

Estimation of above-ground biomass in forest stands from regression on their basal area and height

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Abstract. A generic regression model for above-ground biomass of forest stands was constructed based on published data ($R^2 = 0.88$, $RSE = 32.8$ t/ha). The model was used 1) to verify two allometric regression models of trees from Scandinavia applied to repeated measurements of 275 sample plots from database of Estonian Network of Forest Research (FGN) in Estonia, 2) to analyse impact of between-tree competition on biomass, and 3) compare biomass estimates made with different European biomass models applied on standardized forest structures. The model was verified with biomass measurements from hemiboreal and tropical forests. The analysis of two Scandinavian models showed that older allometric regression models may give biased estimates due to changed growth conditions. More biomass can be stored in forest stands where competition between trees is stronger. The tree biomass calculation methods used in different countries have also substantial influence on the estimates at stand-level. A common database of forest biomass measurements from Europe in similar to pan-tropical tree measurement data may be helpful to harmonise carbon accounting methods.

Key words: forest, biomass, model, competition.

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Introduction

Forests are an integral part of the global carbon (C) cycle and a large C reservoir that is mitigating climate change (Nabuurs *et al.*, 2007). Accurate estimation of the rate of biomass and C accumulation and stor-

age in forest stands is central for understanding how forests have, and will influence climate. However, Neumann *et al.* (2016) found large discrepancies between tree biomass estimates when methods used in different European countries were compared. Typically, forest biomass compo-

nents (stem wood, stem bark, foliage, living branches, stump or root system) are estimated with single tree regression models (Ter-Mikaelian & Korzukhin, 1997; Zianis *et al.*, 2005; Henry *et al.*, 2013) based on tree stem diameter at breast height (d). Only a few of the models are based on extensive sample of trees across different growth conditions (Marklund, 1988; Repola, 2008, 2009; Chave *et al.*, 2005, 2015). However, systematic errors may occur when d -based allometric regression models are applied in stands of different structure (stand density, competition *etc.*) compared to the model data. The structure of forest stands depends on species composition, soil fertility, stand density, management, competition between trees and disturbances (Bormann & Likens, 1992; Schiatti *et al.*, 2016).

When a tree is used as an independent sample then the rate of biomass accumulation increases with the tree size as estimated by Stephenson *et al.* (2014) based on allometric regression models which have d as an independent variable. On the other hand, the fact that bigger trees require each more space compared to small trees for sustaining their growth is well known from allometric theory (Enquist & Niklas, 2002; Nilson, 2005) and also from yield tables used for forest management practice for more than a century (Vanclay, 1994). Forest canopy level feedback loops (Hasenauer, 1997; Frey, 2009) result in a relatively stable crown cover, leaf area index and amount of absorbed photosynthetically active radiation (Nilson & Peterson, 1994) in the absence of stand structure altering disturbances (Schiatti *et al.*, 2016). Hence, tree growth is determined by the potential phytoproductivity of soil (Kölli, 2002; Kölli & Kanal, 2010) and biomass accumulation rate per unit area of a forest is influenced by the tree growth induced competition (Contreras *et al.*, 2011) and limited by tree mortality. An important factor in biomass accumulation, the stem wood density, is influenced by the forest stand basal area increment (Jaakkola *et al.*, 2005) which de-

pends on soil fertility (Kask & Pikk, 2009) and species composition (Lilleleht, 2011). Marklund (1988) found that inclusion of basal area increment improved the accuracy of allometric tree biomass models.

Cannell (1984) showed that a simple model $\ln(B_{AG}) = -0.37 + 0.84 \ln(GH)$ described 91% of the above-ground biomass B_{AG} (t/ha) variation for a dataset of 640 forest stands based on the product of the stand mean height H (m) and basal area per unit area G (m²/ha). In forestry, GH is used to calculate standing wood volume $V=GHF$, where F is stem form factor of the stand (Krigul, 1972). In biomass studies F_{s+sb} is calculated accounting for stems and branches (Nebel *et al.*, 2001). F can be based on parabolic height (Cannell, 1984) or cylinder (Krigul, 1972). Cannell (1984) presented also tree species-specific linear biomass models. However, new measurements of B_{AG} have been published since offering an opportunity to further explore and develop this model. If the model is still valid, it could be used as a standard tool in those geographic areas where forest biomass models are not available. The stand height and basal area-based biomass models are also useful for remote sensing applications (Wulder *et al.*, 2008; Asner *et al.*, 2012; Hayashi *et al.*, 2015; Arumäe & Lang, 2016; Moreno *et al.*, 2016).

Our primary aim was to 1) develop a generic, species independent, above-ground biomass model for forest stands based on stand height and basal area, 2) verify tree biomass models regarding competition, and 3) compare the generic model-based estimates with biomass measurement data and published estimates of biomass. Firstly, we collected published data of stand-level above-ground biomass and stand structure (trees per unit area (N), mean tree stem breast height diameter D , forest height H and G) and estimated parameters for generic biomass model. Secondly, tree biomass regression models were applied to trees growing on Estonian Network of Forest Research (Kiviste *et al.*, 2014) sample

plots and the aggregated estimates were compared to the generic model and analysed in respect to two simple stand structure indices. Additional data sets used in the validation and analysis of the generic model were artificial stands (Neumann et al., 2016), published oak biomass data from Korea, rainforest data from Amazon and Chile, and sample plot measurement data from Poland, Estonia and Brazil.

Material and Methods

Model construction

When the biomass data (Table A1.1) was plotted as a function of GH the relationship was linear with only few outliers. It can be expected that wood density (δ), stand form factor F and biomass allocation into stem, crown and root system will cause some deviations from the general trend. It is possible that stand structure variables e.g. form factor or mean height are not exactly defined and the given values in publications are not compatible. The outliers were two temperate mixed deciduous forests (Newman et al., 2006) showing small biomass for GH and *Quercus* stands in Korea (Son et al., 2004) with big biomass at GH compared to Park et al. (2005).

After excluding the outliers which substantially deviated from the dataset, the relationship between forest above-ground biomass was first estimated with linear model

$$B_{AG} = a_1 + a_2GH \quad (1)$$

(Figure 1, Table 1). The linear model did not work well in young stands and predicted 25.4 t/ha biomass for the stands with $GH = 0$. The second, advanced version of the model with start correction component

$$B_{AG} = a_1 + a_2GH + a_3 \exp(-GH/100) \quad (2)$$

estimates reasonably 7 t/ha at $GH = 0$. For example, above-ground biomass estimate

of a young Norway spruce stand with $H = 100$ cm and $N = 2000$ trees/ha is 2.4 t/ha according to the model from Pastorella & Paletto (2014) and 0.5 t/ha according to Mitt et al. (2014).

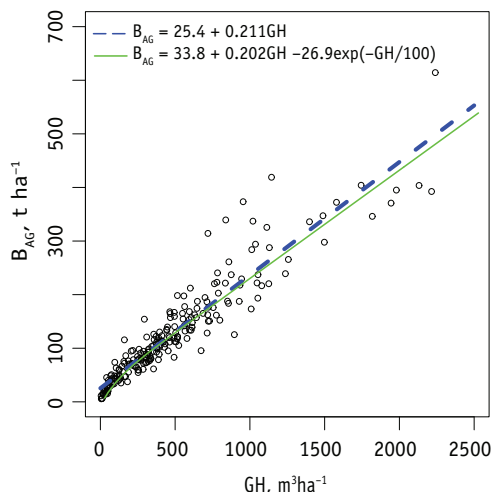


Figure 1. Predictions of model (1) and (2) and the observed values (points) of stand above-ground biomass.

Joonis 1. Puistute maapealse osa biomassi B_{AG} seost rinnaspindala (G) ja kõrguse (H) korrutisega kirjeldavad mudelid (1) ja (2). Algandmed on tabelis A1.1.

Table 1. The parameter estimates for model (1) and (2). All model parameters (standard error in parentheses) were significant at $p < 0.05$. RSE is model standard error.

Tabel 1. Puistute maapealse osa biomassi mudelid. Sulgudes on standardviga.

Model component / Mudeli osa	Model / Mudel	
	(1)	(2)
R_{adj}^2	0.8816	0.8844
Deg. of freedom	225	224
RSE (t/ha)	33.2	32.8
a_1	25.36 (3.3)	33.83 (4.7)
a_2	0.2110 (0.0051)	0.2020 (0.062)
a_3	-	-26.86 (10.6)

To estimate the impact of individual publications and individual stands on the model (2) the biomass data from literature was randomly sampled into 1000 calibration and validation subsets. In Table (2) are the percentiles (t/ha) of the *GH*-based biomass model (2) residual standard error (RSE), root mean squared error (RMSE) and mean estimation error (MEE). MEE and RMSE are based on validation subsets.

Table 2. Dependence of model (2) on sample data. Subsets "rs" are random selection of stands, "rp" are random selection of publications.

Tabel 2. Võimalik algandmete mõju mudelile (2) kui algandmestik jagati 1000 katses mudeli andmeteks ja valideerimise andmeteks. Valiti puistuid "rs" ja teises katses publikatsioonide "rp".

Estimator / Veahinnang	Subset / Valim	Percentiles / Protsentiilid				
		0%	25%	50%	75%	100%
RSE	rs	23	30	33	35	40
RMSE	rs	22	31	34	36	46
MEE	rs	-15	-3	0	3	14
RSE	rp	13	23	29	38	50
RMSE	rp	15	26	35	40	55
MEE	rp	-24	-8	-1	7	23

Model analysis on forest growth study network data

A subset of 275 sample plots (FGN stands) was extracted from the database of Estonian Network of Forest Research (Kiviste *et al.*, 2014). Considering all repeated measurements on the sample plots the total number of observations was 1037. Main tree species in the forests were Scots pine, Norway spruce, silver birch and trembling aspen. All the forests were older than 20 years when first measured (Table A1.2). Tree height for the trees without height measurement was estimated from model of Näslund fitted on the sample trees for each sample plot. The intercept of the model was fixed and two parameters were estimated according to

Padari (1999). Three above-ground biomass b_{AG} estimates for each tree were calculated based on d and h : 1) $b_{AG(R)}$ as sum of fractions (foliage, branches, stem, bark, stump) by using simple multivariate models from Repola (2008, 2009), 2) $b_{AG(M)}$ by using corresponding models from Marklund (1988), and 3) $b_{AG(C)}=0.0673(d^2h\delta)^{0.976}$ with the model (4) from Chave *et al.* (2015) by using wood density given in Table A3.1. Total above-ground biomass B_{AG} G and basal area weighted H (Lorey height) were calculated for each sample plot. By using G and H biomass estimate with Cannell (1984) model $\ln(B_{AG})=-0.37 + 0.84\ln(GH)$ was calculated for each FGN sample plot.

We used two stand structure indices to analyse the single tree-based biomass estimates of FGN stands in respect to the model (2). Hegyi (1974) index of competition for a sample plot

$$CI_{Hegyi} = 1/N_t \sum_{j=1}^{N_t} \sum_{i=1}^n d_i / (d_j s_{ij}) \quad (3)$$

is based on distance s_{ij} from target tree j to the i -th neighbour tree considering competition radius of 8 m. Only the trees from upper layer were included for CI_{Hegyi} . We did not apply sample plot extension as described by Lilleleht *et al.* (2014) in this test; instead, the list of target trees N_t was limited to those located by 8 m towards the centre from sample plot border. Increase in CI_{Hegyi} indicates increase in competition. The second index

$$CI_{LTI}=L/L_{TI} \quad (4)$$

is based on mean distance between trees $L=(100/\sqrt{N})$ and Nilson's model of forest stand self-thinning $L_{TI}=k_1 + k_2D+k_3DH_{100} + k_4H_{100}$ for different species (Sims *et al.* 2009), where H_{100} is site fertility index. The value of $CI_{LTI}=1$ means that the stand is on the self-thinning line and $CI_{LTI}<1$ indicates intensive self-thinning due to insufficient growth space.

To describe the variation in stand-level biomass explained by the competition indi-

ces we constructed general additive models (GAM) with simple isotropic smooth in R computing environment using library “mgcv”. The dependent variables were stand biomass estimates based on $b_{AG(R)}$ and $b_{AG(M)}$ and their normalized values in respect to the model (2).

Model assessment on simulated stand structures

The second dataset (FORMIT stands) contained simulated standardized forest structures for Scots pine, Norway spruce, silver birch, European beech, and common oak. The dataset was adopted from Neumann *et al.* (2016) who compared forest carbon estimation methods across the Europe. The data set was generated with STANDGEN (Kittenberger, 2003), that includes single tree simulation model MOSES (Hasenauer, 1994; Klopff *et al.*, 2011). With STANDGEN, for each of the five selected tree species, three of 0.25 hectare stands were generated which differed in mean and standard deviation of tree diameter and represented forest stands at different ages, allocation, stem number, or stocking density. The generated stands corresponded to a young stand (quadratic mean $D = 10$ cm with standard deviation 1 cm), a middle-aged stand ($30 \text{ cm} \pm 5 \text{ cm}$), and an old stand ($50 \text{ cm} \pm 10 \text{ cm}$). The country-specific above-ground biomass estimates of FORMIT stands were compared to model (2).

Biomass data and published biomass estimates for model assessment

Silver birch biomass data were from 18 stands growing on abandoned agricultural land in the Mazowsze region (central Poland). Scots pine data contain empirical biomass material from 18 managed stands in Bory Lubuskie (western Poland) from different site conditions and age. In each stand the sample plots consisted of approximately 200 trees. More details are given by Zasada *et al.* (2014) and Bronisz & Zasada (2016).

Biomass data from 23 grey alder stands (Uri *et al.*, 2014) from Estonia were used. The stands were not used for parameter estima-

tion for model (2). Foliage mass for the grey alder stands was not measured. For 7-years and older stands constant foliage mass estimate of 3 t/ha was used (Aosaar *et al.*, 2013). Foliage mass for younger stands was scaled linearly starting from 0.5 t/ha for 2-years old grey alder stand. Woody above-ground biomass was calculated from stem and branch volume using biomass density. Also species-specific models for silver birch and grey alder were estimated by using the formulation of model (2).

We also analysed data from three (100×100 m) permanent plots within an old-growth Atlantic moist forest in Vale Natural Reserve, south eastern of Brazil, where all stems with $d \geq 10$ cm were measured at 1.3 m height or above any buttresses (for site description see Rolim *et al.* 2016). For estimates of aboveground biomass except Areaceae species we applied a pantropical allometric model developed by Chave *et al.* ((4): 2015), and species-specific wood density from global database published by Chave *et al.* (2009). For Areaceae species the model from Goodman *et al.* (2013) was used.

Additional published data were from oak forest in Korea (Li *et al.*, 2012) (includes Son *et al.*, 2004; Park *et al.*, 2005) and stand level biomass data by age classes for oak forests in Poland (Orzeł *et al.*, 2006). Rice *et al.* (2004) provided sufficient data in their Table 5 for an old-growth Amazon rainforest to calculate G and they also gave an estimate of the closed canopy height (40 m) and the emergent tree height (55 m). We used an estimate of $H = 45$ m for our calculations. Nebel *et al.* (2001) used $F_{stb}=0.6$ to calculate stem and branch volume for high restinga, low restinga and tahuampa forest in Peruvian Amazon. We calculated initial $GH = V/0.6$ for the stands. We found also biomass data for *Fitzroya cupressoides* forest (Urrutia-Jalabert *et al.*, 2015) and included in comparison the data for trees with $d \geq 10$ cm. Form factor conversion for tropical forest to adopt the data for GH -based model is given in Appendix A2.

Results

Stand-level biomass estimates in FGN forests

The estimated biomass in FGN sample plots was better described by stand height than stand mean diameter for Estonian forests (Figure A4.1). The mean above-ground biomass of FGN forests $B_{AG(R)} = 166.5$ t/ha was estimated with Repola (2008, 2009) model, $B_{AG(M)} = 184.1$ t/ha with Marklund (1988) model and $B_{AG(Ch)} = 204.9$ t/ha with the model from Chave *et al.* (2015). In the FGN forests, $B_{AG(R)}$ was in average 3.2 t/ha less compared to Cannell (1984) model and 10.7 t/ha less compared $B_{AG(2)}$ estimated with model (2) and $B_{AG(M)}$ was respectively 14.4 t/ha and 6.9 t/ha more (Figure 2).

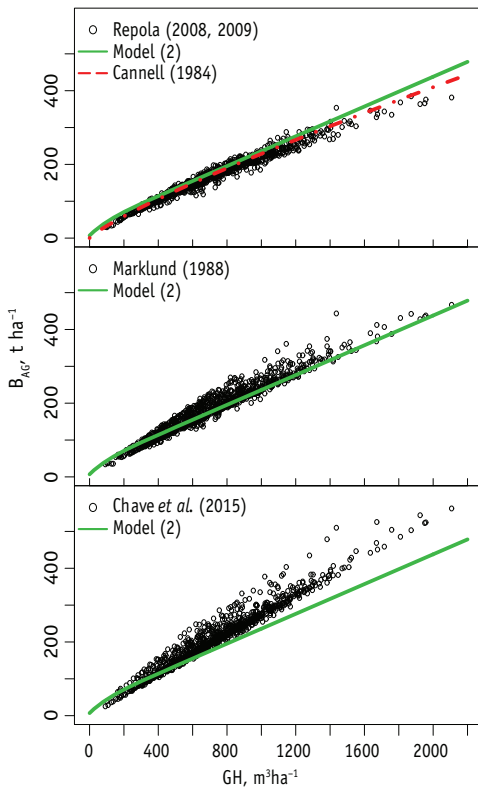


Figure 2. Above-ground biomass of FGN forests as a function of stand basal area and height.

Joonis 2. Puistu maapealse osa biomass metsa kasvukäiguproovitudel hinnatuna erinevate mudelitega.

Biomass as a function of competition indices

The correlation of above-ground biomass estimates was negative with CI-s. However, the numeric values of $CI_{Hegy\ddot{y}i}$ and CI_{LTJ} have opposite meanings i.e. stronger competition is indicated by bigger values of $CI_{Hegy\ddot{y}i}$ and smaller values of CI_{LTJ} , and the conclusions depend on particular CI. Based on $CI_{Hegy\ddot{y}i}$ less biomass is stored in the stands where competition is stronger. This is influenced by the characteristics of $CI_{Hegy\ddot{y}i}$ which becomes a function of L when averaged over all trees in a sample plot. Stronger competition according to $CI_{Hegy\ddot{y}i}$ exists in denser stands which are also younger in Estonia. According to CI_{LTJ} more biomass is stored in stands where intensive self-thinning process indicates stronger competition.

The CI-s explained up to 76% of variation in biomass (Table 3) when no other predictive variables were included in GAM-model. CI_{LTJ} described 34 to 76% of variation in biomass estimates except for $B_{AG(R)} - B_{AG(2)}$ in pine stands. Self-thinning process was intensive in many of the stands according to CI_{LTJ} , but the $B_{AG(R)}$ was substantially less than $B_{AG(2)}$ (Figure 3) and the relationship was scattered. The CI_{LTJ} had strong negative correlation with stand basal area (-0.7) and the correlation was negative (-0.6) with both $B_{AG(R)}$ and $B_{AG(M)}$. This was also reflected in the relationships of CI_{LTJ} with $B_{AG(R)} - B_{AG(2)}$ and $B_{AG(M)} - B_{AG(2)}$ (Figure 3).

$CI_{Hegy\ddot{y}i}$ described 21–66% of variation of the difference $B_{AG(R)} - B_{AG(2)}$ and much less for other analysed biomass variables (Table 3). When $CI_{Hegy\ddot{y}i}$ had smaller values (less competition) then $B_{AG(R)}$ was also less compared to the generic model (2). The relationship was much more scattered for $B_{AG(M)}$ (Figure 3) with most of this scatter in $B_{AG(M)} - B_{AG(2)}$ attributed to deciduous stands and Norway spruce stands. The $CI_{Hegy\ddot{y}i}$ had smaller values in older stands (correlation $(A, CI_{Hegy\ddot{y}i}) = -0.42$) where stand density is less than in younger stands. Both CI-s did, however, improve only marginally B_{AG} estimates when included as third variable

Table 3. Percentage of variation in biomass described by competition indices using general additive models with isotropic smooth function.

Tabel 3. Biomassi variatsioon, mida konkurentsaindeksid (CI) kirjeldavad.

CI	Main species <i>Peapuuliik</i>	Described % of biomass variation / <i>Kirjeldatud variatsioon (%)</i>			
		$B_{AG(R)} - B_{AG(2)}$	$B_{AG(M)} - B_{AG(2)}$	$B_{AG(R)}$	$B_{AG(M)}$
$CI_{Hegyí}$	Spruce	66.3	8.1	2.3	3.6
$CI_{Hegyí}$	Pine	59.3	17.8	24.6	27.4
$CI_{Hegyí}$	Deciduous	21.2	2.4	5.1	3.6
CI_{LTJ}	Spruce	34.4	76.4	65.0	61.7
CI_{LTJ}	Pine	3.8	45.1	55.1	52.3
CI_{LTJ}	Deciduous	46.9	59.7	61.5	61.0

with G and H which described over 94% variation in B_{AG} .

The influence of competition to the relationship of biomass on GH occurs due to form factor decrease (correlation (F , $CI_{Hegyí}$) = 0.65). F decreases with tree height and depends on height to stem diameter ratio. Weaker competition stipulates stem diameter growth more than tree height growth and causes smaller form factor of stems but the share of branches increases (Larson, 1963). In FGN forests $CI_{Hegyí}$ described 38% of variation in branch biomass to total above-ground biomass ratio $B_{branch(R)}/B_{AG(R)}$ in broadleaved-species dominated forests with moderate negative correlation. Less than 11% of $B_{branch(R)}/B_{AG(R)}$ variation in pine and spruce dominated forests was described by $CI_{Hegyí}$ but there was a weak positive correlation. CI_{LTJ} described less than 15% of the $B_{branch(R)}/B_{AG(R)}$. From Marklund (1988) we used models that estimated biomass sum of foliage and branches i.e $B_{crown(M)}$. The variation of $B_{crown(M)}/B_{AG(M)}$ described by $CI_{Hegyí}$ was 32% in spruce stands, 26% in pine stands with moderate positive correlation. There was no correlation between $B_{crown(M)}/B_{AG(M)}$ and $CI_{Hegyí}$ in deciduous stands. CI_{LTJ} had almost no correlation with $B_{crown(M)}/B_{AG(M)}$ in pine stands, but there was moderate positive correlation in deciduous stands and spruce stands although less than 23% of variation in was described. For comparison,

$B_{crown(M)}/B_{AG(M)}$ had strong negative correlation with stand height and moderate negative correlation with basal area.

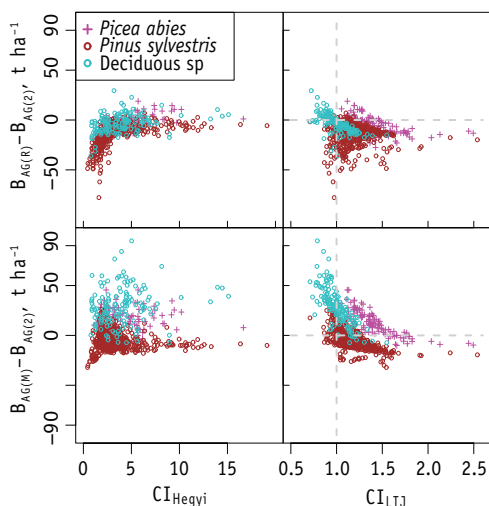


Figure 3. Difference of aggregated single tree biomass B_{AG} in FGN forests from that predicted by model (2) as a function of Hegyi (3) and Nilson's (4) CI-s.

Joonis 3. Üksikuude mudelitel põhineva puistu biomassi erinevus üldisest mudelist (2) sõltuvalt Hegyi (3) ja Nilsoni (4) konkurentsaindeksitest metsa kasvukäiguproovitudükkidel.

Model assessment on simulated stand structures

The biomass estimates for simulated stands with country-specific models had a rather large scatter (Figure 4). Biomass estimate for Norway spruce with Spanish model yielded exceptionally large biomass compared to model (2) and models from other countries. However, on average the biomass estimates for Norway spruce, Scots pine and silver birch were mostly in line with the model (2). The generic model (2) underestimates biomass in beech and oak stands because the species have denser wood (Figure 4). Sample plot data from literature (Cannell, 1982; Bartelink, 1997) show also high biomass density for beech forests. Biomass estimates with model of Orzel *et al.* (2006) are smaller than predicted with national biomass models for FORMIT oak stands, but still greater than estimated with the *GH*-based model (2).

Biomass measurement data and published data for model assessment

The model (2) does not work well on *Quercus mongolica* and *Quercus variabilis* forest data from review by Li *et al.* (2012) (Figure 5). No data was available for form factor assessment. The structure of the forests was described by their arithmetic mean *D* and arithmetic mean *H*, the number of trees per hectare and forest age. We estimated stem biomass density by assuming stem form factor $F = 0.5$ (Lumbres *et al.*, 2014) as $\delta = B_{AG,stem} / (0.5GH)$ and the results ranged from 0.16 to 2.0 g/cm³. By choosing other realistic stand or stem form factor values (Cannell, 1984; Jung *et al.*, 2015) the stem organic matter density estimates did not substantially improve.

However, when sufficient data was available for stand form factor conversion, then it was possible to compare biomass estimates from tropical forests with hemiboreal forests by their *GH* (Figure 5). Compared to the model (2) an old-growth Amazon rainforest (Rice *et al.*, 2004), old-growth Atlantic forest and flood plain forests in the Peruvian Amazon (Nebel *et al.*, 2001)

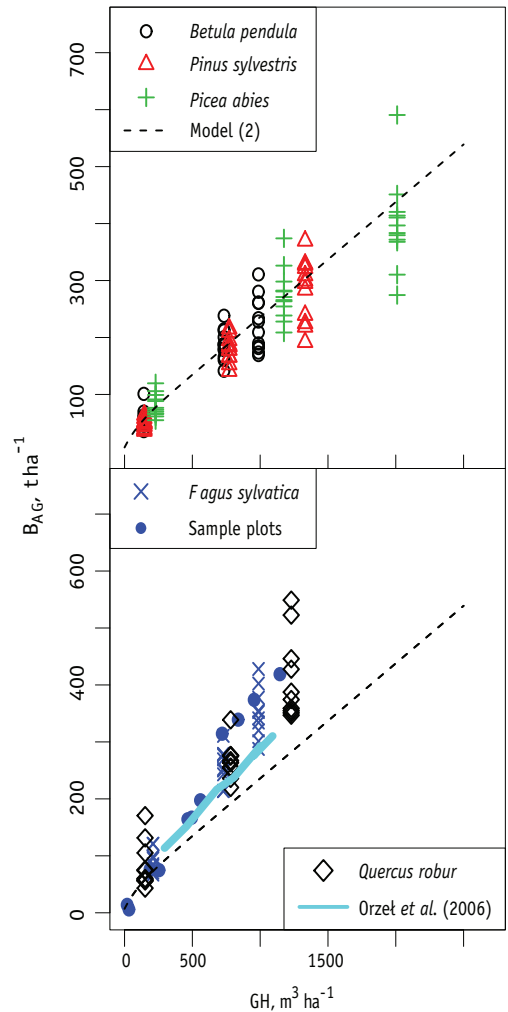


Figure 4. Biomass was predicted with country specific models for five species each forming three stands (Neumann *et al.*, 2016) characterized by different *D*. Sample plots are beech forests from Bartelink (1997) and Cannell (1982).

Joonis 4. Erinevate maade biomassimudelitega (Neumann *et al.*, 2016) saadud hinnangud viie puuliigi kolmele erineva rinnasdiameetriga puistule.

had more biomass for *GH* due to denser wood. *Fitzroya cupressoides* forests from Alerce Costero National Park (Urrutia-Jalabert *et al.*, 2015) had less biomass than predicted by the model (2) due to smaller wood density. After stand form factor conversion the difference in forest biomass in

respect to GH is mainly determined by the organic matter density. For example, the mean wood density in old-growth Atlantic forest sample plots 1, 2 and 3 was 0.655, 0.661 and 0.711 g/cm^3 and the differences of aggregated tree biomass from the stand-level model (2) were respectively 41 t/ha, 39 t/ha and 69 t/ha (Figure 5).

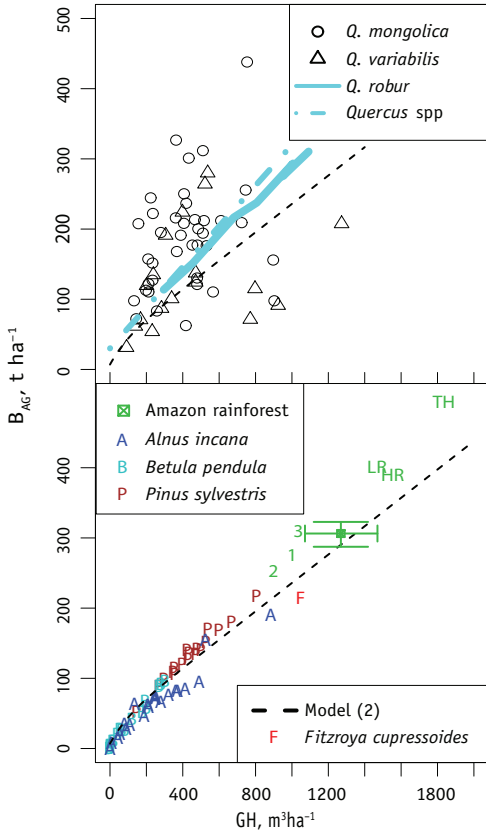


Figure 5. Upper: Oak forest data from Li *et al.* (2012), Orzet *et al.* (2006) model for *Quercus robur* and Cannell (1984) model for *Quercus* spp. Lower: The green point with confidence limits is an old-growth Amazon rainforest (Rice *et al.*, 2004). "1", "2", "3" stand for old-growth Atlantic forest; HR is high restinga, LR is low restinga and TH is tahuampa forest. "F" is plot AC1 from Urrutia-Jalabert *et al.* (2015).

Joonis 5. *Kirjanduses avaldatud puistute maapealse osa biomassi võrdlus mudeliga (2).*

The third set of biomass data was from Estonia and Poland. The model (2) underestimated biomass for Scots pine stands by 10.5 t/ha, overestimated biomass in young silver birch stands by 4 t/ha and overestimated biomass in grey alder stands by 12.8 t/ha. The biomass of grey alder stands had a declining trend with increasing GH except for two sample plots with biggest biomass (Figure 5). Since wood density for the stands was independent on stand structure variables (Uri *et al.*, 2014), the smaller gain is probably caused by changes in form factor. Similar decreasing trend of form factor with increasing forest biomass was observed in FGN forests. Compared to the generic model (Table 1) silver birch model (Table A5.1) predicts more biomass compared to grey alder model for the same GH . The difference is determined mainly by wood density and in some extent by form factor of grey alder and silver birch stands.

Discussion

We estimated parameters for a linear regression model in similar to Cannell (1984) by using biomass data from literature. Wulder *et al.* (2008) constructed a linear biomass model $B_{AG} = 29.2883 + 0.4123V$ for Canada based on stem merchantable volume and presented also individual models for deciduous, coniferous and mixed forests. Wulder *et al.* (2008) do not give confidence intervals for the model parameter estimates. If we assume a mean stem form factor $F \approx 0.5$ (Krigul, 1972; Tappo, 1982) and substitute $GH = V/F$, then the literature data-based model (1) and the model of Wulder *et al.* (2008) are similar. The simple linear model (1) overestimated biomass in young stands, but this was compensated by adding a start correction component and the model was used in further analysis. Biomass measurement data from Estonia and Poland were used to estimate model (2) parameters for grey alder stands

and silver birch stands. The models are convenient to use in practice, since G and H are easy to measure standard variables in forest inventories. A possible application of the proposed generic model could be estimating biomass using remotely sensed tree height (Lefsky, 2010; Simard *et al.*, 2011) and basal area from simple field assessments using for instance Bitterlich samples (Bitterlich, 1948), aggregated and gap-filled forest inventory data (Moreno *et al.*, 2016) or assumptions using management plans or yield tables. However, the stand-level GH -based model structure may need to be enhanced in next studies to explicitly include density (g/cm^3) and stand form factor for better local estimates. The species independent GH -based generic model may yield biased estimates when applied in forests where wood density or form factor differs systematically from the model (2) data set.

The generic model provided a good basis to verify Scandinavian single-tree biomass models in Estonia using forest inventory data from Estonian Network of Forest Research sample plots. From the two examined tree biomass models Repola (2008, 2009) gives slightly smaller estimates compared to Marklund (1988) model. The data for Marklund (1988) model are from 1983 and Repola models (2008, 2009) are based on data from years 1983–2003. With the 20 years, growth conditions are changed due to rising CO_2 level and global warming influencing northern latitudes (Myneni *et al.*, 1997) and the Marklund (1988) models may give already biased estimates in Estonian forests. In similar to boreal forests, old-growth tropical forest trees may be growing also more in the last decades, increasing net primary productivity and altering forest dynamics (Lewis *et al.*, 2009; Brienen *et al.*, 2015), although it is controversial whether it is due to CO_2 level or past disturbances (Clark, 2004; Muller-Landau, 2009) and the rate of growth increase has decreased after 2000 in Amazon forest (Brienen *et al.*, 2015). Probably, the param-

eters for tree-level regression models must be estimated based on new samples to account for changed growth conditions if the models are used for change detection in biomass accounting.

Two competition indices gave somewhat controversial results on dependence of biomass accumulation as a function of competition between trees at stand level when above-ground biomass was used as dependent variable. One reason may be that the FGN-forests are managed and the current structure of the forests does not always reflect past influences of competition. Another reason could be that the CI_{Hegyi} was a function of mean distance between trees, but CI_{LTJ} is dependent mainly on basal area. According to CI_{Hegyi} the more recent model from Repola (2008, 2009) depends more on competition between trees than the older biomass model from Marklund (1988) which seems not to have so much feedback from competition when looking at the difference of aggregated tree biomass from model (2) $B_{AG(2)}$. The shape of the relationship depends somewhat on the reference model, however, the model (2) or the original model of Cannell (1984) as reference give similar results. However, the stronger dependence of $B_{AG(R)}$ on competition was not so evident when absolute values of biomass were analysed. Even more, according to CI_{LTJ} competition described more variability of $B_{AG(M)} - B_{AG(2)}$. However, both of the biomass models produced smaller biomass estimates for stands where self-thinning process was weaker according to Nilson's CI_{LTJ} index. This means for forest management perspective that less trees must be harvested in thinning cuttings to retain sustainable rate of competition for the forest stands where biomass storage is the main purpose. Similar relationship was presented by Luysaert *et al.* (2008) in their Figure 2 regarding stand density and biomass. The share of tree crown mass from total above-ground biomass is also an indicator of competition. Open grown trees have wider crowns than closed canopy

trees (Hasenauer, 1997) as a result of smaller competition. Kilpeläinen *et al.* (2010) analysed biomass data of different Norway spruce provenance clones and found that Repola (2009) model had smaller estimation error compared to Marklund (1988) model which substantially overestimated biomass of branches.

While trees grow, each individual requires more space to survive and mortality (Laarmann *et al.*, 2009) decreases the number of trees per unit area. When the trees grow taller, stand form factor decreases. Both processes influence biomass accumulation in forest stand per unit area. Stand form factor can somewhat be compensated by the increased amount of branches, however, trees with smaller branches and faster self-pruning have better stem-wood quality. Little is known about age dependence of wood density change which may compensate the mortality and forest form factor decrease in some extent. The changes are related to cambial age, plant water distribution maintenance and creating an optimal mechanical structure (Lachenbruch *et al.*, 2011) which results in smaller tree ring width (MacPeak, 1990) and denser outerwood compared to corewood. Wood density is dependent also on site fertility and denser wood e.g. in Scots pine forests can be found on sites of medium fertility (Kask & Pikk, 2009).

Biomass density and tree form factor are accounted directly or indirectly in local allometric regression models, since the properties of sample trees are reflected also in the model parameters. This may be the reason why allometric models of tree biomass from different countries give substantially variable estimates for a tree (Neumann *et al.*, 2016). However, there is no reason why a tree growing near the border of two countries shall have different biomass estimates depending on country. This study showed that the differences between the country-specific tree biomass models propagate further to stand level and the range of estimated biomass is wide (± 30 –50% of mean)

for a stand. Compared to the other models used in Europe, Repola (2008, 2009) models give systematically smaller above-ground biomass estimates for big trees in FORMIT stands. In some cases, a biomass model for individual tree component may fail as appeared with branch mass models used for Norway spruce in Spain. As expected with the denser wood, FORMIT oak and beech stands had more biomass per unit area than predicted by the generic model. However, there was still substantial scatter in biomass estimates made with the country-specific models which can partly explained with the influence of data used for developing the regression models.

The fit of biomass of the simulated stands with the generic model (2) could probably be improved by accounting for possible differences in form factor of the European models. However, not all of the biomass models were based on volume to mass conversion and we did not have volume models for all the countries to estimate form factor for the trees in a uniform manner.

The generic model was well in agreement with data from tropical forests after transforming stand form factor where necessary data for the conversion were available. On the other hand, we found that the stand-level model (2) is not well applicable for the forest data where only arithmetic mean estimates for D and H are available as appeared with oak stand data from Korea. However, we do not know the form factor for each individual stand shown in Li *et al.* (2012). It is also possible that stand form factor, wood density and proportion of bark in oak stands have a large natural variation in Korea. While such published biomass datasets are still informative for further studies, it is recommended to include the descriptive forest variables (V , F , G , N , H_{Lorey}) that allow to apply stand level GH -based models. On the other hand, Chave *et al.* (2015) published a complete database of single tree measurements used to construct pan-tropical biomass model. A similar database with tree biomass sam-

ples from Europe could be useful to improve and harmonize carbon calculation methods.

Conclusions

- If growth conditions change, then single tree biomass models based on more than 20 years old data may be outdated.
- More biomass per unit area basis can be stored into forest stand when competition between trees is strong. Thinning cuttings must be conservative in the forests where carbon storage is main target and aimed to stipulate growth of species with denser wood or longer biological age.
- The introduced generic model allows biomass estimates independent on species using remote sensing data (e.g. height from airborne laser scanning data) and/or simple field measurement or assumptions (e.g. basal area).
- A common database of forest biomass measurements from Europe in similar to pan-tropical tree measurement data may be helpful harmonise carbon accounting methods.

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Appendix A1. Description of stands used for model construction and analysis.

Lisa A1. *Mudeli koostamiseks kasutatud empiirilise andmestik.*

Table A1.1. Description of stands used to estimate parameters for above-ground biomass models (1) and (2). For some stands particular details were collected from several publications.

Tabel A1.1. *Puistute andmed, mille põhjal hinnati mudelite (1) ja (2) parameetrid.*

Source	Species Puuliigid	Count Arv	<i>N</i> (trees/ha)	<i>D</i> (cm)	<i>H</i> (m)	<i>B</i> _{AG} (t/ha)
Uri et al. (2012)*, Varik et al. (2009)	<i>Betula pendula</i>	7	305–100000	1.3–31.0	3.4–30.1	25.7–220.5
Agarmaa (2015) ¹	<i>Pinus sylvestris</i>	3	620–1020	19.0–25.0	20.0–24.0	111.5–217.7
Kadak (2015), Möll (2015) ²	<i>Betula pubescens</i>	5	768–2752	7.9–17.9	12.0–19.3	47.4–122.0
Võsu (2012), Uri et al. (2014)	<i>Alnus incana</i>	11	691–6761	7.4–24.8	11.6–23.1	56.1–184.4
Johansson (2007)	<i>Betula</i> spp.	16	1479–5917	2.7–7.0	4.4–8.2	14.3–63.5
Vares et al. (2004a, 2004b)	<i>Alnus glutinosa</i>	5	1530–7222	7.6–13.9	9.8–15.1	80.2–100.6
Aosaar & Uri (2008)	Betulaceae	4	4850–35600	1.7–4.7	4.4–6.6	15.3–35.4
Johansson (2013)	Hybrid aspens	24	378–2374	8.5–24.4	10.4–25.8	42.3–219.3
Johansson (1999a)	<i>Populus tremula</i>	11	5964–46150	1.9–9.2	3.6–15.8	14.3–162.4
Mikšys et al. (2007) ³	<i>Pinus sylvestris</i>	5	727–2893	6.8–23.2	5.1–20.9	40.0–157.0
Johansson (1999b) ⁴	<i>Picea abies</i>	32	1133–7600	3.2–21.2	4.1–23.4	6.0–237.4
Bartelink (1997)	<i>Fagus sylvatica</i>	6	360–9920	3.1–27.9	3.5–22.5	5.9–166.7
Tobin & Nieuwenhuis (2007)	<i>Picea sitchensis</i>	6	767–2533	6.0–32.0	3.5–26.7	23.3–403.9
Qiong et al. (2011)	<i>Larix olgensis</i>	4	519–2220	9.0–25.7	8.9–26.9	59.2–176.6
Onyekwelu (2004)	<i>Gmelina arborea</i>	10	837–1275	18.4–79.8	15.2–34.3	83.2–394.9
Kilpeläinen et al. (2010)	<i>Picea abies</i>	20	2500 ⁵	7.7–14.9	6.8–13.1	43.1–164.9
Helmsaari et al. (2002)	<i>Pinus sylvestris</i>	3	432–7425	1.6–27.3	2.0–20.1	11.2–121.3
Finè et al. (2003) ⁶	Mixed coniferous	1	1585	10.0–30.8	8.1–20.8	154.0
Forrester et al. (2004)	<i>Eucalyptus</i> spp., <i>Acacia</i> spp.	2	1515	10.0–11.2	10.5–14.1	47.1–75.4
DeBell, et al. (1985)	<i>Eucalyptus</i> spp. mixed	3	2024–2200	9.5–15.0	14–19	37.6–95.3
Cannell (1982)	Various	31	123–12491	3.5–58.5	3.1–43.0	5.7–614.1
Onyekwelu (2007) ⁷	<i>Nauclea diderrichii</i>	9	443–667	9.6–29.3	12.9–28.5	32.5–287.7
Newman et al. (2006) ⁸	Mixed deciduous	2	950–1565	15.7–19.2	26.6–31.9	112.5–113.8
Johnston (1977)	<i>Pinus contorta</i>	2	1483–2764	9.2–9.9	12.8–14.1	48.5–73.7
Peichl & Arain (2006) ⁹	<i>Pinus strobus</i>	3	429–1492	15.6–34.6	9.1–20.1	65.5–176.0
Harrison et al. (2009) ¹⁰	<i>Pseudotsuga menziesii</i>	1	615	35.6	31.6	392.4
Park et al. (2005)	<i>Quercus</i> spp.	3	2450–3175	9.0–10.2	10.3–14.3	70.9–100.7
Son et al. (2004)	<i>Quercus</i> spp.	3	525–1475	12.1–16.6	17.7–18.7	119.9–224.1

* Excluding forest Kooraste2 due to an unrecoverable error causing extremely small wood density according to the data.

¹ Needle mass estimated with Marklund (1988) model for pine based on *d* and *h*.

² Foliage mass (except for stand JS191–40) estimated with Repola (2008) model.

³ Biomass was estimated from Figure 4.

⁴ Mean of model tree height for each stand was used instead *H*₄₀ index.

⁵ Number of trees per unit area was equal in all stands.

⁶ Basal area weighted mean *H* and quadratic mean *D* and total above-ground biomass was used.

⁷ Dominant height was used as estimate of *H*.

⁸ Woody biomass and yearly litter production was used as an estimate for above-ground biomass.

⁹ Description is for upper layer. Basal area of the lower tree layer (*D* < 9 cm) is included into analysis.

¹⁰ Sum of biomass fractions was used from Table 3 instead of given total.

Table A1.2. Overview of FGN forests (Kiviste *et al.*, 2014) used in this study. The range of stand inventory variables is in parentheses.

 Table A1.2. *Uurimuses kasutatud metsa kasvukäiguproovituukkide (Kiviste et al., 2014) valimi üldkirjeldus. Sulgudes on valimi haare.*

Main species Peapuuliik	Count Arv	A, years	N, trees/ha	H, m	G, m ² /ha	H _{100r} , m
<i>Populus tremula</i>	30	55 (35–75)	1337 (549–2920)	23.2 (18.1–28.9)	30.9 (18.6–42.5)	28.9 (22.9–33.5)
<i>Betula pendula</i>	127	55 (20–100)	1151 (168–3236)	21.7 (11.6–32.1)	28.1 (8.6–55.7)	27.8 (19.0–35.0)
<i>Picea abies</i>	193	55 (22–110)	940 (306–2334)	22.0 (11.2–33.0)	29.5 (8.1–48.8)	30.9 (21.5–36.6)
<i>Alnus glutinosa</i>	9	52 (31–70)	1230 (780–2467)	21.6 (18.7–23.5)	29.9 (21.1–37.6)	28.0 (26.1–32.5)
<i>Alnus incana</i>	3	40 (35–44)	2266 (1386–2769)	18.6 (16.4–21.2)	36.1 (34.8–38.1)	27.7 (26.8–29.2)
<i>Pinus sylvestris</i>	675	75 (20–245)	889 (163–3088)	23.1 (10.6–35.3)	31.0 (7.6–62.9)	27.8 (12.0–37.5)

Appendix A2. Form factor conversion for tropical forests.

 Lisa A2. *Troopikametsa vormiarvu teisendus.*

Trees in natural tropical forests tend to have umbrella-like crowns and biomass is not directly comparable by their GH with forests where crowns are more ellipsoidal. Similar problem occurs with substantial differences in stem form factor found e.g. in *Fitzroya cupressoides* forest. To make biomass estimates from rainforests comparable by their GH we applied stand form factor conversion based on volume. For example, by excluding δ from the pan-tropical model (Chave *et al.*, 2015), volume of stem and branches v_{s+b} for a tropical tree is obtained.

If the parameter for GH in the model (1) is expressed as $F\delta$, where F is the stand form factor and $\delta = 0.5 \text{ g/cm}^3$ is mean wood density, then the stand level model

(1) will be $B_{AG} = 25.40 + 0.422\delta GH$. Based on stand volume V_{s+b} and stand form factor $F_{conv} = V_{s+b}/GH$ of a natural tropical forest $GH_{gen} = F_{conv}GH/0.422$. For example, old-growth Atlantic forest sample plot 1 has $B_{AG} = 281 \text{ t/ha}$, $G = 27.9 \text{ m}^2/\text{ha}$, $H=23.5 \text{ m}$, $V_{s+b} = 430 \text{ m}^3/\text{ha}$ and $F_{conv} = 0.6555$. The stand has $GH = 656 \text{ m}^3/\text{ha}$ and corresponding $GH_{gen} = 1019 \text{ m}^3/\text{ha}$.

The data for old-growth Amazon rainforest (Rice *et al.* 2004) did not include live tree volume or wood density data and H was an estimate. By taking an estimate of $\delta = 0.64$ from Table 3 in Rice *et al.* (2004) and taking $H=45 \text{ m}$ and $G=28.2 \text{ m}^2/\text{ha}$ we obtain $F = 0.38$ which is close to the generic model.

Appendix A3. Wood density.

Lisa A3. Puidu tihedus.

The wood density (δ) values (Table A3.1) were used for Chave *et al.* (2015) model applied to trees growing on Estonian Network of Forest Research sample plots (Kiviste *et al.*, 2014) to estimate their above-ground biomass. The values were multiplied by

0.9 to account for bark, since stem diameters were measured over bark. For grey alder 10% of branches were considered (Uri *et al.*, 2014). The constant of 0.45 g/cm³ was used for all other tree species not listed in Table (A3.1).

Table A3.1. Wood density (δ).*Tabel A3.1 Puidu tihedus (δ).*

Tree species	δ	Source
<i>Puuliik</i>	g/cm ³	<i>Allikas</i>
<i>Pinus sylvestris</i>	0.453	Kask (2003)
<i>Picea abies</i>	0.420	Saarmann & Veibri (2006)
<i>Betula pendula</i>	0.650	Uri <i>et al.</i> (2012)
<i>Populus tremula</i>	0.470	Saarmann & Veibri (2006)
<i>Alnus glutinosa</i>	0.490	Saarmann & Veibri (2006)
<i>Alnus incana</i>	0.403	Uri <i>et al.</i> (2014)
<i>Tilia cordata</i>	0.470	Saarmann & Veibri (2006)
<i>Fraxinus excelsior</i>	0.640	Saarmann & Veibri (2006)
<i>Quercus robur</i>	0.650	Saarmann & Veibri (2006)

Appendix 4. Biomass predictor variables of FGN stands.

Lisa 4. Biomassi seosed takseertunnustega metsa kasvukäigu uuringute proovitükkidel.

Above-ground biomass for forest growth network plots (Kiviste *et al.*, 2014) was obtained by estimating first biomass for each tree with Repola (2008, 2009) or Marklund (1988) model and then aggregating tree biomass for each sample plot. Single tree biomass is usually estimated based on the tree

stem diameter. However, stand mean diameter is less informative predictor for biomass in Estonia compared to forest height or stand basal area (Figure A4.1). Mõistus and Lang (2015) found that stand mean height was good predictor for foliage mass in deciduous stands in Järvselja, Estonia.

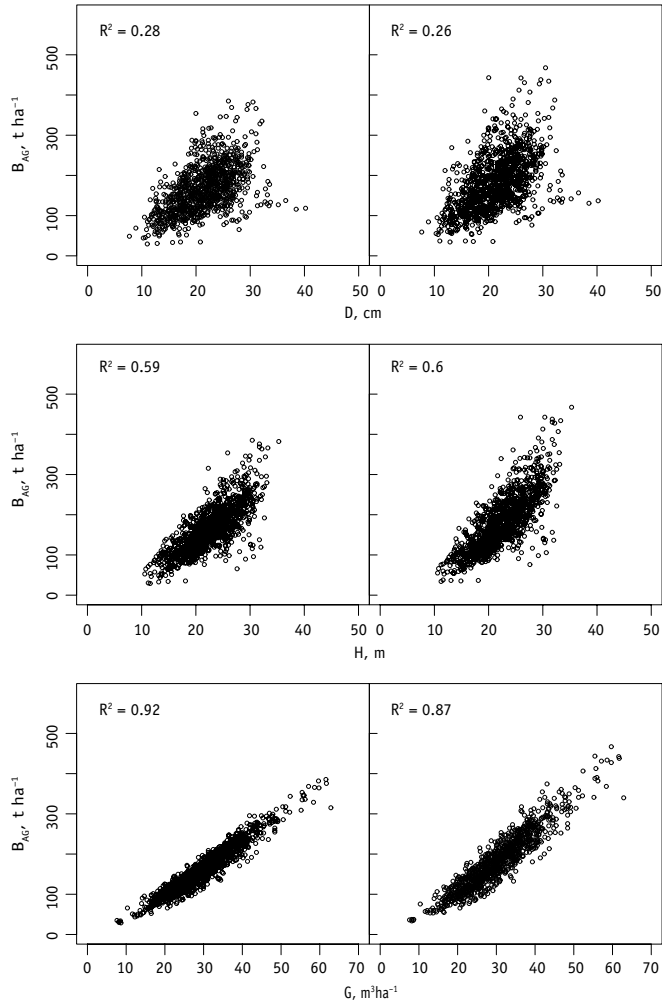


Figure A4.1. Above-ground biomass of forest growth study network (Kiviste *et al.*, 2014) plots as a function of stand mean breast height diameter, stand height and basal area. Left: Repola (2008, 2009), right: Marklund (1988).

Joonis A4.1. Puistu maapealse osa biomassi ja puistu takseertunnuste seosed metsa kasvukäigu proovitükkidel. Biomassi hindamiseks kasutati Repola (2008, 2009) (vasakul) ja Marklundi (1988) mudeleid (paremal).

Appendix 5. Species specific parameters for model (2).

Lisa 5. Puuliigiti lähendatud mudeli (2) parameetrite hinnangud.

We constructed new stand level above-ground biomass models (Table A5.1) for grey alder and silver birch. Data for grey alder stands are from Võsu (2012) and Uri *et al.* (2014). Data for silver birch stands are from Uri *et al.* (2012) (except forest “Kooraste2”), Varik *et al.* (2009) and Zasada *et al.* (2014). The parameter for start correc-

tion term of the grey alder model had a relatively large standard error and $p = 0.1$. However, to avoid biomass overestimation in young stands the start correction was included into the model. The model input values for G and H are regular measures obtained in forest inventory practice.

Table A5.1. Parameters for biomass model (2) in grey alder and silver birch stands. Standard error is in parentheses.

Tabel A5.1. Mudeli (2) parameetrite hinnangud hall-lepikutele ja arukaasikutele. Sulgudes on hinnangute standardvead.

Model component <i>Mudeli osa</i>	Model component values / <i>Hinnangud</i>	
	<i>Alnus incana</i>	<i>Betula pendula</i>
R_{adj}^2	0.9465	0.9868
Deg. of freedom	31	22
RSE (t/ha)	10.2	6.9
a1	19.73 (5.3)	22.37 (4.3)
a2	0.1842 (0.011)	0.2457 (0.010)
a3	-17.1 (10.2)	-17.3 (5.7)

Puistute maapealse osa biomassi hindamine rinnaspindala ja kõrguse järgi

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Kokkuvõte

Puistute biomassi salvestunud süsinik võimaldab mõneti leevendada fossiilsete kütuste kasutamisest tingitud kliima soojenemist (Nabuurs *et al.*, 2007), aga samasuguse puu kohta saadud biomassi hinnangud võivad riigiti oluliselt erineda (Neumann *et al.*, 2016). Tüüpiliselt kasutatakse üksikpuude mudelid (Ter-Mikaelian & Korzukhin, 1997; Zianis *et al.*, 2005; Henry *et al.*, 2013; Marklund, 1988; Repola, 2008, 2009; Chave *et al.*, 2005, 2015), mis põhinevad puu rinnasdiameetril d . Selliste mudelite põhjal jõudsid Stephenson *et al.* (2014) järeldusele, et suuremad puud salvestavad rohkem süsinikku. Puistut tervikuna vaadates tuleb aga arvestada puudevahelise konkurentsi ja suremusega (Enquist & Niklas, 2002; Nilson, 2005). Cannell (1984) näitas, et puistute maapealse osa biomassi B_{AG} (t/ha) hindamiseks sobib lihtne mudel, kus argumendiks on puistu kõrgus H (m) ja rinnaspindala (G) (m^2/ha). Käesolevas töös koostati kirjanduse (tabel A1.1) põhjal mudelid (1, 2, tabel 1, joonis 1), mida kasutati Marklund (1988) ja Repola (2008, 2009) üksikpuude biomassi mudelite hinnangute ja puudevahelise konkurentsi seoste uurimiseks metsa kasvukäigu uuringute proovitükkide (Kiviste *et al.*, 2014) mõõtmisandmetel (tabel A1.2). Analüüsi ka Neumann *et al.* (2016) töös kasutatud standardpuistutele erinevate maade mudelitega saadud biomassi hinnanguid, publitseeritud biomassi andmeid Korea tammikutest, Amazonase vihmametsast ja Tšiilist ning biomassi mõõtmisandmeid Brasiiliast, Eestist ja Poolast. Metsa kasvukäigu uuringute proovitükkidel arutati biomassi hinnangud ka mudeliga (4) Chave *et al.* (2015) tööst kasutades tabelis A3.1 toodud

puidu tihedusi. Puudevahelise konkurentsi hindamiseks arvutati 1) N_t puu alusel, mis asusid 8 m proovitüki piirist seespool, CI_{Hegy} (3), kus s_{ij} on puu j kaugus kuni 8 m olevast naabrist i ja 2) CI_{LTJ} (4) tuginedes puistu hõredusele L (Nilson, 2005) ja piirtihedusele L_{TJ} (Sims *et al.*, 2009).

Selgus, et puistu keskmine kõrgus on Eestis seotud maapealse osa biomassiga paremini kui keskmine tüveläbimõõt D (joonis A4.1). Marklundi (1988) mudelitega saadi süstemaatilisel suuremad biomassi hinnangud $B_{AG(M)}$ kui uuemate Repola (2008, 2009) mudelitega ($B_{AG(R)}$) (joonis 2). Konkurentsiindeksid kirjeldasid eraldi võetuna küll üsna suure osa biomassi variatsioonist (joonis 3, tabel 3), kuid lisaks puistu rinnaspindalale ja kõrgusele nad mudelit oluliselt ei paranda. Puistu hõredusel põhineva konkurentsiindeksi CI_{LTJ} järgi ilmnes aga selgelt, et võrreldes üldise keskmisega on rohkem biomassi puistutes, kus puudevaheline konkurent on suurem.

Üldist mudelit (2) kasutades selgus ka, et erinevate Euroopa maade biomassi hindamise meetodite erinevus ilmneb selgelt ka puistute tasemel (joonis 4). Mudeli (2) abil ei saanud hästi kirjeldada Korea tammikute biomassi (joonis 5), mille vormiarvu ja puidu tiheduse kohta polnud kahjuks konkreetseid andmeid. Samas kirjeldas mudel hästi Brasiilias, Eestis ja Poolas mõõdetud puistute biomassi (joonis 5). Eesti ja Poola andmetel lähendati mudelile (2) eraldi parameetrid hall-lepinkute ja aru-kaasikute jaoks (tabel A5.1).

Kokkuvõtteks: 1) teistsugustest kasvutingimustest pärinevate puude mõõtmisandmetel koostatud biomassimudelid (vanad mudelid) võivad anda süstemaatilise

veaga hinnanguid, 2) puistutes, mille eesmärgiks on süsiniku salvestamine, peab puudevaheline konkurents olema suur ning harvendusraieid tuleks kasutada ainult suurema puidutihedusega või suurema bioloogilise vanusega puuliikidele

eelise andmiseks, ja 3) puude biomassi mõõtmisandmed Eestist ja Euroopast tuleb koondada mudelite arendamiseks ja metoodikate sarnastele alustele viimiseks ühtsesse andmebaasi nagu on tehtud troopikapuude andmetega.

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