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# Estimating the accuracy of site index curves by means of simulation

Skattning av noggrannheten vid höjdbonitering med hjälp av simulering

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#### Abstract

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The aim of this study is to develop a method for estimating the accuracy of site index curves which is independent of the method used for the construction of the curves. Site index curves for Scots pine in Sweden are used to illustrate the method. The investigation deals mainly with prediction errors, i.e. errors caused by actual height development not following the site index curves. Permanent plot data is used to construct a stochastic, autoregressive simulation model, by means of which it is possible to study the variation of site index over age within stands over long periods. A number of simulations are performed and used for estimating prediction errors. Earlier investigations provide figures on sampling and measurement errors which, together with estimated prediction errors, are used to obtain recommendations on the number of plots required within a stand to reach a predefined level of accuracy at site index estimation.

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### **1** Introduction

Site index curves, describing the development of dominant height over age, are widely used for site classification. However, the accuracy of these curves is not always estimated in a reliable way. The reason for this is that the curves are often constructed using methods (e.g. graphical fitting) which do not allow estimation of accuracy by straightforward statistical methods.

The aim of this study is to develop a method for estimating the accuracy of site index curves, which is independent of the method used for constructing the curves. The study was undertaken because there was an urgent need to estimate the accuracy of recently constructed Swedish site index curves (Hägglund, 1972, 1973, 1974). One of these sets of curves (those for Scots pine) is used to illustrate the proposed method.

If unrestricted random sampling of plots within a stand is assumed the accuracy of a site index estimate can, expressed as variance, be written as follows.

$$S_{si}^2 = S_p^2 + (S_m^2 + S_s^2)/n$$

where

 ${\bf S}_{\rm Si}{}^2$  is the total variance of site index within a stand

- $S_p^2$  is the variance caused by "prediction" errors
- $S_m^2$  is the variance caused by measurement errors
- $S_s^2$  is the variance caused by sampling errors n is the number of sampling units within the stand.

Prediction errors might be caused by either bias in the site index curves or by abnormal weather conditions, diseases, vertical variations in soil fertility and other influences which make the actual height development differ from the site index curves.

In a stand  $(S_m^2 + S_s^2)/n$  can be brought near 0 by means of a large number of sampling units, covering different parts of the stand, different operators, different altimeters and so on.  $S_m$  and  $S_s$  have been investigated in some earlier studies, referred to later.

The prediction errors,  $S_p$ , are independent of the number of sampling units within the stand as long as all measurements are made at one occasion. In the investigation carried out here, only prediction errors are studied. Results from the earlier investigations mentioned are used to obtain a more complete picture of the accuracy of site index estimation.

#### 2 Definitions

The definitions used in this study are those used for constructing the Swedish site index curves. These curves describe the development of dominant height over age at breast height. Dominant height is defined as the arithmetic mean height of the 100 largest (by diameter) trees per hectare. The curves are attached to site indices, defined as the dominant height at a total age of 100 years,  $h_{100}$ .

In stand number i at age t years, an "observation" of  $h_{100}$  made without sampling and measurement errors is called  $h_{100}$  (i, t). This observation is obtained by means of the site index curves under investigation. Then, the "true site index"  $h_{100}$  (i), for stand i is defined as

$$h_{100}(i) = 1/(t_2 - t_1) \int_{t_1}^{t_2} h_{100}(i, t) dt$$

where  $t_1$  is a low age, and  $t_2$  is well beyond the normal rotation. In practice, an estimate of  $h_{100}(i)$  is obtained by averaging a sufficiently large number of estimates of  $h_{100}(i, t)$  performed at uniformly distributed ages during the whole rotation. The prediction error is defined as

$$S_p^2(i, t) = E((h_{100}(i, t) - h_{100}(i))^2)$$

where  $h_{100}(i, t)$  is observed without sampling and measurement errors. The symbol E is for expectation. If  $h_{100}$  (i, t) is estimated by  $\hat{h}_{100}(i, t)$ , the sampling and measurement errors are

 $(\mathbf{S}_{m}^{2}(\mathbf{i}, t) + \mathbf{S}_{s}^{2}(\mathbf{i}, t))/n =$ = E(( $\hat{\mathbf{h}}_{100}(\mathbf{i}, t) - \mathbf{h}_{100}(\mathbf{i}, t))^{2}$ )

The definitions are examples and not prerequisites for the method of estimating accuracy. For instance many authors (see e.g. Heger, 1973) define "true site index" as the actual height a stand reaches at a fixed age. This definition may be applied to the method proposed here.

#### **3** Method

A brief description of the method for estimating accuracy follows below. The basic idea is that "true site index" and the variance of site index estimates made at different ages can easily be estimated from permanent plot data where the stand on each plot has been observed during the whole rotation. Observation series of such a length are rare, so shorter observation series are used to simulate series of the desired length. To use this method data from permanent plots consisting of estimates of site index, made in a way that is representative for the normal way of using the site index curves under investigation, must be available. From each plot at least three estimates of site index, made at different ages of the stand, are necessary. The total amount of data must completely cover the range of ages for which estimates of accuracy are of interest. This age range must be so wide that reliable estimates of "true site index" can be obtained from the simulated observation series.

The first step in processing the data is to compute estimated  $h_{100}$  at equidistant ages  $t_1, t_2, \ldots, t_j, \ldots, t_m$  for each plot. The interval between  $t_j$  and  $t_{j+1}$  is of such a length that it is reasonable to assume that the variance about the "true site index" is stable within intervals. The plots or observation series are denoted 1, 2, ..., i, ... k. It is now assumed that  $h_{100}(i, t_j)$  can be described with the following model

 $\beta_1$  and  $\beta_2$  are constants,  $\varepsilon$  the "error"—the stochastic component. The mean of  $\varepsilon$  is 0, the variance is assumed to be constant within age intervals. The correlation between  $\varepsilon(i, t_j)$  and  $\varepsilon(i, t_{j+1})$  where l is an integer is assumed to be 0. The model (1) can be rewritten as follows

$$\begin{split} & h_{100}(i, t_j) = h_{100}(i, t_{j-1}) + (\beta_1(t_j) + \\ & + \beta_2(t_j) - 1) h_{100}(i, t_{j-1}) - \\ & - \beta_2(t_j) (h_{100}(i, t_{j-1}) - h_{100}(i, t_{j-2})) + \epsilon(i, t_j) \end{split}$$

Thus if  $\beta_1$  is 1 and  $\beta_2$  is 0, the process under study is stationary in the mean with independent increments. If  $(\beta_1 + \beta_2)$  is 1 and  $\beta_2$  is not 0, the increments are autocorrelated. If  $(\beta_1 + \beta_2)$  is not 1, the process involves some trend in h<sub>100</sub> over age. Formally (1) is a second-order autoregressive model. See for instance Anderson, 1971.

Assume that  $(\beta_1 + \beta_2)$  is 1. Then the correlation between successive increments is positive if  $\beta_2$  is negative and vice versa.

By means of regression analysis,  $\beta_1$ ,  $\beta_2$  and the standard error of  $\varepsilon$  are estimated as  $b_1$ ,  $b_2$ and the standard error of  $\varepsilon$ . Then (1) is used to simulate successive estimates of  $h_{100}$ . The distribution of  $\varepsilon$  is investigated, and the result of the investigation is used at simulation. By means of the simulation, long "observation series" of  $h_{100}$  are obtained. For each simulated series, "true site index" is estimated as

$$\hat{\mathbf{h}}_{100}(\mathbf{i}) = (1/m) \sum_{j=1}^{m} (\mathbf{h}_{100}(\mathbf{i}, \mathbf{t}_j))$$
 (3)

The accuracy in prediction of  $h_{100}$  at age  $t_j$  can now be estimated as

$$\mathbf{S}_{p}(t_{j}) = \sqrt{(1/k) \sum_{i=1}^{k} (h_{100}(i, t_{j}) - \hat{h}_{100}(i))^{2}} \qquad (4)$$

Formula 4 gives the information wanted in this investigation. Some information about the nature of  $S_p$ —which part is due to "lack of fit" of the site index curves and which is "pure error"—can be obtained from the magnitudes of the constants b. However, in this case, the site index curves are previously tested for "lack of fit" (Hägglund, 1974). The results of these tests were such that the curves are assumed to be unbiased. The magnitude of  $S_p$  is therefore assumed to be entirely dependent on "pure error".

#### 4 Site index curves used in the study

The set of site index curves for Scots pine in Sweden, the accuracy of which is estimated here, is reported in Hägglund, 1974. The curves describe the development of dominant height over age at breast height. To construct the curves stem analysis data from felled trees was used. The trees originate from 213 temporary sample plots. At the time of felling the trees were among the 100 largest per ha. However, this does not guarantee that the felled trees had been dominants during the whole life of the stand. This question was investigated with data from permanent sample plots. The heights of single trees, dominants at some occasion, were followed backwards in time and compared with dominant height. In this way a "rank effect" was detected and quantified with functions. The stem analysis data from single trees was corrected to dominant height development with these functions. A model was fitted to the corrected data set. The model (originating from "Chapman-Richards model"-see e.g. Richards, 1959) can be written as follows.

$$\begin{split} & h(i, t_{j}) - 13 = A(i) \cdot (1 - exp(-c_{1}(i) \cdot t_{j}(i)))^{1/(1 - c_{2}(i))} \\ & c_{1}(i) = b_{10} + b_{11} \cdot A(i)^{b_{12}} \\ & c_{2}(i) = b_{20} + b_{21} \cdot A(i)^{b_{22}} \end{split}$$

h is dominant height in dm for stand i at age  $t_i$ . A,b,c are constants to be estimated. Notice that every stand must have its own, unique A constant. The model (5) was fitted to data according to the least squares principle by means of a three-step process, involving iterative non-linear regression (Gauss-Newton method—see Hartley, 1961) and linear multiple regression. The following estimates of b were obtained

$b_{10} = 1.0002 \ 10^{-4}$	$b_{20} = 6.6074 \ 10^{-2}$
$b_{11} = 9.5953 \ 10^{-6}$	$b_{21} = 4.4189  10^5$
$b_{12} = 1.3755$	$b_{23} = -2.9134$

The function is corrected for logarithmic bias by multiplying with 1.0075.

To relate (5) to  $h_{100}$ , the relationship between total age and age at breast height must be known. When this is the case, (5) can be used to estimate  $h_{100}$ . This can be done for example with graphical site index curves (figure 1) or with a simple step halving algorithm, programmed for computer.

The method for the construction of site index curves, briefly described above, is more thoroughly reported in Hägglund, 1972. It is evident that this method does not permit any straightforward estimating of accuracy. The reasons for this are for example that

- the errors of the observations of age and height from stem analysis are in this case dependent within trees. The reason for this is that the borings were made at points on the stems which were defined as percentages of total height. An error in the measurement of total height will therefore affect all the estimates of heights of boring points.
- the corrections for "rank effects" will probably affect accuracy.
- the fitting of the model (5) to data is complicated. From a statistical viewpoint objections can be made to the method.

One way to estimate the accuracy of the site index curves described is to use the "simulation method".

#### Dominant height, m Övre höjd, m



Figure 1. Site index curves for Scots pine in Sweden. *Höjdutvecklingskurvor för tall i Sverige.* 

#### 5.1 Description of data

The data for the study consists of observation series of total age and dominant height from 203 permanent sample plots. The data was transformed to age at breast height and  $h_{100}$  by means of function (5) and a relationship between total age and age at breast height. These transformed data are illustrated in figure 2. In the figure observation series from plots located in the same stand have been averaged. This averaging of data is only used to make figure 2 clearer. At the further processing no averaging is done.

The permanent plots are 0.1 ha on average. Thus, the dominant height (see "Definitions") is determined from 10 trees. The height measurements are normally performed with the Tirén altimeter. According to Eriksson, 1970, this instrument gives a standard error at single tree measurement of 1.1 % of tree height. This means that the standard error of the dominant height measurement is around 0.35 %. These figures only include random errors. In this investigation other measurement errors cannot be separated from prediction errors.

The plots used are in many respects more homogeneous than average Swedish pine stands. The soil fertility is uniform within plots, the stands are even-aged and of one species, the frequency of diseases is low etc. This might cause differences in site index prediction error between the data set used and "normal" Swedish pine stands. The applicability of the quantitative results from the investigation must therefore be limited to stands with the same properties as the permanent plots.

#### 5.2 Processing of data

The first step at the data processing was to compute  $h_{100}(i, t_i)$  at ages 5, 10, 15, ...., 5 · m years by linear interpolation. The lowest age used here was 15 years, the highest 165 years. According to model (1) a complete "case" in the regression analysis should contain observations from ages  $t_j$  (dependent variable),  $t_{j-1}$  and  $t_{j-2}$ . However, when  $h_{100}(i, 20)$  is the dependent variable, there are no observations of  $h_{100}(i, 10)$ . In this and only in this case the term  $h_{100}(i, t_{j-2})$  is omitted from the model.

Some observation series, which were not long enough to produce a complete case, were omitted from the data set. The total number of cases used in regression analysis is 679. The average number of cases per age is 23, the minimum number 5 and the maximum number 47.

Model (1) was fitted to data for each age separately. The distributions of the residuals from the functions were tested for normality with  $\chi^2$ -tests. The correlations between the residuals  $e(i, t_j)$ ,  $e(i, t_{j-1})$ ,  $e(i, t_{j-2})$  and  $e(i, t_{j-3})$  were studied. The number of observations, the regression coefficients, the standard errors of  $h_{100}(i, t_j)$  and the  $\chi^2$ -tests are reported for each age in Appendix 1.

For all ages the sum of the estimated regression coefficients  $b_1$  and  $b_2$  is close to 1. As mentioned earlier (see "Method") a deviation of  $(b_1+b_2)$  from 1 indicates some trend in

Material till undersökningen. Observationer av  $h_{100}$  från 203 permanenta försöksytor. Observationsserier från näraliggande ytor har sammanslagits till medelförlopp i figuren.

Figure 2. Data for the study. Observations of  $h_{100}$  from 203 permanent sample plots, some of them averaged in this figure.





Figure 3. Standard error of  $e(i, t_j)$ , estimated by regression functions. Medelfelet i  $e(i, t_j)$ , skattat med regressionsanalys.

 $h_{100}$  over age. As reported in Appendix 1  $(b_1 + b_2)$  is somewhat greater than 1 for all intervals up to age 115 years. In a regression performed for the whole material  $(b_1 + b_2)$  is 1.00436, with a standard error of 0.0075. The deviation of  $(b_1 + b_2)$  from 1 is not significant, but a suspicion of a slight trend in  $h_{100}$  over age still remains. The character of the trend is such that  $h_{100}$  increases with age. The reason for this is probably that three plots included in the data (and specially reported in Hägglund, 1974) were extremely overdense at low ages. This resulted in a depressed dominant height growth. After thinning dominant height

growth increased to a level "normal" for the site and age, which resulted in increasing  $h_{100}$ . The trend possibly brought into the data with these plots will, however, be included in the random variation when calculating prediction errors.

For most ages  $b_1$  is positive and  $b_2$  is negative. Hence, the correlation between successive increments is mostly positive. A minor part of this autocorrelation is probably caused by the linear interpolation used at estimating  $h_{100}$  for different ages.

The estimates of the standard errors of  $e(i, t_j)$  over age are illustrated in figure 3.



 $\chi^2$ -tester, några exempel. x = residual/standardavvikelse.

As expected, estimated standard errors decrease over age. Probably because of the low number of observations, the standard error is very unstable for low ages.

The  $\chi^2$ -tests were based on a division of residuals by normalized size (residual/standard deviation) in six classes. Thus, the number of

degrees of freedom is 3. Significant deviations from normal distribution on the 5 %-level are indicated by a  $\chi^2$  greater than 7.81. For only one age was a  $\chi^2$  of that magnitude observed (age 75,  $\chi^2 = 7.88$ ). Some examples of the shape of the distributions are illustrated in figure 4. However the amount of data for the





 $\chi^2$ -test, hela materialet. x = residual/standardavvikelse.

 $\chi^2$ -tests is small. If a  $\chi^2$ -test is performed for the whole material, a significant deviation (5%-level) from the normal distribution is obtained. This deviation is judged as being too small (see figure 5) to be taken into account. At simulation, it is assumed that the residuals e(i, t<sub>j</sub>) are normally distributed.

The correlations between the residuals  $e(i, t_j)$ 

and respectively  $e(i, t_{j-1})$ ,  $e(i, t_{j-2})$  and  $e(i, t_{j-3})$  were investigated in order to check the important assumption of no correlation between residuals. The correlations were respectively 0.02, -0.05 and -0.04, with non-significant t-values of 0.4, -0.6 and -0.3. Thus, there is no reason to suspect correlation between residuals.

#### 6 Simulation of successive site index estimates

The regression functions reported in appendix 1 are used to simulate "observation" series of successive estimates of h<sub>100</sub>. The simulation of a series starts by arbitrary chosing a value of  $h_{100}$  for age 15. Here the values chosen are 14, 16, 18, ..... 28 m. After that, the first regression function is used to calculate a new  $h_{100}$  for age 20. A random deviation  $e(i, t_i)$  is added to the estimate from the regression function. This deviation is calculated as a random number, picked from a normal distribution with mean 0 and a standard deviation in accordance with the standard error calculated for the regression function. Technically, this operation is performed by means of the FORTRAN IV function subroutine RNORM, developed at the Stockholms Datamaskincentral.

The  $h_{100}(i, t_j)$  obtained at ages 15 and 20 are used as independent variables in the second regression function to calculate  $h_{100}(i, t_j)$  at age 25 and so on. In this way observation series with a range from 15 to 165 years are calculated. For an IBM 360/75 computer the CPU-time needed to calculate one such observation series is about 10<sup>-4</sup> minutes. Some simulated series are illustrated in figure 6. In appendix 2, 128 simulated series are reported. There is a tendency of rising  $h_{100}$  with age in these series. The reasons for this have been discussed earlier.





#### 7 Estimating accuracy

From the simulated observation series,  $S_p(t_j)$  were calculated by formula (4)

$$\mathbf{S}_{p}(t_{j}) = \sqrt{(1/k) \sum_{i=1}^{k} (h_{100}(i, t_{j}) - \hat{h}_{100}(i))^{2}} \qquad (4)$$

 $\hat{h}_{100}(i)$  is an estimate of the "true site index" for observation series no i.  $S_p^2(t_i)$  might be divided in two parts, corresponding to "lack of fit" due to the site index curves and "pure error". To do this, the shape of some trend f(t) describing the bias of the site index curves must be known. In principle the bcoefficients can also be used to estimate bias. However, the data set must be large, if reliable estimates of bias are to be calculated from separate five-year intervals. Otherwise there is an evident risk of confusing bias of the site index curves with errors occurring from the fact that the data set is a sample. For example, in the present investigation the data set includes too large a proportion of stands with too slow early dominant height development. To ensure that no depreciation of S<sub>p</sub> is done, these stands are included in the data set. Their variance must however be treated as random.

By performing a number of simulations,  $S_p(t_j)$  is estimated for each age interval. The simulations start at age 15 and end at age 165. At the start,  $h_{100}$  is 14, 16, 18, ..., 28 m. The whole period from age 15 to age 165 is used for calculating "true  $h_{100}$ " by averaging the 31 values per series corresponding to each 5-year interval. It seems reasonable to found the calculation of "true  $h_{100}$ " on as long a series as possible, since  $h_{100}$  is more stable at higher ages. In the present case, the calculation of "true  $h_{100}$ " is based to some extent on extrapolated parts of the site index curves. According to figure 2, this extrapolation does not introduce any bias in the calculations.

At the calculation of  $S_p(t_j)$ , all observation series have the same weight. An alternative way of weighting is to give every series i a weight proportional to the frequency of  $h_{100}(i)$ in the data set. Weighting can also be based on the frequency of  $h_{100}(i)$  in the population where the results shall be applied.

Figure 7 shows an example of the behaviour of  $S_p(t_j)$  for three ages (25, 50 and 100 years) when the number of simulations increases. In the example  $S_p(t_j)$  seems to stabilize after 256 simulations. Here the values of  $S_p(t_j)$ obtained after 8192 simulations are accepted as final. Figure 7 also shows the variation between four sets of 8192 simulations. The difference between the largest and the smallest value of  $S_p(t_j)$  is 3 cm at age 25, 1.5 cm at age 50 and 0.5 cm at age 100.

As mentioned earlier, some measurement errors of a magnitude of 0.35 % of dominant height are included in the simulated series. These errors are, expressed as  $h_{100}$ , around 0.6 dm. The values of  $S_p(t_i)$  obtained by simulation are corrected for these errors. Such corrected  $S_p(t_i)$  are illustrated over age in figure 8 and table 1.

The curve in figure 8 has some irregularities. An attempt to smooth these out with a sixorder polynomial did not give acceptable results. Therefore, the "raw" values are reported as the main results of the investigations. As can be seen, the prediction error  $S_p$  is at the level 20 dm at age 15–20 years. Then the error drops quickly with increasing age. A minimum of around 7 dm is reached at age 80-85. After that, the error increases slowly with increasing age and is around 11 dm at age 155–165.

The relationship between  $S_p$  and age applies to average conditions in the data. No attempt was made to differentiate  $S_p$  for different  $h_{100}$ . Evidently the shape of the relationship depends to some extent on the age interval on which the calculation of "true  $h_{100}$ " is based. The minimum of  $S_p$  will always occur close to the middle of that interval.

As mentioned earlier, the total error of a



Figure 7. Prediction error  $S_p(t_i)$  of  $h_{100}$  over number of simulated observation series.  $S_p(t_i)$  is not corrected for measurement errors. The number of simulations is expressed as  $2^x$ . When for example x = 4, the number of simulations is 16, when x is 13 the number of simulations is 8192. For x = 13, four sets of simulations are illustrated.

Prognosfelet  $S_p(t_j)$  i  $h_{100}$  över antalet simuleringar.  $S_p(t_j)$  är inte korrigerat för mätfel. Antalet simuleringar uttrycks som  $2^x$ . När t ex x är 4 är antalet simuleringar 16. När x är 13 är antalet simuleringar 8192. För x = 13 visas resultaten av fyra uppsättningar simuleringar.



Figure 8. Prediction error of  $h_{100}$  over age at breast height. The figures are based on 8192 simulations and corrected for measurement errors.

Prognosfel i h<sub>100</sub> över brösthöjdsålder. Siffrorna bygger på 8192 simuleringar och är korrigerade för mätfel.

site index estimate for a stand is obtained from the formula

$$S_{si}^2 = S_p^2 + (S_m^2 + S_s^2)/n \tag{6}$$

where p, m and s are for "prediction", "measurement" and "sampling" respectively. The number of sampling units within the stand is called n. Fries, 1974 reported some results concerning the magnitude of  $S_m^2 + S_s^2$ . Some of these are found in table 2 below.

The necessary number of plots within a stand to reach some predefined level of accuracy can be calculated by means of these results and figure 8. Let this level be a standard error  $S_{si}$  of magnitude E dm. Then the number of plots required ( $n_E$ ) is obtained from formula (7) below.

$$n_{\rm E} = (S_{\rm m}^2 + S_{\rm s}^2) / (E^2 - S_{\rm p}^2)$$
(7)

This formula is exemplified in figure 9. The example concerns a stand with "medium" site variation within stand. The plot size chosen is  $254 \text{ m}^2$  and some levels of accuracy, E, from 10 to 20 dm are investigated. As can be seen the number of plots required decreases dramatically when E increases from 10 to 14 dm. It would here seem reasonable to accept an E of 14 dm in most cases. In stands younger than 45 years, a higher E must be accepted.

Tabulated values of  $n_E$  for various combinations of E, plot size and site variation within stand are found in Appendix 3. These figures might be useful at practical site index estimation. However, their application is limited to even-aged, pure stands of Scots pine in Sweden. Table 1. Prediction error of  $h_{100}$  at various ages at breast height. The figures are based on 8192 simulations and corrected for measurement errors.

Prognosfel i  $h_{100}$  vid olika brösthöjdsåldrar. Siffrorna bygger på 8 192 simuleringar och är korrigerade för mätfel.

Age at breast height <i>Bröst- höjds- ålder</i>	Prediction error of $h_{100}$ , dm Prognosfel i $h_{100}$ , dm	Age at breast height Bröst- höjds- ålder	Prediction error of $h_{100}$ , dm <i>Prognosfel</i> <i>i</i> $h_{100}$ , dm
15	20.1	95	7.7
20	18.8	100	8.1
25	19.4	105	8.4
30	18.7	110	8.4
35	16.8	115	8.5
40	15.4	120	8.6
45	12.0	125	9.3
50	10.8	130	9.5
55	9.0	135	9.9
60	8.0	140	9.6
65	8.3	145	9.6
70	7.7	150	10.2
75	7.4	155	11.5
80	6.9	160	11.3
85	6.9	165	11.1
90	7.3		

Table 2. Some examples of the magnitude of  $S_m^2 + S_s^2$  according to Fries, 1974.

Några exempel på storleken av  $S_m^2 + S_s^2$  enligt Fries, 1974.

Plot size	Number of domi-	$S_m^2 + S_s^2$	², dm	
m <sup>4</sup> Ytstorlek m <sup>2</sup>	nant trees per plot used for site estimation	Site vari Bonitets	ation within sta variation inom b	nd estånd
	träd på provytan	low liten	medium <i>måttlig</i>	high <i>stor</i>
1 000	10	196	400	676
254	2	324	576	1 024
154	1	400	729	1 225



Figure 9. Number of plots within stand required to reach a predefined standard error. Behövligt antal provytor inom bestånd för att nå ett förutbestämt medelfel.

Many questions can be discussed in connection with this study. Here however, we limit ourselves to four.

- the simulation model
- division of data by other variables than age
- when should the simulation method be used?
- other applications of simulation in site estimation

For the convenience of the reader, the model (1) is repeated here

$$\begin{aligned} h_{100}(i, t_j) &= \beta_1(t_j) h_{100}(i, t_{j-1}) + \\ &+ \beta_2(t_j) h_{100}(i, t_{j-2}) + \varepsilon(i, t_j) \end{aligned}$$

A more sophisticated model might produce a better description of data. In some cases it might be necessary to introduce a third-order autoregressive term,  $\beta_3 h_{100}(i, t_{j-3})$ , in the model. However, more data is needed because the observation series must be one age interval (in this case five years) longer than when using (1). The model can be made more flexible by bringing terms of higher order than 1 into the model, for instance  $[h_{100}(i, t_{j-1})]^2$ .

It is obvious that no constant term should be brought into the model. If such a constant is not 0, the functions obtained at regression analysis cannot have the following desirable feature for more than two values of  $h_{100}$ : "If  $h_{100}(i, t_{j-1}) = h_{100}(i, t_{j-2})$  then the conditional expectation of  $h_{100}(i, t_j) = h_{100}(i, t_{j-1})$ ." An attempt to perform simulation with regression functions involving constant terms has been made. As expected, the simulated observation series got a preposterous shape when approaching the limits of the data set.

In this investigation data has been divided by age. The aim of this division was to form groups (age intervals) with as constant an interior variance  $(S_p^2)$  as possible. An alternative way of division is to use the dominant height. However, when prediction errors are searched for, it seems more reasonable to use age than height for division. The prediction error is hardly the same in a 12 m stand of age 18 ( $h_{100} = 30$  m) as in a 12 m stand of age 81 ( $h_{100} = 12$  m). This holds for prediction errors only. Measurement errors are probably more constant at constant height than at constant age.

Besides the division by age, also some other divisions, for instance by  $h_{100}$ , might be done. Such a division should be made if there are reasons to believe that  $S_p$  varies with  $h_{100}$  and the data set is large enough. This division by  $h_{100}$  is probably especially valuable if the simulation model will be used for prediction purposes.

The simulation technique used for estimating prediction errors should of course only be used when more straightforward methods are not applicable. For instance when site index curves are constructed as functions

site index = f(height, age)

they can often be investigated for prediction errors by means of standard error of estimate, obtained at regression analysis. The situation is not quite as simple (Heger, 1971) when site index curves are constructed as functions

height = f(site index, age)

For a discussion of these questions, see Curtis, DeMars & Herman, 1974. Whatever model used, one thing must be pointed out. When calculating  $S_p$  from the data used at the construction of the site index curves, there is a risk of underestimating  $S_p$ . This is because "overfitting" often occurs. See Gardner, 1972 for a discussion of this question. In the investigation carried out here, the data used for the calculation of  $S_p$  is not the same as that used in the construction of the site index curves.

An alternative way of calculating the predic-

tion errors is to omit the simulation and use a more traditional statistical approach by calculating expectations for  $S_p$ . This way of solving the present problem seems possible if the regression coefficients and standard errors in Appendix 1 are known. However, the simulation method is more illustrative and much simpler.

The simulation technique described here has been used for estimating prediction errors. It might also be useful for some other purposes. Studying effects of "sampling in time" is closely related to estimating prediction errors. This means using observations from more than one occasion to estimate  $h_{100}$ . In practice this can be done by measuring of whorls. The most proper weighting of such observations in order to form an estimate of site index with maximal accuracy can be studied by simulated observation series. Simulation might also be used to develop aids for prediction of  $h_{100}$ . Assume that in some data set a trend of site index over age, which can be related to some variable x, exists. For simplicity, let us say that x is altitude. It is then possible to include a term  $b_3x$  in model (1). The magnitude of  $b_3$  for different ages can be used to develop different altitudes.

#### 9 Sammanfattning

Syftet med den här genomförda undersökningen är att utveckla en metod för att skatta noggrannheten vid höjdbonitering med höjdutvecklingskurvor, som är oberoende av det sätt på vilket kurvorna konstruerats. Metoden illustreras med hjälp av höjdutvecklingskurvor för tall i Sverige (Hägglund, 1974).

Det totala felet i en skattning av höjdboniteten för ett bestånd kan skrivas som

$$S_{si}^2 = S_p^2 + (S_m^2 + S_s^2)/n$$

där  $S_p$  är prognosfelet,  $S_m$  mätfelet och  $S_s$ samplingfelet. Antalet provytor inom bestånd betecknas n. Med prognosfel menas det fel som uppstår på grund av att beståndets faktiska höjdutveckling mera sällan exakt följer höjdutvecklingskurvorna. Detta kan bero antingen på att kurvorna ej är förväntningsriktiga eller på onormal väderlek, skador, vertikala bördighetsvariationer i markprofilen m fl faktorer. Undersökningen inriktas främst på skattning av prognosfelen, eftersom mätoch samplingfelen studerats i tidigare undersökningar (Eriksson, 1970, Fries, 1974).

Materialet till undersökningen är hämtat från 203 permanenta försöksytor och består av vid olika åldrar upprepade observationer av brösthöjdsålder och övre höjd. Med hjälp av höjdutvecklingskurvorna beräknades  $h_{100}$ , övre höjden vid 100 års total ålder, för varje yta och brösthöjdsålder. Erhållna  $h_{100}$  omräknades med lineär interpolering till att avse vart femte år under den period respektive yta observerats. Till de sålunda erhållna "observationsserierna" av successiva skattningar av  $h_{100}$  anpassades en andra ordningens autoregressiv modell med följande utseende

$$\begin{split} h_{100}(i, t_j) &= \beta_1(t_j) h_{100}(i, t_{j-1}) + \\ &+ \beta_2(t_j) h_{100}(i, t_{j-2}) + \epsilon(i, t_j) \end{split}$$

Beteckningen  $t_j$  står för brösthöjdsålder medan index i avser provyta.  $\beta_1$  och  $\beta_2$  är konstanter medan  $\varepsilon$  är "felet" — den stokastiska komponenten. Modellen anpassades till materialet med regressionsanalys på så sätt att separata funktioner utarbetades för varje ålder t<sub>j</sub>. Härigenom erhölls ett antal funktioner där den beroende variabeln h<sub>100</sub>(i, t<sub>j</sub>) svarar mot åldrarna 15, 20, 25, ....., 165 år. Funktionerna användes för att simulera observationsserier som sträcker sig från 15 till 165 års brösthöjdsålder. Härvid antogs att  $\varepsilon(i, t_j)$  är oberoende av  $\varepsilon(i, t_{j+1})$  där 1 är ett heltal. Vidare antogs att  $\varepsilon(i, t_j)$  har medeltalet 0, är normalfördelat och har en konstant varians inom femårsintervall. Flertalet av dessa antaganden kontrollerades i undersökningen.

För varje simulerad observationsserie skattades det "sanna" värdet på  $h_{100}$  som det aritmetiska medeltalet av de 31 värden på  $h_{100}$ som erhölls vid simuleringen. Prognosfelet beräknades för varje ålder som

$$S_{p}(t_{j}) = \sqrt{(1/k)\sum_{i=1}^{k} (h_{100}(i, t_{j}) - \hat{h}_{100}(i))^{2}}$$

där k är antalet simuleringar och  $\hat{h}_{100}(i)$  skattningen av det "sanna"  $h_{100}$  för den i:te simulerade observationsserien. De slutliga beräkningarna av S<sub>p</sub> grundades på 8 192 simulerade observationsserier. Resultatet av beräkningarna är att prognosfelet är av storleksordningen 2 m vid brösthöjdsåldern 15–20 år. Därefter sjunker felet snabbt med stigande ålder och når ett minimum på ca 7 dm vid 80–85 år. När åldern ökas ytterligare stiger felet långsamt och är ca 11 dm vid 155–165 år.

Undersökningen avslutas med beräkningar avseende det antal provytor som krävs för att skatta  $h_{100}$  för ett bestånd med viss förutbestämd noggrannhet. Härvid användes dels uppgifter rörande storleken av mät- och samplingfel hämtade från Fries, 1974, dels de här gjorda skattningarna av prognosfelet.

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Stockholm, April 1975

Björn Hägglund

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# Appendix 1

Some information about the regression functions

$$\begin{split} h_{100}(i,\,t_j) &= b_1(t_j)\,h_{100}(i,\,t_{j-1}) + b_2(t_j)\,h_{100}(i,\,t_{j-2}) + e(i,\,t_j) \\ h_{100}\text{: site index, dm} \\ i\text{: observation series no} \\ t_j\text{: age at breast height} \\ b_1,\,b_2\text{: estimated coefficients} \\ e\text{: estimated error} \end{split}$$

Dependent variable	No of	Regression	coefficients		Standard error of	Residuals deviation
Dependent variable h <sub>100</sub> at age 20 25 30 35 40 45 50 55 60 65 70 75 80 85 90 95 100 105 110 115 120 125 130 135 140 145 155 160 165	tions	bi	b <sub>2</sub>	$b_1 + b_2$	estimate, dm	from normal distribution $\chi^2$
20	5	1.00589	<u> </u>	1.00589	6.7	0.90
25	5	0.94013	0.07116	1.01129	4.8	1.17
30	7	-0.48922	1.49818	1.00896	4.3	2.24
35	23	0.77101	0.24554	1.01655	7.5	1.49
40	25	1.04847	-0.04476	1.00371	5.0	1.83
45	24	1.44649	-0.44001	1.00648	8.1	3.29
50	25	1.29390	-0.29095	1.00295	4.6	6.57
55	30	1.24068	-0.23872	1.00196	4.8	6.45
60	36	1.47007	-0.46845	1.00112	3.7	4.92
65	36	1.57425	-0.56612	1.00813	2.9	2.59
70	39	1.04606	-0.04456	1.00150	3.4	6.14
75	41	1.33604	-0.32697	1.00907	3.4	7.88*
80	38	1.25789	-0.25065	1.00724	3.4	1.40
85	43	1.41382	-0.41154	1.00228	2.8	2.63
90	47	1.58841	-0.58718	1.00123	2.8	2.50
95	44	1.22279	-0.21754	1.00525	2.4	2.92
100	41	1.31390	-0.31076	1.00314	2.4	1.89
105	28	1.13941	-0.13709	1.00232	2.0	3.45
110	31	0.81972	0.18197	1.00169	2.2	4.18
115	24	1.29629	-0.29914	0.99715	2.6	1.34
120	20	1.10471	-0.10166	1.00305	2.0	3.00
125	11	1.55710	-0.55152	1.00558	2.4	5.46
130	9	1.05536	-0.05616	0.99920	2.5	0.98
135	6	1.33444	-0.33349	1.00095	2.3	1.59
140	7	0.61485	0.38056	0.99541	2.1	0.56
145	8	1.26072	-0.26331	0.99741	1.7	3.56
150	10	1.23046	- 0.22642	1.00404	3.4	2.21
155	7	1.75400	-0.75305	1.00095	2.5	0.56
160	5	0.94351	0.05001	0.99352	1.4	5.87
165	4	0.87931	0.11572	0.99503	0.6	4.45
All	679	1.01140	-0.00704	1.00436	3.9	11.1*

# Appendix 2

# Simulated series of successive site index estimates 128 simulated series of successive $h_{100}$

Age	at br	east l	neight	:																										
15	20	25	30	35	40	45	50	55	60	65	70	75	80	85	90	95	100	105	110	115	120	125	130	135	140	145	150	155	160	165
h <sub>100</sub>	in dn	n																												
140	139	139	141	161	160	158	163	168	172	176	178	180	185	187	187	186	188	189	193	193	195	198	197	194	192	192	194	194	193	192
160	158	156	158	157	161	164	161	156	153	147	147	154	158	160	161	163	161	160	159	160	160	159	156	158	156	153	151	150	147	146
180	178	178	176	185	183	179	183	179	181	189	186	187	182	181	186	187	187	186	187	183	183	187	184	184	188	187	184	182	179	178
200	197	205	197	179	177	181	180	184	190	192	189	189	192	190	197	202	198	195	194	189	188	186	184	181	181	180	176	173	170	169
220	217	220	216	213	202	177	177	180	185	193	195	193	203	207	210	213	220	221	223	221	221	224	224	223	223	220	217	216	214	213
240	239	238	236	250	255	272	281	281	281	286	291	296	305	311	317	322	324	328	328	326	331	335	336	339	337	337	343	344	343	341
260	266	272	262	270	257	247	250	250	251	254	255	266	267	269	266	264	264	268	269	271	273	277	274	275	277	276	277	280	280	278
280	279	293	275	274	272	273	272	265	268	268	266	266	275	277	278	276	280	281	285	284	281	284	286	288	285	285	283	282	279	279
140	130	143	116	127	130	147	151	149	152	155	155	157	166	172	180	184	182	181	176	174	176	178	180	186	181	181	181	181	179	178
160	159	156	161	167	168	163	161	151	150	155	164	167	166	163	162	165	165	164	165	164	166	165	165	168	165	163	160	157	154	153
180	187	188	190	188	185	182	179	190	192	199	193	187	183	183	188	194	195	196	194	194	192	193	194	191	190	191	187	186	183	183
200	203	198	211	211	216	224	220	220	217	219	220	215	216	216	221	223	223	224	229	231	237	239	241	240	239	238	236	239	238	236
220	213	216	206	212	214	205	205	205	205	210	208	206	205	211	215	220	222	224	222	220	220	218	218	217	217	216	214	212	213	211
240	245	249	244	239	239	243	240	237	238	241	237	237	236	233	239	241	247	247	246	245	244	245	249	252	249	250	255	262	259	258
260	265	269	266	273	284	281	279	272	272	272	275	277	279	275	270	270	270	270	268	270	270	273	272	268	263	262	268	272	271	270
280	279	285	283	280	283	284	282	279	276	276	275	273	273	273	281	285	285	287	284	284	283	287	289	289	286	286	290	291	290	288
140	144	144	153	157	160	150	144	140	134	133	133	137	136	140	138	137	138	140	141	142	140	137	135	140	139	139	139	140	139	139
160	166	164	171	177	180	182	185	190	193	195	192	192	195	197	198	198	197	194	190	190	187	191	190	192	191	190	191	195	193	192
180	168	162	176	180	190	196	198	187	180	179	182	181	183	182	181	182	188	193	194	191	189	189	186	186	180	178	178	178	174	174
200	200	196	206	212	225	235	240	234	232	231	240	239	240	240	237	242	246	247	245	246	251	255	253	254	251	250	249	249	246	246
220	225	220	225	227	222	227	228	234	239	240	243	246	248	249	256	261	259	258	258	256	259	264	264	265	262	259	255	251	248	248
240	242	248	240	243	238	228	219	216	216	216	214	217	220	221	221	221	218	218	215	217	219	218	216	213	214	214	214	216	214	213
260	262	266	264	279	286	302	311	317	316	316	316	325	329	330	331	339	349	349	350	348	349	351	349	352	352	352	353	355	354	352
280	277	281	278	280	278	267	264	274	275	277	274	274	271	273	274	278	277	278	275	276	277	277	273	271	271	265	269	269	269	268
140	154	161	150	155	154	157	156	149	141	140	144	140	141	138	141	143	142	140	140	138	140	141	139	138	136	137	140	143	141	140
160	171	177	174	183	192	199	199	199	199	202	202	203	206	210	210	208	210	210	209	206	205	205	204	201	203	201	204	207	205	204
180	189	199	198	212	202	213	224	223	220	216	219	222	219	224	227	227	227	230	229	228	228	229	231	230	225	225	227	232	231	230
200	202	215	206	213	207	213	224	219	223	232	232	237	243	245	248	253	251	254	250	247	249	249	249	247	245	240	241	246	245	243
220	220	218	223	238	237	236	231	236	232	236	238	243	247	249	256	257	257	257	256	256	257	257	260	260	263	263	265	264	264	263
240	244	250	246	237	242	240	245	238	235	234	238	239	238	237	233	235	240	239	240	236	237	237	238	239	235	235	235	237	234	234
260	266	268	262	271	278	261	244	238	234	231	232	238	235	238	240	240	240	239	237	237	235	235	233	232	233	230	224	221	221	220
280	280	291	268	273	266	260	256	262	264	270	266	273	271	272	272	269	269	272	269	267	268	266	263	264	261	258	259	262	261	260

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140 133 137 131 125 129 145 153 159 160 160 159 161 161 156 153 151 153 156 154 149 150 151 152 153 154 153 156 151 148 148 160 160 172 155 160 170 173 178 177 179 175 176 177 182 180 181 182 186 189 187 186 187 189 191 195 191 189 190 188 186 186 180 174 173 176 190 189 192 186 186 183 178 178 180 184 185 185 185 188 181 193 195 194 193 192 188 189 187 185 184 183 182 200 209 218 200 203 203 196 194 195 199 203 208 205 208 208 217 221 225 223 220 223 228 232 230 231 232 231 236 240 238 238 220 216 224 214 220 228 230 222 220 222 220 222 225 221 223 228 227 228 230 234 234 233 235 237 238 240 241 240 238 234 235 237 240 237 237 240 243 253 256 257 264 268 272 274 278 285 290 292 288 291 292 293 288 290 293 290 292 290 293 297 297 298 296 260 260 258 261 269 272 274 270 267 266 270 266 275 277 276 276 276 276 272 270 275 277 279 284 281 282 280 278 282 285 283 282 280 280 284 271 280 278 274 275 272 273 271 274 274 273 273 274 273 273 274 273 273 274 276 277 276 279 278 279 280 279 281 279 279 279 279 140 146 149 142 134 145 153 158 148 146 144 146 151 149 146 142 143 141 142 142 141 141 141 142 145 142 141 142 144 143 143 143 160 153 153 155 155 160 167 180 186 187 188 187 192 193 197 205 204 206 208 209 206 205 205 199 198 196 195 197 204 202 201 180 192 193 191 195 194 187 186 189 196 198 197 199 194 194 194 199 199 202 199 197 199 199 199 197 199 197 192 187 187 185 200 199 198 209 204 199 202 202 200 199 200 204 209 209 206 205 206 208 209 208 207 211 217 214 213 209 207 210 216 216 215 215 220 220 214 224 241 237 239 242 239 241 241 241 240 240 237 231 235 237 235 236 235 237 240 237 234 235 234 237 235 235 235 235 235 240 240 248 230 248 247 237 234 235 234 243 240 246 247 250 252 256 259 260 263 262 262 262 260 261 256 255 254 253 250 250 250 260 260 272 246 252 254 256 258 251 244 239 235 233 230 226 225 226 225 227 228 222 220 220 219 217 216 218 220 220 216 215 280 277 284 274 275 281 272 269 273 273 278 279 280 281 283 289 291 292 292 295 298 298 297 297 297 294 293 298 300 298 297 140 141 138 147 142 144 147 150 148 153 158 158 157 161 163 172 178 178 177 176 174 176 179 181 186 182 184 187 191 187 186 160 170 171 169 177 176 181 181 176 168 165 165 169 174 176 177 181 180 182 182 186 188 185 184 187 185 185 185 185 179 181 179 180 166 171 169 172 169 174 179 178 174 174 168 171 170 165 158 156 154 156 153 152 152 153 152 150 147 143 139 139 135 135 200 206 200 205 190 189 186 185 184 188 192 194 191 197 203 207 210 213 214 210 206 208 212 209 207 204 203 211 221 219 218 220 225 226 219 229 228 233 243 236 236 236 241 244 254 257 256 262 262 262 262 262 261 260 260 262 258 256 257 258 256 255 255 240 240 244 241 237 233 235 235 242 248 253 254 260 260 259 256 261 265 269 271 271 270 270 266 267 267 264 268 274 272 272 260 270 276 272 262 265 279 287 289 289 293 290 288 290 289 288 289 291 291 293 288 290 295 295 297 294 291 294 297 294 294 280 270 276 274 281 290 297 290 284 279 279 280 275 276 273 273 276 272 274 278 278 278 278 279 277 280 279 276 275 275 272 271 140 143 146 148 145 147 125 120 115 118 125 123 126 129 132 138 140 138 141 140 136 134 133 133 135 132 131 133 138 133 134 160 176 171 182 176 183 192 200 201 203 204 200 200 198 199 193 196 197 194 194 192 192 197 200 200 199 196 196 196 196 196 195 180 185 191 183 196 203 201 197 199 198 202 212 217 217 214 213 211 210 212 212 209 208 209 209 208 205 204 199 200 198 200 203 209 208 209 206 197 194 190 192 201 201 205 208 207 204 204 208 210 209 205 204 203 204 207 209 212 217 220 217 217 220 217 226 211 216 217 231 233 241 246 248 251 252 255 259 266 269 269 271 275 274 273 275 272 277 274 275 279 279 278 276 278 276 240 237 240 239 233 239 249 251 247 245 243 243 246 247 247 247 245 245 245 247 247 246 245 245 243 242 241 243 243 242 243 242 243 241 260 257 267 259 268 274 264 266 268 272 276 276 278 277 281 281 284 285 286 289 290 293 295 290 293 290 287 288 284 281 280 280 267 274 270 285 290 285 286 295 293 291 292 292 298 302 306 307 305 306 307 307 309 313 314 316 312 308 314 324 321 320 Age at breast height

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			-																											
15	20	25	30	35	40	45	50	55	60	65	70	75	80	85	90	95	100	105	110	115	120	125	130	135	140	145	150	155	160	165
$h_{10}$	n in d	m																												

140 145 153 140 139 136 128 122 114 117 122 124 125 126 126 126 124 122 122 118 118 119 120 118 114 115 116 115 114 114 113 160 159 162 155 148 146 145 151 151 152 157 163 166 164 164 164 166 166 166 166 168 170 170 169 168 169 167 168 168 171 169 168 180 174 171 173 172 173 172 179 186 189 193 187 182 186 195 202 203 206 209 208 205 205 211 211 211 209 207 209 207 206 205 200 209 205 208 218 222 238 253 259 262 270 269 276 286 286 287 289 289 288 289 287 287 288 289 288 289 288 286 284 285 288 286 286 220 236 241 230 230 223 228 232 239 244 254 250 247 250 251 252 249 251 251 253 251 249 247 245 244 242 242 247 251 248 246 240 235 226 245 242 233 226 218 213 211 214 209 207 207 210 213 216 211 214 210 211 210 206 209 212 208 205 205 210 208 207 260 274 281 273 283 283 280 283 290 298 304 305 308 311 314 311 313 315 316 312 313 312 310 312 314 313 311 307 306 305 303 280 286 292 284 285 284 303 302 301 304 309 308 302 299 296 298 301 304 305 304 301 300 303 300 298 302 300 299 302 299 298 140 141 146 143 146 156 163 158 155 154 152 155 154 159 162 171 172 174 174 175 175 176 181 186 187 188 186 185 186 185 186 185 186 160 157 155 151 158 161 164 159 156 152 152 156 156 151 150 152 155 155 153 154 156 157 156 154 158 155 152 153 158 157 156 180 193 195 194 205 204 203 201 208 210 210 213 216 222 226 224 222 221 221 219 216 217 219 217 217 217 217 219 218 220 217 216 200 211 221 211 207 210 217 214 220 228 236 237 239 235 236 233 234 235 234 235 237 234 238 240 247 245 243 242 240 240 238 218 214 208 209 206 208 210 217 217 218 221 222 222 225 226 230 232 229 226 223 221 220 215 219 213 225 223 222 222 220 220 240 250 249 254 247 250 260 269 268 265 265 267 269 270 272 276 276 279 281 282 283 283 282 280 277 282 282 289 295 294 293 260 254 258 261 272 272 281 290 287 283 276 273 278 283 284 289 290 290 286 283 281 282 281 282 279 277 276 275 273 273 273 292 287 285 281 276 274 264 266 272 271 270 268 269 268 269 271 272 272 273 275 271 269 271 268 268 270 280 287 274 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148 146 160 155 156 152 150 152 141 138 133 128 128 130 135 136 136 135 133 133 133 133 132 137 142 143 139 139 138 134 130 130 130 130 180 177 182 180 181 179 189 190 186 177 174 174 176 185 191 197 202 204 200 201 203 200 199 202 204 201 201 198 194 191 190 200 205 207 210 208 209 206 205 212 218 220 223 226 228 231 236 235 237 235 237 235 234 234 237 236 237 237 239 234 230 230 228 220 218 217 227 229 234 239 238 232 235 239 233 227 224 223 224 224 220 223 223 220 222 224 226 226 223 220 219 219 222 220 240 241 244 240 253 263 271 277 276 270 271 269 266 270 274 278 280 284 284 284 282 284 288 289 288 287 287 291 296 295 294 260 255 255 263 251 258 260 256 255 254 257 260 263 266 270 274 276 277 279 279 279 281 281 277 275 275 275 276 280 278 276 278 276 280 278 289 273 286 292 304 306 296 299 305 304 307 308 308 308 305 304 304 300 299 300 301 301 299 296 297 298 299 297 140 146 140 152 136 129 116 115 110 105 102 103 109 107 106 106 106 105 108 105 102 100 97 92 93 92 90 91 93 91 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155 161 160 166 167 149 138 136 132 130 128 127 124 128 132 134 134 135 134 135 137 140 143 145 144 144 139 136 135 134 180 178 189 177 185 191 192 197 202 207 213 212 222 229 229 230 233 232 234 232 235 239 236 235 232 231 227 224 221 220 200 198 199 198 206 205 199 192 186 187 193 186 184 185 186 187 188 189 191 190 190 192 197 195 200 199 196 195 200 200 198 220 221 222 224 232 236 232 242 245 244 246 249 254 261 260 258 257 258 263 264 268 271 273 272 276 272 267 271 276 274 273 273 240 231 228 239 238 244 240 235 240 245 254 249 253 253 249 245 246 248 248 248 248 248 248 251 250 252 250 249 247 247 247 247 245 260 264 270 266 258 262 273 281 275 273 271 273 272 271 274 273 274 273 274 276 275 277 276 277 278 276 272 270 270 267 267 265 264 280 274 283 271 272 274 287 298 293 299 301 303 307 306 305 307 309 314 313 313 312 316 319 316 317 314 312 318 327 324 322 140 134 135 133 137 146 152 151 147 146 149 146 151 153 154 153 153 150 151 151 155 153 153 156 158 155 156 158 156 155 154 160 152 156 142 154 152 149 153 153 156 158 158 162 161 157 156 156 155 158 156 155 152 154 155 155 155 149 139 139 138 180 194 196 195 200 198 208 211 202 197 198 196 195 192 193 197 196 196 194 195 199 202 202 202 197 199 199 199 200 198 198 200 197 195 200 197 201 207 210 208 205 207 207 215 217 213 207 208 210 214 215 216 221 224 226 226 226 226 226 225 221 221 220 220 216 228 219 235 234 217 217 213 206 206 209 212 217 219 223 228 227 225 225 221 222 221 225 220 220 220 224 228 228 226 226 240 250 257 251 257 252 260 264 269 273 278 278 281 285 289 289 288 289 294 295 297 299 302 302 301 297 294 297 296 294 293 260 253 261 257 249 241 227 222 227 228 231 235 240 242 242 242 243 239 242 247 245 247 250 251 255 254 253 252 255 252 251 255 254 253 252 255 252 251 280 278 281 284 286 285 283 275 271 268 264 264 264 270 271 273 281 285 286 286 287 287 289 288 288 289 287 288 292 291 289

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## Appendix 3

Number of plots required to estimate site index within a stand with predefined standard error. A "0" notation means either that the predefined standard error cannot be reached, or that the necessary number of plots exceeds 99.

Site variation within stand Predefined stand Low Low Low Medium Medium Migh High High Predefined stand Low Low Low Low Medium Medium Medium Medium Medium Medium Medium Medium	Plot size	Age	at breast	height												
	m⁴	15	25	35	45	55	65	75	85	95	105	115	125	135	145	155
		Nece	ssary nu	mber of p	olots											
Predefined stand	lard error 10 d	lm														
Low	1 000	0	0	0	0	10	6	4	4	5	7	7	15	98	25	0
Low	254	0	0	0	0	17	10	7	6	8	11	12	24	0	41	0
Low	154	0	0	0	0	21	13	9	8	10	14	14	30	0	51	0
Medium	1 000	0	0	0	0	21	13	9	8	10	14	14	30	0	51	0
Medium	254	0	0	0	0	30	19	13	11	14	20	21	43	0	73	0
Medium	154	0	0	0	0	38	23	16	14	18	25	26	54	0	93	0
High	1 000	0	0	0	0	36	22	15	13	17	23	24	50	0	86	0
High	254	0	0	0	0	54	33	23	20	25	35	37	76	0	0	0
High	154	0	0	0	0	64	39	27	23	30	42	44	91	0	0	0
Predefined stand	lard error 12 d	m														
Low	1 000	0	0	0	0	3	3	2	2	2	3	3	3	4	4	17
Low	254	0	0	0	0	5	4	4	3	4	4	5	6	7	6	28
Low	154	0	0	0	0	6	5	4	4	5	5	6	7	9	8	34
Medium	1 000	0	0	0	0	6	5	4	4	5	5	6	7	9	8	34
Medium	254	0	0	0	0	9	8	6	6	7	8	8	10	13	11	49
Medium	154	0	0	0	0	12	10	8	8	9	10	10	13	16	14	62
High	1 000	0	0	0	0	11	9	8	7	8	9	9	12	15	13	58
High	254	0	0	0	0	16	14	11	11	12	14	14	18	22	20	87
High	154	0	0	0	0	19	16	14	13	14	17	17	21	27	24	0

#### Predefined standard error 14 dm

Low	1 000	0	0	0	4	2	2	1	1	1	2	2	2	2	2	3
Low	254	0	0	0	6	3	3	2	2	2	3	3	3	3	3	5
Low	154	0	0	0	8	3	3	3	3	3	3	3	4	4	4	6
Medium	1 000	0	0	0	8	3	3	3	3	3	3	3	4	4	4	6
Medium	254	0	0	0	11	5	5	4	4	4	5	5	5	6	6	9
Medium	154	0	0	0	14	6	6	5	5	5	6	6	7	7	7	11
High	1 000	0	0	0	13	6	5	5	5	5	5	5	6	7	7	11
High	254	0	0	0	20	9	8	7	7	7	8	8	9	10	10	16
High	154	0	0	0	24	11	10	9	8	9	10	10	11	13	12	19
Predefined stan	dard error 16	dm														
Low	1 000	0	0	0	2	1	1	1	1	1	1	1	1	1	1	2
Low	254	0	0	0	3	2	2	2	2	2	2	2	2	2	2	3
Low	154	0	0	0	4	2	2	2	2	2	2	2	2	3	2	3
Medium	1 000	0	0	0	4	2	2	2	2	2	2	2	2	3	2	3
Medium	254	0	0	0	5	3	3	3	3	3	3	3	3	4	4	5
Medium	154	0	0	0	7	4	4	4	3	4	4	4	4	5	4	6
High	1 000	0	0	0	6	4	4	3	3	3	4	4	4	4	4	5
High	254	0	0	0	9	6	5	5	5	5	6	6	6	6	6	8
High	154	0	0	0	11	7	7	6	6	6	7	7	7	8	7	10
Predefined stan	dard error 18	dm														
Low	1 000	0	0	5	1	1	1	1	1	1	1	1	1	1	1	1
Low	254	0	0	8	2	1	1	1	1	1	1	1	1	1	1	2
Low	154	0	0	10	2	2	2	1	1	2	2	2	2	2	2	2
Medium	1 000	0	0	10	2	2	2	1	1	2	2	2	2	2	2	2
Medium	254	0	0	14	3	2	2	2	2	2	2	2	2	3	2	3
Medium	154	0	0	17	4	3	3	3	3	3	3	3	3	3	3	4
High	1 000	0	0	16	4	3	3	3	2	3	3	3	3	3	3	4
High	254	0	0	25	6	4	4	4	4	4	4	4	4	5	4	5
High	154	0	0	29	7	5	5	5	4	5	5	5	5	5	5	6

Site variation	Plot size	Age	at breast	height												
within stand	m²	15	25	35	45	55	65	75	85	95	105	115	125	135	145	155
		Nece	ssary nu	mber of p	lots											
Predefined stand	lard error 20 d	m														
Low	1 000	0	8	2	1	1	1	1	1	1	1	1	1	1	1	1
Low	254	0	14	3	1	1	1	1	1	1	1	1	1	1	1	1
Low	154	0	17	3	2	1	1	1	1	1	1	1	1	1	1	1
Medium	1 000	0	17	3	2	1	1	1	1	1	1	1	1	1	1	1
Medium	254	0	24	5	2	2	2	2	2	2	2	2	2	2	2	2
Medium	154	0	31	6	3	2	2	2	2	2	2	2	2	2	2	3
High	1 000	0	29	6	3	2	2	2	2	2	2	2	2	2	2	3
High	254	0	43	9	4	3	3	3	3	3	3	3	3	3	3	4
High	154	0	52	10	5	4	4	4	3	4	4	4	4	4	4	5

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