Article title: Intra and interspecific diversity in a tropical plant clade alter herbivory and

ecosystem resilience

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Data and code used for all analyses and figures can be accessed via the Dryad repository at

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Abstract

Declines in biodiversity generated by anthropogenic stressors at both species and

population levels can alter emergent processes instrumental to ecosystem function and resilience.

As such, understanding the role of biodiversity in ecosystem function and its response to climate

perturbation is increasingly important, especially in tropical systems where responses to changes

in biodiversity are less predictable and more challenging to assess experimentally. Using large

scale transplant experiments conducted at five neotropical sites, we documented the impacts of

changes in intraspecific and interspecific plant richness in the genus *Piper* on insect herbivory,

insect richness, and ecosystem resilience to perturbations in water availability. We found that

reductions of both intraspecific and interspecific *Piper* diversity had dramatic and site specific

effects on herbivory, herbivorous insect richness, and plant mortality. Ecosystem responses to

reduced intraspecific richness were often similar in magnitude to responses to reduced

interspecific richness. Increased water availability reduced herbivory by 4.2% overall, and the

response of herbivorous insect richness and herbivory to water availability was altered by both

intra and interspecific richness in a site contingent manner. Our results underscore the role of

intraspecific and interspecific richness as foundations of ecosystem function, and the importance

of community specific contingencies in controlling function in complex tropical systems.

Keywords: Biodiversity, Ecosystem function, Herbivory, *Piper,* Plant-Insect interactions

Introduction

As climate change and anthropogenic activity alter ecosystems at unprecedented rates, it

has become critical to understand the consequences of biodiversity loss on ecosystem processes

and the maintenance of ecosystem processes through species interactions. A complex mix of

anthropogenic forces are eroding multiple dimensions of global biological diversity, including

plant intraspecific, interspecific, and functional diversity (1,2). Plant diversity affects herbivore

abundance and diversity, thereby influencing biomass allocation and energy fluxes between

trophic levels (3). Because losses of plant diversity can destabilize the flow of resources to

higher trophic levels (3,4), understanding the connection between biodiversity and trophic

interactions is necessary to predict the consequences of the loss of primary producer biodiversity

on ecosystem traits such as resilience to environmental perturbation (5,6). While climate change

is associated with increased absolute precipitation in some regions and decreased precipitation in

others, IPCC models predict mid-century increases in the frequency of extreme precipitation

events in Central and South America (7–9), a phenomenon already observable in many

ecosystems (10). Measuring how precipitation change will affect the relationship between

biodiversity and ecosystem function is therefore increasingly important, particularly in diverse

tropical systems.

Very few multi-site, manipulative diversity experiments have been reported from tropical

areas compared to temperate environments (11), limiting our knowledge of the role of

biodiversity in ecosystem function in the most species-rich regions of the planet (12). Here we

investigate how multiple dimensions of plant diversity affect herbivory and plant mortality at

five neotropical sites and explore how diversity modulates ecosystem responses to changes in

water availability at three of those sites, in the context of several established hypotheses (Table

S1). By altering plant uptake of nutrients and the induction of plant defenses, abnormal levels of

precipitation can alter herbivore pressure, acting to alter the movement of resources into higher

trophic levels. As such, climate extremes of dry and wet periods are expected to strongly perturb

plant-insect interactions and thereby alter ecosystem function (13–15). The insurance hypothesis

suggests that greater biodiversity can act to stabilize ecosystems and improve their resilience to

environmental change (4,16). Despite traditional views that interspecific richness has a greater

impact on ecosystem processes than intraspecific diversity (17), recent research suggests that

plant intra- and interspecific richness can have similar effects on ecosystem productivity and

consumer abundance (18,19), and field studies have demonstrated both positive and neutral

effects of plant diversity on ecosystem resilience (20,21).

To test the insurance hypothesis, the effects of water addition on ecosystem function, and

the relative strength of intra- and interspecific richness in driving ecosystem function, we

conducted common garden experiments at five sites across Central and South America. Using 33

species in the genus *Piper* (Piperaceae) as a model system (22), we manipulated intra- and

interspecific plant richness in Costa Rica, Ecuador, Peru, and two sites in Brazil. We additionally

manipulated water availability in Costa Rica, Ecuador, and Peru (Figs. 1 & S1 A). We predicted

that reduced *Piper* diversity would lead to reduced diversity in higher trophic levels, that water

addition would lead to altered herbivore pressure, and that lower *Piper* diversity would be

associated with more extreme changes in herbivory and plant mortality in response to water

addition. Additionally, we predicted that changes in intraspecific plant richness would affect

ecosystem function with magnitudes similar to interspecific richness.

Methods

Study sites and focal plant genus

We conducted a large-scale transplant experiment replicated across five sites spanning

42° latitude in the Neotropics (Fig. 1 & Table S2) and encompassing a range in annual

precipitation from 1271 mm to 4495 mm (Table S2). At each site, we studied herbivory on

planted individuals in the genus *Piper* (Piperaceae) in response to experimental treatments. Study

sites included lowland equatorial humid forest at La Selva Biological Station, Costa Rica; high

elevation equatorial humid forest in Yanayacu Biological Station, Ecuador; high elevation

equatorial humid forest at El Fundo Génova, Peru; lowland seasonally dry gallery forest in

cerrado within the phytogeographic domain of the Atlantic Forest in Mogi-Guaçu Biological

Reserve, Brazil; and lowland seasonally semideciduous forest in the transition between the

Atlantic Forest and the Cerrado phytogeographic domains in Uaimii State Forest, Brazil. Climate

classifications follow the Köppen-Geiger climate model (Table S2; 22).

Multiple species of *Piper* are found at all sites, ranging from 11 species in Mogi-Guaçu to

50 species in Costa Rica (24). *Piper* is an ideal genus for large-scale comparative studies, as it is

found across the Neotropics and subtropics, from ~10° N to about ~32° S. In addition to being

widespread, *Piper* is abundant and diverse across its range, encompassing ~1000 species in the

Neotropics (25). *Piper* has been the subject of detailed studies of herbivory, and its herbivore

fauna has been surveyed across its range (26–37). Members of the genus host both specialist

caterpillars and beetles as well as generalist caterpillars and Orthopterans (22,26–29,31). Leaf

damage patterns produced by the different classes of herbivores are well documented (31)

Experimental design

At each study site, a factorial experiment was implemented to test the effects of

plant intraspecific and interspecific richness on herbivory, variation in herbivory, and insect

richness. Experimental plants were propagated from cuttings of naturally occurring *Piper* plants

(typically with three nodes and zero to one leaves). In Costa Rica, Ecuador, and Peru a water

addition treatment was also tested. In all sites except Uaimii, experimental plots measured 4 m in

diameter and contained 12 *Piper* individuals, planted either in pots with locally derived soil (for

experiments manipulating water availability) or directly into the ground (for experiments without

water additions in Mogi-Guaçu and Uaimii; Table S2). Plots in Uaimii were 2 m in diameter and

contained six *Piper* individuals, limited by the number of cuttings available. Plants were

cultivated without fertilizer or irrigation.

The interspecific richness treatment had three levels at all sites except Uaimii, where

there were only two levels due to limited numbers of cuttings. At each site a subset of naturally

occurring *Piper* species was selected to act as a species pool for each experimental plot (Table

S3). These species were selected to constitute a breadth of genetic and functional diversity

representative of the *Piper* community present at each site. Combinations of species in each plot

for each level of the interspecific richness treatment were selected randomly from these species

pools. The first level was a monoculture of *Piper* consisting of individuals of a single species. In

the second level, two *Piper* species were randomly selected for planting (e.g. six individuals of

each species). These species combinations were the same when crossed with the low and high

levels of the intraspecific richness treatment (see below). In the high interspecific richness plots,

the number of species varied between sites and included all *Piper* species available to produce

cuttings (Table S2). The minimum number of *Piper* species in a high richness plot was three in

Uaimii, and the maximum was 12 in Costa Rica and Ecuador. Due to the low number of species

available in Uaimii no two-species plots were planted. The interspecific richness treatment was

replicated three times per site. In Mogi-Guaçu and Uaimii, each replicate contained a

monoculture plot for each species.

An intraspecific richness treatment with two levels (clones taken from a single mother

plant or individual cuttings all taken from unique mother plants) was crossed with the

interspecific richness treatment. In Costa Rica and Ecuador, the high interspecific richness plots

only included individuals from unique mother plants because 12 different species were planted.

In Mogi-Guaçu and Uaimii, the experiments were fully crossed. In the high intraspecific richness

treatment, cuttings were from different *Piper* individuals growing at least 10 m apart, so they

were not from the same root system are were most likely genetically unique individuals.

A water addition treatment was crossed with the inter- and intraspecific richness

treatments in Costa Rica, Ecuador, and Peru in order to examine how IPCC predicted regional

mid-century increases in precipitation in these regions (38) may affect herbivory and trophic

interactions. Plants were cultivated in five to six liter pots with drainage holes. Each potted plant

under the water addition treatment was watered with an extra 2 L of water twice per month in

Costa Rica and Ecuador. In Peru, 2 L of water were added to plants every three months. Water

was rapidly applied as a flooding event to completely saturate the soil. All low richness plantings

in Peru were watered, but in Costa Rica and Ecuador the low richness plantings were not watered,

so only two levels of the species richness treatment were crossed with the water treatment (Figs.

1 & S1 A). Plant cultivation periods lasted in excess of one year at all sites, and as such water

application treatments were performed in both the wet season and dry season.

Experimental plots were randomly located in the study sites; in Mogi-Guaçu and Uaimii

they were organized in three replicate blocks. There was a minimum of 20 m between plots in

blocks and 100 m between blocks. Randomly located plots were separated by a minimum of 50

m. The plots were spread across a maximum of 1,500 m in Peru and a minimum of 275 m in

Ecuador. Experimental periods lasted between 1.4 and 2.8 years, depending on site (Table S2),

and all measurements of herbivory were conducted on leaves that were initiated during these

periods. Cuttings were replaced if they died during the first three months of the experiment. One

of the authors was present at all of the sites for multiple visits to ensure as much standardization

of treatments as possible.

To determine the effects of site variation in natural levels of precipitation on the

outcomes of the water treatments, the absolute level of precipitation and precipitation anomaly

relative to climate normals was collected for each month of the experimental periods in Costa

Rica, Ecuador, and Peru (Fig. S2). Climate data for Costa Rica were obtained from La Selva

Biological Station. Data for Ecuador and Peru were obtained using TerraClimate (39), and an

interpolation error in the precipitation for Peru in February 2016, was corrected using data from

the National Service of Meteorology and Hydrology of Peru (SENAMHI).

Measures of herbivory and insect richness

Herbivory was recorded by taking photographs of all the experimental leaves at the end

of the experimental period. Additional photos were taken every three months in Uaimii and

Mogi-Guaçu, and in the first five months of the experiment in Costa Rica. These data were used

to measure herbivory and to determine the types of herbivores feeding on leaves based on

patterns of damage. When herbivores were observed on plants, they were photographed but were

left to continue feeding so as to not interfere with the experiment. Herbivory was quantified by

eye for each type of herbivore on each leaf by a single parataxonomist with extensive experience

measuring herbivory on *Piper* following established protocols (31). The amount of leaf area

consumed was measured in function to the total leaf area by visually dividing the leaf into equal

sized segments to determine the percent area missing. This was measured as a continuous value

to the greatest possible accuracy. Insect herbivores were identified to the lowest taxonomic level

possible based on their damage patterns (genus for specialist Lepidoptera, family for other

Lepidoptera, family for Coleoptera, order for Orthoptera). Direct observations of herbivores were

rare, so only damage patterns were used in analyses. The different types of damage recorded

were used as proxy for insect insect richness on *Piper*. Hereafter, the term "insect richness"

refers to the richness of herbivore damage patterns on plants.

Data analyses

The percentage of leaf tissue consumed by each insect taxon (based on damage patterns)

on individual leaves was summed to determine the total percentage of herbivory on each leaf,

and the variance in herbivory was calculated as the variance in leaf damage within each plant in

a plot. Due to high mortality rates across sites, *Piper* interspecific richness in each plot was the

final number of *Piper* species present in each plot at the end of the experiment (rather than the

number of species planted) and was analyzed as a continuous covariate rather than a categorical

treatment. Because most plant deaths occurred early in the experiments, final *Piper* interspecific

richness more accurately reflects the local plant richness experienced by herbivores. In order to

compare the magnitude of effect of intra and interspecific diversity on predictor variables,

interspecific diversity for each plot was analyzed as the proportion of the interspecific diversity

present in the species pools at each site. The effect of intra and interspecific diversity on each

response variable is reported as the change in response as diversity changes from the lowest

possible value at each site to the highest.

The effects of intraspecific richness, interspecific richness, and water addition on percent

leaf area consumed, the percentage of damaged leaves, variance in herbivory, and insect insect

richness were analyzed across and within sites using hierarchical Bayesian models (HBM). This

framework acts as the Bayesian equivalent of a random-effects model where site is a random

effect, allowing for generalized parameter estimates across sites. Analyses testing *Piper*

intraspecific richness and interspecific richness were conducted using data from all sites; the

effects of water addition and its interactions with richness were conducted using data from

Ecuador, Costa Rica, and Peru. Due to high mortality in Peru, interactions between the water

addition and intraspecific richness treatments could not be modeled for measures of herbivory or

insect richness and were only tested for survival.

We acknowledge that treatment level combinations were not the same across the different

sites, but that is the nature of random-effects models, where different levels of random factors,

such as site or year, consist of treatment levels that are unique to that site or year. In fact, this

type of design goes back to the origins of mixed models (40,41), and are also a strength of the

design, since lack of interactions between fixed and random effects increases generality (42).

Even at a single site, manipulated variables in ecological experiments do not even consist of the

same manipulations across all of the units of replication, as they suffer from problems such as

multiple versions of treatments, interference, and noncompliance (43)

Bayesian structural equation models (BSEMs) were constructed for each site using all

treatments as exogenous variables and insect richness and herbivory as endogenous variables.

Three path models (Table S1) were developed under the assumptions that 1) intraspecific and

interspecific richness may influence herbivory both directly and indirectly by modulating insect

richness 2) water addition may influence both herbivory and insect richness and 3) insect

richness may influence herbivory directly (Table S1). Models I, II, and III were tested in Ecuador,

Costa Rica and Peru, while model III was used to analyze data from Mogi-Guaçu and Uaimii.

Additional models incorporating interactions between intraspecific richness, interspecific

richness, and water availability were tested in Costa Rica, Ecuador, and Peru.

Models were run at the leaf level for herbivory and insect richness, and at the plant level

for the percentage of leaves with damage, variance of herbivory, and *Piper* survival. For all

HBMs and BSEMs, model convergence was estimated visually using traceplots and an $\hat{R}$

discriminatory threshold of 1.1 (44). Model fit was determined via posterior predictive checks

(PPC) using the χ^2 discrepancy of the residuals as the discriminatory function (45). A PPC near

0.5 indicates a high model fit, while values near one or zero indicate poor fit. BSEM models

were further compared using the Deviance Information Criterion (DIC).

All Bayesian models were written in JAGS via the jagsUI package in R (46) using

Markov chain Monte Carlo (MCMC) samplers with weakly-informative priors. For the majority

of models, MCMC runs were conducted for 10,000 iterations using the first 1,000 iterations as a

burn-in phase to generate posterior distributions of parameter estimates for each response

variable. HBMs modeling interactions required 20,000 iterations with the first 5,000 as burn-ins

for all models to consistently converge. Mean parameter estimates and 95% credibility intervals

(CIs) were calculated for all responses. 95% CIs which do not cross the y-axis are typically

associated with less than a 2.5% type S error rate (47). A posterior probability of direction (PD)

was calculated based on the percentage of the posterior distribution responding in the same

direction as the median response. A PD of 95%, for example, indicates that the same direction of

response (e.g. a positive or negative response) was observed in 95% of iterations, regardless of

the magnitude of the response (48). PDs less than 95%, indicate lower confidence that a

relationship exists, but can still be interpreted as the probability that an effect exists.

To analyze the effects of the experimental treatments on plant survival, survivorship

curves were constructed in all sites. Analyses of *Piper* survival were based on the initially

planted interspecific richness treatment of each plot because mortality occurred early in the

experiment. The effect of water addition, intra- and interspecific richness and species identity on

Piper survival were analyzed using Cox proportional hazard models. All data were analyzed

using R software v4.1.0 (49).

Results

Considerable variation in herbivory and plant mortality was observed among study sites.

Percent herbivory was lowest at Mogi; only 9% of leaf tissue was consumed by herbivores

compared to 22% of leaf tissue consumed in Uaimii. *Piper* mortality was highest in Peru (89%),

likely due to an El Niño event, and was lowest in Ecuador (27%; Fig. S3 A). *Piper* mortality

resulted in the reduction of species richness in many plots and the loss of some combinations of

treatments, notably the loss of all *Piper* in unwatered, low intraspecific richness plots in Peru

(Fig. S1 B). Plots originally planted as high interspecific richness treatments had a final richness

of between 1 and 12 species in Costa Rica, 9 and 11 species in Ecuador, 1 and 3 species in Peru,

2 and 4 species in Mogi, and only 2 species in Uaimii. We identified herbivore damage from 13

insect taxa at the five study sites, as well as damage by leaf miners of unknown orders. A total of

10 taxa were observed in Costa Rica, Ecuador, and Peru, 9 taxa were observed in Mogi, and 8

taxa were observed in Uaimii (Fig. S3 B). The majority of taxa in Costa Rica and Ecuador were

Piper specialists, but the proportion of leaf tissue consumed by specialists was only greater than

generalist damage in Costa Rica.

Our experiments revealed pronounced heterogeneity in ecosystem responses to changes

in water and *Piper* diversity between sites (Figs. S4 & S5). PPCs for all HBMs were within 0.03

of 0.5, indicating models fit well. Model II (Table S1) was selected as the most parsimonious

causal model for BSEMs in Costa Rica (PPC = 0.499), Ecuador (PPC = 0.499), and Peru (PPC =

0.498). Fit was high for model III in Mogi (PPC = 0.5) and Uaimii (PPC = 0.497). When all sites

were analyzed together, percent herbivory was 4.2±3.6% (mean ± 95% CI) lower in plots that

received additional water (PD = 98.7%), but we did not detect a cross site effect of water on the

percentage of leaves with damage (Fig. 2 A-B). We did not detect any direct cross-site effects of

water addition, insect richness, or plant diversity on the variance in herbivory (Fig. 2 C), or *Piper*

survivorship (Fig. 2 D). Notably, greater *Piper* interspecific richness was associated with a 15 ±

18.6% increase in the richness of insect herbivores when comparing high diversity plots to low

diversity plots across sites (PD = 95.0%; Fig. 2 E) and an indirect increase in herbivory was

mediated by insect richness (Fig. 3). Insect richness was associated with an 8.8±2.8% increase in

herbivory per insect taxon present (PD = 100%) and a 6.7±6.9% increase in the percentage of

leaves with any damage (PP = 97%) (Fig. 2 A, B). Changes in both intraspecific richness and

interspecific richness altered these ecosystem processes considerably, although effects varied in

strength and direction across sites (Fig. 3, Table S4). Although intraspecific richness was not

quantified, the genetic origin of *Piper* plants (clones vs. independent propagules) affected

herbivory, plant mortality, and insect richness with effect sizes of similar or even greater

magnitude than interspecific richness when standardized as the proportion of available species

richenss at each site. However, we found that intra- and interspecific richness often had

contrasting directions of effect on insect richness and multiple measures of herbivore pressure

(Figs. 3, S4 B-C & S5 B-C). For example, in Costa Rica insect richness was $6.0 \pm 5.9\%$ (PD =

93.7%) lower in plots with high interspecific plant richness compared to low, while high

intraspecific richness increased insect richness by $7.2 \pm 6.9\%$ (PD = 98.1%). In contrast, in

Ecuador high interspecific richness was associated with a $43.6 \pm 6.4\%$ (PD = 100%) increase in

insect richness, and high intraspecific richness decreased insect richness by $13.6 \pm 5.2\%$ (PD =

100%; Fig. S5 B-C).

The effects of *Piper* richness were especially relevant when examining interactions

between *Piper* richness and water addition (Fig. 4). The effects of water addition were altered by

Piper intra- and interspecific richness at all sites. While a reduction in herbivory driven by water

availability was observed in Costa Rica (4.7±2.5%, PD = 100%) and Peru (5.1±3.7%, PD =

100%), this effect was only present in Ecuador when intraspecific richness was high (Fig. S6).

Across sites, water addition had negligible or negative effects on insect richness at low

interspecific richness, but this effect was reduced or reversed when interspecific richness was

high. Insect richness increased by 20.8±18.9% more in interspecifically diverse plots that

received water compared to unwatered plots in Costa Rica (PD = 99%), 15.2±16.8% more in rich,

watered plots in Ecuador (PD = 96%), and 29.3±31.5% more in Peru (PD = 98%) Additionally,

Water addition had a negligible effect on insect richness in Ecuador when intraspecific richness

was low, but increased insect richness by 12.1% when intraspecific richness was high (PD =

99%) (Fig. 4 B). Water addition had a negative effect on *Piper* survival in Costa Rica when

intraspecific richness was low, but improved survival by 12.1 ± 8.5% in high intraspecific

richness plots (PD = 99%; Fig. 4 C). Interactions with water in Peru may have been influenced

by an El Niño event which resulted in abnormally low rainfall and high *Piper* mortality, while

the typically wetter sites in Costa Rica and Ecuador experienced greater precipitation than

average (Fig. S2). As there was 100% mortality in low intraspecific richness, unwatered plots in

Peru, we could not explore interactions between intraspecific richness and water for any response

except *Piper* mortality in this site.

Cox analysis of plant survival

Water addition did not affect *Piper* survival in Costa Rica ($z = -1.2$, $P > 0.2$), or in

Ecuador ($z = 0.2$, $P > 0.8$), but survival was reduced by 48% in plots without water in Peru ($z =$

3.2, $P = 0.001$). Intraspecific richness increased survival by 38% in Costa Rica ($z = -3.3$, $P <$

0.001) and by 32% in Ecuador ($z = -4.9$, $P < 0.001$), but had no effect on survival in Peru ($z = -$

1.55, $P > 0.1$), Mogi ($z = -0.22$, $P > 0.8$), or Uaimii ($z = 0.18$, $P > 0.8$). Interspecific richness had

no effect on survival in Costa Rica ($z = -1.5$, $P > 0.1$), Ecuador ($z = -0.56$, $P > 0.5$), Peru ($z = -$

0.49, $P > 0.6$), or Uaimii ($z = 0.21$, $P > 0.8$). In Mogi, survival was reduced by 13% in plots with

higher interspecific richness ($z = -2.16$, $P = 0.031$). There was an interaction between

intraspecific richness and water addition in Costa Rica and Peru. Survival in Costa Rica

increased in response to water in high intraspecific richness plots, and decreased in response to

water in low intraspecific richness plots ($z = 3.8$, $P < 0.001$), while the opposite pattern was

observed in Peru ($z = -2.4$, $P = 0.02$; Fig. S7). Although plant die-offs caused plot level species

extirpation at several sites, plant species identity was not related to survival rates with the

exception of *P. peltatum*, which had the lowest survival rate of any species planted in Costa Rica.

Further statistical results are available in the supplement.

Discussion

Our results demonstrate two key patterns. First, the strength of effects of intraspecific

richness on higher trophic levels is comparable to that of interspecific richness, supporting our

predictions and corroborating recent studies demonstrating the importance of intraspecific

richness (18,19). However, the effects of intra- and interspecific richness on herbivory, insect

richness, and plant survival can differ dramatically, in contrast to our predictions. Second, we

found that perturbations in water availability can have complex effects on herbivores and plant

survival, and that these effects can be modulated by plant diversity in a context sensitive manner.

While our prediction that water availability would influence herbivory across sites was supported,

our results suggest that biodiversity loss and climatic perturbations may have dramatically

different effects on ecosystem function at local scales, which may diminish our ability to predict

how local communities will change as anthropogenic stressors increase.

Although experimental methods varied somewhat between study sites, we do not believe

this can fully explain the level of heterogeneity observed in ecosystem response. For example,

the methods employed in Costa Rica and Ecuador were nearly identical, and yet the directions of

effect of intra and interspecific diversity on insect richness were reversed in these sites (Fig. S5

B-C). The only meaningful differences between these two sites were the climate and the

composition of the local *Piper* and insect communities. There was considerable variation in both

biotic (Fig. S3 B) and abiotic factors (Table S2) across sites, which may have contributed to the

heterogeneity observed in ecosystem response to *Piper* diversity.

As we did not directly measure plant stress or nutrition, it is difficult to determine the

exact mechanism through which water addition reduced herbivory. The presence of an El Niño

weather pattern during the course of the Peru experiment may have led the water addition

treatment to relieve plants from drought stress, while water addition in Costa Rica and Ecuador

may have added to water stress in treated plants as these sites received an above average level of

rainfall during the course of the experiment. Despite this, water addition consistently suppressed

herbivory in Costa Rica, Ecuador, and Peru under natural levels of *Piper* diversity, suggesting

that predicted increases in precipitation in the next century (38) will dramatically alter the flow

of resources from primary producers. Although we were only able to record the richness of insect

herbivory patterns, this measure is indicative of the functional diversity of insect herbivores on

Piper and any changes to this value represent changes in interactions between *Piper* and higher

trophic levels (31). As such, the additional reduction in effects of water addition on insect

richness when *Piper* richness was low suggest that biodiversity loss in tropical systems will alter

the ability of higher trophic levels to respond to environmental perturbations.

As herbivore taxa can be differently affected by manipulations of diversity (50), variation

in the direction of the effect of intra- and interspecific richness may be due in part to variation in

the composition of insect communities and herbivore pressure measured across sites. Changes in

neighborhood effects when small numbers of plant species dominate a community can lead

either to the reduction or increase of herbivore pressure, dependent both on the nature of plant

species lost and herbivore species present in a given community (51). As such, local variation in

community composition has the potential to greatly alter the effects of both climate change and

biodiversity loss on ecosystem function. Regardless of how biodiversity loss effects ecosystem

function at large scales, variation in abiotic and biotic factors at locals scales can alter these

effects, reducing our ability to predict how anthropogenic activity will alter ecosystem function.

This is especially relevant in tropical systems, which have been the subject of far fewer studies

of ecosystem function. As such, our knowledge of local scale effects on the relationship between

biodiversity and ecosystem function remains limited in these systems.

A long-standing question in ecology has been the extent to which ecosystem function

increases with biodiversity and if this relationship plateaus at a level past which ecological

redundancy predominates. Recent results from less complex temperate grassland systems suggest

that these ecosystems can be described by a mostly linear relationship between richness and

function, where even rare species contribute uniquely to ecosystem function (52). In these

systems, high contingency can be expected, where ecosystem-level effects will depend on most

of the interacting species. In contrast, we might expect that diverse, tropical communities could

be characterized by greater ecological redundancy and thus be subject to less contingency

(53,54). Despite these expectations, our results demonstrate heterogeneity in ecosystem response

to changes in both intraspecific and interspecific richness in five tropical sites, suggesting that

complexity in these systems may not reduce the contingency effects of biodiversity loss.

Understanding the impact of biodiversity loss in tropical forests is fundamental to our ability to

conserve those systems, and our findings highlight the importance of approaching the study of

changes in ecosystem function as context sensitive responses in complex ecosystems.

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L.A.R., L.A.D., A.M.S., T.L.P., E.J.T., M.J.K., and T.J.M. developed the original idea for the

research and secured funding. L.A.S., T.J.M, L.A.D, E.J.T., and H.G.L. collected Peru data.

H.G.L. and L.A.D. collected Costa Rica data. W.R.S., T.R.W., and L.A.D. collected Ecuador

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W.R.S measured herbivory for all sites. A.J.G. analyzed the data and wrote the first draft of the

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443 **Figures**

444 **Figure 1:** *Treatments of intraspecific richness, interspecific richness, and water addition used in*

445 *each of the five study sites.* White tiles represent treatment combinations which were not tested in

446 a given site.

Figure 2. *Hierarchical Bayesian model parameter estimates for the effects of water availability*

intraspecific and interspecific Piper *richness on herbivory, herbivorous insect richness, and*

Piper *survivorship across sites*

Violins represent the cross-site posterior parameter distribution for each relationship in site level

hierarchical Bayesian models. Black lines represent the median posterior estimate and white bars

represent 95% credibility intervals. Percentages above violins indicate the probability of an

effect being positive or negative (as indicated by a negative probability) in response to an

increase of the independent variable. Distributions for water addition compare watered and

control plots; distributions for interspecific richness compare *Piper* species richness standardized

as the proportion of the maximum richness used at a site; distributions for intraspecific richness

compare low and high intraspecific richness treatments; distributions for insect richness compare

responses per insect taxon present on an individual leaf

Figure 3. *Direct and indirect effects of plant diversity and water availability on insect*

herbivores at five study sites. Bayesian structural equation models comparing effects of different

drivers of herbivorous insect richness and herbivory at five sites. Standardized path coefficients

are means of the posterior distribution for the effects estimated at each causal path. Positive

relationships are indicated in blue with triangular heads, and negative relationships are indicated

in red with circular heads. Black arrows indicate path coefficients of zero magnitude. Dot plots

512 summarize the standardized mean of the posterior distribution for each causal path with 95% and

513 80% credibility intervals. Asterisks indicate causal paths where the probability of an effect being

514 positive or negative is >95%.

Figure 4. *Interactions between intraspecific or interspecific richness and water availability on herbivory, herbivorous insect richness, and* Piper *survival*

Bars indicate mean response and standard error of the mean. Percentages indicate the probability that the two slopes are different as calculated using site level hierarchical Bayesian models. Due

556　　to high mortality in Peru, interactions between water and intraspecific richness could not be

557　　compared for any responses except mortality. Single and two species richness plots have been

558　　combined for visualization purposes only.

578   **Supplementary figures and tables**

579   **Table S1:** *Path models and explanatory hypotheses.* Black arrows indicate causal paths between

580   exogenous and endogenous variables.

Hypothesis	Causal paths	Sites tested
Bottom-up diversity: Host plant diversity impacts insect richness through multiple mechanisms, such as reducing host density, masking or amplifying host signals, or altering the proportion of specialist herbivores (1, 2-3)	All models; paths A, C	All sites
Neighborhood effects: Plant diversity directly affects herbivory through mechanisms which do not alter herbivore diversity (1, 2)	All models; paths B, D	All sites
Water affects herbivore diversity: Changes in water availability induce changes to plant nutrition and defenses which can benefit or harm different herbivore taxa, leading to changes in herbivore diversity (4, 5)	Model I; path G	Costa Rica, Ecuador, Peru
Water affects herbivory: Water addition directly affects plant physiology, altering both the nutritive quality of plant tissue and the ability of plants to combat herbivores and leading to changes in herbivory (6)	Models I, II; path F	Costa Rica, Ecuador, Peru

583   **Table S2.** *Study site characteristics and experimental details.*

	La Selva Biological Station	Yanayacu Biological Station	El Fundo Génova	Mogi-Guaçu Biological Reserve	Uaimii State Forest
Location	N9.9400, W84.0433	S0.6008, W77.8903	S11.0948, W75.3516	S32.2510, W47.1558	S30.2957, W43.5731
Mean elevation (m a.s.l.)	35	2124	1133	627	800
Mean annual precipitation (mm)	4495	2900	1767	1271	1600
Mean annual T (°C)	25	16	23	26	19
Date planted	2015-03-01	2015-07-01	2015-10-01	2017-02-01	2018-02-20
Date harvested	2017-02-01	2016-12-01	2017-07-01	2019-12-01	2019-12-20
Number of species in high richness plots	12	12	6	4	3
Planted in pots / ground	Pots	Pots	Pots	Ground	Ground
Water addition treatment applied	Yes	Yes	Yes	No	No
Number of *Piper* planted	360	360	360	432	144
Number of *Piper* surviving till the end of the experiment	200	259	56	182	123

*Climate data from La Selva Biological station are for the experimental period (2015-2018) and

were provided by the Organization for Tropical Studies. Data from El Fundo Génova are based

on nearby San Ramón (https://en.climate-data.org/south-america/peru/junin/san-ramon-28556/).

Data from Ecuador were provided by Yanayacu Biological Station. Data for Mogi-Guaçu

Biological Reserve are from January 2017-December 2019 and are from the Centro Integrado de

Informações Agrometeorológicas of São Paulo. Data for Uaimii State Forest are from the Plano

de Manejo FLOE Uaimii, Instituto Estadual de Florestas of Minas Gerais.

598   **Table S3:** Species of *Piper* used at each study location

Study location				
La Selva Biological Station	Yanayacu Biological Station	El Fundo Génova	Mogi-Guaçu Biological Reserve	Uaimii State Forest
arboreum	grande	arboreum	arboreum	lepturum
peltatum	peludo	reticulatum	crassinervium	corcovadense
biolleyi	ecuadorense	glabribaccum	richardifolium	vicosanum
sancti-felicis	perareolatum	chanchamayanum	miquelianum	
reticulatum	baezanum	armatum		
nudifolium	escabrosa	lechlerianum		
multiplinervium	kelleii			
decurrens	hispidum			
pseudobumbratum	schupii			
imperiale	pubinervulum			
garagaranum	pequenia			
urostachyum				

611   **Table S4.** Mean parameter estimates and probability of direction (PD) for the effects of increases

612   in intraspecific diversity, interspecific richness, water availability and insect richness on

613   measures of herbivory, plant mortality, and insect richness.

Site	Predictor variable	Response variable	Mean parameter estimate	PD
All	Intraspecific richness	Percent herbivory	-0.87	63.59%
All	Intraspecific richness	Presence of damage	-3.26	71.60%
All	Intraspecific richness	Variance in herbivory	-9.97	57.31%
All	Intraspecific richness	Insect richness	-0.02	59.41%
All	Intraspecific richness	Plant survival	6.50	92.48%
All	Insect richness	Percent herbivory	8.78	100.00%
All	Insect richness	Presence of damage	6.63	96.73%
All	Insect richness	Variance in herbivory	17.86	71.13%
All	Insect richness	Plant survival	3.35	66.21%
All	Interspecific richness	Percent herbivory	0.02	50.34%
All	Interspecific richness	Presence of damage	7.40	83.01%
All	Interspecific richness	Variance in herbivory	-52.43	73.46%
All	Interspecific richness	Insect richness	0.20	95.00%
All	Interspecific richness	Plant survival	10.66	80.22%
All	Water availability	Percent herbivory	-4.19	98.73%
All	Water availability	Presence of damage	-6.33	89.46%
All	Water availability	Variance in herbivory	-118.10	91.35%
All	Water availability	Insect richness	-0.07	64.52%
All	Water availability	Plant survival	5.15	79.29%
Costa Rica	Intraspecific richness	Percent herbivory	-0.10	52.68%
Costa Rica	Intraspecific richness	Presence of damage	2.86	79.84%
Costa Rica	Intraspecific richness	Variance in herbivory	37.95	72.01%
Costa Rica	Intraspecific richness	Insect richness	0.10	98.07%
Costa Rica	Intraspecific richness	Plant survival	3.72	80.91%
Costa Rica	Insect richness	Percent herbivory	7.53	100.00%
Costa Rica	Insect richness	Presence of damage	5.60	100.00%
Costa Rica	Insect richness	Variance in herbivory	-6.74	65.69%
Costa Rica	Insect richness	Plant survival	3.60	66.25%
Costa Rica	Interspecific richness	Percent herbivory	-2.43	91.53%
Costa Rica	Interspecific richness	Presence of damage	-4.89	89.55%
Costa Rica	Interspecific richness	Variance in herbivory	-89.88	83.97%
Costa Rica	Interspecific richness	Insect richness	-0.08	93.43%
Costa Rica	Interspecific richness	Plant survival	18.69	99.61%
Costa Rica	Water availability	Percent herbivory	-4.76	99.99%
Costa Rica	Water availability	Presence of damage	-7.96	99.90%
Costa Rica	Water availability	Variance in herbivory	-151.05	98.04%

Costa Rica	Water availability	Insect richness	-0.02	62.33%
Costa Rica	Water availability	Plant survival	12.09	99.74%
Ecuador	Intraspecific richness	Percent herbivory	-3.50	99.87%
Ecuador	Intraspecific richness	Presence of damage	-8.10	99.95%
Ecuador	Intraspecific richness	Variance in herbivory	-80.24	92.45%
Ecuador	Intraspecific richness	Insect richness	-0.18	100.00%
Ecuador	Intraspecific richness	Plant survival	20.46	100.00%
Ecuador	Insect richness	Percent herbivory	8.68	100.00%
Ecuador	Insect richness	Presence of damage	8.23	100.00%
Ecuador	Insect richness	Variance in herbivory	22.77	97.99%
Ecuador	Insect richness	Plant survival	7.46	81.29%
Ecuador	Interspecific richness	Percent herbivory	-1.74	89.56%
Ecuador	Interspecific richness	Presence of damage	8.46	99.73%
Ecuador	Interspecific richness	Variance in herbivory	16.01	57.91%
Ecuador	Interspecific richness	Insect richness	0.58	100.00%
Ecuador	Interspecific richness	Plant survival	7.78	85.86%
Ecuador	Water availability	Percent herbivory	-2.67	99.52%
Ecuador	Water availability	Presence of damage	-5.72	99.78%
Ecuador	Water availability	Variance in herbivory	-28.52	68.94%
Ecuador	Water availability	Insect richness	0.04	85.57%
Ecuador	Water availability	Plant survival	5.51	90.57%
Mogi	Intraspecific richness	Percent herbivory	-2.15	98.59%
Mogi	Intraspecific richness	Presence of damage	-4.29	97.74%
Mogi	Intraspecific richness	Variance in herbivory	-24.33	67.12%
Mogi	Intraspecific richness	Insect richness	0.02	75.41%
Mogi	Intraspecific richness	Plant survival	1.57	66.65%
Mogi	Insect richness	Percent herbivory	11.17	100.00%
Mogi	Insect richness	Presence of damage	12.22	100.00%
Mogi	Insect richness	Variance in herbivory	80.94	99.99%
Mogi	Insect richness	Plant survival	1.95	58.03%
Mogi	Interspecific richness	Percent herbivory	2.02	90.42%
Mogi	Interspecific richness	Presence of damage	6.39	97.05%
Mogi	Interspecific richness	Variance in herbivory	-116.52	90.64%
Mogi	Interspecific richness	Insect richness	0.04	80.94%
Mogi	Interspecific richness	Plant survival	17.14	99.05%
Peru	Intraspecific richness	Percent herbivory	-0.09	51.56%
Peru	Intraspecific richness	Presence of damage	-9.11	94.39%
Peru	Intraspecific richness	Variance in herbivory	-58.63	73.32%
Peru	Intraspecific richness	Insect richness	-0.03	63.89%
Peru	Intraspecific richness	Plant survival	2.84	74.08%
Peru	Insect richness	Percent herbivory	8.09	100.00%
Peru	Insect richness	Presence of damage	4.38	99.51%
Peru	Insect richness	Variance in herbivory	30.07	87.18%
Peru	Insect richness	Plant survival	-2.31	61.10%

Peru	Interspecific richness	Percent herbivory	0.14	50.60%
Peru	Interspecific richness	Presence of damage	16.49	97.23%
Peru	Interspecific richness	Variance in herbivory	-69.46	70.67%
Peru	Interspecific richness	Insect richness	0.16	91.04%
Peru	Interspecific richness	Plant survival	5.70	77.33%
Peru	Water availability	Percent herbivory	-5.15	99.75%
Peru	Water availability	Presence of damage	-5.27	89.79%
Peru	Water availability	Variance in herbivory	-173.87	94.77%
Peru	Water availability	Insect richness	-0.22	99.69%
Peru	Water availability	Plant survival	-2.10	69.02%
Uaimii	Intraspecific richness	Percent herbivory	1.51	71.49%
Uaimii	Intraspecific richness	Presence of damage	2.21	64.77%
Uaimii	Intraspecific richness	Variance in herbivory	74.93	79.39%
Uaimii	Intraspecific richness	Insect richness	-0.01	55.41%
Uaimii	Intraspecific richness	Plant survival	3.92	77.51%
Uaimii	Insect richness	Percent herbivory	8.41	100.00%
Uaimii	Insect richness	Presence of damage	2.83	85.39%
Uaimii	Insect richness	Variance in herbivory	-36.56	83.52%
Uaimii	Insect richness	Plant survival	5.98	75.15%
Uaimii	Interspecific richness	Percent herbivory	2.23	70.86%
Uaimii	Interspecific richness	Presence of damage	10.61	87.21%
Uaimii	Interspecific richness	Variance in herbivory	-3.20	52.59%
Uaimii	Interspecific richness	Insect richness	0.27	98.14%
Uaimii	Interspecific richness	Plant survival	3.53	62.35%

Figure S1. *Treatments and number of plots used across sites*

A) Treatments of intraspecific richness, interspecific richness, and water addition at the

beginning of the experimental period at each of the five study sites, and B) final treatments at the

end of the experimental period in each site. White numerals indicate number of plots used.

Changes in *Piper* species richness and the loss of some treatment combinations was due to *Piper*

mortality over the course of the experimental period.

Figure S2. *Precipitation levels at study sites where the water addition treatment was applied*

A) Monthly precipitation over the experimental period at Costa Rica (CR), Ecuador (EC), and

Peru (PE). B) difference in monthly precipitation from climate normals at the three sites across

the experimental periods for those sites. Bars indicate median values, black points indicate mean

values. C) Cumulative precipitation over the course of the experimental period. Dotted lines

indicate average cumulative precipitation, shaded regions indicate 95%, 80% and 50% quantiles.

Climate normals are based on monthly precipitation for a period from 1958 to 1998.

Figure S3. *Overall herbivory, plant mortality, and insect richness at five study sites*

A) Percent herbivory, percentage of leaves with any damage, herbivorous insect richness, and

percent mortality of *Piper* in Costa Rica (CR), Ecuador (EC), Peru (PE), Uaimii (UI), and Mogi-

Guaçu (MG), across treatments. B) Proportion of herbivorous insect taxa observed at each site

measured by feeding damage patterns. Colored regions indicate the percentage of damage

observations contributed by each taxon. C) Proportion of herbivory attributed to each taxon at

five study sites. Colored regions represent the percentage of each leaf consumed by each taxon at

each site. White tiles represent sites where no leaf damage by that insect taxon was observed.

Figure S4. *HBM parameter estimates of percent herbivory, percentage of leaves with damage,*

variance in herbivory, and percent Piper *survival against levels of A) water addition, B)* Piper

intraspecific richness, C) Piper *interspecific richness, and D) insect richness at each site.*

Violins represent the posterior parameter distribution for each relationship across sites and within

sites in Costa Rica (CR), Ecuador (EC), Peru (PE), Uaimii (UI), and Mogi-Guaçu (MG). Black

lines represent the median posterior estimate while white bars represent 95% credibility intervals.

Percentages above violins indicate the probability of an effect being entirely positive or entirely

negative in response to an increase of the manipulated variable. Distributions for water addition

compare unwatered control and watered plots; distributions for interspecific richness compare

Piper species richness and are standardized as a percentage of the high richness treatment level;

distributions for intraspecific richness compare low and high intraspecific richness plots;

distributions for insect richness compare responses per insect taxon present at the leaf level

Figure S5: *HBM posterior parameter estimates of insect richness compared to levels of A) water*

addition, B) Piper *intraspecific richness, and C)* Piper *interspecific richness*

Violins represent the posterior parameter distribution for each relationship across sites and within

sites at Costa Rica (CR), Ecuador (EC), Peru (PE), Uaimii (UI), and Mogi-Guaçu (MG). Black

lines represent the median posterior estimate while white bars represent 95% credibility intervals.

Percentages above violins indicate the probability of an effect being entirely positive or entirely

negative in response to an increase of the manipulated variable. Distributions for water addition

compare the unwatered control and watered plots; distributions for interspecific richness

compare *Piper* species richness and are standardized as a percentage of the high richness

treatment level; distributions for intraspecific richness compare low and high intraspecific

richness plots

Figure S6: *Bayesian structural equation models for drivers of insect richness, herbivory and*

Piper survival at three sites, including interactions between intraspecific and interspecific

richness, and water addition. Path coefficients indicate the standardized mean of the posterior

distribution for each causal path. Positive relationships are indicated in blue with triangular heads,

and negative relationships are indicated in red with circular heads. Dot plots indicate the

standardized mean of the posterior distribution for each causal path in A with 95% and 80%

credibility intervals. Asterisks indicate causal paths where the probability of an effect being

entirely positive or entirely negative is > 95%.

768 **Figure S7:** *Percent Piper survival over time in five sites, compared to levels of intraspecific*

769 *richness and water addition.*

770   **Supplementary references**

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