

**VALIDATION OF THE INDIVIDUAL-TREE
GROWTH MODEL (ITGM)**

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EXECUTIVE SUMMARY

The Individual-Tree Growth Model (ITGM) is a fully-functional, prototype individual-tree growth model for radiata pine, which implements in combination, tree-level growth and survival equations to project stand- and tree-level growth and yield. ITGM prediction equations require data specific to each of 7 growth modelling regions, and operates on an annual cycle to grow trees from plantation age 15 to rotation age. The ability of this modelling approach has previously been documented using residual analyses with respect to isolated equations, but it has not been validated, as an integrated system of equations, using a fully-functional growth simulator.

During validation, the previously documented modelling approach used to predict static tree height resulted in severely biased prediction of tree height increment. Therefore, a major revision to tree height-growth prediction was undertaken and implemented in ITGM.

During regional tree-level growth and survival analyses, Timberlands West Coast (TWC) was inadvertently overlooked for inclusion. As an interim measure, ITGM was programmed to use Nelson region (GDNS) growth and survival equations. Now, TWC diameter and height growth equations have been fitted.

The objectives of this Report are to document:

- the validation of ITGM to establish the level of confidence at which the ITGM provides useful output and correct inferences about mid-rotation onward growth and yield,
- the new approach for the prediction of tree height increment, and
- the investigation into diameter and height growth equations for TWC.

Regression results confirmed the previous finding, that R^2 is poor (≤ 0.20) for directly predicting tree height increment. Seven regional dummy variables and 4 explanatory variables were able to explain only 17% (R^2) of the variation in tree height increment. Even though the R^2 value is low, the validation exercise established that this prediction approach is superior to the use of the tree height / mean top height ratio prediction approach. The utility of regional dummy variables suggests that separate, specific regional tree height increment equations should be investigated. This regional approach may identify regionally specific explanatory variables, and result in improved R^2 .

Relative to the other regional diameter and height growth equations, the fitted TWC equations have significantly lower R^2 values. Despite repeated attempts at uncovering better explanatory variables unique to TWC, explanatory variables similar to GDNS worked best for TWC, but with poorer results. Validation statistics revealed that the TWC predictions had marginally better results via the use of the GDNS growth equations, rather than the directly fit TWC growth equations. This result is counter-intuitive, but accepted. Until further investigations prove otherwise, TWC will continue to be represented with GDNS growth equations in ITGM, as validation results fully support this.

Overall, validation statistics reveal that, on-average, 5 and 10 year tree- and stand-level diameter and height prediction error is generally $\leq 3\%$ (usually over-predicted), while absolute prediction error is generally $\leq 7\%$. Accompanying predicted tree-diameter distributions generally appear reasonable, and skewed and multi-modal diameter distributions (i.e., other than "normal") are accommodated well.

These results indicate that ITGM, as an iterative prediction system for mid-rotation onward tree and stand growth and survival, operates within acceptable error limits without any obvious anomalies or mechanistic mathematical flaws.

INTRODUCTION

Gordon and Lawrence (1994), Gordon (1996), Lundgren and Gordon (1997), Shula (1997a, b, c) document the development of individual-tree diameter, height, and survival equations applicable to mid-rotation, post-silviculture radiata pine stands, i.e., ≥ 15 years old. Gordon and Shula (1999) document the fully-functional, prototype individual-tree growth model (ITGM), which implements, in combination, the tree-level growth and survival equations to project stand- and tree-level growth and yield.

The individual-tree growth modelling approach has equation forms that are analogous to those used in rotation-age individual-tree growth models (Wykoff et al. 1982; Hann et al. 1997). The ability of this modelling approach has been documented and verified using residual analyses (Shula 1997a, b, c) with respect to isolated equations, but it has not been validated using a fully-functional growth simulator, where the equations work together as a system. The diameter, height, and survival prediction equations require data specific to each of 7 growth modelling regions, and operate on an annual cycle to grow trees from plantation age 15 to rotation age.

During the present validation exercise, the approach and equation form used to predict static tree height (Shula 1997b) subsequently resulted in severely biased prediction of ht_i increment (obtained by subtraction of static tree height at two ages). Therefore, a major revision to tree height-growth prediction was undertaken, implemented in ITGM, and is also described in this Report.

During regional tree-level growth and survival analyses, Timberlands West Coast (TWC) was inadvertently overlooked for inclusion. As an interim measure to represent TWC, ITGM was programmed to use Nelson region (GDNS) growth and survival equations. Now, specific diameter and height growth analyses for TWC have been undertaken, and are also described in this Report.

OBJECTIVES

The objectives of this Report are to document:

- the validation of ITGM, and to establish the level of confidence at which the ITGM provides useful output and correct inferences about mid-rotation onward growth and yield,
- the new approach for the prediction of tree height increment, and
- the investigation into diameter and height growth equations for TWC.

NOTATION

age	= plantation age (years)
ba_i	= sum of basal area of individual trees in a stand larger than tree 'i'
dbh_i	= individual-tree, breast-height (1.4 m) diameter (mm)
ba_dbh	= ba_i ratio, i.e., (ba_i / dbh_i)
chg_pdbh	= change in potential dbh_i ; potential dbh_i growth, as if a MTH tree (Shula 1997a)
dbh_q	= stand, quadratic mean breast-height diameter (mm or cm)
$\exp(x)$	= e^x ; e is the base, 2.71828, of the natural logarithm
G	= stand, basal area (m^2 /hectare)
ht_i	= individual-tree height (m)
h_{pit}	= height potential index of an individual-tree; i.e., ht_i at age 20 (Shula 1997a)
ln	= natural base 2.71828 logarithm
MTD	= stand, mean top breast-height diameter (mm or cm)
MTH	= stand, mean top height (m)
N	= stems per hectare (sph)
S	= radiata pine site index, base age 20 years (m)

$$\begin{aligned} \text{sqdl_dbh} &= \{ [\ln(\text{dbh}_i)]^2 \} \text{ (cm)} \\ \text{sqd_ht} &= (\text{ht}_i)^2 / 1000 \text{ (m)} \\ \text{sph_rdbh} &= \text{stocking (N)} \times (\text{dbh}_i / \text{dbh}_q) \end{aligned}$$

DATA

Origin

The validation analyses are based on a sub-set of the same regional permanent sample plots (PSPs) used to develop the tree-level growth and survival equations. Nonetheless, the validation is considered quasi-independent because:

1. the datasets for the fitting of growth equations were based on low-intensity random selection (~8%, or some 5,300 of 65,500 observations) of tree-level observations from regional PSPs, while
2. the datasets for the validation analyses (actual versus predicted) are based on a complete selection (100%) of tree-level observations on selected regional PSPs.

Furthermore,

3. the fitting of growth equations was based on point-in-time, annulated growth increments, while
4. the validation analysis is based on iterative, cumulative annual growth and yield prediction starting from an initial point-in-time.

Item 1 resulted in a reduction of the likelihood for correlated regression residuals, while item 2 resulted in the provision of necessary PSP input data (complete tree- and stand-level attributes) required to start an ITGM simulation. Thus, the validation analysis is based on many different individual-trees, than was included in the original regression analysis. Furthermore, item 4 resulted in the investigation of accumulated, cascaded prediction errors, while item 3 resulted in the prediction of growth in annual time-steps.

PSPs to develop growth equations for TWC, and subsequent validation, originated from the same database used to develop a TWC stand-level growth model (Kimberley and Hawke, 1999).

Description

PSPs and simulation start- and end-ages used in the validation analyses are listed in **Appendix A**. Summaries of plot data (at simulation start-age), including stand- and tree-level mean, and minimum / maximum values are presented in **Tables 1 and 2**.

Table 1. Summary of plot data, at simulation start-age, providing stand-level mean and (minimum / maximum values).

Region	Plots [No.]	Simulation Start-Age [yrs]	dbh _q [cm]	MTD [cm]	G [m ² /ha]	N [sph]	MTH [m]	S [m]
CNI	21	16 (15/22)	28 (17/55)	36 (28/60)	43 (17/62)	1086 (158/2702)	27 (21/37)	34 (27/37)
GDNS	12	16 (15/21)	26 (21/32)	33 (27/40)	32 (11/56)	681 (180/1630)	21 (17/24)	26 (22/30)
HBAY	12	17 (15/19)	35 (32/41)	40 (36/46)	29 (15/58)	315 (178/721)	25 (20/30)	29 (25/33)
CLAYS	13	18 (15/25)	33 (25/40)	39 (31/46)	37 (16/78)	450 (290/1170)	27 (22/35)	30 (23/33)
SOUTH	10	16 (15/17)	27 (17/35)	32 (25/39)	30 (24/68)	683 (285/2898)	19 (16/20)	24 (21/27)
SANDS	16	19 (15/25)	30 (16/44)	34 (19/49)	22 (9/33)	322 (109/544)	23 (16/31)	24 (20/30)
CANTY	5	17 (15/20)	28 (23/34)	35 (30/40)	38 (31/46)	665 (352/950)	21 (17/23)	25 (22/27)
TWC	6	16 (15/19)	30 (18/41)	41 (39/44)	27 (9/48)	371 (239/553)	22 (16/29)	27 (22/30)

Table 2. Summary of plot data, at simulation start-age, providing tree-level mean and (minimum / maximum values).

Region	Simulation Start-Age [yrs]	No. Observations		dbh _i	ht _i
		dbh _i	ht _i	[cm]	[m]
CNI	16 (15/22)	1662	192	23 (7/69)	26 (15/41)
GDNS	16 (15/21)	425	105	24 (5/44)	20 (11/27)
HBAY	17 (15/19)	383	88	34 (7/51)	24 (17/31)
CLAYS	18 (15/25)	399	101	31 (12/50)	26 (13/40)
SOUTH	16 (15/17)	529	122	23 (5/41)	18 (9/22)
SANDS	19 (15/25)	612	150	29 (9/54)	22 (15/33)
CANTY	17 (15/20)	163	38	28 (11/43)	21 (16/26)
TWC	16 (15/19)	333	109	31 (7/56)	21 (8/30)

METHODS

Background

Validation is an integral step in the process of building a simulation model (Goulding 1979). Validation addresses the ability of a growth simulator, as a whole, to enable users to make correct inferences regarding operational and/or experimental hypotheses of interest based on simulator output. Sharpe (1990) recognised this approach to validation as performance and refutation, in addition to, goodness-of-fit and mathematical rigour. Performance and refutation, by necessity, follow goodness-of-fit and mathematical rigour to fully evaluate the development of isolated equations and the holistic set of equations that comprise a growth simulator. First, equation form and regression parameter estimates are evaluated for goodness-of-fit (e.g., mean square error, parameter significance, constant standard error, and coefficient of determination). Next, the simulator's collective set of equations is evaluated for mathematical rigour, which involves assessment of the simulator's architecture for suitability, logic, and arithmetic functionality. Finally, performance and refutation evaluate a simulator's accuracy and precision of prediction with respect to the dynamic interaction played-out by the holistic set of equations and explanatory variables.

Validation approach

The present validation analyses focus on the evaluation of mathematical rigour and performance. To this end, ITGM was used to simulate growth and yield of selected PSPs to check the simulator's architecture for suitability, logic, and arithmetic functionality; and, to compare simulator output with observed PSP data.

ITGM was programmed in batch-mode to accept comma-delimited ASCII input data, and to make iterative annual growth and yield predictions culminating at future PSP stand ages. ITGM simulations were started with initial PSP starting values (tree- and stand-level variables), but thereafter, iterative annual steps to final projection age were re-initiated with predicted tree- and stand-level values. Predicted stand-level values were obtained by the aggregation of predicted tree-level values based upon predicted probabilities of tree-survival and resultant predicted tree-expansion factors. Predicted values at final PSP stand ages that did not result from exact annual projections (e.g., starting-age 22.05 to ending-age 26.65) were obtained by back-adjustment to exact PSP stand ages via interpolation (e.g., annual prediction from age 22.05 to age 27.05, then interpolation back to age 26.65).

The selection of PSPs (and resultant number of observations) for the validation analyses was based on criteria to provide identically paired, 1x- and 2x-term periods of simulation (1x-term, ~ 5 years; 2x-term, ~ twice the 1x-term, or around 10 years), starting at plantation age ≥ 15 years and without thinning interventions. These strict criteria were chosen to provide exact one-to-one comparative estimates of prediction accuracy and precision for 1x- and 2x-simulation periods considered of most interest to mid-rotation onwards inventory projection. These criteria purposely focused the validation analyses, but also, restricted the scale of the analyses because PSP re-measurements meeting the strict criteria were limited in supply.

Validation attributes and statistics

Validation statistics were calculated for the following stand-level attributes: dbh_q , G , N , MTD , and MTH ; and tree-level attributes: dbh_i and ht_i . There is no single validation statistic that entirely captures model prediction accuracy and precision; therefore, all of the following set of validation statistics were used:

1. R-squared (or modelling efficiency, Soares et al. 1995):

$$R^2 = 1 - [\sum (y_{act_i} - y_{hat_i})^2 / \sum (y_{act_i} - y_{mean_i})^2]$$

2. Mean error (ME):

$$ME = [\sum (y_{act_i} - y_{hat_i})] / n$$

3. Mean percent error (MPE):

$$MPE = \{ [\sum (y_{act_i} - y_{hat_i} / y_{act_i})] / n \} * 100$$

4. Root mean squared error (RMSE):

$$RMSE = [\sum (y_{act_i} - y_{hat_i})^2 / n]^{0.5}$$

5. Mean absolute error (MAE):

$$MAE = [|y_{act_i} - y_{hat_i}| / y_{act_i}] / n$$

6. Mean absolute percent error (MAPE):

$$MAPE = [\sum |y_{act_i} - y_{hat_i}| / y_{act_i} / n] * 100$$

7. Mean actual or predicted (MA, MP):

$$MA = (\sum y_{act_i}) / n$$

$$MP = (\sum y_{hat_i}) / n$$

8. Standard deviation of actual or predicted (STDA, STDP):

$$STDA = \{ [\sum (y_{act_i} - y_{act_mean_i})^2] / n \}^{0.5}$$

$$STDP = \{ [\sum (y_{hat_i} - y_{hat_mean_i})^2] / n \}^{0.5}$$

where, y_{act_i} = actual value of the variable
 $y_{act_mean_i}$ = actual mean value of the variable
 y_{hat_i} = predicted value of the variable
 $y_{hat_mean_i}$ = actual mean value of the predicted variable
 n = number of observations in the validation dataset.

Statistics 1 - 3 represent 'soft' validation assessors, as positive (+) and negative (-) errors accumulate, but have the opportunity to 'average out'. Conversely, statistics 4 - 6 represent 'hard' validation assessors, as errors (absolute or squared) are accumulated without the opportunity to 'average out'. Statistic 4 is a particularly 'hard' assessor, as the squaring of errors, penalises large errors, especially.

The R^2 statistic quantifies the departure of predicted values from actual values relative to the departure of actual values from the mean actual value. This statistic, then, is sensitive to the range in prediction error (variation about zero error) and to the range in the actual data (variation about the mean actual value). A value of 1 identifies a perfect prediction model, while a value ≤ 0 , identifies a prediction model no-better-to-worse than using the mean.

The STDA and STDP statistics quantify the variation about the mean actual and mean predicted values, respectively. The comparison of these validation statistics are particularly meaningful to evaluate tree-level models, as the goal of tree-level models is the accurate projection of the tree-list and tree-size variability in the tree-list.

Height growth prediction

The first height growth investigation (Gordon 1996) concluded that the prediction of individual-tree height increment was not worthwhile, based principally on poor goodness-of-fit statistics. Further investigation (Shula 1997b) accepted this conclusion, and continued to rely on the indirect prediction of ht_i via the direct, static prediction of height ratio ($ht_i:MTH$), as a power function on diameter ratio ($dbh_q:MTD$). Therefore, predicted ht_i increment was a) dependent on predicted MTH at the end of the prediction period, and b) obtained indirectly by subtraction of predicted ht_i at the start and end of the prediction period. Regional specificity was incorporated through the inclusion of regional dummy variables; adjusted R^2 was 0.51.

During the present validation exercise, this height growth modelling approach resulted in severely biased predictions of ht_i increment. This bias was attributed to the inherent reliance on predicted MTH derived from site index, and the instability of the power function on diameter ratio for individual-trees with diameters approaching or greater than stand MTD. To seek a remedy, the first approach to directly model ht_i increment was revisited.

The revised height growth analysis used the same dataset (291 plots) as described in Gordon (1996) and Shula (1997b); however, all observations were utilised ($n=13440$), rather than separating the dataset into regression and validation datasets ($n=5264$ and $n=8176$, respectively). Gordon 1996 (Table 2, page 3) provides the number of observations, and the range in tree height and height increment by Region for the entire dataset.

SAS (SAS Institute Inc. 1989) non-weighted, non-linear regression procedure, NLIN, (method = marquardt) was used to estimate parameter coefficients ($\alpha \leq 0.05$). The equation form that was fit was a generalised Schumacher (1939) log reciprocal:

$$\bullet \quad y = \exp(r_1 + \dots + r_7 + a_1x_1 + \dots + a_nx_n) \quad [1]$$

The dependent variable (y) was annual ht_i increment (m). Where re-measurement did not occur exactly 12 months later, interpolation was used to obtain an annulated observation. Regional specificity was incorporated through the inclusion of regional dummy variables ($r_1 - r_7$). Potential explanatory variables (x_n) common to all regions included all those tried in the first investigation (Gordon 1996), and those newly devised, although used previously in growth and survival analyses (Shula 1997a, b, c). Criteria for judging equation goodness-of-fit, homogeneity of residual variance, and acceptance included adjusted R^2 (Kmenta 1986) and Furnival's Index (Furnival 1961).

TWC diameter and height growth equations

The TWC tree-level diameter and height growth analyses started with the same underlying database (812 PSPs) as available for the contractual development of a TWC stand-level growth model (Kimberley and Hawke 1999). Ultimately, the stand-level modelling exercise used a total of 1802 stand-level measurements from 336 PSPs, resulting from the exclusion of PSPs with: site index $\leq 15m$, very high stocking, and suspect data.

The present tree-level diameter and height growth analyses selected tree-level data from a total of 13187 measurements from 235 PSPs, resulting from the initial exclusion of PSPs with: site index $\leq 15m$, non-annual re-measurements, and suspect data. In the

diameter growth analyses, and in keeping with previous analyses (Gordon and Lawrence 1994 and Shula 1997a), randomly selected data resulted in 778 tree-level observations from 168 PSPs (**Appendix B**). Summaries of plot data (at initial age), including stand- and tree-level mean, and minimum / maximum values are presented in **Tables 3 and 4**. In the height growth analyses, and in keeping with previous analyses (Shula 1997b), keeping all acceptable height data resulted in 2715 tree-level observations from 188 PSPs (**Appendix C**). Summaries of plot data (at initial age), including stand- and tree-level mean, and minimum / maximum values are presented in **Tables 5 and 6**.

SAS (SAS Institute Inc. 1989) weighted, non-linear regression procedure, NLIN, (method = marquardt) was used to estimate parameter coefficients ($\alpha \leq 0.05$). An equation similar to [1] was fit, with the dependent variable (y) as annual dbh_i and ht_i increment (cm and m, respectively). As TWC data was being fit directly, an intercept parameter (a₀) replaced the regional dummy variable (r₁). Potential explanatory variables (x_n) included all those tried in previous investigations (Shula 1997a, b). To better ensure homogenous variance of residuals, a variety of weighting schemes were investigated, including: the reciprocal (or not) of tree-size attributes (e.g., dbh, height), and iterative re-weighting using the reciprocal (or not) of the predicted. Criteria for judging equation goodness-of-fit, homogeneity of residual variance, and acceptance included adjusted R² (Kmenta 1986) and Furnival's Index (Furnival 1961).

Table 3. TWC diameter growth analyses - summary of plot data, at initial age, providing stand-level mean and (minimum / maximum values).

Region	Plots [No.]	Initial Age [yrs]	dbh _q [cm]	MTD [cm]	G [m ² /ha]	N [sph]	MTH [m]	S [m]
TWC	168	20 (15/32)	36 (18/57)	41 (23/63)	27 (6/79)	262 (131/600)	24 (11/39)	25 (14/33)

Table 4. TWC diameter growth analyses - summary of plot data, at initial age, providing tree-level mean and (minimum / maximum values).

Region	Initial Age [yrs]	No. Observations		dbh _i [cm]	ht _i [m]
		dbh _i	ht _i		
TWC	20 (15/32)	778	736	38 (7/84)	23 (5/44)

Table 5. TWC height growth analyses - summary of plot data, at initial age, providing stand-level mean and (minimum / maximum values).

Region	Plots [No.]	Initial Age [yrs]	dbh _q [cm]	MTD [cm]	G [m ² /ha]	N [sph]	MTH [m]	S [m]
TWC	188	20 (15/32)	35 (18/57)	40 (23/63)	26 (6/79)	260 (120/600)	23 (11/39)	24 (14/33)

Table 6. TWC height growth analyses - summary of plot data, at initial age, providing tree-level mean and (minimum / maximum values).

Region	Initial Age [yrs]	No. Observations		dbh _i	ht _i
		dbh _i	ht _i	[cm]	[m]
TWC	20 (15/32)	2715	2715	38 (7/78)	24 (6/43)

RESULTS

Height growth prediction

Regression results confirmed the previous finding (Gordon 1996), that R^2 is poor (≤ 0.20) for predicting ht_i increment. Seven regional dummy variables and 4 explanatory variables (all significantly different than 0 or 1) were able to explain only 17% (R^2) of the variation in ht_i increment. This indicates that the range in squared prediction error is usually greater than the range in squared observed departure from the mean. **Figures 1a and b** present actual ht_i increment by actual dbh_i for the 7 regions, showing that ht_i increment is largely homogenous across tree-size, and quite variable for a given tree-size. This supports the inability to achieve higher R^2 for ht_i increment prediction, i.e., mean ht_i increment is difficult to improve upon via least-squares regression. **Tables 7 and 8** provide fit statistics and parameter coefficients, respectively, from the regression analyses.

Table 7. Mean residual height increment and standard error by Region from the regression analyses ($R^2=0.17$).

Region (no. obs.)	Mean Ht _i Increment Residual (m)	Standard Error (m)
CNI (4449)	0.00883	0.00979
HBAY (1365)	0.00481	0.01607
GDNS (2538)	-0.00492	0.01162
SOUTH (1997)	-0.00827	0.01153
CLAYS (831)	-0.01215	0.01950
CANTY (301)	0.00348	0.03381
SANDS (1959)	-0.01511	0.01134
Combined REGIONS (13440)	-0.00162	0.00507

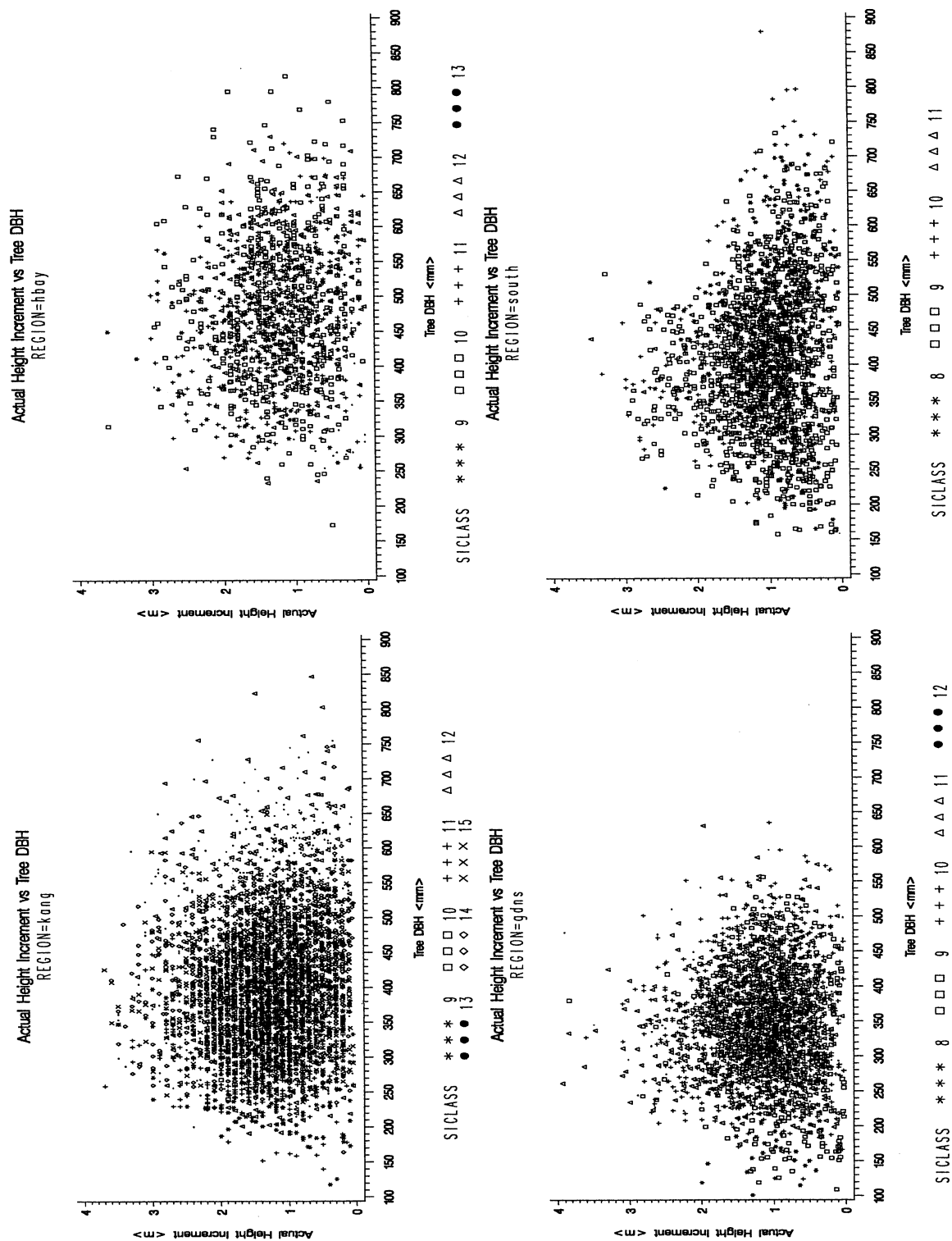


Figure 1a. Actual tree height increment by actual tree dbh for the 7 regions.

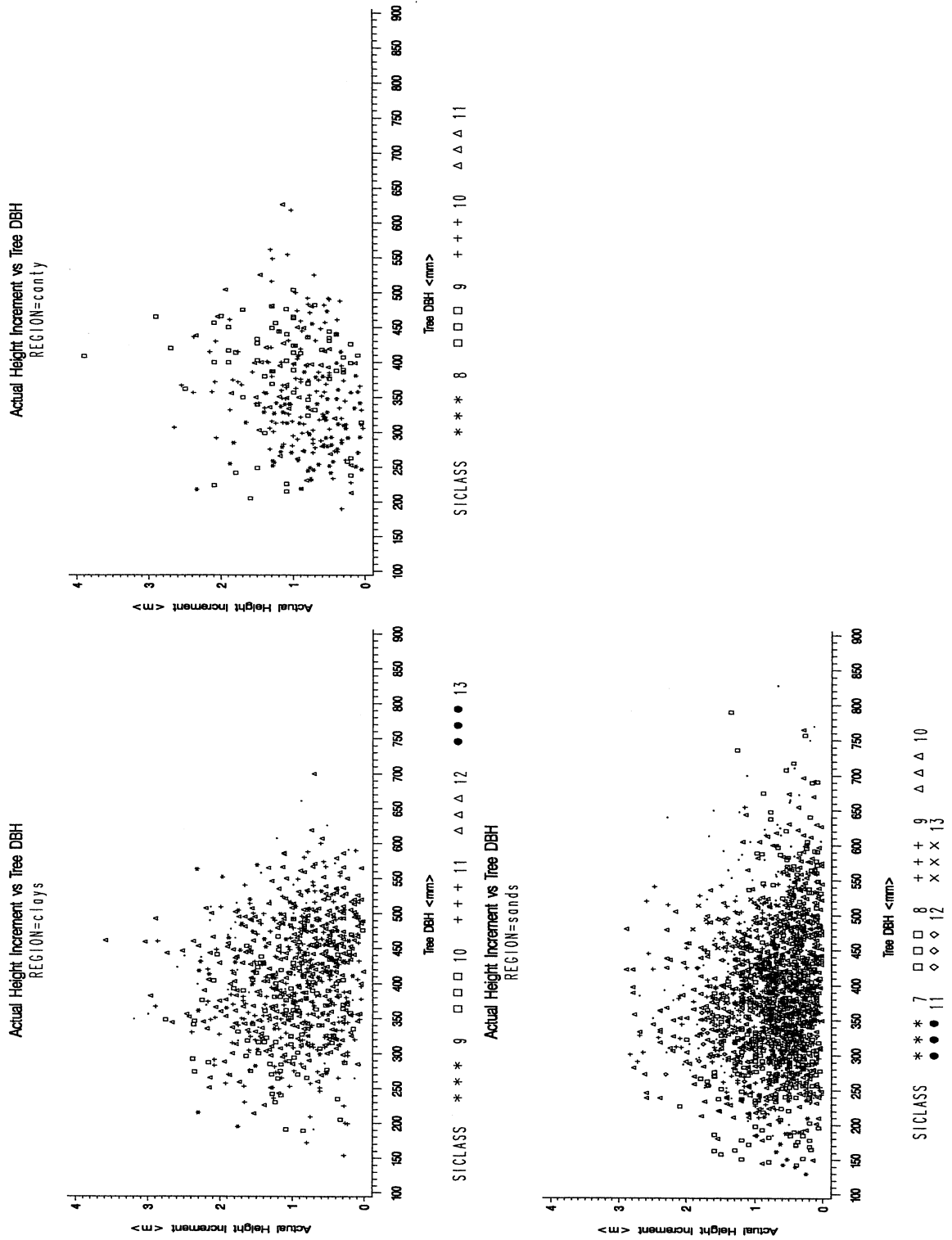


Figure 1b. Actual tree height increment by actual tree dbh for the 7 regions.

Table 8. Parameters and coefficients by Region from the height increment regression analyses ($R^2=0.17$).

$$\text{annual } ht_i \text{ increment} = \exp(r_1 + \dots + r_7 + a_1X_1 + \dots + a_nX_n)$$

Region (no. obs.)	Parameter	Coefficient ($\alpha \leq 0.05$)	Standard Error
CNI (4449)	r1	-0.97793	0.12094
HBAY (1365)	r2	-1.07052	0.12490
GDNS (2538)	r3	-1.11595	0.11640
SOUTH (1997)	r4	-1.14554	0.11879
CLAYS (831)	r5	-1.19604	0.12201
CANTY (301)	r6	-1.29988	0.12276
SANDS (1959)	r7	-1.46447	0.11765
COMMON TO ALL REGIONS (13440)	sqdl_dbh	0.02995	0.00297
	sqd_ht	-0.73821	0.02480
	hpit	0.02942	0.00166
	bal_dbh	-0.36743	0.15484

The ordering of regions by descending magnitude of the coefficient for the regional dummy variable:

	<u>dummy coeff.</u>	<u>correspondent ht_i increment (m)</u>
CNI	-0.98	0.38
HBAY	-1.07	0.34
GDNS	-1.12	0.33
SOUTH	-1.15	0.32
CLAYS	-1.20	0.30
CANTY	-1.30	0.27
SANDS	-1.46	0.23

provides a ranking of largest to smallest regional ht_i base-increment (by solving the prediction equation with simply the regional coefficient). In practice, these regional ht_i base-increments are, then, adjusted up or down by the other explanatory variables in the equation (i.e., diameter, height, height potential index, relative tree-size) with coefficients common to all regions.

Figures 2a and b present ht_i increment residuals by predicted ht_i increment; predictions are unbiased, although residuals range widely about zero error. On average across regions, 62% of the residuals (range, 50% to 70%) are within 0.5m of the actual ht_i increment; while 82% of the residuals (range, 75% to 91%) are within 0.75m of the actual ht_i increment.

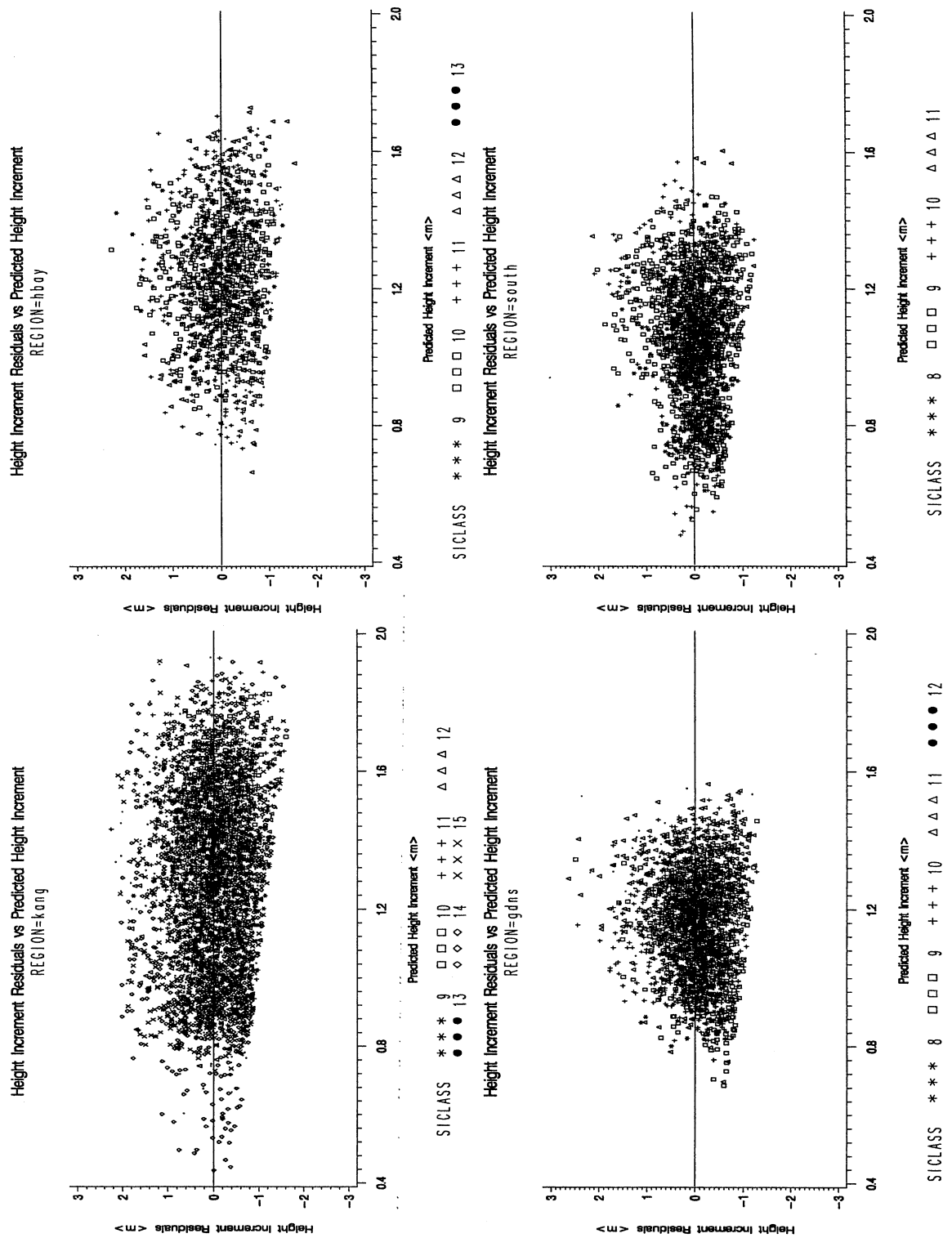


Figure 2a. Tree height increment residuals by predicted tree height increment for the 7 regions.

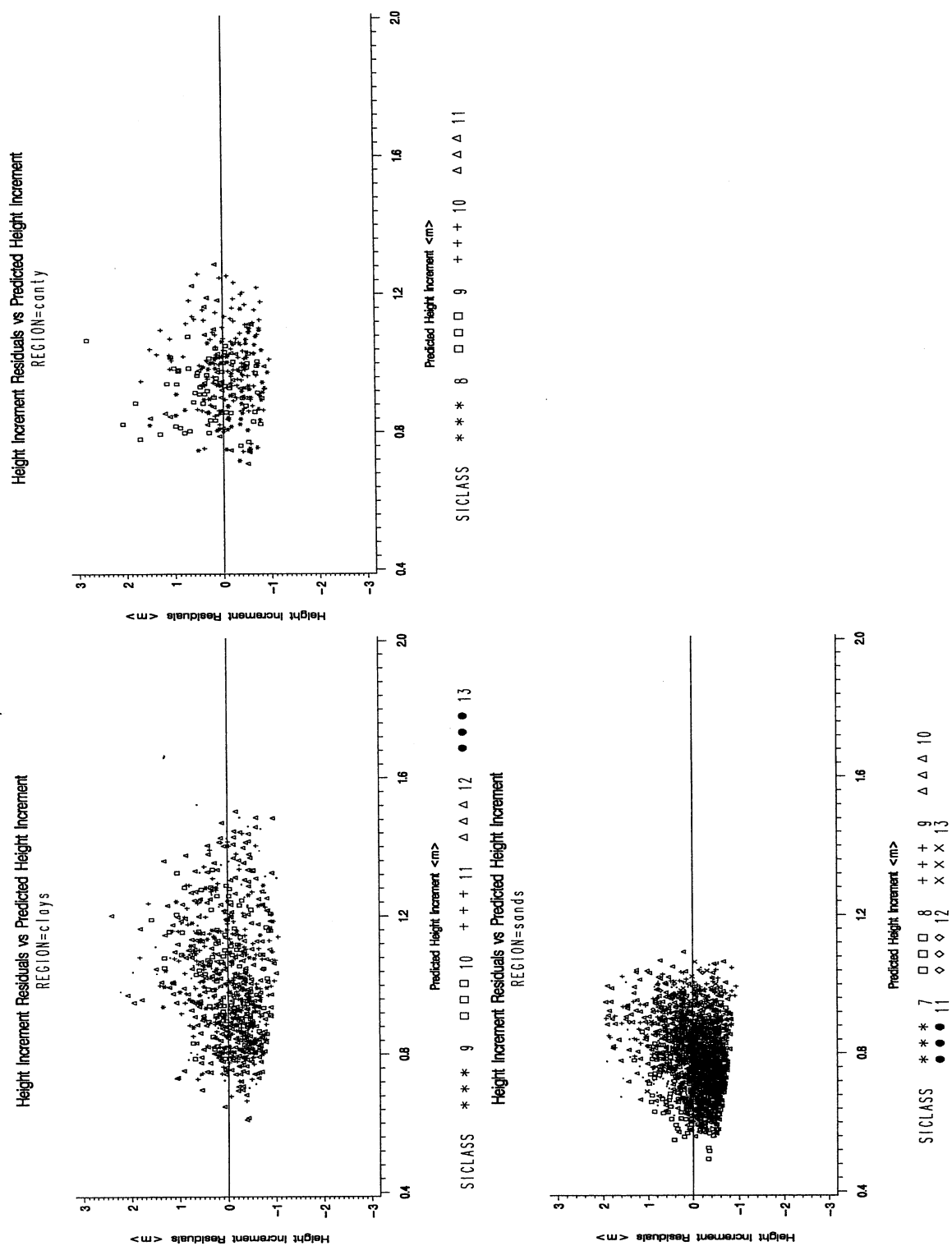


Figure 2b. Tree height increment residuals by predicted tree height increment for the 7 regions.

TWC diameter and height growth equations

Relative to the other regional diameter and height growth equations (Shula 1997a and 1997b), the TWC equations have significantly lower R^2 values. For example, the GDNS diameter and height growth equation R^2 s were 0.56 and 0.17, respectively; but for TWC (Table 9), R^2 s are 0.36, and 0.04, respectively. Despite repeated attempts at uncovering better explanatory variables unique to TWC, explanatory variables similar to GDNS worked best for TWC, but with poorer results. Therefore, it was left to the validation analyses to determine the relative utility to TWC of using either the directly fit TWC equations (Table 9), associated with poorer fit statistics; or in lieu, the GDNS equations (Shula 1997a and 1997b), associated with better fit statistics.

Table 9. TWC diameter and height growth equation parameters, coefficients, and standard errors from the non-linear regression analyses.

$$\text{annual dbh}_i \text{ or ht}_i \text{ increment} = \exp[a_0 + a_1 * x_1 + \dots + a_n x_n].$$

Attribute (no. obs.)	Parameter	Coefficient ($\alpha=0.05$)	Standard Error
dbh _i (778)	intercept	2.14217	0.09031
	chg_pdbh	0.05279	0.00416
	bal_dbh	-4.21678	0.42323
	sph_rdbh	-0.00041	0.00013
ht _i (2715)	intercept	-1.07449	0.11331
	bal_dbh ²	-11.07650	2.60184
	MTD	0.02289	0.00257
	hpit	0.03915	0.00425
	sqd_ht	-1.05357	0.09468

Figures 3 and 4 present dbh_i and ht_i actual increments and residuals by tree size and predicted increments, respectively; predictions are unbiased, although residuals range widely about zero error. For dbh_i and ht_i, 62% of the residuals are within 0.5cm of the actual dbh_i increment; while 50% of the residuals are within 0.5m of the actual ht_i increment.

Validation statistics

Validation statistics revealed that the TWC predictions had marginally better results via the use of the GDNS growth equations, rather than the directly fit TWC growth equations. In this approach, TWC data represented entirely independent data to validate the GDNS equations and to identify the best equations for TWC. While this result is counter-intuitive, the result is accepted, and in the validation statistics that follow, TWC results are based on the ITGM model (as distributed, February 1999) using the GDNS equations, in lieu of, the directly fit TWC equations.

Validation statistics at the stand-level (G, dbh_q, N, MTH, MTD) are provided in Tables 10-14, and at the tree-level (dbh_i and ht_i) in Tables 15-16. The tables present validation statistics by Region for identically paired plots at short (about 5 year) and long (about 10 year) projection periods. Because the Regions, CANTY and TWC, only had low numbers of plots (i.e., 5 and 6 plots) available for this identical pairing, Tables 17-18 (respectively), provide additional validation statistics for these Regions based on the inclusion of all plots (i.e., 11 and 20 plots) suited to a long projection period (i.e., regardless of identical plot pairing with short and long projection periods).

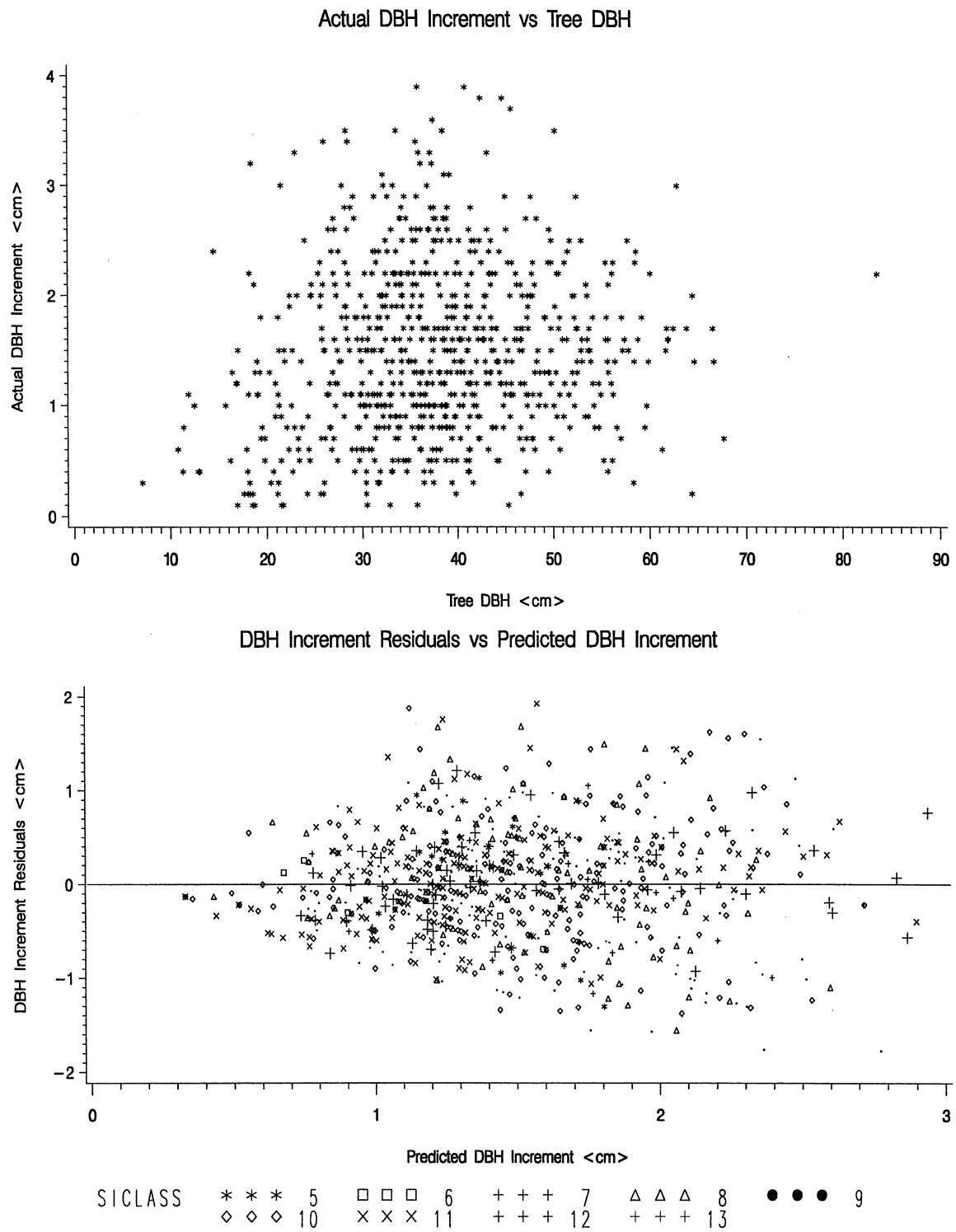


Figure 3. TWC actual tree dbh increments and residuals by tree size and predicted increments.

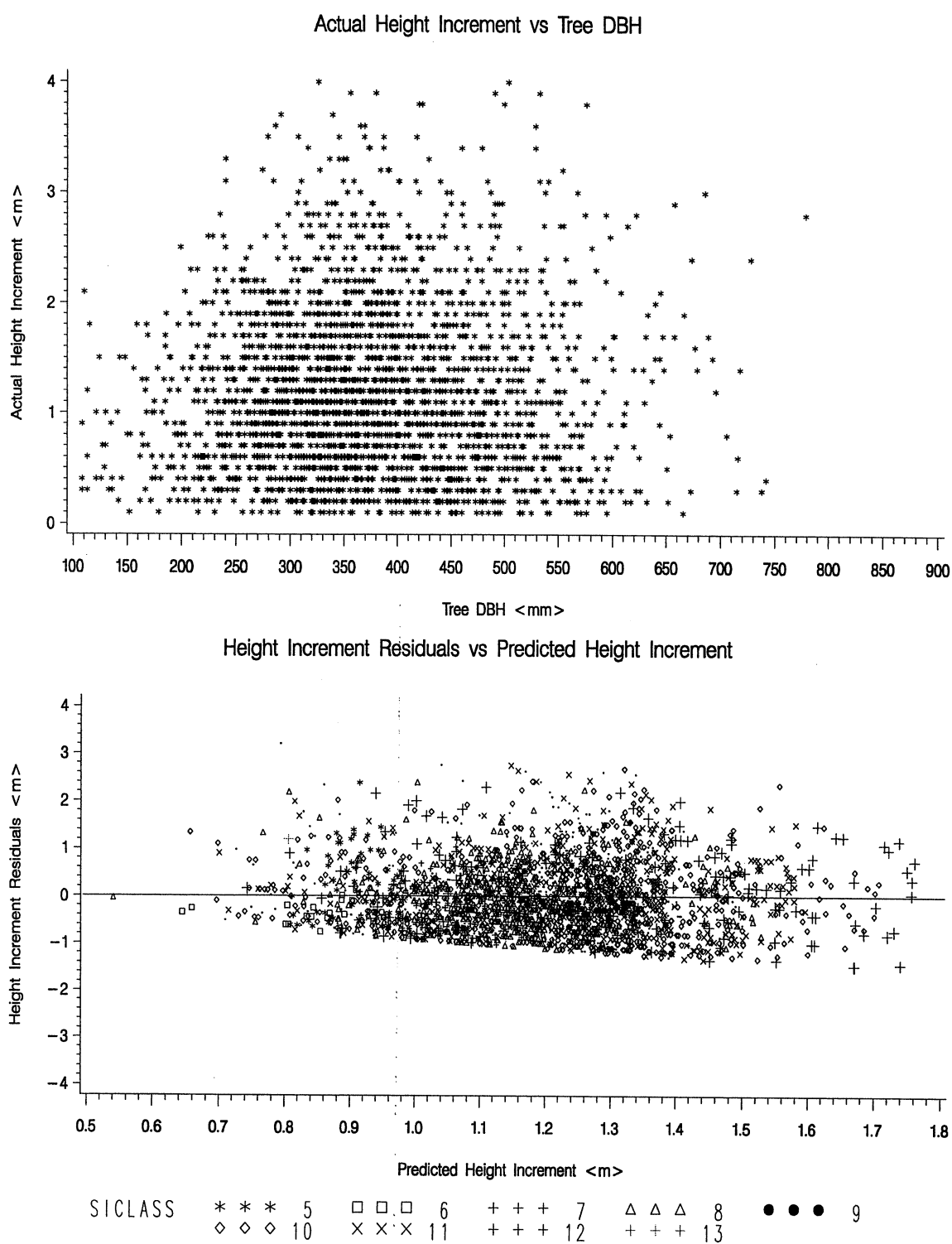


Figure 4. TWC actual tree height increments and residuals by tree size and predicted increments.

Diameter. Generally, tree dbh_i and stand dbh_q (generally, the sub-dominant) are over-predicted, while stand MTD (the dominant) is under-predicted. R-square is usually in the high 80's to mid-90's, and better for the sub-dominant, than for the dominant. Relative to the other Regions, CLAYS and SOUTH, have lower R-square for stand diameter at a long projection period. Mean absolute percent error is usually 3-4% (weighted by the number of plots) for stand diameter, but higher for tree diameter, 5-8% (weighted by the number of trees). In going from the short to long projection period, absolute percent error increases 1-1.5 percent units for stand diameter, but increases 3 percent units for tree diameter. The variability in predicted dbh_i is generally under-represented, but over-represented for MTD.

Height. Generally, tree and stand height are over-predicted. Relative to diameter, R-square is consistently some 20 percent units lower; usually in the low 60's to low-80's, but, better for the dominant (MTH). Relative to the other Regions, CLAYS has a lower R-square for tree ht_i at the long projection period, and a terrible R-square for stand MTH. Mean absolute percent error is usually 3-4% (weighted by the number of plots) for stand MTH, but higher for tree ht_i, 5-6% (weighted by the number of trees). In going from the short to long projection period, absolute percent error increases ≤ 1 percent unit for either tree or stand height. The variability in predicted height is increasingly under-represented, in going from the short to long projection period.

Basal Area. Generally, basal area is under-predicted, except for the Regions, CLAYS and CANTY. R-square is usually in the high 80's-90's, but for SOUTH and CANTY at the short projection period, R-square is some 20 percent units lower. At the long projection period, R-square is negative for SOUTH and CANTY. With the inclusion of additional plots, the R-square for CANTY (Table 17) becomes positive (40's), but is still poor. Mean absolute percent error is usually 4-6% (weighted by the number of plots); and in going from the short to long projection period, absolute percent error, generally, doubles. The variability in predicted basal area is increasingly under-represented, in going from the short to long projection period.

Stocking. Generally, stocking is under-predicted, except for the Region, CANTY. (Note: CANTY is the only Region without a survival function, i.e., assumes 100% survival, because mortality was too scarce to warrant modelling.) R-square is usually in the low to high 90's, but for SOUTH, at the long projection period, R-square is in the low 80's. Mean absolute percent error is usually 3-7% (weighted by the number of plots); and in going from the short to long projection period, absolute percent error, generally, doubles. The variability in predicted stocking is decreasingly under-represented, in going from the short to long projection period.

Tree-size Distribution. **Figures 5-12** present actual and predicted diameter distributions of selected plots from the 8 Regions for the short and long projection periods. "Actual" distributions are based only on surviving trees (i.e., actual mortality is excluded), while "predicted" distributions are based on each tree's predicted probability of survival (i.e., predicted mortality is excluded). **Appendix D** provides projection statistics for the selected Regional plots. To represent each Region, plot selection was based on the long projection period using:

- a) the plot nearest to a dbh_i variance ratio of 1.0 (i.e., variance of predicted / variance of actual; where, $\text{variance} = \Sigma(x_i - \bar{x})^2 / n$), and
- b) the plot nearest to a Region's mean dbh_i variance ratio.

These selection criteria were used, as surrogates, to identify:

- a) the Regional plot most likely to represent the "best ability" to predict a tree-size distribution, and
- b) the Regional plot most likely to represent the "ability on-average" to predict a tree-size distribution.

Note, in Figures 5-12:

- the top set of graphs is the plot nearest to a dbh_i variance ratio of 1.0,
- the left graph is the short projection period,
- the right graph is the long projection period,
- the bottom set of graphs is the plot nearest to the Region's mean dbh_i variance ratio (note: CANTY, one plot represents both, nearest to 1.0 and the mean ratio),
- within any graph, the left and right dbh_i distributions are the actual (a) and predicted (p), respectively,
- the y-axis is stocking (sph) within a dbh_i class, and
- the x-axis is 5 cm dbh_i classes (e.g., 15 cm class is 10 cm < dbh_i ≤ 15 cm).

Table 19 provides mean, standard deviation, minimum, and maximum dbh_i variance ratios by Region, based on the long projection period ($9 \leq \text{period} \leq 13$ years). Predicted dbh_i variance is consistently under-represented relative to actual dbh_i variance (except for the Region, CANTY), as indicated by a weighted (by number of plots) mean variance ratio of 0.78 (exclusive of CANTY's ratio, 1.03) and the minimum/maximum variance ratios being skewed slightly towards minimum.

Table 19. Mean, standard deviation, minimum, and maximum dbh_i variance ratios by Region, based on the long projection period.

Region	No. Plots	Long Projection Period (yrs)	Dbh _i Variance Ratio (predicted : actual)			
			Mean	Std. Dev.	Minimum	Maximum
CNI	21	13	0.77	0.18	0.48	1.14
GDNS	12	9	0.79	0.12	0.66	1.09
HBAY	12	11	0.88	0.17	0.63	1.25
CLAYS	13	10	0.77	0.18	0.53	1.08
SOUTH	10	10	0.62	0.25	0.28	1.02
SANDS	16	11	0.80	0.25	0.40	1.32
CANTY	5	9	1.03	0.23	0.82	1.40
TWC	6	10	0.81	0.12	0.62	0.94

Generally, **Figures 5-12** demonstrate that at low and high stockings, short and long projection periods; the spread in diameter distributions is predicted well. Also, the shape of the distributions are predicted reasonably well, especially with respect to distributions, which start as non-normal, but which become more normally distributed through time (e.g., CNI: Figure 5, top - left to right). The inability to mimic the shape of an actual diameter distribution may be partially attributed to a poor prediction of survival. **Appendix D** reveals that survival is consistently under-predicted (i.e., too much mortality), especially in SOUTH (Figure 9, top-right), where a correspondent inability to mimic the shape of the actual diameter distribution is exhibited.

Table 10. Validation statistics at the stand-level for basal area (m²/ha) by Region.

STATISTIC	CNI	GDNS	HBAY	CLAYS	SOUTH	SANDS	CANTY	TWC
number of plots	21	12	12	13	10	16	5	6
1x projection pd. (yrs)	6	5	6	5	5	5	3	5
2x projection pd. (yrs)	13	9	11	10	10	11	9	10
R-square 1x	.96	.99	.98	.99	.79	.96	.72	.97
R-square 2x	.80	.97	.94	.93	-.38	.87	-.53	.92
Mean error 1x	1.94	.59	.31	-.36	1.46	.34	-1.82	1.48
Mean error 2x	3.98	.93	1.84	-.97	2.93	1.79	-4.27	2.07
Mean % error 1x	3.20	.54	.66	-.87	2.91	.40	-3.57	4.62
Mean % error 2x	5.15	1.20	2.57	-1.81	4.11	3.67	-6.73	3.92
RMSE 1x	2.81	1.55	2.06	1.28	3.94	1.86	2.67	2.63
RMSE 2x	6.05	3.44	4.04	2.89	7.19	3.52	6.04	4.24
Mean abs. error 1x	2.01	1.13	1.77	1.08	3.22	1.63	2.34	2.26
Mean abs. error 2x	4.43	2.50	3.12	2.43	6.30	2.98	4.89	3.64
Mean abs. % error 1x	3.40	2.70	4.49	2.27	6.92	5.61	4.85	6.17
Mean abs. % error 2x	6.15	4.48	6.39	4.22	9.66	7.26	7.88	8.39
Mean actual 1x	55.79	44.42	44.29	49.16	49.22	32.53	46.34	40.08
Mean predicted 1x	53.85	43.84	43.98	49.52	47.76	32.19	48.16	38.60
Mean actual STD 1x	15.04	20.52	16.05	12.52	9.04	8.99	5.64	15.53
Mean predicted STD 1x	14.14	19.71	16.35	12.19	9.66	8.78	7.65	16.50
Mean actual 2x	65.80	53.44	55.41	59.79	65.84	42.98	59.66	49.81
Mean predicted 2x	61.82	52.50	53.57	60.77	62.91	41.20	63.93	47.73
Mean actual STD 2x	13.89	20.90	17.64	11.29	6.46	10.13	5.45	16.23
Mean predicted STD 2x	10.83	19.77	16.89	10.57	7.36	9.50	9.60	16.73

Table 11. Validation statistics at the stand-level for dbh_q (cm) by Region.

STATISTIC	CNI	GDNS	HBAY	CLAYS	SOUTH	SANDS	CANTY	TWC
number of plots	21	12	12	13	10	16	5	6
1x projection pd. (yrs)	6	5	6	5	5	5	3	5
2x projection pd. (yrs)	13	9	11	10	10	11	9	10
R-square 1x	.99	.97	.93	.96	.94	.95	.98	.96
R-square 2x	.98	.94	.87	.70	.89	.90	.96	.88
Mean error 1x	-.25	-.66	.13	-.48	.56	-.79	-.14	.37
Mean error 2x	-1.00	-1.24	.43	-1.35	.56	-1.25	-.32	-.15
Mean % error 1x	-.76	-2.08	.01	-1.18	1.11	-2.13	-.55	1.13
Mean % error 2x	-2.48	-3.64	.44	-3.04	.45	-2.94	-.92	-.41
RMSE 1x	.83	.88	1.38	.71	1.71	1.52	.54	1.37
RMSE 2x	1.62	1.69	2.32	1.74	2.78	2.38	.87	2.32
Mean abs. error 1x	.67	.81	1.08	.62	1.48	1.12	.50	1.23
Mean abs. error 2x	1.43	1.53	1.61	1.55	2.48	1.81	.76	2.05
Mean abs. % error 1x	1.91	2.61	2.38	1.60	4.23	3.03	1.57	3.57
Mean abs. % error 2x	3.53	4.36	3.06	3.57	6.24	4.20	2.15	5.20
Mean actual 1x	34.6	32.2	44.5	38.5	36.1	37.1	31.0	37.9
Mean predicted 1x	34.9	32.9	44.4	39.0	35.6	37.9	31.2	37.5
Mean actual STD 1x	10.8	5.5	5.3	3.5	7.6	6.9	4.8	7.8
Mean predicted STD 1x	10.9	5.6	4.3	3.8	7.3	7.2	4.5	8.2
Mean actual 2x	41.1	36.9	50.5	43.2	42.8	42.9	35.6	43.5
Mean predicted 2x	42.1	38.2	50.1	44.5	42.3	44.2	35.9	43.6
Mean actual STD 2x	11.0	7.0	6.8	3.3	8.8	7.8	4.7	7.4
Mean predicted STD 2x	11.4	6.9	5.3	4.0	8.3	8.2	4.7	7.7

Table 12. Validation statistics at the stand-level for stocking (sph) by Region.

STATISTIC	CNI	GDNS	HBAY	CLAYS	SOUTH	SANDS	CANTY	TWC
number of plots	21	12	12	13	10	16	5	6
1x projection pd. (yrs)	6	5	6	5	5	5	3	5
2x projection pd. (yrs)	13	9	11	10	10	11	9	10
R-square 1x	.99	.98	.99	.99	.97	.98	.99	.99
R-square 2x	.91	.93	.98	.98	.84	.92	.91	.94
Mean error 1x	47	41	5	5	31	15	-15	8
Mean error 2x	74	64	11	18	67	30	-35	17
Mean % error 1x	4.66	4.58	.73	1.54	.63	4.58	-2.02	2.47
Mean % error 2x	9.52	7.84	1.97	4.16	2.58	9.20	-4.67	4.89
RMSE 1x	68	64	13	11	114	17	19	11
RMSE 2x	105	111	22	20	218	33	49	21
Mean abs. error 1x	48	41	9	10	45	15	15	8
Mean abs. error 2x	76	64	14	18	86	30	35	20
Mean abs. % error 1x	4.85	4.58	2.61	2.21	3.29	4.57	2.02	2.47
Mean abs. % error 2x	9.84	7.84	3.95	4.16	6.81	9.20	4.67	5.93
Mean actual 1x	844	654	310	431	632	323	650	356
Mean predicted 1x	796	613	305	426	601	308	665	347
Mean actual STD 1x	581	472	185	159	651	123	205	104
Mean predicted STD 1x	544	425	174	166	540	118	215	104
Mean actual 2x	638	608	303	417	593	322	630	337
Mean predicted 2x	564	545	292	400	526	292	665	320
Mean actual STD 2x	368	424	174	129	565	123	182	96
Mean predicted STD 2x	310	347	156	124	354	111	215	89

Table 13. Validation statistics at the stand-level for MTH (m) by Region.

STATISTIC	CNI	GDNS	HBAY	CLAYS	SOUTH	SANDS	CANTY	TWC
number of plots	21	12	12	13	10	16	5	6
1x projection pd. (yrs)	6	5	6	5	5	5	3	5
2x projection pd. (yrs)	13	9	11	10	10	11	9	10
R-square 1x	.92	.88	.80	.71	.73	.77	.92	.86
R-square 2x	.79	.83	.77	.11	.72	.66	.54	.77
Mean error 1x	-.08	.08	-.51	.01	-.18	-.30	-.38	-.63
Mean error 2x	.15	-.10	-.41	.42	.33	-1.11	.30	-.32
Mean % error 1x	-.31	-.06	-1.79	-.14	-.98	-1.28	-1.69	-2.15
Mean % error 2x	.25	-.83	-1.33	.76	.80	-4.16	.57	-.91
RMSE 1x	.87	1.07	1.32	1.47	1.15	1.68	.71	1.03
RMSE 2x	1.20	1.51	1.54	2.79	1.47	2.37	2.03	1.12
Mean abs. error 1x	.74	.87	1.09	1.18	.96	1.43	.62	.77
Mean abs. error 2x	.88	1.27	1.29	2.50	1.17	2.07	1.50	.88
Mean abs. % error 1x	2.10	3.36	3.50	3.69	4.05	5.39	2.58	2.69
Mean abs. % error 2x	2.05	4.30	3.56	6.78	4.03	7.04	4.77	2.70
Mean actual 1x	35.96	27.41	31.81	32.05	24.66	26.58	24.50	27.88
Mean predicted 1x	36.03	27.33	32.32	32.05	24.84	26.88	24.88	28.52
Mean actual STD 1x	3.09	3.17	3.11	2.84	2.34	3.62	2.77	3.03
Mean predicted STD 1x	2.81	2.51	2.68	2.64	2.05	3.75	2.51	3.56
Mean actual 2x	42.90	32.08	37.53	36.92	30.68	30.08	30.24	33.45
Mean predicted 2x	42.75	32.18	37.94	36.49	30.35	31.18	29.94	33.77
Mean actual STD 2x	2.69	3.82	3.31	3.09	2.93	4.19	3.34	2.58
Mean predicted STD 2x	2.19	2.79	2.53	2.47	2.41	3.78	2.31	3.04

Table 14. Validation statistics at the stand-level for MTD (cm) by Region.

STATISTIC	CNI	GDNS	HBAY	CLAYS	SOUTH	SANDS	CANTY	TWC
number of plots	21	12	12	13	10	16	5	6
1x projection pd. (yrs)	6	5	6	5	5	5	3	5
2x projection pd. (yrs)	13	9	11	10	10	11	9	10
R-square 1x	.97	.96	.81	.93	.80	.95	.96	.96
R-square 2x	.87	.91	.71	.68	.64	.91	.95	.90
Mean error 1x	.56	-.23	.08	.39	2.07	-.41	-.44	.80
Mean error 2x	1.31	-.00	.72	.41	3.03	-.18	-.16	.48
Mean % error 1x	1.29	-.54	-.03	.83	4.72	-.91	-1.21	1.75
Mean % error 2x	2.62	.07	.94	.76	5.78	-.31	-.36	.86
RMSE 1x	1.10	.71	1.62	.82	2.86	1.51	.83	1.42
RMSE 2x	2.26	1.20	2.66	1.49	4.35	2.10	.83	2.04
Mean abs. error 1x	.88	.52	1.24	.70	2.17	1.19	.76	1.27
Mean abs. error 2x	1.97	1.13	1.80	1.27	3.59	1.69	.64	1.58
Mean abs. % error 1x	1.97	1.33	2.40	1.51	4.98	2.85	1.93	2.91
Mean abs. % error 2x	3.70	2.49	2.97	2.47	6.87	3.43	1.37	3.29
Mean actual 1x	45.6	39.5	51.5	46.2	42.6	42.1	38.7	46.0
Mean predicted 1x	45.1	39.7	51.5	45.8	40.6	42.5	39.1	45.2
Mean actual STD 1x	6.8	3.5	3.8	3.3	6.8	6.8	4.5	8.0
Mean predicted STD 1x	7.1	3.8	2.7	3.2	6.6	7.1	4.3	7.9
Mean actual 2x	54.0	45.3	59.1	52.2	50.8	48.9	45.1	52.2
Mean predicted 2x	52.7	45.3	58.4	51.8	47.8	49.1	45.3	51.7
Mean actual STD 2x	6.4	4.2	5.2	2.8	7.6	7.4	4.2	7.1
Mean predicted STD 2x	7.5	4.7	3.4	2.9	7.7	7.8	4.3	7.1

Table 15. Validation statistics at the tree-level for dbh_i (cm) by Region.

STATISTIC	CNI	GDNS	HBAY	CLAYS	SOUTH	SANDS	CANTY	TWC
number of trees	1352	411	375	384	486	611	160	213
1x projection pd. (yrs)	6	5	6	5	4	5	3	5
number of trees	1076	387	368	372	453	610	156	202
2x projection pd. (yrs)	13	9	11	10	9	10	9	10
R-square 1x	.97	.96	.90	.95	.95	.94	.98	.94
R-square 2x	.92	.92	.82	.87	.91	.85	.94	.85
Mean error 1x	.00	-.38	-.27	-.41	.38	-.43	-.20	.44
Mean error 2x	-.18	-.58	.23	-1.04	.35	-.50	-.64	.09
Mean % error 1x	-.23	-1.57	-1.07	-1.41	.53	-1.35	-.88	.82
Mean % error 2x	-1.14	-2.58	-.50	-3.07	-.10	-1.77	-2.33	-.68
RMSE 1x	2.06	1.59	2.83	1.73	2.45	2.01	.99	2.73
RMSE 2x	4.01	2.92	4.86	3.24	4.20	3.61	2.26	4.33
Mean abs. error 1x	1.48	1.25	2.15	1.31	1.73	1.56	.80	2.10
Mean abs. error 2x	2.95	2.32	3.76	2.47	3.15	2.86	1.81	3.31
Mean abs. % error 1x	4.88	4.45	5.27	3.69	5.47	4.58	2.65	6.44
Mean abs. % error 2x	8.15	7.28	8.20	6.24	8.77	7.30	5.43	8.72
Mean actual 1x	30.0	29.3	41.5	37.1	29.3	35.2	30.8	36.5
Mean predicted 1x	30.0	29.7	41.8	37.5	29.0	35.7	31.0	36.0
Mean actual STD 1x	12.3	8.4	9.1	7.9	11.5	8.1	7.9	10.7
Mean predicted STD 1x	12.2	8.4	8.8	7.7	11.0	8.4	7.7	10.7
Mean actual 2x	37.4	33.6	46.8	41.7	34.6	40.8	35.3	42.1
Mean predicted 2x	37.6	34.2	46.6	42.8	34.3	41.3	35.9	42.0
Mean actual STD 2x	14.2	10.2	11.6	8.9	13.9	9.5	9.1	11.3
Mean predicted STD 2x	14.0	10.0	10.7	8.5	13.2	9.5	8.8	11.1

Table 16. Validation statistics at the tree-level for ht_i (m) by Region.

STATISTIC	CNI	GDNS	HBAY	CLAYS	SOUTH	SANDS	CANTY	TWC
number of trees	159	85	57	70	107	133	34	57
1x projection pd. (yrs)	6	5	6	5	5	5	3	5
number of trees	124	81	51	69	72	128	17	47
2x projection pd. (yrs)	13	9	11	9	10	11	9	10
R-square 1x	.83	.84	.76	.74	.71	.77	.78	.74
R-square 2x	.59	.77	.65	.51	.66	.63	.65	.51
Mean error 1x	-.22	.05	-.67	-.04	-.04	.05	-.22	.03
Mean error 2x	.17	-.17	-.39	-.36	.20	-.76	.24	-.05
Mean % error 1x	-.82	-.36	-2.56	-.34	-.65	-.17	-1.02	.44
Mean % error 2x	.11	-1.63	-1.56	-.137	.14	-3.34	.57	-.23
RMSE 1x	1.65	1.53	1.78	1.89	1.49	1.83	1.23	2.02
RMSE 2x	2.44	2.37	2.29	2.58	2.05	2.74	1.86	2.41
Mean abs. error 1x	1.28	1.17	1.40	1.46	1.14	1.45	1.00	1.52
Mean abs. error 2x	1.86	1.91	1.73	2.08	1.57	2.25	1.57	1.95
Mean abs. % error 1x	3.79	4.80	4.76	4.77	4.99	5.50	4.27	6.00
Mean abs. % error 2x	4.51	6.89	4.86	5.99	5.50	7.77	5.24	6.17
Mean actual 1x	34.81	26.12	31.03	31.00	23.40	26.58	23.92	26.74
Mean predicted 1x	35.03	26.07	31.70	31.05	23.44	26.52	24.15	26.71
Mean actual STD 1x	4.07	3.81	3.65	3.74	2.78	3.84	2.64	4.04
Mean predicted STD 1x	3.86	3.31	3.10	3.61	2.34	3.65	2.73	4.92
Mean actual 2x	42.02	30.59	36.83	35.18	29.24	30.17	29.64	32.89
Mean predicted 2x	41.85	30.76	37.22	35.54	29.04	30.93	29.40	32.94
Mean actual STD 2x	3.83	4.95	3.91	3.73	3.54	4.48	3.27	3.50
Mean predicted STD 2x	3.26	3.71	2.82	3.34	2.84	3.75	3.02	4.07

Table 17. Additional validation statistics at the stand- and tree-level for CANTY.

STATISTIC	Stand-Level					Tree-Level	
	G (m ² /ha)	dbh _q (cm)	N (sph)	MTH (m)	MTD (cm)	dbh _i (cm)	ht _i (m)
number of plots or trees	11					387	42
2x projection pd. (yrs)	9					9	8
R-square 2x	.43	.97	.90	.77	.89	.90	.72
Mean error 2x	-2.86	-.15	-24	.46	.62	-.63	.25
Mean % error 2x	-4.52	-.41	-3.66	1.13	1.24	-2.62	.56
RMSE 2x	5.74	.72	42	1.57	1.53	3.05	1.83
Mean abs. error 2x	4.89	.62	33	1.32	1.13	2.32	1.52
Mean abs. % error 2x	7.58	1.68	5.11	4.07	2.29	6.74	4.91
Mean actual 2x	64.57	37.7	596	31.83	48.4	37.0	30.87
Mean predicted 2x	67.43	37.9	620	31.36	47.8	37.7	30.62
Mean actual STD 2x	8.00	4.1	142	3.42	4.7	9.4	3.51
Mean predicted STD 2x	9.08	4.1	161	2.53	4.8	8.8	3.21

Table 18. Additional validation statistics at the stand- and tree-level for TWC.

STATISTIC	Stand-Level					Tree-Level	
	G (m ² /ha)	dbh _q (cm)	N (sph)	MTH (m)	MTD (cm)	dbh _i (cm)	ht _i (m)
number of plots or trees	20					444	178
2x projection pd. (yrs)	10					10	10
R-square 2x	.80	.86	.85	.53	.82	.84	.54
Mean error 2x	.08	.22	-3	-.92	.56	.53	-.36
Mean % error 2x	-2.10	.58	-3.23	-3.46	.99	.85	-1.65
RMSE 2x	5.08	2.16	31	2.50	2.33	4.30	2.99
Mean abs. error 2x	4.44	1.74	24	2.00	1.94	3.31	2.46
Mean abs. % error 2x	13.00	4.28	10.71	6.72	3.99	8.86	8.51
Mean actual 2x	38.71	43.5	265	31.18	49.8	41.9	30.24
Mean predicted 2x	38.63	43.3	269	32.10	49.3	41.4	30.60
Mean actual STD 2x	11.54	5.9	84	3.73	5.6	10.7	4.41
Mean predicted STD 2x	10.59	6.4	78	3.11	5.4	10.9	4.35

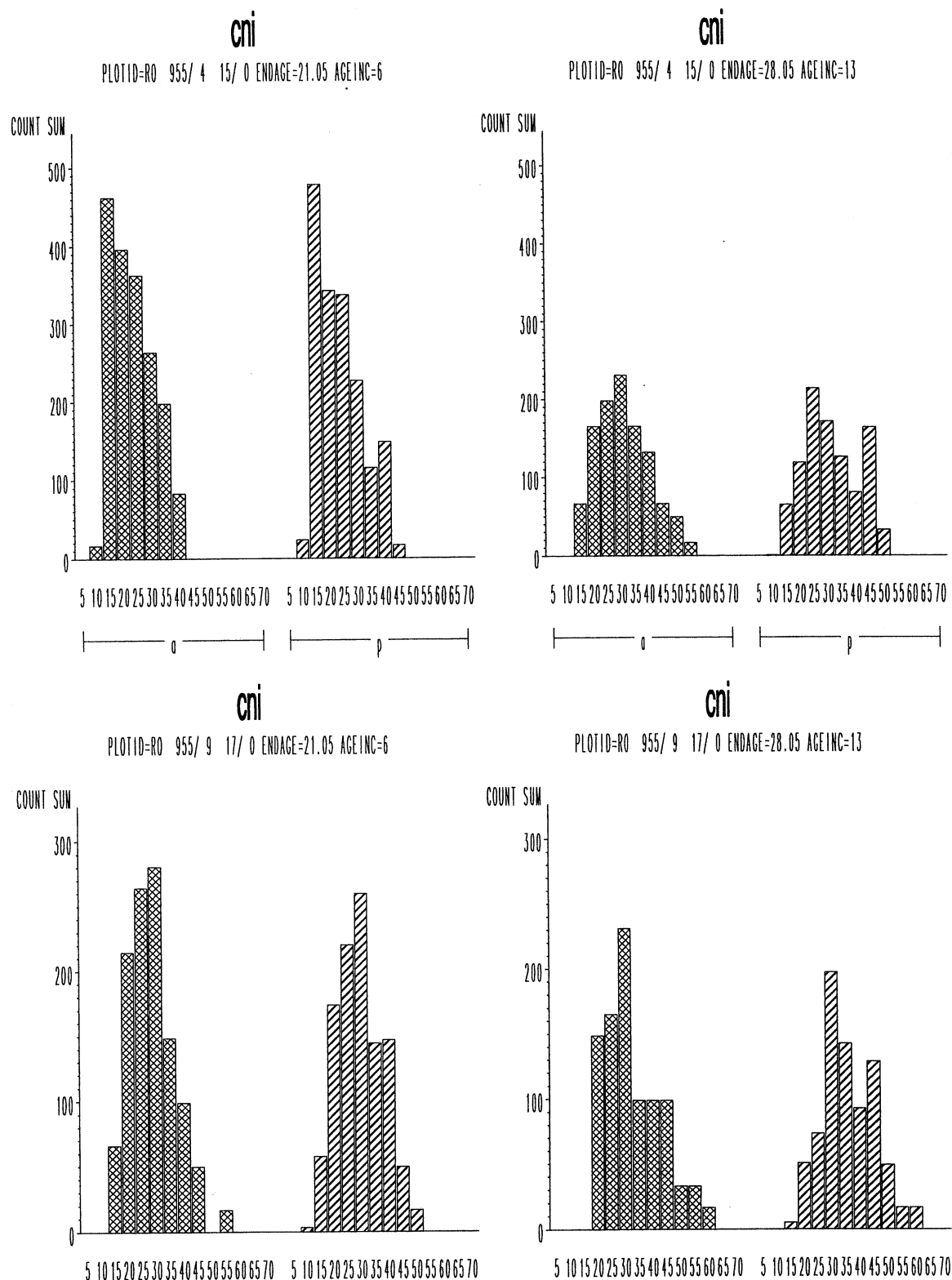


Figure 5. CNI actual (a) and predicted (p) diameter distributions of selected plots for the short and long projection periods. From left to right set: short and long projection periods. From top to bottom set: the plot nearest to a dbh_i variance ratio of 1.0 and the plot nearest to the region's mean dbh_i variance ratio.

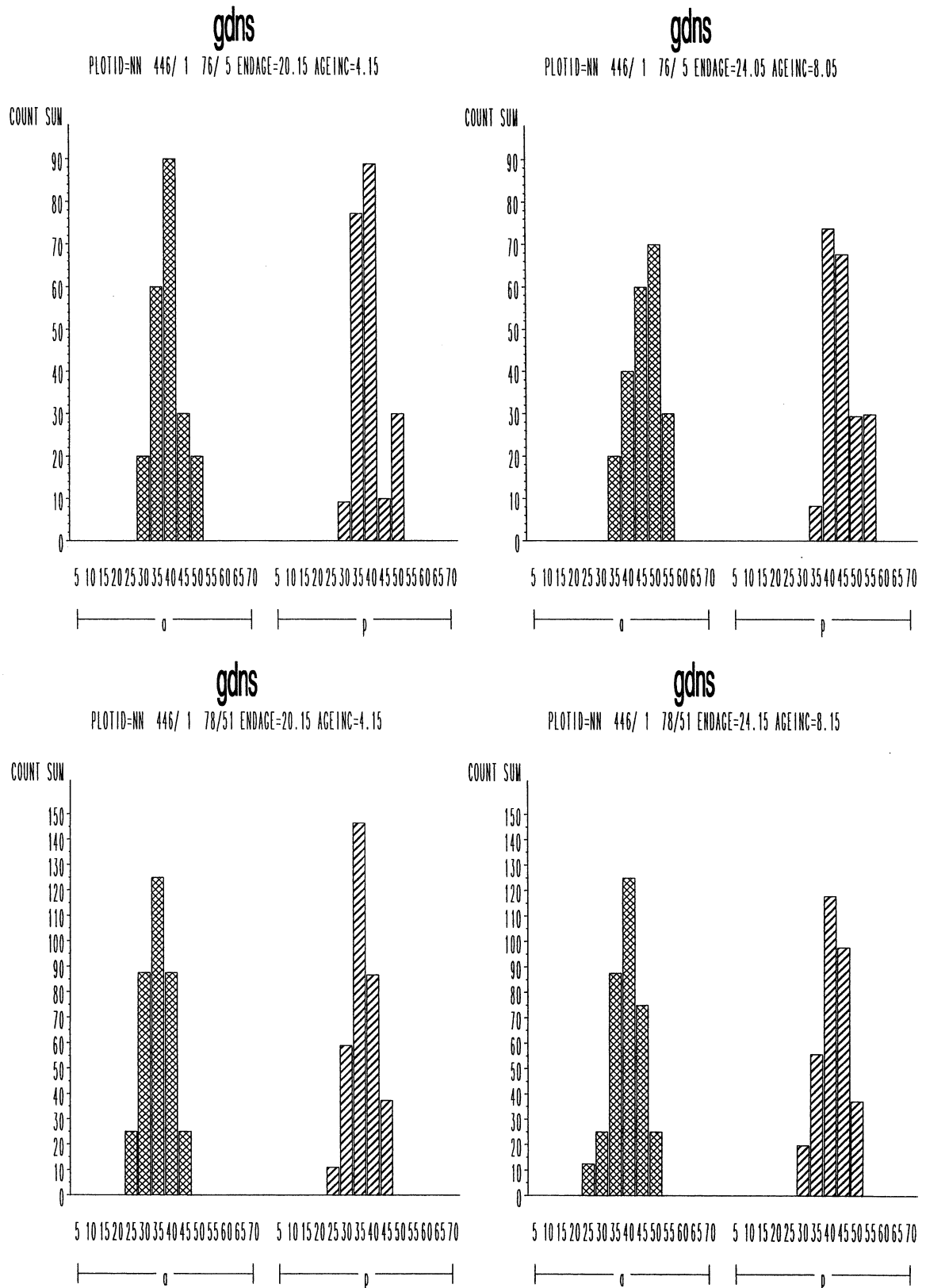


Figure 6. GDNS actual and predicted diameter distributions of selected plots for the short and long projection periods. From left to right set: short and long projection periods. From top to bottom set: the plot nearest to a dbh_i variance ratio of 1.0 and the plot nearest to the region's mean dbh_i variance ratio.

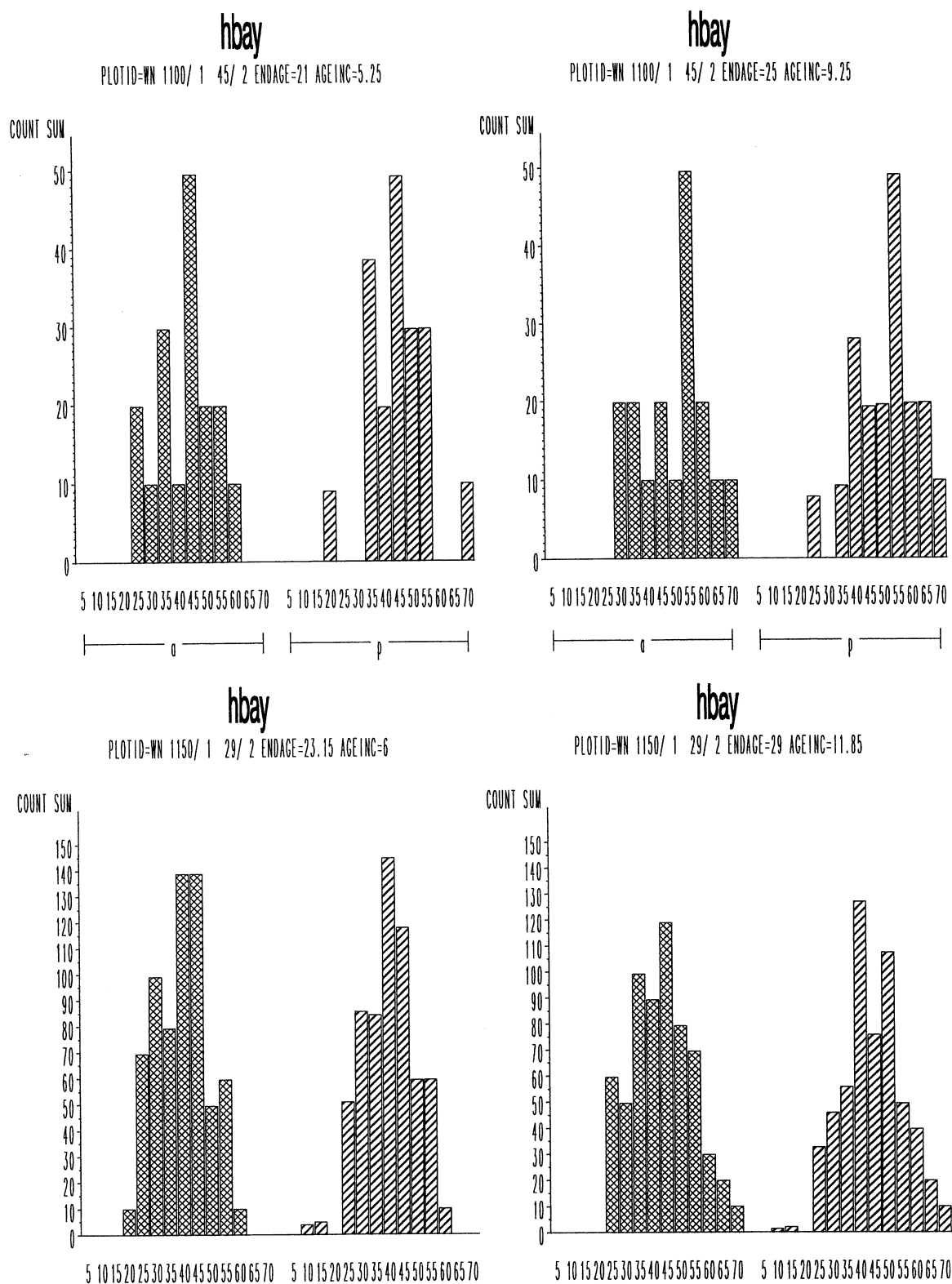


Figure 7. Hbay actual (a) and predicted (p) diameter distributions of selected plots for the short and long projection periods. From left to right set: short and long projection periods. From top to bottom set: the plot nearest to a dbh_i variance ratio of 1.0 and the plot nearest to the region's mean dbh_i variance ratio.

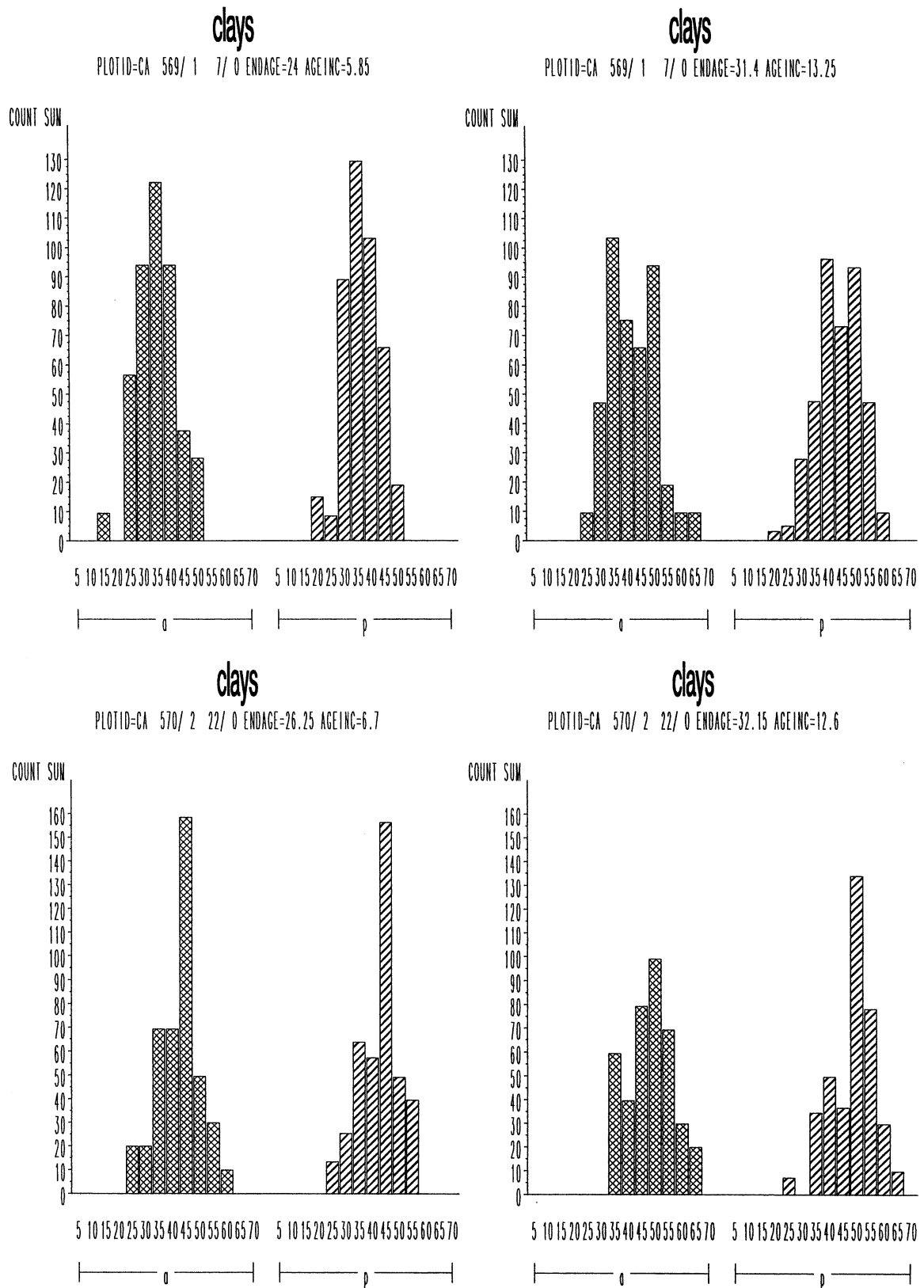


Figure 8. CLAYS actual (a) and predicted (p) diameter distributions of selected plots for the short and long projection periods. From left to right set: short and long projection periods. From top to bottom set: the plot nearest to a dbh_i variance ratio of 1.0 and the plot nearest to the region's mean dbh_i variance ratio.

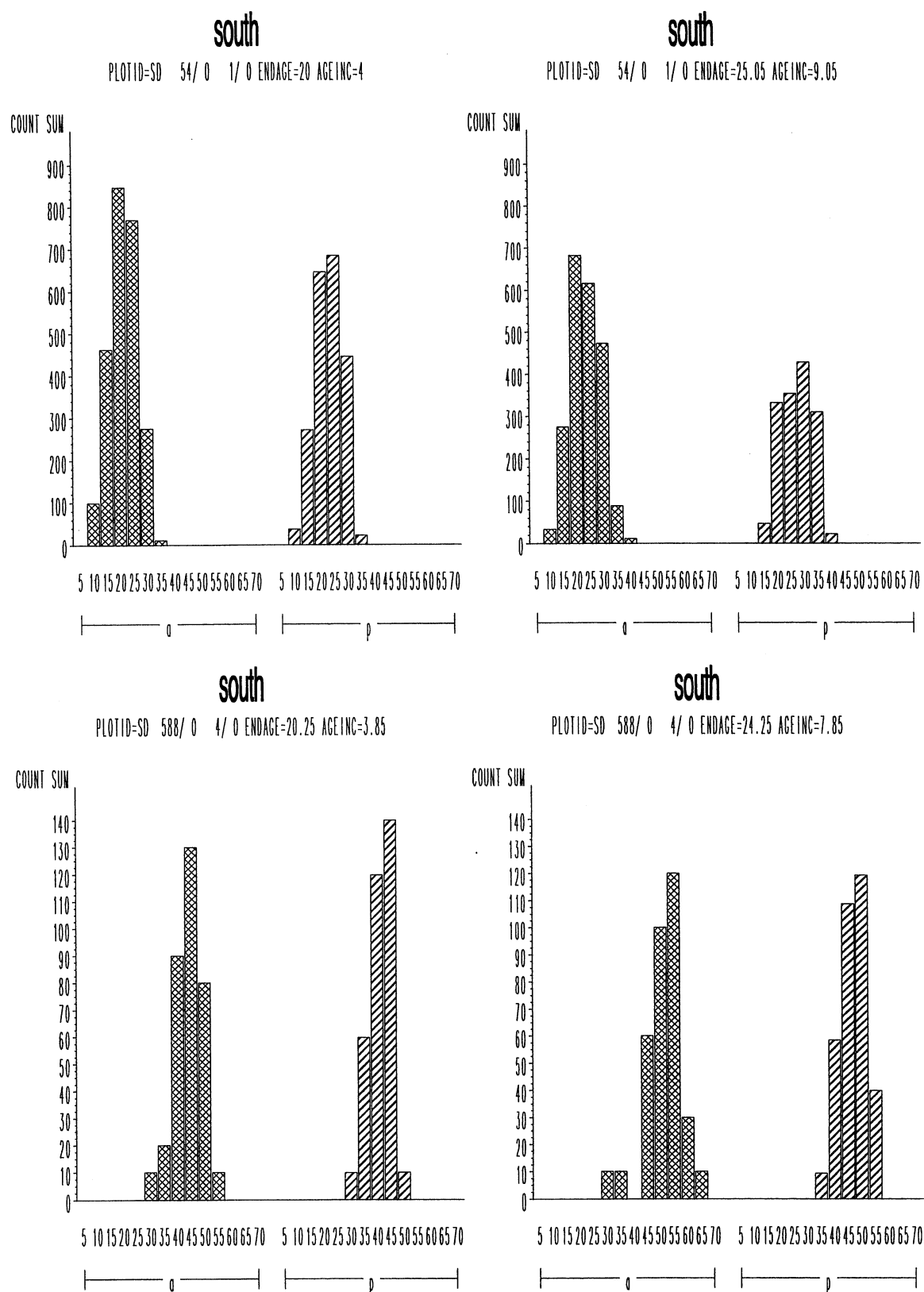


Figure 9. SOUTH actual (a) and predicted (p) diameter distributions of selected plots for the short and long projection periods. From left to right set: short and long projection periods. From top to bottom set: the plot nearest to a dbh_i variance ratio of 1.0 and the plot nearest to the region's mean dbh_i variance ratio.

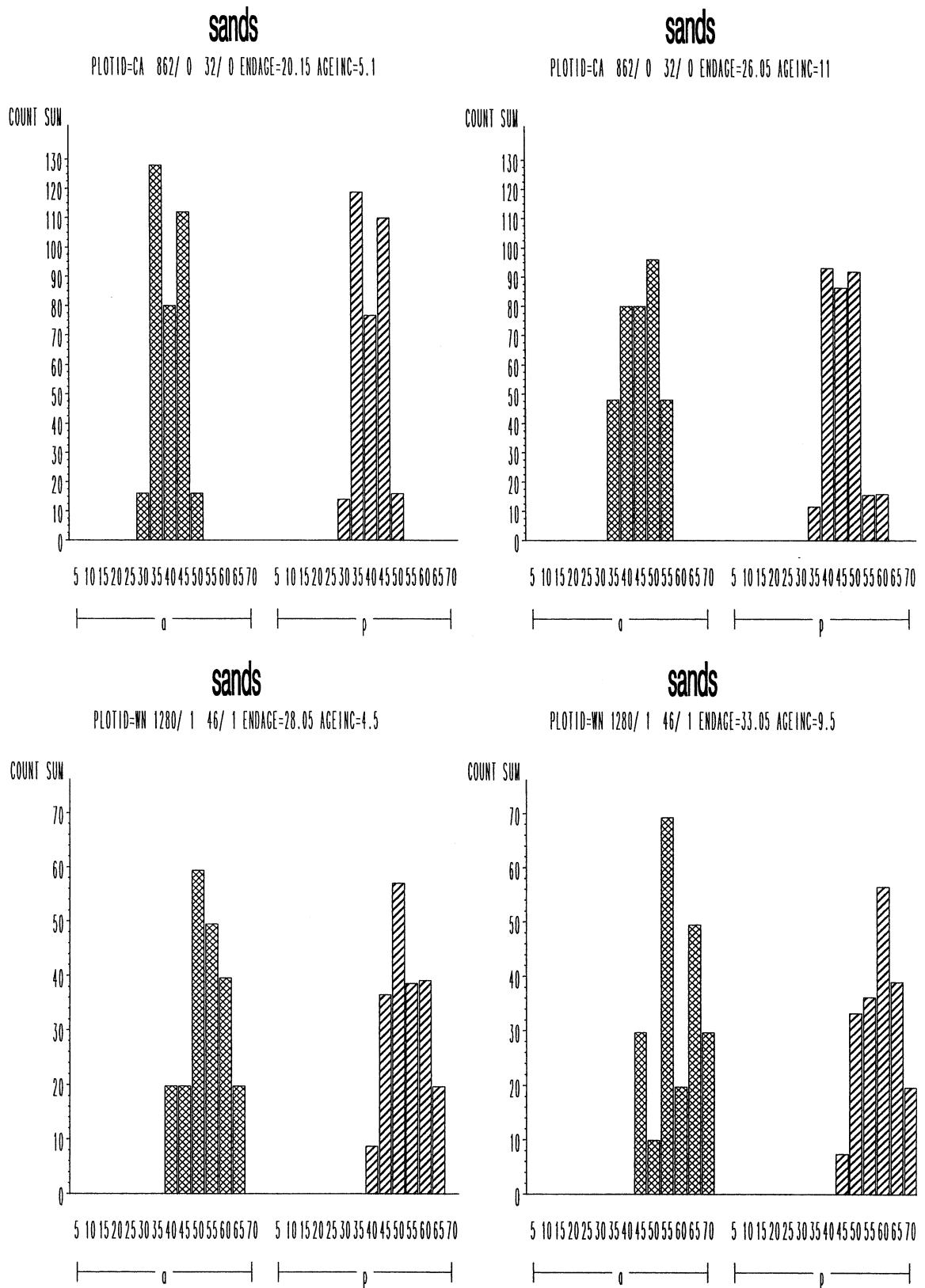


Figure 10. SANDS actual (a) and predicted (p) diameter distributions of selected plots for the short and long projection periods. From left to right set: short and long projection periods. From top to bottom set: the plot nearest to a dbh_i variance ratio of 1.0 and the plot nearest to the region's mean dbh_i variance ratio.

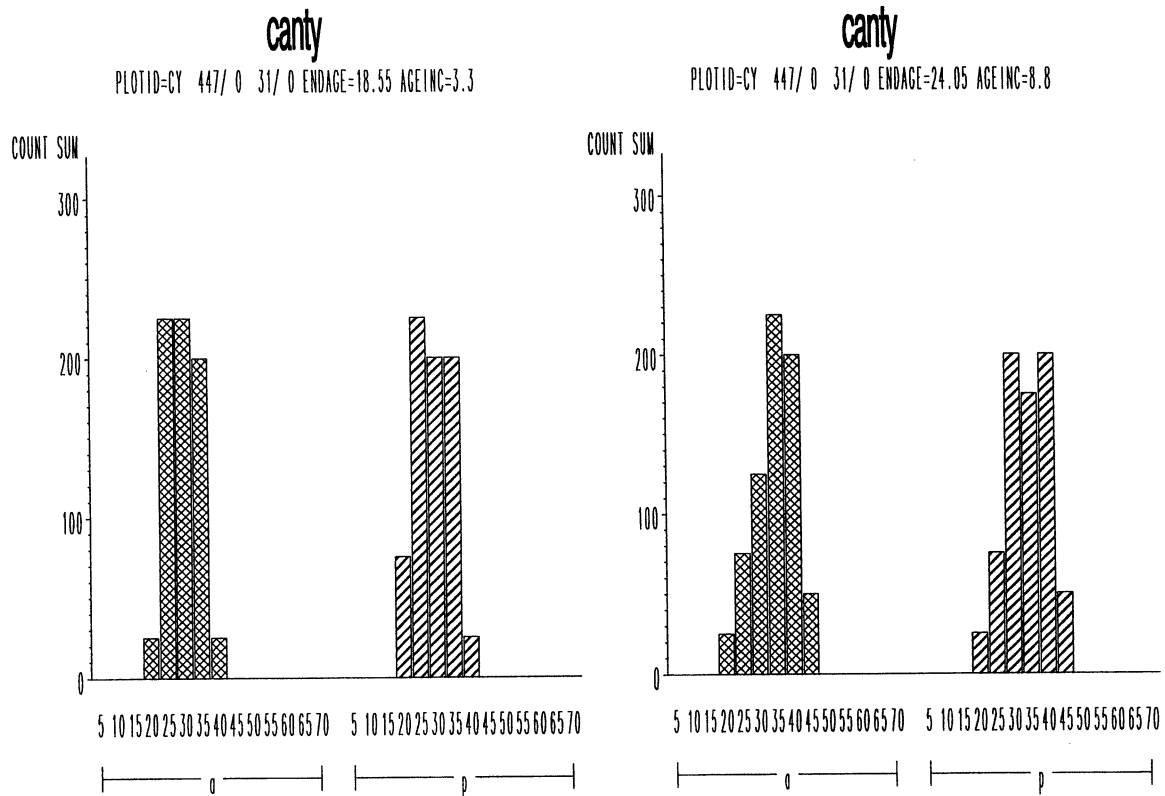


Figure 11. CANTY actual (a) and predicted (p) diameter distributions of selected plots for the short and long projection periods. From left to right set: short and long projection periods. From top to bottom set: the plot nearest to a dbh_i variance ratio of 1.0 and the plot nearest to the region's mean dbh_i variance ratio.

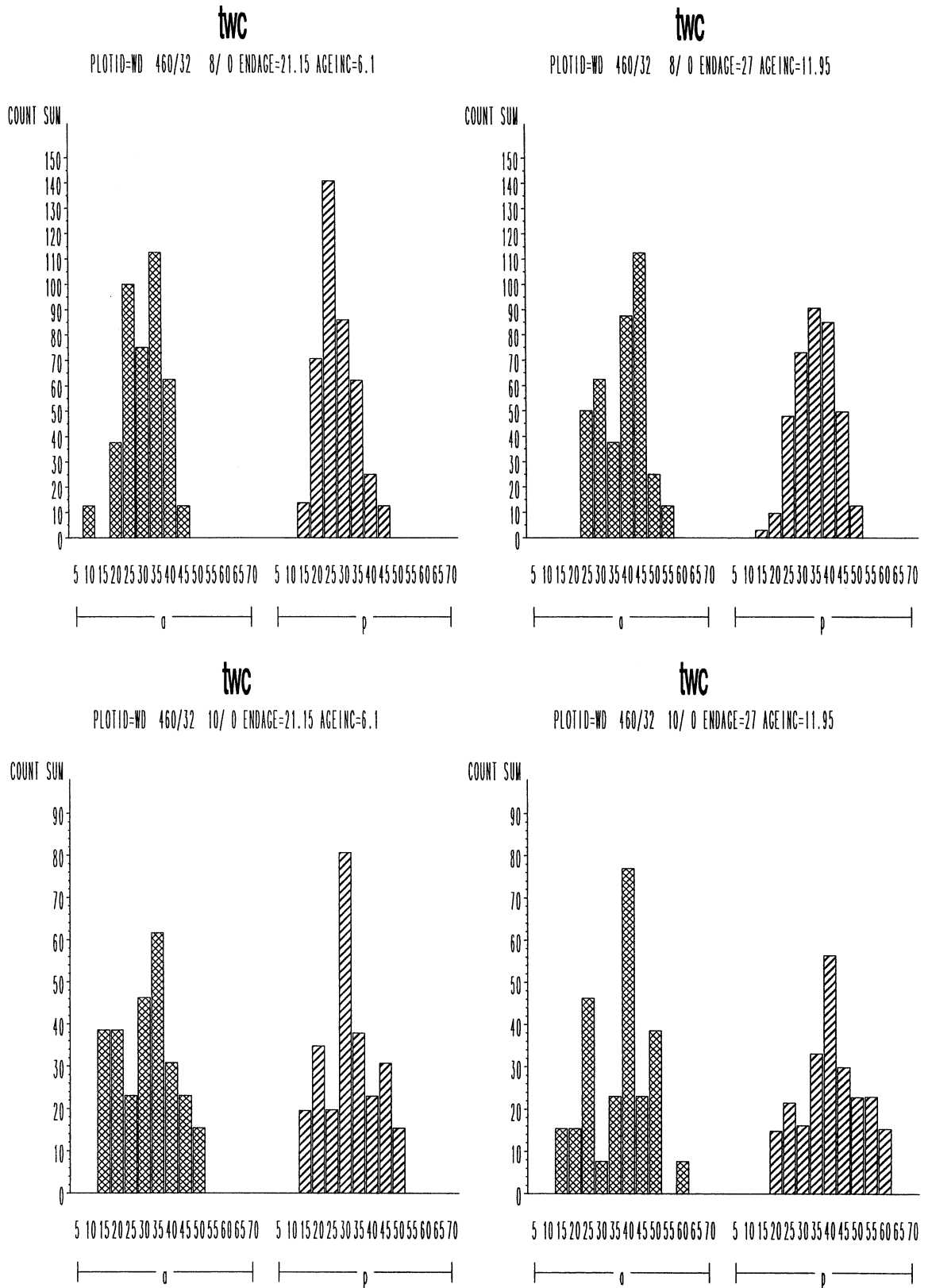


Figure 12. TWC actual (a) and predicted (p) diameter distributions of selected plots for the short and long projection periods. From left to right set: short and long projection periods. From top to bottom set: the plot nearest to a dbh_i variance ratio of 1.0 and the plot nearest to the region's mean dbh_i variance ratio.

DISCUSSION

General

The validation exercise, while not strictly "independent", is at least very much quasi-independent. This is because so few observations were used to develop the prediction equations (both in total number and random selection), relative to the observations contributing to the validation exercise. Also, the validation exercise is based on the prediction equations (diameter and height increment, and survival) working as an iterative prediction system, contrary to the original fitting of separate equations at single, points-in-time.

The validation exercise has focused on very discriminate time periods of prediction, i.e., a given "short" time period (say, 5 years), and a strictly complementary "long" time period (say, twice the length of the "short", i.e., 10 years). This approach allows for critical examination of compounded error of prediction, i.e., cascading error, on a strictly one-to-one comparative basis.

Height prediction

The R^2 for the prediction of annual ht_i increment is significantly less, than that from the prediction of static ht_i / MTH ratio, but this is largely a mathematical artefact. A reduction in the variability of the observed (going from ht_i increment to ht_i / MTH ratio) with a corresponding increase in the variability of the predicted (going from ht_i / MTH ratio to ht_i increment) contrives to mathematically produce, by default, a lower R^2 value.

Unfortunately, the prediction of interest, annual ht_i increment, is by nature, highly variable and less correlated with tree-size attributes, than annual dbh_i increment. Nonetheless, 60-80 percent of predicted observations are within ± 0.50 to $0.75m$ of the actual ht_i increment.

Even though the direct prediction of ht_i increment has low R^2 values, the validation exercise has established that this prediction approach is superior to the use of the ht_i / MTH ratio prediction approach, which resulted in severely biased ht_i increment predictions.

The utility of regional dummy variables in the ht_i prediction equation, suggests that separate, specific regional ht_i increment equations should be investigated, similar to the regionally specific dbh_i increment equations. This regional approach may identify regionally specific explanatory variables, which may result in improved R^2 values.

TWC equations

The results of directly fitting TWC dbh_i and ht_i increment equations suggest that TWC and GDNS can be modelled similarly (using the same explanatory variables), but with less confidence in Westland. But once again, complete reliance on regression R^2 is dodgy, as directly fit TWC equations had significantly poorer R^2 than GDNS, but the validation exercise revealed only marginally better validation statistics with the use of GDNS equations, instead of directly fit TWC equations.

Until further investigations prove otherwise, TWC will be represented with GDNS growth equations in ITGM, as validation results fully support this.

Validation statistics

Validation statistics reveal that, on-average, 5 to 10 year tree- and stand-level diameter and height prediction error is generally $\leq 3\%$ (usually over-predicted), while absolute prediction error is generally $\leq 7\%$. Accompanying predicted tree-diameter distributions generally appear reasonable, and skewed and multi-modal diameter distributions (i.e., other than "normal") are accommodated well. These results indicate that ITGM, as an iterative prediction system for mid-rotation onward tree and stand growth and survival, operates within acceptable error limits without any obvious anomalies or mechanistic mathematical flaws.

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APPENDIX A:

Regional PSPs and Simulation Start- and End-ages (short- and long-term)

CNI

plotid	startage	endage1	endage2
RO 416/ 0 0/ 0	16.00	23.00	31.00
RO 421/ 0 0/ 0	15.15	21.05	27.05
RO 464/ 0 0/ 0	16.05	22.00	29.00
RO 488/ 0 3/ 0	15.40	21.40	28.25
RO 685/ 2 7/ 0	15.25	22.25	30.15
RO 690/ 0 3/ 0	15.15	21.05	27.05
RO 693/ 0 0/ 0	16.00	22.00	28.00
RO 695/ 1 16/ 0	15.15	22.15	30.15
RO 696/ 1 15/ 0	15.15	21.15	27.15
RO 746/ 0 0/ 0	15.15	21.15	27.05
RO 911/ 1 1/ 0	22.05	26.65	32.25
RO 955/ 4 5/ 0	15.05	21.05	28.05
RO 955/ 4 13/ 0	15.05	21.05	28.05
RO 955/ 4 15/ 0	15.05	21.05	28.05
RO 955/ 6 9/ 0	15.05	21.05	28.05
RO 955/ 6 11/ 0	15.05	21.05	28.05
RO 955/ 6 20/ 0	15.05	21.05	28.05
RO 955/ 7 18/ 0	15.05	21.15	28.00
RO 955/ 7 25/ 0	15.05	21.15	28.00
RO 955/ 9 15/ 0	15.05	21.05	28.05
RO 955/ 9 17/ 0	15.05	21.05	28.05

GDNS

plotid	startage	endage1	endage2
NN 183/ 0 1/ 0	20.55	26.40	32.05
NN 278/ 1 9/ 0	16.15	22.00	27.00
NN 278/ 1 13/ 0	16.15	22.00	28.00
NN 376/ 0 1/ 0	15.05	19.00	23.05
NN 446/ 1 76/ 5	16.00	20.15	24.05
NN 446/ 1 77/ 1	15.00	19.25	23.05
NN 446/ 1 78/ 26	16.00	20.15	24.15
NN 446/ 1 78/ 51	16.00	20.15	24.15
NN 446/ 2 77/ 8	15.00	20.25	26.00
NN 462/ 0 69/ 4	16.00	20.00	24.00
NN 462/ 0 69/ 5	16.00	20.00	24.00
NN 462/ 0 69/ 10	17.00	21.00	25.00

HBAY

plotid	startage	endage1	endage2
WN 1100/ 1 35/ 2	15.00	21.00	27.00
WN 1100/ 1 42/ 3	19.25	22.85	26.00
WN 1100/ 1 45/ 1	17.75	24.00	29.00
WN 1100/ 1 45/ 2	15.75	21.00	25.00
WN 1100/ 1 47/ 1	16.95	20.75	24.00
WN 1100/ 1 63/ 1	15.75	22.00	27.00
WN 1150/ 1 29/ 1	17.15	23.15	29.00
WN 1150/ 1 29/ 2	17.15	23.15	29.00
WN 1150/ 1 36/ 7	16.15	22.05	28.00
WN 1150/ 1 39/ 3	15.15	21.15	27.00
WN 1150/ 1 50/ 1	15.55	21.00	27.00
WN 1320/ 1 220/ 1	16.75	23.00	29.00

CLAYS

plotid	startage	endage1	endage2
AK 286/ 1 5/ 0	16.00	20.00	25.15
AK 286/ 2 3/ 0	16.00	20.00	25.15
AK 286/ 3 5/ 0	16.00	20.00	25.15
AK 286/ 3 6/ 0	16.00	20.00	25.15
AK 401/ 0 7/ 0	21.15	26.15	30.15
AK 401/ 0 9/ 0	15.05	19.65	24.15
AK 401/ 0 11/ 0	25.15	29.15	33.00
AK 458/ 1 3/ 0	21.00	24.85	29.15
AK 501/ 4 5/ 0	17.15	22.00	27.55
CA 569/ 1 7/ 0	18.15	24.00	31.40
CA 570/ 2 22/ 0	19.55	26.25	32.15
CA 656/ 0 8/ 0	16.00	20.75	24.25
CA 656/ 0 20/ 0	16.00	20.75	24.25

SOUTH

plotid	startage	endage1	endage2
SD 54/ 0 1/ 0	16.00	20.00	25.05
SD 170/ 0 1/ 0	15.00	21.05	27.05
SD 170/ 0 22/ 0	15.85	20.25	26.00
SD 180/ 0 22/ 0	16.15	21.05	26.05
SD 180/ 0 23/ 0	16.00	20.55	26.00
SD 180/ 0 24/ 0	16.55	23.25	29.00
SD 188/ 0 22/ 0	15.15	21.05	27.25
SD 588/ 0 2/ 0	16.40	20.25	24.25
SD 588/ 0 3/ 0	16.40	20.25	24.25
SD 588/ 0 4/ 0	16.40	20.25	24.25

SANDS

plotid	startage	endage1	endage2
AK 35/ 0 14/ 0	18.05	22.05	27.05
AK 35/ 0 16/ 0	15.15	22.05	30.00
AK 964/ 0 7/ 0	24.00	30.00	37.00
CA 242/ 0 3/ 0	24.00	27.75	31.95
CA 427/ 0 2/ 0	25.25	29.25	33.85
CA 427/ 0 3/ 0	25.25	29.25	33.85
CA 434/ 0 4/ 0	17.85	24.00	31.05
CA 862/ 0 15/ 0	15.00	20.15	26.05
CA 862/ 0 27/ 0	15.15	20.00	26.05
CA 862/ 0 32/ 0	15.05	20.15	26.05
CA 862/ 0 52/ 0	15.00	19.15	23.15
WN 154/ 0 1/ 0	16.00	20.00	25.05
WN 1280/ 1 37/ 6	16.65	23.15	29.00
WN 1280/ 1 46/ 1	23.55	28.05	33.05
WN 1300/ 1 107/ 3	15.40	21.00	26.00
WN 1300/ 1 107/ 4	15.40	21.00	26.00

CANTY

plotid	startage	endage1	endage2
CY 447/ 0 6/ 0	16.25	19.55	25.05*
CY 447/ 0 8/ 0	16.25	19.55	25.05*
CY 447/ 0 17/ 0	16.25	19.55	25.05*
CY 447/ 0 31/ 0	15.25	18.55	24.05*
CY 560/ 2 1/ 0	20.00	23.00	28.00
CY 560/ 2 1/ 0	20.00	27.00*	-
CY 189/ 0 1/ 0	20.00	29.85*	-
CY 432/ 0 1/ 0	17.00	28.15*	-
CY 432/ 0 2/ 0	20.05	28.15*	-
CY 432/ 0 3/ 0	17.00	28.15*	-
CY 432/ 0 4/ 0	15.15	25.05*	-
CY 432/ 0 5/ 0	18.05	25.05*	-

TWC

plotid	startage	endage1	endage2
WD 98/ 0 0/ 0	19.25	23.00	26.05
WD 369/ 0 1/ 0	17.40	23.00	29.15*
WD 417/ 3 4/ 0	15.00	19.00	24.00
WD 417/ 3 4/ 0	15.00	25.00*	-
WD 460/14 83/ 0	15.15	19.15	22.00
WD 460/32 8/ 0	15.05	21.15	27.00*
WD 460/32 10/ 0	15.05	21.15	27.00
WD 460/32 10/ 0	15.05	24.15*	-
WD 345/ 0 1/ 0	15.15	25.25*	-
WD 345/ 0 3/ 0	15.15	25.25*	-
WD 345/ 0 4/ 0	15.05	25.25*	-
WD 345/ 0 5/ 0	15.05	25.25*	-
WD 345/ 0 6/ 0	16.15	25.25*	-
WD 345/ 0 7/ 0	15.15	25.25*	-
WD 345/ 0 8/ 0	15.15	25.25*	-
WD 345/ 0 9/ 0	15.05	25.25*	-
WD 370/ 0 1/ 0	15.00	25.15*	-
WD 373/ 0 1/ 0	15.15	25.25*	-
WD 417/ 1 2/ 0	19.00	29.00*	-
WD 417/ 2 2/ 0	16.00	26.00*	-
WD 417/ 4 5/ 0	16.00	25.00*	-
WD 460/32 5/ 0	15.25	24.25*	-
WD 460/32 6/ 0	15.25	24.25*	-
WD 460/32 7/ 0	15.25	24.25*	-

* Plot and startage / endage included for greater sample size

APPENDIX B:

Diameter Growth Analyses - 168 Timberland West Coast PSPs

<u>Plot id</u>	<u>Initial Age</u>	<u>Plot id</u>	<u>Initial Age</u>	<u>Plot id</u>	<u>Initial Age</u>
FR 258/0 1/0	18.00	WD 109/0 8/0	30.05	WD 460/26 13/0	26.00
FR 258/0 11/0	18.00	WD 109/0 98/0	32.05	WD 460/26 14/0	26.00
FR 258/0 12/0	18.00	WD 109/0 99/0	32.05	WD 460/26 2/0	25.15
FR 258/0 13/0	18.00	WD 337/0 1/0	24.00	WD 460/26 3/0	25.15
FR 258/0 15/0	18.00	WD 337/0 2/0	24.00	WD 460/26 5/0	27.00
FR 258/0 18/0	18.00	WD 337/0 3/0	24.00	WD 460/26 6/0	27.00
FR 258/0 19/0	18.00	WD 337/0 4/0	24.00	WD 460/26 7/0	27.00
FR 258/0 2/0	18.00	WD 337/0 5/0	24.00	WD 460/26 9/0	25.15
FR 258/0 22/0	18.00	WD 337/0 7/0	24.00	WD 460/30 87/0	18.15
FR 258/0 23/0	18.00	WD 337/0 8/0	24.00	WD 460/30 89/0	18.15
FR 258/0 24/0	18.00	WD 337/0 9/0	24.00	WD 460/30 90/0	18.15
FR 258/0 25/0	18.00	WD 345/0 9/0	16.15	WD 460/31 1/0	25.00
FR 258/0 26/0	18.00	WD 367/0 1/0	24.00	WD 460/31 118/0	15.25
FR 258/0 28/0	18.00	WD 368/0 1/0	23.00	WD 460/31 119/0	16.25
FR 258/0 30/0	18.00	WD 369/0 1/0	19.00	WD 460/31 14/0	18.00
FR 258/0 31/0	18.00	WD 370/0 1/0	17.05	WD 460/31 15/0	18.00
FR 258/0 32/0	18.00	WD 371/0 1/0	16.05	WD 460/31 2/0	22.25
FR 258/0 33/0	18.00	WD 372/0 1/0	16.05	WD 460/31 21/0	15.00
FR 258/0 35/0	18.00	WD 417/1 2/0	19.00	WD 460/31 24/0	19.00
FR 258/0 36/0	18.00	WD 417/1 4/0	15.00	WD 460/31 25/0	18.00
FR 258/0 37/0	18.00	WD 417/2 2/0	16.00	WD 460/31 26/0	24.00
FR 258/0 38/0	18.00	WD 417/2 3/0	15.00	WD 460/31 30/0	16.15
FR 258/0 45/0	18.00	WD 417/3 1/0	17.00	WD 460/31 37/0	16.15
FR 258/0 46/0	18.00	WD 417/3 3/0	21.00	WD 460/31 38/0	16.15
FR 258/0 47/0	18.00	WD 417/3 4/0	15.00	WD 460/31 5/0	29.00
FR 258/0 48/0	18.00	WD 417/3 5/0	15.00	WD 460/32 1/0	20.25
FR 258/0 5/0	18.00	WD 417/3 6/0	15.00	WD 460/32 10/0	15.05
FR 258/0 50/0	18.00	WD 417/3 8/0	21.00	WD 460/32 108/0	21.25
FR 258/0 51/0	18.00	WD 417/3 9/0	15.00	WD 460/32 109/0	21.25
FR 258/0 52/0	18.00	WD 417/4 1/0	16.00	WD 460/32 11/0	21.25
FR 258/0 53/0	18.00	WD 417/4 3/0	20.00	WD 460/32 110/0	21.00
FR 258/0 54/0	18.00	WD 417/4 4/0	15.00	WD 460/32 111/0	18.25
FR 258/0 55/0	18.00	WD 417/4 5/0	18.00	WD 460/32 112/0	18.25
FR 258/0 56/0	18.00	WD 417/4 6/0	15.00	WD 460/32 2/0	19.25
FR 258/0 58/0	18.00	WD 417/4 7/0	15.00	WD 460/32 3/0	23.00
FR 258/0 6/0	18.00	WD 417/5 1/0	16.00	WD 460/32 4/0	19.25
FR 258/0 60/0	18.00	WD 417/5 2/0	15.00	WD 460/32 5/0	16.05
FR 258/0 61/0	18.00	WD 460/19 54/0	15.00	WD 460/32 6/0	23.25
FR 258/0 62/0	18.00	WD 460/19 58/0	15.00	WD 460/32 7/0	16.05
FR 258/0 63/0	18.00	WD 460/19 60/0	16.00	WD 460/32 8/0	15.05
FR 258/0 65/0	18.00	WD 460/19 61/0	16.00	WD 460/33 4/0	27.15
FR 258/0 66/0	18.00	WD 460/19 62/0	16.00	WD 460/33 5/0	26.15
FR 258/0 67/0	18.00	WD 460/19 64/0	18.00	WD 460/33 7/0	24.15
FR 258/0 69/0	18.00	WD 460/19 65/0	18.25	WD 460/33 8/0	25.00
FR 258/0 7/0	18.00	WD 460/19 66/0	20.15	WD 460/43 10/0	25.15
FR 258/0 70/0	18.00	WD 460/19 67/0	16.15	WD 460/43 11/0	25.15
FR 258/0 71/0	18.00	WD 460/19 68/0	21.15	WD 460/43 12/0	24.15
FR 258/0 72/0	18.00	WD 460/19 69/0	16.25	WD 460/43 2/0	27.15
FR 258/0 73/0	18.00	WD 460/19 70/0	18.15	WD 460/43 47/0	21.05
FR 258/0 75/0	18.00	WD 460/19 71/0	20.15	WD 460/43 48/0	21.05
FR 258/0 9/0	18.00	WD 460/19 72/0	21.15	WD 460/43 49/0	21.05
WD 109/0 12/0	30.05	WD 460/19 73/0	21.15	WD 460/43 50/0	18.05
WD 109/0 23/0	30.05	WD 460/19 98/0	15.00	WD 460/43 51/0	18.05
WD 109/0 24/0	30.05	WD 460/26 1/0	25.15	WD 460/51 7/0	27.05
WD 109/0 71/0	32.05	WD 460/26 10/0	26.00		
WD 109/0 72/0	32.05	WD 460/26 11/0	26.00		
WD 109/0 73/0	32.05	WD 460/26 12/0	24.15		

APPENDIX C:

Height Growth Analyses - 188 Timberland West Coast PSPs

<u>Plot id</u>	<u>Initial Age</u>	<u>Plot id</u>	<u>Initial Age</u>
FR 258/0 1/0	18.00	FR 258/0 62/0	18.00
FR 258/0 10/0	18.00	FR 258/0 63/0	18.00
FR 258/0 11/0	18.00	FR 258/0 64/0	18.00
FR 258/0 13/0	18.00	FR 258/0 65/0	18.00
FR 258/0 14/0	18.00	FR 258/0 66/0	18.00
FR 258/0 15/0	18.00	FR 258/0 67/0	18.00
FR 258/0 16/0	18.00	FR 258/0 68/0	18.00
FR 258/0 17/0	18.00	FR 258/0 69/0	18.00
FR 258/0 18/0	18.00	FR 258/0 7/0	18.00
FR 258/0 19/0	18.00	FR 258/0 70/0	18.00
FR 258/0 2/0	18.00	FR 258/0 71/0	18.00
FR 258/0 20/0	18.00	FR 258/0 72/0	18.00
FR 258/0 21/0	18.00	FR 258/0 73/0	18.00
FR 258/0 22/0	18.00	FR 258/0 74/0	18.00
FR 258/0 23/0	18.00	FR 258/0 75/0	18.00
FR 258/0 24/0	18.00	FR 258/0 8/0	18.00
FR 258/0 25/0	18.00	FR 258/0 9/0	18.00
FR 258/0 26/0	18.00	WD 109/0 100/0	32.05
FR 258/0 27/0	18.00	WD 109/0 12/0	30.05
FR 258/0 28/0	18.00	WD 109/0 23/0	30.05
FR 258/0 29/0	18.00	WD 109/0 24/0	30.05
FR 258/0 3/0	18.00	WD 109/0 72/0	32.05
FR 258/0 30/0	18.00	WD 109/0 73/0	32.05
FR 258/0 31/0	18.00	WD 109/0 8/0	30.05
FR 258/0 32/0	18.00	WD 109/0 98/0	32.05
FR 258/0 33/0	18.00	WD 109/0 99/0	32.05
FR 258/0 34/0	18.00	WD 345/0 2/0	16.15
FR 258/0 35/0	18.00	WD 345/0 9/0	16.15
FR 258/0 36/0	18.00	WD 367/0 1/0	24.00
FR 258/0 37/0	18.00	WD 368/0 1/0	23.00
FR 258/0 38/0	18.00	WD 369/0 1/0	19.00
FR 258/0 39/0	18.00	WD 370/0 1/0	15.00
FR 258/0 4/0	18.00	WD 371/0 1/0	16.05
FR 258/0 40/0	18.00	WD 372/0 1/0	16.05
FR 258/0 41/0	18.00	WD 417/1 2/0	19.00
FR 258/0 42/0	18.00	WD 417/1 4/0	15.00
FR 258/0 43/0	18.00	WD 417/2 2/0	16.00
FR 258/0 44/0	18.00	WD 417/2 3/0	15.00
FR 258/0 45/0	18.00	WD 417/3 1/0	17.00
FR 258/0 46/0	18.00	WD 417/3 3/0	20.00
FR 258/0 47/0	18.00	WD 417/3 4/0	15.00
FR 258/0 48/0	18.00	WD 417/3 5/0	15.00
FR 258/0 49/0	18.00	WD 417/3 6/0	15.00
FR 258/0 5/0	18.00	WD 417/3 8/0	21.00
FR 258/0 50/0	18.00	WD 417/3 9/0	15.00
FR 258/0 51/0	18.00	WD 417/4 1/0	16.00
FR 258/0 52/0	18.00	WD 417/4 3/0	20.00
FR 258/0 53/0	18.00	WD 417/4 4/0	15.00
FR 258/0 54/0	18.00	WD 417/4 5/0	18.00
FR 258/0 55/0	18.00	WD 417/4 6/0	15.00
FR 258/0 56/0	18.00	WD 417/4 7/0	15.00
FR 258/0 57/0	18.00	WD 417/5 1/0	15.00
FR 258/0 58/0	18.00	WD 417/5 2/0	15.00
FR 258/0 59/0	18.00	WD 460/19 54/0	15.00
FR 258/0 6/0	18.00	WD 460/19 55/0	15.00
FR 258/0 60/0	18.00	WD 460/19 58/0	15.00
FR 258/0 61/0	18.00	WD 460/19 60/0	16.00

<u>Plot id</u>	<u>Initial Age</u>
WD 460/19 61/0	16.00
WD 460/19 62/0	16.00
WD 460/19 64/0	18.00
WD 460/19 65/0	18.25
WD 460/19 66/0	20.15
WD 460/19 67/0	16.15
WD 460/19 68/0	21.15
WD 460/19 69/0	16.25
WD 460/19 70/0	16.25
WD 460/19 71/0	20.15
WD 460/19 72/0	21.15
WD 460/19 73/0	21.15
WD 460/19 98/0	15.00
WD 460/26 1/0	25.15
WD 460/26 10/0	26.00
WD 460/26 11/0	26.00
WD 460/26 12/0	24.15
WD 460/26 13/0	26.00
WD 460/26 14/0	26.00
WD 460/26 2/0	25.15
WD 460/26 3/0	25.15
WD 460/26 5/0	27.00
WD 460/26 6/0	27.00
WD 460/26 7/0	27.00
WD 460/26 9/0	25.15
WD 460/30 87/0	18.15
WD 460/30 89/0	18.15
WD 460/30 90/0	18.15
WD 460/30 92/0	18.15
WD 460/31 1/0	22.25
WD 460/31 118/0	15.25
WD 460/31 119/0	16.25
WD 460/31 14/0	18.00
WD 460/31 15/0	18.00
WD 460/31 2/0	22.25
WD 460/31 21/0	15.00
WD 460/31 23/0	19.00
WD 460/31 24/0	19.00
WD 460/31 25/0	18.00
WD 460/31 26/0	24.00
WD 460/31 30/0	16.15
WD 460/31 37/0	16.15
WD 460/31 38/0	16.15
WD 460/31 5/0	29.00
WD 460/32 1/0	20.25
WD 460/32 10/0	15.05
WD 460/32 108/0	21.25
WD 460/32 109/0	21.25
WD 460/32 11/0	21.25
WD 460/32 110/0	17.25
WD 460/32 111/0	18.25
WD 460/32 112/0	18.25
WD 460/32 2/0	19.25
WD 460/32 3/0	23.00
WD 460/32 4/0	19.25
WD 460/32 5/0	16.05
WD 460/32 6/0	23.25
WD 460/32 7/0	16.05
WD 460/32 8/0	15.05
WD 460/33 4/0	27.15
WD 460/33 5/0	26.15

<u>Plot id</u>	<u>Initial Age</u>
WD 460/33 7/0	24.15
WD 460/33 8/0	25.00
WD 460/43 10/0	25.15
WD 460/43 106/0	19.05
WD 460/43 11/0	25.15
WD 460/43 12/0	24.15
WD 460/43 2/0	27.15
WD 460/43 47/0	21.05
WD 460/43 48/0	21.05
WD 460/43 49/0	21.05
WD 460/43 50/0	18.05
WD 460/43 51/0	18.05
WD 460/51 7/0	27.05

APPENDIX D:

Projection statistics for the selected Regional plots in Figures 5-12

Figure Number & Location ^a	Simulation End-Age (yrs)	Projection Period (yrs)	Stocking (sph)		Dbh _q (cm)		Dbh _i Variance Ratio ^b
			Actual	Predicted	Actual	Predicted	
CNI							
5 t-l	21.05	6.00	1779	1687	22.4	22.7	1.163
5 t-r	28.05	13.00	1087	972	29.7	30.4	1.048
5 b-l	21.05	6.00	1137	1071	27.3	27.9	0.951
5 b-r	28.05	13.00	923	772	32.4	34.2	0.782
GDNS							
6 t-l	20.15	4.15	220	215	37.1	37.0	1.156
6 t-r	24.05	8.05	220	209	44.2	43.0	1.091
6 b-l	20.15	4.15	350	341	33.0	33.8	0.955
6 b-r	24.15	8.15	350	328	37.5	39.0	0.776
HBAY							
7 t-l	21.00	5.25	178	185	42.8	43.8	1.097
7 t-r	25.00	9.25	168	182	48.2	50.2	0.989
7 b-l	23.15	6.00	652	619	38.2	38.9	1.069
7 b-r	29.00	11.85	623	565	42.3	43.1	0.905
CLAYS							
8 t-l	24.00	5.85	443	430	33.5	34.6	0.849
8 t-r	31.40	13.25	433	402	40.4	42.1	0.995
8 b-l	26.25	6.70	425	405	40.6	40.9	0.859
8 b-r	32.15	12.60	395	379	46.2	46.9	0.792
SOUTH							
9 t-l	20.00	4.00	2459	2105	19.5	21.0	1.012
9 t-r	25.05	9.05	2173	1491	21.8	25.4	1.015
9 b-l	20.25	3.85	340	339	41.6	38.9	0.719
9 b-r	24.25	7.85	340	335	49.0	45.0	0.603
SANDS							
10 t-l	20.15	5.10	352	336	37.7	38.0	1.030
10 t-r	26.05	11.0	352	315	43.5	44.2	0.940
10 b-l	28.05	4.50	208	200	51.0	50.7	0.897
10 b-r	33.05	9.50	208	192	56.4	56.8	0.772
CANTY							
11 t-l	18.55	3.30	700	725	27.2	27.0	1.060
11 t-r	24.05	8.80	700	725	32.9	32.2	1.020
TWC							
12 t-l	21.15	6.10	413	411	29.0	26.5	0.920
12 t-r	27.00	11.95	388	371	36.6	33.3	0.943
12 b-l	21.15	6.10	277	262	29.8	30.7	0.823
12 b-r	27.00	11.95	254	233	35.8	39.2	0.821

- a t-l = top left (nearest 1.0^b, short pd.) t-r = top right (nearest 1.0^b, long pd.)
b-l = bottom left (nearest xbar^b, short pd.) b-r = bottom right (nearest xbar^b, long pd.)

- b { variance of predicted dbh_i / variance of actual dbh_i }; variance = $\Sigma(x_i - \bar{x})^2 / n$

