

Mathematical models for cell movement

Part I

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Overview

- Biological background
- Keller-Segel model
- Kinetic models
- Scaling up and down

Overview – Today

- Biological background
 - Cells move — for what ?
 - Life and death of *Dictyostelium discoideum*
- Choosing a problem: the initiation of the aggregation
- The Keller-Segel model.

Chemotaxis

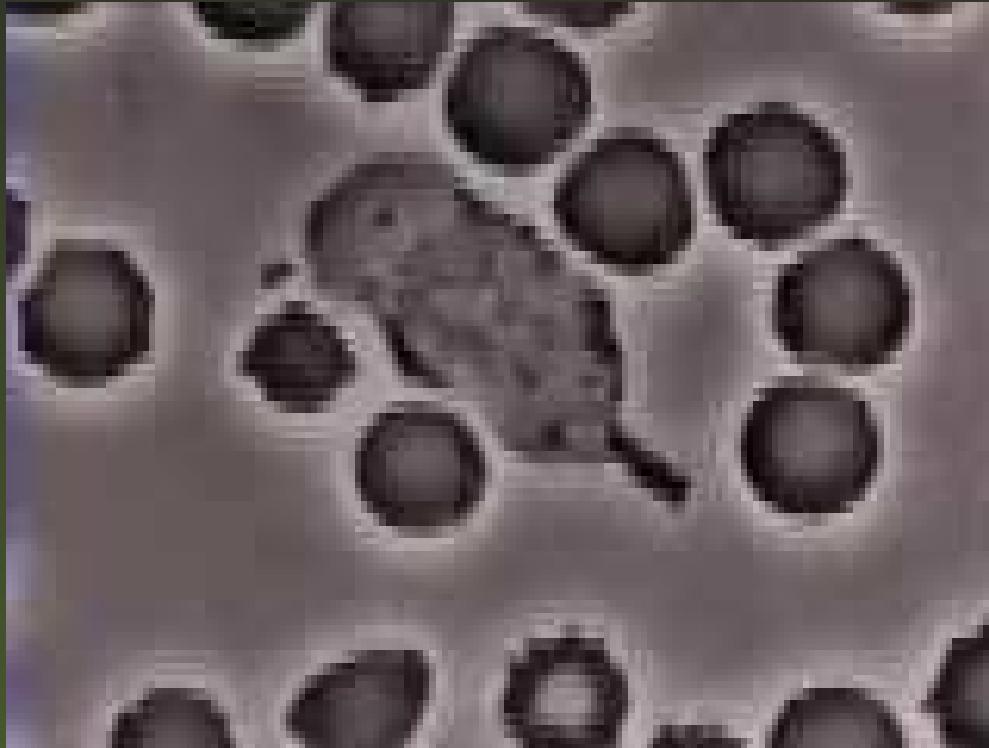


Figure 1: A neutrophil (white blood cell) chasing a bacterium. Film by Peter Devreotes.

Why do cells move?

- In general: looking for better places to live.
- Other reasons are:
 - immunology,
 - embryology and development,
 - aggregation.

One day in the life of *Dd*

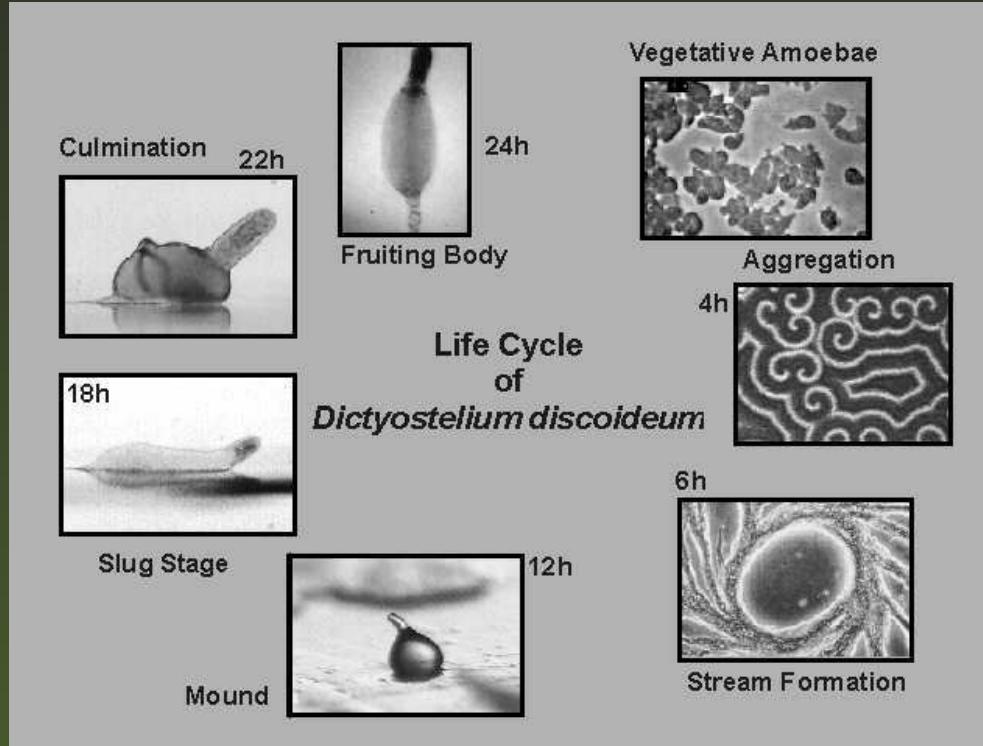


Figure 2: Life cycle of *Dictyostelium discoideum*. Picture made by Florian Seigert and Kees Wiejer (Zoologisches Institut München Ludwig-Maximilians-Universität München).

The initiation of the aggregation

The *Dictyostelium discoideum* moves toward higher concentrations of cAMP.

- Video 1 by Peter Devreotes.
- Video 2 by Peter Devreotes.

The formation of a core

Cells of Dd aggregates in a core.

- Video 1 by Kees Wiejer and Florian Seigert.

The slug phase

The aggregate starts to behave like a slug and migrates.

- Video 1 by Kees Wiejer and Florian Seigert.
- Video 2 by Kees Wiejer and Florian Seigert.
- Video 3 by Kees Wiejer and Florian Seigert.

The Culmination

The *slug* culminates with a spore on the top.

- Video 1 by Kees Wiejer and Florian Seigert.
- Video 2 by Rex Chisolm.

Why it is important to study the *Dd*?

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Many reasons:

- Cell motility,
- Cell communication,
- Cooperation among non-clonal individuals.

Keller-Segel model

“Initiation of Slime Mold Aggregation Viewed as an Instability”, Evelyn Keller and Lee Segel, *J. Theor. Biol.* (1970) **26**, 399–415.

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Variables:

- $\rho(x, t)$ = density of amoebas.

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- $S(x, t)$ = density of cAMP.
- $\eta(x, t)$ = density of phosphodiesterase.

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- $c(x, t)$ = density of a certain instable substance.

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We assume $n = 2$.

Keller-Segel model

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- cAMP is produced by the amoebas at a rate $f(S, \rho)$ per amoeba.
- cAMP is degraded by an extra-cellular enzyme (phosphodiesterase) at rate $g(S, \eta)$ by amoeba.
- cAMP and phosphodiesterase react making a new unstable compound C that decays immediately in phosphodiesterase and certain degenerated product.



Keller-Segel model

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$$\begin{aligned} J &= \text{current ,} \\ \phi &= \text{flux ,} \end{aligned}$$

$$J = D \nabla \phi .$$

- Amoebas concentration varies due to random diffusion and chemotaxis, in the positive direction of cAMP's gradient.

Keller-Segel model

For any substance with density a^i , we have a flux J^i and a creation/destruction term Q^i such that

$$\frac{\partial a^i}{\partial t} = Q^i - \nabla \cdot J^i .$$

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Mass conservation implies that

$$Q^\rho = 0 .$$

Keller-Segel model

In order to obtain Q^S , Q^η and Q^C we shall consider the production by the amoebas and the chemical reaction:

$$\begin{aligned} Q^S &= -k_1 S \eta + k_{-1} c + \rho f(S, \rho) , \\ Q^\eta &= -k_1 S \eta + (k_{-1} + k_2) c + \rho g(S, \eta) , \\ Q^c &= k_1 S \eta - (k_1 + k_2) c . \end{aligned}$$

Keller-Segel model

Fluxes are given by the following expressions:

$$J^\rho = -D_1 \nabla \rho + D_2 \nabla S ,$$

$$J^S = -D_S \nabla S ,$$

$$J^\eta = -D_\eta \nabla \eta ,$$

$$J^c = -D_c \nabla c .$$

Keller-Segel model

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Diffusion coefficients are:

$$D_i = D_i(\rho, S, \eta, c) .$$

Keller-Segel model

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$$\frac{\partial \rho}{\partial t} = \nabla \cdot (D_1 \nabla \rho) - \nabla \cdot (D_2 \nabla S) ,$$

$$\frac{\partial S}{\partial t} = -k_1 S \eta + k_{-1} c + \rho f(S, \rho) + D_S \Delta S ,$$

$$\frac{\partial \eta}{\partial t} = -k_1 S \eta + (k_{-1} + k_2) c + \rho g(S, \eta) + D_\eta \eta ,$$

$$\frac{\partial c}{\partial t} = k_1 S \eta - (k_1 + k_2) c + D_c \Delta c .$$

Keller-Segel model

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- The substance C is in chemical equilibrium:

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(Haldane's assumption)

- Enzyme concentration (in both free and bound forms) is constant

$$\eta + c = \eta_0 .$$

Keller-Segel model

Solving the system, we find

$$\eta = \frac{\eta_0}{1 + K\rho} ,$$

$$K = \frac{k_1}{k_{-1} + k_2} .$$

Keller-Segel model

We write again the system:

$$\begin{aligned}\frac{\partial \rho}{\partial t} &= -\nabla \cdot (D_1 \nabla \rho) + \nabla(D_2 \nabla S) , \\ \frac{\partial S}{\partial t} &= -k(S)S + \rho f(S, \rho) + D_S \Delta S ,\end{aligned}$$

where $k(S) = \eta_0 k_2 K / (1 + KS)$.

Keller-Segel model

We resume the model

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Keller-Segel model

We resume the model

- Amoebas have a random and a chemotactical movement.
- Chemoattractant S diffuses, is created by the amoebas at rate f and decays at rate k .

Keller-Segel model

From now on, we call the following system the Keller-Segel system

$$\begin{aligned}\frac{\partial \rho}{\partial t} &= \nabla \cdot (D \nabla \rho - \chi \rho \nabla S) , \\ \frac{\partial S}{\partial t} &= D_S \Delta S + \varphi(\rho, S) ,\end{aligned}$$

where $D = D(S, \rho)$, $\chi = \chi(S, \rho)$ e $D_S = D_S(S, \rho)$.

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where $D = D(S, \rho)$, $\chi = \chi(S, \rho)$ e $D_S = D_S(S, \rho)$.

$\varphi(\rho, S)$ describes production and decay of the chemoattractant. Typically

$$\varphi = \alpha \rho - \beta S .$$

The Keller-Segel Model

- ρ =density of cells.
- S =density of chemo-attractant.

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$$\begin{aligned}\frac{\partial \rho}{\partial t} &= \nabla \cdot (D \nabla \rho - \chi \rho \nabla S) , \\ \frac{\partial S}{\partial t} &= D_0 \Delta S + \alpha \rho - \beta S ,\end{aligned}$$

where $D = D(S, \rho)$, $\chi = \chi(S, \rho)$, $D = D(S, \rho)$ e
 $D_0 = D_0(S, \rho)$, $\alpha > 0$, $\beta \geq 0$.

The Keller-Segel Model

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- The fast-diffusion limit: elliptic equation for $S.$

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Some important cases:

- The *classical* Keller-Segel model: $\chi, D, D_0 = \text{const.}$
- The no-decay case: $\beta = 0.$
- The fast-diffusion limit: elliptic equation for $S.$

With these assumption the equation for S is

$$\Delta S = -\rho .$$

The Keller-Segel Model

Finally, consider for $x \in \mathbb{R}^2$

$$\begin{aligned}\partial_t \rho &= \nabla \cdot (\nabla \rho - \chi \rho \nabla S) , \\ \Delta S &= -\rho ,\end{aligned}$$

with

$$\rho(\cdot, 0) = \rho^I .$$

The Keller-Segel Model

Properties:

The Keller-Segel Model

Theorem. (*Perthame, Dolbeaut, 2004*) Consider a solution of the KS system, such that

- $\int_{\mathbb{R}^2} |x|^2 \rho(x, t) dx < \infty,$
- $\int_{\mathbb{R}^2} \frac{1+|x|}{|x-y|} \rho(y, t) dy \in L^\infty((0, T) \times \mathbb{R}^2).$

Then

$$\frac{d}{dt} \int_{\mathbb{R}^2} |x|^2 \rho(x, t) dx = 4M \left(1 - \frac{\chi M}{8\pi} \right) .$$

The Keller-Segel Model

Proof:

$$S(x, t) = -\frac{1}{2\pi} \int_{\mathbb{R}^2} \log|x - y| \rho(y, t) dy ,$$

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$$\begin{aligned} S(x, t) &= -\frac{1}{2\pi} \int_{\mathbb{R}^2} \log|x-y| \rho(y, t) dy , \\ \nabla S(x, t) &= -\frac{1}{2\pi} \int_{\mathbb{R}^2} \frac{x-y}{|x-y|^2} \rho(y, t) dy . \end{aligned}$$

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$$\int_{\mathbb{R}^2} \rho(x) \rho(y) \frac{x \cdot (x-y)}{|x-y|^2} dy dx = - \int_{\mathbb{R}^2} \rho(x) \rho(y) \frac{y \cdot (x-y)}{|x-y|^2} dy dx =$$

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The Keller-Segel Model

$$\frac{1}{2} \frac{d}{dt} \int_{\mathbb{R}^2} |x|^2 \rho(x) dx = \frac{1}{2} \int_{\mathbb{R}^2} |x|^2 \nabla \cdot (\nabla \rho - \chi \rho \nabla S) dx$$

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The Keller-Segel Model

Theorem. (*Perthame, Dolbeault, 2004*) Consider the KS system such that

- $M < 8\pi/\chi$,
- $\rho^I \in L^1(\mathbb{R}^2, (1 + |x|^2)dx)$.

Then, the system has a weak solution such that

- $(1 + |x|^2 + \log \rho)\rho \in L_{\text{loc}}^\infty(\mathbb{R}^+, L^1(\mathbb{R}^2))$,
- $\int_0^\infty \int_{\mathbb{R}^2} \rho |\nabla \log \rho - \chi \nabla S|^2 dx dt < \infty$,
- $\rho, \nabla \sqrt{\rho} \in L^2([0, T] \times \mathbb{R}^2), \forall T > 0$.

The Keller-Segel Model

Proof: The proof consists in three steps:

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- Regularization of solutions,
- Estimations,
- Limits.

The Keller-Segel Model

Step 1: (Regularization) Consider

$$K_\varepsilon(z) = \begin{cases} -\frac{1}{2\pi} \log |z| & \text{if } |z| > \varepsilon, \\ -\frac{1}{2\pi} \log \varepsilon & \text{if } |z| \leq \varepsilon. \end{cases}$$

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This means that, from

$$S(x, t) = -\frac{1}{2\pi} \int_{\mathbb{R}^2} \log |x - y| \rho(y, t) dy$$

we change to

$$S_\varepsilon(x, t) = \int_{\mathbb{R}^2} K_\varepsilon(x - y) \rho(y, t) dy = (K_\varepsilon * \rho_\varepsilon)(x, t).$$

The Keller-Segel Model

Step 2: (Estimations) Equation:

$$\partial_t \rho_\varepsilon = \nabla \cdot (\nabla \rho_\varepsilon - \chi \rho_\varepsilon \nabla (K_\varepsilon * \rho_\varepsilon)) .$$

$$\frac{d}{dt}$$

The Keller-Segel Model

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Then:

$$\frac{d}{dt} \int_{\mathbb{R}^2} |x|^2 \rho_\varepsilon(x, t) dx = 4M - \frac{\chi}{2\pi} \int_{|y-x|>\varepsilon} \rho_\varepsilon(x, t) \rho_\varepsilon(y, t) dx dy$$

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We prove similar ε -independent bounds and take the limit $\varepsilon \rightarrow 0$.

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This is in agreement with the fact that aggregation occurs only if the initial density of Dd is above certain threshold.

The Keller-Segel Model

Corollary. *In the two dimensional case, we have:*

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What happens if $M = 8\pi/\chi$?

The Keller-Segel Model

Suppose general dimension n .

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Theorem. (*Corrias, Perthame, Zaag, 2004*) *There exists a constant K such that*

$$\|\rho^I\|_{L^{n/2}(\mathbb{R}^n)} \leq K$$

then the KS system has a global in time weak solution such that

$$\|\rho(\cdot, t)\|_{L^p(\mathbb{R}^n)} \leq \|\rho^I\|_{L^p(\mathbb{R}^n)}, \quad \max\left\{1, \frac{n}{2} - 1\right\} \leq p \leq \frac{n}{2},$$

$$\|\rho(\cdot, t)\|_{L^p(\mathbb{R}^n)} \leq C(t, K, \|\rho^I\|_{L^p(\mathbb{R}^n)}), \quad \frac{n}{2} < p \leq \infty.$$

The Keller-Segel Model

Theorem. (*Corrias, Perthame, Zaag, 2004*) Suppose that $n \geq 3$ and

$$\int_{\mathbb{R}^n} \frac{|x|^2}{2} \rho^I(x) dx \leq CM^{n/(n-2)} ,$$

and assume that

$$M \geq M_0$$

for some $M_0 > 0$. Then the KS system has no global solution with fast decay.

Keller-Segel Models

Consider the following Keller-Segel model (with *prevention of overcrowding*) (Hillen-Painter model):

$$\begin{aligned}\partial_t \rho &= \nabla \cdot (\nabla \rho - \chi \beta(\rho) \rho \nabla S) \\ \Delta S &= -\rho ,\end{aligned}$$

where

$$\beta(\rho) = 0 , \quad \rho \geq \bar{\rho} > 0 .$$

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Theorem. (Hillen, Painter, 2002) *Solutions of the HP model exist globally.*

Keller-Segel Models

Proof: Consider:

$$\begin{aligned}\mathcal{J}_+ &= \{x | \rho(x, t) > \bar{\rho}\} , \\ \mathcal{J}_0 &= \{x | \rho(x, t) = \bar{\rho}\} , \\ \mathcal{J}_- &= \{x | \rho(x, t) < \bar{\rho}\} ,\end{aligned}$$

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and define

$$\rho^+(x, t) = \begin{cases} \rho(x, t) - \bar{\rho} & \text{if } x \in \mathcal{J}_+ , \\ 0 & \text{otherwise.} \end{cases}$$

Keller-Segel Models

$$\frac{1}{2} \frac{d}{dt} \|\rho^+(\cdot, t)\|_{L^2(\mathbb{R}^n)}^2 = \left(\int_{\mathcal{J}_+} + \int_{\mathcal{J}_0} + \int_{\mathcal{J}_-} \right) \rho^+ \rho_t^+$$

Keller-Segel Models

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Keller-Segel Models

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If $\|\rho^I\|_{L^\infty(\mathbb{R}^n)} > \bar{\rho}$, then in a neighbourhood of a point x such that $\rho(x, t) > \bar{\rho}$, the equation is

$$\partial_t \rho = \Delta \rho$$

and the maximum principle holds.

Keller-Segel Models

Define the *non-local* gradient

$$\overset{\circ}{\nabla}_R f(x, t) = \frac{1}{\omega_{n-1} R^{n-1}} \int_{S^{n-1}} f(x + yR) dy .$$

Keller-Segel Models

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Then the Hillen-Schmeiser-Painter model

$$\partial_t \rho = \nabla \cdot \left(\nabla \rho - \chi \rho \overset{\circ}{\nabla}_R S \right) ,$$

has global existence of solutions.

Keller-Segel Models

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or more generally:

$$\beta_\mu(\rho) = \frac{1}{\mu} Q(\mu\rho) ,$$

$$Q(y) \approx y - \alpha y^2 , \quad y \rightarrow \infty ,$$

$$\lim_{y \rightarrow \infty} Q(y) < \infty .$$

Keller-Segel Models

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More precisely:

- For $t < T$, $\lim_{\mu \rightarrow 0} \rho_\mu = \rho_0$.
- This cannot be extended after T because ρ_0 no longer exists (T is the blow up time).
- For any $\mu > 0$, ρ_μ exists for any time t .

Keller-Segel Models

For any $\mu > 0$ the aggregation region is of order $\sqrt{\mu}$. We consider each of the N aggregates as a point particles, $x_j, j = 1, \dots, N$, and consider a regular remainder which is important far from these points:

$$\rho(x, t) \approx \sum_{j=1}^N M_j(t) \delta(x - x_j(t)) + u_{\text{reg}}(x, t) .$$

Keller Segel Models

The chemoattractant concentration is given by

$$S(x, t) \approx -\frac{1}{2\pi} \sum_{j=1}^N M_j(t) \log(|x - x_j(t)|) + S_{\text{reg}}(x, t) ,$$

where

$$S_{\text{reg}} = -\frac{1}{2\pi} \int_{\mathbb{R}^2} \log(|x - y|) u_{\text{reg}}(y, t) dy .$$

Keller-Segel Models

Theorem. (*Velazquez, 2004*) Consider a solution of the V model. If we impose the previous Ansatz, then

$$\frac{\partial \rho_{\text{reg}}}{\partial t} = \Delta \rho_{\text{reg}} + \frac{1}{2\pi} \sum_{j=1}^N M_j(t) \frac{(x - x_j(t))}{|x - x_j(t)|^2} \cdot \nabla \rho_{\text{reg}} \\ - \nabla (\rho_{\text{reg}} \nabla S_{\text{reg}}) ,$$

$$S_{\text{reg}}(x, t) = -\frac{1}{2\pi} \int_{\mathbb{R}^2} \log(|x - y|) \rho_{\text{reg}}(y, t) dy ,$$

$$\dot{x}_i(t) = \Gamma(M(t)) A_i(t) ,$$

Keller-Segel Models

$$\begin{aligned} A_i(t) &= - \sum_{j=1, j \neq i}^N \frac{M_j(t)}{2\pi} \frac{(x_i(t) - x_j(t))}{|x_i(t) - x_j(t)|^2} \\ &\quad + \nabla S_{\text{reg}}(x_i(t), t) , \\ \frac{dM_i(t)}{dt} &= u_{\text{reg}} M_i(t) , \end{aligned}$$

where $\Gamma(M)$ is a function of $M > 8\pi$.

Keller-Segel Models

After certain time, mass will be only in the aggregates, that will interact as

$$\begin{aligned}\dot{x}_i(t) &= \frac{\Gamma(M_i(t))M_j(t)}{2\pi} \cdot \frac{(x_i(t) - x_j(t))}{|x_i(t) - x_j(t)|^2}, \\ \dot{x}_j(t) &= \frac{\Gamma(M_j(t))M_i(t)}{2\pi} \cdot \frac{(x_j(t) - x_i(t))}{|x_j(t) - x_i(t)|^2}\end{aligned}$$

and after certain time they will coalesce in a single aggregate.

Keller-Segel Models

the function $\Gamma(M)$ has the following properties:

$$0 < \Gamma(M) < 1 ,$$

$$\lim_{M \rightarrow 8\pi} \Gamma(M) = 1 ,$$

$$\lim_{M \rightarrow \infty} \Gamma(M) = 0 .$$

The Keller-Segel Model

Is the extension of the Keller-Segel model made by Velazquez general?