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Structure metrics to generalize biomass estimation from lidar across forest types from different continents

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19 Abstract

20 Forest aboveground biomass is a key variable in remote sensing based forest monitoring. Active sensor 21 systems, such as lidar, can generate detailed canopy height products. Relationships between canopy 22 height and biomass are commonly established via regression analysis using information from ground-truth 23 plots. In this way, many site-specific height-biomass relationships have been proposed in the literature 24 and applied for mapping in regional contexts. However, such relationships are only valid within the specific 25 forest type for which they were calibrated. A generalized relationship would facilitate biomass estimation 26 across forest types and regions. In this study, a combination of lidar-derived and ancillary structural 27 descriptors is proposed as an approach for generalization between forest types. Each descriptor is 28 supposed to quantify a different aspect of forest structure, i.e., mean canopy height, maximum canopy 29 height, maximum stand density, vertical heterogeneity and wood density. Airborne discrete return lidar 30 data covering 194 ha of forest inventory plots from five different sites including temperate and tropical 31 forests from Africa, Europe, North, Central and South America was used. Biomass predictions using the 32 best general model (nRMSE = 12.4%, R² = 0.74) were found to be almost as accurate as predictions using five site-specific models (nRMSE = 11.6%, R² = 0.78). The results further allow interpretation about the 33 34 importance of the employed structure descriptors in the biomass estimation and the mechanisms behind 35 the relationships. Understanding the relationship between canopy structure and aboveground biomass 36 and being able to generalize it across forest types are important steps towards consistent large scale 37 biomass mapping and monitoring using airborne and potentially also spaceborne platforms.

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39 Keywords: forest structure; aboveground biomass; canopy height; lidar; generalization

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42 **1. Introduction**

43 Quantifying global carbon stocks of forests as well as their changes over time requires spatially explicit 44 measurements and monitoring (Harris et al., 2012). The primary variable of interest hereby is forest 45 aboveground biomass (AGB). Thus, there is a growing amount of literature about estimating forest 46 biomass from canopy height metrics derived from light detection and ranging (lidar), synthetic aperture 47 radar or photogrammetry (Asner and Mascaro, 2014; Goetz and Dubayah, 2011; Lu et al., 2014; Treuhaft 48 et al., 2015; Zolkos et al., 2013). The majority of these studies investigated data from specific forest sites 49 with the goal to find the best prediction model, i.e., maximizing explained variability (e.g., R²), minimizing 50 prediction error (e.g., RMSE) and minimizing systematic bias, while using the most parsimonious set of 51 predictor variables (Zolkos et al., 2013). Different statistical approaches have been used including multiple 52 linear regression models and machine learning methods (Fassnacht et al., 2014). As a result various site-53 specific and forest type-specific relations for biomass estimation have been proposed and applied 54 successfully for biomass mapping at regional scale.

55 Comparatively few studies have tried to seek for generalization in the estimation approaches. In those studies conducted in the temperate and boreal biomes, field plots ranging from 500 to 1000 m² in size 56 57 were used and reported estimation uncertainties ranged from 2.5 to 23% (Bouvier et al., 2015; Lefsky et 58 al., 2002; Magnussen et al., 2012). In similar studies conducted in the tropical biome, field plots were 59 commonly in the size range from 0.1 to 1 ha and reported estimation uncertainties were around 10% at 60 the 1-ha scale (Gregory P. Asner et al., 2012; Asner and Mascaro, 2014; Vincent et al., 2014, 2012) For 61 consistent global mapping of forest biomass, however, it would be desirable to have more generic 62 relationships which are applicable across different forest types and biogeographic regions. Such an 63 approach would contribute to a better understanding of how structural attributes differ between forest 64 types and how they are related to biomass. It would also facilitate biomass mapping across the globe. 65 Given the variety of metrics that can be derived from lidar data (Næsset, 2002), it would further be

66 desirable to have a minimum set of meaningful metrics, describing different aspects of forest structure, to 67 avoid problems with multicollinearity and extensive model selection procedures (Bouvier et al., 2015). A widely used general approach is the one proposed by Asner et al. (2012) and modified by Asner & 68 69 Mascaro (2014) for pan-tropical application. The function is inspired by individual tree allometry, where 70 tree AGB can be modeled as a multiplicative power law of tree diameter at breast height (DBH; or tree 71 basal area BA), tree height and species-specific wood density (Chave et al., 2014). Hence, as a stand level 72 equivalent for area-based AGB estimation they used a power law of stand BA sum, mean top-of-canopy 73 height (TCH) and average wood density. The approach further assumes a linear relationship between BA 74 and TCH, which may differ between regions, and regional differences in average wood density can be 75 considered. This approach has been established using data from different tropical regions (Hawaii, 76 Panama, Peru, Madagascar) and has been applied successfully in other tropical regions, e.g., Colombia (G. 77 P. Asner et al., 2012), Malaysia (Coomes et al., 2017), Tanzania (Getzin et al., 2017), with uncertainties 78 between 12 and 15% at 1-ha scale.

Bouvier et al. (2015) suggested a different model for generalized AGB estimation. They used several a priori defined lidar-based metrics that captured different aspects of forest structure. Their model was able to produce accurate AGB estimations for different forest types in France. However, site-specific coefficients led to higher prediction accuracies for each site, compared to using only one set of coefficients across all sites.

In this study, we attempted to find a model that is generally applicable throughout different forest types and even different biomes by including structural information on forest stands. Tropical forests in Panama, French Guiana and Gabon were analyzed along with temperate forests in the United States of America and Germany. A generalized approach should also perform well across the gradients from intact to disturbed and natural to managed forests without requiring any stratification. Therefore, our analysis incorporated plots of primary and managed forests. We tried to estimate BA and AGB at the 1-ha scale. Usually, the two are closely correlated. However, BA is a simple inventory-derived metric, which is easily comparable

91 among sites and studies (Vincent et al., 2012). Inventory-based AGB, on the other hand, is more complex 92 to compute. Assumptions about allometric relationships and wood density values are required to derive 93 single tree AGB. It can lead to considerable differences in stand AGB if different assumptions are chosen 94 for the same stand (Duncanson et al., 2017). It has been argued that BA (or its two components stem 95 number and quadratic mean stem diameter) is the essential variable that has to be derived from remote 96 sensing and AGB could then be estimated based on predicted BA and the regional tree diameter-height 97 allometries and wood densities (Vincent et al., 2014, 2012). Here, we chose to analyze both variables - BA 98 for its robustness and comparability and AGB as the major variable of interest in forest carbon mapping 99 efforts.

100 We hypothesized that the following structural forest attributes may contribute to explaining stand BA and 101 AGB: a) mean canopy height, b) maximum possible stand density c) maximum possible tree height, d) 102 vertical canopy heterogeneity, and for AGB, additionally, e) average wood density. Most of these structural 103 attributes can be quantified in several alternative ways. Thus, for one attribute there may be a set of 104 several candidate metrics. Some of the metrics can be derived from lidar, while others require ancillary 105 information. In this analysis, data from 194 ha of temperate and tropical forest from five megaplot sites 106 were combined with the following goals: 1) to find a generic approach for BA and AGB estimation that can 107 be applied across all sites without causing prediction bias at any individual site and 2) to investigate the 108 contributions of the different structural attributes.

109

110 2. Material & Methods

111 *2.1. Study Sites*

112 Data from five forest sites covering different forest types and biogeographical zones were used (Tab. 1). 113 Four study sites are part of the ForestGEO megaplot network (Anderson-Teixeira et al., 2015) and thus 114 they have been inventoried according to a standard protocol. The data structure for the fifth site, Paracou, 115 is similar to the data structure of the ForestGEO sites. For each tree, the diameter at breast height (DBH), 116 spatial position and species identity were recorded. In this study, only trees with a DBH \geq 10 cm were 117 considered, because 10 cm was the minimum DBH recorded at Paracou (Blanc et al., 2009), and all given 118 numbers refer to trees above this size threshold. In the following, each of the five sites is briefly described. 119 1) Barro Colorado Island (BCI), Panama, is a Central American lowland tropical moist forest site with an 120 annual precipitation of 2580 mm and an average temperature of 27.1 °C. The census on the 50-ha plot was 121 conducted in 2010 (Condit, 1998; Condit et al., 2012; Hubbell et al., 1999) and comprised 22,084 trees, 122 which belonged to 223 species. Trees of the six families Malvaceae (17%), Fabaceae (13%), Moraceae (7%), 123 Euphorbiaceae (6%), Rubiaceae (6%) and Meliaceae (6%) account for 56% of AGB with Quararibea 124 asterolepis being the single species with the highest contribution (6%).

125 2) Paracou, French Guiana, is a South American lowland tropical rainforest with an annual precipitation of 126 3040 mm and an average temperature of 26 °C. There are 16 large plots with plots 1 to 15 having an extent 127 of 250 m × 250 m each and plot 16 having an extent of 500 m × 500 m. In 1986 and 1987, selective logging 128 with different treatment intensities (timber logging, fuelwood logging, thinning) was conducted on some 129 of the plots, while others have served as control plots (Hérault and Piponiot, 2018). Since our analysis was 130 conducted for 100 m × 100 m units, only subareas of 200 × 200 m measured from the south-western 131 corners of plots 1 to 15 were used. In total, 85 ha from Paracou were analyzed. Censuses were conducted 132 in 2015 and comprised 53,501 trees of 713 species. Trees of the three families Fabaceae (25%), Lecythidaceae (15%) and Chrysobalanaceae (11%) account for 51% of AGB with *Eperua falcata* being the
 single species with the highest contribution (7%).

3) Rabi, Gabon, is a Central African lowland tropical rainforest site with an annual precipitation of 2300 mm
and an average temperature of 26 °C. The census on the 25-ha plot was conducted from 2010 to 2012
(Labrière et al., 2018) and comprised 12,019 trees, which belonged to 235 species. Trees of the three
families Fabaceae (38%), Ochnaceae (9%) and Simaroubaceae (7%) account for 54% of AGB with *Lophira alata* being the single species with the highest contribution (9%).

4) The Smithsonian Environmental Research Center (SERC) plot, United States of America, is a North
American deciduous broadleaved temperate forest site with an annual precipitation of 1070 mm and an
average temperature of 13.2 °C. The census on the 16-ha plot was conducted in 2014 (Král et al., 2016;
McMahon and Parker, 2015) and comprised 4,719 trees, which belonged to 39 species. Trees of the six
species *Liriodendron tulipifera* (27%), *Liquidamber styraciflua* (16%), *Fagus grandifolia* (11%), *Quercus alba*(8%), *Carya alba* (7%) and *Fraxinus pennsylvanica* (6%) together account for 76% of the AGB.

5) Traunstein, Germany, is a Central European managed mixed temperate forest site, which includes conifer and broadleaf plantations. It has an annual precipitation of 1240 mm and an average temperature of 7.6 °C. The census on the 25-ha plot was conducted from 2015 to 2016. Due to the shape of the plot a rectangular 18-ha subarea was selected for the analysis. It comprised 7,182 trees, which belonged to 25 species. Trees of the four species *Picea abies* (39%), *Fagus sylvatica* (22%), *Acer pseudoplatanus* (19%) and

151 *Abies alba* (15%) together account for 94% of the AGB.

Site	Region	Forest type	Size [ha]	Location	Year of inventory	Year of lidar scan	Basal area range [m² ha ⁻¹]
BCI	Central Panama	Neotropical moist	50	9.15° N, 79.85° W	2010	2009	17.3 – 38.5
Paracou	Northern French Guiana	Neotropical wet	85	5.27° N, 52.92° W	2015	2015	24.8 - 38.7
Rabi	Western Gabon	Afrotropical wet	25	1.92° S, 9.88° E	2010-2012	2015	20.8 - 36.7
SERC	Eastern USA	Nearctic temperate broadleaf	16	38.89° N, 76.56° W	2014	2017	26.2 - 43.5
Traunstein	Southern Germany	Palearctic temperate mixed	18	47.94° N, 12.67° E	2015-2016	2016	7 – 44.6

152 Tab. 1: Information about the study sites

153 2.2. Inventory Data

154 The inventory data was processed to calculate AGB of each tree. Based on species, wood density values 155 were assigned to each tree using the ForestGEO wood density database 156 (http://ctfs.si.edu/Public/Datasets/CTFSWoodDensity/) and, in the case of Paracou, the Global Wood 157 Density Database (Chave et al., 2009; Zanne et al., 2009). Wood density values for 82% of all trees were 158 available at species level. For the remaining trees, median wood densities at genus (14.1%) or family level 159 (3.7%) or the overall median at the site (0.2%) were assigned, respectively. From DBH, the height of each 160 tree was calculated using site specific asymptotic allometric relationships. These relationships were 161 derived by fitting regression models of the Michaelis-Menten type (Equation 1) to the diameter height 162 dataset from Jucker et al. (2017), grouped by biogeographical region and forest type.

163 Equation 1:
$$H = \frac{h_{max} \cdot D}{d_{1/2} + D}$$

164 This equation describes tree height H [m] as a function of DBH D [m] with two parameters: 1) h_{max} [m], 165 which is the asymptotic maximal possible tree height, and 2) $d_{1/2}$ [m], which is the DBH of a tree that has 166 reached a height of half of h_{max}. A verification of whether the derived models describe the DBH-height 167 relations at each site reasonably well, was done by plotting the curves together with the maximal observed 168 DBH and maximal (lidar-derived) height of each hectare (Fig. S1). This showed that the relationships match 169 the observed values for the BCI, Paracou, Rabi and Traunstein plot, but strongly underestimate the tree 170 heights at SERC. Thus, for SERC we discarded the diameter-height relationship obtained from the dataset 171 and instead obtained parameters by directly fitting a regression model to the data points in Fig. S1, 172 representing the lidar-derived maximal heights on each hectare. All height allometry parameters used are 173 listed in Tab. 2.

Site	Species group	h _{max} [m]	d _{1/2} [m]	
BCI	All	57.4	0.43	
Paracou	All	57.4	0.43	
Rabi	All	59.9	0.48	
SERC (discarded)	All	37	0.22	
SERC (used instead)	All	54.7	0.27	
Traunstein	Broadleafs	48.8	0.25	
Traunstein	Conifers	68.9	0.5	

175 Tab. 2: Parameters for the different diameter-height relationships modeled with a Michaelis-Menten equation.

176

Aboveground biomass (AGB [t]) of each tree was calculated according to the general allometric equation
suggested by (Chave et al., 2014) ("Model 5") (Equation 2) with DBH D [m], height H [m] and wood density
WD [t m⁻³].

180 Equation 2: $AGB = 0.559 \cdot D^2 \cdot H \cdot WD$

181

182 2.3. Lidar Data

183 Small footprint discrete return lidar data was collected in BCI in August 2009 using an Optech ALTM Gemini sensor with a mean point density of 21 m⁻² (Lobo and Dalling, 2014), in Paracou in October 2015 using an 184 Riegl LMS Q 780 with a mean point density of 54.6 m⁻², in Rabi in 2015 using a Riegl VQ-480i sensor with a 185 186 mean point density of 2.5 m⁻² (Labrière et al., 2018), at SERC in July 2017 using a Riegl VQ-480i sensor with a mean point density of 54.1 m⁻² (Cook et al., 2013) and in Traunstein in August 2016 using a Riegl LMS Q. 187 680i sensor with a mean point density of 20.8 m⁻². The lidar point clouds were terrain-normalized using 188 189 LAStools (Isenburg, 2011) and rasterized to canopy height models (CHM) with 1-m resolution, by taking 190 the height of the highest return in each 1-m²-cell (Fig. 1). No interpolation was used and cells with no 191 return were filled with value zero (ground height).



192 193 Fig. 1: Canopy height models of the five study sites: a) Barro Colorado Island, b) Rabi, c) Paracou Plot 16 (biodiversity plot), d)

194 Paracou Plot 1 (control plot), e) Paracou Plot 2 (selective logging), f) Paracou Plot 3 (selective logging and timber stand

195 improvement), g) Paracou Plot 4 (selective logging, timber stand improvement and fuelwood collection), h) Smithsonian 196

Environmental Research Center and i) Traunstein. The black grids and numbers represent the 1-ha subplots with each ha 197

representing one record in the analysis.

199 2.4. Forest Structure Metrics

All inventory plots were divided into square-shaped subplots of 1-ha size each. At 1-ha scale a variety of structural metrics was calculated from 1) the inventory data and 2) the lidar data. Inventory-based metrics included basal area sum (BA), number of stems per ha (N), quadratic mean tree diameter (at breast height, Dg), maximum DBH per ha (D_{max}), mean wood density weighted for tree basal area (WD_{BA}) or weighted for tree aboveground volume (WD_{AGV}) and stand density index (SDI, Equation 3), which is a standardized metric for stocking (Reineke, 1933).

206 Equation 3:
$$SDI = N \cdot \left(\frac{25}{D_g}\right)^{-1.605}$$

Additionally to those metrics derived at 1-ha-level, we derived a set of metrics at site-level. Those sitelevel metrics included maximum basal area sum (BA_{smax}) and maximum SDI (SDI_{smax}) of all the 1-ha plots at each site s and basal area-weighted mean wood density (WD_{sBA}) and aboveground volume-weighted mean wood density (WD_{sAGV}) across all trees at each site s.

Lidar-based metrics were maximum canopy height per 1-ha plot H_{max} and per site H_{smax}, mean top-ofcanopy height from CHMs of two different resolutions (1-m and 10-m pixels called TCH₁ and TCH₁₀), standard deviation of the 1-m CHM (SD_{CHM}), coefficient of variation of the 1-m CHM (CV_{CHM}) and Gini index of the 1-m CHM (Gini_{CHM}). The vertical foliage profile (VFP) was derived from the vertical profile of the 1m CHM following the approach described by Harding et al (2001) (Equation 4). Despite being originally developed for large footprint waveforms, the method can also be applied to vertical profiles of CHMs, since the latter have been shown to closely match coinciding waveforms (Blair and Hofton, 1999).

218 Equation 4:
$$VFP(h_i) = \frac{1}{k \cdot \Delta h} \cdot \ln(\frac{GP(h_i)}{GP(h_{i+1})})$$

with k being the light extinction coefficient, Δh the width of one height bin (here 1 m) and GP(h_i) the gap probability (value of the cumulative CHM profile) in height bin h_i (Ni-Meister et al., 2001). All pixels below 5 m height were regarded as ground and k was set to 0.3 for all sites. The parameter k can be described as

- the quotient of a projection coefficient G, which is 0.5 for a random leaf angle distribution, and a clumping index C, which on average is 1.58 for different forest types (Tang et al., 2012). A value of k = 0.3 has been shown to result in good LAI estimations (Getzin et al., 2017). The same vertical distribution metrics as for the CHMs were derived from the VFP, namely SD_{VFP}, CV_{VFP} and Gini_{VFP}. All metrics used for statistical modeling are listed in Tab. 3 and equations to calculate them are detailed in Fischer et al. (2019).
- 227 Tab. 3: List of metrics used in the statistical analysis (CHM = canopy height model).

Acronym	Explanation of metric	Structural aspect	Predictor (P _x)
AGB	Aboveground biomass [t ha ⁻¹]	Target variable	-
BA	Basal area [m ² ha ⁻¹]	Target variable	-
TCH ₁	Mean top-of-canopy height from 1-m pixel CHM [m]	Mean canopy height	P _h
TCH ₁₀	Mean top-of-canopy height from 10-m pixel CHM [m]	Mean canopy height	P _h
BA _{smax}	Maximal basal area at site s [m ² ha ⁻¹]	Maximum stand density	P _d
SDI _{smax}	Maximal stand density index at site s [ha ⁻¹]	Maximum stand density	P _d
H _{smax}	Maximal canopy height at site s [m]	Maximum canopy height	Pm
SD _{CHM}	Standard deviation of the 1-m pixel CHM [m]	Vertical heterogeneity	Pv
СИСНМ	Coefficient of variation of the 1-m pixel CHM [m]	Vertical heterogeneity	Pv
Gini _{снм}	Gini index of the 1-m pixel CHM [m]	Vertical heterogeneity	P _x
SD _{VFP}	Standard deviation of the vertical foliage profile [m]	Vertical heterogeneity	Pv
CV _{VFP}	Coefficient of variation of the vertical foliage profile [m]	Vertical heterogeneity	Pv
Gini _{VFP}	Gini index of the vertical foliage profile [m]	Vertical heterogeneity	Pv
WD _{sAGV}	Mean aboveground volume-weighted wood density at site s [t m ⁻³]	Wood density	Pw
WD _{sBA}	Mean basal area-weighted wood density at site s [t m ⁻³]	Wood density	Pw

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229 2.5. Multivariate Regression Analysis

Regression analysis was conducted to find the best relationship and set of predictor variables for BA and AGB estimation with the main objective to minimize overall root mean squared error (RMSE) across sites. The regression models had the functional form of multivariate power laws. Each predictor was supposed to capture a different structural aspect of the forest. Several candidate metrics were grouped into sets of potential predictors and tested in different combinations. By categorizing metrics into different structural aspects the number of possible metric combinations for testing was reduced. Combinations of redundant metrics, i.e., metrics that capture the same structural aspect, were not tested. For BA, a four predictor
equation was used (Equation 5). For AGB, a five predictor equation was used (Equation 6).

238 Equation 5: 239 Equation 6: $BA = a_0 \cdot P_h{}^{a_h} \cdot P_d{}^{a_d} \cdot P_m{}^{a_m} \cdot P_v{}^{a_v}$ $AGB = b_0 \cdot P_h{}^{b_h} \cdot P_d{}^{b_d} \cdot P_m{}^{b_m} \cdot P_v{}^{b_v} \cdot P_w{}^{b_w}$

240 Each P_x represents a predictor for a certain structural aspect x with a_x being the coefficients for BA 241 estimation and b_x being the coefficients for AGB estimation. For each P_x one single metric from a set of 242 possible predictors was used in each regression model. The predictor sets were defined as follows: The 243 predictor for mean canopy height P_h was either TCH₁ or TCH₁₀. The maximum possible canopy height P_m 244 was exclusively represented by H_{smax}. The predictor for maximum density (stocking) P_d was either BA_{smax} 245 or SDI_{smax}. For vertical heterogeneity of the canopy Pv, SD_{CHM}, CV_{CHM}, Gini_{CHM}, SD_{VFP}, CV_{VFP} and Gini_{VFP} have 246 been explored. Average wood density Pw was only included in the AGB estimation and was either WD_{SBA} 247 or WD_{sAGV}.

248 Correlations among predictors were highest within predictor groups P_x (Fig. S2, e.g., within P_h, P_v and P_w). 249 Thus, by only combining predictors from different groups multicollinearity was reduced. Between groups, 250 the highest correlations were observed between H_{smax} and either of the two density predictors BA_{smax} and 251 SDI_{smax}, while the two among each other were less correlated. We do not expect that maximum possible 252 canopy height and maximum stocking density of forests are generally related. We rather suspect that 253 correlations between these site-level metrics occurred by chance due to the small sample size of five sites. 254 Maximum likelihood parameter estimation in R (R Development Core Team, 2014) was used to derive the 255 coefficients for Equations 5 and 6. All possible metric combinations were tested including all possible 256 subsets discarding one or several predictors Px. The goodness-of-fit was evaluated based on linear 257 regression of predictions against ground-based observations of the dependent variable to quantify R², 258 RMSE and nRMSE (normalized RMSE by dividing it by the mean observed value). Wilcoxon tests were 259 performed to check whether the mean prediction residual at each site deviated significantly from zero, 260 with the goal of identifying unbiased prediction models. For each predictor combination, 1000 261 bootstrapping replicates were performed by resampling the dataset randomly with replacement. Site-level 262 metrics BAsmax, SDIsmax, Hsmax and WDsAGV were recalculated based on the resampled dataset, i.e., if, e.g., 263 the plot with the largest H_{max} of site s was not in the resampled dataset, H_{smax} was set to the largest H_{max} 264 of any plots from site s present in the resampled dataset. Mean bootstrapped statistics (RMSE_b, nRMSE_b 265 and R²_b) served to evaluate the different models. The best predictor combinations for BA and AGB were 266 finally tested in a leave-one-site-out cross validation, in which five different models were fit on data from 267 four sites, respectively, and the fifth site was set aside for testing.

268

269 2.6. Site-specific Reference Regression Models

To assess the performance of the derived general, site-independent, structure-based multi predictor regression models site-specific reference models were required. For this purpose, single predictor regression models were fit. As predictors for these reference models P_h, i.e., TCH₁ or TCH₁₀, were used. These metrics have been most widely used for this purpose (Asner and Mascaro, 2014; Knapp et al., 2018) and showed the highest individual correlations with the target variables BA and AGB (Fig. S2). The models given by Equations 7 and 8 were fit by splitting the dataset into five subsets with each subset containing only records from one site and using the same fitting procedure as described above.

277 Equation 7:
$$BA = a_{0,site} \cdot P_h^{a_{h,site}}$$

278 Equation 8:
$$AGB = b_{0,site} \cdot P_h^{b_{h,site}}$$

As an alternative, also site-specific multi predictor models are possible. However, such models make comparisons more complicated, because the set of best predictors may vary from site to site. In our analysis the predictors P_d , P_m and P_w were all at site level. Thus, the only relevant predictor for site-specific multi predictor models, apart from P_h , was P_v . The best model using a combination of P_h and P_v for each site are listed in the supplements (Tab. S2).

285 **3. Results**

286 3.1. Forest Structure at Different Sites

287 The different structure attributes at 1-ha scale varied within and among the five sites (Fig. 2 and Fig. S3). This section gives an overview on the distributions of the most important ground- and lidar-based 288 structure metrics. BA values ranged from 7 to 44.6 m² ha⁻¹ with a mean of 29.8 m² ha⁻¹. AGB values ranged 289 290 from 76 to 638 t ha⁻¹ with a mean of 354 t ha⁻¹. Mean top-of-canopy height ranged from 5.6 to 38 m when 291 calculated from $1-m \times 1-m$ pixels (TCH₁) and from 15.8 to 41.8 m when calculated from $10-m \times 10-m$ pixels 292 (TCH₁₀). In both cases, the distributions for the tropical sites were similar while TCH were on average higher 293 at SERC and lower at Traunstein. Mean wood densities per hectare were calculated on a BA-weighted and 294 on an AGV-weighted basis. Both were similar in their distributions and mean wood densities calculated for 295 each site across the entire megaplot weighted by either BA or AGV were almost identical with the largest 296 difference being 0.02 t m⁻³ in the case of BCI (WD_{sBA}: BCI: 0.51 t m⁻³, Paracou: 0.69 t m⁻³, Rabi: 0.66 t m⁻³, SERC: 0.48 t m⁻³, Traunstein: 0.5 t m⁻³; WD_{sAGV}: BCI: 0.49 t m⁻³, Paracou: 0.69 t m⁻³, Rabi: 0.66 t m⁻³, SERC: 297 298 0.47 t m⁻³, Traunstein: 0.5 t m⁻³). Due to this similarity, only AGV-weighted wood density at site level 299 WD_{sAGV} was considered in the further analysis. Mean wood densities at Paracou and Rabi exceeded the 300 values from all other sites strongly with WD_{AGV} at the 1-ha scale ranging from 0.6 to 0.74 t m⁻³ at Paracou 301 and Rabi and from 0.42 to 0.55 t m⁻³ at all other sites. Stand density index values ranged from 138 to 778 302 with the lowest values occurring in recently managed parts of Traunstein. The maximal SDIs are proxies 303 for the highest possible stocking density in the different forest types (BCI: 683, Paracou: 749, Rabi: 703, 304 SERC: 708, Traunstein: 778). The maximum canopy height covered a wide range from 27.1 m to 54.7 m. 305 H_{max} at SERC only covered a very narrow range falling inside the range of the tropical sites, while H_{max} at 306 Traunstein were much lower. The maximum canopy heights per site H_{smax} were 54.7 m at BCI, 50 m at 307 Paracou, 52.6 m at Rabi, 46.2 m at SERC and 40.3 m at Traunstein.





Fig. 2: Boxplots of the distributions of a selection of forest structure metrics across the five study sites Barro Colorado Island (B), Paracou (P), Rabi (R), Smithsonian Environmental Research Center (S) and Traunstein (T). Graphics (a) and (b) depict the two target variables basal area and aboveground biomass. Graphics (c) to (h) depict six possible predictor variables. Variables are partly lidar-derived (c, d, f, g) and partly field-derived (a, b, e, h).

313 The vertical heterogeneity was measured in several different ways using standard deviation, coefficient of 314 variation and Gini index of the canopy height model and the vertical foliage profile, respectively. CHM-315 and VFP-based vertical structure metrics showed quite different distribution patterns. For the CHM-based 316 metrics, Paracou and SERC showed the lowest values, due to a homogenous canopy surface, BCI and Rabi 317 showed intermediate values, due to their rough canopy surface with large trees alternating with gaps and 318 Traunstein showed (at least for CV and Gini index) the highest values, due to its heterogeneous structure 319 composed of young and old stands interrupted by forest roads. For the VFP-based vertical structure 320 metrics, Traunstein showed the lowest values, which is in accordance with the fact that large parts of the 321 plot are single-layered stands of different age, while the other sites showed higher values, indicating a 322 more complex, multi-layered canopy.

323

324 3.2. Basal Area Estimation

325 3.2.1. Site-specific Basal Area Estimation

326 Here, basal area was estimated from lidar using a single structural descriptor of stand height Ph. Mean top-327 of-canopy height at 1- and 10-m pixel resolution (TCH₁ and TCH₁₀) were tested as P_h to derive site-specific 328 power law coefficients (a_{0,site} and a_{h,site}). Coefficients for each site are listed in Tab. S1 and a scatterplot 329 with site-specific curves is displayed in Fig. 3. The following goodness-of-fit statistics were derived across 330 all sites using the site-specific relationships: The TCH₁-based basal area predictions resulted in 331 RMSE = 2.5 m² ha⁻¹ (8.3%) and R² = 0.79. The TCH₁₀-based basal area predictions resulted in RMSE = 2.8 m² ha⁻¹ (9.5%) and R² = 0.73. In both cases, the mean residuals were not significantly different 332 333 from zero at any site (Wilcoxon tests with the smallest p-value of all sites being p = 0.54). Using multi 334 predictor models (different metrics of P_h and P_v), site-specific predictions with RMSE = 2.3 m² ha⁻¹ (7.6%) 335 and $R^2 = 0.83$ were achieved. However, the set of best metrics varied from site to site (Tab. S2).



Fig. 3: Site-specific relationships (power laws) between basal area and TCH₁ with each point representing 1 ha.

338

339 3.2.2. Generalized Basal Area Estimation

340 Here, basal area was estimated using several structural descriptors from lidar, which were supposed to 341 capture different aspects of forest structure (Equation 5). In total, 125 regression models consisting of 342 different descriptors and metrics were analyzed. The best models found are listed in Tab. 4. The models 343 are ranked according to increasing mean bootstrapped RMSE_b. The listed models represent the best 344 overall model and the best models with certain structural descriptors P_x removed. For the different P_x in 345 most cases the same metrics were selected. For P_h mostly TCH₁₀ was selected and TCH₁ only occurred once. 346 For Pd always the site specific maximal basal area BAsmax was selected. For Pm the site specific maximal tree 347 height H_{smax} was the only available metric. For P_v, however, four different metrics appear in the list of best 348 models, namely SD_{VFP}, Gini_{VFP}, CV_{VFP} and CV_{CHM}.

Tab. 4: The best basal area estimation models for different predictor combinations ranked for increasing mean bootstrapping root mean squared error (RMSE_b). For explanation of the variable names please refer to the main text.

Mean canopy	Maximal stand	Maximal canopy	Vertical	RMSE	nRMSE	R ²	RMSE _b	nRMSEb	R ² b	Bias
height Ph	density P _d	height P _m	heterogeneity P_{ν}							
TCH10	BA _{smax}	H _{smax}	SD _{VFP}	2.9	9.8%	0.71	2.9	9.6%	0.72	yes
TCH10	BA _{smax}	-	Giniver	3.0	10.0%	0.70	3.1	10.4%	0.67	no
TCH1	-	H _{smax}	-	3.3	10.9%	0.64	3.1	10.4%	0.67	yes
TCH ₁₀	BA _{smax}	-	-	3.4	11.5%	0.60	3.6	12.0%	0.56	yes
TCH10	-	-	CVVFP	3.6	12.2%	0.55	3.6	12.0%	0.56	yes
-	BA _{smax}	-	СV _{снм}	4.2	14.2%	0.40	4.2	14.1%	0.39	no
TCH ₁₀	-	-	-	4.4	14.6%	0.36	4.3	14.5%	0.36	yes

352

353 Equation 9:
$$BA = 9.2 \cdot TCH_{10}^{1.3} \cdot BA_{smax}^{0.359} \cdot H_{smax}^{-1.03} \cdot SD_{VFP}^{-0.305}$$

354 The overall best model was one using all four structural descriptors (nRMSE = 9.8%, Equation 9). The 355 goodness-of-fit decreased only marginally if information on maximal possible height (Pm) was excluded 356 from the predictors (nRMSE = 10%). The third best model was a two-predictor model using only current 357 and maximal possible canopy height (P_h and P_m , nRMSE = 10.9%). Hence, there was no other three-358 predictor model that could exceed this two predictor model in accuracy. It was followed by a model using 359 mean canopy height and maximal possible stand density (P_h and P_d , nRMSE = 11.5%), and one using mean canopy height and vertical heterogeneity (P_h and P_v, nRMSE = 12.2%). At the lower end, the best model 360 361 making no use of mean canopy height (no P_h) was somewhat better (nRMSE = 14.2%) than the one using 362 exclusively mean canopy height (P_h = TCH₁₀, nRMSE = 14.6%). Thus, adding any structural descriptor 363 decreased nRMSE by at least 2.4% compared to a model purely based on canopy height.

364 The additional goal was finding a relationship that is unbiased across all sites. According to the Wilcoxon 365 tests, predictions of the best found model were slightly but significantly biased for BCI (p = 0.0036) and 366 Paracou (p = 0.05), whereas the second best model did not show any significant bias for any of the sites (Tab. S3). Fig. 4 shows the 1:1 plots for site-specific TCH₁-based predictions (a), the generalized TCH₁₀-367 368 based predictions (c) and the predictions using the best model based on structural descriptors (e). Fig. 4 369 also shows the residual distributions resulting from each of the three predictions for the different sites (b, 370 d, f). In the leave-one-site-out cross validation of the best predictor combination, the overall nRMSE 371 increased from 9.8% to 12.3% and moderate but significant biases were observed for three sites (Fig. S4).



Site-specific single predictor models

372

Fig. 4: Scatterplots of predicted basal area against observed basal area using a) site-specific single predictor models, c) a general single predictor model (based on TCH₁₀) and e) the best general multi predictor model (Equation 9). The boxplots on the right hand side show the distribution (quartiles) of prediction residuals at each site with numbers below displaying the mean residual value (bias) and asterisks above indicating whether the means deviate significantly from zero (b, d, f).

378 3.3. Aboveground Biomass Estimation

379 3.3.1. Site-specific Aboveground Biomass Estimation

380 Analogous to BA, AGB was modeled as a power law function of mean canopy height P_h, using either TCH₁ 381 or TCH₁₀ by fitting site-specific coefficients (b_{0,site} and b_{h,site}; Tab. S1). Applying these site-specific 382 relationships, the following goodness-of-fit statistics were derived across all sites: The TCH1-based AGB predictions resulted in RMSE = 41 t ha⁻¹ (11.6%) and R^2 = 0.78. The TCH₁₀-based AGB predictions resulted 383 in RMSE = 41.8 t ha⁻¹ (11.8%) and R^2 = 0.77. In both cases, the mean prediction residuals were not 384 385 significantly different from zero at all sites (Wilcoxon tests with the smallest p-value of all sites being 386 p = 0.39). Using multi predictor models (different metrics of P_h and P_v), site-specific predictions with 387 RMSE = 38.1 t ha⁻¹ (10.8%) and R^2 = 0.81 were achieved. However, the set of best metrics varied from site 388 to site (Tab. S2).



389

390 Fig. 5: Site-specific relationships (power laws) between aboveground biomass and TCH₁ with each point representing 1 ha.

392 3.3.2. Generalized Aboveground Biomass Estimation

393	To derive a generalized AGB estimation model the same structural descriptors as for basal area were used.
394	Additionally, a fifth descriptor for the average wood density P_w was introduced, which resulted in 251
395	models in total. Tab. 5 lists the best models found for different combinations of structural descriptors. The
396	models are ranked according to increasing $RMSE_{\flat}$ (derived from bootstrapping). The listed models
397	represent the best overall model and the best models with a reduced number of structural descriptors P_x .
398	Compared to the basal area estimation more descriptor combinations are possible due to the additional
399	parameter P_w . TCH ₁₀ was selected in most cases for P_h . For P_d maximal basal area BA _{smax} and maximal stand
400	density index SDI_{smax} per site do appear in the models. For P_{ν} three different metrics have been selected:
401	SD _{CHM} , SD _{VFP} and CV _{VFP} .

402Tab. 5: The best aboveground biomass estimation models for different predictor combinations ranked for increasing mean403bootstrapping root mean squared error (RMSEb). For explanation of the variable names please refer to the main text.

Mean canopy	Maximal stand	Maximal canopy	Vertical	Mean wood	RMSE	nRMSE	R ²	RMSE _b	nRMSE _b	R ² _b	Bias
height P _h	density P _d	height P _m	heterogeneity P_{v}	density P_w							
TCH1	SDI _{smax}	H _{smax}	SD _{CHM}	WD _{sAGV}	44.0	12.4%	0.74	46.1	13.0%	0.71	no
TCH ₁₀	BA _{smax}	H _{smax}	SDVFP	-	47.4	13.4%	0.70	47.9	13.5%	0.69	yes
TCH10	SDI _{smax}	H _{smax}	-	WD _{sAGV}	45.6	12.9%	0.73	48.5	13.7%	0.69	no
TCH10	SDI _{smax}	H _{smax}	-	-	46.0	13.0%	0.72	48.5	13.7%	0.69	yes
TCH1	-	H _{smax}	-	-	47.7	13.5%	0.70	48.9	13.8%	0.68	yes
TCH10	BA _{smax}	-	CVVFP	WD _{sAGV}	51.2	14.5%	0.67	49.7	14.0%	0.67	yes
TCH10	BA _{smax}	-	-	WD _{sAGV}	60.1	17.0%	0.55	50.6	14.3%	0.66	yes
TCH10	SDI _{smax}	-	-	-	50.9	14.4%	0.66	50.7	14.3%	0.66	yes
TCH10	-	-	CVVFP	-	54.3	15.3%	0.61	53.9	15.2%	0.61	yes
TCH ₁₀	-	-	-	WD _{sAGV}	57.2	16.1%	0.57	57.2	16.1%	0.57	yes
TCH10	-	-	-	-	58.5	16.5%	0.55	58.3	16.4%	0.55	yes
-	BA _{smax}	-	SD _{VFP}	-	63.0	17.8%	0.48	62.2	17.5%	0.49	yes

404

In the case of aboveground biomass estimation the best model was the one using all five available
structural descriptors (Equation 10; nRMSE = 12.4%).

407 Equation 10:
$$AGB = 1.92 \cdot TCH_1^{-1} \cdot SDI_{smax}^{-0.979} \cdot H_{smax}^{-1.24} \cdot SD_{CHM}^{-0.212} \cdot WD_{sAGV}^{-0.0838}$$

408 Leaving either P_v , P_w or both aside increased the nRMSE by around 1%. The best model that did not rely

409 on any site related ground-based information was the one using only P_h and P_m (nRMSE = 13.5%). Tab. 5

410 documents the results for other descriptor combinations. E.g., a single predictor model based on TCH₁₀

411 only had nRMSE = 16.5%. P_d and P_m were more important than P_v (according to their presence in the best

412 models), which is different from the basal area estimation where P_v was more important than P_d and P_m . 413 The best model without P_h had an nRMSE = 17.8%.

414 With regard to the goal of finding a relationship that is unbiased across all sites, the Wilcoxon tests 415 identified two models for which the mean of residuals at none of the single sites differed significantly from 416 zero (Tab. S4): the models in lines 1 and 3 in Tab. 5. For all other models, predictions were biased for at 417 least one site. Fig. 6 shows the 1:1 plots for site-specific TCH₁-based predictions (a), the generalized TCH₁₀-418 based predictions (c) and the structure guided predictions using the best model (e). Fig. 6 also shows the 419 residual distributions resulting from each of the predictions for the different sites (b, d, f). In the leave-420 one-site-out cross validation of the best predictor combination, the overall nRMSE increased from 12.4% 421 to 24.1% and significant overestimations were observed for all sites but Rabi (Fig. S4).

Site-specific single predictor models



General single predictor model



General multi predictor model





Fig. 6: Scatterplots of predicted aboveground biomass against observed aboveground biomass using a) a site specific single predictor model, c) a general single predictor model (based on TCH₁₀) and e) the best general multi predictor model (Equation 10). The boxplots on the right hand side show the distributions (quartiles) of prediction residuals at each site with numbers below displaying the mean residual value (bias) and asterisks above indicating whether the means deviate significantly from zero.

430 3.4. Comparison of Results

431 Overall, achieved relative errors in BA estimation were somewhat lower than the ones for AGB estimation. 432 The exclusion of different structural descriptors led to an increase in estimation errors. Fig. 7 shows the obtained nRMSE for different sets of P_x in comparison to the nRMSE of site-specific estimations. For BA 433 434 estimation, the best unbiased generic model required four coefficients and resulted in an nRMSE of 10.4%, 435 which is 2.1% higher than the nRMSE of 8.3% obtained from five site-specific models, requiring ten (two 436 per site) coefficients. For AGB estimation, the best unbiased generic model required six coefficients and 437 resulted in an nRMSE of 13%, which is 1.4% higher than the nRMSE of 11.6% obtained from five site-438 specific models, requiring ten coefficients.







Fig. 7: Summary of how the exclusion of certain structural descriptors P_x influence the normalized root mean squared error (nRMSE) of basal area (a) and aboveground biomass estimation (b). The black bar represents the site-specific reference model. The grey bars represent mean bootstrapping nRMSE of the different generic models. The striped bars mark the models which produce unbiased predictions at all sites. For the meaning of the indices of the predictors please refer to the main text.

446

448 **4. Discussion**

The goal of this study was to determine a set of forest structure metrics that can be used for BA and AGB estimation from CHMs at very distinct forest sites, which belong to different biomes. It could be shown that a combination of four metrics capturing mean canopy height, maximal stand density, maximal canopy height and vertical heterogeneity could estimate BA using a generic model across all sites with a high accuracy, which was almost as good as the accuracy achieved by site-specific models. The accuracy for AGB estimation was slightly weaker than the one for BA estimation.

455

456 4.1. The Role of Mean Canopy Height

457 It was found that the mean canopy height (P_h), represented here by TCH₁ and TCH₁₀, was the most 458 important predictor variable, which is in support of its wide use in previous studies (Asner and Mascaro, 459 2014; Duncanson et al., 2015; Lefsky et al., 2002). It was important in BA and AGB estimation, with 460 accuracies decreasing considerably when P_h was dropped from the models. Despite the mathematical 461 simplicity of TCH (the mean height of all CHM pixels) it is a quite comprehensive metric capturing much of 462 the forest structure in a single number. It is influenced by the heights and crown sizes of the trees (which 463 contribute to the CHM) and therefore closely related to Lorey's height (BA-weighted mean tree height) 464 (Asner and Mascaro, 2014). However, TCH also provides information on horizontal vegetation density, if 465 ground pixels, e.g., in canopy gaps, are included in its computation (Lu et al., 2014). There have been studies that tried to separate the "height" and "density" aspect of TCH by calculating mean canopy height 466 467 only from canopy pixels (and excluding ground pixels) and capturing horizontal vegetation density as 468 fractional canopy cover, i.e. the relative proportion of canopy pixels above an arbitrary height threshold, 469 or its inverse, the gap fraction (Bouvier et al., 2015). It has also been shown that fractional canopy cover 470 alone can predict AGB in tropical forests quite well over a range of canopy height thresholds (Meyer et al., 471 2018).

472 It was found that TCH₁₀ derived from a rough 10-m-pixel CHM often performed better than TCH₁ derived 473 from 1-m pixels. This has been observed also in an earlier study at BCI using TCH in single predictor models 474 (Knapp et al., 2018). It might be explained by the ability of TCH₁₀ to capture the canopy structure of the 475 large trees, which also contribute most to BA and AGB, and the larger gaps where such trees are missing. 476 TCH₁ includes more detail and is influenced by the structure of individual tree crowns and small gaps within 477 and between crowns, which might not be relevant or even counterproductive for estimating BA and AGB. 478 In particular, in the context of generalization between different forest types it might be beneficial that 479 TCH₁₀ "ignores" differences in crown shapes.

480

481 *4.2.* The Role of Stand Density

482 Maximal stand density per site (P_d) was of high importance for AGB and BA estimation. Only one in the 483 best eight AGB models did not contain Pd. Asner & Mascaro (2014) pointed out that for many sites BA 484 shows a linear relationship with TCH, but with considerable differences in the slopes, which was therefore 485 an important term in their AGB estimation model. Differences in this relationship can be expected because 486 at different sites different tree species may occur, which have different geometries, in particular regarding 487 the relationships between DBH and height and DBH and crown diameter. Of two stands with the same 488 canopy height, one may contain trees with slender crowns and has a much higher stocking than the other 489 one containing trees with wide crowns. We tried to reduce the necessary information about density as 490 much as possible by only using the maximum observed value per site. As this parameter is not derived 491 from remote sensing data, either inventory data or expert knowledge on the maximum possible density 492 of the forest type would be required. As metrics for P_d, BA_{smax} and SDI_{smax} were used. The two are 493 independent from each other: The highest SDI identifies the stand with the highest stocking according to 494 the self-thinning rule (Reineke, 1933), which is not necessarily the stand with the highest current BA 495 (Fig. S5). Among the five sites investigated, the tropical sites had lower BA_{smax} (38.5, 38.7, and 36.7 m² ha⁻ ¹) than the temperate sites (43.5 and 44.6 m² ha⁻¹). SDI_{smax}, however, was similar at BCI, Rabi and SERC (683, 703, 708) and somewhat higher at Paracou (749) and Traunstein (778). As shown, either of the two metrics could improve the AGB estimation in comparison to the case of missing P_d.

499

500 4.3. The Role of Maximum Height

501 An inclusion of maximum possible height (P_m) was expected to improve estimation models. The reason 502 behind is the same as for stand density, namely the possibility of regionally different DBH-height 503 relationships of trees, that lead to differences in the maximum possible canopy height. Pm can be easily 504 extracted from the remote sensing data (in contrast to P_d). Here, the maximum observed canopy height 505 H_{smax} (in the CHM) was used under the assumption that the plots are large enough to be representative 506 for the maximum possible tree height in the respective forest types. Maximum height showed no 507 relevance in BA estimation, but prediction errors for AGB increased from 11.8% to 14.8% if it was dropped 508 from the model. H_{smax} represents the maximum value that TCH could possibly reach, if the whole area 509 would be fully occupied by trees that all have reached their maximum potential height. Hence, H_{smax} might 510 act as a standardization for TCH. Site-specific relationships between TCH and AGB (Fig. 5) show that, e.g., 511 the forest at Traunstein reaches high AGB values at much lower mean canopy heights than other sites. By 512 additionally providing the information that also H_{smax} at Traunstein is lower than elsewhere, the TCH values 513 are put into the perspective of how high is the forest now and how high could it possibly become. This 514 standardization role of H_{smax} is supported by the fact that all selected models have negative coefficients 515 (a_m, b_m) , commonly close to -1 for H_{smax}, and positive coefficients (a_h, b_h) , commonly close to +1 for TCH, 516 i.e., the ratio TCH / H_{smax} is used in the predictions. Models based on TCH₁ and H_{smax} only were the best 517 two-predictor models in BA and AGB estimation, respectively.

519 4.4. The Role of Vertical Heterogeneity

520 Vertical heterogeneity (P_v) was after mean canopy height (P_h) the second descriptor derived at individual 521 plot-level rather than site-level. It was included in the best BA and AGB models, however, dropping it 522 increased the nRMSE by less than 1%. From the six candidate metrics for P_v , various were chosen in 523 different models.

524 The calculation of the vertical metrics was either based on the canopy height model or on the vertical 525 foliage profile. As visible in Fig. S3, the distributions of SD, CV and Gini index differed strongly depending 526 on whether they were CHM- or VFP-based. CHM-based variability metrics describe the heterogeneity of 527 the canopy surface, including ground pixels, i.e., canopy gaps. VFP-based variability metrics describe the 528 vertical layering of the reconstructed foliage profile, which does not contain any ground component, but 529 up-weights profile parts in the lower heights to compensate for the occlusion by high trees. Hence, their 530 contributions to BA and AGB estimation might be different: CHM-based metrics rather characterize forests 531 in the spectrum from smooth canopy surfaces, as observed for young, dense stands, to rough canopy 532 surfaces, as observed for old or disturbed stands. VFP-based metrics rather account for the overseen trees 533 in the lower canopy. Other studies have also identified the vertical heterogeneity as a component in 534 prediction models. Magnussen et al. (2012) proposed a two-predictor model based on 1) TCH and 2) the 535 variance of the CHM divided by TCH, which is closely related to CV_{CHM} used here. Bouvier et al. (2015) 536 considered two vertical metrics in their four-predictor model: 1) variance of the CHM and 2) CV of the leaf 537 area density in the VFP. In our analysis, vertical heterogeneity was also able to improve site-specific 538 estimation models compared to solely TCH-based models (about 1% decrease in nRMSE) with different 539 vertical metrics being chosen at different sites. To conclude, there is a wide variety of metrics that 540 characterize vertical heterogeneity and they may in fact capture quite different aspects of forest structure. 541 They do contribute in the improvement and generalization of BA and AGB estimation. Future analyses 542 should try to achieve a better understanding of how the different metrics are related to ground-based 543 metrics of forest structure, and whether a combination of several of them could further improve 544 estimation results.

545

546 4.5. The Role of Wood Density

547 Regional differences in average wood density have been suspected to be a main reason behind differences 548 in the height-to-biomass relationship of forests (Asner and Mascaro, 2014; Meyer et al., 2018; Vincent et 549 al., 2014). In our analysis, however, dropping the wood density parameter (P_w) led only to a slight increase 550 in nRMSE of less than 1% for AGB estimations. The values of WD_{sAGV} were very similar for BCI, SERC and 551 Traunstein, but considerably higher for Paracou and Rabi. If region-specific estimates on average wood 552 density are available, they should definitely be considered in AGB estimation models. Nevertheless, our 553 results suggest that compared to other parameters wood density is of minor importance for a generalized 554 AGB estimation. With regard to how average wood density should be calculated, Vincent et al. (2014) 555 argued to use AGV instead of BA as a weighting variable, as AGV of the trees in the ground-truth plots has 556 to be calculated anyway to derive tree AGB, and AGV is the structurally more appropriate weighting 557 variable compared to BA. In this study, for all five study sites WD_{sAGV} and WD_{sBA} were found to be very 558 similar. Thus, only WD_{sAGV} was further used in the analyses.

559

560 4.6. Generalization and Outlook

With the identified structural variables and the fitted coefficients, we propose general prediction models for BA and AGB estimation, which are applicable on temperate and tropical forests under natural and managed conditions. Having such models and also understanding the contribution of different forest structural aspects is important for consistent large scale mapping of forest carbon stocks (Lefsky et al., 2002). This is particularly relevant for upcoming spaceborne missions such as GEDI (Hancock et al., 2019; Stavros et al., 2017), ICESat-2 (Narine et al., 2019), BIOMASS (Le Toan et al., 2011) or Tandem-L (Moreira et al., 2015) which will provide consistent forest height measurements across very different forest types,
not all of which are represented sufficiently in ground-truth datasets.

569 As a next step, the proposed relationships need to be tested at other forest sites to either confirm or, if 570 necessary, adapt them. The results of the leave-one-site-out cross validation suggest that the presented 571 approach for BA estimation is more robust than the one for AGB estimation, with regard to reducing the 572 number of forest types for fitting. The distributions of the various structure metrics have shown that all of 573 the five sites differ in one or another aspect from all others, which apparently makes each of them essential 574 in the calibration of the AGB prediction model. In future analyses, further datasets need to be taken into 575 account. Achieving robust leave-one-site-out cross validation results will require a sufficient degree of 576 structural redundancy among sites, i.e., several sites representing similar forest types.

The Traunstein site was the most distinct site concerning various structural aspects. It remains unclear to which degree this can be explained by ecological differences alone (e.g., only site with large proportion of conifers), and to which degree the intensive management there plays a role. Forest management may alter some of the relationships among structure metrics, compared to natural stands. Future research should try to identify such changes. In case they are significant, management would be an additional aspect which should be considered in generalization approaches. Remote sensing methods for estimating the management regimes and parameters could then complement the biomass estimation.

Furthermore, the influence of spatial scale needs to be investigated, as different sensors produce measurements at different scales (Knapp et al., 2018; Tello et al., 2018). Finally, methods need to be developed for acquiring more of the structural variables entirely from remote sensing and becoming independent from any ground-based input. Individual tree delineation from high resolution canopy height data can be applied to derive stand density information directly from remote sensing (Duncanson et al., 2015; Ferraz et al., 2016). Average wood density can be estimated based on forest type or even species classification using passive optical remote sensing (Fassnacht et al., 2016). These technologies have to be

- combined to derive very detailed estimates from airborne acquisitions at landscape scale. The estimatescan then serve as training areas for wall-to-wall mapping using spaceborne products.
- 593

594 **5. Conclusion**

595 Data from temperate and tropical forest plots was combined to develop a general equation for biomass 596 (and basal area) estimation based on a set of forest structure metrics from lidar remote sensing. The 597 different structural predictors were a priori defined. The results provided insight in the relative importance 598 of mean and maximal canopy height, stand density, vertical heterogeneity and wood density for biomass 599 estimation. Not all of those forest attributes can be derived from lidar data. For maximal stand density and 600 mean wood density field-based information is required at the site level. Alternatively, a model without 601 those attributes can be chosen from the list of models, at the expense of slightly lower prediction 602 accuracies. The found relationships should provide guidance towards a standardized workflow for 603 estimating aboveground biomass for forest carbon mapping and monitoring from remote sensing.

604

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