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INDIVIDUAL TREE MODELS

Individual tree models

 Are systems of equations to simulate stand dynamics by <u>incrementing each tree</u> during a growth period in relation to its growing conditions.



- provide detailed information about stand development and structure, including the distribution of stand volume by size classes
- have inherent flexibility allowing modeling:
 - combinations of species mixtures and stand structures
 - management regimes
 - regeneration methods

Individual tree models types

XNo tree locations known

Also called distance-independent, position-independent, location-independent, non-spatially-explicit

× Usually consist of:

- ✓ Diameter growth equation
- Height growth equation (or h-d relationship to estimate h from d)
- Mortality (stochastically generated or predicted as a function of growth rate and/or tree characteristics)
- Growth predicted as a function of tree size and stand (t and N) and site variables (S)

XTree locations known

Also called distance-dependent, position-dependent, location-dependent, spatially-explicit

× Usually consist of:

- Initialization: trees inputted or generated (incliding their coordinates)
- Growth predicted as a function of tree size, site quality (S) and a measure of competition from neighbouring trees (generally a function of size of the subject tree and (size, distance) to competitors
- Survival controlled stochastically as a function of competition and/or tree attributes

Individual tree models growth and calculus modules



Individual tree models - state variables

× The most common principal variables

- Dominant height (stand level variable)
- Diameter at breast height
- Tree height may also be a principal variable

× Derived variables

- Tree: total height and height to the base of the crown, tree volume, tree biomass (total and per component), sometimes crown width
- Stand: all variables except dominant height

Calculus of stand variables - example for stand volume

× Start with a list of all trees at time t_1 (at least tree diameter/or height):

- Growth module (GM) prediction of:
 - tree mortality
 - tree diameter growth for each tree
 - tree height growth for each tree (eventually)
- Calculus module CM)- estimation of:
 - tree height with a height-diameter curve (if not predicted in the GM)
 - tree volume with a volume equation
 - Calculus of plot volume by summing up the volume of every tree in the plot
 - Expansion to the ha, using the respective expansion factor= 10000/plot area (many models use 1 ha plots therefore this step is not needed)

Modeling individual tree dbh growth

× What factors/variables influence tree dbh growth?

- ✓ site quality
- micro-environment conditions (mainly soil)
- ✓ tree age
- ✓ tree size
- ✓ stand density
- influence of local neighbours
- ✓ genetics

Modeling individual tree dbh growth

- × Several methods have been used to <u>model tree dbh growth</u>, which may be classified as:
 - *** Method I** Linear or nonlinear regression models using i_d or i_g as dependent variable
 - *** Method II** Difference equations (d_{t2} or g_{t2} as dependent variable)
 - Method III Growth potential x modifier type models
 - Dependent variable is usually i_d or i_g

Let's look at each one of the methods

Method I - Linear regression models example



Method II - Difference equations example

× Dbh growth model for dominant cork oak trees without age explicit (200 is an asymptote)



Method III - Potential X modifier type models

× These models are based on the assumption that individual tree growth may be modeled as:

 $i_d = i_d$ potential X modifier

The *i_d* **potential** represents the growth of a tree of the same size that grows without <u>limitations</u>

The modifier is a function that takes values between 0 and 1, <u>defining growth restrictions</u> (usually competition but other factors may also be taken into account)

Method III - Potential X modifier type models

× There are different concepts of <u>potential growth</u> that have been used:

- Maximum growth that a tree of the same species and size/age may attain under "optimum conditions" in terms of water and nutrients
- Maximum observed growth for a tree of the same species and size in "<u>normal</u>" <u>conditions</u>
- Maximum growth of the trees in the same plot (growth of the dominant trees)

Method III - Potential X modifier type models, example

× GLOB-tree model - potential growth



 $ipot_d = ddom_{t2} - ddom_{t1}$

Method III - Potential X modifier type models, example

× GLOB-tree model - modifier



Height estimation - example

- GLOB-tree model:
- × Young stands (t<4 years):

$$h = 1.30 + hdom \left(1 + \left(-0.43487 - 0.0108 \ t + 0.09772 \ hdom - 0.06021 \ dg \right) e^{-0.04864} \ hdom \right) \left(\begin{array}{c} -1.58926 \ \frac{d}{hdom} \end{array} \right)$$

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× Adult stands (t>4 years):

$$h = hdom \left(1 + \left(0.10694 + 0.02916 \frac{N}{1000} - 0.00176 dmax \right) e^{0.03540 hdom} \right) \left(1 - e^{-1.81117 \frac{d}{hdom}} \right)$$

Crown variables - example

× GLOB-tree model - crown ratio



Predicting tree mortality - example

× GLOB-tree model

