MASTER

**User's Manual for FORAR, A Stand
Model for Composition and Growth of
Upland Forests of Southern Arkansas**

D. L. Mielke
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ENVIRONMENTAL SCIENCES DIVISION
Publication No. 1021

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and Growth of Upland Forests of Southern Arkansas

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ENVIRONMENTAL SCIENCES DIVISION
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ABSTRACT

MIELKE, D. L., H. H. SHUGART, and D. C. WEST. 1977. User's manual for FORAR, a stand model for composition and growth of upland forests of Southern Arkansas. ORNL/TM-5767. Oak Ridge National Laboratory, Oak Ridge, Tennessee. 44 pp.

This report is a user's manual for FORAR, a computer model simulating stand growth and composition of upland forests of south central Arkansas. The model computes: (1) the number and biomass of each tree species, and (2) the dbh, age, and species of each individual tree on a 1/12-ha circular plot. FORAR was developed from FORET (Shugart and West 1977), a modified version of the JABOWA model developed by Botkin et. al. (1972a, 1972b). The present version of FORAR considers up to 35 species and 700 trees and has been implemented on IBM 360 series computers.

Once a year trees are stochastically killed or grown and new trees are stochastically planted. Individual trees are killed by a probability function scaled according to the maximum age recorded for the appropriate species. Stocking of new seedlings and sprouts in a given year depends on the computed leaf area index of the plot and the condition of five environmental variables: mineral soil, litter layer, temperature (as expressed by degree-days), wildlife populations, and epidemics. Individual trees grow according to a species-specific optimum growth function modified by climate, available light, competition, and soil moisture. The optimum growth function is a function of the maximum known age, the maximum dbh, and the maximum height recorded for each species.

The driving variable is degree-days. Input consists of 19 general characteristics for each species. Normal output shows total biomass and leaf area on a simulated plot and, for each species, the number of trees on the plot, their dbh's, and their biomass. The model also has options to list all the trees that have died from random causes, logging, and fire.

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INTRODUCTION

Because of the relative ease of estimating the necessary input parameters and because the model retains information on both individuals and species, FORAR (FORest in ARkansas stand simulation model) is versatile enough to be used in several different types of ecological studies. Changes in species composition through succession or because of long-term perturbations can be examined and consequences of different timber and wildlife habitat management techniques can be predicted. Changes in tree species abundance (e.g. to dominance or extinction) can be determined. The model provides dynamic simulations of what is classically referred to as the structure and function of ecosystems. The usefulness of different measures of such indices as diversity, productivity, or stability can be studied.

This user's manual describes the FORAR model, which simulates an upland forest of Union County in southcentral Arkansas. The model was developed from FORET (Shugart and West 1977), a modified version of the JABOWA model used by Botkin et al. (1972a, 1972b). The logging subroutine has been expanded in FORAR. A subroutine to burn the plot and probabilities of trees dying from fire damage has been added. Soil competition is computed using basal area instead of biomass. A fifth environmental variable, species-specific epidemics, has been added to subroutine BIRTH to affect seedling establishment. In addition, variables have been added, deleted, renamed, or shifted to other subroutines to make the program more efficient.

FORAR simulates composition and growth of a forest stand on the basis of each tree greater than 1.27 cm in diameter at breast height (dbh) on a 1/12 ha circular plot. The model computes (1) the numbers and biomass of each species and (2) the dbh, age, and species of each individual tree. Once a year, trees are stochastically killed or deterministically grown and new trees are stochastically planted. Individual trees are killed by a probability function that is scaled according to the maximum age recorded for the appropriate species. The probability that a specific tree will die during a given simulated year is increased if its growth rate (expressed as diameter increment) falls below an acceptable minimum (0.1 cm/yr). Stocking of new seedlings and sprouts in a given simulated year depends on the computed leaf-area index of the plot and the condition of five environmental variables discussed below. The tree species to be stocked are chosen randomly from those species which can germinate under the existing conditions for soil, temperature, wildlife populations, and epidemics. Trees grow according to a species-specific optimum growth function that is modified by soil moisture, competition, available light, and climate. The optimum growth function for each species is a function of the maximum known age, the maximum dbh and the maximum height recorded for the species. The driving variable for the model is degree-days, which is randomly chosen at the beginning of each year from a normal distribution with appropriate mean and variance. FORAR is written in FORTRAN IV and

has been implemented on IBM 360 series computers. The present version of the model considers up to 35 tree species and up to 700 individual trees and can simulate the forest for any prespecified number of years.

PROGRAM DESCRIPTION

The deck setup of FORAR is shown in Figure 1. The control cards shown are those necessary to run the program on the ORNL IBM 360/91 computer. The program is listed in Appendix A-3. In a typical run the maximum computer core used is 270K. Computer time needed will depend on the size of the run, i.e., how many different plots are wanted, how many species are used, and how many years each plot is run. A run of three plots, each potentially having 33 species and running for 600 years, takes approximately two minutes of CPU time.

INPUT

The program currently reads in parameter values for each tree species using the format:

6A4,F6.0,F5.0,F4.3,F5.2,I1,F4.0,F2.0,F5.1,F2.0,F4.1,F4.0,5L1,I4,F3.1 .

This format can easily be changed in subroutine DATA to suit the user. Table 1 lists and describes the input variable names in the order in which they are entered in the above format. Most of the parameters, such as shade tolerance or maximum age, are relatively straight-forward and can be derived from standard textbooks of dendrology or silvics. Harlow and Harrar (1969) and Fowells (1965) were used in the present case for such parameter estimations.

DMAX and DMIN are the maximum and minimum degree-day values associated with the geographic range of each species. They are calculated (using a sine wave, see Appendix A-1) from the January and July mean temperatures at the northern and southern ends of the range of each species (e.g., Figure 2). B2 and B3 are the coefficients of the following equation relating height to diameter (in cm):

$$H = 137 + B2D - B3D^2 . \quad (1)$$

Solving for B2 and B3, using $H=H_{\max}$ and $\frac{dH}{dD} = 0$ when $D=D_{\max}$ Botkin et al. 1972a) results in:

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Fig. 1. Deck structure and job setup for program FORAR.

Table 1. Parameters used in the FORAR model. Table 2 lists the values used for each tree species. See text for explanation of variables.

<u>Parameter Name</u>	<u>Meaning</u>	<u>Field</u>
AAA	Scientific name (up to 24 letters)	xxxxxxxxxxxxxxxxxxxxxxxx
DMAX	Maximum degree-days for species range (see text)	xxxxx.
DMIN	Minimum degree-days for species range (see text)	xxxx.
B3	Derived growth parameter (see text)	.xxx
B2	Derived growth parameter (see text)	xx.xx
ITYPE	Shade tolerance 1 = tolerant 3 = intolerant	x
AGEMX	Maximum age recorded for the species	xxx.
C	Leaf area constant 1 = deciduous 2 = coniferous	x.
G	Derived Growth constant (see text)	xxx.x
STEND	Tendency to sprout Value of 0.,1.,2.,or 3.	x.
SPTMIN	The minimum dbh of a tree that will sprout	xx.x
SPTMAX	The maximum dbh of a tree that will sprout	xxx.
S(1)	Must the seed have a litter layer to germinate? value of T or F	x
S(2)	Must the seed have mineral soil to germinate? value of T or F	x
S(3)	Is the seedling susceptible to hot years? value of T or F	x
S(4)	Is the seedling a highly-preferred food for wildlife? value of T or F	x
S(5)	Is an epidemic seriously damaging the species? value of T or F	x
KTIME	Number of years after a bare plot that a species is liable to be present	xxx
SOIM	The fraction of maximum recorded dbh the species will attain on an upland site	x.x

Fig. 2. Growing degree-day limits DMAX and DMIN for the geographic range of flowering dogwood (*Cornus florida*). Isopleths for growing degree days are calculated from U.S. Geological Survey (1965) and from program DATGEN (Appendix A-1).

$$B2 = \frac{2(H_{\max} - 137)}{D_{\max}} , \quad B3 = \frac{(H_{\max} - 137)}{(D_{\max})^2} . \quad (2)$$

The growth constant G is derived from Equation (4) below and the assumption that $2/3$ of the maximum dbh of the tree is reached at $1/2$ the maximum age. G can be solved for directly (see Botkin et al. 1972b) or found by using a simple program of tree growth on an interactive computer (see Appendix A-2). The variable $KTIME$ is an attempt to handle variation in seed source availability. If there is no parent tree on the plot and if the variable $IMYR$ (the number of years that have passed since the plot had no trees and mineral soil) is greater than $KTIME$, then it is assumed there is no seed source for the species and that it can not be established by seeding. The input values used in the present version of FORAR are listed in Table 2.

The model usually starts a plot with no trees; however, subroutine $LOAD$ can be used to input any desired initial conditions. If a litter layer is wanted at the beginning of a plot, $IMYR$ should be given an initial value greater than 15.

Subroutines

The flowchart for FORAR is shown in Figure 3. There are 14 subroutines in addition to the main program; however, $KILL$, $BIRTH$, and $GROW$ are the main elements in terms of actual computation.

Whenever a random number is needed for any model computation, a call to subroutine $RANDOM$ returns a random number drawn from a flat distribution between 0 and 1. In subroutine $DATA$ the user declares the number of plots wanted, the number of tree species, the number of years each plot will be run, the interval of time between successive outputs, the annual insolation, the maximum basal area a plot can attain, and the mean for $DEGD$. Subroutines $PLOTIN$ and $INIT$ initialize variables, and $PLOTIN$ also updates and prints the plot number.

Subroutine $GAUSS$ randomly chooses a value for $DEGD$ — total degree-days for a year, based on a growing degree-day temperature of $42^{\circ}F$ — from a normal distribution. The mean and standard deviation of $DEGD$ can be calculated from data available from the National Climatic Center, Asheville, North Carolina.

Table 2. Parameter values used in the FORAR model. Scientific binomials follow Little (1971).

Species	DMAX ³	DMIN ³	B3	B2	ITYPE	AGEMX	C ⁸	G	STEND	SPTMIN	SPTMAX	S					KTIME	SOIM
<i>Acer rubrum</i>	13395	1810	0.173	52.58	1 ⁸	150 ²	1	240.2	2 ¹	12.0 ⁸	150 ¹¹	F ¹	T ¹	T	F ¹⁰	F ⁸	9999	0.5
<i>Carpinus caroliniana</i>	10947	2097	0.228	33.56	1 ²	100	1	129.9	2	6.0	70 ¹¹	F	F	F	F ¹⁰	F	9999	0.4
<i>Carya cordiformis</i>	9461	3686	0.306	69.21	3 ¹	300 ⁸	1	117.2	2 ⁵	12.0 ⁸	110 ¹¹	T ⁸	F ²	T ⁸	F ¹	F ⁸	30	0.8
<i>Carya glabra</i>	12652	3686	0.158	49.65	1 ¹	300 ⁸	1	111.3	2 ⁵	12.0 ⁸	145 ¹¹	T ¹	F ²	T ⁸	F ¹⁰	F ⁸	30	0.5
<i>Carya laciniosa</i>	8718	5081	0.318	72.46	1 ¹	300	1	123.2	2 ⁵	12.0	90 ¹¹	T	F	T	F ¹⁰	F ⁵	9999	0.4
<i>Carya ovata</i>	12652	4105	0.312	69.66	1 ¹	300 ²	1	116.5	2 ⁵	2.0 ¹	110 ¹¹	T ⁸	F ²	T ⁸	F ¹⁰	F ⁸	9999	0.7
<i>Carya texana</i>	9461	5526	0.318	57.56	1	300	1	79.6	2 ⁵	12.0	110 ¹¹	T	F	T	F ¹⁰	F ⁵	9999	0.5
<i>Carya tomentosa</i>	10947	3686	0.190	49.41	3 ¹	300 ⁸	1	98.6	2 ⁵	12.0 ²	130 ¹¹	T ¹	F ²	T ⁸	F ¹⁰	F ⁸	9999	0.8
<i>Castanea ozarkensis</i>	7756	5526	0.085	21.72	1	60	1	250.9	2 ¹²	12.0	125 ¹¹	T	F	T	F ¹⁰	T ⁵	9999	0.8
<i>Celtis laevigata</i>	12652	5526	0.336	71.26	1 ¹	350	1	112.7	2 ¹	6.0 ¹	115 ¹¹	T ¹	F	F	F ¹⁰	F	9999	0.6
<i>Cornus florida</i>	10947	3686	0.536	40.81	1 ¹	100 ⁸	1	88.7	3 ¹	12.0	35 ¹¹	F ⁸	F ¹	T ¹	F ¹⁰	F ⁸	9999	0.9
<i>Diospyros virginiana</i>	13395	5526	0.084	35.84	1 ¹	150 ⁸	1	234.1	2 ¹	12.0 ⁸	210 ⁸	F ¹	F ²	T ⁸	F ¹⁰	F ⁸	30 ⁵	0.3
<i>Fagus grandifolia</i>	10204	2097	0.151	46.20	1 ¹	400 ²	1	80.7	2 ¹	6.0 ⁸	30 ⁸	F ¹	F ²	T ⁸	F ¹⁰	F ⁸	9999	0.5
<i>Fraxinus americana</i>	10947	2414	0.080	34.43	3 ¹	300 ⁸	1	113.3	2 ¹	6.0 ¹	20	F ¹	T ²	T ⁸	F ¹⁰	F ⁸	9999	0.9
<i>Ilex opaca</i>	10947	5526	0.156	42.61	1 ²	200 ²	1	133.5	1 ⁵	6.0	35 ¹¹	F ⁸	F	F	F ¹⁰	F	9999	0.5
<i>Juglans nigra</i>	8499	3686	0.074	36.37	3 ¹	250	1	161.5	1 ¹	6.0 ¹	40	T ³	F ²	T ⁸	F ¹⁰	F ⁸	20 ⁵	0.7
<i>Juniperus virginiana</i>	10204	2966	0.236	57.75	3 ¹	300 ¹	2	107.1	0 ¹	12.0 ¹	120 ¹¹	F ¹	T ¹	F ¹	F ¹⁰	F ⁸	20 ⁵	0.9
<i>Liquidambar styraciflua</i>	10947	5526	0.191	55.20	3 ¹	300 ²	1	119.4	2 ¹	12.0 ¹	80 ¹	F ¹	F ¹	F ¹	F ¹⁰	F ⁸	9999	0.7
<i>Morus rubra</i>	13395	3686	0.061	22.15	1 ⁴	75 ⁵	1	260.6	0 ⁵	12.0 ¹	180 ¹¹	T ⁵	F ⁵	T	F ¹⁰	F	30 ⁵	0.3
<i>Nyssa sylvatica</i>	12652	3686	0.202	37.00	1 ⁵	300 ⁸	1	54.8	1 ¹	60.0 ¹	90 ¹	F ⁸	T ⁸	F ⁸	F ¹⁰	F ⁸	9999	0.7
<i>Ostrya virginiana</i>	10204	1810	0.266	49.50	1 ⁷	100	1	219.1	1	6.0	90	F	F	F	F ¹⁰	F	9999	0.4
<i>Pinus echinata</i>	9461	5526	0.290	70.75	3 ¹	300	2	128.4	2 ¹	6.0 ¹	20 ¹	F ¹	T ¹	F ¹	F ¹⁰	F	9999	1.0
<i>Pinus taeda</i>	10947	6391	0.189	60.81	3 ¹	350 ²	2	174.6	0 ¹	12.0 ¹	160 ¹¹	F ¹	T ¹	F ¹	F ¹⁰	F ⁸	9999	1.0
<i>Prunus serotina</i>	10947	3899	0.083	35.57	3 ²	250 ¹	1	139.5	2 ¹	6.0 ¹	210 ¹	F ¹	T	F ⁸	T ¹⁰	F ⁸	9999	0.7
<i>Quercus alba</i>	10204	2966	0.074	36.37	1 ¹	400 ⁹	1	100.7	2 ¹	12.0 ¹	40 ¹	T ¹	F ¹	F ¹	T ¹⁰	F ⁸	9999	0.8
<i>Quercus falcata</i>	10947	5526	0.078	33.57	1 ⁶	400	1	82.9	2 ¹	12.0 ¹	30 ¹	T ⁸	F ⁸	F ⁸	F ¹⁰	F ⁸	9999	0.9
<i>Quercus marilandica</i>	10204	5081	0.111	24.86	3 ⁵	400	1	36.3	2	12.0	40	F ⁵	F	F	F ¹⁰	F	9999	1.0
<i>Quercus shumardii</i>	10947	5081	0.199	65.19	3 ²	400	1	115.9	2 ¹	12.0 ¹	160 ¹	F ¹	F ¹	F	F ¹⁰	F	30 ⁵	0.7
<i>Quercus stellata</i>	10947	5526	0.195	47.75	3 ¹	400	1	67.9	2 ¹	12.0 ¹	40 ¹	T ¹	F ¹	F ⁸	T ¹⁰	F ⁸	9999	1.0
<i>Quercus velutina</i>	9461	3313	0.097	41.57	1 ²	200 ⁹	1	100.1	2 ¹	12.0 ¹	40 ¹	F ¹	T ¹	F ⁸	F ¹⁰	F ⁸	9999	0.8
<i>Sassafras albidum</i>	10947	3686	0.519	75.51	3 ²	200 ⁸	1	108.1	3 ¹	6.0 ¹	60 ¹	T ¹	F ¹	F ⁸	F ¹⁰	F ⁸	20 ⁵	0.8
<i>Ulmus alata</i>	10947	5526	0.230	51.81	3 ¹	125 ⁵	1	212.5	1	6.0	110	F	F ⁵	F	F ¹⁰	F	9999	0.4
<i>Ulmus americana</i>	12652	1522	0.382	39.35	3	300 ¹	1	141.8	1 ¹	6.0 ¹	240 ¹	F ¹	F ¹	T ¹	F ⁵	T ¹⁰	9999	0.4

¹From Fowells (1965)

²From Harlow and Harrar (1969)

³From Little (1971) and U.S. Geological Survey (1965)

⁴From Wigginton (1964)

⁵From Carl Amason (personal communication, 1975)

⁶From Baker (1949)

⁷From Peattie (1950)

⁸From Shugart and West (1977)

⁹From Northeastern Forest Experiment Station (1971)

¹⁰From Martin et al. (1971)

¹¹From Pardo (1973)

¹²From Moore (1960)

¹³All other values not otherwise referenced were developed during the course of this project.

Fig. 3. Flow diagram of the FORAR model.

Subroutine KILL

FORAR, like the Botkin model, assumes that individual tree death can be viewed as a stochastic process, rather than delineating all the different and complex causes of mortality. The probability that a tree will be killed in any one year is $p = 1 - (1 - \epsilon)^n$ (Botkin et al. 1972b). The model assumes that 1% of all seedlings of a species will live to reach the maximum age for that species (Figure 4). Setting p equal to 0.99 and n equal to AGEMX (the maximum known age a species can attain) results in:

$$\epsilon = 4.605/\text{AGEMX} \quad (3)$$

Once a year KILL generates a random number for every tree, and the tree is killed if the random number generated is less than ϵ .

In the present version of FORAR, there are two cases in which the kill probabilities of a tree are changed from the above statement. First, the model uses the array NOGRO to keep track of those trees that grow less than 0.1 cm dbh per year. A tree has only one chance in a hundred of surviving 10 years if such a growth rate is maintained (Botkin et al. 1972b). Accordingly, the random number generated has to be less than 0.368 to kill such a tree. Secondly, a moderate fire (described below) will "damage" those species that are susceptible to fire. In the present version of FORAR, a susceptible tree that is present on the plot when a moderate fire occurs is given a 20% chance of living 5 years (J. Warren Ranney, personal communication 1976); thus, the random number generated must be less than 0.275 to kill the tree.

Trees are also killed deterministically if there is logging or a fire. To log the circular plot, the user specifies in subroutine LOGGING the upper and lower diameter limits of each species of the trees to be logged. Subroutine CUT then removes all trees on the plot within these diameter limits. In subroutine FIRE the user has a choice of specifying a light, a moderate, or a severe burn. A light fire sets the logging limits of each species to a minimum of 0 cm and a maximum of 12.7 cm and then calls subroutine CUT to remove the trees. A severe fire removes all trees from the plot and resets the bare plot counter IMYR to zero, allowing the pioneer species with small values of KTIME to reenter the plot. A moderate fire uses subroutine CUT in the same way as a light burn to remove all the trees of species that are extremely susceptible to fire damage, all the trees less than 25.4 cm dbh of species that are moderately susceptible to fire damage, and all the trees less than 17.8 cm dbh of resistant species. The susceptibility of each species to fire damage is established in the COMPUTED GO TO statements in subroutine FIRE. The values used in FORAR were taken from Davis (1959), Fowells (1965), Harlow and Harrar (1969), or developed in the course of this project. A moderate fire also resets IMYR

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Fig. 4. Functional form for the random death process. The model assumes that 1% of all the seedlings of a species will live to reach the maximum known age. In the example shown, the maximum age of loblolly pine is 350 years.

to zero and changes the survival probabilities of those damage-susceptible trees that were large enough to survive the fire, as discussed above.

Subroutine BIRTH

BIRTH calculates the biomass already on the plot and sets the correct values for the year for the five environmental variables corresponding to the S array in Table 1. As in the FORET model (Shugart and West 1977), the soil is considered to be mineral soil if the biomass on the plot is less than 0.2 metric ton/ha and if IMYR (an additional parameter in the FORAR model) is less than 15. The soil is considered to have a litter layer if the plot biomass is greater than 0.1 metric ton/ha. It is a hot year if DEGD has a value greater than its mean. Wildlife populations are randomly chosen half the time to be large enough to consume the entire seed source of a preferred food species. In the FORAR model, an epidemic is randomly present half the time to destroy the seedlings of certain host species.

BIRTH then separates out those species capable of germinating under existing environmental conditions. For instance, we assume that a species requiring a litter layer to germinate (i.e. S(1) - .TRUE.) cannot be stocked when there is no litter layer on the ground (i.e. when the biomass on the plot is less than 0.1 metric ton/ha). From the subset of possible species, the program randomly chooses from one to three species to actually stock in a given year. A random number of seedlings between zero and eight are then "born" for each of the one to three species and randomly assigned a dbh around 1.27 cm (Tree seedlings are established 137 cm tall in the model in order to have a nonzero dbh.).

After the new seedlings are established, subroutine BIRTH calls subroutine SPROUT, which checks to see if any trees of a species capable of sprouting have died since the last year. If they have, and if their diameter was within the sprouting range of that species ($SPTMIN < dbh < SPTMAX$), a random number of sprouts between zero and three are randomly given dbh's around 0.1 cm and are added to the plot. Where a maximum of three species of seedlings can enter the plot in one pass through subroutine BIRTH, only one randomly chosen tree can sprout in one pass through subroutine SPROUT.

Subroutine BIRTH then calculates whether the biomass on the plot is less than 0.1 metric ton/ha. If it is, the program returns to the top of subroutine BIRTH to go through the process again. Before returning to MAIN, BIRTH updates the age of all the trees.

Subroutine GROW

The basic growth function used by FORAR, FORET (Shugart and West 1977), and JABOWA (Botkin et al. 1972a, 1972b) is:

$$\delta(D^2H) = R LA \left(1 - \frac{D}{D_{\max}} \frac{H}{H_{\max}}\right), \quad (4)$$

where

D = dbh,
 H = height,
 D_{max} = maximum dbh recorded for the species (Pardo 1973),
 H_{max} = maximum height recorded for the species,
 R = a constant, and
 LA = leaf area.

This equation considers the trunk of a tree to be a cylinder and assumes that the change in volume of a tree in one year is proportional to the amount of sunlight the leaves receive times a factor for maintenance of the volume of living tissue already present (Botkin et al. 1972b). The assumptions of the model are as follows (For more details, see Botkin et al. 1972b):

$$H = 137 + B_2D - B_3D^2, \quad (5)$$

$$LA \propto \text{leaf weight}, \quad (6)$$

$$\text{leaf weight} = CD^2,$$

$$\text{and } G = RC, \quad (7)$$

$$\text{where } C = \text{leaf area constant},$$

$$G = \text{a growth constant}.$$

Using the above equations in Equation (4) to solve for D, the annual increment of growth, we find:

$$\delta D = GD \frac{\left(1 - \frac{D}{D_{\max}} \frac{H}{H_{\max}}\right)}{274 + 3B_2D - 4B_3D^2} \quad (8)$$

This is the basic equation for optimum growth on a good site with no competitors. An example of this function for two species is shown in Figure 5.

FORAR modifies the annual growth equation by factors for shading, climate, competition, and soil moisture. In order to modify growth according to the amount of light each tree is receiving, subroutine GROW calculates the amount each tree is shaded, in 0.1 meter increments, by those trees on the plot that are taller. This is done using the equation:

$$AL = PHI e^{0.25SLA} , \quad (9)$$

where

AL = available light for a given tree,
 PHI = annual insolation (in appropriate units), and
 SLA = shading leaf area (Botkin et al. 1972a).

SLA is calculated using Equation (6) as the leaf biomass above a given height (Shugart and West 1977, Sollins et al. 1973). Kramer and Kozlowski (1960) give equations to relate AL to $r(AL)$, the photosynthetic rate:

$$r(AL) = 1 - e^{-4.64(AL-0.05)} \text{ for tolerant species ,} \quad (10)$$

$$r(AL) = 2.24(1 - e^{-1.136(AL-0.08)}) \text{ for intolerant species .} \quad (11)$$

The factor $r(AL)$ is then used as the shading factor multiplying the growth equation. Figure 6 shows $r(AL)$ as a function of AL for tolerant and intolerant species.

FORAR assumes a parabolic functional form for the effect of climate upon tree growth, $T(DEGD)$. Figure 7 shows the relationship between the range of the species and $T(DEGD)$. Optimum climatic effect occurs in the middle of the climatic range of the species. DEGD values at the range extremes reduce $T(DEGD)$, and thus the growth of the trees of that species, to near zero.

The crude competition factor in the model is simply:

$$S(BAR) = 1 - BAR/SOILQ , \quad (12)$$

Fig. 5. Functional form for the optimum growth equation. The slope of the curve is a function of AGEMX. The height of the curve is a function of the maximum dbh recorded for the species (Pardo 1973).

Fig. 6. Functional form relating available light AL to photosynthetic rate $r(AL)$. Equations are from Kramer and Kozlowski (1960).

Fig. 7. Functional form relating degree-days DEGD to the effect of climate on the growth of a tree species. In this example, sassafras can not grow if the degree-days are less than 3686 or greater than 10947.

where

S(BAR) = competition factor,
 BAR = basal area on the plot, and
 SOILO = maximum basal area the plot can sustain.

A plot of S(BAR) is shown in Figure 8.

The last factor modifying tree growth, soil moisture, was added to FORAR when it was found that primarily-bottomland species were comprising too much biomass on the plot. For simplicity, SOIM was assumed to be a constant factor reducing (usually) the growth of a species on an upland site. A more realistic approach would be to use moisture as another driving variable of the model. This, however, would add another level of resolution to an already complex program. Species that are found primarily on dry, poor soils have values of SOIM near 1.0; whereas species that commonly grow well in bottomlands and grow much more poorly on upland sites may have values of SOIM around 0.5.

The final growth equation for the model is thus:

$$\delta D = G D \frac{1 - \frac{D}{D_{\max}} \frac{H}{H_{\max}}}{274 + 3B2D - 4B3D^2} r(AL) T(DEGD) S(BAR) SOIM \quad (13)$$

OUTPUT

A sample output of FORAR is shown in Appendix A-4. Normal output of FORAR shows, for each species, the number of trees on the plot, their dbh's, and the biomass. Total biomass and leaf area on the plot are also given at the end of the species list. Output frequency (KPNT) is specified by the user in subroutine DATA. The model has an option to list all the trees that have died from random causes since the last printout, all the trees that have been removed by logging and the diameter limits for each species, and all the trees that have been killed by fire.

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Fig. 8. Linear function relating basal area on the 1/12 ha plot (BAR) to a general competition factor $S(\text{BAR})$. The maximum basal area the plot can maintain is from Dick Williams (personal communication 1975).

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Appendix A-1

Program DATGEN calculates annual degree-days from January and July mean temperatures. A sine function is used to estimate average daily temperature, with the minimum and maximum values equal to the January and July means, respectively. The area under the sine curve and above the 42°F line is then determined by integration and scaled to the proper value for degree-days.

In the listing shown, the degree-days are calculated and printed out for the matrix of January mean temperatures from 0° to 70° and July mean temperatures from 60° to 90°.

```

ISN 0002      DIMENSION DEGD(100,100)
ISN 0003      DO 6 I=1,71,5
ISN 0004      XI=I-1
ISN 0005      DO 4 J=60,90,5
ISN 0006      KK=1
ISN 0007      XJ=J
ISN 0008      AVE=(XJ+XI)/2.
ISN 0009      IF(XI.GE.XJ) GO TO 39
ISN 0011      R=XJ-XI
ISN 0012      R=.5*R
ISN 0013      DO 11 JJ=1,342
ISN 0014      XJJ=JJ
ISN 0015      HEAT=R*SIN(.017214*(XJJ-91.25))+AVE
ISN 0016      IF(KK.EQ.1.AND.HEAT.GE.42.) GO TO 9
ISN 0018      GO TO 10
ISN 0019      9 D1=XJJ
ISN 0020      KK=2
ISN 0021      10 IF (KK.EQ.2.AND.HEAT.LT.42.) GO TO 12
ISN 0023      GO TO 11
ISN 0024      12 D2=XJJ
ISN 0025      KK=1
ISN 0026      11 CONTINUE
ISN 0027      DEGD(I,J)=COS(.017214*(D1-91.25))-COS(.017214*(D2-91.25))
ISN 0028      DEGD(I,J)=.31831*DEGD(I,J)*(D2-D1)*R-(D2-D1)*(42.-AVE)
ISN 0029      GO TO 4
ISN 0030      39 DEGD(I,J)=10000000.
ISN 0031      4 CONTINUE
ISN 0032      6 TYPE 7,(DEGD(I,J),J=60,90,5)
ISN 0033      7 FORMAT(/,71' ',F8.0))
ISN 0034      STOP
ISN 0035      END

```

Appendix A-2

Program GFIX iteratively determines the value of G, the growth constant, for each tree species. G is calculated using the optimum growth function developed by Botkin et al. (1972a, 1972b) and the assumption that a tree will attain 2/3 of its maximum dbh at 1/2 its maximum age:

$$\delta D = G D \frac{\frac{D H}{1 - \frac{D}{D_{\max}} \frac{H}{H_{\max}}}}{274 + 3B_2D - 4B_3D^2}$$

and

$$D = \frac{2}{3} D_{\max} \text{ when } t = \text{AGEMX}/2$$

Any interactive computer system can be used to run this program. The values for DMAX, HMAX, B2, B3, and AGEMX for the species and the initial guess for G are entered in. The value of DBH typed out can then be compared to 2/3 D_{max}, and a better guess for G can be entered for the next iteration. This is repeated until DBH becomes sufficiently close to 2/3 D_{max}.

ISN 0002	INTEGER AGEMAX
ISN 0003	TYPE 1
ISN 0004	1 FORMAT (2X,'TYPE IN DMAX,HMAX,B2,B3,AGEMAX,4F10.3,I5')
ISN 0005	ACCEPT 2,DMAX,HMAX,B2,B3,AGEMAX
ISN 0006	2 FORMAT (4F10.3,I5)
ISN 0007	3 TYPE 4
ISN 0008	4 FORMAT (2X,'TYPE IN G, F10.3')
ISN 0009	ACCEPT 5,G
ISN 0010	5 FORMAT (F10.3)
ISN 0011	D=.5
ISN 0012	J=IFIX(AGEMAX/2.)
ISN 0013	DO 10 I=1,J
ISN 0014	H=137.+B2*D-B3*D**2
ISN 0015	DELD=(G*D*(1.-(D-H)/(DMAX*HMAX)))/(274.+3.*B2*D-4.*B3*D**2)
ISN 0016	D=D+DELD
ISN 0017	10 CONTINUE
ISN 0018	TYPE 11,DMAX,HMAX,B2,B3,AGEMAX,G
ISN 0019	11 FORMAT (5X,'DMAX',5X,'HMAX',7X,'B2',9X,'B3',4X, A'AGEMAX',4X,'G',/,4F10.3,I5,F10.3)
ISN 0020	TYPE 12,D
ISN 0021	12 FORMAT (6X,'DBH=',F10.4)
ISN 0022	GO TO 3
ISN 0023	STOP
ISN 0024	END

Appendix A-3

Listing of Program FORAR


```

ISN 0002 SUBROUTINE SPROUT(IX,NTREES,DBH)
ISN 0003 COMMON /SDATA/ KSPRT(35),STEND(35),SPTMIN(35),SPTMAX(35)
ISN 0004 COMMON /IRUN/ INDEX
ISN 0005 DIMENSION NTREES(35),ANEW(700),DBH(700),NAGE(700)
ISN 0006 DIMENSION NEW(35)
ISN 0007 COMMON /TRAGE/ IGES(700),IDGE(500)
C      SMALLEST STUMP SPROUT IS .1 CM ON THE AVERAGE.
ISN 0008 SIZE=.1
ISN 0009 NU=1
ISN 0010 DO 230 LL=1,INDEX
ISN 0011 L=LL
ISN 0012 IF(NTREES(L).EQ.0) GO TO 230
ISN 0013 NNU=NU+NTREES(L)-1
ISN 0014 230 NU=NU+NTREES(L)
ISN 0015 KEND=NU-1
ISN 0016 NIX=0
ISN 0017 DO 205 LL=1,INDEX
ISN 0018 L=LL
ISN 0019 IF(STEND(L).LE.0.) GO TO 205
ISN 0020 IF(KSPRT(L).GE.0) GO TO 205
ISN 0021 NIX=NIX+1
ISN 0022 NEW(NIX)=L
ISN 0023 205 CONTINUE
ISN 0024 IF (NIX.EQ.0) GO TO 74
ISN 0025 DO 2 J=1,KEND
ISN 0026 NAGE(J)=IGES(J)
ISN 0027 ANEW(J)=DBH(J)
ISN 0028 2 CONTINUE
ISN 0029 CALL RANDOM(IX,R)
ISN 0030 NIX=NIX*R+1.0
ISN 0031 N=NEW(NIX)
ISN 0032 KSUM=0
ISN 0033 DO 51 I=1,N
ISN 0034 51 KSUM=KSUM+NTREES(I)
C      STEND(I) IS THE TENDENCY FOR THE ITH SPECIES TO STUMP
C OR ROOT SPROUT. THE VALUE OF STEND(I) IS THE AVERAGE
C NUMBER OF SPROUTS (VIABLE) THAT MIGHT OCCUR WITH A GIVEN TREE DEATH.
ISN 0035 CALL RANDOM(IX,R)
ISN 0036 M=R*STEND(N)+1
ISN 0037 IF (M.EQ.0) GO TO 74
ISN 0038 DO 70 J=1,M
ISN 0039 KSUM=KSUM+1
ISN 0040 NTREES(N)=NTREES(N)+1
ISN 0041 NAGE(KSUM)=0
ISN 0042 CALL RANDOM(IX,R)
ISN 0043 ANEW(KSUM)=SIZE+0.1*(1.0-R)**3
ISN 0044 KEND=KEND+1
ISN 0045 K1=KSUM+1
ISN 0046 DO 60 II=K1,KEND
ISN 0047 I=II
ISN 0048 ANEW(I)=DBH(I-1)
ISN 0049 NAGE(I)=IGES(I-1)
ISN 0050 60 CONTINUE
ISN 0051 IF(KEND.LT.701) GO TO 61
ISN 0052 PRINT 98
ISN 0053 STOP
ISN 0054 61 CONTINUE
ISN 0055 98 FORMAT (' <<<<<<<<<LOOK OUT THE NTREES VECTOR GOT TOO BIG>>>>>>>'
ISN 0056 * '>>>>,I3A/' <<<<<<<<<TRY COMPATIBLE SPECIES AND CLIMATE>>>>>>>')
ISN 0057 DO 65 I=1,KEND
ISN 0058 IGES(I)=NAGE(I)
ISN 0059 65 DBH(I)=ANEW(I)
ISN 0060 70 CONTINUE
ISN 0061 N=0
ISN 0062 74 CONTINUE
ISN 0063 DO 9998 I=1,INDEX
ISN 0064 9998 KSPRT(I)=1
ISN 0065 RETURN
ISN 0066 END
ISN 0067

```



```

ISN 0002      SUBROUTINE CUT(DBH,NTREES,LOGS)
ISN 0003      DIMENSION DBH(700),NTREES(35),ANEW(100)
ISN 0004      COMMON /FDATA/ FARDAM(500),ALIMIT(35),BLIMIT(35),IMYR
ISN 0005      DIMENSION KAGES(100),NCUT(35),DBHCUT(100),KKAGE(100)
ISN 0006      COMMON /SDATA/ KSPRT(35),STENO(35),SPTMIN(35),SPTMAX(35)
ISN 0007      COMMON /TRAGE/ IGES(700),IGGE(500)
ISN 0008      COMMON /IRUN/ INDEX
ISN 0009      COMMON /TITLE/ AAA(35,6)
ISN 0010      LOGICAL CUTT,CUTSP,LOGS
ISN 0011      NTOT=0
ISN 0012      DO 1 I=1,INDEX
ISN 0013      NTOT=NTOT+NTREES(I)
ISN 0014      1 CONTINUE
ISN 0015      IF (NTOT.EQ.0) RETURN
ISN 0017      CUTT=.FALSE.
ISN 0018      NEW=0
ISN 0019      NFW=0
ISN 0020      KBEG=1
ISN 0021      KEND=0
ISN 0022      IF (.NOT.LOGS) GO TO 18
ISN 0024      PRINT 15
ISN 0025      15 FORMAT (12X,'DIAMETER LIMITS FOR LOGGING, BY SPECIES'/)
ISN 0026      PRINT 14,(J,ALIMIT(J),BLIMIT(J),J=1,INDEX)
ISN 0027      14 FORMAT (12X,15,2(2X,F5.1))
ISN 0028      DO 2 I=1,INDEX
ISN 0029      IF (NTREES(I).EQ.0) GO TO 2
ISN 0031      CUTSP=.FALSE.
ISN 0032      KEND=KEND+NTREES(I)
ISN 0033      DO 5 JJ=KBEG,KEND
ISN 0034      JJ=JJ
ISN 0035      IF (DBH(J).LT.ALIMIT(I).OR.DBH(J).GT.BLIMIT(I)) GO TO 6
ISN 0037      NTREES(I)=NTREES(I)-1
ISN 0038      NCUT(I)=NCUT(I)+1
ISN 0039      NFW=NFW+1
ISN 0040      DBHCUT(NFW)=DBH(J)
ISN 0041      KKAGE(NFW)=IGES(J)
ISN 0042      IF (DBH(J).GT.SPTMIN(I).AND.DBH(J).LT.SPTMAX(I)) KSPRT(I)=-1
ISN 0044      CUTT=.TRUE.
ISN 0045      CUTSP=.TRUE.
ISN 0046      GO TO 5
ISN 0047      6 NEW=NEW+1
ISN 0048      ANEW(NEW)=DBH(J)
ISN 0049      KAGES(NEW)=IGES(J)
ISN 0050      5 CONTINUE
ISN 0051      IF (CUTSP.EQ.(.FALSE.)) GO TO 72
ISN 0053      II=1
ISN 0054      IF (.NOT.LOGS) GO TO 65
ISN 0056      PRINT 70,II,NFW,(AAA(I,L),L=1,6)
ISN 0057      70 FORMAT (10X,/,2X,'SPECIES',I2,2X,I3,2X,6A4/,X,'DBH LOGS CUT'/)
ISN 0058      GO TO 74
ISN 0059      65 PRINT 71,II,NFW,(AAA(I,L),L=1,6)
ISN 0060      71 FORMAT (/,I2X,'SPECIES',I2,2X,I3,2X,6A4/,X,' DBH TREES BURNED'/)
ISN 0061      74 PRINT 29,(DBHCUT(K),K=1,NFW)
ISN 0062      29 FORMAT (20X,4F8.2)
ISN 0063      DO 80 K=1,NFW
ISN 0064      DBHCUT(K)=C.
ISN 0065      80 CONTINUE
ISN 0066      NFW=0
ISN 0067      72 KBEG=KEND+1
ISN 0068      2 CONTINUE
ISN 0069      IF (CUTT.EQ.(.FALSE.)) GO TO 13
ISN 0071      IF (NEW.EQ.0) GO TO 10
ISN 0073      GO TO 8
ISN 0074      13 PRINT 20
ISN 0075      20 FORMAT (5X,'NO TREES CUT'/)
ISN 0076      DO 7 I=1,NEW
ISN 0077      DBH(I)=ANEW(I)
ISN 0078      IGES(I)=KAGES(I)
ISN 0079      7 CONTINUE
ISN 0080      RETURN
ISN 0081      10 CONTINUE
ISN 0082      PRINT 11
ISN 0083      11 FORMAT (10X,/,10X,'ALL TREES WERE CUT'/)
ISN 0084      DO 12 J=1,700
ISN 0085      DBH(J)=0.0
ISN 0086      IGES(J)=0
ISN 0087      12 CONTINUE
ISN 0088      RETURN
ISN 0089      END

```

```

ISN 0002      SUBROUTINE OUTPUT(DBH,NTREES)
ISN 0003      COMMON /TITLE/ AAA(35,6)
ISN 0004      DIMENSION DBH(700),BDH(500),NTREES(35),IDM(500),BAR(35)
ISN 0005      COMMON /KILD/ NOGRO(100),DOBH(500),KDEAD(35),AREA,KD
ISN 0006      COMMON /TRAGE/ IGES(700),IDGE(500)
ISN 0007      COMMON /RUNNR/ NYR,INYR,KPNT,KAGE,KTIMES,KLAST
ISN 0008      COMMON /HDATA/ B2(35),B3(35),PHI,SOILQ,DEGD
ISN 0009      COMMON /IRUN/ INDEX
ISN 0010      PRINT 11
ISN 0011      11 FORMAT(///' YEAR SPEC.      NUM.      NAME      BIOMA
ZSS')
ISN 0012      PRINT 21,INYR
ISN 0013      21 FORMAT(15)
ISN 0014      96 NDEAD=0
ISN 0015      NTOT=0
ISN 0016      DO 1 I=1,INDEX
ISN 0017      NDEAD=NDEAD+KDEAD(I)
ISN 0018      1 NTOT=NTOT + NTREES(I)
ISN 0019      IF (NTOT.EQ.0) GO TO 100
ISN 0021      N=1
ISN 0022      AREA=0.0
ISN 0023      TBAR=0.0
ISN 0024      DO 20 II=1,INDEX
ISN 0025      I=II
ISN 0026      BAR(I)=0.0
ISN 0027      IF (NTREES(I).EQ.0) GO TO 20
ISN 0029      NN=N+NTREES(I) - 1
ISN 0030      N1=NTREES(I)
ISN 0031      DO 50 JJ=N,NN
ISN 0032      J=JJ
ISN 0033      50 AREA=AREA+1.9283295E-4*DBH(J)**2.129
ISN 0034      DO 97 J=1,N1
ISN 0035      N2=J+N-1
ISN 0036      BDH(J)=DBH(N2)
ISN 0037      BAR(I)=BAR(I)+.1193*DBH(N2)**2.393
ISN 0038      97 CONTINUE
ISN 0039      TBAR=TBAR+BAR(I)
ISN 0040      N2=MINO(12,N1)
ISN 0041      PRINT 2, I,NTREES(I),(AAA(I,K),K=1,6),BAR(I),(BDH(J),J=1,N2)
ISN 0042      IF (N1.GT.12) PRINT 5,(BDH(J),J=13,N1)
ISN 0044      5 FORMAT(20X, 4F8.3,/,20X,4F8.3,/,20X,4F8.3)
ISN 0045      N=N+NTREES(I)
ISN 0046      20 CONTINUE
ISN 0047      TBAR=TBAR*.012
ISN 0048      PRINT 7, NTOT,TBAR,AREA
ISN 0049      7 FORMAT(10X/10X,I7,E12.3,' METRIC TON/HA. LEAF AREA =',F 9.3)
ISN 0050      2 FORMAT(/.5X,I5,' ',I3,' ',.6A4,' ',.E9.3,' ',
Z(/.7,15X,'DBH ',4F8.3,/,20X,4F8.3,/,20X,4F8.3))
ISN 0051      GO TO 200
ISN 0052      100 PRINT 3,INYR
ISN 0053      3 FORMAT(10X,'YEAR',I5,' NO TREES LIVING')
ISN 0054      THEEND=1810.
ISN 0055      IF (DEGD.GT. THEEND) GO TO 296
ISN 0057      298 PRINT 297
ISN 0058      297 FORMAT(10X,'NONE OF THE SPECIES YOU ARE USING CAN EXIST WITH TH
*IS CLIMATE')
ISN 0059      STOP
ISN 0060      296 THEEND=13395.
ISN 0061      IF (DEGD.GT.THEEND) GO TO 298
ISN 0063      200 KD=1
ISN 0064      IF (NDEAD.EQ.0) RETURN
ISN 0066      PRINT 201
ISN 0067      201 FORMAT(' DEAD TREES SINCE LAST PRINTOUT')
ISN 0068      N=1
ISN 0069      DO 99 II=1,INDEX
ISN 0070      I=II
ISN 0071      IF (KDEAD(I).EQ.0) GO TO 99
ISN 0073      NN=N+KDEAD(I) - 1
ISN 0074      N1=KDEAD(I)
ISN 0075      DO 98 J=1,N1
ISN 0076      N2=J+N-1
ISN 0077      BDH(J)=DOBH(N2)
ISN 0078      IDM(J)=IDGE(N2)
ISN 0079      98 CONTINUE
ISN 0080      N2=MINO(7,N1)
ISN 0081      PRINT 4, I,KDEAD(I),(BDH(J),IDM(J),J=1,N2)
ISN 0082      IF (N1.GT.7) PRINT 6,(BDH(J),IDM(J),J=8,N1)
ISN 0084      6 FORMAT(27X,F9.3,I4,F9.3,I4,F9.3,I4,F9.3,I4,F9.3,I4,F9.3,
*,I4)
ISN 0085      4 FORMAT(5X,I5,I7,10X,7(F9.3,I4))
ISN 0086      N=N+KDEAD(I)
ISN 0087      99 CONTINUE
ISN 0088      DO 299 I=1,INDEX
ISN 0089      KDEAD(I)=0
ISN 0090      299 CONTINUE
ISN 0091      RETURN
ISN 0092      END

```

```

ISN 0002      SUBROUTINE GROW(DBH,NTREES)
ISN 0003      DIMENSION DBH(700),NTREES(35),PROF(1000)
ISN 0004      COMMON /IRUN/ INDEX
ISN 0005      COMMON /TDATA/ G(35),C(35),ITYPE(35),AGEMX(35),SOIM(35)
ISN 0006      COMMON /KILD/ NOGRO(100),DOBH(500),KDEAD(35),AREA,KO
ISN 0007      COMMON /HDATA/ B2(35),B3(35),PHI,SOILQ,DEGD
ISN 0008      COMMON /CLIMAT/ DMIN(35),DMAX(35)
ISN 0009      NTOT=0
C             REQUIRES EACH TREE TO ADD A 1.0 MM GROWTH RING EACH YEAR
ISN 0010      AINC=0.1
ISN 0011      DO 1 I=1,INDEX
ISN 0012      1 NTOT=NTOT+NTREES(I)
ISN 0013      IF(NTOT.EQ.0) RETURN
ISN 0015      DO 2 I=1,100
ISN 0016      2 NOGRO(I)=0
ISN 0017      DO 9998 MN=1,1000
ISN 0018      9998 PROF(MN)=0.0
ISN 0019      N2=1
ISN 0020      BAR=0.
ISN 0021      DO 8 LL=1,INDEX
ISN 0022      L=LL
ISN 0023      IF(NTREES(L).EQ.0) GO TO 8
ISN 0025      N3=N2+NTREES(L)-1
ISN 0026      DO 7 KM=N2,N3
ISN 0027      K=KM
ISN 0028      BAR=BAR+.785*(DBH(K)**2)
C
C             THE HEIGHT PROFILE IS CALCULATED IN .1 METER UNITS
ISN 0029      IHT=FIX((B2(L)*DBH(K)-B3(L)*DBH(K)**2)/10.+1.)
C
C             DATA FROM SULLINS ET AL PAGE 41. SCALED TO 1/24 MA.
C
ISN 0030      PROF(IHT)=PROF(IHT)+1.9283295E-4*DBH(K)**2.129
ISN 0031      7 CONTINUE
ISN 0032      8 N2=N2+NTREES(L)
ISN 0033      BAR=BAR/10000.
ISN 0034      DO 9997 MN=1,999
ISN 0035      MN1=1000-MN
ISN 0036      PROF(MN1)=PROF(MN1)+PROF(MN1+1)
ISN 0037      9997 CONTINUE
ISN 0038      N=1
ISN 0039      M=1
ISN 0040      DO 10 I=1,INDEX
ISN 0041      IF(NTREES(I).EQ.0) GO TO 10
ISN 0043      NN=N+NTREES(I)-1
ISN 0044      DO 9 KK=N,NN
ISN 0045      J=KK
ISN 0046      HT=B2(I)*DBH(J)-B3(I)*DBH(J)**2
C             SAME SCALE FACTOR AS ABOVE
ISN 0047      IHT=FIX(HT/10.+2.)
ISN 0048      SLA=PROF(IHT)
ISN 0049      AL=PHI*EXP(-SLA*.25)
ISN 0050      CC=(137.+0.25*B2(I)**2/B3(I))*(0.5*B2(I)/B3(I))
ISN 0051      DINC=G(I)*DBH(J)*((1.0-(137.0*DBH(J)+B2(I)*DBH(J)**2-B3(I)*DBH(J)
* )**3)/CC)/(274.+3.0*B2(I)*DBH(J)
* -4.0*B3(I)*DBH(J)**2)*(1.0-BAR/SOILQ)*SOIM(I)*4.0*(DEGD-DMIN(I))
* (DMAX(I)-DEGD)/(DMAX(I)-DMIN(I))**2
ISN 0052      IF (ITYPE(I).EQ.3) DINC=2.24*(1.0-EXP(-1.136*(AL-.08)))*DINC
C             IF (ITYPE(I).EQ.2) DINC=.5*DINC*(2.24*(1.0-EXP(-1.136*(AL-.08)))+
* (1.0-EXP(-4.64*(AL-.05))))
C             IF (ITYPE(I).EQ.1) DINC=(1.0-EXP(-4.64*(AL-.05)))*DINC
ISN 0054      IF(DINC.LT.AINC) DINC=0.0
ISN 0056      IF(DINC.GT.5.0) PRINT 6,J,I,DBH(J),DINC
ISN 0058      6 FORMAT(' DINC IS .GT. 5.0 FOR TREE',I5,' SPECIES',I3,' DBH',F6,
ISN 0060      *1,' DINC',F8.3)
ISN 0061      IF(DINC.NE.0.0) GO TO 91
ISN 0063      NOGRO(M)=J
ISN 0064      M=M+1
ISN 0065      91 DBH(J)=DBH(J)+DINC
ISN 0066      9 CONTINUE
ISN 0067      10 N=N+NTREES(I)
ISN 0068      RETURN
ISN 0069      END

```

```

SUBROUTINE BIRTH(X,NTREES,DBH)
LOGICAL SWITCH(5),S(35,5)
DIMENSION KTIME(35)
COMMON /FDATA/ FARDAM(500),ALIMIT(35),BLIMIT(35),INMYR
COMMON /RUNNR/ NYR,INYR,KPNT,KAGE,KTIMES,KLAST
COMMON /LOGBLK/ SWITCH,KTIME,S
COMMON /IRUN/ INDEX
DIMENSION NTREES(35),ANEW(700),DBH(700),NAGE(700)
DIMENSION NEW(35)
COMMON /HDATA/ IGES(700),IDGE(500)
COMMON /BZ/ BZ(35),B3(35),PHI,SOILO,DEGD
C   SMALLEST TREE IS 1.27 CM OR .5 INCHES
SIZE=1.27
ISN 0013 CALL RANDOM(I,X,R)
ISN 0014 CALL RANDOM(I,X,RAT)
ISN 0015 MMM=3.*R+1.
ISN 0016 DO 777 MMMMM=1,MMMM
ISN 0017 210 WEIGHT=0.0
ISN 0018 NU=1
ISN 0019 DO 230 LL=1,INDEX
ISN 0020 L=LL
ISN 0021 IF(NTREES(L).EQ.0) GO TO 230
ISN 0022 NNU=NU+NTREES(L)-1
ISN 0023 DO 220 MM=NU,NNU
ISN 0024 M=MM
ISN 0025 BIOM=1.9283295E-4*DBH(M)**2.129
ISN 0026 220 WEIGHT=WEIGHT+BIOM
ISN 0027 EQUATIONS FOR LEAF MASS FROM SOLLIN'S
ISN 0028 C   230 NU=NU+NTREES(L)
ISN 0029 KEND=NU-1
ISN 0030 NIX=0
ISN 0031 DO 44 JKL=1,5
ISN 0032 SWITCH(JKL)=.TRUE.
ISN 0033 C   SWITCH 1 SETS LOWER LIMIT FOR PLANTS NEEDING LITTER
ISN 0034 IF(WEIGHT.GE..1) SWITCH(1)=.FALSE.
ISN 0035 C   SWITCH 2 SETS UPPER LIMIT FOR TREES NEEDING MINERAL SOIL
ISN 0036 IF(WEIGHT.LE..2.AND.IMYR.LE.100) SWITCH(2)=.FALSE.
ISN 0037 C   SWITCH 3 ELIMINATES DROUGHT SENSITIVE SPECIES
ISN 0038 IF(DEGD.LE.8700.0) SWITCH(3)=.FALSE.
ISN 0039 CALL RANDOM(I,X,R)
ISN 0040 C   SWITCH 5 REDUCES SEEDING RATE OF DESIRABLE MAST
ISN 0041 IF(R.GE..5) SWITCH(5)=.FALSE.
ISN 0042 CALL RANDOM(I,X,R)
ISN 0043 C   SWITCH 4 ELIMINATES DEER EATEN SPECIES
ISN 0044 IF(RAT.GT..5) SWITCH(4)=.FALSE.
ISN 0045 DO 205 LL=1,INDEX
ISN 0046 L=LL
ISN 0047 DO 444 JKL=1,5
ISN 0048 444 IF(S(LL,JKL).AND.SWITCH(JKL)) GO TO 205
ISN 0049 THIS SECTION RESTRICTS THE PLANTING OF CERTAIN TREE SPECIES
ISN 0050 THAT TYPICALLY ARE ASSOCIATED WITH OLD FIELD SUCCESSION.
ISN 0051 KTIME(I) IS THE LAST YEAR SINCE A CLEARING THAT THE ITH
ISN 0052 SPECIES CAN BE EXPECTED TO SEED IN. SINCE THE LIMITING FACTOR
ISN 0053 IS POSTULATED TO BE A SEED SOURCE PROBLEM, WE ALLOW SEEDING IN
ISN 0054 IF THERE IS A PARENT TREE ON THE PLOT.
ISN 0055 IF ((IMYR.GT.KTIME(LL)).AND.(NTREES(LL).LE.0)) GO TO 205
ISN 0056 NIX=NIX+1
ISN 0057 NEW(NIX)=L
ISN 0058 205 CONTINUE
ISN 0059 IF (NIX.EQ.0) GO TO 7A
ISN 0060 DO 2 J=1,KEND
ISN 0061 NAGE(J)=IGES(J)
ISN 0062 ANEW(J)=DBH(J)
ISN 0063 2 CONTINUE
ISN 0064 CALL RANDOM(I,X,R)
ISN 0065 C   VALUE OF 5.0 ASSUMES AN AVERAGE OF 12% SEEDLING/HA.
ISN 0066 M=8.0*R
ISN 0067 CALL RANDOM(I,X,R)
ISN 0068 NIX=NIX*R+1.0
ISN 0069 N=NEW(NIX)
ISN 0070 KSUM=0
ISN 0071 DO 51 I=1,N
ISN 0072 51 KSUM=KSUM+NTREES(I)
ISN 0073 DO 70 J=1,M
ISN 0074 KSUM=KSUM+1
ISN 0075 NTREES(N)=NTREES(N)+1
ISN 0076 NAGE(KSUM)=0
ISN 0077 CALL RANDOM(I,X,R)
ISN 0078 ANEW(KSUM)=SIZE*0.3*(1.0-R)**3
ISN 0079 KEND=KEND+1
ISN 0080 K1=KSUM+1
ISN 0081 DO 60 II=K1,KEND
ISN 0082 I=II
ISN 0083 ANEW(I)=DBH(I-1)
ISN 0084 NAGE(I)=IGES(I-1)
ISN 0085 60 CONTINUE
ISN 0086 IF(KEND.LT.701) GO TO 61
ISN 0087 PRINT 98
ISN 0088 STOP
ISN 0089 61 CONTINUE
ISN 0090 98 FORMAT (' <<<<<<<<<<LOOK OUT THE NTREES VECTOR GOT TOO BIG>>>>>>>'
ISN 0091 *),I3IA/' <<<<<<<<<<TRY COMPATIBLE SPECIES AND CLIMATE>>>>>>>')

```

```

ISN 0089      DO 65 I=1,KEND
ISN 0090      IGES(I)=NAGE(I)
ISN 0091      65 DBH(I)=ANEW(I)
ISN 0092      70 CONTINUE
ISN 0093      N=0
ISN 0094      74 CONTINUE
ISN 0095      CALL SPROUT(IX,NTREES,DBH)
ISN 0096      IF(WEIGHT.LT.0.2) GO TO 210
ISN 0098      DO 100 I=1,KEND
ISN 0099      IGES(I)=IGES(I)+1
ISN 0100      100 CONTINUE
ISN 0101      777 CONTINUE
ISN 0102      RETURN
ISN 0103      END

```

```

ISN 0002      SUBROUTINE DATA
ISN 0003      COMMON /TITLE/ AAA(35,6)
ISN 0004      LOGICAL SWITCH(5),S(35,5)
ISN 0005      DIMENSION KTIME(35)
ISN 0006      COMMON /LOGBLK/ SWITCH,KTIME,S
ISN 0007      COMMON /SDATA/ KSPRT(35),SIENU(35),SPTMIN(35),SPTMAX(35)
ISN 0008      COMMON /IRUN/ INDEX
ISN 0009      COMMON /TDATA/ G(35),C(35),ITYPE(35),AGEMX(35),SOIM(35)
ISN 0010      COMMON /RUNNR/ NYR,INYR,KPNT,KAGE,KTIMES,KLAST
ISN 0011      COMMON /HDATA/ B2(35),B3(35),PHI,SOILO,DEGD
ISN 0012      COMMON /CLIMAT/ DMIN(35),DMAX(35)
ISN 0013      INDEX=33
ISN 0014      IX=76123
ISN 0015      KTIMES=1
ISN 0016      KLAST=3
ISN 0017      NYR=500
ISN 0018      KPNT=50
ISN 0019      KAGE=-1
ISN 0020      PHI=1.0
ISN 0021      SOILO=3.83
ISN 0022      DEGD=8700.
ISN 0023      C VALUE FOR SOILO FROM MAX VALUE CITED BY DICK WILLIAMS. GP FORESTER
ISN 0024      DO 23 J=1,INDEX
ISN 0025      23 READ(5,24)(AAA(J,II),II=1,6),DMAX(J),DMIN(J),B3(J),B2(J),ITYPE(J),
ISN 0026      AAGEMX(J),C(J),G(J),STEND(J),SPTMIN(J),SPTMAX(J),
ISN 0027      B(S(J,JI),JI=1,5),KTIME(J),SOIM(J)
ISN 0028      24 FORMAT(6A4,F6.0,F5.0,F4.3,F5.2,I1,F4.0,F2.0,F5.1,F2.0,F4.1,F4.0,
ISN 0029      A5L1,I4,F3.1)
ISN 0030      30 RETURN
ISN 0031      END

```

```
ISN 0002      SUBROUTINE RANDOM(IX,Q)  
ISN 0003      R=FLTRN(IX)  
ISN 0004      RETURN  
ISN 0005      END
```

```
ISN 0002      SUBROUTINE GAUSS(IX,S,AM,DEGD)  
ISN 0003      Z=0.0  
ISN 0004      DO 5 I=1,12  
ISN 0005      5 Z=Z+FLTRN(DUM)  
ISN 0006      Z=Z-6.  
ISN 0007      DEGD=AM+S*Z  
ISN 0008      RETURN  
ISN 0009      END
```

```

ISN 0002      SUBROUTINE INIT(DBH,NTREES,DBHK,NTREEK,JJ)
ISN 0003      COMMON /IRUN/ INDEX
ISN 0004      DIMENSION DBH(1),NTREES(1),DBHK(1),NTREEK(1)
ISN 0005      COMMON /SDATA/ KSPRT(35),STEND(35),SPTMIN(35),SPTMAX(35)
ISN 0006      COMMON /KILD/ NOGRO(100),DDBH(500),KDEAD(35),AREA,KD
ISN 0007      COMMON /TRAGE/ IGES(700),IDGE(500)
ISN 0008      DO 10 J=1,500
ISN 0009      DDBH(J)=0.0
ISN 0010      IDGE(J)=0
ISN 0011      10 CONTINUE
ISN 0012      DO 11 J=1,100
ISN 0013      11 NOGRO(J)=0
ISN 0014      DO 13 J=1,INDEX
ISN 0015      13 KSPRT(J)=1
ISN 0016      IF (JJ.GT.1) GO TO 1
ISN 0018      DO 2 J=1,INDEX
ISN 0019      2 NTREEK(J)=NTREES(J)
ISN 0020      DO 3 K=1,100
ISN 0021      3 DBHK(K)=DBH(K)
ISN 0022      RETURN
ISN 0023      1 CONTINUE
ISN 0024      DO 4 J=1,INDEX
ISN 0025      4 NTREES(J)=NTREEK(J)
ISN 0026      DO 5 K=1,100
ISN 0027      5 DBHK(K)=DBHK(K)
ISN 0028      RETURN
ISN 0029      END

```

```

ISN 0002      SUBROUTINE PLOTIN(IPLLOT,DBH,NTREES)
ISN 0003      COMMON /TRAGE/ IGES(700),IDGE(500)
ISN 0004      COMMON /KILD/ NOGRO(100),DDBH(500),KDEAD(35),AREA,KD
ISN 0005      DIMENSION DBH(700),NTREES(35)
ISN 0006      DO 1 J=1,35
ISN 0007      KDEAD(J)=0
ISN 0008      DBH(J)=0.0
ISN 0009      IGES(J)=0
ISN 0010      1 NTREES(J)=0
ISN 0011      DO 2 J=36,700
ISN 0012      IGES(J)=0
ISN 0013      2 DBH(J)=0.0
ISN 0014      IPLLOT=IPLLOT+1
ISN 0015      PRINT 13,IPLLOT
ISN 0016      13 FORMAT (/, ' PLOT NUMBER ',I4)
ISN 0017      RETURN
ISN 0018      END

```

```

ISN 0002      SUBROUTINE KILL(DBH,NTREES,IX)
ISN 0003      DIMENSION DBH(700),NTREES(35)
ISN 0004      COMMON /FDATA/ FARDAM(500),ALIMIT(35),BLIMIT(35),IMYR
ISN 0005      COMMON /IRUN/ INDEX
ISN 0006      COMMON /SDATA/ KSPRT(35),STEND(35),SPTMIN(35),SPTMAX(35)
ISN 0007      COMMON /TDATA/ G(35),C(35),ITYPE(35),AGEMX(35),SOIM(35)
ISN 0008      COMMON /KILD/ NOGRO(100),DDBH(500),KDEAD(35),AREA,KD
ISN 0009      COMMON /TRAGE/ IGES(700),IDGE(500)
ISN 0010      COMMON /RUNNR/ NYR,INYR,KPNT,KAGE,KTIMES,KLAST
ISN 0011      ID = 1
ISN 0012      IDF=0
ISN 0013      N = 0
ISN 0014      NB=0
ISN 0015      DO 100 J=1,INDEX
ISN 0016      NB=NB+KDEAD(J)
ISN 0017      IF(NTREES(J).EQ.0) GO TO 100
ISN 0019      NA = N + 1
ISN 0020      NN = NTREES(J) + N
ISN 0021      DO 80 II=NA,NN
ISN 0022      I=II
ISN 0023      CALL RANDOM(IX,R1)
ISN 0024      IF (FARDAM(IDF).EQ.I) GO TO 42
ISN 0026      IF(R1.LE.(4.605/AGEMX(J))) GO TO 50
ISN 0028      GO TO 44
ISN 0029      42 IDF=IDF+1
ISN 0030      IF (R1.LE..275) GO TO 50
ISN 0032      44 IF(NOGRO(ID).EQ.I) GO TO 40
ISN 0034      GO TO 80
ISN 0035      40 ID=ID + 1
ISN 0036      CALL RANDOM(IX,R2)
ISN 0037      IF(R2.GT.0.368) GO TO 80
ISN 0039      50 IF(KAGE.LT.0) GO TO 70
ISN 0041      KDEAD(J)=KDEAD(J)+1
ISN 0042      NB=NB+1
ISN 0043      NUP=KD-NB
ISN 0044      IF(NUP.EQ.0) GO TO 56
ISN 0046      DO 55 KK=1,NUP
ISN 0047      K=KK
ISN 0048      IJ=KD-K+1
ISN 0049      IF(IJ.GT.500) GO TO 99
ISN 0051      DDBH(IJ)=DDBH(IJ-1)
ISN 0052      IDGE(IJ)=IDGE(IJ-1)
ISN 0053      55 CONTINUE
ISN 0054      56 DDBH(NB)=DBH(I)
ISN 0055      IDGE(NB)=IGES(I)
ISN 0056      KD=KD+1
ISN 0057      GO TO 70
ISN 0058      99 PRINT 98
ISN 0059      98 FORMAT(' DEAD TREES VECTOR FILLED. RERUN WITH SMALLER PERIOD BE
          *TWEEN PRINTOUTS (KPNT)')
ISN 0060      KAGE=-1
ISN 0061      70 NTREES(J)=NTREES(J)-1
          C THIS IS THE SWITCH TO STUMP SPROUT TREES.
          IF(DBH(I).GT.SPTMIN(J).AND.DBH(I).LT.SPTMAX(J)) KSPRT(J)=-1
ISN 0062      DBH(I)=-1.0
ISN 0064      80 CONTINUE
ISN 0065      N = NN
ISN 0066      100 CONTINUE
ISN 0067      DO 200 II=1,700
ISN 0068      I=II
ISN 0069      IF(DBH(I).EQ.0.0) RETURN
ISN 0070      IF(DBH(I).GT.0.0) GO TO 200
ISN 0072      110 DO 120 KK = 1,699
ISN 0074      K=KK
ISN 0075      DBH(K) = DBH(K+1)
ISN 0076      IGES(K)=IGES(K+1)
ISN 0077      120 CONTINUE
ISN 0078      DBH(700) = 0.0
ISN 0079      IF(DBH(I).LT.0.0) GO TO 110
ISN 0080      200 CONTINUE
ISN 0082      RETURN
ISN 0083      END
ISN 0084

```



```

ISN 0002      SUBROUTINE FIRE (DBH,NTREES)
ISN 0003      DIMENSION DBH(700),NTREES(35)
ISN 0004      COMMON /FDATA/ FARDAM(500),ALIMIT(35),BLIMIT(35),IMYR
ISN 0005      COMMON /SDATA/ KSPRT(35),STEND(35),SPTMIN(35),SPTMAX(35)
ISN 0006      COMMON /IRUN/ INDEX
ISN 0007      LOGICAL LOGS
C
ISN 0008      R=0.
ISN 0009      LOGS=.FALSE.
ISN 0010      IF (R.GE..828) GO TO 20
ISN 0011      PRINT 12
ISN 0012      12 FORMAT (5X, 'LIGHT FIRE'/)
ISN 0013      DO 15 J=1,INDEX
ISN 0014      ALIMIT(J)=0.
ISN 0015      BLIMIT(J)=12.7
ISN 0016      15 CONTINUE
ISN 0017      CALL CUT (DBH,NTREES,LOGS)
ISN 0018      RETURN
ISN 0019      20 IF (R.GE..966) GO TO 30
ISN 0020      PRINT 21
ISN 0021      21 FORMAT (5X, ' MEDIUM FIRE'/)
ISN 0022      DO 25 J=1,INDEX
ISN 0023      ALIMIT(J)=0.
ISN 0024      GO TO (22,22,23,23,22,22,23,22,23,22,22,24,22,23,22,24,22,23,22,23
ISN 0025      A,22,24,24,22,23,23,23,24,24,23,23,23),J
ISN 0026      22 BLIMIT(J)=500.
ISN 0027      GO TO 25
ISN 0028      23 BLIMIT(J)=25.4
ISN 0029      GO TO 25
ISN 0030      24 BLIMIT(J)=17.8
ISN 0031      25 CONTINUE
ISN 0032      CALL CUT (DBH,NTREES,LOGS)
ISN 0033      N=0
ISN 0034      IJ=0
ISN 0035      DO 42 J=1,INDEX
ISN 0036      NA=N+1
ISN 0037      NN=N+NTREES(J)
ISN 0038      GO TO (35,35,40,40,35,35,40,35,40,35,35,35,35,40,35,35,40,35,40
ISN 0039      A,35,35,35,35,40,40,40,40,35,35,40,40,40),J
ISN 0040      40 DO 35 JJ=NA,NN
ISN 0041      IJ=IJ+1
ISN 0042      FARDAM(IJ)=JJ
ISN 0043      35 CONTINUE
ISN 0044      N=NN
ISN 0045      42 CONTINUE
ISN 0046      IMYR=C
ISN 0047      RETURN
ISN 0048      30 DO 36 J=1,INDEX
ISN 0049      IF (NTREES(J).EQ.0) GO TO 36
ISN 0050      KSPRT(J)=-1
ISN 0051      NTREES(J)=0
ISN 0052      36 CONTINUE
ISN 0053      DO 37 J=1,700
ISN 0054      DBH(J)=0.
ISN 0055      37 CONTINUE
ISN 0056      PRINT 38
ISN 0057      38 FORMAT (5X, 'SEVERE FIRE--ALL TREES WERE BURNED'/)
ISN 0058      RETURN
ISN 0059      FNN

```

Appendix A-4

Sample Output of Program FORAR

SAMPLE OUTPUT OF PROGRAM FORAR

PLOT NUMBER 1					
YEAR SPEC.	NUM.	NAME			BIOMASS
0	YEAR	0 NO TREES LIVING			
YEAR SPEC.	NUM.	NAME			BIOMASS
50					
5	3	CARYA LACINIOSA			0.674E 00
	DBH	1.270	1.364	1.271	
6	3	CARYA OVATA			0.251E 02
	DBH	9.260	1.472	1.209	
11	2	CORNUS FLORIDA			0.506E 02
	DBH	9.385	9.378		
12	5	DIOSPYROS VIRGINIANA			0.182E 03
	DBH	10.426	10.307	11.507	11.528
		10.728			
13	6	FAGUS GRANDIFOLIA			0.144E 01
	DBH	1.383	1.270	1.276	1.288
		1.362	1.451		
14	7	FRAXINUS AMERICANA			0.497E 03
	DBH	14.376	14.365	16.780	14.526
		16.222	15.875	0.138	
15	3	ILEX OPACA			0.303E-02
	DBH	0.118	0.172	0.106	
18	11	LIQUIDAMBAR STYRACIFLUA			0.450E 03
	DBH	12.194	13.847	12.539	13.211
		12.277	12.960	13.946	12.633
		1.388	1.270	1.427	
19	1	MORUS RUBRA			0.963E 01
	DBH	6.265			
23	13	PINUS TAEDA			0.692E 04
	DBH	30.851	31.511	39.092	29.708
		35.720	38.035	37.236	34.367
		31.520	33.687	29.159	29.154
		32.003			
24	8	PRUNUS SEROTINA			0.752E 03
	DBH	17.908	17.181	17.240	16.655
		16.766	17.733	16.634	0.168
25	3	QUERCUS ALBA			0.708E 00
	DBH	1.430	1.270	1.270	
26	17	QUERCUS FALCATA			0.531E 03
	DBH	11.952	11.824	12.001	12.693
		12.748	12.252	11.821	12.005
		12.555	12.221	11.820	4.430
		5.289	1.451	1.277	1.321
		1.273			
27	1	QUERCUS MARILANDICA			0.226E 01
	DBH	3.418			
28	7	QUERCUS SHUMARDII			0.170E 03
	DBH	10.143	11.432	10.429	10.143
		10.708	1.371	1.458	
31	8	SASSAFRAS ALBIDUM			0.224E 02
	DBH	8.677	1.503	1.301	0.113
		0.192	1.437	1.389	1.399

32 19 ULMUS ALATA 0.628E 03

DBH 14.744 13.260 12.779 13.492
12.770 12.772 15.100 14.663
14.269 12.770 0.100 1.289
1.287 1.409 0.126 1.270
1.528 1.298 1.270

33 8 ULMUS AMERICANA 0.560E 02

DBH 6.175 6.507 6.619 6.079
5.771 5.768 0.102 1.570

125 0.124E 03 METRIC TON/HA. LEAF AREA = 7.171

YEAR SPEC. 100	NUM.	NAME	BIOMASS
10	3	CELTIS LAEVI GATA	0.773E 00
	DBH	1.271 1.368 1.487	
18	2	LIQUIDAMBAR STYRACIFLUA	0.460E 00
	DBH	1.271 1.359	
21	8	OSTRYA VIRGINIANA	0.191E 01
	DBH	1.316 1.272 1.287 1.392 1.519 1.281 1.299 1.305	
23	6	PINUS TAEDA	0.117E 05
	DBH	63.029 70.297 57.756 46.638 53.833 50.296	
26	5	QUERCUS FALCATA	0.575E 03
	DBH	18.933 19.034 20.430 19.059 1.304	
27	2	QUERCUS MARILANDICA	0.553E 00
	DBH	1.517 1.316	
32	1	ULMUS ALATA	0.211E 00
	DBH	1.270	
27	0.148E 03 METRIC TON/HA. LEAF AREA = 6.904		

YEAR SPEC. 150	NUM.	NAME	BIOMASS
2	1	CARPINUS CAROLINIANA	0.323E 00
	DBH	1.516	
5	1	CARYA LACINIOSA	0.231E 00
	DBH	1.317	
18	1	LIQUIDAMBAR STYRACIFLUA	0.214E 00
	DBH	1.276	
21	3	OSTRYA VIRGINIANA	0.768E 00
	DBH	1.399 1.288 1.435	
23	3	PINUS TAEDA	0.130E 05
	DBH	102.195 58.679 71.723	
25	6	QUERCUS ALBA	0.152E 01
	DBH	1.275 1.409 1.308 1.436 1.492 1.270	
26	11	QUERCUS FALCATA	0.391E 03
	DBH	29.130 3.746 4.503 3.188 0.116 0.110 1.516 1.323 1.291 1.281 1.316	
32	5	ULMUS ALATA	0.142E 01
	DBH	1.284 1.550 1.479 1.541 1.304	
33	2	ULMUS AMERICANA	0.472E 00
	DBH	1.367 1.292	
33	0.161E 03 METRIC TON/HA. LEAF AREA = 6.774		

YEAR SPEC. 200	NUM.	NAME	BIOMASS
2	2	CARPINUS CAROLINIANA	0.534E 00
	DBH	1.502 1.286	
7	2	CARYA TEXANA	0.442E 00
	DBH	1.316 1.270	
10	7	CELTIS LAEVIGATA	0.168E 01
	DBH	1.270 1.327 1.288 1.465 1.289 1.439 1.270	
21	1	OSTRYA VIRGINIANA	0.264E 00
	DBH	1.394	
23	2	PINUS TAEDA	0.157E 05
	DBH	125.903 69.541	
26	10	QUERCUS FALCATA	0.739E 03
	DBH	38.387 1.508 1.273 1.270 1.270 1.302 1.286 1.272 1.516 1.289	
27	11	QUERCUS MARILANDICA	0.274E 01
	DBH	1.325 1.270 1.393 1.276 1.542 1.270 1.540 1.285 1.271 1.350 1.389	
29	7	QUERCUS STELLATA	0.174E 01
	DBH	1.281 1.454 1.271 1.303 1.488 1.314 1.391	
32	4	ULMUS ALATA	0.902E 00
	DBH	1.331 1.270 1.303 1.313	
46		0.197E 03 METRIC TON/HA. LEAF AREA =	7.786

YEAR SPEC. 250	NUM.	NAME	BIOMASS
6	5	CARYA OVATA	0.832E 01
	DBH	3.490 3.159 3.150 3.142 1.600	
8	7	CARYA TOMENTOSA	0.260E 01
	DBH	1.503 1.562 1.574 1.658 1.598 1.808 1.525	
10	8	CELTIS LAEVIGATA	0.476E 01
	DBH	2.083 1.856 1.865 1.856 2.026 1.897 2.172 1.865	
11	4	CORNUS FLORIDA	0.805E 01
	DBH	3.344 3.124 3.418 3.124	
15	13	ILEX OPACA	0.148E 02
	DBH	3.421 3.399 3.563 3.420 2.568 2.617 2.482 2.454 1.274 1.271 1.283 1.270 1.319	
18	7	LIQUIDAMBAR STYRACIFLUA	0.355E 01
	DBH	1.022 1.843 1.812 1.810 1.886 1.841 1.805	
21	6	OSTRYA VIRGINIANA	0.532E 01
	DBH	2.615 2.700 2.030 2.090 2.034 2.273	
26	2	QUERCUS FALCATA	0.147E 04
	DBH	51.207 4.339	

27	15	QUERCUS MARILANDICA	0.985E 01
	DBH	2.676 2.363 2.429 2.417 2.001 2.299 2.035 2.340 1.582 1.534 1.641 1.753 1.656 1.534 1.549	
29	1	QUERCUS STELLATA	0.377E 01
	DBH	4.234	
32	6	ULMUS ALATA	0.311E 02
	DBH	5.226 4.915 5.346 4.915 4.917 3.335	
74		0.188E 02 METRIC TON/HA. LEAF AREA =	0.954

YEAR SPEC. 300	NUM.	NAME	BIOMASS
2	4	CARPINUS CAROLINIANA	0.103E 01
	DBH	1.569 1.386 1.270	1.270
5	3	CARYA LACINIOSA	0.692E 00
	DBH	1.376 1.279 1.293	
6	16	CARYA OVATA	0.283E 03
	DBH	16.100 13.134 9.369 9.139 9.157 6.338 6.737 6.385 6.815 6.083 2.824 2.813 2.962 3.227 2.873 2.909	
7	2	CARYA TEXANA	0.493E 00
	DBH	1.431 1.271	
8	10	CARYA TOMENTOSA	0.229E 03
	DBH	15.015 14.841 8.482 8.763 7.965 7.749 1.270 1.345 1.270 1.352	
10	11	CELTIS LAEVIGATA	0.216E 03
	DBH	11.299 10.962 11.454 8.471 8.459 8.625 8.345 6.664 4.288 3.522 3.508	
11	14	CORNUS FLORIDA	0.540E 02
	DBH	7.389 7.159 7.297 4.481 4.555 1.480 2.127 1.918 1.506 1.480 1.902 1.588 1.486 1.486	
13	1	FAGUS GRANDIFOLIA	0.218E 00
	DBH	1.288	
15	6	ILEX OPACA	0.260E 03
	DBH	15.932 13.160 12.649 12.019 7.631 0.102	
18	11	LIQUIDAMBAR STYRACIFLUA	0.498E 03
	DBH	18.288 18.504 15.747 13.676 8.819 8.557 8.461 8.461 5.405 5.225 0.101	
21	2	OSTRYA VIRGINIANA	0.275E 00
	DBH	0.101 1.417	
25	5	QUERCUS ALBA	0.838E 02
	DBH	11.629 11.258 2.438 2.455 1.477	
26	34	QUERCUS FALCATA	0.482E 04
	DBH	77.409 25.001 15.814 16.623 15.784 15.972 15.789 10.441 6.992 6.904 6.164 5.724 6.056 5.721 5.719 6.029 5.413 5.870 3.911 3.587 3.723 3.592 2.895 2.862 2.862 2.864 2.977 2.862 2.874 2.862 3.148 2.896 2.921 2.995	

27	3	QUERCUS MARILANDICA	0.659E 02
	DBH	9.811 8.891 7.628	
29	20	QUERCUS STELLATA	0.799E 03
	DBH	20.563 13.642 13.651 13.999	
		13.784 13.587 13.728 13.638	
		13.641 14.575 7.393 5.898	
		6.067 5.897 6.470 5.896	
		1.276 1.272 1.270 0.173	
32	11	ULMUS ALATA	0.446E 03
	DBH	22.916 17.756 16.466 4.480	
		4.242 4.289 3.538 3.548	
		1.294 1.665 0.152	
33	1	ULMUS AMERICANA	0.224E 00
	DBH	1.300	
154	0.931E 02	METRIC TON/HA. LEAF AREA =	5.125

YEAR SPEC.	NUM.	NAME	BIOMASS
350			
2	2	CARPINUS CAROLINIANA	0.430E 00
	DBH	1.280 1.278	
5	1	CARYA LACINIOSA	0.223E 00
	DBH	1.299	
6	16	CARYA OVATA	0.792E 03
	DBH	27.565 19.406 15.211 14.672	
		14.207 9.856 9.837 10.460	
		0.136 0.112 1.407 1.329	
		1.303 1.282 1.370 1.270	
7	3	CARYA TEXANA	0.675E 00
	DBH	1.284 1.306 1.320	
8	7	CARYA TOMENTOSA	0.349E 03
	DBH	28.040 1.464 1.452 1.347	
		1.366 1.270 1.497	
10	3	CELTIS LAEVIGATA	0.399E 03
	DBH	19.806 20.004 16.148	
11	3	CORNUS FLORIDA	0.788E 00
	DBH	1.288 1.549 1.521	
16	1	ILEX OPACA	0.342E 00
	DBH	1.553	
18	11	LIQUIDAMBAR STYRACIFLUA	0.192E 04
	DBH	37.444 38.465 29.044 16.818	
		1.425 0.127 1.271 1.276	
		1.412 1.557 1.488	
21	2	OSTRYA VIRGINIANA	0.515E 00
	DBH	1.475 1.273	
26	18	QUERCUS FALCATA	0.386E 04
	DBH	31.984 33.457 31.863 32.422	
		31.914 24.189 19.686 19.530	
		18.363 18.182 17.503 18.110	
		16.980 14.216 13.026 12.965	
		12.952 12.952	
27	2	QUERCUS MARILANDICA	0.450E 00
	DBH	1.332 1.274	
29	5	QUERCUS STELLATA	0.147E 04
	DBH	36.984 25.371 24.921 24.983	
		0.100	
32	1	ULMUS ALATA	0.565E 03
	DBH	34.347	

33	6	ULMUS AMERICANA	0.144E 01				
	DBH	1.276 1.274 1.270	1.309				
		1.310 1.562					
	81	0.112E 03 METRIC TON/HA. LEAF AREA =	6.327				
YEAR SPEC. 400	NUM.	NAME	BIOMASS				
	5	3 CARYA LACINIOSA	0.641E 00				
	DBH	1.270 1.281 1.276					
	6	2 CARYA OVATA	0.263E 03				
	DBH	24.955 1.271					
	7	4 CARYA TEXANA	0.113E 01				
	DBH	1.408 1.491 1.271	1.553				
	8	4 CARYA TOMENTOSA	0.902E 00				
	DBH	1.270 1.270 1.326	1.350				
	10	5 CELTIS LAEVIGATA	0.239E 03				
	DBH	23.952 1.283 1.280	1.285				
		1.532					
	13	1 FAGUS GRANDIFOLIA	0.216E 00				
	DBH	1.282					
	26	12 QUERCUS FALCATA	0.823E 04				
	DBH	48.937 47.770 50.093	48.447				
		37.638 30.663 30.143	29.821				
		27.980 24.674 23.138	22.998				
	27	5 QUERCUS MARILANDICA	0.132E 01				
	DBH	1.271 1.392 1.510	1.473				
		1.296					
	29	7 QUERCUS STELLATA	0.732E 03				
	DBH	38.240 0.101 1.306	1.386				
		1.314 1.273 1.477					
	32	1 ULMUS ALATA	0.218E 00				
	DBH	1.287					
	44	0.114E 03 METRIC TON/HA. LEAF AREA =	5.819				
YEAR SPEC. 450	NUM.	NAME	BIOMASS				
	2	3 CARPINUS CAROLINIANA	0.764E 00				
	DBH	1.422 1.271 1.418					
	6	1 CARYA OVATA	0.635E 03				
	DBH	36.062					
	7	1 CARYA TEXANA	0.215E 00				
	DBH	1.278					
	8	4 CARYA TOMENTOSA	0.911E 00				
	DBH	1.275 1.285 1.382	1.294				
	10	1 CELTIS LAEVIGATA	0.500E 03				
	DBH	32.643					
	15	2 ILEX OPACA	0.466E 00				
	DBH	1.325 1.319					
	25	6 QUERCUS ALBA	0.156E 01				
	DBH	1.461 1.418 1.271	1.564				
		1.280 1.270					

26	11	QUERCUS FALCATA	0.799E 04
	DBH	64.076 69.005 44.239 41.929	
		33.531 2.805 1.480 1.321	
		1.473 1.458 1.273	
27	3	QUERCUS MARILANDICA	0.754E 00
	DBH	1.475 1.270 1.338	
32	3	ULMUS ALATA	0.793E 00
	DBH	1.384 1.305 1.486	
	35	C.110E 03 METRIC TON/HA. LEAF AREA =	5.177
YEAR SPEC. 500	NUM.	NAME	BIOMASS
	2	1 CARPINUS CAROLINIANA	0.227E 00
	DBH	1.309	
	6	1 CARYA OVATA	0.121E 04
	DBH	47.232	
	7	3 CARYA TEXANA	0.848E 00
	DBH	1.271 1.161 1.771	
	10	2 CELTIS LAEVIGATA	0.877E 03
	DBH	41.266 1.278	
	13	3 FAGUS GRANDIFOLIA	0.780E 00
	DBH	1.540 1.271 1.323	
	15	8 ILEX OPACA	0.189E 01
	DBH	1.272 1.565 1.275 1.360	
		1.285 1.270 1.306 1.282	
	21	4 OSTRYA VIRGINIANA	0.990E 00
	DBH	1.511 1.274 1.349	1.272
	26	3 QUERCUS FALCATA	0.119E 05
	DBH	81.316 88.389 60.141	
	32	7 ULMUS ALATA	0.180E 01
	DBH	1.288 1.429 1.335 1.519	
		1.374 1.559 1.293	
	32	0.167E 03 METRIC TON/HA. LEAF AREA =	7.284

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