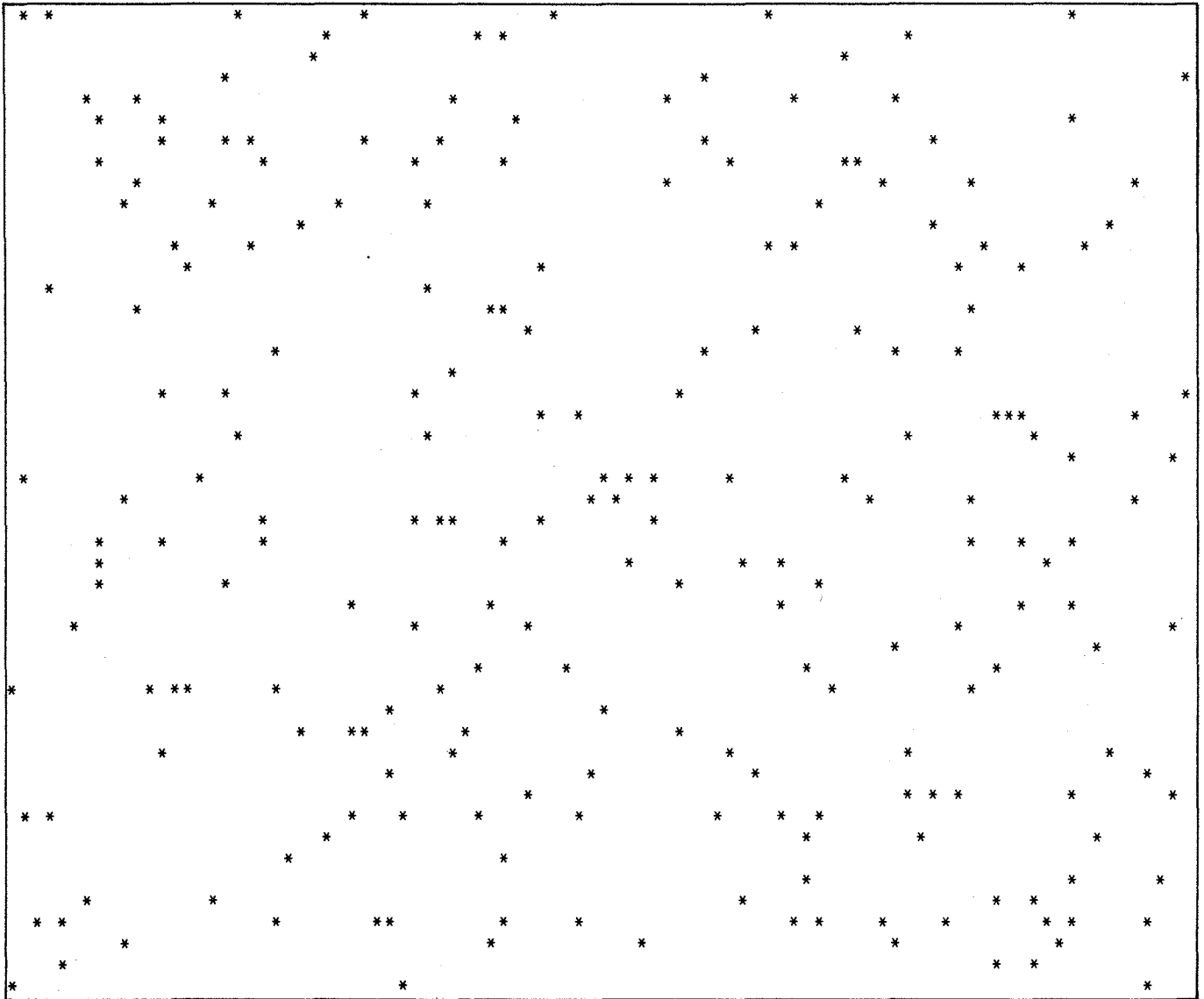


# METHODS FOR MODELING INDIVIDUAL TREE GROWTH AND STAND DEVELOPMENT IN SEEDED LOBLOLLY PINE STANDS



METHODS FOR MODELING  
INDIVIDUAL TREE GROWTH AND STAND DEVELOPMENT  
IN SEEDED LOBLOLLY PINE STANDS

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## ABSTRACT

Methods were developed to model growth and development of seeded loblolly pine (Pinus taeda L.) stands, using individual trees as the basic growth units. Aggregated spatial patterns and individual tree sizes are generated at age 10. Tree diameters and heights are then incremented annually as a function of their size, site quality, competition from neighbors, and stochastic components representing genetic and microsite variability. Individual tree mortality is determined stochastically through Bernouli trials. Subroutines were developed to simulate the effects of hardwood competition and control, thinning, and fertilization. The overall model was programmed in FORTRAN and initial tests were made with published yields. The initial stand generation components were calibrated using a comprehensive set of data from young seeded stands of loblolly pine, but individual tree growth and mortality components relied on previously published relationships developed for plantations. Results indicated that, in order to accurately model stand structure, the growth and mortality relationships must be calibrated for seeded stands. Data collection procedures, calibration methods, and recommendations for further work are discussed.

## ACKNOWLEDGEMENTS

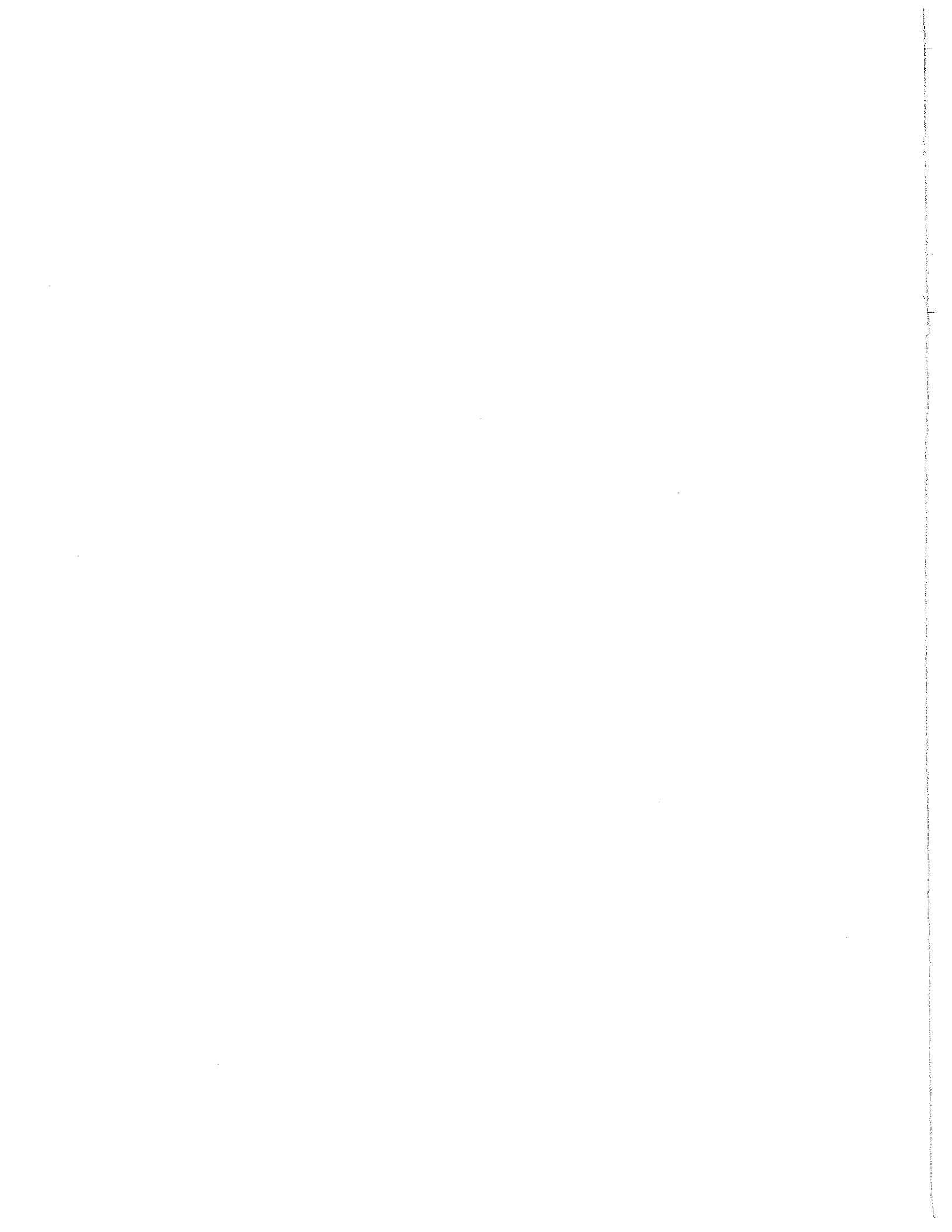
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## COVER

The cover design is a computer-generated spatial pattern for a seeded loblolly pine stand.

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## INTRODUCTION

Loblolly pine (*Pinus taeda* L.) is one of the most commercially important species in the South, with a natural range extending from Maryland through the southeastern and southern states to east Texas. Although recent emphasis has been on plantation management, there exist millions of acres in natural and direct-seeded loblolly pine stands. Increasing loblolly production to meet future demands will require thorough regeneration of all cutover pine sites (Boyce 1975) and natural and direct-seeding should become increasingly attractive regeneration alternatives.

Most recent studies of loblolly pine growth and yield have considered only plantations and those that have considered seeded stands have worked only with natural stands. However, intensive management has reached the point where the forest manager is faced with a number of regeneration alternatives as well as intermediate cultural treatments. Flexible models capable of providing detailed growth and yield information for the range of available management options have been developed for some species, including planted loblolly pine (Daniels and Burkhart 1975), but are badly needed for seeded loblolly pine.

The objectives of this study were to identify, formulate, and where possible quantify individual tree and stand level relationships in natural and direct-seeded loblolly pine stands for the purpose of constructing a flexible tree and stand growth model. In this paper methods are presented for the development and calibration of an individual-tree-based model of stand development for seeded loblolly pine.

The modeling approach taken is drawn from that of Daniels and Burkhart (1975) in their model for managed loblolly pine plantations. Stand development is modeled as the growth and competitive interaction of individual trees. This offers flexibility since it allows use of both tree- and stand-level information and may be closely tied to biological growth processes. Spatial and competitive relationships can be incorporated directly in such a model. Thus, it lends itself to study of intensive management practices such as thinning and fertilization. Because individual tree locations are known, this type of model is naturally suited to the study of stand development in seeded stands where irregular spatial patterns may affect growth.

## RELATED WORK

### Growth and Yield/Stand Modeling

#### Stand Level Models

Yield prediction in natural loblolly pine stands began with classical normal yield tables constructed using graphical techniques from data collected in natural stands of "normal" density (Anon. 1929). Modern quantitative study of growth and yield got its start with MacKinney and Chaiken's (1939) application of multiple regression analysis in constructing a variable density yield equation for loblolly pine. Since that time a number of studies have used multiple regression analysis to construct yield equations for natural and planted southern pine stands (Bennett, et al. 1959, Clutter, 1963, Goebel and Shipman 1964, Burkhart, et al. 1972a, 1972b, and others). Schumacher and Coile (1960) presented a comprehensive study of the growth and yield of natural stands of southern pines which relied on both graphical and regression techniques.

A number of studies have used a diameter distribution analysis procedure for yield prediction in southern pine plantations (Bennett and Clutter 1968, Lenhart and Clutter 1971, Lenhart 1972, Burkhart and Strub 1974, Smalley and Bailey 1974a, 1974b). In this approach a probability density function is used to model the diameter distribution. The number of trees in each diameter class is estimated, total heights are predicted, and volume is calculated by substituting into tree volume equations. Unit area estimates are made by summing over diameter classes of interest. This technique has had very limited application in seeded southern pine stands.

#### Individual Tree Models

Stand models which use the individual tree as the basic growth unit will be denoted individual tree models. Munro (1974) further segregated this class of models into distance dependent and distance independent categories depending on whether or not individual tree locations are required in the list of tree attributes. Distance independent models may simulate tree growth either individually or by size classes, usually as a function of present size and stand level attributes. No general form has been followed in the construction of individual tree distance independent models so it is difficult to make general statements about their structure. Examples of distance independent models are found in the work of Goulding (1972), Stage (1973), Dale (1975), and Botkin, et al. (1970).

Distance dependent models that have been developed, although varying in detail, have, in general, shared a common structure. Initial tree and stand attributes are input or generated and each tree is assigned a coordinate location. The growth of each tree is simulated as a function of its size, the site quality, and a measure of competition from neighbors. The competition index varies from model to model (see e.g., Bella 1971, Gerrard 1969, Keister 1971, Moore, et al. 1973, Daniels 1976, Alemdag 1978) but in general is a function of the tree's size in relation to the size of and distance to competitors (hence, the need for individual tree locations). Mortality may be controlled either probabilistically or deterministically as a function of competition and/or other individual tree attributes.

Individual tree distance dependant models provide very detailed records of stand structure and development and are well suited for inclusion of routines to simulate cultural treatments. Since Newnham and Smith's (1964) original model for Douglas-fir and lodgepole pine a number of advancements have been made which have allowed evaluation of the effects of various management regimes. By varying initial spatial patterns of trees in a stand, the effects of different regeneration alternatives may be evaluated. The ability to generate regular, random, and aggregated patterns was included in Bella's (1970) aspen model, Hatch's (1971) red pine model, and others. Arney (1974) modeled growth along the entire bole of the tree which allowed examination of tree taper and volume relationships. A flexible model capable of simulating development of uneven-aged mixed-species stands was introduced by Ek and Monserud (1974). Thinnings have been studied using distance-dependant models since it is generally felt that response follows directly from the competition relationships included. Response to fertilizer has also been studied (Ek and Monserud 1974, Heygi 1974).

Daniels and Burkhart (1975) developed a model for loblolly pine plantations which includes routines to simulate the effects of site preparation levels, thinning regimes, and fertilizer applications. To date their work represents the only published application of individual tree distance dependent modeling techniques to southern pine species; the model is finding utility in both research and practical industrial applications.

### Spatial Patterns

Interest in quantitative descriptions of forest spatial patterns has increased with the development of distance dependant stand models,

especially when considering the irregular patterns found in seeded stands. Quadrat and distance sampling methods have both been used to quantify departures from random spatial arrangements (see Pielou 1969). Both methods have numerous variations, but almost all published studies involve comparisons of observed spatial characteristics (e.g., plot stem counts in quadrat sampling and distances from random points to nearest plants in distance sampling) with those expected in random populations of the same density, providing both an index and a test for the degree of nonrandomness.

Quadrat sampling is generally easy to apply in the field and can be quite reliable, but estimates of nonrandomness may vary with plot size (Pielou 1969). Distance sampling has been suggested to avoid dependence on plot size, but usually requires an independent density estimate for inferences on spatial patterns. Distances from random points to nearest plants (point-to-plant) and distances from random plants to nearest plants (nearest neighbor) have both been used to quantify spatial patterns. Point-to-plant distances are often preferred since it is difficult to choose plants at random in nonrandom stands (Pielou 1969). After comparing several techniques Payandeh (1970) recommended point-to-plant distance sampling and Pielou's index of nonrandomness for quantifying spatial patterns in natural and computer-generated forest populations.

A number of theoretical frequency distributions have been used in spatial studies. The number of individuals per unit area has been described by the Poisson distribution in random populations and by the negative binomial distribution, the Neyman type A distribution and others in clumped populations (Pielou 1969, Southwood 1966). Ker (1954) demonstrated the utility of the negative binomial distribution in examining spatial patterns in young naturally seeded pine stands. The negative binomial distribution has properties that make it desirable for clumped pattern description. For example, it may be derived as the distribution resulting from any of a number of causal mechanisms which produce clumping (Pielou 1969, Southwood 1966) and its two parameters may be directly interpreted as an overall density parameter and a heterogeneity parameter (loosely, a "clumping factor"). The distribution tends to the Poisson distribution as the heterogeneity parameter tends to infinity. A direct correspondence exists between the discrete quadrat sampling distributions discussed above and continuous distributions of point-to-plant distances. Eberhardt (1967) and others have derived distance distributions for populations in which quadrat sampling would yield Poisson and negative binomial distributions of plot densities.

Daniels (1978) used point-to-plant distance methods and Pielou's (1959, 1969) index of nonrandomness to quantify spatial patterns in 40 5-to-12-year-old loblolly pine stands of seed origin. His work indicated that aggregated, or clumped, patterns were prevalent in all seeding methods studied, including natural (old field), seed tree, broadcast, and aerial methods. Further, nonrandomness index values were not found to be related to seeding method or stand attributes such as age, site index, or stand density.

Distance frequencies were further described by Daniels (1978) using distribution methods. By using squared distance as the variate he derived a form of the Pearson type XI distribution from the aggregated distribution proposed by Eberhardt (1967). The Pearson type XI distribution fit observed values well and was proposed as a general spatial model for seeded stands. Because of its relationship to the negative binomial distribution, its parameters were also interpreted in terms of stand density and heterogeneity. A direct relationship was shown between the heterogeneity parameter and Pielou's index of nonrandomness.

A number of computerized algorithms have been developed to generate spatial arrangements of points. Regular patterns are simple to generate by placing points on a grid. Random patterns may be produced by generating coordinates from a uniform distribution. Aggregated patterns have been generated by concentrating points around clump centers and by establishing density gradients for the placement of points (Newnham 1968, Newnham and Maloley 1970). Wensel (1975) used a method involving a probability matrix which was altered to increase or decrease the probability of future points being located within a certain distance of the point just located.

Although realistic aggregated patterns resulted from the above algorithms, none are related to field measures of spatial pattern mentioned earlier. This prompted Daniels and Spittle (1977) and Stauffer (1978), independently, to develop methods of generating spatial patterns with known spatial parameters (e.g., Pielou's index) by using distributions of point-to-plant distances. This work will be discussed later.

## METHODS

The basic modeling philosophy and framework used by Daniels and Burkhardt (1975) for loblolly pine plantations was adopted in constructing model components for seeded loblolly pine stands. In this approach, stand development is divided into two stages. The first stage involves the generation of an initial stand of trees at the onset of competition. The second deals with the annual growth and development of that stand by simulating the growth, mortality, and competitive interaction of individual trees. Added to this structure are routines to simulate intensive management practices such as thinning and fertilization.

This section provides detailed descriptions for model components in the initial stand generation and stand development stages and for the management routines. Special emphasis has been placed on identifying and quantifying components unique to seeded stands.

### Initial Stand Generation

The initial stand generation stage involves the complete specification of the stand spatial pattern and size distributions including the assignment of individual tree coordinate locations, dbh, height, and crown length. Realistic specification of early stand structure is crucial to subsequent simulation of stand dynamics. The aggregated spatial patterns found in seeded stands are much more complex to model than the simple rectangular patterns of plantations. Size distributions are also more varied. Daniels and Burkhardt (1975) employed a prediction of the age at which intraspecific competition begins to determine the age to generate tree sizes and to begin annual growth computations. This approach was questioned for seeded stands due to the higher degree of variability in size and spatial relationships and even in age itself for some seeding types. These considerations prompted intensive investigations into methods for realistically generating size and spatial relationships in young seeded stands.

### Spatial Patterns

A spatial pattern generator for seeded stands must be capable of generating patterns with varying degrees of aggregation at different levels of stand density. An algorithm was desired which would produce patterns of known aggregation, as measured by an index such as Pielou's. Such an algorithm, which works by essentially inverting the sampling procedures used in point-to-plant distance sampling, was developed.

The Pearson type XI distribution was suggested by Daniels (1978) as a general model for describing squared point-to-plant distances in

seeded stands. This distribution, used here as the basis for generating spatial patterns, may be written with cumulative density function (c.d.f.)

$$F_w(w) = 1 - (1 + \frac{c}{k} w)^{-k}, w > 0$$

where,

w = squared point-to-plant distance

k = heterogeneity parameter

c = density parameter (number of trees per circle of radius = 1 (foot))

Daniels (1978) further noted that the heterogeneity parameter, k, of the Pearson type XI distribution may be estimated by the simple function of Pielou's index of nonrandomness

$$\tilde{k} = \frac{\alpha}{\alpha - 1}$$

where,

$\tilde{k}$  = estimated value of k

$\alpha$  = Pielou's index of nonrandomness

Thus, input to a spatial pattern generator based on this distribution requires only knowledge of the stand density, c, and the nonrandomness value,  $\alpha$ , desired. Such a generator would be applicable to all types of seeded stands including seed tree, natural, aerial, and broadcast seeding.

By inverting the distribution function via the probability integral transformation, values of a Pearson type XI distributed random variable can be generated stochastically. Specifically, squared distances from random points to nearest trees are generated from the following equation:

$$w = \frac{k}{c} [(1-u)^{-1/k} - 1]$$

where,

k = heterogeneity parameter

c = density parameter

$u$  = a random number from the uniform (0,1) distribution

The distance from a random point to the nearest tree,  $r = \sqrt{w}$ , defines a circle of radius  $r$ , centered at the random point, within which no trees are located, but with one tree located on the perimeter. A set of such distances then describes a set of circular open areas. Circles of open area with radius  $r_i$  are generated and then allocated to random points distributed throughout a given area. Actual coordinates of the trees are determined by fixing their positions on the circumference of the generated circles, i.e., by fixing the angles  $\theta_i$  (Figure 1).

In programming this algorithm, steps had to be taken to ensure that no tree be positioned within the open area associated with another tree. This required detailed accounting and mapping of available space on the plot to check, as trees were positioned sequentially, that 1) no new tree location was fixed within the open area of a tree previously positioned, and 2) open areas of new trees contained no previously positioned trees.

Experience with the algorithm indicated that it provided a flexible tool for generating aggregated patterns over a wide range of conditions. However, because of the constant checking for the two conditions mentioned above, computer time and storage demands were judged too high for practical inclusion in a forest stand growth model.

Independently, Stauffer (1978) developed a set of algorithms for aggregating points to fit Pielou's index which was also based on inverting distance sampling methods. He reported biases in his approach; generated aggregation was considerably less than that specified by the input value of Pielou's index. His observed bias is explained by the use of inappropriate squared-distance distributions (e.g., the exponential distribution) and the relaxation of condition 2) above (i.e., no check was made on new tree open areas).

A "hybrid" spatial pattern generator was then developed which used the Pearson type XI distribution to generate squared distances, but in which condition 2) was relaxed. The result provided a generator capable of producing aggregated stands in seconds (rather than minutes) with considerably less aggregation bias than reported by Stauffer (1978). This modified Stauffer algorithm was thus adopted for generating seeded stand spatial patterns.

### Size Distributions

After generating the initial stand spatial pattern and assigning tree coordinates, tree sizes are assigned. A two parameter Weibull

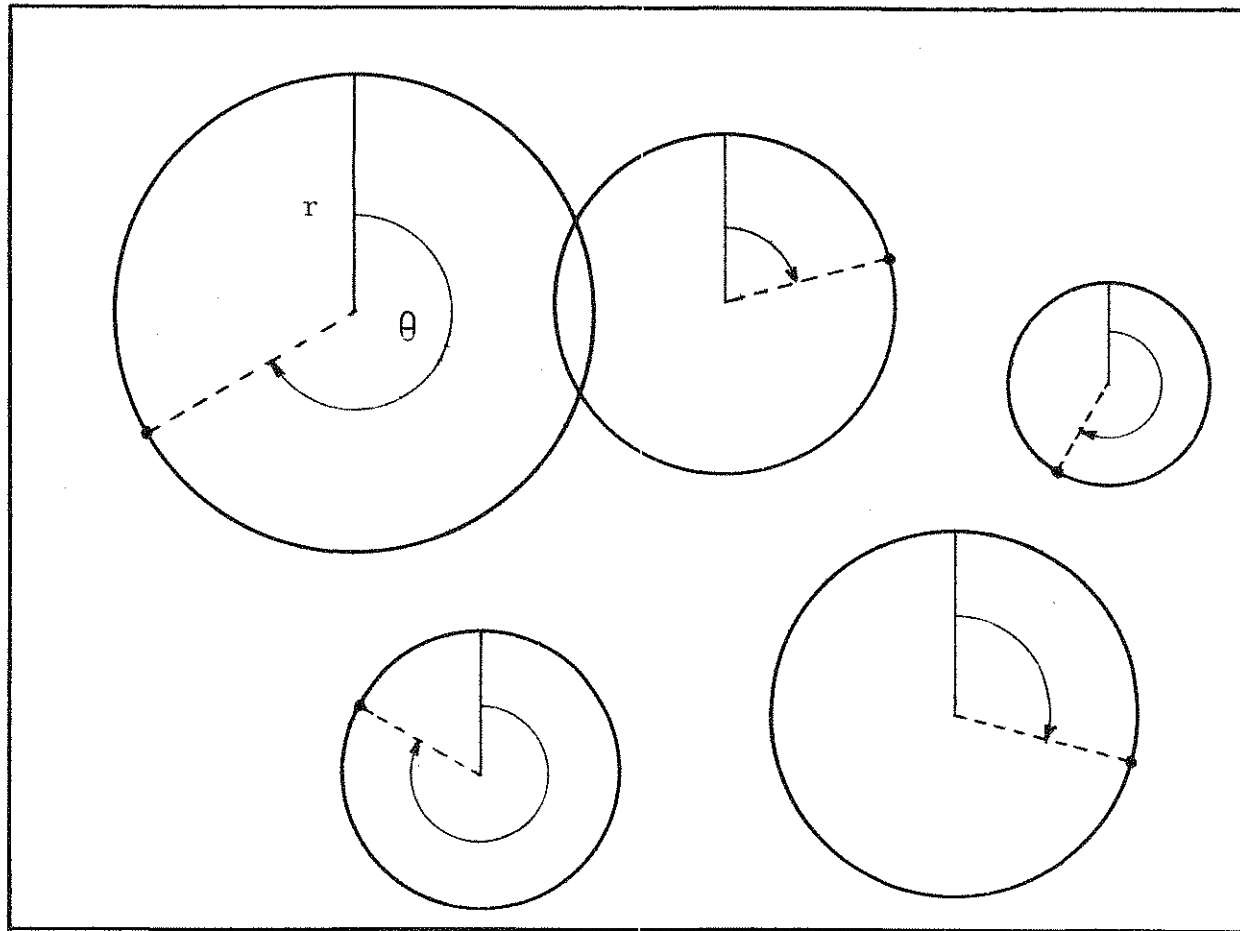


Figure 1. Determining tree positions by fixing distances ( $r$ ) and angles ( $\theta$ ) from random points.

function was chosen to model the diameter distribution of the initial stand. This function can be written with cumulative distribution function (c.d.f.)

$$F_y(y) = 1 - e^{-ay^b} \quad 0 < y < \infty$$

Specifically, diameter at breast height is generated from the function

$$D = \left[ -\frac{1}{a} \ln(1-u) \right]^{1/b}$$

where,

$$D = \text{d.b.h.}$$

$u$  = a random number from the uniform (0,1) distribution

$a, b$  = Weibull parameters

Estimators for parameters  $a$  and  $b$  are

$$\hat{b} = \frac{\ln(N)}{\ln DAVE - \ln DMIN}$$

$$\hat{a} = \left[ \frac{\Gamma(1 + 1/b)}{DAVE} \right]^b$$

where,

DMIN = minimum d.b.h.

DAVE = average d.b.h.

$N$  = number of trees measured for DAVE, DMIN

In conjunction with Daniels' (1978) work, data were collected on size distributions in young seeded stands. Forty 5- to 12-year-old seeded loblolly pine stands were selected from industrial and state ownerships over a wide range of stand conditions in Eastern Virginia and North Carolina (Table 1), to obtain approximately equal numbers in each of the following regeneration categories: 1) seed tree/shelterwood, 2) natural old field, 3) aerial seeded, and 4) broadcast seeded. In each stand, 10 trees were selected for detailed measurements, including d.b.h. total height, crown length, and age. In addition, d.b.h. was determined for all trees in each of three temporary .05-.10 acre plots.

Table 1. Summary of conditions in 40 seeded loblolly pine stands used to derive size relationships for initial stand generation.

Variable	Mean	Range
Age (years)	9	5 - 12
Density (stems/acre)	2067	400 - 6350
Height (feet) <sup>a/</sup>	14.9	7.1 - 30.2
D.B.H. (inches)	1.4	0.1 - 19.1 <sup>b/</sup>

<sup>a/</sup> Average height of dominants and codominants.

<sup>b/</sup> Overstory tree.

Prediction equations were developed to determine DMIN and DAVE in terms of total basal area per acre (BAT) and average height of dominants and codominants (HD) (Table 2). Total height (H) is assigned for each tree using a prediction equation based on d.b.h. (D), HD, surviving number of loblolly pine trees per acre (TS), and age (A) (Table 2). Crown length is determined as total height minus clear bole length (CBL) where CBL is predicted as a function of H, D, TS, and A (Table 2). Coefficients for the equations in Table 2 were solved for using the data summarized in Table 1.

Because of the difficulties involved with determining an age when intraspecific competition begins, a fixed age 10 was chosen for generating the initial stand. It was thought that competition already has begun to affect growth at age 10 in typical seeded stands. To reflect this influence initial diameters are assigned as a function of competition at age 10. For each tree in the stand, d.b.h. is temporarily set equal to DAVE and the competition index is evaluated to provide an index of tree growing space. Actual diameters are then generated, sorted largest to smallest, and assigned to tree locations so that the largest d.b.h. is associated with the smallest competition value, etc. Correlations between tree sizes and spatial measures in young seeded stands were shown by Daniels (1978) to be negligible, but these methods should ensure logical spatial-size relationships.

No attempt was made in the initial stage to project stand conditions to age 10 from some earlier point in time. Input to this stage requires stand information at age 10. Somers, et al.<sup>1/</sup>, derived survivorship curves based on one minus the cumulative density function of the two-parameter Weibull distribution:

$$F(x) = e^{-(x/b)^c}$$

where,

$$F(x) = \text{percent survival}$$

$$x = \text{age}$$

$$c = 2.9561$$

$$b = \text{EXP} [4.9023 - 0.2030 \text{ Log } N_a]$$

$$N_a = \text{initial number of trees at age 3}$$

Then  $F(x)$  times  $N_a$  gives the number surviving at any age  $x$ .

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<sup>1/</sup> Somers, G. L., R. G. Oderwald, W. R. Harms, O. G. Langdon. Predicting mortality with a Weibull distribution. Manuscript submitted to Forest Science.

Table 2. Equations used in generating initial stand in a growth model for seeded loblolly pine.

Equation <sup>a/</sup>	R <sup>2</sup>	S <sub>y·x</sub>
DAVE <sup>b/</sup> = -1.54190 + 1.14324 ln(HD) + 0.0038993 BAT	0.78	0.117
DAVE <sup>c/</sup> = 0.47040 + 0.069485 HD - 0.00000083 A·TS + 5.45478 HD/TS	0.84	0.078
DMIN = -0.067446 + 0.029395 HD - 0.00000112 A·TS + 6.23266	0.75	0.028
ln(H) = 1.44287 + 0.32192 ln(HD) + 0.52118 ln(D) + 0.0026328 BAT + 0.07299/D - 1.08825/A	0.93	0.023
ln(CBL) = -1.43430 + 1.48535 ln(H) - 0.47173 ln(D) + 0.00092034 BAT - 0.10991/D - 3.34385/A	0.96	0.043
ln(TS) = 5.31958 + 0.83535 ln(BAT) + 1.04073 ln(PPINE) - 1.60866 ln(DAVE)	0.85	0.092

<sup>a/</sup> Where DMIN = minimum d.b.h. (inches), DAVE = average d.b.h. (inches), H = total height (feet), CBL = clear bole length (feet), TS = number of loblolly pine trees surviving per acre, BAT = total basal area per acre (ft<sup>2</sup>/acre), HD = average height of dominants and codominants (feet), D = d.b.h. (inches), A = age (years), PPINE = proportion of BAT in pine (pine BA/BAT).

<sup>b/</sup> Used for existing stands only.

<sup>c/</sup> Used in initial stand generation.

The above coefficients were estimated using the data of Harms and Langdon (1976). Briefly, their study consisted of 20, 0.1-acre plots located in the Lower Coastal Plain of South Carolina, all with site index of 105 feet (base age 50). The twenty plots were thinned at age 3 to 5 densities: 1, 2, 4, 8, and 16 thousand trees per acre, with four plots at each density level. Potential users who feel these data are applicable to their stands may wish to use the function above to project stand density at age 3 to that at age 10.

The capacity for simulating existing stands of ages older than 10 years was included. This requires that basal area per acre at the existing age be provided. Basal area is projected back to age 10 using the basal area growth equation of Sullivan and Clutter (1972), average d.b.h. is estimated (Table 2), the number of trees per acre is determined (Table 2), and a stand at age 10 is generated.

### Stand Growth and Development

#### Competition Index

A number of competition indices were evaluated and compared for planted loblolly pine by Daniels (1976). The modified Hegyi index suggested there and used by Daniels and Burkhart (1975) was adopted for seeded loblolly pine stands. It is calculated

$$CI_i = \sum_{j=1}^n (D_j/D_i)/DIST_{ij}$$

where,

D = d.b.h.

DIST = distance between subject tree i and competitor j

$CI_i$  = Competition Index of the tree i

n = the number of neighbors included in a 10 BAF angle gauge sweep with vertex at the subject tree

Competitive stress on border trees is calculated through a translation of plot borders so that border trees compete with border trees on the opposite side of the plot. This technique was suggested by Monserud and Ek (1974) to control plot edge bias.

### Growth Relationships

After generation of the juvenile stand, competition is evaluated and trees are grown individually on an annual basis. In general, growth in height and diameter is assumed to follow some theoretical growth potential. An adjustment or reduction factor is applied to this potential increment based on a tree's competitive status and vigor, and a random component is then added representing microsite and/or genetic variability.

The potential height increment for each tree is considered to be the change in average height of the dominant and codominant trees, obtained as the first difference with respect to age of the following expression, transformed from the site index equation presented by Schumacher and Coile (1960):

$$HD = SI 10^{-6.528(1/A - 1/50)}$$

where,

HD = average height of dominant stand (feet)

SI = site index base 50 (feet)

A = stand age (years)

A tree may grow more or less than this potential, depending on its individual attributes.

Experience in loblolly pine plantations (Daniels and Burkhart 1975) suggested the inclusion of competition index and crown ratio in the height growth adjustment factor with the form

$$(b_1 + b_2 CR^{b_3} e^{-b_4 CI - b_5 CR})$$

where,

CR = crown ratio

CI = competition index

$b_i$  = constants to be estimated from data

The maximum d.b.h. attainable for an individual tree of given height and age was considered to be equal to that when open-grown. An equation describing this relationship was developed from open-grown tree data (Daniels and Burkhart 1975) and is shown below:

$$D_0 = -2.422297 + 0.286583 H + 0.209472 A$$

where,

$D_0$  = open-grown tree d.b.h. (inches)

H = total tree height (feet)

A = age from seed (years)

The first difference of this equation with respect to age was thought to represent a maximum potential diameter increment:

$$PDIN = 0.286583 HIN + 0.209472$$

where,

PDIN = potential diameter increment (inches)

HIN = observed height increment (feet)

This potential diameter increment is reduced by a reduction factor of the form

$$(b_1 + b_2 CL^{b_3} e^{-b_4 CI})$$

where,

CI = competition index

CL = crown length (feet)

The inclusion of measures of photosynthetic potential in the above models plays a key role in determining thinning response. Others have included only competitive effects in such adjustment factors. However, when a tree is released by removing neighboring trees its response will depend not only on the reduction in competition for resources, but the potential it has for using those resources. Both crown length and crown ratio reflect this potential.

Crown length is incremented each year as the difference between height increment and change in clear bole length. Clear bole length is predicted annually as a function of height, d.b.h., age, and basal area per acre (Table 2).

### Mortality

The probability that a tree remains alive in a given year was assumed to be a function of its competitive stress and individual vigor as measured by photosynthetic potential. The probability of survival equation took the form

$$PLIVE = b_1 CR^{b_2} e^{-b_3 CI} b_4$$

where,

PLIVE = probability that a tree remains alive

Survival probability is calculated for each tree and used in Bernouli trials to stochastically determine annual mortality. The calculated PLIVE is compared to a uniform random variate between zero and one. If PLIVE is less than this generated threshold, the tree is considered to have died.

### Management Routines

#### Hardwood Control

Daniels and Burkhart (1975) simulated the effects of competing vegetation and site preparation by including a competition adjustment factor. This factor modified all stand density and competition relationships by, essentially, increasing the number of competing stems. Additional competition was described in terms of "loblolly-equivalent" stems and decreased linearly to a specified age of release.

A similar approach was taken for seeded stands. Three parameters are specified, HDWD, IRLSE, and ARLSE, which determine the proportion of additional competing (loblolly equivalent) stems, the type of release, and the age of release, respectively. If HDWD is set equal to one the number of additional competing stems (in loblolly equivalents) is equal to the number of loblolly stems at age 10. The parameter ARLSE determines the age at which the stand will be released to a pure loblolly stand and IRLSE determines whether the release will be a gradual linear release or a sudden release. The competition adjustment factor (CAF) is

calculated annually from these parameters to obtain the multiplier for competitive relationships.

### Fertilization

The methods used by Daniels and Burkhart (1975) to simulate fertilization were adopted. Fertilizer application was viewed as an adjustment of site quality as measured by site index. A site adjustment factor (SAF) was included which modifies site index for the duration of the fertilizer response. The value of SAF is calculated from three parameters, RESP, LMR, and LR, which specify, respectively, the maximum response in site index, the length of time in years to attain maximum response, and the total length of the response. SAF increases linearly from the time of application until RESP is attained LMR years later, and then decreases linearly until LR.

### Thinning

A thinning routine was constructed which allows thinning from below, by corridors, or in combination. Thinning from below removes trees one at a time, from smallest to largest, until the thinning limit, TLIM is met. The thinning limit may be specified either in terms of residual stand basal area per acre or an upper diameter limit. In either case, a lower diameter limit, DLOW, may be specified below which trees will not be removed. Corridor thinning involves removing a swath of trees. Swaths may be removed in either the x or y direction, or both. Swath widths are controlled by the parameters XCORW and YCORW and swath spacing is controlled by XCORS and YCORS. When used in combination, the corridor thinnings are performed first and the residual stand is then thinned from below to TLIM.

## INITIAL TESTS

A preliminary model, Seed-PTAEDA, based on Daniels and Burkhardt's (1975) plantation model was programmed in FORTRAN IV to include the seeded stand components discussed earlier. The initial stand generation stage was constructed and calibrated using seeded-stand data collected by Daniels (1978) (Table 1). Mapped-stand growth data necessary for calibrating the stand growth and development stage were not available for seeded stands. The individual tree diameter and height growth adjustment factors and the survival probability equation presented by Daniels and Burkhardt (1975) for loblolly pine plantations were used for these initial tests of Seed-PTAEDA. The volume equations used to obtain stand yield estimates are from the natural stand work of Burkhardt et al. (1972a). Input variable definitions, flow charts, and a complete program listing are included in the Appendices.

The natural stand plot data of Burkhardt et al. (1972a) were available for comparisons with simulated yields generated by Seed-PTAEDA. These data consist of stand summary information from 121 temporary plots measured in natural loblolly pine stands located in Virginia and North Carolina (Table 3).

Seed-PTAEDA was used to estimate stand characteristics for each of the 121 observed plots by using the existing stand option mentioned earlier. That is, basal area per acre was projected back in time from the observed age to age 10, when an initial stand is generated. Observed site index was used at age 10. The hardwood control parameter was estimated from observed ratios of basal area in pine to that in hardwood. Growth to the observed age was then simulated.

Early simulations indicated that simulated height and diameter growth were far exceeding observed patterns resulting in large over predictions in total cubic-foot yield and basal area. Moderate over predictions in the number of trees per acre accentuated this bias. Further analysis indicated that bias decreased with decreasing stand age and for young stands close to age 10 bias was negligible. It was concluded that the plantation-derived growth and survival relationships were not well suited for simulating the development of seeded stands. The initial stand generation stage of the model seemed to be working well.

It was thought that perhaps the relative growth patterns of individual trees, once scaled to known average growth curves, could be modeled using the plantation relationships, even if absolute growth predictions were biased. An equation to estimate average height as a function of average dominant height (from the site index curve) was developed from the natural stand data of Burkhardt et al. (1972a) and took the form

Table 3. Summary of stand conditions in 121 natural loblolly pine stands used for testing initial version of seeded stand simulator.

Variable	Mean	Range
Age	29	13 - 77
Density (stems/acre)	476	80 - 1220
Height (feet) <sup>a/</sup>	61.0	39.5 - 90.0
Total basal area (ft <sup>2</sup> /acre)	143.4	35.5 - 269.2

<sup>a/</sup> Average height of dominants and codominants.

$$\text{HAVE} = a + b \text{ HD}$$

where,

HAVE = average height of all trees

HD = average height of dominant and codominant trees

This relationship was used to scale predicted tree heights, after each growth period, so that average height conformed to that expected. Only relative growth allocations for individual trees were then obtained from the plantation equations.

Results from this refinement of the original model were more logical. Height growth was reduced to observed levels and diameter growth, determined from height growth, was also reduced. Over all 121 plots average predicted cubic-foot volume was only 4% greater than the observed average. Basal area per acre was under predicted by 6% on the average.

However, while stand aggregate measures such as total volume and basal area appeared to agree with observed values, predicted stand structure did not agree with that observed. The average predicted number of trees per acre was 27% greater than that observed, whereas average diameter was 12% less than that observed. This indicated that problems still existed in using the plantation-derived survival relationships.

It was again thought that the plantation equations provided accurate relative ratings of survival probabilities. By scaling the predicted survival probabilities downward, numbers of trees were reduced and diameter growth was increased due to decreased competition. Total stand cubic-foot yield and basal area were not greatly affected.

Data were not available to develop a prediction equation for scaling survival probabilities; the above trial was based solely on trial and error simulations. Without quantifying the scaling factor for survival relationships the model, as presented, is somewhat incomplete. Further tests were considered to be of limited usefulness without first calibrating the model.

## CALIBRATION PROCEDURES

Deficiencies in preliminary tests of Seed-PTAEDA indicated the need for detailed calibration of growth and survival relationships after the generation of the initial stand. Calibration will require further data collection specific to growth and survival of individual trees in seeded stands. Data requirements and model fitting techniques for calibration will be discussed.

Complete calibration of Seed-PTAEDA will require refitting three equations: 1) the individual tree height growth adjustment factor, 2) the diameter growth adjustment factor, and 3) the survival probability equation. All three expressions involve competition index and either crown ratio or crown length.

To fit these expressions requires a set of data from remeasured, stem mapped plots. Site index and age must be known. Individual tree measurements must include d.b.h., height, crown length, and a code indicating whether a tree is alive or dead, for at least 2 measurement years. Remeasurements should be close together in time, say one to three years, to avoid insensitivity due to averaging growth over a long period. If possible, the exact year of tree mortality should be known. Plots must be mapped to allow calculation of the competition index, and should be sufficiently large (say greater than .25 acre) to permit a buffer of trees around the interior trees for which the competition index will be calculated.

With these data one may derive the necessary variables for fitting the three equations. The model forms for the equations, as described earlier, should perform well with coefficients specific to seeded stands. The models may be fitted using any non-linear regression routine. However, the availability of new data may offer the potential user an opportunity to investigate new functional relationships, as well. Other competition indices may also be investigated for their applicability to seeded stands, once new data are available. Such modifications from the original model forms may require additional variables to be measured.

## CONCLUSIONS AND RECOMMENDATIONS

Methods have been described for constructing a detailed, flexible model of tree growth and stand development for seeded loblolly pine. The initial stand generation stage was developed and fitted specifically for seeded stands over a wide range of conditions. Preliminary results indicated that this stage of the model described young stand structure quite well. However, subsequent stand development in seeded stands was not well described when plantation-derived growth and survival relationships were used. This is not surprising since stand conditions in the data used for fitting the plantation relationships must be considered a very small subset of conditions found in seeded stands--not just in terms of spatial pattern, but also in age, stand density, site quality, and competition.

Initial attempts to improve predictive ability of the model were moderately successful, but also somewhat inadequate. Methods were used to scale the individual plantation predictions to fit average values for seeded stands. Although this technique was useful in improving predictions, and may be of further interest to some potential users as a means of calibrating the model, it suffers two main drawbacks. First, it serves to fit the model to one specific data set--in this case the test data set. Continued refinement of this type may provide a model that fits the test data set extremely well, but does not ensure flexibility elsewhere. Second, by scaling to stand averages, the model loses its appeal as an individual-tree-based growth model. In effect, after scaling factors were introduced, the model became a series of stand average prediction equations, with the individual tree growth components serving only to allocate stand variability. The computer time and expense incurred by these calculations could not be justified in this context.

As interest grows in seeded stands of loblolly, and as new data become available, it is hoped that complete calibration of the model described here will follow. The development of flexible models, which can provide information for intensive management decisions, is important. The methods described here should help in developing these models for seeded loblolly pine stands.

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## APPENDICES

Appendix I. Input variable definitions for simulation model  
Seed-PTAEDA.

---

Variable Name	Definition
TITLE	A descriptive title up to 80 characters long
NYEARS	Length of simulation in years
SITE	Site index (base age 50)
IX	Random number seed, any odd integer
ALPHA	Pielou's index of nonrandomness
TS	Loblolly pine trees surviving per acre at age 10
AGE	Age of existing stands
BA	Total basal area per acre for existing stands
HDWD	Additional proportion of (loblolly equivalent) competing stems per acre to simulate hardwood competition
IRLSE	Type of release from hardwood competition  1 = gradual release until ARLSE 2 = sudden release at ARLSE
ARLSE	Age at which site will be released from additional competing hardwoods
KIN	Age at next decision period or age of next input
ITHIN	Thinning type:  1 = corridor thinning 2 = low thinning 3 = combination of 1 and 2

---

Appendix I. Input variable definitions for simulation model  
Seed-PTAEDA (continued).

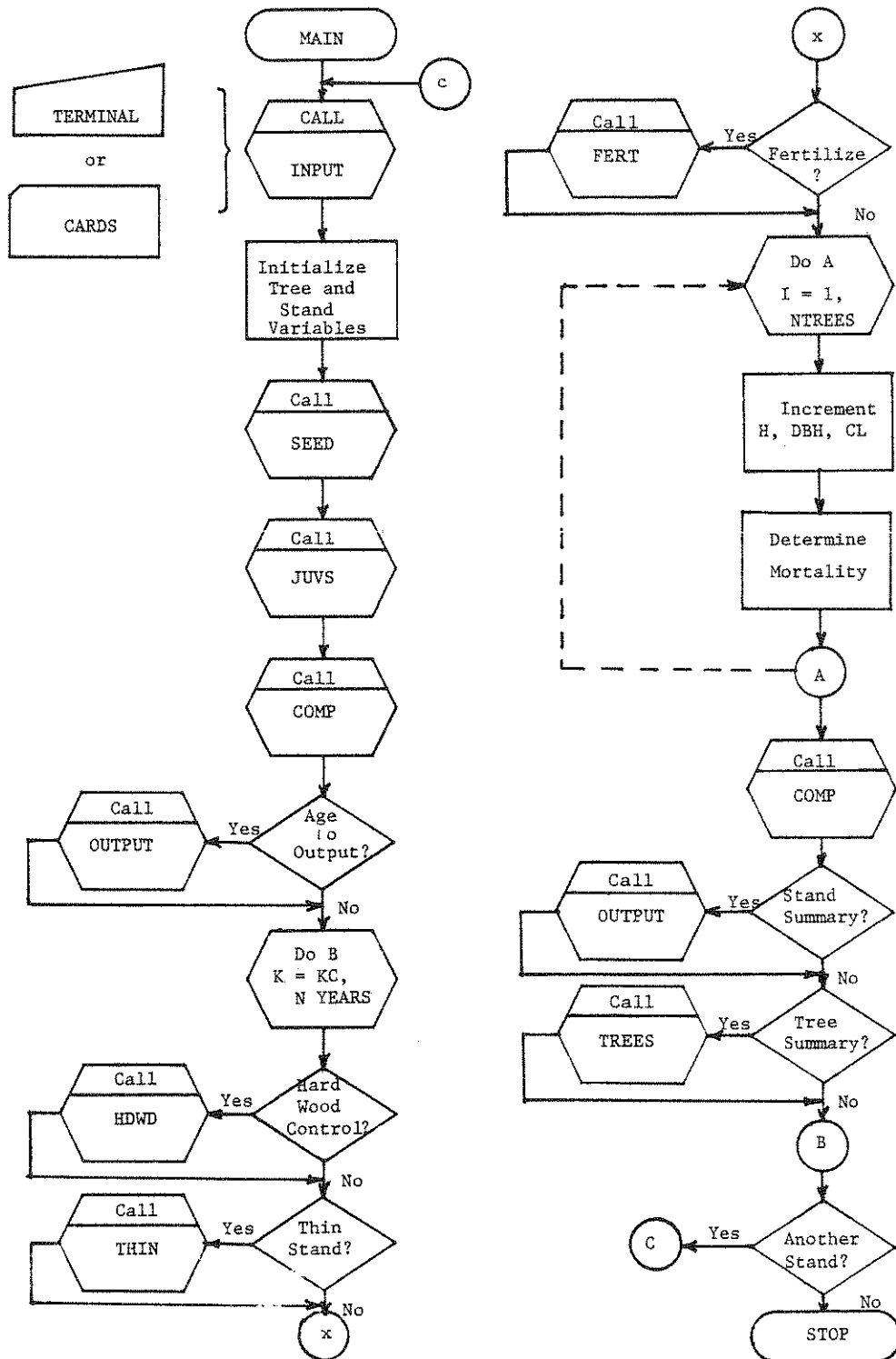
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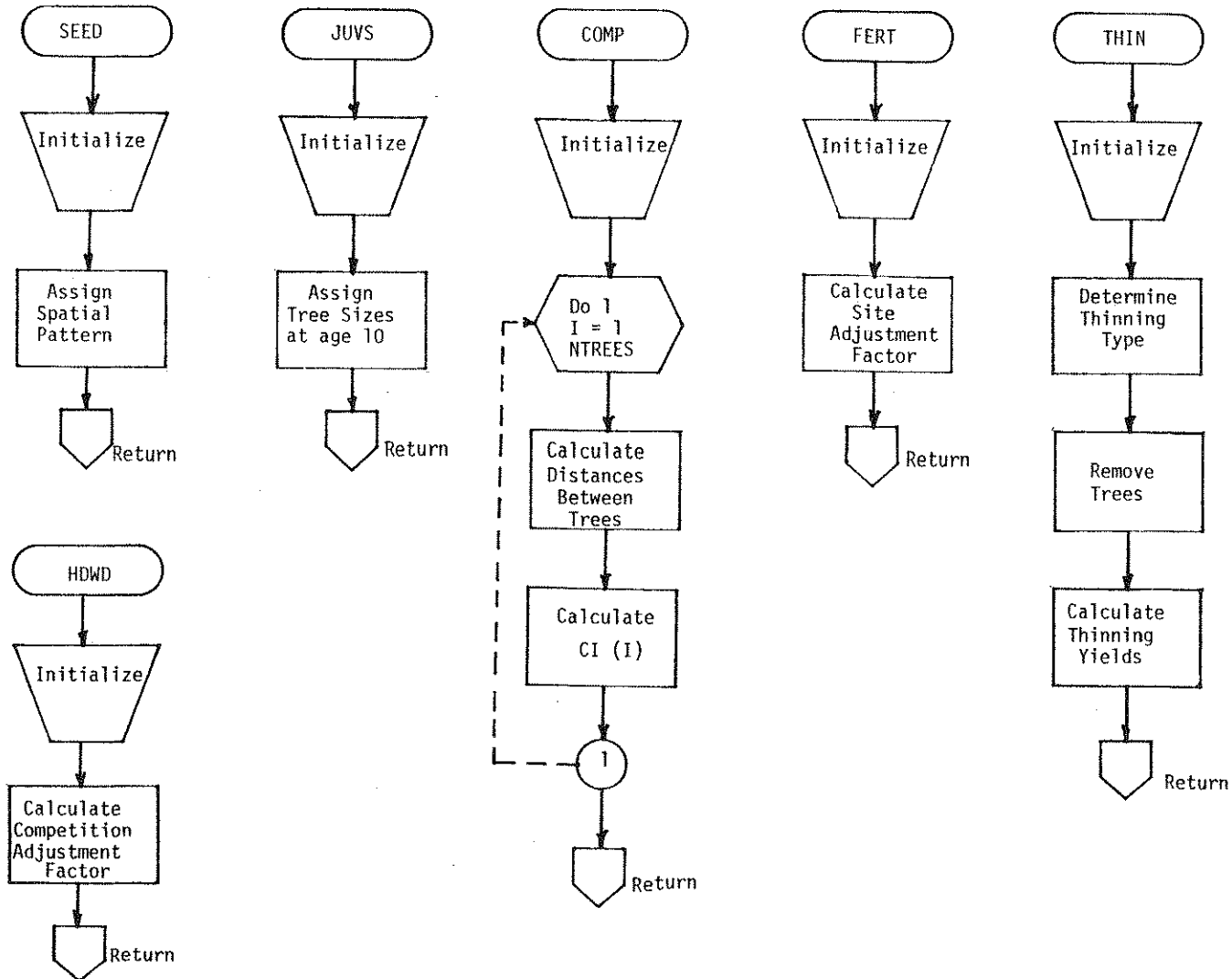
Variable Name	Definition
KTHIN	Age of growing season immediately after thinning
XCORW	Swath width in x direction
YCORW	Swath width in y direction
XCORS	Swath spacing in x direction
YCORS	Swath spacing in y direction
ILOW	Low thinning type  1 = diameter limit 2 = residual basal area limit
DLOW	Lower diameter limit below which trees will not be removed (low thinning option only)
TLIM	Thinning limit: If  ILOW = 1, upper diameter limit above which trees will not be removed ILOW = 2, residual basal to be left after thinning
KFERT	Age of growing season immediately after treatment
RESP	Maximum site index increase (feet) due to fertilization
LMR	Length of time (years) to attain RESP after initially fertilizing
LR	Total length of fertilization response
QAGAIN	To simulate another stand QAGAIN = YES

---

Appendix II. Flowchart of tree and stand growth simulation program Seed-PTAEDA.



Appendix II. Flowchart of tree and stand growth simulation program Seed-PTAEDA (continued).



Appendix III. Source listing of tree and stand growth simulation program  
Seed-PTAEDA.

```
C*****SEE00010
C
C          SEED-PTAEDA                               SEE00020
C
C          SEED-PTAEDA IS A SIMULATION MODEL OF TREE AND STAND GROWTH
C          IN MANAGED, SEEDED LOBLOLLY PINE (PINUS TAEDA L.) STANDS.   SEE00030
C
C          DEVELOPED BY RICHARD F. DANIELS, VPI&SU, 1978.             SEE00040
C
C*****SEE00050
C          DIMENSION VOL(3),S(2)                                       SEE00060
C          COMMON /BLOK1/X(100),Y(100),LMORT(100),KMORT(100),D(100),   SEE00070
C          I H(100),CL(100),CI(100),MID(100),LEDGE(9),ACRES           SEE00080
C          COMMON /BLOK3/YCUFT(75,3),YCUFTM(75,3),BA(75),KJ,K,NLIVE,   SEE00090
C          I NTHIN,HD,NOLD                                             SEE00100
C          COMMON /BLOK4/TITLE(20),NYEARS,SITE,CEXIST,EXAGE,EXBA,     SEE00110
C          I TS,TS10,KCUT,KIN,KTREE,QJUV,QAGAIN                       SEE00120
C          COMMON /BLOK5/HRDWD,CAF,ARLSE,QHDWD,IRLSE                 SEE00130
C          COMMON /BLOK6/KFERT,LMR,LR,RESP,SAF,QFERT                 SEE00140
C          COMMON /BLOK7/KTHIN,ITHIN,ILOW,DLOW,TLIM,XCOR,YCOR,XCORDS,YCORS
C          COMMON /BLOK8/PLOTX,PLOTY,ALPHA                           SEE00150
C          REAL YES/'YES'/,NO/'NO'/                                  SEE00160
C          COMMON /BLOK0/N                                           SEE00170
C          DATA S/0.77093,0.07729/                                  SEE00180
C
C          INPUT INITIAL SIMULATION CRITERIA                          SEE00190
C
C          I CALL INPUTS(IX,NC,NCARDS)                                 SEE00200
C
C          INITIALIZE TREE AND STAND VARIABLES                         SEE00210
C
C          DO 50 K=1,75                                               SEE00220
C          BA(K)=0.                                                   SEE00230
C          DO 50 L=1,3                                                SEE00240
C          YCUFT(K,L)=0.                                             SEE00250
C          50 YCUFTM(K,L)=0.                                         SEE00260
C          DO 60 I=1,N                                               SEE00270
C          D(I)=0.                                                   SEE00280
C          H(I)=0.                                                   SEE00290
C          CL(I)=0.                                                  SEE00300
C          CI(I)=0.                                                  SEE00310
C          KMORT(I)=NYEARS                                           SEE00320
C          60 LMORT(I)=1                                             SEE00330
C          KTHIN=0                                                   SEE00340
C          KOUT=0                                                     SEE00350
C          KTREE=0                                                   SEE00360
C          QFERT=NO                                                  SEE00370
C          NOLD=N                                                    SEE00380
C
C          GENERATE INITIAL STAND                                    SEE00390
C
C          CALL SEED(IX)                                             SEE00400
C          CALL JUVS(IX)                                             SEE00410
C          CALL CEMP                                                SEE00420
C          IF(QJUV.EQ.NO) GO TO 65                                   SEE00430
C          SEE00440
C          SEE00450
C          SEE00460
C          SEE00470
C          SEE00480
C          SEE00490
C          SEE00500
C          SEE00510
C          SEE00520
C          SEE00530
C          SEE00540
C          SEE00550
```

Appendix III. Source listing of tree and stand growth simulation program  
Seed-PTAEDA (continued).

```
CALL OUTPUT SEE00560
KIN=KJ+1 SEE00570
C SEE00580
C COMMENCE ANNUAL TREE GROWTH SEE00590
C SEE00600
65 KC=KJ+1 SEE00610
A=KC SEE00620
DO 200 K=KC,NYEARS SEE00630
A=K SEE00640
C SEE00650
C INPUT MANAGEMENT CRITERIA SEE00660
C SEE00670
C IF(QHDWD.EQ.YES) CALL HDWD(A) SEE00680
IF(KIN .EQ.K) CALL INPUT2 SEE00690
IF(KTHIN.EQ.K) CALL THIN(A) SEE00700
IF(QFERT.EQ.YES) CALL FERT(A) SEE00710
SI=SITE SEE00720
POTH=SI*10.**(-6.528*(1./A-.02)) SEE00730
PHIN=POTH-HD SEE00740
DO 100 I=1,N SEE00750
IF(LMORT(I)-1) 100,10,90 SEE00760
10 CR=CL(I)/H(I) SEE00770
C SEE00780
C DETERMINE TREE MORTALITY SEE00790
C SEE00800
PLIVE=1.086*CR**-.0702826*EXP(-.0281694*(CI(I)*CAF)
1 **1.177809) SEE00810
P=U(IX) SEE00820
IF(P.LT.PLIVE) GO TO 80 SEE00830
NLIVE=NLIVE-1 SEE00840
LMORT(I)=2 SEE00850
KMORT(I)=K SEE00860
GO TO 90 SEE00870
C SEE00880
C SEE00890
C COMPUTE H AND D INCREMENT ON ALL TREES SEE00900
C SEE00910
80 HRED=.54631+CR**1.66254*EXP(4.82722-1.15083*CI(I)
1 *CAF-6.66226*CR) SEE00920
R=STNORM(IX) SEE00930
HIN=PHIN*HRED SEE00940
HINMAX=1.00206*PHIN+.13462026 SEE00950
IF(HIN.GT.HINMAX) HIN=HINMAX SEE00960
PDIN=.28658336*HIN+.2094718 SEE00970
HIN=HIN+R*S(1) SEE00980
IF(HIN.LT.0.) HIN=0. SEE00990
DRED=.086524+.020178*CL(I)**1.179986*EXP(-1.320610
1 *CI(I)*CAF) SEE01000
DIN=PDIN*DRED+R*S(2) SEE01010
IF(DIN.LT.0.) DIN=0. SEE01020
C SEE01030
C SEE01040
C CALCULATE PRODUCTS SEE01050
C SEE01060
C SEE01070
D(I)=D(I)+DIN SEE01080
H(I)=H(I)+HIN SEE01090
90 L=LMORT(I) SEE01100
```

Appendix III. Source listing of tree and stand growth simulation program Seed-PTAEDA (continued).

```
DSQ=D(I)*D(I) SEE01110
IF(L.EQ.1) BA(K)=BA(K)+DSQ SEE01120
YCUFT(K,L)=YCUFT(K,L)+DSQ*H(I)*.00253+.27611 SEE01130
YCUFTM(K,L)=YCUFTM(K,L)+DSQ*H(I)*.00205-.8421 SEE01140
100 CONTINUE SEE01150
BA(K)=BA(K)*.005454/ACRES SEE01160
DO 150 L=1,3 SEE01170
YCUFT(K,L)=YCUFT(K,L)/ACRES SEE01180
YCUFTM(K,L)=YCUFTM(K,L)/ACRES SEE01190
150 CONTINUE SEE01200
C SEE01210
C DETERMINE CROWN LENGTH SEE01220
C SEE01230
T=NLIVE/ACRES SEE01240
DO 101 I=1,N SEE01250
CL(I)=0. SEE01260
IF(LMORT(I).NE.1) GO TO 101 SEE01270
CBL=H(I)**1.48535*D(I)**(-0.47173)*EXP(-1.4343+.92034E-3*BA(K) SEE01280
1 *CAF-0.10991/D(I)-3.34385/A) SEE01290
IF(H(I)-CBL-CL(I).GT.HIN) CBL=H(I)-CL(I)-HIN SEE01300
CL(I)=H(I)-CBL SEE01310
IF(CL(I).LT.0) CL(I)=0. SEE01320
101 CONTINUE SEE01330
HD=POTH SEE01340
CALL COMP SEE01350
C SEE01360
C OUTPUT STAND SUMMARY SEE01370
C SEE01380
IF(KOUT.EQ.K) CALL OUTPUT SEE01390
200 CONTINUE SEE01400
C SEE01410
C HOUSE KEEPING SEE01420
C SEE01430
CALL INPUT3 SEE01440
N=NCLD SEE01450
IF(QAGAIN.EQ.YES) GO TO 1 SEE01460
STOP SEE01470
END SEE01480
C SEE01490
C ***** SEE01500
C SEE01510
SUBROUTINE INPUTS(IX,NC,NCARDS) SEE01520
C SEE01530
C SUBROUTINE INPUT IS DIVIDED INTO 3 MAIN SUB-SECTIONS SEE01540
C DESIGNED TO PROMPT THE USER FOR AND READ INITIAL SIMULATION SEE01550
C CRITERIA, MANAGEMENT CRITERIA, AND PROGRAM CONTINUATION SEE01560
C CRITERIA. THIS SUBROUTINE IS THE ONLY ONE WHICH NEED SEE01570
C BE CHANGED FOR BATCH MODE OPERATION. SEE01580
C SEE01590
C ***** SEE01600
COMMON /BLOK4/TITLE(20),NYEARS,SITE,QEXIST,EXAGE,EXBA, SEE01610
1 TS,TS10,KOUT,KIN,KTREE,QJUV,QAGAIN SEE01620
COMMON /BLOK5/HDWD,CAF,ARLSE,QHDWD,IRLSE SEE01630
COMMON /BLOK6/KFERT,LMR,LR,RESP,SAF,QFERT SEE01640
COMMON /BLOK7/KTHIN,ITHIN,ILOW,DLOW,TLIP,XCCR,YCCR,XCORS,YCORS SEE01650
```







Appendix III. Source listing of tree and stand growth simulation program  
Seed-PTAEDA (continued).

```
DO 1130 J=1,N SEE03310
IF(J.EQ.1) GO TO 1130 SEE03320
IF(SQRT(DISTSQ(XX(I),YY(I),XX(J),YY(J))).GT.(RAD1(I)+RAD1(J))) SEE03330
1 GO TO 1130 SEE03340
CFAC=XX(I)**2+YY(I)**2-XX(J)**2-YY(J)**2 SEE03350
XFAC=2.*XX(J)-2.*XX(I) SEE03360
YFAC=2.*YY(J)-2.*YY(I) SEE03370
IF(XFAC.EQ.0.) GO TO 1050 SEE03380
IF(YFAC.EQ.0.) GO TO 1060 SEE03390
YFAC=-YFAC/XFAC SEE03400
CFAC=-CFAC/XFAC SEE03410
YSQ=YFAC**2+1 SEE03420
YVAL=(CFAC-XX(I))*2.*YFAC-2.*YY(I) SEE03430
CVAL=(CFAC-XX(I))**2+YY(I)**2-RAD1(I)**2 SEE03440
BSQ=YVAL**2 SEE03450
FOURAC=4.*YSQ*CVAL SEE03460
Z=ABS(BSQ-FOURAC) SEE03470
YROOT1=(-YVAL+SQRT(Z))/(2.*YSQ) SEE03480
YROOT2=(-YVAL-SQRT(Z))/(2.*YSQ) SEE03490
XROOT1=YFAC*YROOT1+CFAC SEE03500
XROOT2=YFAC*YROOT2+CFAC SEE03510
GO TO 1070 SEE03520
1050 IF(YFAC.EQ.0.) GO TO 1130 SEE03530
YROOT1=-CFAC/YFAC SEE03540
YROOT2=YROOT1 SEE03550
XSQ=1. SEE03560
XVAL=-2.*XX(I) SEE03570
CVAL=XX(I)**2-RAD1(I)**2+(YY(I)-YROOT1)**2 SEE03580
BSQ=XVAL**2 SEE03590
FOURAC=4.*XSQ*CVAL SEE03600
Z=ABS(BSQ-FOURAC) SEE03610
XROOT1=(-XVAL+SQRT(Z))/(2.*XSQ) SEE03620
XROOT2=(-XVAL-SQRT(Z))/(2.*XSQ) SEE03630
GO TO 1070 SEE03640
1060 XROOT1=-CFAC/XFAC SEE03650
XROOT2=XROOT1 SEE03660
YSQ=1. SEE03670
YVAL=-2.*YY(I) SEE03680
CVAL=YY(I)**2-RAD1(I)**2+(XX(I)-XROOT1)**2 SEE03690
BSQ=YVAL**2 SEE03700
FOURAC=4.*YSQ*CVAL SEE03710
Z=ABS(BSQ-FOURAC) SEE03720
YROOT1=(-YVAL+SQRT(Z))/(2.*YSQ) SEE03730
YROOT2=(-YVAL-SQRT(Z))/(2.*YSQ) SEE03740
1070 THETA1=ATAN2(YROOT1-YY(I),XROOT1-XX(I)) SEE03750
IF(THETA1.LT.0.) THETA1=THETA1+2.*PI SEE03760
THETA2=ATAN2(YROOT2-YY(I),XROOT2-XX(I)) SEE03770
IF(THETA2.LT.0.) THETA2=THETA2+2.*PI SEE03780
THMIN=THETA1 SEE03790
THMAX=THETA2 SEE03800
IF(THETA2.LT.THETA1) THMIN=THETA2 SEE03810
IF(THETA2.LT.THETA1) THMAX=THETA1 SEE03820
I1=360.*THMIN/(2.*PI) SEE03830
I2=360.*THMAX/(2.*PI) SEE03840
IF(I1.EQ.I2) GO TO 1130 SEE03850
```

Appendix III. Source listing of tree and stand growth simulation program  
Seed-PTAEDA (continued).

```
THMED=THMIN+(THMAX-THMIN)/2. SEE03860
XXX=XX(I)+RAD1(I)*COS(THMED) SEE03870
YYY=YY(I)+RAD1(I)*SIN(THMED) SEE03880
IF(SQRT(DISTSQ(XX(J),YY(J),XXX,YYY)).LE.RAD1(J)) GO TC 1110 SEE03890
IF(I1.EQ.0) GO TO 1090 SEE03900
DO 1080 K=1,I1 SEE03910
1080 IDEG(K)=0 SEE03920
1090 DO 1100 K=12,360 SEE03930
1100 IDEG(K)=0 SEE03940
GO TO 1130 SEE03950
1110 IF (I1.EQ.0) IDEG(360)=0 SEE03960
IF(I1.EQ.0) I1=I1+1 SEE03970
DO 1120 K=11,12 SEE03980
1120 IDEG(K)=0 SEE03990
1130 CONTINUE SEE04000
1140 DO 1150 K=1,360 SEE04010
XXX=XX(I)+RAD1(I)*COS(FLOAT(K)*2.*PI/360.) SEE04020
YYY=YY(I)+RAD1(I)*SIN(FLOAT(K)*2.*PI/360.) SEE04030
1150 IF (XXX.LT.0..OR.XXX.GT.PLOTX..OR.YYY.LT.0..OR.YYY.GT. SEE04040
1 PLOTY) IDEG(K)=0 SEE04050
L=0 SEE04060
DO 1160 K=1,360 SEE04070
IF(IDEG(K).EQ.0) GO TO 1160 SEE04080
L=L+1 SEE04090
IDEG(L)=IDEG(K) SEE04100
1160 CONTINUE SEE04110
1170 M=FLOAT(L)*U(IX)+1 SEE04120
IF(M.EQ.(L+1)) M=L SEE04130
IF(L.NE.0) GO TO 1174 SEE04140
C XX(I)=PLOTX*U(IX) SEE04150
C YY(I)=PLOTY*U(IX) SEE04160
C GO TO 1176 SEE04170
M=1 SEE04180
IDEG(M)=360.*U(IX) SEE04190
1174 CONTINUE SEE04200
THETA=2.*PI*IDEG(M)/360. SEE04210
X(I)=XX(I)+RAD1(I)*COS(THETA) SEE04220
Y(I)=YY(I)+RAD1(I)*SIN(THETA) SEE04230
IF(L.EQ.0) X(I)=XX(I) SEE04240
IF(L.EQ.0) Y(I)=YY(I) SEE04250
1190 CONTINUE SEE04260
RETURN SEE04270
END SEE04280
C SEE04290
C***** SEE04300
C SEE04310
SUBROUTINE JUVS(IX) SEE04320
C JUVS SEE04330
C SEE04340
C SUBROUTINE JUVS GENERATES A JUVENILE SEEDS SEE04350
C STAND AT AGE 10 FROM EXISTING STAND INFORMATION. SEE04360
C SEE04370
C***** SEE04380
DIMENSION S(2) SEE04390
COMMON /BLOK1/X(100),Y(100),LMCRT(100),KMCRT(100),D(100), SEE04400
```

Appendix III. Source listing of tree and stand growth simulation program  
Seed-PTAEDA (continued).

```
1 H(100),CL(100),CI(100),MID(100),LEDGE(9),ACRES          SEE04410
COMMON /BLOK3/YCUFT(75,3),YCUFTM(75,3),BA(75),KJ,K,NLIVE,   SEE04420
1 NTHIN,HD,NOLD                                             SEE04430
COMMON /BLOK4/TITLE(20),NYEARS,SITE,QEXIST,EXAGE,EXBA,      SEE04440
1 TS,TS10,KOUT,KIN,KTREE,QJUV,QAGAIN                      SEE04450
COMMON /BLOK5/HDWD,CAF,ARLSE,CHDWD,IRLSE                   SEE04460
COMMON /BLOK8/PLOTX,PLOTY,ALPHA                            SEE04470
REAL YES/'YES'/,NO/'NO'/                                  SEE04480
COMMON /BLOK0/N                                             SEE04490
DIMENSION DUMMY(100)                                       SEE04500
KJ=10                                                       SEE04510
A=KJ                                                         SEE04520
SI=SITE                                                      SEE04530
HD=SI*10**(-6.528*(1./A-.02))                               SEE04540
A=KJ                                                         SEE04550
DAVE= .470401+.069485*HD-.083E-5*A*(TS*CAF)+5.45478*HD/(TS*CAF) SEE04560
DMIN=-.067446+.029395*HD-.112E-5*A*(TS*CAF)+6.23266*HD/(TS*CAF) SEE04570
IF(DAVE.LE.0)DAVE=.0001                                     SEE04580
IF(DMIN.LE.0)DMIN=.0001                                    SEE04590
BHAT=A LOG(TS*.1*CAF)/A LOG(DAVE/DMIN)                     SEE04600
AHAT=(GAMMA(1.+1./BHAT)/DAVE)**BHAT                       SEE04610
ACRES=100./TS                                              SEE04620
NLIVE=N                                                     SEE04630
NMORT=0                                                      SEE04640
NTHIN=0                                                      SEE04650
120 DO 1100 I=1,N                                           SEE04660
    D(I)=DAVE                                               SEE04670
    CI(I)=0.                                                 SEE04680
    LMORT(I)=1                                              SEE04690
1100 DUMMY(I)=(-A LOG(U(I)))/AHAT)**(1./BHAT)              SEE04700
    CALL CCOMP                                              SEE04710
    NTREES=0                                                SEE04720
130 IF(NTREES.EQ.N) GO TO 145                               SEE04730
    DMAX=0.                                                  SEE04740
    CMIN=9.E9                                               SEE04750
    DO 1200 J=1,N                                           SEE04760
        IF(DUMMY(J).LE.DMAX) GO TO 140                      SEE04770
        JD=J                                                 SEE04780
        DMAX=DUMMY(J)                                       SEE04790
140 IF(CI(J).GE.CMIN) GO TO 1200                           SEE04800
    JC=J                                                     SEE04810
    CMIN=CI(J)                                              SEE04820
1200 CONTINUE                                              SEE04830
    D(JC)=DMAX                                             SEE04840
    CI(JC)=9.E9                                             SEE04850
    DUMMY(JD)=0.                                           SEE04860
    NTREES=NTREES+1                                       SEE04870
    DSQ=D(JC)*D(JC)                                        SEE04880
    BA(KJ)=BA(KJ)+DSQ                                       SEE04890
    GO TO 130                                               SEE04900
145 BA(KJ)=BA(KJ)*.005454/ACRES                             SEE04910
    HAV=0.                                                  SEE04920
    DO 1250 I=1,N                                           SEE04930
        H(I)=HD**0.32192*D(I)**0.52118*EXP[1.44287+.263276E-2*BA(KJ)*CAF SEE04940
        I      +0.07299/D(I)-1.08825/A)                     SEE04950
```

Appendix III. Source listing of tree and stand growth simulation program  
Seed-PTAEDA (continued).

```
1250 HAV=HAV+H(I) SEE04960
      HAV=HAV/N SEE04970
      HAVHAT=-1.623476+0.916285*HD SEE04980
      HRAT=HAVHAT/HAV SEE04990
      DO 1300 I=1,N SEE05000
      CI(I)=0. SEE05010
      H(I)=H(I)*HRAT SEE05020
      CBL=H(I)**1.48535*D(I)**(-0.47173)*EXP[-1.4343+.92034E-3*BA(KJ) SEE05030
      *CAF-0.10991/D(I)-3.34385/A] SEE05040
      CL(I)=H(I)-CBL SEE05050
      IF(CL(I).LT.0)CL(I)=0 SEE05060
      DSQ=D(I)*D(I) SEE05070
      IF(D(I).GE.4.55) YCUFTM(KJ,1)=YCUFTM(KJ,1)-.8421+.00205*DSQ*H(I) SEE05080
      YCUFT(KJ,1)=YCUFT(KJ,1)+.27611+.00253*DSQ*H(I) SEE05090
1300 CONTINUE SEE05100
      YCUFTM(KJ,1)=YCUFTM(KJ,1)/ACRES SEE05110
      YCUFT(KJ,1)=YCUFT(KJ,1)/ACRES SEE05120
      RETURN SEE05130
      END SEE05140
C SEE05150
C ***** SEE05160
C SUBROUTINE THIN(A) SEE05170
C SUBROUTINE THIN REMOVES TREES EITHER BY CORRIDORS OR FROM SEE05180
C BELOW. THINNING FROM BELOW MAY BE ACCOMPLISHED BY REMOVING SEE05190
C TREES BELOW A SPECIFIED DBH OR BY THINNING TO A SPECIFIED SEE05200
C RESIDUAL BASAL AREA. CORRIDOR THINNING MAY BE USED IN EITHER SEE05210
C THE X OR Y DIRECTION OR BOTH. SEE05220
C SEE05230
C SEE05240
C SEE05250
C ***** SEE05260
C COMMON /BLOK1/X(100),Y(100),LMORT(100),KMORT(100),D(100), SEE05270
C 1 H(100),CL(100),CI(100),MID(100),LEDGE(9),ACRES SEE05280
C COMMON /BLOK3/YCUFT(75,3),YCUFTM(75,3),BA(75),KJ,K,NLIVE, SEE05290
C 1 NTHIN,HD,NGLD SEE05300
C COMMON /BLOK4/TITLE(20),NYEARS,SITE,QEXIST,EXAGE,EXBA, SEE05310
C 1 TS,TS10,KOUT,KIN,KTREE,QJUV,QAGAIN SEE05320
C COMMON /BLOK7/KTHIN,ITHIN,ILW,DLW,TLIM,XCOR,YCOR,XCORS,YCORS SEE05330
C COMMON /BLOK8/PLOTX,PLOTY,ALPHA SEE05340
C COMMON /BLOK9/N SEE05350
C BATHIN=0. SEE05360
C GO TO (1,2,1),ITHIN SEE05370
C CORRIDOR THINNING SEE05380
C SEE05390
C SEE05400
C 1 IF(YCORS.LE.0) YCORS=1 SEE05410
C IF(XCORS.LE.0) XCORS=1 SEE05420
C NCORY=PLOTY/YCORS+.5 SEE05430
C NCORX=PLOTX/XCORS+.5 SEE05440
C XSTART=XCORS/2.-XCOR/2. SEE05450
C YSTART=YCORS/2.-YCOR/2. SEE05460
C DO 100 I=1,N SEE05470
C IF(LMORT(I).NE.1) GO TO 100 SEE05480
C IF(YCOR.LE.0) GO TO 97 SEE05490
C DO 96 J=1,NCORY SEE05500
```

Appendix III. Source listing of tree and stand growth simulation program  
Seed-PTAEDA (continued).

```
FJ=J
YIN=YSTART*FJ
YAX=YIN+YCOR
IF(YAX.GT.PLOTY) YAX=PLOTY
IF(Y(I).LT.YIN.OR.Y(I).GT.YAX) GO TO 96
NTHIN=NTHIN+1
NLIVE=NLIVE-1
LMORT(I)=3
KMORT(I)=KTHIN
BATHIN=BATHIN+D(I)*D(I)
GO TO 100
96 CONTINUE
97 CONTINUE
IF(XCOR.LE.0) GO TO 99
DO 98 J=1,NCORX
FJ=J
XIN=XSTART*FJ
XAX=XIN+XCOR
IF(XAX.GT.PLOTX) XAX=PLOTX
IF(X(I).LT.XIN.OR.X(I).GT.XAX) GO TO 98
NTHIN=NTHIN+1
NLIVE=NLIVE-1
LMORT(I)=3
KMORT(I)=KTHIN
BATHIN=BATHIN+D(I)*D(I)
GO TO 100
98 CONTINUE
99 CONTINUE
100 CONTINUE
IF(ITHIN.EQ.1) GO TO 3
C
C LOW THINNING
C
2 IF(ILOW.EQ.2) GO TO 22
C
C DIAMETER LIMIT OPTION
C
DO 200 I=1,N
IF(LMORT(I).NE.1) GO TO 200
IF(D(I).LT.DLOW.OR.D(I).GE.TLIM) GO TO 200
NTHIN=NTHIN+1
NLIVE=NLIVE-1
LMORT(I)=3
KMORT(I)=KTHIN
200 CONTINUE
GO TO 3
C
C BA LIMIT OPTION
C
22 BATH=(BA(K-1)-TLIM)*ACRES/.005454
DO 400 IT=1,N
IF(BATHIN.GE.BATH) GO TO 3
DMIN=9.E6
DO 300 I=1,N
IF(LMORT(I).NE.1) GO TO 300
```

Appendix III. Source listing of tree and stand growth simulation program Seed-PTAEDA (continued).

```
IF(D(I).GE.DMIN.OR.D(I).LT.DLOW) GO TO 300
DMIN=D(I)
IMIN=I
300 CONTINUE
BATHIN =BATHIN+D(IMIN)*D(IMIN)
NTHIN=NTHIN+1
NLIVE=NLIVE-1
LMORT(IMIN)=3
KMORT(IMIN)=KTHIN
400 CONTINUE
3 IF(KTHIN.NE.NYEARS-1) GO TO 4
K=K-1
DO 500 I=1,N
IF(KMORT(I).NE.K+1) GO TO 500
DSQ=D(I)*D(I)
BA(K)=BA(K)-DSQ*.005454/ACRES
YCFT=DSQ*H(I)*.00253 + .27611
YCFTM=DSQ*H(I)*.00205-.8421
YCUFT(K,1)=YCUFT(K,1)-YCFT/ACRES
YCUFT(K,3)=YCUFT(K,3)+YCFT/ACRES
YCUFTM(K,1)=YCUFTM(K,1)-YCFTM/ACRES
YCUFTM(K,3)=YCUFTM(K,3)+YCFTM/ACRES
500 CONTINUE
CALL OUTPUT
K=K+1
4 RETURN
END
C
C*****
C SUBROUTINE FERT(A)
C
C SUBROUTINE FERT SIMULATES THE EFFECTS OF
C FERTILIZATION ON SITE QUALITY BY CALCULATING A SITE
C ADJUSTMENT FACTOR (SAF) WHICH ACTS AS A MULTIPLIER OF
C SITE INDEX.
C
C*****
COMMON /BLOK4/TITLE(20),NYEARS,SITE,QEXIST,EXAGE,EXBA,
1 TS,TS10,KCUT,KIN,KTREE,QJUV,QAGAIN
COMMON /BLOK6/KFERT,LMR,LR,RESP,SAF,QFERT
REAL NO/'NO'/
IF(A-KFERT.LE.0) GO TO 50
IF(A-KFERT.GT.LMR) GO TO 20
C
C AGE LE AGE OF MAX RESPONSE (LMR)
C
SAF=RESP*(1.-(KFERT+LMR-A)/LMR)
GO TO 30
20 IF(A-KFERT.GE.LR) GO TO 40
C
C AGE GT AGE OF MAX RESPONSE (LMR)
C
SAF=RESP*(1.+(KFERT+LMR-A)/(LR-LMR))
30 SAF=(SAF+SITE)/SITE
```

Appendix III. Source listing of tree and stand growth simulation program Seed-PTAEDA (continued).

```

      GO TO 50
40 SAF=1
      QFERT=NO
50 RETURN
      END
C
C*****
C
      SUBROUTINE HOWD(A)
C
C      SUBROUTINE HOWD SIMULATES THE INCREASED
C      COMPETITION DUE TO HARDWOODS BY CALCULATING A
C      COMPETITION ADJUSTMENT FACTOR (CAF) WHICH IS USED
C      TO MULTIPLY ALL COMPETITIVE COMPONENTS OF SEED-PTAEDA.
C
C*****
      COMMON /BLOK5/HRDWD,CAF,ARLSE,QHDWD,IRLSE
      REAL NC/'NO'/
      IF(A.GE.ARLSE) GO TO 10
      IF(IRLSE.EQ.2) GO TO 20
      CAF=HRDWD*(1.-A/ARLSE)+1
      GO TO 20
10 CAF=1
      QHDWD=NC
20 RETURN
      END
C
C*****
C
      SUBROUTINE OUTPUT
C
C      SUBROUTINE OUTPUT CALCULATES AND DISPLAYS
C      SUMMARY STATISTICS FOR TREE AND STAND CHARACTERISTICS.
C
C*****
      REAL MAI(3)
      DIMENSION NDC(25,3),HDC(25,3),PROD(3),YINC(3),PA1(3),
1 BAR(4),DMIN(4),DMAX(4),SD(4)
      COMMON /BLOK1/X(100),Y(100),LMORT(100),KMORT(100),D(100),
1 H(100),CL(100),CI(100),MID(100),LEDGE(9),ACRES
      COMMON /BLOK3/YCUFT(75,3),YCUFTM(75,3),BA(75),KJ,K,NLIVE,
1 NTHIN,HD,NOLD
      COMMON /BLOK4/TITLE(20),NYEARS,SITE,QEXIST,EXAGE,EXBA,
1 TS,TS10,KOUT,KIN,KTREE,QJUV,QAGAIN
      REAL YES/'YES'/,NO/'NO'/
      COMMON /BLOK0/N
      IF(QJUV.EQ.NO) GO TO 1
      K=KJ
      QJUV=NC
1 INDEX=1
C
C      CALCULATE STAND SUMMARY STATISTICS
C
      CALL STAT(D,N,LMORT,BAR(1),DMIN(1),DMAX(1),SD(1),INDEX)
      CALL STAT(H,N,LMORT,BAR(2),DMIN(2),DMAX(2),SD(2),INDEX)

```

Appendix III. Source listing of tree and stand growth simulation program  
Seed-PTAEDA (continued).

```
CALL STAT(CI,N,LMORT,BA(3),DMIN(3),DMAX(3),SD(3),INDEX) SEE07160
CALL STAT(CI,N,LMORT,BA(4),DMIN(4),DMAX(4),SD(4),INDEX) SEE07170
INDEX=2 SEE07180
CALL STAT(D,N,LMORT,DUMP1,DMIN2,DMAX2,DUMP2,INDEX) SEE07190
MAXDC=OMAX2+.45 SEE07200
MINDC=DMIN2+.45 SEE07210
IF(MINDC.LT.1) MINDC=1 SEE07220
C SEE07230
C CALCULATE CURRENT, PERIODIC, AND MEAN ANNUAL INCREMENT SEE07240
C SEE07250
DO 100 ID=MINDC,MAXDC SEE07260
DO 100 L=1,3 SEE07270
NDC(ID,L)=0 SEE07280
100 HDC(ID,L)=0 SEE07290
DO 150 M=1,3 SEE07300
YINC(M)=9.E9 SEE07310
150 PAI(M)=9.E9 SEE07320
IF(KJ.EQ.K) GO TO 3 SEE07330
YINC(1)=BA(K)-BA(K-1) SEE07340
YINC(2)=YCUFT(K,1)-YCUFT(K-1,1) SEE07350
YINC(3)=YCUFTM(K,1)-YCUFTM(K-1,1) SEE07360
IF(K-KJ.LT.5) GO TO 3 SEE07370
PAI(1)=(BA(K)-BA(K-5))/5. SEE07380
PAI(2)=(YCUFT(K,1)-YCUFT(K-5,1))/5. SEE07390
PAI(3)=(YCUFTM(K,1)-YCUFTM(K-5,1))/5. SEE07400
3 MAI(1)=BA(K)/K SEE07410
MAI(2)=YCUFT(K,1)/K SEE07420
MAI(3)=YCUFTM(K,1)/K SEE07430
PROD(1)=BA(K) SEE07440
PROD(2)=YCUFT(K,1) SEE07450
PROD(3)=YCUFTM(K,1) SEE07460
TS=NLIVE/ACRES SEE07470
NMORT=N-NLIVE-NTHIN SEE07480
TM=NMORT/ACRES SEE07490
TT=NTHIN/ACRES SEE07500
C SEE07510
C CALCULATE DISTRIBUTION OF SIZES SEE07520
C SEE07530
DO 200 I=1,N SEE07540
L=LMORT(I) SEE07550
IF(L.EQ.0) GO TO 200 SEE07560
ID=D(I)+.45 SEE07570
IF(ID.LT.1) ID=1 SEE07580
NDC(ID,L)=NDC(ID,L)+1 SEE07590
HDC(ID,L)=HDC(ID,L)+H(I) SEE07600
200 CONTINUE SEE07610
DO 300 L=1,3 SEE07620
DO 300 ID=MINDC,MAXDC SEE07630
IF(NDC(ID,L).LE.0) GO TO 300 SEE07640
HDC(ID,L)=HDC(ID,L)/NDC(ID,L) SEE07650
NDC(ID,L)=NDC(ID,L)/ACRES+.5 SEE07660
300 CONTINUE SEE07670
C SEE07680
C DISPLAY TREE AND STAND CHARACTERISTICS SEE07690
C SEE07700
```

Appendix III. Source listing of tree and stand growth simulation program  
Seed-PTAEDA (continued).

```
WRITE(6,6100)(TITLE(M),M=1,20) SEE07710
6100 FORMAT(//' ',20A4/) SEE07720
WRITE(6,6101) K SEE07730
6101 FORMAT('O STAND SUMMARY - AGE',I3//' DIMENSION ', SEE07740
1 'MEAN ST.DEV. MIN MAX') SEE07750
WRITE(6,6102)(BAR(M),SD(M),DMIN(M),DMAX(M), M=1,4) SEE07760
6102 FORMAT(' DBH',6X,4(3X,F5.2)/' HT',5X,4(3X,F5.1)/ SEE07770
1 ' CL',5X,4(3X,F5.1)/' CI',6X,4(2X,F6.4)/) SEE07780
WRITE(6,6103) ACRES,TS10,TS,HD SEE07790
6103 FORMAT('O ACRES SIMULATED ',F10.5/' TREES PER ACRE', SEE07800
1 ' AT AGE 10',F10.0/' TREES SURVIVING PER ACRE',F10.0/ SEE07810
2 ' HEIGHT OF DOMINANT STAND',F11.1/) SEE07820
WRITE(6,6104)(PROD(M),YINC(M),PAI(M),MAI(M),M=1,3) SEE07830
6104 FORMAT('O PRODUCT YIELD INCREM PAI MAI'/ SEE07840
1 ' BASAL AREA',4X,F6.1,3(2X,F6.2)/' CUBIC FEET',3X,F6.0, SEE07850
2 3(2X,F6.1)/' MERCH VOL ',2X,F7.0,3(1X,F7.1)/) SEE07860
IF(NTHIN.LE.0) GO TO 57 SEE07870
WRITE(6,6501) YCUFT(K,3),YCUFTM(K,3) SEE07880
6501 FORMAT(' TOTAL CUBIC FEET THINNED ',F6.0/ SEE07890
1 ' MERCH VOLUME THINNED ',F6.0/) SEE07900
57 CONTINUE SEE07910
WRITE(6,6105) SEE07920
6105 FORMAT('O D CLASS #LIVE MEAN H #MORT MEAN H', SEE07930
1 ' #THIN MEAN H') SEE07940
DO 400 ID=MINDC,MAXDC SEE07950
400 WRITE(6,6106) ID,(NDC(ID,L),HDC(ID,L),L=1,3) SEE07960
6106 FORMAT(' ',I3,3(4X,I5,3X,F6.2)) SEE07970
WRITE(6,6107) TS,TM,TT SEE07980
6107 FORMAT(' TOT ',3(4X,F5.0,9X)/) SEE07990
RETURN SEE08000
END SEE08010
C SEE08020
C ***** SEE08030
C SUBROUTINE COMP SEE08040
C SEE08050
C SUBROUTINE COMP CALCULATES A MODIFIED SEE08060
C HEGYI COMPETITION INDEX ON ALL LIVE TREES IN SEE08070
C A STAND. COMPETITORS ARE FOUND BY SAMPLING SEE08080
C NEIGHBORS BASED ON THEIR SIZE AND DISTANCE AWAY SEE08090
C BY ESSENTIALLY TAKING A POINT SAMPLE AT EACH SEE08100
C SUBJECT TREE WITH A BAF=10 PRISM. SEE08110
C SEE08120
C SEE08130
C ***** SEE08140
DIMENSION JDIS(9),DIST(9),IDIS(4) SEE08150
COMMON /BLOK1/X(100),Y(100),LMORT(100),KMORT(100),D(100), SEE08160
1 H(100),CL(100),CI(100),MID(100),LEDGE(9),ACRES SEE08170
COMMON /BLOK8/PLOTX,PLOTY,ALPHA SEE08180
COMMON /BLOKD/N SEE08190
DATA PLOTR/2.75/,P1/3.14159/,JDIS/1,9,8,7,6,5,4,3,2/ SEE08200
IDIS(1)=1 SEE08210
DMAX=0 SEE08220
DO 100 I=1,N SEE08230
100 IF(D(I).GT.DMAX) DMAX=D(I) SEE08240
DISMAX=PLOTR*DMAX SEE08250
```

Appendix III. Source listing of tree and stand growth simulation program  
Seed-PTAEDA (continued).

```
DISMAY=PLOTR*DMAX                                SEF08260
DO 200 I=1,N                                      SEE08270
MID(I)=2                                          SEE08280
200 IF(X(I).GT.DISMAX.AND.X(I).LT.(PLOTX-DISMAX).AND. SEE08290
1  Y(I).GT.DISMAY.AND.Y(I).LT.(PLOTY-DISMAY)) MID(I)=1
NLESS1=N-1                                       SEE08310
DO 500 I=1,NLESS1                                SEE08320
IF(LMORT(I).NE.1) GO TO 500                      SEE08330
IPLUS1=I+1                                       SEE08340
DO 400 J=IPLUS1,N                                SEE08350
IF(LMORT(J).NE.1) GO TO 400                      SEE08360
INTIOR=MID(I)+MID(J)                             SEE08370
XDIST=X(J)-X(I)                                  SEE08380
YDIST=Y(J)-Y(I)                                  SEE08390
DIST(1)=SQRT(XDIST*XDIST+YDIST*YDIST)           SEE08400
IF(INTIOR.LT.3) GO TO 1                          SEE08410
IF(XDIST) 6,5,5                                  SEE08420
5 DIST(5)=SQRT((XDIST-PLOTX)*(XDIST-PLOTX)+    SEE08430
1  (YDIST      )*(YDIST      ))                SEE08440
IDIS(2)=5                                         SEE08450
GO TO 10                                          SEE08460
6 DIST(6)=SQRT((XDIST+PLOTX)*(XDIST+PLOTX)+    SEE08470
1  (YDIST      )*(YDIST      ))                SEE08480
IDIS(2)=6                                         SEE08490
10 IF(YDIST) 3,8,8                               SEE08500
3 DIST(3)=SQRT((XDIST      )*(XDIST      )+    SEE08510
1  (YDIST+PLOTY)*(YDIST+PLOTY))                SEE08520
IDIS(3)=3                                         SEE08530
ICODE=IDIS(2)+IDIS(3)-7                          SEE08540
GO TO (2,4,11,11,11,7,9),ICCODE                SEE08550
8 DIST(8)=SQRT((XDIST      )*(XDIST      )+    SEE08560
1  (YDIST-PLOTY)*(YDIST-PLOTY))                SEE08570
IDIS(3)=8                                         SEE08580
ICODE=IDIS(2)+IDIS(3)-7                          SEE08590
GO TO (2,4,11,11,11,7,9),ICCODE                SEE08600
2 DIST(2)=SQRT((XDIST-PLOTX)*(XDIST-PLOTX)+    SEE08610
1  (YDIST+PLOTY)*(YDIST+PLOTY))                SEE08620
IDIS(4)=2                                         SEE08630
GO TO 1                                           SEE08640
4 DIST(4)=SQRT((XDIST+PLOTX)*(XDIST+PLOTX)+    SEE08650
1  (YDIST+PLOTY)*(YDIST+PLOTY))                SEE08660
IDIS(4)=4                                         SEE08670
GO TO 1                                           SEE08680
7 DIST(7)=SQRT((XDIST-PLOTX)*(XDIST-PLOTX)+    SEE08690
1  (YDIST-PLOTY)*(YDIST-PLOTY))                SEE08700
IDIS(4)=7                                         SEE08710
GO TO 1                                           SEE08720
9 DIST(9)=SQRT((XDIST+PLOTX)*(XDIST+PLOTX)+    SEE08730
1  (YDIST-PLOTY)*(YDIST-PLOTY))                SEE08740
11 GO TO 1                                         SEE08750
1 RJI=D(J)/D(I)                                  SEE08760
RIJ=1/RJI                                         SEE08770
DO 300 L=1,4                                     SEE08780
LC=IDIS(L)                                       SEE08790
LCC=JDIS(LC)                                     SEE08800
```

Appendix III. Source listing of tree and stand growth simulation program  
Seed-PTAEDA (continued).

```
LEGE(LC)=0 SEE08810
LEGE(LCC)=0 SEE08820
IF(DIST(LC).GE.D(J)*PLOTR) GO TO 20 SEE08830
IF(LEGE(LC).EQ.0) CI(I)=CI(I)+RJI/DIST(LC) SEE08840
20 IF(DIST(LC).GE.D(I)*PLOTR) GO TO 30 SEE08850
IF(LEGE(LCC).EQ.0) CI(J)=CI(J)+RIJ/DIST(LC) SEE08860
30 IF(INTIOR.LE.3) GO TO 400 SEE08870
300 CONTINUE SEE08880
400 CONTINUE SEE08890
500 CONTINUE SEE08900
RETURN SEE08910
END SEE08920
C SEE08930
C***** SEE08940
C SUBROUTINE STAT(X,N,FLAG,XBAR,MIN,MAX,S,INDEX) SEE08950
C SEE08960
C SUBROUTINE STAT CALCULATES THE MEAN, STANDARD SEE08970
C DEVIATION AND RANGE OF INPUT VECTOR. SEE08980
C SEE08990
C SEE09000
C***** SEE09010
REAL X(N),MIN,MAX SEE09020
INTEGER FLAG(N) SEE09030
M=0 SEE09040
SUMX=0. SEE09050
SUMXSQ=0. SEE09060
MAX=0. SEE09070
MIN=1.E10 SEE09080
DO 100 I=1,N SEE09090
IF(FLAG(I).EQ.0) GO TO 100 SEE09100
IF(FLAG(I).NE.1.AND.INDEX.EQ.1) GO TO 100 SEE09110
IF(X(I).GT.MAX) MAX=X(I) SEE09120
IF(X(I).LT.MIN) MIN=X(I) SEE09130
IF(FLAG(I).NE.1) GO TO 100 SEE09140
M=M+1 SEE09150
SUMX=SUMX+X(I) SEE09160
SUMXSQ=SUMXSQ+X(I)*X(I) SEE09170
100 CONTINUE SEE09180
VAR=(SUMXSQ-SUMX*SUMX/M)/(M-1) SEE09190
S=SQRT(VAR) SEE09200
XBAR=SUMX/M SEE09210
RETURN SEE09220
END SEE09230
C SEE09240
C***** SEE09250
C SEE09260
C FUNCTION U(IX) SEE09270
C SEE09280
C GENERATES A UNIFORM(0,1) RANDOM VARIATE SEE09290
C SEE09300
C***** SEE09310
IX=IX*65539 SEE09320
U=.5+IX*.2328306E-9 SEE09330
RETURN SEE09340
END SEE09350
```

Appendix III. Source listing of tree and stand growth simulation program  
Seed-PTAEDA (continued).

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C SEE09360
C***** SEE09370
C FUNCTION STNDRM(IX) SEE09380
C SEE09390
C GENERATES A STANDARD NORMAL RANDOM VARIATE SEE09400
C SEE09410
C SEE09420
C***** SEE09430
C STNDRM=(-2*ALOG(U(IX)))*.5*CGS(6.283*U(IX)) SEE09440
C RETURN SEE09450
C END SEE09460
C SEE09470
C***** SEE09480
C BLOCK DATA SEE09490
C SEE09500
C SEE09510
C***** SEE09520
C COMMON /BLOKD/ N SEE09530
C INTEGER N/100/ SEE09540
C END SEE09550
```