3.3 - Forest growth/dynamics (exercise)

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Objectives

- 1. Learn to perform simulations of forest growth and forest dynamics with medfate
- 2. Evaluate tree growth predictions with tree ring data
- 3. Compare simulated vs observed forest changes between inventories
- 4. Project forest dynamics with/without forest management



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Data

We will use data corresponding to a forest plot of sampled during the third and fourth Spanish National Forest Inventory (SNFI3) in the province of Tarragona (latitude 41° N aprox.).

• The forest plot is dominated by Aleppo pine (*Pinus halepensis*) with an understory of composed of several shrub species.



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- Tree ring data are available for some trees of the forest plot, because it was included in a research project focused on intraspecific variability of functional traits (FUN2FUN, granted to J. Martínez-Vilalta).
- Soil has been already drawn from SoilGrids
- Daily weather data corresponding to the plot location has been obtained with **meteoland**, corresponding to an historical period (SNFI3-SNFI4) and a future period (2015-2100) under scenario RCP 8.5 (from Earth system model MPI-ESM regionalized to Europe using model RCA4).



Step 1. Load Alepo pine forest data

We are given all the necessary data, bundled in a single list:

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alepo <- readRDS("StudentRdata/alepo.rds")</pre>
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Whose elements are...

Element	Description
forest_snfi3	Object of class forest with the stand structure and composition in SNFI3 (yr. 2001)
forest_snfi4	Object of class forest with the stand structure and composition in SNFI4 (yr. 2014)
spt	Object of class SpatialPointsTopography with the coordinates and topography of the plot
soildesc	Data frame with soil properties.
historic_weather	Data frame with daily weather for years 2001-2014.
projected_weather	Data frame with daily weather for years 2015-2100 under RCP8.5 (climate model couple MPIESM/RCA4).
observed_growth	Data frame with annual basal area increments during the 2001-2014 period for four <i>P. halepensis</i> trees in the forest plot (T20_148, T14_148, T25_148 and T3_148).
snfi34_growth	Data frame with density, diameter and height for <i>P. halepensis</i> as measured in SNFI3 and SNFI3.



Step 2. Forest stand metrics

We can use the summary () function for objects of class forest to know the leaf area index and basal area estimated at yr. 2001 (SNFI3):

```
summary(alepo$forest_snfi3, SpParamsMED)

## Tree density (ind/ha): 721.50240945

## Tree BA (m2/ha): 21.5278871

## Cover (%) trees (open ground): 100 shrubs: 100

## Shrub crown phytovolume (m3/m2): 1.04

## LAI (m2/m2) total: 3.6639431 trees: 1.4241149 shrubs: 2.2398282

## Live fine fuel (kg/m2) total: 1.5337579 trees: 0.5444354 shrubs: 0.9893226

## PAR ground (%): 14.5677246 SWR ground (%): 24.0043619
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```

The contribution of the different species to these stand metrics can be known using:

```
species_basalArea(alepo$forest_snfi3, SpParamsMED)
    Pinus halepensis Quercus coccifera Pistacia lentiscus Salvia rosmarinus
                                                                                  Frica multiflora
##
##
             21.52789
                                 0.00000
                                                    0.00000
                                                                        0.00000
                                                                                           0.00000
species_LAI(alepo$forest_snfi3, SpParamsMED)
    Pinus halepensis Quercus coccifera Pistacia lentiscus Salvia rosmarinus
                                                                                  Erica multiflora
##
##
            1.4241149
                               0.2774996
                                                  0.3928101
                                                                      1.3935065
                                                                                         0.1760121
```



Step 2. Forest stand metrics

We repeat the same calculations for yr. 2014 (SNFI4):

```
summary(alepo$forest_snfi4, SpParamsMED)

## Tree density (ind/ha): 707.35530341

## Tree BA (m2/ha): 27.5720378

## Cover (%) trees (open ground): 100 shrubs: 100

## Shrub crown phytovolume (m3/m2): 1.133

## LAI (m2/m2) total: 4.6079012 trees: 1.5995943 shrubs: 3.0083069

## Live fine fuel (kg/m2) total: 1.6496117 trees: 0.6115207 shrubs: 1.038091

## PAR ground (%): 8.6749798 SWR ground (%): 16.3505438
```

There has been an increase of 6 m2/ha in basal area, whereas stand LAI has increased 0.94 m2/m2.



Step 3. Growth simulation between SNFI3 and SNFI4

We were given soil physical characteristics, but we need to build an object of class soil, which we can store in the same alepo list:

```
alepo$soil <- soil(alepo$soildesc)</pre>
```



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```
alepo$soil <- soil(alepo$soildesc)
```

we can check the water holding capacity of the soil using:

```
sum(soil_waterFC(alepo$soil))
## [1] 391.1652
```

which is rather high but we leave it as is.



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which is rather high but we leave it as is.

We now have all the elements to call function forest2growthInput() to generate the input for growth():



Step 3. Growth simulation between SNFI3 and SNFI4

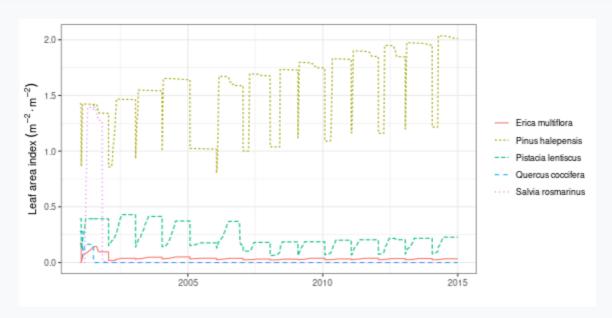
Since the list contains also the historic weather for years 2001-2014 and topography, we are ready to simulate growth:



Step 4. Examine growth results

Many outputs can be inspected using shinyplot() but here we use plot() to display the LAI dynamics of the different species

```
plot(G_34, "PlantLAI", bySpecies = TRUE)
```



The model predicts an increase in LAI for *P. halepensis* (except some years), but shrub species are predicted to lose leaf area.



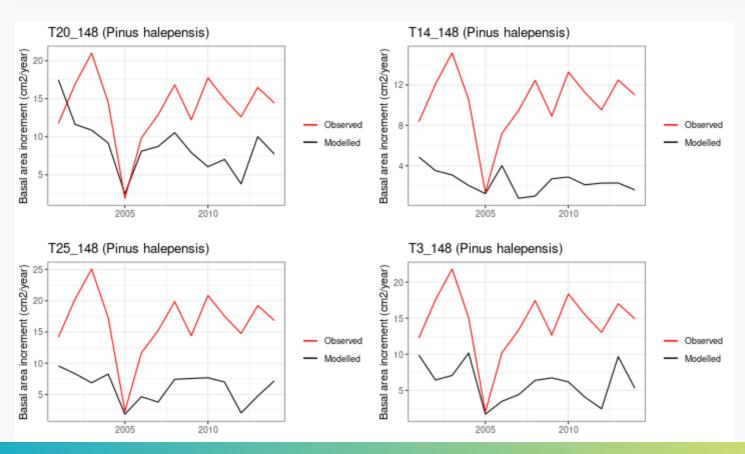
Step 5. Evaluate tree basal area increment

We can use function evaluation_plot() to display the predicted and observed BAI for the four trees with measurements:



Step 5. Evaluate tree basal area increment

When we display the plots we see that the model is overestimating growth in many cases:





Step 5. Evaluate tree basal area increment

Tip: To decide how to proceed when a model fails to fit observations is important to know which model parameters may be responsible for a given result (this is called *sensitivity analysis*).



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In medfate, sapwood (and tree) growth is strongly controlled by parameter RGRcambiummax, which specifies the maximum growth rate of sapwood relative to stem diameter.



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In medfate, sapwood (and tree) growth is strongly controlled by parameter RGRcambiummax, which specifies the maximum growth rate of sapwood relative to stem diameter.

For *P. halepensis* its default value is:

```
SpParamsMED$RGRcambiummax[SpParamsMED$Name=="Pinus halepensis"]
```

[1] 0.003688597



Step 6. Modify maximum growth rate for P. halepensis and repeat simulations

We divide the maximum relative growth rate by two...

```
SpParamsMED$RGRcambiummax[SpParamsMED$Name=="Pinus halepensis"] <- 0.0012</pre>
```



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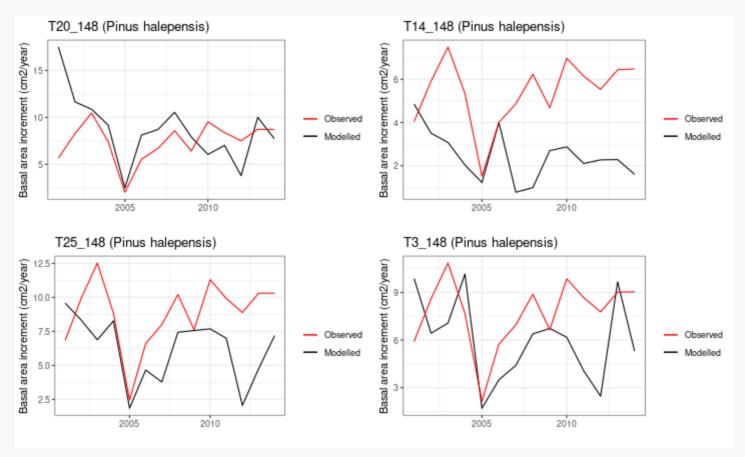
... rebuild the growth input ...

... and launch a new simulation:



Step 6. Modify maximum growth rate for P. halepensis and repeat simulations

We can inspect the fit of the new results to observed data. Overall, we obtain a better fit in terms of the mean BAI, but the model does not capture all observed interannual variation.





Step 7. Reduce the number of tree cohorts

In order to speed-up forest dynamic simulations, we can reduce the number of tree cohorts, which is now:

```
nrow(alepo$forest_snfi3$treeData)
```

[1] 28



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```
nrow(alepo$forest_snfi3$treeData)
## [1] 28
```

Remembering the forest_mergeTrees() function from exercise #1:

```
forest_red = forest_mergeTrees(alepo$forest_snfi3)
```



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Remembering the forest mergeTrees() function from exercise #1:

```
forest_red = forest_mergeTrees(alepo$forest_snfi3)
```

The new forest object has 5 tree cohorts:

2

```
forest_red$treeData
    Species
                           DBH
                                   Height
                                               Z50 Z95
## 1
        148 14.14711 31.60000 1400.0000 522.4242 4000
        148 198.05948 25.38220 1100.2943 522.4242 4000
```

```
148 159.15494 20.49330 936.3455 522.4242 4000
## 4
       148 222.81692 14.62423 809.2011 522.4242 4000
## 5
       148 127.32395 11.85000 820.0000 522.4242 4000
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```

In the following, we will use forest_red to call function fordyn().



Step 8. Run forest dynamics simulation

Remember: unlike spwb() and growth(), we do not need to build an intermediate input object for fordyn() (i.e., there is no function forest2fordynInput()).



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The elements of the output have the following names, which we should be able to understand before moving on (if not, see ?fordyn).

```
names(FD_34)

## [1] "StandSummary" "SpeciesSummary" "CohortSummary" "TreeTable" "DeadTreeTable"

## [6] "CutTreeTable" "ShrubTable" "DeadShrubTable" "CutShrubTable" "ForestStructures"

## [11] "GrowthResults" "ManagementArgs" "NextInputObject" "NextForestObject"
```



Step 9. Compare final stand metrics with the observed stand in SNFI4

In particular, we can examine the stand metrics of the forest object at the end of the simulation...

```
summary(FD_34$NextForestObject, SpParamsMED)

## Tree density (ind/ha): 919.946082936554

## Tree BA (m2/ha): 26.2644408

## Cover (%) trees (open ground): 100 shrubs: 40.3431789

## Shrub crown phytovolume (m3/m2): 0.2695373

## LAI (m2/m2) total: 2.3960396 trees: 1.6132901 shrubs: 0.7827495

## Live fine fuel (kg/m2) total: 1.0456124 trees: 0.6167566 shrubs: 0.4288558

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... and compare them to those obtained in SNFI4 (yr. 2015) for the forest plot:
summary(alepo$forest_snfi4, SpParamsMED)
## Tree density (ind/ha): 707.35530341
## Tree BA (m2/ha): 27.5720378
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The model seems to perform fairly well in terms of final tree density and basal area. However, as expected, it yields too much shrub mortality, resulting in a forest with a low understory biomass.



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Hence, we can use this feature to start our projection from the final state of the previous call to fordyn() and use the projected daily weather:

The predicted final stand basal area is:

```
stand_basalArea(FD_proj$NextForestObject)
## [1] 52.1008
```



Step 11. Management function and management arguments

We will now simulate forest dynamics including forest management.



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However, we need first to understand how the default management function works and the meaning of its parameters:

```
man_args <- defaultManagementArguments()</pre>
names(man_args)
    [1] "type"
                                  "targetTreeSpecies"
                                                            "thinning"
## [4] "thinningMetric"
                                  "thinningThreshold"
                                                            "thinningPerc"
   [7] "minThinningInterval"
                                  "yearsSinceThinning"
                                                            "finalMeanDBH"
## [10] "finalPerc"
                                  "finalPreviousStage"
                                                            "finalYearsBetweenCuts"
## [13] "finalYearsToCut"
                                  "plantingSpecies"
                                                            "plantingDBH"
## [16] "plantingHeight"
                                  "plantingDensity"
                                                            "understoryMaximumCover"
```



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## [13] "finalYearsToCut"
                                  "plantingSpecies"
                                                            "plantingDBH"
## [16] "plantingHeight"
                                  "plantingDensity"
                                                            "understoryMaximumCover"
```

Argument thinningThreshold specifies the stand basal area value that leads to a thinning event. Since our simulation started at 26 m2/ha and increased up to 52 m2/ha, we set the value of thinningThreshold to 30 m2/ha to see some effects during the simulations:

```
man_args$thinningThreshold <- 30
```



Step 12. Projection of forest dynamics with management

The call to fordyn() is similar to the previous one, except for the specification of the management function and parameters:

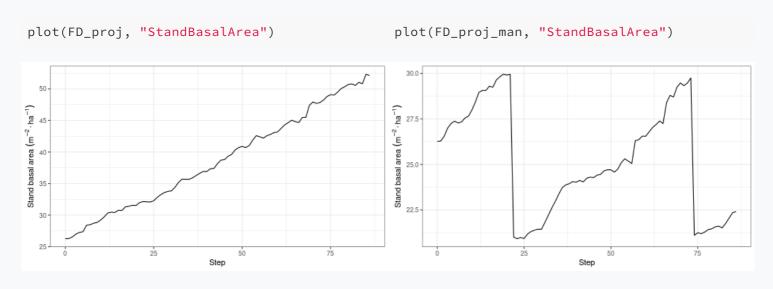


Step 13. Compare forest dynamics with/without management

We can produce plots of stand basal area dynamics to compare the two simulations:

No management

Management





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We can produce plots of stand basal area dynamics to compare the two simulations:

No management plot(FD_proj, "StandBasalArea") plot(FD_proj_man, "StandBasalArea")

Generally speaking, the arguments thinningThreshold and thinningPerc control the frequency and intensity of thinning interventions.



Step 13. Compare forest dynamics with/without management

We can also compare the final tree data frames of the forest objects of the two simulations:

No management

FD_proj\$NextForestObject\$treeData[,1:4]

```
Species
                          Height
##
                   DBH
         148 46.242455 1649.2031
                                   8.459126
## 2
        148 40.009616 1520.4876 118.427760
## 3
        148 35.124450 1463.5371 95.165164
## 4
        148 29.258967 1425.4242 133.231230
## 5
        148 26.490240 1430.9978 76.132131
## 6
        148 21.241281 1338.1708 159.718591
## 7
        148 19.273174 1281.3398 167.667172
## 8
        148 9.760524 768.7657 218.076709
         148 7.587567 577.5869 272.595454
## 9
```

Management

```
FD_proj_man$NextForestObject$treeData[,1:4]
```

```
## Species DBH Height N
## 1 148 47.14512 1657.693 8.459126
## 2 148 40.70770 1525.359 118.427760
## 3 148 36.02478 1469.454 54.184731
```



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```
FD_proj$NextForestObject$treeData[,1:4]
                                                    FD_proj_man$NextForestObject$treeData[,1:4]
    Species
                         Height
                                                        Species
                                                                           Height
##
                  DBH
                                                                     DBH
                                                                                           Ν
        148 46.242455 1649.2031
                                  8,459126
                                                   ## 1
                                                            148 47.14512 1657.693
        148 40.009616 1520.4876 118.427760
                                                            148 40.70770 1525.359 118.427760
## 2
                                                   ## 2
                                                   ## 3
## 3
        148 35.124450 1463.5371 95.165164
                                                            148 36.02478 1469.454 54.184731
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        148 19.273174 1281.3398 167.667172
## 8
        148 9.760524 768.7657 218.076709
        148 7.587567 577.5869 272.595454
## 9
```

The number of tree cohorts is much lower at the end of the simulation with forest management because by default the thinning is specified to be applied to small trees (i.e. thinning = "below").



Step 13. Compare forest dynamics with/without management

Finally, we can use the annual summaries produced by fordyn() to compare the basal area of trees dead or cut during the simulation:

No management

sum(FD_proj\$StandSummary\$BasalAreaDead) ## [1] 17.20003 sum(FD_proj\$StandSummary\$BasalAreaCut) ## [1] 0

Management

```
sum(FD_proj_man$StandSummary$BasalAreaDead)
## [1] 11.26269
sum(FD_proj_man$StandSummary$BasalAreaCut)
## [1] 18.05398
```



No management

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Management

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## [1] 17.20003	## [1] 11.26269
<pre>sum(FD_proj\$StandSummary\$BasalAreaCut)</pre>	<pre>sum(FD_proj_man\$StandSummary\$BasalAreaCut)</pre>
## [1] 0	## [1] 18.05398

The simulation without forest management produced more dead trees than the simulation with management.



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## [1] 17.20003	## [1] 11.26269
<pre>sum(FD_proj\$StandSummary\$BasalAreaCut)</pre>	<pre>sum(FD_proj_man\$StandSummary\$BasalAreaCut)</pre>
## [1] 0	## [1] 18.05398

The simulation without forest management produced more dead trees than the simulation with management.

This arises because:

- Basal mortality rates are multiplied by the current tree density
- Drought stress is decreased in simulations with management

M.C. Escher - Concave and convex, 1955

