

Microbial Ecology and Bioprocess Control : Opportunities and Challenges

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Menu

- Microbial ecology : some basic concepts
- Coexistence or competition
- Recent developments
 - Density dependent growth and coexistence
 - Coexistence in a reactor cascade
 - «Practical» coexistence
- Issues and challenges

Microbial ecology : some basic concepts

- A definition :
Study of the interactions that determine
the abundance and distribution of
organisms
- A keyword : biodiversity
- A basic concept : the competitive
exclusion principle

Competitive exclusion principle

Consider a CSTR (« chemostat ») with two biomasses X_1 and X_2 growing on one limiting substrate S :

$$\frac{dS}{dt} = -\frac{1}{Y_1} \mu_1(S)X_1 - \frac{1}{Y_2} \mu_2(S)X_2 + D(S_{in} - S)$$

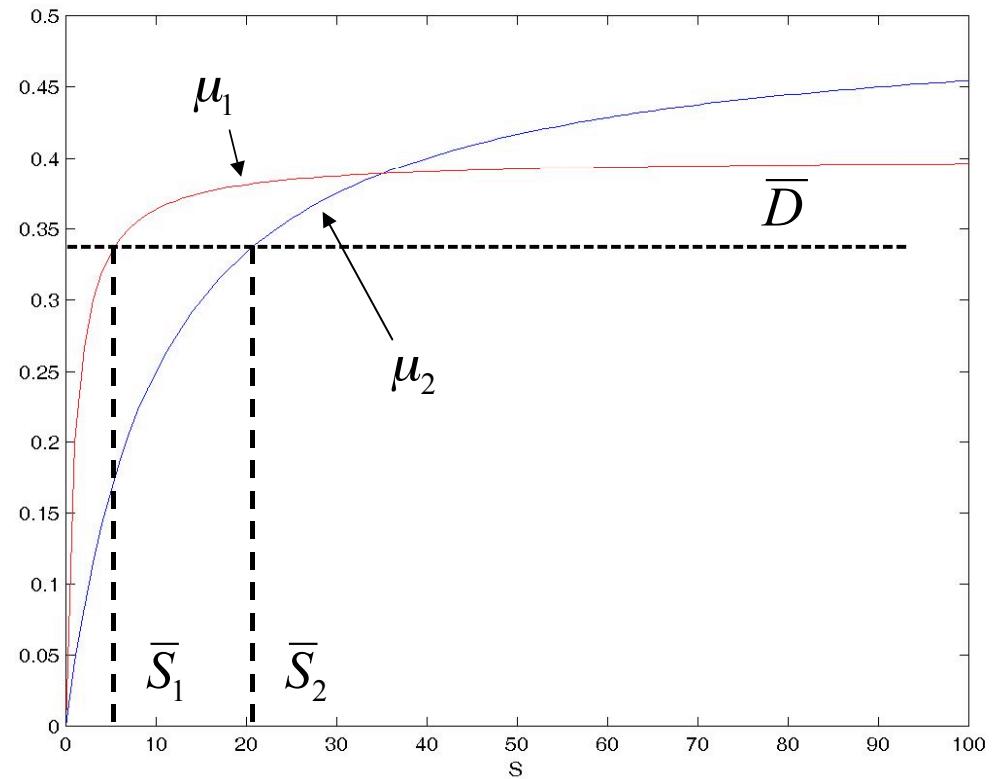
$$\frac{dX_1}{dt} = \mu_1(S)X_1 - DX_1$$

$$\frac{dX_2}{dt} = \mu_2(S)X_2 - DX_2$$

- At steady state : $\bar{\mu}_1(S) = \bar{\mu}_2(S) = \bar{D}$
(only valid for specific values of D)
- In general, only one species will «win the competition and survive» : the one whose growth curve crosses first D («best affinity» or «smallest break-even concentration»)
- Here :

$$\bar{X}_1 = Y_1(S_{in} - \bar{S}), \bar{X}_2 = 0$$

(Hardin, 1960; Butler & Wolkowicz, 1985)



(Extension to n species and other growth curves)

Competitive exclusion principle : experimental validation

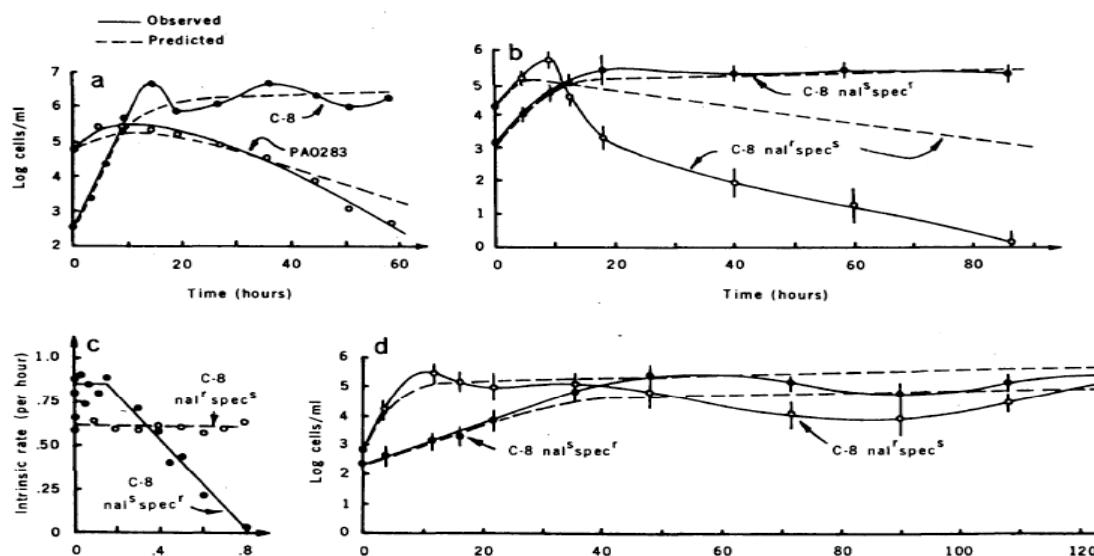
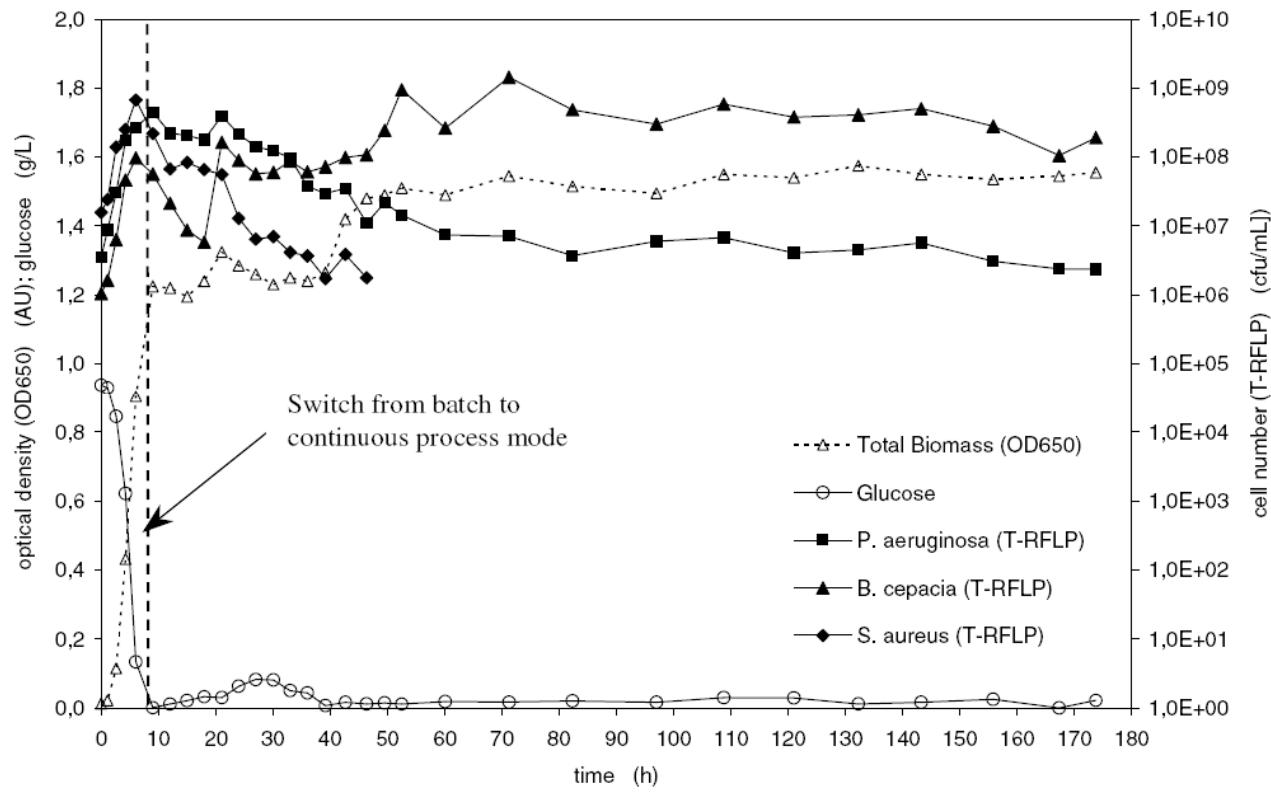


FIG. 5.2 – Validation qualitative expérimentale du comportement du modèle. Les prédictions qualitatives du modèle sont vérifiées pour : a) 2 espèces (*Escherichia coli*, souche C-8 et *Pseudomonas aeruginosa*, souche PAO283) qui diffèrent par leur constante de demi-saturation. b) 2 souches de *Escherichia coli* qui diffèrent par leur taux de croissance maximal. d) Coexistence obtenue avec 2 souches de *Escherichia coli* qui ont le même paramètre J_i . La figure c) représente l'effet de l'acide nalidixique sur le taux de croissance maximal pour les souches considérées C-8. D'après Hansen et Hubbell (1980).

The coexistence of different species is often encountered

experimental evidence :



Schmidt, J. K., B. König et U. Reichl Characterization of a three bacteria mixed culture in a chemostat: Evaluation and application of a quant

Dynamical persistence

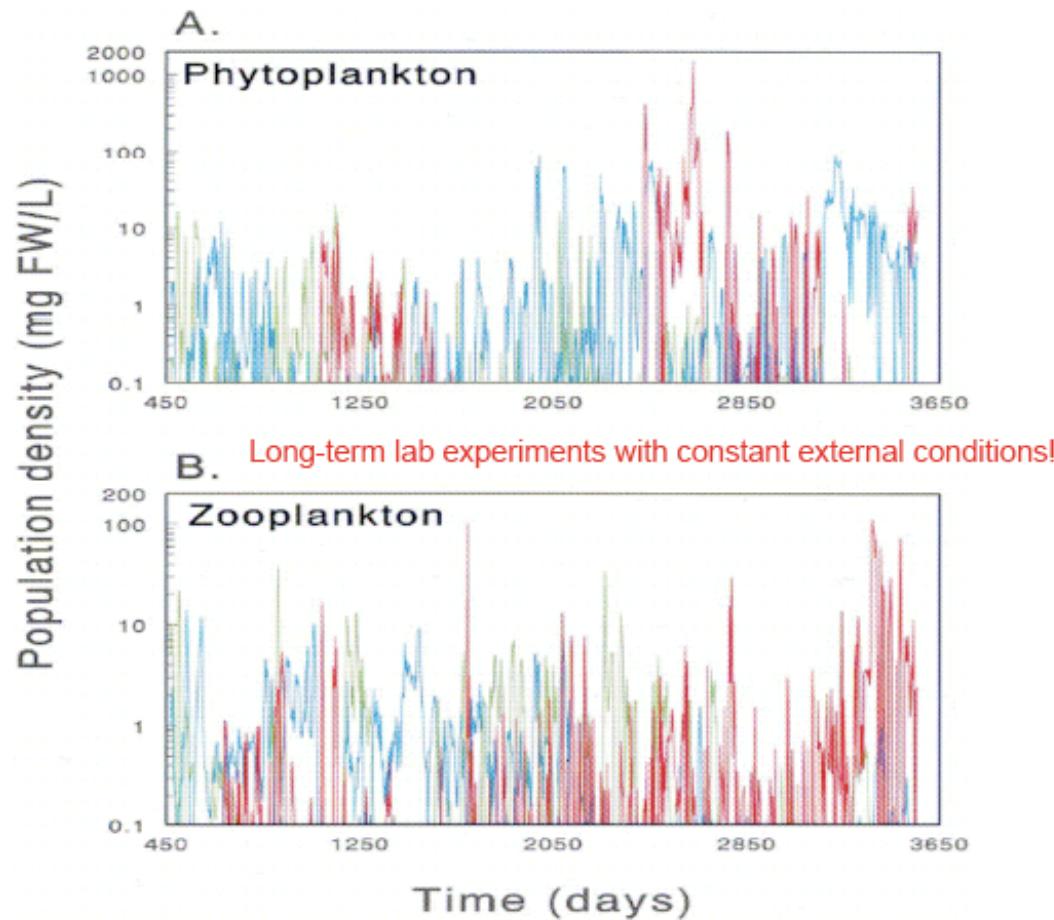


Figure 5. Non-equilibrium dynamics observed in an experimental multispecies community. The community developed in a long-term laboratory experiment under constant external conditions, and consisted of more than 20 different species. Data show the observed time course of (A) the dominant phytoplankton groups (green = green flagellates, blue = prokaryotic pico-phytoplankton, red = the diatom *Melosira*), and (B) the dominant zooplankton groups (green = the rotifer *Brachionus*, blue = the copepod *Eurytemora*, red = protozoans). Data were kindly provided by Heerkloss (unpublished), and by Heerkloss & Klinkenberg (1996), with permission from Schweizerbart'sche Verlagsbuchhandlung.

Coexistence or competition?

- Coexistence can be mathematically emphasized for periodic inlet (D , S_{in}) conditions (e.g. *Smith, 1981*)
- Filamentous backbone theory : coexistence of floc-forming and filamentous bacteria (→ activated sludge) (*Cenens et al, 2000*)
- Recent developments (MERE team) :
 - Density dependent growth and coexistence
 - Coexistence in a cascade of reactors
 - «Practical» coexistence

1. Density dependent growth and coexistence

- Starting point : the specific growth rate depends not only on the substrate concentration S but also on the biomass concentrations X_i : $\mu_i(S, X_j)$, $i, j = 1, \dots, n$
- All species X_i compete for the same substrate S
 - $\mu_i(S, X_j)$ is a decreasing function of X_i
e.g. the growth decreases with the size of the cells
(reduced accessibility to the nutrient)
 - Inter-species vs intra-species competition :
--> $\mu_i(S, X_j)$ is a decreasing function of X_j
- Coexistence if intra-species competition
> inter-species competition

Key mathematical result

- **Dynamical model**

$$\frac{dS(t)}{dt} = D(S_{in} - S(t)) - \sum_{i=1}^n \mu_i(S(t), X_i(t)) X_i(t)$$

$$\frac{dX_i(t)}{dt} = [\mu_i(S(t), X_i(t)) - D] X_i(t) \quad , \quad i = 1, \dots, n$$

$$Y_i = 1 \quad (i = 1, \dots, n) \quad (\text{without loss of generality})$$

- **Assumptions**

A1. $\mu_i(S, X_i) \geq 0$, $\mu_i(0, X_i) = 0$,

$\mu_i(S, X_i)$ is an increasing function of S

A2. For each i , the mapping $X_i \rightarrow \mu_i(S, X_i)$ is decreasing and tends to 0 at infinity

A3. For every i , there exists a $\tilde{S}_i < S_{in}$ such that $\mu_i(\tilde{S}_i, 0) = D$

A4. $\tilde{S} + \sum_{i=1}^n X_i(\tilde{S}) < S_{in}$ with $\tilde{S} = \max\{\tilde{S}_i; i = 1, \dots, n\}$

(from A2, there is a unique $X_i(S)$ such that $\mu_i(S, X_i(S)) = D$)

- **Theorem**

Under assumptions A1 to A4, there exists a unique equilibrium (S, X_1^*, \dots, X_n^*) such that for every i , one has $X_i^* > 0$

and it is globally asymptotically stable.

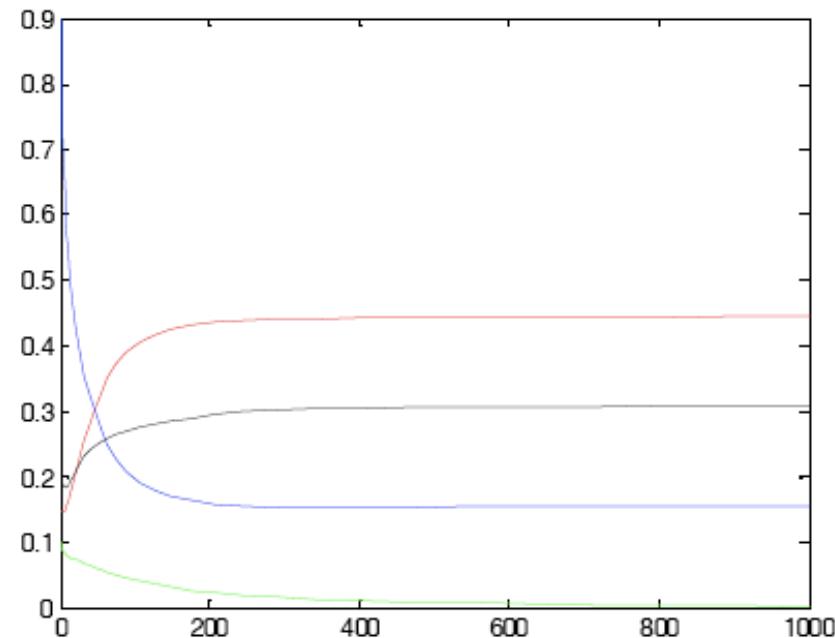
i.e. the system converges towards this equilibrium whatever the initial conditions satisfying $X_i(0) > 0$

Example

$$\left\{ \begin{array}{l} \frac{dS(t)}{dt} = D(S_{in} - S(t)) - \sum_{i=1}^n g\left(X_i(t) + \lambda \sum_{j \neq i} X_j(t)\right) \mu_i(S(t)) X_i(t) \\ \frac{dX_i(t)}{dt} = \left[g\left(X_i(t) + \lambda \sum_{j \neq i} X_j(t)\right) \mu_i(S(t)) - D \right] X_i(t) \\ \mu_i(S) = \frac{a_i S}{b_i + S} \\ g(X) = \frac{1}{1 + c \sqrt[3]{X}} \end{array} \right. , \quad i = 1, \dots, n$$

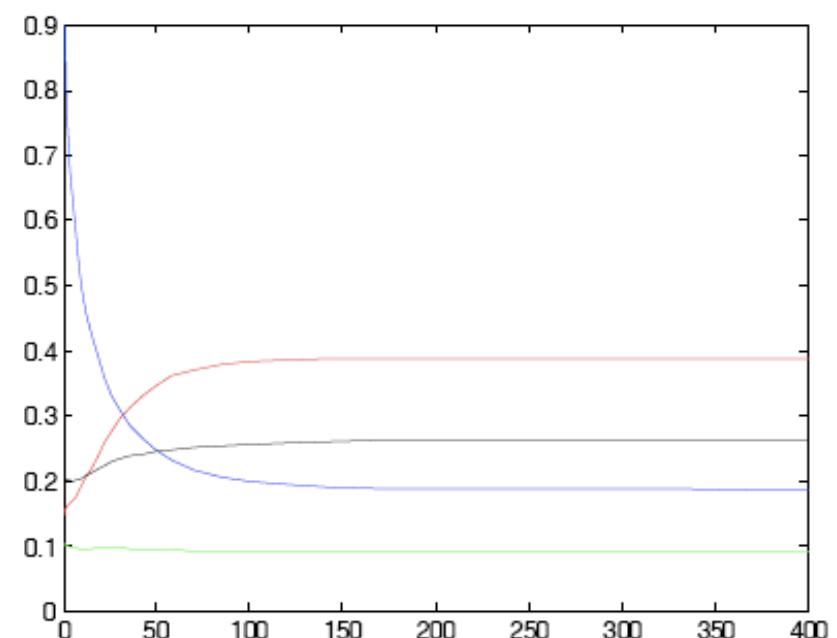
with λ as a measure of the inter-species competition
(dominates if $\lambda > 0.5$)

Exclusion



$$\lambda = 0.55$$

Coexistence



$$\lambda = 0.1$$

Various expressions of the density dependence

1) Ratio dependence, e.g. Contois model :

$$\mu = \frac{\mu_{\max} S}{K_C X + S} \text{ which can be rewritten as : } \mu = \frac{\mu_{\max} \frac{S}{X}}{K_C + \frac{S}{X}}$$

2) Flocculation : $X = \sum_{n=1}^{\infty} n u_n$ with u_n , the density of flocs of size n

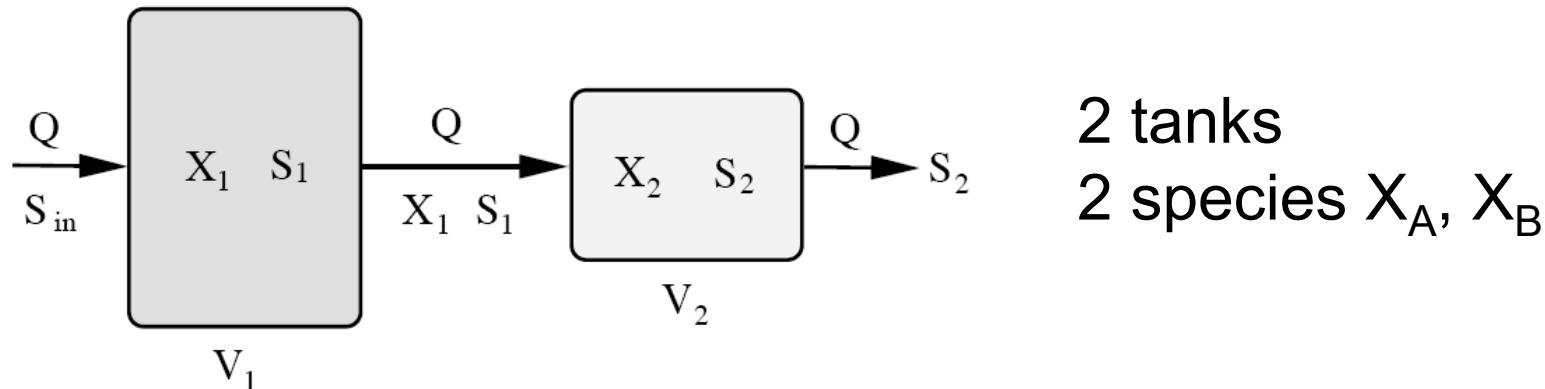
→ Mass balance equation for u_n :

$$\frac{du_n}{dt} = \left(\frac{du_n}{dt} \right)_{\text{bacterial growth}} - Du_n + \left(\frac{du_n}{dt} \right)_{\text{floc interaction}}$$

with (cell division) : $\left(\frac{du_1}{dt} \right)_{\text{bacterial growth}} = -\mu_l(S)u_1$

$$\left(\frac{du_n}{dt} \right)_{\text{bacterial growth}} = \mu_n(S)u_n - \mu_n(S)u_n \quad n \geq 2$$

2. Coexistence in a reactor cascade



- Issue : what happens when a species B starts invading a bioprocess with originally one species A?
 - can we have coexistence/exclusion?
 - is the reactor design appropriate?
- Stephanopoulos & Frederickson (1979) : coexistence in tank 2

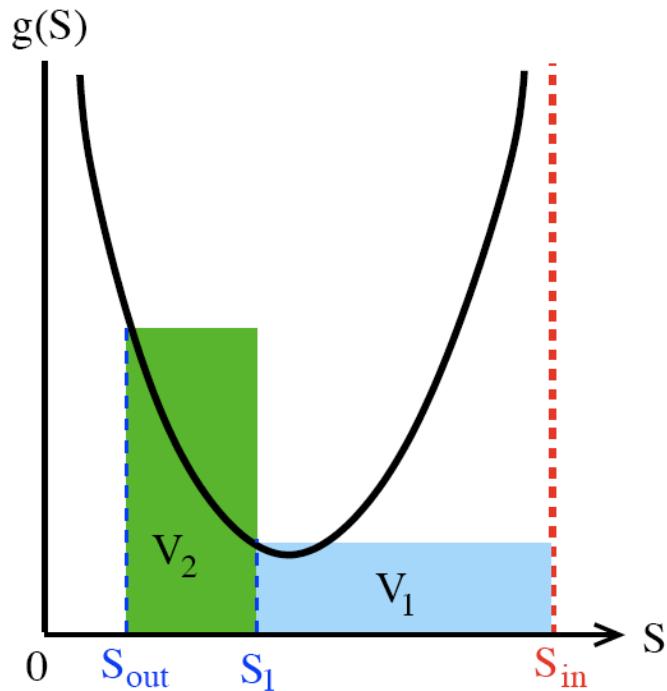
Assumptions ($i=1,2$) :
 $D_i = Q/V_i$
 $Y_i = 1$ (without loss of generality)
 $\mu_i(S)$

→ mass balance equations :

$$\left\{ \begin{array}{l} \frac{dS_1}{dt} = -\mu_A(S_1)X_{A,1} - \mu_B(S_1)X_{B,1} + D_1(S_{in} - S_1) \\ \frac{dX_{A,1}}{dt} = \mu_A(S_1)X_{A,1} - D_1X_{A,1} \\ \frac{dX_{B,1}}{dt} = \mu_B(S_2)X_{B,1} - D_1X_{B,1} \\ \frac{dS_2}{dt} = -\mu_A(S_2)X_{A,2} - \mu_B(S_2)X_{B,2} + D_2(S_1 - S_2) \\ \frac{dX_{A,2}}{dt} = \mu_A(S_2)X_{A,2} + D_2(X_{A,1} - X_{A,2}) \\ \frac{dX_{B,2}}{dt} = \mu_B(S_2)X_{B,2} + D_2(X_{B,1} - X_{B,2}) \end{array} \right.$$

One species in two tanks

Design based on the Modified Inverse Kinetics (MIK) function :



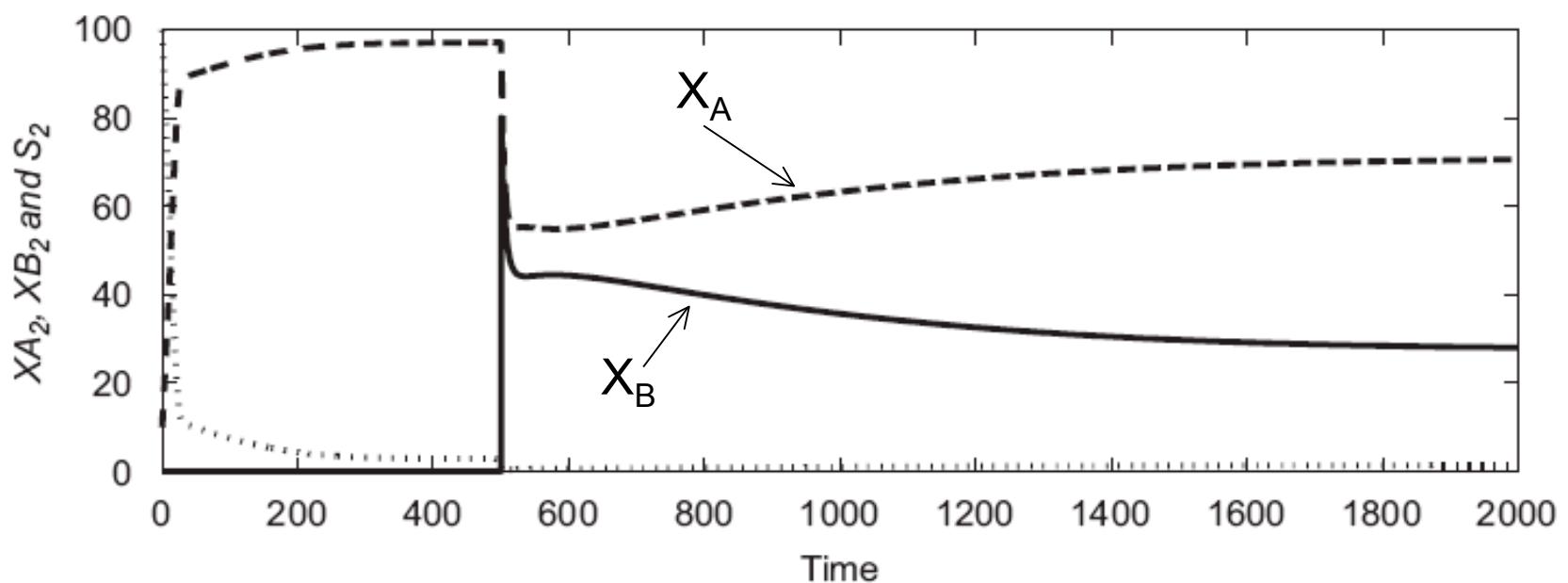
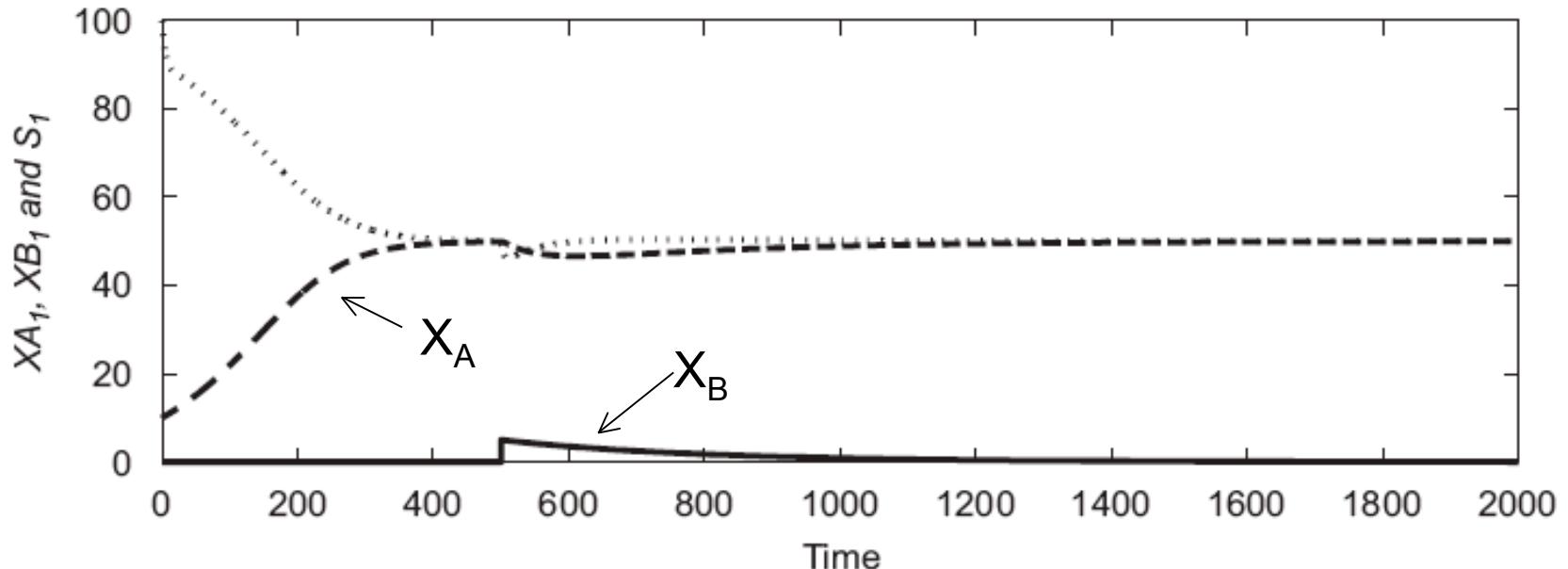
$$g(S) = \frac{1}{\mu(S)(S_{in} - S)}$$

- If $\mu^{-1}(S)$ is convex : MIK has a unique minimum on $]0, S_{in}[$
- If we have also $\frac{\mu(S)}{S}$ non-increasing : $V_1^{\text{opt}} \geq V_2^{\text{opt}}$

Invasion : two species in two tanks

- Break-even concentration («intersection» between D and μ) :
 $\lambda_A(D)$ and $\lambda_B(D)$
- Result :
 - $\lambda_A(D_1) < \lambda_B(D_1)$: B is washed out in tank 1
coexistence in tank 2 if $g_A(\lambda_B(D_2)) (\lambda_A(D_1) - \lambda_B(D_2)) > \frac{1}{D_2}$
 - $\lambda_A(D_1) > \lambda_B(D_1)$: A is washed out in tank 1
coexistence in tank 2 if $g_B(\lambda_A(D_2)) (\lambda_B(D_1) - \lambda_A(D_2)) > \frac{1}{D_2}$
 - Coexistence only possible if $V_1 < V_2$ (non-optimal design)

Coexistence (non optimal reactor design)



3. Practical coexistence

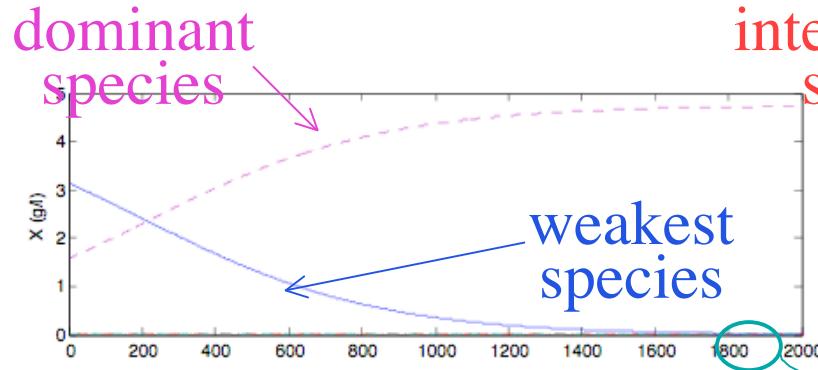
- Competitive exclusion principle :
predicts that only one species will survive
- But : - how long does it take for the dominant species
to take over?
- how will the non-dominant species survive?
- Practical coexistence : conditions for a «long» survival of
the weak species

- New variable : species proportion : $p_i = \frac{X_i}{\sum_{j=1}^n X_j}$
- Dynamics of p_i :

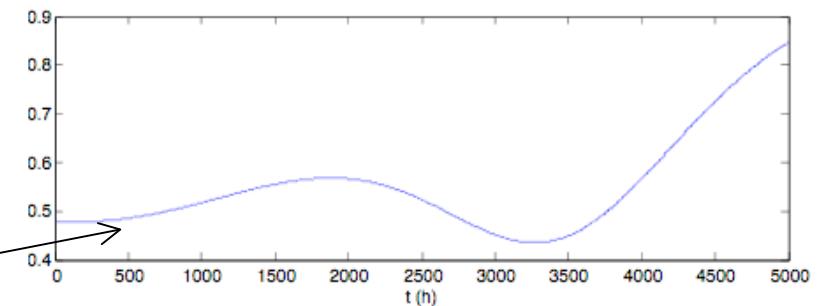
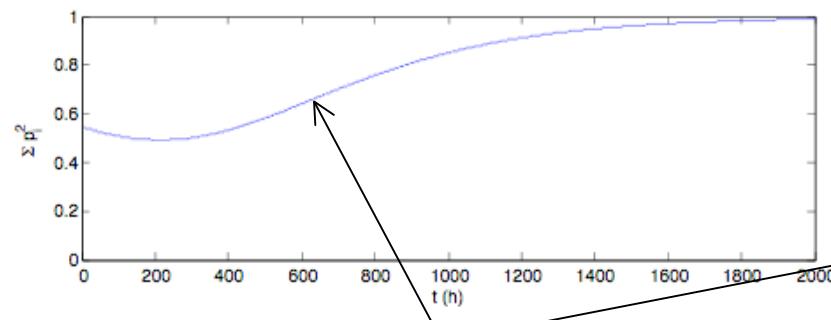
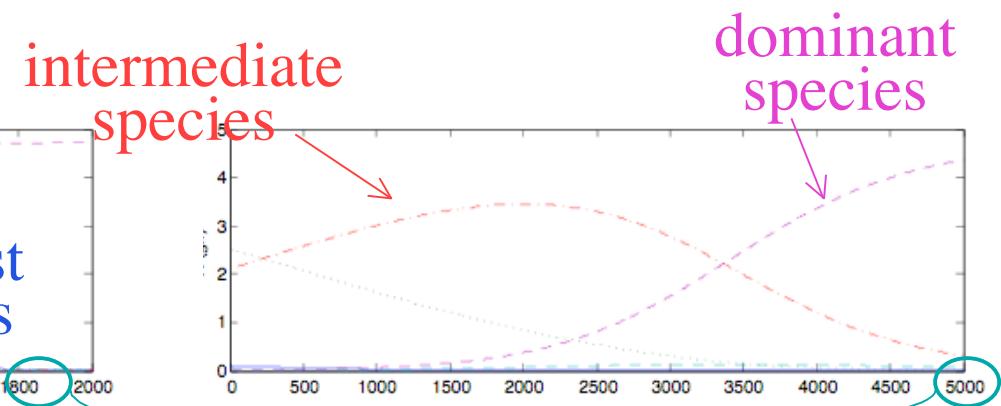
$$\begin{aligned}\frac{dp_i}{dt} &= \left(\sum_{j=1}^n (\mu_i(s) - \mu_j(s)) p_j \right) p_i \\ &= \sum_{j=1}^n A_{ij} p_j p_i \quad \text{with } A_{ij} \text{ skew symmetric}\end{aligned}$$

- **Results :**
 - whatever the initial conditions, the «weakest» species decreases, and the non-dominant increases
 - depending on the initial conditions, the others can possibly increase before decreasing (+ estimate of the increase duration)

Standard exclusion



Practical coexistence

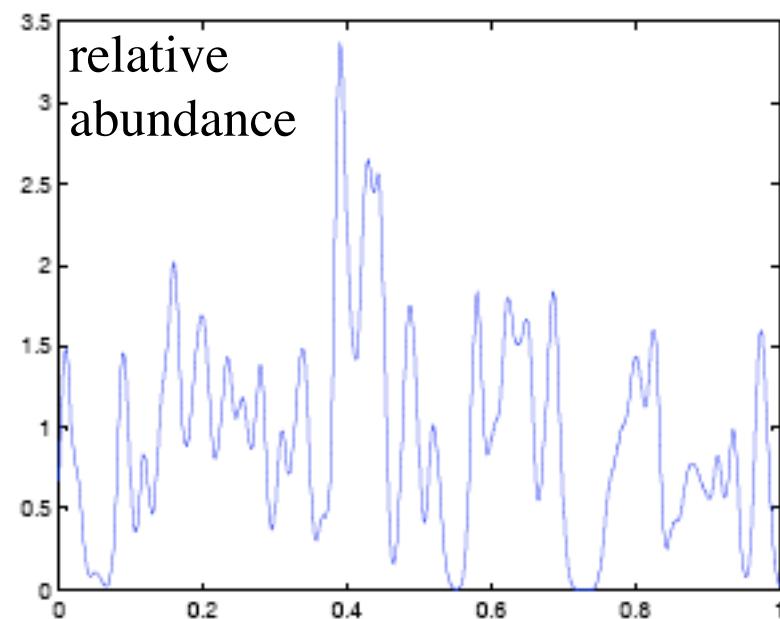
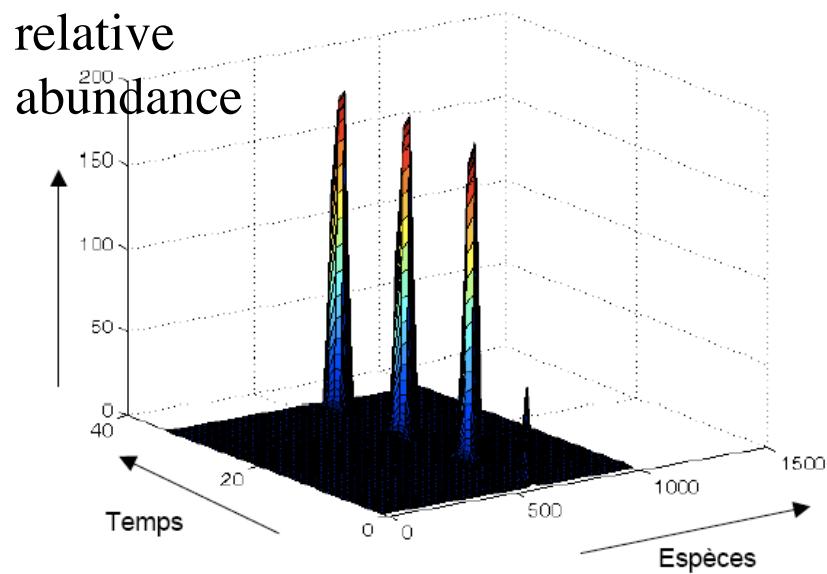


Biodiversity index $\sum p_i^2$ ($\rightarrow 1$ when only species survives)

Issues and challenges

- Coexistence/competition are not just limited to ecology...
- The knowledge of the dynamical mechanisms of coexistence/competition of microbial species can be helpful for improving the running of industrial biological processes, e.g. :
 - Invasion of a culture by a contaminant
(Can we avoid systematic re-inoculation?)
 - Mixed cultures, e.g. :
 - * Lactic fermentation (*L. bulgaricus* vs *S. thermophilus*)
 - * Anaerobic digestion (*thermophilic* vs *mesophilic* bacteria (*Tatarovsky et al*))

- Combination of theoretical/experimental studies
--> Molecular fingerprints via SSCP



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Thank you very much for your attention



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