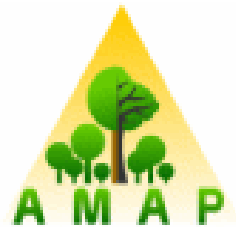


AMAPstudio: a 3D Interactive Software Suite for Plants Architecture Modelling

Sébastien Griffon (Cirad-AMAP)
François de Coligny (INRA-AMAP)



INRA



cirad

December 6th 2013, Montpellier

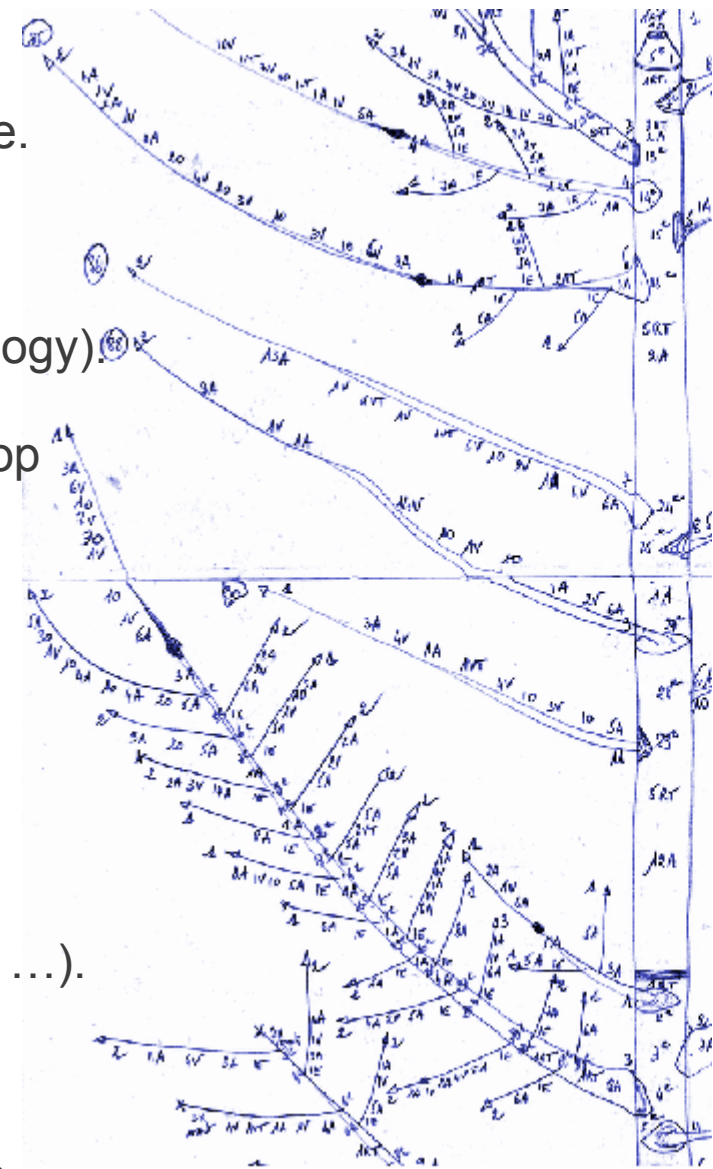
Objectives

Objectives

- Build, simulate, edit, explore and analyse plant architecture.
- User-friendly and end-user oriented.
- Multi-purpose application (botany, agronomy, forestry, ecology).
- Single framework and methodology for modellers to develop their own models.
- Scenario oriented.

Context

- Many existing tools (GroImp, OpenAlea, L-studio, Lignum, ...).
- AMAP develop software for plant modelling since the 80's.
- In 2008, scientists synergy to design all those features in a single software suite to share knowledge and methods with a long-term support.



Y. Caraglio

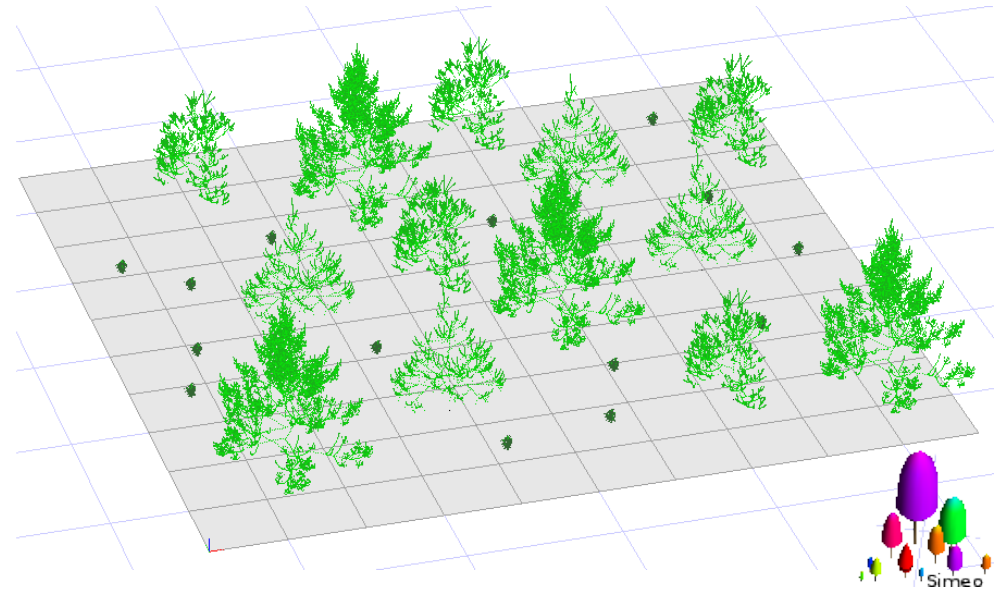
AMAPstudio : features for plant architecture modelling

1- Central plant data structure



2 - Xplo

Individual scale = detailed representation



3 - Simeo

Scene scale = simplified representation

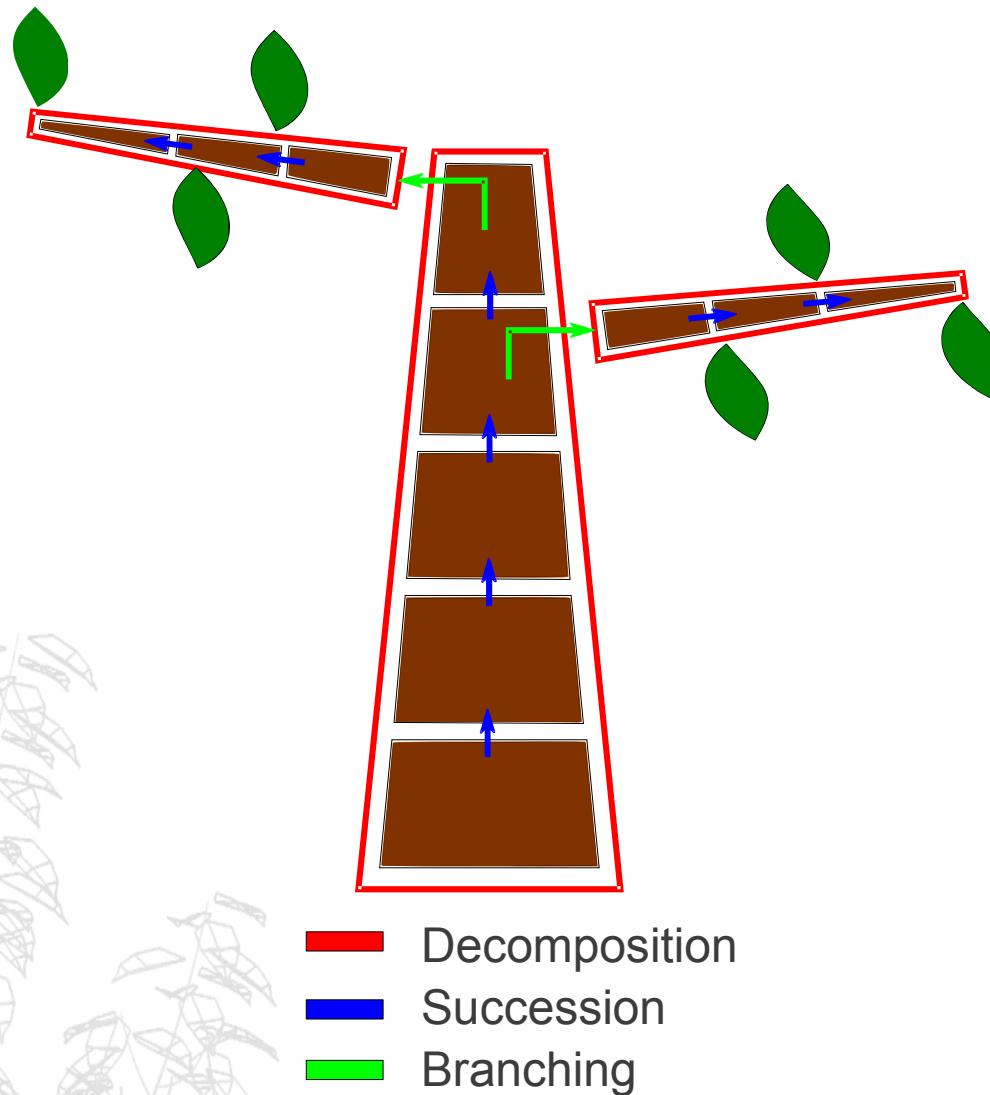
4 - Simulation framework

5 - Common features

I. Central plant data structure 1/3

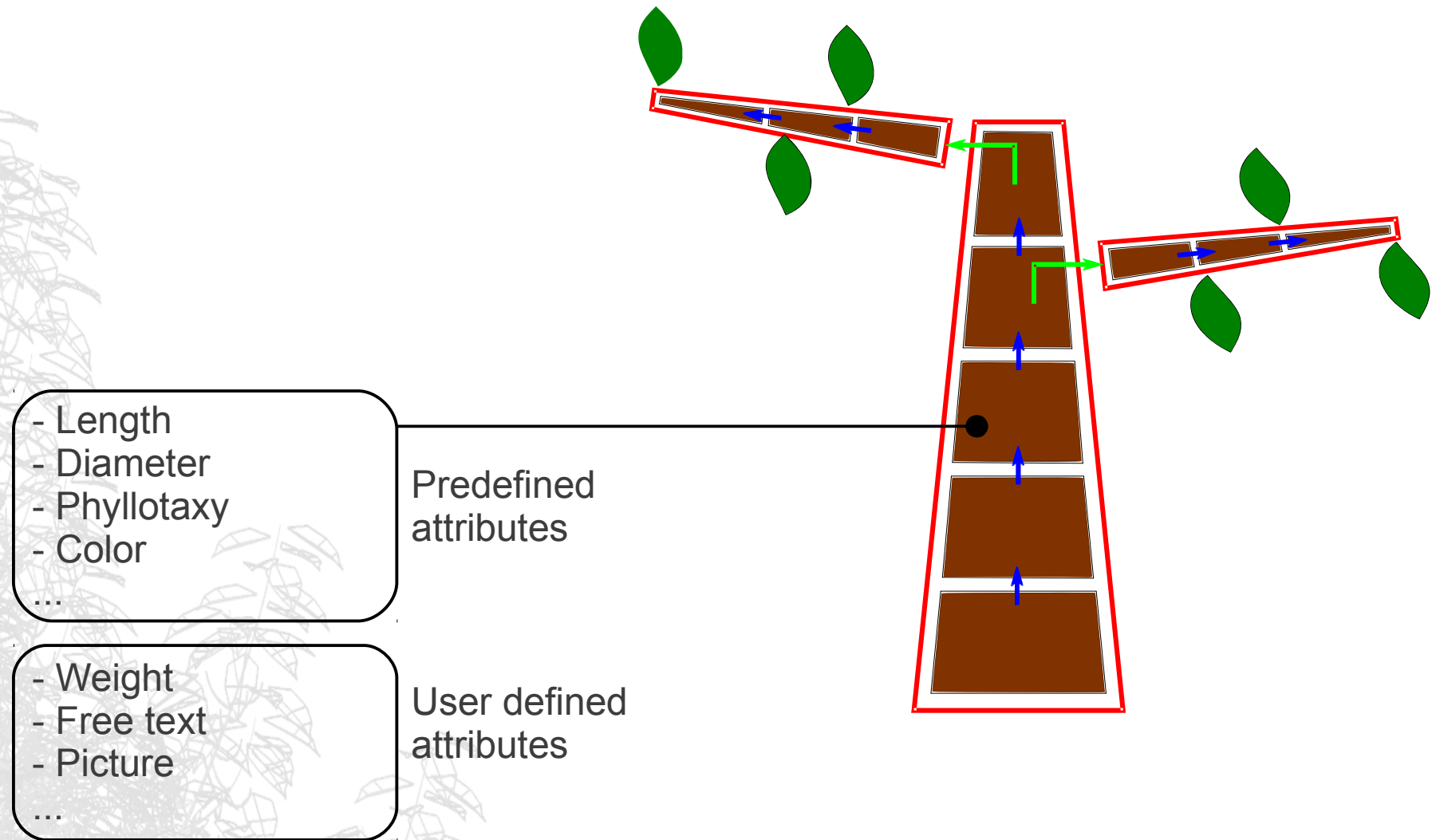
Topology based on the Multiscale Tree Graph (MTG) formalism

C. Godin and Y. Caraglio, "A multiscale model of plant topological structures", 1998 (1)



I. Central plant data structure 2/3

Data base : attributes can be attached to the plant components



I. Central plant data structure 3/3

Topology
+ Geometrical attributes
+ Geometrical additionnal rules
+ Shape builder

→ 3D Mock-up

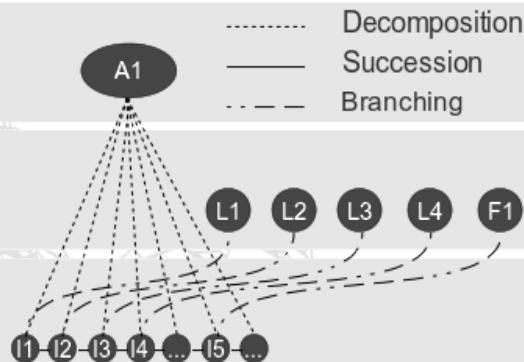
(a) Scales

Axis (A)

Leaf (L)/Fruit (F)

Internode (I)

(b) Topology



(c) Attributes

A1 : Height = 198.6 cm

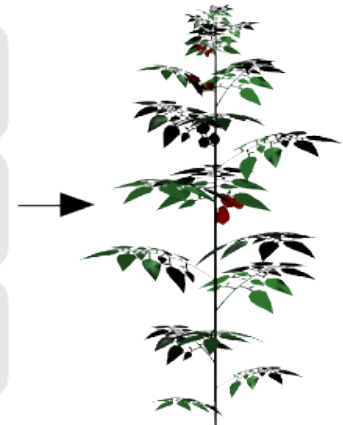
L3 : Length = 17.5 cm
Width = 14.7 cm
Insertion angle = 60 °
...

I2 : Length = 9.2 cm
Diameter = 2.1 cm
...

(d) Geometry



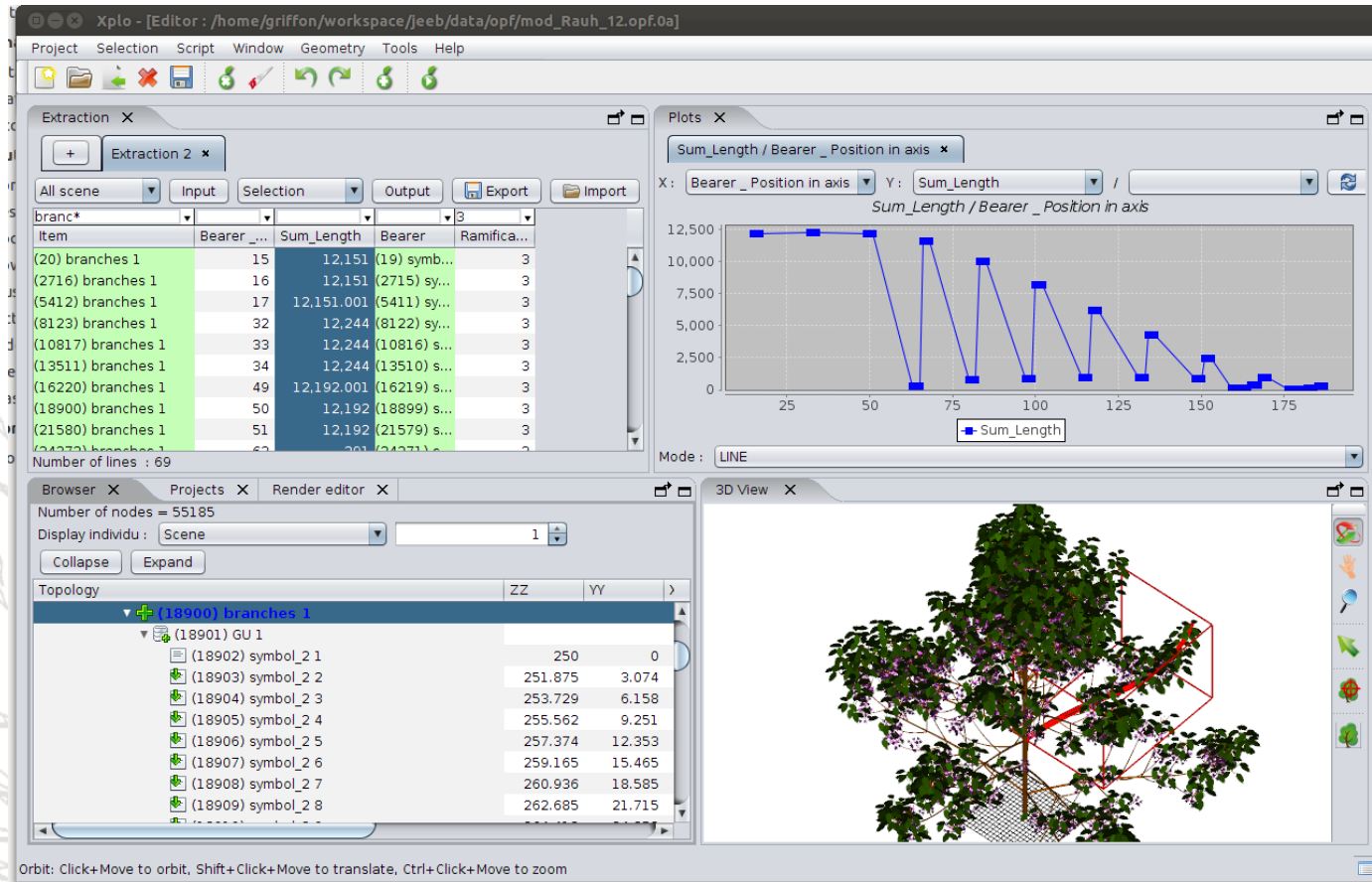
(e) 3D mock-up



II. Xplo : main features



Detailed plant representations in several views



- 1) **Edit and explore** plant architecture
- 2) **Simulate** plant growth with one of the integrated model.

II. Xplo editor : plant edition (table view)

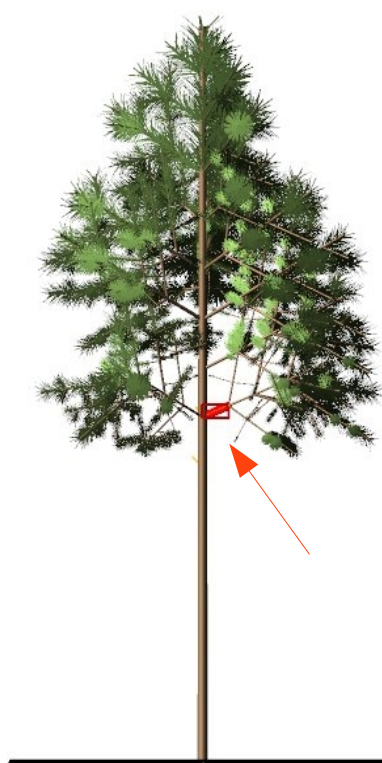
Browser X Render editor X Types editor X 3D View X

Number of nodes = 4404
Display individu : Scene 1

Collapse Expand

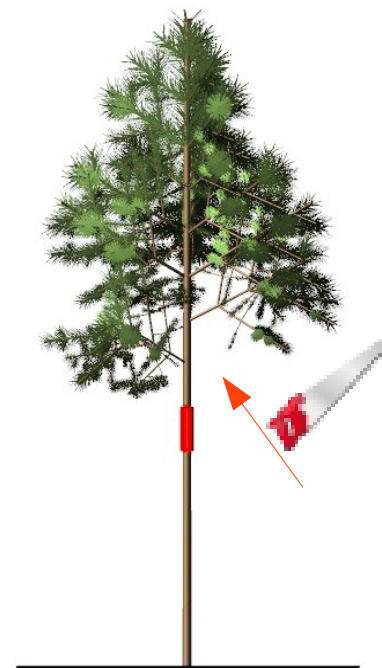
Topology

	Length	Width	InsertionAng
(1) Scene 1			
(2) Plant 1			
+ (3) Axis 1			
(4) Metamer 1			
(5) Internode 1	24.75	5.325	
(6) Metamer 2			
(8) Metamer 3			
(10) Metamer 4			
(12) Metamer 5			
(14) Metamer 6			
(15) Internode 1	25	4.809	
+ (16) Axis 1			69
(17) Metamer 1			
(18) Metamer 7			
(19) Internode 1	25	4.705	
+ (20) Axis 1			67.7
(21) Metamer 1			
(22) Internode 1	14.904	1.3	
+ (23) Axis 1			56.6
+ (136) Axis 1			56.6
(137) Metamer 1			
(165) Metamer 2			
(181) Metamer 3			
(207) Metamer 4			
(229) Metamer 5			
(230) Internode 1	3.6	0.86	



1. Remove organs

e.g : Pruning



3D View X

1

	Length	Width	InsertionAng
(8) Metamer 3			
(10) Metamer 4			
(12) Metamer 5			
(14) Metamer 6			
(15) Internode 1	25	4.809	
+ (16) Axis 1			69
(17) Metamer 1			
(18) Metamer 7			
(19) Internode 1	25	4.705	
+ (20) Axis 1			67.7
(21) Metamer 1			
+ (22) Axis 1			67.7
(23) Metamer 1			
+ (24) Axis 1			67.7
(25) Metamer 1			
(26) Internode 1	14.904	1.3	
+ (27) Axis 1			56.6
(28) Metamer 1			
(56) Metamer 2			
(72) Metamer 3			

II. Xplo editor : plant edition (table view)

Browser X Render editor X Types editor X

Number of nodes = 3620

Display individu : Scene 1

Collapse Expand

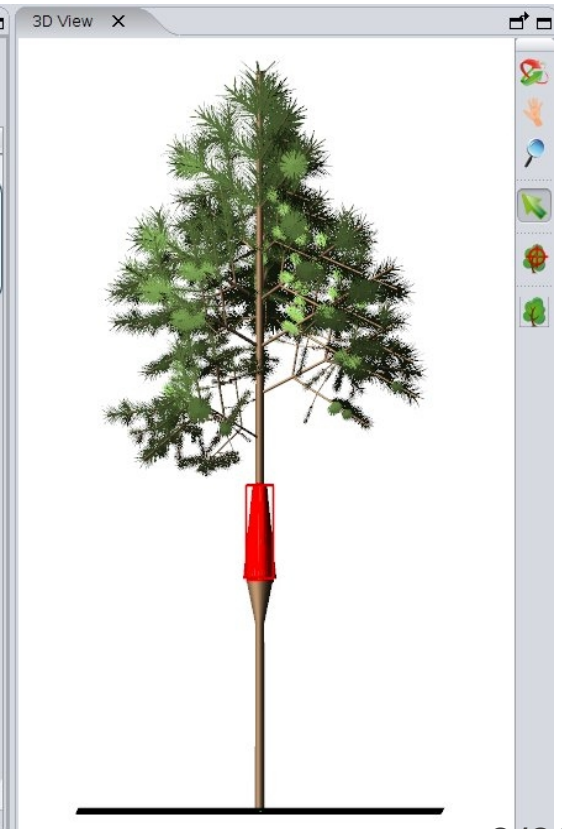
Topology	Length	Width	InsertionAngle
(1) Scene 1			
(2) Plant 1			
(3) Axis 1			
(4) Metamer 1			
(5) Internode 1	24.75	5.325	
(6) Metamer 2			
(8) Metamer 3			
(10) Metamer 4			
(12) Metamer 5			
(14) Metamer 6			
(15) Internode 1	25	4.809	
(16) Axis 1			69
(17) Metamer 1			
(18) Metamer 7			
(19) Internode 1			
(20) Axis 1	25	4.809	67.7
(21) Metamer 1			67.7
(22) Axis 1			67.7
(23) Metamer 1			
(24) Axis 1			
(25) Metamer 1			
(26) Internode 1	14.904	1.3	56.6
(27) Axis 1			
(28) Metamer 1			
(56) Metamer 2			
(72) Metamer 3			



2a. Edit attribute

e.g.
length, diameter

(12) Metamer 5			
(14) Metamer 6			
(15) Internode 1	50	15	
(16) Axis 1			69
(17) Metamer 1			
(18) Metamer 7			
(19) Internode 1			
(20) Axis 1	50	15	67.7
(21) Metamer 1			67.7
(22) Axis 1			67.7
(23) Metamer 1			
(24) Axis 1			
(25) Metamer 1			
(26) Internode 1	14.904	1.3	56.6
(27) Axis 1			
(28) Metamer 1			
(56) Metamer 2			
(72) Metamer 3			



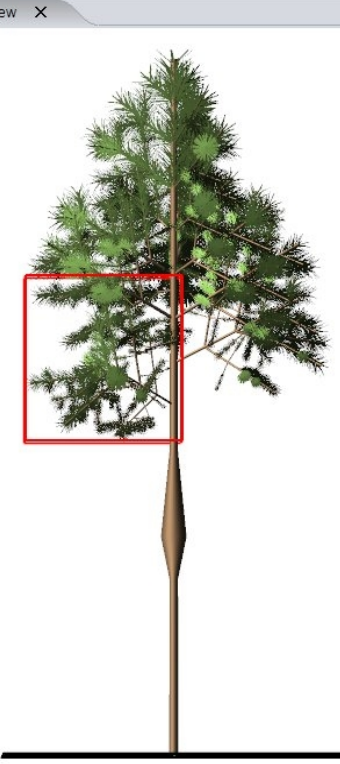
II. Xplo editor : plant edition (table view)

Browser X Render editor X Types editor X 3D View X

Number of nodes = 3620
Display individu : Scene 1


Collapse Expand

ology	InsertionAngle	Length	Phyllotaxy	W
(2) Plant 1				
+ (3) Axis 1	0			
(4) Metamer 1				
(5) Internode 1		24.75		
(6) Metamer 2			112.45	
(8) Metamer 3			112.45	
(10) Metamer 4			112.45	
(12) Metamer 5			112.45	
(14) Metamer 6			112.45	
(15) Internode 1		50		
+ (16) Axis 1	69.48			
(17) Metamer 1				
(18) Metamer 7			112.45	
(19) Internode 1		25		
+ (20) Axis 1	67.788			
(21) Metamer 1				
+ (22) Axis 1	67.788			
(23) Metamer 1				
+ (24) Axis 1	67.788			
(25) Metamer 1				
(26) Internode 1		14.904		
+ (27) Axis 1	67.788			
(28) Metamer 1			90	
(56) Metamer 2			0	
(72) Metamer 3			0	
(98) Metamer 4			0	

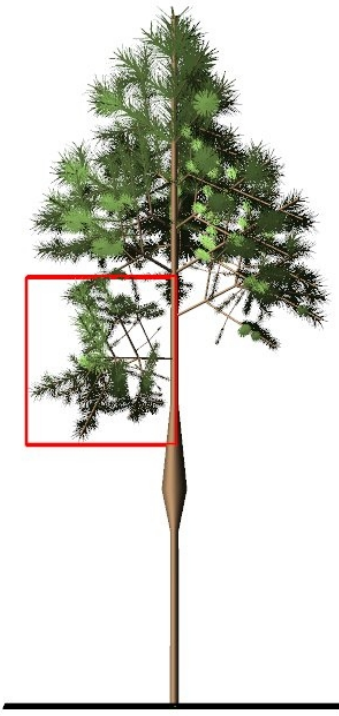


2b. Edit attribute

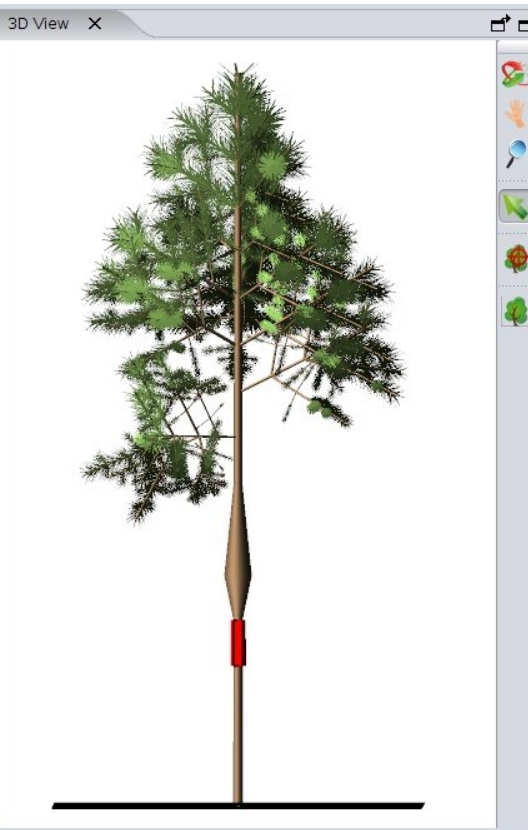
e.g. insertion angle



(5) Internode 1	24.75		
(6) Metamer 2		112.45	
(8) Metamer 3		112.45	
(10) Metamer 4		112.45	
(12) Metamer 5		112.45	
(14) Metamer 6		112.45	
(15) Internode 1	50		
+ (16) Axis 1	69.48		
(17) Metamer 1			
(18) Metamer 7		112.45	
(19) Internode 1	25		
+ (20) Axis 1	67.788		
(21) Metamer 1			
+ (22) Axis 1	67.788		
(23) Metamer 1			
+ (24) Axis 1	90		
(25) Metamer 1			
(26) Internode 1	14.904		
+ (27) Axis 1			
(28) Metamer 1		90	
(56) Metamer 2		0	
(72) Metamer 3		0	
(98) Metamer 4		0	



II. Xplo editor : plant edition

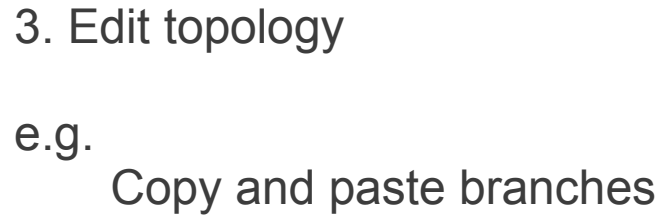


e.g.

[illegible]

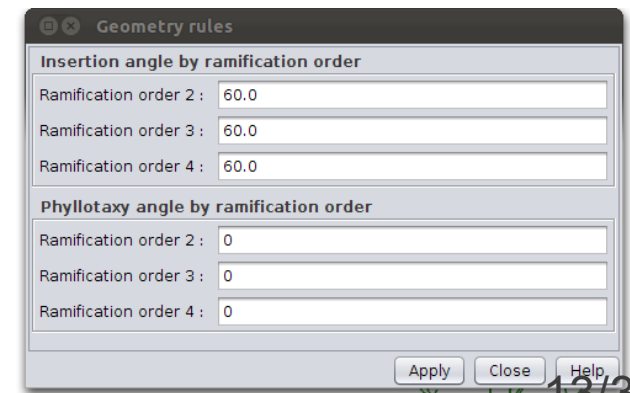
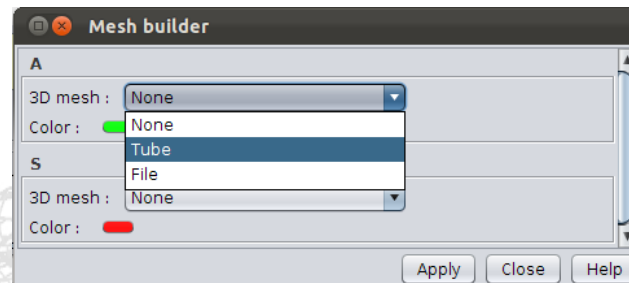
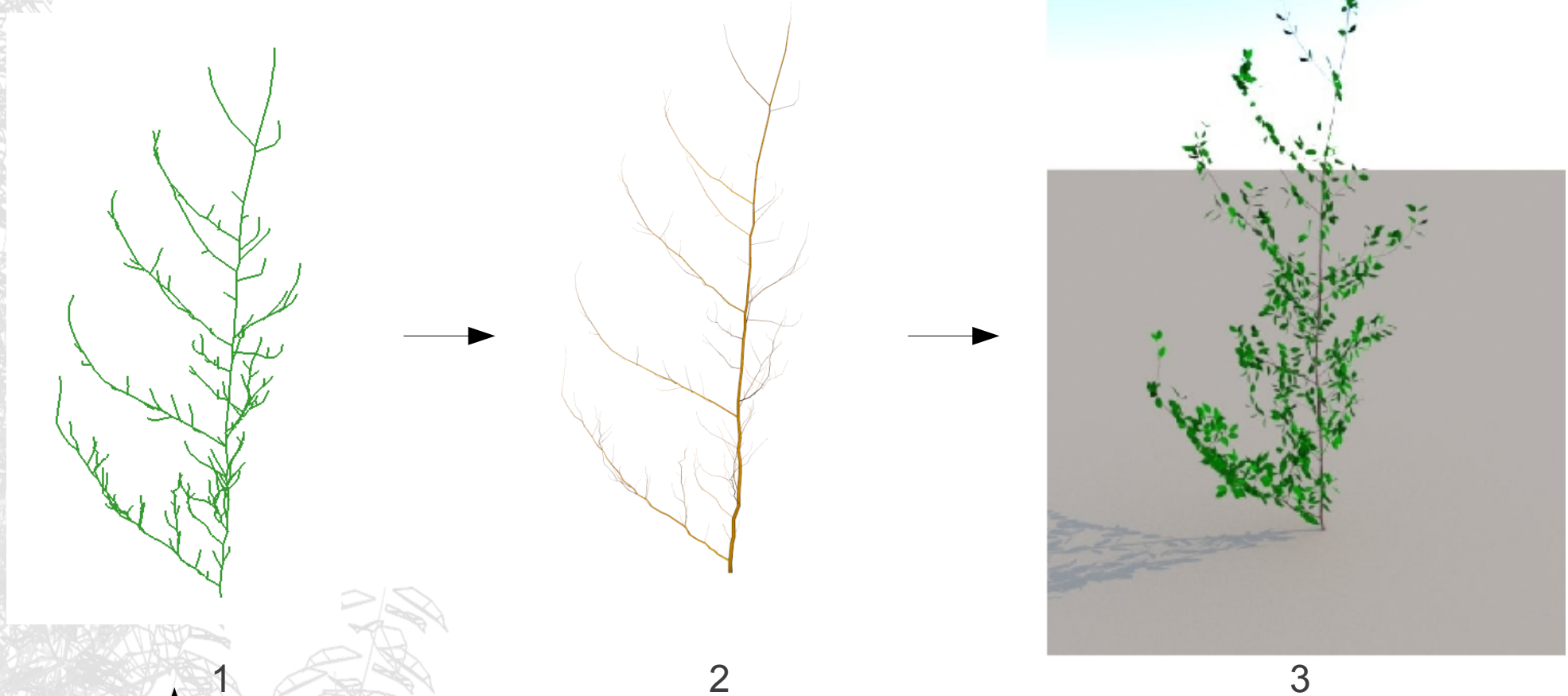
II. Xplo editor : plant

The screenshot displays the Xplo editor interface for plant modeling. The top bar shows three tabs: 'Browser', 'Render editor', and 'Types editor'. Below this, a status bar indicates 'Number of nodes = 3620' and 'Display individu : Scene'. The main workspace is divided into two panels: 'Topology' and 'InsertionAngle'. The 'Topology' panel shows a hierarchical tree of plant components, including Metamers (e.g., (16) Metamer 6, (18) Metamer 7, (810) Metamer 8, (1378) Metamer 9), an Internode ((1379) Internode 1), and Axes ((1380) Axis 1, (1754) Axis 1, (2128) Axis 1). A context menu is open over the selected node (2128) Axis 1, listing various actions such as 'Add an individu', 'Add a successor', 'Add a predecessor', 'Decompose', 'Branch', 'Repeat', 'Remove', 'Cut', 'Copy', 'Paste', 'Change type', and 'Plot attribute'. A red arrow points to the 'Copy' option. Below the main window, a red box highlights a secondary menu with options: 'Remove', 'Cut', 'Copy', and 'Paste'.



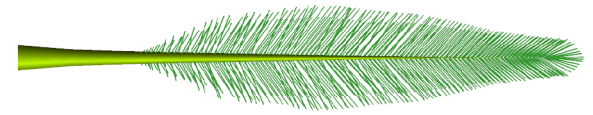
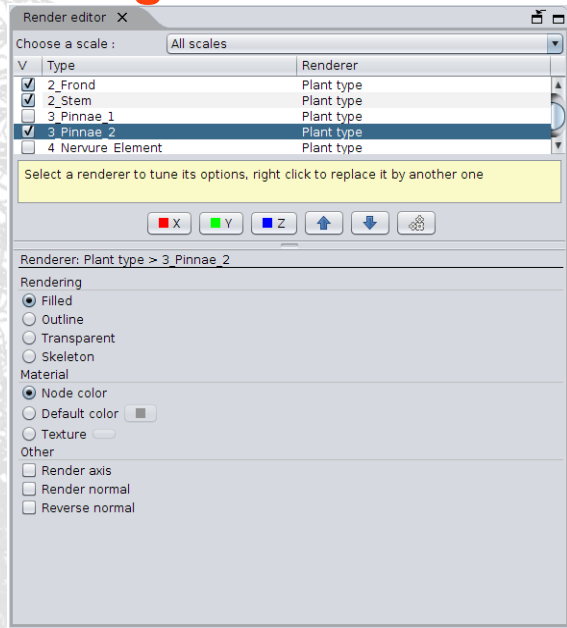
II. Xplo editor : 3D geometry reconstruction

Example : Digitized plant geometry reconstruction

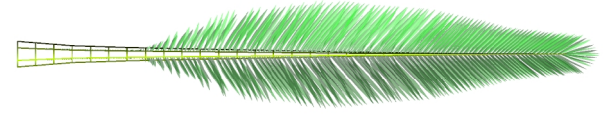
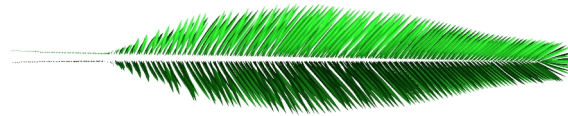


II. Xplo editor : configurable 3D view

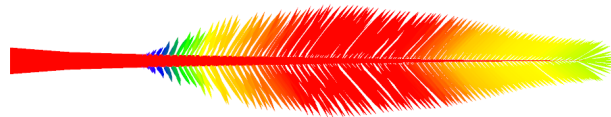
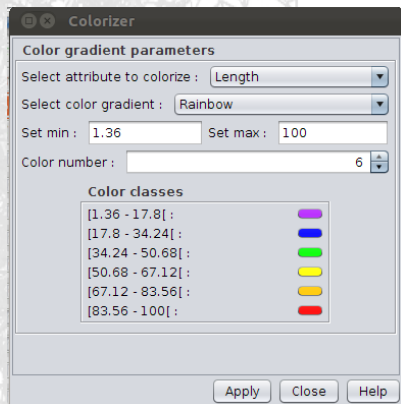
Configurable renderers



Customisable layer rendering



Colorizer tool



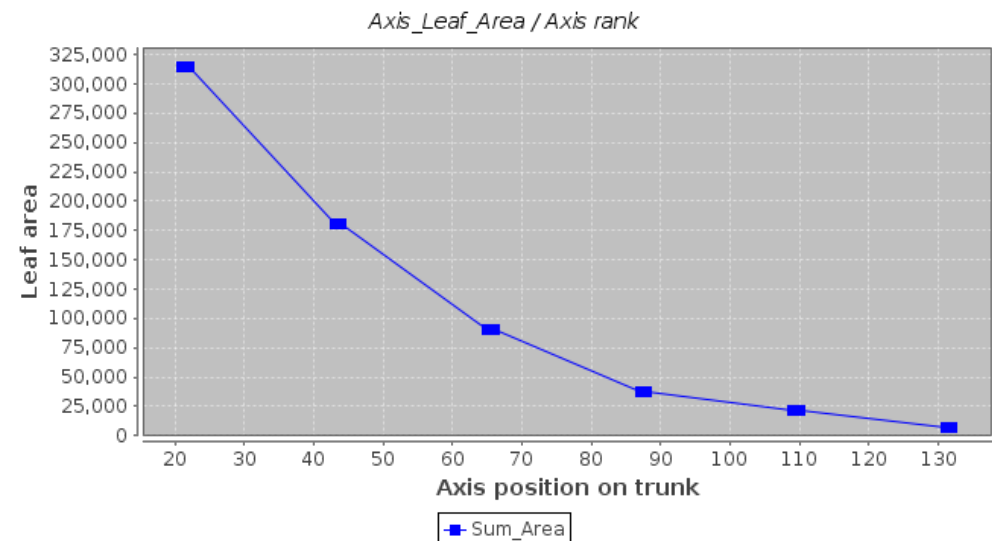
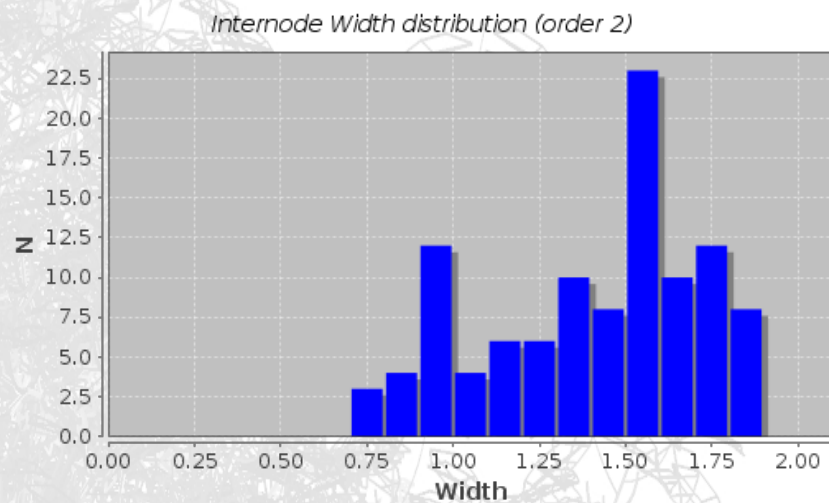
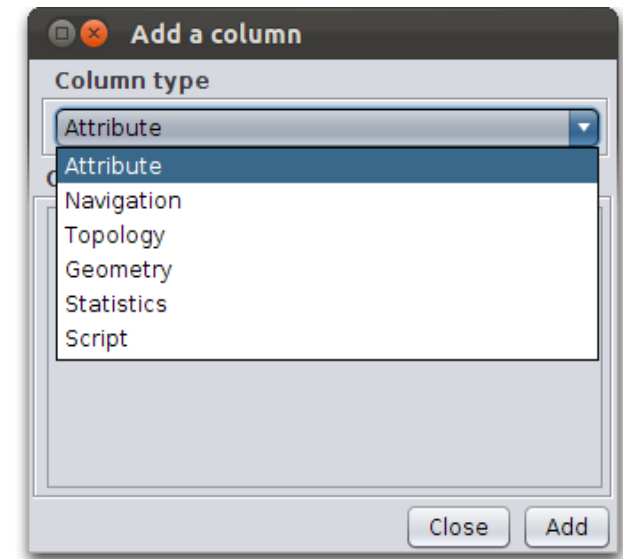
Colors vs attributes

II. Xplo editor : data extraction

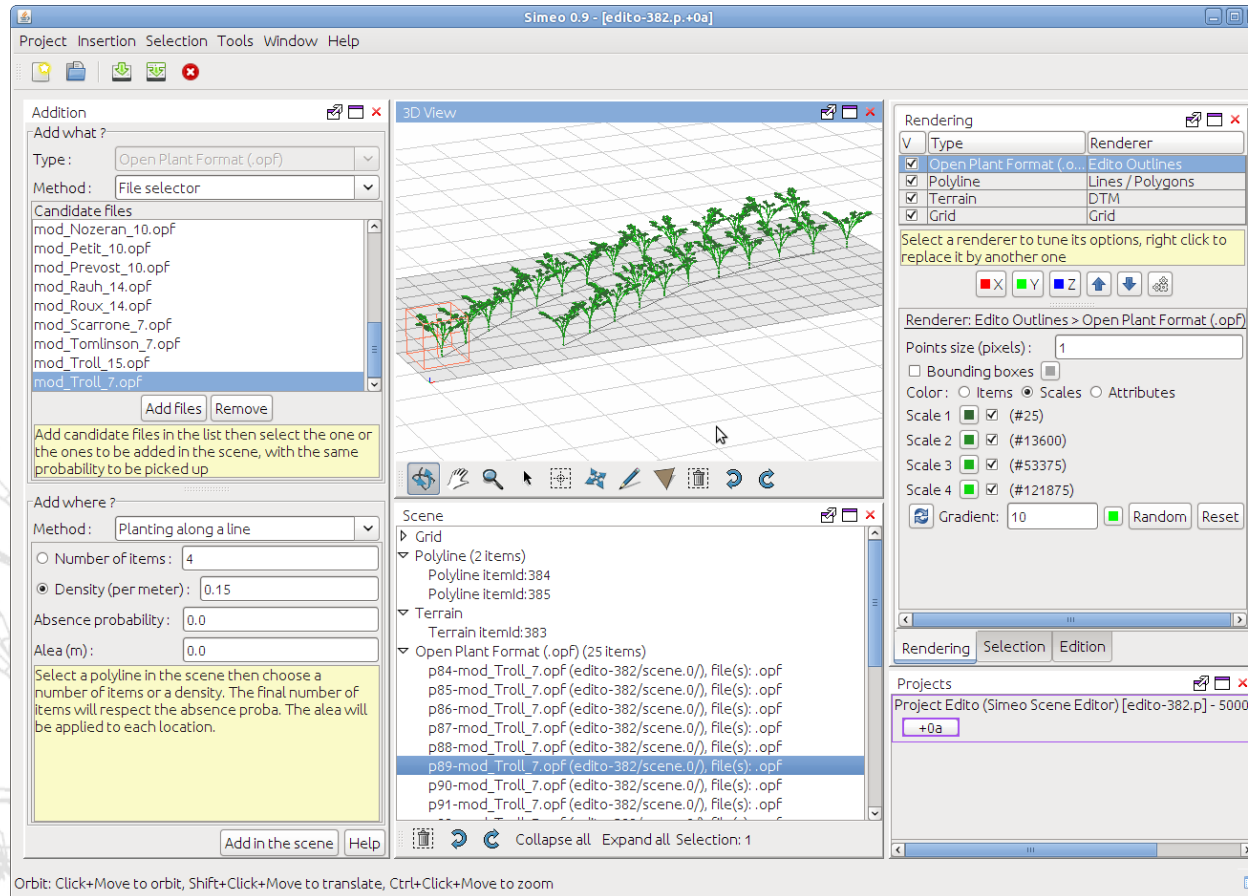
Interactive extraction table

All scene	Input	Selection	Output	Export	Import
branches	2				
Item	Ramification order	Bearer	Bearer _ Position ...	Sum_Area	
branche...	2	branches1 21	21	314966.989	
branche...	2	branches1 22	22	314966.999	
branche...	2	branches1 21	43	180814.383	
branche...	2	branches1 22	44	180814.397	
branche...	2	branches1 21	65	90407.193	
branche...	2	branches1 22	66	90407.203	
branche...	2	branches1 21	87	37912.693	
branche...	2	branches1 22	88	37912.694	
branche...	2	branches1 21	109	21386.648	
branche...	2	branches1 22	110	21386.648	
branche...	2	branches1 21	131	7128.882	
branche...	2	branches1 22	132	7128.882	

Number of lines : 12



III. Simeo : main features



1 – **Edit** high detailed vegetal scenes (adaptative memory management)

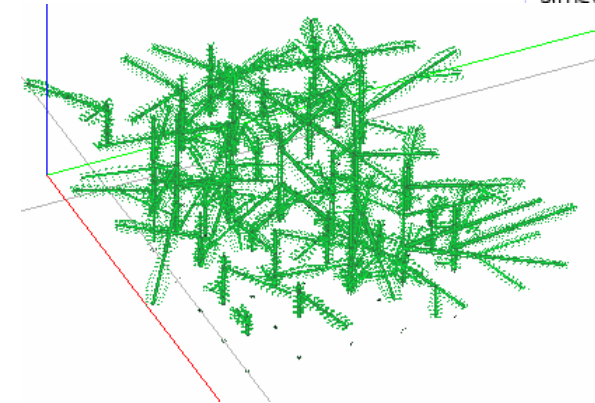
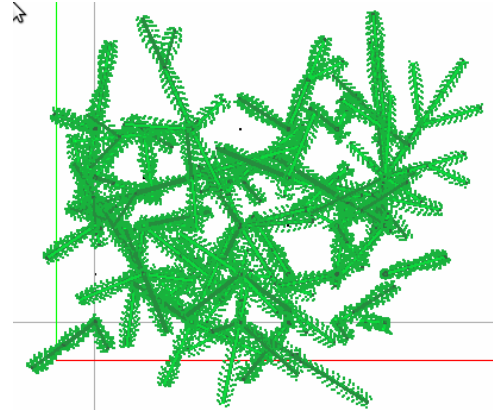
2 – **Connect** with scene level biophysics models (integrated or external) or **simulate** vegetal scene growth.

III. Simeo editor : scene creation and edition

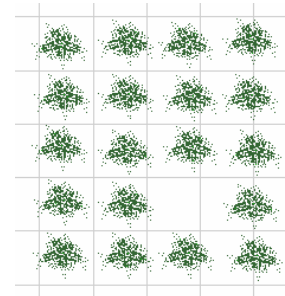
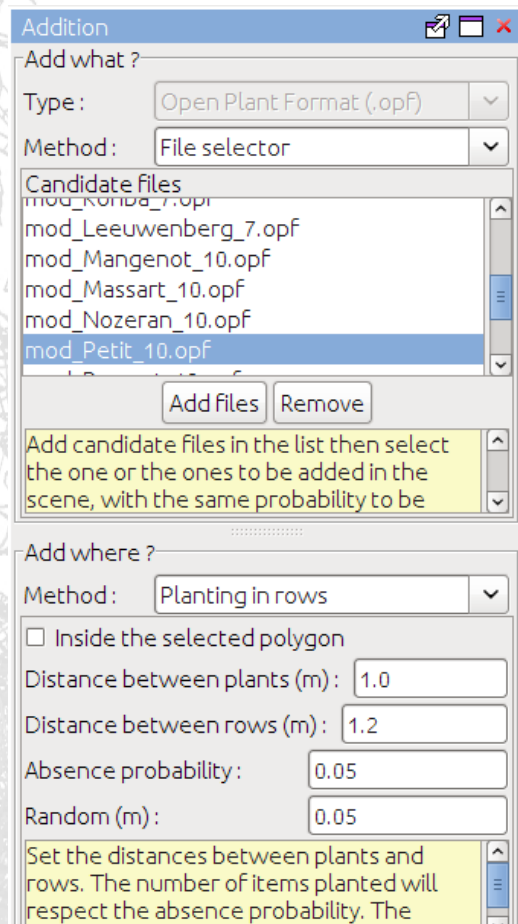


Load scene files

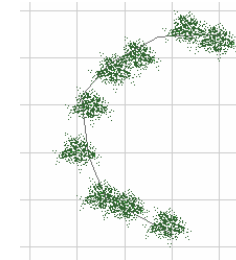
1	1	501.opf	0.04	0.24	0	1	0	0	0
1	2	502.opf	0.04	0.19	0	1	0	0	0
1	3	503.opf	0.04	0.14	0	1	0	0	0
1	4	504.opf	0.04	0.09	0	1	0	0	0
1	5	505.opf	0.04	0.04	0	1	0	0	0
1	6	506.opf	0.09	0.24	0	1	0	0	0
1	7	507.opf	0.09	0.19	0	1	0	0	0
1	8	508.opf	0.09	0.14	0	1	0	0	0
1	9	509.opf	0.09	0.09	0	1	0	0	0



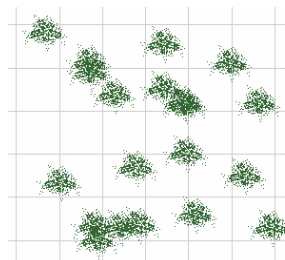
Create scenes with pattern plugins



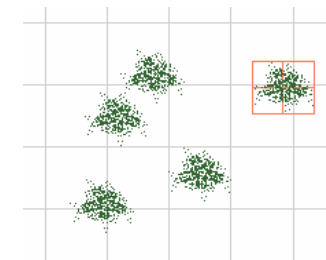
In rows



On a line



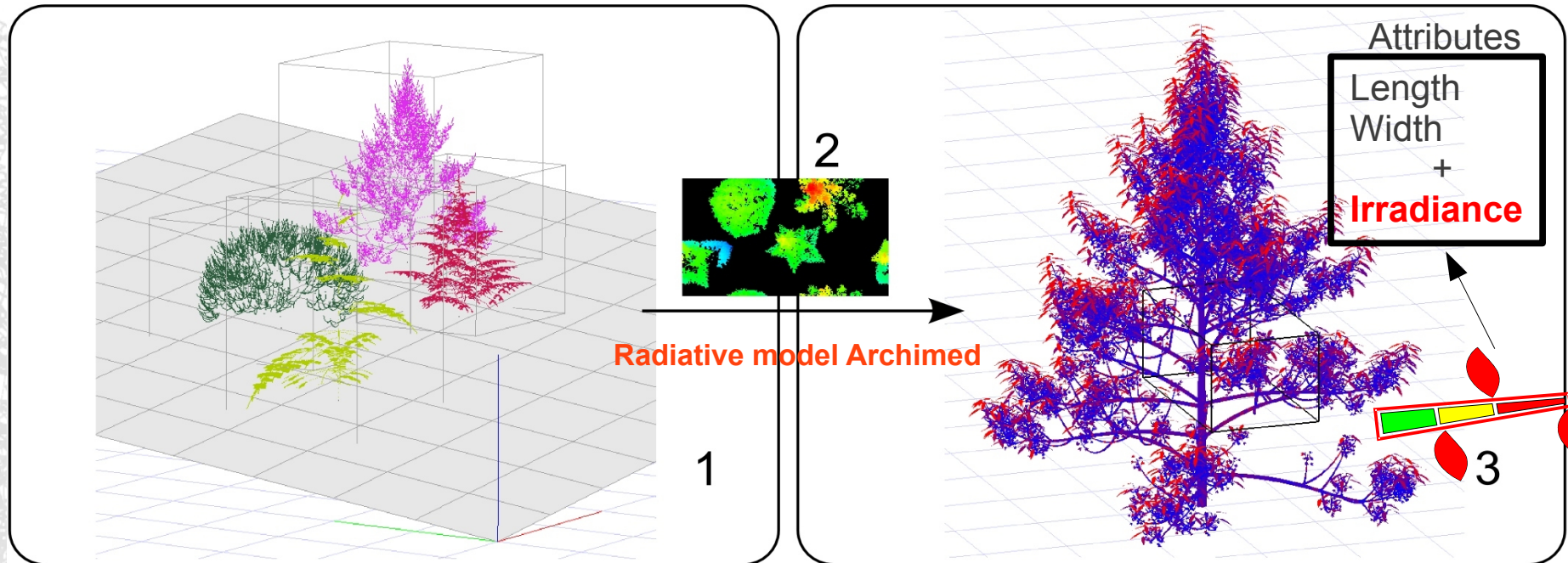
Random



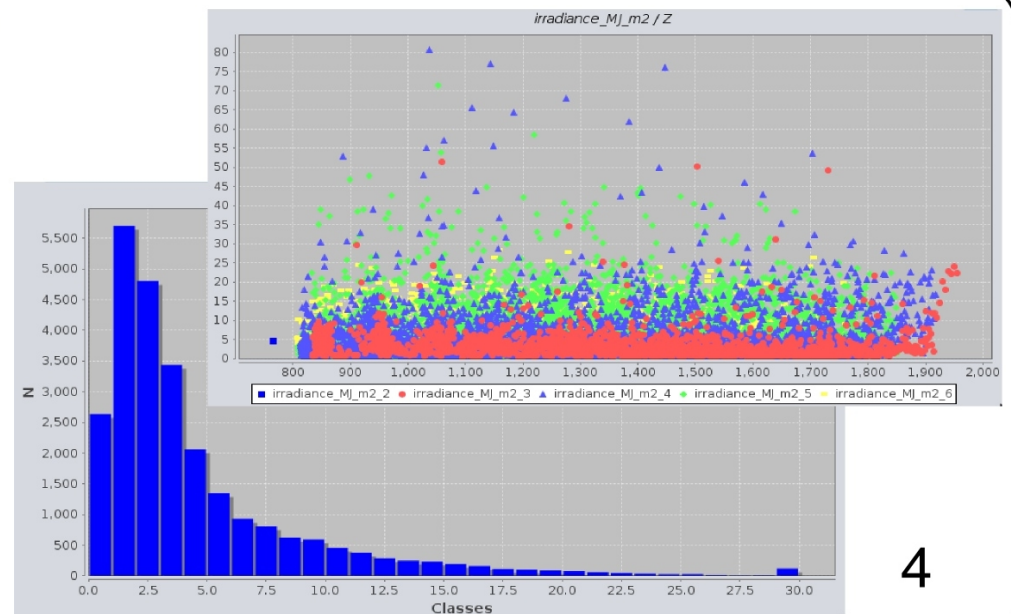
On mouse clicks...

And yours ...

III. Simeo : scene level processing integration



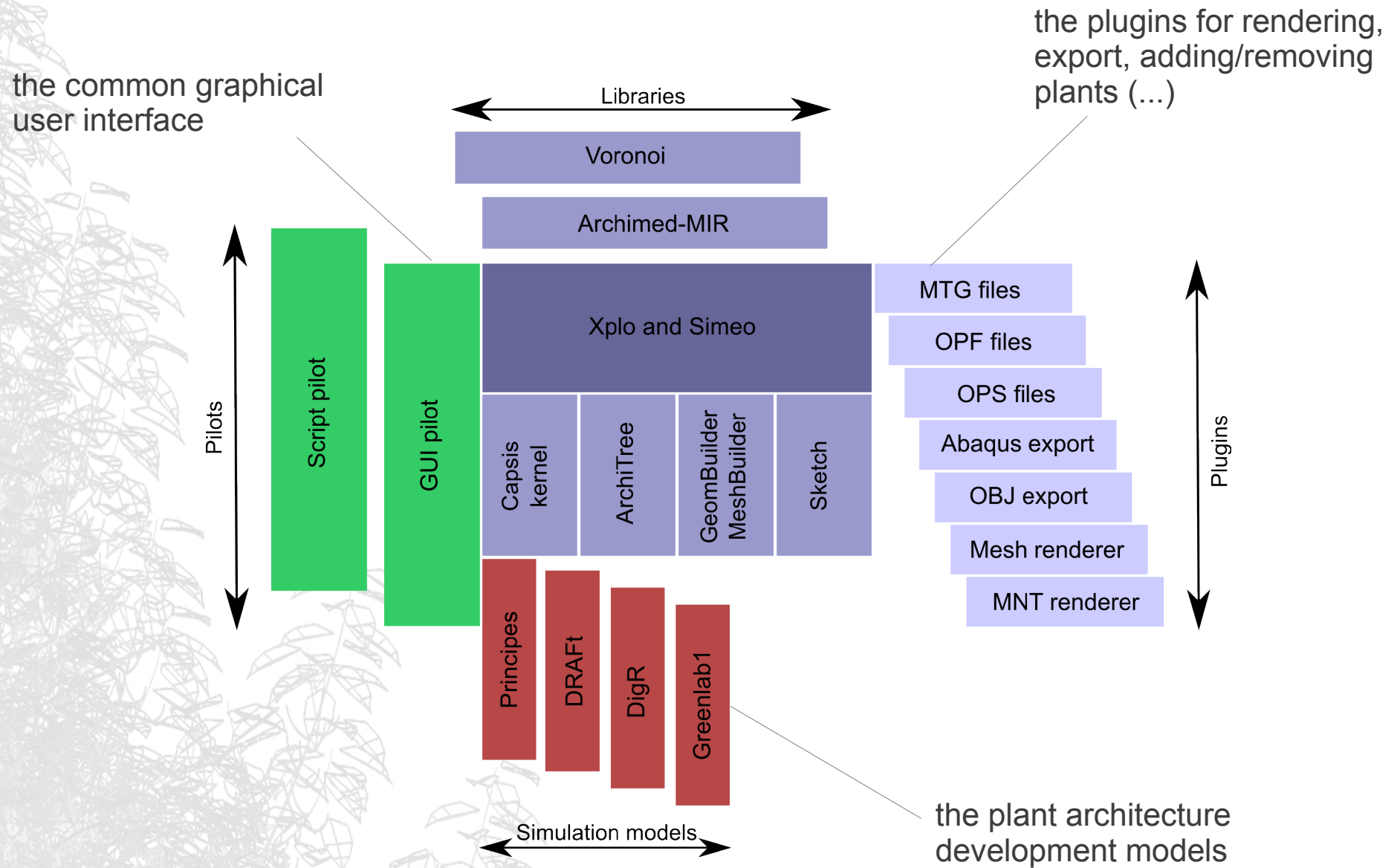
Item	Z	Area	Ramification order	irradiance_MJ_m2
branches1...	765.07	45467.3...	2	4.422
inflo5 1	807.936	98.304	6	5.271
branches4...	808.507	105.562	5	2.976
inflo5 1	808.507	98.306	6	10.148
branches4...	809.527	106.424	5	3.393
inflo5 1	809.527	98.305	6	4.935
inflo5 1	809.637	98.305	6	4.285
branches4...	809.692	31.133	5	1.049
inflo5 1	809.692	98.308	6	3.484
branches4...	809.961	31.133	5	1.987
inflo5 1	809.961	98.308	6	5.687
branches4...	810.018	63.836	5	2.647
branches4...	810.924	32.721	5	1.819
inflo5 1	810.924	98.309	6	1.327
inflo5 1	811.436	98.305	6	2.759
branches3...	811.908	140.948	4	3.334
inflo5 1	812.348	98.309	6	2.15
branches4...	812.385	34.308	5	0.952
inflo5 1	812.385	98.307	6	0.992



1 - Barczi J-F, Rey H, Caraglio Y, de Reffye P, Barthélémy D, Dong Q, Fourcaud T. AMAPsim: an integrative whole-plant architecture simulator based on botanical knowledge, *Annals of Botany* 2008; 101:1125-1138.

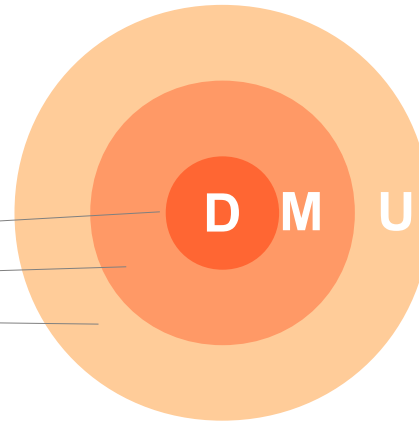
2 - Dauzat J, Clouvel P, Luquet D, Martin P. Using virtual plants to analyse the light-foraging efficiency of a low-density cotton crop, *Annals of Botany* 2008; 101:1153-1166.

IV. Simulation framework : Software design



IV. Simulation framework : Actors and roles

developers
modellers
end-users



The AMAPstudio community:
developers + modellers
co-develop together

Modeller
Christophe Proisy
IRD AMAP
Lollymangrove

Developer
Sébastien Griffon
Cirad AMAP Montpellier

Developer
Francois de Coligny
INRA AMAP Montpellier

Modeller
Hervé Rey
Cirad AMAP
Principes, Sunflower

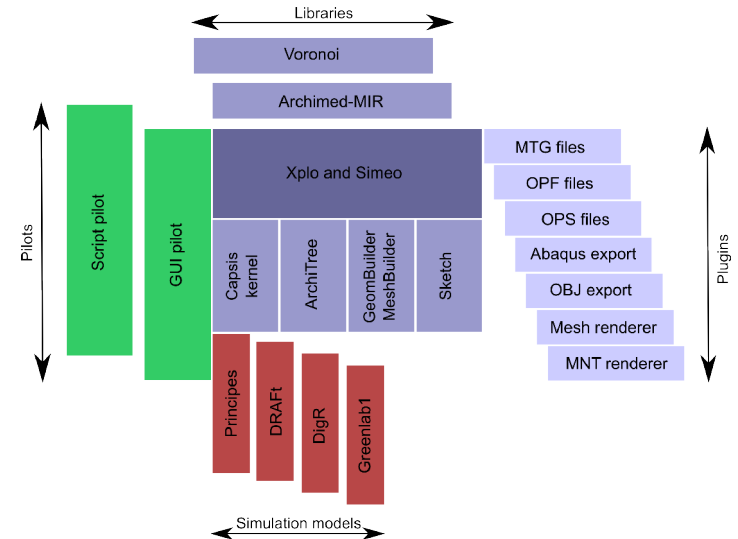
End-user
The MOCAF
network partners

Modeller
Jean Dauzat
Cirad AMAP
Archimed MMR, ART, Lidar

End-user
The StemLeaf
project partners

IV. Simulation framework :

The AMAPstudio Charter

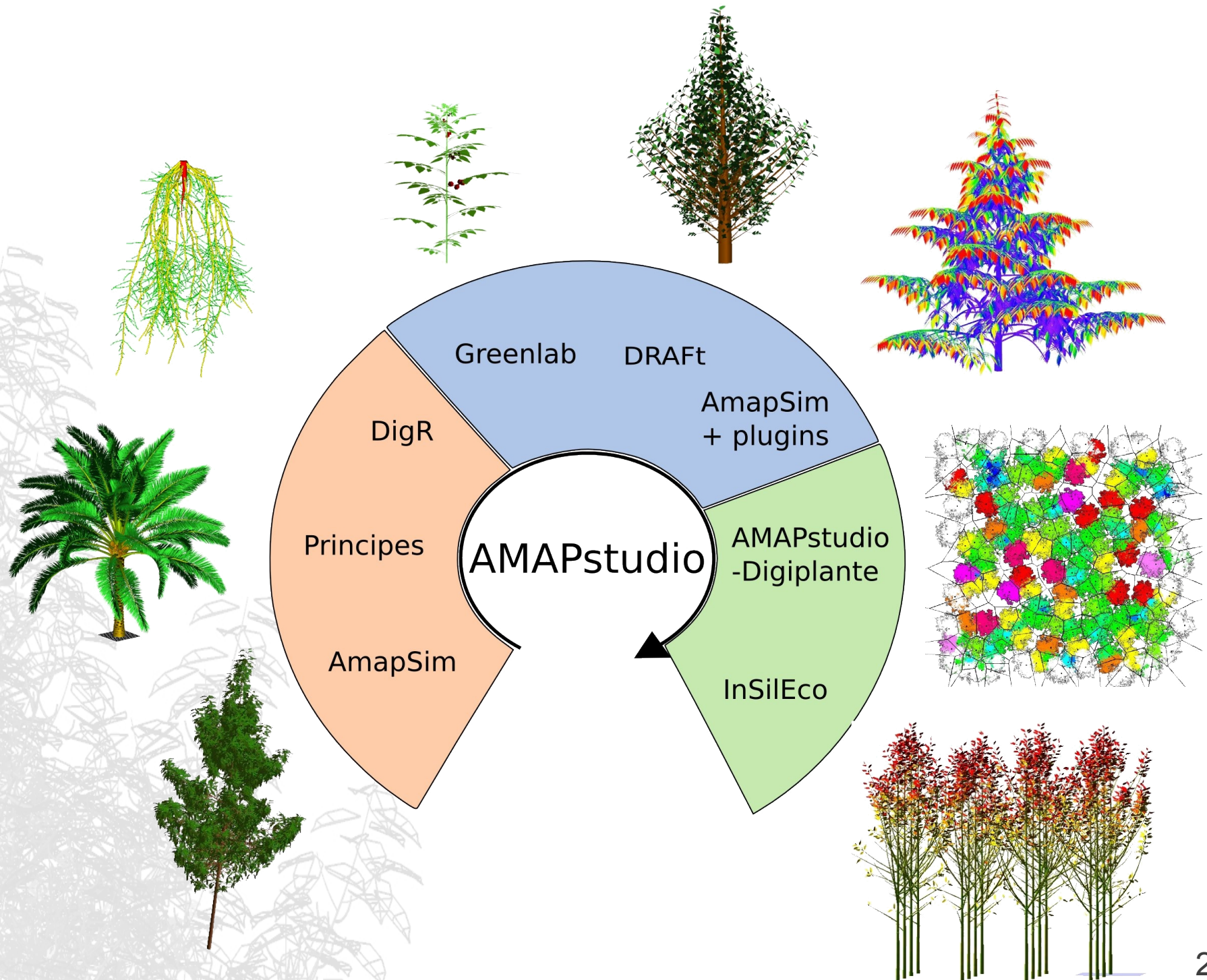


Clear participation rules

All the common parts are free software (LGPL), they are reusable by everyone
-> all **except the workspace/, xplo/module/ and simeo/module/ directories**

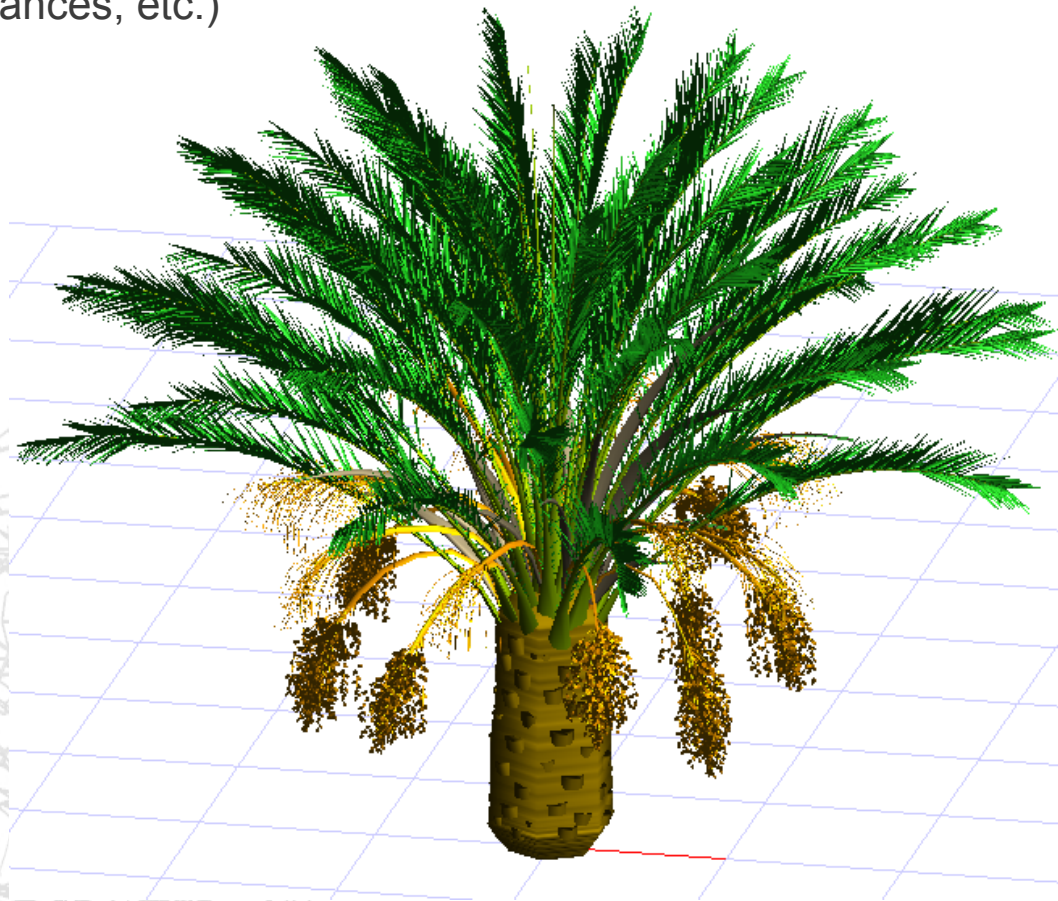
- **Free kernel:** the AMAPstudio kernel is a free software (LGPL licence) : **kernels** + **generic pilots** + **extensions** + **libraries**
- **Development:** the modellers are in charge of the development of their models in AMAPstudio
- **Support:** They can have support from the developers : training sessions, design, starting help, further assistance
- **Free access in the community:** All the source codes are freely accessible by all members in the AMAPstudio community, modules may become the base for new modules, code can be shared...
- **Respect of intellectual property:** all members respect the intellectual property of the other members
- **Validations:** developers deal with technical validation, modellers deal with functional validation
- **Distribution:** the stabilized / validated modules may be distributed when the author decides and chooses a licence (LGPL free license suggested)
- **Decentralization:** modellers manage directly the relations with their end-users: financing, training, assistance, models documentation, contracts...

IV. Simulation framework : various kind of simulators



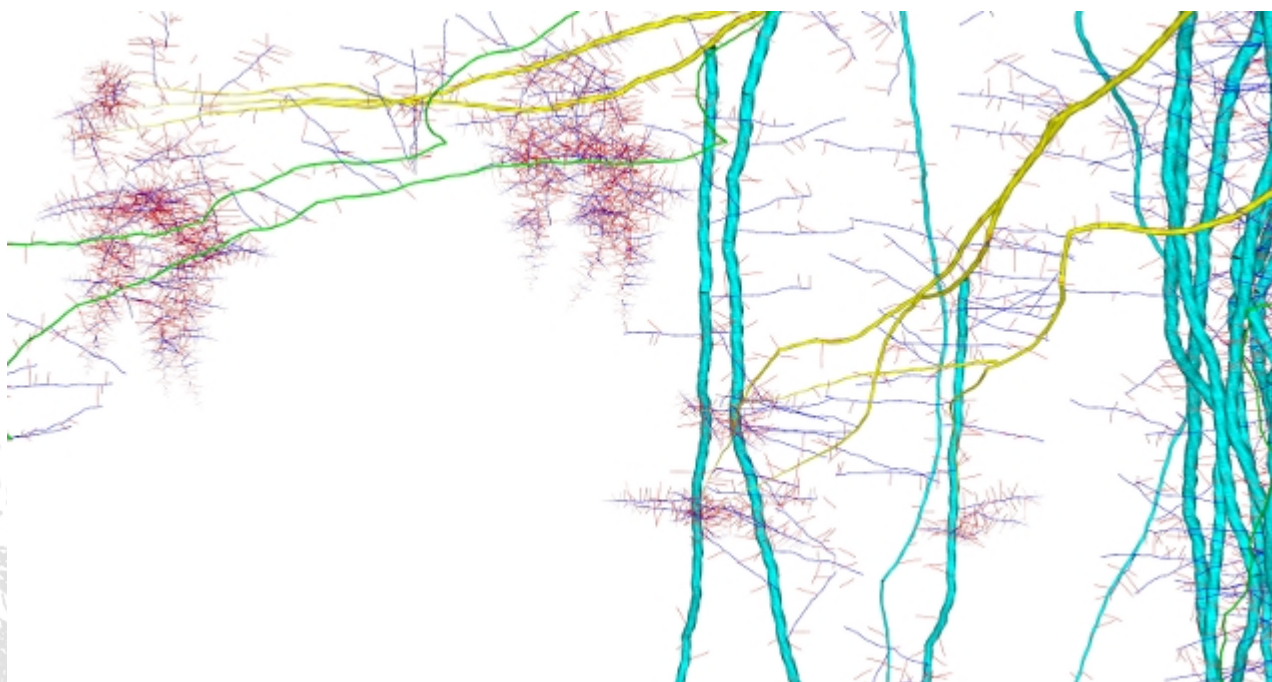
Principes

- models and simules the development and architecture of vegetative and reproductive parts of palm tree from germination to any age
- a continuation of studies that have been carried out for almost 20 years on palm-tree architecture
- Principes brings a generic, multi-scale, structural model for palm-trees based on the organisation of the various organs
- each organ carries its own attributes (lengths, diameters, branching or deviation angles, inter organs distances, etc.)



DigR

- DigR (Rey et al., 2011) is a root architectural model and simulator
- it relies on topological concepts as apical growth, lateral branching, senescence and death, and geometrical features as secondary growth and axes spatial positioning
- each of these properties are sorted into a root typology
- the current version runs without functional processes, however AMAPstudio will help developing further versions including functional – structural interactions during growth simulation and dealing with environmental influence (i.e. soil properties or aerial part contribution)



Greenlab 1

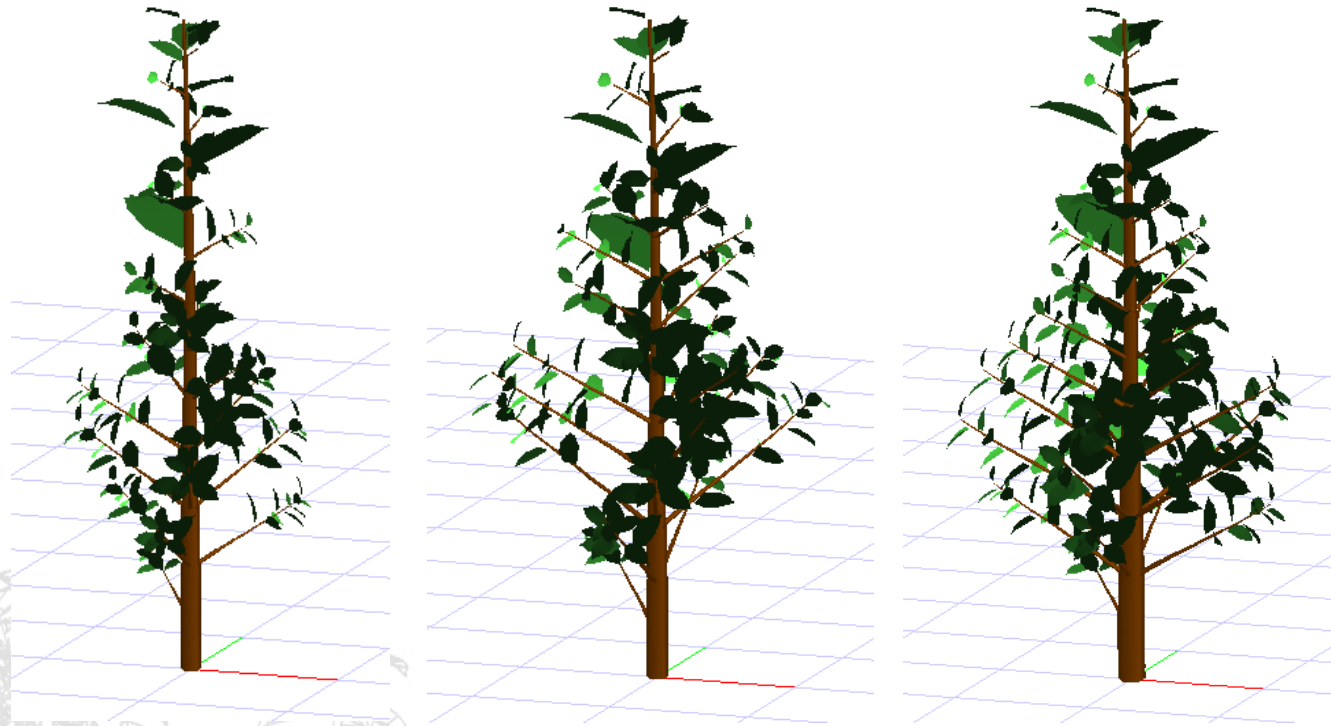
- Greenlab is a mathematical plant model simulating interactions between plant structure and functions
- biomass produced by organs (sources) is allocated to expanding organs (sinks) according to their relative demand
- plant parameters can be adjusted by fitting on real measurements
- Greenlab can compute the plant architecture for various species in interaction with their environment
- this model's underlying concepts can potentially predict the plant's phenotypic plasticity, for instance, pruning affects the internal competition for the resources and may lead to different leaf sizes



de Reffye P, Hu B-G. Relevant qualitative and quantitative choices for building an efficient dynamic plant growth model: GreenLab case. In: Hu B, Jaeger M, eds. Plant Growth Modeling and Applications (PMA03); Proceedings of the 2003 International Symposium on Plant Growth Modeling, Simulation, Visualization and Their Applications; Tsinghua University Press, Springer; 2003. p. 87-107

DRAft (Demand, Resource, Architecture and Functioning at discrete time)

- a minimal FSPM designed to simulate emerging plants morphogenetical gradients
- tree architecture gradients is an emerging property of the interplay between structure function and iterative development
- DRAft simulates the development and functioning of the tree aerial part at a yearly step
- it is based on biomass allocation, and relies on a 6 parameters equations system
- minimalist approach -> possible to use analytical tools to study the model sensitivity and behaviour

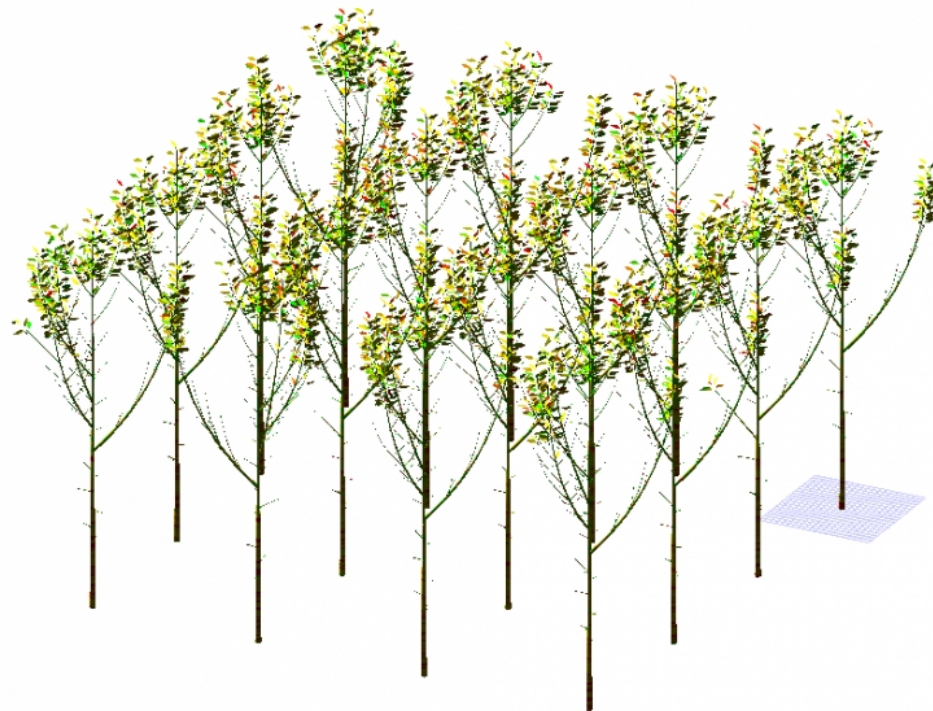


Taugourdeau O, Barczy J-F, Caraglio Y. Simulation of Morphogenetical Gradients Using a Minimal Functional-Structural Plant Model (FSPM). In: Kang M., Dumont Y., Guo Y., eds. Plant Growth Modeling, Simulation, Visualization and Applications. Proceedings of PMA12. Shanghai, China: IEEE press; 2012. p. 379-387.

InSilEco - Architectural plasticity in ecological communities

- simulate the growth of several tree individuals in a forest stand
- the growth of an axis depends on the amount of light that it and its leaves receive
- the simulation process then combines architectural rules defined at species level (AmapSim), an illumination module used to assess light interception by leaves (MIR), and a coupling of axis growth and ramification capabilities with local light interception

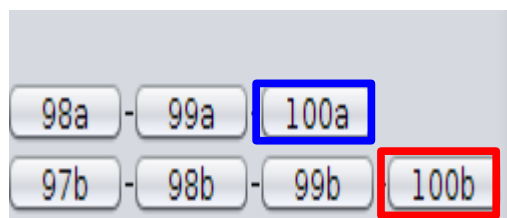
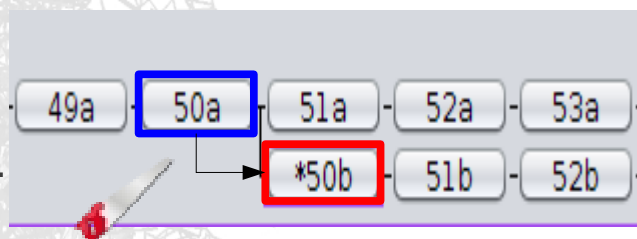
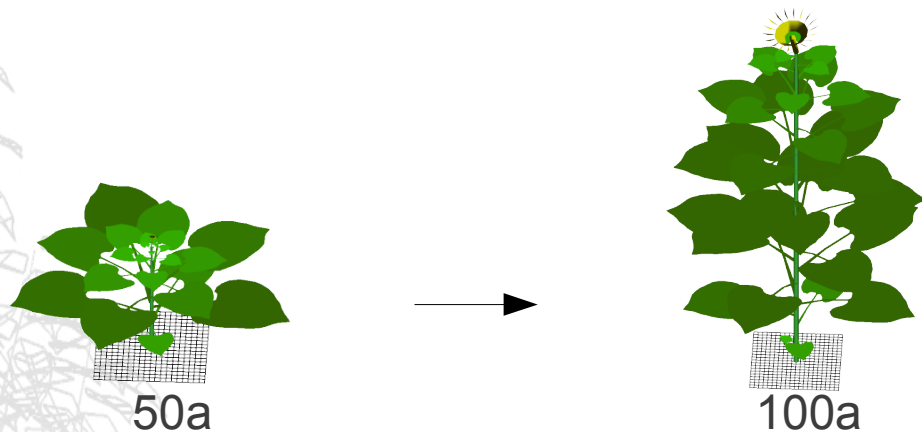
The objective is to assess the effect of competition for light within and between species on the overall structure of the forest stand



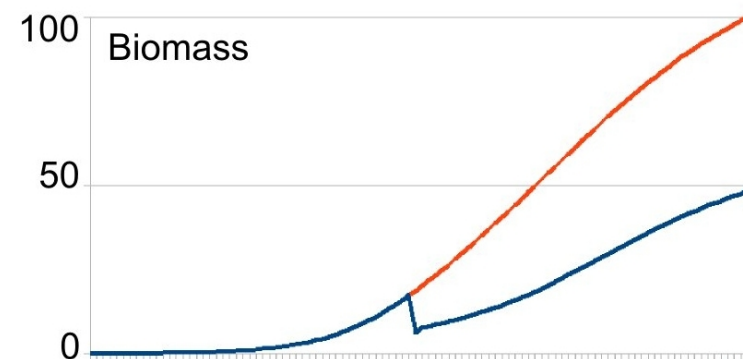
IV. Simulation framework : plant growth scenarios

Greenlab simulator - Sunflower

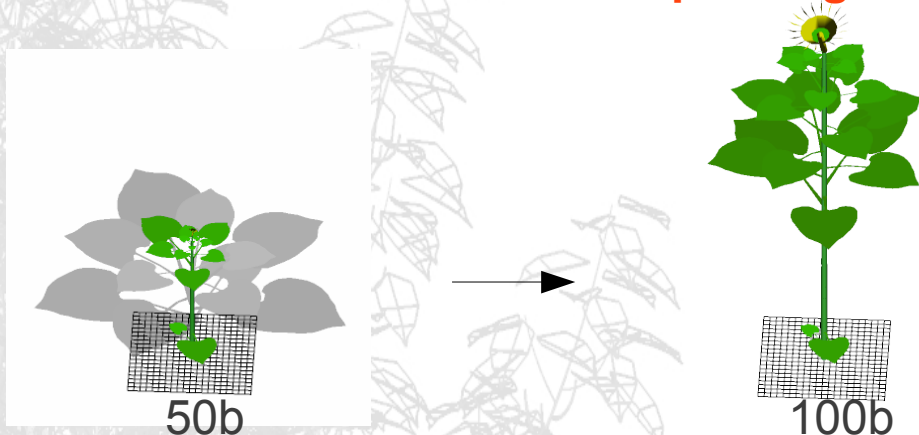
Scenario a : control scenario



Normalized plant growth



Scenario b : leaf pruning at 50



V. Common features

Import/Export various file formats : plug-ins

IMPORT format

AMAPstudio formats :
OPF (Open Plant Format)
OPS (Open Plant Scene)

Other formats :
MTG
LIG + DTA

... Add your formats ...



EXPORT format

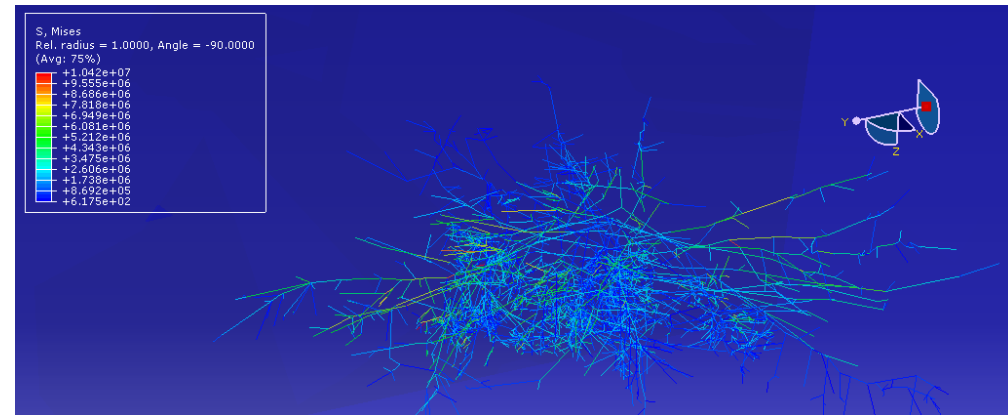
AMAPstudio formats :
OPF (Open Plant Format)
OPS (Open Plant Scene)

Other formats :
MTG
LIG + DTA
Sunflow
Abaqus INP
Obj (Wavefront)

... Add your formats ...



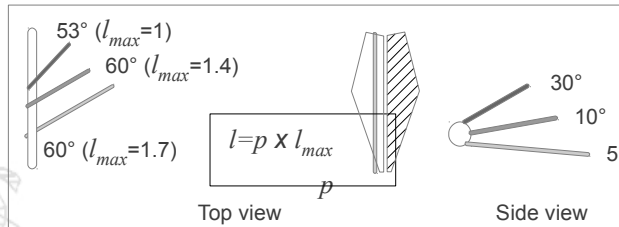
Digitized Maritime pine root system with PiafDigit
(F. Danjon - INRA)



Abaqus export for tree overturning simulation
(T. Fourcaud - CIRAD)

V. Common features

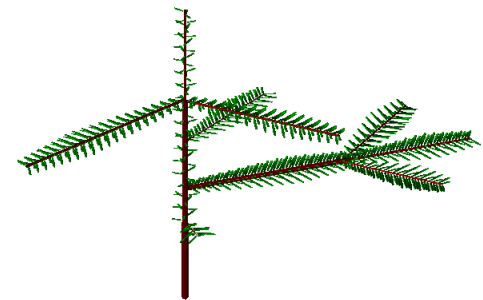
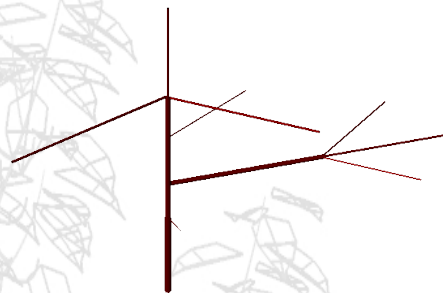
Scripts



```

114
115 SetOrientation= {vtx, X, Y, Z, Len, Xe ->
116
117   if (Length(vtx.getBearer().getComplex())!=null) longPa=( Length(vtx.getBearer().getComplex()) *10 )
118   else longPa=1
119   propLong = Index( vtx.getBearer() ) / longPa
120   LenAdj=Adj (Len,propLong)
121
122   if (vtx.getOrder()==3 | estRad(vtx) ) LenAdj=Len
123
124   vtx.set("RotBearerX",X)
125   vtx.set("RotBearerY",Y)
126   vtx.set("RotBearerZ",Z)
127   vtx.set("Length",LenAdj/coefLongSymboleAig)
128   vtx.set("RotLocalX",Xe)
129   return ""
130 }
131
132 Orientation = { vtx ->
133   NeedleType= Index(vtx.getBearer())%6
134
135   if (vtx.getOrder()==3 | estRad(vtx) ) { NeedleType="A1"}
136   if (vtx.getOrder()==4) {AngleA3=180};//90+
137   else {AngleA3=90};//180
138
139   switch( NeedleType ){
140
141     case "A1" : An=(An+141)%360
142                SetOrientation(vtx,      An              ,      -AngleAigA2, 0,      0.
143                break
144
145     case 0    : SetOrientation(vtx,      AigLayer_1_AngPhyl +AngleA3, AigLayer_1_AngInsert, 0, AigLayer_1_Lengt
146                break
147
148     case 1    : SetOrientation(vtx,      - AigLayer_2_AngPhyl +AngleA3, AigLayer_2_AngInsert, 0, AigLayer_2_Lengt
149                break
150
151

```



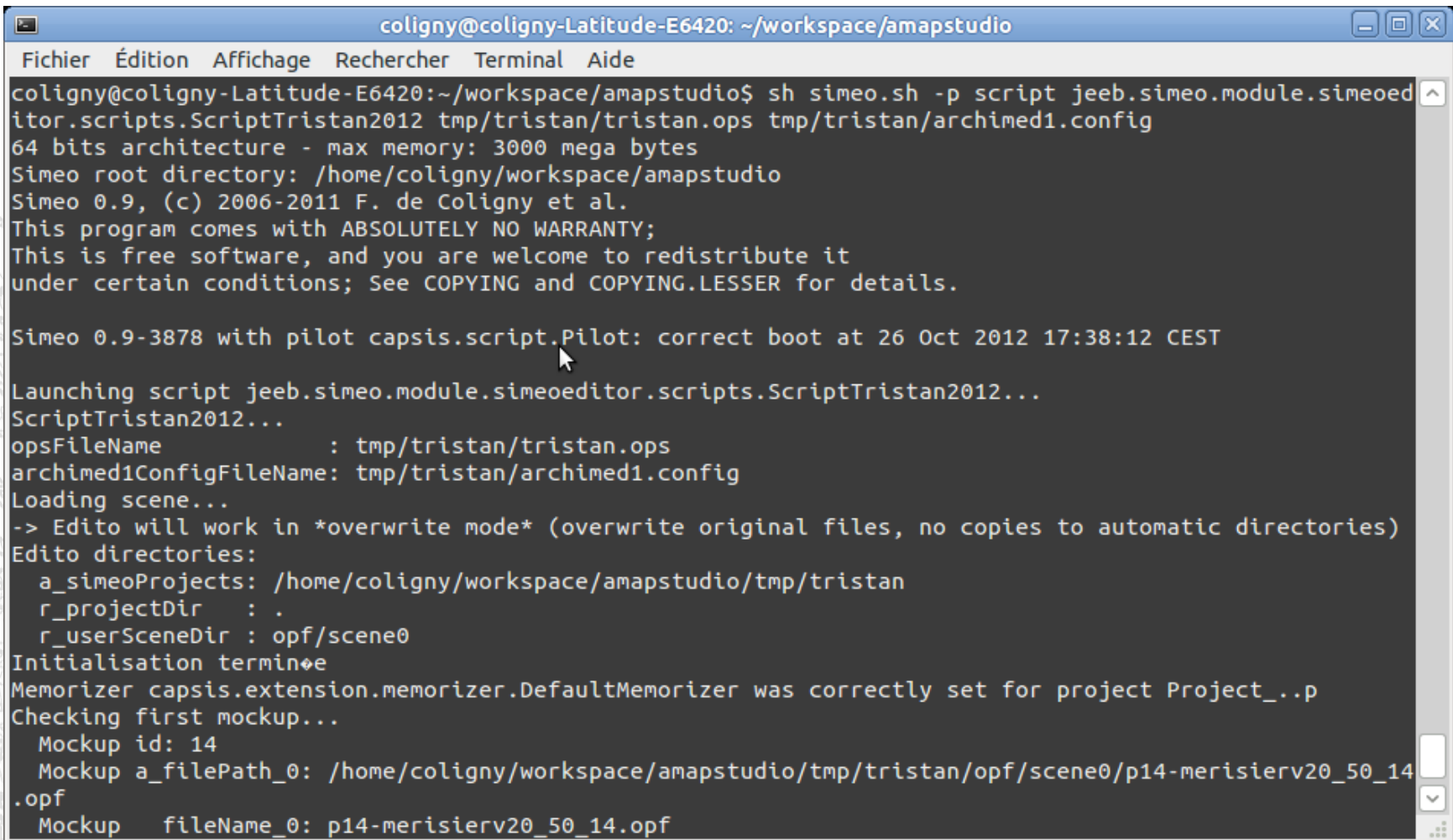
Taugourdeau, O., Dauzat, J., Griffon, S., Sabatier, S., Caraglio, Y., Barthélémy, D., 2012.

Retrospective analysis of tree architecture in silver fir (*Abies alba* Mill.):

ontogenetic trends and responses to environmental variability. *Annals of Forest Science*, 69(6) : 713-721 p.

V. Common features

Batch mode : run without Graphical User Interface



```
coligny@coligny-Latitude-E6420: ~/workspace/amapstudio
Fichier  Édition  Affichage  Rechercher  Terminal  Aide
coligny@coligny-Latitude-E6420:~/workspace/amapstudio$ sh simeo.sh -p script jeeb.simeo.module.simeoeditor.scripts.ScriptTristan2012 tmp/tristan/tristan.ops tmp/tristan/archimed1.config
64 bits architecture - max memory: 3000 mega bytes
Simeo root directory: /home/coligny/workspace/amapstudio
Simeo 0.9, (c) 2006-2011 F. de Coligny et al.
This program comes with ABSOLUTELY NO WARRANTY;
This is free software, and you are welcome to redistribute it
under certain conditions; See COPYING and COPYING.LESSER for details.

Simeo 0.9-3878 with pilot capsis.script.Pilot: correct boot at 26 Oct 2012 17:38:12 CEST

Launching script jeeb.simeo.module.simeoeditor.scripts.ScriptTristan2012...
ScriptTristan2012...
opsFileName      : tmp/tristan/tristan.ops
archimed1ConfigFileName: tmp/tristan/archimed1.config
Loading scene...
-> Edito will work in *overwrite mode* (overwrite original files, no copies to automatic directories)
Edito directories:
  a_simeoProjects: /home/coligny/workspace/amapstudio/tmp/tristan
  r_projectDir   : .
  r_userSceneDir : opf/scene0
Initialisation terminée
Memorizer capsis.extension.memorizer.DefaultMemorizer was correctly set for project Project_..p
Checking first mockup...
  Mockup id: 14
  Mockup a_filePath_0: /home/coligny/workspace/amapstudio/tmp/tristan/opf/scene0/p14-merisierv20_50_14
.opf
  Mockup   fileName_0: p14-merisierv20_50_14.opf
```

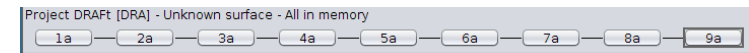
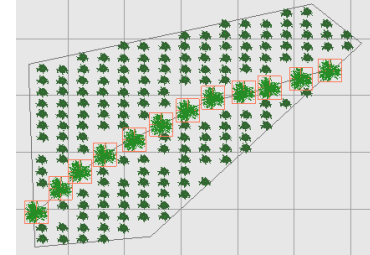
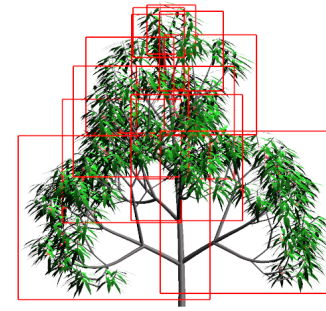
Repetitive simulation
Run on clusters
Sensitivity analyses

...

Conclusion

AMAPstudio assets

- For end-users : interactive editors, 3D view
- For modellers : collaborative framework, methodology and human support
- Scenario oriented : alternate growth and intervention steps to build scenarios
- Long-term mutualisation and capitalisation
- Easy distribution and transfer : free licence kernel, multi-OS (Windows, Linux, Mac)



Communication

- presented in PMA'12 in Shanghai, China
- presented in FSPM2013 in Saariselka, Finland
- a web site
 - documentation for the modellers: <http://amapstudio.cirad.fr/>
 - up to date projects list
- a reference paper accepted in Ecological Modelling



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- Publications
- Charter
- Contact
- AMAP

- ▶ AmapSim
- ▶ Archimed
- ▶ Simeo
- ▶ Xplo



Community news

- Chloé Bourden (Master student, IRD AMAP) used Simeo - Lollymangrove in order to (1) upload and verify her field inventory data, (2) calculate the plots' biomass and carbon with different allometry models and (3) visualise the type of structure for mangrove populations. (C. Bourden, 2.7.2013)
- AMAPstudio was presented by S. Griffon at FSPM2013, the 7th International Conference on Functional – Structural Plant Models in Saariselka, Finland (9-14th june). (F. de Coligny, 14.6.2013)
- A Palm tree workshop related to ecophysiology and modelling has been organized by Cirad last 22nd May 2013. During this workshop, the AMAP lab presented a collective contribution showing the past collaborations with the other teams in Montpellier, the new projects, the goals regarding scientific results and applications, and the possible partnerships to reach them. [2013-05-22-rey-journee-palmier-a-huile-agap.pdf](#) (H. Rey, 23.5.2013)
- René Lecoustre, Jean-Francois Barczi and Hervé Rey attended the First International Meeting on Phylogenetic Resources of Date-Palm from 15 to 17 April 2013 in Djerba, Tunisia. They presented two communication on (i) statistical studies on the allometric relationships for the vegetative part (PRINCIPES model) and (ii) root architecture analysis and modelling (DigR model) of Phoenix dactylifera. After this meeting, a working program was built for the next two years concerning the MOCAF and PHC Maghreb projects. (H. Rey, 29.4.2013)



- A paper about AMAPstudio was published in the IEEE proceedings of the PMA'12 international conference, see the Publications page for more details. (F. de Coligny, 25.11.2012)



<http://amapstudio.cirad.fr/>

Thank you for your attention !



AMAPstudio SDK

AMAPstudio :

- **Libraries :**

- Maths : common math functions
- Formats : OPF, OPS, MTG readers/writers
- Sketch : MVC GUI library
- Structures : ArchiTree and Geometry builder.
- SimulationTools : Scheduler, Listeners, Meristems, Organs.
- Utils : common usefull functions (e.g. AutoUI)

- **Workspace :**

- Sunflower model
- Principes model
- Draft model
- Greenlab model
- ...

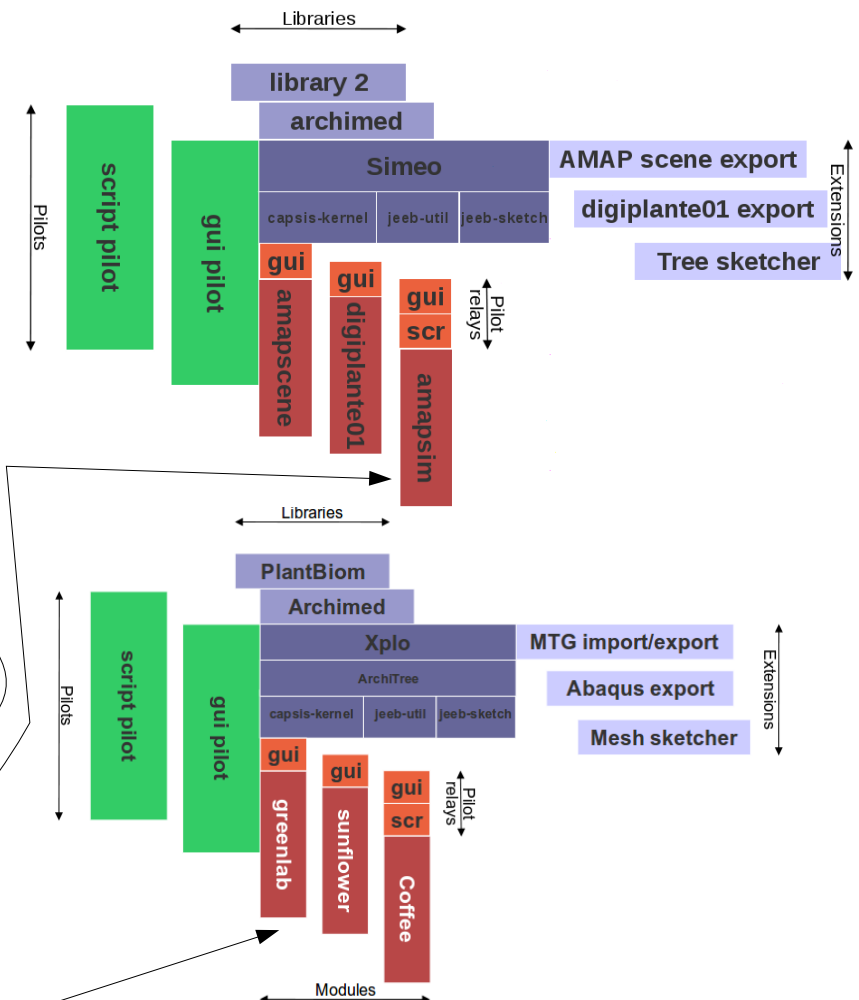
- **Xplo :**

- Kernel
- Gui
- Extensions
- Modules :
 - PlantEditor
 - Sunflower
 - Greenlab
 - ..

- **Simeo :**

- Kernel
- Gui
- Extensions
- Modules :
 - AmapScene
 - Greenlab

Modeller inputs



AMAPstudio singularities

Software aiming at mutualisation in the domain of plants modelling may all have intersections of their scopes.

Here are the AMAPstudio's singularities :

- Editors for end-users, to do simple things quickly (development skills not needed).
- Modellers comfortable with programming can write scripts or plugins.
- Focusing on the plants architecture, from the plant to the forest stand.
- A simulation framework to build and compare scenarios.
- Co-development and charter based methodology.
- Simple procedure to build custom installers for distribution and installation.