Foundation Species Across a Latitudinal Gradient in ² China

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Abstract

Foundation species play important roles in structuring forest communities and 41 ecosystems. Foundation species are difficult to identify without long-term observa-42 tions or experiments and their foundational roles rarely are identified before they are 43 declining or threatened. We used new statistical criteria based on size-frequency dis-44 tributions, species diversity, and spatial codispersion among woody plants to identify 45 potential ("candidate") foundation species in 12 large forest dynamics plots spanning 46 26 degrees of latitude in China. We used these data to identify a suite of candidate 47 foundation species in Chinese forests; test the hypothesis that foundation woody plant 48 species are more frequent in the temperate zone than in the tropics: and compare these 49 results with comparable data from the Americas to suggest candidate foundation gen-50 era in Northern Hemisphere forests. We identified more candidate foundation species 51 in temperate plots than in subtropical or tropical plots, and this relationship was in-52 dependent of the latitudinal gradient in overall species richness. Two species of Acer, 53 the canopy tree Acer ukurunduense and the shrubby treelet Acer barbinerve were the 54 only two species that met both criteria in full to be considered as candidate founda-55 tion species. When we relaxed the diversity criteria, Acer, Tilia, and Juglans spp., 56 and *Corlyus mandshurica* were frequently identified as candidate foundation species. 57 In tropical plots, the tree *Mezzettiopsis creaghii* and the shrubs or treelets *Aporusa* 58 *yunnanensis* and *Ficus hispida* had some characteristics associated with foundation 59 species. Species diversity of co-occurring woody species was negatively associated with 60 basal area of candidate foundation species more frequently at 5- and 10-m spatial grains 61 (scale) than at a 20-m grain. Conversely, Bray-Curtis dissimilarity was positively as-62 sociated with basal area of candidate foundation species more frequently at 5-m than 63 at 10- or 20-m grains. Our data support the hypothesis that foundation species should 64 be more common in temperate than in tropical or boreal forests, and suggest that in 65 the Northern Hemisphere that *Acer* be investigated further as a foundation tree genus. 66

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Keywords: Beta diversity, biodiversity, China, CForBio, codispersion analysis, forest 67 dynamic plots, ForestGEO, latitudinal gradient.

⁶⁹ Introduction

A foundation species is a single species (or a group of functionally similar taxa) that dom-70 inates an assemblage numerically and in overall size (e.g., mass or area occupied), determines 71 the diversity of associated taxa through non-trophic interactions, and modulates fluxes of 72 nutrients and energy at multiple control points in the ecosystem it defines (Ellison, 2019). 73 Because foundation species are common and abundant, they generally receive less attention 74 from conservation biologists, conservation professionals, and natural-resource managers who 75 emphasize the study, management or protection of rare, threatened, or endangered species 76 (Gaston and Fuller, 2007, 2008). However, protecting foundation species before they decline 77 to non-functional levels can maintain habitat integrity, thereby protecting associated rare 78 species at lower cost and less effort (Ellison and Degrassi, 2017; Degrassi et al., 2019). 79

Identifying foundation species is difficult because it can take many years—often decades— 80 to collect enough data to distinguish foundation species from other species that also are 81 common, abundant, or dominant (sensu Grime, 1987) but lack "foundational" characteristics 82 (Baiser et al., 2013; Ellison, 2014, 2019). Rather than investigating one common or dominant 83 species at a time in myriad ecosystems, Ellison and his colleagues have worked with data 84 from individual and multiple large forest dynamics plots within the ForestGEO network¹ to 85 develop statistical criteria that can suggest which tree species might merit further attention 86 as candidate foundation species in forests (Buckley et al., 2016a, b; Case et al., 2016; Ellison 87 et al., 2019). Specifically, Ellison et al. (2019) proposed two statistical criteria for candidate 88 foundation tree species based on their size-frequency and abundance-diameter distributions, 89 and on their spatial effects of on the alpha diversity (as Hill numbers: Chao et al., 2014) and 90 beta diversity (e.g., Bray-Curtis dissimilarity) of co-occurring species. 91

The first criterion is that candidate foundation tree species are outliers from the expected "reverse-J" size-frequency distribution observed in virtually all assemblages of co-occurring species (Loehle, 2006). The departure from expected size-frequency relationships reflects the

¹https://www.forestgeo.si.edu/

⁹⁵ abundance of foundation species and their relatively large sizes that lead to their dispropor⁹⁶ tionate influence on overall community structure. We refer to this criterion as the "outlier
⁹⁷ criterion".

The second criterion (the "diversity criterion") is that the size or abundance of candi-98 date foundation tree species should be negatively associated with species diversity (alpha 99 diversity) of other woody plants at local (small) spatial scales but positively associated with 100 species turnover (beta diversity) across large forest plots or stands (Ellison et al., 2019). The 101 negative spatial association between the size or abundance of foundation tree species with 102 local diversity of co-occurring woody species results simply from the foundation species occu-103 pying most of the available space in a standard 20×20 -m forest plot (or, in fact, any relatively 104 small plot). In contrast, the positive spatial association between the size or abundance of 105 a foundation tree species with beta diversity results from it creating patchy assemblages at 106 landscape scales. For example, forest stands dominated by foundation species such as Tsuqa 107 canadensis in eastern North America or Pseudotsuga menziesii in western North America 108 manifest themselves as distinctive patches on the landscape. When these foundation species 109 decline or are selectively harvested, the landscape is homogenized and beta diversity declines. 110 Indeed, Ellison et al. (2019) suggested that the preservation of landscape diversity may be 111 the most important reason to protect and manage foundation tree species before they decline 112 or disappear. 113

We emphasize that the application of these criteria to identify candidate foundation 114 species leads to the hypothesis that a particular taxon may be a foundation species, not that 115 it is one. Asserting that a species is a foundation species requires additional observational 116 and, ideally, experimental evidence (Ellison, 2014, 2019). Indeed, we derived these two 117 statistical criteria after more than a decade of observational and experimental studies of 118 Tsuga canadensis-dominated forests in New England, USA that lend strong support for the 119 hypothesis that T. canadensis is a foundation species (Orwig et al., 2013; Ellison, 2014). 120 These criteria subsequently were applied to five additional forest dynamics plots in the 121

western hemisphere (Buckley et al., 2016b; Ellison et al., 2019) with encouraging results. 122 Here, we apply these criteria to 12 large forest dynamics plots in China that range from 123 cold-temperate boreal forests to tropical rain forests. These plots are all part of the Chinese 124 Forest Biodiversity Monitoring Network (CForBio)², itself a part of the ForestGEO network. 125 There are two, fundamentally new contributions of this work. First, we test the hypoth-126 esis that foundation tree species should be uncommon or absent in subtropical and tropical 127 forests. Empirical support for particular trees having foundational roles in forests is strongest 128 for temperate forests (Schweitzer et al., 2004; Whitham et al., 2006; Ellison, 2014; Tomback 129 et al., 2016) and low-diversity tropical forests (Ellison et al., 2005), and Ellison et al. (2005) 130 hypothesized that foundation tree species would be less likely in species-rich tropical forests 131 (Ellison et al., 2019). Second, the application of our statistical criteria yield new insights 132 into ecological patterns and processes not only for China, but also concerning similarities 133 between the floras of East Asia and Eastern North America (Tiffney, 1985; Pennington et al., 134 2004). 135

136 Methods

¹³⁷ Forest dynamics plots in China

We used data from 12 of the 17 CForBio plots in our exportation of candidate foundation 138 species in Chinese forests (Table 1). These plots span 26 degrees of latitude and include: 139 the 9-ha broad-leaved Korean pine mixed forest plot at Liangshui in the Xiaoxing'an Moun-140 tains of Heilongjiang Province; the 25-ha Taxus cuspidata-dominated forest in the Muling 141 Nature Reserve, also in Heilongjiang Province; the 25-ha deciduous broad-leaved Korean 142 pine mixed forest plot on Changbai Mountain in Jinlin Province; the 20-ha warm-temperate 143 deciduous broad-leaved forest plot on Dongling Mountain in Beijing; the 25-ha subtropical ev-144 ergreen broad-leaved forest plot on Tiantong Mountain in Zhejiang Province; the 25-ha mid-145

²http://www.cfbiodiv.org

subtropical mountain evergreen and deciduous broad-leaved mixed forest plot on Badagong 146 Mountain in Hunan province; the 24-ha subtropical evergreen broad-leaved forest plot on 147 Gutian Mountain in Zhejiang Province; 20-ha lower subtropical evergreen broad-leaved forest 148 plot on Dinghu Mountain in Guangdong Province; the 25-ha cold-temperate spruce-fir forest 149 plot on Yulong Snow Mountain in Yunnan Province; the 25-ha karst evergreen and deciduous 150 broad-leaved mixed forest plot at Mulun in the Guangxi Zhuang Autonomous Region; the 151 15-ha karst seasonal rain-forest plot at Nonggang, also in the Guangxi Zhuang Autonomous 152 Region; and the 20-ha tropical forest plot at Xishuangbanna in Yunnan Province. 153

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[Table 1 about here.]

The 9-ha Liangshui plot ("LS"; 47.18 °N, 128.88 °E) was established in 2005. This plot 155 is located in the Liangshui National Reserve, which has been spared from logging and other 156 major disturbance since 1952 (Liu et al., 2014), and represents the climax vegetation type 157 of Northeast China (Xu and Jin, 2013). It is considered to be one of the most typical and 158 intact mixed broad-leaved-Korean pine forests in China. The plot has an elevational range 159 from 425 to 508 m a.s.l, a mean annual temperature of -0.3° C, and receives on average 676 160 mm of precipitation annually. In the first census in 2010, 21,355 individuals stems in 48 161 species, 34 genera, and 20 families were recorded. The average age of the overstory trees 162 was approximately 200 years (Liu et al., 2014). The "reverse-J" diameter distribution of 163 all individuals in LS suggested that the forest was regenerating well. The dominant tree 164 species at LS is *Pinus koraiensis*. Major associated tree species include *Tilia amurensis*, *T*. 165 mandshurica, Betula costata, and Fraxinus mandshurica (Xu and Jin, 2013). 166

The 25-ha Muling plot ("MLG"; 43.95 °N, 130.07 °E) was established in 2014 within the Muling Nature Reserve. The elevation within the plot varies from 658–781 m, the average annual temperature is -2° C, and the average annual precipitation is 530 mm. Muling is a typical middle-aged, multi-storied, uneven aged forest. Dominant tree species are *Tilia amurensis, Pinus koraiensis, Acer mono, Abies nephrolepis* and *Betula costata*. 63,877 individuals belonging to 22 families, 38 genera, and 57 woody species were recorded at the

¹⁷³ first census, including the nationally endangered *Taxus cuspidata* (Diao et al., 2016). The ¹⁷⁴ average DBH of all woody stems in MLG at the first census was 7.8 cm.

The 25-ha Changbai Mountain plot ("CB"; 42.28 °N, 128.08 °E), established in 2004, 175 was the first temperate forest dynamics plot in the ForestGEO network. It is considered 176 to be a typical old-growth, multi-storied, uneven-aged forest, and has neither been logged 177 nor suffered other severe human disturbances since 1960 (Wang et al., 2010). The average 178 annual temperature at CB is 3.6 °C and average annual precipitation is 700 mm. The terrain 179 of CB is relatively even, with elevations ranging from 791 to 809 m a.s.l. The height of the 180 main canopy species is ≈ 30 m, and the oldest trees are ≈ 300 years old. In the first census, 181 38,902 individuals in 52 species representing 32 genera and 18 families were recorded. The 182 most common species at CB are *Pinus koraiensis*, *Tilia amurensis*, *Quercus mongolica*, and 183 Fraxinus mandshurica (Hao et al., 2008). The most abundant eight species accounted for 184 83.4% of the total individuals in the plot (Wang et al., 2010). 185

The 20-ha Dongling Mountain plot ("DL"; 39.96 °N, 115.43 °E), established in 2010, is 186 in a warm temperate deciduous broad-leaved forest. The average annual temperature at DL 187 is $4.8 \,^{\circ}\text{C}$ and it receives 500–650 mm of precipitation each year. The mean elevation of the 188 plot is 1395 m, but the terrain is relatively steep with an elevation change of 219 m and 189 slopes ranging from $20-60^{\circ}$ (Liu et al., 2011). In the first census, 52,316 individuals in 58 190 species, 33 genera, and 18 families were recorded. The dominant species are all deciduous 191 trees, and include Quercus wutaishanica, Acer mono, and Betula dahurica (Liu et al., 2011). 192 The most common five species in the plot comprised 61% of all individuals, whilst the most 193 common 20 species comprised 92% of all individuals (Liu et al., 2011). 194

The 20-ha Tiantong plot ("TT"; 29.80 °N, 121.80 °E) represents a typical lower subtropical evergreen broad-leaf forest. It was established in 2009 within the core area of the Ningbo Tiantong National Forest Park. Mean annual temperature at TT is 16.2 °C and mean annual rainfall is 1375 mm. There have been some typhoon-caused landslides in some parts of the plot (Yang et al., 2011), but it is otherwise considered to be free from human disturbance

(Yan et al., 2018). Like Dongling Mountain, TT has a large elevational change across the
plot, ranging from 304 to 603 m a.s.l. In the first census, 94,603 individuals in 152 species,
94 genera, and 51 families were recorded. The dominant species are *Eurya loquaiana*, *Litsea elongata*, and *Choerospondias axiliaris* (Yang et al., 2011).

The 25-ha Badagong Mountain plot ("BDG"; 29.77 °N, 110.09 °E), established in 2011, is 204 located near the center of distribution of the oak genus Fagus. This plot is within the north 205 subtropical mountain humid monsoon climate; the average annual temperature is 11 °C and 206 average annual rainfall is 2105 mm (Lu et al., 2013). The dominant trees are a mixture 207 of evergreen (Cyclobalanopsis multinervis, C. gracilis, and Schima parvflora) and deciduous 208 species (Faque lucida, Carpinus fargesii, and Sassafras tzumu). During the first census, 200 186,556 individuals, belonging to 53 families, 114 genera, and 232 species were recorded 210 (Qin et al., 2018). There were 38 species with 1000 individuals, most in the shrub layer (Lu 211 et al., 2013). 212

The 24-ha Gutian Mountain plot ("GT"; 29.25 °N, 118.12 °E) was established in 2005 as representing a typical mid-subtropical evergreen broad-leaved forest (Legendre et al., 2009). Like the other montane plots, GT has a broad elevational range (446–715 m a.s.l.) with steep topography (slopes 12–62°). Average annual temperature at GT is 15.3 °C and average annual rainfall is 1964 mm. In the first census, 140,700 individuals in 159 species, 104 genera, and 49 families were recorded. Dominant species at GT include *Castanopsis eyrei* and *Schima superba* (Legendre et al., 2009).

The 25-ha Yulong Snow Mountain plot ("YLXS"; 27.14 °N, 100.22 °E), established in 2014, is at the highest elevation (3282 m a.s.l.) of the 12 plots we studied. Although the latitude of this plot is very low, the climate of this coniferous forest plot is cold-temperate because of its high elevation. The average annual temperature at YL is 5.5 °C and annual precipitation is 1588 mm (Huang et al., 2017). In the first census, 47,751 individuals in 62 species, 41 genera, and 26 families were recorded, dominated by *Berberis fallax* and *Abies forrestii* (Huang et al., 2017). The 25-ha Mulun plot ("ML"; 25.80 °N, 108.00 °E), also established in 2014, is within the Mulun National Natural Reserve. The mean annual temperature at ML is 19.3 °C, and the average annual rainfall is 1529 mm. The terrain of the plot is complex and varied. Rock exposure exceeds 60% and soil thickness j30 cm in most areas. In the first census, 108,667 individuals in 227 species, 147 genera, and 61 families were recorded (Lan et al., 2016). The dominant species are *Crytocarya microcarpa, Itoa orientalis, Platycarya longipes*, and *Lindera communis* (Lan et al., 2016).

The 20-ha Dinghu Mountain plot ("DH"; 23.10 °N, 112.32 °E), established in 2005, 234 has an average annual temperature of 20.9 °C and average annual precipitation of 1927 235 mm. This steep, subtropical every forest spans an elevational range of 230–470 m with 236 very steep slopes $(30-50^\circ)$. The first census recorded 71,617 individuals in 210 species, 119 237 genera, and 56 families (Ye et al., 2008). The three canopy-dominant species in the plot are 238 Castanopsis chinensis, Schima superba and Engelhardtia roxburghiana, whilst the sub-canopy 239 is dominated by Syzqium rehderianum and Craibiodendron scleranthum var. kwanqtungense 240 (Ye et al., 2008). 241

The 15-ha Nonggang plot ("NG"; 22.45 °N, 106.95 °E), established in 2011, is in a 242 hot-spot of biodiversity in China. This region is characterized by highly vulnerable and 243 spectacular limestone karst systems. Average annual temperature at NG is 21.5 °C and 244 average annual precipitation is 1350 mm. The first census recorded 66,718 individuals in 245 223 species, 153 genera, and 54 families (Lan et al., 2016). Eight of the recorded species 246 are protected throughout China, 30 are endemic to Guangxi province, and three were new 247 records for China. Representative tree species in NG include *Excentrodendron tonkinense*, 248 Cephalomappa sinensis, Deutzianthus tonkinensis, and Garcinia paucinervis. 249

The 20-ha Xishuangbanna plot ("XSBN"; 21.61 °N, 101.57 °E), established in 2007, is the southernmost CForBio site and is at the northern limit of typical southeast Asian tropical rain forests. It receives 1532 mm of precipitation annually and has an average annual temperature of 21 °C. The tropical seasonal rain forest in XSBN is one of the most

species-rich forest ecosystems in China. At the first census, 95,834 individuals in 468 species,
213 genera, and 70 families were recorded (Lan et al., 2008). The canopy height of this forest
is 50–60 m. The dominant emergent tree species is *Parashorea chinensis*. Subcanopy layers
of the forest are dominated by *Sloanea tomentosa, Pometia pinata*, and *Pittosporopsis kerrii*.

²⁵⁸ Tree census and measurement

Standard ForestGEO procedures (Condit, 1995) have been used to collect data across all CForBio plots. All woody stems (free-standing trees, shrubs, and lianas) at least 1 cm in diameter at breast height ("DBH"; 1.3 m above the ground level) were tagged, measured, identified to species, and mapped. In all of the plots, the individuals have been censused every 5 years (initial census years in these 12 plots varied between 2004 and 2014; Table 1); we used the first census data from each plot in our analysis.

²⁶⁵ The outlier criterion for identifying candidate foundation species

Following Ellison et al. (2019), our first criterion for selecting candidate foundation tree 266 species in each plot was to determine those species that were "outliers" from the typical 267 "reverse-J" distribution of the size-frequency distribution of mean DBH plotted against the 268 number of individuals. We identified outliers by eye rather than fitting a negative exponen-269 tial distribution with an arbitrary number of parameters to the 12 different size-frequency 270 distributions. This initial screen revealed 2–14 candidate foundation tree species in each of 271 the 12 forest dynamics plots (Fig. 1). The largest number of candidate species occurred in 272 MLG and the fewest were in YLXS. To screen species more expansively and avoid missing 273 other possible candidate foundation species, we also included in our first cut those ten species 274 with the highest importance values (IV = relative abundance + relative density + relative275 basal area) in each of the plots. Species that were outliers on the size-frequency plots usually 276 had high importance values, but including the latter did expand our initial pool of candidate 277 species to 10–14 species per plot (Table 2). 278

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[Figure 1 about here.]

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[Table 2 about here.]

²⁸¹ The diversity criterion for identifying candidate foundation species

The second, more stringent criterion for identifying candidate foundation species is a neg-282 ative association between its size (or abundance), and total abundance, three measures of 283 alpha diversity (species richness, Shannon diversity, Inverse Simpson Diversity) of associated 284 woody species and a positive association between its size or abundance and beta diversity 285 (Ellison et al., 2019). The three measures of alpha diversity treat all species identically 286 (species richness), down-weight rare species (Shannon diversity), or down-weight common 287 species (inverse Simpson diversity) within subplots. These associations also should be con-288 sistent across the plots when calculated at a given spatial grain (a.k.a. spatial scale) and at 289 most (ideally all) spatial lags (Buckley et al., 2016*a*; Ellison et al., 2019). 290

²⁹¹ Forest structure and species diversity indices

For each plot, we calculated the total basal area, mean basal area, and total number 292 of individuals of each of the candidate foundation tree and shrub species (Table 3) within 293 contiguous 5×5 , 10×10 , and 20×20 -m subplots. For species other than the candidate foun-294 dation species, we calculated their total abundance, species richness, Shannon and inverse 295 Simpson diversity indices (as Hill numbers: Chao et al., 2014) and mean Bray-Curtis dissim-296 ilarity (overall methods as in Ellison et al., 2019). In all the analysis, we used only the main 297 stem of each individuals (i.e., smaller stems of multi-stemmed individuals were excluded from 298 the analyses). The diversity() and vegdist() functions in the vegan package (Oksanen 299 et al., 2018) of the R software system (R Core Team, 2019) were used for calculating each 300 diversity metric. 301

302 Codispersion analysis

Codispersion describes anisotropic spatial patterns (i.e., different expected values when 303 measured in different directions) of co-occurring variables for given spatial lags and direc-304 tions (Cuevas et al., 2013). The codispersion coefficient ranges from -1 to 1, with positive 305 values indicating a positive spatial association and negative values indicating a negative spa-306 tial association for a given spatial lag and direction. These values can be visualized with 307 a codispersion graph (Vallejos et al., 2015). Buckley et al. (2016a) introduced codispersion 308 analysis to ecologists through an exploration of spatial patterns of species co-occurrence. 300 That paper also provides a basic introduction to the mathematics of codispersion analysis 310 and codispersion graphs. Buckley et al. (2016b) used codispersion analysis to examine spa-311 tial patterns of relationships between environmental characteristics and known or candidate 312 foundation tree species. Ellison et al. (2019) used codispersion analysis to quantify spatial 313 effects of candidate foundation tree species on different measures of diversity of associated 314 woody species in six forest dynamics plots in the Americas. 315

Although we computed codisperison patterns using mean basal area, total basal area, 316 and total abundance of candidate foundation species, we focus our presentation on the 317 codispersion between the total basal area of the candidate foundation species and associated 318 woody plant diversity in the differently-sized $(5 \times 5, 10 \times 10, \text{ and } 20 \times 20\text{-m subplots})$ 319 contiguous subplots in each of the 12 forest dynamics plots (Ellison et al., 2019); qualitatively 320 similar patterns were observed when using mean basal area or total numbers of individuals of 321 candidate foundation species. For each candidate foundation tree species, we first computed 322 the observed codispersion coefficient between its total basal area and abundance, alpha, and 323 beta diversity of the associated woody species in the subplots. The maximum spatial lag 324 examined for each plot ranged from the length of the subplot to one-fourth of the length 325 of the shortest side of each forest plot, which ensured adequate sample sizes for reliable 326 estimation of codispersion coefficients at the largest spatial lag (Buckley et al., 2016a). 327

328 Statistical significance of the codispersion coefficients was determined using null model

analysis (Buckley et al., 2016b; Ellison et al., 2019). Codispersion coefficients for all spatial 329 lags and directions were computed for co-occurrence matrices randomized using a toroidal-330 shift null model, which maintains the autocorrelation structure of the species and spatial 331 patterns caused by underlying environmental gradients while shifting the associated woody 332 species in random directions and distances (Buckley et al., 2016b; Ellison et al., 2019). For 333 each candidate foundation species in each plot, we ran 199 randomizations; significance 334 was determined based on empirical 95% confidence bounds. Calculation of codispersion 335 coefficients and all randomizations were done using custom C and R code written by Ronny 336 Vallejos and Hannah Buckley, respectively. 337

³³⁸ Data and code availability

Each of the CForBio plots were established at different times and are scheduled to be (or already have been) censused every five years. To maximize comparability among datasets, we used data collected at the first census for each plot (Table 1). All datasets are available from the ForestGEO data portal https://ctfs.si.edu/datarequest). R code for all analyses is available on GitHub (https://github.com/buckleyhannah/FS_diversity.

344 **Results**

³⁴⁵ Candidate foundation species in the CForBio plots

Only two species—the shrub *Acer barbinerve* (Fig. 2, 3) and the congeneric tree *Acer ukurunduense* (Fig. 4, 5)—in one plot—MLG—satisfied both the outlier *and* diversity criteria for all diversity measures for candidate foundation species (Figs. 2–5). For these two species in MLG, both criteria were met only at the 5-m spatial grain (Table 3).

[Figure 2 about here.]

[Figure 3 about here.]

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352	[Figure 4 about here.]
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354	[Table 3 about here.]

More species were considered as candidate foundation species when we retained the out-355 lier criterion (Fig. 1) but relaxed the diversity criterion to require only a positive spatial 356 relationship between the size of the candidate foundation species and beta diversity and a 357 negative spatial relationship between the size of the candidate foundation species and at 358 least one of the alpha-diversity measures (species indicated with an asterisk [*] in Table 3). 350 These additional candidate foundation species included two additional Acer species, tree 360 species in the genera Pinus, Taxus, Fraxinus, Quercus, and Tilia, and two shrubs (Corylus 361 mandshurica and Aporusa yunnanensis). However, whether we applied the stringent or re-362 laxed diversity criterion, all but three of the candidate foundation species occurred in plots 363 with cool- or cold-temperate climates. The exceptions were the trees *Pinus massoniana* and 364 Quercus serrata var. brevipetiolata at GT and Aporusa yunnanensis at DH; all three of these 365 species occurred in the subtropical evergreen broad-leaved forest plots. 366

A few of our initial candidate species that had high importance values but were not outliers from the expected size-frequency distributions (unstarred species in Table 2) did partially meet the diversity criterion in both temperate and tropical plots (Table 3). These included Syringa reticulata var. amurensis at CB, Juglans mandshurica at DL, Machilus thunbergii at GT, Brassaiopsis glomerulata at ML, Ficus hispida at NG, and Mezzettiopsis creaghii at XSBN.

³⁷³ Scale-dependence of candidate foundation species

More candidate foundation species—including all species that met at least one of the two criteria—were identified at smaller spatial grains: 15 species at the 5-m grain, 11 at the 10-m grain, and six at the 20-m grain (Table 3). This pattern applied both among and within the

plots. Average codispersion between total basal area of the candidate foundation species and
Bray-Curtis dissimilarity increased significantly with spatial grain (Fig. 6; raw data in Table
4). In contrast, average codispersion between total basal area of the candidate foundation
species and measures of alpha diversity, while generally negative, were more variable and not
scale-dependent (Fig. 6; raw data in Table 4).

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[Figure 6 about here.]

³⁸³ Candidate foundation species across a latitudinal gradient

Both the number of woody species in each plot that were outliers from the expected 384 size-frequency distribution and the number of candidate foundation species increased with 385 increasing latitude (Fig. 7A, C; slopes = 0.3 and 0.15 species/degree of latitude, respectively; 386 P (0.01). As expected, within-plot species richness declined significantly with latitude (slope 387 = -10.2 species/degree of latitude, P i 0.01), but this relationship was unrelated to the 388 latitudinal pattern in either the number of outliers or the number of candidate foundation 389 species. There were no significant relationships between either the number of outliers or the 390 number of candidate foundation species and within-plot species richness (Fig. 7B, D; P =391 0.08 and 0.18 respectively). 392

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[Figure 7 about here.]

Spatial association (expressed as codispersion) within each plot between candidate foundation species and total abundance, mean alpha diversities, and mean beta diversity of associated woody species on average did not vary with latitude at any spatial grain (Fig. 8; raw data in Table 4). Quantile regression (to account for potential extreme effects of foundation species) yielded similar results. There were no observed latitudinal patterns in effects of candidate foundation species except for a slight strengthening of the negative effect of candidate foundation species on associated woody species richness and total abundance

at the 5-m grain (Fig. 8, P = 0.04). Similar results were obtained when understory shrubs were excluded from the analysis (Fig. 9).

[Figure 8 about here.]

[Figure 9 about here.]

405 Discussion

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We applied two new statistical criteria (Ellison et al., 2019) to screen 12 of the 17 CForBio 406 Forest Dynamic plots in China for candidate foundation species. These 12 plots ranged 407 from 47 to 21 °N latitude, represented boreal, conifer-dominated, broad-leaved deciduous, 408 subtropical, and tropical forests (Table 1), and included two forest types referred to by 409 particular species ("Korean pine" mixed forests at Liangshi and Changbai Mountain, and the 410 "Taxus cuspidata" mixed conferous forest at Muling). Such eponyms do suggest traditional 411 or cultural-based knowledge of foundation (or other "important") species (Ellison et al., 412 2005; Ellison, 2019). Whilst both Korean pine (*Pinus koraiensis*) and *Taxus cuspidata* were 413 identified as candidate foundation species (Table 3), they were only candidates in the Muling 414 Taxus cuspidata-dominated forest plot, not in either of the "Korean pine" mixed forests. We 415 also found a strong latitudinal gradient, unrelated to the expected (and observed) underlying 416 latitudinal gradient in woody plant species richness, in the number of candidate foundation 417 species, which were more frequent in temperate than in tropical forest plots (Fig. 7). Where 418 they occurred, candidate foundation species had comparable effects at all latitudes (Figs. 419 8, 9), suggesting that foundation species effects more likely reflect specific combinations of 420 traits and interspecific effects rather than being manifestations of "neutral" (sensu Hubbell, 421 2001) processes (Ellison et al., 2019). 422

423 Candidate foundation species are more common in temperate lati-

424 tudes

Foundation species in forests control species diversity locally within forest stands and at 425 landscape and larger scales by creating habitat for associated flora (e.g., epiphylls, epiphytes, 426 vines, lianas) and modifying soil structure and composition (e.g., Ellison et al., 2005; Brant-427 ley et al., 2013; Baiser et al., 2013; Vallejos et al., 2018; Degrassi et al., 2019; Ellison, 2019). 428 Forest foundation species frequently are common, abundant, large trees (e.g., Schweitzer 429 et al., 2004; Ellison et al., 2005; Whitham et al., 2006; Tomback et al., 2016; Ellison et al., 430 2019), but understory shrubs and treelets also can have foundational characteristics (Kane 431 et al., 2011; Ellison and Degrassi, 2017; Ellison et al., 2019). Ellison et al. (2005) hypoth-432 esized that foundation species would be more likely in temperate forests because of their 433 relatively low species richness and more frequent dominance by one or a small number of 434 taxa. In contrast, tropical forests should lack foundation species as they are species and are 435 dominated less frequently by a small number of taxa. Our data supported this hypothesis: 436 candidate foundation species in the CForBio plots were more common at higher latitudes 437 than in the tropics (Fig. 7; Ellison et al., 2019). This pattern also may reflect the greater 438 importance of deterministic "niche" processes in temperate forests versus the stronger role 439 of "neutral" dynamics in tropical ones (Gravel et al., 2006; Qiao et al., 2015). 440

We hypothesize that tropical forests dominated by a one or a few closely-related species. 441 such as coastal mangrove forests dominated by *Rhizophora* spp. (Tomlinson, 1995) and mon-442 odominant tropical lowland forests dominated by species of Dipterocarpaceae in southeast 443 Asia or species of Leguminosae (subfamily Caesalpinioideae) in Africa and the Neotropics 444 (Torti et al., 2001; Hall et al., 2019) may be structured by foundation species (Ellison et al., 445 2005). Indeed, Gilbertiodendron dewevrei in the Ituri ForestGeo plot in the Democratic 446 Republic of Congo (Makana et al., 2004a, b) has functional characteristics similar to Tsuqa 447 canadensis in northeastern US forests. Gilbertiodendron casts deep shade; produces leaf 448 litter that decomposes very slowly, creating a dense and deep litter layer; creates soils with 440

 $\approx 30\%$ of the available nitrogen (ammonium + nitrate) relative to nearby mixed forests; and 450 has a depauperate (albeit not unique) fauna of leaf-litter ants and mites (Torti et al., 2001). 451 Analysis of species distribution and diversity associated with potential foundation species in 452 Southeast Asian forests dominated by Dipterocarpaceae, such as the ForestGeo 50-ha Pasoh 453 plot in Malaysia (Kochummen et al., 1991; Ashton et al., 2003) versus others lacking abun-454 dant dipterocarps, such as the 30-ha ForestGeo Mo Singto plot in Thailand (Brockelman 455 et al., 2011) or the 2-ha plot in Aluoi, Vietnam (Nguyen et al., 2016) would provide useful 456 comparisons with the analyses of the CForBio plots—especially the 20-ha Xishuangbanna 457 plot—presented here. 458

Conversely, the mid-latitude peak in functional-trait diversity of trees (Lamanna et al., 450 2014) led Ellison et al. (2019) to hypothesize that foundation tree species should be less 460 common in boreal forests at high latitudes or at high elevations in lower latitudes than 461 in more temperate ones. Our data showing no candidate foundation species at the high-462 elevation but low-latitude Yulong Snow Mountain plot support this hypothesis (Table 3). 463 In other high-elevation and high-latitude boreal ecosystems, foundation species tend to be 464 low-growing perennial, cushion- or tussock-forming plants (e.g., Ellison and Degrassi, 2017; 465 Elumeeva et al., 2017). 466

467 Foundation species effects are scale-dependent at landscape, not 468 local scales

Ellison (2019) argued that foundation species increase "patchiness" (beta diversity) at landscape scales, and that this effect of foundation species is of paramount importance when considering whether and how to conserve or otherwise manage them (see also Ellison et al., 2019). Across the 12 CForBio plots, we observed an increase in the strength of foundation species effects on beta diversity, expressed as a significant increase in codispersion between the candidate foundation species and diversity of associated species, at increasingly larger spatial grain (Fig. 6). At the 20-m grain, the magnitude of the codispersion coefficient

approached that of many of the candidate foundation species in ForestGeo plots in the Arra Americas (0.25–0.35; Fig. 6), but still less than the very strong effects of *T. canadensis* in Arra northeastern US forests (Ellison et al., 2019).

Conversely, although foundation species can provide habitat for associated species, thus 479 increasing their local diversity, the opposite pattern and magnitude of effects has been found 480 when analyzing only associated woody plant species in forest dynamic plots (Buckley et al., 481 2016a; Ellison et al., 2019) because foundation species occupy most of the available space. In 482 the CForBio plots, codispersion similarly was negative between candidate foundation species 483 and alpha diversity of associated woody plants (Figs. 2–6), but this relationship did not vary 484 significantly with spatial grain (Fig. 6). Additional data on faunal groups (e.g., Sackett et al., 485 2011; Record et al., 2018) or non-woody plants (e.g., Ellison et al., 2016) could provide a 486 test of whether these candidate foundation species have a positive effect on other associated 487 species that are not competing for space with canopy or subcanopy trees, but such data 488 are collected rarely in forest dynamic plots (but see Schowalter, 1994; Ruchty et al., 2001; 489 Ellison, 2018). 490

⁴⁹¹ Acer as a candidate foundation genus

In this study, four species of *Acer* were candidate foundation species among the three 492 cold-temperate plots in China (Liangshi, Muling, and Changbai: Table 3). Among these, A. 493 ukurunduense and A. barbinerve were the only two of all our candidate foundation species 494 that met the most stringent criteria for consideration. In a comparable study across a 495 latitudinal gradient in the Americas, A. circinatum was identified as a candidate foundation 496 species in the Wind River ForestGeo plot in Washington State, USA (Ellison et al., 2019). 497 We hypothesize that in many forests throughout the Northen Hemishphere, that Acer not 498 only can be a dominant genus in terms of abundance or total basal area, but that it may 499 function as a foundation genus, akin to *Quercus* in the Tyson ForestGEO plot in central 500 North America (Ellison et al., 2019). 501

Acer species often are common and abundant in temperate deciduous broad-leaved, conif-502 erous, and mixed forests throughout the Holarctic (Tiffney, 1985; Pennington et al., 2004), 503 and in subtropical montane forests in China (Xu, 1996). Acer includes 150 species (WFO 504 (World Flora Online), 2020), at least 99 of which (including 61 endemics) occur in China (Xu 505 et al., 2008) and more than a dozen are found in North America (Alden, 1995). Acer species 506 generally are shade tolerant, (i.e., they can regenerate and grow under closed canopies) and 507 have relatively high seedling and sapling survival rates (Tanaka et al., 2008). Some more 508 shade-intolerant ("photophilous") early-successional Acer species create conditions that fa-509 cilitate restoration of both later successional forests and their associated animal assemblages 510 (Zhang et al., 2010). 511

There are several forests named after Acer species in China, including the Acer mono-512 Tilia amurensis-T. mandshurica temperate broad-leaved deciduous forest, the Schima superba-513 Acer caudatum-Toxicodendron succedaneum eastern subtropical forest, and the Cyclobal-514 anopsis multinervis-Castanopsis eyrel var. caudata-Liquidambar acalycina-Acer sinense 515 forest in southwest China (Wu, 1995). Acer also are considered primary "companion" species 516 in Chinese Quercus and mixed broad-leaved-Korean pine forests where multiple Acer species 517 co-occur. For example, six-seven additional *Acer* species were recorded with the three can-518 didate foundation Acer species in the two broad-leaved-Korean pine mixed forests plots (LS, 519 CB). The nine Acer species in the CB plot account for 46% of the total stems (Zhang et al., 520 2010). 521

In North American forests, *Acer* species also define several forest types, including "Sugar Maple" (i.e., *A. saccharum*), "Sugar Maple–Beech–Yellow Birch", "Sugar Maple–Basswood", "Red Maple" (i.e., *A. rubrum*), and "Silver Maple–American Elm" (i.e., *A. saccharinum*) (Eyre, 1980). In forests of the Pacific Northwest of North America, the subcanopy treelet *A. circinatum* not only grows rapidly, has high biomass, and forms broad canopies that suppress other species (Lutz and Halpern, 2006; Halpern and Lutz, 2013), which causes it to have negative codispersion with other woody taxa (Ellison et al., 2019), but it also supports a high

diversity of epiphytes (Ruchty et al., 2001). Another North American species, A. sacchar-529 inum, dominates floodplain forests on well-drained alluvial soils in the eastern U.S. (Gabriel, 530 1990). Although Vankat (1990) subsumed "Silver Maple-American Elm" forests within a 531 "Mixed Hardwood Wetland Forest" type and considered A. saccharinum to be only a minor 532 component of these forests, this species historically was a significant constituent of at least 533 some primary forests in the upper Midwestern U.S. and Canada (Cho and Boerner, 1995; 534 Simard and Bouchard, 1996; Guyon and Battaglia, 2018); supports unique assemblages of 535 birds (Yetter et al., 1999; Knutson et al., 2005; Kirsch and Wellik, 2017); and, among woody 536 species, contributes substantially to carbon fixation in tidal wetlands (Milligan et al., 2019). 537 Acer saccharinum may be similar to other North American (candidate) foundation species 538 whose effects are most pronounced at different successional stages (Ellison et al., 2014, 2019). 530 However, we know of no large plots in either "Silver Maple–American Elm" or "Mixed Hard-540 wood Wetland" forests from which we could derive data to test whether A. saccharinum 541 meets our statistical criteria for candidate foundation species. Whilst it may be premature 542 to establish large forest dynamics plots in floodplains in either the temperate zone or the 543 tropics, or in tropical coastal habitats with low tree diversity, comparable data could be used 544 to test more general ideas about the foundational importance of particular genera, such as 545 Acer or Rhizphora, in forested wetlands worldwide. 546

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558 Author contributions

XQ and AME conceptualized and designed the study and wrote the manuscript; XQ and all the other authors collected the data at the individual CForBio plots. All authors contributed critically to the drafts and gave final approval for publication.

562 Conflict of interest

⁵⁶³ The authors declare no conflicts of interest.

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⁸⁰⁰ Supplementary Information

Table 4. Codispersion statistics for all candidate foundation species listed in Table 3.

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[Table 4 about here.]

Table 1: Geographic data for CForBio forest dynamics plots studied here. Latitude and longitude are in °N and °E, respectively; elevation is in meters above sea level (m a.s.l.); area is in hectares (ha), and census year is the year of the first census of the plot.

\mathbf{Plot}	Province	Lati-	Longi-	Ele-	Vegetation type	Area	Census
		\mathbf{tude}	\mathbf{tude}	va-		(ha)	Year
				\mathbf{tion}			
LS	Hei- longjiang	47.18	128.88	467	Broad-leaved Korean pine mixed forest	9	2010
MLG	Hei- longjiang	43.95	130.07	720	Taxus cuspidata-dominated mixed coniferous forest	25	2014
CB	Jinlin	42.38	128.08	802	Deciduous broad-leaved Korean pine mixed forest	25	2004
DL	Beijing	39.96	115.43	1395	Deciduous broad-leaved forest	20	2010
ΤT	Zhe- jiang	29.80	121.80	454	subtropical evergreen broad-leaved forest	20	2009
BDG	Hunan	29.77	110.09	1412	Mid-subtropical mountain evergreen and deciduous broad-leaved mixed forest	25	2011
GT	Zhe- jiang	29.25	118.12	581	subtropical evergreen broad-leaved forest	24	2005
YLXS	Yunnan	27.14	100.22	3282	Cool-temperate spruce-fir forest	25	2014
ML	Guangxi	25.80	108.00	550	Karst evergreen and deciduous broad-leaved mixed forest	25	2014
DH	Guang- dong	23.10	112.32	350	lower subtropical evergreen broad-leaved forest		2005
NG	Guangxi	22.45	106.95	260	Karst seasonal rain forest	15	2011
XSBN	Yunnan	21.61	101.57	789	Tropical rain forest	20	2007

Plot	Species	Abbreviation	Abundance	DBH	BA	Groups	IV (%)
LS	*Pinus koraiensis	PINKOR	1200	42.8	24.15	Canopy	23.3
	*Corylus mandshurica	CORMAN	4617	2.1	0.38	Shrub	10.5
	*Acer mono	ACEMON	2142	7.7	2.43	Canopy	8.3
	$A can tho panax\ senticos us$	ACASEN	1937	1.6	0.07	Shrub	5.1
	*Abies nephrolepis	ABINEP	905	16.2	3.01	Canopy	5.8
	* Tilia amurensis	TILAMU	728	13.4	3.01	Canopy	5.1
	*Acer ukurunduense	ACEUKU	1262	4.3	0.43	Canopy	4.4
	* Ulmus laciniata	ULMLAC	71	7.7	1.48	Canopy	4.7
	Euonymus pauciflorus	EUOPAU	1279	1.9	0.05	Shrub	4.1
	*Acer tegmentosum	ACETEG	988	5.1	0.49	Canopy	3.4
	*Betula costata	BETCOS	601	1302	2.04	Canopy	3.4
	*Fraxinus mandshurica	FRAMAN	407	12.54	1.27	Canopy	2.4
MLG	* Tilia amurensis	TILAMU	3540	19.4	6.14	Canopy	11.3
	*Acer mono	ACEMON	3893	12.4	3.45	Canopy	7.9
	*Corylus mandshurica	CORMAN	9008	2.0	0.13	Shrub	6.9
	*Abies nephrolepis	ABINEP	3596	14.1	3.07	Canopy	6.9
	*Pinus koraiensis	PINKOR	4820	9.1	2.15	Canopy	6.8
	*Acer barbinerve	ACEBAR	7498	3.2	0.32	Shrub	6.7
	*Acer tegmentosum	ACETAG	3904	8.7	1.42	Canopy	5.8
	*Betula costata	BETCOS	1573	18.0	2.43	Canopy	5.1
	*Acer ukurunduense	ACEUKU	3391	6.1	0.57	Canopy	4.7
	$*Syringa\ reticulata\ var. a murensis$	SYRRET	3926	3.6	0.36	Shrub	4.1
	* Ulmus laciniata	ULMLAC	1361	11.1	.98	Canopy	3.3
	*Fraxinus mandschurica	FRAMAN	961	14.4	1.01	Canopy	2.8
	Acer mandshuricum	ACEMAN	1553	6.9	0.59	Canopy	2.5
	*Populus davidiana	POPDAV	1172	11.8	1.01	Canopy	2.4
	Actinidia kolomikta	ACTKOL	2170	1.8	0.03	Liana	2.3
	Cerasus maximowiczii	CERMAX	1501	5.6	0.27	Canopy	2.2
	$A can tho panax \ senticos us$	ACASEN	1932	1.3	0.01	Shrub	2.1
	Philadelphus schrenkii	PHISCH	1317	1.5	0.01	Shrub	1.8
	Lonicera ruprechtiana	LONRUP	1021	1.5	0.01	Shrub	1.7
	Aralia elata	ARAELA	1506	2.9	.05	Shrub	1.5
	* Taxus cuspidata	TAXCUS	172	39.4	0.93	Canopy	1.5
CB	* Tilia amurensis	TILAMU	2927	31.3	12.31	Canopy	14.8
	*Pinus koraiensis	PINKOR	2468	32.6	9.79	Canopy	12.4
	*Acer mono	ACEMON	6609	7.5	2.69	Canopy	10.6
	*Corylus mandshurica	CORMAN	7834	1.7	0.08	Shrub	9.6
	*Acer pseudosieboldianum	ACEPSE	5984	6.1	1.1	Canopy	8.6
	*Quercus mongolica	QUEMON	926	41.3	6.5	Canopy	8.0
	*Fraxinus mandshurica	FRAMAN	681	47.9	5.81	Canopy	6.7
	*Acer barbinerve	ACEBAR	3911	2.3	0.08	Shrub	5.9
	* Ulmus japonica	ULMJAP	1109	14.1	1.81	Canopy	4.3
	Syringa reticulata var. amurensis	SRYRET	1598	3.8	0.09	Shrub	3.0
DL	*Quercus wutaishanica	QUEWUT	5274	18.9	7.36	Canopy	19.6
	*Acer mono	ACEMON	10539	5.7	1.32	Canopy	12.1
	*Betula dahurica	BETDAH	2536	17.5	3.03	Canopy	9.6
	*Syringa pubescens	SYRPUB	6313	3	0.22	Shrub	6.9
	*Abelia biflora	ABEBIF	5174	2.4	0.23	Shrub	6.0
	*Corylus mandshurica	CORMAN	6192	12.6	0.14	Shrub	5.8
	*Rhododendron micranthum	RHOMIC	71301	2.9	0.04	Shrub	1.8
	Populus davidiana	POPDAV	1967	18.8	1.22	Canopy	5.0
	Betula platyphylla	BETPLA	776	5.5	1.07	Canopy	3.8
	*Fraxinus rhynchophylla	FRARHY	2385	19.5	0.28	Canopy	3.8
	Juglans mandshurica	JUGMAN	576	8.7	0.20	Canopy	2.9
TT	*Eurya loquaiana	EURLOQ	20414	2.6	16.46	Shrub	12.1
1 I	*Litsea elongata	LITELO	10395	2.0 4.8	34.95	Canopy	8.2
	Choerospondias axillaris	CHOERO	1352	4.8 20.8	54.95 76.04	Canopy	0.2 6.5
	-				40.04	Shrub	
	*Distylium myricoides *Lithogarnus henrui	DISMYR LITHEN	6298 2688	6.3			6.4 5.5
	*Lithocarpus henryi *Cuelobalanonsis muhium	LITHEN	2688 2484	12.2	53.91	Canopy	5.5
	*Cyclobalanopsis nubium *Camellia fraterna	CYCNUB CAMFRA	2484 0270	12.4	55.11 6.45	Canopy Shrub	5.5 5.4
	- Oamenia malema	UAMPRA	9279	2.4	6.45	Snrub	5.4
	Schima superba	SCHSUP	1237	17.3	40.67	Canopy	3.8

Table 2: Initial set of candidate foundation species identified as outliers in the abundance-DBH plots (Fig. 1; here marked with an asterisk[*]) and others whose importance values IV) were in the top ten for that plot. Plots are ordered by latitude, and within each plot, species are ordered by IV. Units of diameter (DBH) are cm and units of basal area (BA) are in m^2/ha .

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Plot	Species	Abbreviation	Abundance	DBH	BA	Groups	IV (%)
	Castanopsis fargesii Maabilaa thaabaasii	CASFRA	750	22.7	42.81	Canopy	3.1
BDG	Machilus thunbergii	MACTHU CYCMUL	2807 11503	5.8	18.81	Canopy	2.9
bDG	*Cyclobalanopsis multinervis *Fagus lucida	FAGLUC	2769	13.1	$5.36 \\ 3.84$	Canopy Canopy	o 4.:
	Cyclobalanopsis gracilis	CYCGRA	5322	6.2	2.53	Canopy	3.8
	Carpinus fargesii	CARFAR	2172	11.1	1.97	Canopy	2.4
	Schima parviflora	SCHPAR	1796	9.5	1.86	Canopy	2.5
	Sassafras tzumu	SASTZU	453	30.6	2.19	Canopy	2.5
	Castanea seguinii	CASSEG	471	28.6	1.9	Canopy	1.9
	Betula insignis	BETINS	434	27.7	1.64	Canopy	1.
	Quercus serrata	QUESER	703	20.7	1.53	Canopy	1.
	*Eurya brevistyla	EURBRE	16051	3	0.81	Shrub	5.
	Nyssa sinensis	NYSSIN	805	17.9	1.4	Canopy	1.
	*Rhododendron stamineum	RHOSTA	9549	9.1	4.36	Canopy	6. C
TT.	*Litsea elongata	LITELO	21035	2.6	0.96	Canopy	6.
Τ	*Castanopsis eyrei *Schima superba	CASEYR SCHSUP	$12406 \\ 8514$	$\begin{array}{c} 11.9 \\ 10.4 \end{array}$	$12.52 \\ 6.89$	Canopy Canopy	15. 9.
	*Pinus massoniana	PINMAS	2061	$10.4 \\ 18.7$	4.23	Canopy	9. 4.
	*Cyclobalanopsis glauca	CYCGLA	1620	10.4	0.88	Canopy	4.
	Daphniphyllum oldhami	DAPOLD	2718	6.9	0.00 0.79	Shrub	2.
	Machilus thunbergii	MACTHU	1384	8.6	0.76	Canopy	1.
	Cyclobalanopsis myrsinaefolia	CYCMYR	375	17	0.59	Canopy	0.
	*Quercus serrata var. brevipetiolata	QUESER	3508	10.6	1.63	Canopy	2.
	$*Rhododendron \ ovatum$	RHOOVA	10793	3.9	0.72	Shrub	4
	Loropetalum chinense	LORCHI	4461	5	0.64	Shrub	2
/LXS	*Berberis fallax	BERFAL	28416	1.4	0.2	Shrub	68
	*Abies forrestii	ABIFOR	5207	19.6	15.18	Canopy	51
	Quercus guajavifolia	QUEGUA	1324	45.1	12.92	Canopy	36
	Picea likiangensis	PICLIK	596	50.8	9.03	Canopy	25
	Gamblea ciliata var. evodiifolia	GAMCIL	1065	25.7	2.67	Canopy	14
	Acer pectinatum	ACEPEC SORPRA	$958 \\ 915$	23.3	2.01	Canopy Shrub	11
	Sorbus prattii Viburnum betulifolium	VIBBET	915 1114	$ 11.8 \\ 3.8 $	$\begin{array}{c} 0.49 \\ 0.06 \end{array}$	Shrub	8 8
	Rhododendron yunnanense	RHOYUN	1162	11	$0.00 \\ 0.67$	Shrub	7.
	Padus brachypoda	PADBRA	328	28.5	1.04	Canopy	5
ΛL	*Cryptocarya microcarps	CRYMIC	31939	4.2	2.97	Canopy	15.
	*Itoa orientalis	ITOORI	3050	9.9	1.35	Canopy	4
	$*Lindera \ communis$	LINCOM	4192	4.1	0.33	Shrub	3
	$Eury corymbus\ cavaleriei$	EURCAV	1525	10.6	0.76	Canopy	2
	*Platycarya longipes	PLALON	3520	6.6	0.85	Canopy	2
	Brassaiopsis glomerulata	BRAGLO	2910	4.3	0.27	Canopy	2
	Radermachera sinica	RADSIN	965	9.8	0.43	Canopy	1
	Boniodendron minus	BONMIN DIODUM	$1521 \\ 2590$	$6.2 \\ 3.4$	0.35	Shrub	1
	Diospyros dumetorum Rubovietnamia aristata	RUBARI	2590 1614	$\frac{5.4}{4.7}$	$\begin{array}{c} 0.14 \\ 0.16 \end{array}$	Canopy Shrub	1
н	*Castanopsis chinensis	CASCHI	2311	24.4	9.3	Canopy	12
DH	*Schima superba	SCHSUP	2311 2296	$\frac{24.4}{18.9}$	$9.3 \\ 4.13$	Canopy	6
	*Engelhardtia roxburghiana	ENGROX	737	28.8	3.48	Canopy	4
	*Machilus chinensis	MACCHI	532	16.3	$0.10 \\ 0.85$	Canopy	1
	*Syzygium rehderianum	SYZREH	5990	4.7	0.88	Shrub	4
	$*Grarbiodendron\ scleranthum$	GRASCL	3325	8.6	1.66	Canopy	4
	*Aidia canthioides	AIDCAN	5996	2.3	0.21	Shrub	4
	$*Cryptocarya\ chinensis$	CRYCHI	2557	6.5	1.21	Canopy	3
	Cryptocarya concinna	CRYCON	4478	1.8	0.18	Canopy	3
	*Aporusa yunnanensis	APOYUN	3747	4.7	0.44	Shrub	3
	Ardisia quinquegona	ARDQUI	3702	1.9	0.08	Shrub	2
	Blastus cochinchinensis	BLACOC	4011	1.6	0.06	Shrub	2
	*Acmena acuminatissuma	ACEACU	1484	8.7	1.1	Canopy	2
IG	* Cleistanthus sumatranus	CLESUM	9977	4.8	2	Canopy	9
	*Sterculia monosperma *Vitor kwanasioneje	STEMON VITKWA	$6328 \\ 2470$	5.7 11.4	1.59	Canopy	7
	*Vitex kwangsiensis	VITKWA EXCTON	$2470 \\ 1502$	$ \begin{array}{c} 11.4 \\ 6.5 \end{array} $	$1.44 \\ 0.77$	Canopy Canopy	$6 \\ 2$
	Hacentrodendron tonkingenes		1002	0.0	0.11		2
	Excentrodendron tonkinense					- •	ი
	Diplodiscus trichosperma	DIPTRI	1126	8.1	0.49	Canopy	
						- •	2 2 2

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Plot	Species	Abbreviation	Abundance	DBH	$\mathbf{B}\mathbf{A}$	Groups	IV (%)
	Ficus hispida	FICHIS	2989	3.2	0.28	Shrub	2.1
	Garcinia paucinervis	GARPAU	1684	3.4	0.25	Canopy	2.1
XSBN	*Pittosporopsis kerrii	PITKER	20918	3.2	1.42	Shrub	25.8
	*Parashorea chinensis	PARCHI	7919	5.2	5.68	Canopy	22.4
	$Castanopsis\ echidnocarpa$	CASECH	1679	12.8	2.47	Canopy	7.1
	Garcinia cowa	GARCOW	4333	5.1	0.96	Canopy	7.3
	Mezzettiopsis creaghii	MEZCRE	3300	6.2	1.26	Canopy	6.9
	Baccaurea ramiflora	BACRAM	3212	5.2	0.7	Canopy	5.5
	Knema furfuracea	KNEFUR	3160	4	0.56	Canopy	5.1
	Saprosma ternata	SAPTER	2698	1.9	0.05	Shrub	4.7
	Phoebe lanceolata	PHOLAN	2409	3.6	0.22	Canopy	3.5
	Cinnamomum bejolghota	CINBEJ	1337	5.4	0.44	Canopy	3.9

Table 3: A winnowed list of candidate foundation tree and shrub species (the latter indicated by a plus sign [+]) at three different spatial grains (i.e., subplot size) in 12 Chinese forest dynamics plots. Plots are ordered by latitude, and within each plot, candidate foundation species are ordered alphabetically. The two *Acer* species in **bold type** satisfied all aspects of both the outlier and the diversity criteria for candidate foundation species at the given spatial grain. The starred (*) species satisfied the outlier criterion (Fig. 1) and partially satisfied the diversity criterion at the given spatial grain: a positive spatial relationship between candidate foundation species size and beta diversity, and a negative spatial relationship between candidate foundation species size and at least one measure of alpha diversity. The remaining species did not satisfy the outlier criterion but did meet some aspects of the diversity criterion. No species met either foundation species criterion in the BDGS, TTS and YLXS plots at any spatial grain.

		Spatial grain	
\mathbf{Plot}	5 m	10 m	20 m
LS	*Acer ukurunduense	*Acer ukurunduense	_
	$*Corylus mandshurica^+$		
	*Fraxinus mandshurica		
MLG	$*Acer \ barbinerve^+$	*Acer barbinerve+	
	*Acer tegmentosum		
	*Acer ukurunduense		
	$*Corylus mandshurica^+$		
		*Pinus koraiensis	*Pinus koraiensis
	* Taxus cuspidata		_
	* Tilia amurensis	* Tilia amurensis	* Tilia amurensis
CB	*Acer barbinerve+		
	*Acer pseudosieboldianum	*Acer pseudosieboldianum	
	$*Corylus mandshurica^+$	$*Corylus mandshurica^+$	_
	Syringa reticulata var. $amurensis^+$	Syringa reticulata var. amurensis ⁺	
DL	Juglans mandshurica		
TT			
BDG			
GT	Machilus thunbergii		
			*Pinus massoniana
			*Quercus serrata var. brevipetiolate
YLXS	_		_
ML	Brassaiopsis glomerulata	Brassaiopsis glomerulata	
DH	*Aporusa yunnanensis +	*Aporusa yunnanensis+	$*A porusa yunnan ensis^+$
NG	Ficus hispida+	Ficus hispida ⁺	
XSBN	Mezzettiopsis creaghii	Mezzettiopsis creaghii	Mezzettiopsis creaghii

Table 4: Codisperison statistics for the candidate foundation tree or understory species (the latter indicated by a [+]) in each plot at the spatial grain (**Grain**) at which they were identified (species listed in Table 3). As in Table 3, the two *Acer* species in **bold type** satisfied all aspects of both the outlier and the diversity criteria for candidate foundation species at the given spatial grain. The starred (*) species satisfied the outlier criterion (Fig. 1) and partially satisfied the diversity criterion at the given spatial grain: a positive spatial relationship between candidate foundation species size and beta diversity, and a negative spatial relationship between candidate foundation species size and beta diversity. The remaining species did not satisfy the outlier criterion but did meet some aspects of the diversity criterion. No species me either foundation species are grouped alphatically within increasing grain (subplot) sizes. Values are the minimum (**Min**), median (**Median**), mean (**Mean**), one standard deviation of the mean (**SD**), and maximum (**Max**), computed over all spatial lags, of the codispersion between the basal area of the candidate foundation species and all other woody species in square subplots with the length of a side = the spatial grain.

Plot	Grain	Species	Diversity metric	Min	Median	Mean (SD)	Max
LS	5 m	*Acer ukurunduense	Bray-Curtis	0.03	0.12	0.12(0.02)	0.16
			Richness	-0.13	-0.09	-0.09(0.02)	-0.01
			Shannon	-0.12	-0.08	-0.08(0.02)	-0.01
			Simpson	-0.07	-0.04	-0.04(0.01)	0.01
			Abundance	-0.14	-0.1	-0.10(0.02)	-0.03
		$*Corylus mandshurica^+$	Bray-Curtis	0.01	0.06	$0.06 \ (0.02)$	0.11
			Richness	-0.15	-0.1	-0.10(0.02)	-0.06
			Shannon	-0.14	-0.09	-0.09(0.02)	-0.05
			Simpson	-0.09	-0.05	-0.05(0.02)	0
			Abundance	-0.16	-0.11	-0.11(0.02)	-0.07
		*Fraxinus mandshurica	Bray-Curtis	0	0.05	0.05(0.02)	0.09
			Richness	-0.09	-0.05	-0.05(0.01)	-0.01
			Shannon	-0.09	-0.05	-0.05(0.01)	-0.01
			Simpson	-0.09	-0.05	-0.05(0.02)	0
			Abundance	-0.09	-0.03	-0.03(0.01)	0.01
	10 m	*Acer ukurunduense	Bray-Curtis	0.15	0.21	0.21(0.03)	0.27
			Richness	-0.23	-0.17	-0.17(0.03)	-0.09
			Shannon	-0.19	-0.12	-0.12(0.03)	-0.04
			Simpson	-0.10	-0.05	-0.05(0.02)	0.01
			Abundance	-0.27	-0.21	-0.21 (0.03)	-0.14
MLG	$5 \mathrm{m}$	*Acer barbinerve+	Bray-Curtis	0.12	0.16	0.16(0.01)	0.20
			Richness	-0.22	-0.16	-0.16(0.02)	-0.10
			Shannon	-0.16	-0.13	-0.13(0.01)	-0.08
			Simpson	-0.10	-0.07	-0.07(0.01)	-0.04
			Abundance	-0.24	-0.16	-0.15(0.03)	-0.09
		*Acer tequentosum	Bray-Curtis	0.02	0.05	0.05(0.01)	0.09
		0	Richness	-0.11	-0.07	-0.07(0.01)	-0.04
			Shannon	-0.08	-0.04	-0.04(0.01)	-0.01
			Simpson	-0.06	-0.02	-0.02(0.01)	0.01
			Abundance	-0.15	-0.11	-0.11(0.01)	-0.07
		*Acer ukurunduense	Bray-Curtis	0.03	0.06	0.06(0.01)	0.09
			Richness	-0.11	-0.07	-0.07(0.01)	-0.04
			Shannon	-0.09	-0.05	-0.06(0.01)	-0.01
			Simpson	-0.08	-0.05	-0.05(0.01)	-0.01
			Abundance	-0.11	-0.08	-0.08(0.02)	-0.03
		$Corylus mandshurica^+$	Bray-Curtis	0.12	0.16	0.16(0.01)	0.20
		U U	Richness	-0.14	-0.10	-0.10(0.01)	-0.07
			Shannon	-0.11	-0.08	-0.08(0.01)	-0.04
			Simpson	-0.04	-0.01	-0.01(0.01)	0.02
			Abundance	-0.20	-0.10	-0.11(0.02)	-0.06
		*Taxus cuspidata	Bray-Curtis	0.05	0.09	0.09(0.01)	0.12
		·····	Richness	-0.12	-0.08	-0.08(0.02)	-0.03
			Shannon	-0.09	-0.05	-0.05(0.01)	-0.02
			Simpson	-0.06	-0.03	-0.03(0.01)	0.02
			Abundance	-0.13	-0.07	-0.07(0.02)	-0.04
		* Tilia amurensis	Bray-Curtis	0.02	0.05	0.05 (0.01)	0.01
			Richness	-0.11	-0.06	-0.06(0.01)	-0.00
			Shannon	-0.07	-0.00	-0.04(0.01)	-0.01
			Simpson	-0.07	-0.04	-0.02(0.01)	0.01
			Abundance	-0.05 -0.15	-0.02 -0.09	-0.02(0.01) -0.09(0.02)	-0.01
	10 m	*Acer barbinerve+	Bray-Curtis	-0.13 0.07	-0.03 0.12	-0.03(0.02) 0.12(0.02)	-0.03 0.17
	10 111	ALET OUTOTHETVE	Richness	-0.22	-0.12	-0.12(0.02) -0.11(0.04)	$0.17 \\ 0.03$
			niciniess	-0.22	-0.11	-0.11(0.04)	0.05
			Shannon	-0.14	-0.09	-0.09(0.02)	-0.02

Simpson -0.14 -0.07 -0.07 * Pinus koraiensis Bray-Curtis -0.02 0.13 0.13 * Tilia amurensis Bray-Curtis -0.02 0.13 0.10 * Tilia amurensis Bray-Curtis 0.02 0.05 0.05 0.05 * Tilia amurensis Bray-Curtis 0.02 0.05 -0.06 0.01 Simpson -0.02 0.15 -0.16 0.01 Binamon -0.08 0.02 0.03 -0.08 0.02 0.03 -0.03 -0.08 0.01 Simpson -0.02 -0.18 -0.08 0.01 Simpson -0.02 -0.18 0.01 0.05 0.01 0.05 -0.018 0.01 0.05 -0.018 0.01 0.05 0.06 0.05 0.01 <th>ot</th>	ot
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$\begin{array}{c ccccccccccccccccccccccccccccccccccc$	
$ \begin{array}{c ccccc} {\rm Richness} & -0.09 & -0.03 & -0.03 & (0.0) \\ {\rm Shannon} & -0.06 & -0.01 & -0.01 & (0.0) \\ {\rm Simpson} & -0.04 & 0 & 0 & (0.0) \\ {\rm Abundance} & -0.14 & -0.08 & -0.08 & (0.0) \\ {\rm OL} & 5 \ {\rm m} \ Juglans \ mandshurica & {\rm Bray-Curtis} & 0.06 & 0.24 & 0.24 & (0.0) \\ \end{array} $	
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$ \begin{array}{c ccccc} & Shannon & -0.06 & -0.01 & -0.01 & 0.0.\\ Simpson & -0.04 & 0 & 0 & 0.0.\\ Abundance & -0.14 & -0.08 & -0.08 & 0.0.\\ \hline \\ DL & 5 m Juglans mandshurica & Bray-Curtis & 0.06 & 0.24 & 0.24 & 0.02 \\ \end{array} $	
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Plot	Grain	Species	Diversity metric	Min	Median	Mean (SD)	Max
			Shannon	-0.07	-0.04	-0.04 (0.01)	0.01
			Simpson	-0.06	-0.03	-0.03(0.01)	0.01
			Abundance	-0.08	-0.06	-0.06(0.01)	0.01
	$20 \mathrm{m}$	*Pinus massoniana	Bray-Curtis	0.14	0.20	0.2 (0.04)	0.27
			Richness	-0.28	-0.16	-0.16(0.06)	-0.05
			Shannon	-0.40	-0.27	-0.28(0.06)	-0.12
			Simpson	-0.33	-0.21	-0.21(0.06)	-0.06
			Abundance	0.16	0.26	0.26(0.04)	0.35
		*Quercus serrata var. brevipetiolata	Bray-Curtis	0.17	0.37	0.36(0.06)	0.44
		,	Richness	-0.18	-0.11	-0.11(0.05)	0.04
			Shannon	-0.34	-0.29	-0.28(0.04)	-0.13
			Simpson	-0.31	-0.25	-0.24(0.04)	-0.09
			Abundance	0.39	0.52	0.51 (0.05)	0.58
ML	5 m	Brassaiopsis glomerulata	Bray-Curtis	0.03	0.11	0.10(0.03)	0.17
		1 9	Řichness	-0.09	-0.03	-0.03(0.02)	0.03
			Shannon	-0.07	-0.02	-0.02(0.02)	0.03
			Simpson	-0.05	0	0 (0.02)	0.04
			Abundance	-0.15	-0.08	-0.08(0.04)	0.02
	10 m	Brassaiopsis glomerulata	Bray-Curtis	0.03	0.16	0.15(0.04)	0.02
	10 111	Drassatopsis giomeratata	Richness	-0.11	-0.04	-0.04(0.03)	0.20
			Shannon	-0.11 -0.05	-0.04	1 1	$0.04 \\ 0.05$
						0 (0.02)	
			Simpson	0.03	0.12	0.12(0.04)	0.2
			Abundance	-0.31	-0.19	-0.19(0.07)	0
DH	$5 \mathrm{m}$	$*A porusa yunnanensis^+$	Bray-Curtis	-0.03	0.12	0.11 (0.06)	0.21
			Richness	-0.20	-0.13	-0.12(0.04)	0.01
			Shannon	-0.19	-0.11	-0.11(0.04)	0.02
			Simpson	-0.15	-0.10	-0.10(0.03)	0
			Abundance	-0.22	-0.13	-0.13(0.04)	0
	10 m	$*A porusa yunnanensis^+$	Bray-Curtis	-0.03	0.18	0.17 (0.08)	0.29
			Richness	-0.29	-0.20	-0.19(0.06)	-0.03
			Shannon	-0.21	-0.12	-0.12(0.04)	0
			Simpson	-0.16	-0.09	-0.09(0.03)	-0.02
			Abundance	-0.37	-0.26	-0.24(0.07)	-0.03
	20 m	*Aporusa yunnanensis+	Bray-Curtis	0	0.23	0.21(0.09)	0.35
			Richness	-0.31	-0.24	-0.23(0.05)	-0.11
			Shannon	-0.14	-0.06	-0.06(0.05)	0.06
			Simpson	-0.12	-0.05	-0.04(0.04)	0.05
			Abundance	-0.47	-0.37	-0.34(0.09)	-0.14
NG	5 m	Ficus hispida ⁺	Bray-Curtis	0.03	0.11	0.12(0.04)	0.21
1.0	0 111	1 1000 1110 1000	Richness	-0.18	-0.10	-0.11(0.03)	-0.02
			Shannon	-0.16	-0.10	-0.10(0.03)	-0.02
			Simpson	-0.10 -0.16	-0.10 -0.11	-0.10(0.03) -0.11(0.03)	-0.02 -0.02
			Abundance			· · ·	
	10	Figure biomide+		-0.18	-0.09	-0.09(0.03)	-0.02
	$10 \mathrm{m}$	$Ficus\ hispida^+$	Bray-Curtis	0.08	0.16	0.17 (0.06)	0.3
			Richness	-0.23	-0.15	-0.15(0.04)	-0.06
			Shannon	-0.15	-0.10	-0.10(0.02)	-0.04
			Simpson	-0.1	-0.07	-0.07(0.01)	-0.04
			Abundance	-0.28	-0.15	-0.16(0.05)	-0.06
XSBN	$5 \mathrm{m}$	$Mezzettiopsis\ creaghii$	Bray-Curtis	-0.02	0.09	0.08(0.03)	0.12
			Richness	-0.06	-0.03	-0.03 (0.01)	0.02
			Shannon	-0.05	-0.01	-0.01 (0.01)	0.02
			Simpson	-0.03	-0.01	0(0.01)	0.04
			Abundance	-0.09	-0.06	-0.06(0.02)	0.01
	10 m	Mezzettiopsis creaghii	Bray-Curtis	0	0.20	0.19(0.05)	0.26
			Richness	-0.11	-0.05	-0.05(0.03)	0.03
	10 111						0.14
	10 11		Shannon	-0.06	0.05	0.04 (0.04)	
	10 111		Shannon	-0.06 -0.01	0.05	0.04 (0.04) 0.09 (0.04)	
	10		Simpson	-0.01	0.09	0.09(0.04)	0.21
		Magnettionais areachii	Simpson Abundance	$-0.01 \\ -0.21$	$0.09 \\ -0.15$	$\begin{array}{c} 0.09 \; (0.04) \\ -0.14 \; (0.04) \end{array}$	$\begin{array}{c} 0.21 \\ 0.02 \end{array}$
	20 m	Mezzettiopsis creaghii	Simpson Abundance Bray-Curtis	$-0.01 \\ -0.21 \\ 0.16$	$0.09 \\ -0.15 \\ 0.33$	$\begin{array}{c} 0.09 \ (0.04) \\ -0.14 \ (0.04) \\ 0.31 \ (0.06) \end{array}$	$\begin{array}{c} 0.21 \\ 0.02 \\ 0.38 \end{array}$
		Mezzettiopsis creaghii	Simpson Abundance Bray-Curtis Richness	-0.01 -0.21 0.16 -0.19	$0.09 \\ -0.15 \\ 0.33 \\ -0.09$	$\begin{array}{c} 0.09 \ (0.04) \\ -0.14 \ (0.04) \\ 0.31 \ (0.06) \\ -0.09 \ (0.05) \end{array}$	$0.21 \\ 0.02 \\ 0.38 \\ -0.01$
		Mezzettiopsis creaghii	Simpson Abundance Bray-Curtis Richness Shannon	-0.01 -0.21 0.16 -0.19 -0.06	$\begin{array}{c} 0.09 \\ -0.15 \\ 0.33 \\ -0.09 \\ 0.22 \end{array}$	$\begin{array}{c} 0.09\ (0.04)\\ -0.14\ (0.04)\\ 0.31\ (0.06)\\ -0.09\ (0.05)\\ 0.19\ (0.09) \end{array}$	$\begin{array}{c} 0.21 \\ 0.02 \\ 0.38 \\ -0.01 \\ 0.32 \end{array}$
		Mezzettiopsis creaghii	Simpson Abundance Bray-Curtis Richness	-0.01 -0.21 0.16 -0.19	$0.09 \\ -0.15 \\ 0.33 \\ -0.09$	$\begin{array}{c} 0.09 \ (0.04) \\ -0.14 \ (0.04) \\ 0.31 \ (0.06) \\ -0.09 \ (0.05) \end{array}$	$0.21 \\ 0.02 \\ 0.38 \\ -0.01$

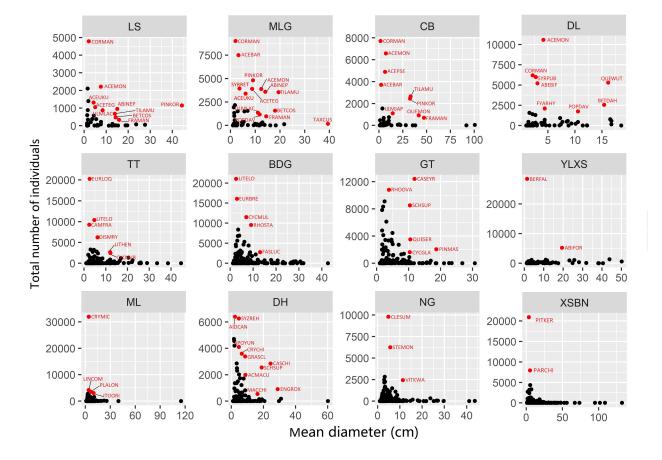


Figure 1: Size (DBH)-frequency distributions of the species in each plot. Species falling outside of the "reverse-J" line (in red) were treated in the first set of candidate foundation species. Plots are ordered left-to-right and top-to-bottom by latitude; species abbreviations are given in Table 2.

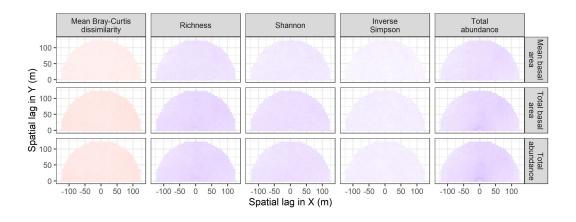


Figure 2: Codispersion between mean basal area, total basal area, or total abundance of *Acer barbinerve* and five different measures of diversity of associated woody species in 5-m subplots in the 25-ha Muling (MLG) plot. Codispersion coefficients were calculated for spatial lags ranging from 0-125 m at 5-m intervals. The values of the codispersion can range from -1 (dark blue) through 0 (white) to 1 (dark red). Statistical significance for codispersion coefficients computed at each spatial lag is shown in Fig. 3.

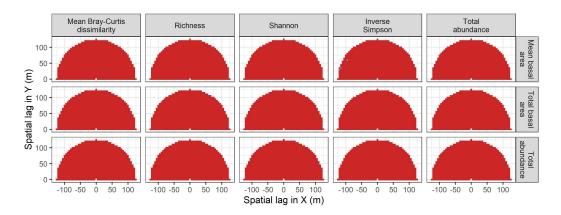


Figure 3: Statistical significance of the codispersion coefficients calculated between basal area or abundance of *Acer barbinerve* and five different measures of diversity of associated woody species in 5-m subplots in the 25-ha Muling (MLG) plot. Statistical significance was determined by comparing observed codispersion at each spatial lag with the distribution of 199 spatial randomizations of a toroidal-shift null model. Red: $P \leq 0.05$; Blue: P > 0.05.

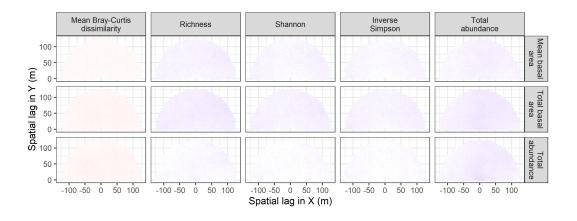


Figure 4: Codispersion between mean basal area, total basal area, or total abundance of *Acer ukurunduense* and five different measures of diversity of associated woody species in 5-m subplots in the 25-ha Muling (MLG) plot. Codispersion coefficients were calculated for spatial lags ranging from 0-125 m at 5-m intervals. The values of the codispersion can range from -1 (dark blue) through 0 (white) to 1 (dark red). Statistical significance for codispersion coefficients computed at each spatial lag is shown in Fig. 5.

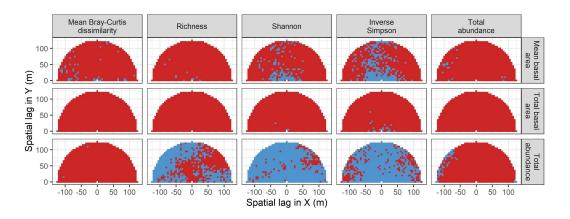


Figure 5: Statistical significance of the codispersion coefficients calculated between basal area or abundance of *Acer ukurunduense* and five different measures of diversity of associated woody species in 5-m subplots in the 25-ha Muling (MLG) plot. Statistical significance was determined by comparing observed codispersion at each spatial lag with a distribution of 199 spatial randomizations of a toroidal-shift null model. Red: $P \leq 0.05$; Blue: P > 0.05.

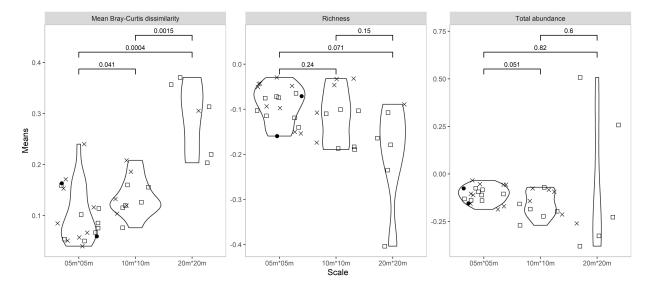


Figure 6: Distribution of average codispersion observed between total basal area of candidate foundation species and Bray-Curtis dissimilarity, species richness, and total abundance of associated woody plant species in continguous 5×5 -, 10×10 -, and 20×20 -m subplots in the twelve CForBio plots. Points indicate mean codispersion values for each candidate foundation species listed in Table 2; solid points indicate the two candidate foundation species in the genus *Acer* that met both the outlier *and* diversity criterion for all indices; hollow squares indicate candidate species that met the outlier criterion and the relaxed diversity criterion; and crosses indicate the remaining candidate foundation species that met only the relaxed diversity criterion. P values for comparisons between groups are shown at the top of each panel.

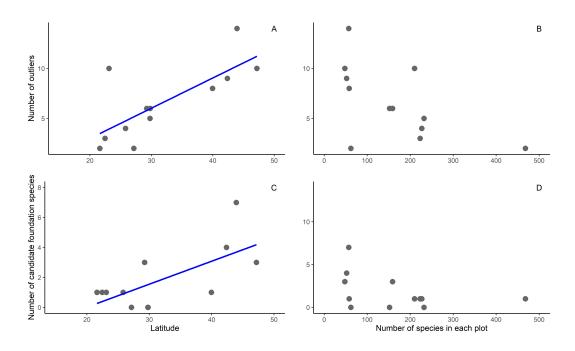


Figure 7: Number of outliers from the expected size-frequency distribution (Fig. 1) and number of candidate foundation species (Table 3) as a function of latitude (\mathbf{A}, \mathbf{C}) or plot-level species richness (\mathbf{B}, \mathbf{D}) . See main text for regression statistics.

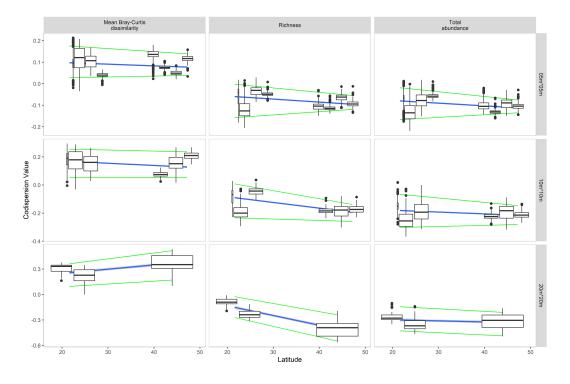


Figure 8: Relationship between latitude and codispersion between candidate foundation species (canopy trees and understory trees and shrubs) and three measures of associated woody-plant diversity at different spatial grains. Box plots illustrate median, upper and lower quartiles, and individual points outside of the upper and lower deciles of average codispersion at each latitude where candidate foundation species occurred (Table 3). Lines are regressions on all the data (blue lines), or on the 5% or 95% quantiles of the data (green lines).

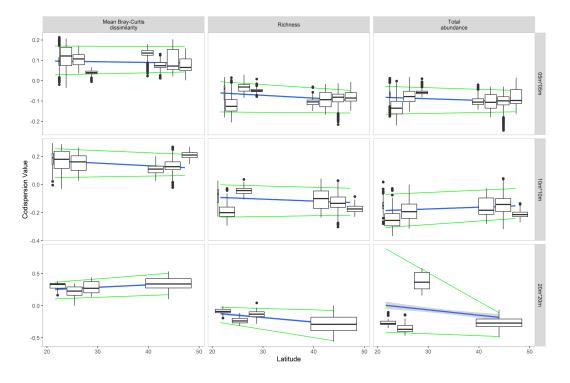


Figure 9: Relationship between latitude and codispersion between candidate foundation canopy tree species and three measures of associated woody-plant diversity at different spatial grains. Box plots illustrate median, upper and lower quartiles, and individual points outside of the upper and lower deciles of average codispersion at each latitude where candidate foundation species occurred (Table 3). Lines are regressions on all the data (blue lines), or on the 5% or 95% quantiles of the data (green lines).