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Title: Developing and Comparing Individual Tree Growth Models of Major Coniferous Species in South Korea Based on Stem Analysis Data

Year: 2023

Version: Published version

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Please cite the original version:

Seo, Y.; Lee, D.; Choi, J. Developing and Comparing Individual Tree Growth Models of Major Coniferous Species in South Korea Based on Stem Analysis Data. *Forests* 2023, *14*, 115.
<https://doi.org/10.3390/f14010115>

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Article

Developing and Comparing Individual Tree Growth Models of Major Coniferous Species in South Korea Based on Stem Analysis Data

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Abstract: Tree growth in Korean red pine (*Pinus densiflora*, hereafter *Pd*), Korean white pine (*Pinus koraiensis*, hereafter *Pk*), and Japanese larch (*Larix kaempferi*, hereafter *Lk*) was modeled using Logistic, Korf, Gompertz, Chapman-Richards, and Weibull equations and stem analysis data from sample trees: 38 trees for *Pd*, 46 trees for *Pk*, and 45 trees for *Lk*. The models were fitted to the total increment of tree size variables, diameter at breast height (DBH), height, basal area, and stem volume, as a function of age. After selecting the best-fit growth function, the current annual increment (CAI) and mean annual increment (MAI) were compared for each variable by species. The optimal growth functions were Chapman-Richards for DBH and stem volume, Korf for height, and Gompertz for basal area. The parameter estimates in the final models were all significant ($p < 0.01$) with best-fit statistics and unbiased residual plots. When plotted with observed values, the growth patterns of each variable were represented properly. The predicted growth curves over age were concave with respect to the Y-axis in DBH and height but lightly convex in basal area, and explicitly convex in stem volume, whereas an asymptote of sigmoid curve in stem volume was not apparent until 100 years. Age with the maximum MAI among variables was arranged similarly to CAI; the age with maximum MAI was earliest for DBH and latest for volume. The maximum growth was achieved earliest in *Lk*, followed by *Pk* and *Pd*. The developed models were able to predict tree size variables and serve as a reference to understand growth characteristics by species.

Keywords: sigmoid growth function; total increment; current annual increment; mean annual increment; dominant trees



Citation: Seo, Y.; Lee, D.; Choi, J. Developing and Comparing Individual Tree Growth Models of Major Coniferous Species in South Korea Based on Stem Analysis Data. *Forests* **2023**, *14*, 115. <https://doi.org/10.3390/f14010115>

Academic Editor: Mark Vanderwel

Received: 30 November 2022

Revised: 23 December 2022

Accepted: 4 January 2023

Published: 7 January 2023



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1. Introduction

Korean red pine (*Pinus densiflora* Siebold & Zucc., hereafter *Pd*), Korean white pine (*Pinus koraiensis* S. & Z., hereafter *Pk*), and Japanese larch (*Larix kaempferi* (Lamb.) Carrière, hereafter *Lk*) are the major plantation species in South Korea because they are fast-growing and commercially used for lumber [1–3]. These species have been intensively planted since the 1970s. In 2020, approximately 36.9% (2,319,832 ha) of forested land in South Korea was occupied by coniferous species [4]. Out of the total coniferous forested area, 68.1% (1,579,787 ha), 6.5% (151,946 ha), and 11.2% (260,255 ha) was occupied by *Pd*, *Pk*, and *Lk*, respectively [4]. Due to the continued importance of these tree species as commercial wood supply, concerns have increased regarding their growing methods and management, as well as future expectations for the growth and development of these species.

In contemporary stand management planning, selecting a tree growth function is an important task of forest science. Growth functions, which are statistical expressions of

biological growth principles, form the basis for the realistic determination of stand productivity level, tree dimensions, and potential structure assortments [5–7]. Growth functions describe the incremental growth changes of an individual or population over time [8]. Selecting the appropriate growth functions for tree and stand modeling is important for developing growth and yield models.

Sigmoid growth functions are more commonly used than non-sigmoid growth functions in forest modeling as the diameter at breast height (DBH), height, and basal area of trees are asymptotic with age [9–11]. The growth functions have been associated with immediate application in regard to forestry management planning. The functions are based on knowledge, definition, and usage of characteristics on the growth curve, and periodic and average increment curves [12].

For these reasons, the models using sigmoid growth functions have been developed to understand growth characteristics and predict tree size in the forestry in advanced countries [10,13–19]. In addition to the total increment of tree size, current annual increment (CAI) and mean annual increment (MAI) are important growth characteristics because those indicate a quantitative amount of increment in relation to the change in age and provide an implication of growth rate, which is also used for economic assessment [8,11]. They can help to evaluate the age at which a tree achieves the maximum size [20–22]. Yet, there is a lack of relevant studies even for the major commercial species such as *Pd*, *Pk*, and *Lk*. Thus, the information on tree growth remains unknown and any comparisons among species incomplete. At present, the variables of tree size in South Korea cannot be predicted because research on growth functions is lacking.

This study aimed to fit the growth models of DBH, height, basal area, and stem volume for *Pd*, *Pk*, and *Lk* using widely used sigmoid growth functions in the field of forest biometrics and to identify the best-fit growth model for each variable of the three species. After determining the best growth model, the total increment, CAI, and MAI of each variable over age were derived from the best growth models of the three species. Based on the simulation, additionally, the growth processes of each variable were compared for the three species.

2. Materials and Methods

2.1. Study Area

2.1.1. Location and Site Characteristics

For this study, *Pd*, *Pk*, and *Lk* plantations were targeted, and the selected experimental plots were located in the national forests of the north-eastern region of South Korea. A total of 129 sites were selected as study areas: 38 sites for *Pd*, 46 sites for *Pk*, 45 sites for *Lk* (Figure 1). The experimental plots were situated between 36°33' and 38°18' N latitude and 127°34' E and 129°21' E longitude and represented the typical forests of South Korea formed in mountain areas. According to the Korea Meteorological Administration [23], the 30-year averaged mean annual temperature was 7.1–13.5 °C, the minimum was 2.3–9.6 °C, and was the maximum 12.2–18.3 °C among municipalities where the plots were located. The mean annual precipitation ranged from 1112.2 to 1695.1 mm in the recent 30-year average (1991–2020), with most of the precipitation occurring during the rainy season in summer, which is a typical climate phenomenon in South Korea.

2.1.2. Plot Characteristic with Stand Density

The experimental plots were established to study the growth and yield according to the thinning intensity. For this purpose, one of the prerequisites of a permanent plot was a stand that was as heavily dense as possible without thinning. The plots were installed in the forest where the most recent thinning was conducted at least more than 7 years ago. The selected target stands had a relatively high stand density before installing the plots and executing the thinning operation [24,25]. The relative density (stand density index divided by maximum stand density index) before thinning was mostly above 0.7 in each stand type by species, according to the models of Lee and Choi [24,25].

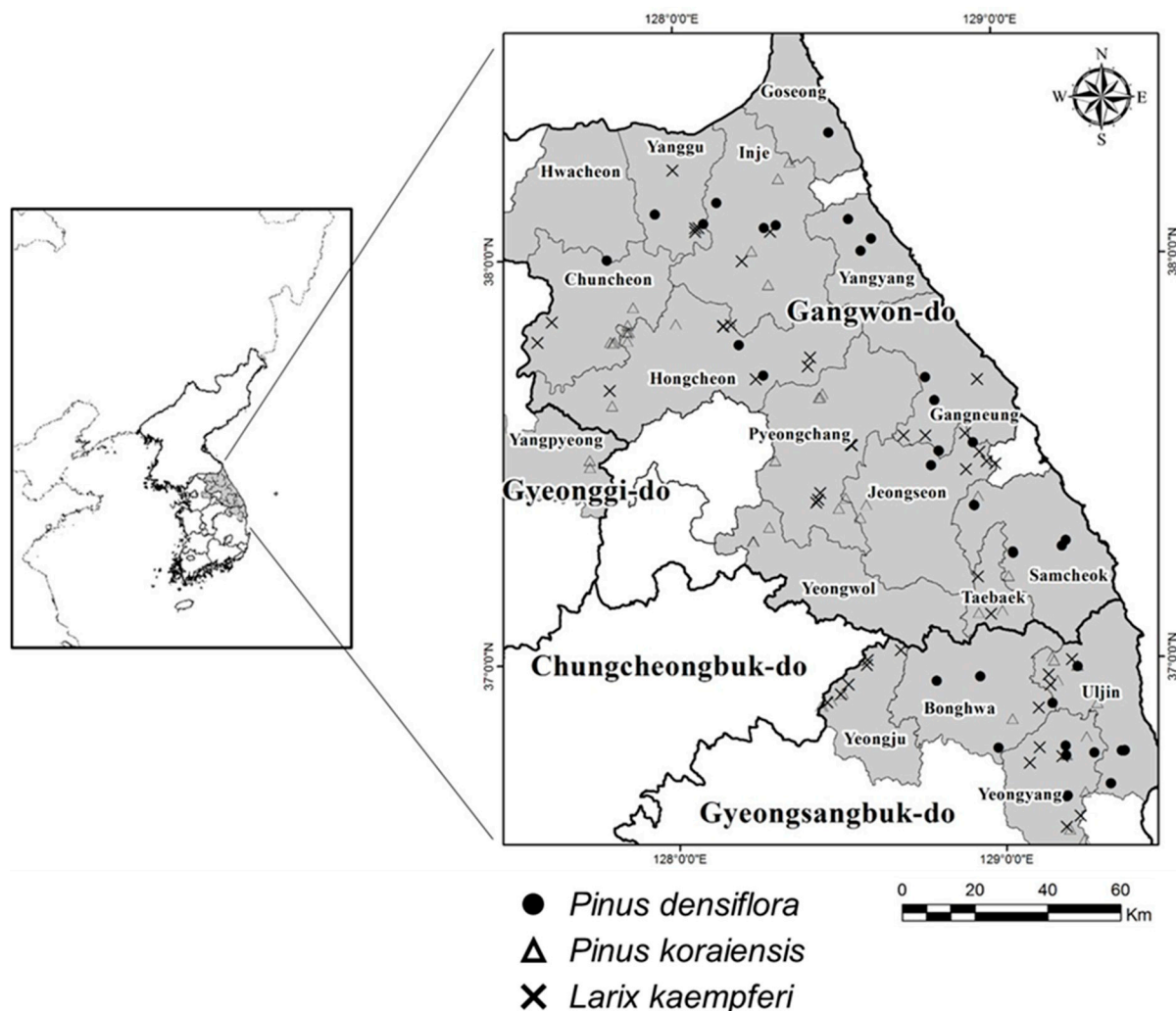


Figure 1. Study sites and plot location by species from which sample trees were collected for stem analysis and model development.

2.2. Data Collection

2.2.1. Sample Tree Selection and Wood Disc Collection

All the analyses in this study were based on sample trees of each site studied for stem analysis. To collect stem profile data, we selected a representative sample tree that was either dominant or codominant in a stand, free of damage, and desirable in terms of stem quality, that is, straight, unforked, and uncrooked stem for sawtimber felled at each of the 129 sites [20,21]. After selecting a sample tree, it was felled, and 3–5 cm thickness of wood discs were collected from the sectioned stem. The discs were collected at a height of 0.2 m for the first wood disc (W_0), 1.2 m for the second one (W_1 ; corresponds to breast height in South Korea), 3.2 m (W_2), and from this point onward at two-meter intervals until the second to the last disc (W_{n-1}). The last wood disc (W_n) was collected 1 or 2 m from the previous (W_{n-1}), depending on the total stem height. For example, the distance between W_{n-1} and W_n was 1 m if the cone length is shorter than 1 m and the distance was 2 m if the cone length is longer than 1 m to collect an appropriate sample disc size, e.g., disc diameter > 1 cm. All the collected wood discs were naturally dried for several days, and then sanded using a sanding machine to prepare the surface for tree ring measurements. The fieldwork was carried out between 2012 and 2017, depending on the experimental site and target species.

2.2.2. Tree Ring Measurements and Stem Analysis

After completing the field works, the number of annual rings were recorded, and the 5-year increment was measured using the SENC 150 Precision Glass Scale Linear Encoder (HEIDENHAIN Corp., Schaumburg, IL, USA), the MEIJI EMZ-5TR microscope (MEIJI TECHNO CO., LTD., San Jose, CA, USA), and the advanced digital readout system Quadra-Chek 10 (METRONICS Inc., Bedford, NH, USA), which allow single-axis measurements at very high levels of precision and accuracy. The tree age was assigned as the number of tree rings on wood disc 0 (W_0) plus 2 as conventionally applied for coniferous tree species in South Korea. The summary statistics of the sample trees are given in Table 1.

Table 1. Summary statistics of sample trees per species at the time of tree felling for stem analysis.

Species	Statistics	Age (Year)	DBH (cm)	Height (m)	Volume (m ³)	No. of Trees	No. of Sample Points at 5-Year Interval
<i>Pinus densiflora</i>	Mean	47	30.5	17.0	0.6707	38	331
	SD	16	8.4	4.0	0.3940		
	Minimum	9	5.2	3.6	0.0067		
	Maximum	99	47.3	23.8	1.6448		
<i>Pinus koraiensis</i>	Mean	40	29.8	17.2	0.6935	46	351
	SD	16	7.9	4.4	0.4743		
	Minimum	15	13.9	7.4	0.0527		
	Maximum	77	45.9	24.6	1.9642		
<i>Larix kaempferi</i>	Mean	38	27.8	22.8	0.7480	45	321
	SD	12	7.9	4.4	0.5160		
	Minimum	19	17.0	12.2	0.1614		
	Maximum	60	47.9	30.6	2.2803		

SD: standard deviation.

The total increment in DBH, height, basal area, and stem volume was measured at 5-year intervals (e.g., age 5, 10, 15, etc.). The 5-year increment of DBH and basal area was calculated based on the wood disc 1 (W_1), and the height increment was computed on a 5-year basis by interpolation method using the number of tree rings on a wood disc and the length between the wood discs. The 5-year increment of the stem volume was the difference between the stem volumes before and after 5-year growth, and the stem volume at each age was a result of the total sum of all log volumes at the corresponding age. The log volume was calculated according to the Smalian's method, which estimates the volume of a log by averaging the areas of the two log ends and multiplying the length of the log (mostly 2 m in our study, except for the 1-m length between W_0 and W_1) by the averaged area. Detailed descriptions of the traditional stem analysis can be found elsewhere [26].

2.3. Modeling Approach

2.3.1. Selection of Candidate Growth Functions

To fit the total increment of DBH, height, basal area, and stem volume of each species over age, growth functions, which are widely used in forest biometrics, were examined for this study: Logistic, Korf, Gompertz, Chapman-Richards, and Weibull [8,10,17,27]. These growth functions present a sigmoid curve, which often hypothesizes an asymptotic biological growth. A different background was developed for each model such as ecology, height growth of the forest stand, age distribution of the human population, animal growth, and probability distribution for the failure rate, and intrinsic characteristics varied according to the equations [10]. The sigmoid pattern is assumed as reasonable and largely applied for tree growth. Thus, the growth functions were compared and tested to find the best fit for this study. The integral and differential forms of each growth function are described in Table 2. In our study, the total increment corresponded to the integral form, and model parameters were fitted using the integral form of each function.

Table 2. Growth functions of integral and differential form used in this study.

Function	Integral Form	Differential Form
Logistic	$Y = a / (1 + ce^{-bt})$	$\Delta Y = abce^{-bt} / (1 + ce^{-bt})^2$
Korf	$Y = ae^{-bt-c}$	$\Delta Y = abct^{-(c+1)}e^{-bt-c}$
Gompertz	$Y = ae^{-be^{-ct}}$	$\Delta Y = abce^{-ct}e^{-be^{-ct}}$
Chapman-Richards	$Y = a(1 - e^{-bt})^c$	$\Delta Y = abce^{-bt}(1 - e^{-bt})^{(c-1)}$
Weibull	$Y = a(1 - e^{-bt^c})$	$\Delta Y = abct^{c-1}e^{-bt^c}$

Note: Y is the diameter at breast height (cm), height (m), basal area (m²), or stem volume (m³) at age t ; ΔY is the current annual increment of tree size at age t ; e , also known as Euler's number, is the base of natural logarithms; a, b, c are parameters to be determined by this study.

2.3.2. Statistical Modeling and Validation

The model fitting was conducted using the R statistical software, and *nls* function in *stats* packages was applied for estimating parameters of nonlinear regression [28]. The response variables of the models were DBH, height, basal area, and volume from all 5-year measurement instances. The explanatory variable was tree age in the corresponding size. Autocorrelation should be considered when several measurements are used from the same sample, although this correlation can be disregarded in stem analysis [9,18]. The 5-year interval is also regarded as less autocorrelated than the 1-year interval; therefore, the modeling can proceed without severe issues [29]. Consequently, the modeling method with only fixed effects was applied without considering autocorrelation and other random effects.

Model validity was evaluated by examining the parameter estimates together with standard error. The significance of p -value with t statistic was also checked to evaluate the stability of parameters. The coefficient of determination, or R-squared (R^2), was calculated as a fit statistic; however, R^2 alone may not be a sufficient metric as the models were fitted based on nonlinear regression. Accordingly, the root mean squared error (RMSE), the mean absolute error (MAE), and the mean absolute percentage error (MAPE) were also used to examine the model performance. The specific equations were as follows:

$$R^2 = 1 - \frac{\sum_{i=1}^n (y_i - \hat{y}_i)^2}{\sum_{i=1}^n (y_i - \bar{y})^2} \quad (1)$$

$$RMSE = \sqrt{\frac{\sum_{i=1}^n (y_i - \hat{y}_i)^2}{n}} \quad (2)$$

$$MAE = \frac{\sum_{i=1}^n |y_i - \hat{y}_i|}{n} \quad (3)$$

$$MAPE = \frac{100\%}{n} \sum_{i=1}^n \left| \frac{y_i - \hat{y}_i}{y_i} \right| \quad (4)$$

where y is a tree parameter such as DBH (cm), height (m), basal area (m²), and stem volume (m³); y_i is the observed value of sample i ; \hat{y}_i is the predicted value of the sample i ; \bar{y} is the observed average value; and n is the number of samples.

For the model validation after model fitting, the residual plots of each growth function were plotted by the variables and species. Additionally, the predicted growth curves with observed sample points were displayed over tree age. The model bias was analyzed using these residual plots and scatterplots. The final model for each variable by species was then selected synthetically, considering the parameter estimates, standard error, p -value, fit statistics, and the degree of bias in residuals.

Using the models selected by the variables, the total increment was demonstrated to show the representative growth curve in the age range of collected trees by species.

Additionally, the CAI and MAI were calculated in simulations by variables. The CAI and MAI were calculated based on the differential equations and on the integral form divided by tree age, respectively (Table 2). The CAI and MAI were described separately and collectively in figures to analyze the validity in terms of the growth pattern and intersecting point between two types of annual increment curves.

3. Results and Discussion

3.1. Parameters Estimates and Fit Statistics

Generally, all models were adequately fitted, and parameters were estimated without any singularity or convergence problem (Tables 3–6). The predicted growth curves were practically identical within the range of observations, but they varied outside the range of fitted tree age according to innate characteristics of the growth functions (Supplementary files). The parameters and standard errors ranged within the logical values in accordance with each growth function characteristic when compared with previous studies [13–17,30]. However, several unstable parameters and standard errors were with insignificant p -value ($p > 0.01$). For example, among DBH models, Korf function of Pd was not significant due to the parameter a ($p > 0.01$) (Table 3), while all parameters of each function in height models were significant (Table 4). In the basal area growth models, Korf for Pd and Pk and Korf, Chapman-Richards, and Weibull for Lk contained insignificant parameters ($p > 0.01$) (Table 5). In the case of volume models, Korf function for all species and Weibull function for Pd and Lk included insignificant parameters ($p > 0.01$) (Table 6). The logistic function performed much worse than all other functions in terms of the predicted curve with an extreme asymptote. Therefore, these models with unstable parameter estimates and/or unreliable prediction behavior were excluded from the best function (Tables 3–6 and Figures S1–S12).

Table 3. Model parameters of DBH growth functions and indicators of fit.

Species	Variables	Parm a		Parm b		Parm c		Fit Statistics			
		Estimate	S.E.	Estimate	S.E.	Estimate	S.E.	R ²	RMSE	MAE	MAPE
<i>Pinus densiflora</i> ($n = 303$)	Logistic	35.7659 (<0.0001)	1.3117	0.0847 (<0.0001)	0.0053	11.8485 (<0.0001)	1.2272	0.8291	4.0989	3.2799	0.5316
	Korf	134.3419 (0.0113)	52.7221	10.8703 (<0.0001)	1.6634	0.5042 (<0.0001)	0.1035	0.8412	3.9507	3.1056	0.3499
	Gompertz	39.8321 (<0.0001)	1.9710	3.3558 (<0.0001)	0.1921	0.0499 (<0.0001)	0.0040	0.8375	3.9962	3.1779	0.4498
	Chapman-Richards	47.4990 (<0.0001)	4.7042	0.0278 (<0.0001)	0.0053	1.6201 (<0.0001)	0.1748	0.8410	3.9537	3.1232	0.3843
	Weibull	44.1380 (<0.0001)	4.2293	0.0047 (<0.0001)	0.0010	1.4011 (<0.0001)	0.0886	0.8405	3.9590	3.1357	0.3984
<i>Pinus koraiensis</i> ($n = 324$)	Logistic	37.5633 (<0.0001)	0.6866	0.0980 (<0.0001)	0.0040	12.7995 (<0.0001)	1.0008	0.9125	3.2541	2.5803	0.4549
	Korf	82.1652 (<0.0001)	9.9198	14.0460 (<0.0001)	1.8092	0.7005 (<0.0001)	0.0657	0.9315	2.8781	2.1464	0.1926
	Gompertz	40.3021 (<0.0001)	0.9038	3.5817 (<0.0001)	0.1525	0.0606 (<0.0001)	0.0028	0.9243	3.0261	2.3421	0.3319
	Chapman-Richards	44.5393 (<0.0001)	1.6455	0.0393 (<0.0001)	0.0035	1.8676 (<0.0001)	0.1294	0.9294	2.9217	2.2029	0.2484
	Weibull	42.2732 (<0.0001)	1.5261	0.0047 (<0.0001)	0.0007	1.4882 (<0.0001)	0.0567	0.9280	2.9509	2.2372	0.2784

Table 3. Cont.

Species	Variables	Parm a		Parm b		Parm c		Fit Statistics			
		Estimate	S.E.	Estimate	S.E.	Estimate	S.E.	R ²	RMSE	MAE	MAPE
<i>Larix kaempferi</i> (n = 314)	Logistic	30.9468 (<0.0001)	0.8779	0.1130 (<0.0001)	0.0069	10.8427 (<0.0001)	1.1272	0.8507	3.6823	2.8871	0.5809
	Korf	86.7510 (0.0002)	22.6673	9.5077 (<0.0001)	1.2776	0.5696 (<0.0001)	0.0951	0.8744	3.3769	2.5151	0.2813
	Gompertz	33.3797 (<0.0001)	1.2273	3.2992 (<0.0001)	0.1944	0.0708 (<0.0001)	0.0051	0.8634	3.5213	2.7202	0.4476
	Chapman-Richards	39.1338 (<0.0001)	3.0142	0.0398 (<0.0001)	0.0068	1.5779 (<0.0001)	0.1638	0.8712	3.4201	2.5994	0.3629
	Weibull	37.5837 (<0.0001)	3.1412	0.0091 (<0.0001)	0.0016	1.3459 (<0.0001)	0.085	0.8701	3.4339	2.6253	0.3937

S.E.: standard error. R²: coefficient of determination. RMSE: root mean squared error. MAE: mean absolute error. MAPE: mean absolute percentage error.

Table 4. Model parameters of height growth functions and indicators of fit.

Species	Variables	Parm a		Parm b		Parm c		Fit Statistics			
		Estimate	S.E.	Estimate	S.E.	Estimate	S.E.	R ²	RMSE	MAE	MAPE
<i>Pinus densiflora</i> (n = 331)	Logistic	18.7523 (<0.0001)	0.3992	0.1097 (<0.0001)	0.0055	12.763 (<0.0001)	1.2794	0.8825	2.0430	1.5882	0.3479
	Korf	38.8374 (<0.0001)	5.6742	12.832 (<0.0001)	2.1414	0.7140 (<0.0001)	0.0871	0.8848	2.0225	1.5436	0.2555
	Gompertz	20.1584 (<0.0001)	0.5500	3.5208 (<0.0001)	0.1950	0.0674 (<0.0001)	0.0039	0.8883	1.9921	1.5135	0.2848
	Chapman-Richards	21.7626 (<0.0001)	0.9170	0.0460 (<0.0001)	0.0049	1.8879 (<0.0001)	0.1715	0.8886	1.9890	1.5016	0.2519
	Weibull	20.5390 (<0.0001)	0.7875	0.0055 (<0.0001)	0.0011	1.5140 (<0.0001)	0.0718	0.8890	1.9857	1.4970	0.2588
<i>Pinus koraiensis</i> (n = 351)	Logistic	20.5229 (<0.0001)	0.2727	0.1128 (<0.0001)	0.0037	11.6502 (<0.0001)	0.7722	0.9348	1.6225	1.3120	0.2559
	Korf	41.1095 (<0.0001)	3.2831	11.3164 (<0.0001)	1.0129	0.6949 (<0.0001)	0.0485	0.9510	1.4074	1.0828	0.1383
	Gompertz	21.7925 (<0.0001)	0.3299	3.3721 (<0.0001)	0.1156	0.0706 (<0.0001)	0.0024	0.9461	1.4761	1.1706	0.1881
	Chapman-Richards	23.6574 (<0.0001)	0.5455	0.0459 (<0.0001)	0.0029	1.7276 (<0.0001)	0.0914	0.9504	1.4154	1.0956	0.1468
	Weibull	22.8045 (<0.0001)	0.5183	0.0082 (<0.0001)	0.0009	1.4080 (<0.0001)	0.0417	0.9492	1.4326	1.1195	0.1621
<i>Larix kaempferi</i> (n = 321)	Logistic	24.6783 (<0.0001)	0.4069	0.1341 (<0.0001)	0.0063	9.8793 (<0.0001)	0.8678	0.8995	2.4378	1.9596	0.2788
	Korf	48.1994 (<0.0001)	5.3799	9.2619 (<0.0001)	1.0353	0.6928 (<0.0001)	0.0697	0.9156	2.2340	1.7319	0.1806
	Gompertz	26.0162 (<0.0001)	0.5230	3.1811 (<0.0001)	0.1575	0.0873 (<0.0001)	0.0044	0.9095	2.3140	1.8123	0.2247
	Chapman-Richards	28.1977 (<0.0001)	0.9276	0.0570 (<0.0001)	0.0055	1.6441 (<0.0001)	0.1279	0.9142	2.2528	1.7540	0.2001
	Weibull	27.2797 (<0.0001)	0.9147	0.0133 (<0.0001)	0.0019	1.3617 (<0.0001)	0.0599	0.9132	2.2658	1.7651	0.2093

Table 5. Model parameters of basal area growth functions and indicators of fit.

Species	Variables	Parm <i>a</i>		Parm <i>b</i>		Parm <i>c</i>		Fit Statistics			
		Estimate	S.E.	Estimate	S.E.	Estimate	S.E.	R ²	RMSE	MAE	MAPE
<i>Pinus densiflora</i> (<i>n</i> = 303)	Logistic	0.1222 (<0.0001)	0.0075	0.0824 (<0.0001)	0.0054	40.6221 (<0.0001)	6.0108	0.7974	0.0132	0.0098	9.3810
	Korf	1.2308 (0.2652)	1.1027	20.7981 (0.0001)	5.3120	0.5068 (0.0005)	0.1447	0.8092	0.0128	0.0090	1.5625
	Gompertz	0.1578 (<0.0001)	0.0161	5.3828 (<0.0001)	0.3859	0.0384 (<0.0001)	0.0039	0.8067	0.0129	0.0093	4.8037
	Chapman-Richards	0.2152 (<0.0001)	0.0486	0.0215 (0.0002)	0.0056	2.6596 (<0.0001)	0.3882	0.8088	0.0128	0.0091	2.3751
	Weibull	0.1632 (<0.0001)	0.0281	0.0002 (0.0081)	0.0001	2.0586 (<0.0001)	0.1441	0.8079	0.0128	0.0092	3.0804
<i>Pinus koraiensis</i> (<i>n</i> = 324)	Logistic	0.1300 (<0.0001)	0.0043	0.0902 (<0.0001)	0.0043	37.9451 (<0.0001)	4.1735	0.9000	0.0112	0.0082	9.7820
	Korf	0.7276 (0.0153)	0.2985	22.6280 (<0.0001)	4.5783	0.6046 (<0.0001)	0.1010	0.9141	0.0104	0.0068	0.6608
	Gompertz	0.1581 (<0.0001)	0.0085	5.3982 (<0.0001)	0.2984	0.0449 (<0.0001)	0.0030	0.9103	0.0106	0.0073	4.2698
	Chapman-Richards	0.1992 (<0.0001)	0.0226	0.0272 (<0.0001)	0.0042	2.7702 (<0.0001)	0.2860	0.9131	0.0104	0.0070	1.6581
	Weibull	0.1597 (<0.0001)	0.0145	0.0003 (0.0002)	0.0001	2.0530 (<0.0001)	0.1011	0.9118	0.0105	0.0072	2.5677
<i>Larix kaempferi</i> (<i>n</i> = 314)	Logistic	0.1048 (<0.0001)	0.0093	0.0976 (<0.0001)	0.0086	35.2623 (<0.0001)	5.8297	0.7502	0.0137	0.0089	10.6329
	Korf	9.3665 (0.7741)	32.6058	14.8994 (<0.0001)	0.6978	0.2927 (0.1204)	0.1880	0.7567	0.0135	0.0082	2.1481
	Gompertz	0.1420 (<0.0001)	0.0238	4.9745 (<0.0001)	0.3850	0.0440 (<0.0001)	0.0066	0.7550	0.0136	0.0085	6.1127
	Chapman-Richards	0.2583 (0.1026)	0.1578	0.0178 (0.1013)	0.0108	2.1727 (<0.0001)	0.4391	0.7567	0.0136	0.0083	2.9777
	Weibull	0.1616 (0.0201)	0.0691	0.0004 (0.0039)	0.0001	1.9114 (<0.0001)	0.2065	0.7567	0.0135	0.0083	3.3699

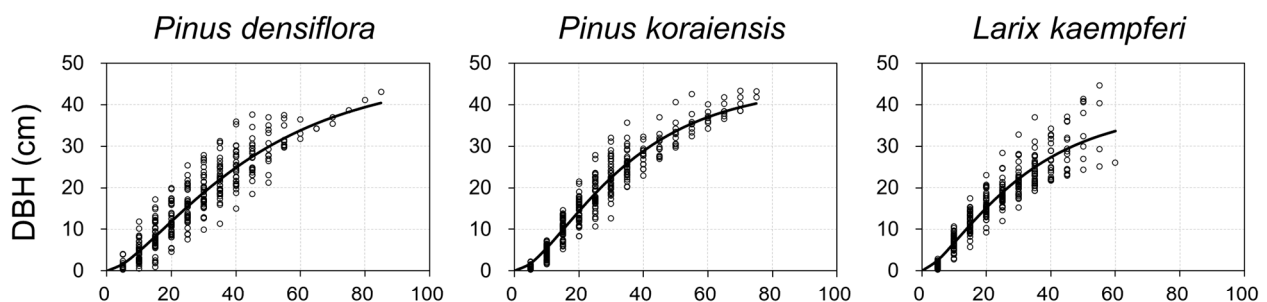
In addition, goodness-of-fit was examined with R², RMSE, MAE, and MAPE. Some of the models presented the best-fit statistics (Tables 3–6). The functions with best-fit statistics differed among variables and species, even though the metrics were comparable in some cases. Residual plots were also plotted over the predicted values and independent variable, which was tree age. No model showed an unusual or biased residual trend in variables and species (Supplementary files). When the predicted growth curve regarding total increment was graphically illustrated with observed sample points, the predicted lines over age generally traversed in the center of observations, and this characteristic represented the reasonable model fit. The biological concepts were examined assuming that tree size followed a sigmoid growth, and the CAI and MAI intersected at a mature stage. The prediction suitability, such as asymptote, growth rate, and inflection point, was evaluated for the final model selection in addition to the statistical assessment (Supplementary files). Considering all these evaluations, namely parameter estimates, fit statistics, residuals, and prediction suitability, we selected the best growth function for each variable. Consequently, the selected models were Chapman-Richards function for DBH and stem volume, Korf function for height, and Gompertz function for basal area.

Table 6. Model parameters of stem volume growth functions and indicators of fit.

Species	Variables	Parm <i>a</i>		Parm <i>b</i>		Parm <i>c</i>		Fit Statistics			
		Estimate	S.E.	Estimate	S.E.	Estimate	S.E.	R ²	RMSE	MAE	MAPE
<i>Pinus densiflora</i> (<i>n</i> = 303)	Logistic	1.2895 (<0.0001)	0.085	0.0899 (<0.0001)	0.0061	88.0312 (<0.0001)	16.9867	0.7866	0.1326	0.0891	6.8688
	Korf	8.2089 (0.1461)	5.6334	43.3596 (0.0265)	19.4449	0.7223 (0.0001)	0.1831	0.7957	0.1298	0.0817	1.0726
	Gompertz	1.7361 (<0.0001)	0.204	7.1821 (<0.0001)	0.6973	0.0394 (<0.0001)	0.0043	0.7946	0.1301	0.0832	2.6467
	Chapman-Richards	2.1471 (<0.0001)	0.4443	0.0268 (<0.0001)	0.0060	4.0338 (<0.0001)	0.7078	0.7955	0.1298	0.0820	1.4566
	Weibull	1.5350 (<0.0001)	0.2069	0.00002 (0.0767)	0.00001	2.6495 (<0.0001)	0.1816	0.7944	0.1302	0.0832	2.1675
<i>Pinus koraiensis</i> (<i>n</i> = 323)	Logistic	1.6238 (<0.0001)	0.0694	0.0844 (<0.0001)	0.0037	64.6695 (<0.0001)	6.4897	0.9200	0.1030	0.0756	12.2879
	Korf	45.0916 (0.299)	43.3502	22.8314 (<0.0001)	3.4632	0.444 (<0.0001)	0.0999	0.9337	0.0937	0.0591	0.8214
	Gompertz	2.3206 (<0.0001)	0.1952	6.2032 (<0.0001)	0.2834	0.0358 (<0.0001)	0.0026	0.9299	0.0964	0.0649	4.8052
	Chapman-Richards	3.8923 (0.0001)	0.9583	0.0175 (<0.0001)	0.0040	2.9132 (<0.0001)	0.2992	0.9328	0.0944	0.0613	1.7015
	Weibull	2.5051 (<0.0001)	0.4821	0.00005 (<0.0001)	0.00001	2.2834 (<0.0001)	0.1106	0.9318	0.0951	0.0635	2.5439
<i>Larix kaempferi</i> (<i>n</i> = 313)	Logistic	1.2817 (<0.0001)	0.1067	0.1126 (<0.0001)	0.0092	81.0024 (<0.0001)	17.1789	0.7870	0.1495	0.0902	12.8173
	Korf	21.2675 (0.5182)	32.8801	25.1394 (0.0083)	9.4589	0.5363 (0.0164)	0.2222	0.7885	0.1489	0.0834	0.5679
	Gompertz	1.8400 (<0.0001)	0.3157	6.7796 (<0.0001)	0.7206	0.0475 (<0.0001)	0.0068	0.7892	0.1487	0.0850	4.3490
	Chapman-Richards	2.5000 (0.0066)	0.9134	0.0293 (0.0048)	0.0103	3.5893 (<0.0001)	0.7982	0.7891	0.1487	0.0835	1.2202
	Weibull	1.4885 (<0.0001)	0.2979	0.00003 (0.1138)	0.00002	2.6557 (<0.0001)	0.2351	0.7899	0.1484	0.0837	1.9647

3.2. Growth Simulation and Characteristics

After determining the most suitable model for each variable, the total increment, CAI, and MAI were visually displayed to examine growth patterns over age and compare them between species (Figures 2–5). Although the predicted sigmoid growth curves traversed the observed sample points properly, it should be noted that some trees could have a larger tree size than the estimated asymptote because the parameters were fitted for unbiased, accurate predictions (Figure 2).

**Figure 2.** Cont.

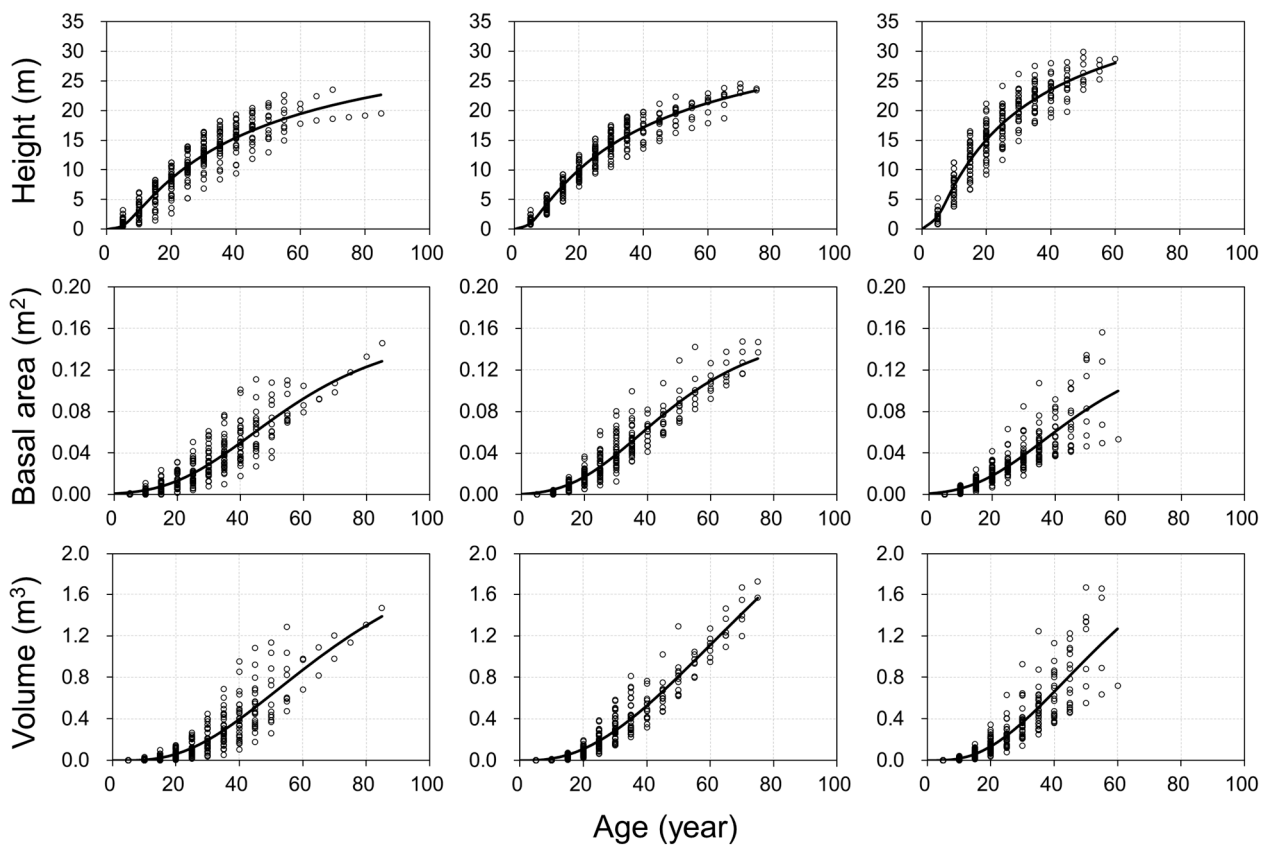


Figure 2. Predicted growth curves and observations of tree size variables by species. The best models by variables were based on fit statistics and model performance; the selected growth functions were Chapman-Richards for DBH and stem volume, Korf for height, and Gompertz for basal area.

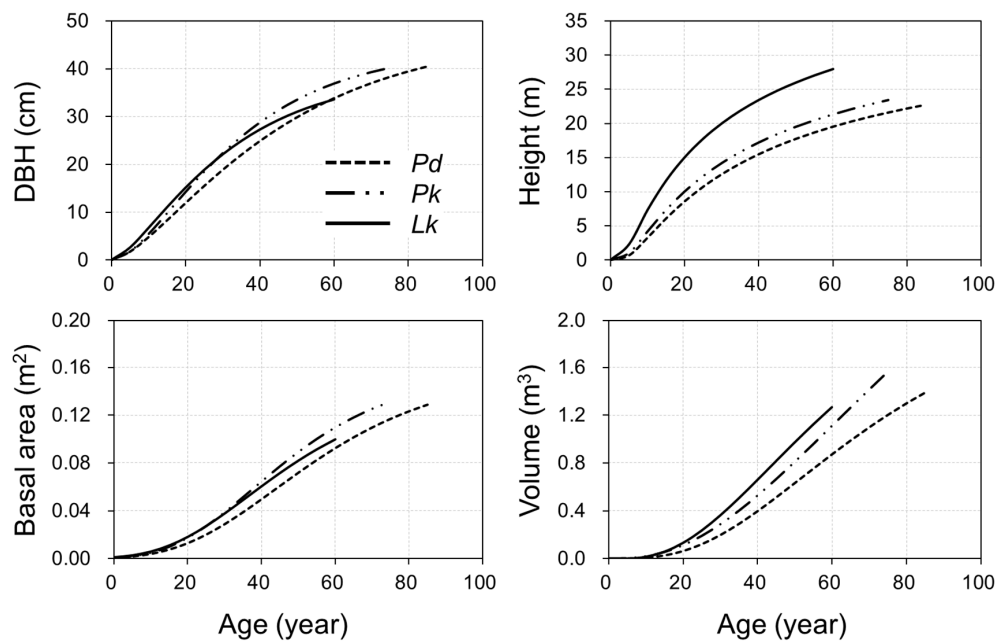


Figure 3. Predicted growth curves of DBH, height, basal area, and stem volume for species within the age range of modeling data. The best models by variables are based on fit statistics and model performance of the results; the selected growth functions are Chapman-Richards for DBH and stem volume, Korf for height, and Gompertz for basal area. *Pd*: *Pinus densiflora*. *Pk*: *Pinus koraiensis*. *Lk*: *Larix kaempferi*.

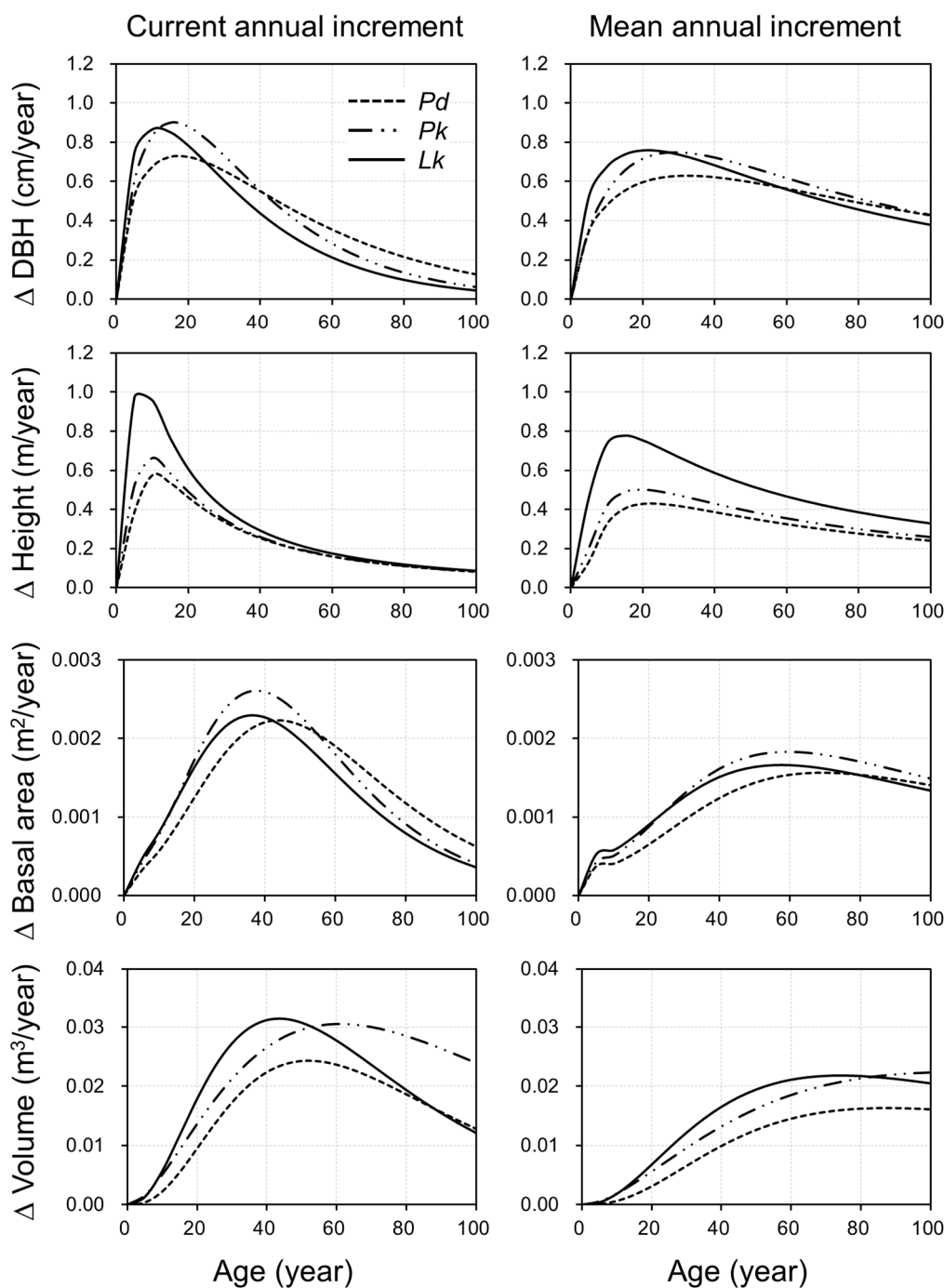


Figure 4. Simulated current annual increment and mean annual increment of DBH, height, basal area, and stem volume in different species. The best models by variables are based on fit statistics and model performance of the results; the selected growth functions are Chapman-Richards for DBH and stem volume, Korf for height, and Gompertz for basal area.

The age at maximum growth differed according to the variable and species. The total increment in DBH in *Lk* followed a similar pattern to that in *Pk* until 40 years of age, but later, *Pk* was the largest among species (Figure 3). The age at which the DBH reached a diameter of 30 cm was 50 years for *Pd*, 42 years for *Pk*, and 47 years for *Lk*. In contrast, *Lk* constantly showed a superior growth in height compared to *Pd* and *Pk*. The height at 40 years of age, an average felling age for the species in South Korea, was 15 m for *Pd*, 17 m for *Pk*, and 23 m for *Lk*. The inflection point in both the DBH and height curve was situated at an early age, and then the growth decreased reaching the asymptote, which is

concave with respect to the Y-axis as reported in many other studies on DBH and height growth [30–34].

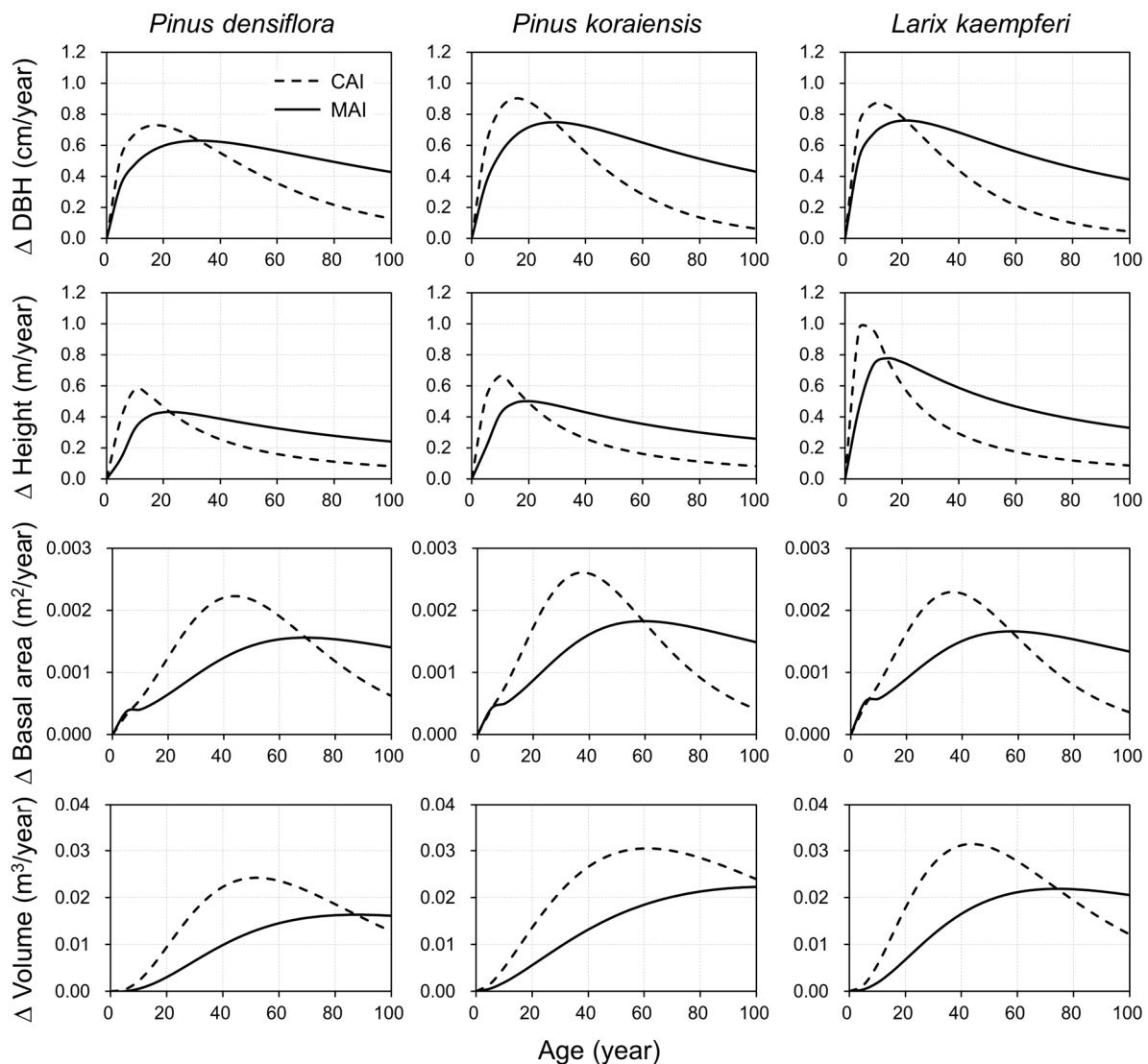


Figure 5. Simulated annual increment by tree size variables and species to examine current annual increment (CAI) and mean annual increment (MAI). The best models by variables are based on fit statistics and model performance of the results; the selected growth functions are Chapman-Richards for DBH and stem volume, Korf for height, and Gompertz for basal area.

The order of basal area by species was similar to that of DBH, but the basal area of *Pk* was largest after 35 years of age, followed by that of *Lk* and *Pd*. However, the volume curve was distinct from that of the other variables. The volume was larger in *Lk* than in *Pd* and *Pk* from an early age, and this trend continued (Figure 3). The age with 1 m³ of stem volume was the shortest, with 51 years for *Lk*, followed by 56 years for *Pk* and 65 years for *Pd*. Moreover, the volume presented vigorous growth even after the age at the inflection point of other variables, which stood for concave shape with respect to the Y-axis. In other words, the asymptote pattern of the basal area appeared later, at nearly 80 years of age, and that of the volume was not clearly defined until 100 years of age (Figure 3).

This concurs with the results of previous studies on volume growth [35–40]. Additionally, the one-dimensional growth pattern, such as DBH and height, was similar to that reported in previous studies [13–16]. However, the growth of a two- (basal area) or three-dimensional size (stem volume) exhibited a constant incremental pattern as the

sigmoidal asymptote was not apparent and the pattern was further prolonged. It should be noted that the growth characteristics in our study were analyzed at the tree level, and carrying capacity was not considered at the stand level.

The growth characteristics, such as maximum growth and inflection point, became clearer when the CAI and MAI were analyzed (Figure 4). In the CAI analysis, the age with maximum increment differed among species, and it was in the range of 10–17 years for DBH, 5–10 years for height, 36–44 years for basal area, and 45–61 years for stem volume (Figure 4). The order of age with maximum increment was the same in MAI, but the culmination of MAI for each variable occurred at an older age than that of CAI in all species (Figure 4). Thus, the age with maximum MAI among species was 21–32 years for DBH, 15–22 years for height, 58–69 years for basal area, and 74–106 years for stem volume.

When CAI and MAI were analyzed by variables and species, the growth pattern and characteristics were evident in regard to the culmination, intersecting point, and its age (Figure 5). For DBH and height growth, the age of the intersection point was earliest in *Lk*, followed by that in *Pk* and *Pd*. For basal area growth, the age intersection points by species were in the same order as that for DBH and height growth, although these points were reached at a later age. Lee et al. [41] reported that CAI of *Pd* in DBH was mostly less than 0.5 cm year^{-1} after 40 years of age, which was analogous to our results. In volume growth, the age at the intersection point was different from the other variables: 86 years for *Pd*, 106 years for *Pk*, 74 years for *Lk*. This indicated that stem volume growth before 70 years of age was incremental at individual trees level (Figure 5). Similar results were reported in studies on the stem volume growth of Japanese cedar (*Cryptomeria japonica* (Thunb. ex L.f.) D. Don) [42] and basal area growth of Norway spruce (*Picea abies* (L.) H. Karst.) [39]. This resulted from the three-dimensional growth of stem volume being influenced by both basal area and height.

The CAI and MAI of the stem volume can be different from those of the basal area because the stem volume is a three-dimensional growth that combines DBH and height. Consequently, the age at the intersection of CAI and MAI varied by tree size variable and species in our study (Figure 5). This is considered the usual increment trend of the major coniferous species in South Korea. Nonetheless, it is noteworthy that the age for maximum growth of an individual tree does not conform with the age at the stand level. This is because the stand dynamics, including between-tree competition for resources, mortality, and ingrowth, were not taken into account [11]. Therefore, long-term monitoring with repeated measurements using permanent plots is considered indispensable and must be carried out persistently to study the growth and yield of a forest stand.

3.3. Model Evaluation and Applicability

Although the best models fit well with significant parameters and are statistically supported (Tables 3–6), tree ages should be verified when predicting the dependent variables of tree size. The best model was validated as unbiased and accurate within the sample range. However, there is still uncertainty beyond the modeling range because the age range of samples collected was covered restrictively and differed among species (Table 1, Figures 4 and 5). The growth curves could be changed with additional data on the age range; thus, they should be carefully handled for extrapolation. The spatial range should be referenced for application as the samples were collected from the stands of each species in Gangwon and North Gyeongsang provinces of South Korea. Those regions are known as suitable and favorable sites in South Korea for *Pd*, *Pk*, and *Lk* [43].

Moreover, tree parameters, especially those related to diameter growth such as DBH, basal area, and stem volume, were regarded to be linked to stand density. Thus, the stand density should be referenced as the background when examining the growth status of sample trees. The trees sampled in the present study were dominant or codominant in the plots where the relative density was higher than 0.7. These plots represent dense stands, where the growth of biometric features, for example, diameter vs. height, may differ from that in low-density stands [24,25]. Nevertheless, the study plots cannot be

considered as unthinned stands because the history of silvicultural treatments and stand density during the entire period of tree growth is unknown. Even though some variations cannot be removed due to the inherent background of sample trees, the selected models were assessed to offer the representative growth curve for each tree size variable by species. Thus, the developed models in this study were considered to be helpful for understanding and managing the target species.

4. Conclusions

This study aimed to fit the model parameters and examine growth characteristics for major tree size variables of Pd , Pk , and Lk using the widely used sigmoid growth functions in forest biometrics. The selection of applied growth functions for all species was based on variables such as Chapman-Richards for DBH and stem volume, Korf for height, and Gompertz for the basal area. The final models revealed the applicable parameter estimates and properly depicted the overall growth pattern and characteristics of the trees. Contrary to DBH and height, the total increment of stem volume showed convex growth curves with respect to the Y-axis, suggesting that the volume pattern in three-dimensional growth was different from the one-dimension growth of DBH and height. Additionally, the age with maximum CAI and MAI was different among species and variables. The earliest age by variable was height, followed by DBH and basal area, and volume.

When using the models, especially for DBH, basal area, and volume, the applied stand density should be referenced because the size can vary with stand density. The temporal and spatial range should be noted for unbiased estimation, for example, for tree age and study area in the data, because it could lead to extrapolation and possibly unstable prediction outside the modeling range. The developed models were based on the growth characteristics of a tree in each stand; thus, they do not represent growth and yield characteristics at the stand level. Even though the sample data may not include tree growth characteristics from all stand conditions, our results provided the general growth patterns appropriately for the target species. Therefore, our model parameters and simulated growth patterns can be used as references to predict tree size by variables and understand growth characteristics for Pd , Pk , and Lk , particularly in South Korea.

Supplementary Materials: The following supporting information can be downloaded at: <https://www.mdpi.com/article/10.3390/f14010115/s1>, Figure S1. Residual plots of DBH models over predicted values by species; Figure S2. Residual plots of height models over predicted values by species; Figure S3. Residual plots of basal area models over predicted values by species; Figure S4. Residual plots of stem volume models over predicted values by species; Figure S5. Total increment of DBH by model type in each species; Figure S6. Total increment of height by model type in each species; Figure S7. Total increment of basal area by model type in each species; Figure S8. Total increment of stem volume by model type in each species; Figure S9. Comparison of current annual increment (CAI) and mean annual increment (MAI) by DBH model type in each species; Figure S10. Comparison of current annual increment (CAI) and mean annual increment (MAI) by height model type in each species; Figure S11. Comparison of current annual increment (CAI) and mean annual increment (MAI) by basal area model type in each species; Figure S12. Comparison of current annual increment (CAI) and mean annual increment (MAI) by stem volume model type in each species.

Author Contributions: Conceptualization, Y.S., D.L. and J.C.; methodology, Y.S. and D.L.; investigation, Y.S. and D.L.; data curation, D.L.; formal analysis, Y.S. and D.L.; software, Y.S. and D.L.; Resources, Y.S., D.L. and J.C.; Validation, Y.S., D.L. and J.C.; Visualization, D.L.; writing—original draft, Y.S. and D.L.; writing—review and editing, Y.S., D.L. and J.C.; Supervision, J.C.; funding acquisition, J.C.; project administration, J.C. All authors have read and agreed to the published version of the manuscript.

Funding: This study was carried out with the support of R&D Program for Forest Science Technology (Project No. 2022464B10-2224-0201) provided by Korea Forest Service (Korea Forestry Promotion Institute) and with the support of the Basic Science Research Program through the National Research Foundation of Korea (NRF) funded by the Korean Ministry of Education (Grant No. NRF-2016R1D1A1B02011648) in 2023.

Data Availability Statement: Not applicable.

Acknowledgments: Financial support for the data collection was provided by the National Forest Management Division of the Korea Forest Service. The Forest Resource Monitoring Center on Climate Change (FRMCCC) at the Kangwon National University provided data maintenance support.

Conflicts of Interest: The authors declare no conflict of interest. The funders had no role in the design of the study; in the collection, analyses, or interpretation of data; in the writing of the manuscript; or in the decision to publish the results.

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