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Tree seedling trait optimization and growth in response to local-scale soil and light variability

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ABSTRACT

At local scales, it has been suggested that high levels of resources lead to increased tree growth via trait optimization (highly peaked trait distribution). However, this contrasts with **This is the author manuscript accepted for publication and has undergone full peer review but has not been through the copyediting, typesetting, pagination and proofreading process, which may lead to differences between this version and the [Version of Record](#). Please cite this article as [doi: 10.1002/ECY.3252](https://doi.org/10.1002/ECY.3252)**

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33 (i) theories that suggest that trait optimization and high growth occur in the most common
34 resource level and (ii) empirical evidence showing that high trait optimization can be also
35 found at low resource levels. This raises the question of how are traits and growth optimized
36 in highly diverse plant communities? Here, we propose a series of hypotheses about how
37 traits and growth are expected to be maximized under different resource levels (low, the most
38 common, and high) in tree seedling communities from a subtropical forest in Puerto Rico.
39 We studied the variation in the distribution of biomass allocation and leaf traits and seedlings
40 growth rate along four resource gradients: light availability (canopy openness) and soil K,
41 Mg, and N contents. Our analyses consisted of comparing trait kurtosis (a measurement of
42 trait optimization), community trait means, and relative growth rates at three resource levels
43 (low, common, and high). Trait optimization varied across the three resource levels
44 depending on the type of resource and trait, with leaf traits being optimized under high N and
45 in the most common K and Mg conditions, but not at any of the light levels. Also, seedling
46 growth increased at high light conditions and high N and K but was not related to trait
47 kurtosis. Our results indicate that local-scale variability of soil fertility and understory light
48 conditions result in shifts in species ecological strategies that increase growth despite a weak
49 trait optimization, suggesting the existence of alternative phenotypes that achieve similar
50 high performance. Uncovering the links between abiotic factors, functional trait diversity and
51 performance is necessary to better predict tree responses to future changes in abiotic
52 conditions.

53

54 **Keywords:** biomass allocation traits, canopy openness, kurtosis, leaf area, specific leaf area,
55 Puerto Rico, soil nutrients

56 INTRODUCTION

57 Understanding how abiotic factors drive functional trait distributions and growth of local
58 communities is important for determining community responses to future changes in
59 environmental conditions (Violle et al. 2007, Estrada et al. 2016, Sakschewski et al. 2016).
60 Trait distributions of plant communities are assumed to be the outcome, to some degree, of
61 environmental conditions exerting selective forces and favoring species that make a good fit
62 for given conditions. In other words, traits are assumed to reflect optimal or, at least,
63 optimized ecological strategies for any given environment (although this set of optimal traits
64 may change due to temporal changes in conditions) (Southwood 1977, Grime 1979, Keddy
65 1992, McGill et al. 2006). As a result, local species composition should cover a relatively
66 narrow range of trait values that fit the requirements of the environment there. However, this

67 expectation contradicts the often-observed pattern of a wide diversity of forms and functions
68 in tropical plant communities and suggests the existence of alternative phenotypes with
69 equivalent performance (Marks and Lechowicz 2006, Worthy et al. 2020).

70 At local scales, the distribution of a given resource is typically unimodal, with some
71 resource values being more frequent than values located at the tails of the distribution (Fig. 1,
72 Appendix S1: Fig. S1) (Abrams 1995, Brigatti et al. 2007). For example, levels of canopy
73 openness (used as a proxy for understory light availability) are often around 10% in tropical
74 forests. However, light availability values, even in a closed forest, can range between < 1%
75 and 35%, with these extreme values being less frequent than mean values (Chazdon and
76 Fetcher 1984). Similarly, soil resources may exhibit a similar peaked hump-shaped
77 distribution, with soils showing low and high levels of fertility towards the tails of the
78 distribution (Appendix S1: Fig. S1). As traits reflect the interaction between the organisms
79 and their environment, local variation in abiotic factors is expected to have an impact on trait
80 distributions in plant communities and on growth. There are at least three potential scenarios
81 describing trait distributions at the community level, and plant growth in response to resource
82 gradients:

83 *Scenario 1: Peaked trait distributions towards common conditions.* In the first
84 scenario, uncommon conditions are too rare to lead to peaked trait distributions –trait
85 optimization– given that resources are not suitable and abundant enough to maintain stable
86 populations that represent a good fit for those conditions, while common conditions represent
87 suitable areas and allow trait optimization (Fig. 1A) (McGill et al. 2006, Violle et al. 2007).
88 These two different levels of trait optimization would imply better performance (e.g., growth)
89 for individuals present in the most common conditions and lower performance of those
90 individuals in places with uncommon resource levels.

91 *Scenario 2: Peaked trait distributions towards rare conditions.* Uncommon
92 conditions that represent extremes of the resource distribution (characterized by either low or
93 high resource levels) could lead to trait optimization due to strong selective forces that
94 operate under these conditions (Fig. 1B). For example, in high-light conditions species that
95 acquire resources fast and efficiently outcompete other species leading to optimization in leaf
96 traits. Similarly, in nutrient-poor soils only a handful of specialized species may persist,
97 representing a (relatively) narrow range of traits. Empirical evidence has shown that trait
98 values are more strongly selected at the extremes of the resource distribution characterized by
99 either low or high resource levels, with trait optimization occurring at these extremes rather
100 than at most common ones (Weiher et al. 1998, Cornwell and Ackerly 2009, de Bello et al.

101 2009, Kraft and Ackerly 2010, Swenson et al. 2011, Bernard-Verdier et al. 2012, Bruelheide
102 et al. 2018). For instance, a previous study found that open sites (i.e., treefall gaps) exert
103 strong selective forces on tree communities allowing only light-demanding species to
104 successfully colonize these forest gaps (trait optimization), while intermediate and low light
105 level sites exhibit greater diversity of life-forms (lower trait optimization) (Hubbell 2005).
106 Other studies have also shown that across environmental gradients, low soil nutrient levels
107 tend to favor stress-tolerant species, resulting in trait optimization towards lower extremes of
108 the resource distribution (Pinho et al. 2018). These patterns in trait distributions have been
109 well documented at the landscape and regional scales (Cornwell and Ackerly 2009, Enquist
110 et al. 2015, Šimová et al. 2015). At local scales, similar trends are expected if extreme abiotic
111 conditions select for particular traits that maximize organisms' performance (e.g., high
112 growth) (Fig. 1B).

113 *Scenario 3: Low peaked trait distributions occur in both common and rare resource*
114 *levels.* Alternatively to these two scenarios, traits might not be strongly optimized towards a
115 single or a narrow range of values. Instead, plants may display contrasting phenotypes that
116 constitute alternative solutions for the given local conditions (Marks and Lechowicz 2006,
117 Muscarella and Uriarte 2016, Worthy et al. 2020). For example, Hirose and Werger (1995)
118 found that in a tropical forest different species of trees exhibit contrasting strategies for
119 capturing light that result in similar performance. In this case, we should observe a lack of
120 trait optimization across different resource levels, yet species may still achieve high
121 performance anywhere along the local gradient of conditions.

122 Here, we examine both biomass allocation and non-biomass allocation traits. Biomass
123 allocation traits represent a group of traits that have shown great variation within species and
124 reflect the amount of biomass that plants allocate to a particular organ depending on the
125 resources that are the most limiting for plant growth (Bloom, Chapin, & Mooney, 1985;
126 Poorter et al., 2012; Umaña, Zhang, Cao, Lin, & Swenson, 2018). For example, according to
127 optimal allocation theory, strong limitation in soil nutrient content should result in increased
128 biomass allocation to roots at the expense of lower biomass allocation to leaves and stems
129 (Bloom et al. 1985). Another group includes non-biomass allocation traits, such as specific
130 leaf area or leaf area, which have shown strong responses to shifts in a variety of abiotic
131 factors (Chapin, 1980; Craine, Froehle, Tilman, Wedin, & Chapin, 2001; Oksanen, Fretwell,
132 Arruda, & Niemela, 1981; Reich, Walters, & Ellsworth, 1997; Wright et al., 2004). For
133 example, environments with low resource levels tend to be dominated by species with
134 conservative traits such as low specific leaf area and low leaf area (Le Bagousse-Pinguet et

135 al. 2017). Combined, biomass allocation and non-biomass allocation traits capture a wide
136 range of species responses to local-scale resource variation.

137 We study trait optimization in response to local-scale resource variation and how that
138 results in differences in seedling growth in a subtropical forest. We use growth and trait
139 information collected from all individuals present in 200 seedling plots (~1800 individuals)
140 in combination with local-scale abiotic information on soil nutrient content (K, Mg, and N)
141 and canopy openness (a proxy for light availability in the forest understory) (Appendix S1:
142 Fig. S1). While most trait-based studies have used species' mean traits values, which assumes
143 that all individuals within a species are identical in their traits, here, we have trait and
144 seedling growth information for all individuals. With this information, we can accurately
145 study trait and growth distributions across local-scale resource gradients. Specifically, we ask
146 the following questions:

147

148 1. How does trait optimization differ in extreme vs. common resource levels? We propose to
149 examine the three alternative scenarios described above by quantifying trait optimization
150 under different resource levels.

151

152 2. How do community mean trait values change across resource levels? We expect changes in
153 trait means across different resource levels depending on the type of trait. Specifically, we
154 expect biomass allocation traits to vary according to the optimal allocation theory (i.e., higher
155 allocation towards most limiting resources), while non-biomass allocation traits will be more
156 conservative (e.g. low SLA and LA) towards the most limiting extremes of resource levels.

157

158 3. How does seedling growth vary between extreme vs. common resource levels? Does trait
159 optimization lead to high seedling growth? We predict variation in seedling growth across
160 common and extreme levels of soil nutrient and light conditions, with at least two potential
161 and non-exclusive outcomes: 1) at high resource levels, seedling growth rates are high; 2)
162 seedling growth rates are high in resource levels that show high trait optimization.

163

164 **METHODS**

165 *Field surveys and trait data collection*

166 The study was conducted in El Yunque National Forest, in Puerto Rico (65°47' W, 18°19' N).
167 This forest, classified as a subtropical wet forest, has a mean annual rainfall of 3548 mm and
168 an average temperature of 23°C (Ewel and Whitmore 1973). The vegetation is dominated by

169 the tree species *Dacryodes excelsa* (Burseraceae) and the palm *Prestoea acuminata*
170 (Arecaceae). In 2013, 200 1x1m seedling plots were established to monitor seedling growth
171 for one year. Seedling plots were arranged in a regular grid separated by 10 m. All tree
172 seedlings ≤ 50 cm in height ($n = 1771$) were measured, tagged, and identified to species
173 (Umaña et al. 2015). Between June and July 2014, all surviving individual seedlings were
174 collected for trait measurements. We measured fresh leaf area (LA in cm^2) for 1-3 fully
175 expanded leaves. The leaves were then dried and weighed to calculate specific leaf area (SLA
176 = LA/dry mass; cm^2/g). Roots were cleaned and separated from the main stem. Leaves and
177 roots were dried in an oven for 72 hours at 70°C and measured for dry mass (g) to calculate
178 traits related to biomass allocation. Leaf mass fraction (LMF) was calculated as: LMF = leaf
179 dry mass/total plant dry mass. Root mass fraction (RMF) as: root dry mass/total plant dry
180 mass. Leaf area ratio (LAR) was calculated as: LAR = leaf area/total plant dry mass (Poorter
181 et al. 2012). Leaf traits included leaf area (LA) and specific leaf area (SLA). LA reflects the
182 area displayed to capture light (Rozendaal et al. 2006), while SLA describes a range of
183 strategies (conservative to acquisitive) for carbon assimilation (Reich et al., 1997; Wright et
184 al., 2004).

185

186 *Relative growth rate*

187 We defined seedling relative growth rate (RGR) as the change in log-transformed total height
188 (cm) from 2013 to 2014. The total height of each seedling was measured from the base to the
189 most distant part of the main stem. We standardized relative growth values by subtracting the
190 mean of each species and dividing by the standard deviation per species. This allows
191 comparisons across species with different inherent growth rates, as those involved in the
192 survival-growth trade-off (Arellano et al. 2019).

193

194 *Soil nutrient content and canopy openness*

195 Soil samples were extracted below the organic litter layer (0-10 cm depth) in July 2014. Each
196 soil sample consisted of mixed soil from each of the plot corners and from the center of the
197 seedling plots, and we collected a total of 200 soil samples. Soil samples were sent to
198 Brookside Laboratories Inc. (Ohio, USA) for chemical analyses. Magnesium and potassium
199 (Mg, mg/kg; and K, mg/kg) were extracted using the Mehlich-III solution. Magnesium is an
200 essential macronutrient found in the chlorophyll of green plants and potassium is essential for
201 photosynthesis and protein synthesis as well as carbohydrate transport and storage (Maathuis
202 2009). Potassium has been found to limit seedling growth in tropical and temperate forests

203 (Tripler et al. 2006, Wright et al. 2011). Total soil N concentration was obtained using the
204 total combustion method. Nitrogen is an essential constituent of proteins found in chlorophyll
205 and is needed in substantial amounts (Maathuis 2009). While soil P is an important limiting
206 resource for tropical forests, most of the P concentrations in our samples fell below the
207 detection threshold for the method and we could not use it for the present study.

208

209 To assess light conditions, we took hemispherical photographs using a camera Nikon
210 Coolpix5000 with an FC-E8 Nikon fisheye lens. The photographs were taken at 1m height in
211 the center of each seedling plot at uniform light conditions at dawn with homogeneous light
212 conditions (Glatthorn and Beckschäfer 2014). The photographs were analyzed using the Gap
213 Light Analyzer software (GLA) (Frazer et al. 2000) (available at
214 [https://www.caryinstitute.org/science-program/our-scientists/dr-charles-d-canham/gap-light-](https://www.caryinstitute.org/science-program/our-scientists/dr-charles-d-canham/gap-light-analyzer-gla)
215 [analyzer-gla](https://www.caryinstitute.org/science-program/our-scientists/dr-charles-d-canham/gap-light-analyzer-gla)). Appendix S1: Table S1 reports ranges and mean values for all the studied
216 abiotic variables.

217

218 *Plot selection at different resource levels*

219 Our approach consisted of classifying plots into three resource levels (low, common, and
220 high) and then calculating trait kurtosis per each resource level. This approach allows us to
221 use seedling communities large enough at each resource level in order to have robust
222 measurements of kurtosis, otherwise, our plots are too small to have a high number of trait
223 values (seedling abundance range: [1, 39]; mean = 9). Prior to the analyses, we examined
224 correlations between abiotic variables across the 200 plots (Appendix S1: Table S2) and their
225 distribution. All abiotic variables showed unimodal distribution with the most common
226 resource level around the average resource value (Appendix S1: Fig. S1).

227 Next, we calculated the relative frequency of each resource level using a kernel
228 density estimation for each abiotic variable using the R function "density" from package
229 "stats" (R Development Core Team 2017). Given that the density estimation is affected by
230 the bandwidth, we used two methods to select the appropriate bandwidth. The first method is
231 based on a linear diffusion process (Botev et al. 2010), a non-parametric density estimator
232 independent of a normal model and less sensitive to outliers (Botev et al. 2010). This method
233 was implemented by using the function "botev" from the package "provenance" in R
234 (Vermeesch 2020). The second method uses the asymptotic mean integrated squared method
235 error (AMISE) and consists of evaluating the mean integrated squared error of a density
236 estimate based on a normal distribution (see Appendix S2).

237 We classified subsets of plots into “low resource” plots, “high resource” plots, and
238 “the most common resource” plots, for each environmental variable (hereafter referred to as
239 low, high, and common, respectively). The low resource plots corresponded to the 10% with
240 the lowest level of the resource. The high resource plots corresponded to the 10% with the
241 highest level of the resource. The most common resource plots corresponded to the 10%
242 closest to the most frequent conditions, inferred from the location of the peak in the unimodal
243 estimation of density. Since we have some plots with repeated values for the different
244 resources studied, the number of plots selected per resource level varied between 10-16% of
245 the total number of plots; Appendix S1: Table S3 contains the specific number of plots
246 selected per resource level, and Appendix S1: Figure S1 indicates the plots that were selected
247 in each resource level (of the 200 plots established, some were excluded from the analyses to
248 focus on the most extreme and common values). To determine whether our results were
249 sensitive to the number of selected plots for each resource level, we repeated all the analyses
250 using the $p\%$ plots with the lowest, highest, or the most common levels of the resource, p in
251 $\{5, 6, \dots, 14, 15\}$. The results (Appendix S3) were not sensitive to the value of p , except in
252 low ranges (likely due to the low sample size, the inherent variability in the tails and/or the
253 skewed distribution of some of the resources studied).

254 255 *Measurement of trait optimization at each resource level*

256 To examine how trait distribution varied with different resource levels (question 1), we
257 compared trait kurtosis values using information from seedlings located in the low, high, and
258 the most common resource plots. Very peaked distributions (high kurtosis) indicate strong
259 trait optimization, while very flat distributions (low kurtosis) indicate weak trait optimization
260 (Enquist et al. 2015, Le Bagousse-Pinguet et al. 2017). We calculated kurtosis for individual
261 traits per resource level. We choose kurtosis over variance because we are interested in
262 detecting a pointed peak (or lack of it) in trait distributions beyond the information about how
263 spread is the data (Figure S2 in Appendix S1 shows that the degree to which a given
264 distribution is peaked is independent of the spread of the data) (Enquist et al. 2015).

265 Given that kurtosis values are difficult to compare with each other, at least in terms of
266 how much they differ from the expected by chance, we used a more interpretable metric than
267 raw kurtosis, by calculating the Standardized Effect Sizes (SES) for the kurtosis values
268 obtained in all resource level classes, for all environmental variables and traits. To calculate
269 the SES for kurtosis, we created null models where each trait value per individual within a
270 community matrix (including all 200 plots) was randomized 999 times and we re-calculated

271 kurtosis for each resource level for each randomized dataset. We then obtained a mean null
272 value and standard deviation from the null distribution that were used to calculate the kurtosis
273 SES:

274

$$275 \quad \text{SES Kurtosis} = \frac{\text{Observed kurtosis} - \text{mean null kurtosis}}{\text{SD null kurtosis}}$$

276

277 SES kurtosis values > 1.96 indicate higher than expected kurtosis in a certain trait
278 given the number of seedlings present in plots at each resource level and SES kurtosis values
279 < -1.96 values indicate a lower than expected kurtosis for a certain trait given the number of
280 seedlings present in plots at each resource level.

281

282 *Change in mean trait values in each resource level*

283 To determine how community mean trait values change across resource levels (question 2),
284 we calculated and compared mean community trait values at each resource level. When
285 calculating mean trait values, we used all individual seedlings present at each resource level.
286 To allow comparisons between traits measured in different units, we used standardized trait
287 values at the community level (mean = 0, standard deviation = 1). Next, to determine whether
288 trait means at each resource level differ from zero, we performed a bootstrap sampling with
289 replacement. To do this, we randomly selected seedlings (with different trait values) for each
290 resource level 999 times and calculated the mean per trait. If 95% of the values of the
291 bootstrap distribution overlapped zero, then it was considered not significant. In other words,
292 this test examines whether the trait mean at a given resource level differs from the expected
293 for the species.

294

295 *Variation in seedling growth rates at each resource level*

296 To address how seedling growth varies across resource levels (question 3), we performed a
297 similar analysis from the one used for mean traits, but instead of using functional traits, we
298 used RGR (see raw RGR distributions at each resource level in Appendix S1: Fig. S3). We
299 also performed a bootstrap sampling with replacement to assess the significance of changes
300 in RGR at each resource level (*i.e.* deviations in RGR from the expected for the species).
301 Furthermore, we evaluated if kurtosis was related to mean RGR at each resource level
302 (question 4) using a linear mixed-effects model (function "lmer", package "lme4" (Bates et
303 al. 2015) in R (R Development Core Team 2017)). In these models, the response variable was

304 mean RGR at each resource level. The independent variable was trait kurtosis per resource
305 level; abiotic factors (canopy openness, soil N, Mg, and K) and resource levels (high, low and
306 the most common) were included as intercept-specific random effects, to account for the
307 effects of variation of both variables on growth. The sample size for this model was $n = 60$.
308 To calculate the marginal and conditional coefficients of determination described by
309 Nakagawa and Schielzeth (2013), we used the function "r2" in the package "performance"
310 (Lüdecke 2020).

311

312 **RESULTS**

313 *Variation in trait distributions with different resource levels (question 1)*

314 Observed kurtosis patterns based on different traits showed a variation across the different
315 studied resource levels (low, common, and high), but the patterns were resource- and trait-
316 specific (Appendix S1: Fig. 2). For canopy openness, observed kurtosis values for all traits
317 did not differ from a random expectation, but there was a generalized trend for higher SES
318 kurtosis in the most common light conditions, except for RMF (Fig. 2). For soil K content,
319 SLA had higher than expected kurtosis at high resource values while LAR had higher than
320 expected kurtosis at the most common level, the rest of the traits had kurtosis that was not
321 significantly different than expected by the null model. For soil Mg content, LAR had higher
322 than expected kurtosis at the most common resource level, indicating trait optimization and
323 RMF had higher than expected kurtosis at the low resource level. The rest of the traits had
324 kurtosis levels that were not significantly different from the null expectation for soil Mg. For
325 soil N content, SLA, LAR, and LMF had a higher than expected kurtosis at the high resource
326 level indicating trait optimization at high N levels. The rest of the traits had kurtosis that was
327 not significantly different than expected by the null model for soil N. Results obtained using
328 the AMISE method were largely consistent with the results described above, especially for
329 results based on canopy openness and soil N (Appendix S2). For Mg and K, kurtosis values
330 of LAR, LMF, and SLA were less consistent (Appendix S2).

331

332 *Variation in mean trait values with different resource levels (question 2)*

333 In addition to the kurtosis analyses, we also examined shifts in mean trait values across the
334 three resource levels. For canopy openness, LA and LAR were significantly higher at high
335 light levels, and LAR, RMF, and SLA were significantly lower at the most common resource
336 level. The means of the rest of the traits were not significantly different from their
337 community-wide means at any level of canopy openness (Fig. 3). For soil K content, LA was

338 significantly higher at low and high resource levels but it was significantly lower at the most
339 common resource level. Also, LAR was significantly higher when soil K was high. The rest
340 of the traits showed means that were not significantly different from the mean community
341 value for soil K (Fig. 3). For soil Mg content, RMF and LA were significantly higher, while
342 SLA was significantly lower at the most common resource level. The rest of the traits showed
343 means that were not significantly different from the mean community value for soil Mg (Fig.
344 3). For soil N content, LAR and LMF were significantly higher at the low resource level,
345 while LMF and LAR were significantly lower and RMF was significantly high at the most
346 common resource level, and LA, RMF, and LAR were significantly higher at the high
347 resource level. The rest of the traits showed means that were not significantly different from
348 the mean community value for soil N (Fig. 3).

349

350 *Community growth rates in three resource levels (question 3)*

351 Mean seedling relative growth rate (RGR) was significantly higher at high levels of canopy
352 openness, soil N, and K content (Fig. 4). For the rest of the resource levels, standardized
353 RGR showed no significant differences. In addition, in the linear mixed-effects model
354 relating trait kurtosis and growth, the kurtosis had virtually no effect on RGR (coefficient =
355 0.01, 95% CI: [-0.01, 0.03]); the model had a low fit (marginal $R^2 = 0.02$ and conditional R^2
356 = 0.53).

357

358 **DISCUSSION**

359 Understanding how resource variation at local scales determines trait distributions and plant
360 performance remains a major question in ecology. To examine this question, we combined
361 individual trait and growth information collected from tropical seedling communities along
362 local-scale light and soil fertility gradients. We found that trends in trait optimization are not
363 generalizable across soil and light gradients. Specifically, trait optimization was observed for
364 the most common levels of K and Mg as well as in high soil N sites, while traits were not
365 particularly optimized at any level of canopy openness. Variation in growth rates was also
366 resource-specific, with seedling growth being the highest when canopy openness and soil K
367 and N content were high but was not related to trait optimization. Overall, these findings
368 suggest that peaked trait distributions do not necessarily match the most common resource
369 levels nor lead to an increase in growth as it is commonly assumed for local-scales studies.
370 Further, we show that seedling growth can increase despite a lack of trait optimization.
371 Below, we discuss in more detail our findings.

372

373 *Traits are optimized under low, high and common resource levels*

374 We observed a strong variation of trait optimization across three resource levels. However,
375 these shifts in trait optimization were highly dependent on the type of resource and studied
376 trait. For instance, LA, LMF, and LAR showed peaked distributions when soil N content was
377 high supporting the scenario showed in Fig. 1B. Similar results have been reported in studies
378 including broader environmental gradients than the one included in this study, in which high
379 soil fertility leads to high trait kurtosis (Enquist et al. 2015, Le Bagousse-Pinguet et al. 2017).
380 However, when considering soil Mg or K content, peaks in trait distributions (for LAR) were
381 found at sites with the most common resource level, supporting the scenario showed in Fig.
382 1A and suggesting strong selecting forces on trait distributions under the most common
383 conditions. These results are concordant with previous theoretical studies predicting a high
384 frequency of trait values under the most common conditions (Austin 1986, McGill et al.
385 2006, Enquist et al. 2015), yet empirical evidence has remained scarce. Surprisingly, we also
386 found a lack of trait optimization for any of the studied traits when considering canopy
387 openness that can be related to high temporal variability in light conditions. Overall, the
388 discrepancies in trait distributions across different resource levels suggest that selective
389 forces operate with variable intensity at different resource availabilities and may even result
390 in a prevalence of low peaked trait distributions. These results are consistent with another
391 study of tree communities across the entire island of Puerto Rico that examined the variation
392 in community weighed mean trait values with shifts in species occurrence to test for trait
393 optimization (Muscarella and Uriarte 2016). Their results showed that while some traits
394 support the optimization hypothesis (LMA, wood density, and maximum height), there were
395 many other cases where trait optimization was not found.

396

397 *Mean trait values are more acquisitive under high resource levels*

398 We expected a variation in mean trait values in response to different resource levels (Fig. 3).
399 Specifically, we predicted acquisitive strategies (i.e., high SLA and LA) to be predominant at
400 high resource levels and increased biomass allocation to organs involved in the acquisition of
401 limited resources (low resource levels). For leaf allocation traits, our results did not follow
402 the expected trend and instead, showed that seedlings tend to allocate a greater area to leaves
403 (LAR) rather than allocating biomass to roots in habitats experiencing high light levels.
404 These findings contradict the optimal allocation theory in which plants are expected to
405 allocate greater leaf biomass in low light conditions than at higher light availability (Mooney

406 1972, Bloom et al. 1985). However, for non-biomass allocation leaf traits, our results were
407 consistent with our expectations with LA increasing with high understory light availability
408 and indicating that seedling communities located in more open environments are
409 characterized by acquisitive traits. These findings are concordant with previous studies
410 conducted at a Neotropical forest where species displayed high leaf sizes values at high light
411 levels contributing to their carbon gain (Lambers and Poorter 1992, Poorter and Rozendaal
412 2008). It is important to note that similar trends observed for LAR and LA can be also
413 explained by the correlation between both traits (Fig. S4, Appendix S1).

414 By taking together results of mean and kurtosis trait values for variable light
415 conditions, we infer that trait distributions across the three light levels show to some degree
416 shifts in mean functional strategies, yet under each light level, there is no evidence of trait
417 optimization. This indicates that trait selection operates differently across low, most common,
418 and high light levels but within each resource level there is a substantial variation in
419 phenotypes that result in platykurtic trait distributions. This agrees with results from a
420 previous study from the El Yunque forest that showed that during the seed-to seedling
421 transition, functional divergence tends to be higher than expected by chance (Umaña et al.,
422 2016).

423 Mean trait values also varied in response to soil resource levels. We expected greater
424 biomass allocation to roots in poor soil conditions as suggested by the optimal allocation
425 theory (Bloom et al., 1985; Garnier, 1991; Marschner, Kirkby, & Cakmak, 1996; Santiago et
426 al., 2012; Wright et al., 2011) and an increase in SLA as soils become more fertile (Ordoñez
427 et al. 2009). However, these expectations were not supported in this study as we found no
428 general trends for patterns of biomass allocation or distribution of leaf-related acquisitive
429 strategies. One potential explanation is that patterns of biomass allocation and trait variation
430 depend on the interaction of multiple abiotic factors (Blonder et al. 2018). For example, root
431 biomass responds to both soil fertility and water availability that does not always covary at
432 the local scales included here and may result in conflicting trait responses. Another potential
433 explanation is that trait-trait correlations may potentially mask plant responses to variation in
434 resource availability. For example, the significant correlation between leaf and root mass
435 fractions may limit the ability to identify and separate the responses of below and
436 aboveground traits (Fig. S4).

437

438 *Seedling growth increases under high light and high soil N and K content but is not related to*
439 *trait optimization*

440 We expected that seedling growth would increase in areas of high resource levels and/or in
441 areas of high trait optimization (high trait kurtosis). Our results show that high light
442 availability and soil N and K content resulted in high seedling growth rates (Fig. 4); however,
443 growth was not related to trait optimization. Our findings related to light suggest that
444 understory light conditions are a key limiting factor in this forest and that at high understory
445 light availability, plants might enhance their growth in the absence of a trait optimization.
446 Our results are concordant with results from other plant community studies that have shown
447 species exhibiting contrasting phenotypes that increase plant performance and conform
448 alternative phenotypes under particular resource conditions (Hirose and Werger 1995, Pistón
449 et al. 2019, Worthy et al. 2020). Our results also agree with previous studies conducted in the
450 studied forest showing light as a key factor influencing performance at early ontogenetic
451 stages (Comita et al. 2009, Uriarte et al. 2018).

452

453 For soil nutrients, the direct effects of soil N and K should be carefully interpreted as both
454 variables were significantly correlated making it difficult to determine the independent
455 contribution of each factor to seedling growth ($r = 0.56$, $p < 0.001$, Table S2 in Appendix S1).
456 Although it has been suggested that soil N should not be considered as a limiting factor for
457 tropical forests (Hedin et al. 2009), previous studies in tropical regions have found that soil N
458 may have positive significant effects on plant biomass accumulation and productivity
459 (Vitousek and Sanford 1986, Mirmanto et al. 1999, LeBauer and Treseder 2008, Graefe et al.
460 2010, Santiago et al. 2012). Similarly, soil K has shown significant effects on seedling
461 growth, as found in our study (Santiago et al. 2012, Wright 2019).

462

463 *Conclusion*

464 Our findings suggest that high plant growth does not necessarily result from trait
465 optimization. Furthermore, we did not find support for the expectation that the most common
466 resource level translates into highly peaked trait distributions and high seedling growth.
467 Instead, we found that seedling growth at the Luquillo forest is limited by light, soil K, and N
468 content and that increase in growth can be the outcome from co-occurring contrasting
469 phenotypes. Overall, our study suggests that the high functional diversity found in tropical
470 forests partially results from a variety of phenotypic designs that perform well under given
471 resource levels. Also, an important portion of the trait variation observed in our study comes
472 from within species and highlights the need to incorporate this level of information in future

473 studies to gain a better understanding of community responses to local-scale resource
474 gradients.

475

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481 designed the fieldwork. MNU conducted fieldwork. MNU analyzed the data with input from
482 GA and NGS. MNU wrote the manuscript with input from GA, NGS, and JZ.

483

484 **Supporting Information**

485 Additional supporting information may be found online at: [link to be added in production]

486

487 **Data Availability**

488 Data is available from the Dryad Digital Repository (Umaña et al. 2021):

489 <https://doi.org/10.5061/dryad.sqv9s4n29>

490

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691

692 FIGURE LEGENDS

693 **Figure 1.** Conceptual diagram depicting changes in trait distributions (kurtosis) between rare
694 and the most common resource levels. In the center, the plot depicts the variation in a given
695 resource that describes a unimodal distribution with values at the two extremes being less
696 frequent than the values in the central section. Based on these three resource levels, we show
697 two scenarios (A and B) of how trait distribution is expected to vary between rare and
698 common resource levels.

699

700 **Figure 2.** SES kurtosis of trait values for three resource levels (low, common, and high).
701 Positive SES values above indicate kurtosis values higher than expected by chance. Negative
702 SES values below indicate significantly lower than the expected kurtosis in a given
703 community. Dashed horizontal lines represent ± 1.96 indicating significantly higher/lower
704 kurtosis than expected by chance. Color codes represent individual traits. LA– leaf area,
705 SLA– specific leaf area, LAR–leaf area ratio, LMF–Leaf mass fraction, RMF– root mass
706 fraction.

707

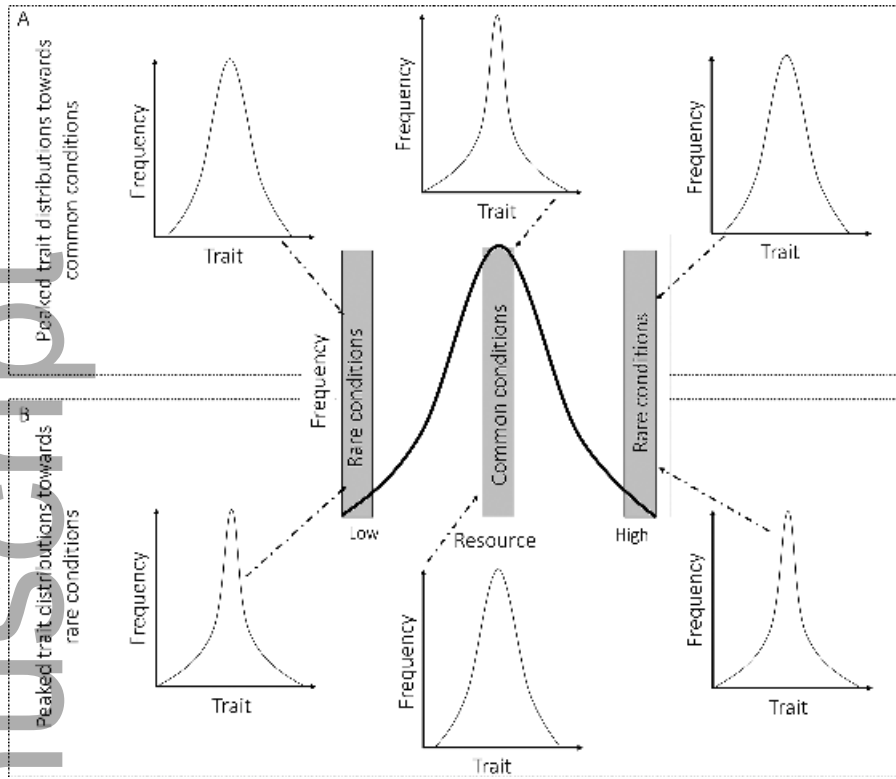
708 **Figure 3.** Differences in mean trait values for three resource levels (low, common, and high).
709 All traits were standardized at the community level (mean = 0, standard deviation = 1) to
710 make them comparable. Positive values indicate that for a given resource level, mean traits

711 are higher than the whole-community mean. Negative values indicate that for a given
712 resource level, mean traits are lower than the whole-community mean. Triangles indicate
713 95% confidence intervals not overlapping zero when estimating the distribution of the mean
714 after 999 bootstrap resampling, while circles represent 95% confidence intervals that overlap
715 zero. Color codes represent individual traits. Acronyms for traits are the same as in Fig. 2.

716

717 **Figure 4.** Differences in mean seedling relative growth rates (RGR standardized at the
718 species level) for three resource levels (low, common, and high). Color codes represent
719 resources (canopy openness and three soil nutrients). Positive values indicate that mean RGR
720 at a given resource level is higher than the whole-community mean. Negative values indicate
721 that mean RGR at a given resource level is lower than the whole-community mean. Triangles
722 indicate that 95% confidence intervals of bootstrap sampling do not overlap zero, circles
723 indicate that the 95% confidence intervals overlap zero.

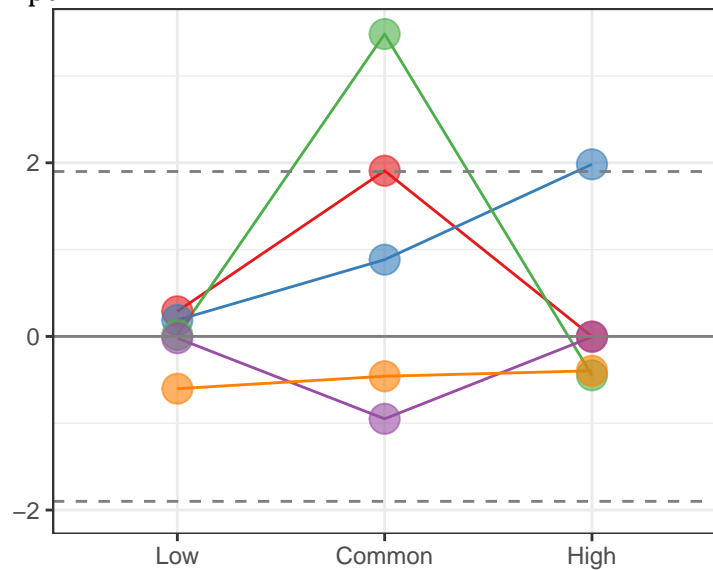
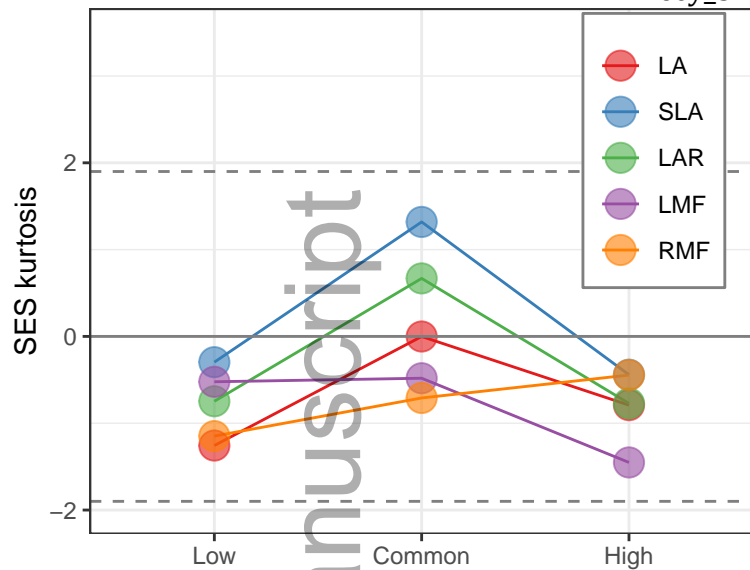
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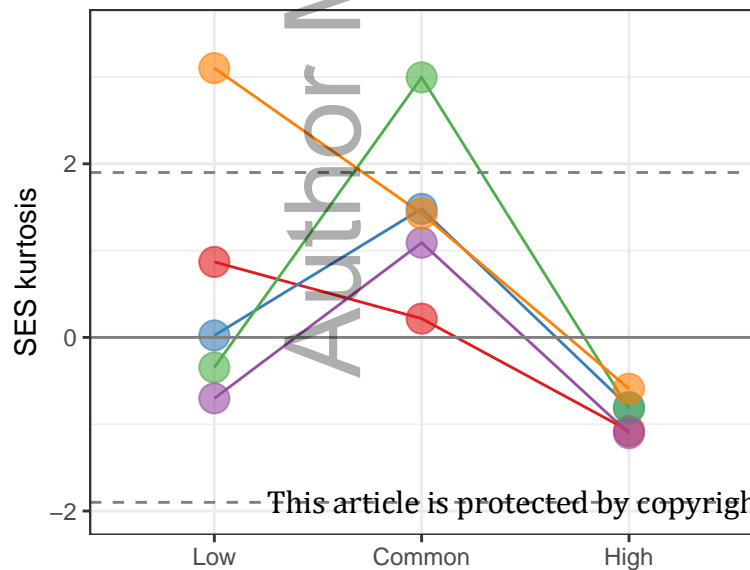
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Canopy openness

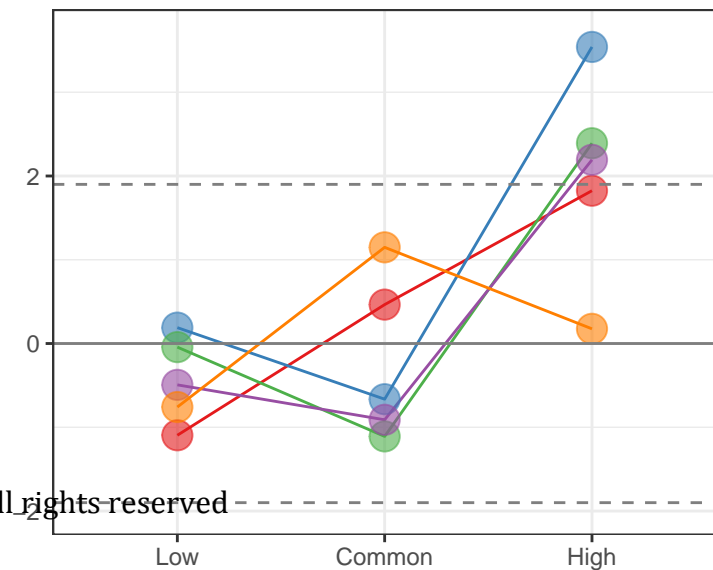
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Mg

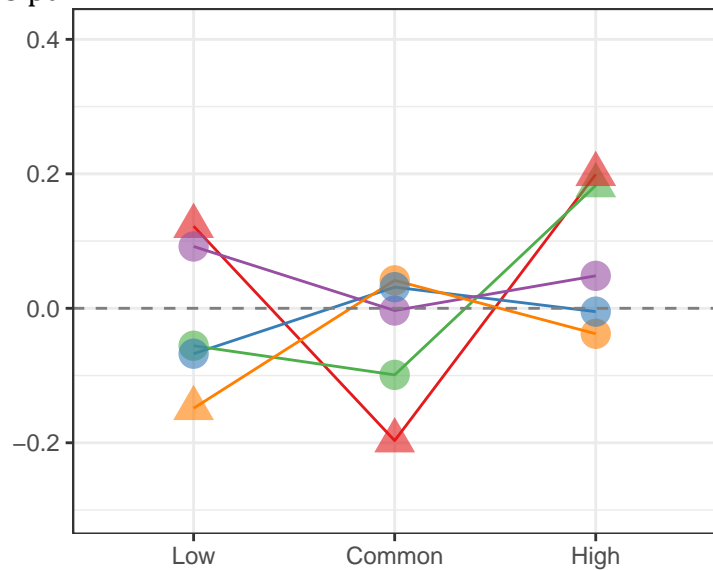
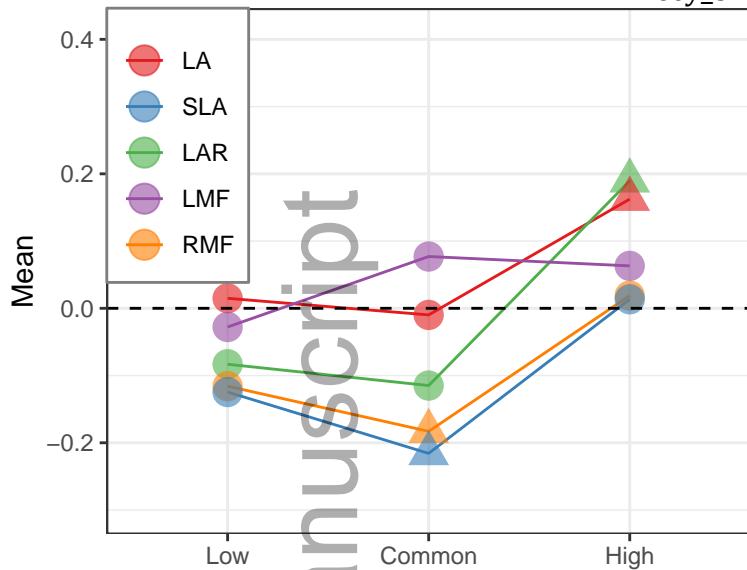


N



Canopy openness

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