

Modèles mathématiques pour des écosystèmes complexes

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Outline

1.  Optimisation of biofilm productivity,
 - ▶ Context,
 - ▶ Model,
 - ▶ Simulations,
 - ▶ Perspectives.
2.  Contribution of (epi)genetic in periodic environment,
 - ▶ Context,
 - ▶ Model & Analysis,
 - ▶ Simulations,
 - ▶ Perspectives.

Phototrophic biofilm

Phototrophic?

Biofilm using light and inorganic carbon source to growth.



(a) Rotating microalgae biofilm device
Hans C. Bernstein et al. 2014

Motivation:

Credible alternative for biofuels

Why?

- ▶ High production yield for lipids,
- ▶ Easy to harvest (just scalp),
- ▶ A wide variety of species,
- ▶ Can develop in sea and oceans,
- ▶ Combined with wastewater treatment?

Objective:

Quantify the influence of growing conditions and harvest on productivity.



Fig. 2 The vertical phototrophic biofilm reactor (**a**), a close-up of the biofilm before harvesting (**b**), harvesting part of the biofilm with the adhesive comb (**c**), and the biofilm setup after harvesting (**d**)

Figure: Boelee et al. 2014

- ▶ **Observation:** Harvesting pattern impact biofilm productivity
- ▶ **Question:** What is the optimal harvesting strategy?
- ▶ **Context:**
 - Collaboration with biologist and experimentalist:
 - O. Bernard (Lov & Inria),
 - F. Lopes (CentraleSupélec),
 - A. Fanesi (CentraleSupélec),
 - M. Ribot (Institut Denis Poisson)

Theoretical framework for mixture theory

Mixture of $K \geq 1$ constituents: \mathbf{C}_k , each constituent is described by:

- ▶ Its volume fraction: $\phi_k(t, x) := \lim_{\mathbb{V} \rightarrow 0} \frac{\text{volume of } \mathbf{C}_k \text{ in } \mathbb{V}}{\text{volume of } \mathbb{V}}$
- ▶ Its speed $V_k(t, x)$
- ▶ Its mass density ρ_k (assumed constant)

Fundamental properties:

- ▶ Total volume conservation: $\sum_{k=1}^K \phi_k = 1$
- ▶ Mass balance equation:

$$\underbrace{\partial_t(\phi_k \rho_k) + \nabla_x \cdot (\phi_k \rho_k V_k)}_{\text{transport}} = \underbrace{\nabla_x \cdot (D_k \nabla_x(\phi_k \rho_k))}_{\text{diffusion}} + \underbrace{\Gamma_k}_{\text{exchanges}}$$

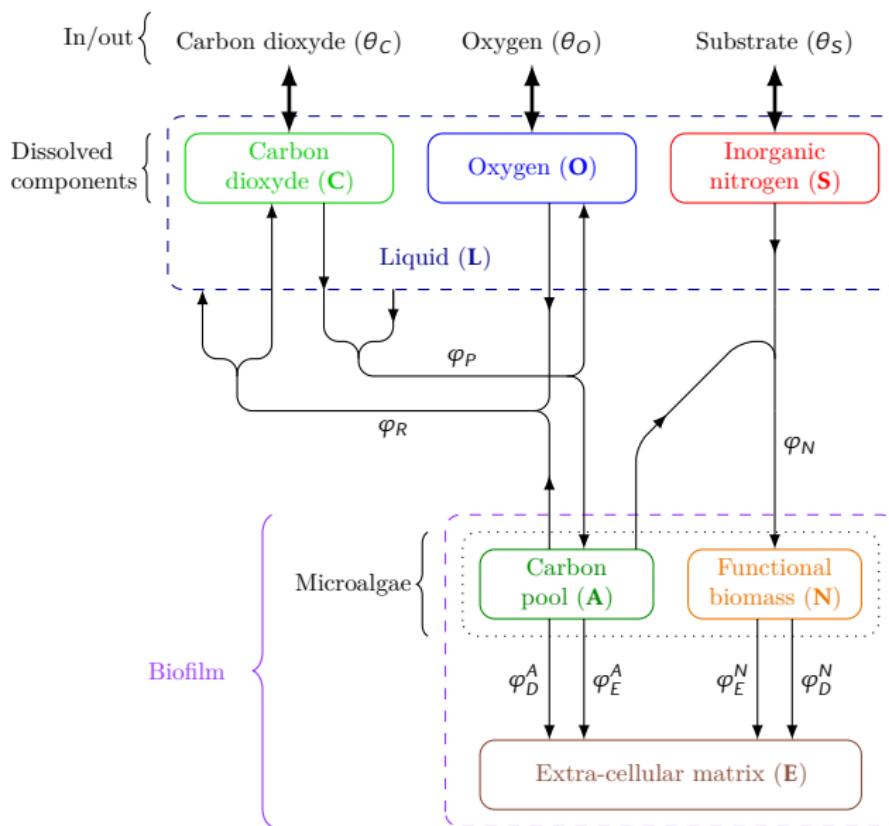
- ▶ Momentum conservation (Force balance equation):

$$\underbrace{\partial_t(\phi_k \rho_k V_k) + \nabla_x \cdot (\phi_k \rho_k V_k \otimes V_k)}_{\text{inertial terms}} = \underbrace{-\phi_k \nabla_x P}_{\text{pressure}} + \underbrace{F_{\text{fric}} + F_{\text{visc}} + \dots}_{\text{other forces}}$$

Advantages:

- Mesoscale
- Physical constraints included
- Different properties for each \mathbf{C}_k
- Interfaces without free boundary

Schematic representation of the system



Mixture framework – Mass balance

- Total volume conservation: $A + N + E + L = 1$
- Mass conservation: Transport

$$\text{Microalgae} \quad \left\{ \begin{array}{ll} \text{Carbon pool:} & \partial_t A + \nabla_x \cdot (A \mathbf{V}_M) = \Gamma_A / \rho \\ \text{Functional biomass:} & \partial_t N + \nabla_x \cdot (N \mathbf{V}_M) = \Gamma_N / \rho \\ \text{Extracellular matrix:} & \partial_t E + \nabla_x \cdot (E \mathbf{V}_E) = \Gamma_E / \rho \\ \text{Liquid:} & \partial_t L + \nabla_x \cdot (L \mathbf{V}_L) = \Gamma_L / \rho \end{array} \right.$$

- Pseudo incompressibility: Mass transfers \Rightarrow Pressure

$$\nabla_x \cdot ((A + N) \mathbf{V}_M + E \mathbf{V}_E + L \mathbf{V}_L) = \frac{1}{\rho} (\Gamma_A + \Gamma_N + \Gamma_E + \Gamma_L)$$

- Dissolved components: Transport by the liquid & Diffusion

$$\theta = \begin{cases} S & \text{Substrate} \\ C & \text{Carbon dioxide} \\ O & \text{Oxygen} \end{cases} \quad \partial_t(\theta L) + \nabla_x \cdot (\theta L \mathbf{V}_L) = \underbrace{\nabla_x \cdot \left(D_\theta \mathbf{L} \nabla_x \theta \right)}_{\text{diffusion}} + \frac{\Gamma_\theta}{\rho_L}$$

Mixture framework – Force balance

- Biological phases: $\phi = A + N, E$

$$\partial_t(\phi \rho_\phi v_\phi) + \nabla_x \cdot (\phi \rho_\phi v_\phi \otimes v_\phi) =$$
$$- \underbrace{\phi \nabla_x P}_{\text{Pressure}} + \underbrace{\nabla_x(\gamma_\phi \phi)}_{\text{Elastic}} + \underbrace{\sum_{\ell \neq \phi} m_{\ell, \phi} (v_\phi - v_\ell)}_{\text{Friction}} + \underbrace{\Gamma_\phi v_\phi}_{\text{Exch.}}$$

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- Hypothesis: Conservation of total momentum supply

Mixture framework – Force balance

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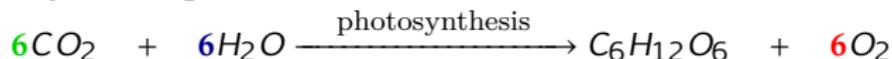
- Hypothesis: Conservation of total momentum supply
- Liquid phase:

$$\partial_t(L \rho_L V_L) + \nabla_x \cdot (L \rho_L V_L \otimes V_L) =$$
$$- \underbrace{L \nabla_x P}_{\text{Pressure}} - \underbrace{\sum_{\phi \neq L} m_{k, L} (V_L - V_\phi)}_{\text{Friction}} - \underbrace{\sum_{\phi \neq L} \Gamma_\phi V_\phi}_{\text{Exch.}}$$

Source terms modelling

► Construction of source terms:

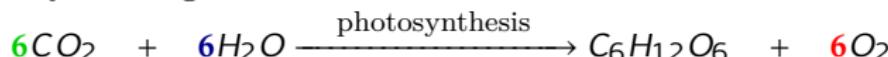
1. Identify a biological mechanism



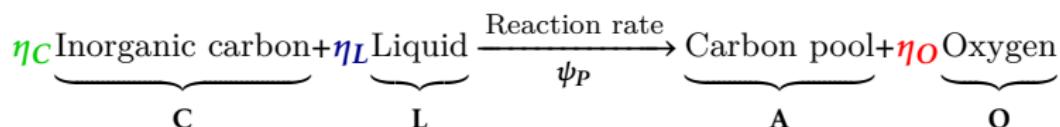
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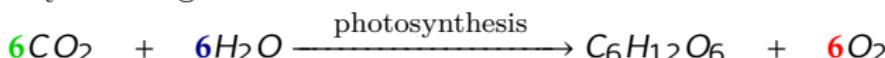
2. Translate in term of considered components



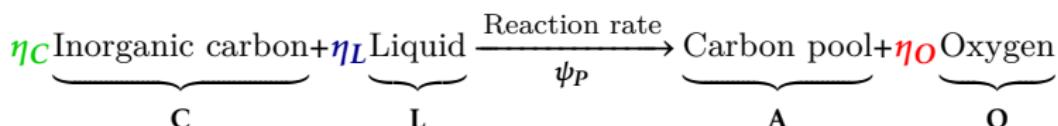
Source terms modelling

► Construction of source terms:

1. Identify a biological mechanism



2. Translate in term of considered components



3. Express the information in the source terms:

$$\Gamma_C = -\eta_C \psi_P + \dots$$

$$\Gamma_A = \psi_P + \dots$$

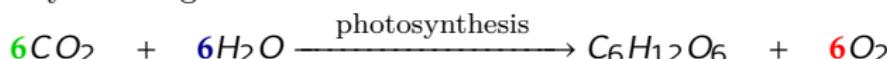
$$\Gamma_L = -\eta_L \psi_P + \dots$$

$$\Gamma_O = \eta_O \psi_P + \dots$$

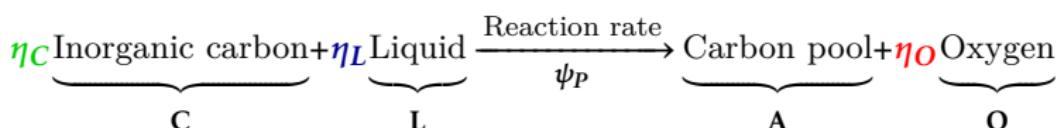
Source terms modelling

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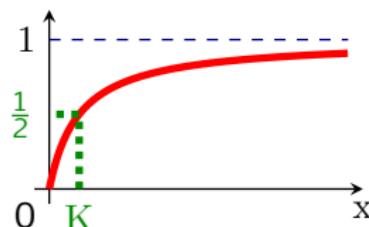
$$\Gamma_O = \eta_O \psi_P + \dots$$

► Considered mechanisms:

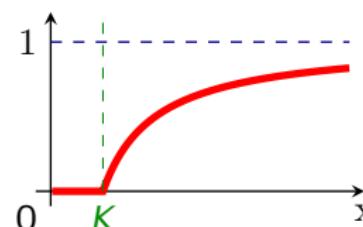
1. Photosynthesis
2. Respiration
3. Functional biomass synthesis
4. Extra-cellular matrix excretion
5. Mortality

Reaction rates modelling: ψ

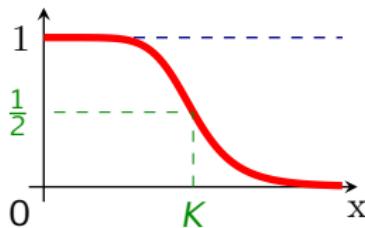
$$\psi := \prod_{i \geq 0} f_i(\phi) \quad \begin{cases} f_i & \text{elementary functions} \\ \phi & \text{volume or mass fraction} \end{cases}$$



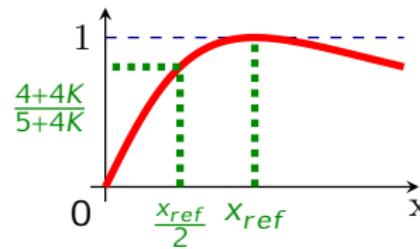
(a) Monod's law $f(x) = \frac{x}{K+x}$



(b) Droop's law $f(x) = \max\left\{0, 1 - \frac{K}{x}\right\}$



(c) Sigmoidal law $f(x) = \frac{1}{1+(x/K)^\alpha}$



(d) Haldane's law $f(x) = \frac{2(1+K)\bar{x}}{\bar{x}^2 + 2K\bar{x} + 1}$, $\bar{x} = x/x_{ref}$

Reaction rates modelling

- Highly nonlinear reaction rates:
Example:

$$\psi_P = \mu_P \rho_M N \frac{C}{\mathcal{K}_C + C} \frac{(1 + \mathcal{K}_L)L}{\mathcal{K}_L + L} \frac{2(1 + \mathcal{K}_I)\hat{I}}{\hat{I}^2 + 2\mathcal{K}_I\hat{I} + 1} \frac{\max\left\{0, 1 - \frac{Q_{min}}{\min\{Q, Q_{max}\}}\right\}}{Q_{max} - Q_{min}} \frac{1}{1 + \left(\frac{O}{\mathcal{K}_O}\right)^\alpha},$$

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- Received light intensity:

$$\hat{I}(z) = \frac{I_0}{I_{opt}} \exp\left(- \int_z^H \tau_L L + \tau_M (A + N + E) d\xi\right)$$

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- Functional biomass quota: $Q = \frac{N}{N+A}$.

Reaction rates modelling

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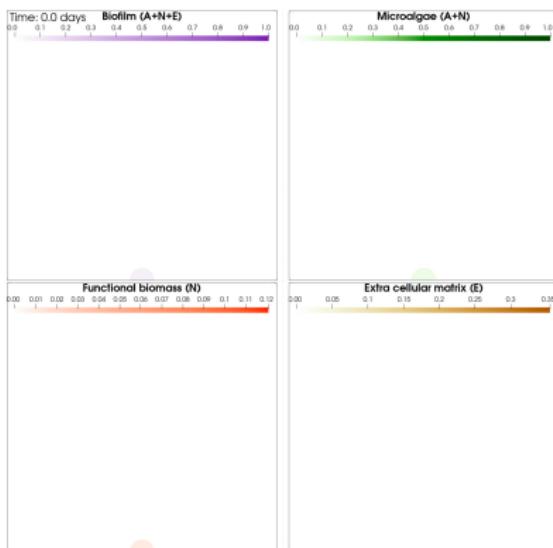
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- Functional biomass quota: $Q = \frac{N}{N+A}$.
- Coupled mass balances:

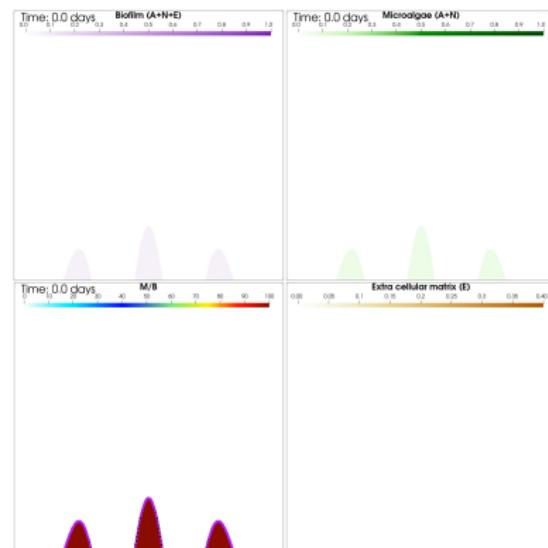
$$\partial_t A + \nabla_x \cdot (AV_M) = \frac{1}{\rho_M} (\psi_P - \psi_R - \eta_N^A \psi_N - \psi_E^A - \psi_D^A)$$

$$\partial_t (CL) + \nabla_x \cdot (CLV_L) - \nabla_x \cdot (\mathcal{D}_C L \nabla_x C) = \frac{1}{\rho_L} (\eta_R^C \psi_R - \eta_P^C \psi_P),$$

2D numerical simulations

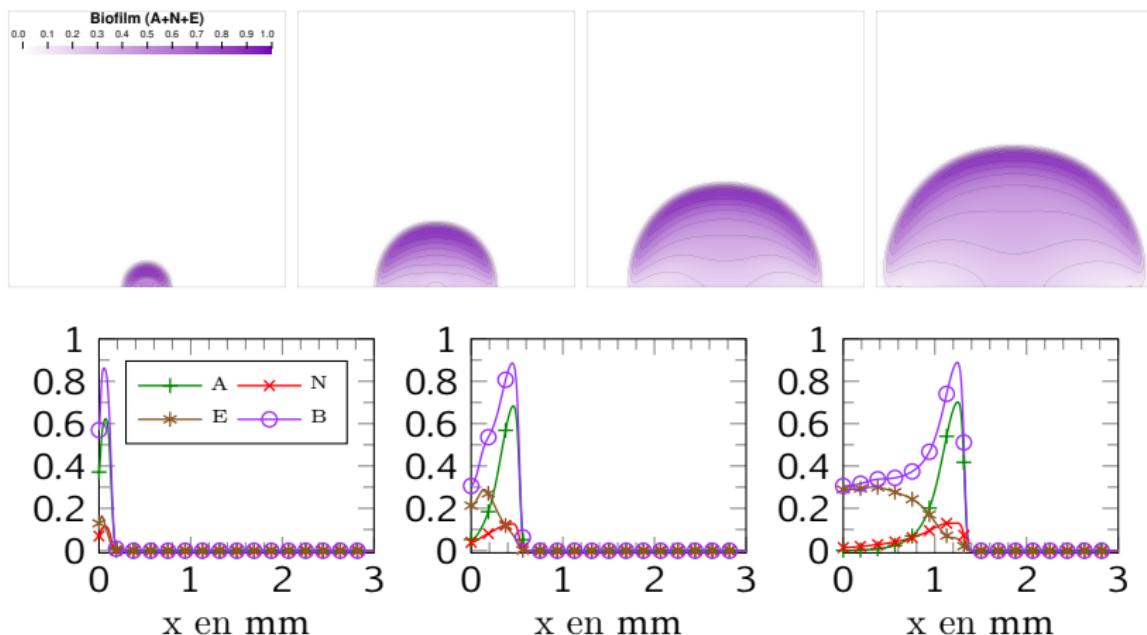


(a) Single spot colonie

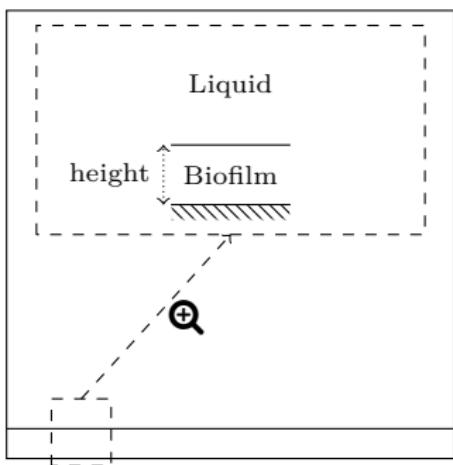


(b) Three spots colonie

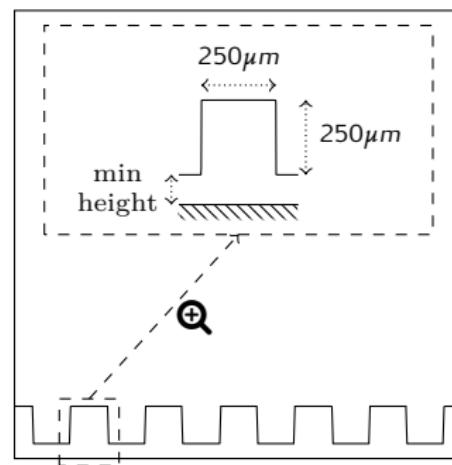
1D versus 2D numerical simulations



Harvest policy: Battlement versus uniform



(a) Uniform harvesting



(b) Battlement harvesting

Example: Battlement harvesting video.

Harvest policy: Impact of height and frequency?

Objective

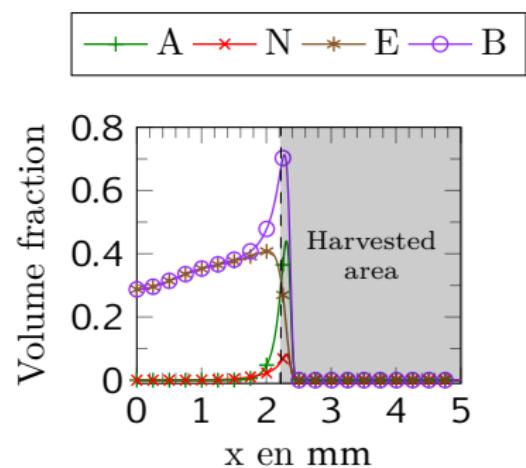
Quantify the impact of harvesting parameters on productivity.

Limitation

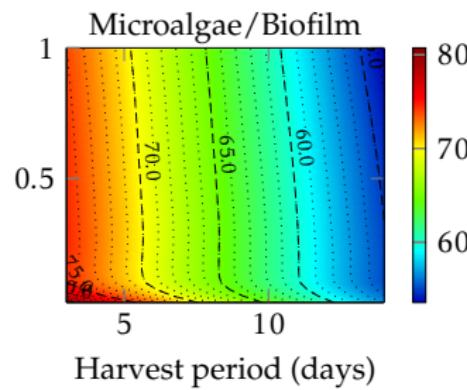
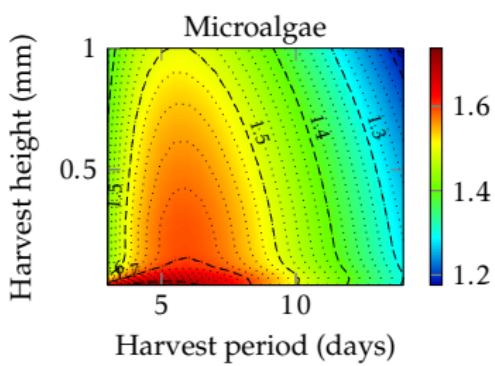
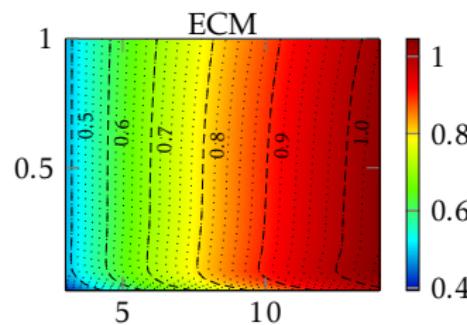
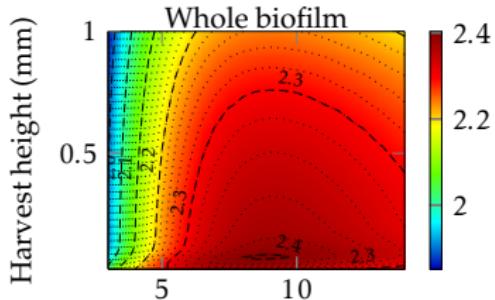
2D numerical simulation of ~ 50 days takes about two weeks on 2.40GHz Xeon.

Methodology

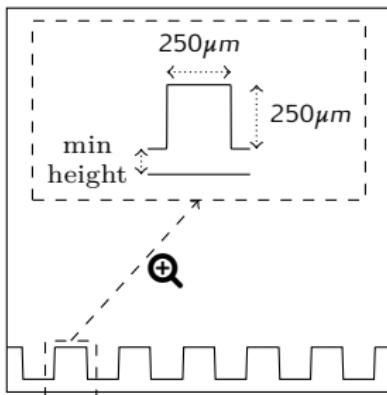
Investigate the impact of harvest height and frequency with 1D numerical simulation (about ~ 1 h for 90 days).



Harvest policy: Impact of height and frequency



Harvest policy: Impact of scraping pattern

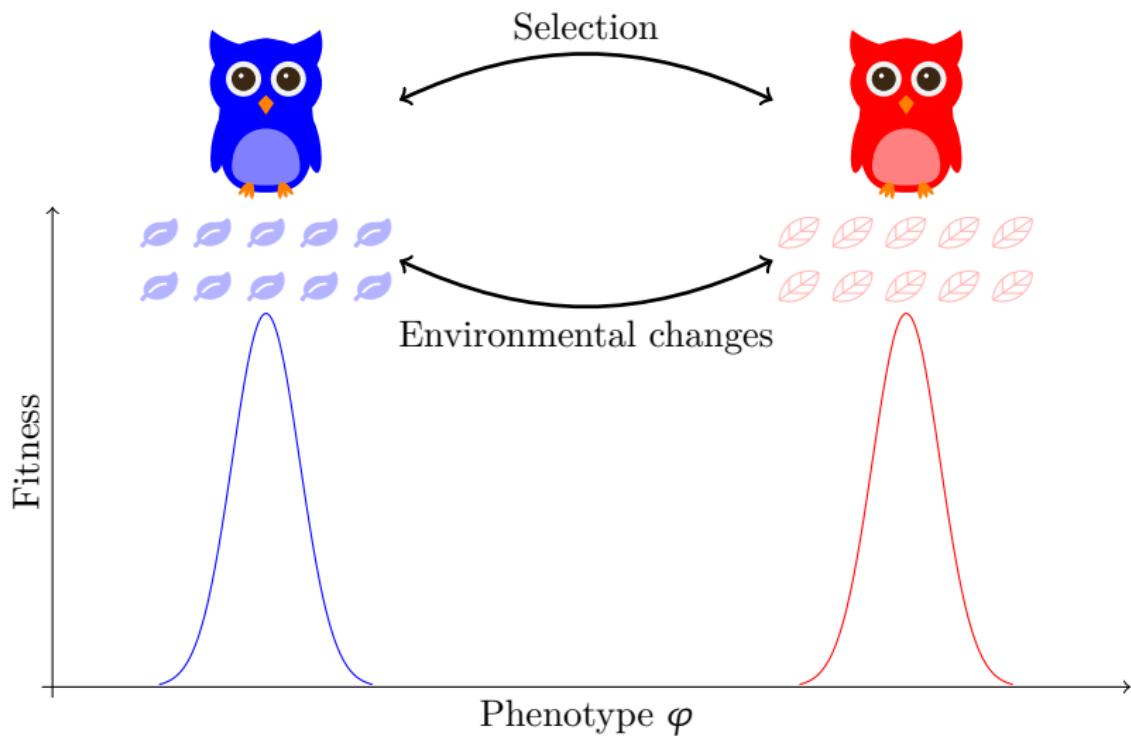


Battlement harvest shape:

- ▶ significantly increases productivity,
- ▶ modifies components productivity.

Period Min height	(days) (µm)	4 100	6.5 100	9 75
A	(%)	10.33	9.98	1.90
M	(%)	8.87	8.97	1.77
E	(%)	-19.77	-2.44	10.58
B	(%)	-1.31	4.92	4.90

Evolution: process of mutation and selection



General context

- ▶ **Observation:** Heredity is not just genetics
- ▶ **Debate:** What is the contribution of non-genetic heredity to evolution?
- ▶ **Difficulties:**
 - Non-genetic inheritance is multiple and heterogeneous,
eg. methylation \neq culture,
 - It is complex to formalize and analyze (\neq genetic),
 - Evolution theory is based on population genetic
whereas others mode of heredity is unclear and under
debate.

Approach from E. Rajon & S. Charlat, 2018

Philosophy

Idea: Expand the concept of mutation rate to epimutation rate,

Epigenetic: Heritable phenotype changes that do not involve alterations in the DNA sequence,

Advantages:

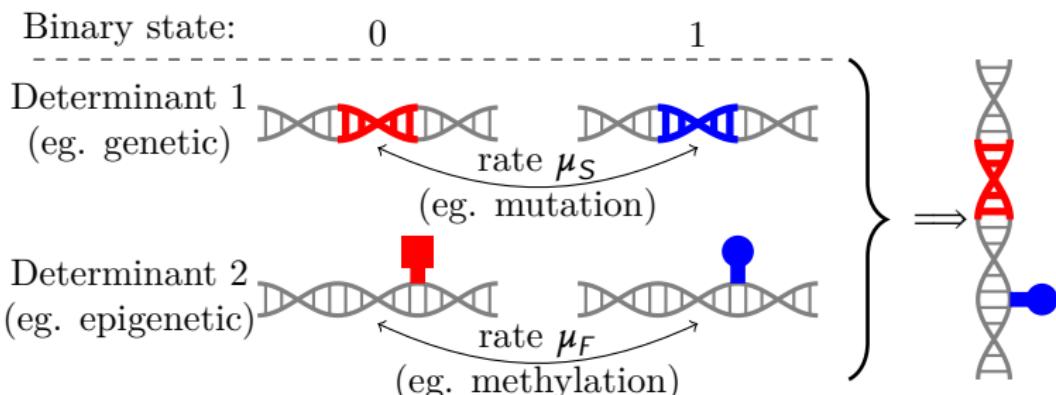
- ▶ Provides a comparable parameter between heterogeneous systems,
- ▶ Clear mathematical formalisms: epigenetic mutation rate \gg genetic mutation rate

Collaborators:

- ▶ E. Rajon & S. Charlat from Laboratoire de Biométrie et Biologie Évolutive, Université de Lyon, Université Lyon 1,
- ▶ V. Calvez from Institut Camille Jordan (ICJ), Université Claude Bernard Lyon 1

Approach from E. Rajon & S. Charlat, 2018

Schematic description of individuals

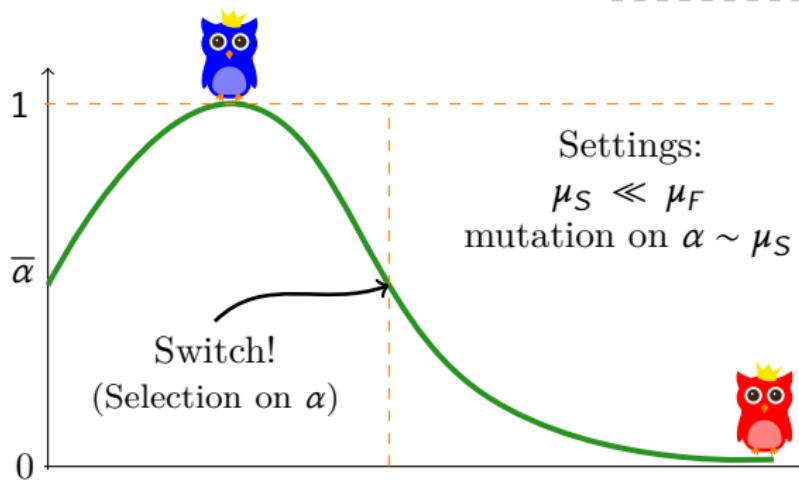
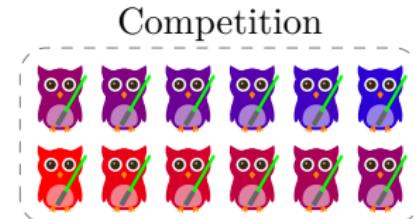
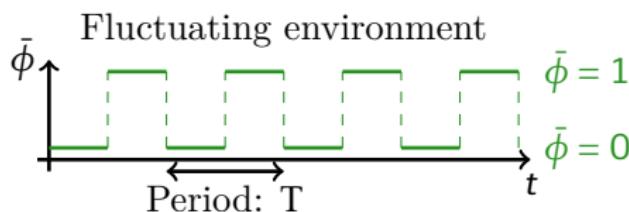


Questions:

- What is the contribution of non-genetic inheritance to evolution?
- Is the contribution of epigenetic (ie. α) under selection?

Approach from E. Rajon & S. Charlat, 2018

Selection in fluctuating environment



Mathematical model

The model is an adaptation of E. Rajon & S. Charlat, 2018.

$$\underbrace{\partial_t G_i}_{\text{Evolution}} = \underbrace{\mu_\alpha \int_0^1 \mathcal{K}(\alpha, \alpha') G_i(t, \alpha') d\alpha'}_{\text{Modifier locus mutations}} + \underbrace{\eta \left(1 - \frac{\rho}{N}\right) G_i}_{\text{Limited growth}} - \underbrace{s |\phi_i - \bar{\phi}(t)|^\gamma G_i}_{\text{Fitness selection}} + \underbrace{(M \times \mathcal{G})_i}_{\text{Déterminant's mutation}}$$

Groups	G_1	G_2	G_3	G_4
Determinant set	{0, 0}	{0, 1}	{1, 0}	{1, 1}
Phenotypes: ϕ_i	0	α	$1 - \alpha$	1

- μ_α mutation rate on α (very rare $\implies \mu_\alpha \sim \mu_S$),
- Mutation kernel: $\mathcal{K}(\alpha, \alpha')$ is gaussian law centered in α' ,
- Maximal growth rate: η ,
- Total population size: $\rho(t) = \sum_{i=1}^4 \int_0^1 G_i(t, \alpha) d\alpha$,

Mathematical model

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$$\underbrace{\partial_t G_i}_{\text{Evolution}} = \underbrace{\mu_\alpha \int_0^1 \mathcal{K}(\alpha, \alpha') G_i(t, \alpha') d\alpha'}_{\text{Modifier locus mutations}} + \underbrace{\eta \left(1 - \frac{\rho}{N}\right) G_i}_{\text{Limited growth}} - \underbrace{s |\phi_i - \bar{\phi}(t)|^\gamma G_i}_{\text{Fitness selection}} + \underbrace{(M \times \mathcal{G})_i}_{\text{Déterminant's mutation}}$$

- ▶ Selection rate: s ,
- ▶ Phenotype of subpopulation: $\phi_i \in \{0, \alpha, 1 - \alpha, 1\}$,
- ▶ Phenotype environment promoted: $\bar{\phi}(t) \in \{0, 1\}$,
- ▶ Mutations:

$$\mathcal{M} \times \mathcal{G} = \begin{pmatrix} -\mu_F - \mu_S & \mu_F & \mu_S & 0 \\ \mu_F & -\mu_F - \mu_S & 0 & \mu_S \\ \mu_S & 0 & -\mu_F - \mu_S & \mu_F \\ 0 & \mu_S & \mu_F & -\mu_F - \mu_S \end{pmatrix} \times \begin{pmatrix} G_1 \\ G_2 \\ G_3 \\ G_4 \end{pmatrix}.$$

Mathematical model analysis

Existence of solution

Theorem:

Under mild assumptions, the system has a unique nonnegative solution.

Proof idea: Banach–Picard fixpoint method (Cf. Transport Equations in Biology - B. Perthame).

Mathematical model analysis

Lyapunov exponent approach in infinite population

Hypothesis: Asymptotically (ie. on a long range of repeated periods) the population is monomorphic with respect to α .

Equilibrium characterisation: Floquet spectral problem, where the Floquet eigenvalue is the average of the fitness over a period :

$$\omega(\alpha) = \exp(\lambda(\alpha)) = \left(\text{eigenvalue} \left(\exp\left(\frac{T}{2}A\right) \exp\left(\frac{T}{2}B\right) \right) \right)^{\frac{1}{T}}$$

where A (resp. B) is the matrix of the evolutionary dynamics in environment \mathcal{A} (resp. \mathcal{B}) :

$$A = \begin{pmatrix} 1 & \mu_F & \mu_S & 0 \\ \mu_F & 1 - s\alpha^\gamma & 0 & \mu_S \\ \mu_S & 0 & 1 - s(1 - \alpha)^\gamma & \mu_F \\ 0 & \mu_S & \mu_F & 1 - s \end{pmatrix} - \frac{\rho(t)}{N} \text{Id} \quad (1)$$

Mathematical model analysis

Lyapunov exponent approach in infinite population

Interpretation: In an infinite population setting, where all possible architectures are present in the population, we expect that architectures which are defined via some α will predominate over another architecture β if $\lambda(\alpha) > \lambda(\beta)$.

Assume that the trait fitness function is convex. Then, the Lyapunov exponent $\lambda(\alpha)$ is a convex function of the trait architecture α .

Remark: This result generalises for any number of determinants.

Mathematical model analysis

Interpretation

1. Composite mutation rates are selected against: selection of extremal values of $\alpha \implies \alpha = 0$ or $\alpha = 1$.

Question: When does it switch from $\alpha = 0$ to $\alpha = 1$?

Mathematical model analysis

Interpretation

1. Composite mutation rates are selected against: selection of extremal values of $\alpha \implies \alpha = 0$ or $\alpha = 1$.

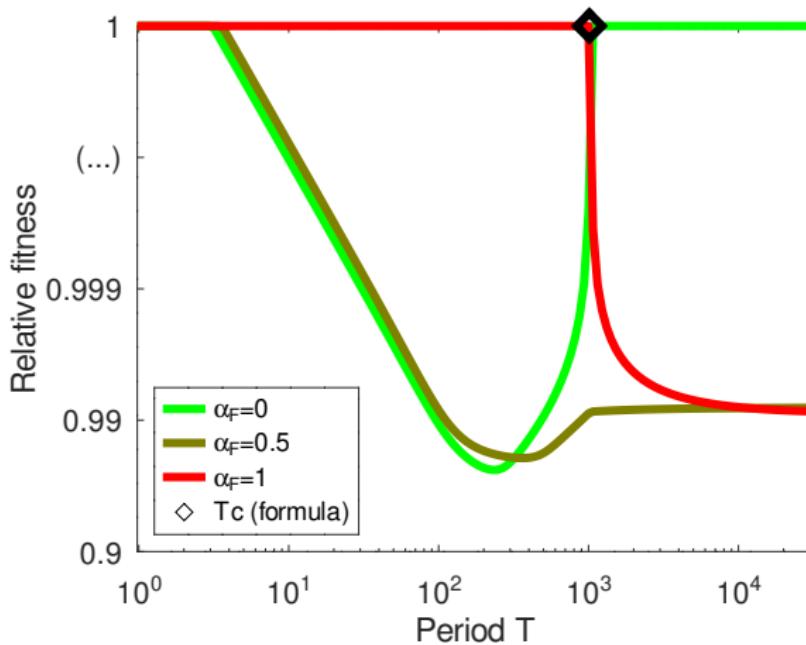
Question: When does it switch from $\alpha = 0$ to $\alpha = 1$?

2. In the regime $\mu_S \ll \mu_F$ the value of T_c can be computed analytically :

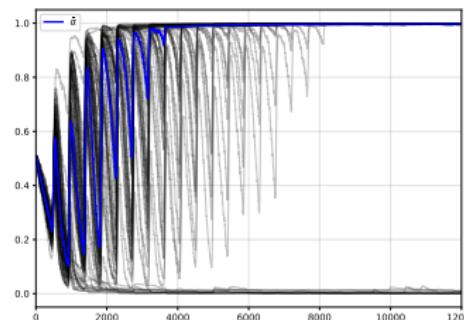
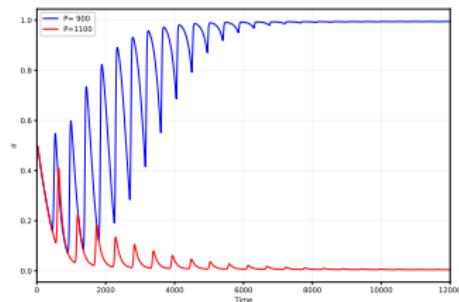
$$\frac{T_c}{2} \approx \frac{1}{\mu_F} \ln\left(\frac{\mu_F}{\mu_S}\right) \left(1 + \frac{\mu_F}{s}\right).$$

Numerical simulation

Relative fitness of the clones & estimation of the switch



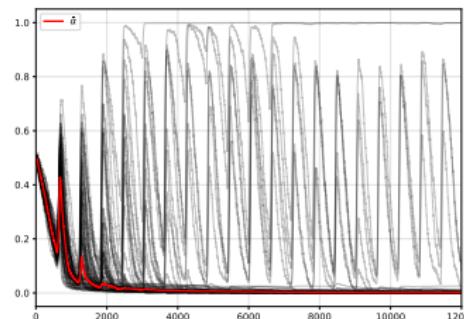
Selection on α for polyclonal finite and infinite population Size



(a) $P = 900$

Figure: Simulation of the PDE system \Rightarrow infinite population

Right side: Monte-Carlo simulations \Rightarrow finite population.



(b) $P = 1200$

Selection on α for large polyclonal population

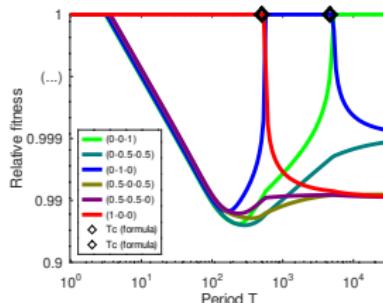


Figure: Relative fitness for 3 media

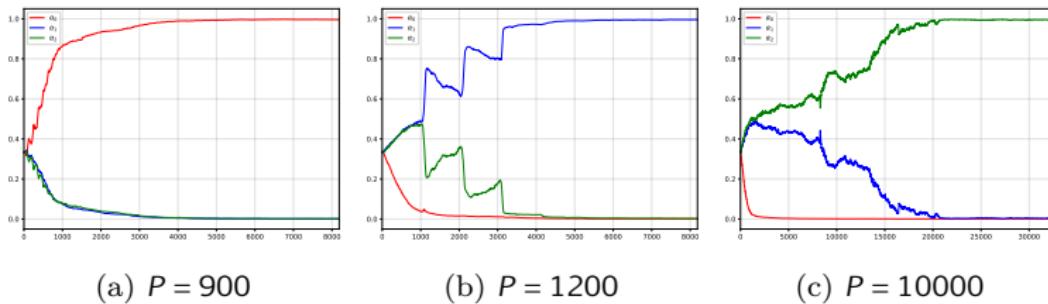


Figure: Monte-Carlo simulations \Rightarrow population of finite size

Numerical simulation

what happens if the fitness is non-convex?

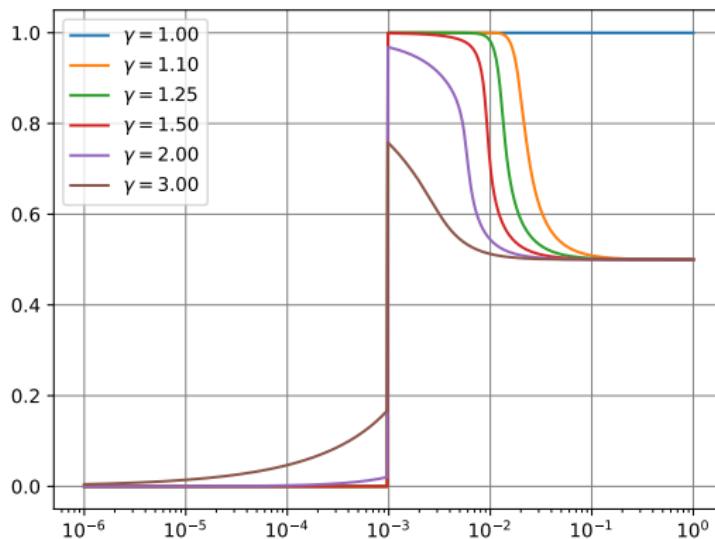
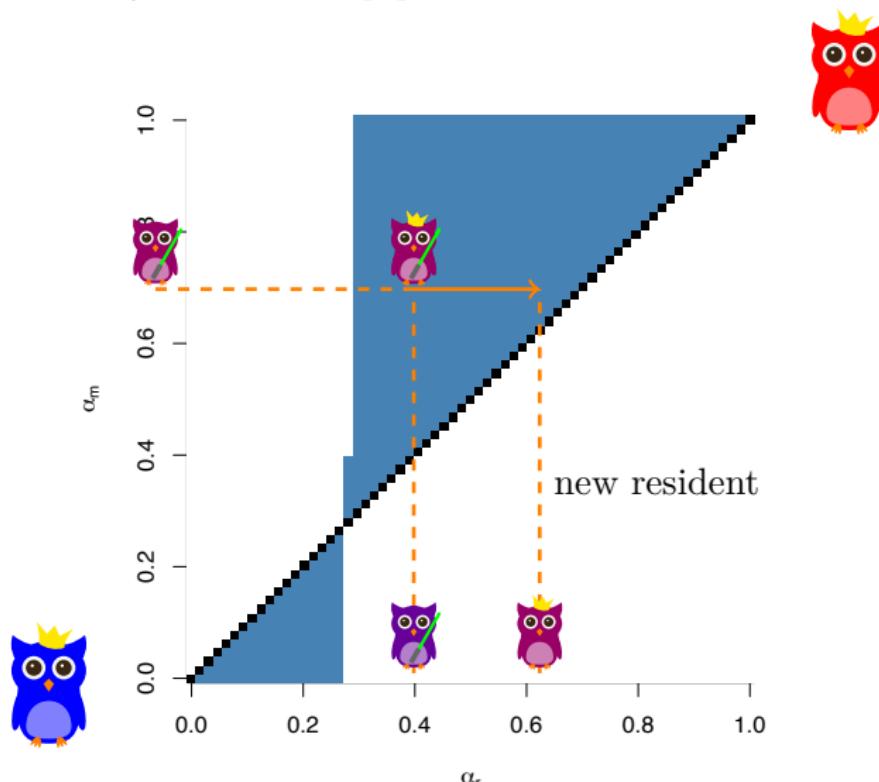


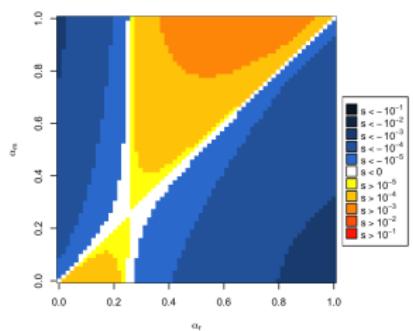
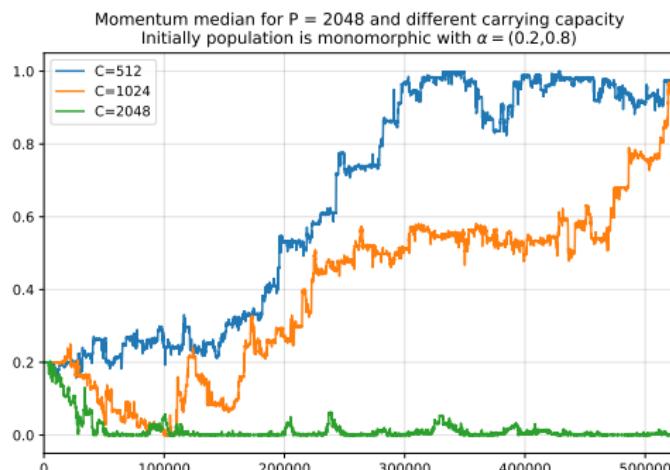
Figure: Selected α with respect to convexity of the selection law:
 $-s|\phi_i - \bar{\phi}(t)|^\gamma$

Adaptative dynamic & Invasion

What about the stability of monoclonal populations



Adaptative dynamic & Invasion

(a) $P = 2048$ (b) Median dynamic for monoclonal populations
(24 simulations for each period).



Conclusion & Perspectives

Results:

- ▶ Structure formation,
- ▶ Role of dissolved components in the development of structures,
- ▶ Quantification of limiting factors: light, Oxygen, ...
- ▶ Optimal strategy quantification for uniform harvest,
- ▶ Battlement scrapping harvest induce productivity increase.

Perspectives:

- ▶ Include the water flow,
- ▶ Take into account the viscosity,
- ▶ Multi-species biofilm,
- ▶ Calibration and comparison with experimental data.

 Conclusions & Perspectives

Conclusions:

- ▶ Asymptotic behaviour characterisation,
- ▶ Composite mutation rates are selected against (ie. $\alpha = 0$ or 1),
- ▶ Analytic expression for T_c .

Perspectives:

- ▶ Extend the results for random fluctuation periods,

Open questions:

- ▶ What is the dynamic for small populations?

Thank you for your attention!

Questions?

References:

- ▶ Understanding photosynthetic biofilm productivity and structure through 2D simulation, B. Polizzi, A. Fanesi, F. Lopes, M. Ribot, O. Bernard, Plos Computational Biology (accepted).
- ▶ A time-space model for the growth of microalgae biofilms for biofuel production, B. Polizzi, O. Bernard, M. Ribot, Journal of Theoretical Biology, Volume 432, 7 November 2017, Pages 55-79.
- ▶ (In)exhaustible suppliers for evolution? Epistatic selection tunes the adaptive potential of nongenetic inheritance,
Rajon Etienne and Charlat Sylvain,
The American Naturalist, 2019.
- ▶ The evolution of composite mutation rates,
Bastien Polizzi, Vincent Calvez, Sylvain Charlat and Etienne Rajon,
In preparation.

Details about the property proof

$$\partial_t G_i = \mu_\alpha \int_0^1 K(\alpha, \alpha') G_i(t, \alpha') d\alpha' + \eta(1 - \rho) G_i - s|\phi_i - \bar{\phi}(t)|G_i + (M \times \mathcal{G})_i$$

- ▶ $\mu_\alpha \sim \mu_S \implies \partial_t G_i = +\eta(1 - \rho)G_i - s|\phi_i - \bar{\phi}(t)|G_i + (M \times \mathcal{G})_i$,
- ▶ The system to solve is equivalent to $Q \exp(AT)\mathcal{G} = \lambda\mathcal{G}$ with $A = M - s\text{diag}(0, \alpha, 1 - \alpha, 1)$,
- ▶ The eigenvalue of interest is $\lambda = \exp\left(\eta \int_t^{t+T} (\rho(x) - 1) dx\right)$ and the ratio in each subpopulation is given by the eigenvector,
- ▶ Approximate eigen elements using Taylor expansions.

Remark: Numerically it is easy to extract the largest eigenvalue and its eigenvector.

NUMERICAL METHOD OVERVIEW

- Semi-implicit approach for the mass balance equations:

$$\frac{(\theta L)^{n+1} - (\theta L)^n}{dt} + \nabla_x \cdot (\theta L V_L)^n = \nabla_x \cdot \left(D_\theta L^n \nabla_x \frac{(\theta L)^{n+1}}{L^n} \right) + f(U^n) - g(U^n) (\theta L)^{n+1}$$

$f(U)$ production terms & $g(U)\theta L$ consumption terms

- Chorin-Temam's projection method for the conservation of momentum:

- 1 Projection step for V_ϕ , $\phi = M, E, L$:

$$\begin{aligned} \frac{(\phi V_\phi)^{n+\frac{1}{2}} - (\phi V_\phi)^n}{dt} + \nabla_x \cdot (\phi V_\phi \otimes V_\phi)^n \\ = \frac{1}{\rho_\phi} \left(-\nabla_x (\gamma_\phi \phi^n) + \sum_{\phi' \neq \phi} m_{\phi, \phi'} (V_\phi - V_{\phi'})^{n+\frac{1}{2}} + (\Gamma_\phi V_\phi)^n \right) \end{aligned}$$

- 2 Elliptic equation for P : Variable coefficients & Nonhomogeneous
- 3 Correction step: $V_\phi^{n+1} = V_\phi^{n+\frac{1}{2}} - \frac{\Delta t}{\rho_E} (\nabla_X P)^{n+1}$