

Individual-tree distance-dependent growth models for uneven-sized Norway spruce

Nils Fagerberg*, Jan-Ola Olsson, Peter Lohmander, Martin Andersson and Johan Bergh

Department of Forestry and Wood Technology, Linnaeus University, Universitetsplatsen 1, Växjö 351 95, Sweden

*Corresponding author Tel: +46702828945; E-mail: nils.fagerberg@lnu.se

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Individual tree selection (ITS) is one option to manage uneven-sized forest ecosystems. However, scientifically based field guidelines adapted to ITS and economic profitability are rare, often because there is a lack of suitable tree models to use in growth and treatment simulations. The objective of this study is to develop individual-tree distance-dependent growth models focusing on Norway spruce dominated uneven-sized stands. Three models of different complexity, but with the same structural basis, are presented, followed by some examples of growth patterns for the subject trees. The data include 1456 trees (307 sample trees) collected from five sites in southern Sweden. The basic model (S) depends on subject tree size as the predictor, the second model (SD) adds distance to competitors as a predictor, and the third model (SDC) adds crown ratio as a predictor to the structure. R^2_{Adj} increases with number of predictors from 0.48 to 0.58 to 0.62. The levels of RMSE improve accordingly from $5.02 \text{ cm}^2 \text{ year}^{-1}$ (S) to $4.43 \text{ cm}^2 \text{ year}^{-1}$ (SD) and $4.26 \text{ cm}^2 \text{ year}^{-1}$ (SDC). The present calibration range and model structures primarily make the models suitable for management simulation of individual-tree selection of Norway spruce in southern Sweden. The format of the models allows for further extension with additional predictors and calibration data with greater coverage.

Introduction

Uneven-sized forest management with individual tree selection (ITS) is one way to accommodate diverse expectations from various stakeholders. However, increased application of ITS is, in many regions, hindered by the paucity of scientifically based field guidelines. Currently there are relatively few tree growth models available that are appropriate for the ITS scenario analyses required to develop optimal field recommendations.

ITS is generally understood as a practice in which trees are individually selected for harvest in a compromise between set management aims, e.g. silvicultural, economic or ecological (Pommerening and Murphy 2004; Spinelli et al. 2016). In ideal case, the corresponding field operations follow a science-based protocol of selection guidelines or rules, defined in accordance with the management goals for the forest.

In the context of resource use efficiency (Binkley 2004), ITS guidelines that optimize round wood value production are of interest. This aim implicates that the field selection analysis shall evaluate whether a tree group, which is competing for the same resources as the subject tree, will perform better or worse if the subject tree is selected for harvest or not. This decision involves many dimensions, of which a central part is the ability to forecast individual tree growth of residual trees depending on the decision taken.

The task of defining an appropriate structure for an individual-tree growth model, adapted to the specified requirements, involves compromises between conflicting interests, including model complexity to achieve biological realism (Buchman and Shifley 1983), and model simplicity to allow for flexibility and relevance in field applications (Pacala et al. 1996; Robinson and Monserud 2003). Theory-oriented approaches support more flexible representation of data (Weiskittel et al. 2011) and are potentially less sensitive to extrapolation beyond the observed data range (Pretzsch 2009; Weiskittel et al. 2011) compared with more pure statistical methods. At the other end, theoretical complexity should be constrained to the level at which input and output variables are compatible with units available for the field user (Pukkala and Miina 1998; Pretzsch 2009).

Distance-dependent models (Munro 1974) are adequate tools for quantifying the variation in stand structure, which enables precise prediction of single tree growth (Bella 1971; Mitchell 1975; Daniels et al. 1986; Pukkala 1989; Canham et al. 2004; Pretzsch 2009, p. 310). With distance-dependency incorporated, the model can be used to analyze spatial impact on ITS. Results from distance-dependent models are also easier to transform into useable field guidelines than those from distance-independent models (Pukkala and Miina 1998).

Tree age should not be included as a predictor for growth in uneven-sized stands, due to large variances, both within stands

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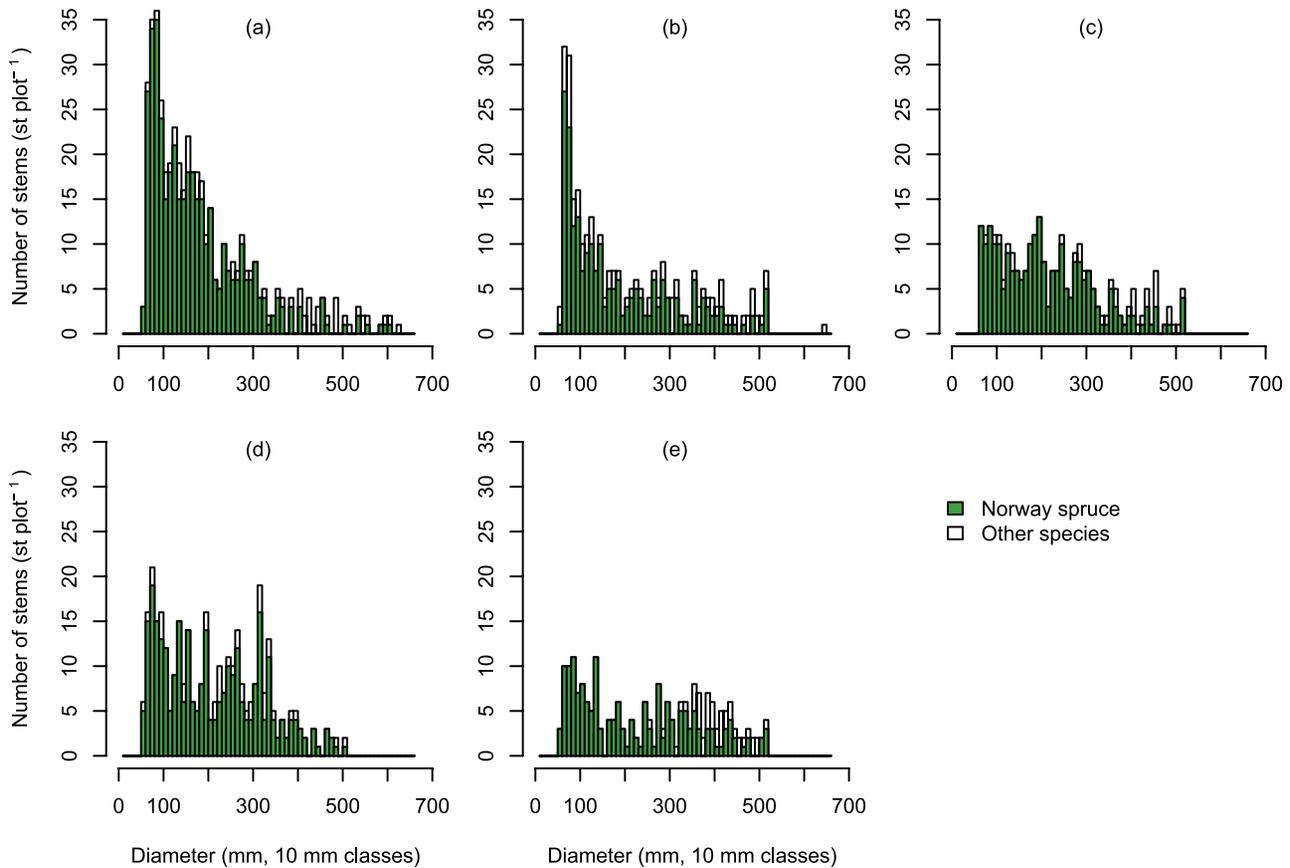
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Table 1 Model training data sets.

| | Measure | Mosshult | Romperöd | Simontorp | Öveshult | Lilla Norrskog |
|--------------|---|-----------|-----------|-----------|----------------------|----------------|
| Site | Latitude (°) | 57°08 | 56°19 | 56°21 | 56°37 | 57°18 |
| | Altitude (m) | 190 | 100 | 120 | 210 | 270 |
| | Site index (SI) ^a | G31 | G30 | G31 | G30 | G29 |
| Stand | Number of stems (st ha ⁻¹) | 920 | 654 | 612 | 726 | 488 |
| | Basal area (m ² ha ⁻¹) | 38 | 30 | 30 | 32 | 32 |
| | Mean dbh (cm) | 19.3 | 20.1 | 22.1 | 20.8 | 25.6 |
| | Proportion of Norway Spruce | 0.63 | 0.58 | 0.74 | 0.83 | 0.62 |
| | Time since last cutting (years) | >50 | 44 | 10 | >30 | >10 |
| | Self-thinning ratio | 0.18 | 0.07 | 0.05 | 0.1 | 0.03 |
| | Number of sample trees | 57 | 43 | 68 | 74 (71) ^b | 65 |
| Sample trees | Mean dbh (cm) | 24.2 | 23.2 | 22.6 | 23.6 | 20.9 |
| | SD dbh (cm) | 13.4 | 12.6 | 12.2 | 10.8 | 13.3 |
| | Max dbh (cm) | 60.5 | 51.5 | 51.5 | 50.9 | 51.2 |
| | Min dbh (cm) | 6.8 | 6.1 | 6.4 | 5.9 | 6.0 |
| | Mean growth (cm ² year ⁻¹) | 6.50 | 8.07 | 8.14 | 6.80 | 9.52 |
| | SD growth (cm ² year ⁻¹) | 6.13 | 7.13 | 7.14 | 5.07 | 8.85 |
| Revision | Period | 2012–2016 | 2012–2016 | 2013–2017 | 2013–2017 | 2013–2017 |

Stand measures based on trees with dbh (diameter at breast height) > 6 cm. Species proportions based on basal area. Self-thinning ratio represents number of trees that have died within the last 10 years divided by number of living trees. SD = standard deviation. ^aSI = height at age 100, according to the Swedish system (Hägglund and Lundmark 1977). ^bThe model that incorporates crown ratio included 71 of the sample trees.


Figure 1 Diameter distribution histograms for the calibration data plots, (a) Mosshult, (b) Romperöd, (c) Simontorp, (d) Öveshult and (e) Lilla Norrskog.

sticks placed along the sub-plot borders. As a consequence of the circular shape of the sub-plots, some additional trees, adjacent to the 60 × 60 m square, were recorded and used for the calibration. The resulting total data set, from the five sites, consisted of 1456 live tree positions. Data on species and cross-calipered dbh was collected for all trees. At one of the sites, tree positioning was also undertaken using a terrestrial laser (Leica P40), which indicated a systematic underestimation of the Postex distances by 1–3 per cent and a random variation of 4 cm (1 SD).

Sample trees ($n = 307$) were selected for additional data collection and later use as subject trees in the model calibration. The selection used stratified random sampling, based on tree variables within four fixed sub-divisions of dbh-classes. The selection within each strata was made subjectively on the basis of two-dimensional plots of preliminary competition variables and maximizing the coverage in these plots. Examples of such variables are tree diameter, summed basal areas of all trees within a certain radius or basal area of larger spruce within a certain radius. Only trees with complete records of competitor tree positions within 10 m were selected as sample trees. The additional data from the sample trees covered tree height, crown length, crown width and annual ring width from the last 5 years. Ring widths were measured from increment cores extracted from the north side at breast height (corresponding calendar years are referred to as revision period in Table 1), using an LINTAB 6 device (RinnTech). Mean measurement error was estimated to 0.016 mm, of which 38 per cent is explained by technical error and the remaining part is due to intra ring width variations and shrinkage. To be included in the data set, the sample trees also had to meet the following criteria: (1) no detected root rot within the sapwood, (2) a live top shoot, and (3) no visible damage that could impair growth and that has been caused by factors other than tree competition.

To prepare the data set for calibration of the model the tree diameters at the start of the growth period had to be estimated. To do this, the ring widths of the last 5 years were used to calculate the total diameter growth for each sample tree. From this, simple linear regression was used to estimate diameter growth as function of tree diameter. The regression model was then used to reduce the diameter of all trees except the sample trees. For the sample trees, the year ring measurements were used directly.

Modelling

The selected model structure can be described as of ‘grey box’ type (Pretzsch 2009, p. 428), built up by structures that can, to some extent, be explained by biological or physical phenomena. The first part represents the internal growth potential of the subject tree and follows the structure suggested by Lohmander (2017), see equation (1). The expression consists of two terms describing the biotic potential and the constraints due to increasing size (Zeide 1993).

$$\frac{dx_i}{dt} = a_1 x_i^{0.5} + a_2 x_i^{1.5} \quad (1)$$

where $\frac{dx_i}{dt}$ is the basal area increment of subject tree i ($\text{cm}^2 \text{ year}^{-1}$), x is basal area (m^2) and the parameters a are regression coefficients. The exponent 0.5 of the first term (the biotic

potential) is defined by proportions between tree diameter (square root of the basal area), vertical crown projection area and basal area growth. The second term, the internal growth constraint, is a relative share multiplier, defined by the difference between 1 and a fitted proportion of the basal area, $1 - bx$. Hence, $\frac{dx_i}{dt} = a_1 x^{0.5} (1 - bx) = a_1 x^{0.5} - a_1 b x^{1.5}$ (Lohmander 2017).

The second part, the competition index (CI), adds external growth restriction to the first internal growth part due to inter-tree competition based on size-distance indices (see equation (2) for the fundamental structure). The formula is a development from the general structure described by Burkhardt and Tomé (2012, p. 210).

$$CI_i = \sum_{j \neq i}^n x_j g(x_i, x_j) f(R_{ij}) \quad (2)$$

where CI is defined as the sum of the basal area x of all competitor trees, weighted by a size weighting function ($g(x_i, x_j)$) and by a distance weighting function ($f(R_{ij})$), in relation to each competitor tree j . From the assumptions above, candidate formulae for CI were created and evaluated separately. For the distance weighting function, $f(R_{ij})$, three requirements were set:

1. The effect of competition shall decrease with distance, i.e. $\frac{df(R)}{dR} \leq 0$.
2. The decrease with distance must be strong enough so that the competition from trees close to the subject tree is more important than the total competition from trees far away. Expressed formally, this means that there should exist a distance, d , probably in the range of 4–10 m, for which the area integral $\int_0^d f(R) \cdot 2\pi R \cdot dR$ is greater than $\int_d^\infty f(R) \cdot 2\pi R \cdot dR$. In other words, for a tree in a forest with reasonably even stem distribution, the summed competition from trees within the distance of d is more important than the sum of the competition from all other trees in the forest.
3. Getting closer to the subject tree, the weighting function shall approach a maximum value asymptotically, i.e. $\lim_{R \rightarrow 0} f(R) = k$ and $\frac{df(0)}{dR} = 0$.

Once the basic CI formula was selected, it was further developed and fitted simultaneously with the complete growth model. Please refer to Olsson and Fagerberg (2019) for further details.

The candidate formulae of the total model expressions were evaluated in the first development phase with adjusted R^2 , statistical properties of the regression parameters and residual analysis. In a second phase, when the main part of the model structure was formulated, 10-fold cross-validation was also employed. For the cross-validation, the total sample was randomly divided into ten groups of equal numbers of subject trees. From this grouping, ten different training samples were created by omitting one test group in each sample; i.e. each test group contained unique observations where each observation is used only once for validation. The model was then fitted to each of these training samples and the performance was studied with the excluded validation group (residual plots and RMSE), see Figure 2. Furthermore, the sample-to-sample-variation of the parameter values were standardized and analyzed (Figure 3). Empirical studies have shown that 5- to 10-folds is optimal to yield test error rate

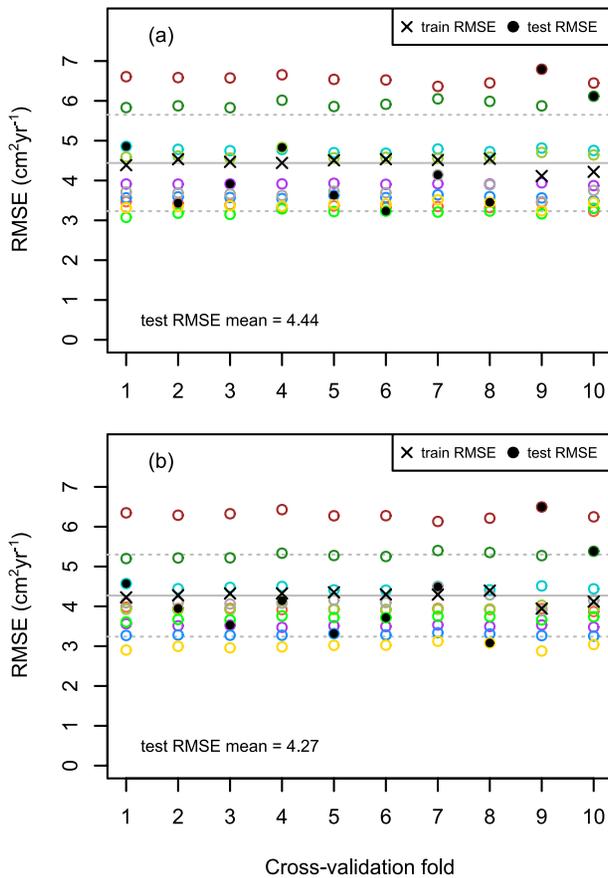


Figure 2 RMSE per cross-validation fold with (a) the SD-model and (b) the SDC-model (included predictors; S=tree size, D=inter-tree distance, and C=crown ratio). Lines represent test RMSE values; grey solid = mean, grey dotted = standard deviation. Test sets are indicated as empty circles with a unique colour per sample when they are part of the training set. Cross indicates training set mean value. Differences between training and test RMSE means are 0.01 for the SD-model and 0.01 for the SDC-model (training RMSE d.f. ≥ 273 , test RMSE d.f. ≥ 30).

estimates with a balanced trade-off between high bias and high variance effects (James et al. 2013).

Three model alternatives were developed with different numbers of predictor variables: (1) with the predictor subject tree size (S); (2) with the predictors tree size and distance to competitors (SD); and (3) with the predictors tree size, distance to competitors and crown ratio of the subject tree (SDC), see equations (3)–(5).

$$S: \frac{dx_i}{dt} = a_1 x_i^{0.5} + a_2 x_i^{1.5} = x_i^{0.5} (a_1 + a_2 x_i) \quad (3)$$

$$\begin{aligned} SD: \frac{dx_i}{dt} &= a_1 x_i^{0.5} + a_2 x_i^{1.5} + a_3 x_i^{0.5} \left(\sum_{j \neq i} x_j \left(\frac{x_j}{x_i} \right)^{k_2} e^{-\left(\frac{R_{ij}}{k_3} \right)^2} w_j \right)^{k_1} \\ &= x_i^{0.5} \left(a_1 + a_2 x_i + a_3 \left(\sum_{j \neq i} x_j \left(\frac{x_j}{x_i} \right)^{k_2} e^{-\left(\frac{R_{ij}}{k_3} \right)^2} w_j \right)^{k_1} \right) \end{aligned} \quad (4)$$

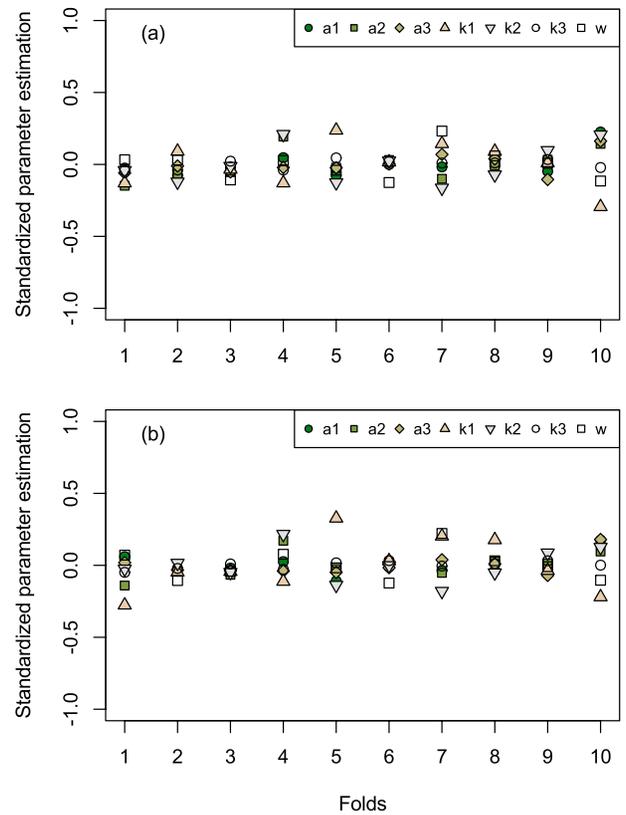


Figure 3 Fitted parameter estimates per cross-validation fold, standardized as relative deviation from mean, (a) SD-model and (b) SDC-model (included predictors; S=tree size, D=inter-tree distance and C=crown ratio).

$$\begin{aligned} SDC: \frac{dx_i}{dt} &= CR \cdot a_1 x_i^{0.5} + CR \cdot a_2 x_i^{1.5} + CR \cdot a_3 x_i^{0.5} \\ &\quad \times \left(\sum_{j \neq i} x_j \left(\frac{x_j}{x_i} \right)^{k_2} e^{-\left(\frac{R_{ij}}{k_3} \right)^2} w_j \right)^{k_1} \\ &= CR \cdot x_i^{0.5} \left(a_1 + a_2 x_i + a_3 \left(\sum_{j \neq i} x_j \left(\frac{x_j}{x_i} \right)^{k_2} e^{-\left(\frac{R_{ij}}{k_3} \right)^2} w_j \right)^{k_1} \right) \end{aligned} \quad (5)$$

where CR is the living crown ratio of total tree height (living crown length is defined by the lowest living branch). w is a parameter used for competitor tree species other than spruce (if spruce then $w=1$). R_{ij} is the distance between the subject tree i and competitor tree j (m). Linear parameters are indicated by a , while non-linear parameters are indicated by k .

The fitting of the models was accomplished in two steps. The first step applied a hierarchical setup using the Matlab function `nlinfit` (a numerical iterative function for non-linear least square regression) to estimate the non-linear parameters. For each call by `nlinfit`, the linear parameters, a_1 to a_3 , were estimated analytically by standard linear least squares inside the call-back function. In this step, each site was allocated a unique value of

Table 3 Regression statistics of the S-, SD- and SDC-models (included predictors; S = tree size, D = inter-tree distance, and C = crown ratio) based on the complete calibration set.

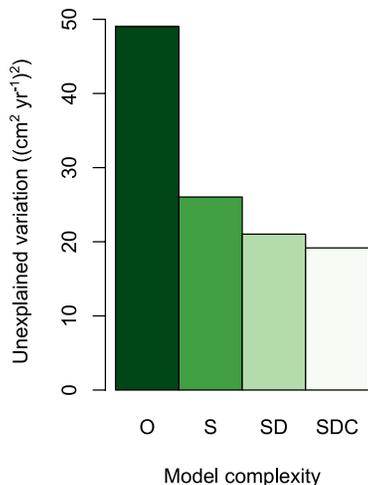
| Parameter | S | | | SD | | | SDC | | |
|-------------|----------|---------|---------|----------|---------|---------|----------|---------|---------|
| | Estimate | t-value | P-value | Estimate | t-value | P-value | Estimate | t-value | P-value |
| a_1 | 43.76 | 33.7 | <0.001 | 106.6 | 92.8 | <0.001 | 154.4 | 104.5 | <0.001 |
| a_2 | -18.13 | -0.95 | 0.344 | -149.1 | -5.5 | <0.001 | -212.2 | -5.7 | <0.001 |
| a_3 | | | | -118.2 | -7.7 | <0.001 | -142.7 | -4.1 | <0.001 |
| k_1 | | | | 0.501 | 2.2 | 0.029 | 0.357 | 1.5 | 0.128 |
| k_2 | | | | 0.276 | 3.1 | 0.002 | 0.325 | 2.7 | 0.008 |
| k_3 | | | | 5.033 | 10.5 | <0.001 | 4.918 | 9.6 | <0.001 |
| w | | | | 0.482 | 3.5 | <0.001 | 0.637 | 3.6 | <0.001 |
| RMSE | 5.02 | | | 4.43 | | | 4.26 | | |
| MSE | 25.2 | | | 19.7 | | | 18.2 | | |
| R^2_{Adj} | 0.481 | | | 0.588 | | | 0.622 | | |

RSME and MSE were calculated consistently for all models by dividing the residual sum of squares by n . R^2_{Adj} = coefficient of determination adjusted for the number of predictors.

Table 4 Site and model specific statistics of the S-, SD- and SDC-models (included predictors; S = tree size, D = inter-tree distance, and C = crown ratio).

| Site | Mean residual | | | Adj R^2 | | |
|----------------|---------------|------|------|-----------|------|------|
| | S | SD | SDC | S | SD | SDC |
| Mosshult | -2.3 | 0.1 | 0.1 | 0.45 | 0.73 | 0.76 |
| Romperöd | 0.4 | -1.1 | -0.8 | 0.52 | 0.55 | 0.66 |
| Simontorp | 0.1 | -0.4 | -0.5 | 0.51 | 0.58 | 0.55 |
| Öveshult | -1.7 | -1.5 | -1.2 | 0.26 | 0.4 | 0.38 |
| Lilla Norrskog | 2.3 | 2.2 | 1.9 | 0.5 | 0.61 | 0.68 |

Mean residuals ($\text{cm}^2 \text{ year}^{-1}$) and adjusted R^2 (Adj R^2).


Figure 4 Unexplained variation depending on model complexity. O represents variance (S^2) of the observations. S, SD, SDC represent test mean square error (MSE) of models S, SD and SDC, respectively (included predictors; S = tree size, D = inter-tree distance and C = crown ratio). Degrees of freedom are equal to n in all estimates.

(see section Modelling). If the first two terms, represented by model S, are solved as a differential equation, see [Lohmander \(2017\)](#), the explicit solution shows that annual diameter increment is almost constant as long as the trees are moderately sized. This increment capacity then decreases as the tree grows larger and the internal processes require an increasing proportion of the collected resources. Since the model is expressed as basal area growth this behaviour may not be obvious when just looking at the expression.

The third term, the competition index, is regarded as the sum of the basal area of competitor trees, but with weighting factors for the different competitors and an exponent (k_1) on the summed competition. When the third predictor variable (CR) was added, the best result was achieved when it was included as a multiplier in all three terms. This can be viewed as a scaling of the complete model, but of course, the values of parameters change to adopt for this.

The competition index

Basal area was selected as the unit to quantify competition since it produced less parameter variation compared with using dbh. The competition term also includes the square root of the basal area of the subject tree ($x_i^{0.5}$, corresponding to dbh). This is because it was found that the summed competition

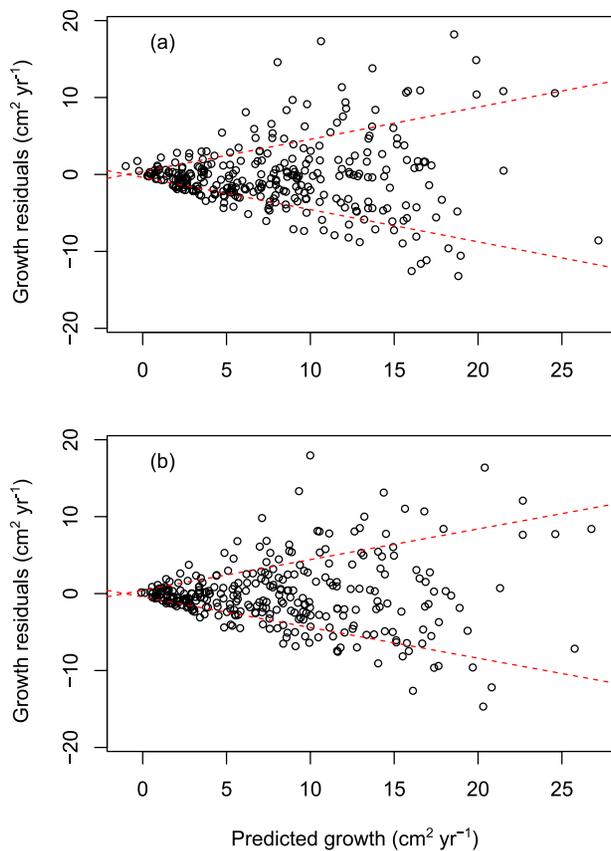


Figure 5 Growth residuals plotted against predicted growth, with (a) the SD-model ($n = 307$) and (b) the SDC-model ($n = 304$) (included predictors; S = tree size, D = inter-tree distance, and C = crown ratio). Lines indicate ± 1 SD (red dotted), calculated with linear regression of accumulated RMSE values where each accumulation step is represented by the mean value within a window of 15 observations.

resulted in a certain reduction of the annual ring width, rather than a certain reduction of the basal area growth. The effect of competitor tree species on the inter-tree competition was considered by the weight (w) (≤ 1) which reduces this effect from species other than spruce. The data available do not support unique weighting factors for the different species. The estimates of w support previous findings stating that competition from other species results in less growth reduction than competition from spruce (Pukkala et al. 2009, 2013). In this study, competition influence from other species was allocated weights of 48 per cent (SD) and 64 per cent (SDC) compared with spruce.

The parameter k_1 (0.50 and 0.36) exhibits a decreasing competition impact per unit (basal area) with increasing levels of competition. This effect was less significant when crown ratio was added to the model. The parameter k_2 adjusts the relative size estimate of competitors. In some other studies, only competition from larger trees is included (Lorimer 1983; Pukkala and Kolström 1987). This kind of approach assumes a one-sided competition (Burkhardt and Tomé 2012, p. 215) that results in a step

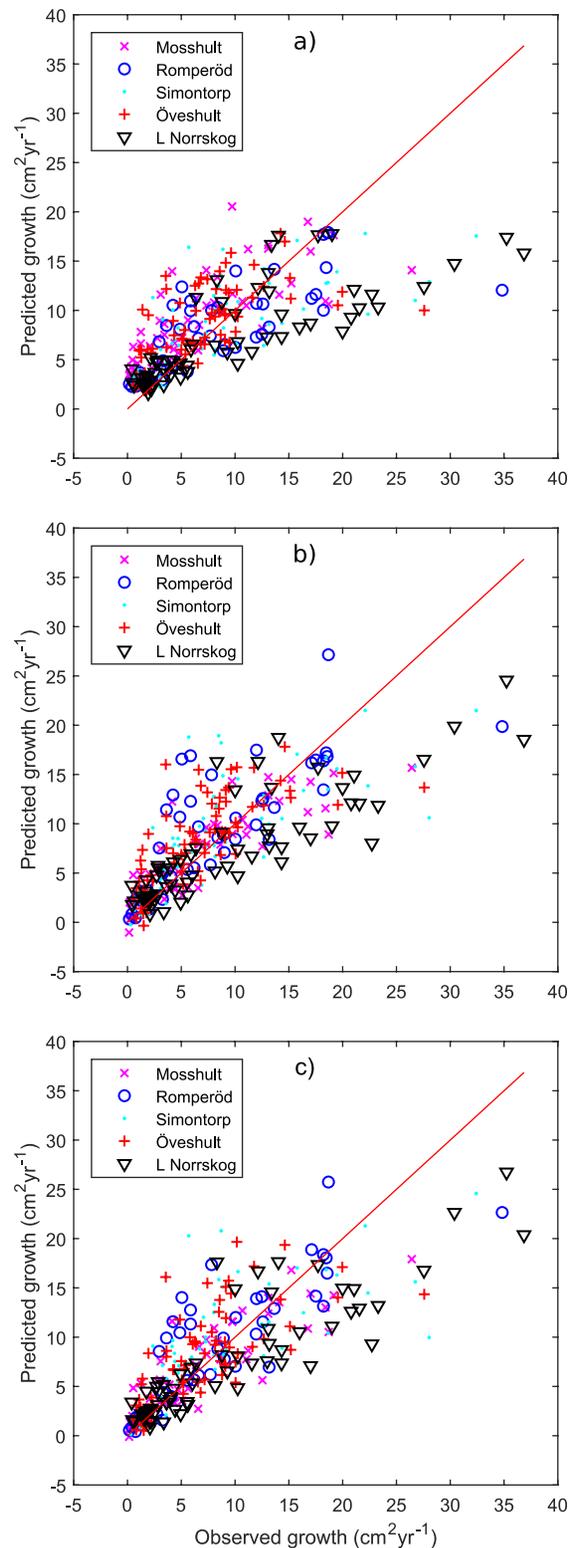


Figure 6 Predicted growth plotted against observed growth presented per model and site (a) S-model, (b) SD-model and (c) SDC-model (included predictors; S = tree size, D = inter-tree distance and C = crown ratio). The red line indicates perfect prediction.

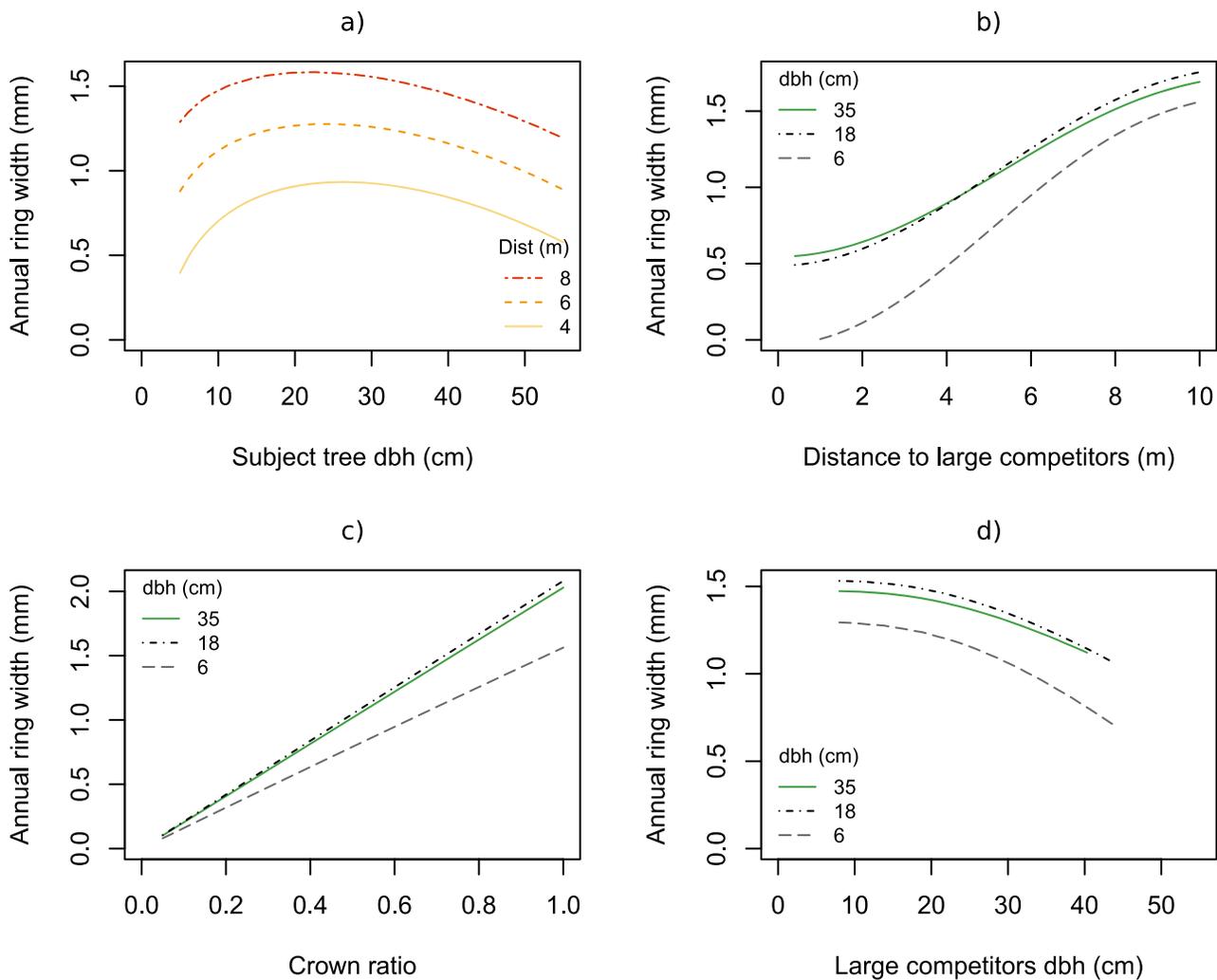


Figure 8 Subject tree annual ring widths from an uneven-sized growth competition case, calculated with the SDC-model (included predictors; S = tree size, D = inter-tree distance, and C = crown ratio). All diagrams represent the same group of competing trees with settings kept constant except for the respective x variable displayed, i.e. (a) subject tree dbh (diameter at breast height), (b) distance to large competitors, (c) crown ratio and (d) dbh of large competitors. Tree group start setting; total basal area = $20 \text{ m}^2/\text{ha}$, crown ratio = 0.6, number of large competitors = 4, dbh of large competitors = 35 cm, dbh of small competitors = 8 cm, all trees are Norway spruce and all competitors are located 6 m from the subject tree. Group basal area is kept constant by adjusting the number of small competitors according to the sizes of the large trees (initial number of small competitors depending on subject tree dbh are 29, 43 and 48, respectively). Dist is distance between large competitors and the subject tree.

ultimately motivated by spatial ITS optimization aims, the performance appears to be satisfactory compared with existing models.

Some research suggests that distance-dependent models are able to reliably predict growth in stand types outside the range of the calibration data (Clutter et al. 1983; Vanclay 1994). This study, however, indicates systematic simulation bias for applications in even-sized structures when calibration is done with uneven-sized data.

Given the data used for the calibration, the model can be expected to be valid for uneven-sized Norway Spruce in Swedish boreal conditions at latitude 56° – 58° and medium to fertile site qualities. The proportion of other species should be moderate (<40 per cent). Since the models depend on spatial information, general use is hampered by limited access to coordinate set

data, but the rapid technical development within forest mensuration can be expected to remove that shortage in the future. The spatial model format, in which competition is defined with both distance and tree size, enables unlimited ITS management analyses since all the options for creating cost-efficient field recommendations with different combinations of metrics are still available. With access to spatial models, future research will resolve whether distance-dependent information is necessary for optimal ITS management, or if it is sufficient to rely on non-spatial growth models.

Future model development

This study shall primarily be considered as a pilot study based on a limited data set. Considering the presented results and

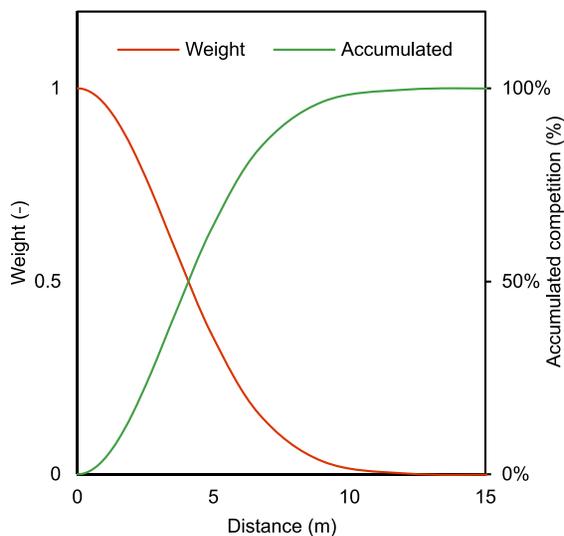


Figure 9 The distance weighing function and its area integral displayed as relative accumulated competition, assuming an even spatial distribution of competitor basal area.

the partly theoretical approach, the models should be suitable for further extension with additional predictors and calibration data with greater coverage. Following the results of Forrester (2021), the candidates for a fourth and a fifth predictor are, in the same order, variables indicative of site quality and treatment. Apart from broadening the geographical area and developing models for additional species, species-specific calibration of the parameter (w) would also make the models applicable to a wider range of situations.

Model performance could potentially be refined if a more representative measure for the crown ratio was applied. Since crown length was measured from the height of the lowest living branch, an inevitable consequence is that stem sections with only fragmented coverage of green needles are in many cases included in the living crown. Estimates that more accurately represent the capacity of the canopy to absorb light could probably improve model prediction, e.g. by defining living crown base as height to the lowest whorl with at least three living branches (Burkhardt and Tomé 2012, p. 100). Another option for developing the model structure is to incorporate subject tree size as a predictor for the distance weighting function. Pretzsch (2009, p. 296) points out that a defined search radius is only adequate for a certain tree size. In the current distance function, weight is solely decided by the distance regardless of subject tree size.

Conclusion

The distance-dependent models presented are well suited for application in simulation studies of uneven-sized stand structures. Specifically, the calibration range and model structure make them appropriate for use within management simulation of individual-tree selection of Norway spruce dominated stands in southern Sweden. The model structure allows for management optimization of influential selection criteria such as subject tree target diameter, subject tree crown ratio, competitor

distance and competitor tree size. The format of the models should make them suitable for further extension with additional predictors and calibration data with greater coverage.

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Conflict of interest statement

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Data availability

The data underlying this article are available in SND [Swedish National Data Service], at <https://doi.org/10.5878/6tkj-tb97>.

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