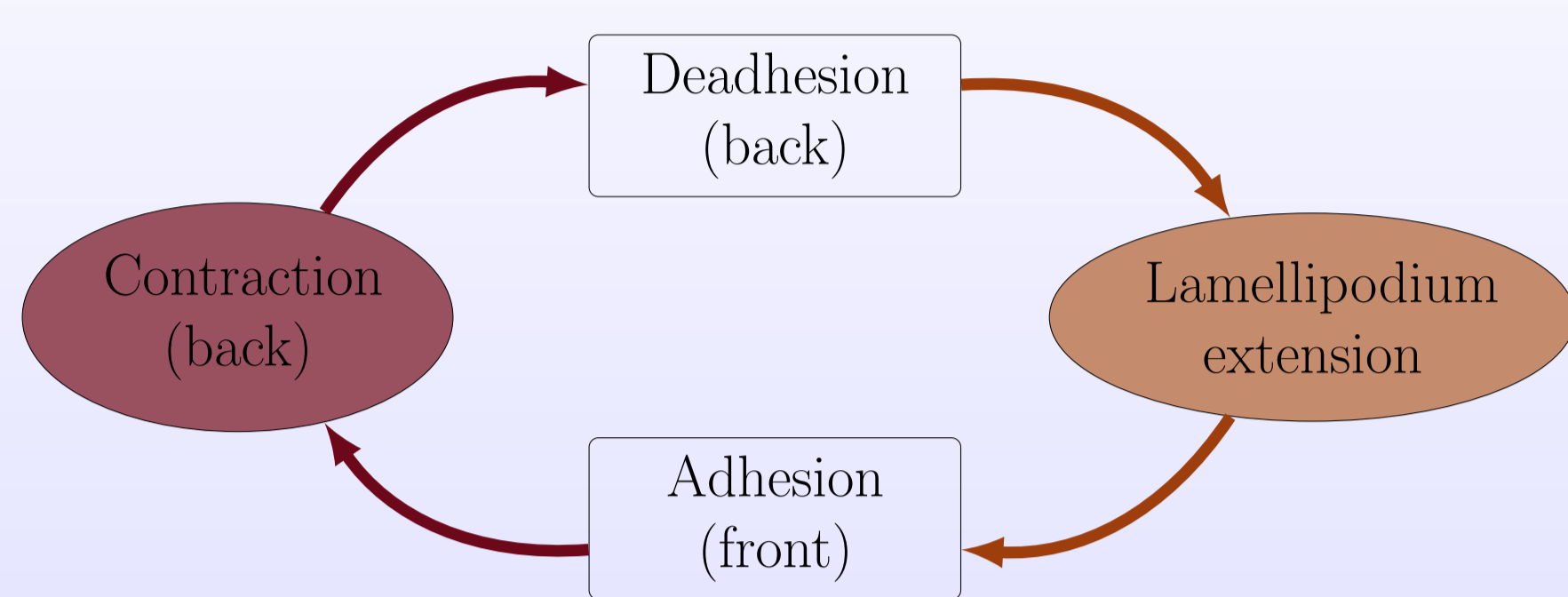


Biological Context

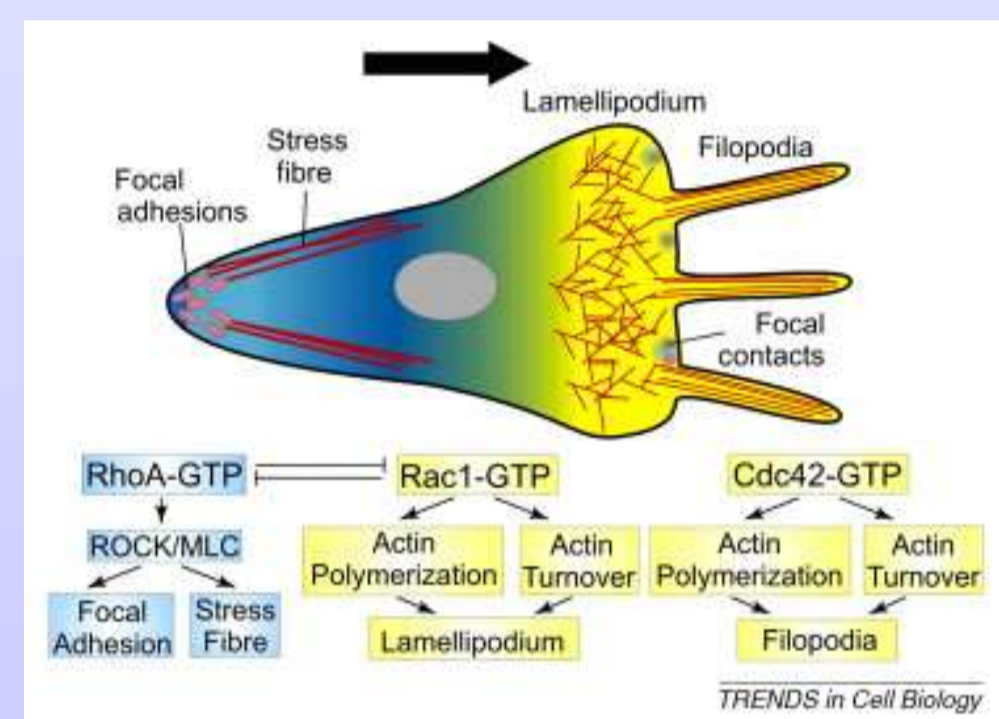
Cell migration is a complex biological process involving a succession of different events :



Many proteins and components of the cell are involved in cell migration. We focus on three important **actors** that act directly on cell motility :

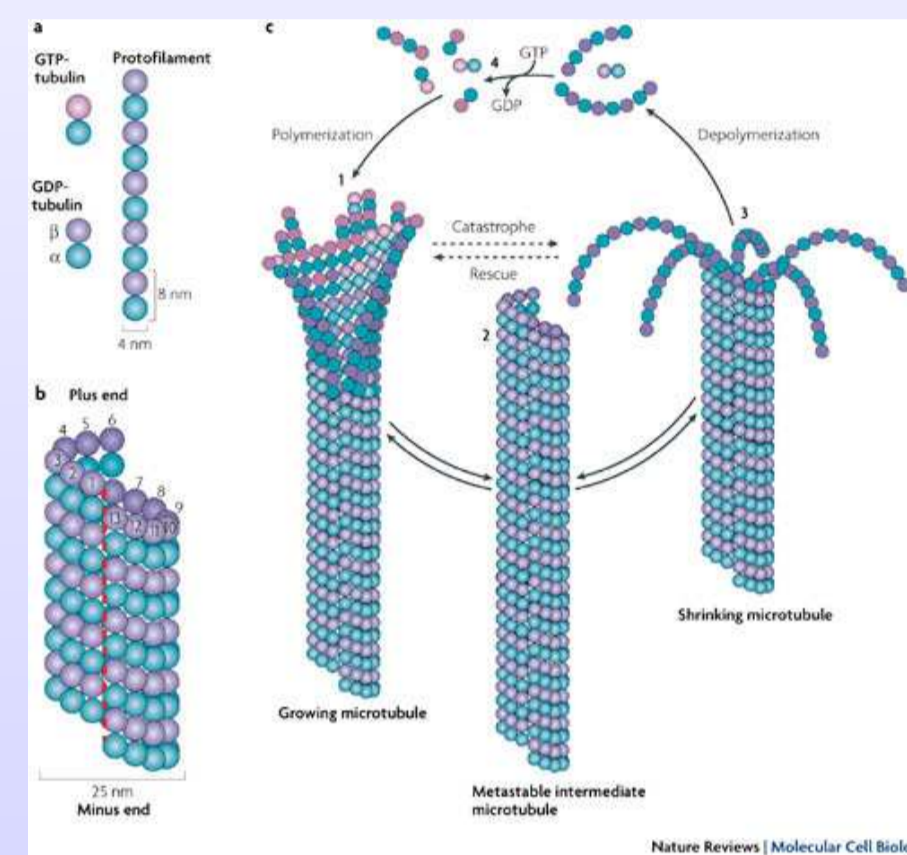
- **Actin**. Part of cytoskeleton, Creation of lamellipodium, Contraction.
- **Rac**. Encourage lamellipodium extension.
- **Rho**. Promote contraction.

Among those actors, Rac and Rho can be active or inactive inside the cytoplasm. Their activity depends strongly on **microtubules**.



Microtubules Dynamics

Microtubules (MT) are part of the cytoskeleton, constituted of long tubes polymers of tubulin.



MT have a highly dynamic behavior called **dynamic instability**. Their plus ends alternate between phases :

- polymerization
- depolymerization

Role of Microtubules

Cell Division :

Microtubules play a crucial role in **mitosis**. The mitotic spindle, that **segregate the chromosomes** during mitosis is mainly composed of microtubules.

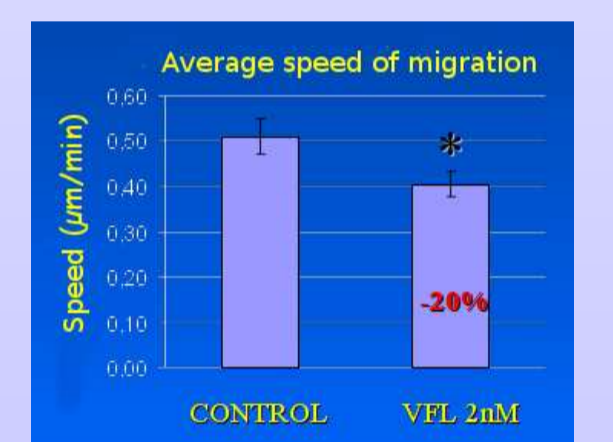
Transport :

Motor protein, like dynein or kinesin can attach to microtubules. They are involved in vesicles and organelles transportation throughout the cytoplasm.

Cell Migration :

Microtubules have an effect on proteins that regulate migration. Their dynamic activity regulate the activation or inactivation of Rac and Rho proteins. **The polymerization of the microtubule activates Rac and the depolymerization activates Rho.**

It has been shown that microtubules can be a target for anticancer therapies. In particular during angiogenesis and metastatic process, even at low doses.



One protein Model

Variables :

u -velocity ; p -pressure ; Rac -concentration of active Rac ; \overline{Rac} -concentration of inactive Rac

Mechanical Model :

$$-\mu\Delta u + \nabla p = F_{el} + F_{net}, \quad x \in \mathbb{R}^2$$

$$\nabla \cdot u = 0$$

Biochemical Model :

$$\frac{\partial Rac}{\partial t} + u \cdot \nabla Rac - D_{Rac} \Delta Rac = g(Rac) \overline{Rac} - \delta Rac, \quad x \in \Omega(t), t > 0$$

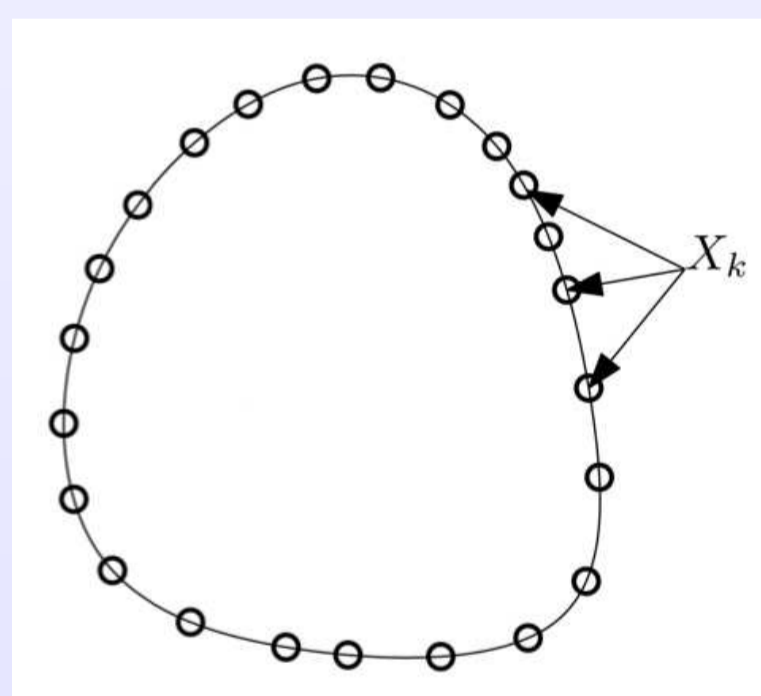
$$\frac{\partial \overline{Rac}}{\partial t} + u \cdot \nabla \overline{Rac} - D_{\overline{Rac}} \Delta \overline{Rac} = -g(Rac) \overline{Rac} + \delta Rac$$

$$g(Rac) = \frac{\tau_{Rac \rightarrow Rac}}{\tau_{Rac \rightarrow Rac} + \frac{\tau_{Rac}^2}{K^2 + Rac^2}}$$

Interface Representation :

We use a **Lagrangian Marker Points(LMP)** method, based on a parametrization of the interface.

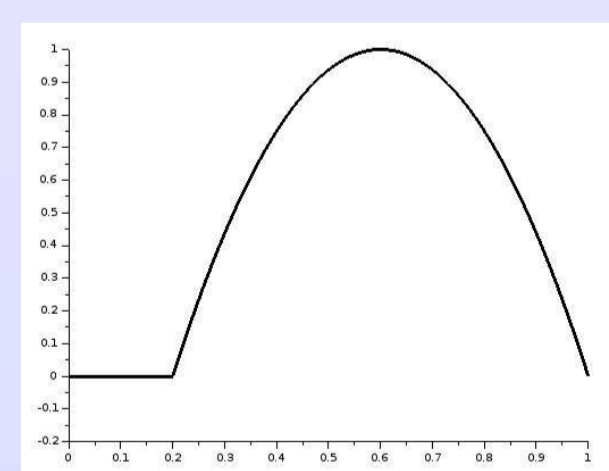
$$\frac{\partial X}{\partial t}(s, t) = u(X(s, t), t)$$



Forces : F_{el} , elastic force ; F_{net} , protrusion force

$$F_{el} = \frac{\partial}{\partial s} \left[T_0 \left(\left| \frac{\partial X}{\partial s} \right| - 1 \right) \frac{\partial X}{\partial s} \right]$$

$$F_{net} = h(Rac) \left(\frac{\partial X}{\partial s} \right)^\perp$$



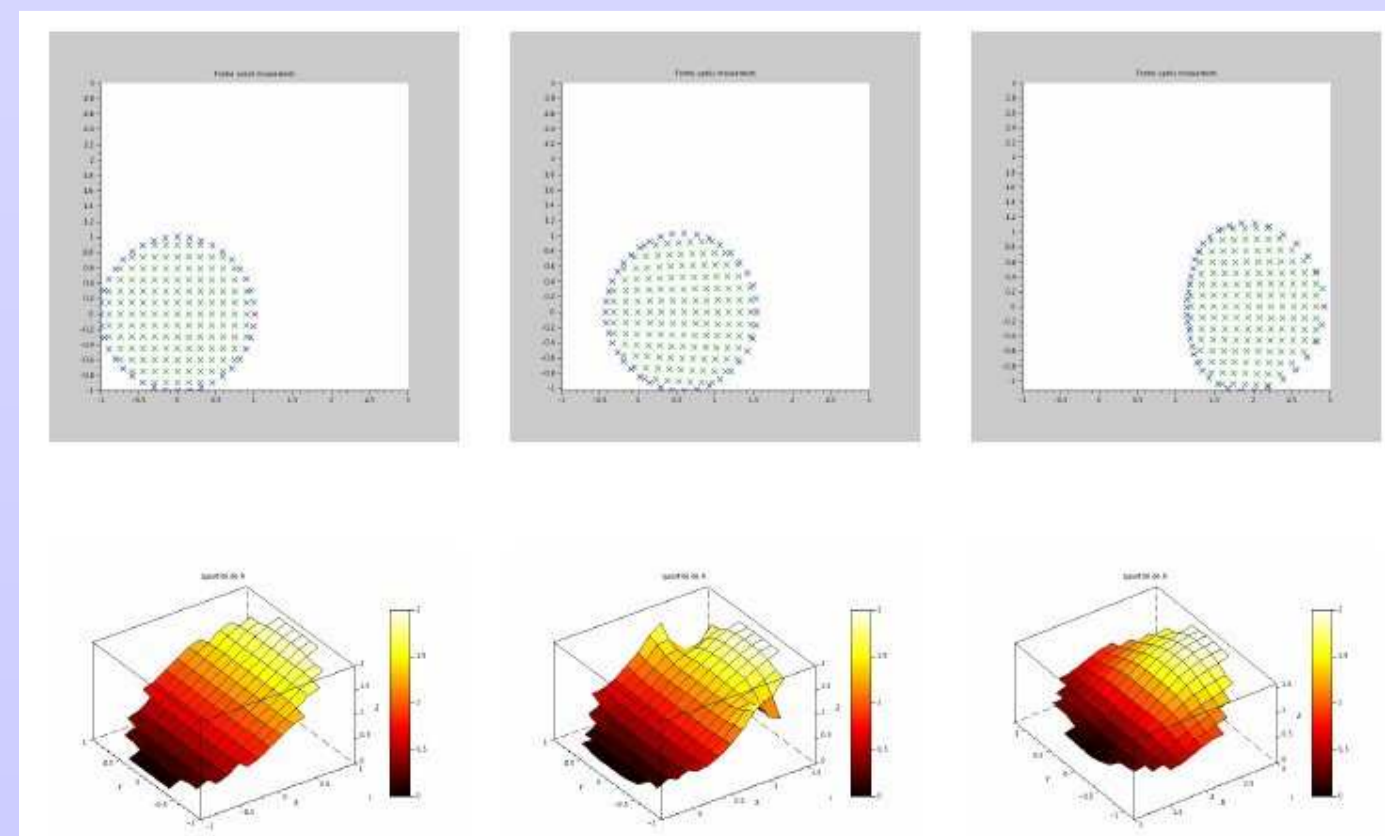
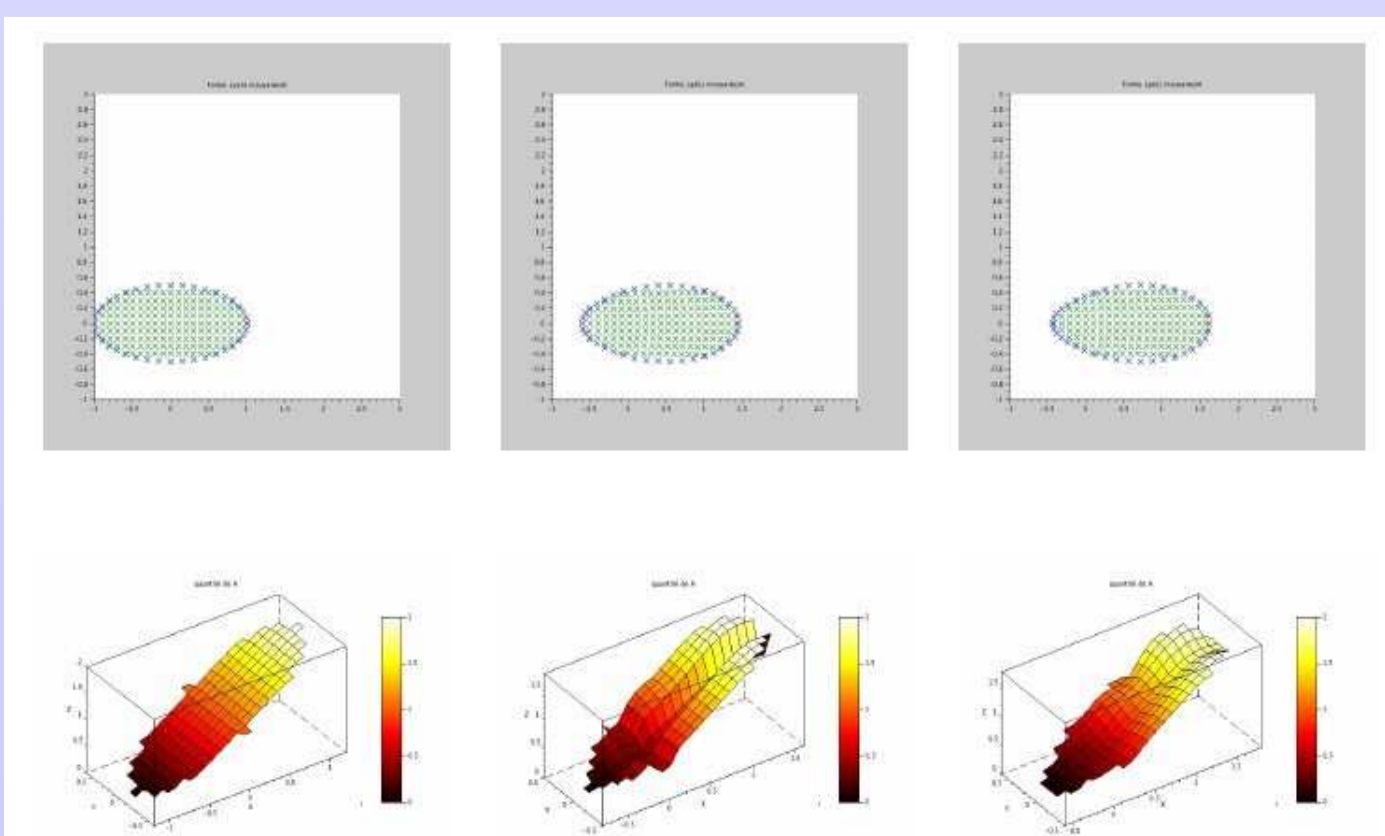
Function h

Numerical Results :

Difficulties :

- The cell is a moving domain
- Interpolation Lagrangian-Eulerian
- Limitation on the deformation of the membrane

Methods : Stokeslet - Finite Volume - Euler Scheme



Two Protein Model with MT regulation

Variables :

u -velocity ; p -pressure ; Rac -concentration of active Rac ; \overline{Rac} -concentration of inactive Rac ; Rho -concentration of active Rho ; \overline{Rho} -concentration of inactive Rho ; MT_i -plus end of MT number i ; L_i -length of MT number i ; Tub -concentration of Tubulin

Mechanical Model :

$$-\mu\Delta u + \nabla p = F_{el} + F_{net}, \quad x \in \mathbb{R}^2$$

$$\nabla \cdot u = 0$$

Biochemical Model :

$$\frac{\partial Rac}{\partial t} + u \cdot \nabla Rac - D_{Rac} \Delta Rac = \sum_i [g(Rac) k_0(x) \overline{Rac} - \tau_{Rac \rightarrow \overline{Rac}} (1 - k_0(x)) Rho Rac] \mathbf{1}_{B(MT_i, d_{MT})}$$

$$\frac{\partial \overline{Rac}}{\partial t} + u \cdot \nabla \overline{Rac} - D_{\overline{Rac}} \Delta \overline{Rac} = - \sum_i [g(Rac) k_0(x) \overline{Rac} - \tau_{Rac \rightarrow \overline{Rac}} (1 - k_0(x)) Rho Rac] \mathbf{1}_{B(MT_i, d_{MT})}$$

$$\frac{\partial Rho}{\partial t} + u \cdot \nabla Rho - D_{Rho} \Delta Rho = \sum_i [g(Rho) (1 - k_0(x)) \overline{Rho} - \tau_{Rho \rightarrow \overline{Rho}} k_0(x) Rho Rac] \mathbf{1}_{B(MT_i, d_{MT})}$$

$$\frac{\partial \overline{Rho}}{\partial t} + u \cdot \nabla \overline{Rho} - D_{\overline{Rho}} \Delta \overline{Rho} = - \sum_i [g(Rho) (1 - k_0(x)) \overline{Rho} - \tau_{Rho \rightarrow \overline{Rho}} k_0(x) Rho Rac] \mathbf{1}_{B(MT_i, d_{MT})}$$

$$\frac{\partial Tub}{\partial t} + u \cdot \nabla Tub - D_{Tub} \Delta Tub = - \sum_i d \frac{\partial L_i}{\partial t} \delta_0(x - MT_i)$$

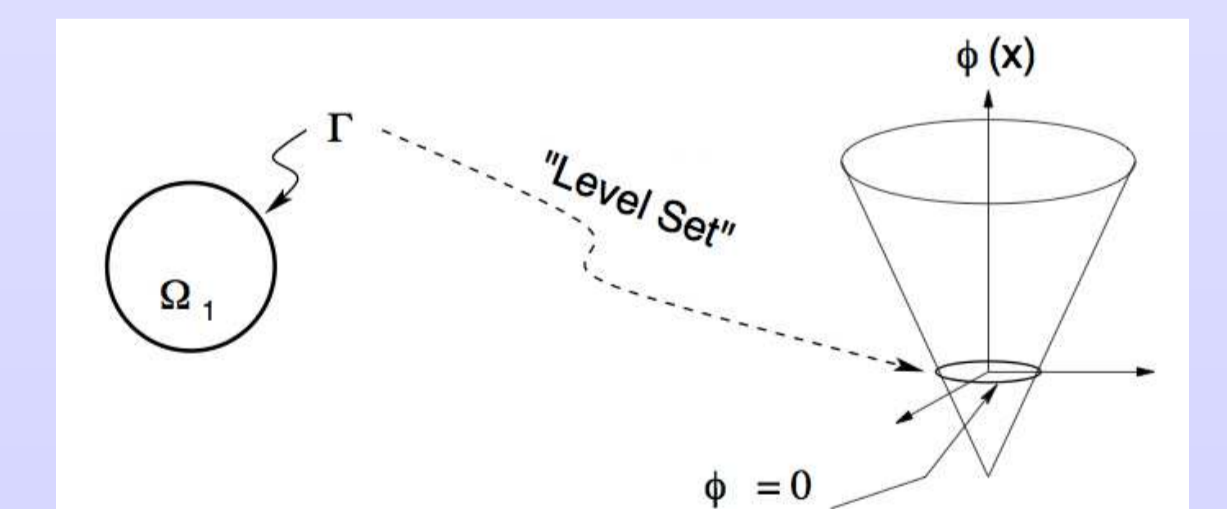
$$\frac{\partial L_i}{\partial t} = \alpha (Tub - c_c)$$

$$\frac{\partial MT_i}{\partial t} = \alpha (Tub - c_c) \left(\frac{\eta \nabla Tub \pm u}{\|\eta \nabla Tub \pm u\| + \varepsilon} \right) + u$$

Interface Representation :

We use a **Level-Set method**, based on an implicit representation of the interface as the zero level curve of a function ϕ called the Level-Set function, in order to avoid problem of interpolation between Lagrangian and Eulerian coordinates.

$$\frac{\partial \phi}{\partial t} + u \cdot \nabla \phi = 0$$



Forces : F_{el} , elastic force ; F_{net} , protrusion and contraction force

$$F_{el} = -\lambda \nabla \cdot \left(\frac{|\nabla \phi| - 1}{|\nabla \phi|} (\nabla \phi \otimes \nabla \phi) \frac{1}{\varepsilon} \zeta \left(\frac{\phi}{\varepsilon} \right) \right)$$

$$F_{net} = h(Rac) \frac{\nabla \phi}{|\nabla \phi|} - h(Rho) \frac{\nabla \phi}{|\nabla \phi|}$$

Numerical Tools : Finite Volume on locally refined meshes (DDFV) - WENO

Perspectives : Development of numerical tools - Comparison with experiments

References

- [Mai08] E. Maitre. *Equations de transport, Level Set et mécanique eulérienne. Application au couplage fluide-structure*. Habilitation à diriger des recherches, Université de Grenoble, November 2008.
- [PHM⁺12] A. Pagano, S. Honoré, R. Mohan, R. Berges, A. Akhmanova, and D. Braguer. Epithilone B inhibits migration of glioblastoma cells by inducing microtubule catastrophes and affecting EB1 accumulation at microtubule plus ends. *Biochemical Pharmacology*, pages 432-433, May 2012.
- [VFEK11] B. Vanderlei, J.J. Feng, and L. Edelstein-Keshet. A computational model of cell polarization and motility coupling mechanics and biochemistry. *Multiscale Model Simul.*, pages 1420-1443, October 2011.