

# 3.2 - Forest growth/dynamics (practice)

Miquel De Cáceres, Victor Granda, Aitor Ameztegui

Ecosystem Modelling Facility

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# Outline

1. Forest growth inputs
2. Running forest growth
3. Evaluation of growth predictions
4. Forest dynamics

# 1. Forest growth inputs

## Creating the forest growth input object

We assume we have an appropriate forest object:

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data(exampleforestMED)
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and simulation control list:

```
control <- defaultControl("Granier")
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a soil input object:

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examplesoil <- soil(defaultSoilParams(4))
```

and simulation control list:

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

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

```
x <- forest2growthInput(exampleforestMED, examplesoil, SpParamsMED, control)
```

# 1. Forest growth inputs

## Structure of the growth input object (1)

The growth input object is a `list` with several elements:

```
names(x)
```

```
## [1] "control"          "soil"             "canopy"           "cohorts"
## [5] "above"            "below"            "belowLayers"      "paramsPhenology"
## [9] "paramsAnatomy"    "paramsInterception" "paramsTranspiration" "paramsWaterStorage"
## [13] "paramsGrowth"     "paramsAllometries" "internalPhenology" "internalWater"
## [17] "internalCarbon"   "internalAllocation" "internalMortality"
```



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## [13] "paramsGrowth" "paramsAllometries" "internalPhenology" "internalWater"
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```

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*:

```
x$above
```

```
##           SP           N   DBH Cover   H           CR           SA   LAI_live LAI_expanded LAI_dead
## T1_148 148 168.0000 37.55   NA 800 0.6605196 437.032040 0.96734365 0.96734365 0
## T2_168 168 384.0000 14.60   NA 660 0.6055642 57.407064 0.86167321 0.86167321 0
## S1_165 165 749.4923   NA 3.75 80 0.8032817 1.251072 0.03928201 0.03928201 0
```

# 1. Forest growth inputs

## Structure of the growth input object (2)

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

```
x$paramsGrowth
```

```
##          RERleaf RERsapwood  RERfineroot CCleaf CCsapwood CCfineroot RGRleafmax RGRsapwoodmax
## T1_148 0.01210607  5.18e-05 0.0009610199 1.5905      1.47      1.3      0.03      NA
## T2_168 0.01757808  5.18e-05 0.0072846640 1.4300      1.47      1.3      0.03      NA
## S1_165 0.02647746  5.18e-05 0.0072846640 1.5320      1.47      1.3      0.03      0.002
##          RGRcambiummax RGRfinerootmax  SRsapwood  SRfineroot  RSSG fHDmin fHDmax  WoodC
## T1_148  0.002475167          0.1 0.0001261398 0.001897231 0.3725000    80    160 0.4979943
## T2_168  0.000650000          0.1 0.0001261398 0.001897231 0.9500000    40    100 0.4740096
## S1_165          NA          0.1 0.0001261398 0.001897231 0.7804035    NA     NA 0.4749178
##          MortalityBaselineRate
## T1_148          0.0050
## T2_168          0.0000
## S1_165          0.0015
```

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##          MortalityBaselineRate
## T1_148          0.0050
## T2_168          0.0000
## S1_165          0.0015
```

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`:

```
x$internalCarbon
```

```
##          sugarLeaf starchLeaf sugarSapwood starchSapwood
## T1_148 0.4029239 0.00925123  0.5738487    3.276375
## T2_168 0.3585751 0.00925123  1.0741383    3.280965
## S1_165 0.7223526 0.00925123  0.2857655    3.445161
```

## 2. Forest growth

### Forest growth run

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

```
G <- growth(x, examplemeteo, latitude = 41.82592, elevation = 100)
```

```
## Initial plant cohort biomass (g/m2): 7109.27
## Initial soil water content (mm): 291.257
## Initial snowpack content (mm): 0
## Performing daily simulations
##
## [Year 2001]: .....
##
## Final plant biomass (g/m2): 7347.27
## Change in plant biomass (g/m2): 238.004
## Plant biomass balance result (g/m2): 238.004
## Plant biomass balance components:
##   Structural balance (g/m2) 147 Labile balance (g/m2) 117
##   Plant individual balance (g/m2) 264 Mortality loss (g/m2) 26
## Final soil water content (mm): 268.687
## Final snowpack content (mm): 0
## Change in soil water content (mm): -22.57
## Soil water balance result (mm): -22.57
## Change in snowpack water content (mm): 0
## Snowpack water balance result (mm): 0
## Water balance components:
##   Precipitation (mm) 513
##   Rain (mm) 462 Snow (mm) 51
##   Interception (mm) 95 Net rainfall (mm) 368
##   Infiltration (mm) 409 Runoff (mm) 9 Deep drainage (mm) 75
##   Soil evaporation (mm) 19 Transpiration (mm) 338
```

## 2. Forest growth

### Growth output object

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

```
class(G)
```

```
## [1] "growth" "list"
```

... whose elements are:

Elements	Information
latitude, topography, weather, growthInput	Copies of the information used in the call to <code>growth()</code>
growthOutput	State variables at the end of the simulation (can be used as input to a subsequent one)
WaterBalance, Soil, Stand, Plants	[same as <code>spwb</code> ...]
LabileCarbonBalance	Components of the labile carbon balance
PlantBiomassBalance	Components of individual- and cohort-level biomass balance
PlantStructure	Structural variables (DBH, height, sapwood area...)
GrowthMortality	Growth and mortality rates
subdaily	Sub-daily outputs (not relevant here)

## 2. Forest growth

### Plots and summaries

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

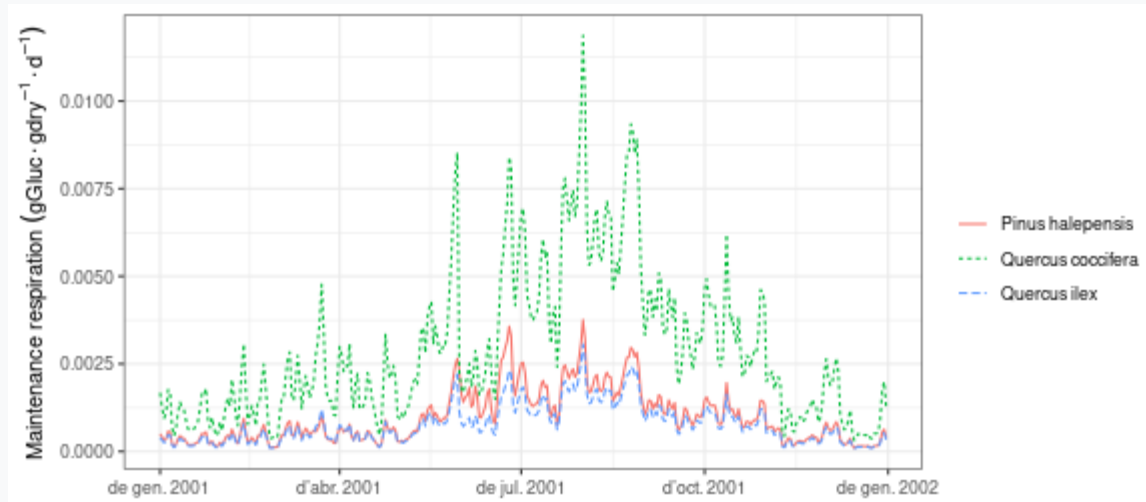
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Users can inspect the output of `growth()` simulations using functions `summary()` and `plot()` on the simulation output.

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

```
plot(G, "MaintenanceRespiration", bySpecies = TRUE)
```



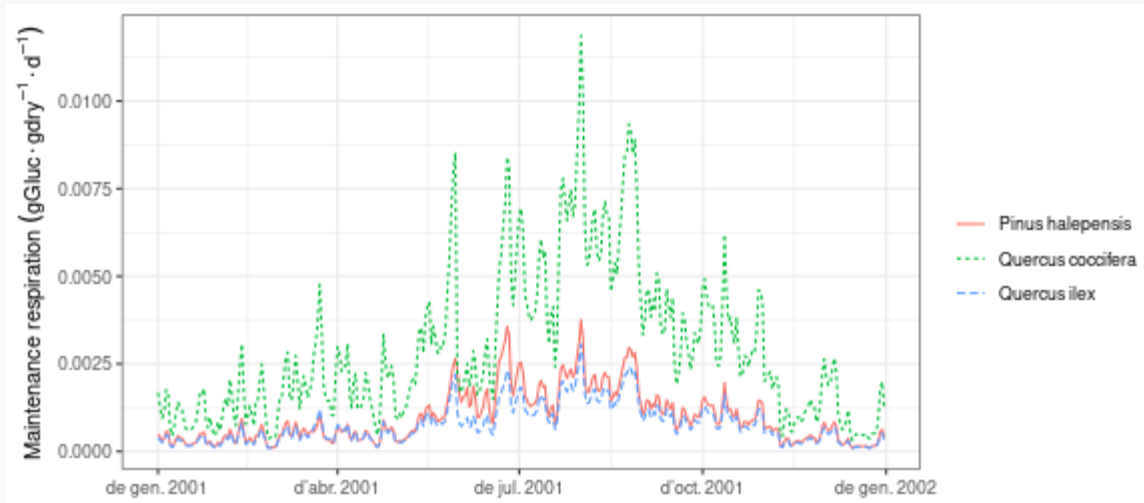
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```
plot(G, "MaintenanceRespiration", bySpecies = TRUE)
```



... 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.

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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.

```
data(exampleobs)
head(exampleobs)
```

##		SWC	ETR	E_T1_148	E_T2_168	FMC_T1_148	FMC_T2_168	BAI_T1_148	BAI_T2_168
##	2001-01-01	0.3114584	2.4128514	0.1213818	0.13682385	125.9553	93.13606	6.108618e-06	0
##	2001-01-02	0.2886956	2.2369541	0.3445222	0.21002440	125.8921	93.12522	8.984836e-10	0
##	2001-01-03	0.2939884	0.5408668	0.4552857	0.16751167	125.8828	93.11057	1.601437e-13	0
##	2001-01-04	0.3018085	1.7802127	0.3788494	0.04146642	125.7638	93.11051	1.363496e-11	0
##	2001-01-05	0.2957380	2.0220037	0.4057144	0.22997694	125.9627	93.04916	1.976577e-03	0
##	2001-01-06	0.3010738	2.8295695	0.1988711	0.23597595	125.8718	92.97068	1.705353e-03	0
##		DI_T1_148	DI_T2_168						
##	2001-01-01	1.105870e-07		0					
##	2001-01-02	3.029737e-11		0					
##	2001-01-03	0.000000e+00		0					
##	2001-01-04	2.948372e-13		0					
##	2001-01-05	4.457414e-05		0					
##	2001-01-06	3.254593e-05		0					

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##		DI_T1_148	DI_T2_168						
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##	2001-01-03	0.000000e+00		0					
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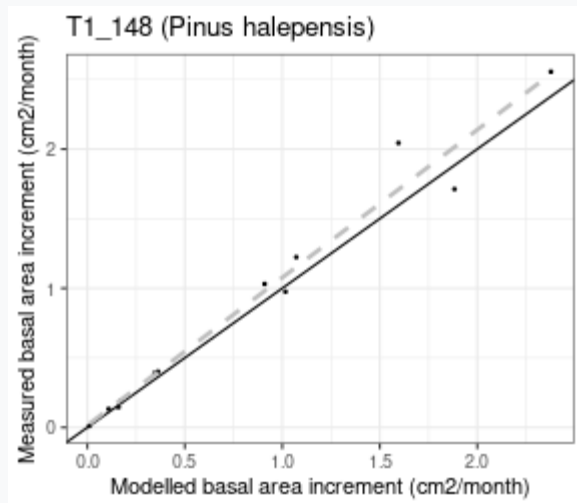
To specify observed growth data at *monthly* or *annual scale*, you should specify the first day of each month/year (e.g. 2001-01-01, 2002-01-01, etc for years) as row names in your observed data frame.

### 3. Evaluation of growth predictions

#### 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()`:

```
evaluation_plot(G, exampleobs, "BAI",  
               cohort = "T1_148",  
               temporalResolution = "month",  
               plotType = "scatter")
```

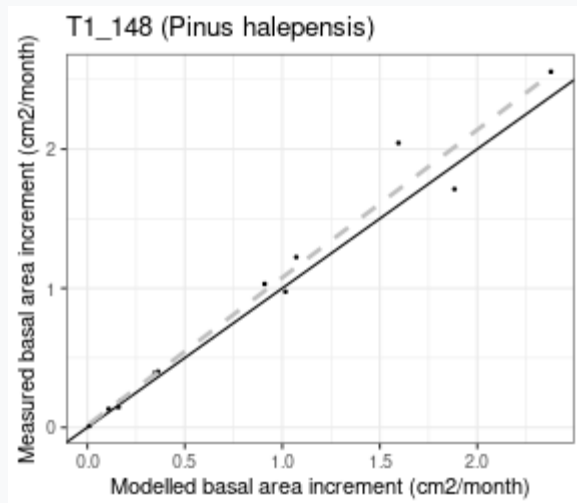


### 3. Evaluation of growth predictions

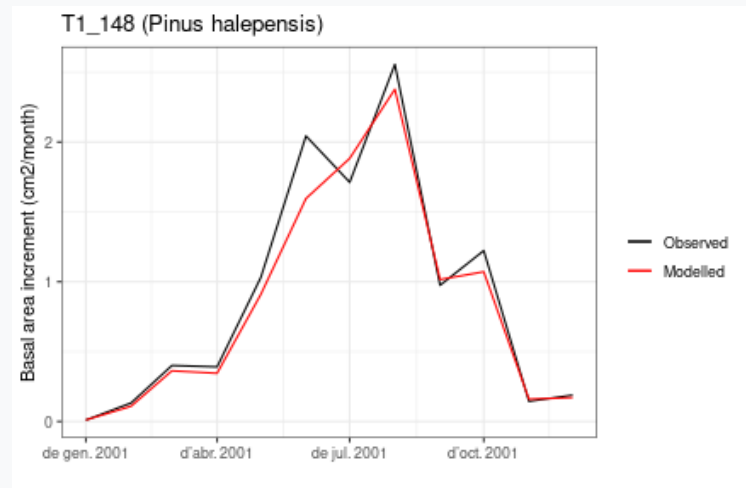
#### Evaluation plot

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()`:

```
evaluation_plot(G, exampleobs, "BAI",
               cohort = "T1_148",
               temporalResolution = "month",
               plotType = "scatter")
```



```
evaluation_plot(G, exampleobs, "BAI",
               cohort = "T1_148",
               temporalResolution = "month",
               plotType = "dynamics")
```

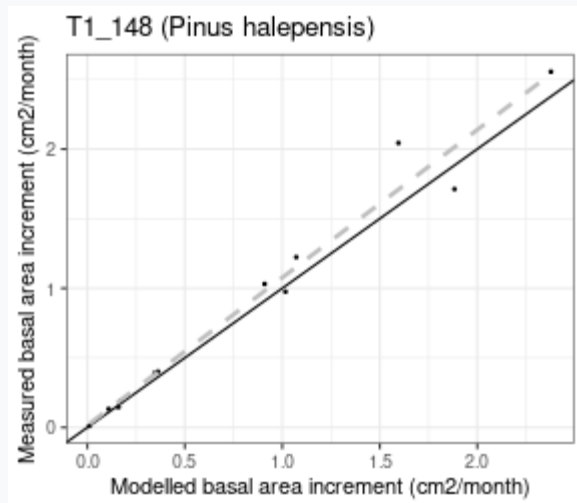


### 3. Evaluation of growth predictions

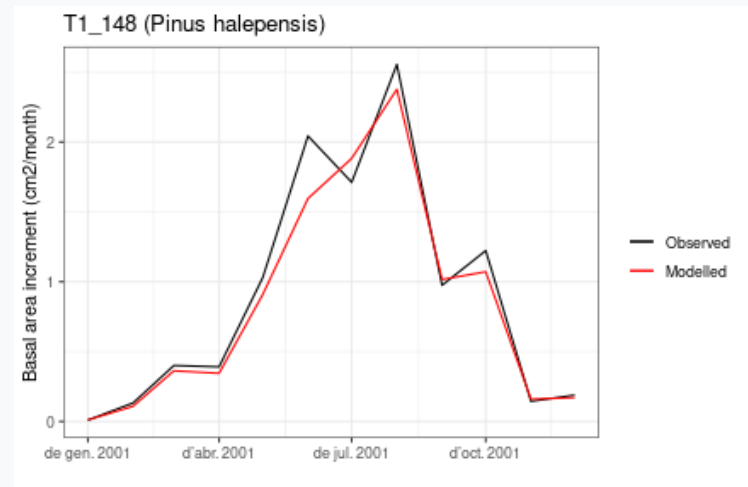
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```



```
evaluation_plot(G, exampleobs, "BAI",
               cohort = "T1_148",
               temporalResolution = "month",
               plotType = "dynamics")
```



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

### 3. Evaluation of growth predictions

#### Evaluation metrics

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

```
evaluation_stats(G, exampleobs, "BAI", cohort = "T1_148",
                temporalResolution = "month")
```

##	n	Bias	Bias.rel	MAE	MAE.rel	r	NSE	NSE.abs
##	12.000000000	-0.06622203	-7.35231796	0.10432221	11.58240068	0.98533974	0.96085838	0.84864074

## 4. Forest dynamics

### Weather preparation

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

```
meteo = rbind(examplemeteo, examplemeteo, examplemeteo)
```





## 4. Forest dynamics

### Simulation

**Remember:** `fordyn()` operates on forest objects directly, instead of using an intermediary object (such as `spwbInput` and `growthInput`).

```
fd<-fordyn(exampleforestMED, examplesoil, SpParamsMED, meteo, control,  
           latitude = 41.82592, elevation = 100)
```

```
## Simulating year 2001 (1/3):  (a) Growth/mortality, (b) Recruitment  
## Simulating year 2002 (2/3):  (a) Growth/mortality, (b) Recruitment  
## Simulating year 2003 (3/3):  (a) Growth/mortality, (b) Recruitment
```

## 4. Forest dynamics

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**Important:** `fordyn()` calls function `growth()` internally for each simulated year.

## 4. Forest dynamics

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## Simulating year 2003 (3/3):  (a) Growth/mortality, (b) Recruitment
```

**Important:** `fordyn()` calls function `growth()` internally for each simulated year.

The verbose option of the control parameters only affects function `fordyn()`, i.e. all console output from `growth()` is hidden.

## 4. Forest dynamics

### Forest dynamics output

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

Elements	Information
StandSummary, SpeciesSummary, CohortSummary	<i>Annual</i> 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 <code>growth()</code> at each time step.
ManagementArgs	Management arguments for a subsequent call to <code>fordyn()</code> .
NextInputObject, NextForestObject	Objects <code>growthInput</code> and <code>forest</code> to be used in a subsequent call to <code>fordyn()</code> .

## 4. Forest dynamics

### Forest dynamics output

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

exampleforestMED

```
## $ID
## [1] "1"
##
## $patchsize
## [1] 10000
##
## $treeData
##   Species    N   DBH Height  Z50   Z95
## 1     148 168 37.55    800 100   600
## 2     168 384 14.60    660 300 1000
##
## $shrubData
##   Species Cover Height  Z50   Z95
## 1     165   3.75    80 200 1000
##
## $herbCover
## [1] 10
##
## $herbHeight
## [1] 20
##
## attr(,"class")
## [1] "forest" "list"
```

fd\$NextForestObject

```
## $ID
## [1] "1"
##
## $patchsize
## [1] 10000
##
## $treeData
##   Species      DBH    Height      N Z50   Z95
## 1     148 38.05782 827.3037 165.4297 100   600
## 2     168 14.82244 667.2506 384.0000 300 1000
##
## $shrubData
##   Species Height    Cover Z50   Z95
## 1     165 73.51455 3.045061 200 1000
##
## $herbCover
## [1] 10
##
## $herbHeight
## [1] 20
##
## attr(,"class")
## [1] "forest" "list"
```

## 4. Forest dynamics

### Forest dynamics output

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

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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:

```
fd$StandSummary
```

##	Step	NumTreeSpecies	NumTreeCohorts	NumShrubSpecies	NumShrubCohorts	TreeDensityLive	
## 1	0	2	2	1	1	552.0000	
## 2	1	2	2	1	1	551.1388	
## 3	2	2	2	1	1	550.2821	
## 4	3	2	2	1	1	549.4297	
##		TreeBasalAreaLive	DominantTreeHeight	DominantTreeDiameter	QuadraticMeanTreeDiameter		
## 1		25.03330	800.0000	37.55000		24.02949	
## 2		25.17084	809.1730	37.71939		24.11423	
## 3		25.30857	818.2975	37.88910		24.19893	
## 4		25.44491	827.3037	38.05782		24.28284	
##		HartBeckingIndex	ShrubCoverLive	BasalAreaDead	ShrubCoverDead	BasalAreaCut	ShrubCoverCut
## 1		53.20353	3.750000	0.00000000	0.00000000	0	0
## 2		52.64148	2.737354	0.09623075	0.004097987	0	0
## 3		52.09501	2.887898	0.09660091	0.004305995	0	0
## 4		51.56784	3.045061	0.09696355	0.004541953	0	0



## 4. Forest dynamics

### Forest dynamics output

... and *species-level* statistics are shown using:

```
head(fd$SpeciesSummary)
```

##	Step	Species	Name	NumCohorts	TreeDensityLive	TreeBasalAreaLive	ShrubCoverLive
## 1	0	148	Pinus halepensis	1	168.0000	18.604547	NA
## 2	0	165	Quercus coccifera	1	NA	NA	3.750000
## 3	0	168	Quercus ilex	1	384.0000	6.428755	NA
## 4	1	148	Pinus halepensis	1	167.1388	18.676546	NA
## 5	1	165	Quercus coccifera	1	NA	NA	2.737354
## 6	1	168	Quercus ilex	1	384.0000	6.494295	NA
##		BasalAreaDead	ShrubCoverDead	BasalAreaCut	ShrubCoverCut		
## 1		0.00000000	NA	0	NA		
## 2		NA	0.00000000	NA	0		
## 3		0.00000000	NA	0	NA		
## 4		0.09623075	NA	0	NA		
## 5		NA	0.004097987	NA	0		
## 6		0.00000000	NA	0	NA		

## 4. Forest dynamics

### Forest dynamics output

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

```
fd$TreeTable
```

##	Step	Year	Cohort	Species	Name	N	DBH	Height	Z50	Z95
## 1	0	NA	T1_148	148	Pinus halepensis	168.0000	37.55000	800.0000	100	600
## 2	0	NA	T2_168	168	Quercus ilex	384.0000	14.60000	660.0000	300	1000
## 3	1	2001	T1_148	148	Pinus halepensis	167.1388	37.71939	809.1730	100	600
## 4	1	2001	T2_168	168	Quercus ilex	384.0000	14.67423	662.4111	300	1000
## 5	2	2002	T1_148	148	Pinus halepensis	166.2821	37.88910	818.2975	100	600
## 6	2	2002	T2_168	168	Quercus ilex	384.0000	14.74849	664.8315	300	1000
## 7	3	2003	T1_148	148	Pinus halepensis	165.4297	38.05782	827.3037	100	600
## 8	3	2003	T2_168	168	Quercus ilex	384.0000	14.82244	667.2506	300	1000

**Note:** The NA values in Year correspond to the initial state.

## 4. Forest dynamics

### Forest dynamics output

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

```
fd$TreeTable
```

##	Step	Year	Cohort	Species	Name	N	DBH	Height	Z50	Z95
## 1	0	NA	T1_148	148	Pinus halepensis	168.0000	37.55000	800.0000	100	600
## 2	0	NA	T2_168	168	Quercus ilex	384.0000	14.60000	660.0000	300	1000
## 3	1	2001	T1_148	148	Pinus halepensis	167.1388	37.71939	809.1730	100	600
## 4	1	2001	T2_168	168	Quercus ilex	384.0000	14.67423	662.4111	300	1000
## 5	2	2002	T1_148	148	Pinus halepensis	166.2821	37.88910	818.2975	100	600
## 6	2	2002	T2_168	168	Quercus ilex	384.0000	14.74849	664.8315	300	1000
## 7	3	2003	T1_148	148	Pinus halepensis	165.4297	38.05782	827.3037	100	600
## 8	3	2003	T2_168	168	Quercus ilex	384.0000	14.82244	667.2506	300	1000

**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:

```
fd$DeadTreeTable
```

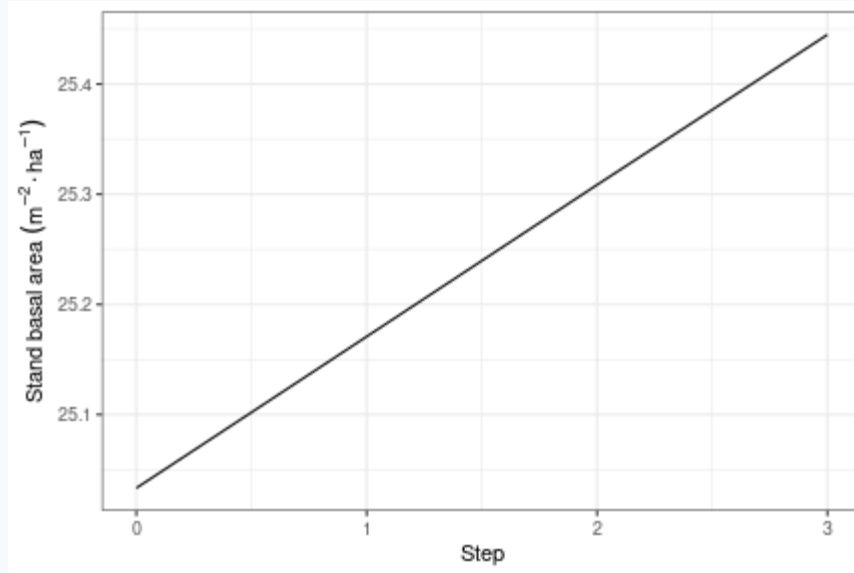
##	Step	Year	Cohort	Species	Name	N	DBH	Height	Z50	Z95
## 1	1	2001	T1_148	148	Pinus halepensis	0.8611814	37.71939	809.1730	100	600
## 2	2	2002	T1_148	148	Pinus halepensis	0.8567669	37.88910	818.2975	100	600
## 3	3	2003	T1_148	148	Pinus halepensis	0.8523751	38.05782	827.3037	100	600

## 4. Forest dynamics

### Summaries and plots

The provides a `plot` function for objects of class `fordyn`. For example, we can show the year-to-year variation in stand-level basal area using:

```
plot(fd, type = "StandBasalArea")
```

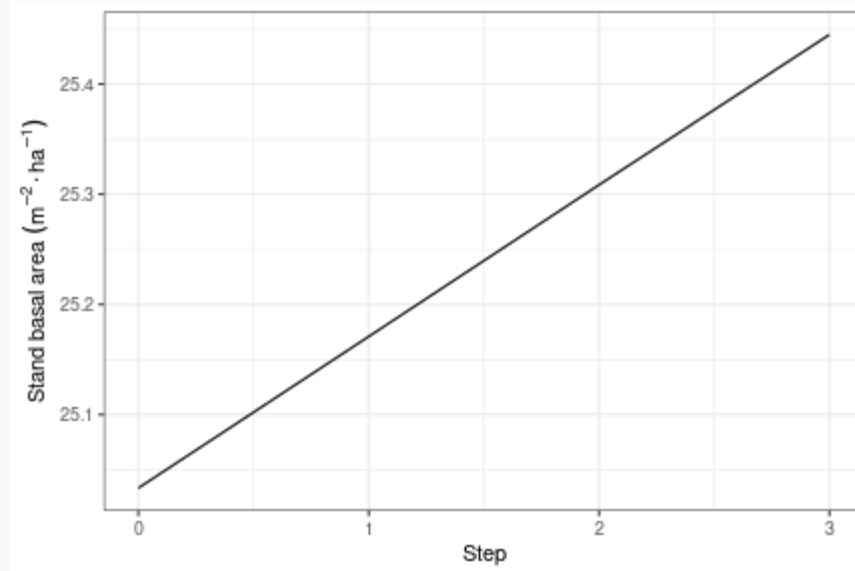


## 4. Forest dynamics

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```



These plots are based on the *annual summaries* included in the output.

## 4. Forest dynamics

### Summaries and plots

**Remember:** Function `fordyn()` makes internal calls to function `growth()` and stores the result in a vector called `GrowthResults`.

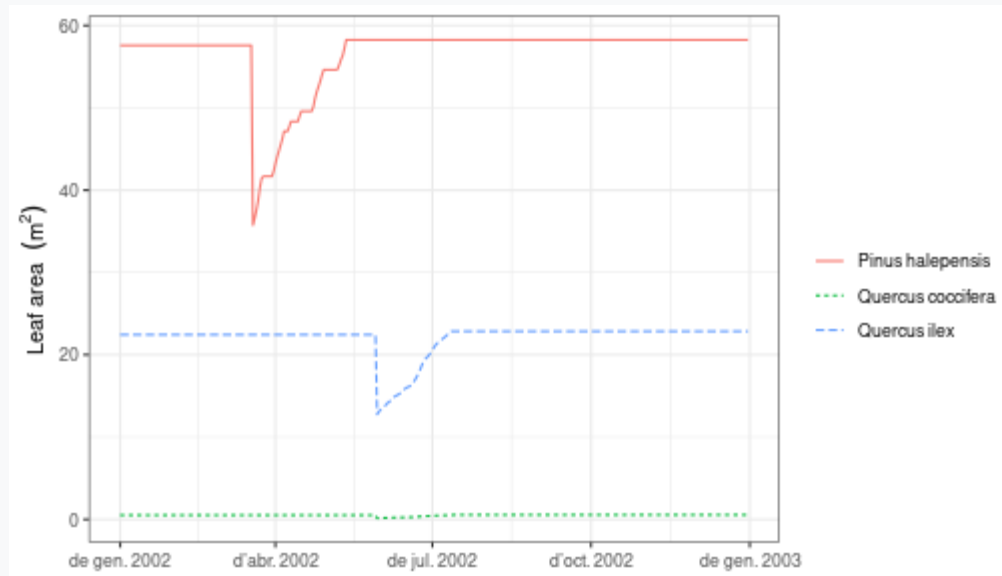
# 4. Forest dynamics

## Summaries and plots

**Remember:** Function `fordyn()` makes internal calls to function `growth()` and stores the result in a vector called `GrowthResults`.

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

```
plot(fd$GrowthResults[[2]], "LeafArea", bySpecies = T)
```

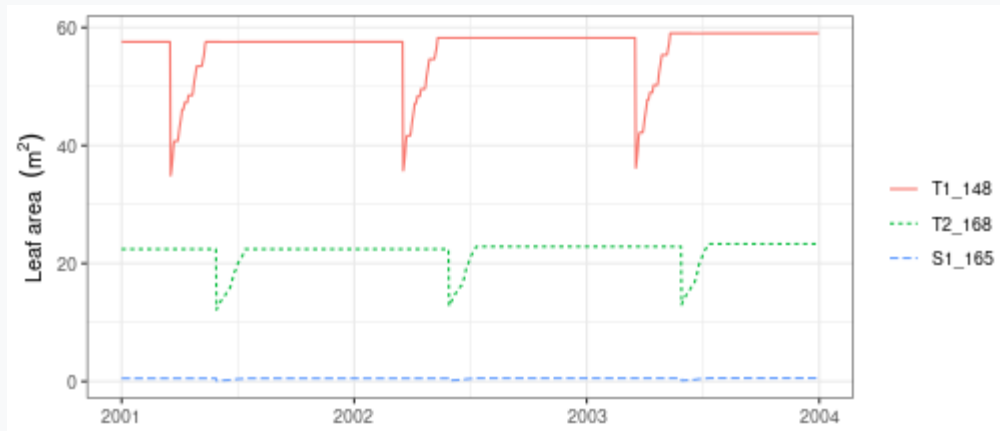


# 4. Forest dynamics

## Summaries and plots

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

```
plot(fd, "LeafArea")
```



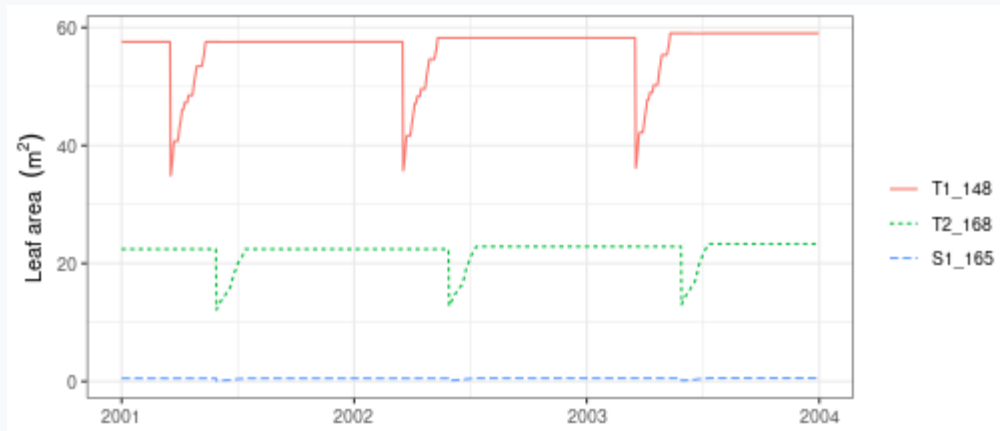


## 4. Forest dynamics

### Summaries and plots

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```



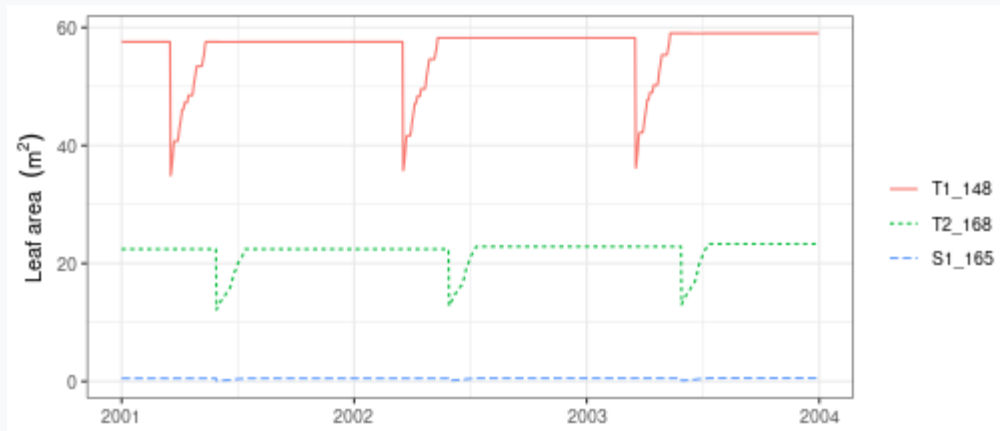
In this case, the `plot()` function assembles all the information from `GrowthResults` (accounting for cohort additions/deletions) and draws the plot.

## 4. Forest dynamics

### Summaries and plots

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

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```



In this case, the `plot()` function assembles all the information from `GrowthResults` (accounting for cohort additions/deletions) and draws the plot.

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

## 4. Forest dynamics

### Forest dynamics including management

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

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## 4. Forest dynamics

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To run simulations with management we need to define (and modify) management arguments (see ? `defaultManagementArguments()`).

```
# Default arguments
args <- defaultManagementArguments()
# Here one can modify defaults before calling fordyn()
#
```

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args <- defaultManagementArguments()
# Here one can modify defaults before calling fordyn()
#
```

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

```
fd<-fordyn(exampleforestMED, examplesoil, SpParamsMED, meteo, control,
           latitude = 41.82592, elevation = 100,
           management_function = defaultManagementFunction,
           management_args = args)
```

## 4. Forest dynamics

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```

When management is included, two additional tables are produced, e.g.:

```
fd$CutTreeTable
fd$CutShrubTable
```

## 4. Forest dynamics

### Forest dynamics including management

Function `defaultManagementArguments()` returns a list with default values for *management parameters*:

Element	Description
<code>type</code>	Management model, either 'regular' or 'irregular'
<code>thinning</code>	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
<code>thinningMetric</code>	The stand-level metric used to decide whether thinning is applied, either "BA" (basal area), "N" (density) or "HB" (Hart-Becking index)
<code>thinningThreshold</code>	The threshold value of the stand-level metric causing the thinning decision
<code>thinningPerc</code>	Percentage of stand's basal area to be removed in thinning operations
<code>minThinningInterval</code>	Minimum number of years between thinning operations
<code>finalMeanDBH</code>	Mean DBH threshold to start final cuts
<code>finalPerc</code>	String with percentages of basal area to be removed in final cuts, separated by '-' (e.g. "40-60-100")
<code>finalYearsBetweenCuts</code>	Number of years separating final cuts



## 4. Forest dynamics

### Forest dynamics including management

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 occurred
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.

## 4. Forest dynamics

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finalYearsToCut	State variable to count the years to be passed before new final cut is applied.

**Remember:** Besides using the in-built management function, you could program your own management function and specify its own set of parameters.

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

