Traits linked with species invasiveness and community invasibility vary with time, stage and indicator of invasion in a long-term grassland experiment.

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Author contributions: JAC conceived the idea for the paper; DT established the experiment; ATC, JAC, MK, DT & PW surveyed the experiment; JAC, PW, JC-B & PR collected trait data; JAC & ALS designed the analytical approach; ALS undertook the modeling; JAC wrote the first draft of the paper with contributions from ALS; all authors helped write and edit subsequent drafts.

Data availability: Data sets are archived as experiment ‘e093’ and can be obtained from: http://www.cbs.umn.edu/explore/field-stations/cedarcreek/research/data. Trait data, and detailed methods about their collection, are available through TRY (Dataset ID 354, owned by the authors): https://www.try-db.org.

Running head: Traits linked with invasion vary with context

Article type: Letter

Number of words in abstract: 155

Number of words in main text: 5063
Number of references: 60

Number of figures: 4

Number of tables: 2

Keywords: invasion stages, functional traits, ecosystem invasibility, species invasiveness, indicators of invasion, seed addition experiment, Cedar Creek Ecosystem Science Reserve, long-term grassland experiment
Abstract

Much uncertainty remains about traits linked with successful invasion – the establishment and spread of non-resident species into existing communities. Using a 20-year experiment, where 50 non-resident (but mostly native) grassland plant species were sown into savannah plots, we ask how traits linked with invasion depend on invasion stage (establishment, spread), indicator of invasion success (occupancy, relative abundance), time, environmental conditions, propagule rain, and traits of invaders and invaded communities. Trait data for 164 taxa showed that invader occupancy was primarily associated with traits of invaders, traits of recipient communities, and invader-community interactions. Invader abundance was more strongly associated with community traits (e.g. proportion legume) and trait differences between invaders and the most similar resident species. Annuals and invaders with high specific leaf area were only successful early in stand development, whereas invaders with conservative carbon capture strategies persisted long-term. Our results indicate that invasion is context-dependent and long-term experiments are required to comprehensively understand invasions.
Introduction

There has been a considerable research effort to identify the characteristics of successful invaders – non-resident species that can establish, become abundant and spread in communities into which they are introduced. Much uncertainty remains about traits linked with invasion. Seed mass, for example, has been found to be positively, negatively and not linked at all with plant invasion (Pyšek & Richardson 2007; Catford et al. 2016). Acknowledging that different traits are linked with different processes of invasion (Dawson et al. 2009; Blackburn et al. 2011), recent work has highlighted the importance of considering both the stage of invasion (e.g. colonisation, survival, naturalisation, dominance; Dawson et al. 2009; Moravcová et al. 2015; Divíšek et al. 2018) and the type of invasion (e.g. wide geographic range vs high local abundance; Catford et al. 2016). These findings have led to widespread acknowledgement that the broader context of invasion – e.g. propagule pressure, time since invasion, resource availability, environmental conditions, biotic interactions, and characteristics of recipient communities (Kempel et al. 2013; Carboni et al. 2016; Conti et al. 2018) – likely influences the traits that are associated with successful invaders.

Such context dependence of relationships between traits and invasion success should not come as a surprise: diversity and multi-species coexistence are commonly thought to rely on interspecific tradeoffs where no one life history strategy is optimal under all circumstances, allowing many species to coexist (Tilman 1990; Chesson 2000). A wide diversity of life history strategies and trait syndromes should therefore be viable across sites and systems (Tilman 2011; Díaz et al. 2016). This diversity does not mean that attempting to predict invader occupancy and abundance is futile; rather, accurate prediction requires that the context of invasion be adequately considered.
We used a 20-year seed addition experiment to ask:

1) How are seven traits – lifespan, growth form, N-fixing ability, specific leaf area (SLA), leaf dry matter content (LMDC), plant height, seed mass – associated with invasion of non-resident species?

2) Do trait-based relationships vary with time, invasion stage (establishment in seeded areas, spread to unseeded areas) and indicator of invasion success (invader occupancy, invader relative abundance when present)?

3) Do traits of invaders, traits of recipient communities (means weighted by abundance), or invader-community trait-based interactions best explain the occupancy and abundance of invaders in seeded and unseeded areas?

Seeds of 50 grassland plant species, hereafter “invaders”, were sown into grassland openings in an established temperate oak savannah at Cedar Creek Ecosystem Science Reserve in Minnesota, USA (Tilman 1997). The ~2 ha savannah has been maintained through controlled spring burns in roughly every two years out of three for the last 50 years, has been free of livestock for >70 years, and has never been ploughed or cultivated. The majority of our invaders were rare or absent in plots to which they were added. Only 4 of the 50 invaders are non-native to Minnesota, reducing the relevance of issues around the transfer of species across biogeographical boundaries (e.g. enemy release; Fridley & Sax 2014; Buckley & Catford 2016). Although most invasion studies focus on exotic species, native species invasions can provide insights into the process of invasion and importance of context (e.g. Tilman 1997; Davis & Pelsor 2001; Mwangi et al. 2007). We used trait data, collected locally, to characterize 164 resident and invader taxa that were observed in the 20-year
experiment. The seven study traits were selected based on hypothesised relationships with invasion success (see Appendix S1 for rationale).

Methods

Study design

Cedar Creek (45.4°N, 93.2°W) has nitrogen-limited sandy soils, annual precipitation of ~770 mm, mean summer temperatures of 27°C, and winter lows of -14°C. In August 1991, 30 2 m x 2 m permanently marked plots were haphazardly positioned throughout the experimental savannah (Fig. S1; 45.397° N, 93.181° W; Tilman 1997). Each plot was divided into four 1 m x 1 m subplots, and percent cover of all vascular plant species, bare mineral soil, and litter were estimated (see Appendix S3 for additional details about all methods).

In June 1992, ten seed treatments were randomly assigned to two subplots per plot (hereafter “seeded subplots”, cf. “unseeded subplots”, Fig. S1). These involved the addition of no species (a control), 5, 10, 15, 20, 25, 30, 35, 40, and 54 species (Fig. S1). There were six replicates per treatment, with combinations of species in each replicate randomly drawn from a pool of 54 total species. 4.5 g/m² seeds of each species were used. Seeds were gathered on site or obtained from a local company (Prairie Restorations, Minnesota).

After 1991, subplots were surveyed eight times using the same method as in 1991. 164 plant taxa were recorded across all subplots and years, of which 85 were observed in 1991, prior to seeding (Table S1, Fig. S2). 54 non-sown taxa colonised the subplots naturally after 1991, but high turnover meant that some of these gains were offset by losses (total plant richness varied from 84 species in 1992 to 113 species in 2012).

Our analytical approach accounted for the changing recipient community by updating
the community each timestep, such that community changes would not have affected our findings.

Twenty-seven of the sown species were present in at least one plot in 1991. To account for this, sown species were considered target invaders only if they were absent from a given 2 m x 2 m plot in 1991, before seed addition. We define the recipient community as all species present in a given subplot at a given point in time (whether sown or not), excluding the target invader. Two sown invaders that were never observed (Symphyotrichum novae-angliae and Mimulus ringens) were excluded from analysis. We treated the two sown woody species (Celastrus scandens and Rosa arkansana) as members of recipient communities because our focus was on invasion of herbaceous species. Of the 50 resulting target invaders, 46 are native to MN, 43 are perennial, 12 are grasses (Poaceae), 10 are leguminous forbs (Fabaceae) and 28 are non-leguminous forbs (Table S1).

Four soil cores (20 cm deep, 2.5 cm diameter) were collected in each plot in 1991, extracted with 0.01 M KCl, and analyzed to give extractable soil nitrogen (NO$_3$) concentration and soil moisture (Tilman 1997). Though only for a single year, these variables were used to indicate local environmental conditions: they corresponded with site topography, which ranged from drier, elevated, nutrient-poor soils, to low-lying, occasionally saturated soils that were comparatively nitrogen rich (correlation between elevation at 5 m x 5 m resolution and: soil moisture, $r = -0.570$; soil NO$_3$, $r = 0.462$).

**Plant traits**
We classified all taxa based on lifespan (annual and biennial OR perennial), growth form (grass OR forb) and ability to fix nitrogen (legume OR non-legume) (Tilman 1997; USDA-NRCS 2018; Table S1).

In the summers of 2014 and 2015, we measured maximum vegetative height, leaf dry matter content (LDMC) and specific leaf area (SLA) of 143 taxa growing naturally at Cedar Creek following Pérez-Harguindeguy et al. (2013). Dried seed masses were determined from plants at Cedar Creek or from local suppliers (Prairie Moon, Prairie Restorations). These measurements provided numerical trait data for 96.3% of the vegetation cover in all subplots across all years. Of 570 species-trait measurements used in the study, 531 were measured on plants <1 km of the study site; 30 used means of congeneric or confamilial species at Cedar Creek (Table S2); and 9 were from Kew Seed Information Database (Liu et al. 2008).

We examined five forms of each individual numerical trait (Fig. 1), which we hypothesised would relate to occupancy and abundance in different ways (Appendix S1):

i. **Invader** mean trait value indicates the intrinsic invasiveness of invaders (van Kleunen et al. 2010; Richardson & Pyšek 2012).

ii. **Community-weighted mean** (CWM) trait value, calculated as mean trait value weighted by proportional cover, indicates recipient community’s intrinsic invasibility (Richardson & Pyšek 2012).

iii. **Invader-community hierarchical distance** (Invader - Community) indicates differences between invader and CWM traits. Depending on context, higher (e.g. height) or lower (e.g. SLA in resource-limited plots) hierarchical distance
can indicate when invaders have higher fitness than recipient communities (Lai et al. 2015).

iv. Invader-community dissimilarity, or absolute trait distance ($|\text{Invader} - \text{Community}|$), indicates net differences between invader and CWM traits (similar to Lai et al. 2015). This may relate to the extent of environmental sorting (trait underdispersion) or niche differentiation (trait overdispersion).

v. Invader-community gap size, calculated as the difference between trait values of resident species that fall immediately above and below invaders’ trait values ($\bar{a} < i < \bar{b}$; gap size = $\bar{b} - \bar{a}$; Fig. 1), indicates potential degree of niche differentiation between invaders and their nearest “functional neighbours” in recipient communities (similar to Gallien et al. 2014).

**Statistical analyses**

Four response variables of invasion

We used hierarchical linear regression models to identify variables associated with invasion of non-resident species (Jamil et al. 2012; Pollock et al. 2012). We examined two indicators of invasion success (occupancy and abundance) at both of two stages of invasion (establishment and spread), giving four response variables in total. All analyses relate to standing live vegetation.

*Occupancy* was the presence or absence of a target invader in seeded subplots (for establishment) or unseeded subplots (for spread).

*Relative abundance* was the proportion of total vascular plant cover in seeded subplots (for establishment) or unseeded subplots (for spread) made up of the target taxon when it was present (Catford et al. 2012).
Establishment (sensu Blackburn et al. 2011) encompasses recruitment and/or survival and/or reproduction in the site of introduction. Establishment was defined as a target invader’s occupancy or abundance when present in the subplots into which they were sown, calculated for each survey year (Fig. S1b). Establishment therefore reflects both successful invasion, and the dynamics of establishment and persistence, because an invader could be present, absent, and present again in subsequent years.

Spread (sensu Blackburn et al. 2011) encompasses dispersal and establishment beyond the site of introduction – albeit only over a short distance in our study. Spread was defined as a target invader’s occupancy or abundance when present in unseeded subplots that were adjoining seeded subplots in which invaders had established (Fig. S1b). As such, spread includes both vegetative and seed-mediated expansion of target invaders. Because the spatial arrangement of the subplots within plots meant invaders in unseeded subplots could have arrived from either of the seeded subplots, spread was analysed at the plot level (30 plots), combining data from both unseeded subplots (Fig. S1).

Regression model structure

In all models: Species traits and year were included as fixed effects (Table 1, Appendix S2). For each numerical trait (plant height, LDMC, SLA, seed mass), we used each of the five trait forms as regressors (labelled i to v in Plant traits, above, Fig. 1). For each categorical trait (lifespan, growth form, legume), we ran models that included i) invader trait category only, ii) proportional cover of that trait category in recipient community only, and iii) interactions between invader and proportional cover in community. We included target invaders and plot (for spread) or subplot nested within plot (for establishment) as random intercept terms to account for similar responses within invader species and for repeated measurements, respectively. By
including year as a fixed effect, we explicitly modelled temporal trends in trait
terms. The random intercepts accounted for repeated measurements within plots
and subplots (Pinheiro & Bates 2000).

In establishment models for both occupancy and abundance: we included the total
number of other invader species sown into each subplot to account for potential
interactions among invaders and their seed rain. We represented environmental
conditions with variables indicating soil moisture (range = 2.49–16.46%) and soil
nitrate concentration (range = 0.013–1.85 mg kg\(^{-1}\)), both of which were measured in
1991 only. We allowed the slope of the NO3 term to vary (as a random effect) by
species, as this is a primary limiting resource at our site, and species-level responses
to NO3 are known to vary greatly (Tilman 1997). We did not fit fixed effect estimates
by species because we lacked sufficient power.

In spread models for both occupancy and abundance: we used cumulative abundance
of target invaders in seeded subplots from each survey year and all previous surveys
to indicate potential cumulative seed rain into unseeded subplots. Preliminary
analyses indicated that environmental variables were not significantly related to
spread, so we did not consider them further.

Probability of occupancy was modelled using binomial generalized linear mixed
models in the lme4 package (Bates et al. 2013) in R version 3.5 (R Core Team 2014).
Abundance data were zero-inflated, so we modelled abundance conditional on
presence (i.e. removed zeros) and used mixed-effects beta regression in the
glmmADMB R package (Fournier et al. 2012; Skaug et al. 2013). All models used a
logit link function. Apart from year, which was coded as an integer varying from 0
(1992) to 20 (2012), all non-binary numeric variables were centered and scaled
(Appendix S6). For details on models and convergence criteria, see Appendices S2, S3, S4 and S6.

**Model selection**

To identify the “best” regression forms, we conducted model selection in two steps. In Step 1, we compiled a series of models that varied in structure and complexity (Appendix S2). We did not mix trait forms in Step 1, so we had five subsets of models – one for each trait form. We used the structure of the top-ranked model for each response variable from Step 1 as our template for models analysed in Step 2. Into that template, we sequentially substituted different trait forms that had important effects in Step 1. We considered trait forms important if they were significantly related (P<0.05) to response variables in the top-ranked model from each of the five model subsets analysed in Step 1. The models in Step 2 could potentially include different forms for different traits (e.g. SLA hierarchical distance and CWM seed mass), but only one form of each trait was used in each model (e.g. CWM SLA was never used with invader SLA); this was because different trait forms were highly correlated and so we could identify trait forms that best explained invasion, thereby addressing Question 3.

We ranked all models using Bayesian Information Criterion (BIC, Jones 2011). We considered all models that differed in BIC from the best-supported model by ≤ 5 to have support from the data (Hegyi & Garamszegi 2011). We present results of the best-supported models in the main paper and all models with ΔBIC ≤ 5 in Fig. S3. We considered effects of explanatory variables to be significant if 95% confidence intervals did not cross zero in at least one of the models with ΔBIC ≤ 5. We calculated marginal and conditional R² values for the occupancy models to indicate variance explained respectively by fixed and random effects, and by fixed effects only.
We could not calculate $R^2$ values for the abundance models because there is currently no available method for mixed-effect beta regression (Nakagawa, pers. comm, Nov 2017; Nakagawa & Schielzeth 2016).

**Results**

**General trends**

Relative abundance of invaders increased with time in seeded subplots and in unseeded subplots (Figs 2m, 3h). Invaders that established and accumulated high abundance in seeded subplots over the duration of the experiment were highly likely to spread to, and reach high abundance in, adjacent unseeded subplots (Figs 3a, 3i).

The number of sown invaders was not related to establishment of target invaders, suggesting that the presence of other invaders in seeded subplots did not strongly affect invasion of individual species in our experiment.

**Question 1: Traits linked with invasion success**

Invader N-fixing ability, life span and growth form were linked with invader occupancy but not abundance for both establishment (seeded subplots) and spread (unseeded sublots; Table 2; Fig. 4). Legumes had higher occupancy than non-legumes when directly sown (Fig. 2l). By the end of the experiment, perennial and grass invaders had higher occupancy than annuals and forbs in both seeded and unseeded subplots (Figs 2j, 2k, 3c, 3g).

Recipient communities with higher perennial cover had higher invader occupancy and abundance in seeded subplots (Figs 2d & 2o), but the opposite was true for invader abundance in unseeded subplots (Fig. 3l). Invader abundance was higher in seeded
subplots with low proportional legume cover (Fig. 2p), but – in long-term – the
opposite was true in unseeded subplots (Fig. 3m).

Seed mass and LDMC had the most consistent long-term invasion trends, relating to
three of our four response variables (Table 2). Apart from the first few years of
invasion, invader occupancy and – in seeded subplots – invader abundance was
highest in communities with heavier seeds (Figs 2h, 2r, 3e) and higher LDMC (Figs
2i, 2s, 3f), whether in absolute terms (i.e. based on CWM) or relative to invaders (i.e.
based on hierarchical trait distances). SLA and plant height were associated with
invasion, but trends varied over time and depending on invasion stage and indicator
(discussed below).

**Question 2: Trait-based relationships varied with time, invasion stage and indicator
of invasion success**

Factors related to invasion were not uniform across invasion stages or indicators of
invasion success, and were not uniform over time (Figs 2, 3 & 4, Table 2, Appendix
S3).

**Time:** Relationships between invader occupancy in seeded subplots and invader
lifespan, invader growth form and community LDMC varied over time (Fig. 2i-k). In
unseeded subplots, invaders with higher SLA than recipient communities initially had
higher occupancy, and invader abundance was initially higher where communities had
higher SLA CWM; these trends reversed with time (Figs 3d, 3j, S2).

**Stage:** Invaders with dissimilar heights to recipient communities (Fig. 2c) or their
nearest functional neighbours (Fig. 2q) were more likely to occupy and reach high
abundance in seeded subplots respectively, but the opposite was true in unseeded
subplots (except for short-term abundance; Figs 3b, 3k). Invaders were more abundant
in seeded subplots that had low legume cover (Fig. 2p), but – from 10 years onwards
– the opposite trend was true in unseeded subplots (Fig. 3m). Invader occupancy and
abundance in seeded subplots were positively related to community perennial cover
(Figs 2d, 2o) – in contrast to long-term abundance trends in unseeded subplots (Fig.
3l). Invader occupancy was higher when invaders had lower SLA values than
recipient communities in seeded (Fig. 2b) and unseeded (long-term only, Fig. 3d)
subplots, but invader abundance was higher in unseeded subplots with low SLA
communities (long-term only, Fig. 3j).

Indicator: Invader occupancy was lower but abundance higher in seeded subplots
with higher initial soil NO₃ (Figs 2a, 2n).

**Question 3: Relative importance of invader traits, community traits, and invader-
community trait-based interactions**

Based on our best-supported models, invader occupancy was related to four of the
five trait forms (all but gap size, Fig. 1) – albeit in different ways and via different
numerical traits (Figs 2, 3 & 4, Appendix S4), indicating that invader traits,
community traits and invader-community interactions all influence invader
occupancy. In contrast, invader abundance was related to community traits (CWM)
and gap size; intrinsic invader traits were unrelated to invader abundance in any of
best-supported (or competing) models (Figs 2, 3 & S3, Appendix S4).

**Model performance**

The best-supported models for occupancy in seeded and unseeded subplots accounted
for about 63% and 59% of observed variance respectively (conditional R²), and their
fixed effects accounted for 15% and 44% (marginal R², Appendix S3). Results of the
best model and competing models for each response variable had consistent
ecological interpretations (Fig. S3, Appendix S4); the only exception was LDMC in the establishment-abundance model, suggesting that LDMC is not strongly linked with invader abundance in seeded subplots (Fig. S3b).

Discussion

Our 20-year experiment showed that factors associated with invasion of non-resident species are not uniform across time, stages, or indicators of invasion (Table 2, Fig. 4). Despite the variation in observed relationships, our study accords with existing theories that propagule pressure, environmental conditions, invader and community traits and their interactions, time, and indicator and stage of invasion all collectively affect invasion trends (Catford et al. 2009; Blackburn et al. 2011; Richardson & Pyšek 2012). All seven traits that we examined were related to at least one indicator or stage of invasion, but the strength, direction and consistency of their relationships varied. Invader occupancy was related to four trait forms, while invader abundance was mostly associated with CWM plus invader differentiation from functionally similar residents. We discuss the main findings to our three research questions after briefly discussing general trends.

Invaders’ cumulative abundance in seeded subplots was positively associated with their greater occupancy and relative abundance in adjacent unseeded subplots. This was presumably because of higher propagule rain (or clonal spread) and higher suitability of the local ecological conditions for population growth. Given that abundances of invader species increased over 20 years in both seeded and unseeded areas, it not only seems likely that some invaders have formed self-sustaining populations in vicinities where sown, but also that their local population sizes will continue to increase – at least until density-dependent effects become stronger.
**Question 1: Traits associated with invasion**

Legumes had higher occupancy than non-legumes where directly sown (Fig. 2l), as expected (Appendix S1), but N-fixing ability of invaders was unrelated to their abundance or spread. This was a surprising result given the study system is nitrogen-limited (Tilman 1997), but previous work has shown that deer herbivory can severely constrain the abundance of legumes in the study savannah (Knops *et al.* 2000).

Invaders that spread into unseeded subplots were more abundant in the long-term when recipient communities had high proportional cover of legumes and low perennial cover, likely because of greater N availability in these communities – consistent with findings elsewhere (Mwangi *et al.* 2007). Importantly, relationships between invasion and community legume and perennial cover varied, perhaps reflecting temporal variation in the extent of light, N and moisture limitation (Mwangi *et al.* 2007; Willis *et al.* 2010).

We were surprised that grass-, perennial- or legume-dominated communities did not have lower abundances of invading grasses, perennials or legumes respectively. Studies at Cedar Creek (Symstad 2000; Fargione *et al.* 2003) and elsewhere (Emery 2007; Mwangi *et al.* 2007) have found that invader abundance was greatest when invaders were functionally distinct from invaded communities. Biotic resistance from limiting similarity is not universal though (Price & Pärtel 2013). For example, competition experiments revealed that *Phragmites australis* invasion was more strongly inhibited by annual species rather than by species similar to *P. australis* (Byun *et al.* 2013). Our findings may differ from other studies at Cedar Creek because our study used more detailed trait-based information and did not rely so heavily on functional group classifications, and because it occurred in a more heterogenous
environment where effects of environmental sorting would be stronger (Kraft et al. 2015).

Higher long-term occupancy of invaders that were grasses, perennials or had lower SLA than recipient communities (Figs 2k,j,b, 3c,d,g) may reflect the longer lifespans of plants with these characteristics (Wright et al. 2004; Lauenroth & Adler 2008), and the greater ability of grasses (cf forbs, though there are exceptions) to spread clonally (Lauenroth & Adler 2008). Despite the apparent logic of these results, the lifespan and SLA results nevertheless seem to contradict common findings where annual species and those with fast leaf economics (indicated by high SLA) are typically more successful as invaders (van Kleunen et al. 2010). We provide two explanations for this. First, most seed addition experiments are of much shorter durations. Had we run the experiment for five and not 20 years, and had we only examined species occupancy, we too would have concluded that annuals and invaders with higher relative SLA posed a higher invasion risk than perennials or low SLA species. However, our full set of results show that such conclusions are true only under certain conditions. Second, in observational studies of invasions, which largely focus on exotic species, invading species are typically found in modified ecosystems (Catford et al. 2011; Moles et al. 2012) or disturbed sites characterized by high resource availability; under these conditions, the importance of resource conservation and longevity are secondary to high growth rates and short generation times (Kuster et al. 2008; Moravcová et al. 2015). In our study savannah, resource availability was low and disturbance matched historical forms and levels. It is thus fitting that longer-lived invaders and those with a more conservative carbon capture strategy persisted longer in the experimental plots. The potential benefits of being non-native (Buckley & Catford 2016; Catford et al. 2018) may also explain why our results – based on invasion of mostly native, non-resident species – differ from observational studies of
exotic invasions.

Long-term invader occupancy and, in seeded subplots, abundance were greater in communities that had seed masses and LDMCs that were high or higher than invaders (Figs 2h-i, 2r-s, 3e-f). If we ignore invader abundance in unseeded subplots, which was unrelated to seed mass and LDMC, these results suggest that communities dominated by species with heavier seeds and longer-lived, tougher leaves are more invasible in the long-term than communities with lighter seeds and low LDMC.

Reflecting the well-established tradeoff between seed mass and number (Moles 2018), communities dominated by species with fewer, heavier seeds may be more seed-limited than lighter-seeded communities (Tilman 1997; Turnbull et al. 2000). As such, heavier-seeded communities may thus have greater numbers of unoccupied safe sites, increasing the relative recruitment success of invaders in these communities (Duncan et al. 2009; Wandrag et al. under revision). Because a set mass (rather than number) of seeds was added to seeded subplots, the seed mass-number tradeoff remained intact in our experiment, such that light-seeded invaders would have had more seeds than heavier seeded species, and thus more opportunities to recruit (Duncan et al. 2009; Wandrag et al. under revision). The relative advantage of lighter, more numerous seeds was apparent for occupancy (where hierarchical distance was significant), but not abundance, the latter of which was mostly strongly related to seed mass CWM. That invader abundance was only linked with seed mass in seeded (cf unseeded) subplots may stem from the high rate of seed addition to seeded subplots (4.5 g/m² per species): the high seeding rate may have enabled invaders to occupy most, if not all, available safe sites, whereas the unseeded subplots were presumably still seed-limited (Wandrag et al. under revision).
We hypothesised that invaders with slower leaf economics than recipient communities (higher LDMC, lower SLA) would be more competitive and thus more abundant than their opposite counterparts in the long-term – reflecting resource economics (Reich 2014), and consistent with long-standing ideas of $K$-selected species replacing $r$-selected ones (Pianka 1970). Trends related to SLA were consistent with our hypotheses, but trends related to LDMC were not (Appendix S1). The entire study savannah is burnt every two in three years such that the occurrence of fire does not explain invasion trends. However, communities with higher LDMC may be more flammable (Pérez-Harguindeguy et al. 2013) and thus experience hotter, more damaging fires, potentially making them more vulnerable to invasion by increasing resource availability and opportunities for colonisation (Davis & Pelsor 2001).

When seeded, invaders were most successful when their heights were distinct from the CWM and their nearest neighbours in recipient communities (Figs 2c, 2q). These trends suggest that height differentiation from residents facilitates invader establishment (Fig. 1, Herben & Goldberg 2014), but the cause is unclear; competition for light alone would presumably favour invasion by species that are taller than (indicated by hierarchical distance), rather than distinct from, the recipient community (Willis et al. 2010). The opposite trends were observed in unseeded subplots (Figs 3b, 3k) – possibly reflecting the greater importance of environmental sorting or other selective pressures (which would select functionally similar cf. novel invaders; Kraft et al. 2015; Divíšek et al. 2018) for invader spread. Trait data for 53 species in the study savannah indicate that there is no correlation between clonality and relative plant height ($r = 0.11$, Cleland et al. 2008), so clonal spread does not explain these height-based trends.
Question 2: Trait-based relationships vary with time, stage and indicator of invasion

Our 20-year experiment showed that factors associated with invasion success are not uniform across time, stages, or indicators of invasion (Table 2, Fig. 4). Conditions that were favourable for some stages and indicators of invasion, and at some times, were disadvantageous or had negligible effects at others. These results highlight that how, where and when invasion is measured can affect ecological inferences. Our findings indicate that invasion events observed during the first few years after seed addition only tell part of the invasion story; comprehensive understanding of invasion requires long-term studies in natural, intact ecosystems.

Recent studies have shown that traits linked with invasion depend on invasion stage (Dawson et al. 2009; Moravcová et al. 2015; Conti et al. 2018; Divišek et al. 2018), recipient community characteristics (Lai et al. 2015; Carboni et al. 2016; Conti et al. 2018), and the way invasion is measured (Carboni et al. 2016). However, we believe that our study is among the first to reveal that invasion trait-based trends also vary with time (see Kempel et al. 2013): even when examining a single indicator of invasion, traits of invaders or characteristics of recipient communities positively associated with early invasion could be negatively associated with later invasion.

Question 3: Relative importance of invader traits, community traits, and invader-community trait-based interactions differ for invader occupancy and abundance

Invader occupancy was associated with all five trait forms we examined except gap size – more or less in line with our expectations (Appendix S1). In contrast, invader abundance was most strongly linked with recipient community traits (CWM) and differentiation from invaders’ nearest functional neighbours (gap size; Table 2, Fig.
Unlike occupancy, species that reach high relative abundance must be able to acquire a high share of available resources. Species can do this by outcompeting other species for limited resources (Tilman 1982), occupying different niches (Chesson 2000), or becoming abundant in sites with high levels of resource availability (Davis et al. 2000). It is therefore logical that community invasibility and invader-community interactions – both linked with biotic resistance (Conti et al. 2018) – strongly influence invader abundance, whereas intrinsic invader traits are of less importance.

By examining occupancy and relative abundance of 50 non-resident plant species in seeded and unseeded areas over 20 years, and characterising 164 taxa with seven traits, our study highlights the complexity and context dependence of biological invasions. Invasion success varied with time, invasion stage and indicator of success, and was influenced by traits of invaders, traits of recipient communities, and trait-based interactions between the two.

Acknowledgements: We thank Troy Mielke for coordinating research at Cedar Creek, Mara Sagedahl and Christine Carroll for help with trait data collection, Cedar Creek interns for assisting with vegetation surveys, and Wade Blanchard for statistical advice. Comments from three reviewers helped us improve the paper. We acknowledge funding from the Australian Research Council (DE120102221) and the ARC Centre of Excellence for Environmental Decisions (to JAC), and the US National Science Foundation Long-Term Ecological Research Program, including DEB-0620652 and DEB-1234162. Cedar Creek Ecosystem Science Reserve and the University of Minnesota provided further support.
References


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57.


Supporting Information

Appendix S1: Hypotheses and rationale for selection of seven study traits and use of multiple trait forms [SI text document].

Appendix S2: Structure of the candidate models for all four response variables (invader occupancy and abundance in seeded and unseeded subplots) examined in Step 1 and Step 2 of the model selection procedure, and description of all variables [excel file].

Appendix S3: Additional details about statistical analyses [SI text document].

Appendix S4: Model selection results and top model coefficients for all four response variables (invader occupancy and abundance in seeded and unseeded subplots) [excel file].

Appendix S5: Pairwise correlations for all terms used in the models. [excel file].

Appendix S6: R code used in data processing and analysis:
https://github.com/annabellisa/Catford_E93 [github repository].

Table S1: Characteristics of all 164 taxa observed in the study, including functional groups, frequency of occurrence and mean trait values. Descriptions of data are provided within the file [excel file].

Table S2: Source of numerical trait data for the 164 taxa observed in this study [SI text document].

Figure S1: Map of oak savannah study site showing location of experimental plots. (a) Locations of the 30 2 m x 2 m experimental plots; (b) number of sown invaders (see colour legend) added to seeded subplots (NW and SE corners of the plots) in
1992 and number of invaders still present in seeded subplots in 2012 (integers next to subplots). The establishment models used data from 54 of the 120 subplots into which seed were added. The spread models used combined data from the two unseeded control subplots within each of the 30 plots (i.e. 2 unseeded subplots were summed and analysed as a single data point, giving 30 plots) [SI text document].

**Figure S2** Number of taxa observed across the 20-year experiment [SI text document].

**Figure S3** Coefficients for all models with a ΔBIC of <5 for a) establishment probability, b) establishment abundance, c) spread probability and d) spread abundance [SI text document].

**Figure S4** Mean standardised cover of functional groups over time. “Other grass” includes grasses identified at the genus level that could not be classified as C3 or C4 [SI text document].
Table 1: Main explanatory variables considered (= “yes”) in occupancy and abundance models for invader establishment in seeded subplots and invader spread into unseeded subplots. These variables were available for selection during the model selection process; *variable included in spread-occupancy models but not spread-abundance models. A full list of specific variables is provided in Appendix S2.

<table>
<thead>
<tr>
<th>Variable</th>
<th>Description</th>
<th>Variable type</th>
<th>Establishment</th>
<th>Spread</th>
</tr>
</thead>
<tbody>
<tr>
<td>Year</td>
<td>Survey year</td>
<td>temporal</td>
<td>yes</td>
<td>yes</td>
</tr>
<tr>
<td>No_seeded</td>
<td>Number target invader species added to given subplot</td>
<td>propagule pressure</td>
<td>yes</td>
<td>no</td>
</tr>
<tr>
<td>Seeded subplot</td>
<td>Seeded subplot cumulative abundance</td>
<td>propagule pressure</td>
<td>no</td>
<td>yes</td>
</tr>
<tr>
<td>Moisture</td>
<td>Soil moisture in 1991</td>
<td>environmental</td>
<td>yes</td>
<td>no</td>
</tr>
<tr>
<td>Soil NO3</td>
<td>Soil NO3 concentration in 1991</td>
<td>environmental</td>
<td>yes</td>
<td>no</td>
</tr>
<tr>
<td>SLA</td>
<td>Absolute or relative specific leaf area of invader and/or community</td>
<td>numerical trait (5 forms)</td>
<td>yes</td>
<td>yes</td>
</tr>
<tr>
<td>Height</td>
<td>Absolute or relative height of invader and/or community</td>
<td>numerical trait (5 forms)</td>
<td>yes</td>
<td>yes</td>
</tr>
<tr>
<td>Seed</td>
<td>Absolute or relative seed mass of invader and/or community</td>
<td>numerical trait (5 forms)</td>
<td>yes</td>
<td>yes</td>
</tr>
<tr>
<td>LDMC</td>
<td>Absolute or relative leaf dry matter content of invader and/or community</td>
<td>numerical trait (5 forms)</td>
<td>yes</td>
<td>yes</td>
</tr>
<tr>
<td>Prop_peren</td>
<td>Proportion of community cover that is perennial</td>
<td>categorical trait</td>
<td>yes</td>
<td>yes</td>
</tr>
<tr>
<td>Prop_grass</td>
<td>Proportion of community cover that is grass</td>
<td>categorical trait</td>
<td>yes</td>
<td>yes*</td>
</tr>
<tr>
<td>Prop_legume</td>
<td>Proportion of community cover that is legume</td>
<td>categorical trait</td>
<td>yes</td>
<td>yes</td>
</tr>
<tr>
<td>Lifespan</td>
<td>Lifespan of invader (annual &amp; biennial OR perennial)</td>
<td>categorical trait</td>
<td>yes</td>
<td>yes</td>
</tr>
<tr>
<td>Grass</td>
<td>Growth form of invader (grass OR forb)</td>
<td>categorical trait</td>
<td>yes</td>
<td>yes*</td>
</tr>
<tr>
<td>Legume</td>
<td>N-fixing ability of invader (legume OR non-legume)</td>
<td>categorical trait</td>
<td>yes</td>
<td>yes</td>
</tr>
</tbody>
</table>
Table 2: Observed conditions under which the occupancy and abundance of target invaders in seeded and unseeded subplots are **highest** in the short-term (<5 years) and long-term (>10 years). For example, occupancy in seeded subplots was highest when soil NO$_3$ was low in both the short- and long-term, and when invaders were forbs (short-term) or grasses (long-term). I indicates non-resident (target) invader; C indicates recipient community; I = C indicates that successful invaders are similar to community for that given trait (trait underdispersion), whereas I ≠ C indicates that successful invaders are dissimilar to community for that trait (trait overdispersion); I > C and I < C indicate cases where successful invader has higher and lower trait values than the recipient community respectively; neighbours refers to invader’s two nearest functional neighbours in the recipient community, i.e. trait gap size (Fig. 1). Hyphens indicate no statistically significant relationship; NA indicates that the relationship was not formally tested. Functional group similarity refers to interactions between invaders and the recipient communities based on species’ growth form, lifespan and N-fixing ability.

<table>
<thead>
<tr>
<th>Invasion stage</th>
<th>Establishment in seeded subplots highest when…</th>
<th>Spread to unseeded subplots highest when…</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>Occupancy</td>
<td>Abundance</td>
</tr>
<tr>
<td>Short- or long-term trends</td>
<td>Short</td>
<td>Long</td>
</tr>
<tr>
<td>Soil NO$_3$ in plot in 1991</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Invader growth form</td>
<td>Low</td>
<td>Low</td>
</tr>
<tr>
<td>Invader lifespan</td>
<td>Forb</td>
<td>Grass</td>
</tr>
<tr>
<td>Proportion perennial in community</td>
<td>Annual</td>
<td>Perennial</td>
</tr>
<tr>
<td>Invader N-fixing ability</td>
<td>Forb</td>
<td>Grass</td>
</tr>
<tr>
<td>Proportion legume in community</td>
<td>High</td>
<td>High</td>
</tr>
<tr>
<td>Functional group similarity$^\text{a}$</td>
<td>I = C</td>
<td>I = C</td>
</tr>
<tr>
<td>Plant height</td>
<td>I ≠ C</td>
<td>I ≠ C</td>
</tr>
<tr>
<td>LDMC</td>
<td>C = Low</td>
<td>C = High</td>
</tr>
<tr>
<td>SLA</td>
<td>C &gt; I</td>
<td>C &gt; I</td>
</tr>
<tr>
<td>Seed mass</td>
<td>-</td>
<td>C &gt; I</td>
</tr>
<tr>
<td>Invader abundance in seeded subplots</td>
<td>NA</td>
<td>NA</td>
</tr>
<tr>
<td>Trends through time</td>
<td>Conditional</td>
<td>Increased with time</td>
</tr>
</tbody>
</table>
**Figure 1** Sketch showing the five trait forms linked with our hypotheses of invasion success that relate to invader traits, community traits, and invader-community trait-based interactions (see Appendix S1 for rationale of predictions). The solid blue line shows the hypothetical distribution of trait values for the recipient community, the dashed blue lines indicate trait values for two species in the recipient community (Residents a & b), and the pink line shows the trait distribution of a target invader. i) Invader trait indicates the “intrinsic invasiveness” of the target invader, represented by mean trait value across individuals of the target invader; ii) Community-weighted mean (CWM) trait value indicates recipient community’s “intrinsic invasibility”, calculated as mean trait value weighted by proportional cover; iii) Invader-community hierarchical trait distance indicates whether invaders had a higher or lower trait value than the CWM of recipient communities, and the magnitude of this difference ($iii = i - ii$); iv) Invader-community dissimilarity (absolute trait distance) indicates how different the invader is to the community regardless of the direction of that difference ($iv = |i - ii|$); v) Invader-community trait gap size indicates the degree of niche differentiation between the invader and its nearest “functional neighbours”, calculated as the difference between the trait values of the resident species that fall immediately above and below the invader’s trait value in the recipient community ($v = \tilde{b} - \tilde{a}$).
Figure 2 The establishment stage of invasion, modelled as occupancy and abundance of 50 target invaders, sown into seeded subplots. Plots show results for the best supported model for invader: a-l probability of occupancy (top panels); and m-s) proportional abundance, conditional on presence (bottom panels); variables that were significantly related to invader occupancy and abundance are displayed. Each observation in the data represented the response of an individual invader in a given year, in a given subplot, and observations were grouped by species with a random effect. Effects of explanatory variables are shown holding all other explanatory variables in each of the two models constant, such that each response can be interpreted independently of other variables in that model. When a numeric variable interacted with another numeric variable, the response was displayed for the minimum and maximum values in the data.
Figure 3 The spread stage of invasion, modelled as occupancy and abundance of 50 target invaders, which spread to unseeded subplots. Plots show results for the best supported model for invader: a-g) probability of occupancy (top panels); and h-m) proportional abundance, conditional on presence (bottom panels); variables that were significantly related to invader occupancy and abundance are displayed. Each observation in the data represented the response of an individual invader in a given year, in a given subplot and observations were grouped by species with a random effect. Effects of explanatory variables are shown holding all other explanatory variables in each of the two models constant, such that each response can be interpreted independently of other variables in that model. When a numeric variable interacted with another numeric variable, the response was displayed for: d-f) the minimum and maximum values in the data; and j-m) the first and third quartile in the data. The values at which to plot interactions were chosen based on the distribution of the data. The y-axis is displayed with a different range on panel (i).
Figure 4 Summary of overall results showing conditions and traits associated with A) occupancy and abundance of sown, target invaders that established in seeded subplots, and B) occupancy and abundance of established invaders that spread from seeded subplots to adjacent unseeded subplots. Short-term refers to trends within the first 5 years of the experiment; long-term refers to trends in the last 10 years of the experiment.