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1 Title

2 Pan-tropical prediction of forest structure from the largest trees

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195 Abstract

Aim. Large tropical trees form the interface between ground and airborne observations, offering a unique opportunity to capture forest properties remotely. However, despite rapid development of metrics to characterize the forest canopy from remotely sensed data, a gap remains between aerial and field inventories. To close this gap, we propose a new pan-tropical model to predict plot-level forest structure properties and biomass from just the largest trees, as a proxy for the whole plot inventory.

202 **Location.** Pan-tropical

Method. Using a dataset of 867 plots distributed among 118 sites across the tropics, we tested the ability to predict quadratic mean diameter, basal area, Lorey's height and community wood density from the ith largest trees, i.e. testing the cumulative information gathered from these i trees ranked by decreasing diameter. These tests served as a basis to select the optimal number of the largest trees and further predict plot-level biomass from a single model.

208 **Result.** Focusing on readily available information captured by airborne remote sensing, we 209 show that measuring the largest trees in tropical forests enables unbiased predictions of plot 210 and site-level forest structure. The 20 largest trees per hectare predicted quadratic mean 211 diameter, basal area, Lorey's height and community wood density with 12%, 16%, 4% and 4% of relative error. Building on this result, we developed a new model to predict plot-level AGB 212 213 from measurements of the 20 largest trees. This model allows an independent and unbiased 214 prediction of biomass with 17.7% of error compared to ground estimates. Most of the remaining 215 error is driven by differences in the proportion of total biomass held in medium size trees (50-70 cm), which shows some continental dependency with American tropical forests presenting 216 217 the highest levels of total biomass share in these intermediate diameter classes.

Conclusion. Our approach provide new information on tropical forest structure and can be employed to generate accurately field estimates of tropical forest carbon stocks to support the calibration and validation of current and forthcoming space missions. It will reduce the cost of programs to monitor, report, and verify forest resources, and will contribute to scientific understanding of tropical forest ecosystems and response to climate change.

223 Introduction

224 The fundamental ecological function of large trees is well established for tropical forests. They 225 offer shelter to a multiple organisms (Remm & Lõhmus, 2011; Lindenmayer et al., 2012), 226 regulate forest dynamics, regeneration (Harms et al., 2000; Rutishauser et al., 2010) and total biomass (Stegen et al., 2011), and are important contributor to the global carbon cycle 227 (Meakem et al., 2017). Being major components of the canopy, the largest trees also suffer 228 more than sub-canopy and understory trees from climate change, as they are directly exposed 229 230 to variations in solar radiation, wind strength, temperature seasonality and relative air humidity (Laurance et al., 2000; Nepstad et al., 2007; Lindenmayer et al., 2012; Thomas et al., 2013; 231 Bennett et al., 2015; Meakem et al., 2017). Because they are visible from the sky, large trees 232 are ideal for monitoring forest responses to climate change via remote sensing (Bennett et al., 233 234 2015; Asner et al., 2017).

235 Large trees encompass a disproportionate fraction of total above-ground biomass (AGB) in 236 tropical forests (Chave et al., 2001), with some variations in their relative contribution to the 237 total AGB among the tropical regions (Feldpausch et al., 2012). In Central Africa, the largest 238 5% of trees, i.e. the 5% of trees with the largest diameter at 130 cm per area, store 50% of forest aboveground biomass on average (Bastin et al., 2015). Consequently, the density of 239 large trees largely explains variation in AGB at local (Clark & Clark, 1996), regional (Malhi et 240 al., 2006; Saatchi et al., 2007), and continental scales (Stegen et al., 2011; Slik et al., 2013). 241 242 Detailing the contribution of each single tree to the diameter structure, we showed previously 243 that plot-level AGB can be predicted from a few large trees (Bastin et al., 2015), with the measurement of the 20 largest trees per hectare being sufficient to estimate plot-level biomass 244 with less than 15% errors in reference to ground estimates. These findings opened the 245 246 possibility of measuring the largest trees to cost-effectively monitor forest biomass in Central Africa, rather than conducting full inventories of all size classes. Similarly, they suggested that 247 248 remote sensing (RS) approaches should focus on the measurement of the largest trees, instead of properties of the entire forest. 249

Several efforts are underway to close the gap between remote sensing and field surveys (e.g. 250 Jucker et al. 2016a, Coomes et al. 2017). However, field inventories still rely on exhaustive 251 data collection, while remote sensing surveys provide a limited alternative for the following 252 253 reasons. Existing RS approaches that provide predictions of biomass with less than 20% error 254 for 1 ha plot size are either specific to the relationship between forest type and image/scene properties (Barbier et al., 2011; Asner et al., 2012; Barbier & Couteron, 2015), or require 255 ground measurement of all trees above or equal to 10 cm of D for calibration (Asner et al., 256 257 2012; Asner & Mascaro, 2014). Using mean canopy height extracted from active sensors (Mascaro et al., 2011; Ho Tong Minh et al., 2016), or canopy grain derived from optical images 258 (Proisy et al., 2007; Ploton et al., 2012, 2017; Bastin et al., 2014), the biomass is predicted 259 from remote sensing with a typical error of only 10-20% compared to ground-based estimates, 260 but is limited to the extent of the scene used. An interesting development to alleviate this spatial 261 restriction lies in the 'universal approach', proposed by Asner et al. (2012) and further adapted 262 in Asner and Mascaro (2014), in which plot-level biomass is predicted by a linear combination 263 264 of ground-based and remotely-sensed metrics. The 'universal approach' relies upon canopy 265 height metrics derived from radar or LiDAR (top of canopy height, TCH), and basal area (BA, 266 i.e. the cross-sectional stem area) and community wood density (i.e. weighted by basal area, WD_{BA}) derived from full field inventories. AGB is then predicted as follows (Asner *et al.*, 2012): 267 268 $AGB = aTCH^{b1}BA^{b2}WD_{BA}^{b3}(1)$

While generally performing better than approaches based solely on remote sensing of tree height (Coomes *et al.*, 2017), this model largely relies on exhaustive ground measurements (i.e. wood density and basal area of all trees above 10 cm of diameter at 130 cm, neither of which is measured using any existing remotely sensed data).

Recent advances in remote sensing allow the identification of single trees in the canopy (Ferraz *et al.*, 2016), estimation of adult mortality rates for canopy tree species (Kellner & Hubbell,
2017), description of the forest diameter structure (Stark *et al.*, 2015), depiction of crown and
gap shapes (Coomes *et al.*, 2017), and even identification of some functional traits of canopy
species (Asner *et al.*, 2017). Building upon this work, we test the capacity of metrics from the

278 largest trees that can be potentially derived using remote sensing to predict plot-level biomass 279 (i.e. the summed AGB of all live trees $D \ge 10$ cm in a plot). To this end, we tested the following 280 model:

281 AGB = $a(Dg_{LT}, H_{LT}, WD_{LT})^{b1}$ (2)

282 Where for the ith largest trees, Dg_{LT} is the quadratic mean diameter, H_{LT} the mean height, and 283 WD_{LT} the mean wood density averaged among the ith largest trees.

Using a large database of forest inventories gathered across the Tropics (Figure 1), including 284 285 secondary and old growth forest plots, we test the ability of the largest trees to provide information on various metrics estimated at 1-ha plot level, such as the mean guadratic 286 diameter, the basal area (BA), the Lorey's height (i.e. plot-average height weighted by BA), the 287 community wood density (i.e. plot-average wood density weighted by BA) and mean above-288 ground live biomass (AGB) (supplementary figure 1). While previous work focused on 289 estimating biomass in Central African forests (Bastin et al., 2015), the present study aims at 290 generalizing the potential of large trees in predicting these different plot metrics at continental 291 292 and pan-tropical scales. Taking advantage of a unique dataset gathered across the tropics (XX 293 ha, YYY plots), we also investigate major differences in forest structure across the three main 294 tropical regions, South America, Africa and South East Asia. We further discuss how this approach can be used to guide innovative RS techniques and increase the frequency and 295 296 representativeness of ground data to support global calibration and validation of current and 297 planned space missions. These include the NASA Global Ecosystem Dynamics Investigation 298 (GEDI), NASA-ISRO Synthetic Aperture Radar (NISAR), and ESA P-band radar (BIOMASS). This study is a step forward in bringing together remote sensing and field sampling techniques 299 for quantification of terrestrial C stocks in tropical forests. 300

301 Material & Methods

302 Database

For this study, we compiled standard forest inventories conducted in 867 1-hectare plots from 303 304 118 sites across the three tropical regions (Figure 1), including mature and secondary forests. Each site comprises all the plots in a given geographical location, i.e. within a 10 km radius 305 and collected by a PI and its team. These consisted of 389 plots in America (69 sites), 302 306 plots in Africa (35 sites) and 176 plots in Asia (14 sites). Data were provided by Principal 307 308 Investigators (see supplementary Table 1), and through datasets available at ForestPlots (https://www.forestplots.net/), TEAM (http://www.teamnetwork.org/) CTFS 309 and

310 (<u>http://www.forestgeo.si.edu/</u>) networks.

We selected plots located between 23°N and 23°S, including tropical islands, with an area of 311 312 at least 1-ha to ensure stable intra-sample variance in basal area (Clark & Clark, 2000). Plots 313 in which at least 90% of the stems were identified to species, and in which all stems with the diameter at 130 cm greater than or equal to 10 cm had been measured were included. Wood 314 315 density, here recorded as the wood dry mass divided by its green volume, was assigned to 316 each tree using the lowest available taxonomic level of botanical identifications (i.e. species or 317 genus) and the corresponding average wood density recorded in the Global Wood Density Database (GWDD, Chave et al., 2009; Zanne et al., 2009). Botanical identification was 318 319 harmonized Taxonomic Names Resolution Service through the 320 (http://tnrs.iplantcollaborative.org), for both plot inventories and the GWDD. For trees not 321 identified to species or genus (~5%), we used plot-average wood density. We estimated heights of all trees using Chave et al.'s (2014) pan-tropical diameter-height model which 322 accounts for heterogeneity in the D-H relationship using an environmental proxy: 323

324 $Ln(H) = 0.893 - E + 0.760 ln(D) - 0.0340 ln(D)^{2} (3)$

Where *D* is the diameter at 130cm and *E* is a measure of environmental stress (Chave *et al.*, 2014). For sites with tree height measurements (N=20), we developed local D-H models, using a Michaelis-Menten function (Molto *et al.*, 2014). We used these local models to validate the

328 predicted Lorey's height (i.e. plot average height weighted by BA) from the largest trees, of 329 which height has been estimated with a generic H-D model (equation 3, Chave et al. 2014).

We estimated plot biomass as the sum of the biomass of live tree with diameter at 130 cm

superior or equal to 10 cm, using the following pan-tropical allometric model (Réjou-Méchain

et al., 2017):

333 AGB=exp(-2.024-0.896E+0.920ln(WD)+2.795ln(D)-0.0461(ln(D²))) (4)

Plot-level metric estimation from the largest trees

335 The relationship between each plot metric, namely basal area (BA), the quadratic mean diameter (Dg), Lorey's height (H_{BA:} the mean height weighted by the basal area) and the 336 community wood density (WD_{BA}; the mean wood density weighted by the basal area), and 337 those derived from largest trees was determined using an iterative procedure following Bastin 338 339 et al. (2015). Trees were first ranked by decreasing diameter in each plot. An incremental procedure (i.e. including a new tree at each step) was used to sum or average information of 340 the *i* largest trees for each plot metric. Specifically, each plot-level metric was predicted by the 341 342 respective metric derived from the ith largest trees. For each increment, the ability (goodness 343 of fit) of the *i* largest trees to predict a given plot-metric was tested through a linear regression. To avoid overfitting, a Leave-One-Out procedure was used to develop independent site-344 specific models (N=118). Specifically, the model to be tested at a site was developed with data 345 from all other sites. Errors were then estimated as the relative root mean square error (rRMSE) 346 347 computed between observed and predicted values (X):

348 $rRMSE = \bar{X} \sum \sqrt{\frac{(Xobs - Xpred)^2}{n}}$ (5)

The form of the regression model (i.e. linear, exponential) was selected to ensure a normal distribution of the residuals.

To estimate plot basal area, we used a simple power-law constrained on the origin, as linear model resulted in non-normal residuals. Plot-level basal area (BA) was related to the basal area for the *i* largest trees (BA_i) using:

354 BA =
$$b_1 \Sigma BA_i^{\gamma 1}$$
 (6)

To estimate the quadratic mean diameter, Lorey's height and the wood density of the community, we used simple linear models relating the plot-level metrics and the value of the metrics for the *i* largest trees:

358 $D_g = a_2 + b_2 D_{gi}$ (7)

359
$$H_{BA} = a_3 + b_3 H_i$$
 (8)

360 $WD_{BA} = a_4 + b_4 \overline{WD_i}$ (9)

Both Lorey's height (H_{BA}) and the average height ($\overline{H_i}$) of the ith largest trees depend on the same D-H allometry, which always contains uncertainty whether we use a local, a continental or a pan-tropical model. To test the dependence of the prediction of H_{BA} from $\overline{H_i}$ on the allometric model, we used measurement from Malebo in the Democratic Republic of the Congo, where all heights were measured on the ground (see supplementary figure 2).

The quality of the predictions of plot-level metrics from the largest trees is quantified using the relative root mean square error (rRMSE) between measured and predicted values, and displayed along the cumulated number of largest trees (Figure 2). Model coefficients are estimated for each metric derived from the largest trees (N_{LT}) and averaged across the 118 models (see supplementary table 2).

371 Mean rRMSE is plotted as a continuous variable, while its variation is presented as a 372 continuous area between 5th and the 95th percentiles of observed rRMSE (Figure 2).

373 The optimal number of largest trees for plot-level biomass estimation

The optimal number of largest trees N_{LT} was determined from the prediction of each plot-level metric considered above, i.e. keeping a small number of trees while ensuring a low level of error for each structural parameter. We then predicted plot-level biomass from the N_{LT} model (equation 2). The final error was calculated by propagating the entire set of errors related to equation 4 (Réjou-Méchain *et al.*, 2017) in the N_{LT} model (i.e. error associated to each allometric model used). The model was then cross-validated across all plots (N=867).

380 Investigating residuals: what the largest trees do not explain

To understand the limits of predicting AGB through NLT, we further investigated the relationship 381 between AGB residuals and key structural and environmental variables using linear modelling. 382 383 Forest structure was investigated through the total stem density (N), the guadratic mean 384 diameter (Dg), Lorey's height (H_{BA}) and community wood density (WB_{BA}). As environmental 385 data, we used the mean annual rainfall and the mean temperature computed over the last 10 years at each site using the Climate Research Unit data (New et al., 1999, 2002), along with 386 rough information on soil types (Carré. et al., 2010). Major soil types were computed from the 387 388 soil classification of the Harmonized World Soil Database into IPCC (intergovernmental panel on climate change) soil classes. In addition, considering observed differences in forest 389 structure across tropical continents (Feldpausch et al., 2011) and recent results on pan-tropical 390 floristic affinities (Slik et al., 2015), we tested for an effect of continent (America, Africa and 391 Asia) on the AGB residuals. 392

393 The importance of each variable was evaluated by calculating the type II sum of squares that 394 measures the decrease in residual sum of squares due to an added variable once all the other variables have been introduced into the model (Langsrud, 2003). Residuals were investigated 395 396 at both plot and site levels, the latter analyzed to test for any influence of the diameter structure, 397 which is usually unstable at the plot level due to the dominance of large trees on forest metrics at small scales (Clark & Clark, 2000). Here we use a principal component analysis (PCA) to 398 summarize the information held in the diameter structure by ordinating the sites along the 399 400 abundance of trees in each diameter class (from 10 to +100 cm by 10 cm bins).

402 Results

403 **Plot-level metrics**

404 Plot metrics averaged at the site level (867 plots, 118 sites) present important variations within 405 and between continents. In our database, the quadratic mean diameter varies from 15 to 42 cm²ha⁻¹, the basal area from 2 to 58 m²ha-1, Lorey's height from 11 to 33 m and the wood 406 density weighted by the basal area from 0.48 to 0.84 gcm⁻³ (Supplementary figure 1). Such 407 important differences between minimal and maximal values are observed because our 408 409 database cover sites with various forest types, from young forest colonizing savannas to old growth forest. However, most of our sites are found in mature forests, as shown by relatively 410 high average and median value of each plot metric (average aboveground biomass = 302 411 Mgha⁻¹; supplementary figure 1). In general, highest values of aboveground biomass are found 412 in Africa, driven by highest values of basal area and highest estimations of Lorey's height. 413 414 Highest values of wood density weighted by basal area are found in America.

415 **Plot-level estimation from the i largest trees**

Overall, plot metrics at 1 ha scale were well predicted by the largest trees, with qualitativeagreement among global and continental models (Figure 2).

418

When using the 20 largest trees to predict basal area (BA) and quadratic mean diameter (Dg), 419 420 the mean rRMSE was < 16% and 12%, respectively (Figs 3a and 3b). Lorey's height (H_{BA}) and 421 wood density weighted by basal area (WD_{BA}) were even better predicted (Figs 3c and 3d), with mean rRMSE of 4% for the 20 largest trees. The prediction of Lorey's height from the largest 422 trees using local diameter-height model (supplementary Figure 2a) yielded results similar to 423 those obtained using equation 3 of Chave et al. (2014). More importantly, it also yielded similar 424 425 results to prediction of Lorey's height from the largest trees using plots where all the trees were measured on the ground (supplementary figure 2b). This suggests that our conclusions are 426 robust to the uncertainty introduced by height-diameter allometric models. 427

428 AGB prediction from the largest trees

- 429 We selected "20" as the number of largest trees to predict plot metrics. The resulting model
- 430 predicting AGB (Mg ha⁻¹) based on the 20 largest trees is:
- 431 AGB = $0.0735 \times (Dg_{20}H_{20}WD_{20})^{1.1332}$ (rRMSE=0.179; R²=0.85; AIC= -260.18) (10)
- 432 Because the exponent was close to 1, we also developed an alternative and more operationa
- 433 I model with the exponent constrained to 1, given by:
- 434 AGB = $0.195 \times (Dg_{20}H_{20}WD_{20})$ (rRMSE=0.177; R²=0.85; AIC=-195) (11)
- Ground measurements of plot AGB were predicted by our N_{LT} model with the exponent constrained to 1, with a total error of 17.9% (Figure 4), a value which encompass the error of the N_{LT} model and the error related to the allometric model chosen. The Leave-One-Out crossvalidation procedure yielded similar results (rRMSE=0.19; R²=0.81), validating the use of the model on independent sites.

440 **Determining the cause of residual variations**

The explanatory variables all together explain about 37% of the variance in AGB both at plot 441 and site levels when omitting the diameter structure, and about 63% at site level when included 442 443 (Fig. 5). In general, forest structure and particularly the stem density explained most of the 444 residuals (table 1; weights: 79% and 54% at plot- and site-level respectively). The stem density 445 was followed by a continental effect (weights: 18%, 28% and 1%, respectively for Africa, South America and Asia) and by the effect of H_{BA} and WD_{BA} (respective weights: 1% and 0% at the 446 447 plot level, 0% and 11% at the site level, and 23% and 0% when accounting for the diameter 448 structure at the site level). Inclusion of the diameter structure provided the best explanation of 449 residuals, with 63% of variance explained, and a weight of 69% for the first axis of the PCA (supplementary figure 3). This first axis of the PCA was related to the general abundance of 450 trees at a site, and in particular medium-sized trees (40-60cm). Among environmental 451 452 variables, only rainfall was significantly related to the residuals at the site level when the diameter structure was considered (2%). 453

454 **Discussion**

455 The largest trees, convergences and divergences between continents

456 Sampling a few largest trees per hectare generally allows an unbiased prediction of four key 457 descriptors of forest structures across the Tropics. There is generally no improvement in predicting basal area, quadratic mean diameter, Lorey's height (H_{BA}) or community wood 458 density beyond the first 10-to-20 largest trees (Figure 2, Figure 3a). In some cases, e.g. when 459 a forest plot presents an abundant number of large trees (Figure 5d), increasing the number 460 461 of trees sampled improves the model's accuracy. This is the case for BA for which rRMSE continues to decrease up to 100 largest trees (Figure 2a). In contrast, Lorey's height 462 predictions are altered when a large number of trees are included (Figure 2c), i.e. when 463 smaller, often suppressed, trees draw the average down (Farrior et al., 2016). This might 464 explain why the prediction of AGB does not mirror that of basal area (Figure 2b, Figure 3a), 465 466 and suggest that the number of largest trees shall be set independently to each predictor considered. Interestingly, the evolution of relative error in AGB prediction as a function of the 467 468 number of largest trees considered does not follow the same path between continents. For 469 instance, the error of prediction saturates more quickly in Africa and Asia than Asia, where 470 high variations of residuals are observed. Investigation of residuals showed that the diameter structure (Figure 5c, supplementary Figure 3b), and in particular the number of medium size 471 472 trees (Figure 5d), drives variability in AGB predictions. It is therefore not surprising to see that 473 in our dataset the site with higher levels of underestimations is the one with the highest number 474 of medium size trees, which is found in Asia in the Western Ghats of India.

The good performance of models based on the 20 largest trees in predicting Lorey's height and community wood density at site level was not surprising. Both metrics were indeed weighted by basal area, driven *de facto* by the largest trees. Their consistency across sites and continents was not expected though. This suggests that the relationship between the 20 largest trees and descriptors of forest structures is stable across the tropics, and prove the generality of our approach. Slight differences are however noticeable when comparing the distribution of the pan-tropical model residuals across continents (Figure 6, supplementary figure 4).In America, our pan-tropical model tend to slightly underestimate basal area (mean: -5%) and overestimate Lorey's height (mean: +3%) (supplementary figure 4), suggesting peculiar forest structures (i.e. higher tree height for a given diameter, and lower fractions of large trees, supplementary figure 2). In Asia, and in particular in Africa, large (i.e. DBH > 50 cm) trees are more abundant and encompass a large fraction of plot biomass. The basal area tends to be slightly overestimated in Africa, resulting in average to a 3% overestimation of AGB (Figure 6a).

Interestingly, while a recent global phylogenetic classification of tropical forest groups American with African forests vs. Asian forests (Slik *et al.*, 2018), our results tend more to single out American forests. Although this deserves further investigations, it might reveal a lack of close relationship between forest structure properties and phylogenic similarity, which echoes recent results on the absence of relationship between tropical forest diversity and biomass (Sullivan *et al.*, 2017).

495 Largest trees, a gateway to global monitoring of tropical forests

496 Revealing the predictive capacity held by the largest trees, our results constitute a major step 497 forward to monitor forest structures and biomass stocks. The largest trees in tropical forests 498 can therefore be used to accurately predict and efficiently infer various ground-measured properties (i.e. the quadratic mean diameter, the basal area, Lorey's height and community 499 500 wood density), while previous work has predicted only biomass "estimates" (e.g. Slik et al., 501 2013; Bastin et al., 2015). This approach allows us to (i) describe forest structure independently 502 of any biomass allometric model (ii) and cover local variations in D-H relationship, known to vary locally (Feldpausch et al., 2011; Kearsley et al., 2013;). It is also (iii) relatively insensitive 503 504 to differences in floristic composition and community wood density (Poorter et al., 2015).

505 Furthermore, the "largest trees" models were developed for each plot-level metric and for any 506 number of largest trees. Thus, they do not rely on any arbitrary threshold of tree diameter. Note 507 that the optimal number of largest trees to be measured (i.e. 20) was set for demonstration 508 and can vary depending on the needs and capacities of each country or project (see 509 supplementary table 2). In the same way, local models could integrate locally-developed 510 biomass models, when available. Consequently our approach (i) can be used in young or 511 regenerating un-managed forests with a low "largest tree" diameter threshold and (ii) is 512 compatible with recent remote sensing approaches able to single out canopy trees and 513 describe their crown and height metrics (Ferraz *et al.*, 2016; Coomes *et al.*, 2017).

514 Aboveground biomass model from the largest trees, a multiple opportunity

Globally, the N_{LT} model for the 20 largest trees allows plot biomass to be predicted with 17.9%
error. This result is a pan-tropical validation of results obtained in Central Africa (Bastin *et al.*,
2015). It opens new perspectives towards cost-effective methods to monitor forest structures
and carbon stocks through largest trees metrics, i.e. metrics of objects directly intercepted by
remote-sensing products.

Developing countries willing to implement a Reduction of Emissions from Deforestation and 520 521 Forest Degradation (REDD+), shall also report on their carbon emissions (CE) and develop a national CE reference level (IPCC, 2006; Maniatis & Mollicone, 2010). However, most tropical 522 countries lack capacities to assume multiple, exhaustive and costly forest carbon assessment 523 524 (Romijn et al., 2012). By measuring only a few large trees per hectare, our results show that 525 it is possible to obtain unbiased estimates of aboveground C stocks in a time and cost-efficient 526 manner. Assuming that 400 to 600 trees D > 10 cm are measured in a typical 1-ha sample plot, monitoring only 20 trees is a significant improvement. Although finding the 20 largest trees 527 in a plot of several hundred individuals requires evaluating more than 20 trees, in practice, a 528 529 conservative diameter threshold could be defined to ensure that the 20 largest trees are 530 sampled. An alternative approach could also be found in the development of relascope-based approach adapted to detection of the largest trees in tropical forests. Using such approach 531 would facilitate rapid field sampling in extensive areas to produce large scale AGB estimates. 532 Those could fullfil the needs in calibration and validation of current and forthcoming space 533 missions focused on aboveground biomass. 534

535 Our findings also points towards the potential effectiveness of using remote sensing 536 techniques to characterize canopy trees. Here, remote sensing data could be used for direct 537 measurement (e.g. tree level metrics such as height, crown width, crown height) of the largest

trees instead of indirect development of complex metrics (e.g. mean canopy height, texture) 538 used to extrapolate forest properties. While some further refinements are needed, most of the 539 tools required to develop "largest trees" models are readily available. In particular, Ferraz et 540 541 al. (2016) developed an automated procedure to locate canopy trees based on airborne LiDAR data, to measure their height and crown area. Crown area could further be linked to basal area, 542 as the logarithm of crown area is consistently correlated with a slope of 1.2-1.3 to the logarithm 543 of tree diameter across the tropics (Blanchard et al., 2016). Regarding wood density, 544 545 hyperspectral signature offers a promising way to assess functional traits remotely (e.g. Asner et al., 2017) which could potentially be tested to detect variability in wood properties. 546 Alternative approaches could focus on the development of plot-level AGB prediction by 547 replacing the basal area of the largest trees with their crown metrics. While the measurement 548 549 of crown areas have yet to be generalized when inventorying plots, several biomass allometric models already partition trunk and crown mass (Jucker et al., 2016; Ploton et al., 2016; 550 551 Coomes et al., 2017).

552 The main limitation of our approach lies in the understory and sub-canopy trees. We show that 553 most of the remaining variance is explained by variations in diameter structures, and in particular among the total stem density. Interestingly, stem density was generally identified as 554 a poor predictor of plot biomass in tropical forests (Slik et al., 2010; Lewis et al., 2013). 555 556 However, our results show that stem density explains most of the remaining variance (Table 557 S1). This suggests that, in addition to trying to understand large-scale variations in large trees 558 and other plot metrics, which can be directly quantified from remote sensing, we should also put more effort into understanding variation in smaller trees, which mainly drives total stem 559 density and the total floristic diversity. Smaller trees are also essential to characterize forest 560 561 dynamics and understand changes in carbon stocks. Several options are nonetheless possible from remote sensing, considering the variation in lidar point density below the canopy layer 562 (D'Oliveira et al., 2012), the distribution of leaf area density (Stark et al., 2012, 2015; Tang & 563 Dubayah, 2017) or the use of multitemporal lidar data to get information on forest gap 564

generation dynamics and consequently on forest diameter structure (Kellner *et al.*, 2009;
Farrior *et al.*, 2016).

567 Large trees in degraded forests

568 If large trees are a key feature of unmanaged forests, they are conspicuously absent from managed or degraded forests. Indeed, large trees are targeted by selective or illegal logging, 569 and are the first to disappear or to suffer from incidental damages when tropical forests are 570 exploited for timber (Sist et al., 2014). The loss of largest trees drastically changes forest 571 572 structures and diameter distributions, and their loss is likely to counteract the consistency in forest structures observed through this study. Understanding how, or whether, managed 573 forests deviate from our model predictions could help characterize forest degradation, which 574 accounts for a large fraction of carbon loss worldwide (Baccini et al., 2017), acknowledging 575 576 that rapid post-disturbance biomass recovery (Rutishauser et al., 2015) will remain hard to 577 capture.

578 Conclusion – towards improved estimates of tropical forest biomass

579 The acquisition, accessibility and processing capabilities of very high spatial, spectral and 580 temporal resolution remote sensing data has increases exponentially in recent years (Bastin 581 et al., 2017). However, to develop accurate global maps, we will have to obtain a greater number of field plots and develop new ways to use remote sensing data. Our results provide 582 583 a step forward for both by (i) decreasing drastically the number of individual tree measurements 584 required to get an accurate, yet less precise, estimate of plot biomass and (ii) opening the door 585 to direct measurement of plot metrics measured from remote sensing to estimate plot biomass. As highlighted by Clark and Kellner (2012), new biomass allometric models relating plot-level 586 biomass measured from destructive sampling and plot-level metric measured from remote-587 588 sensing products should be developed, as an alternative to current tree-level allometric models. Such an effort will lead largely to lower operational costs and uncertainties surrounding 589 terrestrial C estimates, and consequently, will help developing countries in the development of 590 national forest inventories and aid the scientific community in better understanding the effect 591 592 of climate change on forest ecosystems.

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608 **Contributions**

J.F.Bastin and E.Rutishauser conceptualized the study, gathered the data, performed the
analysis and wrote the manuscript. All the co-authors contributed by sharing data and
reviewing the main text. A.R.Marshall, J.Poulsen and J.Kellner revised the English.

612 **Conflict of interest**

The authors declare there is no conflict of interest associated to this study.

614 Figures

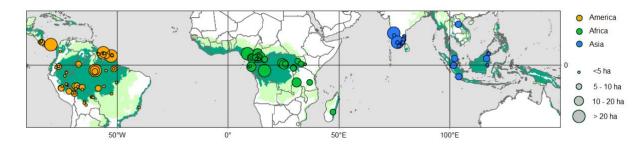
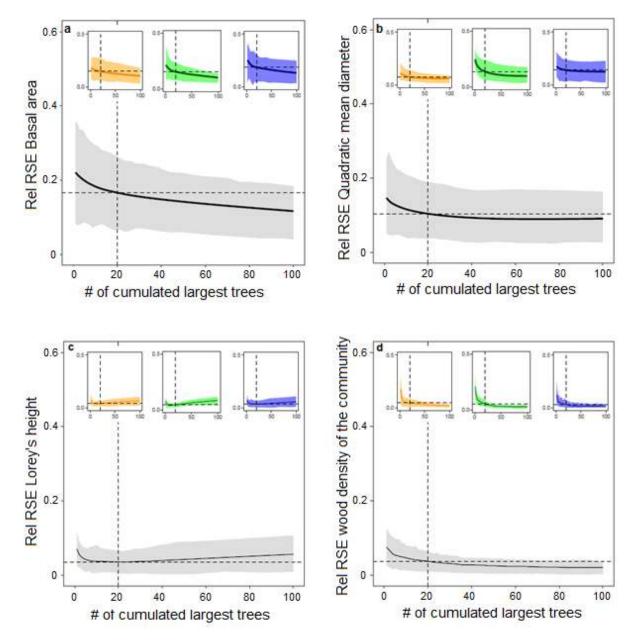


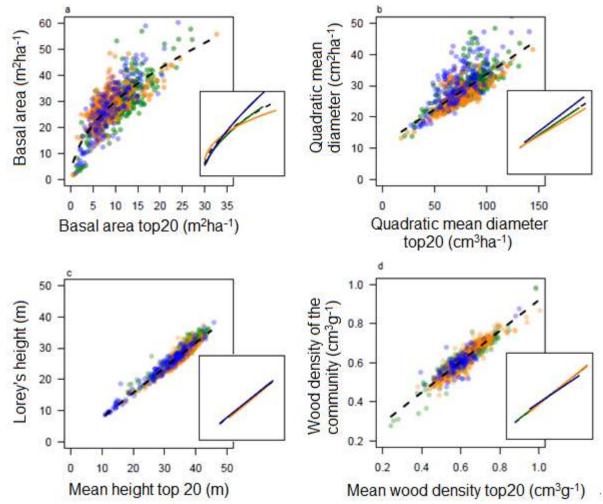
Figure 1. Geographic distribution of the plot database. We used 867 plots of 1 hectare

- from 118 sites. Dots are colored according to floristic affinities (Slik et al. 2015), with America,
- Africa and Asia respectively in orange, green and blue. They are also sized according the total
- 619 area surveyed in each site.



620

Figure 2. Quality of the prediction of plot metrics from largest trees. Variation of the 621 622 relative Root Mean Square Error (rRMSE) of the prediction of plot metric from i largest trees versus the cumulative number of largest trees for (a) basal area, (b) quadratic mean diameter, 623 (c) Lorey's height and (d) wood density weighted by the basal area. Results are displayed at 624 the pan-tropical level (main plot in grey) and at the continental level (subplots; orange = 625 626 America; green = Africa; blue = Asia). The solid line and shading shows the mean rRMSE and the 5th and the 95th percentiles. Dashed lines represent the mean rRMSE observed for each 627 model, when considering the 20 largest trees. 628



629

Figure 3. Prediction of plot metrics (y-axis) from the 20 largest trees (x-axis). Results are 630 shown for (a) basal area, (b) quadratic mean diameter, (c) Lorey's Height and (d) wood density 631 weighted by the basal area. Each dot corresponds to a single plot, colored in orange, green 632 and blue for America, Africa and Asia respectively. Both pan-tropical (black dashed lines) and 633 634 continental (coloured lines) regression models are displayed. These results show that substantial part of remaining variance, i.e. not explained by largest trees, is found when 635 636 predicting the basal area and the quadratic mean diameter, with slight but significant 637 differences between continents.

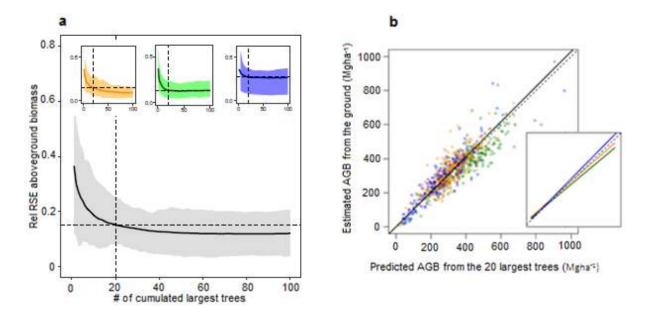




Figure 4. Prediction of AGB from plot metrics of the 20 largest trees. Results are shown for the 867 plots, among the three continents colored orange, green and blue for America, Africa and Asia respectively. The regression line of the model is shown as a continuous black line while the dashed black line shows a 1:1 relationship. The figure shows an unbiased prediction of AGB across the 867 plots, with slight but significant differences between the 3 continents.

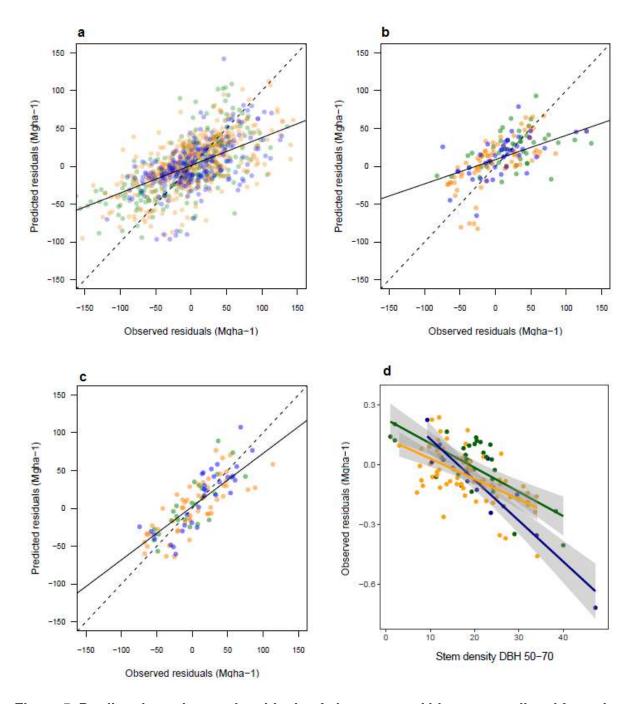
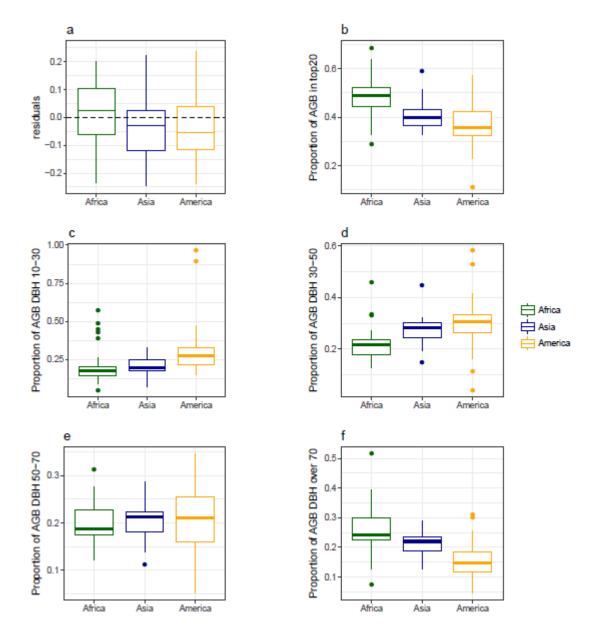


Figure 5. Predicted vs. observed residuals of above ground biomass predicted from the 20 largest trees. Residuals are explored at three different levels: (a) plot, (b) site [without considering the diameter structure as an explanatory variable], (c) site [considering the diameter structure] and (d) along the stem density of medium size trees. America, Africa and Asia are colored in orange, green and blue respectively. The figures show a good prediction of residuals in (a) and (b), driven by stem density, anda less biased prediction in (c), driven by

- the diameter structure. Variance of observed residuals are also well explained by the stem
- density of medium size trees (d), which mainly drive the first axis of the PCA.



654

Figure 6. Comparison across continents of aboveground biomass prediction per site and their 655 contribution to different share of the diameter structure. Africa, Asia and America, are colored 656 657 in green, blue and orange, respectively. The distribution of the residuals of pan-tropical 658 aboveground biomass prediction from the 20 largest trees (a) shows predictions are slightly 659 overestimated in Africa (+2%), and slightly underestimated in Asia (-2%) and America (-6%). 660 The proportion of aboveground biomass in the 20 largest trees (b) is highest in Africa (48%), followed by Asia (40%) and America (35%). The decomposition across four diameter classes 661 (c-f, i.e. from 10 to 30, 30 to 50, 50 to 70 and beyond 70 cm) of their relative share of the total 662 biomass shows that most of the biomass is found in the large trees in Africa, and in the small 663

- to medium trees in America. Asia presenting a more balanced distribution of biomass across
- the diameter structure.

666 Tables

Table 1. Weight of each variable retained for the explanation of AGB residuals. Weights are calculated as a type II sum of squares, which measures the decreased residual sum of squares due to an added variable once all the other variables have been introduced into the model. Results are shown for the exploration of residuals at the plot and at the site level, with and without consideration of the diameter structure. Weights are dominated by structural variables, and in particular the stem density and the diameter structure. Height, wood density and continent have also a non-negligible influence on residuals.

| Level of residual | Parameter | Weight | 674 |
|---------------------|---|----------|-----|
| Plot | | | |
| | Stem density* | 79 | |
| | Continent* | 18 | |
| | Lorey's height* | 1 | |
| | Major soil types | 1 | |
| | Temperature | 1 | |
| | Wood density weighted by the basal area | 0 | |
| | Rainfall | 0 | |
| Site without | | | |
| diametric structure | | - / | |
| | Stem density* | 54 | |
| | Continent* | 28 | |
| | Wood density weighted by the basal area* | 11 | |
| | Rainfall | 3 | |
| | Major soil types | 3 | |
| | Temperature | 2 | |
| | Lorey's height | 0 | |
| | Lorey Sheight | 0 | |
| Site with diametric | | | |
| structure | | | |
| | PCA axis 1* | 69 00 | |
| | Lorey's height* | 23 | |
| | Rainfall* | 3 | |
| | Major soil types | 3 | |
| | Continent | 1 | |
| | Temperature | 1 | |
| | Wood density weighted by the basal area | 0 | |
| | PCA axis 2 | 0 | |

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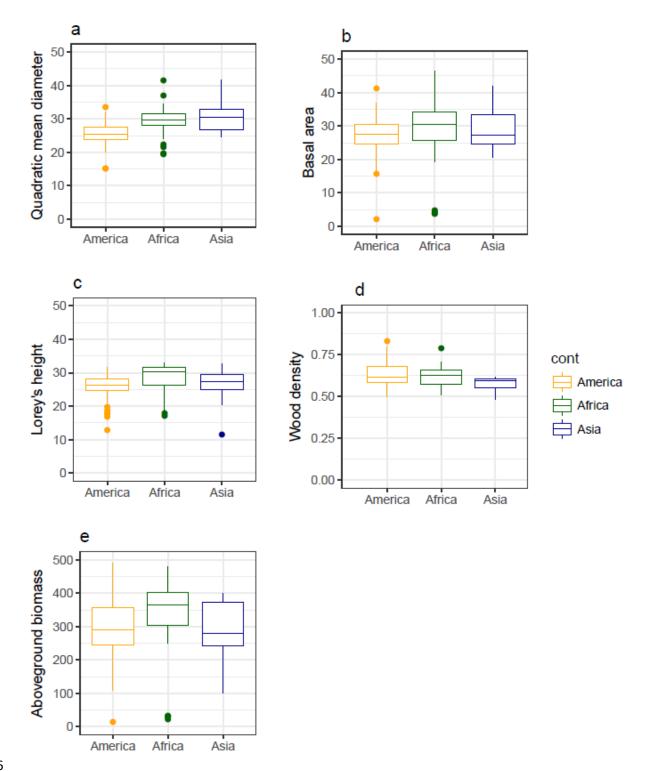
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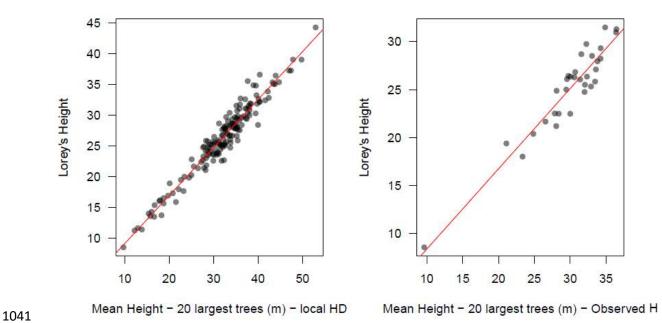
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- 1033 Supplementary table 1. Plot, Site and Pls
- 1034 Supplementary table 2. Coefficients of plot level structure prediction from the *ith*
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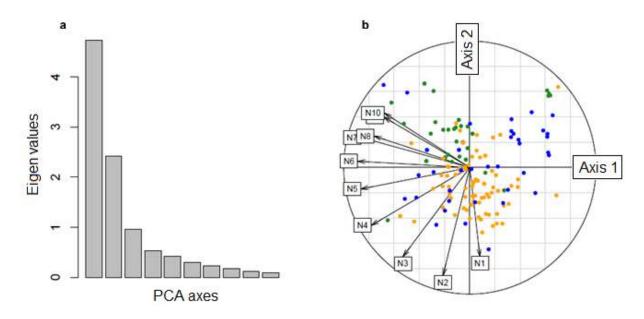


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Supplementary figure 1. Cross-continent comparison of plot-metrics distribution averaged at the site level. Figures illustrates respectively the distribution of the values for 1038 1039 the quadratic mean diameter (a), basal area (b), Lorey's height (c), wood density (d) and 1040 aboveground biomass (e).

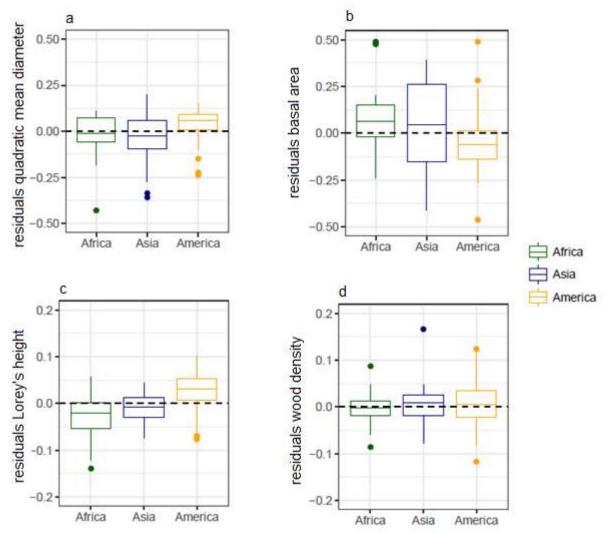


Supplementary figure 2. Lorey's Height prediction from the 20 largest trees. Figures
show the results using (i) local D-H allometries for 20 sites (left subfigure) and (ii) using plots
where height is measured on all trees in Malebo site in the Democratic Republic of the Congo
(right subfigure).



Supplementary figure 3. PCA on the diameter structure and corresponding mean 1047 1048 distribution for high contributions of axis 1 and axis 2. (A) Illustration of top and low 1049 percentile observed for each axis, with diameter distributions represented as the relative 1050 difference with the average observed distribution.(B) Biplot with contribution to the PCA of all 1051 the diameter classes, with the respective position of each site in the space defined by axis1 1052 and 2. Axis 1 is driven by differences in global abundance of trees and axis 2 is driven by a 1053 difference of balance between abundance of small vs. large trees. Colors represent continent, with Africa, America and Asia respectively in green, orange and blue. 1054

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Supplementary figure 4. Cross-continent comparison of the relative residuals from the prediction of plot-metrics from the 20 largest trees. The relative residuals are generally low (<10%). Systematic small differences can however be found in America, where the quadratic mean diameter and Lorey's height tend to be slightly overestimated and the basal area slightly underestimated.</p>