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1 **Title**

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

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195 **Abstract**

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

202 **Location.** Pan-tropical

203 **Method.** Using a dataset of 867 plots distributed among 118 sites across the tropics, we tested
204 the ability to predict quadratic mean diameter, basal area, Lorey's height and community wood
205 density from the i^{th} largest trees, i.e. testing the cumulative information gathered from these i
206 trees ranked by decreasing diameter. These tests served as a basis to select the optimal
207 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%
212 of relative error. Building on this result, we developed a new model to predict plot-level AGB
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-
216 70 cm), which shows some continental dependency with American tropical forests presenting
217 the highest levels of total biomass share in these intermediate diameter classes.

218 **Conclusion.** Our approach provide new information on tropical forest structure and can be
219 employed to generate accurately field estimates of tropical forest carbon stocks to support the
220 calibration and validation of current and forthcoming space missions. It will reduce the cost of
221 programs to monitor, report, and verify forest resources, and will contribute to scientific
222 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 & Löhmus, 2011; Lindenmayer et al., 2012),
226 regulate forest dynamics, regeneration (Harms et al., 2000; Rutishauser et al., 2010) and total
227 biomass (Stegen et al., 2011), and are important contributor to the global carbon cycle
228 (Meakem et al., 2017). Being major components of the canopy, the largest trees also suffer
229 more than sub-canopy and understory trees from climate change, as they are directly exposed
230 to variations in solar radiation, wind strength, temperature seasonality and relative air humidity
231 (Laurance et al., 2000; Nepstad et al., 2007; Lindenmayer et al., 2012; Thomas et al., 2013;
232 Bennett et al., 2015; Meakem et al., 2017). Because they are visible from the sky, large trees
233 are ideal for monitoring forest responses to climate change via remote sensing (Bennett et al.,
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
239 forest aboveground biomass on average (Bastin et al., 2015). Consequently, the density of
240 large trees largely explains variation in AGB at local (Clark & Clark, 1996), regional (Malhi et
241 al., 2006; Saatchi et al., 2007), and continental scales (Stegen et al., 2011; Slik et al., 2013).

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
244 measurement of the 20 largest trees per hectare being sufficient to estimate plot-level biomass
245 with less than 15% errors in reference to ground estimates. These findings opened the
246 possibility of measuring the largest trees to cost-effectively monitor forest biomass in Central
247 Africa, rather than conducting full inventories of all size classes. Similarly, they suggested that
248 remote sensing (RS) approaches should focus on the measurement of the largest trees,
249 instead of properties of the entire forest.

250 Several efforts are underway to close the gap between remote sensing and field surveys (e.g.
251 Jucker et al. 2016a, Coomes et al. 2017). However, field inventories still rely on exhaustive
252 data collection, while remote sensing surveys provide a limited alternative for the following
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
255 properties (Barbier et al., 2011; Asner et al., 2012; Barbier & Coutron, 2015), or require
256 ground measurement of all trees above or equal to 10 cm of D for calibration (Asner et al.,
257 2012; Asner & Mascaro, 2014). Using mean canopy height extracted from active sensors
258 (Mascaro et al., 2011; Ho Tong Minh et al., 2016), or canopy grain derived from optical images
259 (Proisy et al., 2007; Ploton et al., 2012, 2017; Bastin et al., 2014), the biomass is predicted
260 from remote sensing with a typical error of only 10-20% compared to ground-based estimates,
261 but is limited to the extent of the scene used. An interesting development to alleviate this spatial
262 restriction lies in the ‘universal approach’, proposed by Asner et al. (2012) and further adapted
263 in Asner and Mascaro (2014), in which plot-level biomass is predicted by a linear combination
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,
267 WD_{BA}) derived from full field inventories. AGB is then predicted as follows (Asner et al., 2012):
268 $AGB = aTCH^{b1}BA^{b2}WD_{BA}^{b3}(1)$

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

273 Recent advances in remote sensing allow the identification of single trees in the canopy (Ferraz
274 et al., 2016), estimation of adult mortality rates for canopy tree species (Kellner & Hubbell,
275 2017), description of the forest diameter structure (Stark et al., 2015), depiction of crown and
276 gap shapes (Coomes et al., 2017), and even identification of some functional traits of canopy
277 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 $D \geq 10$ cm in a plot). To this end, we tested the following
280 model:

$$281 \text{ AGB} = a(D_{\text{gLT}i}H_{\text{LT}i}W_{\text{LT}i})^{b1} \quad (2)$$

282 Where for the i^{th} largest trees, D_{gLT} is the quadratic mean diameter, H_{LT} the mean height, and
283 W_{LT} the mean wood density averaged among the i^{th} largest trees.

284 Using a large database of forest inventories gathered across the Tropics (Figure 1), including
285 secondary and old growth forest plots, we test the ability of the largest trees to provide
286 information on various metrics estimated at 1-ha plot level, such as the mean quadratic
287 diameter, the basal area (BA), the Lorey's height (i.e. plot-average height weighted by BA), the
288 community wood density (i.e. plot-average wood density weighted by BA) and mean above-
289 ground live biomass (AGB) (supplementary figure 1). While previous work focused on
290 estimating biomass in Central African forests (Bastin et al., 2015), the present study aims at
291 generalizing the potential of large trees in predicting these different plot metrics at continental
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
295 approach can be used to guide innovative RS techniques and increase the frequency and
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).
299 This study is a step forward in bringing together remote sensing and field sampling techniques
300 for quantification of terrestrial C stocks in tropical forests.

301 **Material & Methods**

302 **Database**

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

311 We selected plots located between 23°N and 23°S, including tropical islands, with an area of
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
314 diameter at 130 cm greater than or equal to 10 cm had been measured were included. Wood
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
318 Database (GWDD, Chave et al., 2009; Zanne et al., 2009). Botanical identification was
319 harmonized through the Taxonomic Names Resolution Service
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
322 heights of all trees using Chave et al.'s (2014) pan-tropical diameter-height model which
323 accounts for heterogeneity in the D-H relationship using an environmental proxy:

$$324 \ln(H) = 0.893 - E + 0.760 \ln(D) - 0.0340 \ln(D)^2 \quad (3)$$

325 Where D is the diameter at 130cm and E is a measure of environmental stress (Chave et al.,
326 2014). For sites with tree height measurements (N=20), we developed local D-H models, using
327 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).
 330 We estimated plot biomass as the sum of the biomass of live tree with diameter at 130 cm
 331 superior or equal to 10 cm, using the following pan-tropical allometric model (Réjou-Méchain
 332 et al., 2017):

$$333 \text{ AGB} = \exp(-2.024 - 0.896E + 0.920 \ln(\text{WD}) + 2.795 \ln(D) - 0.0461(\ln(D^2))) \quad (4)$$

334 **Plot-level metric estimation from the largest trees**

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

$$348 \text{ rRMSE} = \bar{X} \sum \sqrt{\frac{(X_{\text{obs}} - X_{\text{pred}})^2}{n}} \quad (5)$$

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

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

$$354 \text{ BA} = b_1 \sum BA_i^{Y_1} \quad (6)$$

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

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

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

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

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

366 The quality of the predictions of plot-level metrics from the largest trees is quantified using the
367 relative root mean square error (rRMSE) between measured and predicted values, and
368 displayed along the cumulated number of largest trees (Figure 2). Model coefficients are
369 estimated for each metric derived from the largest trees (N_{LT}) and averaged across the 118
370 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**

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

380 **Investigating residuals: what the largest trees do not explain**

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

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
395 variables have been introduced into the model (Langsrud, 2003). Residuals were investigated
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
398 at small scales (Clark & Clark, 2000). Here we use a principal component analysis (PCA) to
399 summarize the information held in the diameter structure by ordinating the sites along the
400 abundance of trees in each diameter class (from 10 to +100 cm by 10 cm bins).

401

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
406 $\text{cm}^2\text{ha}^{-1}$, the basal area from 2 to 58 m^2ha^{-1} , Lorey's height from 11 to 33 m and the wood
407 density weighted by the basal area from 0.48 to 0.84 gcm^{-3} (Supplementary figure 1). Such
408 important differences between minimal and maximal values are observed because our
409 database cover sites with various forest types, from young forest colonizing savannas to old
410 growth forest. However, most of our sites are found in mature forests, as shown by relatively
411 high average and median value of each plot metric (average aboveground biomass = 302
412 Mgha^{-1} ; supplementary figure 1). In general, highest values of aboveground biomass are found
413 in Africa, driven by highest values of basal area and highest estimations of Lorey's height.
414 Highest values of wood density weighted by basal area are found in America.

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

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

418

419 When using the 20 largest trees to predict basal area (BA) and quadratic mean diameter (Dg),
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
422 mean rRMSE of 4% for the 20 largest trees. The prediction of Lorey's height from the largest
423 trees using local diameter-height model (supplementary Figure 2a) yielded results similar to
424 those obtained using equation 3 of Chave et al. (2014). More importantly, it also yielded similar
425 results to prediction of Lorey's height from the largest trees using plots where all the trees were
426 measured on the ground (supplementary figure 2b). This suggests that our conclusions are
427 robust to the uncertainty introduced by height-diameter allometric models.

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:

$$431 \text{ AGB} = 0.0735 \times (\text{Dg}_{20}\text{H}_{20}\text{WD}_{20})^{1.1332} \text{ (rRMSE=0.179; R}^2\text{=0.85; AIC= -260.18) (10)}$$

432 Because the exponent was close to 1, we also developed an alternative and more operationa
433 l model with the exponent constrained to 1, given by:

$$434 \text{ AGB} = 0.195 \times (\text{Dg}_{20}\text{H}_{20}\text{WD}_{20}) \text{ (rRMSE=0.177; R}^2\text{=0.85; AIC=-195) (11)}$$

435 Ground measurements of plot AGB were predicted by our N_{LT} model with the exponent
436 constrained to 1, with a total error of 17.9% (Figure 4), a value which encompass the error of
437 the N_{LT} model and the error related to the allometric model chosen. The Leave-One-Out cross-
438 validation procedure yielded similar results (rRMSE=0.19; R²=0.81), validating the use of the
439 model on independent sites.

440 **Determining the cause of residual variations**

441 The explanatory variables all together explain about 37% of the variance in AGB both at plot
442 and site levels when omitting the diameter structure, and about 63% at site level when included
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
446 America and Asia) and by the effect of H_{BA} and WD_{BA} (respective weights: 1% and 0% at the
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
450 (supplementary figure 3). This first axis of the PCA was related to the general abundance of
451 trees at a site, and in particular medium-sized trees (40-60cm). Among environmental
452 variables, only rainfall was significantly related to the residuals at the site level when the
453 diameter structure was considered (2%).

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
458 predicting basal area, quadratic mean diameter, Lorey's height (H_{BA}) or community wood
459 density beyond the first 10-to-20 largest trees (Figure 2, Figure 3a). In some cases, e.g. when
460 a forest plot presents an abundant number of large trees (Figure 5d), increasing the number
461 of trees sampled improves the model's accuracy. This is the case for BA for which rRMSE
462 continues to decrease up to 100 largest trees (Figure 2a). In contrast, Lorey's height
463 predictions are altered when a large number of trees are included (Figure 2c), i.e. when
464 smaller, often suppressed, trees draw the average down (Farrion et al., 2016). This might
465 explain why the prediction of AGB does not mirror that of basal area (Figure 2b, Figure 3a),
466 and suggest that the number of largest trees shall be set independently to each predictor
467 considered. Interestingly, the evolution of relative error in AGB prediction as a function of the
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
471 structure (Figure 5c, supplementary Figure 3b), and in particular the number of medium size
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.

475 The good performance of models based on the 20 largest trees in predicting Lorey's height
476 and community wood density at site level was not surprising. Both metrics were indeed
477 weighted by basal area, driven de facto by the largest trees. Their consistency across sites
478 and continents was not expected though. This suggests that the relationship between the 20
479 largest trees and descriptors of forest structures is stable across the tropics, and prove the
480 generality of our approach. Slight differences are however noticeable when comparing the
481 distribution of the pan-tropical model residuals across continents (Figure 6, supplementary

482 figure 4). In America, our pan-tropical model tend to slightly underestimate basal area (mean:
483 -5%) and overestimate Lorey's height (mean: +3%) (supplementary figure 4), suggesting
484 peculiar forest structures (i.e. higher tree height for a given diameter, and lower fractions of
485 large trees, supplementary figure 2). In Asia, and in particular in Africa, large (i.e. DBH > 50
486 cm) trees are more abundant and encompass a large fraction of plot biomass. The basal area
487 tends to be slightly overestimated in Africa, resulting in average to a 3% overestimation of AGB
488 (Figure 6a).

489 Interestingly, while a recent global phylogenetic classification of tropical forest groups
490 American with African forests vs. Asian forests (Slik et al., 2018), our results tend more to
491 single out American forests. Although this deserves further investigations, it might reveal a lack
492 of close relationship between forest structure properties and phylogenic similarity, which
493 echoes recent results on the absence of relationship between tropical forest diversity and
494 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
499 properties (i.e. the quadratic mean diameter, the basal area, Lorey's height and community
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
503 vary locally (Feldpausch et al., 2011; Kearsley et al., 2013;). It is also (iii) relatively insensitive
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**

515 Globally, the N_{LT} model for the 20 largest trees allows plot biomass to be predicted with 17.9%
516 error. This result is a pan-tropical validation of results obtained in Central Africa (Bastin et al.,
517 2015). It opens new perspectives towards cost-effective methods to monitor forest structures
518 and carbon stocks through largest trees metrics, i.e. metrics of objects directly intercepted by
519 remote-sensing products.

520 Developing countries willing to implement a Reduction of Emissions from Deforestation and
521 Forest Degradation (REDD+), shall also report on their carbon emissions (CE) and develop a
522 national CE reference level (IPCC, 2006; Maniatis & Mollicone, 2010). However, most tropical
523 countries lack capacities to assume multiple, exhaustive and costly forest carbon assessment
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
527 plot, monitoring only 20 trees is a significant improvement. Although finding the 20 largest trees
528 in a plot of several hundred individuals requires evaluating more than 20 trees, in practice, a
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
531 approach adapted to detection of the largest trees in tropical forests. Using such approach
532 would facilitate rapid field sampling in extensive areas to produce large scale AGB estimates.
533 Those could fulfil the needs in calibration and validation of current and forthcoming space
534 missions focused on aboveground biomass.

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

538 trees instead of indirect development of complex metrics (e.g. mean canopy height, texture)
539 used to extrapolate forest properties. While some further refinements are needed, most of the
540 tools required to develop “largest trees” models are readily available. In particular, Ferraz et
541 al. (2016) developed an automated procedure to locate canopy trees based on airborne LiDAR
542 data, to measure their height and crown area. Crown area could further be linked to basal area,
543 as the logarithm of crown area is consistently correlated with a slope of 1.2-1.3 to the logarithm
544 of tree diameter across the tropics (Blanchard et al., 2016). Regarding wood density,
545 hyperspectral signature offers a promising way to assess functional traits remotely (e.g. Asner
546 et al., 2017) which could potentially be tested to detect variability in wood properties.
547 Alternative approaches could focus on the development of plot-level AGB prediction by
548 replacing the basal area of the largest trees with their crown metrics. While the measurement
549 of crown areas have yet to be generalized when inventorying plots, several biomass allometric
550 models already partition trunk and crown mass (Jucker et al., 2016; Ploton et al., 2016;
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
554 particular among the total stem density. Interestingly, stem density was generally identified as
555 a poor predictor of plot biomass in tropical forests (Slik et al., 2010; Lewis et al., 2013).
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
559 put more effort into understanding variation in smaller trees, which mainly drives total stem
560 density and the total floristic diversity. Smaller trees are also essential to characterize forest
561 dynamics and understand changes in carbon stocks. Several options are nonetheless possible
562 from remote sensing, considering the variation in lidar point density below the canopy layer
563 (D’Oliveira et al., 2012), the distribution of leaf area density (Stark et al., 2012, 2015; Tang &
564 Dubayah, 2017) or the use of multitemporal lidar data to get information on forest gap

565 generation dynamics and consequently on forest diameter structure (Kellner et al., 2009;
566 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
569 managed or degraded forests. Indeed, large trees are targeted by selective or illegal logging,
570 and are the first to disappear or to suffer from incidental damages when tropical forests are
571 exploited for timber (Sist et al., 2014). The loss of largest trees drastically changes forest
572 structures and diameter distributions, and their loss is likely to counteract the consistency in
573 forest structures observed through this study. Understanding how, or whether, managed
574 forests deviate from our model predictions could help characterize forest degradation, which
575 accounts for a large fraction of carbon loss worldwide (Baccini et al., 2017), acknowledging
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
582 number of field plots and develop new ways to use remote sensing data. Our results provide
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.
586 As highlighted by Clark and Kellner (2012), new biomass allometric models relating plot-level
587 biomass measured from destructive sampling and plot-level metric measured from remote-
588 sensing products should be developed, as an alternative to current tree-level allometric
589 models. Such an effort will lead largely to lower operational costs and uncertainties surrounding
590 terrestrial C estimates, and consequently, will help developing countries in the development of
591 national forest inventories and aid the scientific community in better understanding the effect
592 of climate change on forest ecosystems.

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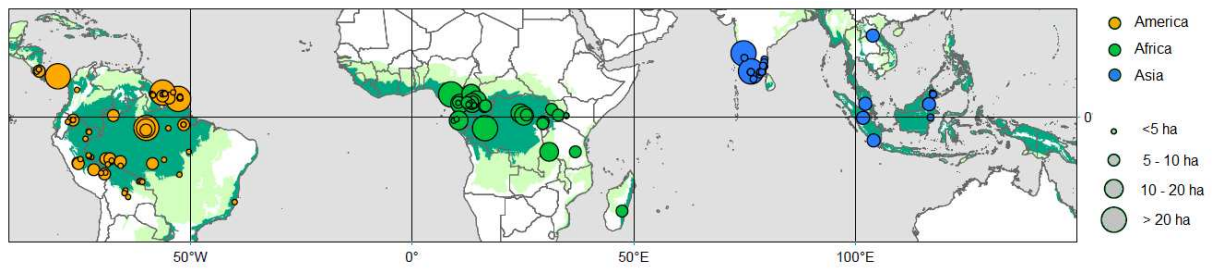
608 **Contributions**

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

612 **Conflict of interest**

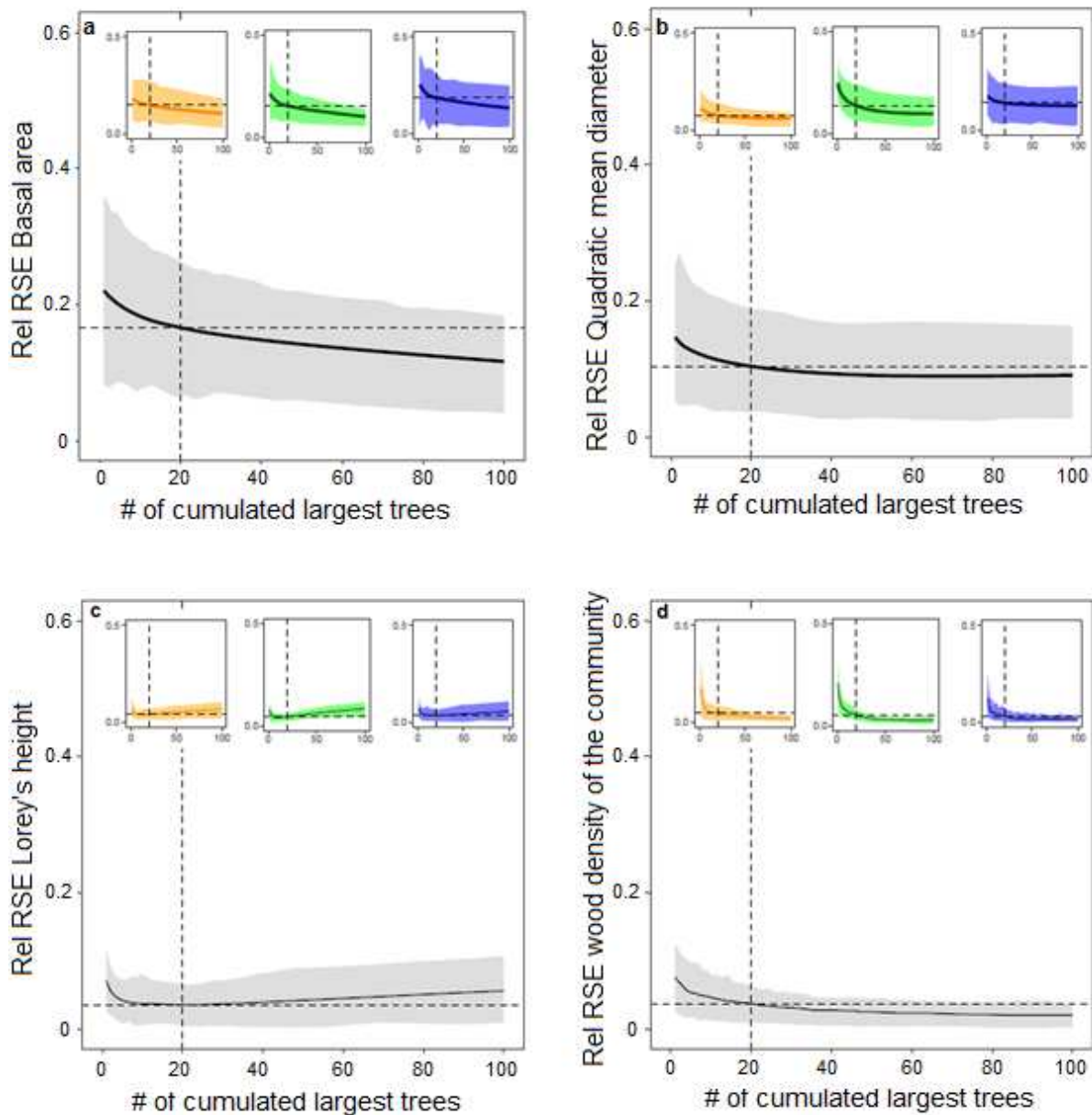
613 The authors declare there is no conflict of interest associated to this study.

614 **Figures**



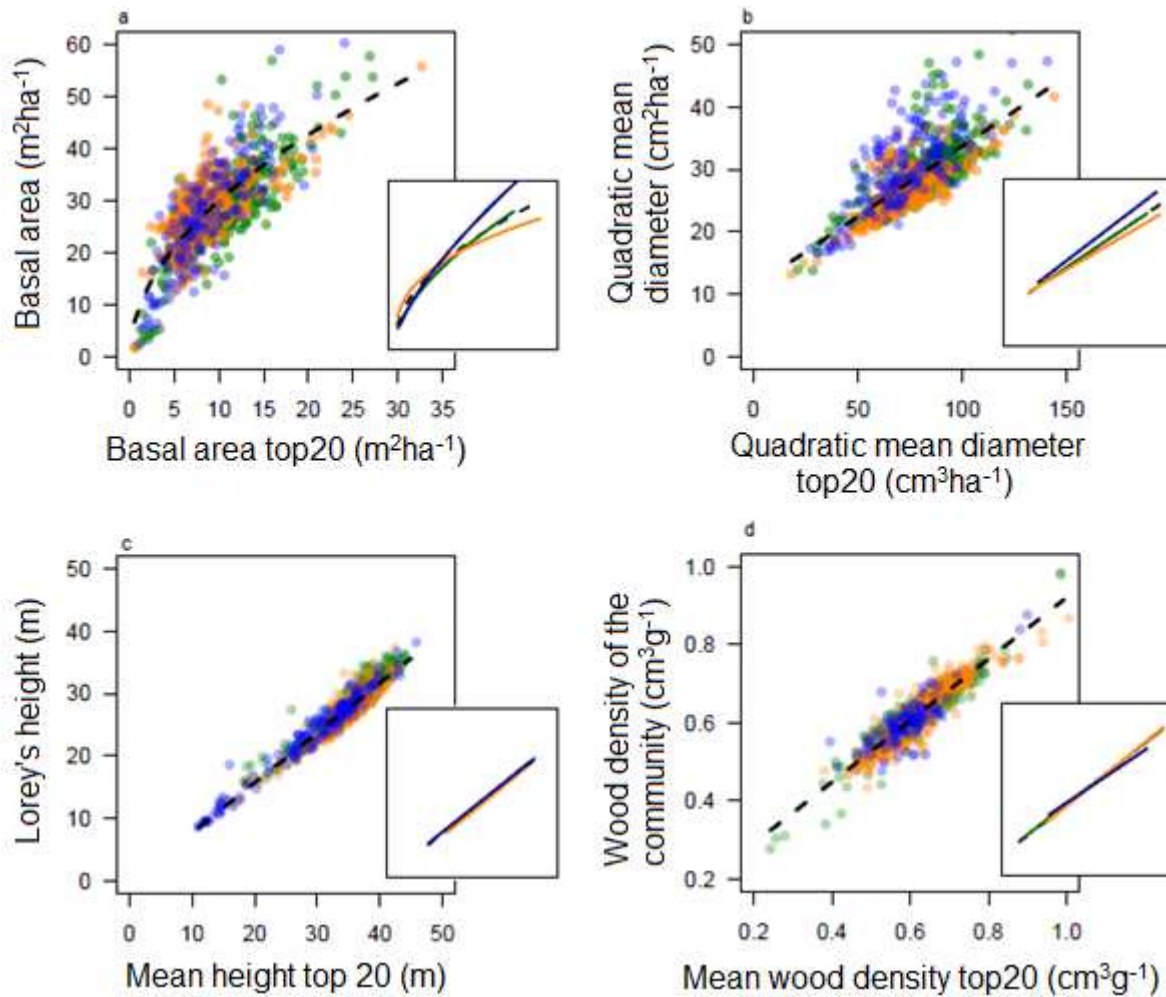
615

616 **Figure 1. Geographic distribution of the plot database.** We used 867 plots of 1 hectare
617 from 118 sites. Dots are colored according to floristic affinities (Slik et al. 2015), with America,
618 Africa and Asia respectively in orange, green and blue. They are also sized according the total
619 area surveyed in each site.



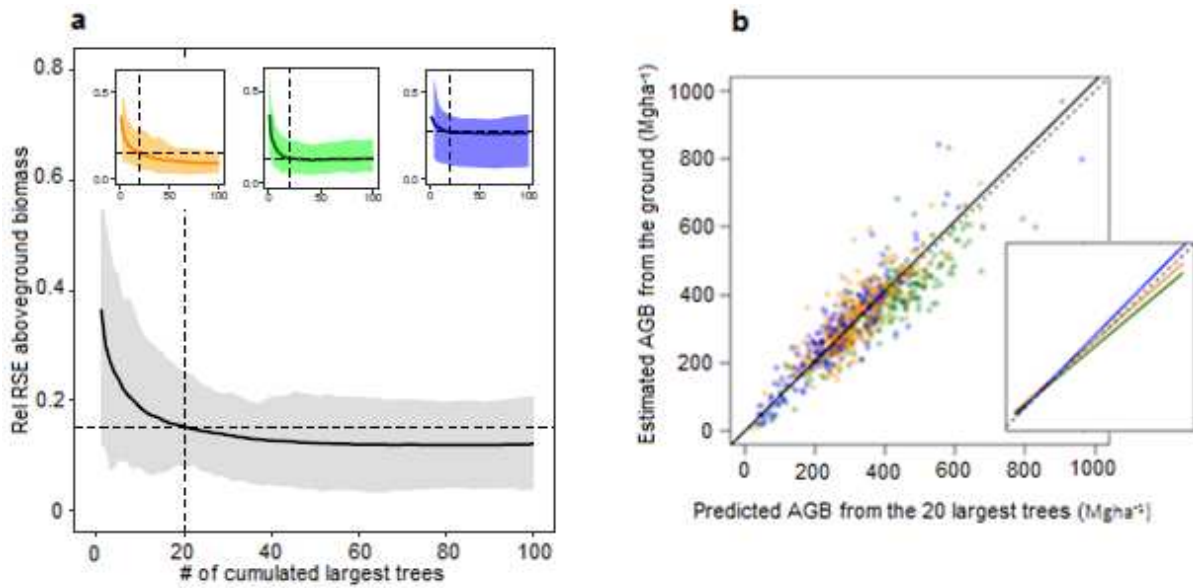
620

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



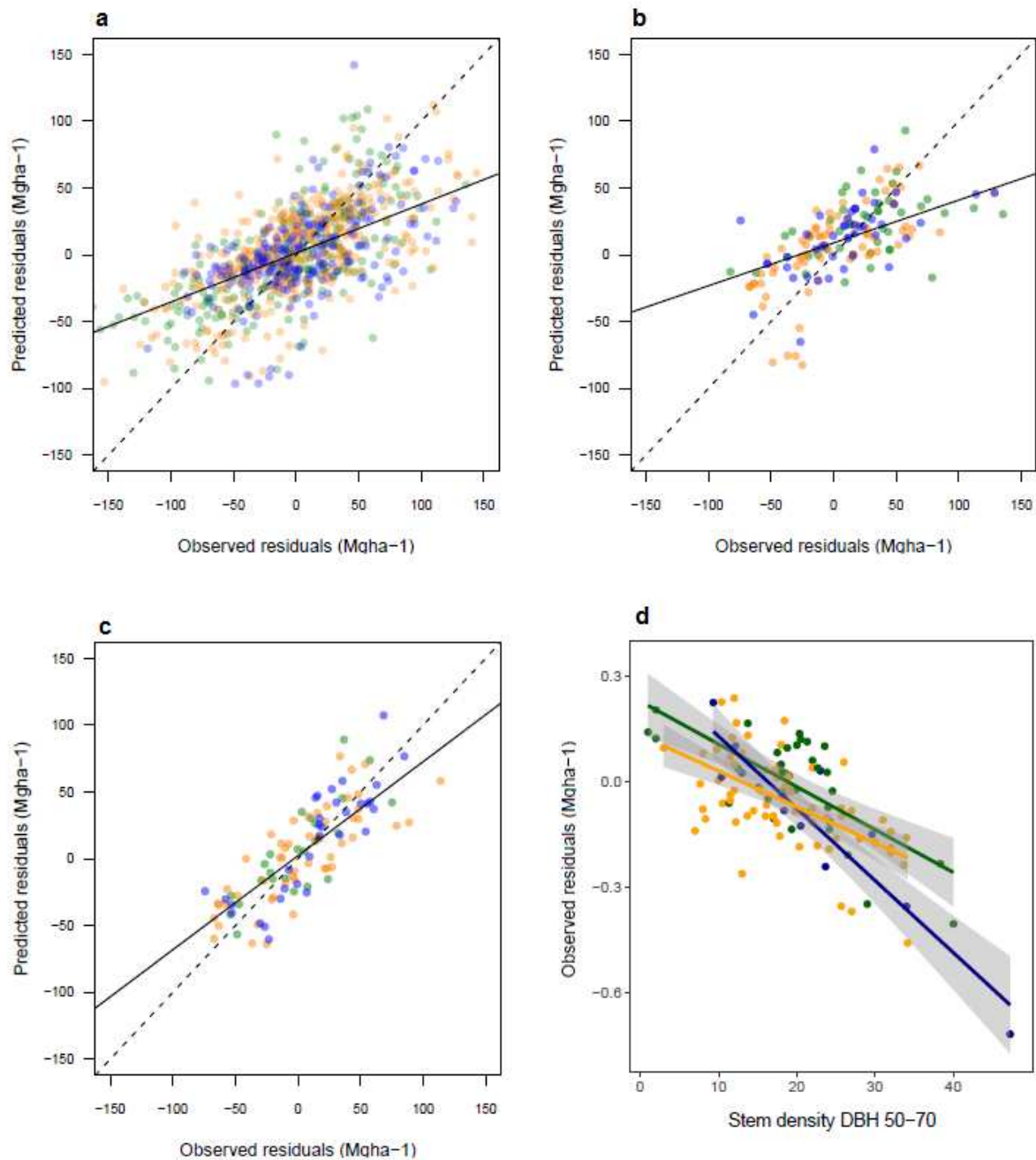
629

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



638

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



645

646 **Figure 5. Predicted vs. observed residuals of above ground biomass predicted from the**

647 **20 largest trees.** Residuals are explored at three different levels: (a) plot, (b) site [without

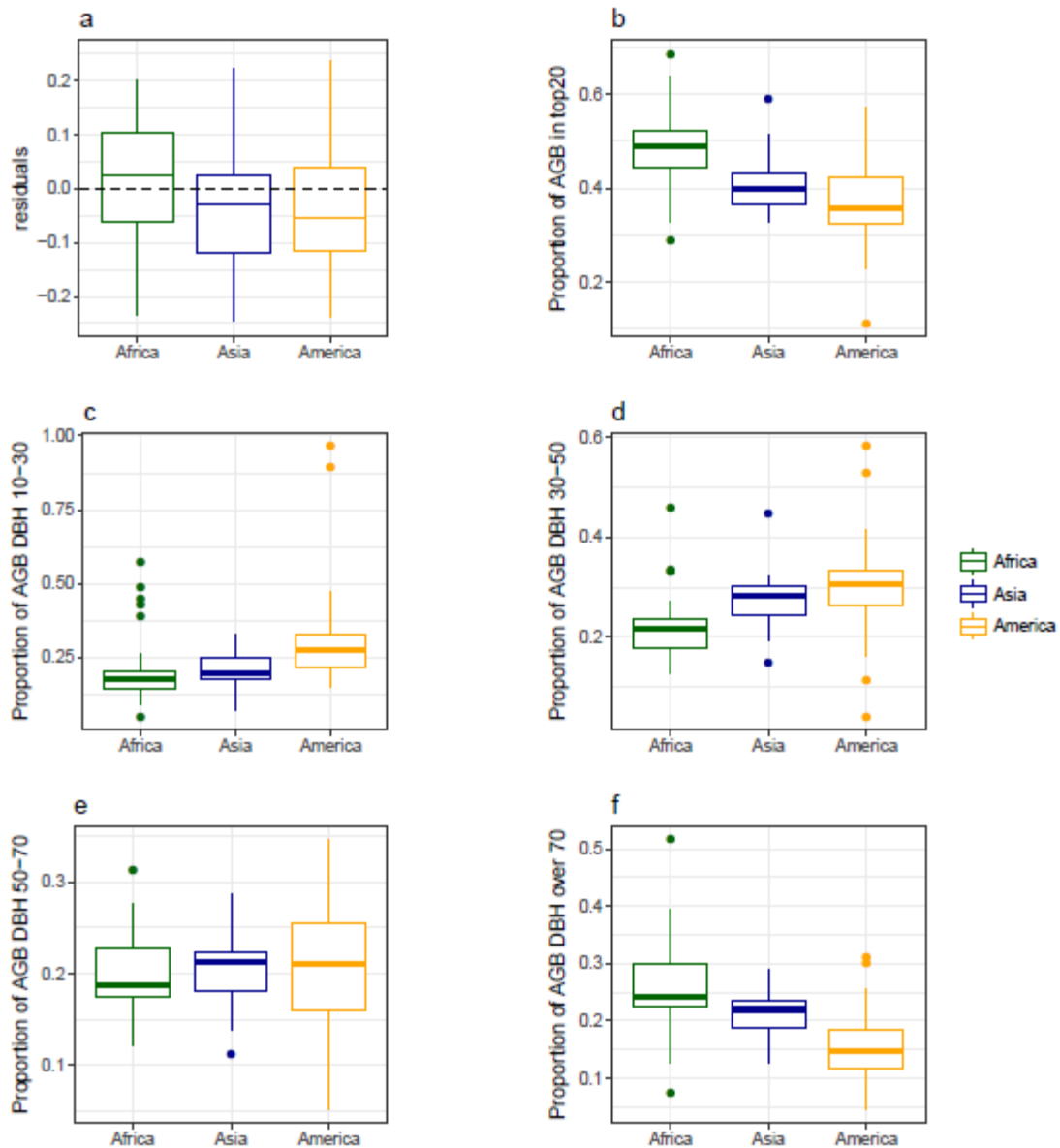
648 considering the diameter structure as an explanatory variable], (c) site [considering the

649 diameter structure] and (d) along the stem density of medium size trees. America, Africa and

650 Asia are colored in orange, green and blue respectively. The figures show a good prediction

651 of residuals in (a) and (b), driven by stem density, and a less biased prediction in (c), driven by

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



654

655 Figure 6. Comparison across continents of aboveground biomass prediction per site and their
 656 contribution to different share of the diameter structure. Africa, Asia and America, are colored
 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%),
 661 followed by Asia (40%) and America (35%). The decomposition across four diameter classes
 662 (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
 663 biomass shows that most of the biomass is found in the large trees in Africa, and in the small

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

666 **Tables**

667 **Table 1. Weight of each variable retained for the explanation of AGB residuals.** Weights
 668 are calculated as a type II sum of squares, which measures the decreased residual sum of
 669 squares due to an added variable once all the other variables have been introduced into the
 670 model. Results are shown for the exploration of residuals at the plot and at the site level, with
 671 and without consideration of the diameter structure. Weights are dominated by structural
 672 variables, and in particular the stem density and the diameter structure. Height, wood density
 673 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	
Site with diametric structure	PCA axis 1*	69	
	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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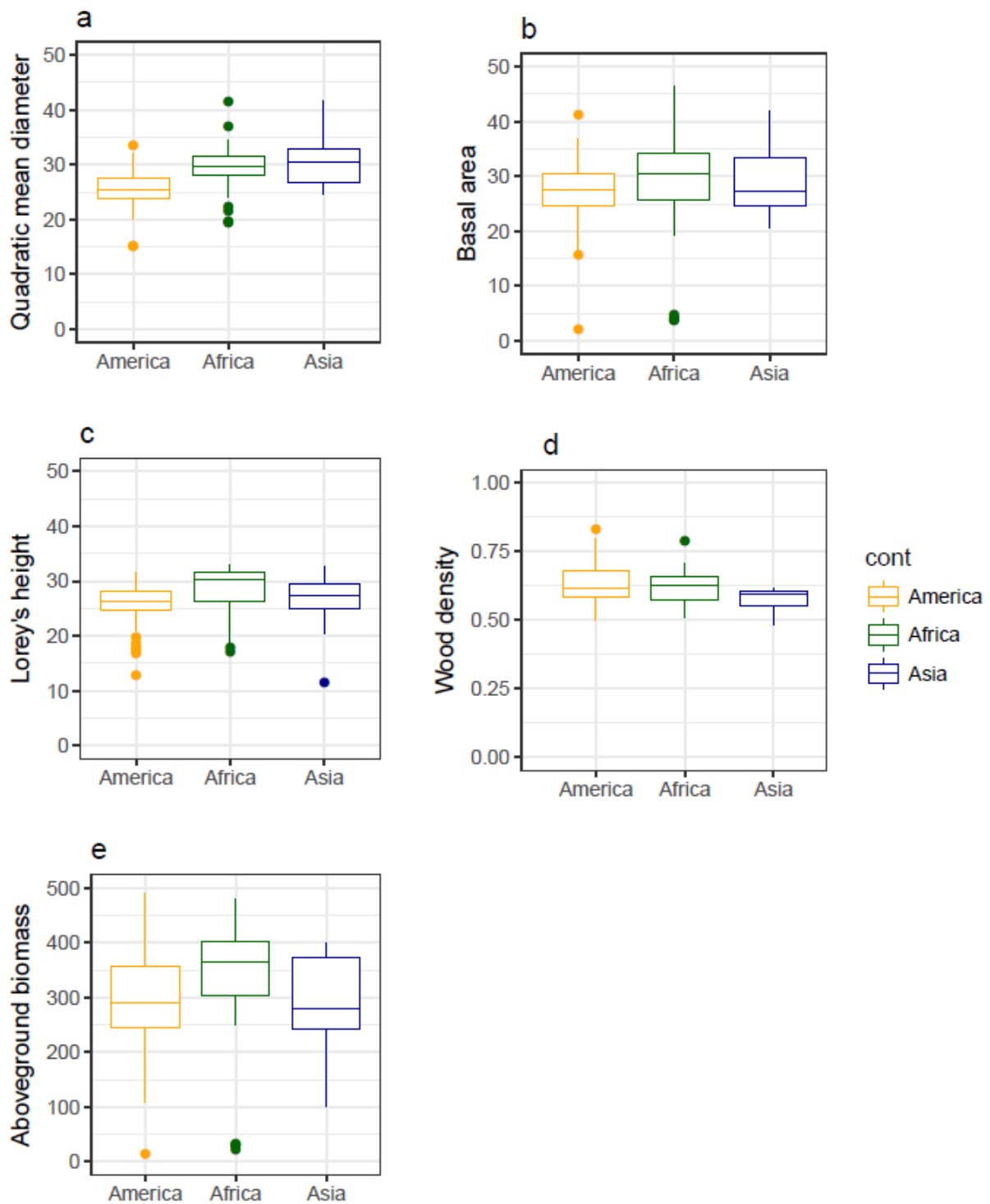
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1032 **Supplementary information.**

1033 **Supplementary table 1. Plot, Site and Pls**

1034 **Supplementary table 2. Coefficients of plot level structure prediction from the ith**

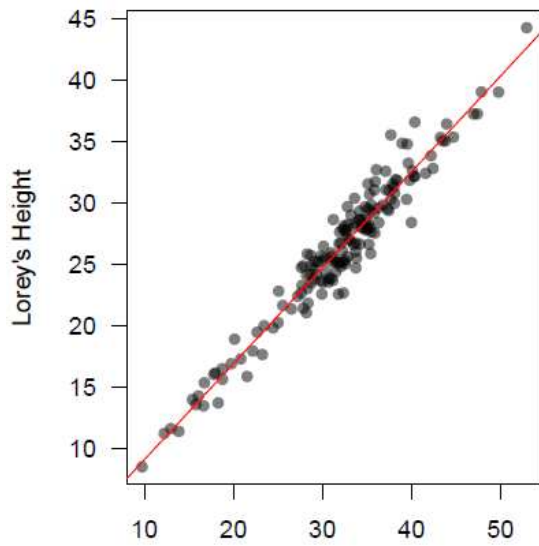
1035 **largest trees.**



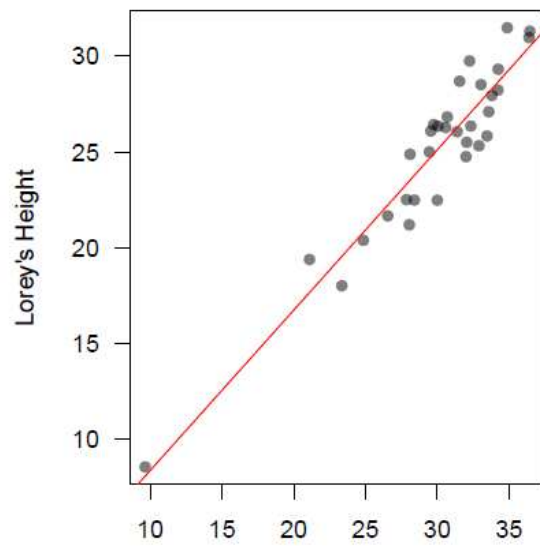
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Supplementary figure 1. Cross-continent comparison of plot-metrics distribution

1038 **averaged at the site level.** Figures illustrates respectively the distribution of the values for
1039 the quadratic mean diameter (a), basal area (b), Lorey's height (c), wood density (d) and
1040 aboveground biomass (e).



Mean Height - 20 largest trees (m) - local HD



Mean Height - 20 largest trees (m) - Observed H

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Supplementary figure 2. Lorey's Height prediction from the 20 largest trees. Figures

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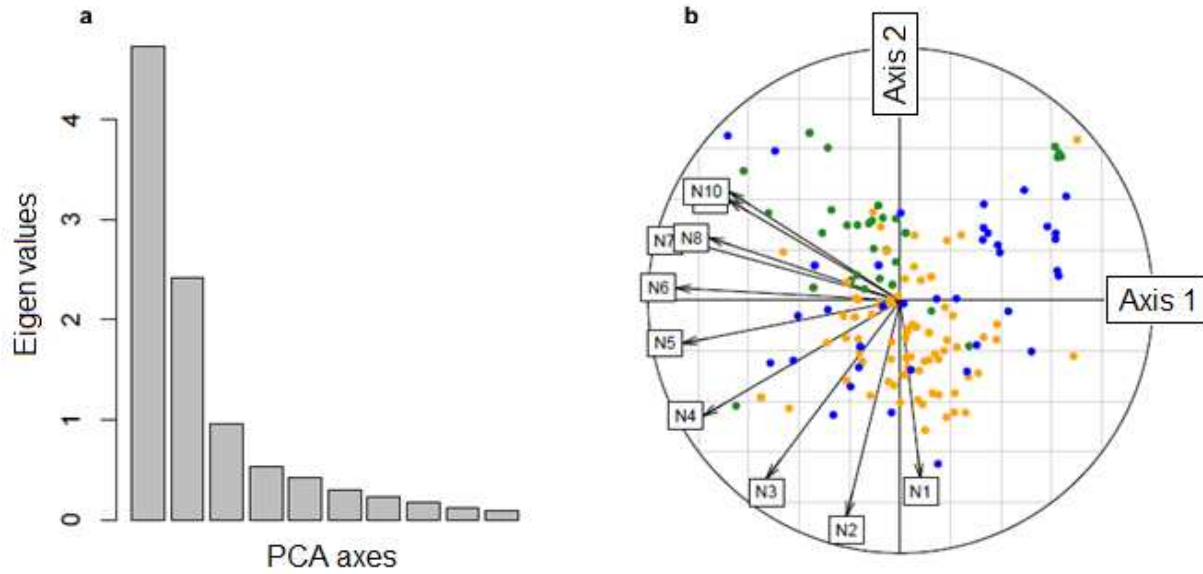
show the results using (i) local D-H allometries for 20 sites (left subfigure) and (ii) using plots

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where height is measured on all trees in Malebo site in the Democratic Republic of the Congo

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(right subfigure).



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1047 **Supplementary figure 3. PCA on the diameter structure and corresponding mean**

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

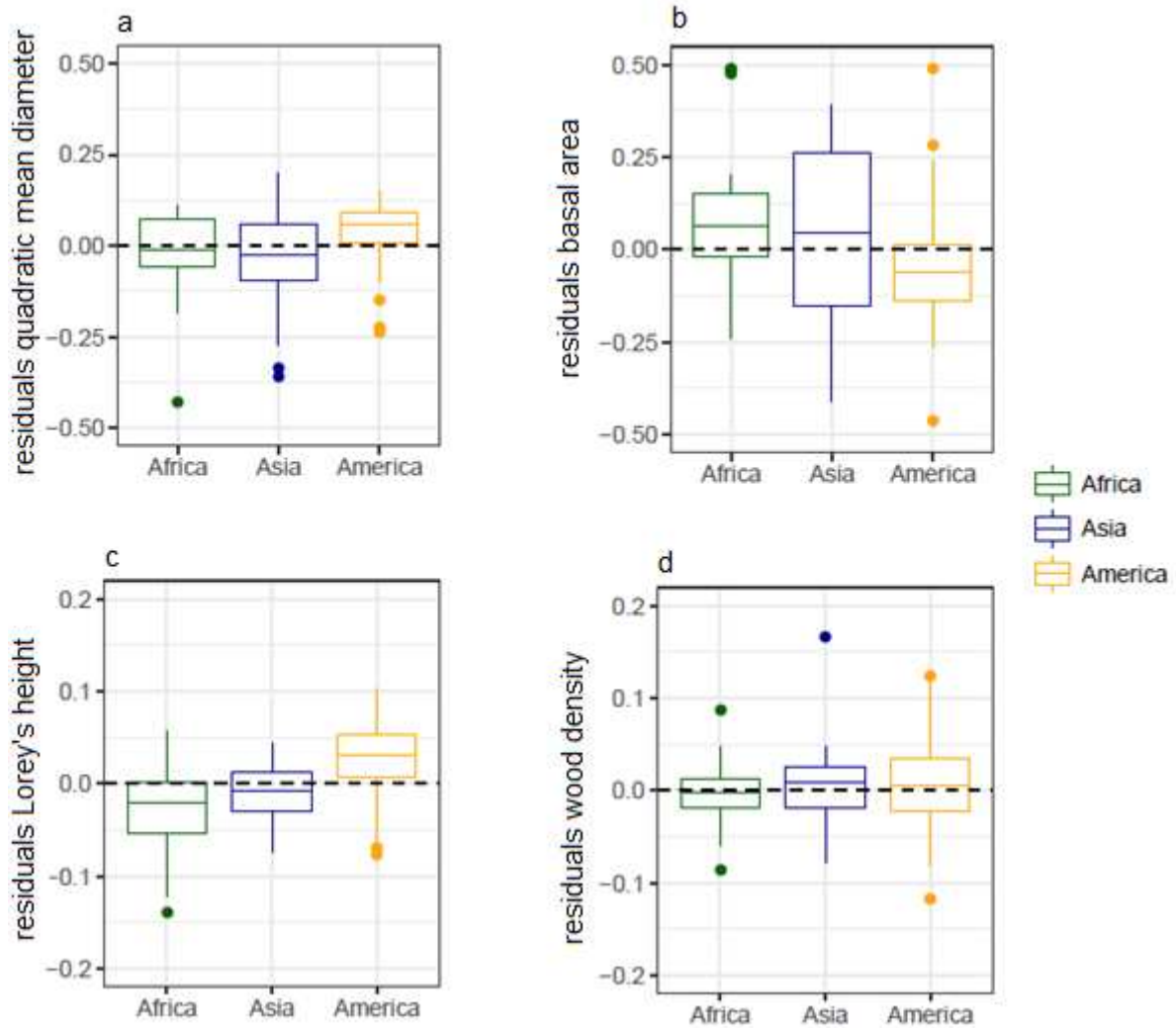
1050 difference with the average observed distribution.(B) Biplot with contribution to the PCA of all

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,

1054 with Africa, America and Asia respectively in green, orange and blue.



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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.

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