

Mathematical modeling of a growing tissue

Our basis: mechanical model for one tissue

Individual Based Models



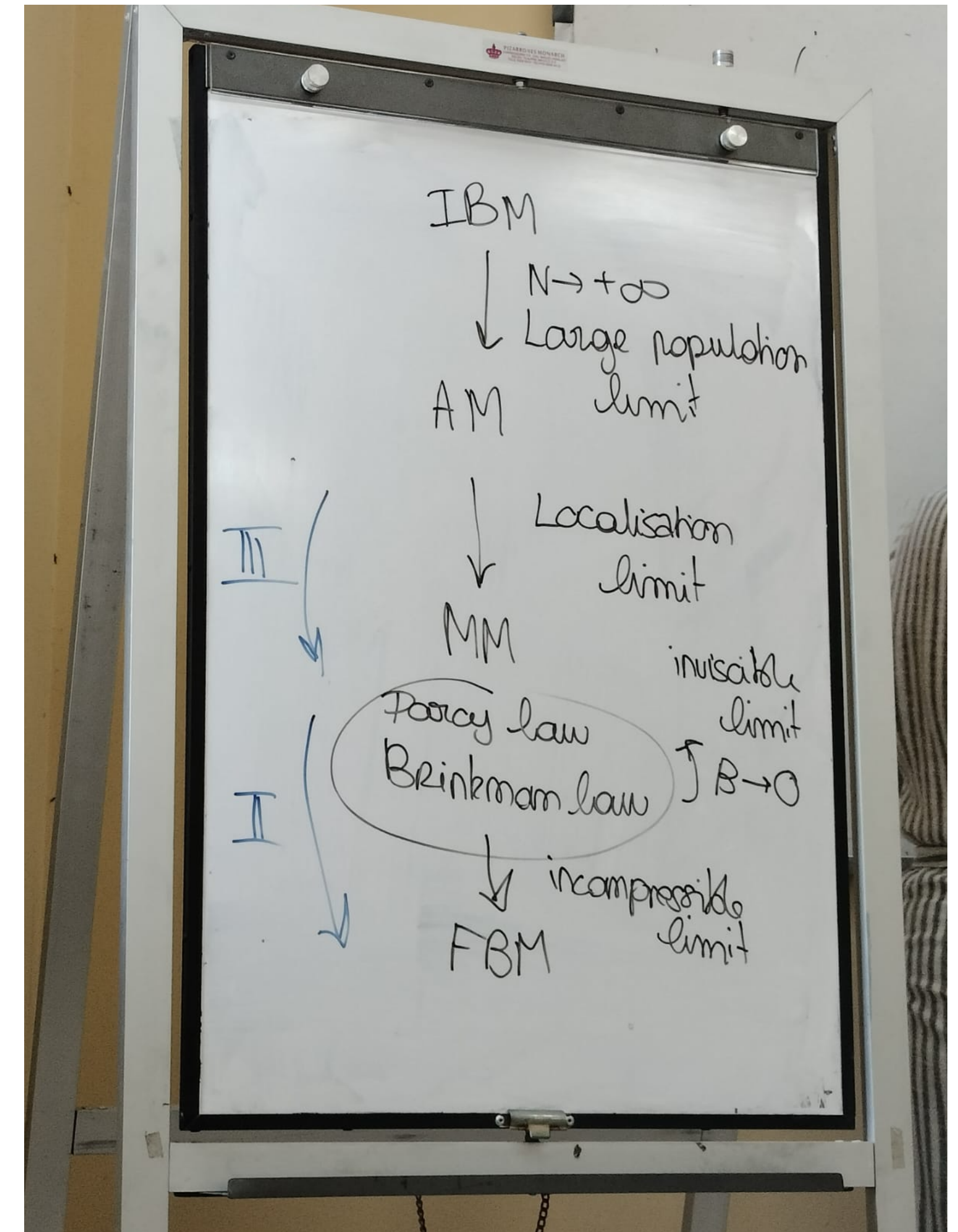
Aggregation Models



Mechanical Models



Free Boundary Models



Our basis: mechanical model for one tissue

$$\partial_t n + \nabla \cdot (nv) = nG(p),$$

Growth and migration

$$v = -\nabla p,$$

Darcy's law

$$p = p(n) = \epsilon \frac{n}{1-n},$$

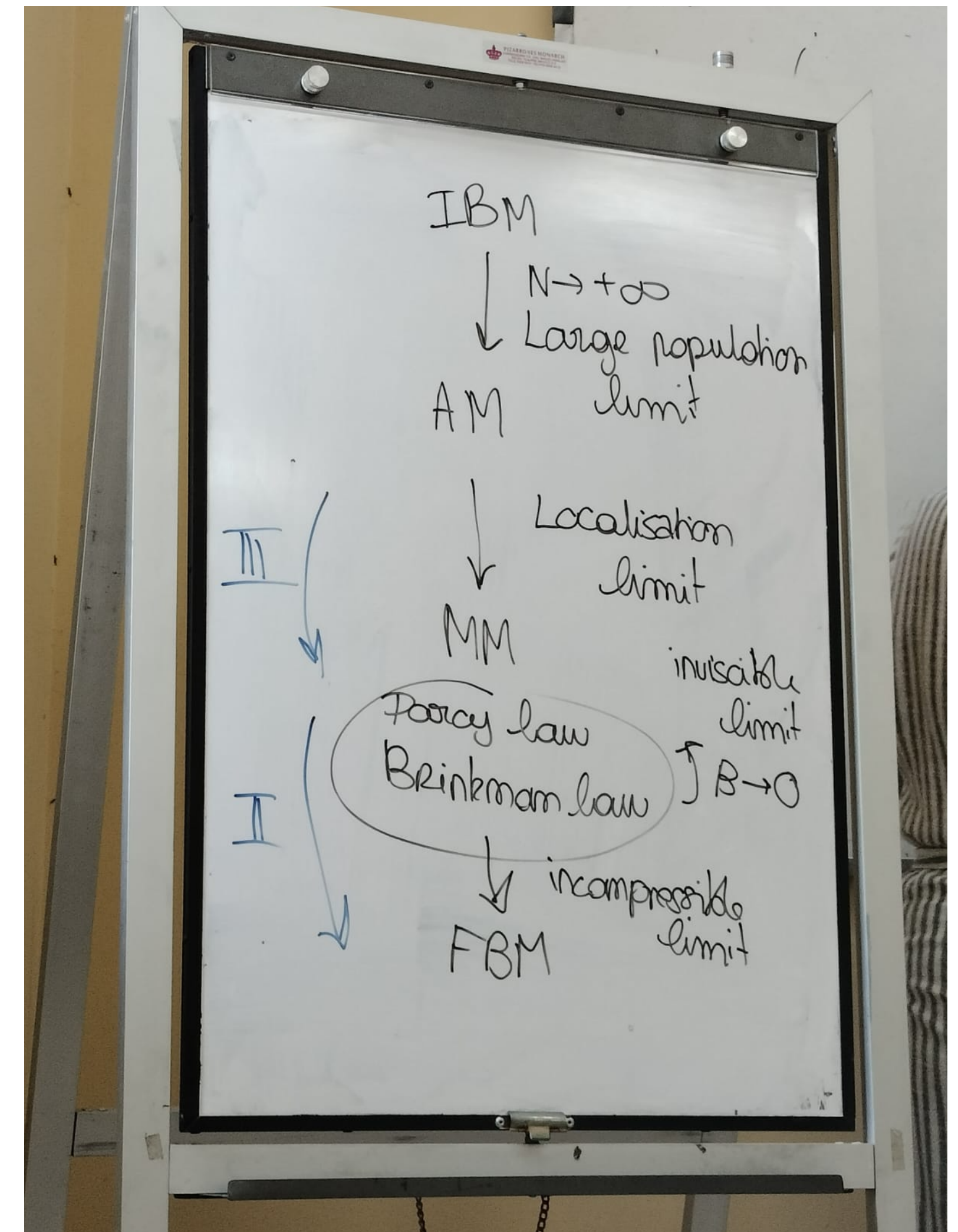
Pressure law (singular)

$n(t, x)$ cell density of the tissue

$v(t, x)$ velocity of the tissue

$p(n)$ pressure due to congestion

G pressure-dependent growth function



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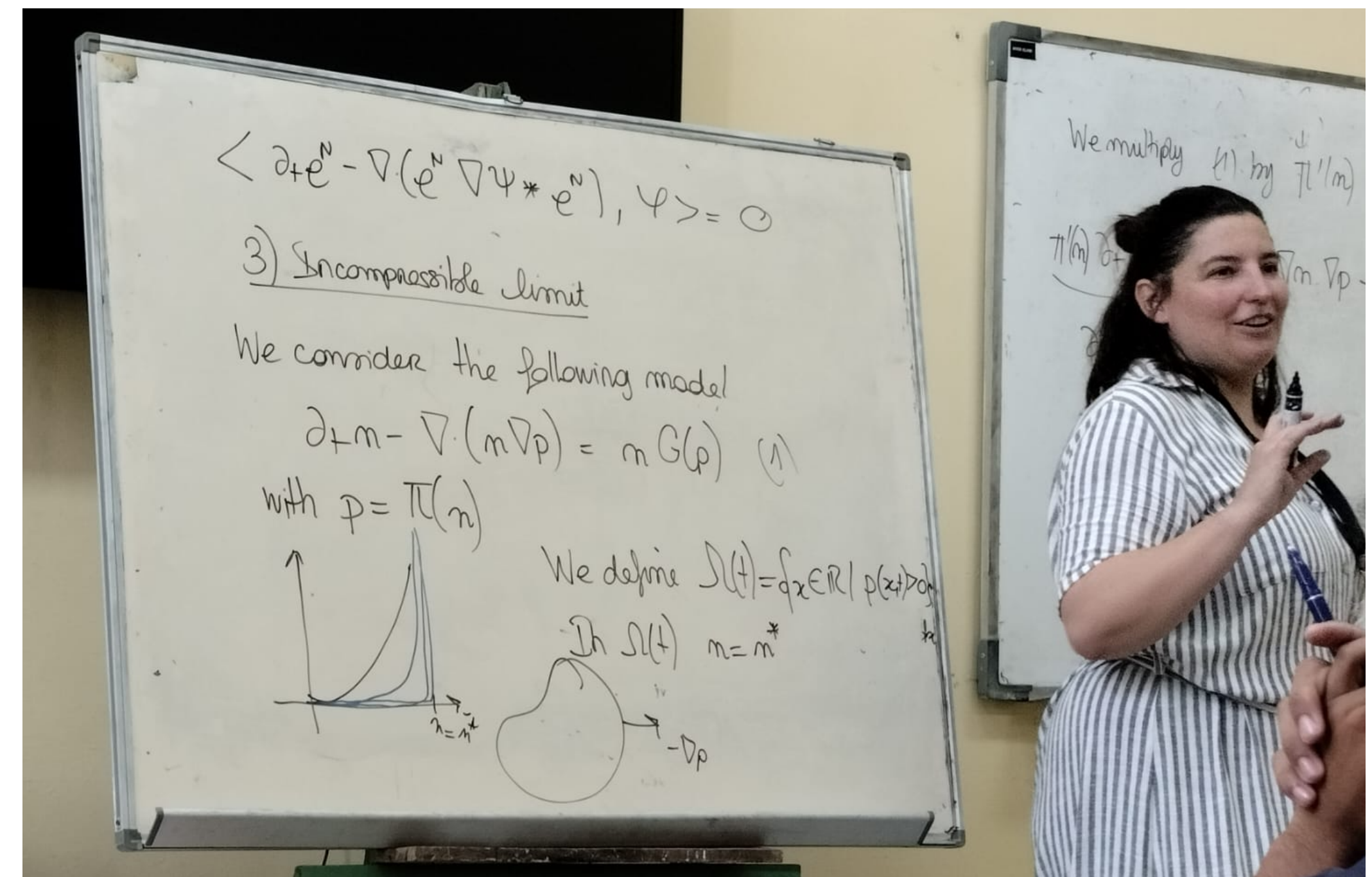
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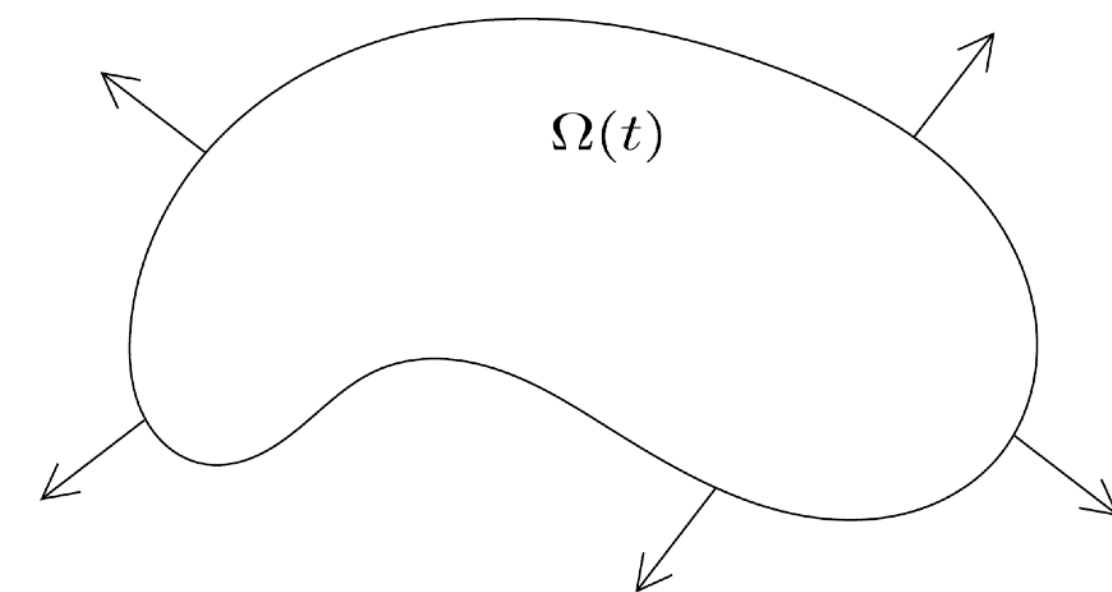
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When the pressure parameter vanishes:



$$n = 1, \text{ in } \Omega(t)$$

$$-\Delta p = G(p), \text{ in } \Omega(t)$$

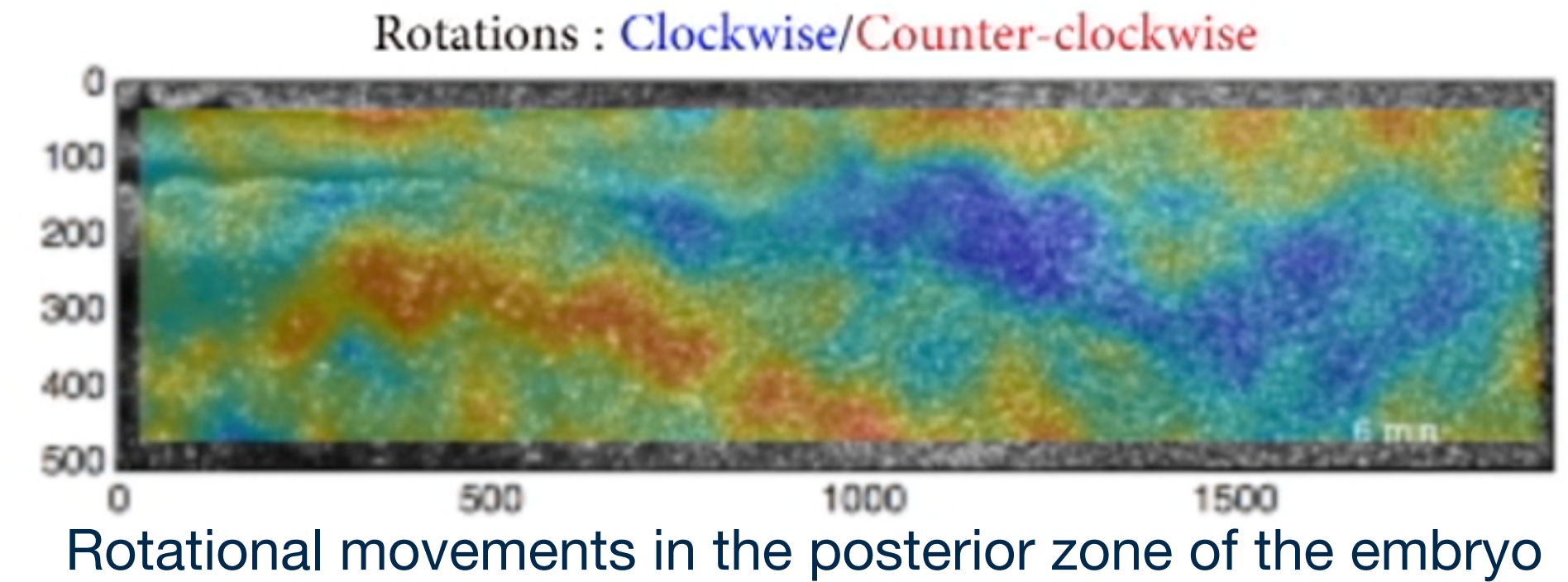
$$v_{\partial\Omega} = -(\nabla p \cdot \nu)\nu, \text{ on } \partial\Omega(t)$$

Viscosity and rotational

Using **Darcy's law**

$$\mathbf{v} = -\nabla p,$$

we necessarily have a null rotational !

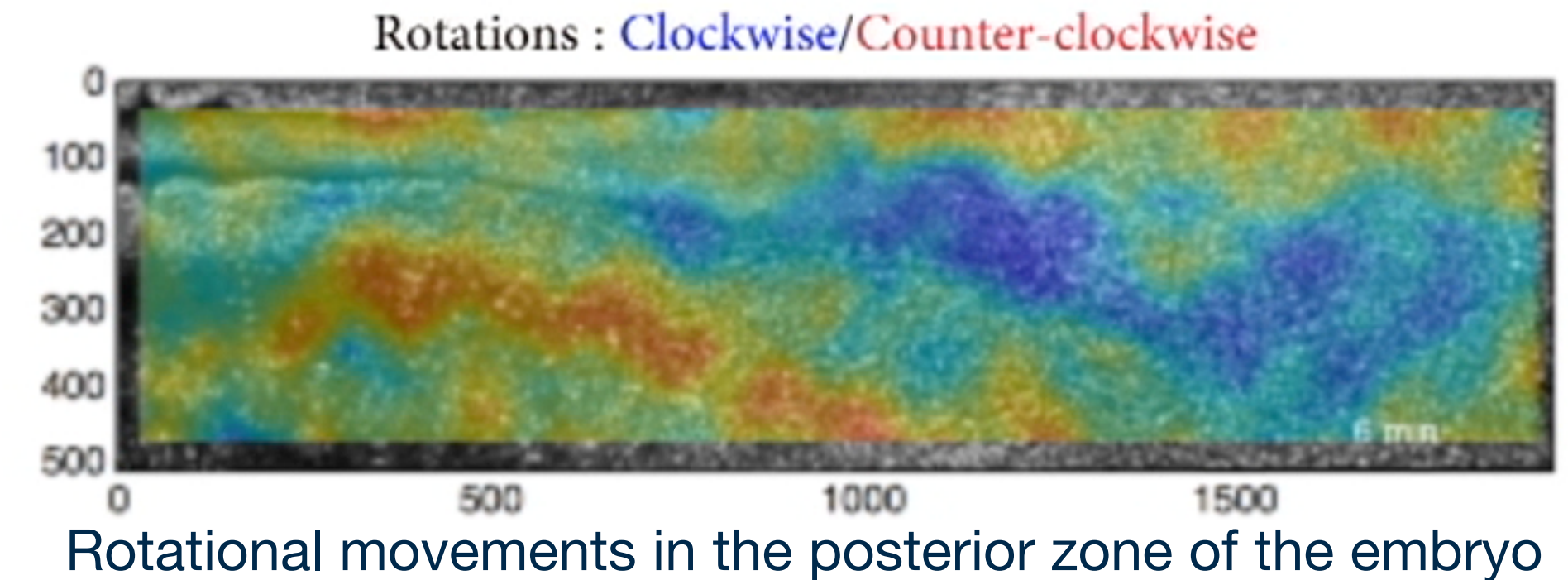


Viscosity and rotational

Using **Darcy's law**

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we necessarily have a null rotational !



To reproduce the rotational movements observed in the embryo, and take into account the (differential) viscosities of the tissues, we choose instead the **Brinkman form**:

$$-\beta \Delta \mathbf{v} + \mathbf{v} = -\nabla p$$

β viscosity of the tissue

Multi-tissue models

Mechanical model for two tissues (1)

$$\left\{ \begin{array}{l} \partial_t n_1 + \nabla \cdot (n_1 v_1) = n_1 G_1(p_1), \\ \partial_t n_2 + \nabla \cdot (n_2 v_2) = n_2 G_2(p_2), \\ -\beta_1 \Delta v_1 + v_1 = -\nabla p_1, \\ -\beta_2 \Delta v_2 + v_2 = -\nabla p_2, \\ p_1 = p_\epsilon(n_1 + n_2), \\ p_2 = p_\epsilon(n_1 + n_2), \\ p_\epsilon(n) = \epsilon \frac{n}{1-n}. \end{array} \right.$$

n_i : density of each tissue

v_i : velocity of each tissue

p_i : pressure of each tissue

β_i : viscosity of each tissue

p_ϵ : congestion pressure, parametrized by $\epsilon > 0$

G_i : growth functions

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?

mixing of the two tissues ?

Mechanical model for two tissues (1)

Propagated segregation: starting from initial segregated tissues

$$n_1^{ini} n_2^{ini} = 0$$

we get that the tissues remain segregated for all times

$$n_1(t, \cdot) n_2(t, \cdot) = 0, \quad t > 0.$$

Mechanical model for two tissues (1)

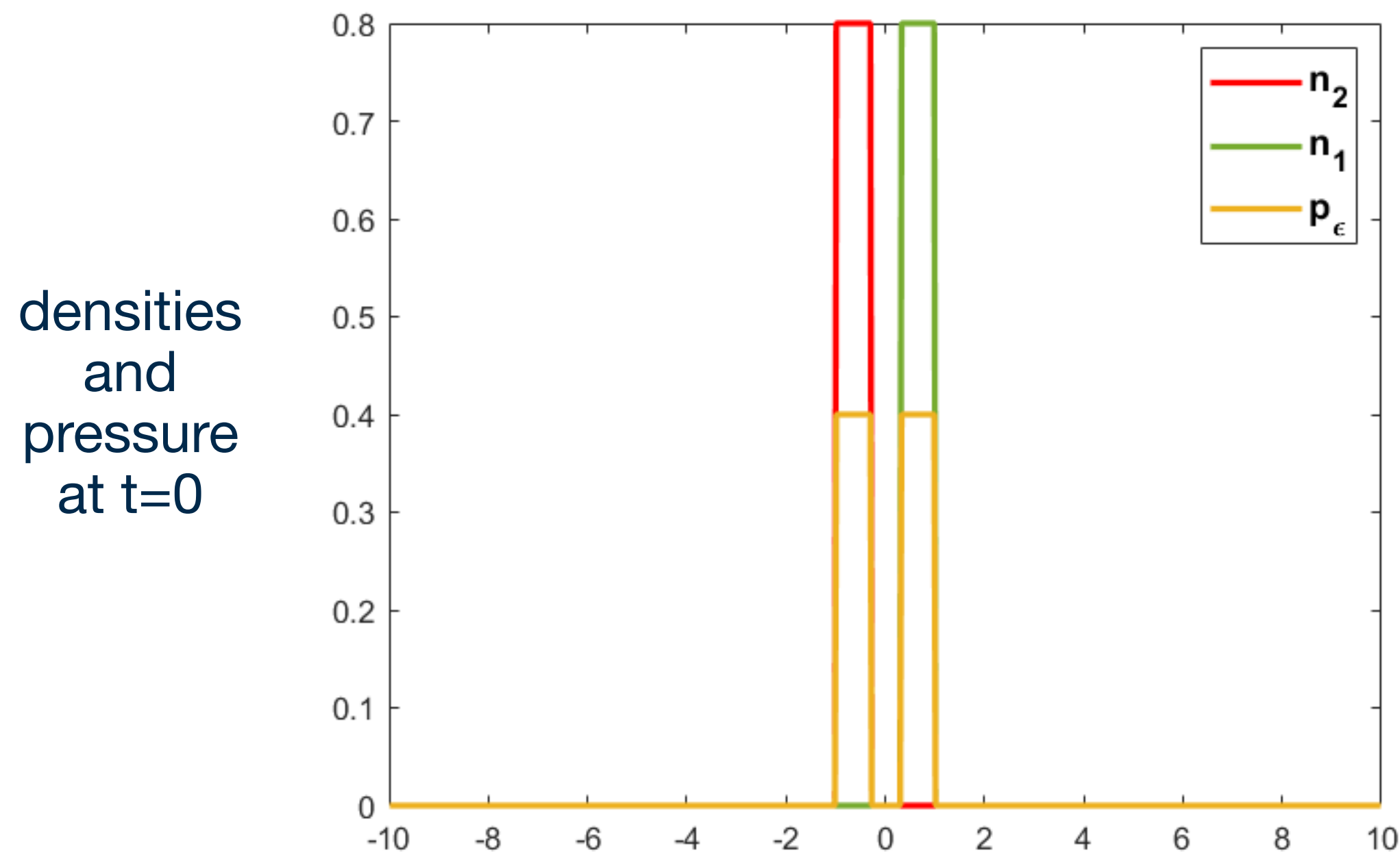
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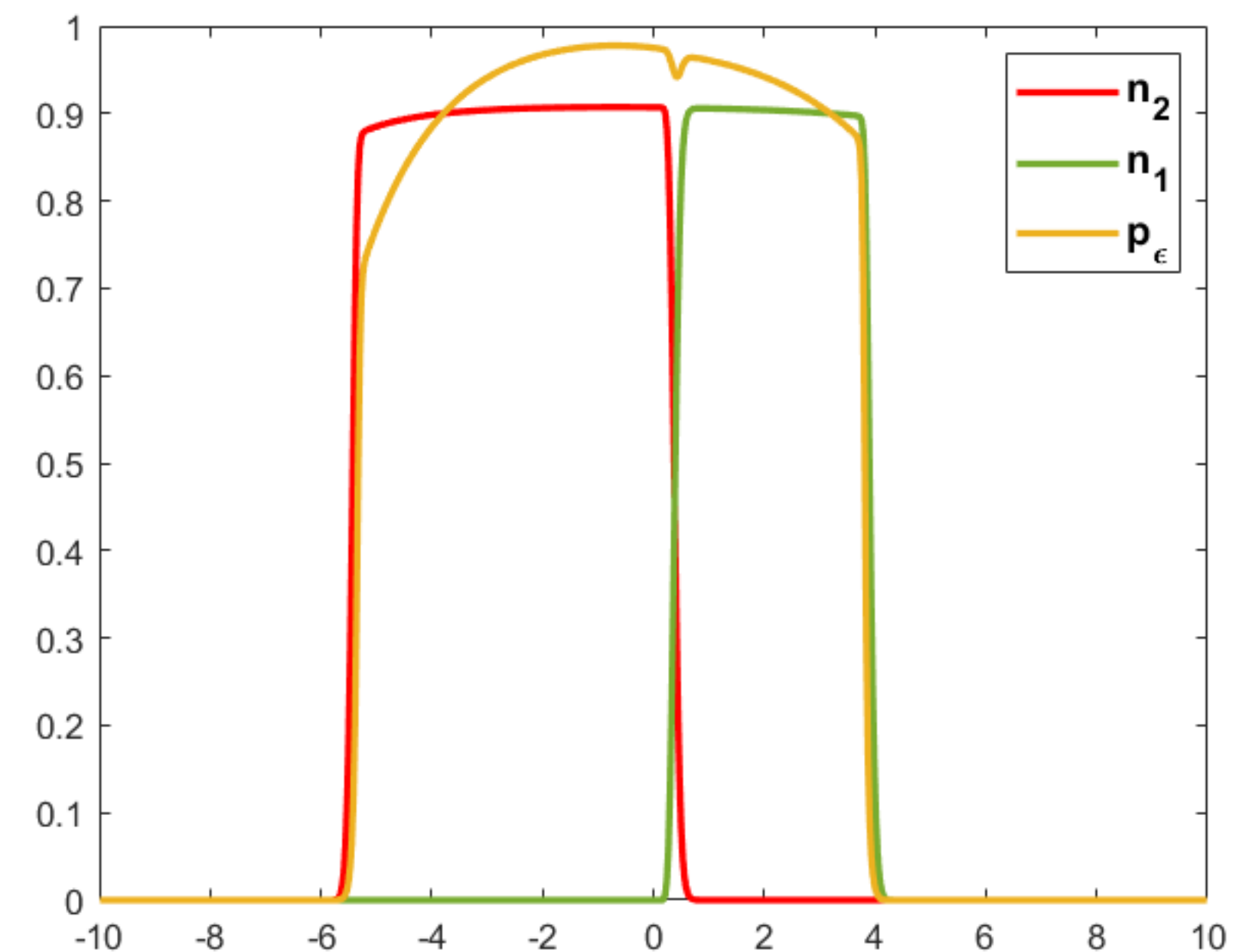
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$$n_1(t, \cdot) n_2(t, \cdot) = 0, \quad t > 0.$$

This property is *proved* for equal viscosities, and *numerically observed* for different viscosities.



$$\beta_1 = 3, \beta_2 = 1.$$



Mechanical model for two tissues (1)

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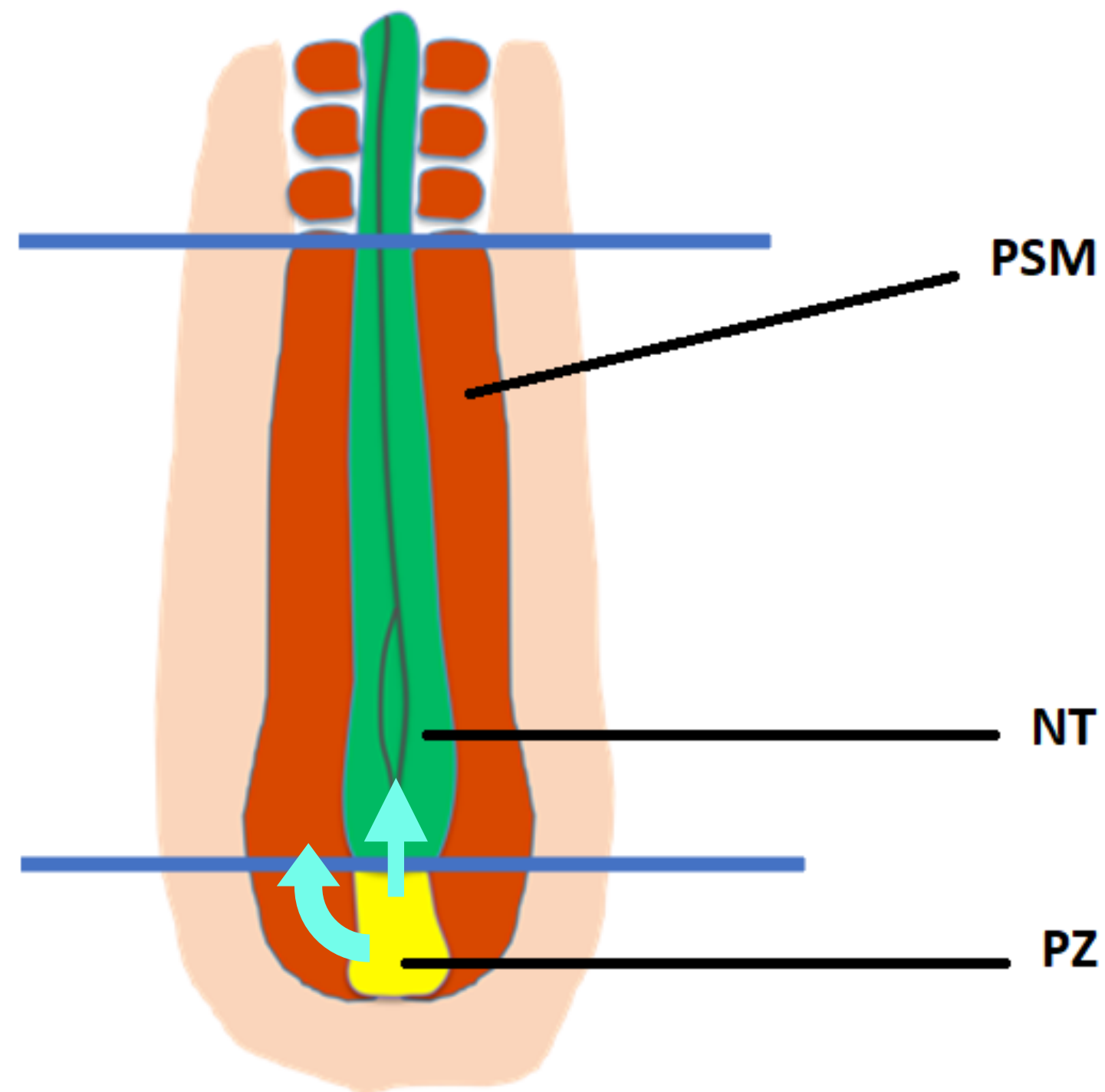
β_i : viscosity of each tissue

p_ϵ : congestion pressure, parametrized by $\epsilon > 0$

G_i : growth functions

Segregation is **propagated** from the initial data (**passive segregation**).

Mechanical model: origin of segregation ?



Cells of NT and PSM types migrate from the PZ towards the two other tissues

If the segregation is only propagated, then, what is its origin ?

What happens at the interface with the PZ, which is a mixed zone ?

Biological hypothesis: could it be that some active segregation is at play ?

We keep our first model with *passive segregation* but also build a model with *active segregation* in order to test this hypothesis.

Mechanical model for two tissues (2)

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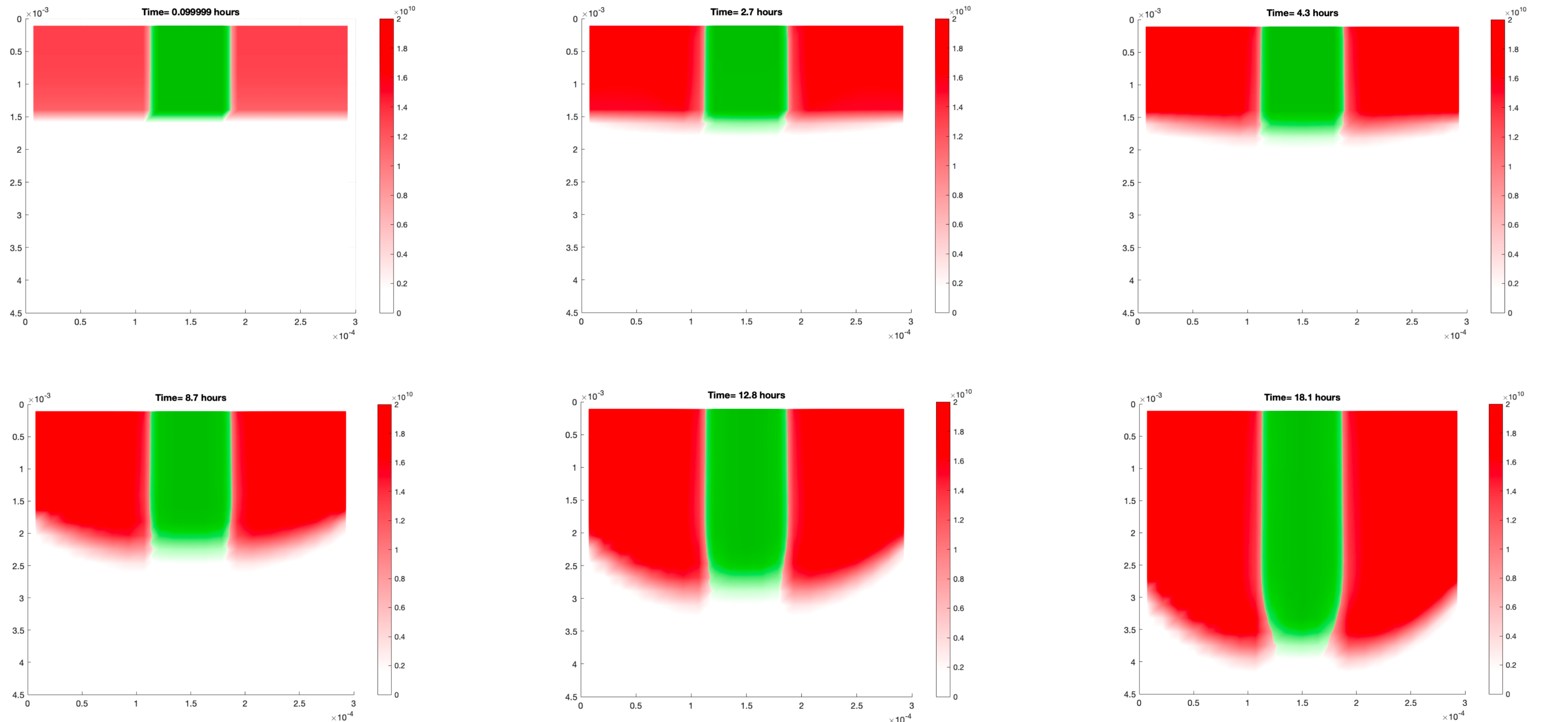
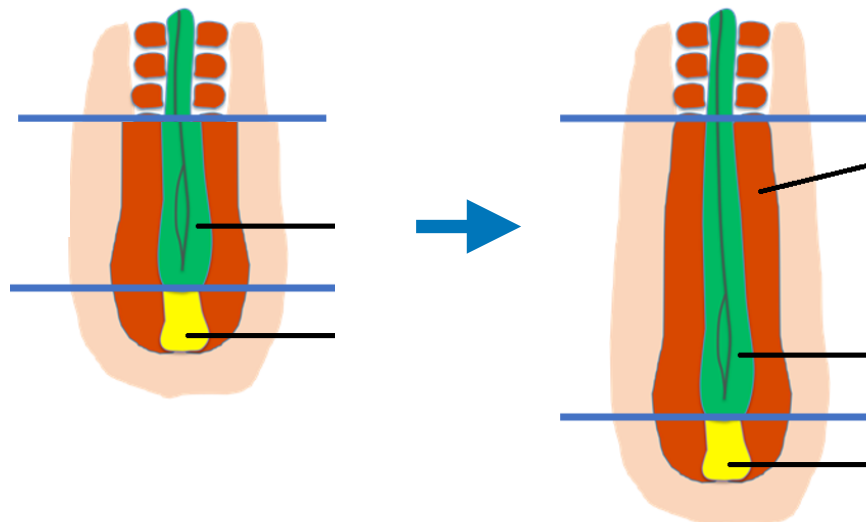
q_m : repulsion pressure, parametrized by $m > 0$

$\alpha > 0$: regularization parameter

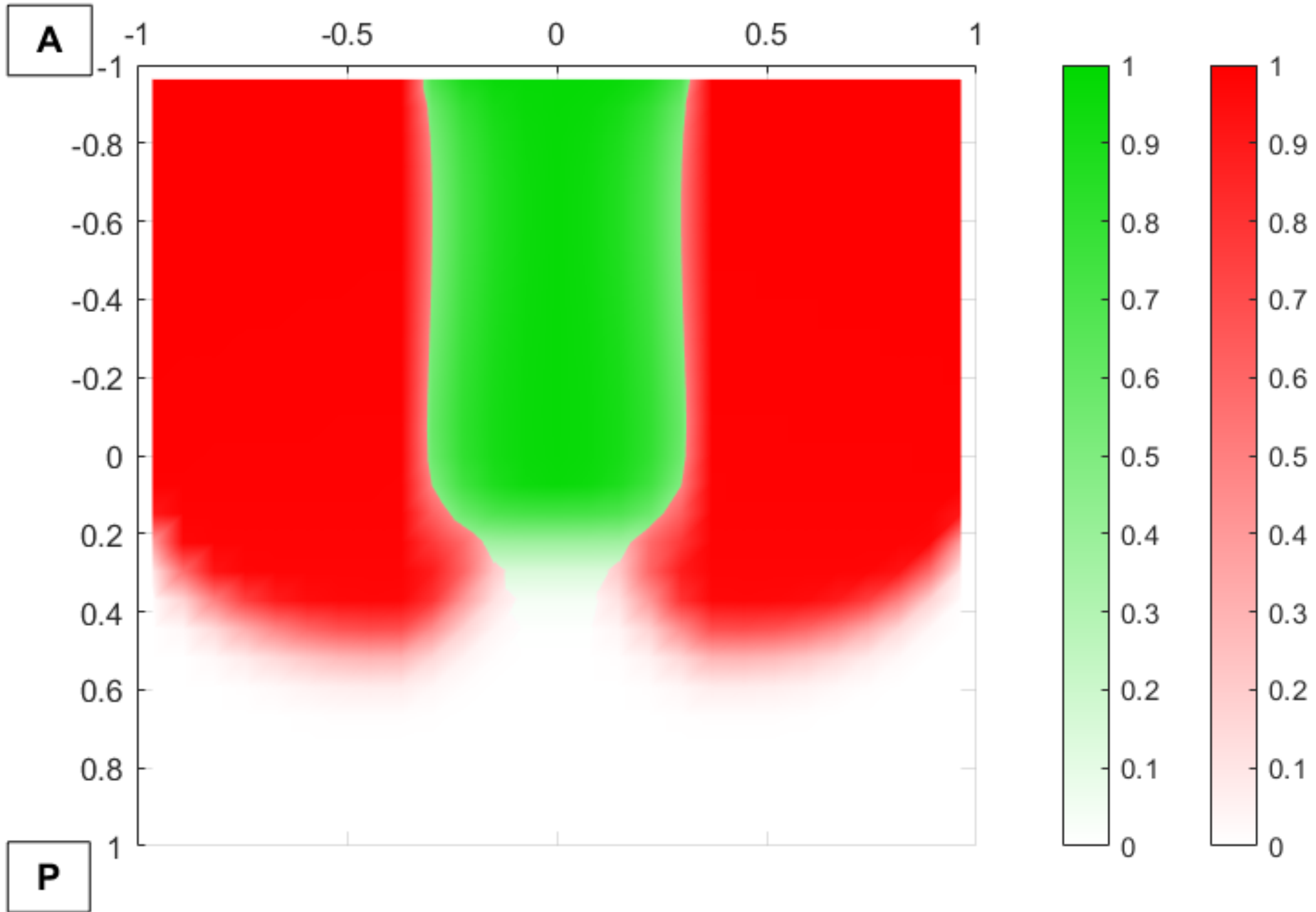
Active segregation

Numerical simulation and comparison with data

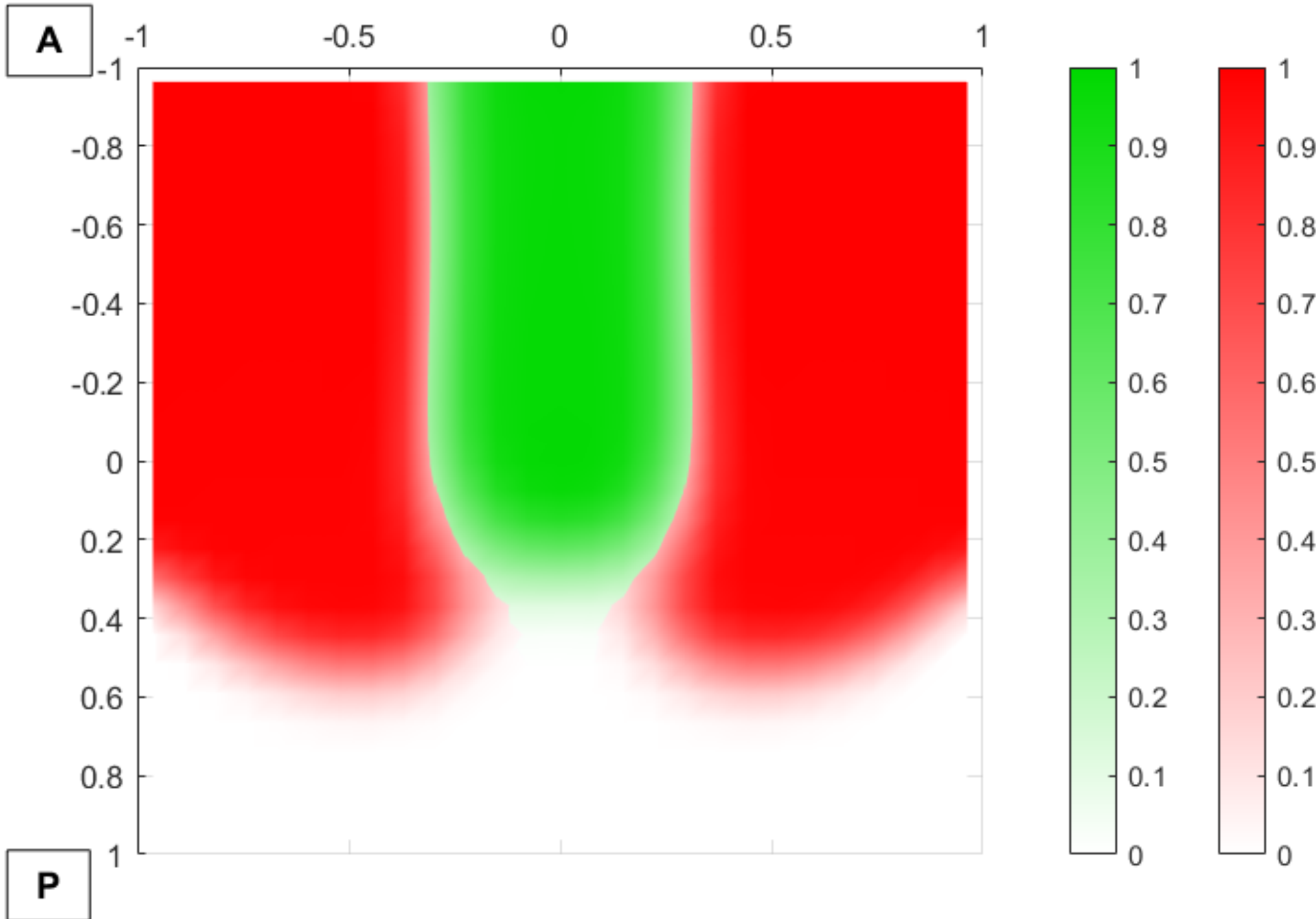
Model 1 (fluid, passive segregation)



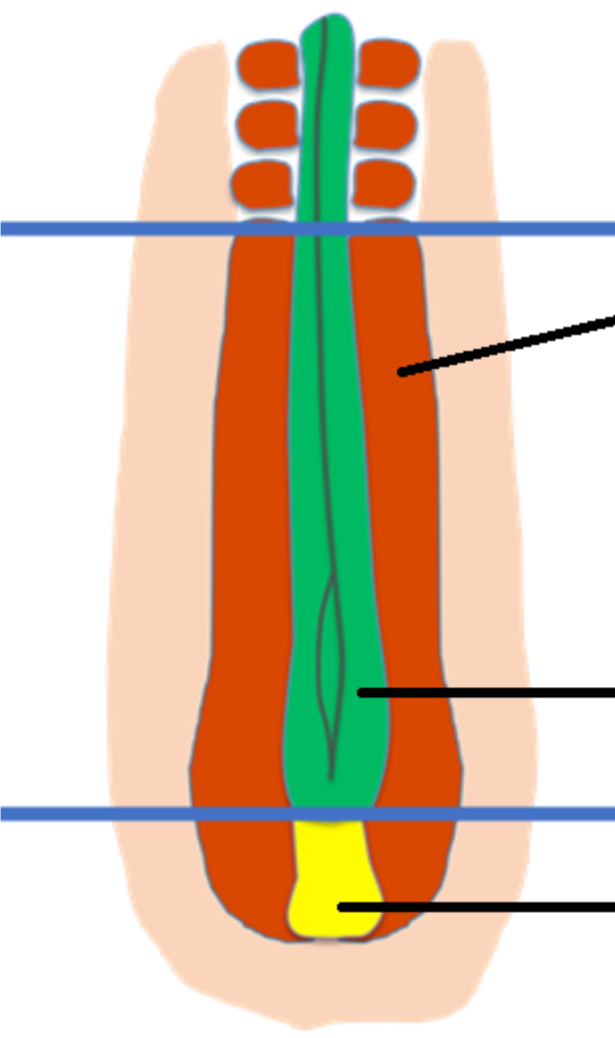
Comparison between models: densities



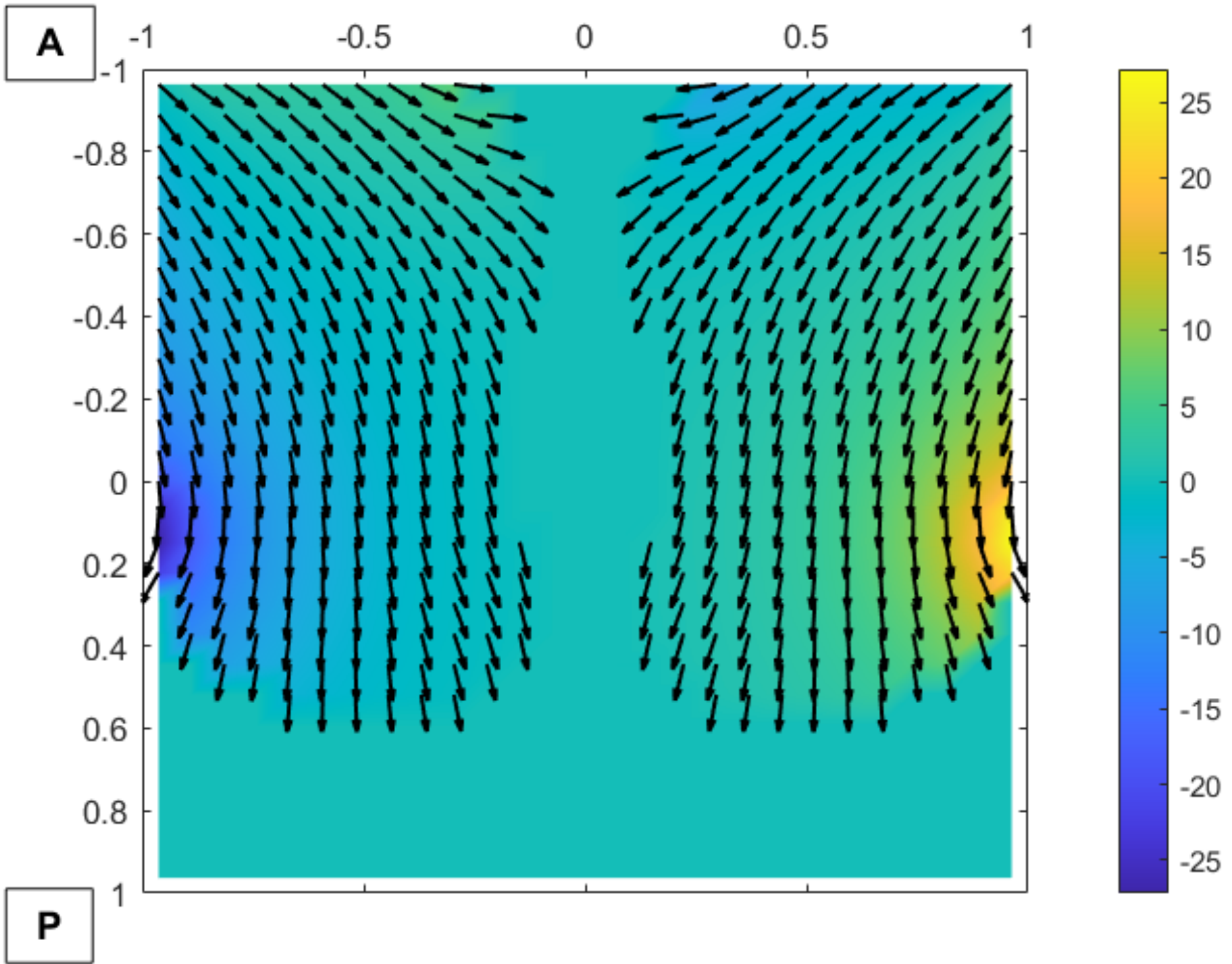
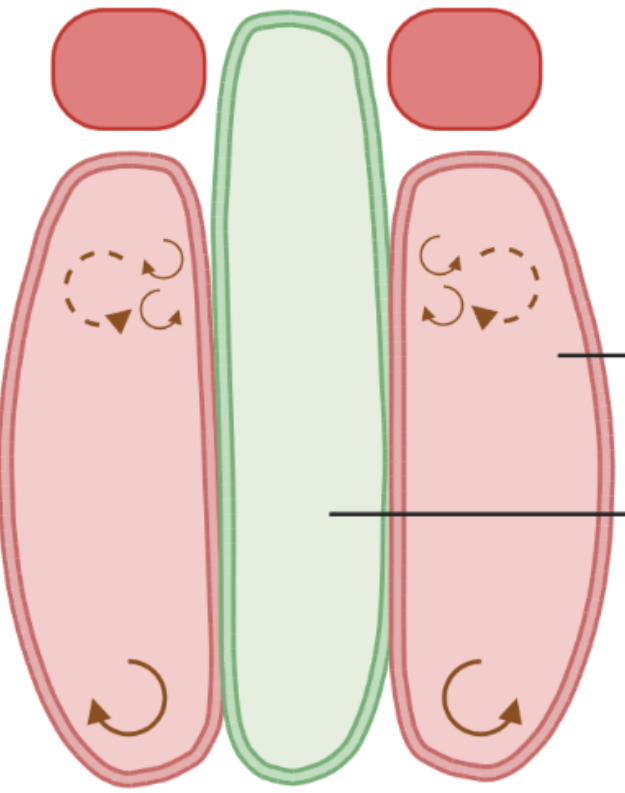
Fluid model,
Passive segregation



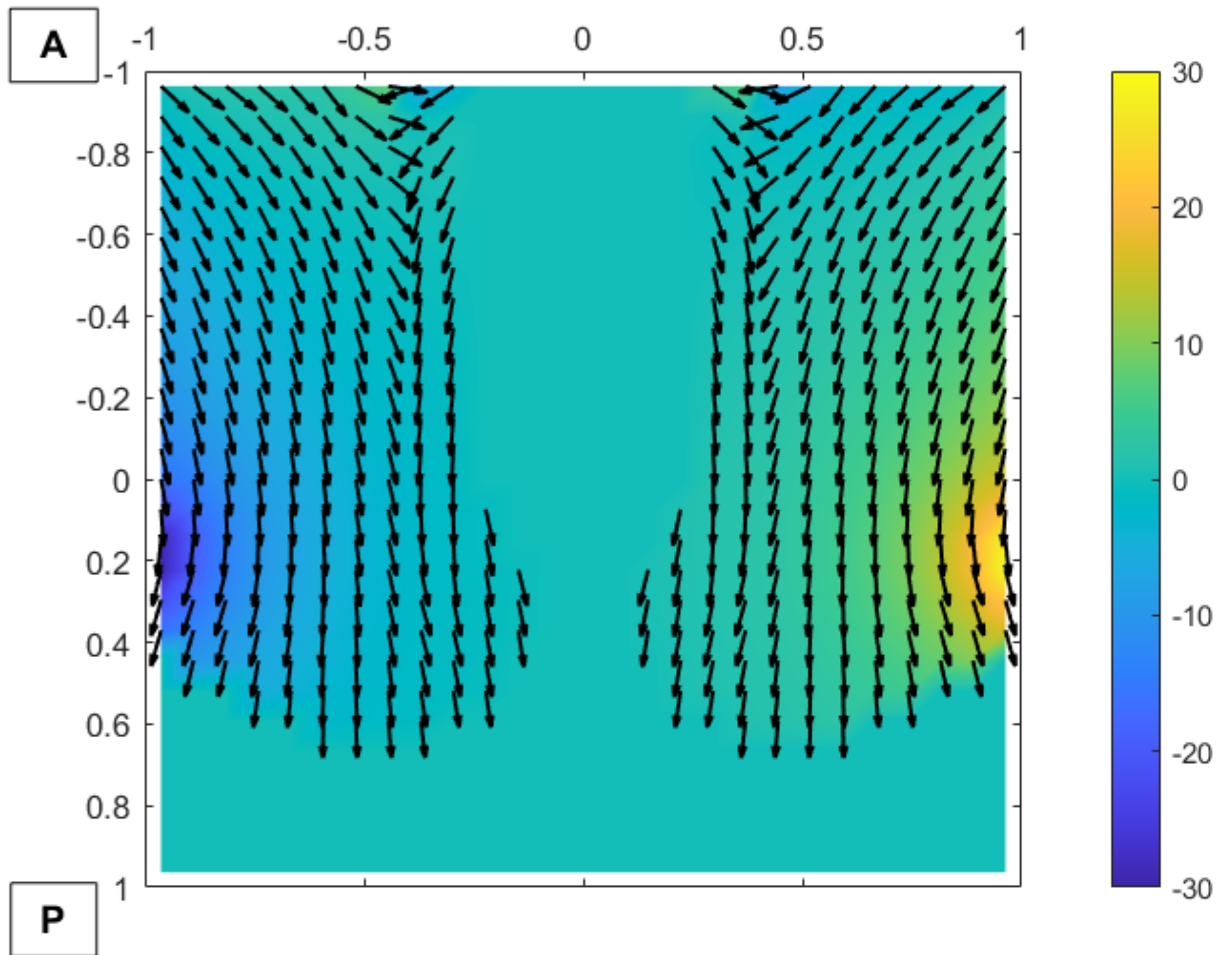
Fluid model,
Active segregation



Comparison between models: rotational

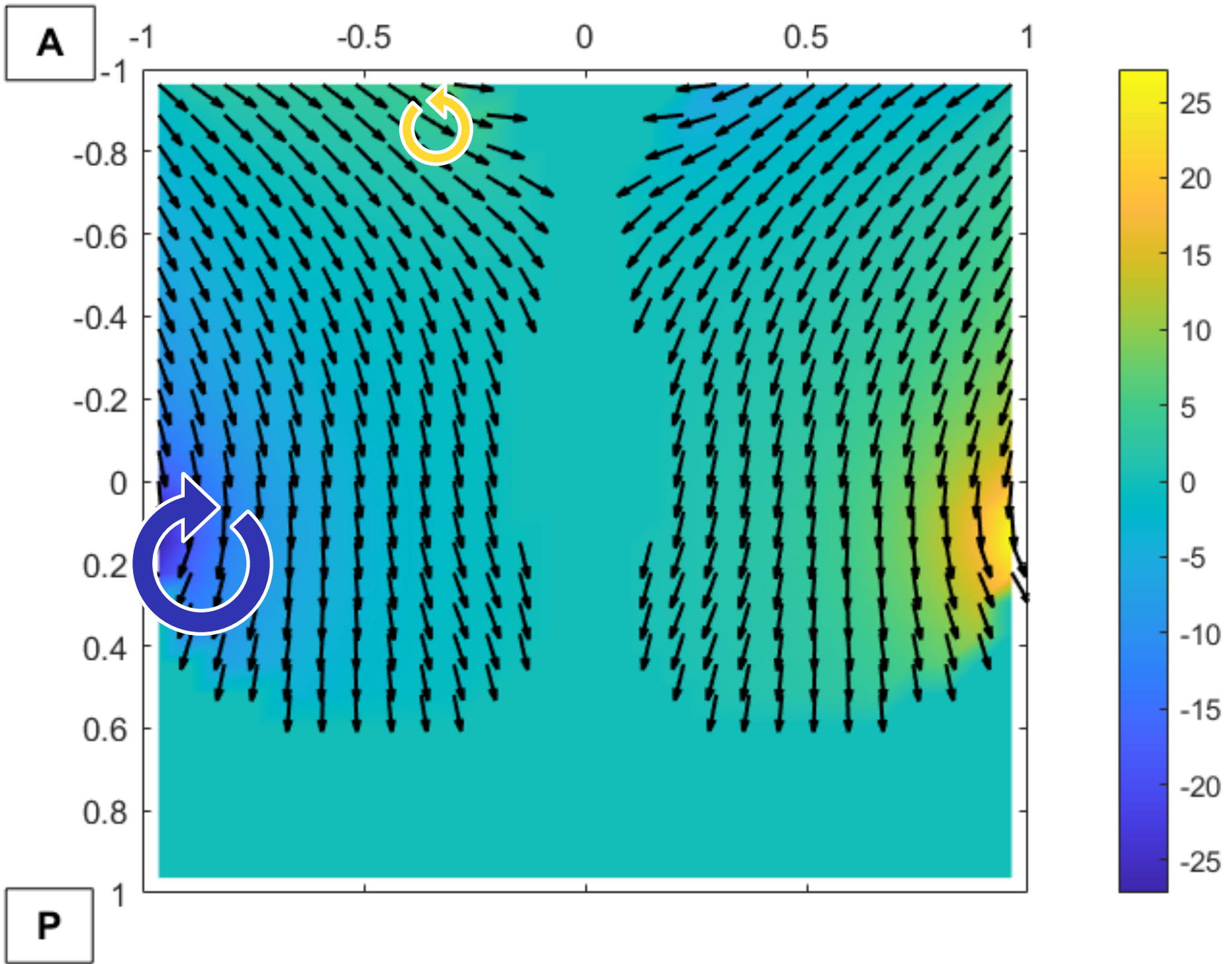
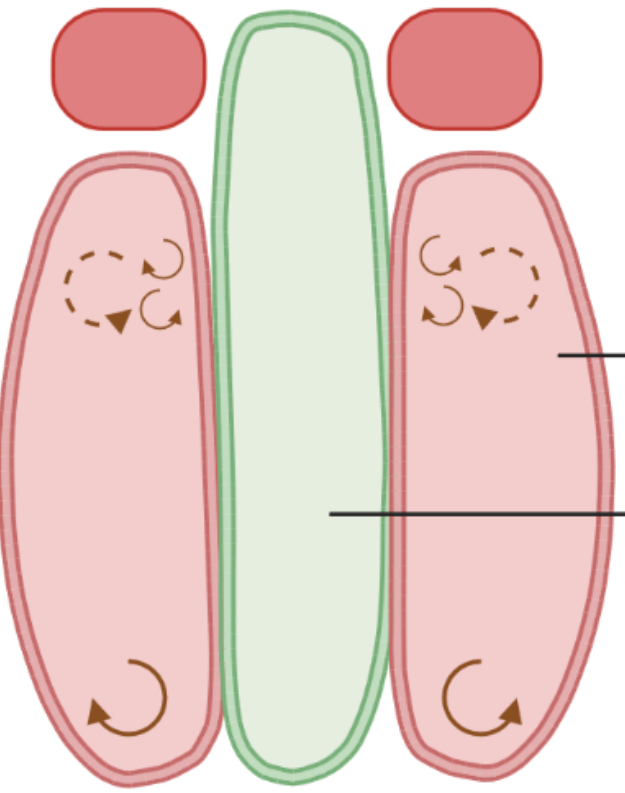


Fluid model,
Passive segregation

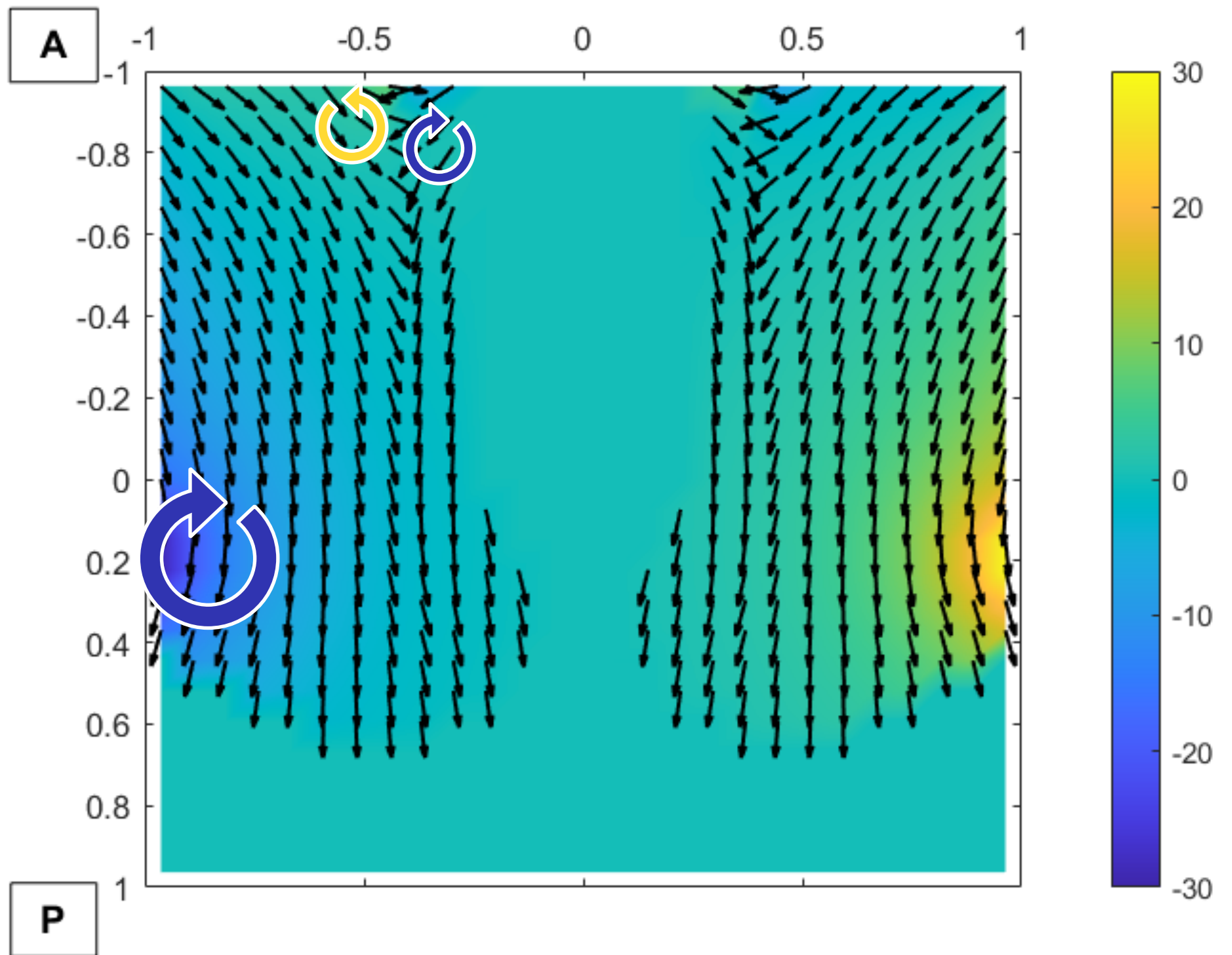


Fluid model,
Active segregation

Comparison between models: rotational

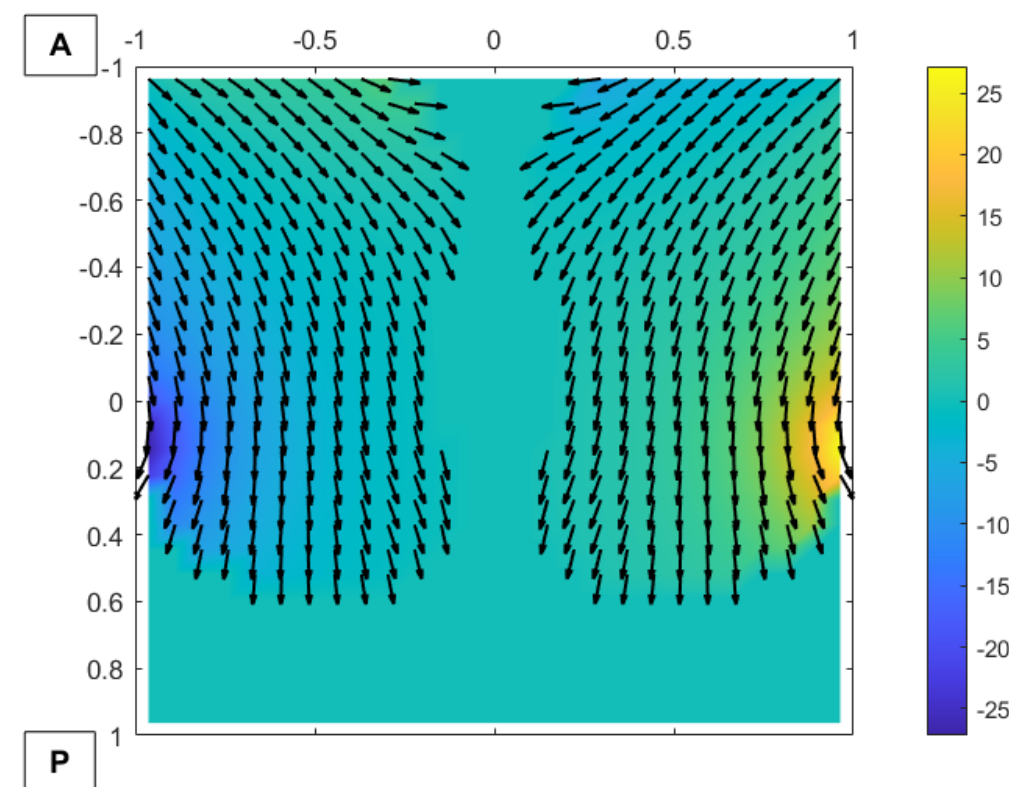
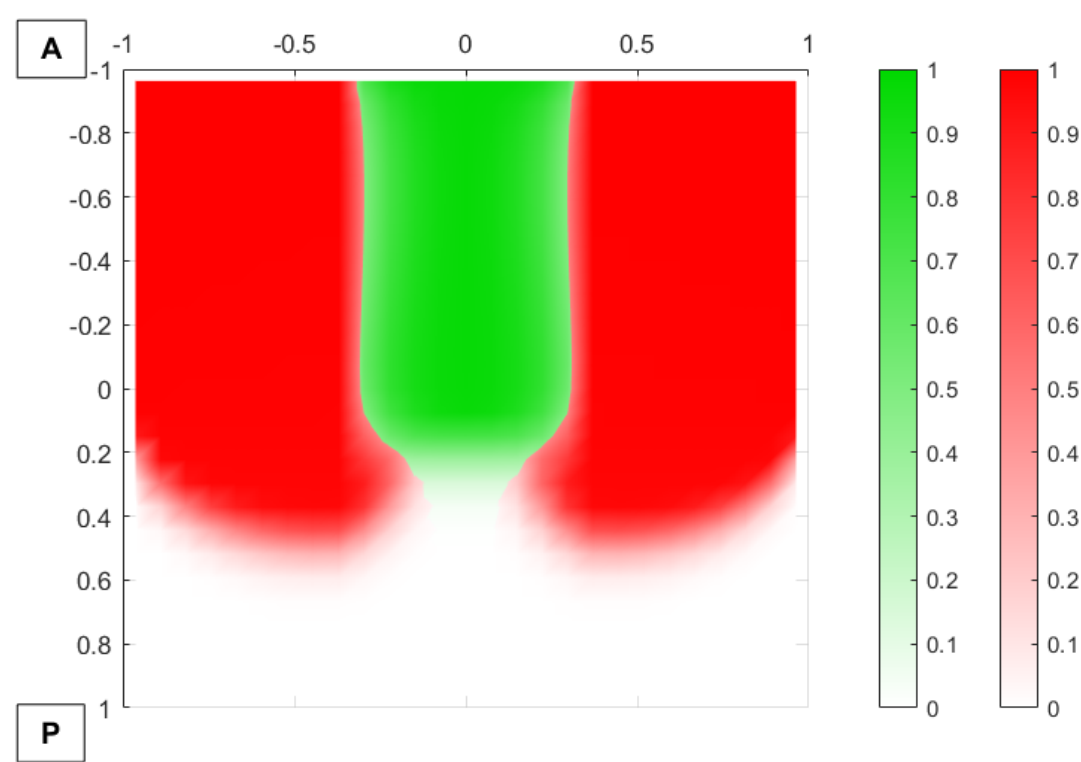


Fluid model,
Passive segregation

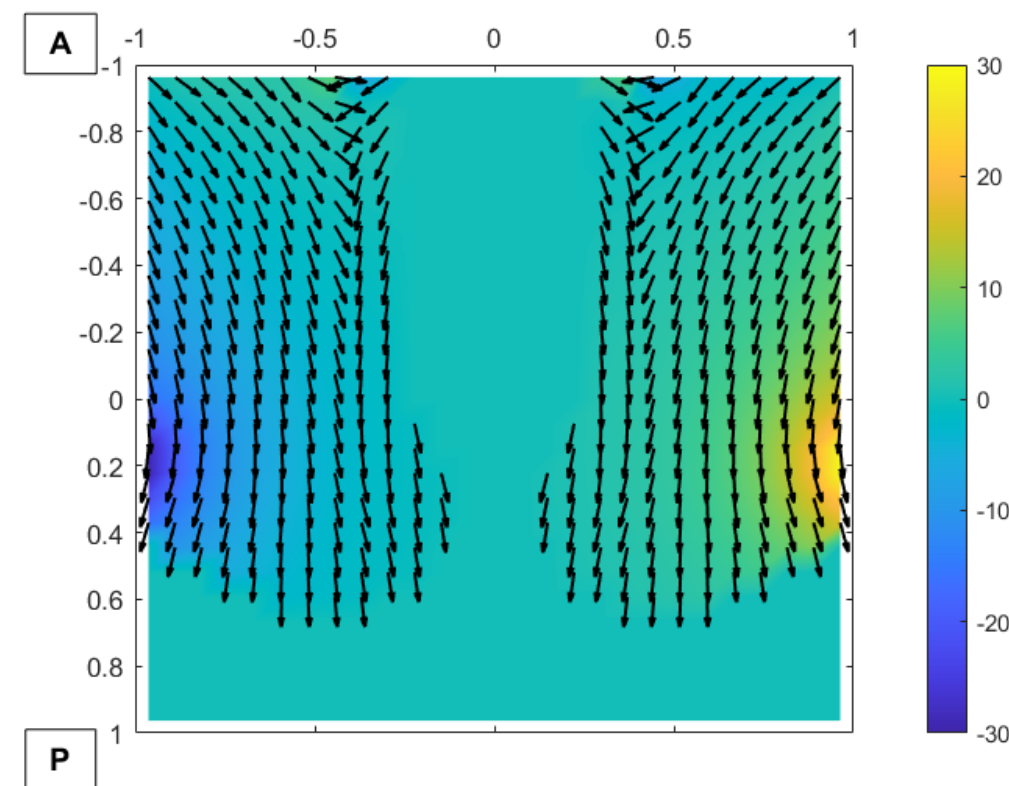
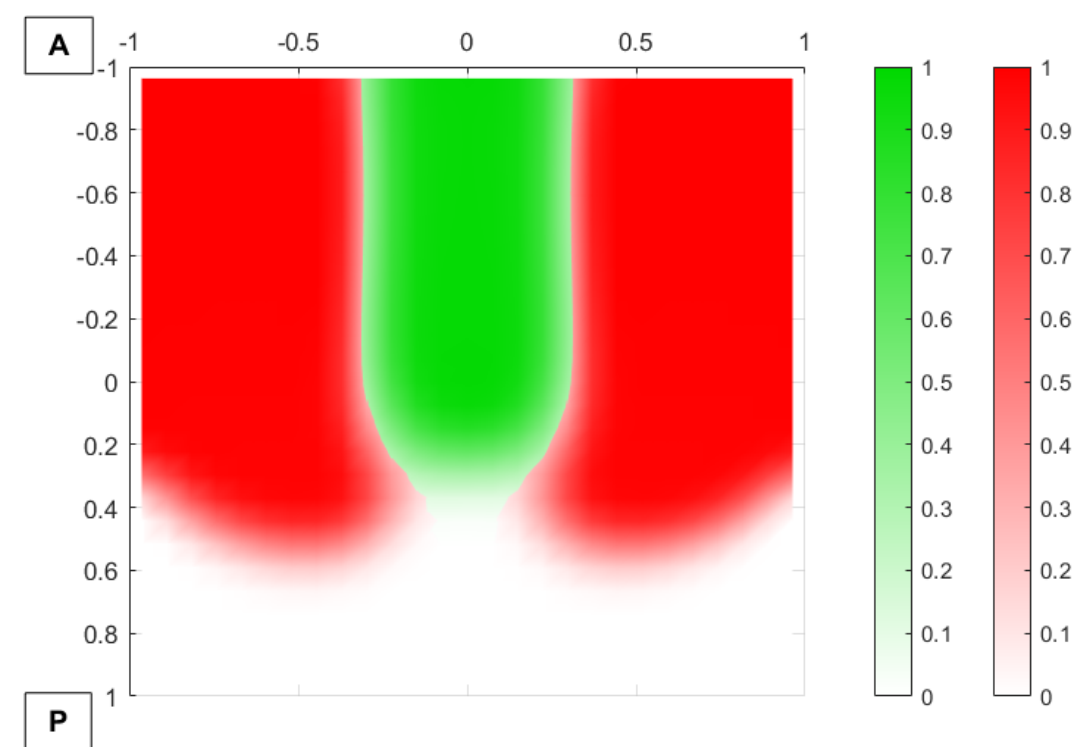


Fluid model,
Active segregation

Comparison between models



Fluid model,
Passive segregation



Fluid model,
Active segregation

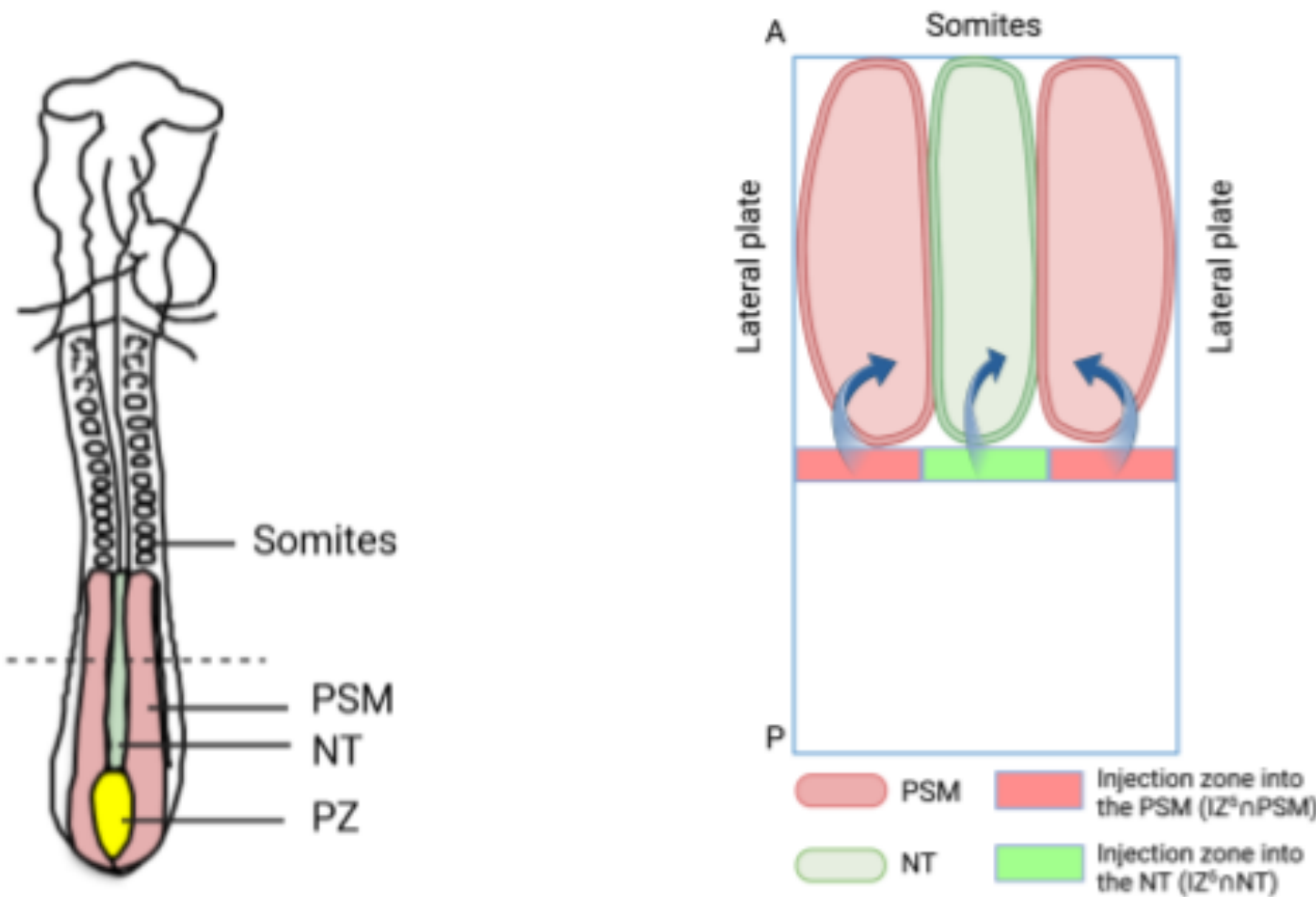


Our model suggests that some active segregation is at play.

Diving deeper into the biological data

One step forward towards biology:

- we add injection (duly *quantified*)
- other parameters taken from the biological literature



Parameter	Source	Value	Unit
NT viscosity (β_1)	Michaut 2018	$10^4 - 10^5$	Pa · s
PSM viscosity (β_2)	Mahadevan 2017, Michaut 2018	$10^4 - 10^5$	Pa · s
Proliferation rate of the NT (g_1)	Bénazéraf 2017	1/10.83	1/ hour
Proliferation rate of the PSM (g_2)	Bénazéraf 2017	1/8.75	1/ hour
Injection rate from the PZ into the NT (κ_{NT})	Measured experimentally	5	Cells/ hour/ 10 μ m
Injection rate from the PZ into the PSM (κ_{PSM})	Measured experimentally	10	Cells/ hour/ 10 μ m
Total maximal density (n_{max})	Bénazéraf 2017	2800	Cells/ 100 μ m ³
Pressure sensing (ϵ)	Estimated free parameter	1	Pa
Tissue friction with surroundings (μ)	Mahadevan 2017, Michaut 2018	$10^{12} - 10^{13}$	Pa · s/m ²
Growth sensitivity near the maximal density (α)	Estimated free parameter	8	Dimensionless
Width of injection zone (δ)	Measured experimentally	214	μ m

Tissue growth

Cell proliferation

$$\partial_t n_1 + \nabla \cdot (n_1 v_1) = n_1 g_1 \left(1 - \left(\frac{n_1}{n_{max}} \right)^\alpha \right)_+ + \frac{\kappa_{NT}}{\delta} \chi_{IZ^\delta \cap NT},$$

Cell Injection

$$\partial_t n_2 + \nabla \cdot (n_2 v_2) = n_2 g_2 \left(1 - \left(\frac{n_2}{n_{max}} \right)^\alpha \right)_+ + \frac{\kappa_{PSM}}{\delta} \chi_{IZ^\delta \cap PSM},$$

$$-\beta_1 \Delta v_1 + \mu v_1 = -\nabla p,$$
$$-\beta_2 \Delta v_2 + \mu v_2 = -\nabla p,$$
$$p = \epsilon \frac{n}{n_{max} - n}, \quad n = n_1 + n_2.$$

Time-space evolution of the NT density and of the PSM density

Brinkman's law for the NT and PSM velocities

Congestion pressure

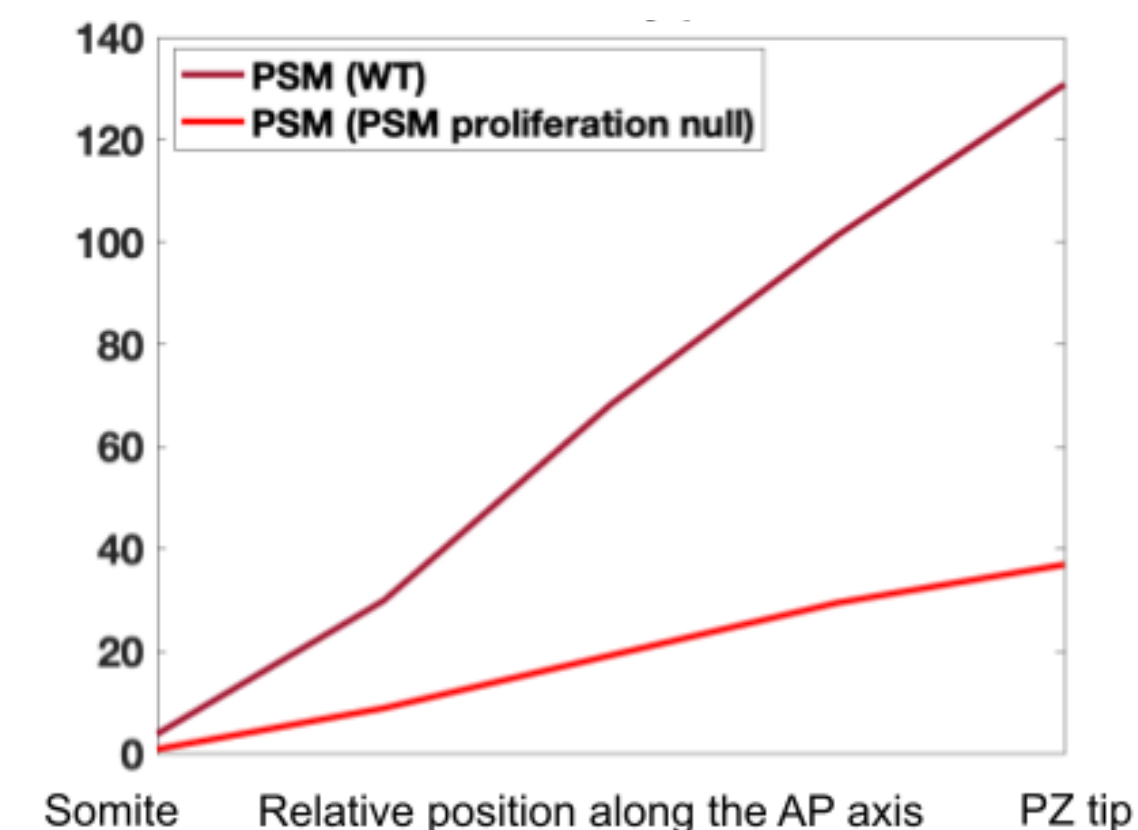
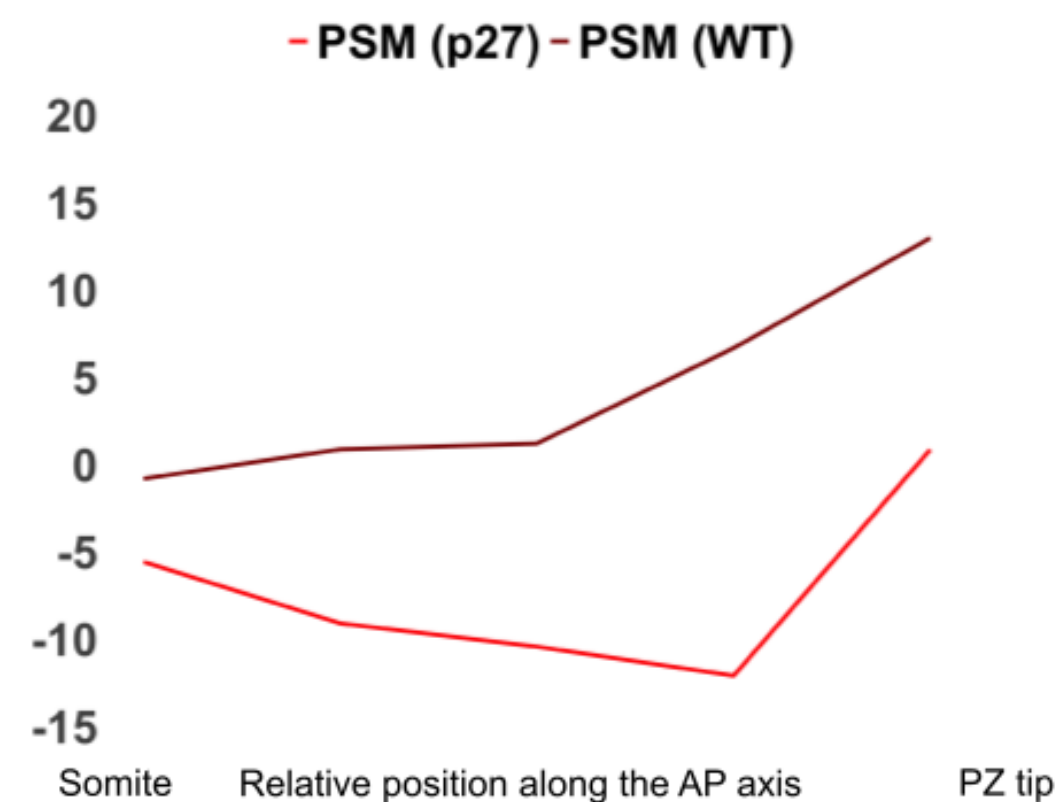
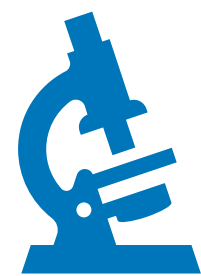
Diving deeper into the biological data

One step forward towards biology:

- we add injection (duly *quantified*)
- other parameters taken from the biological literature

Validation of the model:

- with « natural » parameters, compared with *wild type* data (« healthy » embryo)
 - with pathological parameters, compared with experiments *in vivo* (proliferation suppressed, ...)
- In particular, the model reproduces well the differential elongation and the sliding between tissues.



Diving deeper into the biological data

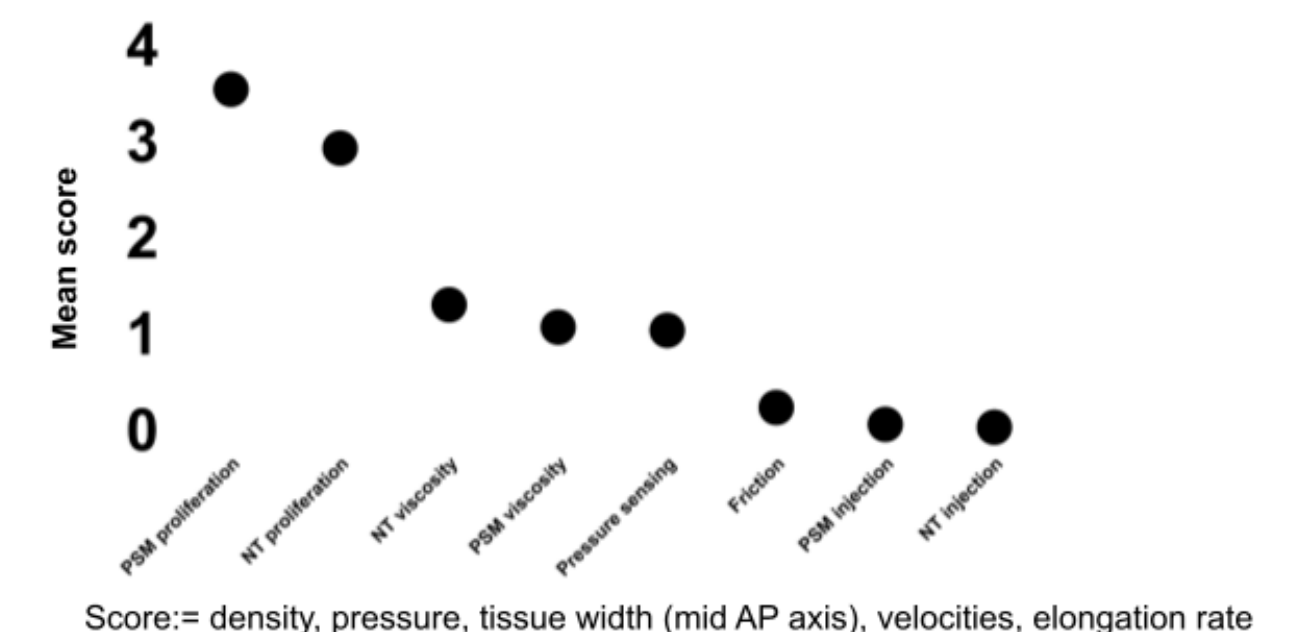
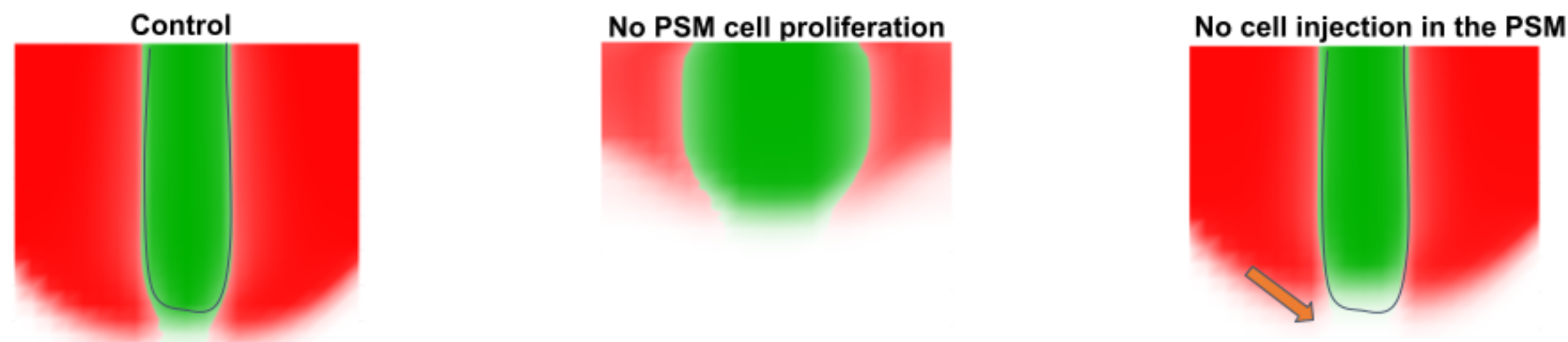
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Output of the model: we perform a sensitivity analysis based on relevant outputs characterizing shapes and velocities of the tissues. This analysis reveals the underestimated importance of (differential) tissue proliferation in the embryogenesis.



Mathematical analysis: a few words

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Can we compute the incompressible limit ?

$$\epsilon \rightarrow 0, \quad m \rightarrow \infty \quad \mathbf{and} \quad \alpha \rightarrow 0$$

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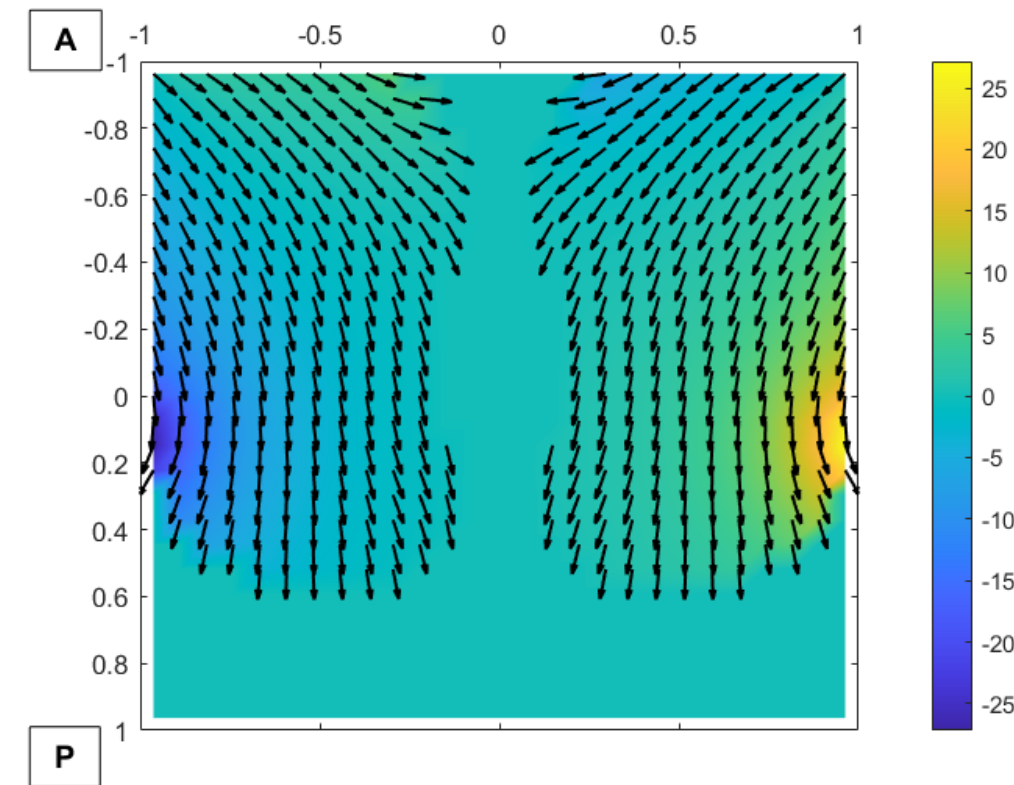
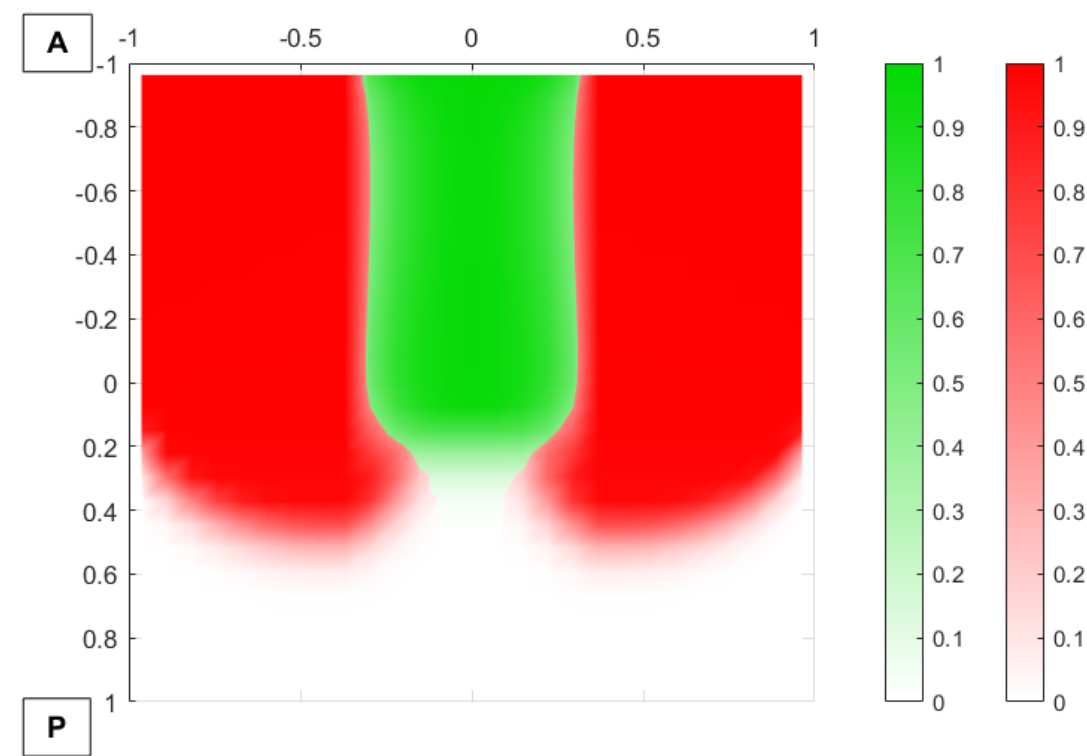
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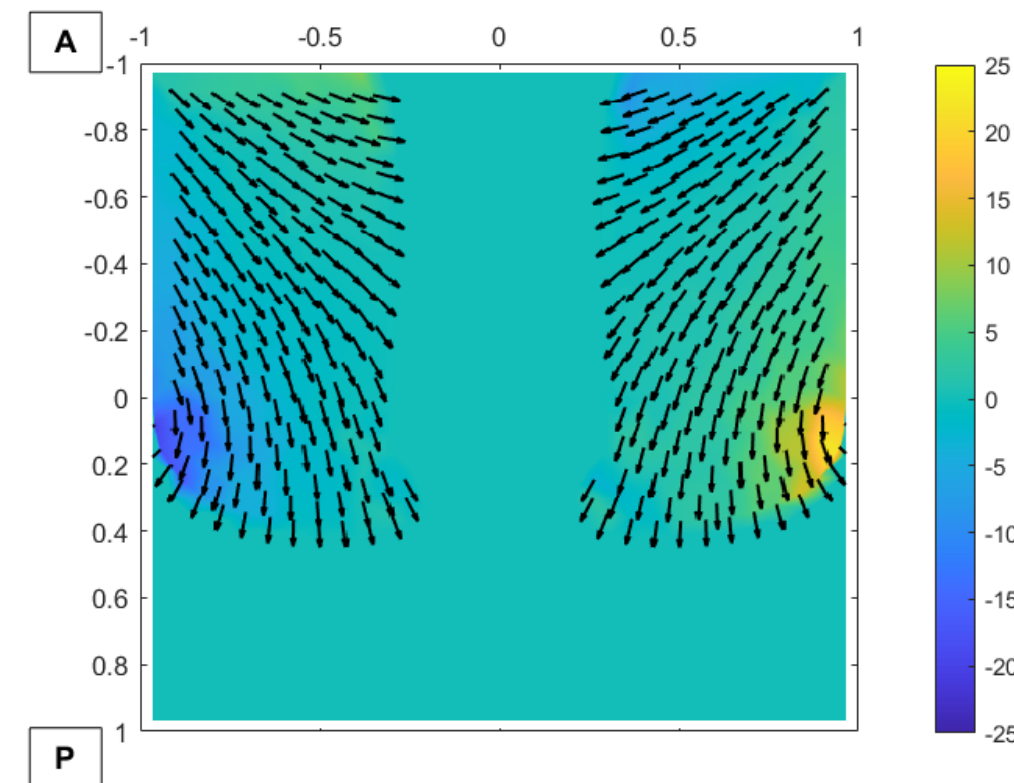
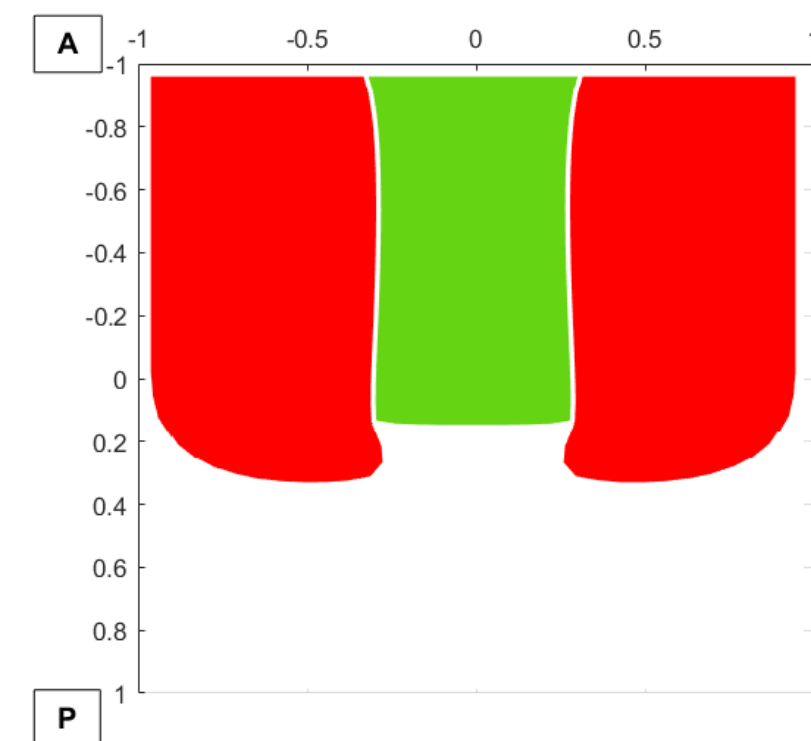
$$\epsilon \rightarrow 0, \quad m \rightarrow \infty \quad \mathbf{and} \quad \alpha \rightarrow 0$$

Formally, YES.

Simulation before and at the limit



Fluid model,
Passive segregation



Geometrical model,
Passive segregation

Analysis at the incompressible limit

Under simplifying assumptions (in a bounded domain in 2D, stationary case, linearly decreasing growth function, smooth subdomains, given segregation pressure in L^2), and under an *ellipticity condition*, we prove:

- well-posedness (weak solution in H^1)
- smoothness inside each subdomain
- equivalence with a transmission problem
- pressure jump at the interface (when the segregation pressure is null)

Transmission problem

$$\text{on } \Omega_1 : (T_1) \begin{cases} -\beta_1 \Delta v_1 + v_1 - \frac{1}{g_1} \nabla \nabla \cdot v_1 = 0, \\ -\beta_2 \Delta v_2 + v_2 - \frac{1}{g_1} \nabla \nabla \cdot v_1 = 0, \end{cases}$$

$$\text{on } \Omega_2 : (T_2) \begin{cases} -\beta_1 \Delta v_1 + v_1 - \frac{1}{g_2} \nabla \nabla \cdot v_2 = 0, \\ -\beta_2 \Delta v_2 + v_2 - \frac{1}{g_2} \nabla \nabla \cdot v_2 = 0, \end{cases}$$

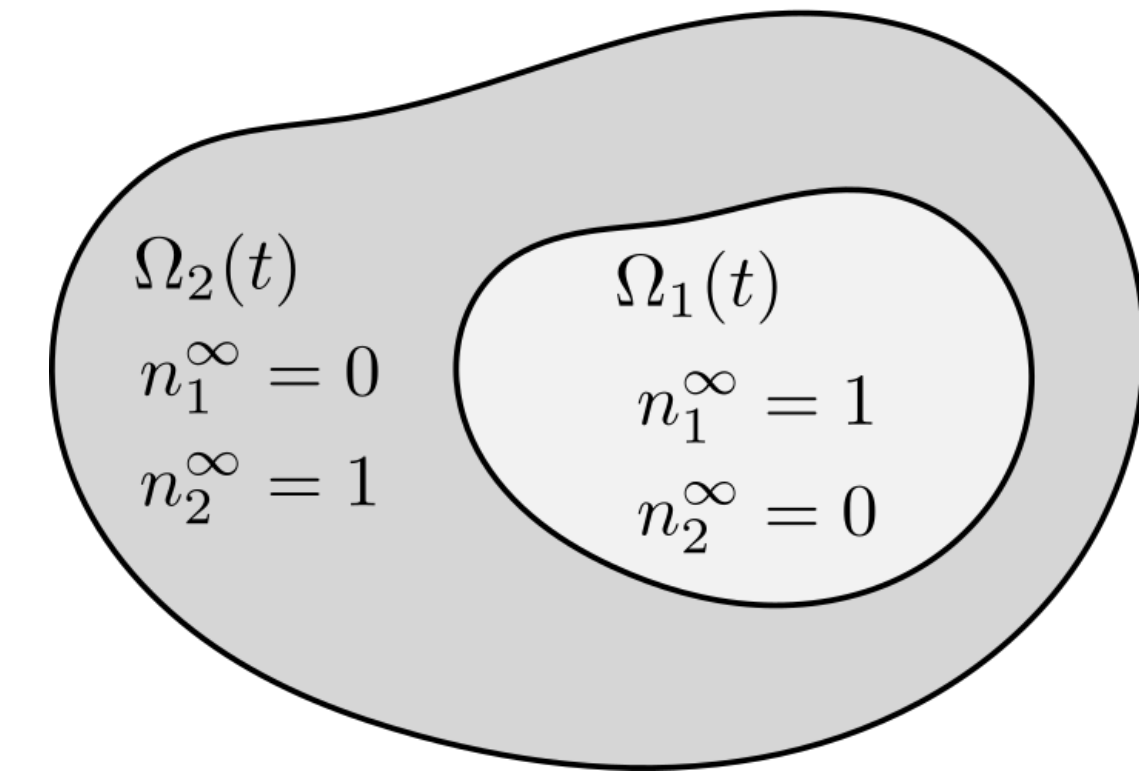
$$\text{on } \Omega^c : (T_3) \begin{cases} -\beta_1 \Delta v_1 + v_1 = 0, \\ -\beta_2 \Delta v_2 + v_2 = 0. \end{cases}$$

$$\text{on } \Gamma_1 : (C_1) \begin{cases} \beta_1 [(\nabla v_1)_{\Omega_1} - (\nabla v_1)_{\Omega^c}] \cdot \vec{\nu} = [p_1^* - \frac{1}{g_1} (\nabla \cdot v_1)_{\Omega_1}] \vec{\nu}, \\ \beta_2 [(\nabla v_2)_{\Omega_1} - (\nabla v_2)_{\Omega^c}] \cdot \vec{\nu} = [p_1^* - \frac{1}{g_1} (\nabla \cdot v_1)_{\Omega_1}] \vec{\nu}, \\ (v_1)_{\Omega_1} = (v_1)_{\Omega^c}, \quad (v_2)_{\Omega_1} = (v_2)_{\Omega^c}, \end{cases}$$

$$\text{on } \Gamma_2 : (C_2) \begin{cases} \beta_1 [(\nabla v_1)_{\Omega_2} - (\nabla v_1)_{\Omega^c}] \cdot \vec{\nu} = [p_2^* - \frac{1}{g_2} (\nabla \cdot v_2)_{\Omega_2}] \vec{\nu}, \\ \beta_2 [(\nabla v_2)_{\Omega_2} - (\nabla v_2)_{\Omega^c}] \cdot \vec{\nu} = [p_2^* - \frac{1}{g_2} (\nabla \cdot v_2)_{\Omega_2}] \vec{\nu}, \\ (v_1)_{\Omega_2} = (v_1)_{\Omega^c}, \quad (v_2)_{\Omega_2} = (v_2)_{\Omega^c}, \end{cases}$$

$$\text{on } \Gamma : (C_3) \begin{cases} \beta_1 [(\nabla v_1)_{\Omega_1} - (\nabla v_1)_{\Omega_2}] \cdot \vec{\nu} = [(p_1^* - p_2^*) + \frac{1}{g_2} (\nabla \cdot v_2)_{\Omega_2} - \frac{1}{g_1} (\nabla \cdot v_1)_{\Omega_1}] \vec{\nu}, \\ \beta_2 [(\nabla v_2)_{\Omega_1} - (\nabla v_2)_{\Omega_2}] \cdot \vec{\nu} = [(p_1^* - p_2^*) + \frac{1}{g_2} (\nabla \cdot v_2)_{\Omega_2} - \frac{1}{g_1} (\nabla \cdot v_1)_{\Omega_1}] \vec{\nu}, \\ (v_1)_{\Omega_1} = (v_1)_{\Omega_2}, \quad (v_2)_{\Omega_1} = (v_2)_{\Omega_2}, \quad v_1 \cdot \vec{\nu} = v_2 \cdot \vec{\nu}. \end{cases}$$

$$\Omega^c(t) \quad p^\infty = 0$$



The rigorous incompressible limit

For the passive segregation model:

- recently proven for viscous tissues *in the whole domain*

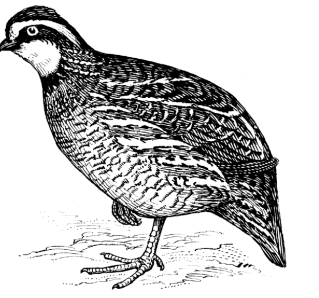
refs: [Perthame Vauchelet 2015] (one tissue); [Dębiec Schmidtchen 2020] and [Dębiec Perthame Schmidtchen Vauchelet 2021] (multi-species, *same viscosities*)

- still open in a bounded domain
- still open for different viscosities

The active segregation model is another challenge..!

Conclusion

Conclusion



Summary

- We developed multi-tissue mechanical models with *passive* or *enforced* segregation
- We derived from it (formally) a geometrical model
- We analyzed the geometrical model: stationary state, pressure jump
- We simulated the model and compared it with biological data

Biological conclusions

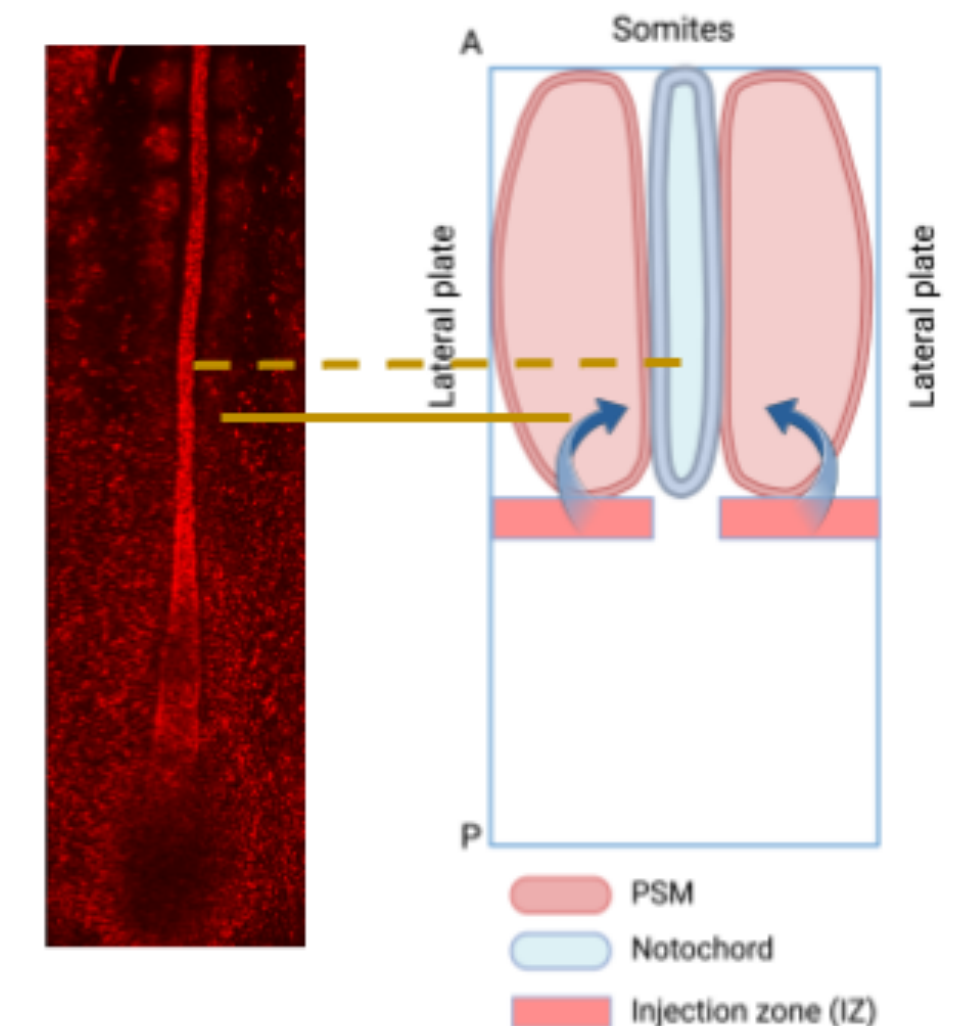
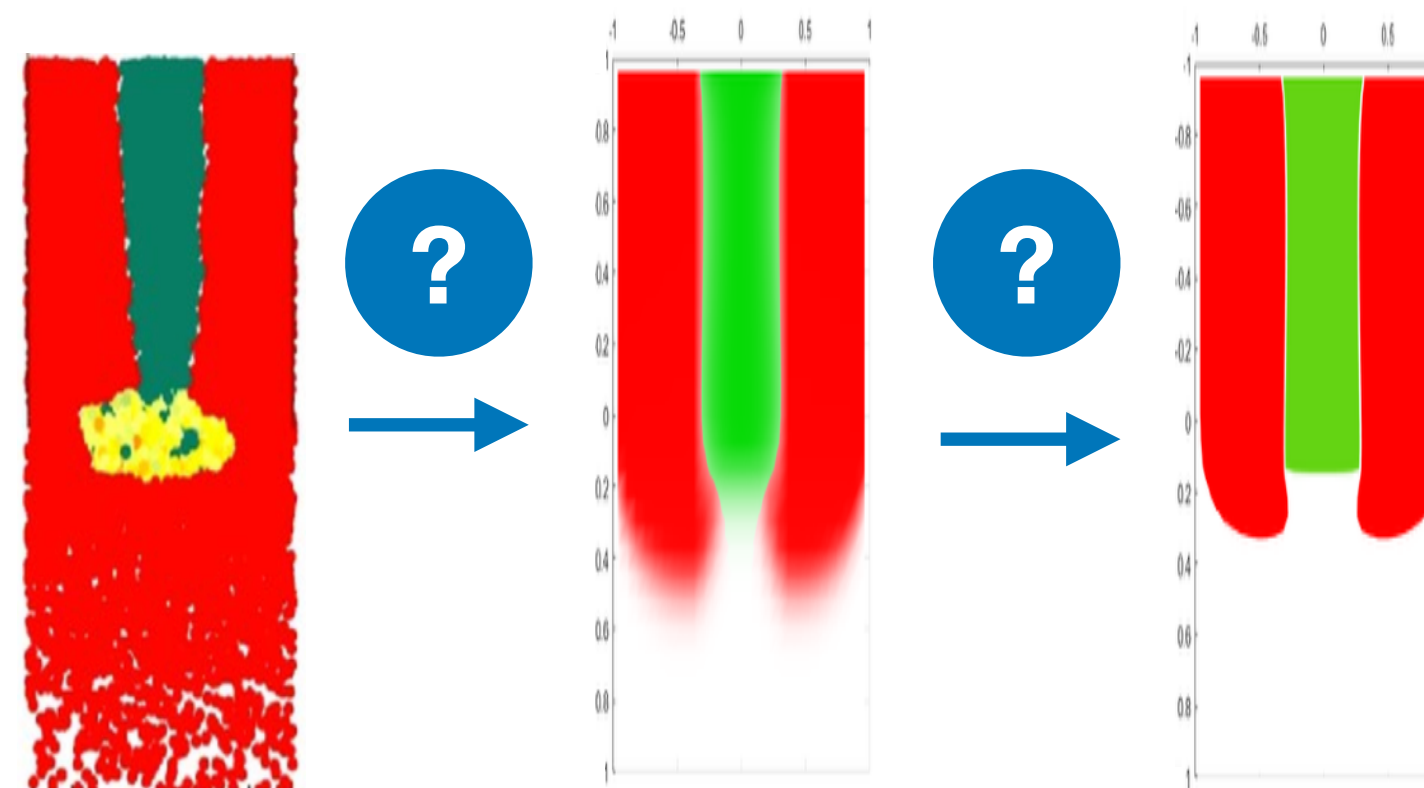
- Suggests the existence of a segregation force
- Highlights the underestimated role of differential tissue proliferation rates

A detailed black and white illustration of a quail bird, likely a Japanese quail, shown in profile facing left. The bird has a rounded body, a small head with a prominent eye, and a short, thick beak. Its plumage is intricately patterned with dark spots and streaks over a lighter background. The legs are short and stout. The bird is standing on a small patch of ground indicated by simple lines.

-

- Suggests the existence of a segregation force
- Highlights the underestimated role of differential tissue proliferation rates

- More tissues, 3D, ...
- Diverse species
- Link with microscopic models
- Rigorous incompressible limit

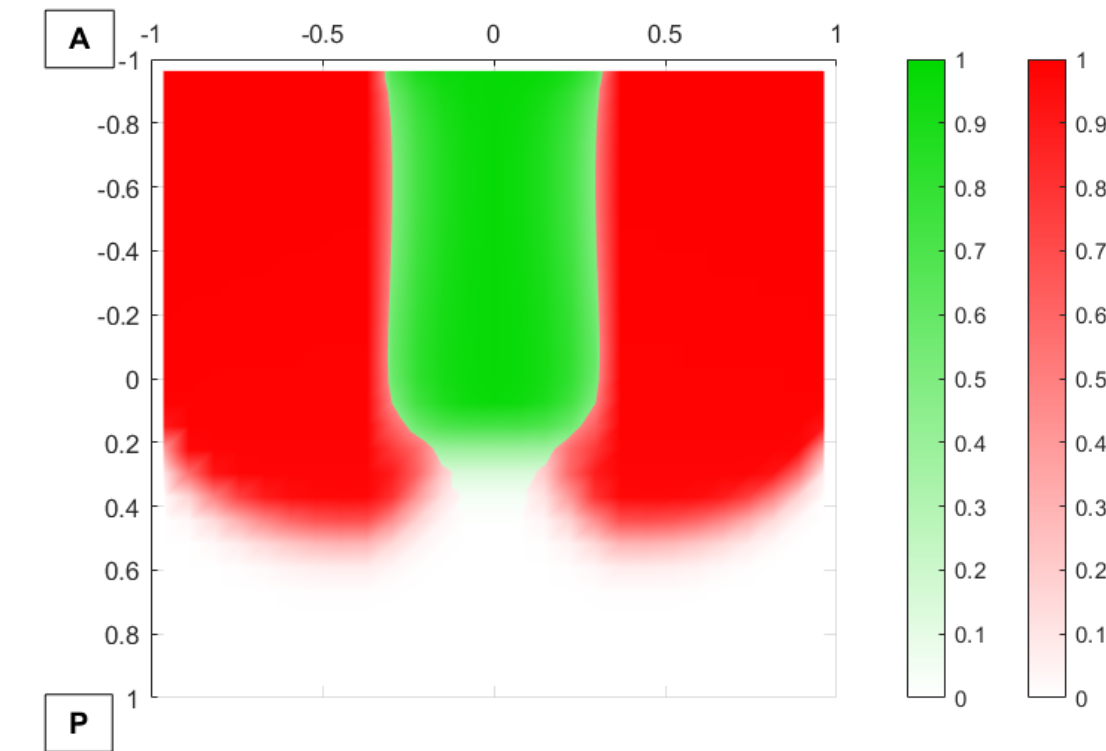


Thank you for your attention

Numerical schemes

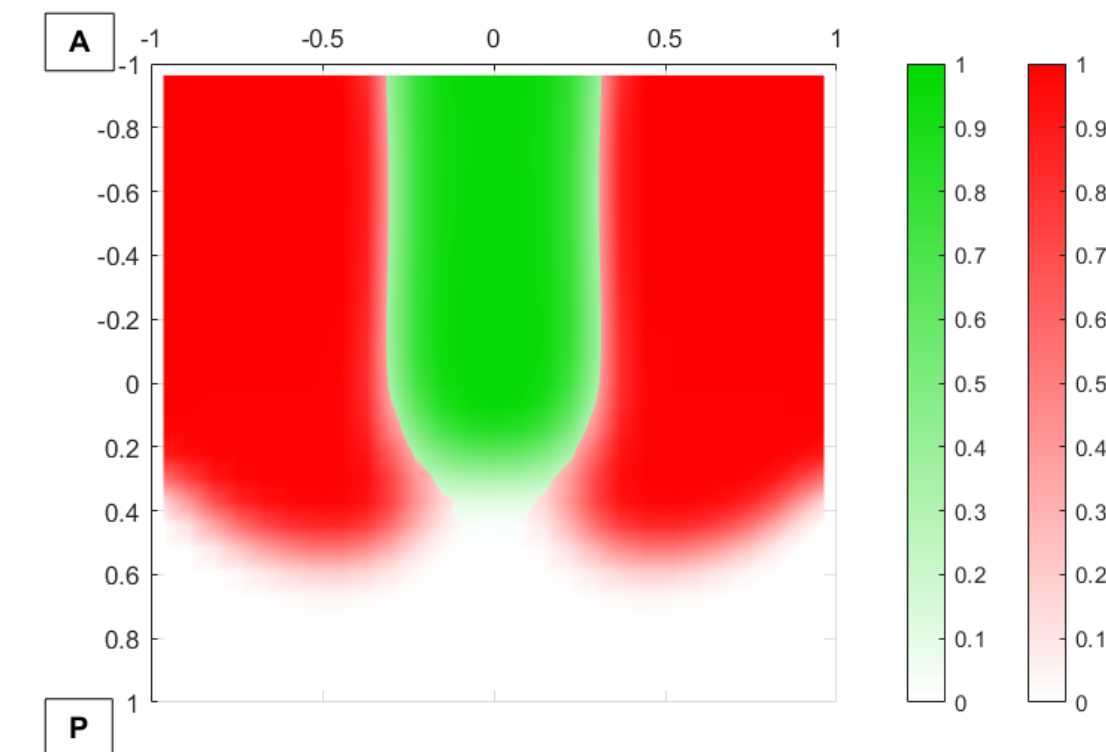
Mechanical model (1):

Finite volume semi-implicit scheme - Matlab



Mechanical model (2):

Finite volume semi-implicit scheme, with a relaxation method (order reduction) - Matlab



Geometrical model:

Finite elements - Freefem++

