Supporting Information

Methods

5 *Study sites*

Our forest plots were located in Sarapiquí County, Heredia Province, Costa Rica (Table S1). The regional life zone is tropical wet forest with annual temperature and rainfall averaging 26° C and \sim 3800 mm, respectively (1). Soils in the study

10 areas are derived from weathered basalt and are primarily classified as ultisols (2). Conditions at plots prior to abandonment were largely consistent although plots varied in the abundance of remnant trees and surrounding vegetation (3).

Trait data

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Whenever possible we sampled fully expanded sun-lit leaves with low levels of herbivory or epiphyll cover. For each tree, we collected small branches from the field and transported them to the lab in plastic bags. In the lab, we re-cut the stems and stored them in water to ensure that all tissues were equally hydrated. Samples

20 were stored in the dark at 4°C until measurements were made. Fresh weight, leaf size, leaf thickness and leaf toughness were measured within 24 hr. Leaf size was measured on a digital leaf area meter (LI-3100, LiCor Environmental, Lincoln, Nebraska). Leaf thickness was measured with a digital micrometer, and we used a leaf penetrometer to measure leaf toughness (punch force; Chatillion push-pull

- 25 gauge, Chatillion, USA). Dry leaf weight was measured after drying for ~72 hours at 60°C. Leaf density was calculated as the inverse of leaf thickness*leaf dry matter content*specific leaf area (4). Trait values were measured on two leaves per tree, and averaged prior to analyses. We measured wood specific gravity (WSG) on 1-51 individuals of 176 study species. We used a 5.15 mm
- 30 increment borer (Suunto, Finland) to core each tree from the bark to the pith. Samples were transported to the lab in plastic bags. After removing the bark, we measured wood core volume with the water displacement method and dry weight after \sim 72 hrs at 105 \degree C (5).

Leaf traits were measured on a total of 1,984 individuals (Table S2).

35 Wood specific gravity (WSG, unitless) was measured on 1,281 individuals of 176 species. We log-transformed leaf size, leaf density, SLA, leaf thickness, and leaf toughness because all were strongly right-skewed.

Demographic models

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Diameter effects

In order to control for spurious correlations between species identity and mean

DBH variation among species we standardized DBH to the mean for each species.

45 *Change in neighborhood trait diversity (NTD)*

In the main text we presented results on total change in NTD, i.e. change in diversity due to growth, mortality and recruitment. To further investigate the potential causes of changes in NTD, we also partitioned NTD into that arising from growth and stem turnover. NTD change due to neighbor growth is calculated

- 50 as (*NCIS* / *NCI* for a stems' neighbors at time *t* + 1) (*NCIS* / *NCI* for a stems' neighbors at time *t*), restricted to those neighbors surviving the interval. Thus when growth of neighbors increases neighbor trait diversity, NTD change is positive. Change in NTD due to growth was averaged for all stems in each plot, and plot averages were then averaged across years. Thus the unit of observation in
- 55 the t-test was each plot. Change in NTD due to stem turnover was computed similarly, with the exception that it was calculated as (*NCIS* / *NCI* for a stems' neighbors recruiting between time *t* and *t* + 1) - (*NCIS* / *NCI* for a stems' neighbors dying between time t and $t + 1$). Thus a negative value for change in NTD due to turnover indicates that recruiting stems were less functionally diverse
- 60 (standardized to their *NCI*) than dying stems.

Results

Survival model overview – We fit survival models for each of the traits (see

- 65 Methods). Mean annual stem survival varied between 96.1 and 96.4%, depending on the species included for each trait model. Posterior predictions had 93.4–94.0% accuracy in predicting survival vs. mortality, predicting survival accurately in 96.6–96.9% of cases (Table S3). Survival was significantly greater for large DBH individuals of a species compared to small individuals in all models (Table S4).
- 70 Survival also tended to be greater for individuals in less crowded neighborhoods (Table S5)

1. Trait relationships with average survival

75 Species with high leaf toughness and low leaf size had significantly greater average survival while other traits were not significant (Table S6).

2. Neighborhood crowding and changes in trait relationships with survival

- 80 No additional traits of focal trees significantly influenced species response to crowding (*β²* parameter, Table S7).
	- *3. Trait-mediated competitive dominance hierarchy*

85 Other traits did not show significant hierarchical dominance effects (Table S8, *β³* parameter).

4. Traits and niche variation among neighbors

- 90 Two other traits exhibited significant evidence for niche-based neighbor interactions. Focal trees exhibited significantly greater survival as *absolute* trait differences with neighbors (*NCIS*: trait difference weighted by *NCI*) increased for leaf size and leaf thickness (*β⁴* parameter, Table S9). Neighbor differences for other traits had no significant effects on survival. However, estimates for the
- 95 effect of *NCIS* on survival were positive in all cases.

5. Successional differences in trait diversity

Neighborhood trait diversity (*NTD*) increased with stand age for the remaining

100 functional traits. Leaf size (linear mixed effects model; t=5.8, p<0.0001), leaf thickness (t = 5.0, p<0.0001), leaf toughness (t = 6.6, p<0.0001) and leaf density $(t=3.9, p=0.0002)$ all showed significant increases in NTD from early- to midsuccessional stands to old-growth stands.

105 *Sources of change in neighborhood trait diversity (NTD)*

Increases in NTD with stand age were partly due to annual growth of surviving neighbors, with leaf size and thickness showing significant annual increases in NTD due to growth (one sample t-tests, see Table S10, Figure S2). In contrast, no

110 traits showed significant positive annual change in NTD due to stem turnover, and NTD for LDMC and WSG showed significant annual decreases due to stem turnover (Table S10, Figure S2).

References

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Tables

Table S1. Stand characteristics of eight 1-ha monitoring plots in northeastern

145 Costa Rica. The number of unique stems whose survival was modeled is included. This table is modified from Table 1 of Chazdon *et al.* (2010).

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Table S2. Summary statistics for traits and number of individuals and species

sampled.

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Table S3. Accuracy of posterior predictive simulations of survival (averaged across posterior simulations). Accuracy is further split into proportion of surviving trees predicted accurately and proportion of dying trees predicted

170 accurately. Trait abbreviations: LDMC = leaf dry matter content, SLA = specific leaf area, WSG = wood specific gravity.

Table S4. The slope of DBH effect on survival for the full model (eqn. 1 in the

- 180 main text), incorporating different traits, and showing median of posterior distributions with 95% CI in parentheses. β_5 is the average DBH effect across species; note that in our model the DBH effect varied randomly across species β_{5s} $\sim N(0, \sigma^2)$]. Also note that DBH effects were not modeled as a function of traits, or neighbors, so that similar DBH effects were expected across models. DBH
- 185 values were standardized within species. Note that all 95% CIs exclude zero. Trait abbreviations: LDMC = leaf dry matter content, SLA = specific leaf area, WSG = wood specific gravity.

195 **Table S5.** The slope of *NCI* effect, *β2.0* (see eqn. 4), on survival for models incorporating different traits, showing median of posterior distributions with 95% CI in parentheses. Entries in bold indicate 95% CIs that exclude zero. Trait abbreviations: LDMC = leaf dry matter content, SLA = specific leaf area, WSG = wood specific gravity.

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Table S6. Effect of interspecific trait variation on average survival rates β ^{*I*} (see eqn. 2). Posterior medians and 95% CIs are shown. Entries in bold indicate 95%

210 CIs that exclude zero. Trait values were standardized to mean zero and unit standard deviation. Trait abbreviations: LDMC = leaf dry matter content, SLA = specific leaf area, WSG = wood specific gravity.

*Model β¹ survival intercept*trait effect*

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Table S7. Effect of the interaction between NCI and traits on survival rates, β_2

- 225 (see eqn. 4). Negative values indicate greater sensitivity to NCI, *i.e.* reduced performance as NCI and the trait increase or reduced performance as NCI increases for specific habitat groups. Posterior medians and 95% CIs are shown. Entries in bold indicate 95% CIs that exclude zero. Trait values were standardized to mean zero and unit standard deviation. Trait abbreviations: LDMC = leaf dry
- 230 matter content, SLA = specific leaf area, WSG = wood specific gravity.

Model β² survival NCI sensitivity Leaf size 0.028 (-0.053, 0.131) LDMC -0.103 (-0.216, 0.02) SLA $0.013 (-0.089, 0.115)$ Leaf thickness $-0.006 (-0.098, 0.089)$ Leaf toughness $-0.008 (-0.097, 0.08)$ Leaf density $-0.016 (-0.103, 0.075)$ WSG 0.054 (-0.047, 0.155)

Table S8. Effect of crowding mediated by trait hierarchy of neighboring trees (NCH) , β_3 (see eqn. 1). Positive values indicate that as a neighbor's value of the trait decreases relative to the focal tree, the effect of crowding is reduced.

240 Posterior medians and 95% CIs are shown. Entries in bold indicate 95% CIs that exclude zero. Trait values were standardized to mean zero and unit standard deviation. Trait abbreviations: $LDMC = leaf$ dry matter content, $SLA = specific$ leaf area, $WSG =$ wood specific gravity.

Model β³ survival NCIH effect

Table S9. Effect of crowding mediated by trait similarity of neighboring trees (*NCIS*), β_4 (see eqn. 1). Positive values indicate that as neighbor trait differences

250 increase, the effect of crowding is reduced. Posterior medians and 95% CIs are shown. Entries in bold indicate 95% CIs that exclude zero. Trait values were standardized to mean zero and unit standard deviation. Trait abbreviations: $LDMC = leaf dry matter content, SLA = specific leaf area, WSG = wood specific$ gravity.

Model β⁴ survival NCIS effect Leaf size **0.157 (0.03, 0.301)** LDMC **0.166 (0.035, 0.328)** SLA **0.184 (0.075, 0.308)** Leaf thickness **0.147 (0.041, 0.262)** Leaf toughness 0.056 (-0.039, 0.169) Leaf density 0.098 (-0.008, 0.228) WSG 0.052 (-0.041, 0.164)

Table S10. Average plot change in NTD due to neighbor growth or turnover tested versus a null expectation of zero (one sample t-test, $N = 8$ plots, $df = 7$).

- 260 NTD change due to neighbor growth is calculated as: (*NCIS* / *NCI* for a stems' neighbors at time $t + 1$) - (*NCIS* / *NCI* for a stems' neighbors at time t), restricted to those neighbors surviving the interval. Thus when growth of neighbors increases neighbor trait diversity, NTD change is positive. Change in NTD due to growth was averaged for all stems in each plot, and plot averages were then
- 265 averaged across years. Thus the unit of observation in the t-test was each plot. Change in NRD due to stem turnover was computed similarly, with the exception that it was calculated as: (*NCIS* / *NCI* for a stems' neighbors recruiting between time *t* and $t + 1$) - (*NCIS* / *NCI* for a stems' neighbors dying between time *t* and *t* + 1). Thus a negative value for change in NTD due to turnover indicates that
- 270 recruiting stems were less functionally diverse (standardized to their *NCI*) than dying stems. Entries in bold indicate significant results. Trait abbreviations: $LDMC = leaf dry matter content, SLA = specific leaf area, WSG = wood specific$ gravity

Figures

Figure S1. Relationship between species traits and survival. Each species is

285 represented by a dot with size proportional to the square root of number of observations. A) Species (N=176) mean survival increases (y-axis) as WSG increases (x-axis). B) As leaf dry matter content (LDMC) increases (shown as dot color), species (N=214) mean survival increases (x-axis). Parameter values are plotted in model units. Lines show 95% CI for each species parameter, which is

290 drawn from a hyperdistribution.

- 295 **Figure S2.** The change in NTD due to (A) the growth of surviving neighbors or (B) replacement of dying neighbors by recruiting neighbors. NTD change due to neighbor growth is calculated as $(NCIS / NCI$ for a stems' neighbors at time $t + 1$) - (*NCIS* / *NCI* for a stems' neighbors at time *t*), restricted to those neighbors surviving the interval. Thus when growth of neighbors increases neighbor trait
- 300 diversity, NTD change is positive. Change in NTD due to growth was averaged for all stems in each plot, and plot averages were then averaged across years. Thus the unit of observation in the t-test was each plot. Change in NRD due to stem turnover was computed similarly, with the exception that it was calculated as $(NCIS/NCI$ for a stems' neighbors recruiting between time *t* and $t + 1$) - $(NCIS/NCI)$
- 305 *NCI* for a stems' neighbors dying between time *t* and *t* + 1). Thus a negative value for change in NTD due to turnover indicates that recruiting stems were less functionally diverse (standardized to their *NCI*) than dying stems.

A. Yearly change in NTD due to neighbor growth

