

# Consequences of plant functional traits and plant-microbe interactions on species coexistence

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## Abstract:

How do interactions between organisms and their environment influence the dynamics of ecological communities? Ecologists have long understood that species performance can be impacted by both their response to the abiotic variation and by their interactions with other organisms. Despite this long history, we still lack a clear understanding of how the ecological strategies of species mediate their responses to environmental variation, whether individual variation within species can change the outcome of competition, and how multi-trophic interactions contribute to the structure of natural communities. To address this, in my dissertation I aim to experimentally investigate how species coexistence is impacted by plant ecological strategies (as summarized by functional traits) and species-specific interactions with soil microorganisms in an annual plant community in coastal California. To do so, I will apply recent advances in theoretical ecology to plant community ecology and integrate the complementary perspectives on coexistence from the fields of plant competition and plant-microbe interactions.

A longstanding assumption has been that the coexistence of species with dissimilar functional traits is stabilized due to their distinct demographic responses to environmental heterogeneity. For Chapter 1 I plan to test this assumption using recent advances in coexistence theory that quantify how variation in demographic responses to spatial heterogeneity can promote coexistence. Specifically, I propose to test whether functional trait differences are predictive of the similarity in the demographic responses of species to environmental gradients.

Another striking observation from nature is that individuals within a species can vary substantially in their functional traits. Despite considerable interest, the drivers of this variation among individuals ('intra-specific trait variation', or ITV) and its influence on species coexistence and community dynamics remain largely unclear. For Chapter 2, I aim to investigate the drivers of ITV and ask how it might influence the outcome of competitive environments.

Finally, over the past two decades, a great deal of research has highlighted the role of soil microorganisms in shaping terrestrial plant communities. Microbes have been shown to promote plant performance in stressful conditions and have wreaked lasting ecological damage in other communities. In nature, such biotic interactions unfold concurrently with resource competition between plants; however in the literature, their effects are generally considered independently of one another. For Chapter 3, I aim to demonstrate how the effects of resource competition and plant-microbe interactions can be jointly considered under an established framework in modern coexistence theory via a critical review of existing plant-soil feedback literature.

## Background

Plant communities can house remarkable amounts of diversity. What are the fundamental processes that allow so many species to coexist with each other? Modern coexistence theory, emerging from the work of Chesson (2000), frames species coexistence as an outcome of two types of differences between species. *Fitness differences* are differences between species that confer one species a density-independent competitive advantage over the other. For example, differential tolerance of a generalist predator can cause a fitness difference: in the absence of other processes, the species that is less susceptible to the predator will outperform the more susceptible species. *Stabilizing niche differences* are differences between species that cause a species to limit its own population growth rate more than that of its competitor. Variation between species in the timing of resource use can act as stabilizing niche differences. Stabilizing niche differences must overcome any fitness differences between species to permit their coexistence (Figure 1). In other words, coexistence is promoted by species differences that minimize fitness differences and those that increase stabilizing niche differences (although fitness equality alone is insufficient for coexistence without some stabilizing niche differences) (Chesson 2000, Adler et al. 2007).

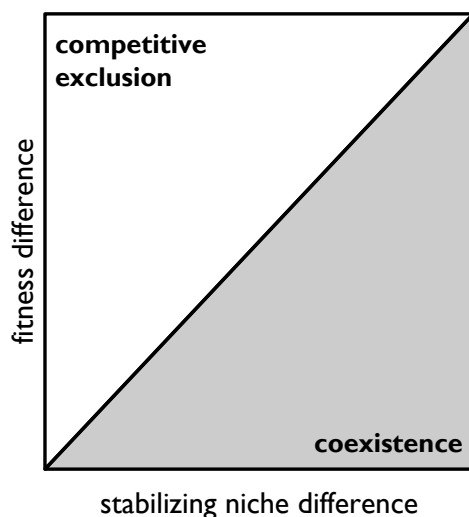


Figure 1: Coexistence is possible when stabilizing niche differences overcome fitness differences. Figure after Adler et al. (2007)

**Trait based tests of coexistence:** What determines the magnitude of fitness and niche differences between species? One way to progress towards a general answer to this question is to understand the relationship between demographic dissimilarity and differences in species functional traits, which summarize their ecological strategy (Westoby et al. 2002, McGill et al. 2006). Is coexistence more likely between species with more dissimilar ecological strategies? Or are functionally similar species pairs more likely to coexist due to minimal fitness differences? Do the same traits predict the strength of coexistence via various mechanisms? Understanding species coexistence as a result of their functional traits can help interpret patterns of trait variation in natural communities and make ecology a more predictive discipline (McGill et al. 2006).

As a first pass, ecologists extended the principle of limiting similarity (MacArthur and Levins 1967) into the modern coexistence framework to hypothesize that functional dissimilarity predicts the probability of coexistence. Recent advances in the field (Mayfield and Levine 2010, Kraft et al. 2015) caution against such simplification. For example, Kraft et al. (2015) found that in an annual plant system, species differences in fruiting phenology are well correlated with both fitness and niche differences: species with later fruiting phenology tend to outcompete those with earlier phenology, but species experience a demographic advantage when rare when in competition in species that differ in phenology.

To understand coexistence as a result of species ecological strategies, we need a better understanding of how species traits are linked to the demographic drivers that determine the outcome of competition, and

how individual variation in functional traits can affect the outcome of competitive interactions. As spatial heterogeneity is thought to promote diversity in many plant communities by introducing stabilizing niche differences (Chesson 1994), understanding the relationship between functional traits and species' demographic responses to environmental variation can provide a stronger foundation for future studies linking plant functional variation to coexistence in variable landscapes.

**The microbial revolution:** While functional traits promise to improve our understanding of how interactions among plants shape distribution and abundance patterns, most studies in this area have ignored the role that belowground interactions play in shaping plant communities. A recent revolution in ecology has highlighted the ubiquity of microbes across ecosystems and the role they may play in generating and maintaining the diversity of larger organisms (Bever et al. 2015, Fierer 2017). In fact, the colonization of terrestrial habitats by plants may itself have been catalyzed by a mutualism with arbuscular fungi (Humphreys et al. 2010). Evidence that soil microbes influence the dynamics of contemporary plant communities is mounting. For instance, Mangan et al. (2010) experimentally grew seedlings of six tropical tree species with microbes of various origins and showed plant species abundance may be driven by their susceptibility to soil microorganisms. Laliberté et al. (2015) have suggested that a tradeoff between plant phosphorus acquisition efficiency and susceptibility to pathogens may explain hyperdiversity in shrublands and tropical rainforests.

Interactions between plants and microbes play out simultaneously with the resource competitive dynamics that have been the focus of plant ecology for decades. For example, a specialist pathogen may introduce stabilizing niche differences to promote the persistence of a competitive inferior in a system with large fitness differences. However, these two mechanisms are most often studied independently of one another (van der Putten et al. 2016), and the relative importance of microbial effects on plant coexistence is still unknown. Synthesizing these mechanisms in a common theoretical framework will be key to moving towards a better understanding of their combined effects on community structure and dynamics.

In my thesis, I propose to take a holistic view of plants from a perspective of their own physiology (as summarized by functional traits), their responses to environmental variation, and their interactions with microbes to ask three questions:

**Q1:** How do differences in functional traits relate to the differences in species' demographic response to environmental variation?

**Q2:** What are the drivers of functional trait variation within species, and how might ITV affect the outcome of plant competition?

**Q3:** How can we jointly consider the effects of plant-microbe interactions and resource competition on plant species coexistence?

I will address the first two questions by performing experiments using ~20 species from the annual plant community in Sedgwick Reserve, Santa Barbara County, California. These species occur across a heterogeneous landscape with diversity in topography and edaphic properties (Gram et al. 2004). For Question 1, I have estimated the demographic response of annual plant species to environmental variation by planting seeds into 24 plots across the reserve. I plan to ask whether changes in species demography along envi-

ronmental gradients are predicted by commonly measured plant functional traits. For Question 2, I again planted seeds across the reserve to test whether ITV in functional traits is linked to species demographic variation across the landscape, and whether ITV due to competitive neighborhoods can affect the outcome of competition. Finally, I propose to address Question 3 by showing how plant-microbe interactions can be considered in light of recent advances in coexistence theory that jointly consider effects of competition and natural enemies (Chesson and Kuang 2008). I plan to conduct a critical review of the plant-soil feedback literature and to extend Chesson and Kuang (2008)'s framework to include the consequences of mutualistic interactions. My thesis will combine recent advances in coexistence theory with empirical advances in plant and microbial ecology to contribute towards a holistic and mechanistic understanding of plant diversity maintenance.

## **Chapter 1: How do functional traits relate to species responses to variable environments?**

*Additional collaborators: Emily Curd (UCLA); Andy Kleinhesselink (UCLA); Jonathan Levine (ETH-Zurich)*

A foundational principle of trait-based ecology is that traits are functional to the extent that they influence population demography by directly influencing growth, survival, and reproduction (Violle et al. 2007). As variation in demographic properties is at the heart of species coexistence (Chesson 2000), understanding the links between species traits and their demography is a critical step in using traits to understand coexistence. Although ecologists have long understood that trait differences can reduce the niche overlap between species to promote coexistence (MacArthur and Levins 1967), recent advances have shown that trait differences can also promote competitive exclusion (Adler et al. 2013, Kraft et al. 2015). In this chapter, I will focus on one of the most important drivers of species coexistence- spatial environmental heterogeneity (Chesson 1994). In particular, I will assess if traits are likely to influence species coexistence by mediating species demographic responses to spatially variable environments.

Despite being fundamental to interpreting patterns of trait variation in nature, the links between trait differences and species' demographic responses remain largely unresolved (Shipley et al. 2016, but see Adler et al. 2014 and Wright et al. 2010). Many studies have shown non-random associations between trait values and the environment (Cornwell and Ackerly 2009, Fortunel et al. 2014). However, it is difficult to interpret these results in the context of the demographic drivers of coexistence, in large part because observed species abundances may not reflect the intrinsic response of species to the environment (Adler et al. 2013). Indeed, the most abundant species at a site need not have the traits that confer the highest fitness at that site (Fox 2012). Therefore, isolating the impact of environment while keeping the competitive environment constant is a critical step in building stronger links between functional trait differences and variation in species responses to variable environments (Adler et al. 2013). Interpreting these results in light of ecological first principles regarding trait-environment relationships can give generality to our findings.

In addition to the competitive neighborhood, the interactions between plants and microbes may also change across space to influence plant performance. Variation of soil microbes can have important effects on plant responses to abiotic environmental variation. For example, Lau and Lennon (2012) found that the fruit

production of *Brassica rapa* was less impacted by a drought treatment when the plants were grown with a drought-adapted soil microbial community than when grown with a soil microbial community that had not been previously exposed to drought. Therefore, the soil microbial community may function as an additional environmental variable that to date has received relatively little attention from plant ecologists. Recent advances in sequencing technology have made it feasible to characterize variation in soil microbial communities, and we can begin to assess their role in driving plant environmental preferences.

To address these gaps, I propose to use experimental data to estimate the contribution of trait differences to species-specific demographic responses to environmental variation, including turnover in soil microbial communities, for the first chapter of my dissertation.

## Hypotheses

**H0:** Species may respond identically to environmental variation, irrespective of functional traits. We may expect this pattern if the environmental gradient spans a single stress gradient axis such that all species perform poorly at some sites and well at others (Figure 2a). In this case, observed patterns of species turnover across the landscape at Sedgwick may be driven by poor dispersal.

**H1:** Species may have idiosyncratic responses to the environmental gradient, but the response may be unrelated to functional traits. There is potential for environmental variation to promote species coexistence, but the stability of coexistence would be uncorrelated to species' functional similarity (Figure 2a).

**H2:** Variation in responses of species to environmental gradients may be explained by their functional traits. This pattern might arise if the functional traits are indicative of species' environmental preference along the gradient sampled (Figure 2a).

**H3:** Similarity in some traits may correlate with similarity in response to certain environmental variables. For example, drought tolerance may predict demographic responses to an aridity gradient but not to a light gradient, and seed size may predict responses to a soil texture gradient but not an aridity gradient (Figure 2b).

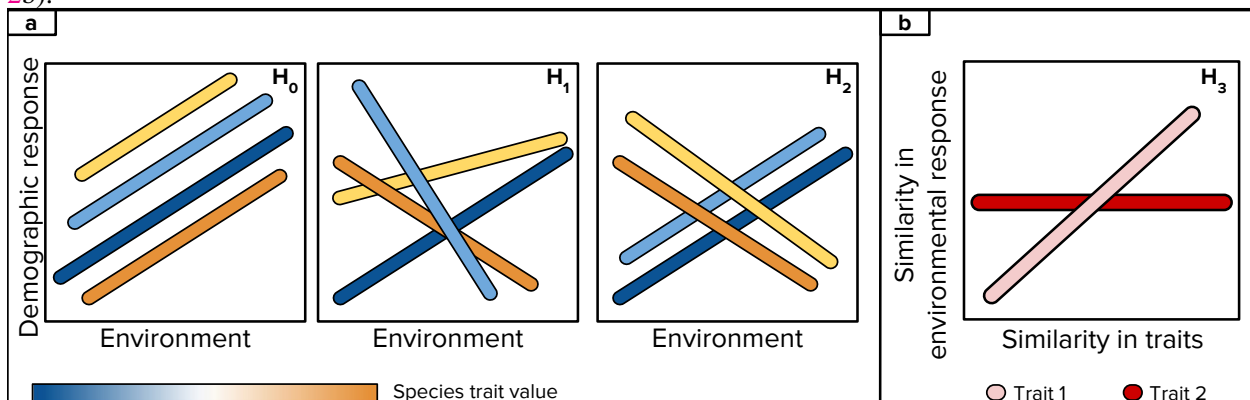


Figure 2: Potential ways in which trait differences may correlate with differences in species' demographic response.

## Methods

**Demographic variation:** In 2015-2016, we planted seeds of 17 annual plant species (Table 1) at 24 plots in a setup that allowed us to parameterize a simplified annual plant population growth model. In this model, the population size of a species  $j$  at a site  $x$  grows according to its seed bank survival rate ( $s_j$ ), its germination rate ( $g_{j,x}$ ), and its intrinsic seed production response at each site ( $\lambda_{j,x}$ ), but the rate of growth is reduced by how much competition it faces at each site ( $\eta_x$ ) and how sensitive it is to competitors ( $r_{j,x}$ ):

$$\frac{N_{t+1,j,x}}{N_{t,j,x}} = (1 - g_{j,x}) * s_j + \frac{g_{j,x} * \lambda_{j,x}}{1 + r_{j,x} * \eta_x}$$

In this framework, environmental response can be parameterized in three ways: germination rate, the per-capita seed production in the absence of competitors, and a composite index that yields the expected population growth rate in the absence of competitors.

**Trait variation:** A set of plant, seed, root, and whole-plant functional traits were measured for twelve of the seventeen species in this experiment by Kraft et al. (2015) (Table 2). I have measured the same set of traits for the five remaining species and have additionally measured leaf osmotic potential (Bartlett et al. 2012), which is a strong predictor of leaf drought tolerance.

Table 2: List of functional traits for Chapter 1 study

Trait
Leaf size
Specific Leaf Area (SLA)
Leaf Dry Matter Content (LDMC)
Osmotic Potential
Leaf Nitrogen Concentration
Seed Mass
Specific Root Length (SRL)
Rooting Depth
Max. Height
Canopy Shape Index
Fruiting Phenology
Carbon Isotope Ratio

**Abiotic environmental variation:** We characterized various environmental properties at each of our experimental sites during the 2015-16 growing season. Soil was sent to Western A&L Labs for chemical and physical analysis. Air temperature at each of the plots was measured with temperature data loggers (iButtons) that were placed inside PVC tubing to shield from solar radiation, and gravimetric water content was measured for twice for each plot - once each in the peak growing season for the early and later phenology species. In analyses, I will use both raw environmental variables and NMDS scores to characterize the environment.

**Biotic environmental variation:** We characterized the variation in soil microbiota among our plots by using metagenomic sequencing to profile the diversity of bacteria, fungi, and small eukaryotes. Soil was pooled from three equally spaced points within each experimental plot into a single sterilize 15 mL tube and stored at -80°C until further processing.

Total DNA was extracted using the MoBio Power Soil Extraction kit and PCRs were conducted using 16s rRNA primers for bacteria, fungal ITS primers, and 18s rRNA primers for small eukaryotes. Pooled samples

Table 1: Species list for demography experiment

Family	Species
Asteraceae	<i>Uropappus lindleyi</i>
	<i>Centaurea melitensis</i>
	<i>Chaenactis glabriuscula</i>
	<i>Hemizonia congesta</i>
	<i>Lasthenia californica</i>
	<i>Micropus californicus</i>
Boraginaceae	<i>Amsinckia menziesii</i>
Onagraceae	<i>Clarkia botatae</i>
	<i>Clarkia purpurea</i>
Euphorbiaceae	<i>Euphorbia spathulata</i>
Fabaceae	<i>Acmispon wrangelianus</i>
	<i>Medicago polymorpha</i>
Poaceae	<i>Bromus madritensis</i>
	<i>Hordeum murinum</i>
	<i>Vulpia microstachys</i>
Lamiaceae	<i>Salvia columbariae</i>
Plantaginaceae	<i>Plantago erecta</i>

were sequenced on the Illumina MiSeq platform at the UC-Berkeley sequencing facilities. Sequence data will be used to ordinate sites according to their soil microbial communities in multidimensional space.

**Analyses:** I propose to assess the influence of functional traits on the demographic responses of species to environmental gradients with a generalized linear mixed modeling (GLMM) framework modified from Jamil et al. (2012). In this framework, the performance of a species  $i$  across a landscape  $j$  is modeled as a function of species traits  $z$ , environmental properties  $x$ , and crucially, an interaction term between the two:

$$p_{i,j} \sim \underbrace{a_0}_{\text{species effect}} + \underbrace{a_1 z_i}_{\text{trait effect}} + \underbrace{b_0 x_j}_{\text{environment effect}} + \underbrace{b_1 z_i x_j}_{\text{interaction}} + \underbrace{\varepsilon_{\alpha i} + \varepsilon_{\beta i} + \gamma_j}_{\text{error terms}}$$

The species effect term  $a_0$  is a fixed effect that can be thought of as capturing unmeasured species traits, and the random effect term  $\gamma_j$  represents unmeasured environmental variation. The contribution of trait differences to interspecific variation in environmental response (i.e. the slope parameter  $\beta_1$ ) can be quantified as a function of the variance explained ( $\hat{\sigma}_b$ ) by a model that includes a trait:environment interaction term and one that lacks it:

$$C_\beta = 1 - \frac{\hat{\sigma}_{\beta, \text{no interaction term}}^2}{\hat{\sigma}_{\beta, \text{with interaction term}}^2}$$

### Preliminary results, implications, and further analyses

Preliminary analyses show that most of the species pairs had uncoupled demographic responses- in other words, most species pairs did not respond in the same way to the multivariate changes in environment across sites (Figure 3a), which suggests against  $\mathbf{H}_0$ . Thus, variable demographic responses of species to environmental heterogeneity may contribute to their coexistence across this landscape. The environmental variation among our sites stems primarily from differences in soil  $[\text{Ca}^{2+}]:[\text{Mg}^{2+}]$  ratio and in soil texture and depth (Figure 3b). The species in this experiment differed primarily in leaf economics spectrum traits on the first axis of variation and on whole plant traits on the second axis of variation. Preliminary analyses conducted with simple linear regressions hint at some trait differences that may explain variation in species responses to environment. For example, four out of the five smallest-seeded species in this experiment perform better at sites with higher values of soil NMDS 2 (shallower and sandier soils), whereas all other species prefer deeper soils with higher clay content.

Modeling variation in demographic response to environment as a function of traits in this framework above will address a fundamental gap of trait-based plant ecology: the assumption that functional traits have direct and predictive relationships with measurable environmental gradients (Adler et al. 2013, Shipley et al. 2016). Due to strong separation of sites along axes of soil chemistry and physical structure, we may find that root and seed traits that directly link to species' belowground ecological strategies are more relevant



to explaining variation in species response in our system than traits that primarily capture aboveground strategies. Alternatively, physiological traits with a clear link to an environmental axis (e.g. leaf osmotic potential to water availability) may show stronger links to plant performance than integrative traits like SLA that are related to various environmental axes. An important step will be to conduct a power analysis to assess the extent to which the potentially subtle but important influences of individual traits (as might be expected for some of the integrative functional traits measured for this study) can be recovered given the probabilistic nature of many demographic models (Snyder & Ellner, *in prep*). A number of studies have also found phylogenetic structure in plant communities; therefore, incorporating evolutionary non-independence of our samples by using phylogenetically informed GLMMs (Ives and Helmus 2011) is another promising next step. I plan to interpret all results in light of ecological first-principles and in the framework of modern coexistence theory to yield generality to these findings. Armed with a more robust understanding of the link between traits and demography, trait-based plant ecology can move towards its potential to be a predictive science in a rapidly changing world (McGill et al. 2006)

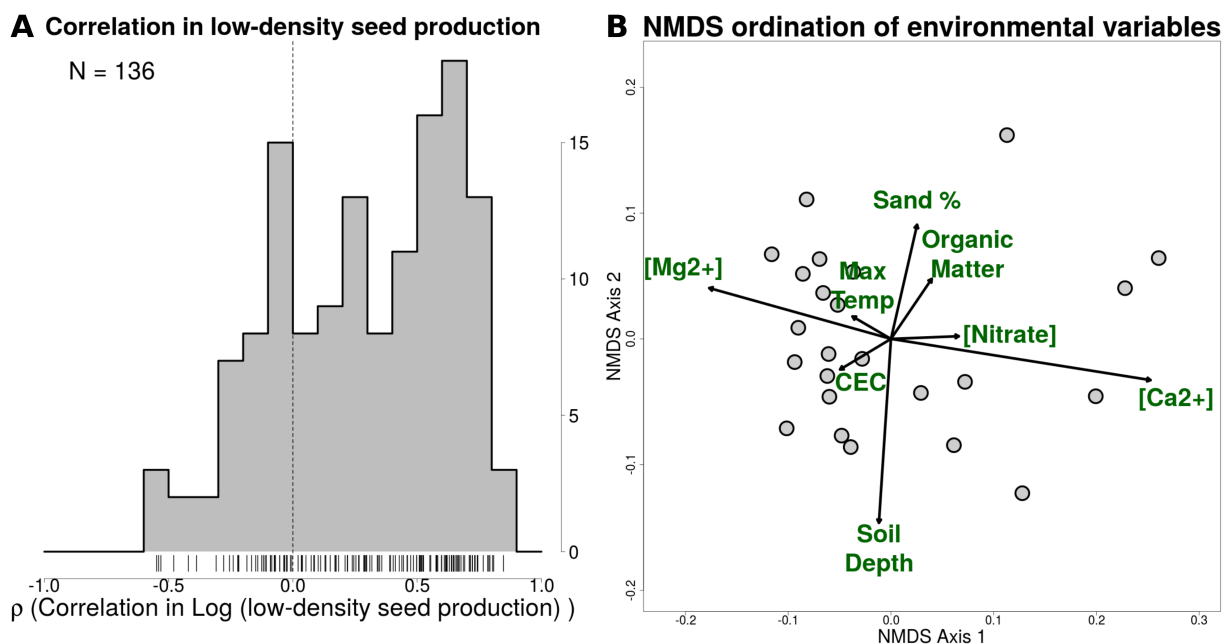


Figure 3: **A.** A histogram of how correlated species are in their environmental response shows that species pairs tend to have decoupled weakly or negatively correlated responses (all pairwise correlations in species responses <1). **B.** Soil chemical and physical properties were the primary axes of variation among sites.

## Chapter 2: The nature and consequences of intra-specific trait variation

Individuals within a species can have vastly different phenotypes- indeed, genetic variation within populations is the fundamental basis for natural selection. Intra-specific variation (ITV) in plant functional traits can be quite high: in a recent meta-analysis, Siefert et al. (2015) showed that 25% of functional trait variation within communities is driven by ITV. Variation in functional traits within species has many causes. Among other drivers, traits can vary due to genotype (Brouillette et al. 2013), ontogeny (Spasojevic et al. 2014), fine-scale environmental variation (e.g. between shaded vs. sun-exposed leaves) (Nobel 1976),



regional environmental variation (Cornwell and Ackerly 2009), and local competitive neighborhoods (Turcotte and Levine 2016).

How important is this ITV to understanding the structure of natural communities? Although community dynamics fundamentally emerge from interactions between individuals, ecologists tend to rely on species averages to study the drivers of species coexistence, leaving the question of ITV impacts on coexistence largely unexplored (Violle et al. 2012, Turcotte and Levine 2016). The ability to infer community-wide dynamics from species-mean values measured on a small number of individuals per species is at the core of the generalizability of trait-based ecology (McGill et al. 2006, Shipley et al. 2016); therefore, it is important to understand how ITV may affect these processes.

Whether ITV promotes or hinders coexistence, and whether observed levels of ITV affect our predictions of the outcome of competitions, still remains unknown. One line of reasoning suggests that variation among individuals of a species may decrease species-level fitness differences (Fridley et al. 2007) or promote stabilizing niche differences (Clark 2010) and thus make coexistence more likely. On the other hand, Hart et al. (2016) demonstrated with a theoretical model incorporating viability in demographic rates that ITV is more likely to destabilize coexistence due to non-linear averaging. However, an important limitation of this model was that it assumed a well-mixed population with no structure in ITV- in other words, the demographic traits of an individual were randomly drawn from a population distribution and were not a function of their local environment.

ITV is unlikely to be random in nature, and we need a better understanding of how abiotic and biotic environments shape ITV to better understand its potential effects on coexistence (Turcotte and Levine 2016). However, mapping the structure of ITV alone is not sufficient to understanding its influence on coexistence. An understanding of how functional traits relate to the demographic drivers of coexistence is needed to incorporate ITV into coexistence predictions. As Kraft et al. (2015) have shown how functional traits relate to fitness differences and stabilizing niche differences in the Sedgwick annual plant system, this community is an ideal one in which to ask these questions:

**Q1:** How do functional traits shift in response to spatial environmental variation?

**Q2:** Do species shift their ecological strategies depending on their competitive neighborhoods? How frequently do such shifts change the predicted outcome of competition between two species?

## **Hypotheses**

**Q1H0:** Species functional traits may not change across strict environmental gradients. In this case, observed patterns of community-wide trait turnover across gradients may be driven by species turnover or due to trait responses to competitive neighborhoods.

**Q1H1:** Functional traits may shift in the same way across species in response to environmental gradients (e.g. trait values may increase across the gradient for all species).

**Q1H2:** Traits may shift across environmental gradients, but the slope and magnitude of trait shifts may differ between species.

**Q1H2a:** Species that are less demographically sensitive to the environmental variation (i.e. species with consistently high population growth rates) may show higher ITV than species that are more sensitive to the environment.

**Q2H0:** Species functional traits may not shift in response to competitive backgrounds.

**Q2H1:** Functional traits may shift in the same way across species in response to competitive backgrounds (e.g. trait values may be higher in competition than when grown alone, irrespective of species).

**Q2H2:** Traits may shift across competitive backgrounds, but the slope and magnitude of trait shifts may differ between species. In this case, ITV might have the potential to shift the outcome of competition by influencing the magnitude of niche and fitness differences between species.

**Q2H2a:** Traits of competitive inferiors may respond more strongly than those of competitive superiors in a way that decreases fitness differences or increases niche differences (Kraft et al. 2015); alternatively, trait changes may shift in ways that reinforce the dominance of the competitive superior.

## Methods

**Trait sampling:** In 2016-2017 we planted seeds of 20 species in the same locations we had used for our demography experiment in Chapter 1. Seeds were planted 15 cm away from one another to ensure that plants were growing with a very low density of competition so that we could estimate the impact of environment alone on traits for **Q1**. Additionally, at one site, we planted all species in a high competitive density to estimate the effects of competitive neighborhoods on traits for **Q2**. We measured leaf size, SLA, and LDMC on three leaves from each individual. We additionally measured plant height and canopy shape index on the same individuals and stored root tissue in ethanol to measure specific root length (SRL). In 2017-2018, I plan to sample these same traits on individuals growing alone, with conspecific competitors, and with heterospecific competitors to test if trait response to competitors depend on the identity of the competitor species (Turcotte and Levine 2016).

**Analysis:** I propose to perform GLMM analyses similar to those being used in Chapter 1 to describe trait shifts in response to environmental gradients. Specifically, I propose model the traits of species  $i$  at site  $j$  as a function of the overall trait mean, the effect of species and the local environment, and an interaction between species and site:

$$trait_{i,j} \sim \underbrace{a}_{\text{trait mean}} + \underbrace{b_0 x_i}_{\text{species effect}} + \underbrace{c_0 z_j}_{\text{environment effect}} + \underbrace{c_1 x_i z_j}_{\text{interaction}} + \underbrace{\epsilon_{\alpha j} + \epsilon_{\beta i} + \gamma_i}_{\text{error terms}}$$

As a preliminary test of whether functional traits change with competitive background and whether such a change is species-specific (**Q2**), I performed a two-factor ANOVA testing for the influence of species identity, competition background, and their interaction on the three leaf traits (leaf area, SLA, LDMC). If species have idiosyncratic responses to competition, then trait shifts in response to competition might have potential to change competitive outcomes. Finally, I will use the trait-based models of fitness and niche

differences from Kraft et al. (2015) to test whether the predicted outcome of coexistence changes after incorporating ITV.

## Preliminary results, future analyses, and implications

In total, we measured traits on >800 individuals to date for this study, with an average of 40 individuals per species. In our preliminary analysis for **Q2**, we found species- and trait-specific consequences of competitor density on traits. Specifically, we found that individuals growing in dense competition tended to have higher SLA and larger leaves than individuals of the same species grown at low-density ( $p < .0001$ ). Competitive density had no clear effect on LDMC. There was a significant interaction between species and competitive background for all three traits, indicating species-specific responses to competition (Figure 4). Low germination of some species resulted in an unbalanced dataset; therefore, it will be important to perform further nonparametric tests (e.g. REML F-tests (Bolker et al. 2009)) to verify the robustness of these results.

These preliminary results suggest that shifts in traits in response to competitors have the potential to change the predicted outcome of competitive interactions. For example, species with more resource conservative leaf traits (e.g. low SLA, high LDMC) tend to be competitive superiors in this system (Kraft et al. 2015); therefore, idiosyncratic responses of these traits to competition may serve to either increase or decrease fitness differences. As a first test of this possibility, I will use the trait-based models of species interactions from Kraft et al. (2015) to ask whether accounting for all of the observed intraspecific variation significantly changes the magnitude of fitness and niche differences. This approach would be similar to that of Hart et al. (2016) in assuming a well-mixed community with no structure in ITV. I will then perform more targeted analyses that ask if ITV in response to competition specifically changes the predictions of coexistence.

Although functional traits can vary significantly within species, the consequences of this variation on species coexistence are unclear. By quantifying the impact of ITV in functional traits to variation in species demography, this study will begin to address an important gap between the theoretical work that focuses on ITV in demographic traits and the large body of empirical work that has documented ITV in functional traits.

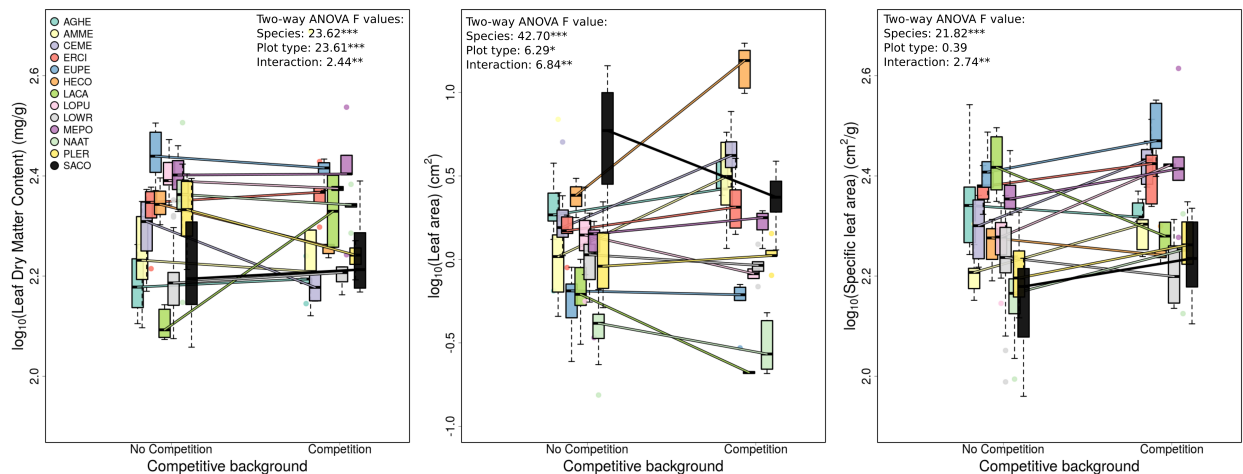


Figure 4: Two-way ANOVAs suggest that leaf functional traits have idiosyncratic responses to competitive backgrounds across species.

### Chapter 3: Considering the consequences of soil microbial communities on plant species dynamics under the modern coexistence framework

Microbes are ubiquitous in nature and can have dramatic impacts on the performance of larger organisms. Redman (2002) found that when colonized by endophytic fungi in the genus *Curvularia*, the Panic grass *Dichanthelium lanuginosum* could withstand sustained soil temperatures greater than 50°C. This interaction is critical to understanding the persistence of the grass in geothermal soils where few other plants occur. Conversely, microbes can severely harm plant populations. For instance, the oomycete *Phytophthora ramorum* caused dramatic mortality of tanoak (*Lithocarpus densiflorus*) and coast live oak (*Quercus agrifolia*) in coastal California and Oregon in the late 1990s (Rizzo et al. 2005). This continues to have profound impacts on the community and ecosystem dynamics in those systems (Cobb et al. 2013). Theoretical and empirical advances in the last two decades have highlighted that microbes can also have subtler mechanisms for effecting profound changes in plant communities. For example, van der Putten et al. (1993) suggested that plant succession in foredune communities is driven by the promotion of late-successional plant species in soils that harbor a microbial community cultivated by early-successional species. In a generalized theoretical framework that has stimulated a great deal of research, Bever et al. (1997) showed how feedback loops between plants and soil microbes could permit the coexistence of competing plant species. In short, plant-microbe interactions have the potential to dramatically influence the outcomes of plant species interactions and community assembly.

But how important are the effects of belowground plant-microbe interactions in comparison to the competitive effects that are the subject of traditional coexistence and community assembly theory? We cannot answer this question without a framework that simultaneously considers the effects of both mechanisms. For the third chapter of my dissertation, I propose to demonstrate that the effects of plant-microbe interactions can be considered jointly with the effects of resource competition under the competition-predation framework of Chesson and Kuang (2008) and critically review the empirical work conducted under the plant-soil feedback framework of Bever et al. (1997) in this new context. Fundamental to this reframing is the fact that microbes can influence plants in ways that are independent of any feedbacks (i.e. some microbial contexts always benefit or always harm plant species, regardless of plant species density) to influence fitness differences, and via feedbacks to influence stabilizing niche differences. Here I briefly review the state of PSF research, summarize Chesson and Kuang (2008)'s coexistence framework, and begin to show how the former can be integrated into the latter.

**State of plant-soil feedback literature:** Plants can alter both biotic and abiotic characteristics of the soil. Plants that later grow in those soils will be responding to the changes- this cyclical relationship is the basis of plant-soil feedbacks (PSFs). Bever et al. (1997) showed that if the effects of the soil community cultivated by a plant species *A* are more negative to the population growth rate of species *A* than the growth rate of plant species *B*, and vice-versa (i.e. more negative intra-specific effects than inter-specific effects), then PSFs can promote species coexistence. Alternatively, positive PSFs can drive competitive exclusion by preferentially promoting the performance of the abundant species. This framework has inspired many studies (a Web of Science search for titles containing "plant-soil feedback\*" yielded 171 results overall and 73 in 2015-2017

alone) which have documented a diversity of positive, negative, and neutral effects of PSFs.

Despite such extensive experimentation, it remains difficult to predict the importance of PSFs: their impacts seem to be highly species- and environment-specific (Casper and Castelli 2007, van der Putten et al. 2016). Recently, Smith-Ramesh and Reynolds (2017) predicted that PSFs are less important for coexistence in higher nutrient sites where plants can have high immunity and produce more defense compounds, whereas van der Putten et al. (2016) predict that the higher performance of pathogens in higher nutrient sites will increase the importance of negative PSFs. Additionally, most experimental work is conducted under highly controlled greenhouse settings, and it is difficult to translate these results into the importance of PSFs in nature (van der Putten et al. 2016). Finally, an important limitation of this framework from the perspective of coexistence theory is that it focuses solely on one set of processes that might promote coexistence (namely, stabilizing niche differences) but fails to incorporate potential effects of microbes on fitness differences (for example, by acting as a generalist enemy that drives fitness differences). Reframing plant-soil interactions in modern coexistence theory may highlight heretofore overlooked parameters and suggest avenues of research to start building generalities.

**Overview of competition/predation framework:** That shared natural enemies can change the outcome of strict resource competition has been long understood in ecology as “apparent competition” (Holt 1977). Chesson and Kuang (2008) integrated apparent competition into modern coexistence theory to show how natural enemies can affect coexistence in much the same way as resource competition. If natural enemies are specialists, then intra-specific feedback loops are stronger than inter-specific feedback loops and coexistence is promoted via lower niche overlap. On the other hand, physiological tradeoffs between natural enemy defense and rates of reproduction can promote coexistence via equalizing fitness differences (Chesson 2012). Starting from a system that modeled the dynamics of enemies, prey, and resources, Chesson and Kuang (2008) derived equations to quantify niche overlap ( $\rho$ ) and fitness ( $\kappa_j$ ) as a joint function of resource competition and natural enemies<sup>1</sup>:

$$\rho = \frac{\sum_l \frac{c_{jl}v_l c_{kl}}{r_l^R \alpha_l^R} + \sum_m \frac{a_{jm}w a_{km}}{r_m^P \alpha_m^P}}{\sqrt{\left( \sum_l \frac{c_{jl}^2 v_l}{r_l^R \alpha_l^R} + \sum_l \frac{a_{jm}^2 w}{r_m^P \alpha_m^P} \right) \left( \sum_l \frac{c_{kl}^2 v_l}{r_l^R \alpha_l^R} + \sum_l \frac{a_{km}^2 w}{r_m^P \alpha_m^P} \right)}}$$

$$\kappa_j = \frac{\sum_l \frac{c_{jl}v_l}{\alpha_l^R} - \sum_m \frac{a_{jm}}{\alpha_m^P} - \mu_j}{\sqrt{\left( \sum_l \frac{c_{jl}^2 v_l}{r_l^R \alpha_l^R} + \sum_m \frac{a_{jm}^2 w}{r_m^P \alpha_m^P} \right)}}$$

<sup>1</sup>The parameters are as follows:  $c_{jl}$  represents the consumption rate of resource  $l$  by species  $j$ ;  $v_l$  is the unit value of resource  $l$ ,  $\alpha_l^R$  is the reciprocal of resource carrying capacity,  $r_l^R$  is the rate or resource replacement,  $a_{jm}$  is the attack rate of enemy  $m$  on species  $j$ ,  $w$  is the unit value of the focal species to the enemy,  $\alpha_m^P$  is the reciprocal of enemy carrying capacity, and  $r_m^P$  is the enemy's intrinsic population growth rate.

All boxed terms indicate the effects of natural enemies, whereas all unboxed terms indicate the effects of resource competition (save  $\mu_j$ , which represents maintenance costs of species  $j$ ). **The key takeaway** is that in this framework, niche differences and fitness differences are both simultaneously influenced by resource competition as well as natural enemies.

### **Preliminary results, further analyses, and implications**

I have implemented the model demographic model from Chesson and Kuang (2008) as an R webapp (<https://gauravsk.shinyapps.io/ck-2008/>). Analysis of this model shows that the net impacts of plant-microbe feedbacks can be influenced by several parameters. For example, consider the consequences of adjusting two parameters in the model: **1)** the rate of resource regeneration ( $r_l^R$  and  $r_g^R$ ), and **2)** the degree to which enemy populations can grow independently of the focal species (i.e. due to predation on other species in the system) ( $r_m^P$  and  $r_n^P$ ). The relative importance of PSFs is higher in more fertile sites (higher resource replacement rates  $r^R$ ) and when microbes interact weakly with species outside of the focal group (low  $r^P$ , Table 3). The insight regarding the increased importance of PSFs in sites with high resource replacement rates is in contradiction with the prediction of Smith-Ramesh and Reynolds (2017), who suggest that PSFs should be less relevant in more fertile sites but fail to consider the **relative importance** of PSFs and resource competition. However, it will be important to analyse both the steady-state and transient dynamics of the model (e.g. time to competitive exclusion), as many empirical studies focus on the effects of varying state variables (e.g. total resources level at time  $t$ ), rather than model parameters (e.g. resource replacement rate).

I plan to continue exploring the behavior of this model and find generalities to help interpret the many contradictory ideas regarding PSFs and their role in shaping plant communities. To extend the framework, I plan to modify the model to incorporate potential mutualistic impacts of microbes on plant population growth. This can be done by setting the attack rate  $a_{jm}$  to be a negative number. After exploring the dynamics of the model, I propose to conduct a thorough review of the literature since Bever et al. (1997) and interpret the results in light of the joint competition-natural enemies model. This review will highlight areas that have received little attention but are important to understanding the role of PSFs in nature. Finally, I will use this framework to inform and conduct experimental work at the UCLA greenhouses that rigorously tests the role of PSFs relative to that of resource competition in natural systems using the annual plant species and soil from Sedgwick. For example, in 2016 I conducted a growth chamber experiment to test for whether fitness differences (as estimated by germination rate and vegetative biomass) were impacted by soil microbial context across five sites with varying soil chemistry. Such experiments can be used to test specific predictions regarding the *net effects* of resource competition and microbial dynamics on plant communities. Reframing PSFs under the modern coexistence framework will help the field meet its new frontier of understanding variation in the relative importance of PSFs in nature.

$a_{jm}$	$a_{jn}$	$a_{km}$	$a_{kn}$	$r_l$	$r_g$	$r_m$	$r_n$	$\rho_{net}$	$\rho_{competition}$	$\rho_{enemies}$	$\frac{\kappa_j}{\kappa_k}$	coexist?
0.001	0.00	0.00	0.001	2	2	0.1	0.1	1.00	1	1.00	1	no
<b>0.015</b>	0.00	0.00	<b>0.015</b>	2	2	0.1	0.1	0.95	1	0.13	1	yes
0.015	0.00	0.00	0.015	<b>0.4</b>	<b>0.4</b>	0.1	0.1	0.99	1	0.13	1	yes- but lot less stable
0.015	0.00	0.00	0.015	2	2	<b>1</b>	<b>1</b>	0.99	1	0.13	1	yes- but lot less stable

Table 3: Contribution of niche differentiation due to natural enemies ( $\rho_{enemies}$ ) to net niche differentiation ( $\rho_{net}$ ) increases with resource replacement rates (higher  $r_l$  and  $r_g$ ) and when enemies depend strongly on the focal species (lower  $r_m$  and  $r_n$ )

The values of other parameters were as follows:  $c_{jl} = .04, c_{jg} = .02, c_{kl} = .02, c_{kg} = .02, v_g = v_l = .2, u_j = u_k = .1, W = .01, \alpha_g = \alpha_l = .05, \alpha_m = \alpha_n = .1$

## Timeline

Year	Season	Plan
2017	Fall- Winter	Analysis and writeup for Chapter 1; Literature reviews for Chapter 3
2018	Spring- Summer	Writeup and submission for Chapter 1; Additional data collection for Chapter 2; Literature reviews & modeling for Chapter 3
2018	Fall- Winter	Analysis for Chapter 2, Modeling & analysis for Chapter 3
2019	Spring- Summer	Writeup and Analysis for Chapter 2, Writeup for Chapter 3
2019	Fall- Winter	Writeup for Chapter 2; Submission for Chapter 3
2020	Spring	Submission for Chapter 2



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