



Generic allometric models including height best estimate forest biomass and carbon stocks in Indonesia



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ABSTRACT

The choice of an appropriate allometric model is a critical step in reducing uncertainties in forest biomass stock estimates. With large greenhouse gases emissions due to deforestation, a systematic assessment and comparison of the models available in Indonesia is crucial for accurate assessments of forest carbon stocks and implementing REDD+ projects. In the present study, we compared the ability of two regional and two generic (pan-tropical) allometric models to estimate biomass at both tree and plot levels. We showed that regional models had lower performance in estimating tree biomass, with greater bias (−31–8%) and higher AIC (177–204), compared to generic models (bias: −2–2%; AIC: 57–67). At the plot level, the regional models underestimated biomass stocks by 0–40% compared to the best generic model. The error in plot biomass stocks associated to models relying solely upon DBH ranged between −5 and +15%. The integration of tree height estimated regionally resulted in an overestimate of 5–10% in unmanaged forests. Despite the difficulty to accurately assess tree heights in tropical forests, integrating all or part of them in biomass assessment can reduce uncertainties.

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1. Introduction

Indonesian tropical forests have been extensively logged from 2000 and 2010 (Miettinen et al., 2011), contributing to c. 80% of yearly emissions of greenhouse gases of the country (PEACE, 2007). The ability to accurately estimate forest carbon stocks is essential in Reducing Emissions from Deforestation and Forest Degradation (REDD+) mechanisms in order to establish reliable National Reference Emission Levels (NREL) and to estimate carbon stock changes. However, forest biomass stocks are still poorly estimated in most tropical regions and remain a major uncertainty in our understanding of the potential of tropical forests in mitigating climate change (Houghton, 2005). Several research efforts are under way to fill this gap, relying upon a combination of large-scale remotely-sensed imagery and ground-based measurements (Houghton et al., 2009; FAO, 2010). However, despite strong commitment of the Indonesian Government, its capacity to report car-

bon stocks from forest inventories remains low (Romijn et al., 2012). More generally, the main source of uncertainty in biomass estimates lies in the choice of a particular allometric model (Molto et al., 2013). To date, only two studies have developed biomass models in unmanaged Dipterocarp forests of Borneo (Yamakura et al., 1986; Basuki et al., 2009). However, the range of application of these models have hardly been tested and compared with more generic ones (but see Laumonier et al., 2010). Harvesting trees and weighing their components is time-consuming and most local allometric models encompassed only a small number of trees, likely not to reflect the full tree size distribution (Chave et al., 2005). To avoid this bias and to fill the lack of site-specific allometric equations, two major studies developed generic models and overcame these caveats in accounting for large pan-tropical datasets and large trees (DBH > 50 cm) (Brown, 1997; Chave et al., 2005). However the use of generic models may introduce errors in biomass stock estimates (Chave et al., 2004; Melson et al., 2011) and in Indonesia, site-specific models showed less bias in biomass estimates than generic ones (Basuki et al., 2009; Kenzo et al., 2009b). Depending on the model used, individual tree above-ground biomass (AGB) can vary by as much as a factor two (Basuki et al., 2009), introducing considerable uncertainties in forest biomass stocks computation (Nogueira et al., 2008; Laumonier et al., 2010). Although the use of generic models relies upon the

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assumption that tree-level errors average out at plot level, bias is rarely assessed for forest stands across landscapes (van Breugel et al., 2011). Height and diameter relationship (H–DBH) greatly varies among forest types and regions (Feldpausch et al., 2011). Hence, at sites where no data were used to calibrate the generic models and with different H–DBH relationship, only generic models accounting for both H and DBH are expected to give reliable results (Henry et al., 2010; Vieilledent et al., 2011). Globally, accounting for tree height resulted in more accurate estimate of biomass at both tree and plot levels (Chave et al., 2005; Feldpausch et al., 2012).

Despite the vivid interest for carbon accounting in the region, no study has yet compared how the choice of allometric models affects biomass estimates in Dipterocarp forests. This study is divided into two parts. First, we compared the general accuracy of available peer-reviewed allometric models on an original destructive sample of 108 trees. Second, we investigated how these models affected carbon stock estimates across 12 forest plots representing a total area of 12 ha, focusing on the impact of tree height inclusion in these models.

Our aim was to provide guidance on estimating forest carbon stocks, in order to develop realistic scenarios of GHG emissions from land use change in Indonesia. We are notably addressing: (1) whether site-specific models better predict biomass at both tree and plot levels than generic models; (2) whether the inclusion of tree height improves biomass stock estimates at our sites and (3) how does the inclusion of tree height affect biomass estimates in forests with different H:DBH relationship.

2. Material and methods

2.1. Destructive sampling

We compiled data from destructive measurements made between 2007 and 2012 across East Kalimantan province in Indonesia, mainly from unmanaged lowland Dipterocarp forests (Noor'an, unpublished and Samalca, 2007). These trees did not come from one particular forest site and were hence not suitable to develop a local allometric model. However, we used them to test for the goodness of fit of published models. The DBH distribution ranged from 6 to 129.3 cm, not different from the average DBH distribution of primary forest plots used in this study ($X^2 = 89.9167$, $df = 80$, $P = 0.21$). The main families were Dipterocarpaceae (65%), Malvaceae (3%) and Fabaceae (3%).

2.2. Study sites and forest inventories

We used plots established in unmanaged lowland Dipterocarp forests in Sumatra and East Kalimantan, Borneo (Table 1). The climate at the Kalimantan sites is equatorial with a mean annual rainfall at Tanjung Redeb (Berau District, East Kalimantan) of 2105 mm from 1987 to 2007. All sites were classified as Ultisols (i.e. Xanthic Hapludox, Arenic Kanhapludults). Two sites were established in Community Protected Areas, where local communities historically harvested a few large trees for their own needs (1–5 trees ha⁻¹).

Those plots were classified as old logged over forests. In each plot, all trees were tagged, diameter was measured at breast height (130 cm, DBH) or above buttresses and identified by a professional botanist in the field or at Bogor Herbarium. Dry wood specific gravity (WSG) was determined using the lowest level of botanical identification possible (Chave et al., 2006) and taking the appropriate value reported in the Global Wood Density Database (Zanne et al., 2009). When no botanical identification was available, we used plot-averaged WSG.

Total tree height, referred henceforth to as 'height', in the plots located in Kalimantan was systematically measured using a laser rangefinder, with a possible error of a few meters (Nikon, Forestry 550). In the plots of Sumatra, heights were estimated with a Blume Leiss hypsometer and cross-checked with measurements done by climbing trees (accuracy ± 0.5 m for small and medium trees, ± 3 m for large emergent and canopy trees, Y.Laumonier pers.com). In all the other sites, a single operator did all the measurements to avoid inter-operator variability (Larjavaara and Muller-Landau, 2013).

2.3. Comparison of allometric models at tree level

Despite the importance of Dipterocarp forests in terms of area and carbon stocks, only a few suitable allometric models were found in the literature (Table 2). Two studies (Yamakura et al., 1986; Basuki et al., 2009) proposed site-specific allometric models. Two others (Ketterings et al., 2001; Kenzo et al., 2009a) developed allometric models in secondary logged-over forests. Ketterings et al. (2001) worked in a forest regrowing after slash and burn, in which cultivated species (i.e. *Artocarpus* or *Hevea*) were still present. The second study took place in an industrial logged-over forest concession, where the abundance of pioneer species such as *Macaranga spp.* or *Gluta spp.* indicated a much higher intensity of disturbance (2nd or 3rd rotation). As our study considers 'old-growth secondary forest' i.e. forest stands that have been selectively logged for at least 30 years and have not been clearcut, these last two models were judged irrelevant and were discarded. We also used the generic pan-tropical allometric models developed by Brown (1997), updated by Pearson et al. (2005), and by Chave et al. (2005). These models have been widely used, notably in the context of REDD+, and were recommended by the IPCC guidelines (IPCC 2003, 2006) for estimating carbon stocks in tropical forests.

Using the destructive sample, we compared the performance of prediction of the six models using four *ad hoc* indices, as reported in Vieilledent et al. (2011). We computed the residual standard error RSE, defined as the standard deviation of the residual errors ε_i (with $\varepsilon_i = \log(\text{AGB}_i) - \log(\text{AGB}_{\text{est}})$, where AGB_i and AGB_{est} represent the actual and estimated biomass of a tree i). Large RSE values indicate poor regression models. Second, we computed the coefficient of determination of each model, defined as:

$$R^2 = 1 - \frac{\sum_i \varepsilon_i^2}{\sum_i [\log(\text{AGB}_i) - \log(\text{AGB})_{\text{mean}}]^2} \quad (1)$$

with $\log(\text{AGB})_{\text{mean}}$ being the mean of log-transformed observed values. Models with a high number of parameters generally result in a

Table 1
Plots location, surface, average elevation, average number of stems per hectare and basal area (BA).

| Plot_ID | Forest type | Location | Long | Lat. | Surface (ha) | Elevation (m) | Stems (ha ⁻¹) | BA (m ² /ha) |
|---------|---------------|---------------------------------|---------|--------|--------------|---------------|---------------------------|-------------------------|
| BM_PF | Unmanaged | Batu Majang, East Kalimantan | 115.222 | 0.565 | 2 | 286 | 577 | 33.8 |
| BM_SF | Old secondary | Batu Majang, East Kalimantan | 115.220 | 0.559 | 2 | 213 | 534 | 24.7 |
| BT_PF | Unmanaged | Barong Tongkok, East Kalimantan | 115.415 | -0.024 | 1 | 289 | 496 | 36.5 |
| BT_SF | Old secondary | Barong Tongkok, East Kalimantan | 115.548 | -0.185 | 1 | 180 | 700 | 39.8 |
| PMY_PF | Unmanaged | Pasir Mayang, Sumatra | 102.093 | 1.083 | 6 | 100 | 669 | 30.1 |

Table 2

Published allometric models for mixed-species forests, sample size, DBH range of applicability and coefficient of determination (R). DBH = diameter at breast height, WSG = wood specific gravity or wood density, H = tree height.

| Authors | Year | Region | Reference | Model | Sample size | Range (cm) | R^2 |
|---------------------|------|-----------------|------------|--|---|------------|-------|
| Yamakura & Hagiwara | 1986 | East Kalimantan | Yamakura | Weight stem = $0.02909 \cdot (\text{DBH}^2 \cdot H)^{0.9813}$ | 76 | 4.5–150 | 0.99 |
| | | | | Weight branch = $0.1192 \cdot (\text{weight stem})^{1.059}$ | 191 | | 0.9 |
| | | | | Weight leaves = $0.09146 \cdot (\text{weight stem} + \text{weight branch})^{0.7266}$ | 191 | | 0.92 |
| Basuki et al. | 2009 | East Kalimantan | Basuki.DBH | $\ln(\text{AGB})_{\text{est}} = 1.201 + 2.196 \cdot \ln(\text{DBH})$ | 122 | 6.5–200 | 0.96 |
| | | | | Basuki.WSG | $\ln(\text{AGB})_{\text{est}} = (-0.744) + 2.188 \cdot \log(\text{DBH}) + 0.832 \cdot \log(\text{WSG})$ | | 122 |
| Chave et al. | 2005 | Pan-tropical | Chave.DBH | $\ln(\text{AGB})_{\text{est}} = (-1.499) + 2.148 \cdot \ln(\text{DBH}) + 0.207 \cdot \ln(\text{DBH})^2 - 0.0281 \cdot \ln(\text{DBH})^3 + \ln(\text{WSG})$ | 2410 | 5–156 | 1 |
| | | | | Chave.H | $\ln(\text{AGB})_{\text{est}} = (-2.977) + \ln(\text{WSG} \cdot \text{DBH}^2 \cdot H)$ | | 2410 |
| Pearson et al. | 2005 | Pan-tropical | Pearson | $\ln(\text{AGB})_{\text{est}} = (-2.289) + 2.649 \cdot \ln(\text{DBH}) - 0.021 \cdot \ln(\text{DBH})^2$ | 226 | 5–148 | 0.98 |

better fit to the data and R^2 should be interpreted considering the degrees of freedom of the model $df = n_{\text{obs}} - n_{\text{par}}$, with n_{obs} the number of observations and n_{par} the number of parameters. Third, we computed the Akaike Information Criterion for each model, $\text{AIC} = -2\log(L) + 2n_{\text{par}}$, L being the model likelihood. The best model minimizes the value of AIC. Finally, we computed the overall model bias by summing each individual error (ε_i) expressed as a percentage of tree biomass and taking the median value.

$$\text{Bias} = \text{median} \left(100 \cdot \left(\frac{1}{\exp(\varepsilon_i) - 1} \right) \right) \quad (2)$$

2.4. Inclusion of tree height in allometric models

We investigated the importance of height (H) integration in biomass computation by comparing Chave's equations (Table 2, Eqs. (4), (5)) with and without height. In addition to height measurements ($N = 7389$), we developed regional H :DBH relations (the two regions here are Sumatra and East Kalimantan) in order to test the minimal sample size to accurately estimate tree height. We used here a Weibull function of the form:

$$H_{\text{regional}} = a \times (1 - \exp(-b \times \text{DBH}^c)) + \varepsilon, \text{ with } \varepsilon \sim N(0, 1) \quad (3)$$

Feldpausch et al. (2012) showed that the Weibull- H function lowered the relative error in the small diameter classes ($\text{DBH} < 50$ cm) compared to other usual functions, and was therefore more adapted to skewed diameter distributions. In their study, the authors developed a continental model for South East Asia and Borneo (Table 3A).

We examined how the inclusion of tree height in biomass allometric models affected plot-level biomass estimates. We compared Chave's equation (Table 2, Eq. (5)) including height (1) measured in the field, (2) estimated regionally, (3) estimated continentally and (4) Chave's equation without height (Table 2, Eq. (4)). In addition, we investigated the minimal sample size required to accurately infer H from DBH for each forest type. We developed a Weibull- H function for different sample sizes (1%, 5%, 10%, 20% and 50% of initial population) and tested its ability to predict height of a given pool of trees (20% of initial population). To ensure convergence of the model, the DBH distribution of the sample was similar to the original one. We computed the average error of prediction

$(100 \cdot (H_{\text{predicted}} - H_{\text{measured}}) / H_{\text{measured}})$ using 500 simulations per sample size.

2.5. Plot biomass computation and confidence intervals

For each tree, we computed 1000 biomass estimates for each allometric model using two error terms for both WSG and H following the methodology developed by Feldpausch et al. (2012), assuming no error for the DBH measurements. The error terms were estimated as

$$\widehat{\text{WSG}}_i = \text{WSG}_i + \varepsilon_i, \text{ with } \varepsilon \sim N(0, \sigma_{\text{WSG}}) \text{ and } \widehat{\text{WSG}}_i \in [0.1, 1.1] \quad (4)$$

$$\widehat{H}_i = H_i + \varepsilon_i, \text{ with } \varepsilon \sim N(0, \sigma_H) \text{ and } \widehat{H}_i \in [5, 70] \quad (5)$$

where the “hat” symbol indicates estimates that include an error term randomly chosen in a Normal distribution of mean = 0 and of standard deviation (σ) of WSG or H computed per plot. Biomass stocks were computed at plot level by summing a randomly chosen estimate (for a given allometric model) among 1000 realisations for each tree present in the plot. The 95% confidence interval was calculated as the 2.5th and 97.5th percentiles of the 1000 realisations of each estimate.

All computation and analyses were carried out using R statistical software (R Development Core Team, 2013) and the code is freely available on www.runmycode.org.

3. Results

3.1. Comparison of allometric models

All models accurately predicted the biomass of our sample of felled trees (Table 4), explaining between 90 and 96% of the variation observed. But overall, regional models had lower performance, with greater bias (−31–8%) and higher AIC (177–204), compared to generic models (bias: −2–2% and AIC: 57–67).

The generic allometric model developed by Chave et al. (2005) including height was the best model with the highest coefficient of determination (0.964) and the lowest residual standard error (0.309) and AIC (56.6). On the contrary, the model developed locally in the same region by Basuki et al. (2009) greatly underesti-

Table 3

Continental (A) and regional (B, C) tree height models of the form: $H_{\text{predicted}} = a \cdot (1 - \exp(-b \cdot \text{DBH}^c))$.

| Source | Region | Scale | a | b | c | RSE | N | |
|--------|--------------------------|-----------------|-------------|----------|--------|--------|-------|------|
| A | Feldpausch et al. (2012) | South East Asia | Continental | 57.122 | 0.0332 | 0.8468 | 5.69 | 2948 |
| B | This study | East Kalimantan | Regional | 1989.144 | 0.0018 | 0.5306 | 5.29 | 3192 |
| C | This study | Sumatra | Regional | 56.703 | 0.0547 | 0.739 | 3.417 | 4013 |

Table 4

Comparison of regional (Yamakura, Basuki.DBH and Basuki.WSG) with generic (Chave.DBH, Chave.H and Pearson) models. R^2 is the coefficient of determination, df is the degree of freedom, N par the number of parameters, RSE the residual standard error, AIC is the Akaike Information Criterion, Bias (%) is the median model relative error.

| Equation | R^2 | df | N par | RSE | AIC | Bias |
|------------|-------|------|---------|-------|-------|-------|
| Yamakura | 0.959 | – | – | 0.325 | – | 7.8 |
| Basuki.DBH | 0.912 | 106 | 2 | 0.37 | 177.2 | –29.8 |
| Basuki.WSG | 0.903 | 105 | 3 | 0.359 | 204.6 | –31.3 |
| Chave.DBH | 0.963 | 104 | 4 | 0.319 | 66.9 | –1.7 |
| Chave.H | 0.964 | 107 | 1 | 0.309 | 56.6 | 2.0 |
| Pearson | 0.962 | 105 | 3 | 0.322 | 66.6 | –0.8 |

mated individual tree biomass, resulting in very low aggregated biomass estimates at the plot level (average = –30%, Table 5).

3.2. Including tree height in biomass estimates

Chave.H returned slightly better fit than the one relying solely upon DBH (Chave.DBH, Table 4). Based on this comparison, we considered Chave.H as the most accurate model and served as reference.

The minimum sample size to accurately estimate tree height was low and did not vary significantly among sites (Fig. 1). Measuring only 1% (40–90 trees) reflecting the actual DBH distribution in a plot enabled to accurately estimate tree height at each site. For instance at Barong Tongkok (BT-PF), measuring 1% of the population resulted in an average error of prediction of tree height of 5.6% and 2.75% of the biomass stock versus 4.8% and 2.3% respectively for a sample size of 50%.

Additionally, we developed two regional models (Table 3B and C) to estimate tree heights, and used the continental height estimates developed by Feldpausch et al. (2012). Both regional and continental H-models were compared to the actual heights. Overall, regional models showed smaller bias in height estimates (Fig. 2A) compared to continental models (Fig. 2B). In unmanaged forest, the former showed a bias roughly constant across diameter classes with height overestimated by 10–20%. Overestimation was exacerbated in secondary forest plots, where height estimates of trees $70 < DBH < 120$ cm nearly doubled.

In most plots, the overestimation of tree height by regional or continental H-models resulted in a general biomass overestimation among almost all diameter classes (Fig. 3). The only marked difference was found in Sumatra (PMY-PF) were the continental

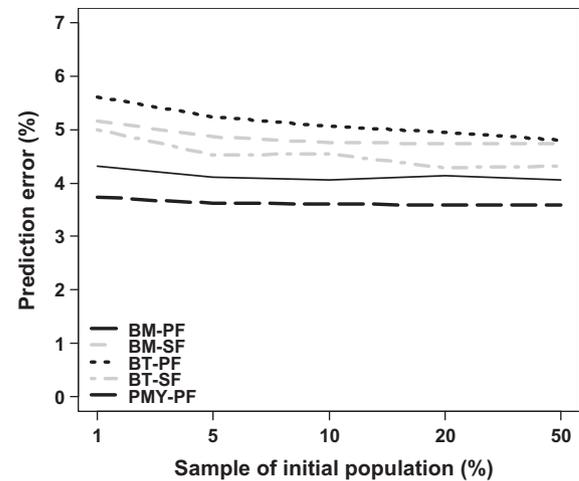


Fig. 1. Error of prediction of tree height ($(H_{\text{predicted}} - H_{\text{measured}}) / H_{\text{measured}}$) using a given fraction (in %) of the initial population to develop a H:DBH model. BM_PF (unmanaged forest in Batu Majang), BM_SF (old secondary forest in Batu Majang), BT_PF (unmanaged forest in Barong Tongkok), BT_SF (old secondary forest in Barong Tongkok), PMY_PF (unmanaged forest in Pasir Mayang).

H-model slightly underestimated tree heights (Fig. 2B) and subsequent biomass estimates (Fig. 3).

3.3. Plot biomass stocks

Plotting all models and confidence intervals reviewed in this study revealed a large range in biomass stock estimates (Fig. 4), with differences greater than 100 Mg ha^{-1} depending on the models and site compared.

When compared with Chave.H model, the regional models developed by Basuki et al. (2009) and Yamakura et al. (1986) underestimated biomass stocks by 25–40% and 0–10% respectively (Table 4). Contrastingly, all generic models relying upon BDH only overestimated in average biomass stocks when compared to the best predictive model (Chave.H). For instance, using Chave.DBH resulted in an AGB overestimation of 15% in the Eastern Kalimantan unmanaged forest plots (Chave.DBH, Table 5), and in an underestimation of 5% in the Sumatran plot. The second generic model (Pearson, Table 5) showed only small differences with Chave.H model, except in the Sumatran plot. The integration of regional height estimates into Chave.H model resulted in a slight overestimate of 5–10% in unmanaged forest, and almost no departure in secondary forest (–3–0%).

Table 5

Mean, 95% confidence interval (CI) and relative difference in percent (diff.) with the equation of Chave et al. (2005) integrating height (Chave.H) by forest types. BM_PF: Unmanaged forest in Batu Majang, BM_SF: old secondary forest in Batu Majang, BT_PF: unmanaged forest in Barong Tongkok, BT_SF: old secondary forest in Barong Tongkok, PMY_PF: unmanaged forest in Pasir Mayang.

| | Yamakura | | | Basuki.DBH | | | Basuki.WSG | | | Chave.DBH | | |
|--------|----------|-----------|-------|--------------|-----------|-------|-------------|-----------|-------|-----------|-----------|-------|
| | Mean | CI | Diff. | Mean | CI | Diff. | Mean | CI | Diff. | Mean | CI | Siff. |
| BM-PF | 347 | (307–392) | –2 | 256 | – | –28 | 256 | (240–290) | –28 | 407 | (352–453) | 15 |
| BM-SF | 226 | (221–259) | –4 | 175 | – | –26 | 178 | (174–199) | –25 | 262 | (239–284) | 11 |
| BT-PF | 388 | (318–450) | –9 | 283 | – | –34 | 295 | (266–348) | –31 | 491 | (407–572) | 15 |
| BT-SF | 397 | (346–452) | 0 | 292 | – | –26 | 293 | (277–337) | –26 | 445 | (387–504) | 12 |
| PMY-PF | 325 | (319–349) | –9 | 215 | – | –40 | 227 | (224–242) | –36 | 340 | (315–349) | –5 |
| | Chave.H | | | Chave.Hregio | | | Chave.Hcont | | | Pearson | | |
| | Mean | CI | Diff. | Mean | CI | Diff. | Mean | CI | Diff. | Mean | CI | Diff. |
| BM-PF | 354 | (293–427) | 0 | 370 | (307–441) | 5 | 406 | (328–477) | 15 | 397 | – | 12 |
| BM-SF | 236 | (219–285) | 0 | 228 | (217–283) | –3 | 255 | (234–306) | 8 | 242 | – | 2 |
| BT-PF | 426 | (314–548) | 0 | 449 | (350–567) | 5 | 492 | (371–601) | 16 | 450 | – | 6 |
| BT-SF | 397 | (339–501) | 0 | 396 | (341–511) | 0 | 438 | (368–539) | 10 | 429 | – | 8 |
| PMY-PF | 357 | (334–386) | 0 | 393 | (359–417) | 10 | 333 | (314–365) | –7 | 302 | – | –15 |

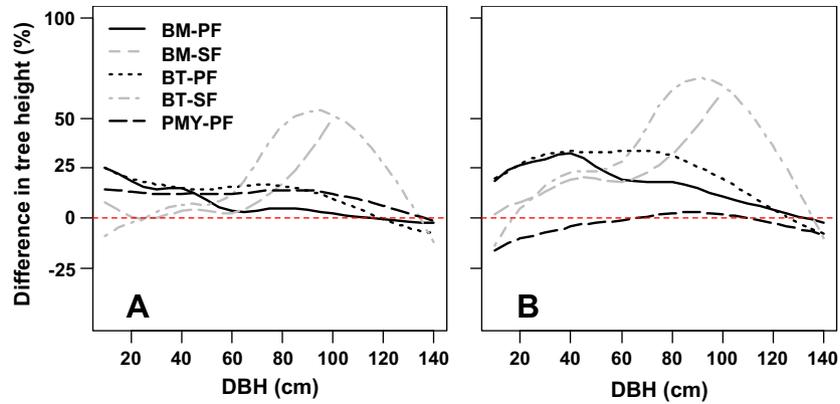


Fig. 2. Difference between measured tree height and (A) regionally estimated height, (B) continentally estimated height plotted as a function of tree DBH by forest types. The curves were smoothed by a lowess method. At BM-SF, the upper DBH limit was 100 cm. BM_PF: unmanaged forest in Batu Majang, BM_SF: old secondary forest in Batu Majang, BT_PF: unmanaged forest in Barong Tongkok, BT_SF: old secondary forest in Barong Tongkok, PMY_PF: unmanaged forest in Pasir Mayang.

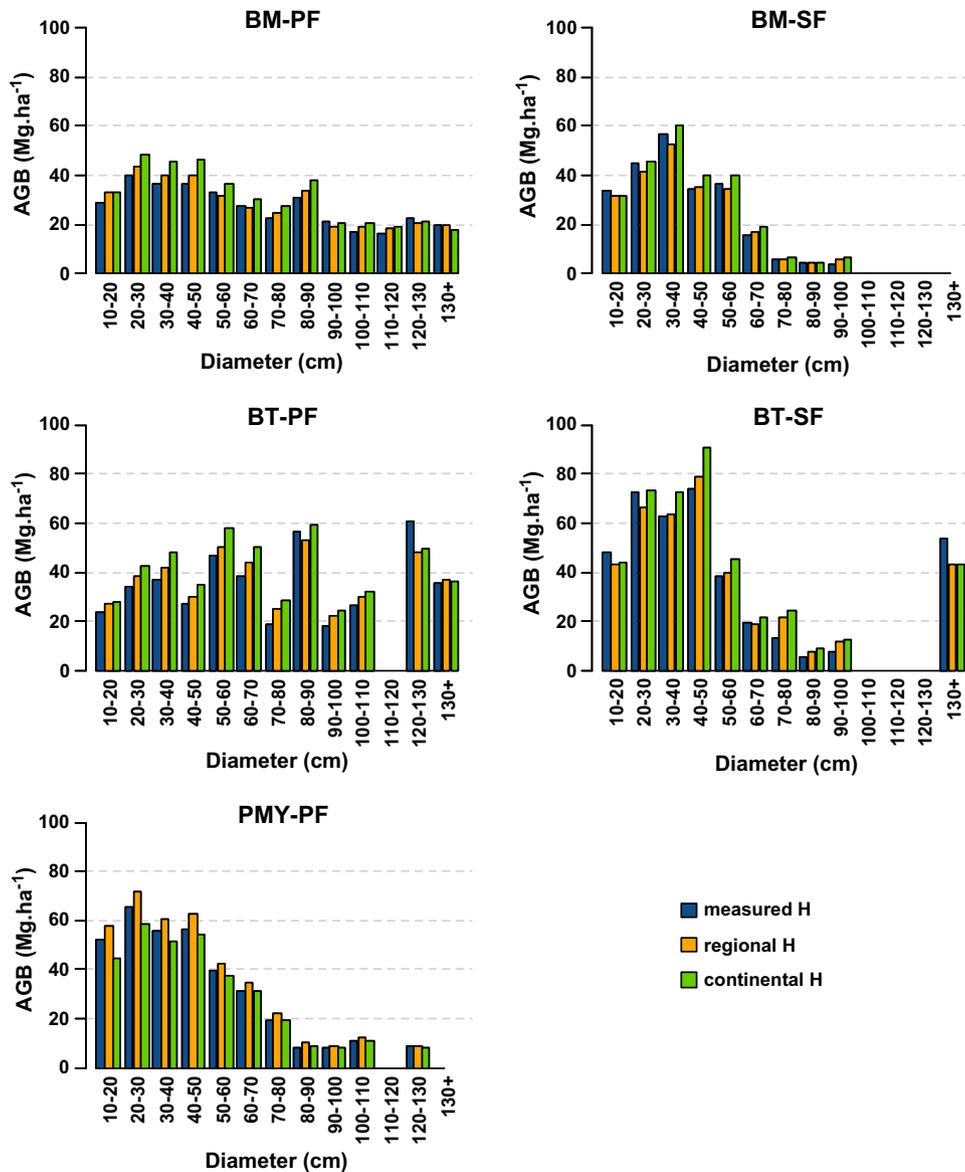


Fig. 3. Comparison of biomass stocks computed using Chave's equation (Chave.H) by diameter class per site, using measured height (blue), regional height estimates (orange) and continental height estimates (green). BM_PF: unmanaged forest in Batu Majang, BM_SF: old secondary forest in Batu Majang, BT_PF: unmanaged forest in Barong Tongkok, BT_SF: old secondary forest in Barong Tongkok, PMY_PF: unmanaged forest in Pasir Mayang. (For interpretation of the references to color in this figure legend, the reader is referred to the web version of this article.)

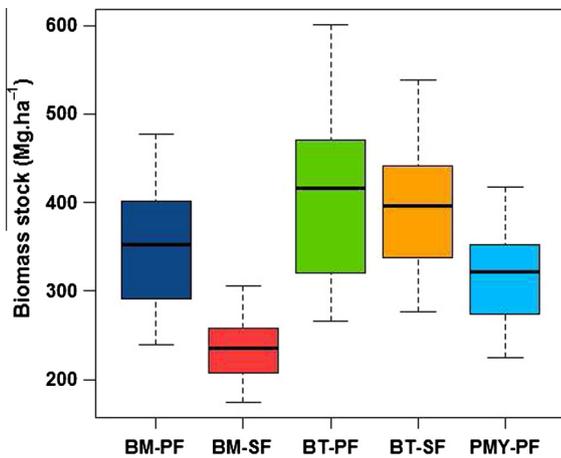


Fig. 4. Biomass stocks range combining six allometric models (Table 3) and their bootstrapped confidence intervals per forest type. Boxes represent 25th and 75th percentiles, whiskers represent 10th and 90th percentiles, line represents median. BM_PF: unmanaged forest in Batu Majang, BM_SF: old secondary forest in Batu Majang, BT_PF: unmanaged forest in Barong Tongkok, BT_SF: old secondary forest in Barong Tongkok, PMY_PF: unmanaged forest in Pasir Mayang.

4. Discussion

4.1. Recommendations on the use of allometric models

The generic pantropical model developed by Chave et al. (2005) including tree height provided the best biomass estimates when applied to our destructive samples (Table 3). This result was expected as the Indonesian sites used in their study were both located in East Kalimantan, about 200 km from where trees used in this study were collected. Additionally, Chave et al. (2005) showed that H-models had smaller departure from observed values compared to DBH-models in most tropical forests, but with a notable exception in East Kalimantan. Our results are consistent with this finding, showing that generic models relying solely upon DBH and WSG (Chave.DBH or Pearson) were also very good at predicting biomass at our sites. However, these last models can result in an error of $\pm 15\%$ of the actual biomass stock in certain forests.

Despite the fact that Dipterocarp forests represent the dominant vegetation in Borneo, it is most likely that accounting for other forest types with different structures (i.e. kerangas forests, peat swamp forests, forests on limestone) would have given different results. For instance, in African forests where H:DBH relationship is very different and from which no data were used to calibrate those generic models, Chave.DBH model largely overestimated biomass while Chave.H gave very good fit (Henry et al., 2010; Vieilledent et al., 2011; Fayolle et al., 2013).

As tree height is generally not recorded in forest inventories, models relying solely upon DBH are likely to remain widely used by foresters. We showed here that the generic model developed by Brown (1997), updated by Pearson et al. (2005), showed similar performance to the model integrating WSG and DBH (Chave.DBH), but with slightly smaller bias. Both models outperformed the regional models developed in East Kalimantan. In conclusion, generic models relying solely upon DBH and WSG remain appropriate, but should be used with caution as they generally overestimate biomass.

4.2. Measuring tree height in tropical forests

With advances in laser instruments, it has become easier to accurately and rapidly assess tree height in the field. In a tropical forest, direct vertical measurements of the last branch was found to underestimate of actual tree height by 20% (Larjavaara and Mul-

ler-Landau, 2013). It is likely that the error remains proportional to tree height, affecting primarily emergent trees. Rapid advances in LiDAR-derived mean canopy height might help to overcome this caveat and seems to be a promising way of integrating average forest stand height into plot carbon stocks measurements (Asner et al., 2011).

Despite similar continental H:DBH relationships found across Asia, Africa and the Guyana Shield (Feldpausch et al., 2011), we found that continental H:DBH models only poorly explained the variance observed at our sites, notably in old-growth secondary forests (Fig. 2). We highlight here that the continental model proposed by Feldpausch et al. (2012) was originally developed for unmanaged forests and should be used with caution in secondary forests. For instance, trees growing in logged forests in the Amazon were found to be shorter with larger crowns (Nogueira et al., 2008). This phenomenon might explain our results in secondary forests, where large trees had much smaller heights than expected. We showed that H:DBH model can be fitted with only a small fraction of the forest stand (Fig. 1), as long as the sample is equally distributed along the actual DBH distribution. In a first attempt, trees were randomly chosen, embedding the model to converge in most cases. This result is encouraging and shows that integrating tree height into carbon stock assessment would not require a lot of additional field work.

4.3. Above-ground biomass stock in trees of Dipterocarp forests

Using the best predictive model (Chave.H), we found an average value of 378 Mg ha^{-1} in unmanaged and 316 Mg ha^{-1} in secondary forests. These values are lower than those previously reported for Dipterocarp forests (Paoli et al., 2008; Slik et al., 2010). Both studies used Chave's equation based on DBH and WSG, with AGB stocks ranging from 457 to 606 Mg ha^{-1} . Our study shows that these figures are likely to be overestimated by at least 10%. Lower AGB stock in secondary forests was mainly explained by the absence of very large trees (DBH > 100 cm) that usually encompass a large fraction of AGB in tropical forests (Paoli et al., 2008; Rutishauser et al., 2010). However, these figures remained relatively high compared to forests recovering from conventional logging that range between 150 and 300 Mg ha^{-1} (Berry et al., 2010; Saner et al., 2012). This strengthens our initial postulate of considering these plots as mature secondary forests and constitutes one of the reasons we decided not to use allometric models developed in logged-over forests of Sumatra (Ketterings et al., 2001) or Borneo (Kenzo et al., 2009a). At one site (BT_SF), no logging activity was carried out over the last 40 years, while none was carried at the second site (BM_SF).

Such systematic assessment should be performed in other forest types and ecoregions across Indonesia in order to determine the validity and the choice of the appropriate allometric model.

4.4. On the choice of a suitable allometric model

The choice of a particular allometric model will remain mainly driven by data availability. Due to time and costs constraints, most forest inventories are restricted to DBH measurements and DBH-models will remain widely used. However, accounting for tree heights can reduce uncertainties surrounding biomass estimates in Dipterocarp forests. Overall, the average absolute difference in biomass stocks between H-models and DBH-models was 10–15%, in the same magnitude of what was found in Malaysian Borneo and Brunei (Feldpausch et al., 2012). Despite the fact that local H:DBH allometry can be obtained from a small sample (45–90 individuals) of the stand, regional H-models can provide a fair alternative (Fig. 2A). In the field, measuring tree heights do not represent a heavy extra-cost and required on average 3–5 min at our

sites. Most models overestimate the biomass of large trees, what could be considerably reduced in measuring systematically their height. In addition to a representative sample of the DBH distribution, focusing on large trees might help improving biomass estimate and represent a good compromise between time constraints and accuracy.

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