- 1 *Title*: Biomass and demographic dynamics of the Brazil-nut family (Lecythidaceae) in a mature
- 2 Central Amazon rain forest
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- 8 * Deceased
- 9 Keywords: biomass, growth, mortality, recruitment, habitat associations, flood disturbance
- 10 Highlights:
 - The number of Lecythidaceae individuals decreased by >1000, but basal area and biomass did not change significantly over 30 years.
- Mortality is highest in the lower-lying, wetter habitats.
- The most abundant species in the plot do not overlap in their habitat associations, nor do sister species.
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18 Abstract

19 In this study, we analyze the demographic rates and biomass dynamics of Lecythidaceae

- 20 in a 100-ha permanent forest plot in the Central Amazon. We characterize habitats in the plot,
- 21 analyze the effect of habitat on tree density, species richness, and biomass and demographic
- 22 dynamics, and quantify significant species habitat associations. The plot experienced a 13.2 %
- 23 net loss of Lecythidaceae individuals (n = 1023) but aboveground biomass (AGB; Mg ha⁻¹) did
- 24 not differ between the two censuses. Growth rates averaged 1.2 mm year⁻¹ with high intraspecific
- 25 variation and differed significantly among species height classes. Stem density and species
- 26 richness decreased in all six terra firme habitats and were significantly lower in both years in the
- 27 flood disturbed habitat. Only the flood disturbed areas experienced a significant loss in AGB,
- along with the highest annual mortality, recruitment, and growth rates across the plot. The lower-
- 29 lying habitats experienced higher mortality rates than the slopes and plateaus, suggesting too
- 30 much moisture is a higher risk to Lecythidaceae individuals than drought in this plot. There were
- 31 no overlapping habitat associations among sister species, or among the seven most abundant

species, which suggests that *terra firme* habitat association provides a niche partitioning
mechanism for closely related Lecythidaceae species. As intended with its establishment, this
plot continues to provide integral baseline data for of one of the most ecologically dominant and
culturally significant tree families in the Amazon region.

36 Introduction

37 Long term research on the demography and biomass dynamics of tropical tree species, 38 and on how these dynamics relate to environmental conditions, is critical to predict how tropical 39 forests will respond to global change. Such data are also needed to sustainably manage these 40 forests for forest products and biodiversity conservation (Davies et al., 2021). Baseline data for 41 trees in the Amazon basin is particularly important considering recent land use and climate 42 changes, combined with the region's outsize role in global carbon budgets (Laurance et al., 2000; 43 Longo et al., 2016). In permanent forest inventory plots tagged and geo-located trees provide 44 ecological information about demographic rates, life stage transitions (Pinero et al., 1984), 45 phenology (Lepsch-Cunha and Mori, 1999a), spatial dynamics, and responses to extreme 46 weather events (Feldpausch et al., 2016; Mori and Becker, 1991), among other processes. Forest 47 inventory plots can illuminate ecological processes that underlie forest composition and 48 dynamics at local scales (Zuleta et al., 2017), while providing opportunities for in depth studies 49 of focal clades (Baker et al., 2017; Lepsch-Cunha and Mori, 1999b; Mori and Lepsch-Cunha, 50 1995).

51 The Lecythidaceae family (Order Ericales) has considerable economic, cultural, and 52 ecological significance in the Amazon basin (Da Cunha and De Albuquerque, 2006; Mori and 53 Prance, 1990; Perreault, 2005; Prance et al., 1987) and is emblematic of old growth Amazon rain 54 forest (Guevara et al., 2017; Pitman et al., 2008; ter Steege et al., 2013). The Central Amazon has 55 particularly high Lecythidaceae diversity, resulting from a combination of higher endemism than 56 other regions and migration from other areas (Mori et al., 2016). Lecythidaceae ranks third in 57 family-level abundance in Amazon forests (following Fabaceae and Sapotaceae); it contains one 58 of the most abundant Amazon tree genera, *Eschweilera*, and one of Amazonia's most widespread 59 and abundant species, Eschweilera coriacea (Duque et al., 2017; ter Steege et al., 2020). The family has a disproportionately high number of hyperdominant species given its richness (ter 60 61 Steege et al., 2013). Lecythidaceae are dominant among a variety of forest strata, and E. coriacea 62 is the only species to be hyperdominant in all size categories and in all regions of the Amazon 63 (Draper et al., 2021). Four Lecythidaceae species are in the top 20 species contributing to 64 biomass in lowland Amazonia, including two of the top three - E. coriacea and the Brazil nut tree Bertholletia excelsa (Fauset et al., 2015). Because of its contribution to biomass, 65 66 Lecythidaceae is important in global biogeochemical processes such as carbon sequestration. 67 In the past several decades, the Amazon is thought to have acted as a carbon sink, with 68 aboveground biomass (AGB) of trees ≥10cm DBH (diameter at breast high) increasing in mature 69 forest (Baker et al., 2004a; De Castilho et al., 2010; Phillips et al., 1998). Additionally, large-70 stature, fast-growing trees, including two Lecythidaceae genera, Eschweilera and Corythophora, 71 have increased in density in some studies (Laurence et al. 2004). Many large-scale analyses 72 using networks of permanent plots and averaging across all clades have concluded that AGB is, 73 on average, increasing in the Amazon basin (Phillips and Lewis 2014; but see Wright 2005). 74 Information on how particular clades change in biomass, and what factors influence those 75 dynamics at the species level, is less well reported. For clades that have such an outsize 76 contribution to Amazon biomass dynamics, however, taxon-specific information can be 77 informative for conservation decisions and management.

78 Habitat factors may strongly influence demography and biomass dynamics at the species 79 level. Species may be associated with favored habitats based on physiological traits and 80 tolerances. An example is topographically influenced hydrologic variation (Zuleta et al., 2020), 81 which is one way in which species partition niche space, and may therefore contribute to 82 coexistence of closely related species (Harms et al., 2001). Species may be competitively 83 dominant and therefore more relatively abundant if they are better able to perform in certain 84 conditions (Harms et al., 2001). Habitat associations are known to change with life stage (Comita 85 et al., 2007), and demographic performance may not align particularly well with distribution-86 based habitat associations (Bin et al., 2016; Kenfack et al., 2014). This suggests that adult habitat 87 associations may not be stable, although the extent of their dynamism remains unclear. 88 Differences in dynamism are well documented at the Amazon-wide scale as a function of climate 89 and edaphic conditions (Phillips et al., 2004; Quesada et al., 2009). At the local scale, if certain 90 species, particularly those with higher wood density, are associated with habitats that are 91 changing due to precipitation and drought variation in the Amazon, an overall change in biomass 92 trends may occur.

93 Small-scale disturbance can also play a significant role in forest dynamics (Negrón-94 Juárez et al 2018). In terra firme forest, blow-downs (Negrón-Juárez et al., 2010; Nelson et al., 95 1994), lightning (Gora et al., 2020), and anomalous weather events (Mori and Becker, 1991) can 96 influence forest dynamics by increasing mortality and creating gaps which increase available 97 light. Flooding plays a large role in structuring tropical várzea and igapó forest, which are 98 defined by predictable annual flood pulses that filter species adapted to tolerate anoxic 99 conditions (Parolin et al., 2004). Flood disturbance in *terra firme* is uncommon, but standing 100 water in terra firme depressions has been documented (Kahn, 1987; Mori and Becker, 1991;

101 Myster, 2007; Rossano Mendes Pontes et al., 2012). The extent to which terra firme flooding 102 influences forest demographic and biomass dynamics remains an open question.

103 In this paper, we report on the dynamics of Lecythidaceae a large (100-hectare) 104 permanent forest plot north of Manaus, Brazil in which all individuals in the family were first 105 studied in the late 1980s and recensused for this study after 30 years. We analyze the effect of species, tree size, and height class on demographic rates. Additionally, we characterize habitats 106 107 in the plot, and analyze the effect of habitat on tree density, species richness, and biomass and 108 demographic dynamics. We quantify significant species habitat associations, and determine 109 whether they have remained stable over the past 30 years.

110 **Methods:**

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111 Study Site and plot:

112 The study site is part of the Biological Dynamics of Forest Fragments Project (BDFFP), 113 located approximately 70 km north of Manaus, Brazil. The 100-ha Lecythidaceae plot is located 114 in continuous forest Reserve 1501, or "KM 41", which is located on terra firme, well-drained 115 Tertiary sediments with primarily clayey latosols (oxisols), that are poor in nutrients (Fearnside and Filho, 2001; Mori et al., 2001). Mean annual temperature is 26.7° C, and mean annual 116 117 rainfall is 2325mm, with a dry season from July to September generally with less than 100mm of 118 rainfall (Bierregaard et al., 2001). Low-lying areas within the plot accumulate standing water 119 during years of unusually heavy precipitation; a flood was recorded in 1989, when >1m of 120 standing water was observed in 7 areas of the plot (Mori and Becker, 1991). 121 The Lecythidaceae plot was established to study the ecology and systematics of an

important Central Amazon tree clade, and to obtain precise baseline information on frequency,

123 density, dominance, and species richness that could be used to compare changes in composition and demography as a result of climate change and forest fragmentation (Mori et al., 2001). All
Lecythidaceae individuals ≥10 cm DBH were identified, measured, tagged, and mapped, and
vouchers were collected for each species and for any individual for which the determination was
uncertain. Detailed taxonomic work has been done on all species in the plot (Mori and LepschCunha, 1995), therefore identification errors are likely to be uncommon compared to other
locations.

130 Mori and colleagues recorded 38 species of Lecythidaceae in the initial census of the 131 100-ha plot. The most abundant species in the plot was the hyperdominant *Eschweilera coriacea*, 132 followed by *E. truncata*. The rarest species was the Brazil nut tree *Bertholletia excelsa*, which 133 was represented by a single individual. Eight genera are represented in the plot (Allantoma, 134 Bertholletia, Cariniana, Corythophora, Couratari, Eschweilera, Gustavia, and Lecythis), 135 although Eschweilera and Lecythis are non-monophyletic (Mori et al., 2007). All species were 136 assigned to one of three adult height classes by Mori & Lepsch Cunha (1995) determined by species maximum height at maturity: understory, which we will refer to as subcanopy (≤ 25 m), 137 138 canopy (25-35m), and emergent (>35m).

139 2018-2019 Census and taxonomic determinations:

We performed a complete recensus of the Lecythidaceae plot between February 2018 and March 2019. We attempted to locate all tagged Lecythidaceae trees from the original census, and DBH was measured at 1.3 or above buttresses. Trees were recorded as missing if their tags could not be found, unless a conspecific tree lacking a tag and with DBH equal to or larger than the missing tree was found in the same location (x, y coordinates of the 20x20 plot), in which case it was assumed to be the missing individual. 146 When a tree new to the 10cm size class was found, this recruit was identified, tagged, its 147 DBH was measured, and the tree was mapped by measuring the distance and the angle from the



Fig. 1: Top: map of the BDFFP, modified from Laurance et al. 2011. Red square shows approximate location of the "KM 41" Lecythidaceae plot. Bottom: map of the 6 habitats in the Lecythidaceae plot.

nearest 20x20m quadrat corner post. If a recruit could not be identified in the field, an herbarium 148

- 149 specimen was made and deposited at BDFFP for subsequent verification. Fertile specimens were
- 150 deposited at the INPA (Instituto Nacional de Pesquisas da Amazônia) herbarium.

Eighty individuals in the plot were sampled for DNA for studies on hybridization (Larson et al., 2021) and systematics (Vargas et al., in prep) of Lecythidaceae. The taxonomic

153 identification of eleven individuals in the plot, including three recruits (13.8% of the total



154 individuals sampled for molecular work) were changed. If a DNA collection was made from an

Fig. 2: Cladogram, with tips showing taxon names and species codes in parentheses. Clade shown on the right (Vargas et al., in prep). Height class shown by color of tip label.

155 individual in the plot, sequenced, and the subsequent phylogenetic analysis showed with high

confidence that that individual was in a clade other than its original identification, the
identification was changed. Two *Lecythis* species in the plot have not yet been formally
described because reproductive material has not yet been collected (Mori and Lepsch-Cunha,
1995). These undescribed species are designated *Lecythis sp.01* and *Lecythis sp.05* in a
Lecythidaceae monograph that outlines their vegetative characteristics (Mori and Lepsch-Cunha
1995; Mori et al. 2001).

Lecythidaceae are currently in the process of taxonomic re-circumscription based on
recent phylogenetic analyses, particularly *Lecythis* and *Eschweilera*, which are polyphyletic
lineages (Vargas et al. in prep). All *Eschweilera* species in the plot are in the Parvifolia clade
apart from *E. amazoniciformis*, which will form a monotypic genus in the revised taxonomy
(Vargas et al. in prep). The *Lecythis* species in the plot fall into three clades – Chartaceae (*L. gracieana*, *L. parvifructa*, *L. retusa*, *L. sp05*), Pisonis (*L. Pisonis*, *L. zabucajo*), and Poiteaui (*L. barnebyi*, *L. prancei*, *L. sp01*) (Fig. 2).

169 Demographic and biomass analyses:

A randomization analysis was used to determine whether stem density (individuals ha⁻¹), 170 species richness (species ha⁻¹), basal area (m² ha⁻¹) and biomass (Mg ha⁻¹) differed between the 171 172 two censuses. Hectare values were resampled with replacement 1000 times, and the mean values 173 of each resample were calculated. The differences between means in 1989 and in 2019 were 174 calculated, and if $\leq 2.5\%$ (two-tailed, $\alpha=0.05$) of the mean differences overlapped 0 then the 175 years were considered to be significantly different. All individuals surviving from the first to the 176 second census were used in growth analyses apart from those that decreased by more than 25% 177 of their initial DBH, which we considered to be measurement error (30 individuals), and the few broken trees for which only a resprout could be measured. Growth rate was calculated as (DBH_{t2}) 178

179 $-DBH_{tl}$ *10/(t_2 - t_1) and is expressed in (mm yr⁻¹). Relative growth rate was calculated as $(ln(DBH_{t2})-ln(DBH_{t1})/t$, expressed in cm cm⁻¹ yr⁻¹. For species-specific demographic and 180 181 biomass analyses, individuals not identified to species (Lecythis sp., Eschweilera "sagotiana") 182 were excluded, as were the two presumed hybrid individuals (E. wachenheimii x coriacea) (Larson et al., 2021). Annual percent mortality was calculated as $m = (1 - (N_s/N_0)^{1/t}) * 100$, where 183 N_s is the number of surviving stems, N_0 is the number of stems at the beginning of the census 184 185 interval, and t is the length of the census interval in years (Sheil et al., 1995). Annual percent recruitment was calculated as $r = 1 - (N_s/N_t)^{1/t} - 1) * 100$ where Ns is the number of surviving 186 187 stems, N_t is the number of stems in the second census, and t is the length of the census interval in 188 years. This is the final-density-based annual recruitment equation, in which recruits are assumed 189 to suffer the same mortality as pre-existing individuals, which offers symmetric comparisons to 190 annual mortality (Kohyama et al., 2018).

191 A linear model with post-hoc Tukey HSD tests to determine among which height classes 192 (subcanopy, canopy, emergent) and habitats mean annual growth rates differed. Linear 193 regression models were used to determine whether species percent mortality and recruitment 194 differed as a function of mean species DBH_{t1} and whether percent recruitment changed as a 195 function of abundance_{t1}. A Kruskal-Wallis test was used to determine whether there were 196 significant differences in annual percent mortality or recruitment among species height classes. 197 Individual probability of survival (1 = survival, 0 = mortality) as a function of size was analyzed 198 using a binomial generalized linear mixed effects model, with DBH_{t1} and adult height class as 199 fixed effects and species as a random effect.

200 Aboveground biomass was estimated using the R package "BIOMASS" (Rejou-Mechain 201 et al., 2017), which includes inputs of species-specific wood density, where possible, as well as 202 latitude/longitude coordinates which account for variation in height-diameter relationships as a 203 function of environment where height is not available (Chave et al., 2014). For 27 of the species, 204 these data were available from the TRY database (Adler et al., 2014; Chave et al., 2009; Falster 205 et al., 2015; Kattge et al., 2020; Paine et al., 2015; Wright et al., 2007; Zanne et al., 2009). When 206 more than one value was available for a given species, values were averaged, and the mean value was used to calculate biomass. For one species (Couratari tauari) where TRY values weren't 207 208 available but the "BIOMASS" package had species-level wood density values for the South 209 American Tropics, that value was utilized. When a species-level value was not available, the 210 generic average was used (13 species including all individuals not identified to species, hybrids, 211 and Eschweilera bracteosa, E. rankiniae, E. romeu-cardosoi, E. cyathiformis, E. 212 pseudodecolorans, E. laevicarpa, Lecythis barnebyi, L. retusa, L. sp01, and L. sp05). Analyses 213 were carried out in R version 3.6.3 (R CoreTeam, 2020). 214 Habitat definition and analyses: 215 Subplots of size 10x10m were assigned to one of six habitat types; flood disturbed

216 habitat, ravine, slope, concave, convex, and plateau, as modified from Kenfack et al (2014) based 217 on *a priori* knowledge of the topography, soil characteristics, and past flooding in the plot. 218 Cutoffs of 60m and 99m, 15°, and zero were used for elevation, slope, and convexity, 219 respectively. There is a distinctive plateau in the eastern part of the plot with geomorphological 220 characteristics separate from the rest of the plot (Mori and Becker, 1991), therefore is 221 categorized as its own habitat. There are also sandy ravines with small seasonal streams that 222 occur below 60m, the largest of which flows north to south in the middle of the eastern half of 223 the plot (Mori et al., 2001). Subplots were assigned to the flood disturbed habitat if more than half of the subplot fell within areas mapped as flooded by Mori & Becker (2001). While the 224

225 flood disturbed areas are largely nested within the concave habitat, we chose to categorize them 226 as a distinct habitat to examine the way that these heavily disturbed areas differ in demographic 227 and biomass dynamics from other low-lying areas in the plot. Concave and convex habitats are 228 distinguished by whether water is likely to be retained in a subplot (concave), or runoff into a 229 nearby subplot (convex). Ravine subplots have an elevation of <60m; slope subplots have a mean elevation \geq 60 and < 99m and a slope of \geq 15 degrees. Concave subplots have a mean 230 231 elevation of \geq 60m and <99m, slope < 15 degrees, and convexity \leq 0. Convex subplots have a 232 mean elevation of \geq 60m and < 99m, slope < 15 degrees, and convexity > 0. Plateau subplots 233 have a mean elevation of \geq 99m. Subplot slope and convexity were determined using the 234 fgeo_topography function in the "fgeo" package in R (Lepore et al., 2019).

235 Linear mixed effects models were used to analyze change in stem density, species 236 richness, and biomass over time among habitats. Density, richness, and biomass were modeled as 237 a function of fixed effect factors time and habitat, and a random effect of 10x10 subplot. A 238 bootstrapping analysis was used to detect differences in percent mortality and recruitment among 239 habitats. The number of 10x10 subplots that make up a given habitat were sampled randomly 240 with replacement from 10x10 subplots in that habitat, and annual percent mortality and 241 recruitment were calculated. This was done 10,000 times, and 95% confidence intervals for 242 mortality and recruitment were generated for each habitat. If these confidence intervals were not 243 overlapping, recruitment and mortality were determined to be significantly different in those 244 habitats.

A torus translation analysis (Harms et al., 2001) was used to identify species habitat associations for both censuses. Random translated coordinates which maintained the spatial relationships between individuals but shifted the x and y coordinates by a randomly generated number between the minimum and maximum plot coordinates were generated 9999 times. When an edge was reached, the coordinates wrapped around to the alternate side of the plot. For each translation, the relative abundance of each species in each habitat was calculated. If the observed relative abundance of a species in a habitat was greater than or less than the torus-translated relative abundance in that habitat \geq 97.5% of the time (α =0.5, two-tailed), there was determined to be a significant positive or negative habitat association, respectively.





Fig. 3: Mean density of individuals ha⁻¹ in DBH bins in 1989 (dark gray bars) and 2019 (light gray bars) for all individuals (top left), subcanopy species (top right), canopy species (bottom left), and emergent species (bottom right). Black bars represent standard error. Asterisks indicates a significant difference: p < 0.0005 (***), p < 0.005 (**), and p < 0.05 (*).

255 We recovered 6741 Lecythidaceae trees in the 2019 census, representing 37 species 256 (Table S1). Eschweilera coriacea remained the most abundant species, with 1369 individuals, 257 followed by E. truncata with 1166 individuals, and E. wachenheimii with 894 individuals. As 258 few as 33 and as many as 134 individuals were found in a hectare, and the average density of 259 Lecythidaceae per hectare was 67.3 ± 16.3 (mean \pm SD). This is a significant decrease (p<0.0005) from 1989, which had a mean density of 77.6±19.9 trees per ha. Species richness per hectare 260 261 ranged from 9 to 23, and the average species richness per hectare was 16.3 ± 2.5 , a significant (p < 262 .05) decline from 17.3±2.6 species in 1989. Two species (Cariniana micrantha and Allantoma 263 *lineata*) increased in abundance, *Lecythis sp.05* maintained its abundance, and the rest 264 experienced a net loss of individuals. Total plot basal area decreased slightly from 376.3 m^2 to 366 m². There was not a significant difference between mean basal area in 1989 (3.76±.086 m² 265 ha⁻¹) and in 2019 (3.66±0.92 m² ha⁻¹). 266

The total number of individuals in all DBH classes (10-15cm, 15-20cm, 20-30cm,>30cm) decreased between the two censuses (Fig. 3). The largest decrease was seen in the smallest DBH class across all species. Among all subcanopy species, there was an increase in individuals in all bins except 0-15cm. There were no increases among all canopy or emergent species in any DBH bin (Fig. 3).

272 Demography

Growth rates on average were low, with high intraspecific variation. Average growth
across all individuals (5681) during the 30 years between censuses was 36.9±32.6 mm
(mean±SD), or 1.2±1.1mm year ⁻¹, and average relative growth rate was .0056 cm cm⁻¹ yr⁻¹.
Average species annual growth rate ranged from 0.55±0.99 mm year ⁻¹ (*Lecythis barnebyi*,
subcanopy) to 3.12±2.4 mm year ⁻¹ (*Cariniana micrantha*, emergent), and annual relative growth



Fig. 4. Left: boxplot of annual relative growth (cm cm⁻¹ year⁻¹) by height class, non-shared letters represent significant differences. Right: boxplots of annual relative growth per species, all individual data points shown behind boxplots. Species code color represents height class: light gray = subcanopy, medium gray = canopy, black = emergent.

ranged from 0.009±0.006 cm cm⁻¹ yr⁻¹ (Allantoma decandra, emergent) to 0.003±0.005 cm cm⁻¹ 278 279 yr⁻¹ (L. barnebyi) (Fig. 4). Five out of the six species with the lowest relative growth rates were 280 subcanopy species, and the subcanopy height class had significantly lower mean annual relative 281 growth than either canopy or emergent height classes. There was not a significant difference in relative growth between canopy and emergent heigh classes, while emergents had higher mean 282 283 annual growth (cm yr⁻¹). 284 A total of 960 recruits (into the 10cm size class) were found during the 2019 census, 285 which is 0.51% annual recruitment. The riparian specialist Allantoma lineata had the highest 286 percent recruitment, with an initial abundance of 5 individuals and 3 recruits. Eight species had

287 zero recruits (that survived to the second census, *Eschweilera. micrantha, E. collina, E.*

288 rankiniae, E. carinata, Lecythis. barnebyi, L. sp01, Couratari tauari, Bertholletia. excelsa).

289 There was no relationship between species plot abundance in 1989 and percent recruitment

290 (p>0.1, $R^2 = -0.02$), and there are no significant differences in percent recruitment among height

classes (Fig. 5). Additionally, there is no relationship between annual percent recruitment and
mean DBH in 1989 (Fig. 5). There was no relationship between percent recruitment and percent
mortality among species (Fig. S1).

294 One quarter of the trees (1982 individuals, 25.54%) died in the thirty years between 295 censuses; an annual mortality rate of 0.98 percent. Mortality was more than double the



Fig 5. Top left: Species annual mortality as a function of mean DBH_{t1} . Regression indicates a significant relationship (p<0.05). Top right: Probability of survival as a function of DBH_{t1} for each height class. Binary survival data for each individual (survived = 1, died = 0) shown as points. Bottom Mean and standard error percent mortality (left) and recruitment (right) by height class.

recruitment, resulting in a net loss of 1023 individuals \geq 10cm DBH (13.2%). Percent mortality decreased as a function of mean species DBH_{t1} (R² = .16, p < 0.05, linear regression), while percent recruitment did not (Fig. 5).

A generalized linear mixed model was used to determine how the probability of survival changes as a function of DBH, and whether that relationship differs among height classes. The effect of DBH_{t1} on survival is more dramatic for subcanopy species than for canopy or emergent species (Fig. 5). There was no significant effect of DBH on survival for emergent species, and the relationship between size and survival differs significantly between subcanopy and both canopy and emergent species.



Fig. 6: Change in relative abundance (grayscale bars) and relative biomass (red hatched bars) by height class (left), and species (right). Species ordered left to right by decreasing abundance.

305 Biomass:

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306 Total aboveground living biomass (AGB) of Lecythidaceae during the first census (1989)
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307 was 5143.33 Mg, and during the second census (2019), it was 5129.13 Mg. There was not a
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- 308 significant decrease in mean biomass ha⁻¹, which was 51.4±14.2 Mg ha⁻¹ in 1989 and 51.3±15.5
- 309 Mg ha⁻¹ in 2019. Growth contributed an increase of 1430.87 Mg, mortality contributed a

310 decrease of 1544.95 Mg, and recruitment contributed an increase of 99.89 Mg. The aboveground 311 biomass of subcanopy species increased by 2.53 Mg (0.59% of initial subcanopy AGB), canopy 312 AGB increased by 14.61 Mg (0.39% of initial canopy AGB), and emergent AGB decreased by 313 33.45 Mg (3.45% of initial emergent AGB). Species with unknown height class contributed an 314 increase of 2.53 Mg of AGB. The highest changes in relative abundance were seen among abundant species in the plot (Fig. 6). While there was a net loss of both subcanopy and canopy 315 316 individuals, subcanopy species increased in relative abundance, in large part due to an increase in 317 relative abundance of E. wachenheimii. Canopy and emergent height classes experienced a 318 decrease in relative abundance. Relative biomass change, however, was distributed across rare 319 and common species, and changed most in emergent species. The largest increase in biomass 320 was contributed by E. coriacea, a canopy species and the most common species in the plot, 321 followed by Allantoma decandra, a relatively rare emergent species. The species with the largest 322 decrease in relative biomass was L. zabucajo (emergent), largely due to loss of a few very large 323 individuals, followed by E. truncata (canopy), the second most abundant species in the plot. 324 Lecythis zabucajo, while having the largest decrease in relative biomass in the plot, saw only a 325 small decrease in relative abundance.

Emergent species tended to have a much more significant change in relative biomass than relative abundance, while subcanopy species and most canopy species had the opposite. Subcanopy species had an overall increase in relative abundance and very small increase in relative biomass, while canopy and emergent species had a decrease in relative abundance. Canopy species increased in relative biomass, and emergents were the only height class to decrease in relative biomass. Emergent species account for only ~8% of individuals, but for ~ 18% of AGB in the plot.



Fig. 7. Change in density (individuals), richness (species), and AGB (Mg) per 10x10m subplot between the two censuses across habitats.

333 *Habitat*:

334 The concave habitat was the most common (32.66 ha), followed by the slope (25.07 ha), 335 convex (24.81 ha), plateau (10.17 ha), and flood disturbed (3.07 ha) habitats. Density of 336 individuals and species richness per 10x10m subplot decreased significantly between 1989 and 337 2019 in all habitats. Density in the flood disturbed habitat was significantly lower than the 338 ravine, concave, and convex habitats in 1989, and significantly lower than all habitats in 2019. 339 Richness was significantly lower in the flood disturbed habitat than all other habitats except for 340 the plateau in1989, and significantly lower than all habitats except for the ravine in 2019. The decrease between 1989 and 2019 in density and species richness was not different among 341 342 habitats. Biomass in 1989 was similar among all habitats. In 2019, the flood disturbed habitat 343 had significantly lower biomass than the slope, convex, and plateau habitats. The flood disturbed 344 habitat is the only one to have experienced a significant change in biomass – biomass decreased 345 significantly by 73.8 Mg AGB, or 60% of initial AGB.



Fig. 8: Annual percent recruitment, mortality, and relative growth across habitats. Mean and 95% confidence intervals shown. Habitats that do not share a letter represent significant differences among groups.

346 In the whole plot, relative mortality of the three height classes was proportional to their 347 initial relative abundances, while in the flood disturbed habitat a disproportionately high number of deaths were emergent individuals, and a disproportionately low number were subcanopy. 348 349 Mortality in the flood disturbed habitat was higher than in any other habitat. Additionally, 350 mortality in the ravine was significantly higher than on the plateau, and mortality in the concave 351 habitat was significantly higher than both slope and plateau habitats (Fig. 8). Bootstrapped 352 confidence intervals generated for annual percent recruitment of the flood disturbed habitat do 353 not overlap with any other habitat, showing that the flood disturbed habitat had significantly 354 higher recruitment than any other habitat (Fig. 8). The flood disturbed habitat also showed 355 significantly higher relative growth rates than all other habitats (Fig. 8). 356 All species with \geq 100 individuals (n=18) were analyzed for habitat associations in 1989 357 and in 2019 (Table S2). There were a total of 22 significant habitat associations in 1989 (12 358 positive, 10 negative), and 19 significant habitat associations in 2019 (10 positive, 9 negative). 359 The plateau had the largest number of habitat associations both in 1989 (7) and in 2019 (6), and 360 the ravine had the smallest, with none in 1989, and one in 2019. There were 7 changes in habitat 361 association between the two time points, 3 of which occurred in the plateau: three species that

362 were positively associated with a habitat in 1989 demonstrated no association with that habitat in 363 2019, two negative associations became neutral, one neutral relationship changed to a negative 364 association, and one neutral relationship changed to a positive association. In five of the seven 365 changes, however, the species relative abundance was greater or less than the null relative 366 abundance 95% of the time (α =0.1, two-tailed) (Figs. S2 & S3). The two above this significance level were Eschweilera pseudodecolorans losing a negative association with the concave habitat, 367 368 and E. pedicellata gaining a positive association with the plateau. Eschweilera coriacea, E. 369 cyathiformis, and E. pseudodecolorans had the highest numbers of significant associations in 370 1989 (3), and E. coriacea and E. cyathiformis remained the species with the highest number of 371 significant associations in 2019 (3). Eschweilera coriacea is the only species with two significant positive associations, which were consistent across years. The convex habitat is the only to show 372 373 no differences in associations over time (Table S2).

374 Discussion:

This study reports on the demography, biomass changes, and habitat associations of Lecythidaceae in mature Amazon rain forest. The intention with the initial establishment of the Lecythidaceae plot, over 30 years ago, was to provide baseline data to be monitored over time and compared to other plots (Mori et al., 2001). We determined how demographic and biomass dynamics vary among species, height classes, and habitats in the plot, and how species are partitioning environmental niche space.

381 Demographic dynamics

Growth rates in the Lecythidaceae plot are slow, averaging 1.2±1.1 (mean±SD) mm yr⁻¹.
In BDFFP, the average growth rates of trees above 10cm DBH in mature forests is 1.6 mm yr⁻¹
(Nascimento et al., 2005), which is slightly higher than the Lecythidaceae plot, but includes a

385 wider range of life history strategies including pioneer species. Long-term annual growth and 386 maximum (upper decile) growth is similar to Lecythidaceae species in other BDFFP inventory 387 plots (Chambers et al., 1998; Laurance et al., 2004). Growth is slower than the global wet tropical forest average of 4.7 mm yr⁻¹ (Locosselli et al., 2020), as this Central Amazon forest has 388 389 high wood density and slow turnover relative to forests with higher nutrient soils (Muller-390 Landau, 2004). There was large variation in relative growth among individuals of a given 391 species. This is expected as growth is highly dependent on site conditions of light, moisture, 392 herbivory, and soil (Chambers et al., 1998; Nascimento et al., 2005; Rüger et al., 2011). Growth 393 differs significantly among height class with subcanopy species showing slower relative growth 394 rates than canopy and emergent species. Canopy and emergent species did not differ in relative 395 growth, but emergent species had higher absolute growth.

396 Total recruitment in the plot lagged mortality, causing a net decrease in individuals. 397 Although there is evidence of increasing demographic turnover in the Amazon over the past 398 several decades (Phillips et al., 2008), a lower relative increase in recruitment than mortality has 399 caused a net decrease in number of stems since 1983 (Brienen et al., 2015), consistent with our trend. Plot recruitment was lower than Lecythidaceae recruitment in five hectares of undisturbed 400 401 upland forest in the same area over five years between 1981 and 1986 (Rankin-de-Merona et al., 402 1990). The same study also found higher recruitment than mortality among Lecythidaceae, while 403 we found the opposite to be true. The area in that study was much smaller, and therefore more 404 sensitive to environmental and demographic stochasticity. Alternatively, it may be possible that 405 Lecythidaceae are, on average, recruiting less successfully than they were 35 years ago, although 406 additional monitoring is required to confirm this. There is no difference in recruitment among 407 height classes, nor does percent recruitment change as a function of mean species DBH,

408 therefore a stature-recruitment trade-off is not detected among Lecythidaceae species, likely
409 because the smaller-statured species are still relatively long-lived – no species exhibit a short410 lived breeder functional type (Rüger et al., 2020).

411 Mortality was high relative to recruitment in our study; there was a net loss of >1000 412 individuals, in contrast to recent findings of increasing prevalence of the genera Eschweilera, Corythophora, and Couratari (Esquivel-Muelbert et al., 2019; Laurance et al., 2004). The 413 414 increase in other studies was attributed to taller statured and fast-growing species in these genera, 415 however many of the Eschweilera in our study are slow-growing subcanopy and canopy species 416 and therefore not likely to drive this increase. At the species level, mortality decreased as a 417 function of mean DBH, thus tends to be lower in larger-statured species (Fig. 5). At the 418 individual level, the decreased probability of survival with increasing DBH is characteristic of 419 tropical forests, where the highest risk of mortality occurs at the seedling stage (Uriarte et al. 420 2004; Iida, Poorter, et al. 2014; Ma et al. 2014) and is often U-shaped, with an uptick in 421 mortality risk at large sizes. The smallest individuals in the plot are 10 cm DBH, thus it would be 422 expected that survival probability decreases with DBH having already made it to the adult stage 423 (King et al. 2006; Rüger, Huth, et al. 2011; Iida, Kohyama, et al. 2014). Large trees may be more 424 prone pathogenic damage, experience increased water limitation, be at increased risk of 425 structural instability and windthrows, and experience higher reproductive costs (Coomes and 426 Allen, 2007; Iida et al., 2014; Mencuccini et al., 2005). Size has been shown not to have a strong 427 effect on probability of survival in larger-statured, non-pioneer species (Uriarte et al., 2004), and 428 mortality has been shown to be low at larger DBH for emergent species (Clark and Clark, 1992), 429 in line with the negligible decrease in probability of survival with increasing DBH for emergent 430 species that we found.

431 Habitat and demography:

432 A particularly impactful La Niña year occurred in 1989 after the plot was established, in 433 which standing water of up to 1m was seen in six areas in the plot (Mori and Becker 1991). 434 Relative growth rates in the flood disturbed habitat were significantly higher than in any other 435 habitat. The flooded habitat also showed the highest rates of recruitment and mortality, signifying increased dynamism in these disturbed areas. Relative to sandier valleys, terra firme 436 437 plateaus tend to have higher soil clay content, and therefore increased nutrient levels (Laurance 438 et al., 2006), which may account for the higher mean DBH and annual growth on plateaus than 439 the lower-lying habitats, although relative growth rates were similar among all non-flood-440 disturbed habitats. Alternatively, there may be more large-statured species on the plateau, 441 however only two emergent species were abundant enough to include in the habitat association 442 analysis. While the low concave habitat is likely more similar to the flood disturbed habitat in 443 soil properties, it experienced less disturbance and therefore did not have the same increased 444 light availability leading to high relative growth rates. 445 In addition to the flood disturbed habitat, the other habitats associated with increased

moisture or standing water, the ravine and the concave habitats, saw higher mortality than the 446 447 plateau, and the concave habitat had higher mortality than the slope (Fig. 8). While plateaus tend 448 to be at increased risk of seasonal moisture limitation, lower-lying streams and swampy areas 449 have been shown to increase mortality risk of seedlings (Johnson et al., 2017), and the same risks 450 of hypoxic conditions and soil erosion may similarly increase adult mortality. Additionally, there 451 may be an increased number of drought-resistant species on the plateau, mitigating moisture 452 limitation risk. In this plot, among Lecythidaceae individuals, too much moisture seems to be a 453 larger risk to survival than not enough.

454 High plot mortality cannot be explained entirely by the flooding event, however it did 455 increase overall mortality in these areas and differential relative mortality among size classes. 456 Rainfall data suggest 2008-2009, another intense La Niña year, may have been comparable to 457 1989 (Marengo et al., 2012). Water accumulation in low-lying *terra firme* is known to occur 458 elsewhere (Pearson and Derr 1986; Myster 2007), however, the effects on long-term 459 demographic and biomass dynamics are not well understood. El Niño and associated droughts 460 are often discussed as a significant source of disturbance in Amazonia (Laurance et al., 2004; 461 Wright, 2005), and while La Niña is also known to increase mortality in Amazonian forests 462 (Aleixo et al., 2019), La Niña-associated flooding warrants future study given its potential to 463 influence demography and biomass dynamics of Amazon forests.

464 Biomass dynamics:

Although there was a decrease in individuals in the plot, AGB remained relatively stable – growth and recruitment nearly balanced AGB loss from mortality. Lecythidaceae can be estimated to account for about 5.5% of species richness, 10.5% of individuals (De Oliveira and Mori, 1999), and about 15% of AGB in a hectare at BDFFP (Baker et al., 2004b). Thus, the contribution of Lecythidaceae to the density of individuals in the forest is disproportionately high given its species richness, and its biomass contribution is disproportionately high given its density.

472 Species that have changed the most in relative abundance are different from those that 473 have changed the most in relative biomass, the latter which tend to be emergent species. Biomass 474 change is due largely to mortality and growth of large trees. Although many small trees died and 475 recruited into the plot, these contributed relatively little overall to biomass dynamics over the 476 past 30 years. Biomass is highly correlated with stem density in the plot, with the most abundant 477 species contributing the most biomass. An exception is the large contribution of the one B. 478 excelsa to plot biomass in 1989, and the outsize contribution of some emergent species such as L. 479 zabucajo. Emergent species accounted for only 12.4% of the decrease in individuals, but for the 480 entire net decrease in ABG. There are no emergent species in the top 10 most common 481 Lecythidaceae species in the plot, and no emergent species have abundances above 200 individuals, but they account for nearly 20% of ABG in the plot. This outsize contribution to 482 483 biomass dynamics points to the importance of accounting for height class and life history when 484 considering conservation tactics, selective logging, and a forest's capacity to sequester carbon.

485 While the biomass decrease seen in the plot was small, it is contrary to a current trend of 486 positive net biomass change in trees ≥ 10 cm DBH in the Amazon over the past several decades 487 (Baker et al., 2004a). This trend varies, however, including at other BDFFP plots (Baker et al., 488 2004a), and the biomass increase has decelerated in the recent past (Brienen et al., 2015). Natural 489 disturbance regimes can lead to a decrease in AGB, including increased mortality in waterlogged 490 areas (Baker et al., 2004a; Ferry et al., 2010). The net decrease in AGB in flood-disturbed areas 491 is larger than the total net loss of AGB in the plot, and only flooded quadrats have significantly 492 decreased in biomass.

493 Sources of disturbance additional to flooding have been documented in the surrounding 494 areas during the time between censuses. These include blowdown events that caused high 495 mortality and biomass loss (Negrón-Juárez et al., 2010; Nelson, 2005; Nelson et al., 1994), as 496 well as two El Niño drought years. Given the large time interval (30 years) without intermediate 497 biomass observations, the magnitude of other disturbances is unknown, however they may have 498 contributed to the higher-than-average mortality and lack of increase in biomass across the 499 censuses. Additionally, if disturbance resulted in differential mortality of smaller trees, or subsequent decline in long-lived understory trees as blowdowns might be expected to, a decrease
in recruitment may be seen due to mortality of recruits before they enter the 10-cm size class
(Nelson, 2005). Given the slow growth of Lecythidaceae, it is likely that the time since
disturbance has not been long enough to cause the subsequent increase in biomass seen
elsewhere.

505 Individuals above 10cm contribute over 95% of AGB, therefore the biomass dynamics 506 that we observed based on individuals greater than 10cm should largely parallel overall 507 Lecythidaceae biomass dynamics in the plot (Chave et al., 2003). Because Lecythidaceae is a 508 dominant family in many parts of the Amazon, the decrease in biomass in this study, even if 509 slight, underlies the importance of continued monitoring.

510 Habitat Associations

511 A torus translation analysis illuminated 22 significant habitat associations in the first 512 census, and 19 in the second. There is no significant relationship between species abundance and 513 number of significant habitat associations, thus being common in the plot is not a function of 514 habitat generalization or specialization. Five of the 6 generalist species (no significant positive or 515 negative habitat associations) in 1989, and 4 of the 6 in 2019, were subcanopy species. This 516 suggests that species in the subcanopy height class may be less likely to specialize than taller 517 statured species, however we are limited in this analysis by the rarity of many species in the plot; 518 only two emergent species were abundant enough to include in the habitat association analysis. 519 The plateau located on the eastern part of the plot is the most dynamic habitat – it experienced 520 the highest number of changes in habitat associations. Habitat associations in the flood disturbed 521 habitat are similarly stable as other habitats (only one difference between the two censuses). This 522 similarity suggests associations in 2019 were not a function of shifting species composition due

to a single disturbance between censuses, thus flooding in these areas may be less anomalous than previously assumed. No associations changed from positive to negative, or vice versa – the changes were between neutral associations and a significant association, and 5 of the 7 neutral associations associated with those shifts were nearly significant, thus we conclude that habitat associations are generally stable in the plot.

528 *Eschweilera coriacea*, the most abundant species in the plot, is significantly positively 529 associated with the flood disturbed and the concave habitat and has a significant negative 530 association with the plateau. This demonstrates a preference for high moisture terra firme 531 habitats and an ability to tolerate flooding. It is known to occur in seasonally inundated várzea 532 habitat (Mori et al., 2010) therefore adaptations to low-oxygen conditions may contribute to its 533 ability to dominate in occasionally flooded terra firme forest. Eschweilera pseudodecolorans and 534 L. prancei were found to be plateau species in a forest divided into two habitat types - valley and 535 plateau/slope (Cosme et al., 2017; Oliveira et al., 2019), consistent with our finding them 536 positively associated with the plateau. Eschweilera coriacea has been defined as a generalist (not 537 restricted to either the valley or the plateau) (Oliveira et al., 2019), however its P_{50} (the water potential at which 50% loss of conductivity occurs) is similar to the mean P_{50} value among valley 538 539 individuals. Additionally, E. coriacea, analyzed as part of a species complex in the Parvifolia 540 clade, was found to dominate over other species in areas with increased topographic wetness 541 (Schmitt et al., 2021). While currently one of the most locally and regionally dominant lineages, 542 increasing drought risk associated with climate change may alter the composition of 543 competitively dominant species.

544 The second most abundant species in the plot, *E. truncata*, previously defined as a 545 plateau/slope species (Oliveira et al., 2019), is positively associated with the convex habitat, which has a higher mean elevation, higher runoff and likely less soil moisture than the flood disturbed or concave habitats. The two most abundant species in the plot, therefore, show nonoverlapping positive habitat associations. Additionally, none of the sister species pairs in the plot share habitat associations, and none of the seven most abundant species in the plot share positive habitat associations. This is in line with other studies demonstrating habitat specialization as a niche partitioning mechanism (de Oliveira et al., 2014), and likely contributes to the coexistence of many closely related species in the plot.

553 Conclusions

554 Lecythidaceae is a clade of slow growing, mature forest trees that have significant 555 economic and ecological value. Accurate demographic information for species, some which are 556 exploited for timber, is integral to sustainable logging operations. In a *terra firme* forest in the Brazilian Amazon, a post-logging growth increment of 4-5 mm yr⁻¹ with 1% mortality was 557 558 deemed necessary to sustain a 40 year reduced-impact logging cycle (Sist and Ferreira, 2007). Of 559 the three species in our study that are commonly logged (Cariniana micrantha, Eschweilera *coriacea* and *Couratari guianensis*) only *C. micrantha* has a 90th percentile growth rate (a proxy 560 for light gap growth) > 5 mm yr⁻¹, and none have a mean growth rate of > 4 mm yr⁻¹. 561 *Eschweilera coriacea*, has a 90th percentile growth rate $< 3 \text{ mm yr}^{-1}$. While dynamics and 562 563 therefore requirements for sustainability and regeneration may differ by forest, the slow growth 564 rates in our study suggests a 30-year logging cycle is not sustainable, even under high light 565 conditions. Our height class- and habitat-specific demographic information, particularly 566 mortality, may also be of use to improve the parameterization of simulation models used for 567 forest management (van Gardingen et al., 2006).

568	The Amazon, like many places in the world, is experiencing more extreme weather
569	events as a function of climate change. This may alter forest abiotic conditions such as moisture
570	availability; understanding species-habitat associations can allow us to consider how habitat
571	changes might influence forest composition and biomass dynamics. As intended with its
572	establishment, this plot continues to provide integral baseline data for a large area of mature,
573	undisturbed forest, that can be used to compare dynamics to the nearby fragmented forest, and to
574	other Amazon forests that are being impacted by human land use and climate change.
575	
576	
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