemisiifolia* may be influenced by different mycorrhizal functions, leading to mutualism in less fertile roadside habitats and parasitism in more fertile cornfield soils.
Moreover, the study highlights the great importance of the soil context for plant responses to mycorrhizal inoculation, and that mycorrhizal functions may be unpredictable when AM fungal communities are introduced to novel soils. Further, the findings show that adaptation to soil conditions may play a crucial role in the early stages of the spread of *A. artemisiifolia* along the road.

**Manuscript II** (Chapter 3) Non-native *Ambrosia artemisiifolia* are more influenced by relative density and identity of neighboring plant species than arbuscular mycorrhiza.

**Background and Aims** One way by which invasive plants can interact with AM fungi has been described as the enhanced mutualisms hypothesis or facilitation, whereby invasive plants are positively influenced by the AM fungal association of the new range to the detriment of native species (Reinhart and Callaway 2006; Shah et al. 2009). In this context, several studies have shown that AM fungi may contribute to the dominance of invasive over native plants by altering competitive interactions (e.g. Marler et al. 1999; Callaway et al. 2004b; Shah et al. 2008). In this chapter, I investigated the effects of natural AM fungal communities on the competitive ability of *A. artemisiifolia* in two greenhouse experiments always maintaining the ecological context of soil and AM fungi. I studied *A. artemisiifolia* grown together with one of four co-existing mycorrhizal plant species in a 1:4 (target) or 4:1 (challenger) relative density. As neighbor species I selected *Conyza canadensis, Artemisia vulgaris, Daucus carota* and *Tanacetum vulgare*, which I found co-occurring with *A. artemisiifolia* in ruderal communities. Moreover, I studied the influence of a roadside AM fungal community on *A. artemisiifolia* and *D. carota* grown in pairwise situations of intra- and interspecific competition.

**Results** Regardless of presence/absence of AM fungal communities, *A. artemisiifolia* was highly dominant in all interspecific competitive arrangements under the nutrient poor soil conditions tested. Divergent AM fungal effects on biomass of *A. artemisiifolia* as a function of neighbor plant were only indicated as trends: target *A. artemisiifolia* tended to either increase in shoot biomass (with *C. canadensis* or *T. vulgare*), decrease (with *D. carota*) or was unaffected (with *A. vulgaris*) under presence of mycorrhiza (Figure II.3a). In pairwise competitive situations, roadside AM fungi had an amplifying (negative) effect on *A. artemisiifolia* in intraspecific competition, and a neutral effect in
mixture with *D. carota*. Moreover, *A. artemisiifolia* experienced strong competition by conspecifics, which caused decreases in shoot biomass and number of male inflorescences, but earlier flowering of female flowers. Among the mycorrhizal plant competitors, *C. canadensis* performed poorly compared to the other neighboring species tested.

**Conclusion** The results of the experiments demonstrate that *A. artemisiifolia* is an exceptionally good competitor both at low and high relative density in comparison to co-existing mycorrhizal plant species in the new European range under nutrient poor soil conditions. Moreover, my findings suggest that the competitive ability of *A. artemisiifolia* – a successful pioneer plant and a species with a strongly ruderal life history – is very weakly influenced by natural AM fungal communities in the presence of other mycorrhizal plants in low fertility soils. Therefore, the invasive success of *A. artemisiifolia* in Central Europe may not be related to facilitation by natural AM fungal communities.

**Manuscript III** (Chapter 4) Distinct seed morphs of *Galinsoga parviflora* (Asteraceae) give rise to different soil feedbacks.

**Background and Aims** Heterocarpy is the phenomenon that a single plant produces two or more distinct fruit types, which often differ in dispersal mechanisms and ecological behavior (*sensu* Tanowitz et al. 1987). Beside fruit, the terms diaspore or seed are also used: I refer to seed *sensu lato*. Heterocarpy has been extensively studied from the perspective of seed size, dormancy and germination behavior (Mandák 1997). Some studies, further, report about differences in competitive ability of plants arising from different seed morphs (e.g. Rai and Tripathi 1987; Imbert et al. 1997) or the influence of environmental conditions (urban habitat fragmentation) on the ratio of non-dispersing to dispersing seeds of a heterocarpic plant species (Cheptou et al. 2008). In this chapter, I investigated another aspect of ecological behavior. I asked if plants arising from distinct seed types differ in their interaction with soil biota, i.e., exhibit divergent responses of soil feedback. I studied the non-native plant *G. parviflora*, which produces seeds both equipped with a pappus for long-distance dispersal and seeds without a pappus (non-pappus) for maintaining the existing population. The hypothesis was that plants arising from non-pappus seeds would exhibit better performance, i.e., less negative soil feedback, in soil trained by the mother plant than plants grown from pappus seeds. To test this, I trained soil
over two plant generations with plants either arising from non-pappus or pappus seeds, and studied feedback responses of pappus and non-pappus progeny as ‘trained versus sterile’ soil contrasts (Figure III.1).

**Results** Progeny grown from distinct seed morphs of *G. parviflora* differed in soil feedback. Plants grown from pappus seeds had greater root biomass and produced more flower heads, as well as total number of seeds and pappus seeds per plant in sterile compared to ‘self-cultivated’ (trained) soil conditions. For plants arising from non-pappus seeds the differences between sterile and trained soil treatment were less pronounced. Hence, negative feedback contrasts for the above mentioned response variables were always larger for plants grown from pappus seeds than those from non-pappus seeds (Figures II.2a–c). Progeny of non-pappus seeds, moreover, showed a significant shift towards a higher portion of pappus seeds in trained soil compared to all other combinations of soil treatment and seed type tested (Figure II.2d). Further, the data indicated that plants from pappus seeds produced significantly more root biomass when they had a non-pappus history, i.e. came from non-pappus mother/grandmother plants. For non-pappus progeny, we did not find such a history influence. Soil Feedback differences could not be correlated with differences in root infection with pathogenic fungi or colonization by AM fungi.

**Conclusion** Consistent with the hypothesis, the results demonstrate that progeny from non-pappus seeds, which are produced for *in situ* persistence, are less negatively affected by conditions of ‘self-cultivated’ (trained) soil than progeny from pappus seeds, which easily disperse from the capitula and represent the long-distance dispersal type. Therefore, the magnitude of negative feedback corresponds to the dispersal potential of the different seed types giving an advantage to non-pappus progeny in the existing population. Plants grown from non-pappus seeds, moreover, showed proportionally greater investment in long-distance dispersal under unfavorable conditions of ‘self-cultivated’ soil, which may allow escape from sib competition and negative density effects.

**Synthesis**
Chapter 2 and 3 of this dissertation highlight the importance of the ecological context in AM research. Recently, *A. artemisiifolia* has been repeatedly cited as an example of a plant whose invasive success is facilitated by AM fungi in the new range (e.g. Shah et al. 2009;
This statement is based on a study by Fumanal et al. (2006) showing that *A. artemisiifolia* is colonized by AM fungi in different habitats in France and, moreover, demonstrating that single-grown plants produced more biomass when soil was inoculated with the AM fungus *Glomus intraradices*. In this regard, my research was motivated to find further evidence for *A. artemisiifolia* being promoted by AM fungi in the new European range under conditions of a maintained ecological context of the soil, natural AM fungal community and plant origin. However, when testing AM fungal communities from a roadside and a cornfield habitat, the presence of AM fungi increased performance of *A. artemisiifolia* in less fertile roadside soil only (Chapter 2). Moreover, natural AM fungal communities did not increase the competitive ability of *A. artemisiifolia* in the presence of co-existing mycorrhizal plant competitors in the new range (Chapter 3). Furthermore, the positive effect of the roadside AM fungal community on *A. artemisiifolia* in isolation was reversed when *A. artemisiifolia* grew with a conspecific (Chapter 3). Therefore, *A. artemisiifolia* is an unconvincing example of how an invasive plant is promoted by AM fungal associations in the new range. Contrary to Fumanal et al. (2006), the results of this dissertation, which are always based on experiments maintaining the ecological context of soil and AM fungi, show that the successful spread of *A. artemisiifolia* may not be related to the impact of AM fungi in Central Europe. These findings further illustrate that a plant species’ response in isolation to an AM fungal isolate does not necessarily predict its response to natural AM fungal communities under comparatively high realism and its response to AM fungal communities in competitive situations.

Chapter 4 of this dissertation demonstrates that the role of propagules in invasive plant processes can be ecologically very complex, in particular when invasive plants are heterocarpic, i.e., produce distinct types of seeds on a single plant. *G. parviflora* represents such a non-native, heterocarpic plant species, which produces pappus and non-pappus seeds. The findings of this dissertation, both experimental and literature researched, illustrate the different strategies of the two seed types of *G. parviflora*: greater dispersal ability away from maternal plants of lighter pappus seeds is accompanied with shorter survival in the seed bank, and reduced performance of plants arising from pappus seeds in home soil, as well as lower competitive ability in mixed populations under moderate soil fertility compared to plants from non-pappus seeds, which experience less negative soil feedback. Therefore, heterocarpy in non-native species *G. parviflora* may help that plant to
successfully cope with disadvantageous conditions of ‘self-cultivated’ soil in existing populations (non-pappus seed type) and to colonize new habitats (pappus seed type).

**Future perspectives**

Further research into the successful spread of non-native *A. artemisiifolia* should focus on questions other than facilitation by AM fungi. The diametrically opposed results reported on the competiveness of *A. artemisiifolia* in the new range to date – competitively dominant to co-existing mycorrhizal plant species (Chapter 3, present thesis) and strongly competitively inferior to *Lolium multiflorum* (Leskovšek et al. 2012) – highlight the need for additional competition studies in different ecosystems. Evolutionary aspects, such as adaptation of *A. artemisiifolia* to soil conditions, as indicated by the present dissertation for a roadside habitat, may also be interesting for further research and may give insights into rapid evolution of local adaptation of non-native plants in harsh environmental conditions of the new range. In this context, the spread of *A. artemisiifolia* along the road may also be studied from the perspective of adhesive seed transport by vehicles and propagule pressure (von der Lippe and Kowarik 2012).

Moreover, results presented in this dissertation strongly suggest further research into soil feedback of progeny from heterocarpic plants. It would be interesting to test responses of progeny from the different seed types of *G. parviflora* to different soil fractions, and to disentangle what kind of soil biota may have caused the greater negative feedback of plants grown from pappus compared to those from non-pappus seeds. Aside from *G. parviflora*, other annual plant species, like *Centaura solstitialis* L., *Crepis foetida* L., *Crepis sancta* (L.) Babc., *Hedypnios cretica* (L.) Dum. Cour., *Leontodon saxatilis* Lam. or *Picris echioides* L., which also produce seeds both equipped with a pappus and without, may represent appropriate study species. In addition, maternal history effects, i.e., influence of the seed type of mother/grandmother plants on heterocarpic progeny, need further evaluation and may be more common in heterocarpic plant species.
Zusammenfassung


In dieser Dissertation berichte ich über die Interaktion von invasiven Pflanzen mit Bodenorganismen, insbesondere mit arbuskulären Mykorrhizapilzen (AM). Ich habe eine Reihe von Experimenten durchgeführt, um die Bedeutung der AM-Pilz-Assoziation für die erfolgreiche Ausbreitung von *Ambrosia artemisiifolia* im neuen europäischen Gebiet zu prüfen; *A. artemisiifolia* wurde vermutet, durch die Symbiose mit AM-Pilzen in Mitteleuropa gefördert zu sein (Fumanal et al. 2006). Ferner habe ich mich dem Phänomen der Heterokarpie der nicht einheimischen Pflanze *Galinsoga parviflora* in einer Studie zum Boden-Feedback gewidmet. Ich untersuchte, ob die unterschiedlichen Ausbreitungsfähigkeiten der zwei distinkten Samentypen mit unterschiedlichen Boden-Feedback
Reaktionen korrelieren, was zum Erfolg von *G. parviflora* im neuen Gebiet beitragen könnte. In allen Experimenten gewährte ich stets den ökologischen Kontext von Boden und AM-Pilzgemeinschaft; meine Ergebnisse haben daher einen vergleichsweise hohen Bezug zur Realität. Im Folgenden fasse ich die wichtigsten Ergebnisse der drei Manuskripte zusammen.

**Manuskript I** (Kapitel 2) Unterschiedliche Antworten von *Ambrosia artemisiifolia* auf natürliche AM-Pilzgemeinschaften im neuen europäischen Gebiet.


**Ergebnisse** In der geringen Bodenfruchtbarkeit des Straßenrand-Bodens wirkten die natürlichen AM-Pilzgemeinschaften mutualistisch auf *A. artemisiifolia*, jedoch nicht im Ackerboden (Abbildung I.3a). Das geringere Pflanzenwachstum im Ackerboden in Verbindung mit der wenig kooperativen Ackerboden-AM-Pilzgemeinschaft ging mit Veränderungen des pflanzlichen Wurzelsystems zu größerer Feinheit einher. Ich habe keinen Hinweis auf lokal adaptierte Interaktionen zwischen Pflanzen und AM-Pilzen gefunden, aber Adaptation der Pflanzen vom Straßenrand zur lokalen Umgebung des


**Manuskript II** (Kapitel 3) Nicht-einheimische *Ambrosia artemisiifolia* sind mehr beeinflusst von relativer Dichte und Identität der Nachbarpflanze als durch arbuskuläre Mykorrhiza.


Manuskript III (Kapitel 4) Distinkte Samenmorphe von *Galinsoga parviflora* (Asteraceae) geben Anlass zu unterschiedlichen Boden-Feedbacks.


Ergebnisse Die Nachkommen aus den distinkten Samenmorphe von *G. parviflora* unterschieden sich im Boden-Feedback. Pflanzen, die aus Samen mit Pappus wuchsen, hatten mehr Wurzel-Biomasse und produzierten mehr Blütenköpfchen sowie eine größere Anzahl an totalen Samen sowie Samen mit Pappus unter sterilen verglichen zu ‚selbstkultivierten‘ (trainierten) Bodenbedingungen. Für Pflanzen, die aus Samen ohne Pappus hervorgingen, waren die Unterschiede zwischen der sterilen und trainierten

101


103
‚selbstkultivierten‘ Bodens in bestehenden Populationen (Samentyp ohne Pappus) und dem Kolonisieren neuer Habitate (Samentyp mit Pappus) erfolgreich zurechtzukommen.

**Zukunftsperspektiven**


heterokarpe Nachkommen vermittelt werden, weitere Beurteilung und könnten möglicherweise mehr verbreitet sein in heterokarpen Pflanzenarten.
REFERENCES TO GENERAL INTRODUCTION AND SUMMARY


Kowarik I, Pyšek P (200X) The first steps towards unifying concepts in invasion ecology were made one hundred years ago: revisiting the work of the Swiss botanist Albert Thellung. Divers Distrib, doi: 10.1111/ddi.12009 (online early).


CONTRIBUTION TO THE PUBLICATIONS


Own contributions: I conceived and designed the experiment, performed the experiment and laboratory work including seed collection in the field, soil sampling and sieving, soil analyses, preparation of mycorrhizal inocula, biomass measurements, root scanning, root staining, as well as fungal root colonization assessments. I analyzed the data, interpreted the results, and I wrote the manuscript.

II  Bäucker C, Rillig MC (2012) Non-native Ambrosia artemisiifolia are more influenced by relative density and identity of neighboring plant species than arbuscular mycorrhiza. (In preparation for submission)

Own contributions: I conceived and designed the experiments, performed the experiment and laboratory work including seed collection in the field, soil sampling and sieving, preparation of mycorrhizal inocula, set-up, run and harvest. I analyzed the data, interpreted the results, and I wrote the manuscript.

III  Bäucker C, Rillig MC (2012) Distinct seed morphs of Galinsoga parviflora (Asteraceae) give rise to different soil feedbacks. (Submitted to Acta Oecologica)

Own contributions: I conceived and designed the experiment, performed the experiment including the two soil training stages and the final feedback experiment, performed laboratory work including biomass measurements, seed counting, root staining, and fungal root colonization assessments. I analyzed the data, interpreted the results, and I wrote the manuscript.
CONGRESS CONTRIBUTIONS

September 7, 2011

Talk: ‘Divergent responses of an invasive plant to native AM fungal assemblages’
41st Annual Meeting of the Ecological Society of Germany, Austria and Switzerland,
Oldenburg, September 5–9, 2011

December 3, 2008

Talk: ‘Die Konkurrenzfähigkeit der Beifuß-Ambrosie in Abhängigkeit von AM-Pilzen’
4. Nationale interdisziplinäre Workshop zu Ambrosia artemisiifolia des Julius Kühn-
Instituts, Braunschweig, December 3–4, 2008

July 3, 2008

Poster: ‘The competitive ability of common ragweed as influenced by its relative density,
soil biota and neighboring plant species’
Conference Plant-Microbial Interactions, Kraków, July 2–6, 2008
APPENDIX A

Overview on existing theories mentioned in the thesis

The following table refers to hypotheses in invasion ecology, which are mentioned in the present thesis. The theories are ordered by three foci/categories that might be derived from hypotheses explaining the success of invasive plants in terrestrial ecosystems. The categories are:

i) features of the invasive species  
ii) characteristics of the new environment/habitat  
iii) interactions of invasive species with their new environment

Bibliographic references cited in Table A.1 are listed under point ‘References to general introduction and summary’.
### Supplemental Table A1 Overview on existing theories mentioned in the thesis.

**Category: features of the invasive species**

<table>
<thead>
<tr>
<th>Hypothesis</th>
<th>Definition/Notes</th>
<th>References</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>Ideal weed</strong></td>
<td>Characteristics and life history traits of the invading species facilitate invasion by enabling them to outcompete native species. <em>Notes:</em> Traits of an ideal weed are related to ‘r-strategists’ (ruderal life history, small seed size, high and early fecundity and fertility, rapid growth), as well as high phenotypic and genotypic plasticity.</td>
<td>Elton 1958; Baker 1965, 1974; Rejmánek and Richardson 1996; Sakai et al. 2001; Pyšek and Richardson 2007</td>
</tr>
<tr>
<td><strong>Propagule pressure</strong></td>
<td>High supply and frequency (number) of plant propagule introductions increase chance of successful invasion due to seed swamping, continual supplementation, high genetic diversity, greater probability of introduction to favorable environments, compensation of high mortality rates and bottleneck-situations, respectively. <em>Notes:</em> Propagules are seeds, adult plants or reproductive vegetative fragments.</td>
<td>Williamson and Fitter 1996b; Lonsdale 1999; Pyšek and Richardson 2006; Colautti et al. 2006</td>
</tr>
<tr>
<td><strong>Lag phase</strong></td>
<td>Many invasions are characterized by a lag phase followed by exponential range expansion. Species success may be delayed several decades or even centuries. <em>Notes:</em> Factors associated with time lags might be intrinsic or extrinsic. Intrinsic factors are rate of population increases (e.g. late fecundity) or occurrence of evolutionary changes. Extrinsic factors are related to lacks of favorable environmental conditions (‘invasion windows’ (Johnstone (1986))).</td>
<td>Kowarik 1995</td>
</tr>
</tbody>
</table>
Supplemental Table A.1 continued.

<table>
<thead>
<tr>
<th>Category: features of the invasive species</th>
<th>Hypothesis</th>
<th>Definition/Notes</th>
<th>References</th>
</tr>
</thead>
<tbody>
<tr>
<td>Evolution of increased competitive ability (EICA)</td>
<td>Invasive success of individuals of a species in the new range is related to re-allocation of resources from defense mechanisms into growth and enhanced competitive abilities. (Individuals of a species taken from an area where they have been introduced produce more biomass than individuals taken from the species native range.)</td>
<td>Notes: EICA assumes absence of herbivores; hence, it is based on enemy release. First study species was <em>Lythrum salicaria</em> L. (purple loosestrife).</td>
<td>Blossey and Nötzold 1995; Bossdorf et al. 2005; Joshi and Vrieling 2005</td>
</tr>
<tr>
<td>Tens rule</td>
<td>The theory estimates potential that a non-native species becomes invasive. It says that 10% of imported species become introduced, 10% of those introduced species become established, and 10% of established species become a pest (i.e. cause high economic costs); hence, 1 in 10 of established species becomes highly invasive.</td>
<td>Note: The theory is based on mathematical models regarding non-native species in Britain (both animals and plants). It refers to terms, which are related to environmental or economic impact (e.g. pest).</td>
<td>Williamson and Brown 1986; Williamson and Fitter 1996a</td>
</tr>
<tr>
<td>Category: characteristics of the new environment</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>-----------------------------------------------</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Hypothesis</td>
<td>Definition/Notes</td>
<td>References</td>
<td></td>
</tr>
</tbody>
</table>
| Fluctuating resource availability | A plant community becomes more susceptible to invasion whenever there is an increase in the amount of unused resources. This can be due to increase in supply (e.g. abiotic disturbance such as rainfall) or a decrease in resource use (e.g. predation or die back of resident plants) or both.  
  *Notes:* The theory rests upon that an invading species must have access to available resources and that a species will have greater success in invading a community if it does not encounter intense competition for these resources from resident species. Competition intensity, therefore, should be inversely correlated with the amount of unused resources. | Davis et al. 2000 |
| Disturbance                     | Invasion success is related to disturbance events, which increase or decrease resource availability, which involves change in historical disturbance regimes. Non-native species have an equal chance of success at colonization and establishment compared to resident species because of changes in the rate or intensity of the turnover rate/flux of resources in a habitat.  
  *Notes:* Disturbance can be natural (e.g. floods, fires, hurricanes) or anthropogenic (e.g. introduction of cattle grazing, damming and straightening of rivers). Resources can include space, nutrients, or light. Disturbance alone is not always followed by invasion. | Sher and Hyatt 1999; Colautti et al. 2006 |
Supplemental Table A.1 continued.

<table>
<thead>
<tr>
<th>Category: characteristics of the new environment</th>
</tr>
</thead>
<tbody>
<tr>
<td>Hypothesis</td>
</tr>
<tr>
<td>-------------</td>
</tr>
<tr>
<td>Opportunity window</td>
</tr>
<tr>
<td>Notes: The theory assumes that most species have periods of relative activity and relative inactivity during a year. Opportunities arise during times when resident species are relatively inactive and are not placing high demands on resources, or when invaders and residents differ in their responses to varying factors.</td>
</tr>
<tr>
<td>Shea and Chesson 2002</td>
</tr>
<tr>
<td>Dynamic equilibrium model</td>
</tr>
<tr>
<td>Notes: The theory is based on the dynamic equilibrium model of species diversity (Huston 1979). Since disturbance can be both abiotic and biotic, increasing frequency and intensity of invasive species also alter disturbance regimes resulting in lower diversity of the community and dominance by the invasive species</td>
</tr>
<tr>
<td>Huston 2004</td>
</tr>
</tbody>
</table>
### Category: characteristics of the new environment

<table>
<thead>
<tr>
<th>Hypothesis</th>
<th>Definition/Notes</th>
<th>References</th>
</tr>
</thead>
<tbody>
<tr>
<td>Environmental heterogeneity</td>
<td>Predicts that heterogeneity both increases invasion success and reduces the impact to native species in the community, because it promotes invasion and coexistence mechanisms that would not possible in homogeneous environments. Notes: Heterogeneity can result from abiotic heterogeneity, i.e. variation in the physical environment or biotic heterogeneity, i.e. variation in the occurrence and abundance of organisms. It can occur both at temporal and spatial scales (interaction neighborhood, local or regional metacommunity). Invasion processes depend on heterogeneity at local and regional metacommunity scales.</td>
<td>Melbourne et al. 2007</td>
</tr>
</tbody>
</table>

### Category: interactions between invasive species with their new environment

<table>
<thead>
<tr>
<th>Hypothesis</th>
<th>Definition/Notes</th>
<th>References</th>
</tr>
</thead>
<tbody>
<tr>
<td>Enemy release</td>
<td>Introduced to new ranges, non-native species experience a decrease in regulation by herbivores and other natural enemies, which results in a rapid increase in distribution and abundance. It assumes that generalists in the new range have greater impact on native than non-native species, and host switching events by specialist enemies of native congeners are rare. Notes: Mitchell and Power (2003) studied 473 species and found support for escape from aboveground enemies: on average 84% fewer fungi and 24% fewer viruses infected the plants in their introduced range in North America compared with their native range in Europe.</td>
<td>Keane and Crawley 2002; Mitchell and Power 2003; Colautti et al. 2004 (for arguments against enemy release)</td>
</tr>
</tbody>
</table>
### Category: interactions of invasive species with their new environment

<table>
<thead>
<tr>
<th>Hypothesis</th>
<th>Definition/Notes</th>
<th>References</th>
</tr>
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<tr>
<td>Novel weapons</td>
<td>Invasive plants profit from differences in the species composition of competitors in the new range compared to their native range because non-native species possess/release root exudates (allelochemicals), which are highly inhibitory to naïve plants in the new range. In the native range, these allelopathic compounds are relatively ineffective against natural neighbors due to co-evolutionary processes. Notes: Root exudates may also function as mediators of new plant–soil microbial interactions. The selective advantage of possessing an allelopathic compound in the new range may result in rapid evolution of the allelochemical, so that root exudates have greater toxicity or are produced in greater quantities. Famous example is <em>Centaurea stoebe</em> (former wrongly termed <em>C. maculosa</em>), which produces (+/−)-Catechin; (−)-Catechin is phytotoxic (e.g. Bais et al. 2003).</td>
<td>Callaway and Ridenour 2004; Hierro et al. 2005</td>
</tr>
<tr>
<td>Increased nitrogen cycling</td>
<td>The hypothesis assumes that non-native plants interact with soil microbes in a biogeographically explicit way resulting in greater nitrogen availability in the soil, from which invasive plants profite more than native plants. Notes: The theory is based on the observation that invaded ecosystems show higher plant productivity and increased soil nitrogen pools/total ecosystem nitrogen stocks (Liao et al. 2008). Invasive plants often produce higher litter quality (contains a higher concentration of nitrogen).</td>
<td>Rout and Callaway 2009; Rout and Callaway 2012</td>
</tr>
</tbody>
</table>
## Supplemental Table A.1 continued.

### Category: interactions of invasive species with their new environment

<table>
<thead>
<tr>
<th>Hypothesis</th>
<th>Definition/Notes</th>
<th>References</th>
</tr>
</thead>
</table>
| Enhanced mutualisms      | Non-native species experience greater positive effects from associations with mutualists, e.g. arbuscular mycorrhizas in the new range than in the native range. The impact of antagonists (pathogens) is similar in both ranges. The theory is based on the assumption that invasive species are able to associate with soil biota in the new range.  
**Notes:** The mechanism behind the positive effect of arbuscular mycorrhizal (AM) fungi on plants in the new range could also be exploitation, because AM fungi associate with multiple plants simultaneously over a mycelial network. For species that associate with (EM), however, invasion success highly depends on introduction of the appropriate EM fungi (Richardson et al. 1994; Richardson et al. 2000b). | Reinhart and Callaway 2006        |
| Mycorrhizal degradation  | The hypothesis predicts that disturbance events that disrupt mutualistic relationships such as mycorrhizas could facilitate the establishment of non-native species. After disturbance non-native species may become dominant, if they are less dependent on mutualism (e.g. AM fungi) and invest little in maintaining the soil community structure. The degraded soil community structure hinders successful re-establishment of native species.  
**Notes:** The theory refers to ecosystems where native plants have strong mutualistic relationships with soil AM fungi. It assumes that i) increased disturbance intensity causes a decline in arbuscular mycorrhizas, ii) non-native species are often less dependent on mutualists (e.g. AM fungi), and iii) plants have different AM fungal hosting abilities. | Vogelsang et al. 2004            |
### Category: interactions of invasive species with their new environment

<table>
<thead>
<tr>
<th>Hypothesis</th>
<th>Definition/Notes</th>
<th>References</th>
</tr>
</thead>
<tbody>
<tr>
<td>Invasional meltdown</td>
<td>In invaded ecosystems, success of non-native species is enhanced by presence of other non-native species due to synergistic interactions among invaders ('invasion domino effect'). Notes: Synergistic effects may arise from direct or indirect facilitative interactions that increase likelihood of survival and/or of ecological impact, and possibly the magnitude of impact of a non-native species.</td>
<td>Simberloff and Von Holle 1999</td>
</tr>
</tbody>
</table>
APPENDIX B

Supplemental Tables A.I.1–A.I.3 to Chapter 2

Supplemental Table A.I.1  Analyses of variance (ANOVA) on the second principal component score (PC2) of the Principal Component Analyses (PCAs) on plant biomass, root traits and mycorrhization of *Ambrosia artemisiifolia* in the experiment. Values in bold indicate significance at $P < 0.05$.

<table>
<thead>
<tr>
<th></th>
<th>PCA</th>
<th>Plant Biomass</th>
<th>Root traits</th>
<th>Mycorrhization</th>
</tr>
</thead>
<tbody>
<tr>
<td>Factors</td>
<td>d.f.</td>
<td>$F$</td>
<td>$P$</td>
<td>$F$</td>
</tr>
<tr>
<td>P.ori</td>
<td>1</td>
<td>0.18</td>
<td>0.672</td>
<td>9.52</td>
</tr>
<tr>
<td>Soil</td>
<td>1</td>
<td>0.20</td>
<td>0.654</td>
<td>3.90</td>
</tr>
<tr>
<td>Myc.treat</td>
<td>2</td>
<td>0.44</td>
<td>0.647</td>
<td>0.39</td>
</tr>
<tr>
<td>P.ori x Soil</td>
<td>2</td>
<td>0.18</td>
<td>0.668</td>
<td>0.84</td>
</tr>
<tr>
<td>P.ori x Myc.treat.</td>
<td>2</td>
<td>2.73</td>
<td>0.069</td>
<td>0.66</td>
</tr>
<tr>
<td>Soil x Myc.treat.</td>
<td>2</td>
<td>0.75</td>
<td>0.473</td>
<td>0.80</td>
</tr>
<tr>
<td>P.ori x Soil x Myc.treat.</td>
<td>2</td>
<td>0.69</td>
<td>0.503</td>
<td>0.81</td>
</tr>
<tr>
<td>Residuals</td>
<td>132</td>
<td>54</td>
<td>132</td>
<td></td>
</tr>
</tbody>
</table>

d.f., degree of freedom; P.ori., plant origin; Myc.treat., mycorrhizal treatment
**Supplemental Table A.1.2** Responses of *Ambrosia artemisiifolia* to soil and mycorrhizal treatment in the experiment.

<table>
<thead>
<tr>
<th>Variables</th>
<th>roadside soil</th>
<th>cornfield soil</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>local soil inoc.</td>
<td>foreign soil inoc.</td>
</tr>
<tr>
<td><strong>Biomass traits</strong></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Shoot (mg)</td>
<td>98.6 ± 7.0&lt;sup&gt;b&lt;/sup&gt;</td>
<td>106.3 ± 6.4&lt;sup&gt;b&lt;/sup&gt;</td>
</tr>
<tr>
<td>Root (mg)</td>
<td>106.9 ± 11.4&lt;sup&gt;b&lt;/sup&gt;</td>
<td>115.0 ± 11.2&lt;sup&gt;b&lt;/sup&gt;</td>
</tr>
<tr>
<td>Reproduction 5 weeks</td>
<td>10 ± 2&lt;sup&gt;b&lt;/sup&gt;</td>
<td>10 ± 2&lt;sup&gt;b&lt;/sup&gt;</td>
</tr>
<tr>
<td>Total seed number</td>
<td>91 ± 9&lt;sup&gt;b&lt;/sup&gt;</td>
<td>82 ± 6&lt;sup&gt;bc&lt;/sup&gt;</td>
</tr>
<tr>
<td>Ripe seed (mg)</td>
<td>43.9 ± 6.7&lt;sup&gt;b&lt;/sup&gt;</td>
<td>43.9 ± 7.4&lt;sup&gt;b&lt;/sup&gt;</td>
</tr>
<tr>
<td>Immat. seed (mg)</td>
<td>49.8 ± 5.2&lt;sup&gt;b&lt;/sup&gt;</td>
<td>51.8 ± 6.0&lt;sup&gt;b&lt;/sup&gt;</td>
</tr>
<tr>
<td>Male flower (mg)</td>
<td>26.8 ± 3.3&lt;sup&gt;b&lt;/sup&gt;</td>
<td>32.2 ± 3.4&lt;sup&gt;b&lt;/sup&gt;</td>
</tr>
<tr>
<td><strong>Root traits</strong></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Length per vol. (km/m&lt;sup&gt;2&lt;/sup&gt;)</td>
<td>54.9 ± 5.1&lt;sup&gt;c&lt;/sup&gt;</td>
<td>62.1 ± 6.5&lt;sup&gt;b&lt;/sup&gt;</td>
</tr>
<tr>
<td>Root surface area (cm&lt;sup&gt;2&lt;/sup&gt;)</td>
<td>92.5 ± 8.6&lt;sup&gt;c&lt;/sup&gt;</td>
<td>113.6 ± 13.0&lt;sup&gt;c&lt;/sup&gt;</td>
</tr>
<tr>
<td>Fine root volume (mm&lt;sup&gt;3&lt;/sup&gt;)</td>
<td>9.1 ± 1.4&lt;sup&gt;a&lt;/sup&gt;</td>
<td>8.1 ± 1.1&lt;sup&gt;a&lt;/sup&gt;</td>
</tr>
<tr>
<td>Fine root length (m)</td>
<td>5.2 ± 0.9&lt;sup&gt;a&lt;/sup&gt;</td>
<td>4.3 ± 0.7&lt;sup&gt;a&lt;/sup&gt;</td>
</tr>
<tr>
<td>Coarse root volume (mm&lt;sup&gt;3&lt;/sup&gt;)</td>
<td>284.4 ± 30.8&lt;sup&gt;b&lt;/sup&gt;</td>
<td>399.4 ± 57.8&lt;sup&gt;b&lt;/sup&gt;</td>
</tr>
<tr>
<td>Coarse root length (m)</td>
<td>11.2 ± 1.1&lt;sup&gt;c&lt;/sup&gt;</td>
<td>14.3 ± 1.5&lt;sup&gt;bc&lt;/sup&gt;</td>
</tr>
<tr>
<td>Root diameter (mm)</td>
<td>0.180 ± 0.005&lt;sup&gt;b&lt;/sup&gt;</td>
<td>0.192 ± 0.004&lt;sup&gt;b&lt;/sup&gt;</td>
</tr>
</tbody>
</table>
Supplemental Table A.1.2 continued.

<table>
<thead>
<tr>
<th>Variables</th>
<th>roadside soil</th>
<th>foreign soil inoc.</th>
<th>non-myc. control</th>
<th>cornfield soil</th>
<th>foreign soil inoc.</th>
<th>non-myc. control</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>local soil inoc.</td>
<td></td>
<td></td>
<td></td>
<td>local soil inoc.</td>
<td></td>
</tr>
<tr>
<td>Total colonization (%)</td>
<td>79 ± 3&lt;sup&gt;a&lt;/sup&gt;</td>
<td>76 ± 2&lt;sup&gt;a&lt;/sup&gt;</td>
<td>0&lt;sup&gt;c&lt;/sup&gt;</td>
<td>45 ± 3&lt;sup&gt;b&lt;/sup&gt;</td>
<td>49 ± 4&lt;sup&gt;b&lt;/sup&gt;</td>
<td>0&lt;sup&gt;c&lt;/sup&gt;</td>
</tr>
<tr>
<td>AMF hyphae (%)</td>
<td>21 ± 4&lt;sup&gt;c&lt;/sup&gt;</td>
<td>72 ± 2&lt;sup&gt;a&lt;/sup&gt;</td>
<td>0&lt;sup&gt;d&lt;/sup&gt;</td>
<td>40 ± 3&lt;sup&gt;b&lt;/sup&gt;</td>
<td>22 ± 3&lt;sup&gt;c&lt;/sup&gt;</td>
<td>0&lt;sup&gt;d&lt;/sup&gt;</td>
</tr>
<tr>
<td>AMF arbuscules (%)</td>
<td>6 ± 1&lt;sup&gt;c&lt;/sup&gt;</td>
<td>28 ± 3&lt;sup&gt;a&lt;/sup&gt;</td>
<td>0&lt;sup&gt;d&lt;/sup&gt;</td>
<td>19 ± 2&lt;sup&gt;b&lt;/sup&gt;</td>
<td>7 ± 2&lt;sup&gt;c&lt;/sup&gt;</td>
<td>0&lt;sup&gt;d&lt;/sup&gt;</td>
</tr>
<tr>
<td>AMF vesicles (%)</td>
<td>2 ± 1&lt;sup&gt;b&lt;/sup&gt;</td>
<td>16 ± 2&lt;sup&gt;a&lt;/sup&gt;</td>
<td>0&lt;sup&gt;b&lt;/sup&gt;</td>
<td>2 ± 1&lt;sup&gt;b&lt;/sup&gt;</td>
<td>0.7 ± 0&lt;sup&gt;b&lt;/sup&gt;</td>
<td>0&lt;sup&gt;b&lt;/sup&gt;</td>
</tr>
<tr>
<td>FE hyphae (%)</td>
<td>69 ± 5&lt;sup&gt;a&lt;/sup&gt;</td>
<td>8 ± 2&lt;sup&gt;c&lt;/sup&gt;</td>
<td>0&lt;sup&gt;d&lt;/sup&gt;</td>
<td>5 ± 3&lt;sup&gt;cd&lt;/sup&gt;</td>
<td>32 ± 5&lt;sup&gt;b&lt;/sup&gt;</td>
<td>0&lt;sup&gt;d&lt;/sup&gt;</td>
</tr>
<tr>
<td>FE arbuscules (%)</td>
<td>45 ± 4&lt;sup&gt;a&lt;/sup&gt;</td>
<td>5 ± 2&lt;sup&gt;c&lt;/sup&gt;</td>
<td>0&lt;sup&gt;c&lt;/sup&gt;</td>
<td>2 ± 1&lt;sup&gt;c&lt;/sup&gt;</td>
<td>22 ± 4&lt;sup&gt;b&lt;/sup&gt;</td>
<td>0&lt;sup&gt;c&lt;/sup&gt;</td>
</tr>
<tr>
<td>FE vesicles (%)</td>
<td>25 ± 3&lt;sup&gt;a&lt;/sup&gt;</td>
<td>3 ± 1&lt;sup&gt;c&lt;/sup&gt;</td>
<td>0&lt;sup&gt;c&lt;/sup&gt;</td>
<td>1 ± 1&lt;sup&gt;c&lt;/sup&gt;</td>
<td>9 ± 2&lt;sup&gt;b&lt;/sup&gt;</td>
<td>0&lt;sup&gt;c&lt;/sup&gt;</td>
</tr>
</tbody>
</table>

Local soil inoc., local soil inoculum; foreign soil inoc., foreign soil inoculum; non-myc. control, non-mycorrhizal control; Reproduction 5 weeks, number of seeds produced after five weeks; Ripe seed, weight of ripe seeds; Immature seed, weight of immature seeds; Male flower, weight of male inflorescences; Root diameter, average root diameter; Total colonization, total AM fungal root colonization; AMF, coarse AM fungi; FE, Fine endophytes.

Different lower case letters indicate significant differences (P < 0.05) among treatment groups within a row according to Tukey’s HSD test.
**Supplemental Table A.1.3** Biomass variables and root traits in response to soil and plant origin of *Ambrosia artemisiifolia* in the experiment.

<table>
<thead>
<tr>
<th>Variables</th>
<th>Roadside Soil (local plant origin)</th>
<th>Roadside Soil (foreign plant origin)</th>
<th>Cornfield Soil (local plant origin)</th>
<th>Cornfield Soil (foreign plant origin)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Shoot (mg)</td>
<td>96.3 ± 6.0^b^</td>
<td>95.2 ± 5.0^b^</td>
<td>339.3 ± 11.9^a^</td>
<td>330.3 ± 14.8^a^</td>
</tr>
<tr>
<td>Root (mg)</td>
<td>104.6 ± 9.1^b^</td>
<td>98.2 ± 8.9^b^</td>
<td>311.9 ± 17.8^a^</td>
<td>349.8 ± 32.8^a^</td>
</tr>
<tr>
<td>Reproduction 5 weeks</td>
<td>9 ± 1^b^</td>
<td>10 ± 2^b^</td>
<td>33 ± 6^a^</td>
<td>35 ± 6^a^</td>
</tr>
<tr>
<td>Total seed number</td>
<td>82 ± 5^b^</td>
<td>75 ± 7^b^</td>
<td>181 ± 12^a^</td>
<td>193 ± 12^a^</td>
</tr>
<tr>
<td>Ripe seed (mg)</td>
<td>47.1 ± 5.5^b^</td>
<td>41.6 ± 5.5^b^</td>
<td>89.0 ± 11.5^a^</td>
<td>100.6 ± 13.6^a^</td>
</tr>
<tr>
<td>Immat. seed (mg)</td>
<td>47.6 ± 4.1^b^</td>
<td>42.0 ± 5.3^b^</td>
<td>96.0 ± 6.8^a^</td>
<td>108.5 ± 9.5^a^</td>
</tr>
<tr>
<td>Male flower (mg)</td>
<td>32.7 ± 2.7^c^</td>
<td>25.0 ± 2.7^c^</td>
<td>89.3 ± 6.0^a^</td>
<td>66.0 ± 4.1^b^</td>
</tr>
<tr>
<td>Length per volume (km/m^3)</td>
<td>53.2 ± 4.1^b^</td>
<td>56.7 ± 7.2^b^</td>
<td>90.6 ± 3.9^a^</td>
<td>106.3 ± 5.1^a^</td>
</tr>
<tr>
<td>Root surface area (cm^2)</td>
<td>87.9 ± 7.5^b^</td>
<td>101.9 ± 13.5^b^</td>
<td>216.8 ± 10.5^a^</td>
<td>252.9 ± 21.1^a^</td>
</tr>
<tr>
<td>Fine root volume (mm^3)</td>
<td>9.2 ± 0.9^a^</td>
<td>7.9 ± 1.3^a^</td>
<td>8.7 ± 0.7^a^</td>
<td>13.2 ± 0.7^a^</td>
</tr>
<tr>
<td>Fine root length (m)</td>
<td>5.3 ± 0.6^a^</td>
<td>4.3 ± 0.8^a^</td>
<td>4.8 ± 0.4^a^</td>
<td>7.8 ± 0.5^a^</td>
</tr>
<tr>
<td>Coarse root volume (mm^3)</td>
<td>265.0 ± 28.3^b^</td>
<td>352.0 ± 53.7^b^</td>
<td>1286.2 ± 93.6^a^</td>
<td>1790.5 ± 347.6^a^</td>
</tr>
<tr>
<td>Coarse root length (m)</td>
<td>10.6 ± 1.0^b^</td>
<td>12.6 ± 1.5^b^</td>
<td>22.3 ± 1.0^a^</td>
<td>23.8 ± 1.5^a^</td>
</tr>
<tr>
<td>Average root diameter (mm)</td>
<td>0.173 ± 0.005^b^</td>
<td>0.189 ± 0.003^b^</td>
<td>0.253 ± 0.005^a^</td>
<td>0.247 ± 0.010^a^</td>
</tr>
</tbody>
</table>

Reproduction 5 weeks, number of seeds produced after five weeks; Length per volume, root length per volume of soil.

Different lower case letters indicate significant differences (P < 0.05) among treatment groups according to Tukey’s HSD test.
### APPENDIX C

**Supplemental Tables A.II.1–A.II.5 to Chapter 3**

**Supplemental Table A.II.1** Variables of the target–challenger experiment in response to relative density of *A. artemisiifolia* (grown as target vs. challenger), neighboring plant species and soil treatment (AM fungi vs. non-mycorrhizal control). Values represent mean ± SE.

<table>
<thead>
<tr>
<th>Neighbor</th>
<th><em>C. canadensis</em></th>
<th><em>D. carota</em></th>
<th><em>A. vulgaris</em></th>
<th><em>T. vulgare</em></th>
</tr>
</thead>
<tbody>
<tr>
<td>Soil treatment</td>
<td>AM fungi</td>
<td>non-myc.</td>
<td>AM fungi</td>
<td>non-myc.</td>
</tr>
<tr>
<td>A. artemisiifolia grown as target</td>
<td>Shoot biomass of</td>
<td>5.601 ± 0.318</td>
<td>4.345 ± 0.264</td>
<td>2.076 ± 0.190</td>
</tr>
<tr>
<td>Shoot biomass of neighboring species (g)</td>
<td>0.191 ± 0.025</td>
<td>0.138 ± 0.013</td>
<td>0.546 ± 0.057</td>
<td>0.400 ± 0.042</td>
</tr>
<tr>
<td>A. artemisiifolia</td>
<td>0.143 ± 0.132</td>
<td>0.429 ± 0.397</td>
<td>1.000 ± 0.670</td>
<td>0.286 ± 0.171</td>
</tr>
</tbody>
</table>
Supplemental Table A.II.1 continued.

<table>
<thead>
<tr>
<th>Neighbor</th>
<th>Conyza canadensis</th>
<th>Daucus carota</th>
<th>Artemisia vulgaris</th>
<th>Tanacetum vulgare</th>
</tr>
</thead>
<tbody>
<tr>
<td>Soil treatment</td>
<td>AM fungi</td>
<td>non-myc.</td>
<td>AM fungi</td>
<td>non-myc.</td>
</tr>
<tr>
<td><strong>A. artemisiifolia</strong> grown as challenger</td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Shoot biomass of</td>
<td>1.189</td>
<td>1.101</td>
<td>1.163</td>
<td>1.179</td>
</tr>
<tr>
<td><strong>A. artemisiifolia</strong> (g)</td>
<td>± 0.048</td>
<td>± 0.042</td>
<td>± 0.051</td>
<td>± 0.057</td>
</tr>
<tr>
<td>Shoot biomass of neighboring species (g)</td>
<td>0.073</td>
<td>0.072</td>
<td>0.159</td>
<td>0.204</td>
</tr>
<tr>
<td>± 0.013</td>
<td>± 0.007</td>
<td>± 0.013</td>
<td>± 0.017</td>
<td>± 0.033</td>
</tr>
<tr>
<td>Number of male flowers of</td>
<td>2.893</td>
<td>2.536</td>
<td>2.75</td>
<td>3.714</td>
</tr>
<tr>
<td><strong>A. artemisiifolia</strong></td>
<td>± 0.394</td>
<td>± 0.305</td>
<td>± 0.295</td>
<td>± 0.572</td>
</tr>
<tr>
<td>Number of female flowers</td>
<td>1.643</td>
<td>2.429</td>
<td>1.679</td>
<td>2.5</td>
</tr>
<tr>
<td>of A. artemisiifolia</td>
<td>± 0.391</td>
<td>± 0.683</td>
<td>± 0.562</td>
<td>± 0.539</td>
</tr>
</tbody>
</table>

**Supplemental Table A.II.2** Response variables of the target–challenger experiment indicating a significant effect of neighboring species. Within a row, different lower case letters indicate significant differences ($P < 0.05$) among treatment groups according to Tukey’s HSD test.
**Supplemental Table A.II.3** Shoot biomass (mean ± SE) of *A. artemisiifolia* in response to its relative density (i.e. grown as target or challenger), and neighboring plant species tested in the target–challenger experiment. Different lower case letters indicate significant differences (*P* < 0.05) among treatment groups according to Tukey’s HSD test.

<table>
<thead>
<tr>
<th>Shoot biomass of <em>A. artemisiifolia</em> (g) grown as</th>
<th>target</th>
<th>challenger</th>
</tr>
</thead>
<tbody>
<tr>
<td>In presence of <em>Conyza canadensis</em></td>
<td>4.973 ± 0.266&lt;sup&gt;a&lt;/sup&gt;</td>
<td>1.145 ± 0.034&lt;sup&gt;c&lt;/sup&gt;</td>
</tr>
<tr>
<td>In presence of <em>Daucus carota</em></td>
<td>2.556 ± 0.219&lt;sup&gt;b&lt;/sup&gt;</td>
<td>1.171 ± 0.038&lt;sup&gt;c&lt;/sup&gt;</td>
</tr>
<tr>
<td>In presence of <em>Artemisia vulgaris</em></td>
<td>2.694 ± 0.190&lt;sup&gt;b&lt;/sup&gt;</td>
<td>1.180 ± 0.046&lt;sup&gt;c&lt;/sup&gt;</td>
</tr>
<tr>
<td>In presence of <em>Tanacetum vulgare</em></td>
<td>2.512 ± 0.255&lt;sup&gt;b&lt;/sup&gt;</td>
<td>1.137 ± 0.049&lt;sup&gt;c&lt;/sup&gt;</td>
</tr>
</tbody>
</table>

**Supplemental Table A.II.4** Shoot biomass of *A. artemisiifolia* (mean ± SE) in response to neighboring plant species tested and soil treatment (AM fungi vs. non-mycorrhizal control) in the target–challenger experiment. Different lower case letters indicate significant differences (*P* < 0.05) among treatment groups according to Tukey’s HSD test.

<table>
<thead>
<tr>
<th>Shoot biomass of <em>A. artemisiifolia</em> (g)</th>
<th></th>
</tr>
</thead>
<tbody>
<tr>
<td>Soil treatment</td>
<td>AM fungi</td>
</tr>
<tr>
<td>In presence of <em>Conyza canadensis</em></td>
<td>3.395 ± 0.611&lt;sup&gt;a&lt;/sup&gt;</td>
</tr>
<tr>
<td>In presence of <em>Daucus carota</em></td>
<td>1.619 ± 0.157&lt;sup&gt;c&lt;/sup&gt;</td>
</tr>
<tr>
<td>In presence of <em>Artemisia vulgaris</em></td>
<td>1.914 ± 0.247&lt;sup&gt;bc&lt;/sup&gt;</td>
</tr>
<tr>
<td>In presence of <em>Tanacetum vulgare</em></td>
<td>2.080 ± 0.287&lt;sup&gt;bc&lt;/sup&gt;</td>
</tr>
</tbody>
</table>

**Supplemental Table A.II.5** Percentages colonization by AM fungal structures (hyphae, arbuscules and vesicles) in roots of *Ambrosia artemisiifolia* and *Daucus carota* in the pairwise competition experiment with situations of intra- and interspecific competition.

<table>
<thead>
<tr>
<th>A. artemisiifolia</th>
<th>D. carota</th>
</tr>
</thead>
<tbody>
<tr>
<td>Competition</td>
<td></td>
</tr>
<tr>
<td></td>
<td>intraspecific</td>
</tr>
<tr>
<td>hyphae (%)</td>
<td>23.2 ± 8.9</td>
</tr>
<tr>
<td>arbuscules (%)</td>
<td>15.0 ± 5.8</td>
</tr>
<tr>
<td>vesicles (%)</td>
<td>0.6 ± 0.4</td>
</tr>
</tbody>
</table>
APPENDIX D

Supplemental Tables A.III.1–A.III.2 to Chapter 4

Supplemental Table A.III.1 Mixed effect model analysis on percentages of AM fungal structures (hyphae, arbuscules, vesicles) and percentage root colonization by non-AM fungi in soils inoculated with trained soil. In the model plant genotype was treated as random effect. Values in bold indicate significance at \( P < 0.05 \).

<table>
<thead>
<tr>
<th>Factors</th>
<th>Root colonization by</th>
<th>AM hyphae</th>
<th>AM arbuscules</th>
<th>AM vesicles</th>
<th>non-AM fungi</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td></td>
<td>( P )</td>
<td>( P )</td>
<td>( P )</td>
<td>( P )</td>
</tr>
<tr>
<td>Soil</td>
<td></td>
<td>0.222</td>
<td>(&lt;0.001)</td>
<td>0.3419</td>
<td>1.000</td>
</tr>
<tr>
<td>Seed.hist</td>
<td></td>
<td>0.083</td>
<td>0.121</td>
<td>0.7520</td>
<td>0.362</td>
</tr>
<tr>
<td>Seed.type</td>
<td></td>
<td>0.127</td>
<td>0.515</td>
<td>0.8702</td>
<td>1.000</td>
</tr>
<tr>
<td>Soil x Seed.hist</td>
<td></td>
<td>1.000</td>
<td>0.685</td>
<td>0.5998</td>
<td>0.520</td>
</tr>
<tr>
<td>Soil x Seed.type</td>
<td></td>
<td>0.462</td>
<td>0.386</td>
<td>0.9217</td>
<td>0.308</td>
</tr>
<tr>
<td>Seed.hist x Seed.type</td>
<td></td>
<td>0.416</td>
<td>0.581</td>
<td>0.3786</td>
<td>0.505</td>
</tr>
<tr>
<td>Soil x Seed.hist x Seed.type</td>
<td></td>
<td>0.080</td>
<td>0.210</td>
<td>0.5742</td>
<td>0.832</td>
</tr>
</tbody>
</table>

Seed.hist, seed history; Seed.test, seed type tested.
**Supplemental Table A.III.2** Analyses of variance (ANOVA) on seed weight of seeds produced by *Galinsoga parviflora* during the second training round of the experiment. Values in bold indicate significance at $P < 0.05$.

<table>
<thead>
<tr>
<th>Factors</th>
<th>d.f.</th>
<th>F</th>
<th>$P$</th>
</tr>
</thead>
<tbody>
<tr>
<td>Soil</td>
<td>1</td>
<td>3.053</td>
<td>0.0939</td>
</tr>
<tr>
<td>Seed.hist</td>
<td>1</td>
<td>0.933</td>
<td>0.3441</td>
</tr>
<tr>
<td>Seed.type</td>
<td>1</td>
<td><strong>38.272</strong></td>
<td>&lt;0.001</td>
</tr>
<tr>
<td>Soil x Seed.hist</td>
<td>1</td>
<td>0.148</td>
<td>0.7043</td>
</tr>
<tr>
<td>Soil x Seed.type</td>
<td>1</td>
<td>3.861</td>
<td>0.0616</td>
</tr>
<tr>
<td>Seed.hist x Seed.type</td>
<td>1</td>
<td>0.227</td>
<td>0.6381</td>
</tr>
<tr>
<td>Soil x Seed.hist x Seed.type</td>
<td>1</td>
<td>2.133</td>
<td>0.1577</td>
</tr>
</tbody>
</table>

Residuals                   | 23   |       |        |

d.f., degree of freedom; Seed.hist, seed history; Seed.test, seed type tested.