

Étapes de conception d'un modèle de simulation d'épidémies fongiques aérienne en fonction de l'architecture du couvert.

Projet ANR Archidemio 2009-2012

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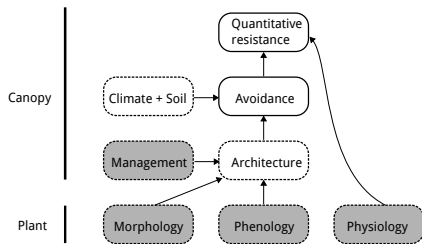
UMR 875 BIA et 1248 AGIR, INRA, France



Context

The canopy architecture as a lever to reduce diseases severity

- ▶ Chemical-based crop protection for air-borne diseases
- ▶ How to reduce treatments? (economy, ecology, policies)
 - specific resistance : eliminate epidemic
 - quantitative resistance : control (moderate crop injuries)



⇒ Adaptation of the canopy architecture to control the disease

Outline

Biology

- ▶ How much does the canopy architecture impacts the epidemic process ?

Methods

- ▶ Field experiments and process-based modelling to unravel crop × pathogen interactions.
- ▶ Four architecturally contrasted crop - pathogen systems

Conception

- ▶ How to deal with different disciplinary points of view ?
- ▶ Which level of architectural representation to adopt ?

Design methodology

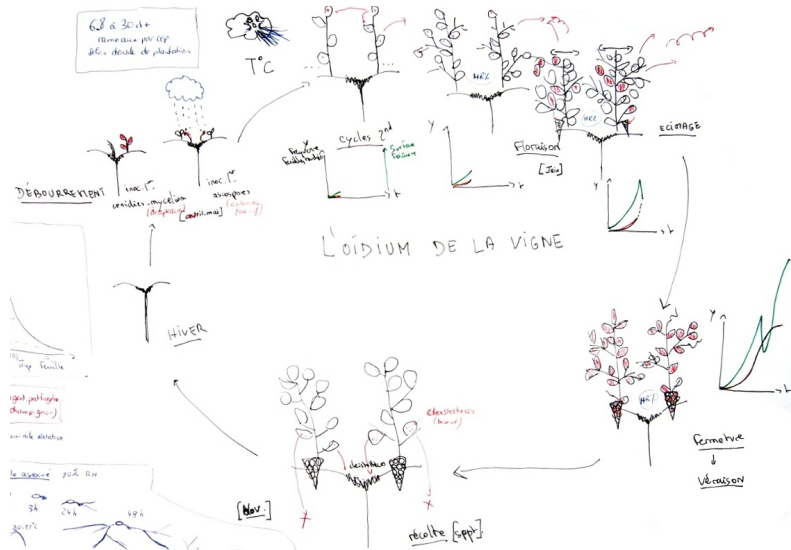
Define different modelling roles

- ▶ Applied mathematics modellers :
 - propose model framework (conceptual model)
 - propose equation shapes (formal model)
 - build interaction tool (software + formation)
- ▶ Thematic modellers :
 - define processes to include
 - adjust, parametrize equations (data or expertise)
 - propose elements for qualitative (behaviors) et quantitative (data) evaluations

Analogy with *top-down* systemic approaches :
observations \rightarrow [share model] \leftarrow concepts

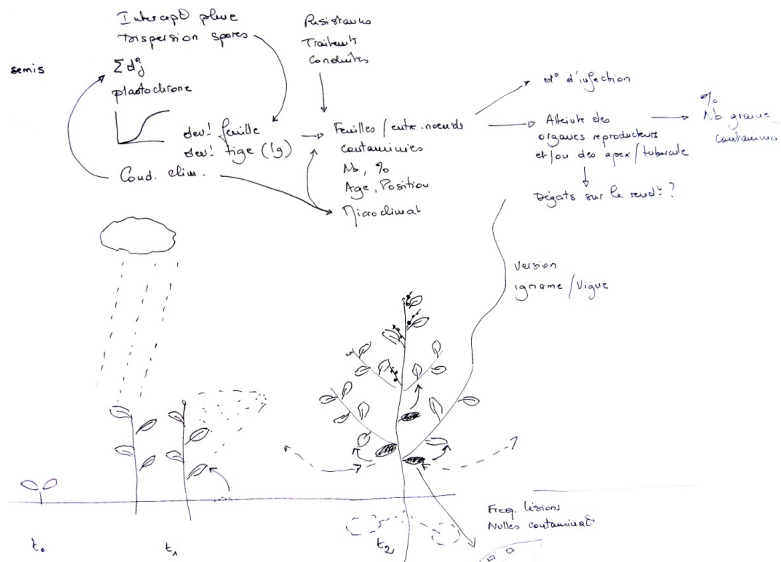
Grapevine conceptual model

focus : grapevine - powdery mildew disease cycle ; pathologist view



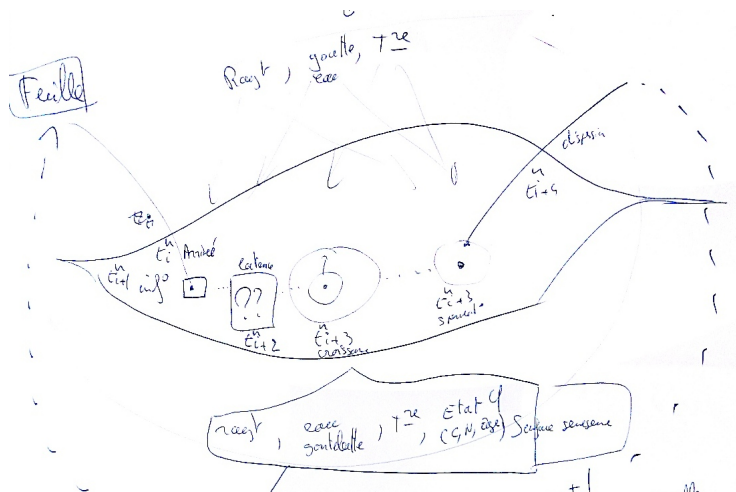
Yam conceptual model

focus : integration of epidemic processes ; ecophysiological view



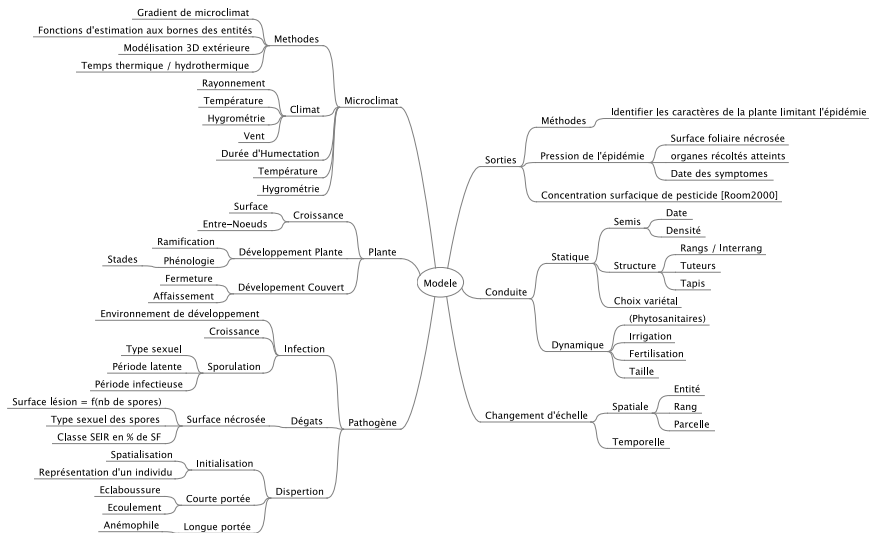
Disease conceptual model

focus : local scale disease development ; bioclimatologist view



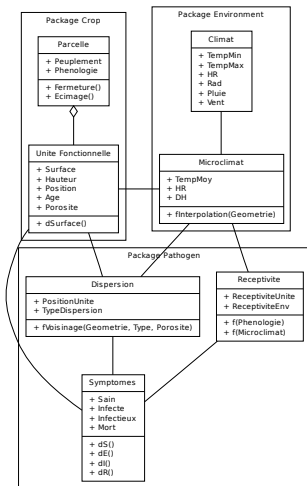
General conceptual model

global view on pathosystems



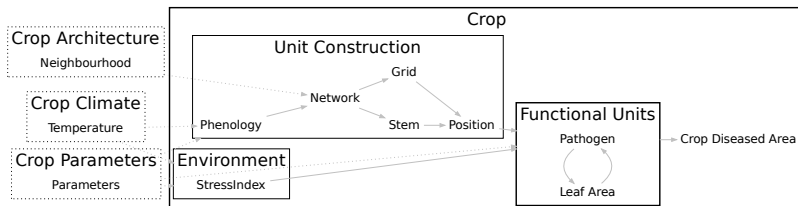
General UML conceptual model

focus : identification of main processes



Formal model : general structure

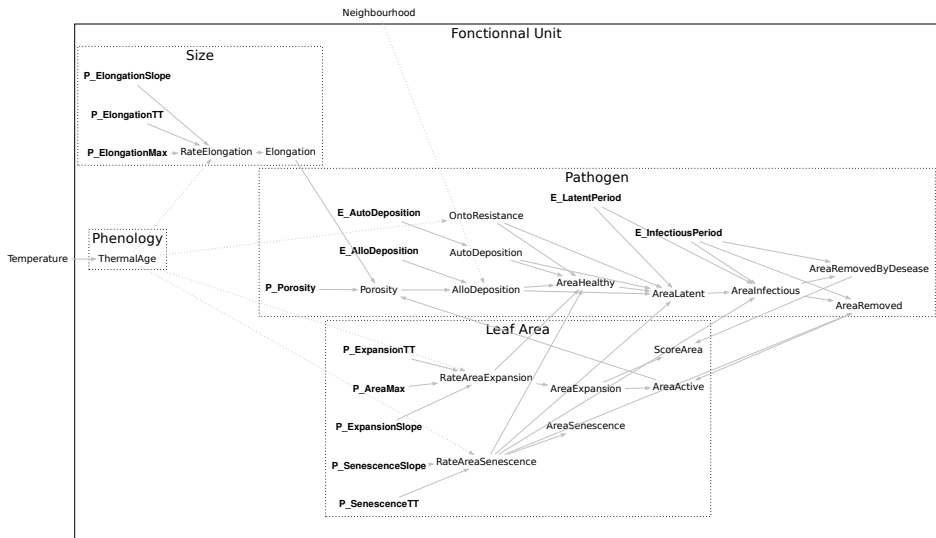
Setting a generic description of 4 pathosystems



- ▶ Controller model (structure) + individual model (fonction)
- ▶ Pathosystem-specific parametrization
 - functional units : homogeneous architectural element
 - network of connections : crop architecture and dispersion type

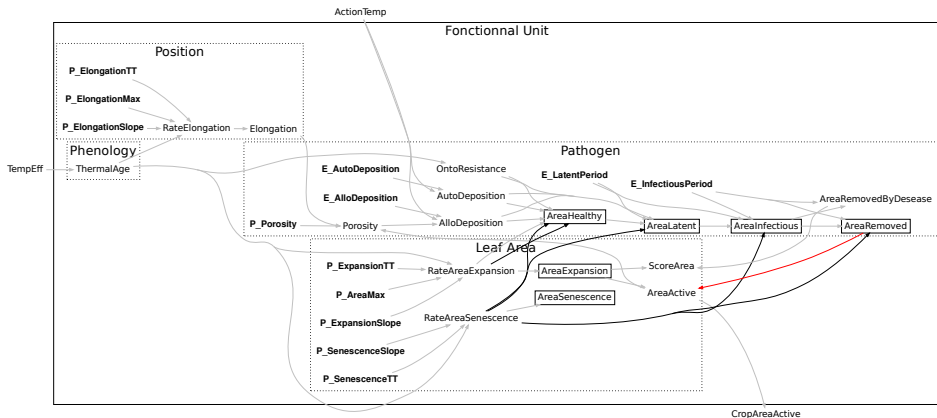
Formal model : structure of functional unit

crop (logistic function) + epidemiological (SEIR) models



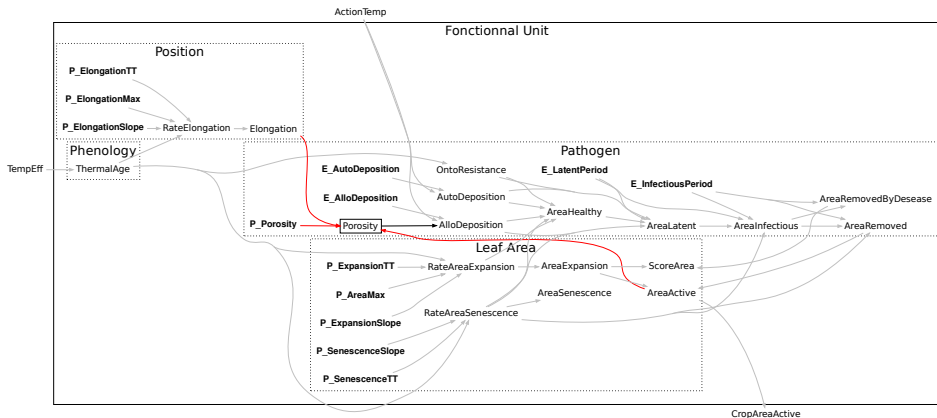
Formal model : structure of functional unit

interaction between crop and pathogen growth



Formal model : structure of functional unit

leaf area density (LAD) affects pathogen dispersion



Software model

A link between model levels of abstraction

conceptual

healthy = expansion – local infection – remote infection – senescence

formal

$$\Delta H = \Delta E - \left(\tau_u \cdot I \cdot \frac{H}{E} \cdot \rho_d \cdot \theta_d \right) - \left(\delta i_d \cdot \frac{H}{E} \cdot \rho_d \cdot \theta_d \right) - \left(\Delta S \cdot \frac{H}{A} \right)$$

code

```
// Healthy area dynamic
AreaHealthy = fmax(
  AreaHealthy(-1)
  -InitQuantity()
  +RateAreaExpansion()
  -(RateAutoDeposition() * AreaInfectious(-1) * AreaHealthy(-1)/AreaExpansion(-1)) * Receptivity()
  -(InDeposition() * AreaHealthy(-1)/AreaExpansion(-1)) * Receptivity()
  -(RateAreaSenescence() * AreaHealthy(-1)/AreaActive(-1)),0);
```

Design constraints

Scope statement governing the design choices

- aims
 - ▶ focus on the system functioning : comprehensive approach
 - ▶ focus on environmental and management uncertainties : analytic approach
- knowledge
 - ▶ differences in the modelling states among pathosystems : "lowest common denominator"
- complexity
 - ▶ link physiology and structure at the organ level : physical pathogen dispersion
 - ▶ bypass explicit plant architecture : graph-based dispersion
- evaluation
 - ▶ define specific system behaviours to be reproduced by the model

Design constraints

Chosen design options, test-driven development

- aims
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 - ▶ **define specific system behaviours to be reproduced by the model**

A model to promote discussion on epidemic management at the canopy scale

A simple model with complex behaviors regarding plant diseases

- ▶ minimizing computing time (low complexity, efficient software implementation)
- ▶ suitable for large numerical exploration (e.g. sensitivity analysis)

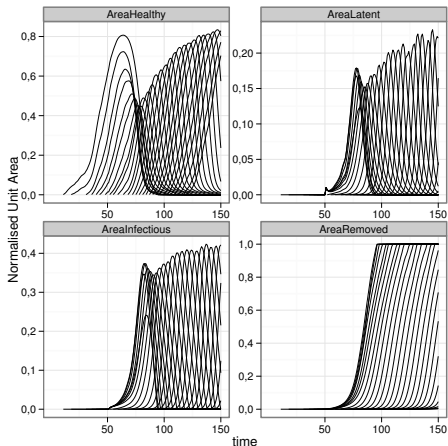
Identification of shared concepts among case studies

- ▶ comparative epidemiology : common view, distinct systems

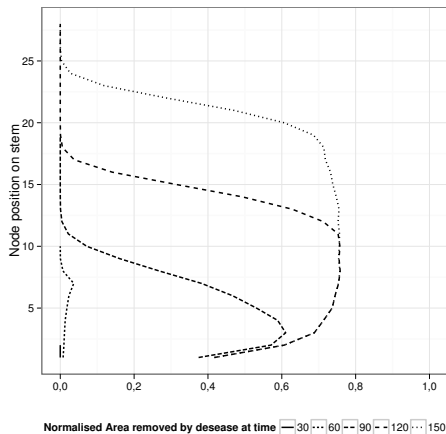
Qualitative model evaluation : vertical epidemics

Impact of crop development on disease profiles

Individual growth



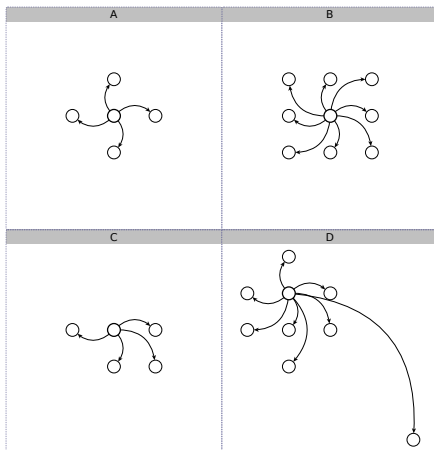
Disease profiles



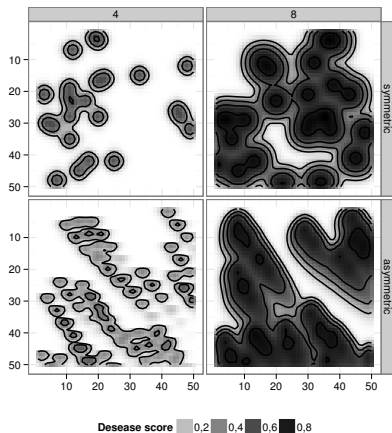
Qualitative model evaluation : spatial epidemics

Impact of local neighborhood parametrization on disease severity

Local neighborhood



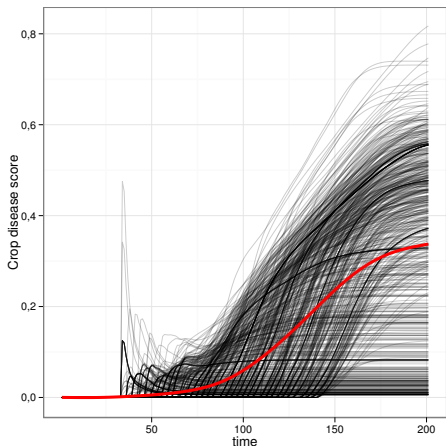
Resulting disease map



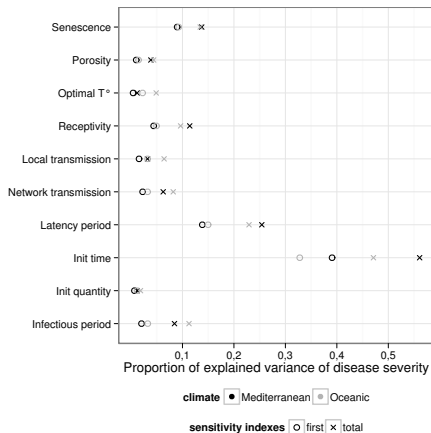
Quantitative model evaluation : sensitivity analysis

Impact of canopy characteristics (18 - 24%) on the dynamic of disease severity

Epidemic variability (n=1k)



Sensitivity analysis (n=10k)



Ideotype design : towards a model-assisted breeding ?

Crop ideotype \approx plant architecture + crop management | environment

Connection with quantitative genetics (session 5)

- ▶ Plant architectural traits
- ▶ Physiological traits

Connection with integrated pest management (session 6)

- ▶ Representation of canopy architecture is too crude.
 - mainly temporal effects (avoidance, senescence, latency)
 - limited management actions (sowing, spatial organisation)

⇒ Combine *in vivo* and *in silico* experiments

⇒ Participative modelling

A challenging integration of different scales

Sub-plot scale : process representation and simplification

- ▶ Light and spore interception : 3D architecture (e.g. L-Systems)
- ▶ Dispersion : wind and splash dispersal physical models
- ▶ Microclimate : energy balance models

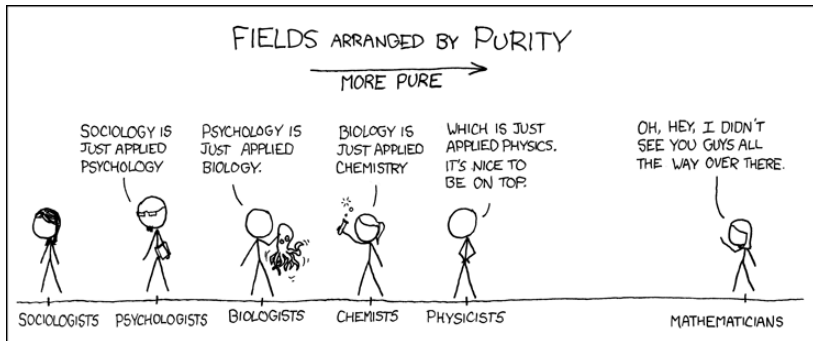
Landscape scale : robustness assessment

- ▶ Spatialisation (GIS, decision modelling)
- ▶ Aggregation and optimisation methods

⇒ Modelling platform should facilitate model coupling between scales or formalisms

Integration in real work condition

Why modellers should not (only) be mathematicians



source : xkcd.com, a web comic about romance, sarcasm, math and language

Thank you all !

- ▶ Amélioration des Plantes et Biotechnologies Végétales (APBV) - INRA GAP, Ploudaniel
- ▶ Agrosystèmes tropicaux (ASTRO) - INRA EA, Petit-Bourg
- ▶ Biologie des Organismes et des Populations Appliquée à la Protection des Plantes (BIO3P) - INRA SPE, Rennes
- ▶ Biométrie et Intelligence Artificielle (BIA) - INRA MIA, Toulouse
- ▶ Institut Mathématique de Bordeaux (IMB), Univ. Bordeaux 1
- ▶ Santé Végétale (SV) - INRA SPE, Bordeaux
- ▶ Agrosystèmes et Agricultures, Gestion de ressources, Innovations et Ruralités (AGIR) - INRA EA, Toulouse