# Forest growth and dynamics (practice)

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### Outline

- 1. Forest growth inputs
- 2. Running forest growth
- 3. Evaluation of growth predictions
- 4. Forest dynamics
- 5. Forest dynamics including management

M.C. Escher - Up and down, 1947





# 1. Forest growth inputs



### Creating the forest growth input object

We assume we have an appropriate forest object:

```
1 data(exampleforest)
```

a species parameter data frame:

```
1 data(SpParamsMED)
```

a soil data frame:

```
1 examplesoil <- defaultSoilParams(4)</pre>
```

and simulation control list:

```
1 control <- defaultControl("Granier")
```

With these four elements we can build our input object for function growth():

```
1 x <- forest2growthInput(exampleforest, examplesoil, SpParamsMED, control)</pre>
```



#### Structure of the growth input object (1)

The growth input object is a list with several elements:

```
1 \text{ names}(x)
 [1] "control"
                                     "soil"
 [3] "snowpack"
                                     "canopy"
     "herbLAI"
                                     "herbLAImax"
     "cohorts"
                                     "above"
     "below"
 [9]
                                     "belowLayers"
[11] "paramsPhenology"
                                     "paramsAnatomy"
[13] "paramsInterception"
                                     "paramsTranspiration"
[15] "paramsWaterStorage"
                                     "paramsGrowth"
[17] "paramsMortalityRegeneration" "paramsAllometries"
[19] "internalPhenology"
                                     "internalWater"
[21] "internalLAIDistribution"
                                    "internalCarbon"
[23] "internalAllocation"
                                     "internalMortality"
[25] "internalFCCS"
```

Element above contains the above-ground structure data that we already know, but with an additional column SA that describes the estimated initial amount of *sapwood area*:

```
LAI live
        SP
                      DBH Cover
                                           CR
T1 148 148 168,0000 37,55
                             NA 800 0.6605196 383.4520992 0.84874773
T2_168 168 384.0000 14.60
                             NA 660 0.6055642 47.0072886 0.70557382
S1_165 165 749.4923
                       NA 3.75 80 0.8032817
                                                0.9753929 0.03062604
       LAI expanded LAI dead LAI nocomp
                                           Loading ObsID
         0.84874773
                           0 1.29720268 0.32447403
                                                    <NA>
T1 148
T2 168
         0.70557382
                           0 1.01943205 0.20102636
                                                    <NA>
         0.03062604
                           0 0.04412896 0.01407945
S1 165
                                                    <NA>
```



#### Structure of the growth input object (2)

Elements starting with params\* contain cohort-specific model parameters. An important set of parameters are in paramsGrowth:

```
RERleaf RERsapwood RERfineroot CCleaf CCsapwood CCfineroot
T1_148 0.01210607 5.15e-05 0.0009610199 1.5905
                                                    1.47
                                                                1.3
T2_168 0.01757808 5.15e-05 0.0072846640 1.4300
                                                    1.47
                                                               1.3
S1_165 0.02647746 5.15e-05 0.0072846640 1.5320
                                                    1.47
                                                               1.3
      RGRleafmax RGRsapwoodmax RGRcambiummax RGRfinerootmax SRsapwood
T1 148
            0.09
                                 0.002628095
                                                       0.1 0.000135
                            NA
            0.09
T2 168
                                 0.002500000
                                                      0.1 0.000135
                            NA
            0.09
                         0.002
                                                       0.1 0.000135
S1 165
                                         NA
       SRfineroot
                       RSSG fHDmin fHDmax
                                             Upood
T1_148 0.001897231 0.3725000
                                     160 0.4979943
T2 168 0.001897231 0.9500000
                               40
                                   100 0.4740096
S1_165_0.001897231_0.7804035
                               NA
                                      NA 0.4749178
```

Elements starting with internal\* contain state variables required to keep track of plant status. For example, the metabolic and storage carbon levels can be seen in internalCarbon:

```
sugarLeafstarchLeafsugarSapwoodstarchSapwoodT1_1480.40292390.009251230.57384873.201897T2_1680.35857510.009251231.07413833.100817S1_1650.72235260.009251230.28576552.654773
```



# 2. Forest growth



#### Forest growth run

The call to function growth() needs the growth input object, the weather data frame, latitude and elevation:

```
1 G <- growth(x, examplemeteo, latitude = 41.82592, elevation = 100)
Initial plant cohort biomass (g/m2): 5068.34
Initial plant water content (mm): 4.73001
Initial soil water content (mm): 290.875
Initial snowpack content (mm): 0
Performing daily simulations
 Year 2001:.....
Final plant biomass (g/m2): 5256.53
Change in plant biomass (g/m2): 188.193
Plant biomass balance result (g/m2): 188.193
Plant biomass balance components:
  Structural balance (g/m2) 104 Labile balance (g/m2) 92
  Plant individual balance (g/m2) 196 Mortality loss (g/m2) 8
Final plant water content (mm): 4.73794
Final soil water content (mm): 274.787
Final snowpack content (mm): 0
Change in plant water content (mm): 0.00793033
Plant water balance result (mm): -0.00116903
Change in soil water content (mm): -16.0878
Soil water balance result (mm): -16.0878
Change in snowpack water content (mm): 0
Snowpack water balance result (mm): 7.10543e-15
Water balance components:
  Precipitation (mm) 513 Rain (mm) 462 Snow (mm) 51
  Intercention /mm\ 00 Not reinfell /mm\ 070
```



#### **Growth output object**

GrowthMortality

Function growth() returns an object of class with the same name, actually a list:

```
1 class(G)
[1] "growth" "list"
... whose elements are:
  1 names(G)
                          "topography"
                                               "weather"
 [1] "latitude"
 [4] "growthInput"
                          "growthOutput"
                                              "WaterBalance"
 [7] "CarbonBalance"
                          "BiomassBalance"
                                              "Soil"
                          "Stand"
                                              "Plants"
    "Snow"
[13] "LabileCarbonBalance" "PlantBiomassBalance" "PlantStructure"
[16] "GrowthMortality"
Elements
                                     Information
                                     Copies of the information used in the call to growth()
latitude, topography, weather,
growthInput
                                     State variables at the end of the simulation (can be used as input to a subsequent one)
growthOutput
                                     [same as spwb ...]
WaterBalance, Soil, Snow, Stand,
Plants
                                     Stand-level carbon blance
CarbonBalance
                                     Components of the individual-level labile carbon balance
LabileCarbonBalance
                                     Components of indvidual- and cohort-level biomass balance
PlantBiomassBalance
                                     Structural variables (DBH, height, sapwood area...)
PlantStructure
```

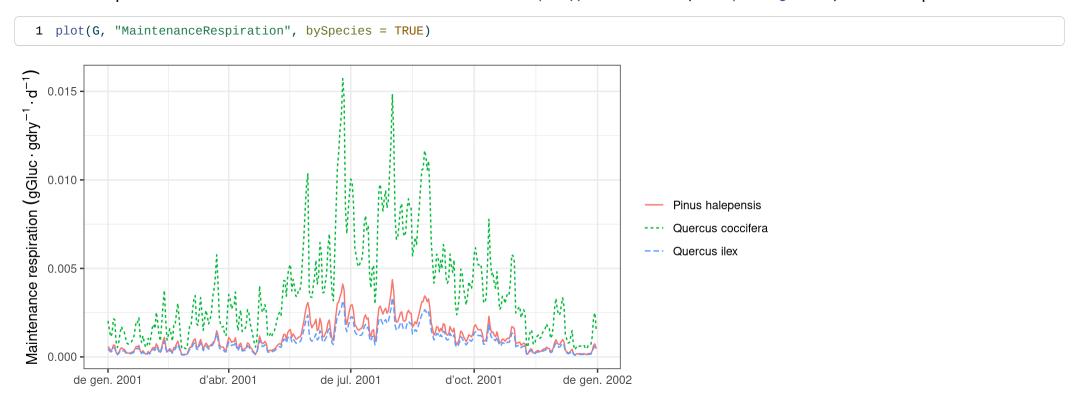
Growth and mortality rates



### **Post-processing**

Users can inspect the output of growth() simulations using functions extract(), summary() and plot() on the simulation output.

Several new plots are available in addition to those available for spwb() simulations (see ?plot.growth). For example:



... but instead of typing all plots, we can call the interactive plot function shinyplot().



# 3. Evaluation of growth predictions



#### **Observed data frame**

Evaluation of growth simulations will normally imply the comparison of predicted vs observed **basal area increment** (BAI) or **diameter increment** (DI) at a given temporal resolution.

Here, we illustrate the evaluation functions included in the package using a fake data set at *daily* resolution, consisting on the predicted values and some added error.

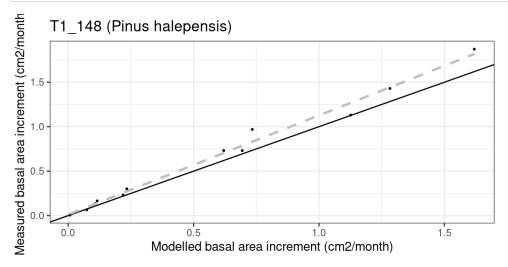
```
1 data(exampleobs)
  2 head(exampleobs)
       dates
                                   E_T1_148
                                              E_T2_168 FMC_T1_148 FMC_T2_168
1 2001-01-01 0.3007733 2.2436218 0.09187857 0.14142950
                                                         125.9071
                                                                     93.07915
2 2001-01-02 0.3091627 2.3236565 0.26480973 0.19095008
                                                         125.9137
                                                                     93.07863
3 2001-01-03 0.2996498 0.7409083 0.15345643 0.17546363
                                                         125.8760
                                                                     93.10512
                                                                     93.07022
4 2001-01-04 0.3042764 1.7173522 0.23470647 0.04643454
                                                         125.8643
5 2001-01-05 0.3054886 2.0002562 0.37687792 0.10623552
                                                         125.8493
                                                                     93.08487
6 2001-01-06 0.3062005 2.0722706 0.16342360 0.05550329
                                                         125.9367
                                                                     93.07343
    BAI_T1_148 BAI_T2_168
                             DI_T1_148 DI_T2_168
1 6.222625e-06
                        0 9.948881e-08
2 3.091274e-10
                        0 1.071090e-11
3 1.298482e-13
                        0 0.000000e+00
4 2.886195e-11
                        0 5.552753e-13
5 1.287020e-03
                        0 1.367289e-05
6 1.471202e-03
                        0 1.000411e-05
                                                0
```

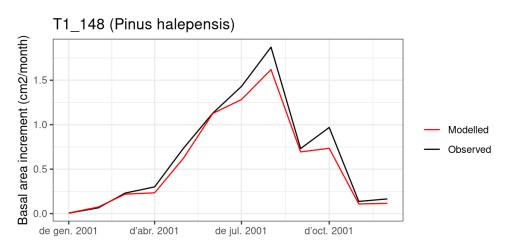
To specify observed growth data at *monthly* or *annual scale*, you should specify the first day of each month/year (e.g. 2001-01-01, 2002-01-01, etc for years) as row names in your observed data frame.



#### **Evaluation plot**

Assuming we want to evaluate the predictive capacity of the model in terms of monthly basal area increment for the *pine cohort* (i.e. T1\_148), we can plot the relationship between observed and predicted values using evaluation\_plot():





Using temporalResolution = "month" we indicate that simulated and observed data should be temporally aggregated to conduct the comparison.

The following code would help us quantifying the *strength* of the relationship:



# 4. Forest dynamics



### Forest dynamics run

#### Weather preparation

In this vignette we will fake a three-year weather input by repeating the example weather data frame four times:

```
1 meteo <- rbind(examplemeteo, examplemeteo, examplemeteo, examplemeteo)</pre>
```

we need to update the dates in row names so that they span four consecutive years:

#### Simulation

The call to fordyn() has the following form:

#### **Important**

- fordyn() operates on forest objects directly, instead of using an intermediary object (such as spwbInput and growthInput).
- fordyn() calls function growth() internally for each simulated year, but all console output from growth() is hidden.



As with other models, the output of fordyn() is a list, which has the following elements:

Elements	Information
StandSummary, SpeciesSummary, CohortSummary	Annual summary statistics at different levels
TreeTable, ShrubTable	Structural variables of <b>living</b> cohorts at each annual time step.
DeadTreeTable, DeadShrubTable	Structural variables of <b>dead</b> cohorts at each annual time step
CutTreeTable, CutShrubTable	Structural variables of <b>cut</b> cohorts at each annual time step
ForestStructures	Vector of forest objects at each time step.
GrowthResults	Result of internally calling growth() at each time step.
ManagementArgs	Management arguments for a subsequent call to fordyn().
NextInputObject, NextForestObject	Objects growthInput and forest to be used in a subsequent call to fordyn().



For example, we can compare the initial forest object with the final one:

```
1 exampleforest
                                                                          1 fd$NextForestObject
$treeData
                                                                       $treeData
           Species N DBH Height Z50 Z95
                                                                                               DBH Height
                                                                                                                   N Z50 Z95
                                                                                  Species
1 Pinus halepensis 168 37.55
                               800 100 600
                                                                       1 Pinus halepensis 38.01410 824.7217 166.7796 100 600
      Quercus ilex 384 14.60
                                                                             Quercus ilex 15.04679 673.8471 382.6513 300 1000
                              660 300 1000
$shrubData
                                                                       $shrubData
                                                                                   Species Height
            Species Cover Height Z50 Z95
                                                                                                      Cover Z50 Z95
1 Quercus coccifera 3.75
                              80 200 1000
                                                                       1 Quercus coccifera 75.4552 3.237287 200 1000
$herbCover
                                                                       $herbCover
                                                                       [1] 10
[1] 10
$herbHeight
                                                                       $herbHeight
[1] 20
                                                                       [1] 20
$seedBank
                                                                        $seedBank
[1] Species Percent
                                                                                   Species Percent
<0 files> (o «row.names» de longitud 0)
                                                                        1 Pinus halepensis
                                                                                               100
                                                                              Quercus ilex
                                                                                               100
attr(,"class")
                                                                       3 Quercus coccifera
                                                                                               100
[1] "forest" "list"
                                                                       attr(,"class")
                                                                       [1] "forest" "list"
```



The output includes **summary statistics** that describe the structural and compositional state of the forest corresponding to *each* annual time step.

For example, we can access *stand-level* statistics using:

	1 fd\$StandS	ummary			
	Step NumTree	Species NumTreeCoho	orts NumShrubSp	ecies NumShrub	Cohorts
1	0	2	2	1	1
2	1	2	2	1	1
3	2	2	2	1	1
4	3	2	2	1	1
5	4	2	2	1	1
	TreeDensityLive TreeBasalAreaLive DominantTreeHeight DominantTreeDiameter				
1	552.0			800.0000	37.55000
2	551.3	663 25.208	314 8	306.2122	37.66571
3	550.7	269 25.383	313 8	312.4144	37.78185
4	550.0	818 25.558	325 8	318.5877	37.89804
5	549.4	309 25.733	303 8	324.7217	38.01410
	QuadraticMea	nTreeDiameter HartE	BeckingIndex Sh	rubCoverLive E	BasalAreaDead
1		24.02949	53.20353	3.750000	0.0000000
2		24.12711	52.82391	3.092051	0.03917375
3		24.22476	52.45105	3.139858	0.03983747
4		24.32243	52.08602	3.188231	0.04050957
5		24.41991	51.72923	3.237287	0.04118879
	ShrubCoverDe	ad BasalAreaCut Shr	rubCoverCut		
1	0.0000000	00 0	0		
2	0.0053088	98 0	0		
3	0.0047844	68 0	0		
4	0.0048583	42 0	0		
5	0.0049331	17 0	0		
5	0.00-3331	_, 0	0		



Another useful output of fordyn() are tables in long format with cohort structural information (i.e. DBH, height, density, etc) for each time step:

```
1 fd$TreeTable
   Step Year Cohort
                             Species
                                           DBH
                                                 Height
                                                               N Z50
                                                                      Z95 ObsID
          NA T1_148 Pinus halepensis 37.55000 800.0000 168.0000 100
                                                                      600
                                                                           <NA>
                         Quercus ilex 14.60000 660.0000 384.0000 300 1000
          NA T2_168
                                                                           <NA>
      1 2001 T1_148 Pinus halepensis 37.66571 806.2122 167.6992 100
                                                                           <NA>
                         Quercus ilex 14.71218 663.4915 383.6671 300 1000
      1 2001 T2_168
                                                                           <NA>
      2 2002 T1_148 Pinus halepensis 37.78185 812.4144 167.3956 100
                                                                           <NA>
      2 2002 T2_168
                        Quercus ilex 14.82389 666.9588 383.3314 300 1000
                                                                           <NA>
      3 2003 T1_148 Pinus halepensis 37.89804 818.5877 167.0890 100
                                                                           <NA>
      3 2003 T2_168
                        Quercus ilex 14.93552 670.4135 382.9928 300 1000
      4 2004 T1_148 Pinus halepensis 38.01410 824.7217 166.7796 100
                                                                           <NA>
10
      4 2004 T2_168
                        Quercus ilex 15.04679 673.8471 382.6513 300 1000
                                                                           <NA>
```

Note

The NA values in Year correspond to the initial state.

The same information can be shown for trees that are predicted to die during each simulated year:

```
1 fd$DeadTreeTable
Step Year Cohort
                          Species
                                       DBH
                                             Height
                                                            N N_starvation
   1 2001 T1_148 Pinus halepensis 37.66571 806.2122 0.3007828
                     Quercus ilex 14.71218 663.4915 0.3328893
   1 2001 T2_168
   2 2002 T1_148 Pinus halepensis 37.78185 812.4144 0.3036481
                     Quercus ilex 14.82389 666.9588 0.3357419
   2 2002 T2_168
   3 2003 T1_148 Pinus halepensis 37.89804 818.5877 0.3065253
                     Quercus ilex 14.93552 670.4135 0.3386082
   3 2003 T2_168
   4 2004 T1_148 Pinus halepensis 38.01410 824.7217 0.3094089
   4 2004 T2_168
                     Quercus ilex 15.04679 673.8471 0.3414828
N_dessication N_burnt Z50 Z95 ObsID
            0
                    0 100
                           600
                                <NA>
                    0 300 1000
                                <NA>
```

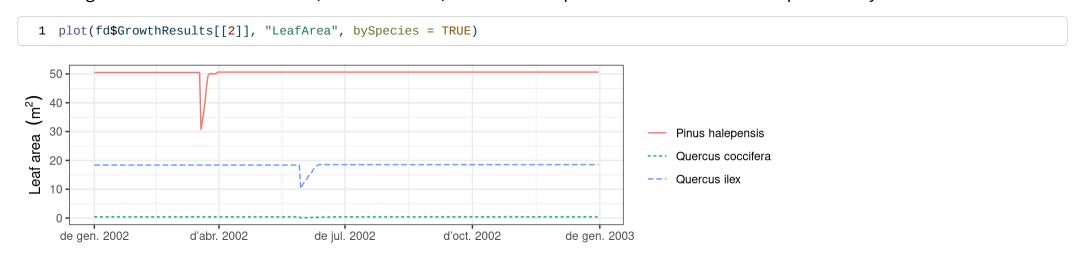


3	0	0	100	600	<na></na>
4	0	0	300	1000	<na></na>
5	0	0	100	600	<na></na>
6	0	0	300	1000	<na></na>
7	0	0	100	600	<na></na>
8	0	0	300	1000	<na></na>

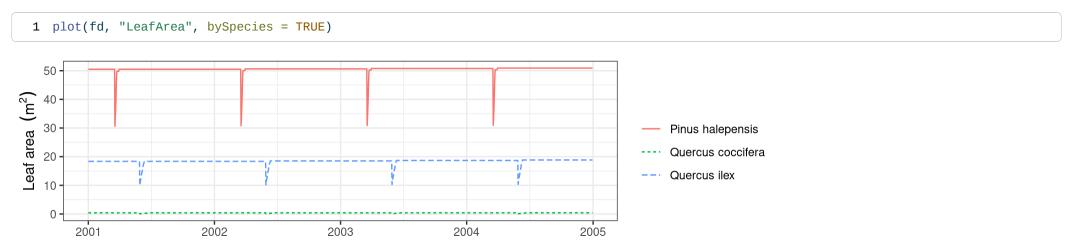


### **Post-processing**

Accessing elements of GrowthResults, we can extract, summarize or plot simulation results for a particular year:



It is also possible to plot the whole series of results by passing a fordyn object to the plot() function:



Finally, we can create interactive plots using function shinyplot(), in the same way as with other simulations.



5. Forest dynamics including management



#### Running simulations with management

fordyn() allows the user to supply an *arbitrary* function implementing a desired management strategy for the stand whose dynamics are to be simulated.

The package includes an in-built default function called defaultManagementFunction() along management parameter defaults provided by function defaultManagementArguments().

To run simulations with management we need to define (and modify) management arguments...

```
1 # Default arguments
2 args <- defaultManagementArguments()
3 # Here one can modify defaults before calling fordyn()
4 #</pre>
```

... and call fordyn() specifying the management function and its arguments:

When management is included, tables CutTreeTable and CutShrubTable will contain extraction data:

```
1 fd$CutTreeTable
2 fd$CutShrubTable
```



## Forest management parameters (1)

Function defaultManagementArguments() returns a list with default values for management parameters to be used in conjunction with defaultManagementFunction():

Element	Description
type	Management model, either 'regular' or 'irregular'
targetTreeSpecies	Either "all" for unspecific cuttings or a numeric vector of target tree species to be selected for cutting operations
thinning	Kind of thinning to be applied in irregular models or in regular models before the final cuts. Options are "below", "above", "systematic", "below-systematic", "above-systematic" or a string with the proportion of cuts to be applied to different diameter sizes
thinningMetric	The stand-level metric used to decide whether thinning is applied, either "BA" (basal area), "N" (density) or "HB" (Hart-Becking index)
thinningThreshold	The threshold value of the stand-level metric causing the thinning decision
thinningPerc	Percentage of stand's basal area to be removed in thinning operations
minThinningInterval	Minimum number of years between thinning operations
finalMeanDBH	Mean DBH threshold to start final cuts
finalPerc	String with percentages of basal area to be removed in final cuts, separated by '-' (e.g. "40-60-100")
finalYearsBetweenCuts	Number of years separating final cuts



## Forest management parameters (2)

The same list includes *state variables* for management (these are modified during the simulation):

Element	Description
yearsSinceThinning	State variable to count the years since the last thinning ocurred
finalPreviousStage	Integer state variable to store the stage of final cuts ('0' before starting final cuts)
finalYearsToCut	State variable to count the years to be passed before new final cut is applied.

Tip

Instead of using the in-built management function, you could code your own management function and specify its own set of parameters!



## M.C. Escher - Up and down, 1947

