

Article



# **Evaluation of Stand Biomass Estimation Methods for Major Forest Types in the Eastern Da Xing'an Mountains, Northeast China**

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Abstract: Currently, forest biomass estimation methods at the regional scale have attracted the greatest attention from researchers, and the development of stand biomass models has become popular a trend. In this study, a total of 5074 measurements on 1053 permanent sample plots were obtained in the Eastern Da Xing'an Mountains, and three additive systems of stand biomass equations were developed. The first additive system (M-1) used stand variables as the predictors (i.e., stand basal area and average height), the second additive system (M-2) utilized stand volume as the sole predictor, and the third additive system (M-3) included both stand volume and biomass expansion and conversion factors (BCEFs) as the predictors. The coefficients of the three model systems were estimated with nonlinear seemingly unrelated regression (NSUR), while the heteroscedasticity of the model residuals was solved with the weight function. The jackknifing technique was used on the residuals, and several statistics were used to assess the prediction performance of each model. We comprehensively evaluated four stand biomass estimation methods (i.e., M-1, M-2, M-3 and a constant BCEF (M-4)). Here, we showed that the (1) three additive systems of stand biomass equations showed good model fitting and prediction performance, (2) M-3 significantly improved the model fitting and performance and provided the most accurate predictions for most stand biomass components, and (3) the ranking of the four stand biomass estimation methods followed the order of M-3 > M-2 > M-4 >M-1. Our results demonstrated these additive stand biomass models could be used to estimate the stand aboveground and belowground biomass for the major forest types in the Eastern Da Xing'an Mountains, although the most appropriate method depends on the available data and forest type.

Keywords: forest inventory; stand biomass; additive equations; nonlinear seemingly unrelated regression

# 1. Introduction

Among the studies on global climate change and the carbon cycle, research on the quantity, distribution, and dynamics of forest carbon stocks is popular and remains a high priority for predicting the growth and yield of forests [1–3]. Since the carbon concentrations in a tree or stand components are relatively constant (approximately 50%), most studies focus on forest biomass estimations rather than carbon storage estimations. Thus, the calculation of accurate forest biomass estimations has become one of the most crucial steps for successfully implementing the Reducing Emissions from Deforestation and Forest Degradation (REDD+) project as well as for the conservation and enhancement of forest carbon stocks and the sustainable management of forests. These initiatives provide a framework that benefits developing countries by rewarding them financially to reduce carbon emissions [4,5]. To date,

different countries have used various methods for their national-scale carbon assessments, which differ in the degree of accuracy and the amount of field inventory [6–10].

Forest stand biomass can be estimated at either the tree or stand level. At the tree level, the most commonly used method is the allometric equation, which estimates tree biomass using simple tree variables, such as the tree diameter at breast height (D) and/or tree height (H) [11–14]. Tree allometric equations require inventories that have high cost and time requirements, although they yield high accuracies for estimating tree biomass. The sum of the biomasses of individual trees provides the biomass for a plot or stand. Alternatively, the stand biomass can be estimated using either stand biomass models or biomass expansion and conversion factors (BCEFs), which represent the ratio of the stand biomass to stand volume. Historically, most biomass studies in the literature have focused on tree biomass models, while efforts related to stand biomass models have been limited or lacking [6,7,9,10,15]. For the stand biomass estimations, some researchers believe that the use of stand biomass models can more easily estimate the stand biomass and prevent having to address complex error propagation procedures at different spatial scales [6,7].

In recent years, with the rapid accumulation of forest biomass data throughout the world, researchers have attempted to improve forest biomass estimations and proposed various stand biomass estimation methods [7,10,16–18]. Studies have shown that stand biomass is closely related to some easily measured stand variables, such as the quadratic mean diameter, average height, and basal area of the stand [7,8,19]. In addition, the total and individual components of stand biomass have a strong correlation with stand volume, which has been used as a predictor in stand biomass models [20–22]. Recent methodology guidelines from the IPCC [23] provided a set of species-specific default values for BCEFs. It is well known that using a constant BCEF to quantify stand biomass produces certain biases or errors because BCEFs vary depending on the growth conditions and stand development stage, such as the stand age, stand size and stand density [15,16,21]. Studies have noted that BCEFs are not constant, and linear models have been established between stand biomass and volume [20,24,25]. Other researchers constructed nonlinear models (e.g., hyperbolic functions, reciprocal equations, and power functions) to express the relationships between stand biomass and BCEFs [7,10,15–17,26]. Overall, using stand biomass models with stand volume as the sole predictor or expanding the stand volume with an available BCEF is a relatively simple method for estimating stand biomass [7]. However, few comparative evaluations have been performed for different stand biomass estimation methods in the literature, especially for predicting stand biomass across a large geographic region.

When more than one biomass component is found in the same sample plot, the stand biomass equation is used to fit the total and component biomass data simultaneously, which explains the inherent correlations among the biomasses of the stand components in the same sample plot [6,12,13]. Consequently, the sum of stand biomass predictions from the component biomass models and the total biomass model are the same. To achieve the additivity of stand biomass equations, various parameter estimation methods and model specifications are used in linear and nonlinear models [11,12,27–29]. Among these methods of parameter estimation, nonlinear seemingly unrelated regression (NSUR) and seemingly unrelated regression (SUR) are the most widely used. An advantage of SUR and NSUR is the low variance of the total stand biomass model because of their own predictor variables and weighting function account for heteroscedasticity, which makes SUR and NSUR popular methods for parameter estimation in nonlinear and linear stand biomass equations [11–14,30]. Although several researchers have proposed the inclusion of additivity, it has often been ignored in some stand biomass models [8,10].

Additionally, similar to tree biomass models, stand biomass models commonly show heteroscedastic model residuals. To overcome the heteroscedasticity of the stand biomass model residuals, logarithmic transformation or a weighted regression should be performed before the construction of each carbon model [12,13]. To acquire an ideal result from logarithmic regression, a correction is necessary after the antilog transformation, i.e., the predicted values are multiplied by a correction factor [31,32]. However, when determining the total and component equations

of stand biomass, after applying the correction factor to the logarithmic equations of the additive system, realizing additivity is difficult [32]. Thus, the weighted regression successfully overcomes the heteroscedasticity of the total and component biomass model residuals in an additive system [12,13].

Asia is one of three primary locations of temperate mixed forests worldwide (i.e., northeastern North America, Europe, and eastern Asia), and they are mainly distributed in the forest regions of northeastern China. Chinese temperate forests are widely distributed in the Eastern Da Xing'an Mountains. These temperate forests play a crucial role in the Chinese national carbon budget and climatic system. Unfortunately, the forest resource inventory data in this region have not been fully utilized, and only limited stand biomass models are available. Thus, the objectives of this study were to (1) develop three alternative additive systems of stand biomass equations (i.e., stand biomass models using stand variables (M-1), stand biomass models using stand volume (M-2), and stand biomass models using both stand volume and BCEF (M-3)) for estimating the stand biomass of major forest types (including white birch forest, larch forest, poplar-birch forest) in the Eastern Da Xing'an Mountains, Northeast China, (2) use the jackknife method to validate the performance of those stand biomass models, and (3) evaluate the predictive ability of four alternative methods (i.e., M-1, M-2, M-3, and constant BCEF (M-4)) to estimate the total and component biomasses in each forest stand.

# 2. Materials and Methods

#### 2.1. Study Area and Data

The Eastern Da Xing'an Mountains in Northeast China are defined as in our previous study [14], and they cover an area of 83,517 km<sup>2</sup> (Figure 1). The terrain, soil, and climate of the region were described in Dong et al. [14] and, thus, are not detailed here. According to the classification standard of forest types in the Eastern Da Xing'an Mountains, the forest types in this study were mainly divided as follows: white birch forest, larch forest, poplar-birch forest, deciduous broadleaf mixed forest, coniferous and broadleaf mixed forest, and coniferous mixed forest. Among these forest types, white birch forest have the largest proportions of the forest area and stand volume in the Eastern Da Xing'an Mountains, occupying 37.44% and 32.93% of the forest area and 41.42% and 28.56% of the forest volume, respectively.



Figure 1. Geographical location of the study area in the Eastern Da Xing'an Mountains, Northeast China.

Our stand biomass data were collected from 1,053 permanent sample plots in the National Forest Continuous Inventory (NFCI) for the region. Most of the initially established plots were remeasured approximately four times (1990–2010, 5-year intervals), providing a total of 5074 measurements (Table 1). Those permanent sample plots were located throughout the species distributions across the Eastern Da Xing'an Mountains (Figure 1), which were set at the cross points of the kilometer networks (8 × 8 km). The sample plots were square with a size of 600 m<sup>2</sup>. The diameter at breast height (*D*) of the tree for each sample plot was measured at  $D \ge 5$  cm by the diameter tape. The average tree height in each sample plot was measured using a Blume-Leiss hypsometer.

Forest Types	Measurements	Statistics	Dq	$H_a$	G	V	N	$W_r$	$W_s$	W <sub>b</sub>	W <sub>f</sub>
		Min	5.3	5.0	0.5	1.7	200.0	0.49	1.12	0.10	0.05
Mathita himle famat	0(2	Max	23.2	24.0	30.8	226.0	3533.0	52.49	137.80	29.84	5.57
Forest Types         Measurements         Statistics $D_q$ $H_a$ $G$ $V$ $N$ $W_r$ White birch forest $P63$ Min         5.3         5.0         0.5         1.7         20.0         0.49           Max         23.2         24.0         30.8         226.0         353.0         52.49           Mean         11.1         11.9         11.5         68.6         122.67         15.18           SD         3.3         3.4         6.5         43.4         684.0         9.46           Min         6.0         5.0         0.6         2.3         20.00         0.46           Max         36.7         32.0         39.6         340.0         395.0         96.33           Max         36.7         32.0         39.6         340.0         395.0         96.33           Max         36.7         32.0         39.6         340.0         395.0         96.33           Max         31.9         14.0         15.6         107.6         1130.9         25.91           SD         4.1         4.1         7.5         57.6         62.4         15.66           <	15.18	40.51	7.58	1.76							
		SD	3.3	3.4	6.5	43.4	684.0	9.46	25.38	$\begin{array}{c ccccccccccccccccccccccccccccccccccc$	1.10
		Min	6.0	5.0	0.6	2.3	200.0	0.46	1.36	0.16	0.09
Laugh found	1740	Max	36.7	32.0	39.6	340.0	3950.0	96.33	203.40	30.86	6.35
Larch lorest	1749	Mean	13.9	14.0	15.6	107.6	1130.9	25.91	62.86	7.74	2.10
		SD	4.1	4.1	7.5	57.7	662.4	15.66	35.79	4.78	1.06
Poplar-birch forest	293	Min	5.3	5.1	0.7	2.6	200.0	0.47	1.66	0.12	0.05
		Max	24.1	23.2	36.2	288.2	3433.3	57.59	168.53	31.49	5.48
		Mean	12.1	13.1	16.9	111.9	1543.8	19.25	62.36	10.20	2.24
		SD	3.8	4.1	7.5	58.8	712.4	10.54	32.25	6.55	1.16
Deciduous broadleaf	501	Min	5.4	5.0	0.8	3.2	200.0	0.83	1.95	0.20	0.08
		Max	27.0	23.1	37.3	298.8	3150.0	59.91	175.93	45.44	5.72
mixed forest		Mean	12.7	11.1	14.0	78.5	1207.7	18.65	54.41	13.18	2.31
		SD	4.0	3.6	6.4	43.5	603.4	9.78	28.67	9.45	1.19
Coniformus and		Min	6.9	5.0	1.0	4.5	217.0	0.88	2.40	0.29	0.13
broadloaf mixed	1263	Max	27.0	32.3	36.5	291.6	3350.0	60.32	170.78	27.78	6.70
forest	1205	Mean	12.8	12.8	16.1	106.0	1365.2	24.10	62.41	9.90	2.37
Iotest		SD	3.0	3.5	6.6	47.8	683.5	11.36	28.65	5.13	1.04
		Min	6.3	5.4	1.2	10.3	200.0	0.55	1.90	0.30	0.18
Coniferous mixed	205	Max	33.1	23.6	33.3	288.8	3933.0	69.77	158.62	22.14	8.50
forest	505	Mean	14.1	13.7	17.2	123.2	1219.7	21.92	64.87	8.99	3.47
		SD	4.4	3.9	6.8	56.8	601.3	11.62	$\begin{array}{cccccccccccccccccccccccccccccccccccc$	1.57	

**Table 1.** Summary statistics of stand variables and biomass components for the six forest types in the Eastern Da Xing'an Mountains, Northeast China.

where  $D_q$  represents the quadratic mean diameter,  $H_a$  represents the average tree height, G represents the stand basal area per hectare, V represents the stand volume, N represents the number of trees per hectare,  $W_r$  represents the stand root biomass,  $W_s$  represents the stand stem biomass,  $W_b$  represents the stand branch biomass, and  $W_f$ represents the stand foliage biomass.

In the NFCI, the specific calculation methods for the stand variables are as follows: (1) the average tree height ( $H_a$ ) of the stand was the average height of 3–5 standard trees for dominant species in a plot, (2) the quadratic mean diameter ( $D_q$ ) of the stand was calculated using the formula  $\sqrt{\sum D_i^2/n}$ , (3) the stand basal area per hectare (*G*) was the accumulation of all individual tree basal areas, (4) the number of trees per hectare (*N*) was the ratio of the number of trees in a plot to the plot area, and (5) the individual tree volume was calculated by the volume equation based on the *D* for only the Eastern Da Xing'an Mountains (because the tree height was not measured in the NFCI), and the accumulation of all individual tree volumes for each sample plot was the stand volume (*V*).

We previously developed species-specific tree biomass allometric equations with only tree D as the predictor for the tree total and component biomass (i.e., the stand total biomass ( $W_t$ ), the stand root biomass ( $W_r$ ), the stand stem biomass ( $W_s$ ), the stand branch biomass ( $W_b$ ), and the stand foliage biomass ( $W_f$ )) [12,32,33], and they were applied to each tree within the permanent sample plots. The stand biomass (Mg·ha<sup>-1</sup>) was determined on an area basis for each sample plot. Thus, the descriptive statistics of the stand biomass components (i.e., root, stem, branch, and foliage) and stand variables (i.e.,  $D_q$ ,  $H_a$ , G, V, and N) were summarized for each forest type, as shown in Table 1.

#### 2.2. Stand Biomass Estimation Models

In this study, three alternative models for stand biomass estimation were proposed: (1) the stand variables were the predictors for estimating the stand total and component biomasses (namely, Method 1 or M-1), (2) the stand volume was the sole predictor for estimating the stand total and component biomasses (namely, Method 2 or M-2), and (3) both stand volume and appropriate biomass conversion and expansion factors (BCEFs) were used to estimate the stand total and component biomasses (namely, Method 3 or M-3). For the above three methods, the additive systems of biomass equations that consider the inherent correlations among the biomasses of the stand components in the same sample plot were defined to predict the stand total and component biomasses for each of the major forest types.

#### 2.2.1. Stand Biomass Models with Stand Variables (M-1)

The stand total and component biomasses (i.e., root, stem, branch, and foliage) were regressed on the stand variables using the following power function:

$$W_{i} = e^{\beta_{i0}} X_{1}^{\beta_{i1}} X_{2}^{\beta_{i2}} \dots X_{j}^{\beta_{ij}} + \varepsilon_{i}$$
(1)

where  $W_i$  represents the stand total biomass and the biomass of each component (*i* = branch (*b*), foliage (*f*), root (*r*), stem (*s*), and total (*t*)) at the stand level (Mg ha<sup>-1</sup>),  $X_j$  represents the stand variables (*j* = 1, ..., *k*), such as the stand basal area (*G*) and average height ( $H_a$ ),  $\beta_{ij}$  represents the parameters estimated by the model; and  $\varepsilon_i$  is the additive error term of the model.

A visual inspection of the stand component biomass data indicated that the stand component biomass could be well expressed by *G* and *H<sub>a</sub>*. Therefore, the equation  $W_i = e^{\beta_{i0}}G^{\beta_{i1}}H_a^{\beta_{i2}}$  was used to estimate the stand total and component biomasses for each of the major forest types.

According to the description by Parresol [28], the additive system of five equations with cross-equation error correlations for the stand total, root, stem, branch, and foliage biomass are listed as follows:

$$\begin{pmatrix}
W_r = e^{\beta_{r0}}G^{\beta_{r1}}H_a^{\beta_{r2}} + \varepsilon_r \\
W_s = e^{\beta_{s0}}G^{\beta_{s1}}H_a^{\beta_{s2}} + \varepsilon_s \\
W_b = e^{\beta_{b0}}G^{\beta_{b1}}H_a^{\beta_{b2}} + \varepsilon_b \\
W_f = e^{\beta_{f0}}G^{\beta_{f1}}H_a^{\beta_{f2}} + \varepsilon_f \\
W_t = W_r + W_s + W_b + W_f + \varepsilon_t
\end{cases}$$
(2)

where  $W_r$ ,  $W_s$ ,  $W_b$ ,  $W_f$ , and  $W_t$  represent the root, stem, branch, foliage, and total stand biomass (Mg ha<sup>-1</sup>), respectively.

2.2.2. Stand Biomass Models with Stand Volume (M-2)

The primary results indicated that there was a strong correlation between stand biomass and stand volume (*V*) [20–22]. Thus, the equation  $W_i = e^{\beta_{i0}} V^{\beta_{i1}}$  can be used to calculate the stand total and component biomasses for each of the major forest types. The additive system of five equations was specified as follows:

$$W_{r} = e^{\beta_{r0}} V^{\beta_{r1}} + \varepsilon_{r}$$

$$W_{s} = e^{\beta_{s0}} V^{\beta_{s1}} + \varepsilon_{s}$$

$$W_{b} = e^{\beta_{b0}} V^{\beta_{b1}} + \varepsilon_{b}$$

$$W_{f} = e^{\beta_{f0}} V^{\beta_{f1}} + \varepsilon_{f}$$

$$W_{t} = W_{r} + W_{s} + W_{b} + W_{f} + \varepsilon_{t}$$
(3)

The symbols used in the additive system are the same as in Equations (2).

2.2.3. Stand Biomass Models with Both Stand Volume and BCEFs (M-3)

In this study, the stand biomass estimation was also calculated using BCEFs, which are commonly defined as follows:

$$BCEF_i = \frac{W_i}{V} \tag{4}$$

where  $W_i$  represents the stand total biomass and the biomass of each component (*i* = branch (*b*), foliage (*f*), root (*r*), stem (*s*), and total (*t*)) at the stand level (Mg ha<sup>-1</sup>); and *V* represents the stand volume.

We attempted to elaborate multiple regression models for stand biomass and BCEF using the stand variables. The adjusted coefficient of determination ( $R_a^2$ ), root mean squared error (RMSE), and AIC revealed that models with one or two predictors were the best choices. For this reason, we decided to present only the following models (Equations (5)–(7)), which allowed for the computation of stand

biomass and BCEF using available stand variables. To evaluate the relationships between BCEF and stand variables, we used the following model types:

$$BCEF_{i} = f_{i}(X_{j}) = \beta_{i0}X_{j}^{\beta_{ij}} + \varepsilon_{i}$$
(5)

$$BCEF_{i} = f_{i}(X_{j}) = X_{j}/(\beta_{i0} + \beta_{ij}X_{j}) + \varepsilon_{i}$$
(6)

$$BCEF_i = f_i(X_j) = \beta_{i0} + \beta_{ij}/X_j + \varepsilon_i$$
(7)

where  $X_j$  represents the stand variable (j=1, ..., k), such as  $H_a$ ,  $D_q$ , and others.  $f_i(X_j)$  is the equation of BCEF<sub>i</sub>,  $\beta_{ij}$  represents the parameters estimated by the model; and  $\varepsilon_i$  is the additive error term of the model

Our results showed that the BCEF model types for each biomass component varied for different forest types (as listed in Table 5). In general, the additive system of five equations was specified as follows:

$$W_{r} = f_{r}(X_{j})V + \varepsilon_{r}$$

$$W_{s} = f_{s}(X_{j})V + \varepsilon_{s}$$

$$W_{b} = f_{b}(X_{j})V + \varepsilon_{b}$$

$$W_{f} = f_{f}(X_{j})V + \varepsilon_{f}$$

$$W_{t} = W_{r} + W_{s} + W_{b} + W_{f} + \varepsilon_{t}$$
(8)

The symbols used in the system are the same as in the above equations.

Due to the heteroscedasticity in the model residuals shown by the stand biomass data, a weighting function was defined and applied for each stand biomass model. Following previous applications for modeling residual heteroscedasticity, the variances, or the squares of residuals ( $\varepsilon^2$ ), in the *i*th observation were functionally related to other predictor variables, such as  $\varepsilon_i^2 = \sigma^2(x_i)^p$ , where  $x_i$  is the stand variable and  $\varepsilon_i$  is the unweighted model residual. Hence, we chose  $1/(x_i)^p$  as the weight function. In this study,  $1/G^p$  for M-1 and  $1/V^p$  for M-2 and M-3 were chosen as the weight functions, and *p* was confirmed based on each stand biomass model. In the computations, the weight function for heteroscedasticity  $1/G^p$  and  $1/V^p$  was multiplied and programmed using the PROC MODEL procedure in SAS by specifying resid. $W_i = \text{resid.}W_i/\sqrt{1/(x_i)^p}$  [12,13,34,35].

The above three additive systems of stand biomass in Equations (2), (3) and (8) fit the data of each forest type with NSUR under the SAS/ETS model [35].

# 2.3. Model Evaluation and Validation

The three additive systems (Equations (2), (3), and (8)) suit the entire data set and were tested with the jackknife technique, in which all the observations except one (sample size *N*-1) were used to construct the stand biomass equation, and the dependent variable for the excluded observation was predicted with the fitted model. The five statistics of each system equation based on the jackknifing technique were used to evaluate the fitting performance (adjusted coefficient of determination ( $R_a^2$ ) and root mean square error (RMSE)) and the predictive performance (mean prediction error (MPE), mean absolute error (MAE), and mean absolute percent error (MAE%)) of each stand biomass prediction equation. The mathematical expressions of the four statistics are as follows:

$$R_{a}^{2} = 1 - \frac{\sum_{i=1}^{N} (W_{i} - \hat{W}_{i})^{2}}{\sum_{i=1}^{N} (W_{i} - \overline{W})^{2}} \left(\frac{N-1}{N-p}\right)$$
(9)

RMSE = 
$$\sqrt{\frac{\sum_{i=1}^{N} (W_i - \hat{W}_i)^2}{N - p}}$$
 (10)

$$MAE = \frac{\sum_{i=1}^{N} |W_i - \hat{W}_{i,-i}|}{N}$$
(12)

$$MAE\% = \frac{\sum_{i=1}^{N} |W_i - \hat{W}_{i,-i} / \hat{W}_{i,-i}|}{N}$$
(13)

where  $W_i$  is the value of the *i*th observed stand biomass;  $\hat{W}_i$  is the *i*th stand biomass prediction from the model fit to all the data (sample size *N*),  $\overline{W}$  is the average value of the stand biomass,  $\hat{W}_{i,-i}$  is the prediction of the *i*th observation of the model fitted using *N*-1 observations excluding the usage of the *i*th observation; and *p* is the total number of model parameters.

# 2.4. Evaluation of Several Stand Biomass Estimation Methods

In this study, once the stand biomass estimate was calculated for each component, the component estimates were summed to produce the total stand biomass estimate. We used the following four methods to calculate the stand biomass of each tree component: (1) stand biomass models with the stand variables as the predictors (M-1), (2) stand biomass models with the stand volume as the sole predictor (M-2), (3) stand biomass models with stand volume and BCEF as the predictors (M-3), and (4) a constant BCEF (M-4), in which a constant BCEF value for each component for major forest types was applied to calculate the stand biomass of each component by Equations (4), and the total estimated stand biomass was the sum of the component estimates. The constant BCEF values of each component are listed for the major forest types in Table 2.

**Table 2.** Constant (average value) and standard deviation (SD) of biomass expansion and conversion factors (BCEFs) at the stand level (Mg·m<sup>-3</sup>) for the six forest types in the Eastern Da Xing'an Mountains, Northeast China.

Forest Types	Statistics	BCEFr	BCEFs	BCEFb	BCEF <sub>f</sub>
TATE is the instance of	Constant	0.2255	0.5944	0.1056	0.0257
white birch forest	SD	0.0214	0.0552	0.0346	0.0039
Laugh (anna)	Constant	0.2338	.338         0.5754         0.0699           0290         0.0409         0.0126           05755         0.5751         0.0557	0.0205	
Larch forest	SD	0.0290	0.0409	0.0126	0.0041
Poplar birch forest	Constant	0.1725	0.5651	0.0856	0.0202
i opiai-biten iorest	SD	0.0327	0.0529	0.0324	$\begin{array}{c ccccc} 0.1056 & 0.0257 \\ 0.0346 & 0.0039 \\ 0.0699 & 0.0205 \\ 0.0126 & 0.0041 \\ 0.0856 & 0.0202 \\ 0.0324 & 0.0040 \\ 0.1639 & 0.0305 \\ 0.0865 & 0.0076 \\ 0.0926 & 0.0229 \\ 0.0260 & 0.0040 \\ 0.0731 & 0.0299 \\ \end{array}$
Desidence has all of active different	SD         0.0327         0.0529           Constant         0.2438         0.6994	0.6994	0.1639	0.0305	
Deciduous broadlear mixed forest	SD	0.0512	0.1269	0.0865	0.0076
Coniformus and broadloof missed format	Constant	0.2265	0.5872	0.0926	0.0229
Conferous and broadlear mixed forest	SD	0.0269	0.0462	0.0260	BCEF <sub>f</sub> 0.0257 0.0039 0.0205 0.0041 0.0202 0.0040 0.0305 0.0076 0.0229 0.0040 0.0229 0.0040 0.0299 0.0097
Conifornation missed format	Constant	0.1775	0.5152	0.0731	0.0299
Confirerous mixed forest	SD	0.0410	0.0621	BCEFb         BCEFf           0.1056         0.0257           0.0346         0.0039           0.0699         0.0205           0.0126         0.0041           0.0856         0.0202           0.0324         0.0040           0.1639         0.0305           0.0865         0.0076           0.0926         0.0229           0.0260         0.0040           0.0731         0.0299           0.0130         0.0097	0.0097

An analysis of variance was used to test the differences between the four methods (treatment) for estimating stand biomass with the sample plots as the blocks. Furthermore, a percentage difference (MPD%) was defined to quantify the biases in the four methods used to estimate stand biomass.

$$MPD\% = \frac{\sum_{i=1}^{N} \left| \frac{W_i - \hat{W}_i}{W_i} \right|}{N} \times 100$$
(14)

where  $W_i$  is the value of *i*th observed stand biomass,  $\hat{W}_i$  is the *i*th stand biomass predicted by the four methods, and *N* is the sample size.

#### 3. Results

# 3.1. Model Fitting for Stand Biomass Models

The coefficient estimates, standard errors (SEs), and goodness-o*F*-fit statistics ( $R_a^2$  and RMSE) of M-1 (Equations (2)) are shown for the six forest types in Table 3. The results indicated that most of the biomass equations in M-1 fit the stand biomass data well, with  $R_a^2 > 0.70$  and RMSE < 20.0 Mg·ha<sup>-1</sup>. The model fit the stand total and stem biomass data the best, while relatively small  $R_a^2$  values were observed in the branch and foliage equations (Table 3).

We fit the M-2 (Equations (3)) for the six forest types. The results indicated that *V* had a good correlation with stand biomass. The stand total and stem biomass equations had larger  $R_a^2$  and smaller RMSE values than the stand foliage and branch biomass equations (Table 4). Compared with M-1, the M-2 for most forest types had large  $R_a^2$  and small RMSE values, e.g., the stand stem biomass equation of coniferous mixed forest presented a 12.6% increase in  $R_a^2$  and a 44.9% decrease in RMSE. When the biomass equation had the sole predictor variable *V*, the majority of stand biomass equations in M-2 had large  $R_a^2$  and small RMSE values (Table 4).

The constant values of BCEF<sub>r</sub>, BCEF<sub>s</sub>, BCEF<sub>b</sub> and BCEF<sub>f</sub> are listed for the six forest types in Table 2. There were some variations among the BCEF<sub>i</sub> values across those forest types, which explained why the BCEF values were considered stand dependent. We analyzed the relationship between the BCEF and stand variables. The results indicated that depending on the forest type and biomass component, the predictor variables  $D_q$  and  $H_a$  best interpreted the variations in BCEF. It was evident that the BCEF model types for each component were different for the different forest types (Table 5). Based on the BCEF models, we fitted the third additive systems using both stand volume (*V*) and BCEF (M-3, Equations (8)). For most forest types, the proportion of variation interpreted by the additive system with *V* and BCEF was analogous to that interpreted by the additive systems including stand variables, such as *G* and  $H_a$  or stand volume only (Tables 4–6).

Forest Types	Components	nts Equations	$\beta_{i0}$		$\beta_i$	1	$\beta_i$	2	$R_a^2$	RMSE	Weight Function	
<b>51</b>	1	1	Estimate	SE	Estimate	SE	Estimate	SE	u		0	
White birch forest	Root	$W_r = e^{\beta_{r0}} G^{\beta_{r1}} H_a^{\beta_{r2}}$	-0.3681	0.0211	1.0134	0.0038	0.2417	0.0107	0.9670	1.7191	G <sup>2.3111</sup>	
	Stem	$W_s = e^{\beta_{s0}} G^{\beta_{s1}} H_a^{\beta_{s2}}$	0.3658	0.0243	1.0138	0.0047	0.3378	0.0118	0.9720	4.2478	G <sup>1.8435</sup>	
	Branch	$W_b = e^{\beta_{b0}} G^{\beta_{b1}} H_a^{\beta_{b2}}$	-2.5802	0.0755	0.9553	0.0143	0.8883	0.0333	0.8398	2.1523	G <sup>1.3376</sup>	
	Foliage	$W_f = e^{\beta_{f0}} G^{\beta_{f1}} H_a^{\beta_{f2}}$	-3.0214	0.0379	0.9788	0.0075	0.4702	0.0173	0.9443	0.2592	G <sup>1.2728</sup>	
	Total	$W_t = W_r + W_s + W_b + W_f$	-	-	-	-	-	-	0.9644	7.7291	G <sup>1.7861</sup>	
	Root	$W_r = e^{\beta_{r0}} G^{\beta_{r1}} H_a^{\beta_{r2}}$	-0.9170	0.0387	1.0639	0.0075	0.4644	0.0166	0.8827	5.3645	G <sup>2.4562</sup>	
	Stem	$W_s = e^{\beta_{s0}} G^{\beta_{s1}} H_a^{\beta_{s2}}$	0.2552	0.0290	1.0538	0.0057	0.3689	0.0124	0.9254	9.7728	G <sup>2.3102</sup>	
Larch forest	Branch	$W_b = e^{\beta_{b0}} G^{\beta_{b1}} H_a^{\beta_{b2}}$	-1.8547	0.0417	1.1227	0.0073	0.3007	0.0197	0.8412	1.9034	G <sup>2.8988</sup>	
	Foliage	$W_f = e^{\beta_{f0}} G^{\beta_{f1}} H_a^{\beta_{f2}}$	-1.9338	0.0270	1.0276	0.0059	-0.0552	0.0114	0.9498	0.2383	G <sup>1.5773</sup>	
	Total	$W_t = W_r + W_s + W_b + W_f$	-	-	-	-	-	-	0.9175	16.3387	G <sup>2.3770</sup>	
	Root	$W_r = e^{\beta_{r0}} G^{\beta_{r1}} H_a^{\beta_{r2}}$	-0.6818	0.0709	1.0891	0.0145	0.2072	0.0361	0.8644	3.8810	G <sup>1.9913</sup>	
	Stem	$W_s = e^{\beta_{s0}} G^{\beta_{s1}} H_a^{\beta_{s2}}$	0.3915	0.0357	1.0531	0.0072	0.2872	0.0179	0.9645	6.0795	G <sup>1.9630</sup>	
Poplar-birch forest	Branch	$W_b = e^{\beta_{b0}} G^{\beta_{b1}} H_a^{\beta_{b2}}$	-2.2658	0.1651	1.0131	0.0442	0.6504	0.0669	0.7546	3.2449	G <sup>1.2065</sup>	
	Foliage	$W_f = e^{\beta_{f0}} G^{\beta_{f1}} H_a^{\beta_{f2}}$	-2.6787	0.0948	0.9712	0.0274	0.2830	0.0385	0.8820	0.3969	$G^{0.7684}$	
	Total	$W_t = W_r + W_s + W_b + W_f$	-	-	-	-	-	-	0.9376	12.4215	G <sup>1.5947</sup>	
	Root	$W_r = e^{\beta_{r0}} G^{\beta_{r1}} H_a^{\beta_{r2}}$	-0.0292	0.0457	1.1270	0.0111	-0.0225	0.0211	0.9011	3.0743	G <sup>2.0893</sup>	
Deciduous broadloaf	Stem	$W_s = e^{\beta_{s0}} G^{\beta_{s1}} H_a^{\beta_{s2}}$	0.7085	0.0336	1.1494	0.0084	0.0885	0.0151	0.9375	7.1641	G <sup>1.9423</sup>	
mixed forest	Branch	$W_b = e^{\beta_{b0}} G^{\beta_{b1}} H_a^{\beta_{b2}}$	-1.0325	0.1065	1.4481	0.0260	-0.1398	0.0511	0.5196	6.5472	G <sup>1.7915</sup>	
hixed forest	Foliage	$W_f = e^{\beta_{f0}} G^{\beta_{f1}} H_a^{\beta_{f2}}$	-2.0889	0.0478	1.1408	0.0119	-0.0519	0.0225	0.8434	0.4704	$G^{1.5845}$	
	Total	$W_t = W_r + W_s + W_b + W_f$	-	-	-	-	-	-	0.8838	16.2834	G <sup>1.9621</sup>	
	Root	$W_r = e^{\beta_{r0}} G^{\beta_{r1}} H_a^{\beta_{r2}}$	-0.2980	0.0460	1.0499	0.0097	0.2189	0.0179	0.8520	4.3714	G <sup>1.9153</sup>	
Coniforous and	Stem	$W_s = e^{\beta_{s0}} G^{\beta_{s1}} H_a^{\beta_{s2}}$	0.6282	0.0341	1.0590	0.0071	0.2181	0.0132	0.9244	7.8758	G <sup>1.9069</sup>	
broadleaf mixed forest	Branch	$W_b = e^{\beta_{b0}} G^{\beta_{b1}} H_a^{\beta_{b2}}$	-1.6087	0.0614	1.0813	0.0122	0.3442	0.0237	0.7140	2.7444	G <sup>1.4223</sup>	
bioucieur nixeu forest	Foliage	$W_f = e^{\beta_{f0}} G^{\beta_{f1}} H_a^{\beta_{f2}}$	-2.3012	0.0329	1.0264	0.0067	0.1207	0.0129	0.8902	0.3458	$G^{1.4523}$	
	Total	$W_t = W_r + W_s + W_b + W_f$	-	-	-	-	-	-	0.9097	13.6576	$G^{1.8568}$	
	Root	$W_r = e^{\beta_{r0}} G^{\beta_{r1}} H_a^{\beta_{r2}}$	-0.2841	0.1084	1.1484	0.0233	0.0343	0.0520	0.6721	6.6523	G <sup>2.8070</sup>	
	Stem	$W_s = e^{\beta_{s0}} G^{\beta_{s1}} H_a^{\beta_{s2}}$	0.3872	0.0814	1.1543	0.0196	0.1862	0.0363	0.8468	12.6796	G <sup>2.1331</sup>	
Coniferous mixed forest	Branch	$W_b = e^{\beta_{b0}} G^{\beta_{b1}} H_a^{\beta_{b2}}$	-1.3594	0.0849	1.0907	0.0209	0.1701	0.0373	0.8421	1.7045	$G^{2.0005}$	
	Foliage	$W_f = e^{\beta_{f0}} G^{\beta_{f1}} H_a{}^{\beta_{f2}}$	-1.7837	0.1034	0.8950	0.0262	0.1876	0.0434	0.7893	0.7194	G <sup>1.2489</sup>	
	Total	$W_t = W_r + W_s + W_b + W_f$	-	-	-	-	-	-	0.8420	19.2912	G <sup>2.3796</sup>	

**Table 3.** Model coefficient estimates, standard errors (SEs), model fitting statistics, and weight functions for the additive system of stand biomass equations using stand variables (namely, M-1) for the six forest types in the Eastern Da Xing'an Mountains, Northeast China.

Forest Types	Components	Equations	$\beta_i$	0	$\beta_i$	1	$R_{-}^{2}$	RMSF	Weight Function	
51	1		Estimate	SE	Estimate	SE	- n <sub>u</sub>	RIVIOL		
	Root	$W_r = e^{eta_{r0}} V^{eta_{r1}}$	-1.3278	0.0140	0.9580	0.0033	0.9799	1.3392	V <sup>1.5306</sup>	
White birch forest	Stem	$W_{ m s}=e^{eta_{ m s0}}V^{eta_{ m s1}}$	-0.4863	0.0118	0.9893	0.0026	0.9849	3.1148	V <sup>1.1589</sup>	
	Branch	$W_b=e^{eta_{b0}}V^{eta_{b1}}$	-2.7349	0.0371	1.1142	0.0083	0.8192	2.2864	V <sup>1.2267</sup>	
	Foliage	$W_f = e^{eta_{f0}} V^{eta_{f1}}$	-3.6432	0.0193	0.9928	0.0044	0.9444	0.259	V <sup>1.0907</sup>	
	Total	$W_t = W_r + W_s + W_b + W_f$	-	-	-	-	0.9783	6.0348	V <sup>1.3182</sup>	
	Root	$W_r = e^{\beta_{r0}} V^{\beta_{r1}}$	-1.7833	0.0146	1.0742	0.0032	0.9486	3.5507	V <sup>1.9175</sup>	
	Stem	$W_s=e^{eta_{s0}}V^{eta_{s1}}$	-0.7210	0.0091	1.0381	0.0020	0.9797	5.0940	V <sup>1.8483</sup>	
Larch forest	Branch	$W_b=e^{eta_{b0}}V^{eta_{b1}}$	-3.0835	0.0216	1.0941	0.0048	0.8776	1.6705	V <sup>2.3133</sup>	
	Foliage	$W_f = e^{eta_{f0}} V^{eta_{f1}}$	-3.3961	0.0277	0.8887	0.0060	0.8796	0.3691	V <sup>1.4197</sup>	
	Total	$W_t = W_r + W_s + W_b + W_f$	-	-	-	-	0.9741	9.1558	V <sup>1.9543</sup>	
	Root	$W_r = e^{eta_{r0}} V^{eta_{r1}}$	-1.7307	0.0645	0.9913	0.0135	0.8870	3.5422	V <sup>1.5588</sup>	
	Stem	$W_s=e^{eta_{s0}}V^{eta_{s1}}$	-0.4645	0.0216	0.9738	0.0046	0.9647	6.0568	V <sup>1.5025</sup>	
Poplar-birch forest	Branch	$W_b=e^{eta_{b0}}V^{eta_{b1}}$	-3.4476	0.1073	1.2090	0.0221	0.7181	3.4773	V <sup>1.5094</sup>	
	Foliage	$W_f = e^{eta_{f0}} V^{eta_{f1}}$	-3.7540	0.0539	0.9639	0.0111	0.8568	0.4373	V <sup>0.9094</sup>	
	Total	$W_t = W_r + W_s + W_b + W_f$	-	-	-	-	0.9414	12.0418	V <sup>1.5480</sup>	
	Root	$W_r = e^{eta_{r0}} V^{eta_{r1}}$	-1.2042	0.0318	0.9457	0.0077	0.7794	4.5912	V <sup>1.4114</sup>	
Desiduous breedloof	Stem	$W_s=e^{eta_{s0}}V^{eta_{s1}}$	-0.2496	0.0423	0.9723	0.0096	0.8465	11.2301	V <sup>1.1079</sup>	
mixed forest	Branch	$W_b=e^{eta_{b0}}V^{eta_{b1}}$	-1.9774	0.1601	1.0408	0.0354	0.3156	7.8146	V <sup>1.0047</sup>	
inixed forest	Foliage	$W_f = e^{eta_{f0}} V^{eta_{f1}}$	-2.9984	0.0658	0.8809	0.0149	0.6615	0.6916	V <sup>0.9736</sup>	
	Total	$W_t = W_r + W_s + W_b + W_f$	-	-	-	-	0.752	23.7878	V <sup>1.0922</sup>	
	Root	$W_r = e^{eta_{r0}} V^{eta_{r1}}$	-1.5946	0.0124	1.0234	0.0029	0.9072	3.4624	V <sup>1.9507</sup>	
Conifornus and broadloof	Stem	$W_s=e^{eta_{s0}}V^{eta_{s1}}$	-0.5931	0.0107	1.0129	0.0024	0.9668	5.2164	V <sup>1.7036</sup>	
mixed forest	Branch	$W_b=e^{eta_{b0}}V^{eta_{b1}}$	-2.4238	0.0725	1.0102	0.0152	0.7265	2.6839	V <sup>1.2166</sup>	
inixed forest	Foliage	$W_f = e^{eta_{f0}} V^{eta_{f1}}$	-3.2728	0.0504	0.8894	0.0105	0.8447	0.4113	V <sup>0.8493</sup>	
	Total	$W_t = W_r + W_s + W_b + W_f$	-	-	-	-	0.9512	10.0404	V <sup>1.3309</sup>	
	Root	$W_r = e^{eta_{r0}} V^{eta_{r1}}$	-1.9686	0.0854	1.0505	0.0184	0.7036	6.324	V <sup>2.0523</sup>	
	Stem	$W_s=e^{eta_{s0}}V^{eta_{s1}}$	-1.1586	0.0463	1.1051	0.0097	0.9535	6.9835	V <sup>1.9406</sup>	
Coniferous mixed forest	Branch	$W_b=e^{eta_{b0}}V^{eta_{b1}}$	-2.5850	0.0885	0.9942	0.0178	0.8994	1.3603	V <sup>1.0203</sup>	
	Foliage	$W_f = e^{eta_{f0}} V^{eta_{f1}}$	-2.3607	0.0899	0.7545	0.0187	0.7225	0.8256	V <sup>0.8568</sup>	
	Total	$W_t = W_r + W_s + W_b + W_f$	-	-	-	-	0.9294	12.8935	V <sup>1.8315</sup>	

**Table 4.** Model coefficient estimates, standard errors (SEs), model fitting statistics, and weight functions for the additive system of stand biomass equations using stand volume (namely, M-2) for the six forest types in the Eastern Da Xing'an Mountains, Northeast China.

Forest Types	Components	Equations	$\beta_{i0}$		$\beta_{i1}$		$\beta_i$	2	R <sub>2</sub> <sup>2</sup>	RMSE	Weight Function	
<b>91</b>		-1	Estimate	SE	Estimate	SE	Estimate	SE	a	in the second se	8	
	Root	$W_r = H_a / (\beta_{r0} + \beta_{r1} H_a) V$	-3.8705	0.3870	4.8137	0.0377			0.9806	1.3183	V <sup>1.7159</sup>	
	Stem	$W_{\rm s} = \beta_{\rm s0} D_a^{\beta_{\rm s1}} H_a^{\beta_{\rm s2}} V$	0.5195	0.0098	0.1354	0.0079	-0.0803	0.0078	0.9866	2.9346	V <sup>1.3715</sup>	
White birch forest	Branch	$W_b = \beta_{b0} D_a^{\beta_{b1}} H_a^{\beta_{b2}} V$	0.0179	0.0008	0.9309	0.0239	-0.1979	0.0219	0.8945	1.7465	V <sup>2.1459</sup>	
	Foliage	$W_f = \beta_{f0} D_q^{\beta_{f1}} V$	0.0160	0.0005	0.1952	0.0133	-	-	0.9476	0.2515	V <sup>1.7707</sup>	
	Total	$W_t = W_r + W_s + W_b + W_f$	-	-	-	-	-	-	0.9825	5.4113	V <sup>1.5780</sup>	
	Root	$W_r = D_q / (\beta_{r0} + \beta_{r1} D_q) V$	13.8235	0.4135	3.2953	0.0314	-	-	0.9573	3.2350	V <sup>2.2507</sup>	
	Stem	$W_s = \beta_{s0} D_q^{\beta_{s1}} H_a^{\beta_{s2}} V$	0.4419	0.0052	0.1069	0.0049	-0.0086	0.0025	0.9816	4.8573	V <sup>2.2398</sup>	
Larch forest	Branch	$W_b = D_q / \left(\beta_{b0} + \beta_{b1} D_q\right) V$	22.8657	2.2582	12.5955	0.1868	-	-	0.8729	1.7027	V <sup>2.2522</sup>	
	Foliage	$W_f = D_a / (\beta_{f0} + \beta_{f1} D_a) V$	-289.2900	4.4326	71.4847	0.4036	-	-	0.9232	0.2948	V <sup>1.2022</sup>	
	Total	$W_t = W_r + W_s + W_h + W_f$	-	-	-	-	-	-	0.9743	9.1126	V <sup>2.2993</sup>	
	Root	$W_r = \beta_{r0} D_a^{\beta_{r1}} H_a^{\beta_{r2}} V$	0.2137	0.0161	0.2494	0.0509	-0.3270	0.0509	0.8884	3.5205	V <sup>1.7272</sup>	
	Stem	$W_s = \beta_{s0} D_q^{\beta_{s1}} H_a^{\beta_{s2}} V$	0.6663	0.0208	0.0725	0.0239	-0.1369	0.0237	0.9618	6.2994	V <sup>1.4822</sup>	
Poplar-birch forest	Branch	$W_b = \beta_{b0} D_a^{\beta_{b1}} H_a^{\beta_{b2}} V$	0.0173	0.0021	1.1033	0.0680	-0.4538	0.0693	0.8114	2.8447	V <sup>2.1804</sup>	
	Foliage	$W_f = \beta_{f0} D_q^{\beta_{f1}} H_a^{\beta_{f2}} V$	0.0196	0.0017	0.3182	0.0539	-0.3008	0.0533	0.8575	0.4361	V <sup>1.5919</sup>	
	Total	$W_t = W_r + W_s + W_b + W_f$	-	-	-	-	-	-	0.9451	11.6543	V <sup>1.8216</sup>	
	Root	$W_r = \beta_{r0} D_q^{\beta_{r1}} H_a^{\beta_{r2}} V$	0.2863	0.0116	0.3425	0.0183	-0.4423	0.0194	0.8800	3.3866	V <sup>1.7508</sup>	
Deciduous broadloaf	Stem	$W_s = \beta_{s0} D_q^{\beta_{s1}} H_a^{\beta_{s2}} V$	0.5869	0.0167	0.3837	0.0134	-0.3406	0.0146	0.9247	7.8648	V <sup>2.0869</sup>	
mixed forest	Branch	$W_b = \beta_{b0} D_q^{\beta_{b1}} H_a^{\beta_{b2}} V$	0.0296	0.0028	1.2145	0.0328	-0.5961	0.0410	0.6560	5.5406	V <sup>1.9637</sup>	
nuxed forest	Foliage	$W_f = \beta_{f0} D_q^{\beta_{f1}} H_a^{\beta_{f2}} V$	0.0328	0.0017	0.4190	0.0228	-0.4854	0.0256	0.7798	0.5578	V <sup>2.0743</sup>	
	Total	$W_t = W_r + W_s + W_b + W_f$	-	-	-	-	-	-	0.8783	16.6652	V <sup>2.2717</sup>	
	Root	$W_r = \beta_{r0} D_q^{\beta_{r1}} H_a^{\beta_{r2}} V$	0.1808	0.0058	0.1817	0.0176	-0.0930	0.0153	0.9119	3.3740	V <sup>2.3133</sup>	
Coniferous and	Stem	$W_s = \beta_{s0} D_q^{\beta_{s1}} H_a^{\beta_{s2}} V$	0.4878	0.0085	0.1637	0.0097	-0.0915	0.0084	0.9711	4.8728	V <sup>1.9186</sup>	
broadleaf mixed forest	Branch	$W_b = \beta_{b0} D_q^{\beta_{b1}} H_a^{\beta_{b2}} V$	0.0280	0.0014	0.7682	0.0258	-0.3026	0.0243	0.7655	2.4851	V <sup>2.1090</sup>	
broudlear milled forest	Foliage	$W_f = H_a / \left(\beta_{f0} + \beta_{f1} H_a\right) V$	-66.2815	7.3367	49.8705	0.6624	-	-	0.8360	0.4227	V <sup>1.6276</sup>	
	Total	$W_t = W_r + W_s + W_b + W_f$	-	-	-	-	-	-	0.9575	9.3737	V <sup>2.1261</sup>	
	Root	$W_r = D_q / \left(\beta_{r0} + \beta_{r1} D_q\right) V$	-0.9311	2.9416	5.7384	0.2412	-	-	0.7110	6.2451	V <sup>2.2076</sup>	
	Stem	$W_s = D_q / (\beta_{s0} + \beta_{s1} D_q) V$	4.3149	0.4804	1.6180	0.0360	-	-	0.9492	7.3040	V <sup>1.9316</sup>	
Coniferous mixed forest	Branch	$W_b = \beta_{b0} D_a^{\beta_{b1}} H_a^{\beta_{b2}} V$	0.0589	0.0042	0.1876	0.0316	-0.1121	0.0321	0.9015	1.3461	V <sup>1.2137</sup>	
	Foliage	$W_f = \beta_{b0} H_a^{\beta_{b1}} V$	0.0455	0.0053	-0.1797	0.0443	-	-	0.6828	0.8826	V <sup>1.1395</sup>	
	Total	$W_t = W_r + W_s + W_b + W_f$	-	-	-	-	-	-	0.9282	13.0000	V <sup>2.2442</sup>	

**Table 5.** Model coefficient estimates, standard errors (SEs), model fitting statistics, and weight functions for the additive system of stand biomass equations using stand volume and BCEF (namely, M-3) for the six forest types in the Eastern Da Xing'an Mountains, Northeast China.

Forest Types	Source of Variation	Root Biomass		Stem Biomass		Branch Biomass		Foliage Biomass		Total Biomass	
J1	Source of variation	F-Value	<i>p</i> -Value	F-Value	<i>p</i> -Value	F-Value	<i>p</i> -Value	F-Value	<i>p</i> -Value	Total Biomass           F-Value $p$ -Val $503.58$ < 0.00 $12.33$ < 0.00 $312.42$ < 0.00 $273.76$ < 0.00 $273.76$ < 0.00 $94.89$ < 0.00 $6.94$ < 0.00 $255.41$ < 0.00 $0.1$ $0.957$	<i>p</i> -Value
	Block (Plot)	641.65	< 0.0001	593.86	< 0.0001	84.73	< 0.0001	406.9	< 0.0001	503.58	< 0.0001
white birch forest	Treatment (4 Methods)	48.64	< 0.0001	15.15	< 0.0001	16.81	< 0.0001	25.62	< 0.0001	12.33	< 0.0001
	Block (Plot)	245.83	< 0.0001	326.91	< 0.0001	268.58	< 0.0001	142.58	< 0.0001	312.42	< 0.0001
Larch forest	Treatment (4 Methods)	99.07	< 0.0001	44.82	< 0.0001	110.01	< 0.0001	123.24	< 0.0001	70.36	< 0.0001
D 1 1:16 (	Block (Plot)	245.88	< 0.0001	306.33	< 0.0001	66.16	< 0.0001	221.05	< 0.0001	273.76	< 0.0001
Poplar-birch lorest	Treatment (4 Methods)	2.84	0.0369	7.34	0.0001	10.87	< 0.0001	8.83	< 0.0001	3.04	0.0284
Deciduous broadleaf	Block (Plot)	92.43	< 0.0001	124.58	< 0.0001	32.7	< 0.0001	77.67	< 0.0001	94.89	< 0.0001
mixed forest	Treatment (4 Methods)	16.26	< 0.0001	6.36	< 0.0001	6.76	0.0003	22.48	< 0.0001	6.94	< 0.0001
Coniferous and	Block (Plot)	269.22	< 0.0001	277.81	< 0.0001	99.47	< 0.0001	239.67	< 0.0001	255.41	< 0.0001
broadleaf mixed forest	Treatment (4 Methods)	269.22	0.4381	0.13	0.9425	1.87	0.1329	59.34	< 0.0001	0.1	0.9572
C :( · 1( ·	Block (Plot)	139.22	< 0.0001	132.51	< 0.0001	134.67	< 0.0001	83.76	< 0.0001	140.9	< 0.0001
Coniferous mixed forest	Treatment (4 Methods)	2.21	0.0859	6.64	0.0002	1.26	0.2864	30.48	< 0.0001	3.4	0.0173

Table 6. Comparison of the four estimation methods of stand biomass of the six forest types in the Eastern Da Xing'an Mountains, Northeast China.

# 3.2. Model Validation for Stand Biomass Models

The model validation statistics (Equations (9)–(13)) for M-1, M-2, and M-3 were computed. For most stand biomass models, the MPE was close to 0, the MAE was relatively small (< 20 Mg·ha<sup>-1</sup>), the MAE% was less than 30%, and M-2 and M-3 seemed preferable to M-1. For the stand total biomass, relatively small model prediction errors were reported for three of the systems in all forest types, except deciduous broadleaf mixed forest, and M-3 and M-2 seemed to be better than M-1, which was also found for the stand stem biomass (Figure 2). On the other hand, the stand biomass equations for root, branch, and foliage had less accurate predictions than the total and stem models, especially the branch biomass models.



**Figure 2.** Mean prediction error (MPE), mean absolute error (MAE) and mean absolute percent error (MAE%) among the total and component biomasses for each forest type. WBF, LF, PBF, DBMF, CBMF and CMF stand for white birch forest, larch forest, poplar-birch forest, deciduous broadleaf mixed forest, coniferous and broadleaf mixed forest and coniferous mixed forest, respectively. TB, RB, SB, BB and FB represent the total, root, stem, branch, and foliage biomass, respectively.

To quantify the contributions of the variations in the observed stand biomass to the model predictions, the residual and approximate confidence bands of the observed data containing approximately 90% of the average curve were derived for all additive systems of stand biomass equations. The specific methods used to calculate the approximate confidence band were detailed in Bi et al. [36]. Since the relative proportion of stand stem biomass was large for the six forest types, Figure 3 shows only the observed stand total and stem biomasses for the six forest types plotted against their predicted values from M-1, M-2 and M-3. The incorporation of *V* and BCEF as the predictors in M-3 led to slightly narrower confidence bands for the stand biomass than in M-1 (with *G* and  $H_a$  as the predictors). In general, M-3 predicted the stand total biomass and all components very well for the six



**Figure 3.** Multipanel display of observed total and stem biomass for each forest type plotted against their predicted values from M-1, M-2 and M-3. The diagonal line of unity is shown with the 90% upper and lower confidence limits of prediction error in each panel.

# 3.3. Comparison of Methods for Estimating Stand Biomass

The analysis of variance was performed to compare the performance of the four estimation methods (i.e., M-1, M-2, M-3, and M-4) in estimating stand biomass, where the four methods were the treatment, and the sample plots served as the blocks (Table 6). The results showed that there were remarkable differences in the stand total and component biomasses among the four methods for white birch forest, larch forest, poplar-birch forest, and deciduous broadleaf mixed forest (Table 6), although significant differences were not observed between the four methods for the stand total, root, stem, and

branch biomass models of coniferous and broadleaf mixed forest or for the stand root and branch biomass models of coniferous mixed forest (Table 6).

Furthermore, the percentage difference (MPD%) between the stand total biomass and component biomasses estimated by the four methods was calculated (Figure 4). The results showed that the MPD% between the stand total biomass and components in M-3 was small. However, the MPD% of M-1 was slightly larger than that of the other three methods in all forest types except deciduous broadleaf mixed forest. The overall ranking based on MPD% obeyed the following order: M-3 > M-2 > M-4 > M-1.



**Figure 4.** The mean percentage differences of the four methods used to estimate stand biomass for each forest type. WBF, LF, PBF, DBMF, CBMF and CMF stand for white birch forest, larch forest, poplar-birch forest, deciduous broadleaf mixed forest, coniferous and broadleaf mixed forest and coniferous mixed forest, respectively. TB, RB, SB, BB and FB represent the total, root, stem, branch and foliage biomass, respectively.

# 4. Discussion

Stand total and component biomasses are frequently estimated using stand variables, which are usually easy to obtain via field investigations. In recent studies on stand biomass estimations, typical allometric equations based on the power-law model are often applied to increase the prediction accuracy of stand biomass [6–8,16]. In our M-1 model system, the stand basal area *G* was a significant

and important predictor for the stand total and component biomasses of the six forest types, which confirmed its role in previous studies [6,7]. The estimated coefficients of the stand basal area were positive, indicating that the stand biomass increased as the stand basal area increased. However, for the same stand basal area, large variations occurred among the stand total and component biomasses. Therefore, to improve the prediction accuracy of the stand biomass models, other stand variables should be considered. Previous studies have verified that the dominant height of the stand was the second most important stand variable because it reflected the site quality [7,19]. Unfortunately, the stand dominant height was not available in our data from the permanent sample plots of the NFCI. Instead, we used the average stand height  $H_a$ , which was statistically significant for most stand root, stem, branch, and foliage biomass models. As a result, we strongly suggest that future studies on stand biomass consider and investigate the effects of the stand average height or dominant height in the stand biomass models. Other stand variables, such as stand density and stand age, were also included in some stand component biomass equations, and they were commonly considered to represent the competition within a stand and the stage of stand development [6]. The effects of stand density and stand age on some biomass components, such as stem or foliage, are statistically significant and widely recognized, which is likely due to their impacts on branching characteristics and biomass partitioning among the tree components [6,37]. However, the six forest types in this study were natural forests in the Eastern Da Xing'an Mountains. Therefore, they were not good choices for estimating stand biomass. In addition, information on silvicultural practices, such as thinning and tending, was not available, which may affect stand biomass accumulation and allocation and result in biased stand biomass estimations [38].

Many studies indicate that there is a strong correlation between stand biomass and stand volume [20–22]. For most forest types, a linear relationship between stand biomass and stand volume is obviously insufficient and controversial. Thus, many researchers have used nonlinear models to describe the relationship and improve model fitting [7,15–17,24]. As expected, BCEF was not constant over stand development in different forest types (Table 2). Thus, using constant BCEF values would provide biased estimations of stand biomass. In this study, Equations (5) and (6) were used to model the BCEF variations, which is consistent with the methods used in other studies in the literature. In addition, some studies used stand age as a predictor for modeling BCEF [15,39–41]. Although the stand age was unavailable in this study, the stand variables (e.g.,  $H_a$  and  $D_q$ ) used in our model systems were indirect surrogates for stand development to a certain extent.

Parresol [28] noted that the aggregation approach is the standard method for ensuring the additivity of estimates of stand total and component biomasses. In Parresol's method, a nonlinear model is assigned to each of the stand biomass components before aggregating the biomasses of these stand components into the stand total biomass. Notably, if the same predictor variables are used for modeling all stand biomass components, and the same weights are chosen for heteroscedasticity in the model residuals, NSUR will produce singular covariance matrices that do not guarantee a unique solution. Thus, when the same or different predictor variables are considered, the heteroscedasticity of the model residuals should be overcome by different sets of weights, and NSUR is a feasible parameter estimation method [12,42]. The aggregative models in this study were estimated using a weighted NSUR with different weight functions to explain the inherent correlations among the stand component biomasses in the same sample plot [11–13,27,28]. Unfortunately, some nonadditive stand biomass models are still published because of the use of the ordinary least-squares regression (OLS) estimation method [8,10].

We evaluated different methods for quantifying stand biomass, and the results were nonconclusive. The results indicated that, except for deciduous broadleaf mixed forest, M-3 performed better than other models, especially for stand total and stem biomass. The reason for this finding may be that there is a close relationship between stand volume and stem biomass, and a higher proportion of stem biomass is included in the total biomass [7,43]. If the stand volume is not available, M-1 will be an effective method to accurately estimate the stand total and component biomasses. For the deciduous

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broadleaf mixed forest, the stand biomass models with stand variables (M-1) were generally less biased than those obtained from the stand biomass models with stand volume (M-2) or stand volume and BCEF (M-3). The poor performance of the latter two additive models may be the result of the biases after modeling BCEF by using the stand variables. Many studies have shown that using a constant BCEF to quantify stand biomass introduces certain biases or errors in stand biomass or carbon estimations so that the development of BCEF models is recommended when the data are available [16]. Generally, BCEFs vary depending on the growth conditions and stand development stage expressed by the age and size of the stand [10,15–17,41]. Thus, it is easy to understand that M-4 exhibited a worse prediction accuracy than M-2 and M-3. In general, both M-2 and M-3 are stand biomass estimations method based on stand volume, which is essentially different from M-1 [44–46]. Overall, the use of M-1, M-2, and M-3 in conjunction with stand growth and yield models would be useful for the growth predictions of the six forest types in response to changes in stand conditions and is appropriate for sustainable forest ecosystem management studies.

To date, few studies have developed individual tree biomass equations for the major forest species in the Eastern Da Xing'an Mountains, Northeast China [14]. However, the data required to apply these equations are not always available. In addition, some growth models for the six forest types in the region were previously developed at Northeast Forestry University, China, which mainly calculated the stand variables, such as *G* and *V*. For the stand biomass estimations, the use of stand variables can likely avoid the need to address complex error propagation procedures for stand biomass estimations at different spatial scales [7]. Although estimating the stand biomass using stand variables may produce some differences in the tree-based biomass estimation, the differences seem acceptable at the forest or stand level for a large geographic region. Thus, the stand biomass models using stand variables (e.g., *G*, *V*, *H*<sub>a</sub>, and *D*<sub>a</sub>) would be useful, convenient, and efficient.

# 5. Conclusions

Our study provided a comprehensive overview of the methods used to estimate the aboveground and belowground stand biomass of the six major forest types in the eastern Da Xing'an Mountains, Northeast China. Three additive systems of stand biomass equations were developed and evaluated: the stand biomass models using stand variables (M-1), the stand biomass models using stand volume (M-2), and the stand biomass models using both stand volume and BCEF (M-3). In these three model systems, the inherent correlation between stand total biomass and component biomass was constrained by applying NSUR for the model parameter estimation. We also comprehensively evaluated four stand biomass estimation methods (i.e., the three additive systems and constant BCEF (M-4)). The results indicated that the model fitting and validation of M-3, which included both stand volume and BCEF, were better than those of M-1 and M-2. Overall, the four methods showed good accuracy in estimating stand total and component biomasses for the six major forest types in the region. However, the choice of methods depends on the available data. If the stand volume is not known, then the stand biomass can be estimated using stand variables, such as *G* and  $H_a$ . However, caution should be used when predicting stand total and component biomasses outside the data range used in this study.

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