

## Selection of Growth Projection Intervals for Improving Parameter Estimation of Stand Growth Model<sup>1</sup>

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林分生長 모델의 母數 推定 能力 向上을 爲한 生長 測定間隔의 選擇<sup>1</sup>

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### ABSTRACT

This study aimed to provide a strategy for selecting an adequate combination of growth intervals(i. e. times between age  $T_1$  and age  $T_2$ ) to be used to improve the reality of the growth equation through obtaining better precision of parameter estimates. Variety of growth functions were fitted to the data and one equation which best fitted the data was chosen for the analysis. A modified Schumacher projection equation, selected as a best equation, that included dummy variables representing locality as a predictor variable was fitted for basal area and height equations with nonoverlapping growth interval and all possible growth interval data sets of Douglas-fir(*Pseudotsuga menziesii* Mirb.Franco). The data were measured in all parts of the South Island of New Zealand. It was found that the precision of parameter estimates was increased in both basal area and height equations by using data set which contained a range of measurement intervals from short to long term.

*Key words*: Dummy variables, Basal area model, Mean top height model, *Pseudotsuga menziesii* Mirb.Franco, Schumacher projection equation.

### 要 約

본 연구는 보다 정확한 母數 추정을 통한 生長모델의 현실성을 향상시키는데 이용되는 生長 측정 간격(임목의 측정 초기 연령  $T_1$ 과 재측정 연령  $T_2$ 의 기간)의 적합한 조합을 선택하기 위한 계획을 제 공하는데 목적이 있다.

다양한 生長식을 데이터에 적용한 후 가장 적합한 것으로 판정된 生長식을 분석에 이용하였다. 여 러 生長식을 분석한 결과 최적의 生長식으로 판명된 더미 변수를 포함하는 변형 Schumacher 방정식 을 임분 胸高斷面積 生長식과 平均樹高 生長식을 얻기 위하여 이용하였다. 그리고 사용된 자료는 뉴 질랜드 남섬 전역에서 측정된 美松(*Pseudotsuga menziesii* Mirb.Franco)의 生長 측정기간이 변형되지 않은 데이터와 모든 가능한 生長 측정기간을 포함하는 변형된 2종류의 데이터이었다. 단기의 측정기 간에서부터 장기의 측정기간의 범위를 포함하는 데이터(모든 가능한 生長 측정기간을 포함하는 데이 터)를 사용할 때 흉고단면적 生長식과 임분 평균수고 生長식에서 모수 추정의 정확성이 증가되는 것 이 발견되었다.

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**INTRODUCTION**

Forest growth is defined as the increase in diameters of one or more individuals in a forest stand over a given period time(e. g. growth in m<sup>3</sup>/ha/year). Yield refers to the total production over a given time period(e. g. volume in m<sup>3</sup>/ha). Yield is the directly related to growth in that it is the accumulated growth to a specified point in time, and can be derived mathematically by integrating a growth function. The derivative of a yield function will therefore give a growth function(Vanclay, 1994).

The general purpose of growth and yield models, despite their complexity of structure, can be explained simply as ; given a set of stand or tree characteristics, such as basal area and stems per hectare at one point in time (T<sub>1</sub>), to predict by how much these characteristics will have changed at a future time (T<sub>2</sub>) given specified stand or tree treatments.

Growth interval data obtained from remeasured permanent plots or trees are referred to as repeated measurements or a real growth series. The term means that *N* experimental subjects are observed in each of *k* successive occasions that possibly correspond to different experimental conditions, the *i*<sub>th</sub> subject yielding *y*<sub>*ij*</sub> on the *j*<sub>th</sub> occasion. Such repeated measurements data can be used for investigating tree dynamics as well as for modeling growth.

The effective expression of the existing relationship between growth and yield was first presented by Clutter(1963) and has been referred to algebraic difference equation(Borders et al., 1984). That is

$$Y_2 = f(Y_1, T_1, T_2, \theta, MR) \tag{1}$$

where

*Y*<sub>2</sub> = value of a continuous variable derived for a tree or stand at age *T*<sub>2</sub>

*Y*<sub>1</sub> = value of the same variable at initial measurement

*T*<sub>1</sub> = tree or stand age at initial measure-

ment

*T*<sub>2</sub> = tree or stand age at next re-measurement

*θ* = set of parameters of equation

and MR = management regime.

In this approach, at a given time, the future state of the variables and the transition functions or changes in the state variable are a function of the initial state of the variable, time elapsed, management inputs and prevailing environment. In using projection equation of this function form, growth series data can be used in fitting routines to obtain sample estimates of parameters of equation that best described the growth and yield of the selected stand variables.

The most common procedure for establishing parameters in algebraic differential equations is to use only the nonoverlapping growth interval (i.e. A<sub>1</sub> to A<sub>2</sub>, A<sub>2</sub> to A<sub>3</sub>, --- A<sub>*n*-1</sub> to A<sub>*n*</sub>) (Borders et al., 1984 ; Cluter, 1963 ; Sullivan and Clutter, 1972). Another possibility is to use all possible intervals for each unit. If permanent sample plots have been measured *n* times there are *<sup>n</sup>C*<sub>2</sub> combinations of different intervals between time *T*<sub>1</sub> and *T*<sub>2</sub> that can be derived and used to build equations.

This paper was intended to compare and contrast the two data sets formulations namely, Nonoverlapping growth interval and all possible growth interval, to provide a strategy for selecting an appropriate combination of projection intervals.

**MATERIALS AND METHODS**

Data for this study came from a large database of Douglas-fir(*Pseudotsuga menziesii* Mirb. Franco) permanent sample plots maintained by New Zealand Forest Research Institute. The data were measured in all parts of the South Island of New Zealand. All of 355 permanent sample plots ranging from 0.02 to 0.04 ha and 1844 sets of measurements were used in this analysis. Table 1 is the summary of the mean and extreme values of age, mean top height and basal area per hectare of the raw data.

**Table 1.** Summary of mean and extreme values extracted from permanent sample plots data

Region	No. of observations	Variable*	Mean	Minimum	Maximum
Canterbury	241	Age(years)	32.8	9.0	61.0
		H <sub>100</sub> (m)	22.9	2.9	39.3
		Altitude(m)	326.1	150.0	790.0
		G(m <sup>2</sup> /ha)	46.3	0.43	116.2
Nelson	929	Age(years)	27.5	7.0	58.0
		H <sub>100</sub> (m)	22.9	5.6	47.8
		Altitude(m)	438.1	183.0	625.0
		G(m <sup>2</sup> /ha)	42.2	1.2	109.4
Southland	449	Age(years)	33.6	7.0	78.0
		H <sub>100</sub> (m)	24.1	4.1	47.4
		Altitude(m)	251.1	50.0	625.0
		G(m <sup>2</sup> /ha)	51.3	1.1	141.7
Westland	225	Age(years)	26.9	5.0	59.1
		H <sub>100</sub> (m)	18.8	1.9	37.5
		Altitude(m)	229.0	0.0	330.0
		G(m <sup>2</sup> /ha)	29.9	0.01	123.8
Sum	1844	-	-	-	-

\* H<sub>100</sub> = mean top height of the stand

G = net basal area of the stand

The two data sets were created from raw data base namely, nonoverlapping interval data set that consisted of about 1600 sets of measurements and all possible interval data set which had 6173 sets of measurements in through lag and put statements in Statistical Analyses System(SAS).

Prior to any model estimation the data were verified and screened to ascertain the reliability of the data. Examples of data validation included that ensuring number of stems/ha at age<sub>2</sub> were not greater than that of at age<sub>1</sub>, basal area/ha at age<sub>2</sub> were greater than that of at age<sub>1</sub> and age<sub>2</sub> were greater than age<sub>1</sub>. Residuals were also used to detect outliers. Outliers were observations that had residuals greater than  $\pm 3.5$  standard deviation from zero(Xu, 1990). Standard residuals S<sub>i</sub> defined as

$$S_i = \frac{\text{Residual}}{\text{MSE}} \quad (2)$$

where in (2)

Residual=(Observed value) - (Fitted value) and  
MSE = Residual mean square

As a rule, in all the model fitting routines, observations which have value of S<sub>i</sub> greater than  $\pm 3.5$  were considered to be outliers.

The main standard analytical procedures used was non-linear ordinary least-squares regression. Regression equation can be fitted variables sets of any sort, but in this study it was ensured that the dependent and independent variables conform to biologically and mathematically sound relationship, the functions used were of appropriate form to represent the independent relationship and parameter estimates were free of apparent bias.

In order to apply any regression to data sets of variables, it is assumed that the residual errors were independent, had a mean of zero and constant variance, and plot of residuals followed a normal distribution(Sokal and Rohlf, 1981; Draper and Smith, 1981). Variety of sigmoid shaped functions were applied to the data sets using the

PROC NLIN procedure of the SAS package(SAS institute Inc., 1990) and the derivative free algorithm method for non-linear least squares(Ralston and Jennrich, 1978) to find which function could be used to compare the two data sets.

Anamorphic and polymorphic forms of Schumacher(Clutter *et al.*, 1983), Chapman-Richard(Pienaar and Turnbull, 1973), Hossfeld(Xu, 1990) and Gompertz equations were fitted to two data sets. The equation formulations used are listed in Table 2.

The residuals resulting from fitting each of these equations were analyzed using following methods :

1. comparison of mean square errors(MSE) ;
2. examination of plots of residuals against predictor variables and predicted values to provide ocular estimates of their normality of errors ; and
3. comparison of extreme deviation and moments of the residual on the assumption that they should be normally and independently distributed with mean zero and constant variance  $\delta^2$ .

PROC UNIVARIATE(SAS institute Inc., 1990) procedure was also used to ascertain the goodness of the equations and normality because this procedure provides a wide range of statistics to supplement the residual patterns inferences. After fitting equation chosen as a best model

with two data sets, coefficients of parameters were compared.

## RESULTS AND DISCUSSION

Two compatible projection equations for basal area/ha and mean top height were derived and compared.

### 1. Basal Area Model

Most of the anamorphic equations were found large bias in residuals pattern. The Chapman-Richard functions displayed difficulty in convergence and once parameters were estimated bias was noted in the graphical representation of residuals. The coefficients of the general equations fitted to the data are presented in Table 3 and 4 with respective mean square error(MSE) value.

Since the equation with the least biased residual plots was found the lowest MSE value, the values of this in Table 3 and 4 were used to indicate the best fitting equation. The Schumacher polymorphic function with MSE of 4.15 and 19.10 for nonoverlapping and all possible data sets respectively, were found to give a better fit than the rest of the equations. This equation, therefore, was considered for further examination.

After trying numerous modifications to the Schumacher equation, with the addition and sub-

**Table 2.** Equation forms applied to data sets

Equation name	Equation Forms*
Hossfeld Polymorphic	$Y_2 = 1/((1/Y_1) (T_1/T_2)^\gamma + (1/\alpha) (1 - (T_1/T_2)^\gamma))$
Hossfeld Anamorphic	$Y_2 = 1/((1/Y_1) + \theta (1/T_2^\beta - T_1^\beta))$
Schumacher Polymorphic	$Y_2 = \exp(\ln(Y_1) (T_1/T_2)^\beta + \alpha (1 - (T_1/T_2)^\beta))$
Schumacher Anamorphic	$Y_2 = Y_1 \exp(\beta (1/T_1^\alpha - 1/T_2^\alpha))$
Chapman-Richards Polymorphic	$Y_2 = (\alpha/\gamma)^{1/(1-\beta)} (1 - (1 - (\gamma/\alpha) Y_1^{(1-\beta)}) \exp(-\gamma(1-\beta)(T_2 - T_1)))^{1/(1-\beta)}$
Chapman-Richards Anamorphic	$Y_2 = Y_1 ((1 - \exp(-\beta T_1)) / (1 - \exp(-\beta T_2)))^{1/(1-\gamma)}$
Gompertz polymorphic	$Y_2 = \exp(\ln(Y_1) \exp(-\beta(T_2 - T_1) + \gamma(T_2^2 - T_1^2) + \alpha(1 - \exp(-\beta(T_2 - T_1) + \gamma(T_2^2 - T_1^2))))$

\*  $Y_1$  = net basal area of the stand or mean top height at age  $T_1$   
 $Y_2$  = net basal area of the stand or mean top height at age  $T_2$   
 $\alpha, \beta, \gamma$  and  $\theta$  are parameters to be estimated

**Table 3.** Coefficients for general equation fitted to nonoverlapping basal area data

Model Name	Parameters				MSE
	$\alpha$	$\beta$	$\gamma$	$\theta$	
Hossfeld Polymorphic	112.65	-	2.53	-	4.77
Hossfeld Anamorphic	-	1.48	-	1.36	21.79
Schumacher Polymorphic	5.11	1.08	-	-	4.15
Schumacher Anamorphic	0.81	19.94	-	-	8.60
Chapman-Richards Polymorphic	1.05	0.36	0.04	-	8.87
Chapman-Richards Anamorphic	-	0.002	1.79	-	24.20
Gompertz Polymorphic	4.93	0.09	0.006	-	5.83

**Table 4.** Coefficients for general equation fitted to all possible basal area data

Model Name	Parameters				MSE
	$\alpha$	$\beta$	$\gamma$	$\theta$	
Hossfeld Polymorphic	100.45	-	2.84	-	21.11
Hossfeld Anamorphic	-	1.66	-	1.72	189.35
Schumacher Polymorphic	4.98	1.17	-	-	19.10
Schumacher Anamorphic	0.82	18.62	-	-	81.88
Chapman-Richards Polymorphic	0.91	0.48	0.07	-	29.12
Chapman-Richards Anamorphic	-	0.004	1.59	-	191.97
Gompertz Polymorphic	4.79	0.09	0.006	-	20.02

traction of various predictor variables and alteration to the equation form, the modified Schumacher polymorphic equation (3) that included dummy variables representing locality was found to give the best fit for the both two data sets.

$$G_2 = G_1(T_1/T_2)^\beta ((\alpha + \beta_1k_1 + \beta_2k_2 + \beta_3k_3) / (1 - (T_1/T_2)^\beta)) \quad (3)$$

where in (3)

- $G_2$  = net basal area of the stand( $m^2/ha$ ) at  $T_2$
- $G_1$  = net basal area of the stand( $m^2/ha$ ) at  $T_1$
- $T_1$  = age in years at the beginning of a growth periods
- $T_2$  = age in years at the end of a growth period
- $K_1, K_2$  and  $K_3$  = dummy variables for region
- $\alpha, \beta, \beta_1, \beta_2$  and  $\beta_3$  = parameters to be estimated.

The goodness of the fit was evaluated through plots of residuals against predicted values as shown Fig. 1 and 2. And both are acceptable with no apparent bias.

Parameter estimates were slightly different between the two cases as shown in Table 5. The total regression and residual sums of square are seen to be much less for nonoverlapping intervals. This arises because there was only an annual interval in that case, while the all possible intervals had much wider scatter.

The t statistic, estimated by dividing the parameter estimate by asymptotic standard error, was much greater for all parameters included in the all possible interval model. That t statistic test provides indicative guidelines on the relative precision. In this case the all possible interval model estimates provide a better fit to the data. In addition, the confidence intervals for both models show that all estimates are significant at  $p < 0.05$  as none includes zero, the 95% confidence intervals for the all possible model are much tighter.

## 2. Mean Top Height Model

A range of growth equations in different form were fitted to the data sets and analyzed. Anamorphic Schumacher, polymorphic Schumacher and polymorphic Hossfeld equations were found

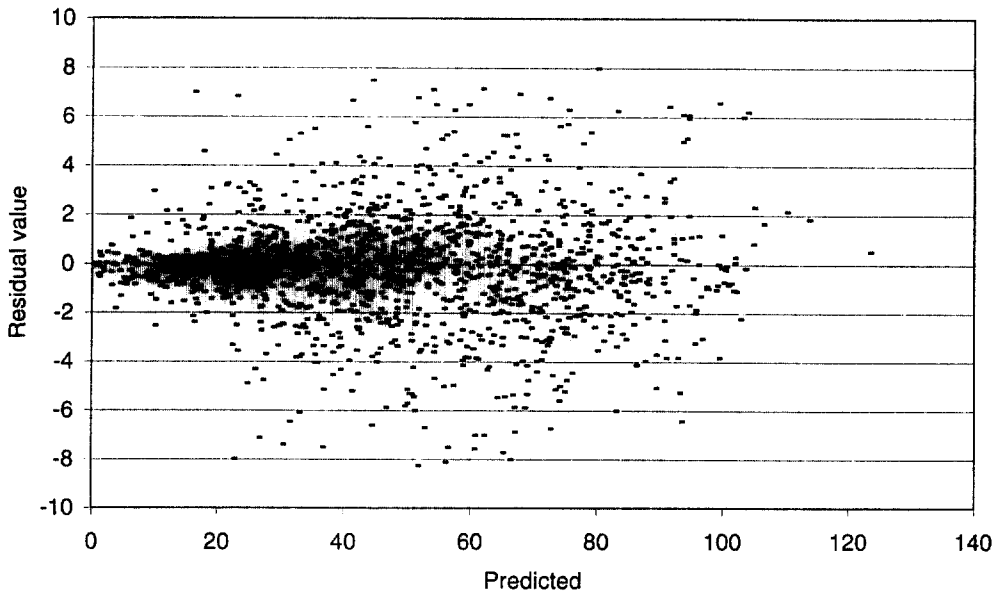


Fig. 1. Plot of residual Vs predicted for nonoverlapping intervals basal area equation

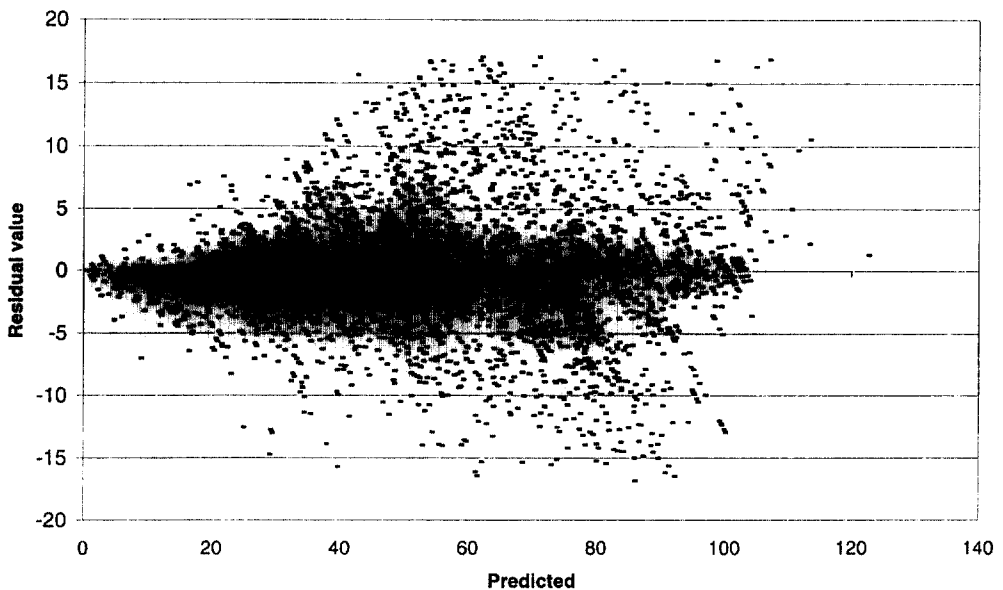


Fig. 2. Plot of residual Vs predicted for all possible intervals basal area equation

to give a better fit than rest of the equations based on analysis of residuals. The coefficients and the residual mean square error for seven candidate equations are presented in the Table 6 and 7.

The Schumacher polymorphic equation with mean square error(MSE) of 0.59 and 0.96 for

nonoverlapping and all possible data sets respectively, were found to give the best fit after comparing residual patterns, mean residual error and PROC UNIVARIATE statistics. Therefore, this equation was chosen for further improvement by incorporating several explanatory variables in a logical manner. After modification of this equa-

**Table 5.** Non linear least squares summary statistics for basal area models

SOURCE	Degree of Freedom		Sums of Squares		Mean Squares	
	Nonoverlapping	All possible	Nonoverlapping	All possible	Nonoverlapping	All possible
Regression	5	5	4,508,417	1,7528,522	901,683	3,505,704
Residual	1,794	6,105	6,414	104,319	3.5755	17.088
Uncorrected total	1,799	6,110	4,514,831	1,7632,842		
Corrected total	1,798	6,109	1,017,708	3,650,294		

Parameter	Estimates		Std. Error / t statistics		95% Confidence Interval			
	Nonoverlapping	All possible	Nonoverlapping	All possible	Nonoverlapping		All possible	
					Lower	Upper	Lower	Upper
$\alpha$	4.9149	4.7274	0.0268 / 183.12	0.0166 / 284.05	4.86	4.97	4.69	4.76
$\beta$	1.0801	1.1904	0.0183 / 59.08	0.0083 / 143.95	1.04	1.12	1.17	1.20
$\beta_1$	0.1293	0.2143	0.0225 / 5.75	0.0158 / 13.55	0.09	1.17	0.18	0.24
$\beta_2$	0.3201	0.3506	0.0224 / 14.26	0.0167 / 21.02	0.28	0.36	0.32	0.38
$\beta_3$	0.3120	0.4063	0.0296 / 10.53	0.0210 / 19.39	0.25	0.37	0.37	0.45

**Table 6.** Coefficients for general equation fitted to nonoverlapping mean top height data

Model Name	Parameters				MSE
	$\alpha$	$\beta$	$\gamma$	$\theta$	
Hossfeld Polymorphic	74.80	-	1.39	-	0.61
Hossfeld Anamorphic	-	1.27	-	2.25	0.96
Schumacher Polymorphic	5.63	0.38	-	-	0.59
Schumacher Anamorphic	0.32	8.67	-	-	0.64
Chapman-Richards Polymorphic	0.56	0.32	0.03	-	0.69
Chapman-Richards Anamorphic	-	0.001	1.83	-	3.08
Gompertz Polymorphic	4.16	0.05	0.0003	-	0.67

**Table 7.** Coefficients for general equation fitted to all possible mean top height data

Model Name	Parameters				MSE
	$\alpha$	$\beta$	$\gamma$	$\theta$	
Hossfeld Polymorphic	77.04	-	1.44	-	0.98
Hossfeld Anamorphic	-	1.27	-	2.12	2.05
Schumacher Polymorphic	5.69	0.39	-	-	0.96
Schumacher Anamorphic	0.31	8.90	-	-	1.10
Chapman-Richards Polymorphic	0.71	0.28	0.04	-	1.43
Chapman-Richards Anamorphic	-	0.003	1.64	-	21.37
Gompertz Polymorphic	4.19	0.05	0.003	-	1.22

tion through adding and subtracting of explanatory variables, a modified Schumacher polymorphic equation (4) that include dummy variables representing locality gave the best fit for two data sets.

$$H_{100,2} = H_{100,1} (T_1/T_2)^{\beta} ((\alpha + \beta_1 k_1 + \beta_2 k_2) (1 - (T_1/T_2)^{\beta})) \quad (4)$$

where in (4)

$H_{100,2}$  = mean top height in meters at age  $T_2$   
 $H_{100,1}$  = mean top height in meters at age  $T_1$   
 $T_1$  = age in years at the beginning of a growth periods  
 $T_2$  = age in years at the end of a growth period  
 $K1, K2$  and  $K3$  = dummy variables for region  
 $\alpha, \beta, \beta_1$  and  $\beta_2$  = parameters to be estimated.

Altitude and dummy variables were found to improve the model when they were introduced independently to the basic Schumacher equation. Though altitude has been found to be an important variable for explaining variations in mean top height growth(Woollons and Hayward, 1985 ; Mason, 1992), it was not included in the final formulations. The reason is that the modification of the model through adding and subtracting of these two variables was not superior to equation including only dummy variables. Parameter estimates for both equations are summarized in Table 8.

All the parameter estimates were significant at least 5% level. Fig. 3 and 4 show the plot of residuals against predicted values. Both models were tested in terms of actual observation minus predictions for the data sets. The precision

achieved in those overall equations was better than any other models.

The residuals about the predicted values never exceed  $\pm 4.0m$ . The mean residual values for the nonoverlapping intervals model was 0.0335 about one third greater than that for all possible intervals model at -0.011. The ideal would be a mean residual value of 0. Skewness for the all possible data form had a value of 0.1492 compared with 0.2497 for the nonoverlapping form. The ideal would, of course, be a skewness value of 0. The t statistic of all possible intervals equation was also much greater than that of nonoverlapping intervals equation.

**CONCLUSION**

This research showed that use of the nonoverlapping data set form to build growth and yield models was efficient, appropriate and resulted in estimates of parameters that were generally precise. However, The more precise estimates of parameters were achieved when more efficient mix of projection intervals was used. This research provides some positive evidences that by using data sets which contains a range of  $Age_{(T2)} - Age_{(T1)}$  time intervals, as opposed to the more common method of using annual or bi-

**Table 8.** Non linear least squares summary statistics for mean top height model

SOURCE	Degree of Freedom		Sums of Squares		Mean Squares	
	Nonoverlapping	All possible	Nonoverlapping	All possible	Nonoverlapping	All possible
Regression	4	4	1,018,458	3,896,357	254,614	974,089
Residual	1,681	5,790	990	5,325	0.5894	0.9197
Uncorrected total	1,685	5,794	1,019,449	3,901,682		
Corrected total	1,684	5,793	132,944	412,275		

Parameter	Estimates		Std.Error / t statistics		95% Confidence Interval			
	Nonoverlapping	All possible	Nonoverlapping	All possible	Nonoverlapping		All possible	
					Lower	Upper	Lower	Upper
$\alpha$	5.2913	5.5352	0.1250 / 42.34	0.0415 / 133.39	5.05	5.54	5.45	5.62
$\beta$	0.4101	0.3867	0.0210 / 19.95	0.0058 / 67.02	0.37	0.45	0.38	0.40
$\beta_1$	0.2178	0.1956	0.0392 / 5.56	0.0161 / 12.13	0.14	0.29	0.15	0.23
$\beta_2$	0.1693	0.0408	0.0443 / 3.82	1.0187 / 2.17	0.08	0.26	0.01	0.08



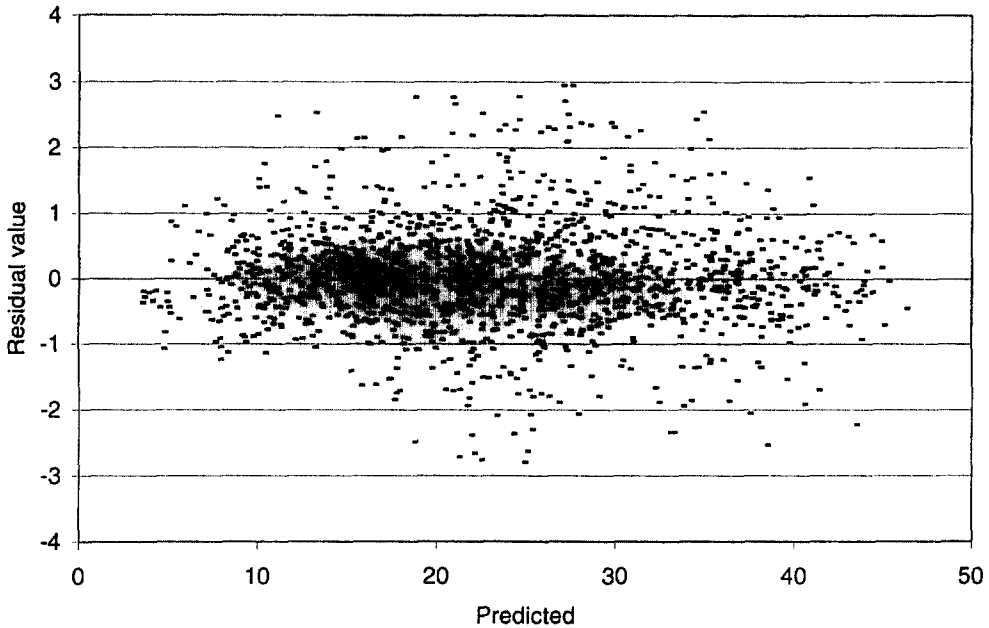


Fig. 3. Plot of residual Vs predicted for nonoverlapping intervals mean top height equation

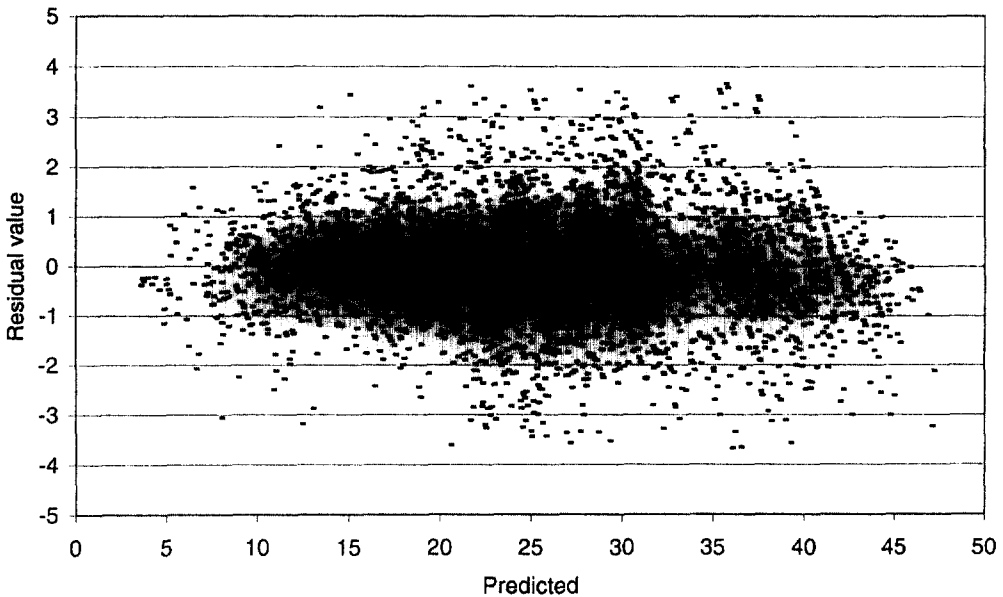


Fig. 4. Plot of residual Vs predicted for all possible intervals mean top height equation

annual intervals, the precision of parameter estimates could be increased.

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