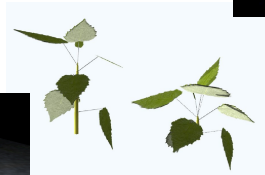
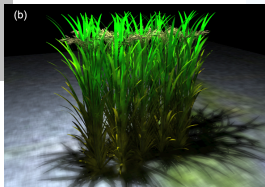


FSPM-P: A general FSPM prototype model

A first step towards a generic FSPM

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"Modelling and Simulation with GroIMP"
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Initial situation

- Developer with:
 - No or little programming experience → challenge of programming
 - Little knowledge of biological systems → challenge of modelling
- Common way of modelling:
 - Ad hoc
 - Usually starting "from scratch"
 - (initially) no (clear) concept/design
 - Unsystematic
 - Extend/change existing models
 - Getting overview of code more difficult
- ⇒ "Reinventing the wheel"

But...

- Same components and recurring parts
- Subsystems can be re-used
- Benefit from former models

⇒ Prototype

Advantages of a general FSPM-Prototype

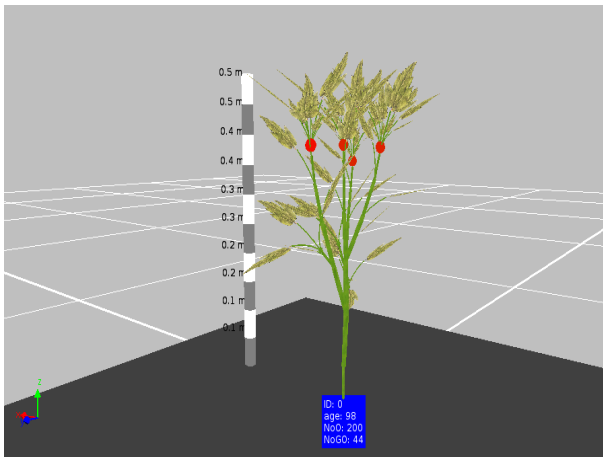
- Includes the (meta)knowledge of several models
- Predefined:
 - (Clear) proved design (software-technical)
 - Model concept: (init → control flow → output ...)
- Easy to parameterize, extend and use
- Rationalizes the development process and use of FSPMs
- Makes models comparable and combinable
- Facilitates communication between modeller, experimentator and programmers → can help make it a more accessible tool

FSPM-Prototype

Since Nov. 2009:

- Base model: rapeseed model by C. Groer
- Reworking the concept and the design
 - Strictly object-oriented
 - Modular design
- Extending by several parts
 - Model control
 - Photosynthesis rate models (differing in complexity from simple light-response curves to biochemical Farquhar-type models)
 - Source-Sink Ratio for model regulation
- June 2010: FSPM-P V0.02

FSPM-P Structure



FSPM-P Design

Plant definition

SpeciesParameterS1.rgg

OrganS1.rgg

RulesS1.rgg

Model control

main.rgg

```
#init(): void  
+growDaily(): void  
+growHourly(): void
```

Global constants

Parameter.rgg

```
+initParameters(): void  
-readClimateData(): void  
-readClimateDataFile(): void
```

Photosynthesis models

LEAFC3 / LEAFC3N

LEAFC3.java

```
+compute(): void
```

Species.java

LEAFC3N.java

```
+compute(): void
```

SpeciesN.java

Environment.java

Functions.rgg

```
+LiethPasian(): double  
+Thornley(): double
```

KimLieth.java

```
+estimatePS(): double
```

Others

Charts.rgg

```
-INDI_ID_A: int  
-INDI_ID_B: int  
-CHARTS: int[]  
-CHARTS_VS: int[]  
#initCharts(): void  
#updateCharts(): void
```

Tools.rgg

ToDo.txt

readme.txt

Changes.txt

MyFileChooser.java

```
+getFile(): File
```

Organ definitions

- Same organ interface for all organs
- Different hierarchical scales within the plant:
 - Organs: seed, root, bud, leaf, internode, flower, fruit, ect.
 - organ with predefined state variables and methods representing internal processes: photosynthesis, growth, maintenance and growth respiration
 - Organs aggregations: individual, shoot
 - containing summary functions based on XL-queries (e.g. for whole-plant photosynthesis)

Model work flow

- Initialisation
- Growth step (daily / hourly)
 - ① Update sky and sun
 - ② Run the light model
 - ③ Apply rules:
 - Update Organs (Functions, shape, etc.)
 - Morphology and Cut
 - Transport rules (currently not used but predefined)
 - ④ Update Output (Charts / Files)
- Summary / Final output
 - Harvest / Yield Chart

Model work flow

Main loop:

```
1 public void growDaily() {  
  ...  
3  if (dayofyear<MAX_STEPS) {  
    dayofyear++;  
5  
    Tools.INSTANCE.updateLight();  
7    RADIATION.compute();  
    RulesS1.INSTANCE.applyRules();  
9    updateCharts();  
11   ...  
   }  
13 }
```

Model work flow

Rules:

```
1 protected void applyRules() {  
3     morphologyRules();  
     cutRules();  
5     //transportRules();  
     organUpdates();  
7     otherRules();  
9 }
```

Model work flow

Morphology rules:

```
1 private void morphologyRules() [  
    s:Seed, (s.isGerminateConditions()) ==>  
3     Root(s.getIndiID()) s Bud(1,1, s.getIndiID());  
  
5     b:Bud(rank, order, indiID), (b.isGrowingConditions()) ==>  
        RV(-0.15)  
7     Internode(rank, order, indiID)  
        [Leaf(rank, order, indiID)]  
9     if(b.isBranchingConditions()) (  
        [RL(30) Shoot0(indiID) Bud(rank+1, order+1, indiID)]  
11    )  
        Rotate(random(-5,5), random(-5,5), PHYLLLOTAX + random(-5, 5))  
13    Bud(rank+1, order, indiID);  
  
    ...  
15 ]
```

Model work flow

Growing Conditions:

```
1 public boolean isGrowingConditions() {  
    ...  
3 return  
    // architectural conditions  
5 rank<11 && order<=2 &&  
    // absorbed sensed power  
7 absorbedSensedPower &&  
    // average SSR as growth regulation  
9 (indi.getAverageSourceSinkRatio())>0.66) &&  
    // plastochron  
11 plastochron<=0 ||  
    // average SSR as growth regulation: building new sinks,  
13 // if avgSSR is to big and the bud is old enough  
    (indi.getAverageSourceSinkRatio())>MAX_AVERAGE_SOURCE_SINK_RATIO  
    && tempsum>200);  
15 }
```

Carbon budget

- Source: a portfolio of photosynthesis models (Farquhar, Thornley, KimLieth, LiethPasian, ...)
 - Collect produced carbon to a central pool CP in the individual (Partial softening of Central Pool concept: growth of leaves with local pool, remainder of local pool fed into the central pool.)

$$CP_t = CP_{t-1} + \Sigma(PS_{all \text{ growing organs}})$$

- Sink: Based on modified relative sink strength concept:
 - For each organ i :
 - Potential growth rate PGR_i (derivative of logistic or Beta distribution)
 - Relative sink strength $RSS = PGR_i / \Sigma(PGR_{all \text{ growing organs}})$
 - Actual growth (rate) $AG_i = RSS * CP_{t-1}$
 - At the individual scale:
 - $CP_t = CP_{t-1} - \Sigma(AGR_{all \text{ growing organs}})$

Regulation of processes

- \implies Observation: Source-sink ratio (SSR)¹
- Dynamic average SSR , exhibits a range, depending on source or sink-limitation of 0.1 to 1.5
 - Processes regulation
 - If $avgSSR > 1.1$:
 - decrease source strength (via PS efficiency)
 - increase number of sinks (bud break)
 - increase organ PGR
 - < 0.9 : vice versa

¹Marcelis, L.F.M. 1996. Sink strength as a determinant of dry matter partitioning in the whole plant.

FSPM-P - Project

Features:

- FSPM-P Model
- User manual
 - Model description
 - Building A New Model
 - Experimental Settings
 - Measurement protocol
 - 2D Digital Measurements
 - Data processing
 - Results
 - Templates
 - ...

FSPM-Prototype V0.02 for GroIMP
User Manual

Michael Henke and Gerhard H. Buck-Sorlin

September 24, 2010

What can be modelled

- Individual plants
- Plant stands (canopies)
- Temperature-sensitive processes
- Variations in morphology under certain growth conditions
- ...

Application areas

- Efficient model development
- Intercropping
- Demonstration purpose
- Education and understanding
- Decision support; yield equation

Planned/Started Projects

- Rapeseed model:
Tian Tian, G. Buck-Sorlin, M. Henke; 2010/11, Hangzhou, China
- Rice model:
Wu Lingtong, M. Henke; 2010/11, Hangzhou, China
- Cotton model:
G. Buck-Sorlin, Zhang Lizhen; 2010/11, Beijing, China
- Wheat/maize intercropping model:
Zhu Junqi, G. Buck-Sorlin, 2010-14, Wageningen, NL

Next steps

Generally

- Fine tuning of parameters
- Implement a species specific model
- Extend and complete the user manual

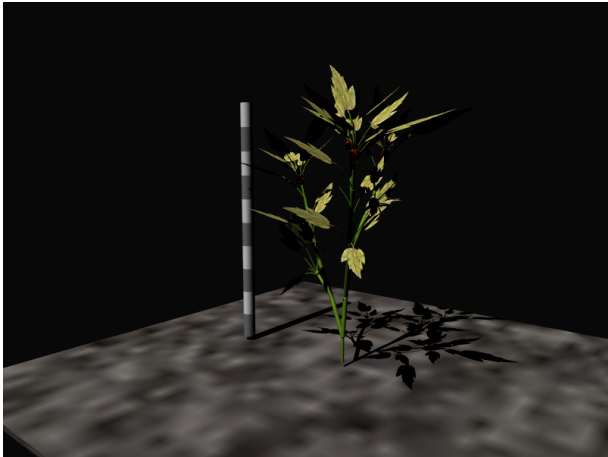
Improvements

- Modularize the photosynthesis models (Standardisation, identify submodels)

Outlook

Extensions

- Soil model (→ Root model)
- QTL genotypes as model parameters
- Breeder (Genetic Algorithm)
- Transport phenomena: water, hormones, signals (→ water stress, control bud breaks,)
- Central carbon pool concept \implies local carbon pool and transport based concept
- Generalize rules



Thank you for your attention!

Ďakujem!

Dank je!

Danke!

Merci!

谢谢!

