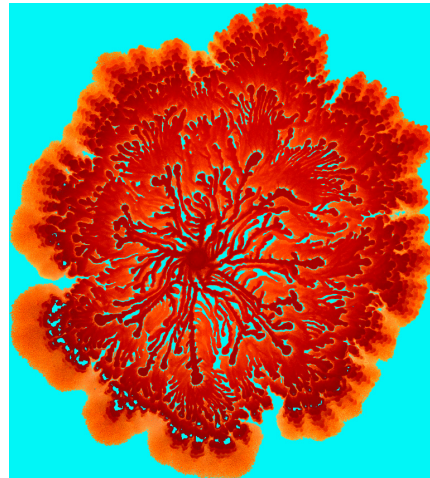
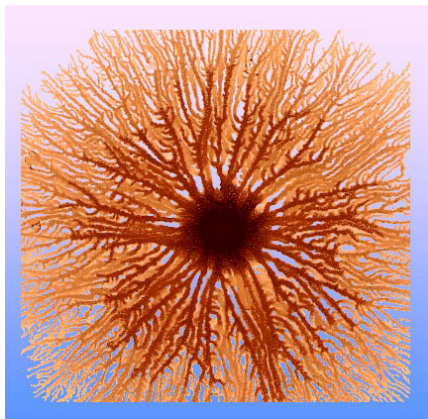


CHEMOTAXIS MODELS : mathematical analysis

Benoît Perthame, ENS, Paris



OUTLINE OF THE LECTURE

- I. How and why do cell move
- II. Keller-Segel model
- III. dentritic bacterial growth
- IV. Angiogenesis
- V. Hyperbolic and kinetic models

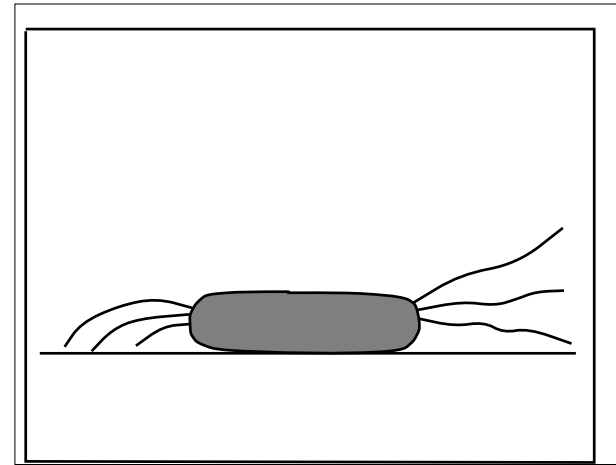
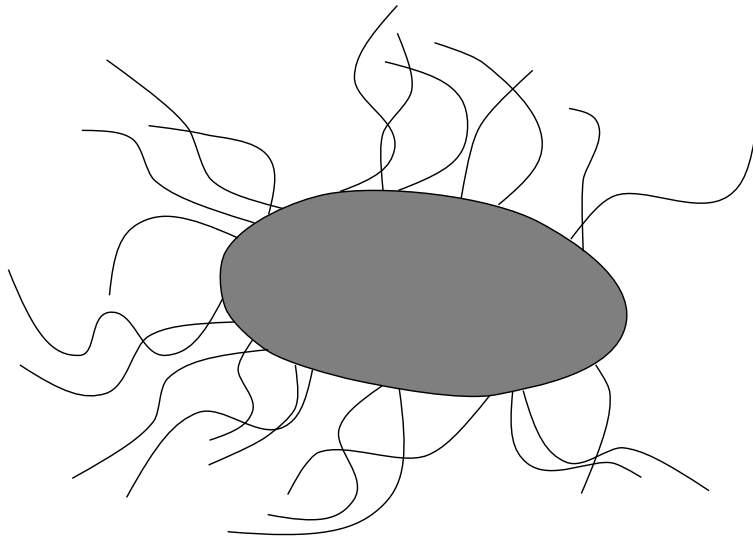
OUTLINE OF THE LECTURE

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COLLABORATORS

J. Dolbeault, L. Corrias, H. Zaag
F. Chalub, P. Markowich, C. Schmeiser
V. Calvez, F. Filbet, P. Laurencot
A. Marrocco, A. Vasseur

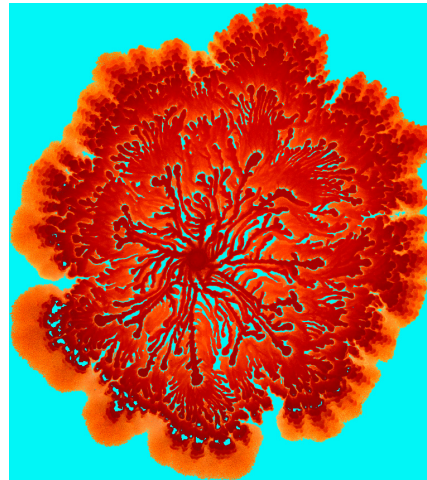
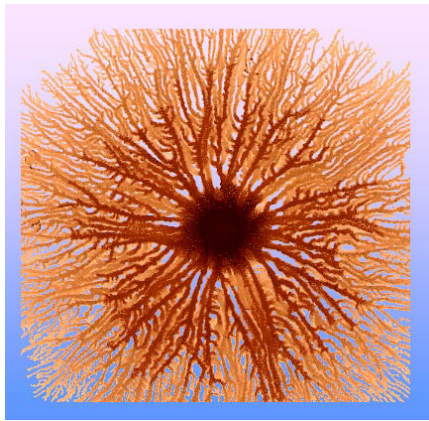
HOW DO CELLS MOVE : bacteria



E. Coli and Micrococcus are equipped with external devices ; they emit chemicals and can react to these chemicals

HOW DO CELLS MOVE : bacteria

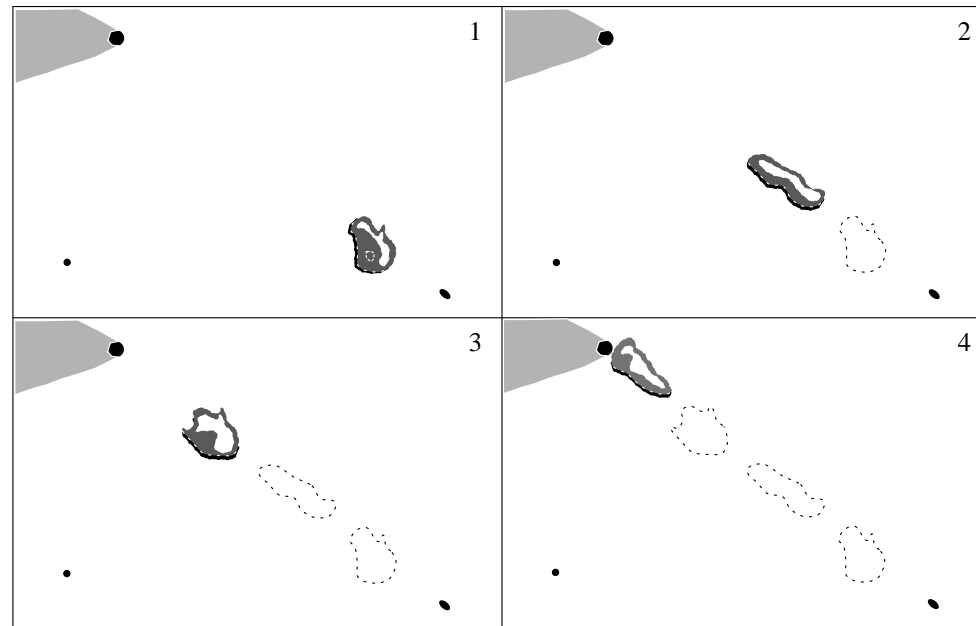
This creates a collective motion and results in patterns



By [D. Jukowska](#), [S. Seror](#), [B. Holland](#) (Institut de Génétique et Microbiologie)

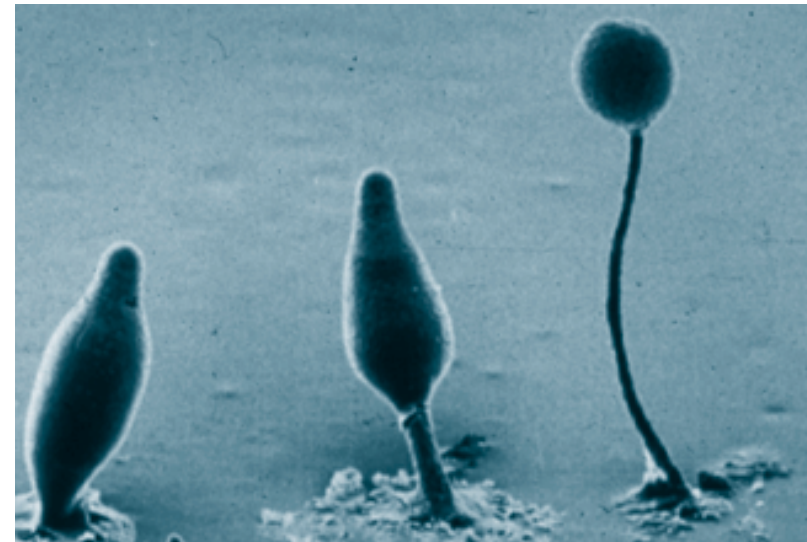
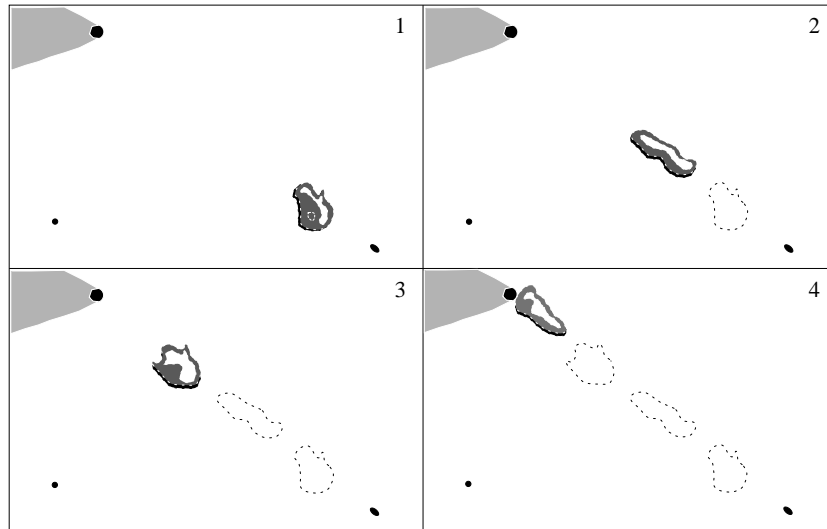
How to characterize these colonies ?

HOW DO CELLS MOVE : amoebia



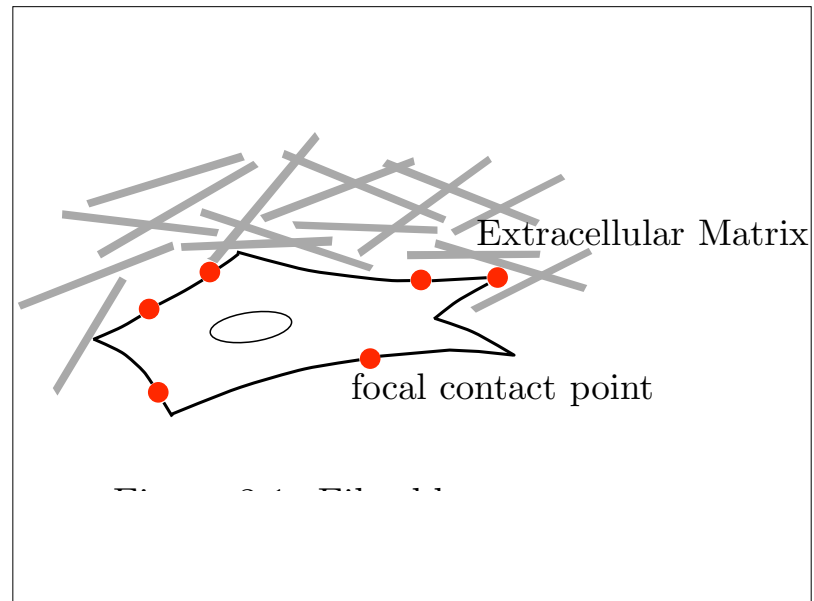
Dictyostelium Discoideum uses an internal pseudopod

HOW DO CELLS MOVE : amoebia

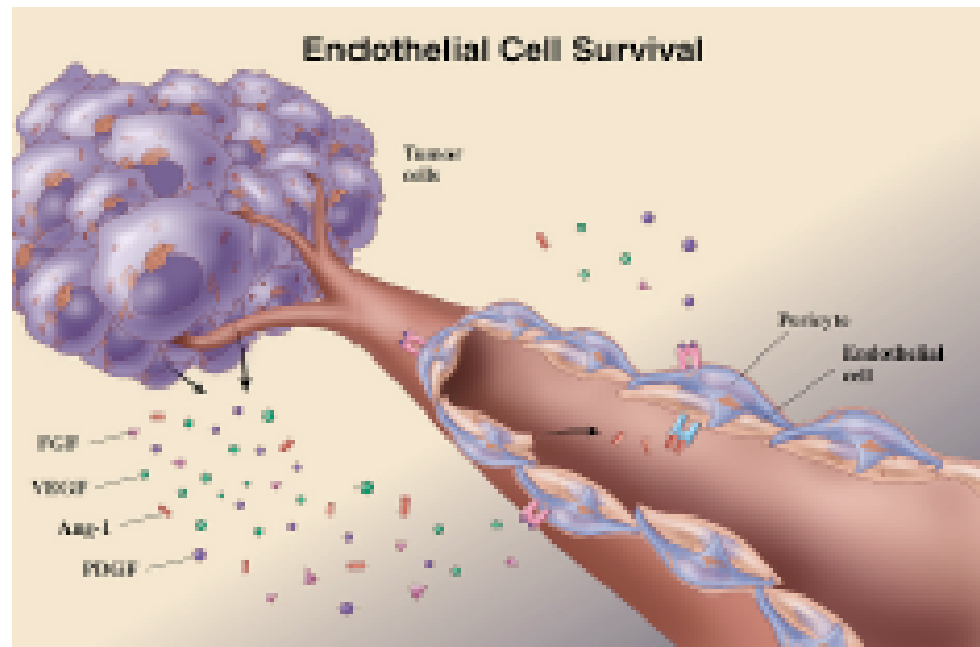
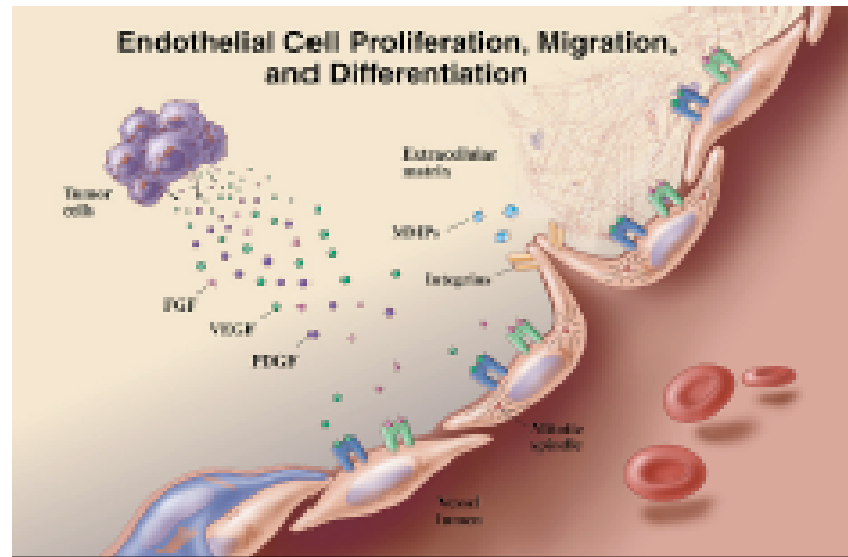
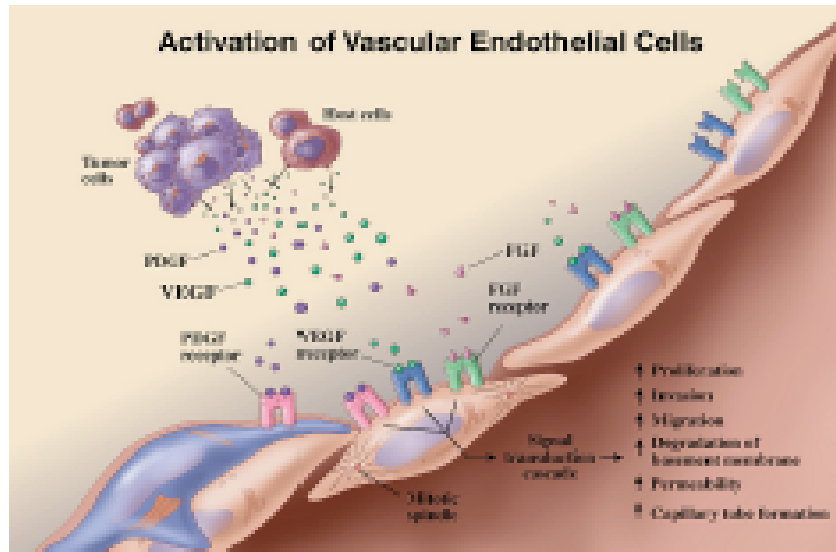


Dictyostelium Dyscoideum can create a fruiting body

HOW DO CELLS MOVE : fibroblast



In vivo, fibroblast can move within the extracellular matrix



HOW DO CELLS MOVE : tumor growth

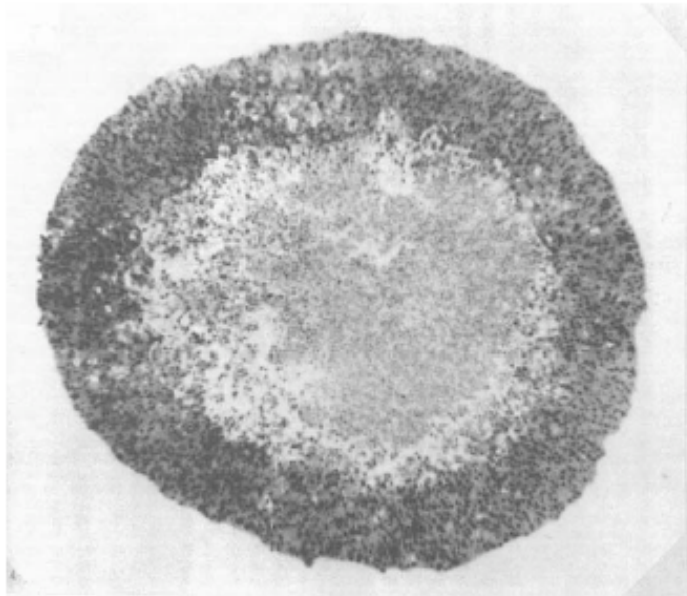
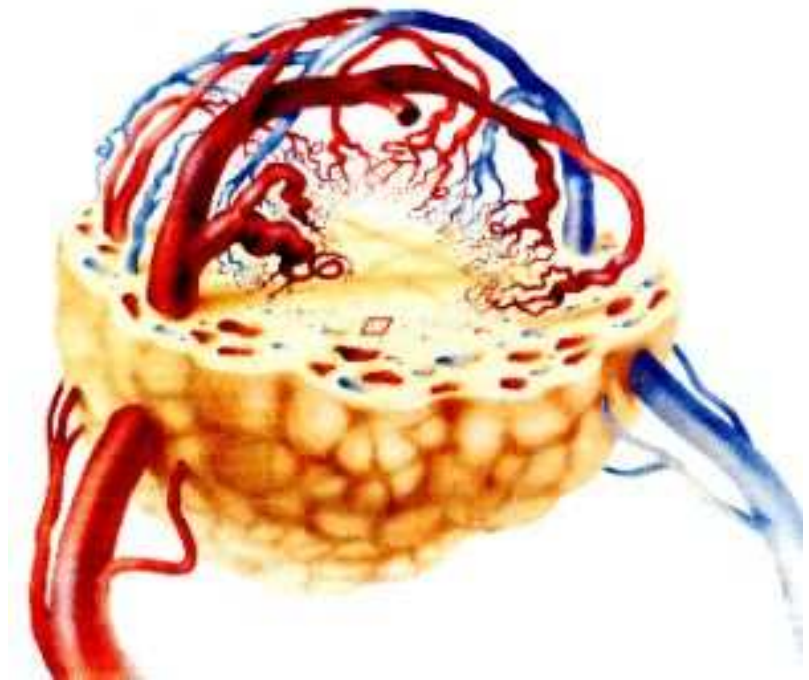


Fig. 1. An illustration of the structure of a multicellular tumour spheroid, with an outer rim of proliferating cells and an inner necrotic core; these are separated by a layer of quiescent cells. The spheroid diameter is 1.4 mm. [Reproduced from Sutherland et al. (Cancer Res.

Tumor spheroid



vascularized tumor through angiogenesis

HOW DO CELLS MOVE : endothelial cells

Animation from M. Mirshahi

(INSERM E 355 faculté de Médecine Paris VI)

CHEMOTAXIS : Keller-Segel model

The mathematical modelling of cell movement goes back to Patlak (1953), E. Keller and L. Segel (70's)

$$\begin{aligned} n(t, x) &= \text{density of cells at time } t \text{ and position } x, \\ c(t, x) &= \text{concentration of chemoattractant,} \end{aligned}$$

In a collective motion, the chemoattractant is emitted by the cells that react according to biased random walk.

$$\begin{cases} \frac{\partial}{\partial t} n(t, x) - \nu_{\text{bact}} \Delta n(t, x) + \text{div}(n \chi \nabla c) = 0, & x \in R^d, \\ \frac{\partial}{\partial t} c(t, x) - \nu_{\text{chem}} \Delta c(t, x) + \tau c = n(t, x), \end{cases}$$

The parameter χ is the sensitivity of cells to the chemoattractant.

CHEMOTAXIS : Keller-Segel model

$$\begin{cases} \frac{\partial}{\partial t}n(t, x) - \Delta n(t, x) + \operatorname{div}(n\chi\nabla c) = 0, & x \in R^d, \\ -\Delta c(t, x) = n(t, x), \end{cases}$$

This model, although very simple, exhibits a deep mathematical structure and mostly only dimension 2 is understood, especially "chemotactic collapse".

This is the reason why it has attracted a number of authors.

CHEMOTAXIS : Keller-Segel model

$$\begin{cases} \frac{\partial}{\partial t}n(t, x) - \Delta n(t, x) + \operatorname{div}(n\chi\nabla c) = 0, & x \in \mathbb{R}^d, \\ -\Delta c(t, x) = n(t, x). \end{cases}$$

- Childress, Percus (84); Jäger, Luckhaus (92),
- Rascle, Zitti (95); Nagai (95); Biler, Nadzieja(93),
- Herrero, Medina, Velazquez (96-04);
- Brenner, Constantin, Kadanoff, Schenkel, Venkatarami (98);
- Horstmann (00); Corrias, Dolbeault, Perthame, Zaag (04);

CHEMOTAXIS : Keller-Segel model

Theorem (dimensions $d \geq 2$) - (method of Sobolev inequalities)

(i) for $\|n^0\|_{L^{d/2}(R^d)}$ small, then there are global weak solutions,

(ii) these small solutions gain L^p regularity,

(iii) $\|n(t)\|_{L^\infty(R^d)} \rightarrow 0$ with the rate of the heat equation,

(iii) for $(\int |x|^2 n^0)^{(d-2)} < C \|n^0\|_{L^1(R^d)}^d$ with C small, there is blow-up in a finite time T^* .

CHEMOTAXIS : Keller-Segel model

In dimension 2, for Keller and Segel model :

$$\begin{cases} \frac{\partial}{\partial t} n(t, x) - \Delta n(t, x) + \operatorname{div}(n \chi \nabla c) = 0, & x \in \mathbb{R}^2, \\ -\Delta c(t, x) = n(t, x), \end{cases}$$

Theorem (dimension $d=2$) (Method of energy)

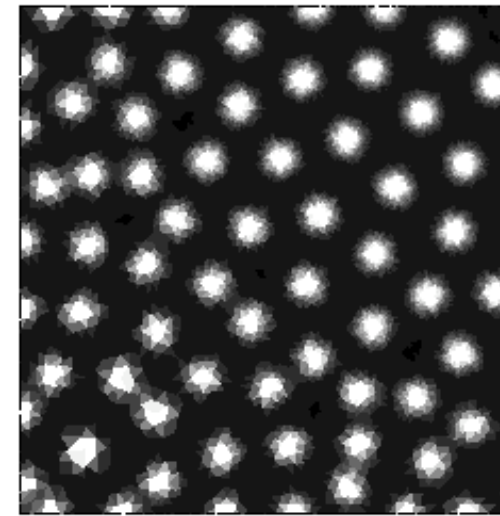
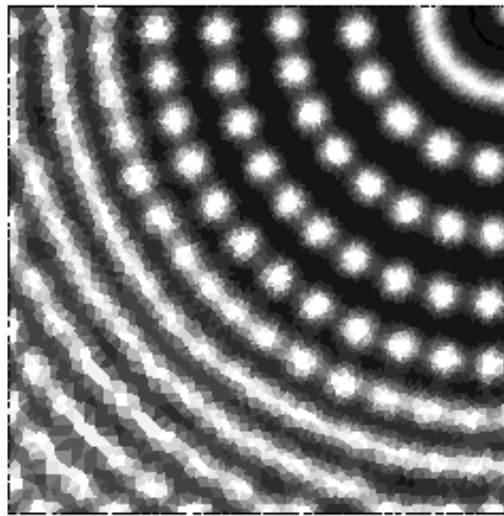
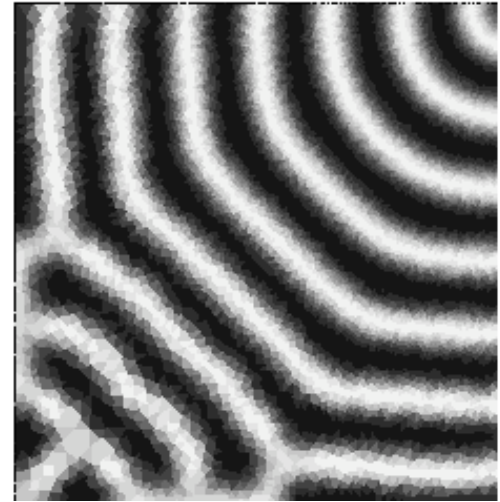
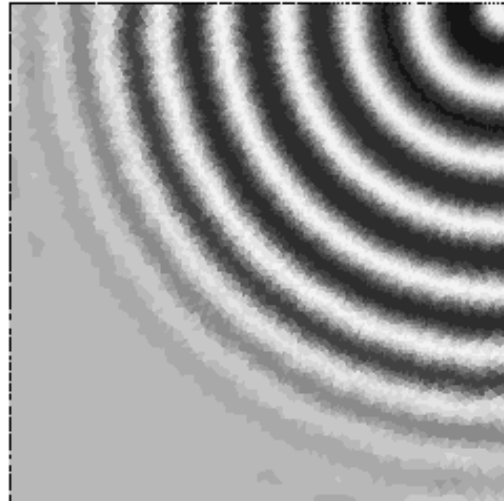
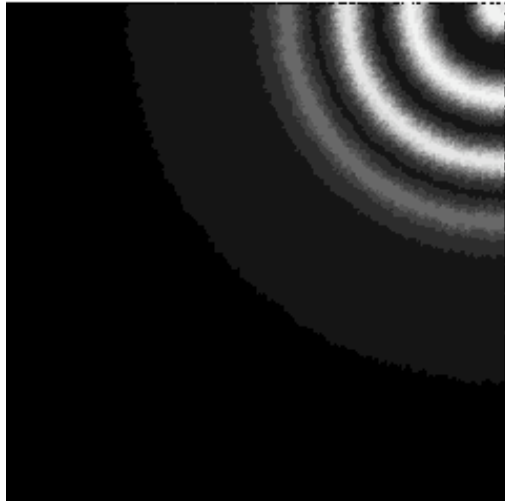
- (i) for $\|n^0\|_{L^1(\mathbb{R}^2)} < \frac{8\pi}{\chi}$, there are smooth solutions,
- (ii) for $\|n^0\|_{L^1(\mathbb{R}^2)} > \frac{8\pi}{\chi}$, there is creation of a singular measure (blow-up) in finite time.
- (iii) For radially symmetric solutions, blow-up means

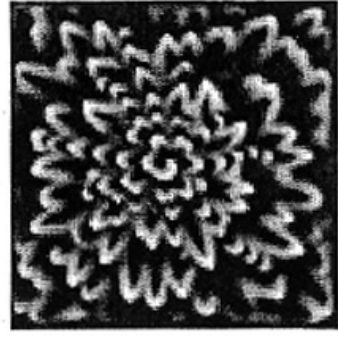
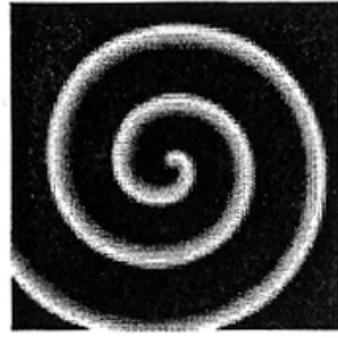
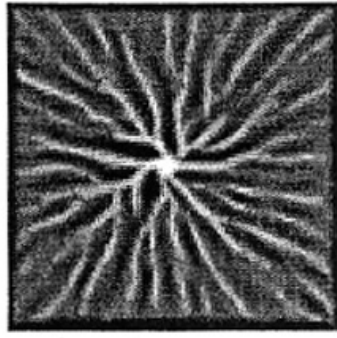
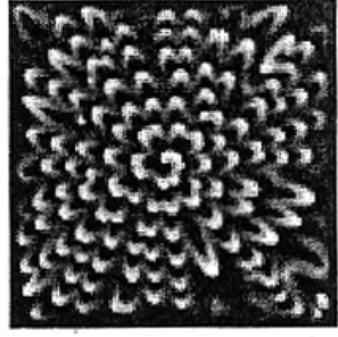
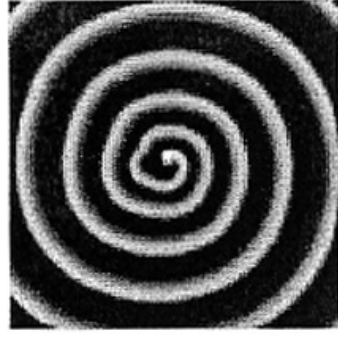
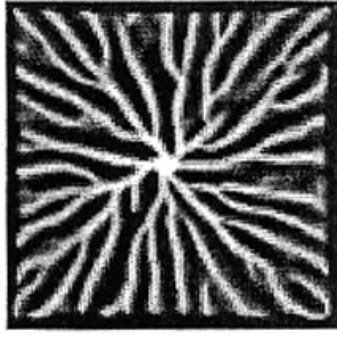
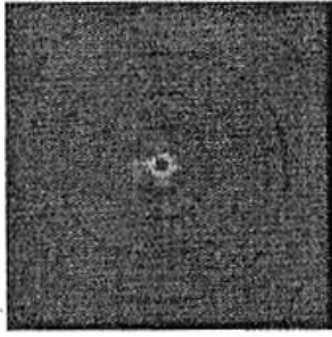
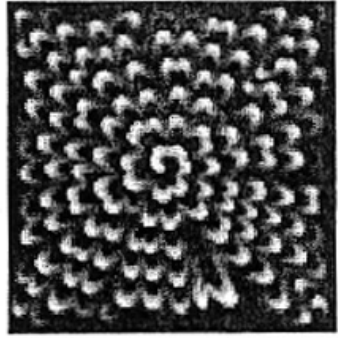
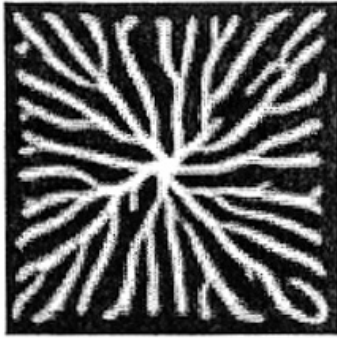
$$n(t) \approx \frac{8\pi}{\chi} \delta(x=0) + \text{Rem.}$$

CHEMOTAXIS : dimension 2

Interest : The Keller-Segel model seems successful to describe aggregation, some 'ring patterns', threshold are observed experimentally.

Mathematically a variety of methods are used ; Sobolev embeddings, energy methods, refined Hardy-Littlewood-Sobolev estimates (from 1994), convolution estimates, DeGiorgi method, asymptotic analysis...etc





(a)

(a)

Dentritic bacterial growth



Is chemotaxis involved in the bacterial movement of *Bacillus Subtilis* and in the development of such colonies? Another theory is proposed by Mimura

Dentritic bacterial growth

The simplest model is due to Mimura

$$\left\{ \begin{array}{l} \frac{\partial}{\partial t} n(t, x) - \Delta n = n \left(S - \frac{n}{(1+n)(1+S)} \right), \\ \frac{\partial}{\partial t} S(t, x) - \Delta S = -nS, \\ \frac{\partial}{\partial t} f(t, x) = n \frac{n}{(1+n)(1+S)} \end{array} \right.$$

More elaborated models are due to BenJacob, Kitsunezaki, Shikezada... and are based on the 'nutrient gradient' principle.

ANR Project with Institut de Génétique et Microbiologie (Paris-Sud), Ecole Polytechnique (M. Plapp)

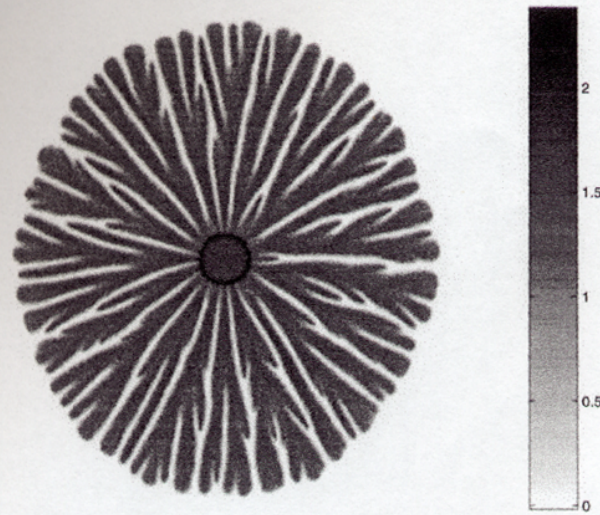


Fig. 28. 2D growth pattern ($b + s$) of the Kitsunzaki model with repulsive chemotactic signaling included. Parameters are: $\chi_{0r} = 1$, $D_r = 1$, $\Gamma_r = 0.25$, $\Omega_r = 0$, $A_r = 0.001$. Other parameters as in Fig. 19.

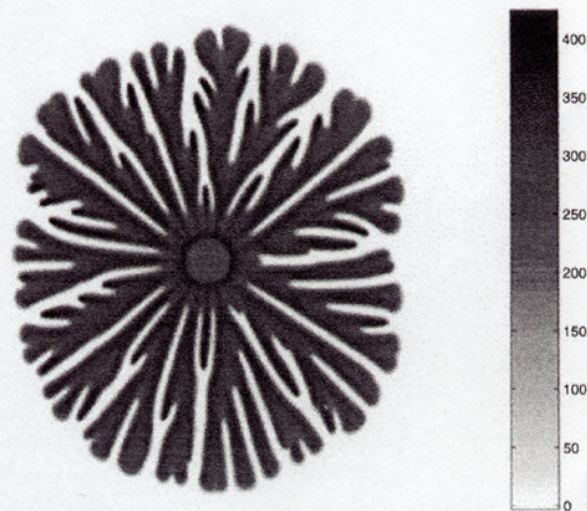
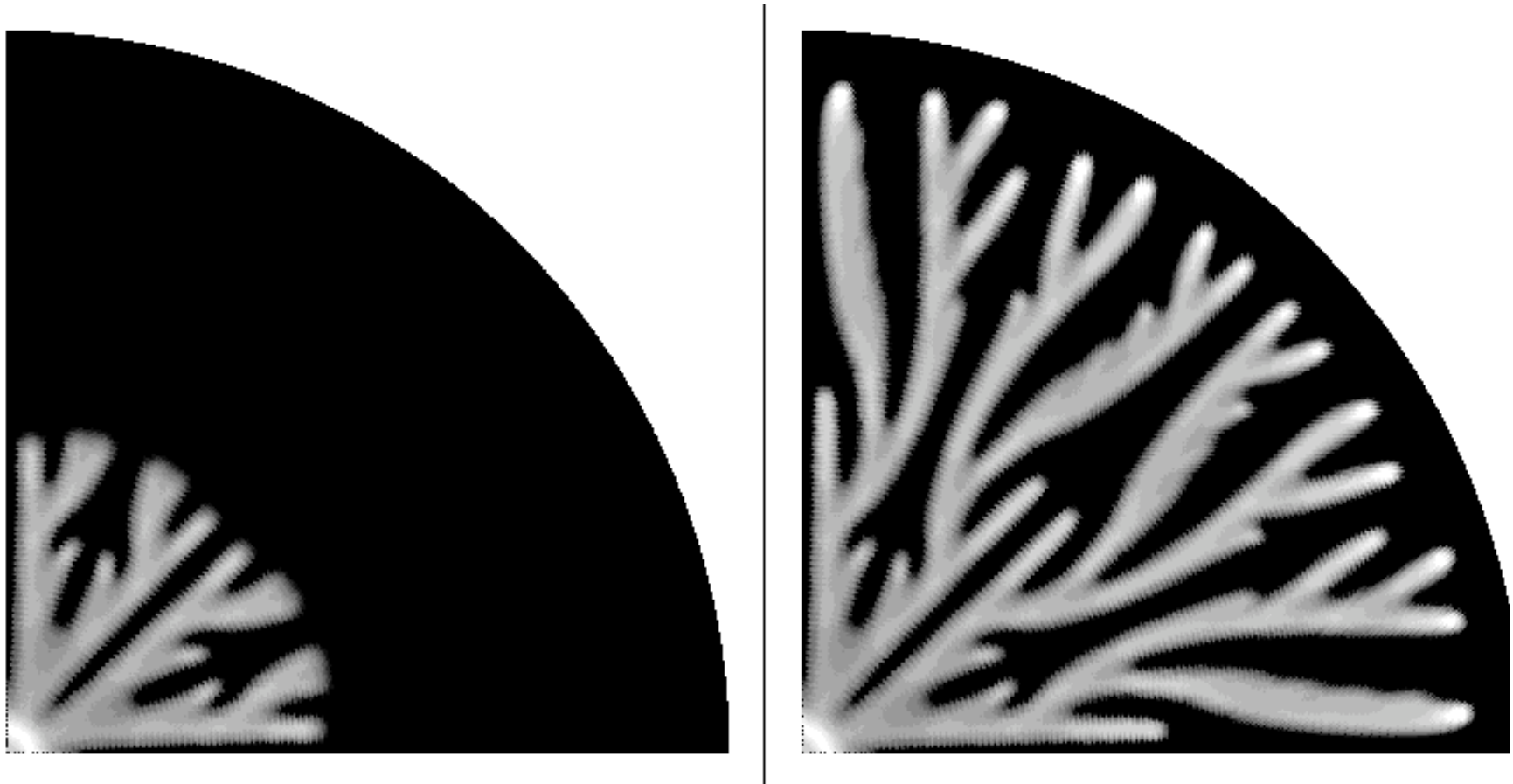
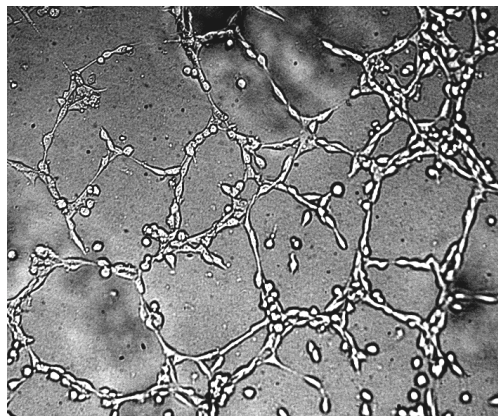
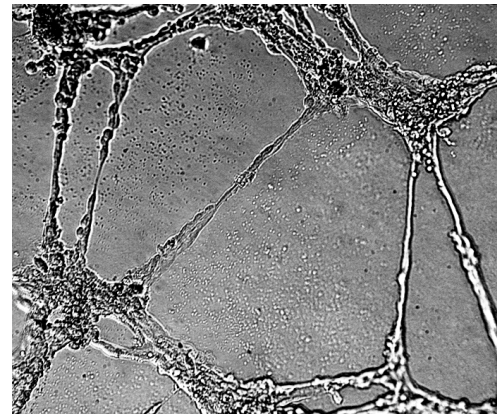
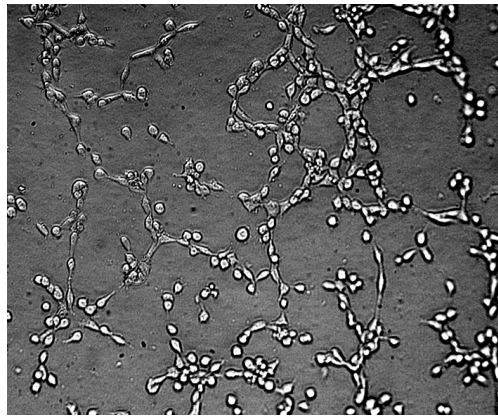
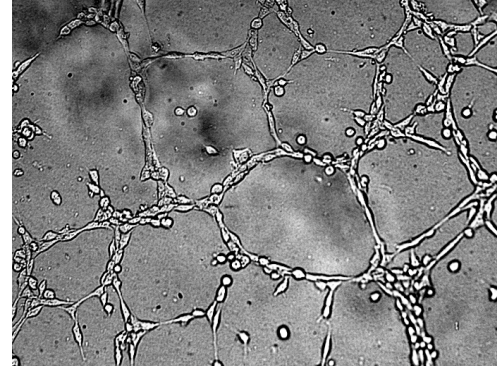
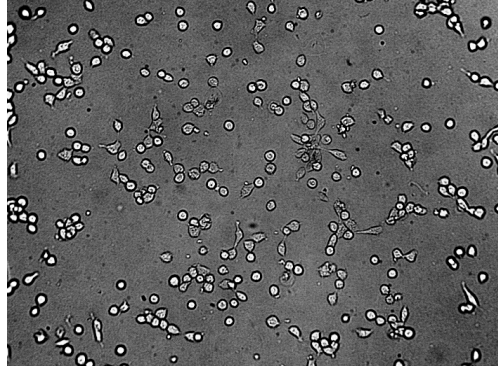


Fig. 29. 2D growth pattern ($b + s$) of the Mimura et al. model with food chemotaxis included. $\chi_{0f} = 0.06$. Other parameters as in Fig. 23a.



Simulation of Mimura's model by A. Marrocco (inria, bang)

Networks and hyperbolic models



**HBMEC SUR
MATRIGEL
T=0 ,2H,4H,6H,20H**

Networks and hyperbolic models

A group of Torino **Ambrosi, Gamba, Preziosi et al** proposed a *hydrodynamics model*

$$\begin{cases} \frac{\partial}{\partial t} n(t, x) + \operatorname{div}(n u) = 0, & x \in \mathbb{R}^2, \\ \frac{\partial}{\partial t} u(t, x) + u(t, x) \cdot \nabla u + \nabla n^\alpha = \chi \nabla c - \mu u, \\ \frac{\partial}{\partial t} c(t, x) - \Delta c(t, x) + \tau c(t, x) = n(t, x). \end{cases}$$

Networks and hyperbolic models

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Keller-Segel model can be viewed as a special case where the acceleration term is neglected

$$\frac{\partial}{\partial t} u(t, x) + u(t, x) \cdot \nabla u = 0.$$

Networks and hyperbolic models

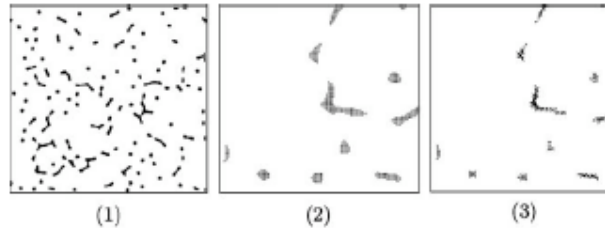


Fig. 4. Formation of network: (1) density and zoom on the left-bottom corner of (2) the density and (3) velocity field obtained with 50 cells/mm².

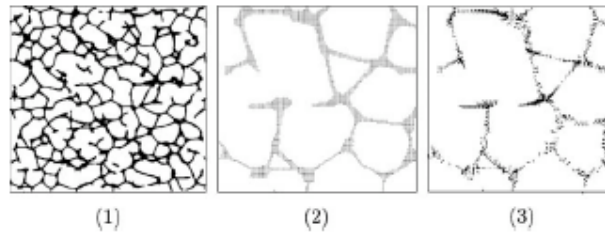
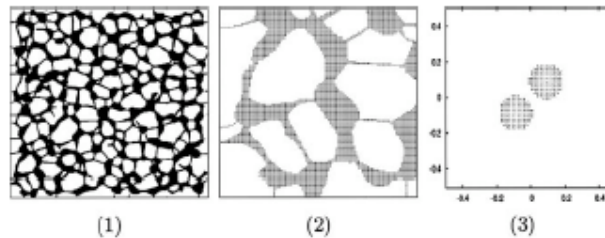
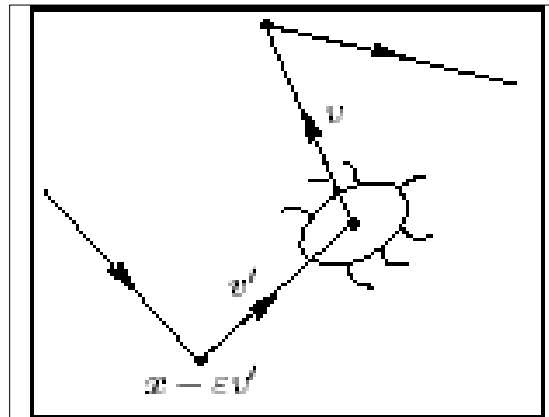


Fig. 5. Formation of network: (1) density and zoom on the left-bottom corner of (2) the density and (3) velocity field obtained with 100 cells/mm².



KINETIC MODELS

E. Coli is known (since the 80's) to move by run and tumble depending on the coordination of motors that control the flagella



See [Alt, Dunbar, Othmer, Stevens.](#)

KINETIC MODELS

Denote by $f(t, x, \xi)$ the density of cells moving with the velocity ξ .

$$\frac{\partial}{\partial t} f(t, x, \xi) + \underbrace{\xi \cdot \nabla_x f}_{\text{run}} = \underbrace{\mathcal{K}[f]}_{\text{tumble}},$$

$$\mathcal{K}[f] = \int K(c; \xi, \xi') f(\xi') d\xi' - \int K(c; \xi', \xi) d\xi' f,$$

$$-\Delta c(t, x) = n(t, x) := \int f(t, x, \xi) d\xi,$$

$$K(c; \xi, \xi') = k_-(c(x - \varepsilon \xi')) + k_+(c(x + \varepsilon \xi)).$$

Nonlocal, quadratic term on the right hand side for $k_{\pm}(\cdot, \xi, \xi')$ sublinear.

$x + \varepsilon \xi$ represents a (fundamental) memory term.

KINETIC MODELS

Theorem (Chalub, Markowich, P., Schmeiser)

Assume that $0 \leq k_{\pm}(c; \xi, \xi') \leq C(1 + c)$ then there is a GLOBAL solution to the kinetic model and

$$\|f(t)\|_{L^{\infty}} \leq C(t) [\|f^0\|_{L^1} + \|f^0\|_{L^{\infty}}]$$

-) Open question : Is it possible to prove a bound in L^{∞} when we replace the specific form of K by

$$0 \leq K(c; \xi, \xi') \leq \|c(t)\|_{L_{loc}^{\infty}} ?$$

-) Hwang, Kang, Stevens : $k(\nabla c(x - \varepsilon \xi'))$ or $k(\nabla c(x + \varepsilon \xi))$

KINETIC MODELS : diffusion limit

One can perform a parabolic rescaling based on the memory scale

$$\frac{\partial}{\partial t} f(t, x, \xi) + \frac{\xi \cdot \nabla_x f}{\varepsilon} = \frac{\mathcal{K}[f]}{\varepsilon^2},$$

$$\begin{aligned} \mathcal{K}[f] &= \int K(c; \xi, \xi') f' d\xi' - \int K(c; \xi', \xi) d\xi' f, \\ -\Delta c(t, x) &= n(t, x) := \int f(t, x, \xi) d\xi, \\ K(c; \xi, \xi') &= k_-(c(x - \varepsilon \xi')) + k_+(c(x + \varepsilon \xi)). \end{aligned}$$

Theorem (Chalub, Markowich, P., Schmeiser) With the same assumptions, as $\varepsilon \rightarrow 0$, then locally in time,

$$f_\varepsilon(t, x, \xi) \rightarrow n(t, x), \quad c_\varepsilon(t, x) \rightarrow c(t, x),$$

$$\begin{cases} \frac{\partial}{\partial t} n(t, x) - \operatorname{div}[D \nabla n(t, x)] + \operatorname{div}(n \chi \nabla c) = 0, \\ -\Delta c(t, x) = n(t, x). \end{cases}$$

and the transport coefficients are given by

$$D(n, c) = D_0 \frac{1}{k_-(c) + k_+(c)},$$

$$\chi(n, c) = \chi_0 \frac{k'_-(c) + k'_+(c)}{k_-(c) + k_+(c)}.$$

The drift (sensitivity) term $\chi(n, c)$ comes from the memory term.

Interpretation in terms of random walk : memory is fundamental.

Conclusion

Several open questions are

-) Is it possible to prove blow-up for

$$\|n^0\|_{L^{d/2}} \text{ large (without moment condition),}$$

-) Despite the knowledge of possible blow-up modalities, proofs are not rigorous,

-) More elaborate systems (see V. Calvez, A. Marrocco),

-) Mathematical theory for dendritic growth,

-) Mathematical models for dendritic growth,

-) Modeling : genome, networks, coefficients.