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A plant pathogen multiscale problem

dedicated to Chris Cosner

M. Langlais

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Multi-scale problem



Courtesy from Pixels et Grains d'argent, and from INRA

Host-parasite system

- *Vitis vinifera*, vine stock & plot,
 - ***Spatial structures*** : leaves, plants, plots, ***no dispersal***,
 - ***Seasonal variations*** : growing season, climate (temperature),
 - ***Highly anthropized system*** : wide diversity of cropping systems driving *host dynamics* (secondary shoots / leaves appearance),
 - No specific resistance of the host.

Host-parasite system

- *Vitis vinifera*, vine stock & plot,
 - ***Spatial structures*** : leaves, plant, plot scales, **no dispersal**,
 - ***Seasonal variations*** : growing season, climate (temperature),
 - ***Highly anthropized system*** : wide diversity of cropping systems driving *host dynamics* (secondary shoots / leaves appearance),
 - No specific resistance of the host.
- *Erysiphe necator*, powdery mildew,
 - ***Airborne dispersal of conidia (spores)***,
 - ***Sensitive to the climate*** : wind speed & direction (?),
 - ***Sensitive to the quality of its host*** : e.g., age of leaves, ...
 - Hard to detect and quantify at the vineyard level,
 - No reliable prevention tool : routine chemical sprays,
 - ❖ 30% world market in fungicides dedicated to vine, (F: 60 M€/year),
 - Over-wintering in the bark of vine stocks & within dormant buds,

Outline : multi-scale problem

- (1) **Architectural discrete 3D** plant - pathogen model,
discrete w.r.t. time & space, *over a single growing season*,

$$X(n+1) = F(X(n), n, \pi(n), \theta(n)), \quad n \geq 0.$$

- (2) **Continuous aggregated** plant - pathogen model,
ODEs SEIRT-like, *for a perennial plant*,

$$d_t X(t) = F^{year}(X(t), t, \pi(t)), \quad t \geq 0.$$

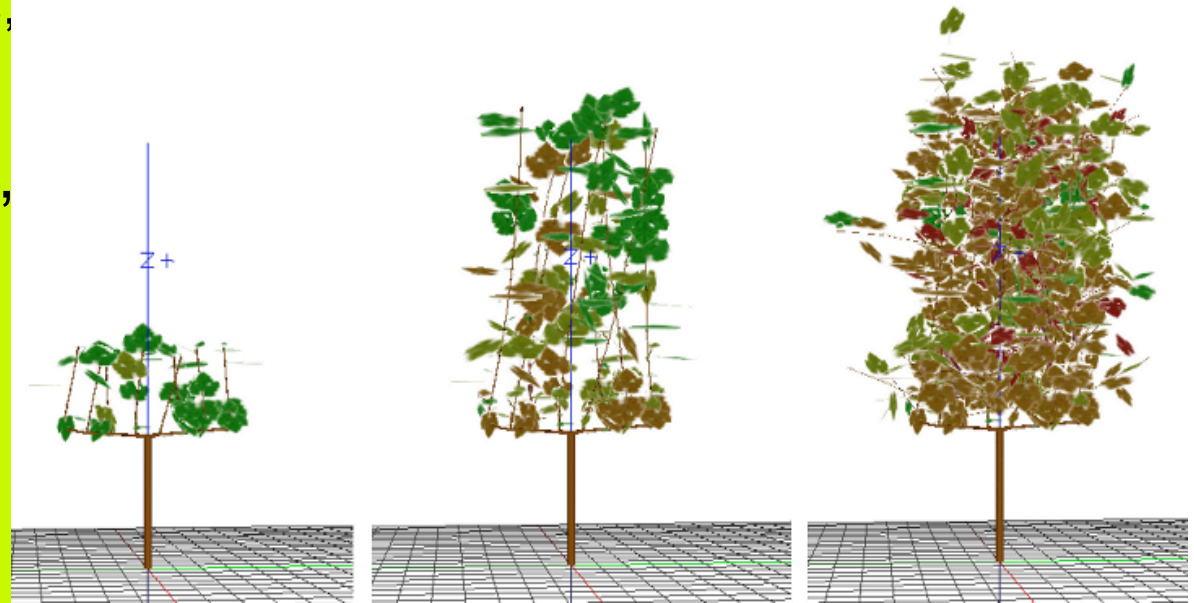
- (3) **Spatially structured** plot - pathogen model,
RD - SEIR(T) like,

$$\partial_t U = \nabla \cdot (D(x, N) \nabla U) + V \cdot \nabla U + G^{year}(U, x, \pi(t)), \quad t \geq 0, x \in D.$$

Discrete 3D

plant - pathogen scale

- ***Host dynamic***: complex 3D architectural model,
 - Appearance, growth, size, age, 3D localization and surface area of primary,
 - Leaves, internodes,
 - Shoots, grapes,
 - ***Climatic scenario***,
 - Daily basis,
 - Secondary organs,
 - ***Anthropization***,
 - Shoot topping,

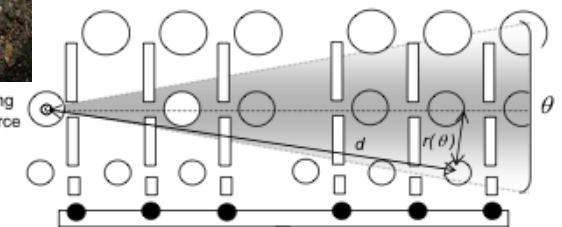


Discrete 3D plant - pathogen scale

- ***Coupled host-pathogen dynamic*** : complex 3D architectural model,
- Primary infection :
 - Calendar date,
 - 3D localization on host,
- Lesion growth,
 - Latency period,
 - Spore production period,
 - Spore release,
- *Within* vine stock spore dispersal,
 - Speed of wind (daily),
- Secondary infections,



sporulating
leaf = source

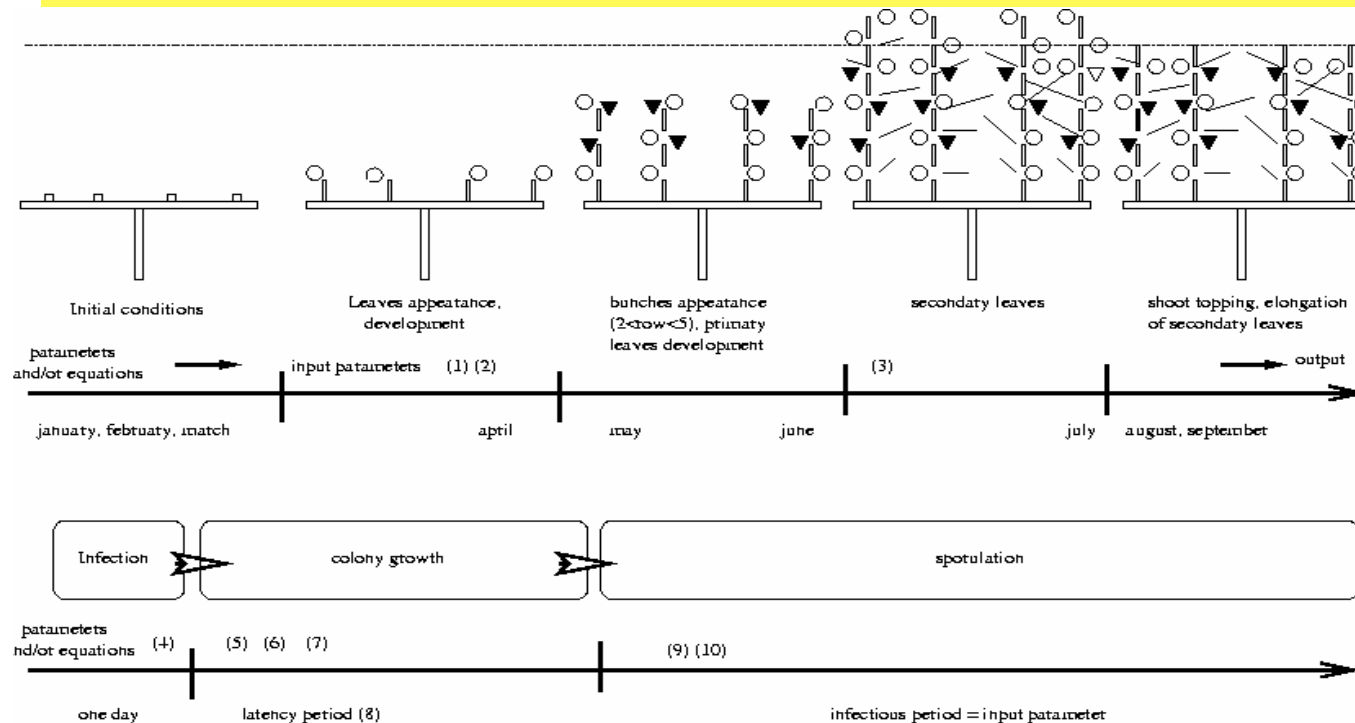


Field data,
INRA Cabernet Sauvignon,
Digitalization of vine stocks,

Discrete 3D plant - pathogen scale

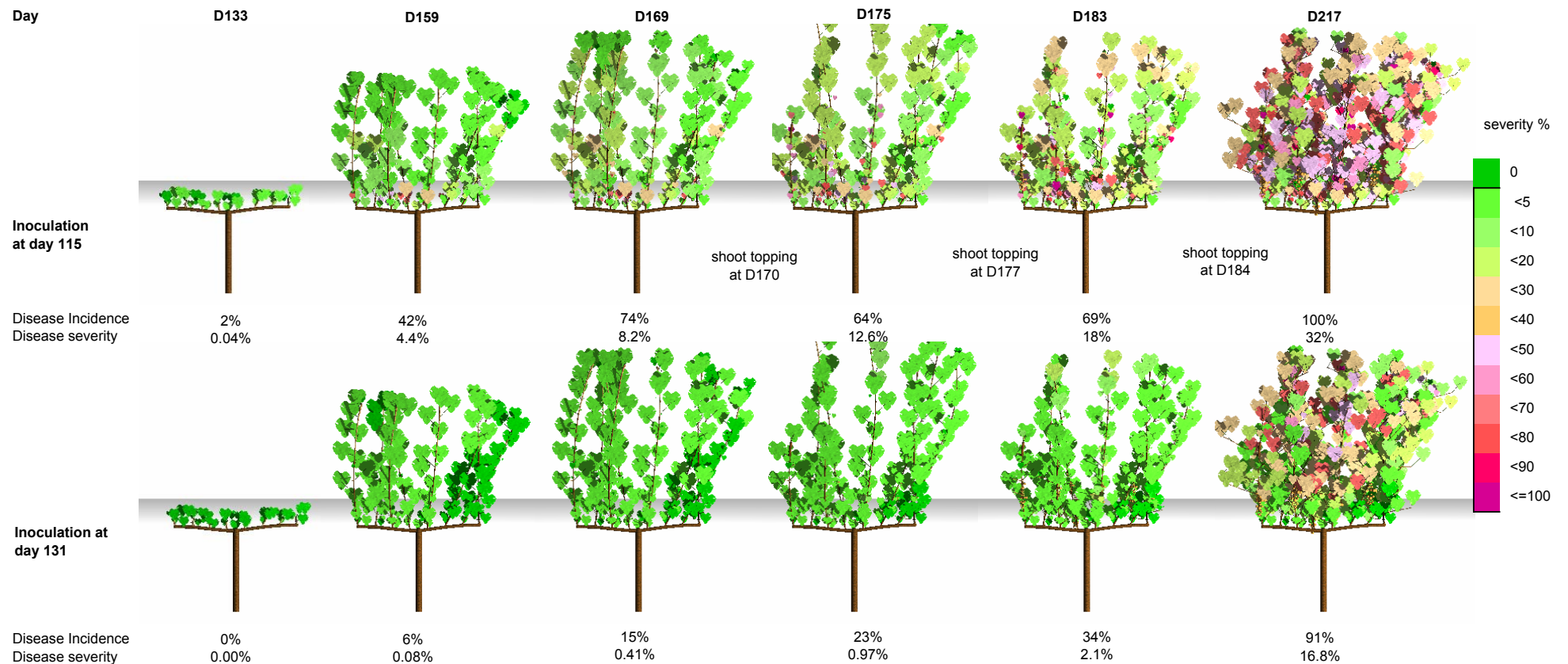
- Plant / pathogen dynamics for a growing season, complex 3D model, more than 30 parameters ...

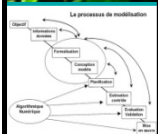
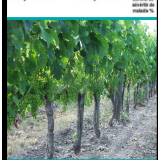
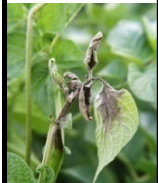
$$X(n+1) = F(X(n), n, \pi(n), \theta(n)), \quad n \geq 0.$$



Discrete 3D plant - pathogen scale

- Scenario 1998, inoculation day : 115 vs 131,



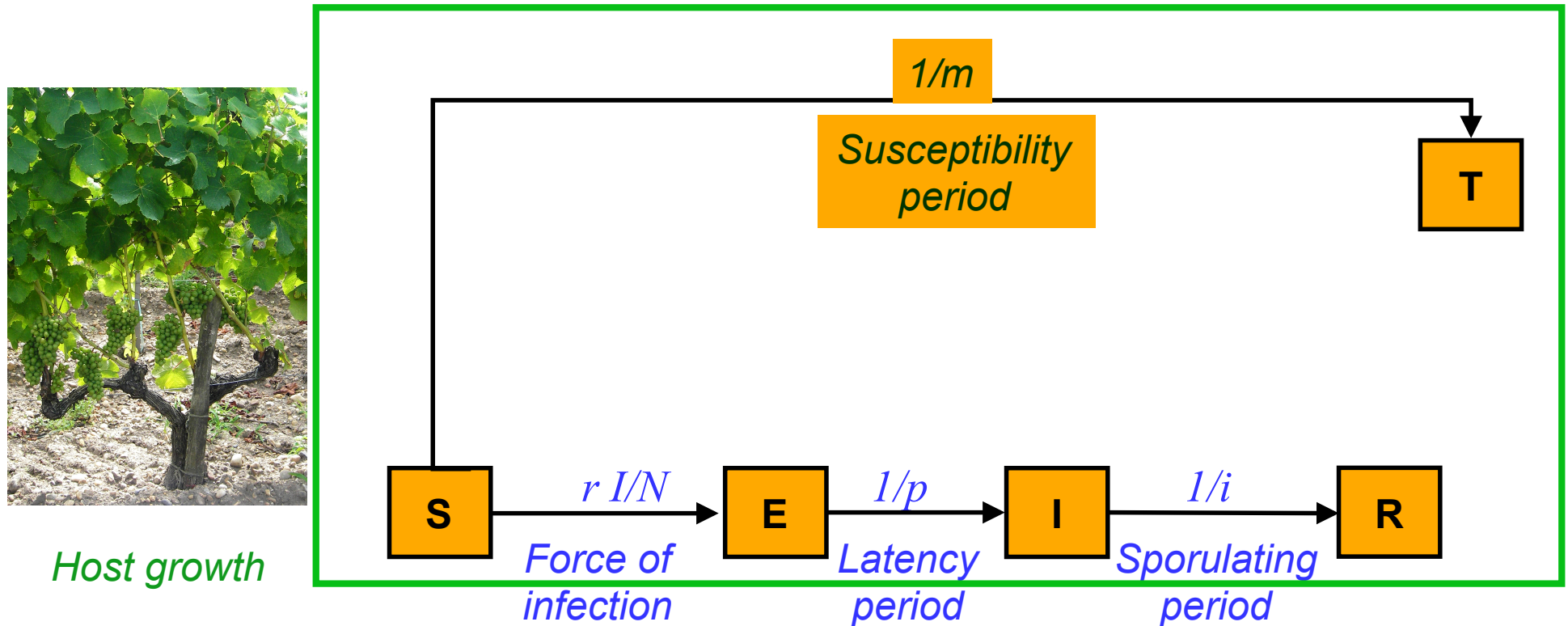


Discrete 3D plant - pathogen scale

- PCA PLS analysis, Climatic scenarios and crop management do not impact the same variables,
 - Year (climate) is better correlated to early disease,
 - Crop management is more correlated to later disease,
- Sensitivity and elasticity analysis,
 - Age of leaves : susceptibility, ontogenic resistance,
 - Vigour of the host,
 - Sporulation,
 - Inoculation date,
 - Temperature, dominant wind, ...
- Qualitative analysis ?

Continuous aggregated plant - pathogen “minimal” model

- SEIRT compartmental model w.r.t. leaf surface area,



– incidence : $r \times S \times I/N$, Vanderplank, etc.

Continuous aggregated plant - pathogen “minimal” model

- SEIRT model : **S**usceptible, **E**xposed, **I**nfectious, **R**etired, **T** (ontogenic resistance),

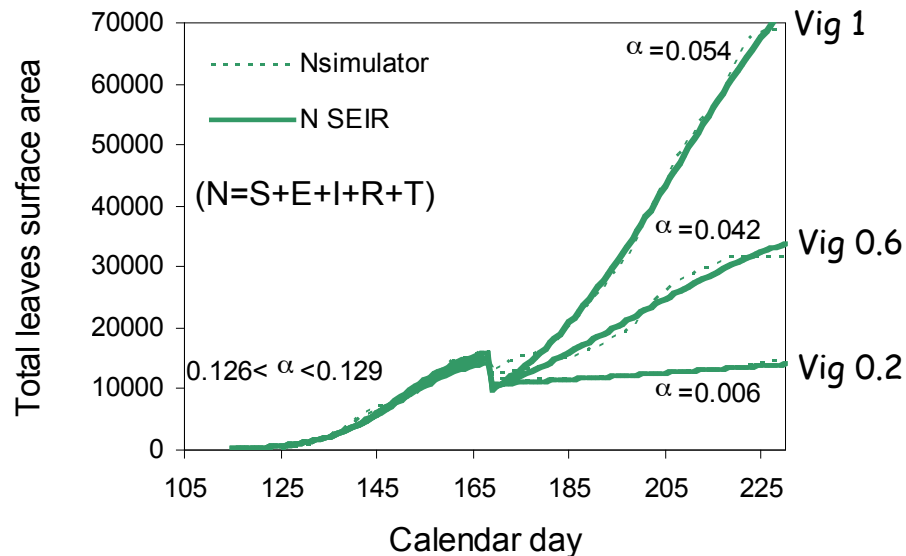
$$\left\{ \begin{array}{l} d_t S = \Lambda - r I \frac{S}{N} - \frac{1}{m} S \\ d_t E = +r I \frac{S}{N} - \frac{1}{p} E \\ d_t I = +\frac{1}{p} E - \frac{1}{i} I \\ d_t R = +\frac{1}{i} I \\ d_t T = +\frac{1}{m} S \end{array} \right. \quad \Rightarrow d_t N = \Lambda = \alpha \left(1 - \frac{N}{K} \right) N.$$

– Find a set of parameters (r, α, K) yielding consistent dynamics w.r.t. the discrete model ?

Continuous aggregated plant - pathogen “minimal” model

- « Aggregated » ODEs model fitted with discrete model,
 - Good fitting **to aggregate time-dependent processes** provided, \neq sets of parameters (r, α, K) at shoot topping,
 - LSM : total leaf / diseased surface area,
 - Problem at shoot topping : status of remaining leaves ?

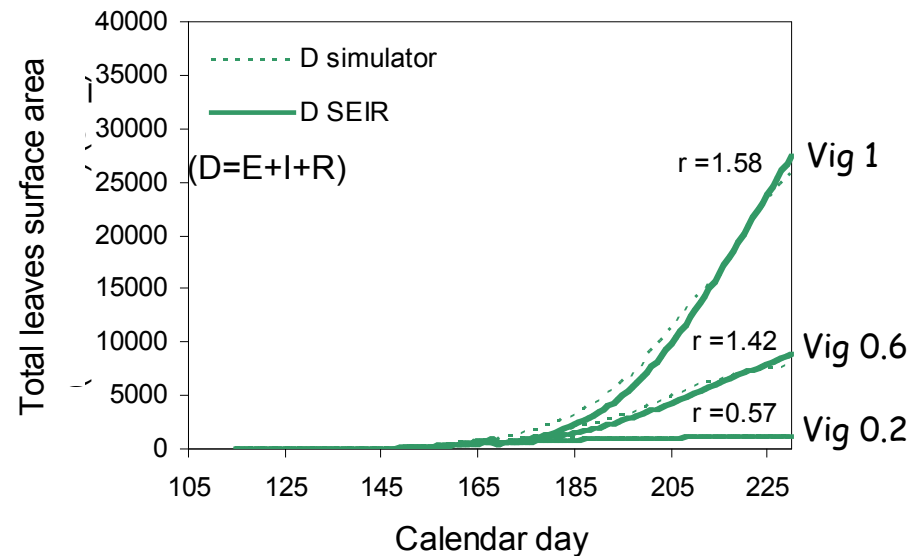
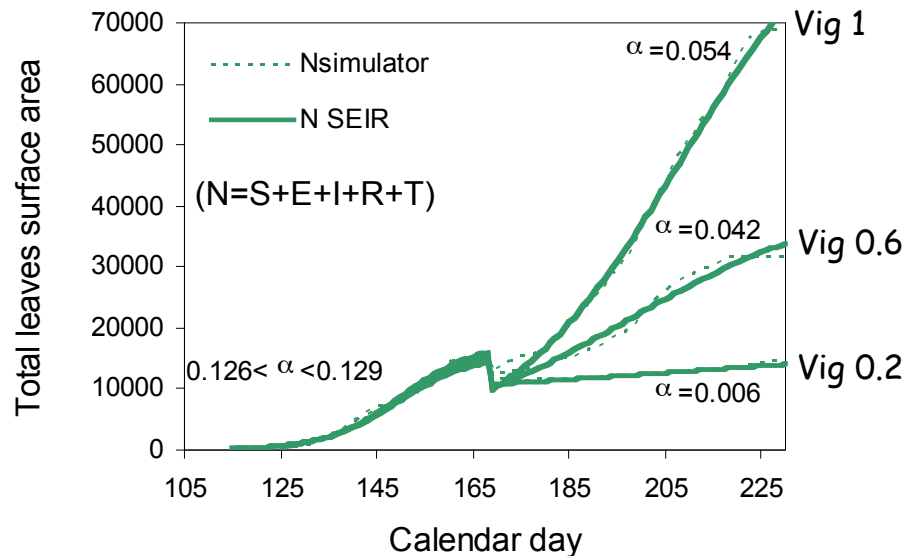
1998



Continuous aggregated plant - pathogen “minimal” model

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Continuous aggregated **plant - pathogen “minimal” model**

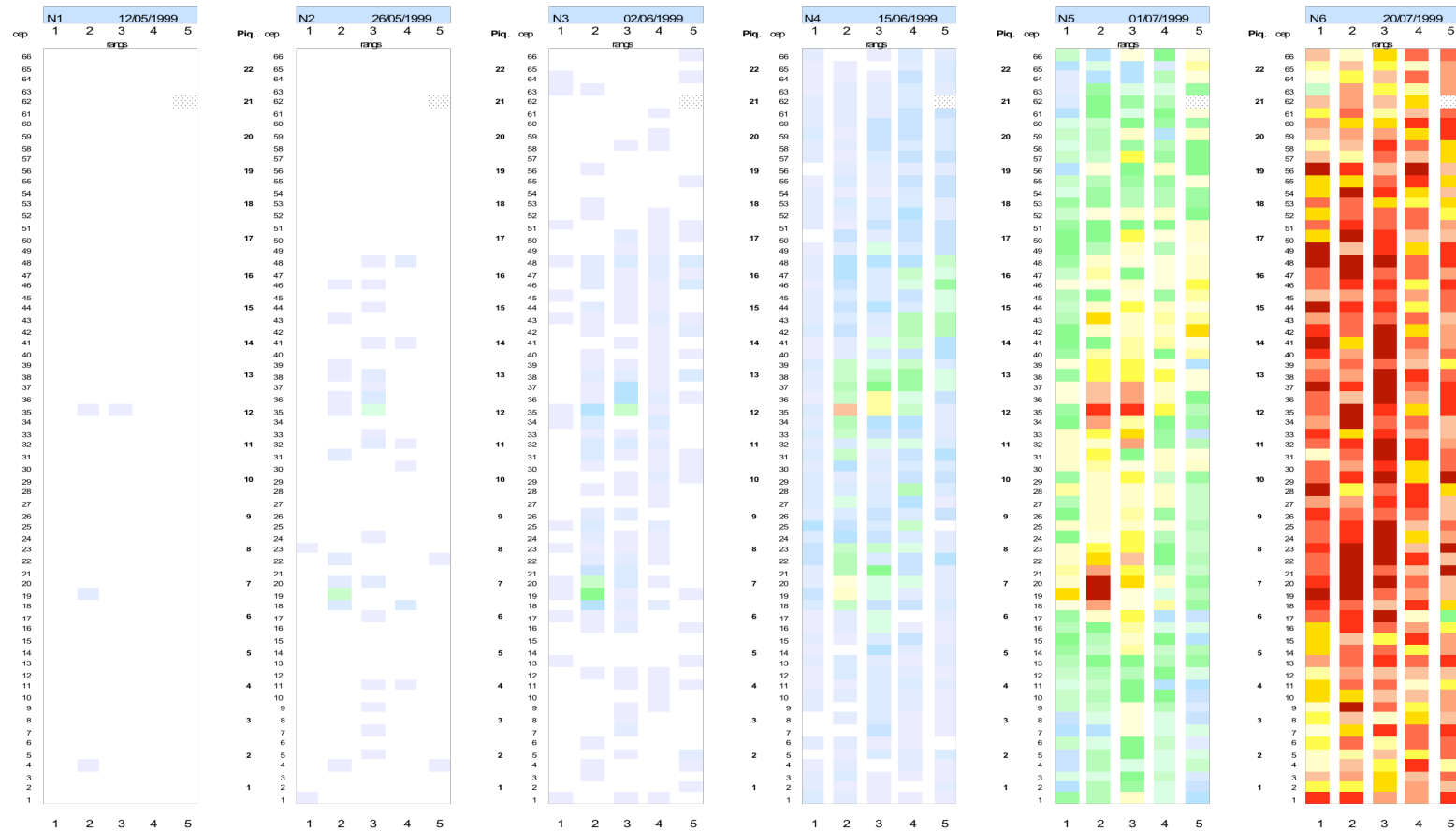
- Qualitative analysis wrt standard epidemiology,
 - Computing the reproductive number R_0 , $R_0 = r \times i$,
 - Computing the effective reproductive number $R_{\text{effective}}$,
 - ❖ $R_{\text{effective}} = r i S(t)/N(t)$,
 - ❖ $t \rightarrow R_{\text{effective}}(t)$ « decreasing outside shoot-topping » ?

Continuous aggregated **plant - pathogen “minimal” model**

- Qualitative analysis wrt standard epidemiology,
 - Computing the reproductive number R_0 , $R_0 = r^*i$,
 - Computing the effective reproductive number $R_{\text{effective}}$,
 - ❖ $R_{\text{effective}} = r i S(t)/N(t)$,
 - ❖ $t \rightarrow R_{\text{effective}}(t) \ll \text{decreasing outside shoot-topping} \gg ?$
 - Computing the asymptotic state in closed form ($+\infty = \text{winter}$) ?
 - ❖ without ontogenic resistance ($1/m = 0$ or $T = 0$),
$$S = S^{\text{infinity}}, E = I = 0, R = R^{\text{infinity}}.$$
 - ❖ with ontogenic resistance,
$$S = 0, E = I = 0, R = R^{\text{infinity}}, T = T^{\text{infinity}}.$$

Field data (1999), INRA Bordeaux

- Experimental plot with 5 rows & 66 vine stocks per row,



Spatially structured **plot - pathogen model**

Pathogen dispersal :
conceptual analysis.

Long and short distance dispersal
type 1 : short alone,
type 2 : both (coalescence),
type 3 : (scaling effect)
reduced short / large long.

Shigesada, N., Kawasaki. K.

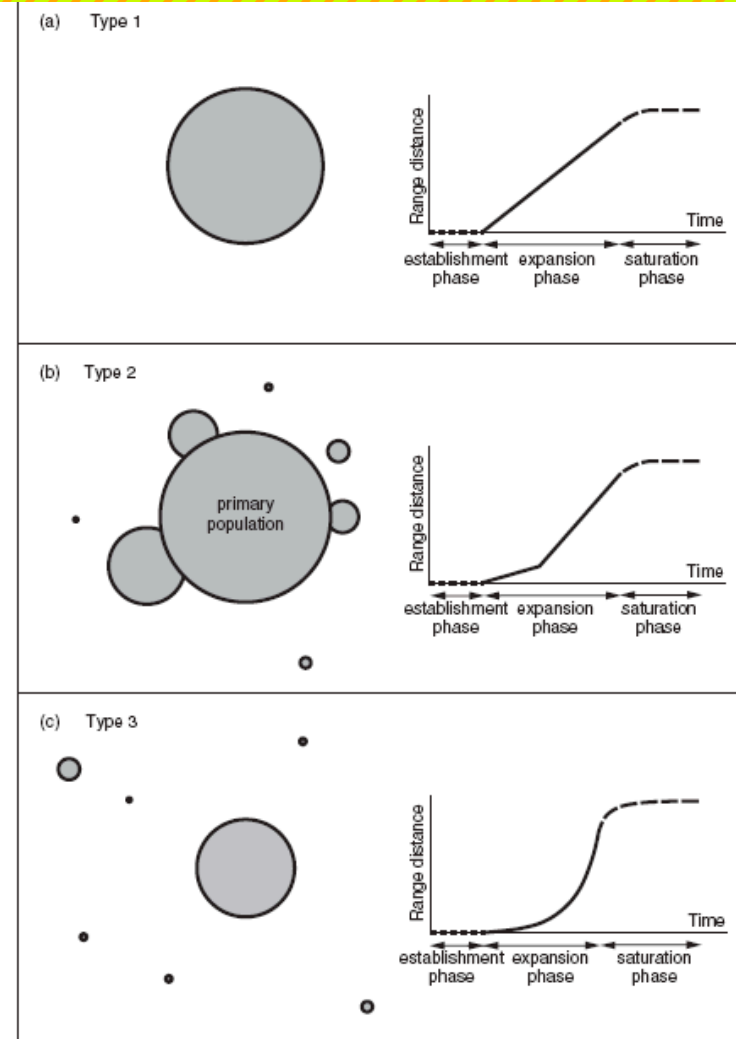
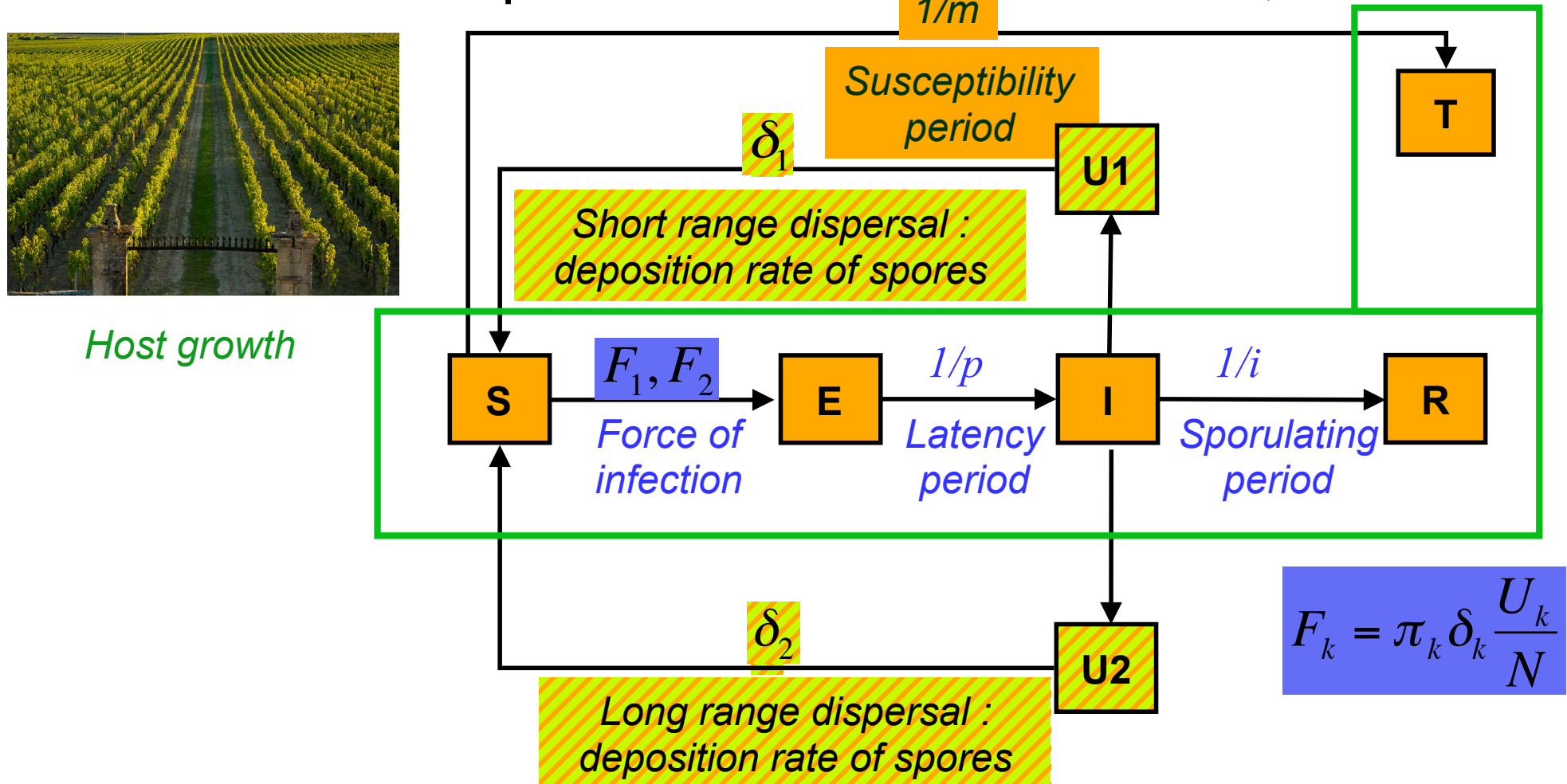


Figure 17.1 Three types of range expansions: (a) type 1, (b) type 2, and (c) type 3. For each type, the spatial pattern and range-versus-time curve are shown on the left and right, respectively. See text for detail. (Adapted from Shigesada & Kawasaki 1997, with permission of Oxford University Press.)

Spatially structured plot - pathogen model

- SEIRT compartmental model over rows,



Spatially structured **plot - pathogen model**

- **Reaction-Diffusion / SEIRT model :**

(1) Reaction-Diffusion system for dispersal of spores,

$$\begin{cases} \partial_t U_1 = \nabla \cdot (d_1(x, N) \nabla U_1) - \delta_1 U_1 + \gamma f(N) I, \\ \partial_t U_2 = \nabla \cdot (d_2(x, N) \nabla U_2) - V(x, t) \cdot \nabla U_2 - \delta_2 U_2 + \gamma (1 - f(N)) I. \end{cases}$$

- ❖ U_1 & U_2 , short (plant scale) and long range dispersed spores,
 - Fick's law diffusion, $0 < d_1(x, N) \leq d_2(x, N)$,
 - Convection term $V(x, t)$, for average vs strong winds (?),
- ❖ δ_i , deposition rate of spores U_i , $i=1, 2$, $\delta_2 \leq \delta_1$,
- ❖ γ , production rate of spores per infectious unit,
- ❖ $f(N)$, % of spores dispersed at short range, Aylor (1999),

Spatially structured plot - pathogen model

- **Reaction-Diffusion / SEIRT model :**

(2) SEIRT ODEs models over rows : production of spores by I and contamination of S by U_1 and U_2

$$\left\{ \begin{array}{l} d_t S = \Lambda - (\pi_1 \delta_1 U_1 + \pi_2 \delta_2 U_2) \frac{S}{N} - \frac{1}{m} S \\ d_t E = +(\pi_1 \delta_1 U_1 + \pi_2 \delta_2 U_2) \frac{S}{N} - \frac{1}{p} E \\ d_t I = +\frac{1}{p} E - \frac{1}{i} I \\ d_t R = +\frac{1}{i} I \\ d_t T = +\frac{1}{m} S \end{array} \right.$$

$\pi_1 \gamma \approx r$, ODEs model,
 $\pi_2 \leq \pi_1$, inoculum eff.

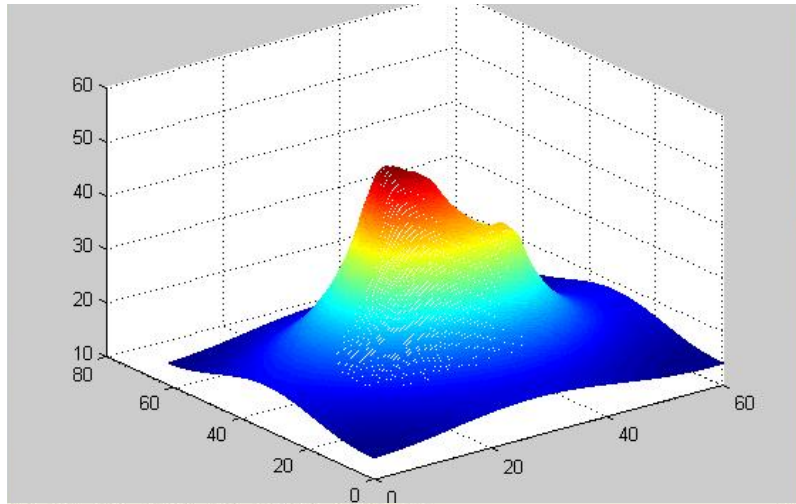
$\Rightarrow d_t N = \Lambda = \alpha \left(1 - \frac{N}{K} \right) N.$

Spatially structured **plot - pathogen model (a)**

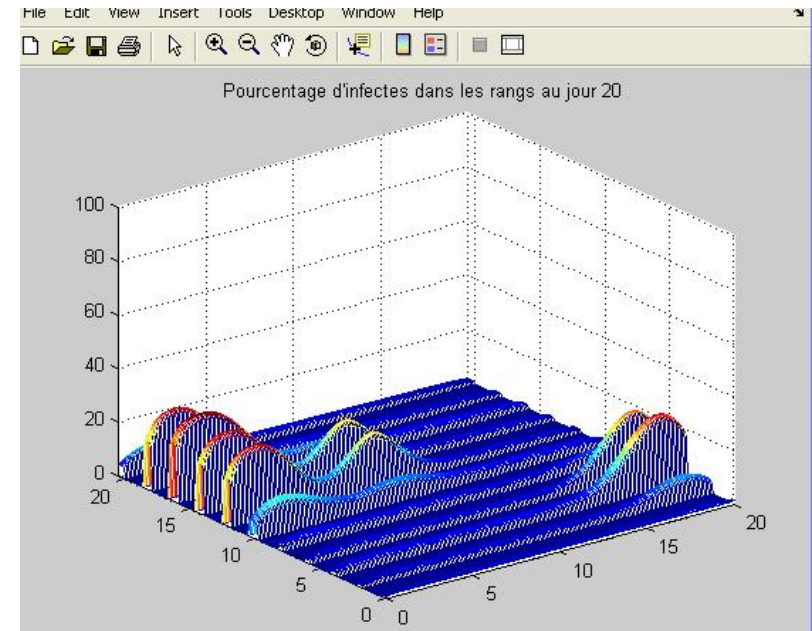
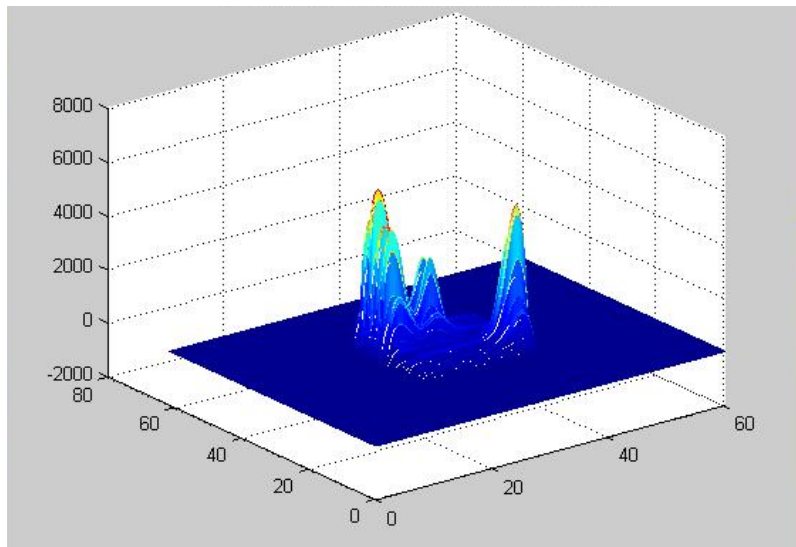
- Basic model, R-D / SEIRT :
 - R-D system,
 - ❖ constant diffusivities, no convection term,
 - ❖ kinetic terms : homogeneous / heterogeneous,
 - ❖ cf. Zawolek & Zadoks (1992), Vanderplank (1963),
 - *Boundary conditions ?*
 - *homogeneous Dirichlet conditions* set on a much larger domain than the spatial range of interest,
 - Mathematical analysis : IBVP.
 - Numerical experiments (Matlab interface).
 - vs
 - TW analysis.

Spatially structured **plot - pathogen model (a)**

Spores : long range dispersal



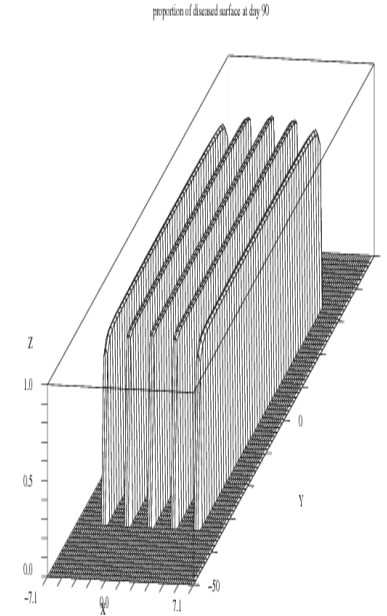
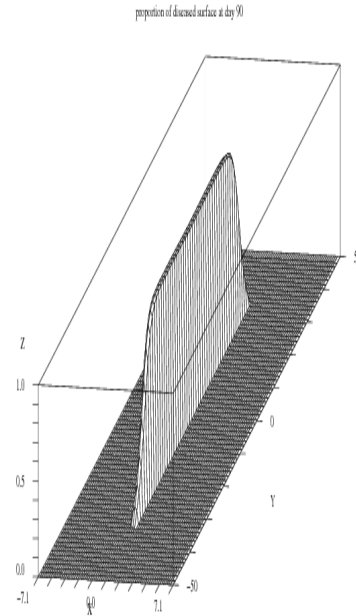
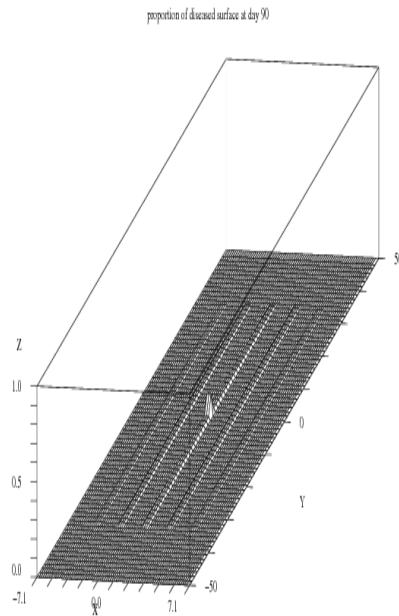
Spores : short range dispersal



Spatial spread of infection

Spatially structured plot - pathogen model (a)

- Proportion of disease lesion / dispersal, Day 90,
 - Long range dispersal only : weak uniform contamination,
 - Short range dispersal only : central row is contaminated,
 - Both : contamination spreads all over the five rows,



Spatially structured **plot - pathogen model (b)**

- ***Qualitative analysis :***

- Dimensionless model, R-D / SEIR (*no T*),

$$\frac{\partial U_1}{\partial t} = \Delta U_1 - \eta U_1 + \eta F I$$

$$\frac{\partial U_2}{\partial t} = d\Delta U_2 - \eta U_2 + \eta(1 - F)I$$

$$\frac{\partial S}{\partial t} = -R_0 (U_1 + U_2) S$$

$$\frac{\partial E}{\partial t} = +R_0 (U_1 + U_2) S - \frac{1}{\tau} E$$

$$\frac{\partial I}{\partial t} = \frac{1}{\tau} E - I$$

Spatially structured **plot - pathogen model (b)**

- ***Qualitative analysis (no T) :***
 - Wave fronts, $\varphi(x-c.t)$, 1D, pulses / infection, ***linking***,
 - ***Pre-infection state***,
 - ❖ $U_1 = U_2 = 0, S = K, E = I = R = 0.$
 - ***Post-infection state***,
 - ❖ $U_1 = U_2 = 0, S = K^{infinity}, E = I = 0, R = R^{infinity}.$
 - When $R_0 > 1$, minimal speed TW, c^* ,
 - implicit closed form ($K^{infinity}$),
 - $K^{infinity}$ a root of, $h = 1 - e^{-R_0 h}$

Spatially structured **plot - pathogen model (b)**

- The value of c^* is given by solving ($F < 1$) the characteristic equation:

Find (λ^*, c^*) such that $0 < \lambda^* < \frac{c + \sqrt{c^2 + 4d\eta}}{2d}$ and
$$\begin{cases} \psi(\lambda^*, c^*) = 1 \\ \frac{\partial \psi}{\partial \lambda}(\lambda^*, c^*) = 0 \end{cases}$$

with

$$\psi_{SEIR}(\lambda, c) = \frac{R_0}{(1 + c\lambda)(1 + c\tau\lambda)} \left(\frac{\eta F}{\eta + c\lambda - \lambda^2} + \frac{\eta(1 - F)}{\eta + c\lambda - d\lambda^2} \right)$$

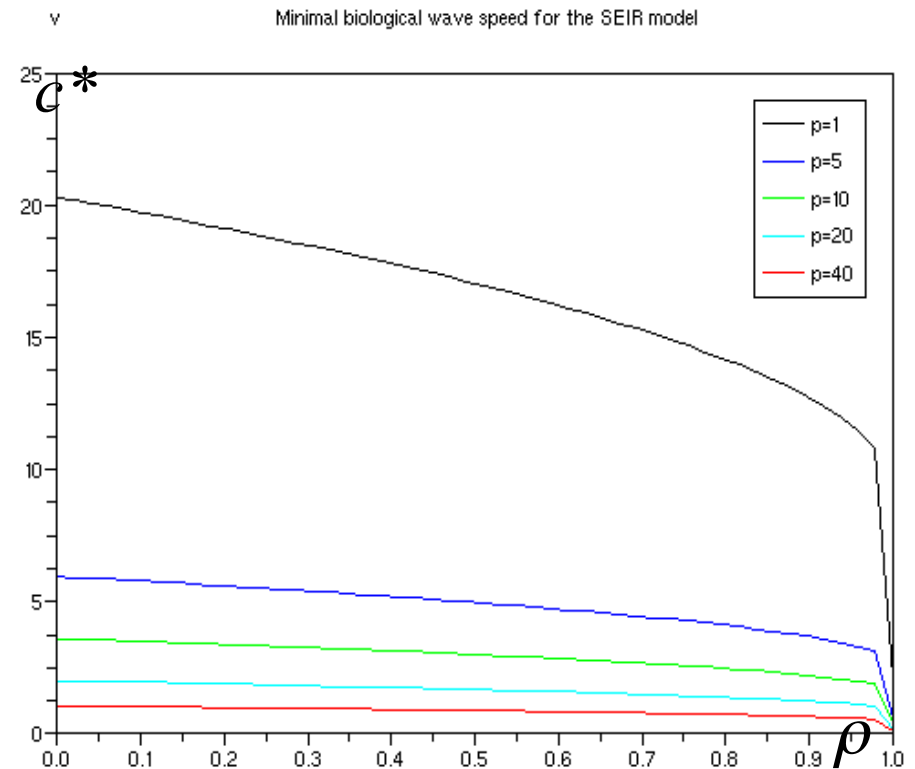
Spatially structured plot - pathogen model (b)

- *Dimensionless minimal speed, c^* ,*
 - Assuming : $\delta_2 = \delta_1$, $\pi_2 = \pi_1$,
 - $F(N)=\rho$, % of short range dispersed spores,

Decreasing w.r.t. ρ .

Discontinuity at $\rho = 1$.

cf. Shigesada & Kawasaki.



Spatially structured plot - pathogen model (b)

- *Dimensionless minimal speed, c^* ,*
 - Assuming : $\delta_2 < \delta_1$, $\pi_2 < \pi_1$,
 - $F(N)=\rho$, % of short range dispersed spores,

maximum achieved

at a suitable optimal ρ ,

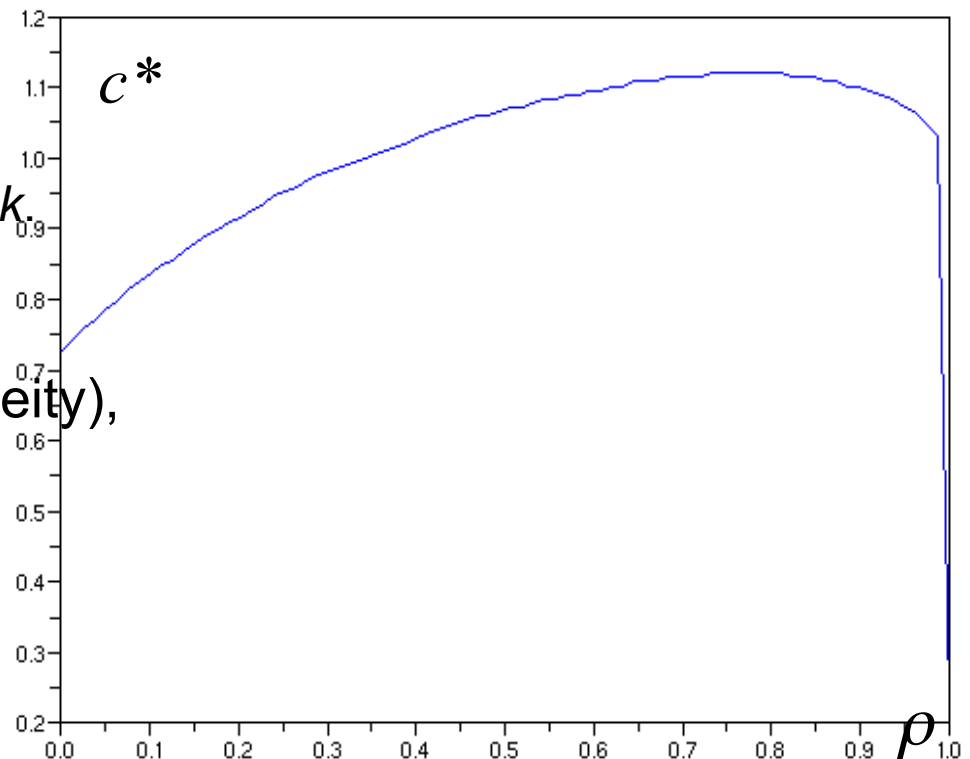
cf. Zawolek & Zadok.

actual speed (*slightly, 50%*)

over-estimated (?, heterogeneity),

Discontinuity at $\rho = 1$.

cf. Shigesada & Kawasaki.

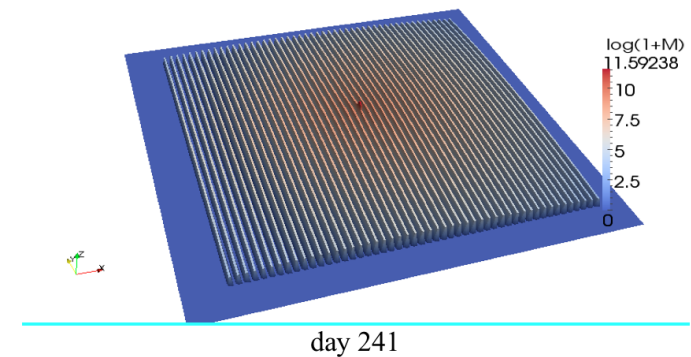
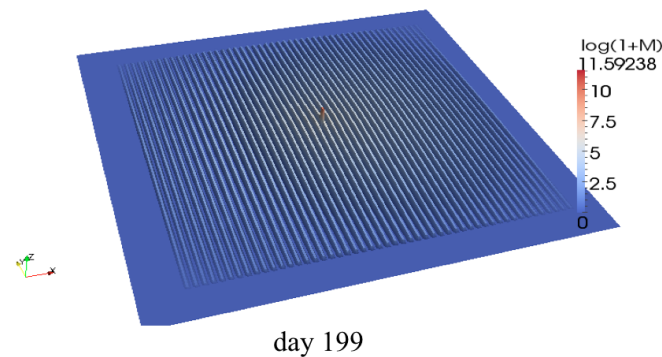
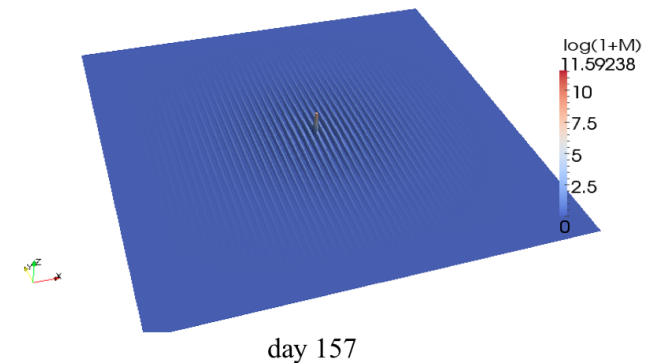
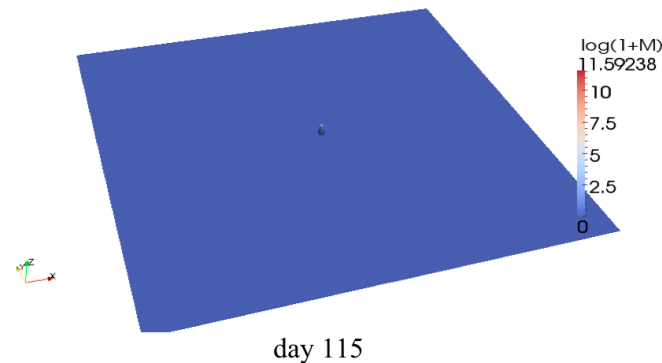


Spatially structured plot - pathogen model (b)

- 2D spatio-temporal periodical WF collaborative work A. Ducrot & H. Matano,
- ***in which direction is the speed faster ?***
numerics : along rows,
vs
data : across rows, but ...
 - *Numerical work in progress (turbulence, nonlinear diffusion, architecture & $f(N)$, ...).*

Spatially structured plot - pathogen model (b)

- 2D spatio-temporal dynamics,
- *Nonlinear short range dispersal, data fitting ?*



Further developments ...

- Collecting data and calibrating 2D and 3D models, including transport,
 - no known markers for *Erysiphe necator*, powdery mildew, compared to pollen dispersal,
 - Secondary infections from alien sources ...
- Control problems using fungicides,
 - when ?
 - coupling with shoot topping driving host dynamics ?
 - reducing or delaying the pathogen speed of propagation if coupled to ontogenic resistance ?

Spatially structured plot - pathogen model (c)

- Spatially periodic spatial plot,
 - reference cell, Y , “a single plant” (short range scale),
 - small parameter, $\varepsilon > 0$,
 - spatial domain, Ω ,
a set of cells of size $\varepsilon * Y$,
 - available vegetal tissue density, function χ , period Y ,
 $\chi(x, t, x/\varepsilon)$.
- Simpler to handle as $\varepsilon \rightarrow 0$?
 - spatially periodic: ill-conditioned



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Specific fundings

- INRA SPE, post-doc 2002-04,
- Région Aquitaine, doc 2005-2008,
- ARC INRIA, 2009-10, & INRA 2009,
 - Apple scab,
 - Montpellier CIRAD-INRA-INRIA & INRA Clermont et Gautheron,
 - post-doc 01/08 2010, M1 2009 et M2R 2010,
- ANR SYSTERRA ARCHIDEMIO, 2009-12,
 - 3 more patho-systems, Bretagne & Guadeloupe,
 - M1 2009 et M2R 2010, 2011.

