

Translation of SSM into R

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Objectives

- no dependency on Excel (R is open-source and cross-platform), except for input files that come from the original version (to allow easy re-use of existing inputs and comparisons between versions)... but possibility to enter inputs in another format
- possibility to run from a website
- possible to run in batch mode, and to run several simulations (i.e. several locations-climates-crop managements) at the same time

Desired improvements of the code over the original SSM.xls code

- use the same code for all crops (in excel version, the code is repeated 3 times (wheat, maize and legumes), with subtle differences, which makes the code (i) difficult to read and understand, (ii) difficult to maintain (one has to make the same changes in several places in the code)=> use switches given in the crop parameter file to activate or not crop-specific parts, the rest of the code stays the same
- remove all hard-coded parameters (including species names and stage names) => put them in the appropriate parameter files (crops or GeneralPhysicalParameters)
- use more readable names of variables to facilitate code understanding =>use full names instead of acronyms.
- add modularity (to help code maintenance and future developments) => divide the code into different procedure (put them in different files?).

Architecture of the model

The model is "encapsulated" so that function definitions and model variables are not laying around in the user's workspace. The setup function creates an object (here, named MyModel, but it could be anything else), that contains an instance of the model and that can be manipulated by handler functions (defined in file handlers.R), e.g. functions to set the simulation options, to run the model for n time steps, to plot variables etc... Within the object, there are several "global" variables (named in uppercase), procedures (i.e. R functions that have no arguments and return no value, but access and modify global variables), and functions (which are called by procedures to compute different intermediary variables that are used by procedures). All intermediary variables and state variables, as well as input parameters (e.g. climate variables and crop parameters (which vary according to the crop currently present)), are saved at each time step (the current time step is a dataframe named ALLDAYDATA, which is updated by each procedure, and at the end of the timestep, this dataframe is appended to a list (ALLSIMULATEDDATA) with one element per timestep.

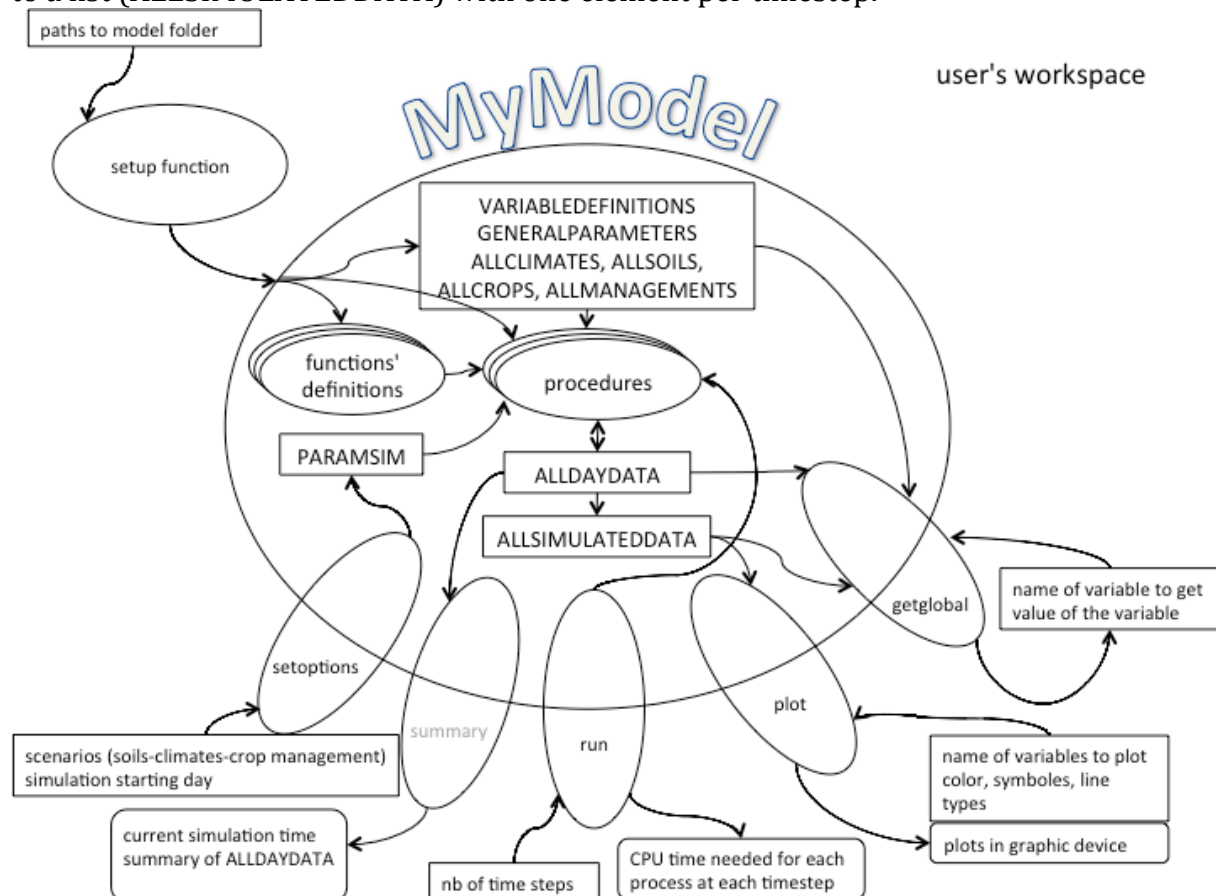


Figure 1: the setup function takes as argument the path to the model folder and returns an R object with its own environment, separated from the user's workspace. This object (here called "MyModel") can be manipulated by handler functions, for example MyModel\$run(100) will run the model for 100 timesteps. Ovals: functions; squares: vectors, dataframes or lists; rounded squares: outputs to the console or graphic device

Naming conventions

UPPERCASE: "global" (i.e. within the model) variables, accessible to all procedures

lowercase: local (i.e. within each function or procedure) variable

#variable names

#iXXXX: Input variable (e.g. iPr : Precipitation)

#pXXXX: Parameter (e.g. pPhyllochron)

#cXXXX: Computed variable (e.g. cDeltaLAI): variable that is computed from scratch at each time step
 #sXXXX: State variable (e.g. sLAI): variable that keeps its value from one time step to the beginning of the next
 #function names
 #fXXXX: Function (e.g. fPhotosynthesis)
 #rXXXX: procedure (e.g. rUpdateLAI)
 #eXXXX: read input scenarios from External files (e.g. eReadInInputs)
 #mXXXX: Model interface, i.e. handler functions (e.g. mRun, mPlotDynamics)

Model algorithm

- At the beginning of the simulation, a first element (corresponding to day 0) is created in ALLSIMULATEDDATA, with state variables initialised at 0.
- At the beginning of each timestep, a new ALLDAYDATA dataframe is created, with initial values as specified in ALLVARIABLES (read from sheet "savedeachday" of the excel file allvariables.xlsx, which contains the definition, unit, original SSM name and initial value of all variables, usually NA for computed variables). Then state variables (name starting with s) are initialised at their value from the previous time step. Then the procedure (name starting with r) corresponding to each module is called sequentially.
- In each procedure, local computed variables (name starting with c) are created using different functions (name starting with f), using as arguments either state or computed variables (from ALLDAYDATA). At the end of the procedure, ALLDAYDATA is updated with all state variables (which will be used to initialise their values for the next time step) as well as parameters and computed variables (to allow access to intermediary variables when analysing the model outputs).
- Each function corresponds to the computation of one intermediary variable in the SSM code, we tried to keep the code as similar to the original SSM code as possible... notable deviations from the initial code are given in section "Details on the deviations from the original SSM model, by module", page 12

Structure of the global variables

PARAMSIM

List of 8

```
$ simustart : Date[1:1], format: "1997-11-01"
$ cases : 'data.frame': 2 obs. of 4 variables:
..$ climatename: chr [1:2] "Ain Hamra - Meknes" "Ain Hamra - Meknes"
..$ soilname : chr [1:2] "325_-35" "325_-35"
..$ lat : num [1:2] 35 45
..$ long : num [1:2] -5 -5
$ directory : chr "/Users/user/Documents/b_maison/congeMat/D4DECLIC/runSSM"
$ climateformat: chr "standardSSM"
$ cropformat : chr "standardSSM"
$ soilformat : chr "standardSSM"
$ managformat : chr "standardSSM"
$ Neffect : logi TRUE
```

simustart is the day when the simulation should start (used to find the climate data for this day) ; cases is a data.frame with n rows (=n cases= n combinations of climate, soil, geographic coordinates (and in the future crop management) ; directory is the path to

the folder where your input folder, containing your input (soil, climate, crop parameter, crop management) files are placed (and where the output folder will be created to store graphs or export files) ; climateformat, cropformat, soilformat and managformat are the formats in which these input files are (for now only "standardSSM" is available, it was planned to include other formats to allow automatic link with external climate and soil databases, but not done yet) ; Neffect is a boolean indicating which version of the model to use: with or without N effect on LAI decrease (for now, only the version with N effect is coded).

ALLDAYDATA

'data.frame': 2 obs. of 111 variables:

```
$ iDate          : Date, format: "1997-11-04" "1997-11-04"
$ sAccumulatedLeafDryMatter      : num  0 0
$ sAccumulatedStemDryMatter      : num  0 0
$ sAccumulatedVegDryMatter       : num  0 0
$ sLAI              : logi  NA NA
$ sMainstemNodeNumber           : num  1.72 1.72
$ sPlantdensity             : num  280 280
$ sCrop                  : chr  "WHEAT" "WHEAT"
$ sCultivar              : chr  "durum wheat" "toto"
$ sBiologicalDay           : num  2.88 2.88
$ sGrowthStage            : chr  "germination" "germination"
$ sThermalUnit            : num  79.2 79.2
$ sRootFrontDepth         : num  0 0
```

etc... (all variables listed in sheet "savedeachday" of file allvariables.xlsx)

rows are cases (=combinations of climate-soil-management)

ALLSIMULATEDDATA

list of n data.frames (ALLDAYDATA at the end of the time step, one for each timestep (plus day 0)).

VARIABLEDEFINITIONS

this data.frame is read from sheet "savedeachday" in file "allvariables.xlsx" from the model folder.

'data.frame': 111 obs. of 9 variables:

```
$ name          : chr  "sAccumulatedLeafDryMatter" "sAccumulatedStemDryMatter"
"sAccumulatedVegDryMatter" "sLAI" ...
$ typeinthemodel : chr  "stateVariable" "stateVariable" "stateVariable" "stateVariable"
...
$ typeR         : chr  "numeric" "numeric" "numeric" "numeric" ...
$ module        : chr  "DMDistribution" "DMDistribution" "DMDistribution" "LAI" ...
$ unit          : chr  "g m-2" "g m-2" "g m-2" "m2 m-2" ...
$ definition     : chr  "Accumulated leaf dry matter" "Accumulated Steam dry matter"
"Accumulated vegetative (leaf + stem) dry matter" "Leaf area index" ...
$ translationSSM : chr  "WLF" "WSF" "WVEG" "LAI" ...
```

```
$ french      : chr "Stock de matière sèche dans les feuilles" "Stock de matière sèche
dans les parties végétatives" "Stock de matière sèche dans les tiges" "Index de surface
foliaire" ...
$ defaultInitialvalue: num 0 0 0 0 0 0 0 0 0 ...
```

ALLCROPS

data.frame with rownames crop.cultivar, and with parameters in columns

It is created by reading sheet allcrops of file crops.xlsx in the input folder of the simulation folder (NOT the model folder, which contains example input files, which are not read). Warning: compared with the sheet Crop in GW_SSM_iCrop_TOTAL_modiflog.xlsm, I had to add parameters TBVER, TP1VER, TP2VER, TCVER, VDSAT from the excel file where crops are in different sheets, and the parameters for stage durations (and stage names) and filters for case-specific processes. Parameters in the R version have more readable names than in the excel version, and the translation is done thanks to the file allvariables.xlsx, which contains the R names and the SSM names. In the following export, some parameters have their R name and type, and some are still as they were read from the crop parameter file, the later are the ones that are not yet in allvariables.xlsx because they aren't used in the code yet (code not finished). And the filters are not correct yet.

'data.frame': 6 obs. of 85 variables:

```
$ CROP      : chr "WHEAT" "WHEAT" "WHEAT" "Chickpea" ...
$ Cultivar   : chr "Ble_Dur_1" "Ble_Tendre_1" "Ble_Tendre_2" "Ghab2" ...
$ pPhyllochron : num 118 110 110 46 0 0
$ pcoefPlantLeafNumberNode : num 1 1 1 1 0 0
$ pExpPlantLeafNumberNode : num 2.5 2.34 2.34 2 0 0
$ a_plapow_d : chr "1" "1" "1" "1" ...
$ b_plapow_d : chr "0" "0" "0" "0" ...
$ pSpecificLeafArea : num 0.02 0.02 0.02 0.0161 0 0
$ pFreezeThresholdTemp : num -5 -5 -5 -5 0 0
$ pFreezeFracLeafDestruction : num 0.01 0.01 0.01 0 0 0
$ pHeatThresholdTemp : num 30 30 30 0 0 0
$ pHeatFracLeafDestruction : num 0.1 0.1 0.1 0 0 0
$ pTbasRUE : num 0 0 0 2 0 0
$ pTopt1RUE : num 15 15 15 14 0 0
$ pTopt2RUE : num 22 22 22 30 0 0
$ plethalRUE : num 35 35 35 38 0 0
$ pKPAR : num 0.65 0.65 0.65 0.65 0 0
$ pRadEfficiencyOptimal : num 2.2 2.2 2.2 1.8 0 0
$ CO2RUE : chr "0.8" "0.8" "0.8" "0.8" ...
$ pParCoefLeafAeraLow : num 0.6 0.6 0.6 0.53 0 0
$ pParCoefLeafAeraHigh : num 0.3 0.3 0.3 0.3 0 0
$ pTotalCropMassFromCoefLeafAreaLowHigh : num 160 160 160 180 0 0
$ pParCoefLeafAeraTerminationLeafGrow : num 0.1 0.1 0.1 0.13 0 0
$ pFractionCropMassTranslocateable : num 0.22 0.22 0.22 0.22 0 0
$ pGrainConversionCoefficient : num 1 1 1 1 0 0
$ pPotentialSlopeHarvestIndex : num 0.008 0.01 0.01 0.01 0 0
$ pPotentialSlopeHarvestIndexCriticalPoint1: num 0 0 0 0 0 0
$ pPotentialSlopeHarvestIndexCriticalPoint2: num 600 600 600 0 0 0
$ pPotentialSlopeHarvestIndexCriticalPoint3: num 1200 1200 1200 9999 0 ...
```

\$ pPotentialSlopeHarvestIndexCriticalPoint4: num 3200 3200 3200 9999 0 ...
 \$ iDEPORT : chr "200" "200" "200" "200" ...
 \$ pMaxDepthWaterExtraction : num 1000 1000 1000 1000 0 0
 \$ pPotentialRootGrowth : num 30 30 30 17 0 0
 \$ pTranspirationEfficiencyLinkedToCO2 : num 5.8 5.8 5.8 5 0 0
 \$ TEC700 : chr "0" "0" "0" "0" ...
 \$ WSSG : chr "0.3" "0.3" "0.3" "0.3" ...
 \$ WSSL : chr "0.5" "0.40000000000000000008"
 "0.40000000000000000008" "0.4" ...
 \$ WSSD : chr "0.4" "0.40000000000000000008"
 "0.40000000000000000008" "0" ...
 \$ WSSN : chr "0" "0" "0" "0.5" ...
 \$ FLDKL : chr "20" "20" "20" "20" ...
 \$ pTbaseVernalization : num -1 -1 -1 NA -1 -1
 \$ pTopt1Vernalization : num 0 0 0 NA 0 0
 \$ pTopt2Vernalization : num 8 8 8 NA 8 8
 \$ pTlethalVernalization : num 12 12 12 NA 12 12
 \$ pVDSAT : num 50 50 50 NA 50 50
 \$ pSpecLeafNGreenLeaf : num 1.8 1.5 1.5 2.3 0 0
 \$ pSpecLeafNSenescenceLeaf : num 0.4 0.4 0.4 0.68 0 0
 \$ SNCG : chr "1.4999999999999999E-2" "1.4999999999999999E-2"
 "1.4999999999999999E-2" "1.37E-2" ...
 \$ SNCS : chr "5.10000000000000004E-3" "5.00000000000000001E-3"
 "5.00000000000000001E-3" "3.8999999999999998E-3" ...
 \$ GNCmin : chr "2.0899999999999998E-2" "2.1299999999999999E-2"
 "2.1299999999999999E-2" "3.37000000000000001E-2" ...
 \$ GNCmax : chr "2.0899999999999998E-2" "2.1299999999999999E-2"
 "2.1299999999999999E-2" "3.37000000000000001E-2" ...
 \$ MXNUP : chr "0.25" "0.25" "0.25" "0.45" ...
 \$ pTbasedev : num 0 0 0 2 0 0
 \$ pTopt1dev : num 27.5 27.5 27.5 21 0 0
 \$ pTopt2dev : num 27.5 27.5 27.5 30 0 0
 \$ pTlethaldev : num 40 40 40 40 0 0
 \$ pVernalizationSensitivity : num 0.002 0.008 0.008 NA 0 0
 \$ pCriticalPhotoPerdiod : num 14 21 21 0 0 0
 \$ pPhotoPeriodSensitivity : num 0.17 0 0 0.05 0 0
 \$ bdSOWEMR : chr "6" "4" "4" "8.5" ...
 \$ bdEMRTIL : chr "5" "9.4925465837500003" "11.6" "36" ...
 \$ bdTILSEL : chr "8" "7.7740683243499999" "9.5" "5.7" ...
 \$ bdSELBOT : chr "6" "2.5822646555555551" "3.1555555555" "4.3" ...
 \$ bdBOTEAR : chr "6" "0.64556843888871" "0.788888888875" "22" ...
 \$ bdEARANT : chr "15" "3.1555555555555555" "3.1555555555" "7" ...
 \$ bdANTPM : chr "43" "40.4" "37.6" "0" ...
 \$ bdPMHM : chr "8" "8" "8" "0" ...
 \$ bdBRP : chr "0" "0" "0" "8.5" ...
 \$ bdTRP : chr "0" "0" "0" "76.5" ...
 \$ bdBSG : chr "0" "0" "0" "54.5" ...
 \$ bdTSG : chr "0" "0" "0" "76.5" ...
 \$ bdTLM : chr "0" "0" "0" "52.5" ...

```

$ bdTLP : chr "0" "0" "0" "60.1" ...
$ bdBLS : chr "0" "0" "0" "54.5" ...
$ bdBNF : chr "0" "0" "0" "12" ...
$ thresholds :List of 6
..$: Named num 6 5 8 6 6 ...
..- attr(*, "names")= chr "germination" "emergence" "tillering" "stemElongation" ...
..$: Named num 4 9.493 7.774 2.582 0.646 ...
..- attr(*, "names")= chr "germination" "emergence" "tillering" "stemElongation" ...
..$: Named num 4 11.6 9.5 3.156 0.789 ...
..- attr(*, "names")= chr "germination" "emergence" "tillering" "stemElongation" ...
..$: Named num 8.5 36 5.7 4.3 22 ...
..- attr(*, "names")= chr "germination" "emergence" "tillering" "stemElongation" ...
..$: Named num 0 0 0 0 0 ...
..- attr(*, "names")= chr "germination" "emergence" "tillering" "stemElongation" ...
..$: Named num 0 0 0 0 0 ...
..- attr(*, "names")= chr "germination" "emergence" "tillering" "stemElongation" ...
$ vernalisation.filter : chr "is.after('emergence', 0) &
is.before('stemElongation', 0)" "is.after('emergence', 0) & is.before('stemElongation', 0)"
"is.after('emergence', 0) & is.before('stemElongation', 0)" "FALSE" ...
$ photoperiod.filter : chr "FALSE" "FALSE" "FALSE" "FALSE" ...
$ waterstress.filter : chr "is.after('emergence', 0) & is.before('senescence',
10)" "is.after('emergence', 0) & is.before('senescence', 10)" "is.after('emergence', 0) &
is.before('senescence', 10)" "is.after('emergence', 0) & is.before('senescence', 10)" ...
$ LAI_Mainstem.filter : chr "is.after('germination', 0) & is.before('Booting',
0)" "is.after('germination', 0) & is.before('Booting', 0)" "is.after('germination', 0) &
is.before('Booting', 0)" "is.after('germination', 0) & is.before('Booting', 0)" ...
$ LAI_Secondary.filter : chr "is.after('Booting', 0) & is.before('earring', 5)"
"is.after('Booting', 0) & is.before('earring', 5)" "is.after('Booting', 0) & is.before('earring',
5)" "is.after('Booting', 0) & is.before('earring', 5)" ...
$ DMDistribution_SeedGrowing.filter : chr "is.after('anthesis', 5) &
is.before('anthesis', 41.5)" "is.after('anthesis', 5) & is.before('anthesis', 38.9)"
"is.after('anthesis', 5) & is.before('anthesis', 36.1)" "is.after('anthesis', 5) &
is.before('anthesis', -1.5)" ...
$ DMProduction.filter : chr "is.after('germination') & is.before('anthesis',
41.5)" "is.after('germination') & is.before('anthesis', 38.9)" "is.after('germination') &
is.before('anthesis', 36.1)" "is.after('germination') & is.before('anthesis', -1.5)" ...
$ rRootDepth.filter : chr "is.after('emergence',0) & is.before('anthesis',5)"
"is.after('emergence',0) & is.before('anthesis',5)" "is.after('emergence',0) &
is.before('anthesis',5)" "is.after('emergence',0) & is.before('anthesis',5)" ...
$ name : chr "WHEAT.Ble_Dur_1" "WHEAT.Ble_Tendre_1"
"WHEAT.Ble_Tendre_2" "Chickpea.Ghab2" ...

```

ALLCLIMATES

data.frame with one row for each date in each climate. It is created by reading all sheets of file climates.xlsx in the input folder of the simulation folder (NOT the model folder, which contains example input files, which are not read).

'data.frame': 15336 obs. of 8 variables:

```
$ YEAR : num 1995 1995 1995 1995 1995 ...
```



```

$ DOY      : num 1 2 3 4 5 6 7 8 9 10 ...
$ iRSDS    : num 11.3 9.1 11.7 13.1 7.3 12 11.4 14 13.9 13.5 ...
$ iTASMax   : num 21.2 12.8 15 12.8 13.5 15.4 12.8 12.8 14.2 13.5 ...
$ iTASMin   : num 7.8 8.2 4.2 5.5 4 7.8 6.4 0.1 -2.3 2.7 ...
$ iPr       : num 0 2 0 0 0 0 0 0 0 0 ...
$ date      : Date, format: "1995-01-01" "1995-01-02" "1995-01-03" ...
$ climatename: chr "Ain Hamra - Meknes" "Ain Hamra - Meknes" "Ain Hamra - Meknes"
"Ain Hamra - Meknes" ...

```

ALLSOILS

not done yet.

the watermodule is being coded, for now it uses a list named PARAMSSOILS, but it might change in the future

List of 7

```

$ pNLayer      : num [1:3] 2 2 3
$ pDrainLayer   : num [1:3] 2 2 3
$ pSoilAlbedo   : num [1:3] 0.12 0.13 0.15
$ U             : logi NA
$ pSoilCurveNumber: num [1:3] 60 70 80
$ pVPDcoef      : num [1:3] 0.65 0.65 0.65
$ paramlayers   : List of 3
..$ :'data.frame':   2 obs. of 6 variables:
.. ..$ layer      : int [1:2] 1 2
.. ..$ pLayerThickness: num [1:2] 300 700
.. ..$ pSaturation   : num [1:2] 0.36 0.4
.. ..$ pFieldCapacity : num [1:2] 0.24 0.25
.. ..$ pWiltingPoint : num [1:2] 0.1 0.12
.. ..$ pSoilDryness  : num [1:2] 0.03 0.04
..$ :'data.frame':   2 obs. of 6 variables:
.. ..$ layer      : int [1:2] 1 2
.. ..$ pLayerThickness: num [1:2] 200 800
.. ..$ pSaturation   : num [1:2] 0.36 0.4
.. ..$ pFieldCapacity : num [1:2] 0.24 0.25
.. ..$ pWiltingPoint : num [1:2] 0.1 0.14
.. ..$ pSoilDryness  : num [1:2] 0.04 0.04
..$ :'data.frame':   3 obs. of 6 variables:
.. ..$ layer      : int [1:3] 1 2 3
.. ..$ pLayerThickness: num [1:3] 300 200 400
.. ..$ pSaturation   : num [1:3] 0.36 0.4 0.38
.. ..$ pFieldCapacity : num [1:3] 0.24 0.25 0.22
.. ..$ pWiltingPoint : num [1:3] 0.1 0.12 0.09
.. ..$ pSoilDryness  : num [1:3] 0.03 0.04 0.035

```

ALLMANAGEMENTS

It is created by reading soils.xlsx in the input folder of the simulation folder (NOT the model folder, which contains example input files, which are not read).

List of possible crop managements plans (named "row XX" based on the row number of <--- MangRowNo in the excel file).

List of 3

```
$ row 13:List of 11
```



```

..$ dfCode      : 'data.frame':      1 obs. of 2 variables:
.. ..$ Code      : chr "ROTATION_BLE"
.. ..$ Description: num 0
..$ dfSowing     : 'data.frame':      1 obs. of 11 variables:
.. ..$ FixFind: num 4
.. ..$ Fyear  : num 2007
.. ..$ yrno   : num 1
.. ..$ SimDoy : num 20
.. ..$ Fpdoy  : num 288
.. ..$ Lpdoy  : num 20
.. ..$ StopDAP: num 280
.. ..$ SowTmp : chr "-"
.. ..$ SowWat : num 0.3
.. ..$ Pden   : num 280
.. ..$ STBLW  : num 0.01
..$ nitrogenScenario: num 2
..$ nitrogenNumber : num 1
..$ nitrogenDatetype: num 1
..$ nitrogendf      : 'data.frame':      1 obs. of 4 variables:
.. ..$ NapplNumber: num 1
.. ..$ DAPorCBD   : num 1
.. ..$ amount     : num 5
.. ..$ FracVol    : num 7
..$ waterLevel    : num 0
..$ waterScenario : num 2
..$ waterNumber    : num 0
..$ waterDatetype  : num 0
..$ waterdf        : 'data.frame':      0 obs. of 2 variables:
.. ..$ DAPorCBDorDOY: logi(0)
.. ..$ amount       : logi(0)
$ row 36:List of 11
idem
$ row 59:List of 11
idem

```

GENERALPARAMETERS

this data.frame is read from sheet " generalPhysicalParameters" in file "allvariables.xlsx" from the model folder.

'data.frame': 5 obs. of 9 variables:

```

$ name      : chr "pMinimalSoilEvaporation" "pSoilWettingWaterQuantity"
"pCanopyExtinctionCoefficient" "pCropAlbedo" ...
$ typeinthemodel : chr "parameter" "parameter" "parameter" "parameter" ...
$ typeR        : chr "numeric" "numeric" "numeric" "numeric" ...
$ module       : chr "rWaterBudget" "rWaterBudget" "rWaterBudget" "rWaterBudget"
...
$ unit        : chr "mm" "mm" "-" "-" ...
$ definition   : chr "Minimal Soil Evaporation" "amount of rainfall and/or irrigation
required to wet the top soil layer to return soil evaporation from Stage II to Stage I"
"canopy extinction coefficient" "Crop albedo" ...
$ translationSSM : chr "EOSMIN" "WETWAT" "KET" "CALB" ...

```

```
$ french      : chr "evaporation minimale du sol" NA NA NA ...  
$ defaultInitialvalue: num 1.5 10 0.5 0.23 0.48
```

Structure of the files in the model folder

- allvariables.xlsx : excel file with the name, module where it is used, definition, unit, original SSM name and initial value of all variables
- exampleRunSSM.R : R script giving an example of how to run the model (and the definition of the setup function : this is the script that the user will run in the console)
- externalFilesReadingSSM.R : function definitions for reading in external files (does not actually read anything, the reading is done when the handler function mRun() is run for the first time)
- functionsSSM.R : definition of all functions and procedures that are at the core of the SSM model (to be done: split this file by module to facilitate maintenance)
- handlersSSM.R : definition of handler functions (mRun, mPlotDynamics...)
- headersSSM.R : loading of required packages
- HousekeepingFunctionsSSM.R : definitions of functions that are not specific of SSM but are usefull for the model algorithm.
- initialisationEnvironmenVariablesSSM.R : creation of all the encapsulated "global" variables.
- input : folder with example input files
 - climates.xlsx : excel file with one sheet per possible climate (same content as in SSM.xlsx)
 - crops.xlsx : excel file with one sheet per species, one column per cultivar (same content as in SSM.xlsx except extra parameters at the bottom: thresholds for the different stages (obtained with a formula using the bdXXX parameters, warning: for the formula to give R-readable values, the decimal separator in excel must be the point, not coma as is the default in French language setting) and filters (text in R language that will be parsed by the model) for the different modules... see "filters", page 11.
 - managementPlans.xlsx not used yet (but read anyway): excel file with one sheet per crop Management plan (rule for sowing and sowing density, nitrogen and water applications): (same content as in SSM.xlsx).
 - soils.xlsx excel file with one sheet per possible soil (same content as in SSM.xlsx)
- LICENSE: GNU licence, I don't know why it's here
- progR_essaiSSMfiltres.R : to be removed, it's a remnant of old stuff
- README.md : file used by github to display on <https://github.com/mariegosme/SSM.R>
- SSMWaterModule.R : temporary file with the water module partially coded... will eventually be merged into functionsSSM.R (or kept as separate file if modules are separated into different files).

Function setup() sources, locally within itself, the files headersSSM.R, initialisationEnvironmenVariablesSSM.R, externalFilesReadingSSM.R, HousekeepingFunctionsSSM.R, functionsSSM.R, handlersSSM.R in this order.

How to run the model

1) run the function definition of function setup

2) build the model

```
mymodel<-setup("/Users/user/Documents/b_maison/congeMat/D4DECLIC/SSM/")
```

3) prepare cases (these will be the rows of ALLSIMULATEDDATA):

```
mycases<-data.frame(climatename="Ain Hamra - Meknes", soilname="325_-35",  
lat=c(35, 45), long=-5)
```

#climatename: one of the sheet names of file climates (if climate in standard SSM format)

#soilname: one of the sheet names of file soils (if in standard SSM format)

#to do: define crop rotation and management (once management procedure is completed)

```
rownames(mycases)<-c("Meknes35degres", "Meknes45degres") #these will be the  
cases names used in the plots, outputs etc...
```

4) define the options of the simulation

```
paramsim<-list(  
  simustart=as.Date("1997-11-01"), #date of start of the simulation  
  cases=mycases, #cases (e.g. combinations of soil-climate-management-location)  
  directory="/Users/user/Documents/b_maison/congeMat/D4DECLIC/runSSM", #folder  
  where your input (with climates and soils files) and output folders are  
  climateformat="standardSSM",  
  cropformat="standardSSM",  
  soilformat="standardSSM",  
  managformat="standardSSM",  
  Neffect=TRUE  
)
```

5)use this to set the simulation options

```
mymodel$setoptions(paramsim)
```

6)run the model for 4 timesteps

```
mymodel$run(4)
```

7)plot the dynamics of some variables

```
dynamiques<-mymodel$plot(c("iTASMin", "iTASMax", "iRSDS"),  
  colors=c(iTASMin="blue", iTASMax="red", iRSDS="black"),  
  whatcolors="variables",  
  linetypes=c(iTASMin=1, iTASMax=1, iRSDS=2), whatlinetypes="variables",  
  symbols=c(Meknes35degres=1, Meknes45degres=8), whatsymbols="cases")
```

Details on the deviations from the original SSM model, by module

all modules

We use **"filters"** to know to which cases to apply time-dependent functions or procedures: in the original SSM code, the conditions were hard-coded for each crop (e.g. for wheat:

bdBRV = bdEMR 'Beg. response to vernalization

bdTRV = bdSEL 'End of response to vernalization

If CBD >= bdBRV And CBD <= bdTRV do vernalisation calculations

In SSM.R, these conditions are part of the crop parameterisation:

`vernalisation.filter = "is.after('emergence', 0) & is.before('stemElongation', 0)"` which means that vernalisation is applied only to wheat crops that are already emerged but haven't started the stem elongation stage. The syntax of `is.before` and `is.after` is the following: `is.before(stage, bd=-Inf)`, which means that if `bd` is not specified, the stage itself is not included (`is.before('tillering')` means that it happens strictly before the beginning of tillering stage, while `is.before('tillering', 5)` means that it happens until the 5th biological day within the tillering stage. For `is.after`, the syntax is `is.after(stage, bd=Inf)`, which means that if `bd` is not specified, the stage itself is not included (`is.after('tillering')` means that it happens strictly after the end of tillering stage, while `is.after('tillering', 5)` means that it happens starting on the 5th biological day within the tillering stage. Note that the stage names are not hard-coded: they come from the names given to the thresholds for each stage provided in the crop parameter file.

Phenology

In the original excel version, phenology is determined by comparing the cumulated "biological days" from sowing with stage-change biological days computed at the beginning based on the biological days necessary to complete each stage (found in the crop parameter file, or, for maize, from unulated thermal units between the two previous stages). It seemed more logical to cumulate biological days within each stage (resetting the cumulated sum to almost 0 (actually, the "extra" biological days beyond the threshold obtained on the day of the stage change are used to initialise the sum for the next day), and compare this number directly with the stage duration of each stage.

Another complicated aspect was the determination of when each process took place (e.g. vernalization, secondary growth of leaves etc...): these were hard-coded in the code for each species, based on phenological stages (e.g. for wheat `bdTLM=bdBOT`) or on parameters (e.g. for legume, `bdTLM` is a parameter found in the crop parameter file). The use of filters (see "filters", page 11) is supposed to simplify this: R-readable code in the crop parameter file tells under which condition each conditional process takes place.

For example, the following phenological scale (as I understood it from the code) can be simply expressed as the following crop parameters (note that the filters are obtained with excel formulas from the original stage durations so it's easy to create these filters for new cultivars from their original parameters):

N102		="is.after('ANT', 5) & is.before('ANT', "& N85-1.5 & ")"		
	A	N	O	P
4	CropColNo -->	14		
5	CROP:	WHEAT		
6	Cultivar:	Cocorit		
96	thresholds	c(SOW=6,EMR=5,TIL=8,SEL=6,BOT=6,EAR=15,ANT=43, PM=8, MAT=Inf)		
97	vernalisation.filter	is.after('EMR', 0) & is.before('SEL')		
98	photoperiod.filter	FALSE		
99	waterstress.filter	FALSE		
100	LAI_Mainstem.filter	is.after('EMR', 0) & is.before('BOT')		
101	LAI_Secondary.filter	is.after('BOT', 0) & is.before('ANT', 5)		
102	DMDistribution_Se	is.after('ANT', 5) & is.before('ANT', 41.5)		
103	DMPProduction.filter	FALSE		
104	rRootDepth.filter	is.after('EMR',0) & is.before('ANT',5)		
105	TBVER	-1		
106	TP1VER	0		
107	TP2VER	8		
108	TCVER	12		
109	VDSAT	50		
110	pPhotoperiodFunc	fComputeCoefPhotoperiodWheat		

Figure 4: possible specification of wheat phenology and conditioning of different crop processes to phenology, following the original version's names

For maize, the following scale can be translated in the following parameters:

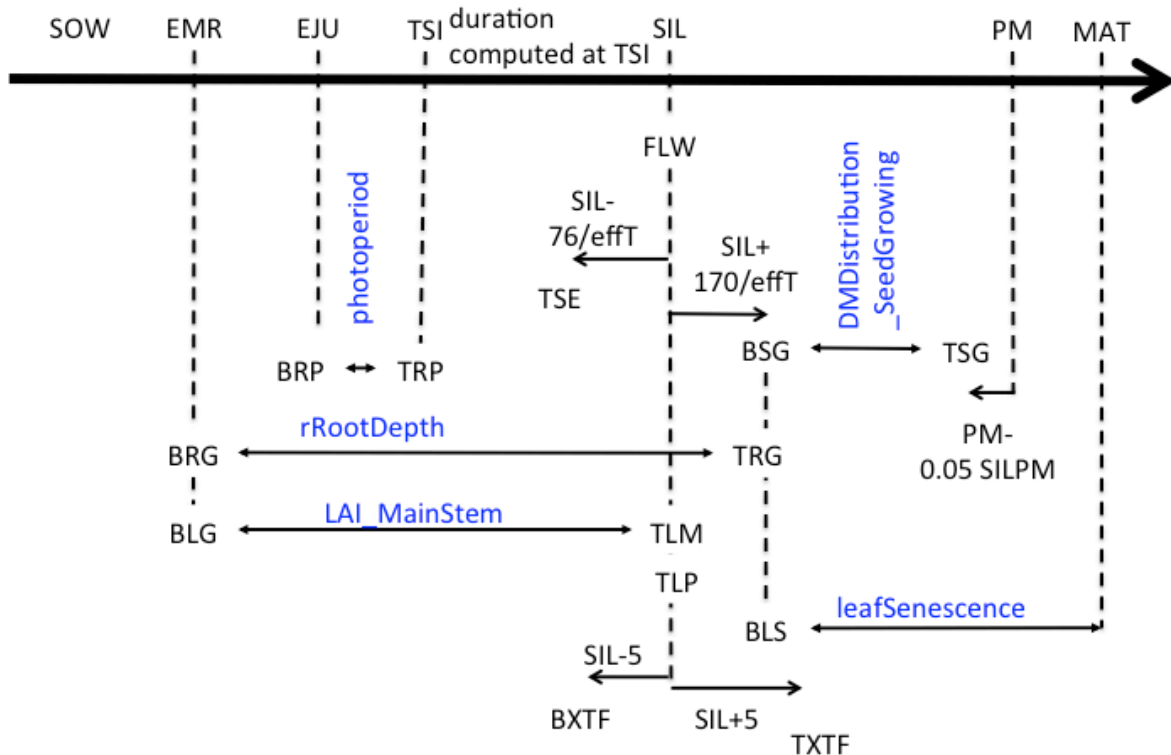


Figure 5: phenological scale of maize. Names in black are the original version's names (except "bd" was removed), names in blue are the names of the filters. effT is TP1D-TBD

P102		= "is.after('SIL', 170) & is.before('SIL', "& 0.95*250&")"			
	A	P	Q	R	S
4	CropColNo -->				
5	CROP:	MAIZE			
6	Cultivar:	bidule			
96	thresholds	c(SOW=6, EMR=10, EJU=4, TSI=NA, SIL=200, PM=4, MAT=Inf)			
97	vernalisation.filter	FALSE			
98	photoperiod.filter	is.after('EJU', 0) & is.before('TSI')			
99	waterstress.filter	FALSE			
100	LAI_Mainstem.filter	is.after('EMR', 0) & is.before('SIL')			
101	LAI_Secondary.filter	FALSE			
102	DMDistribution_Se	is.after('SIL', 170) & is.before('SIL', 237.5)			
103	DMPProduction.filter	FALSE			
104	rRootDepth.filter	is.after('EMR', 0) & is.before('SIL', 170)			
105	TBVER				
106	TP1VER				
107	TP2VER				
108	TCVER				
109	VDSAT				
110	pPhotoperiodFunct	fComputeCoefPhotoperiodMaize			

Figure 6: possible specification of maize phenology and conditioning of different crop processes to phenology. With dummy numbers because I don't have examples of existing maize parameters.