University of Strasbourg Master CSMI Academic Year 2024/2025



Mathematical Modeling and Simulations of Organ-on-Chip (OOC)

Internship Presentation

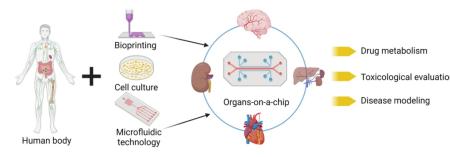
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August 21, 2025

Organ-on-Chip: Context and Motivation

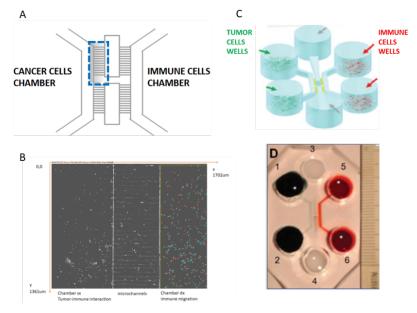
 What is OOC? A 3D microfluidic platform that mimics human tissue microenvironments to simulate cellular dynamics and interactions.



Motivation:

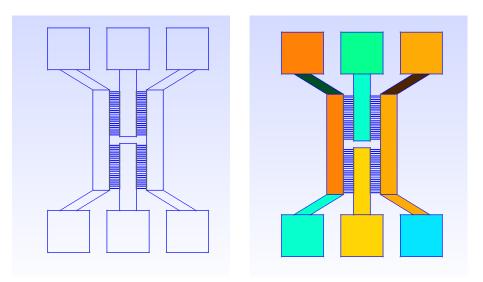
- Understand the mechanisms governing cell dynamics in microfluidic systems.
- Investigate interactions between immune and tumor cells.

Illustration of the Organ-on-Chip Device and Geometries



Typical

Illustration of the Organ-on-Chip device and geometries



Typical Organ-on-Chip geometry: microchannels and chambers(made with GMSH tool)

Goal and Applications

Goal:

- Develop numerical tools to simulate cellular dynamics (e.g., chemotaxis) in Organ-on-Chip systems.
- Build a mathematical model based on chemotaxis, adapted to complex OOC geometries (2D chambers and 1D channels);
- Use Hybrid Discontinuous Galerkin (HDG) method or feelpp toolboxes for solving the models, focusing on stability and accuracy;
- Test and validate the method on simple analytical cases and realistic OOC configurations;
- Explore improvements such as adaptive meshes and integration of experimental data for model calibration;

• Applications:

- Test biological hypotheses.
- Optimize experimental conditions.
- Predict cellular behaviors.

Why Use the HDG Method?

- **Context:** Discontinuous Galerkin (DG) methods are powerful for solving nonlinear hyperbolic systems, offering:
 - High-order accuracy for convection and diffusion.
 - Flexibility on arbitrary meshes.
 - Simple boundary condition handling.
 - Good parallelization and adaptivity.
- **Limitation:** DG methods are often computationally expensive due to many globally coupled unknowns.
- HDG Solution:
 - Reduces global degrees of freedom compared to DG.
 - Provides optimal approximation for both primal and flux variables.
 - Enables local postprocessing for improved accuracy and conservation.

HDG Method: Overview and Stabilization

- Hybridizable Discontinuous Galerkin (HDG):
 - Solves PDEs on polyhedral meshes, reduces global unknowns to numerical trace p̂_h ∈ M_h.
 - Local spaces: $\mathbf{V}(K) = [P_k(K)]^n$, $W(K) = P_k(K)$, $M_h = P_k(\mathcal{E}_h)$.
- Formulation:

$$\begin{split} (\boldsymbol{\Lambda} \mathbf{u}_h, \mathbf{v})_{\mathcal{T}_h} - (p_h, \nabla \cdot \mathbf{v})_{\mathcal{T}_h} + \langle \hat{p}_h, \mathbf{v} \cdot \mathbf{n} \rangle_{\partial \mathcal{T}_h} &= 0, \\ \langle \hat{\mathbf{u}}_h \cdot \mathbf{n}, \mu \rangle_{\partial \mathcal{T}_h \setminus \Gamma} &= 0, \\ \hat{\mathbf{u}}_h &= \mathbf{u}_h + \tau (p_h - \hat{p}_h) \mathbf{n}. \end{split}$$

- Stabilization:
 - Stabilization parameter τ affects convergence:
 - Table of convergence rates:

au	\mathbf{u}_h	p_h	$ar{p}_h$	k
O(h)	k+1	k	k+2	≥ 1
O(1)	k+1	k+1	k+2	≥ 1
O(1/h)	k	k+1	k+1	≥ 1

HDG Methods for Convection-Diffusion Problems

- Problem Setup:
 - Second-order elliptic problem on polyhedral domain $\Omega \subset \mathbb{R}^n$:

$$\begin{cases} \Lambda \mathbf{u} + \nabla p - \Lambda \alpha p = \mathbf{0}, & \text{in } \Omega, \\ \nabla \cdot \mathbf{u} + dp = f, & \text{in } \Omega, \end{cases}$$

- Boundary conditions: $p = h_D$ on Γ_D , $\mathbf{u} \cdot \mathbf{n} = h_N$ on Γ_N .
- Notations:
 - Mesh \mathcal{T}_h , faces $\mathcal{E}_h = \mathcal{E}_h^o \cup \mathcal{E}_h^{\partial}$.
 - Spaces: V_h , W_h for flux u_h and scalar p_h ; M_h for numerical trace \hat{p}_h .
- Weak Formulation on Elements:

$$(\Lambda \mathbf{u}_h, \mathbf{v})_K - (p_h, \nabla \cdot \mathbf{v})_K - (\Lambda \alpha p_h, \mathbf{v})_K + \langle \hat{p}_h, \mathbf{v} \cdot \mathbf{n} \rangle_{\partial K} = 0, - (\mathbf{u}_h, \nabla w)_K + \langle \hat{\mathbf{u}}_h \cdot \mathbf{n} w \rangle_{\partial K} + (dp_h, w)_K = (f, w)_K.$$

• Foundation for HDG: Local solvers on each element, global coupling via \hat{p}_h .

HDG Methods for Convection-Diffusion Problems

Matrix Formulation and Static Condensation

- **Goal:** Reduce the HDG system to a linear system for the numerical trace \hat{p}_h (static condensation).
- System with Numerical Flux:

$$\begin{split} (\Lambda \mathbf{u}_h, \mathbf{v})_{\mathcal{T}_h} - (p_h, \nabla \cdot \mathbf{v})_{\mathcal{T}_h} - (\Lambda \alpha p_h, \mathbf{v})_{\mathcal{T}_h} + \langle \hat{p}_h, \mathbf{v} \cdot \mathbf{n} \rangle_{\partial \mathcal{T}_h} = 0, \\ (\nabla \cdot \mathbf{u}_h, w)_{\mathcal{T}_h} + \langle \tau p_h, w \rangle_{\partial \mathcal{T}_h} + (dp_h, w)_{\mathcal{T}_h} - \langle \tau \hat{p}_h, w \rangle_{\partial \mathcal{T}_h} = (f, w)_{\mathcal{T}_h}. \end{split}$$

- Static Condensation Process:
 - Eliminate \mathbf{u}_h and p_h locally on each element K.
 - Solve globally for \hat{p}_h .
 - Reconstruct \mathbf{u}_h and p_h element-wise using local equations.

HDG: Boundary Conditions, Local and Global solvers

Dirichlet Boundary Conditions:

- On Γ_D : $\langle \hat{p}_h, \mu \rangle_{\Gamma_D} = \langle h_D, \mu \rangle_{\Gamma_D}$.
- Compute L^2 -projection of h_D onto $M_h|_{\Gamma_D}$ to set \hat{p}_h values.

Neumann Boundary Conditions:

• On Γ_N : Use numerical flux to contribute to the global system via $\langle h_N, \mu \rangle_{\Gamma_N}$.

Global System Assembly:

- Enforce flux continuity across interior faces.
- Assemble global matrix \mathbb{M} and vector **F** from local D^K , D_f^K .
- System: $\mathbb{M}\widehat{\mathbf{p}} = \mathbf{F} + \mathbf{G}_N$.
- Solve for \hat{p}_h , then recover \mathbf{u}_h , p_h locally.

Keller-Segel Model

Mathematical Model

- Variables: Describes chemotaxis: cell density u, chemoattractant φ .
- Equations:

$$\begin{cases} \partial_t u - \nabla \cdot (\mu \nabla u - \chi u \nabla \varphi) = f_u, \\ \partial_t \varphi - \nabla \cdot (\nu \nabla \varphi) + a \varphi - b u = f_\varphi \end{cases} \begin{cases} \nu \nabla \varphi \cdot n = 0, \\ \mu (\nabla u - \chi u \nabla \varphi) \cdot n = 0 \end{cases} \text{ on } \partial \Omega$$

• Auxiliary Variables:

$$\begin{cases} \boldsymbol{j} = -\mu \nabla \boldsymbol{u} + \chi \boldsymbol{u} \nabla \varphi \\ \boldsymbol{\psi} = -\nu \nabla \varphi \end{cases} \qquad \begin{cases} \boldsymbol{j} \cdot \boldsymbol{n} = 0, \\ \boldsymbol{\psi} \cdot \boldsymbol{n} = 0 \end{cases} \text{ on } \partial \Omega$$

• Expected:

- Preserve non-negativity of u, φ .
- Capture blow-up phenomenon in test cases.
- Mass conservation under zero-flux conditions.

Keller-Segel Model: HDG Discretization

• Numerical fluxes:

$$\begin{cases} \widehat{j}^{K} \cdot \mathbf{n}^{K} = j^{K} \cdot \mathbf{n}^{K} + \gamma_{u}(\mathbf{u}^{K} - \widehat{\mathbf{u}}^{K}) \\ \widehat{\psi}^{K} \cdot \mathbf{n}^{K} = \psi^{K} \cdot \mathbf{n}^{K} + \gamma_{\varphi}(\varphi^{K} - \widehat{\varphi}^{K}) \end{cases}$$

• Stabilization parameters:

$$\gamma_u = \frac{ au_D}{h} + au_C$$
, $au_C = \max(\psi \cdot n, 0)$, au_D constant $= 1$ or 10

Problem: Find $U_h = (u_h, \widehat{u_h}, j_h) \in V_h$ and $\Phi_h = (\varphi_h, \widehat{\varphi_h}, \psi_h) \in S_h$ such that, $\forall W = (w, \widehat{w}, q) \in V_h$ and $\Theta = (\tau, \widehat{\tau}, \theta) \in S_h$, it holds:

$$\left(\frac{\partial u_h}{\partial t}, w\right)_{\Omega} + (\widehat{j_h} \cdot n, w)_{\mathcal{E}_h} - (j_h, \nabla w)_{\mathcal{T}_h} = (f_u, w)_{\Omega},$$

$$\left(\frac{\partial \varphi_h}{\partial t}, \tau\right)_{\Omega} + (\widehat{\psi_h} \cdot n, \tau)_{\mathcal{E}_h} - (\psi_h, \nabla \tau)_{\mathcal{T}_h} + (a\varphi_h, \tau)_{\Omega} - (bu_h, \tau)_{\Omega} = (f_{\varphi}, \tau)_{\Omega},$$

$$(\beta \psi_h, \theta)_{\Omega} - (\varphi_h, \nabla \cdot \theta)_{\mathcal{T}_h} + (\widehat{\varphi_h}, \theta \cdot n)_{\mathcal{E}_h} = 0,$$

$$(\alpha j_h + \beta \chi u_h \psi_h, q)_{\Omega} - (u_h, \nabla \cdot q)_{\mathcal{T}_h} + (\widehat{u_h}, q \cdot n)_{\mathcal{E}_h} = 0,$$

OOC model: One chamber model Description

- Domain: $\Omega = [0, 500] \times [0, 1000] (\mu m)$.
- Variables:
 - u(x, y, t): Density of immune cells.
 - v(x, y, t): Density of tumor cells.
 - $\varphi(x, y, t)$: Chemotactic factor secreted by immune cells.
 - $\psi(x, y, t)$: Chemotactic factor secreted by tumor cells.
- Main equations:

$$\begin{split} \partial_t v - \nabla \cdot (D_v \nabla v) + \lambda(\varphi) v &= 0, \\ \partial_t u - \nabla \cdot (D_u \nabla u - \chi u \nabla \psi) &= 0, \\ \partial_t \varphi - \nabla \cdot (D_\varphi \nabla \varphi) + \beta \varphi - \alpha u &= 0, \\ \partial_t \psi - \nabla \cdot (D_\psi \nabla \psi) + a_\psi \psi - b_\psi v &= 0, \end{split}$$

• $\lambda(\varphi) = \frac{k_1 \varphi}{k_2 + \varphi}$: Tumor destruction by immune cells.

Coefficient descriptions and modelling Choices

Symbol	Description	Value	Units
$\frac{1}{D_{V}}$	Tumor-cell diffusivity	5.6×10^{1}	$\mu \mathrm{m}^2/\mathrm{s}$
D_{μ}	Immune-cell diffusivity	2.0×10^{2}	$\mu \mathrm{m}^2/\mathrm{s}$
D_{φ}	Diffusivity of φ	$9.0 imes 10^2$	$\mu \mathrm{m}^2/\mathrm{s}$
$D_{\psi}^{^{r}}$	Diffusivity of $\dot{\psi}$	$9.0 imes 10^2$	$\mu \mathrm{m}^2/\mathrm{s}$
χ	Chemotactic sensitivity	$10^6 - 10^{10}$	$M\mu \mathrm{m}^2/(\mathrm{s}\cdot\mathrm{cell})$
α	Production rate of φ per immune cell	$1.0 imes 10^{-1}$	
β	Decay rate of φ	$1.0 imes 10^{-4}$	s^{-1}
a_{ψ}	Decay rate of ψ	$1.0 imes 10^{-4}$	s^{-1}
b_{ψ}	Production rate of ψ per tumor cell	$1.0 imes 10^{-1}$	$s^{-1}\cdotcell^{-1}$
k_{arphi_1}, k_{arphi_2}	Killing parameters	1.0, 1.0	
k_1	cellular drift velocity	1.0	$M\cdot \mu$ m^2/s
k_2	receptor dissociation constant	1.0	М
k_3	drift velocity	1.56×10^{10}	$M \cdot \mu$ m^2/s
L_x, L_y	chambers dimensions	500, 1000	$ m \mu m$
c_L	Channel length	600	$ m \mu m$
c_w	Channel width	12	$\mu\mathrm{m}$
c_d	Width beetween channels	100	$\mu\mathrm{m}$

Initial and Boundary Conditions and Experimental Setup

- Initial Conditions:
 - Gaussian distributions:

$$v(0,x,y) = 10^{-3} \exp\left(-\frac{x^2 + (y - 500)^2}{1000}\right) + 10^{-3} \exp\left(-\frac{x^2 + (y - 1000)^2}{1000}\right)$$

$$+ 10^{-3} \exp\left(-\frac{x^2 + y^2}{1000}\right),$$

$$u(0,x,y) = 5.10^{-3} \exp\left(-\frac{(x - 400)^2 + (y - 500)^2}{1000}\right)$$

$$+ 4.10^{-3} \exp\left(-\frac{(x - 400)^2 + (y - 900)^2}{1000}\right) + 2.10^{-3} \exp\left(-\frac{(x - 400)^2}{1000}\right)$$

$$\phi(0,x,y) = 0, \quad \psi(0,x,y) = 0.$$

• **Boundary Conditions:** Homogeneous Neumann ($\partial\Omega$): $\nabla v \cdot n = 0$, $\nabla u \cdot n = 0$, $\nabla \phi \cdot n = 0$, $\nabla \psi \cdot n = 0$.

• Mesh: Uniform triangulation of $[0,500] \times [0,1000]$ with $h=5 \,\mu\text{m}$.

• Time Integration: Θ -scheme with $\Delta t = 1\,\mathrm{s}$, up to $T = 10\,000\,\mathrm{s}$.

OOC model: Two chamber model Description

- **Domains:** Ω^1 , Ω^2 : chambers,
- Governing equations in chambers $(\Omega^*, * = 1, 2)$:

$$\begin{split} \partial_t v^* - \nabla \cdot \left(D_v^* \nabla v^* \right) + \lambda^* (\varphi^*) v^* &= 0, \\ \partial_t u^* - \nabla \cdot \left(D_u^* \nabla u^* - \chi u^* \nabla \psi^* \right) &= 0, \\ \partial_t \varphi^* - \nabla \cdot \left(D_\varphi^* \nabla \varphi^* \right) + \beta \varphi^* - \alpha u^* &= 0, \\ \partial_t \psi^* - \nabla \cdot \left(D_\psi^* \nabla \psi^* \right) + a_\psi^* \psi^* - b_\psi^* v^* &= 0, \end{split}$$

• Governing equations in the channel (ω) :

$$\begin{split} \partial_t v^c - \partial_s \big(D_v^c \partial_s v^c \big) + & \lambda^c \big(\varphi^c \big) v^c = 0, \\ \partial_t u^c - \partial_s \big(D_u^c \partial_s u^c - \chi \partial_s \psi^c u^c \big) = 0, \\ \partial_t \varphi^c - \partial_s \big(D_\varphi^c \partial_s \varphi^c \big) + \beta \varphi^c - \alpha u^c = 0, \\ \partial_t \psi^c - \partial_s \big(D_\psi^c \partial_s \psi^c \big) + a_\psi^c \psi^c - b_\psi^c v^c = 0, \end{split}$$

• Boundary conditions: Homogeneous Neumann on $\partial \Omega^* \setminus \gamma^*$:

$$\begin{cases} D_{v}^{*} \nabla v^{*} \cdot \mathbf{n}^{*} = 0, \\ (D_{u}^{*} \nabla u^{*} - \chi u^{*} \nabla \psi^{*}) \cdot \mathbf{n}^{*} = 0, \end{cases} \begin{cases} D_{\varphi}^{*} \nabla \varphi^{*} \cdot \mathbf{n}^{*} = 0, \\ D_{\psi}^{*} \nabla \psi^{*} \cdot \mathbf{n}^{*} = 0. \end{cases}$$

Chemotactic Sensitivity and Killing Functions

• Basic model: directional movement up a spatial gradient of chemoattractant

$$\chi := k_3$$

with drift velocity k_3 (see table 14).

 Receptor saturation: dependence on concentration of chemoattractant in a cell's local environment

$$\chi(\varphi) := k_1 \frac{1}{(k_2 + \varphi)^2}$$

where k_1 is the cellular drift velocity and k_2 the receptor dissociation constant (see table 14).

• Overcrowding: ability to move freely reduces at high densities

$$\left|\chi(u,arphi) := k_1 rac{1}{(k_2 + arphi)^2} \left(1 - rac{u}{u_{\mathsf{max}}}
ight)
ight|$$

Killing function: for the tumor cells, the killing function is defined as

$$oxed{\lambda(arphi) := rac{k_{arphi_1}arphi}{k_{arphi_2} + arphi}}$$

Initial and Boundary Conditions and Experimental Setup

- Initial Conditions:
 - Gaussian distributions:

$$v(0,x,y) = 10^{-3} \exp\left(-\frac{x^2 + (y - 500)^2}{1000}\right) + 10^{-3} \exp\left(-\frac{x^2 + (y - 1000)^2}{1000}\right)$$

$$+ 10^{-3} \exp\left(-\frac{x^2 + y^2}{1000}\right),$$

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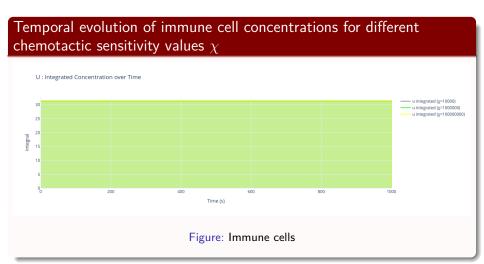
$$+ 4.10^{-3} \exp\left(-\frac{(x - 400)^2 + (y - 900)^2}{1000}\right) + 2.10^{-3} \exp\left(-\frac{(x - 400)^2}{1000}\right)$$

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Results for the one chamber model: Statistics and Visualization



Explanation...

Statistics and Visualization



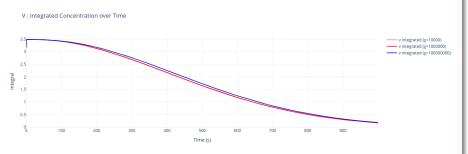


Figure: tumor cells

Explanation...

Initial Distributions: Tumor and Immune Cells

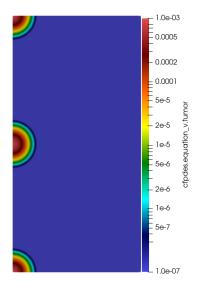


Figure: Tumor v

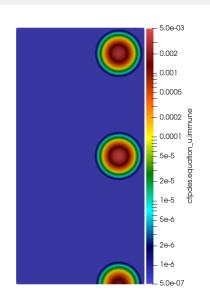


Figure: Immune u

Initial Distributions: Chemoattractants

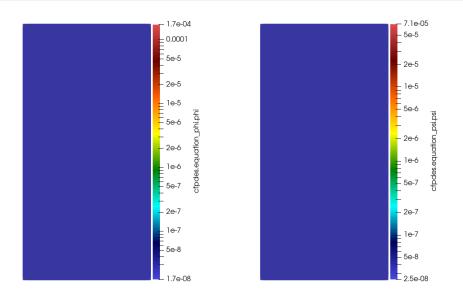


Figure: Chemoattractant φ

Figure: Chemoattractant ψ

Immune Cell Density Evolution (I)

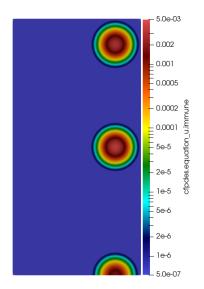


Figure: Immune u at t = 0

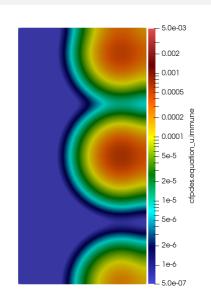


Figure: Immune u at t = 10

Immune Cell Density Evolution (II)

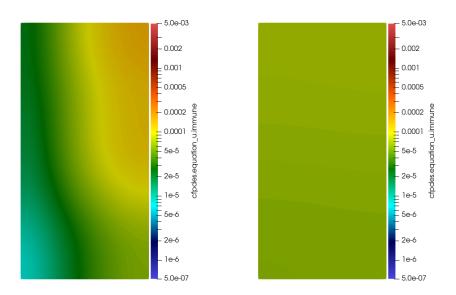


Figure: Immune u at t = 100

Figure: Immune u at t = 1000

Tumor Cell Density Evolution (I)

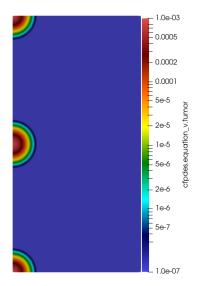


Figure: Tumor v at t = 0

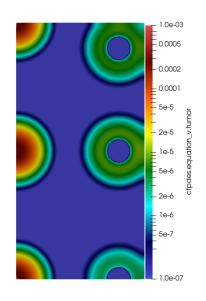


Figure: Tumor v at t = 10

Tumor Cell Density Evolution (II)

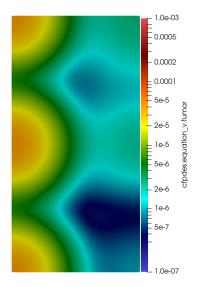


Figure: Tumor v at t = 100

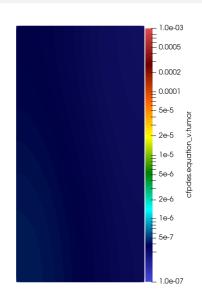
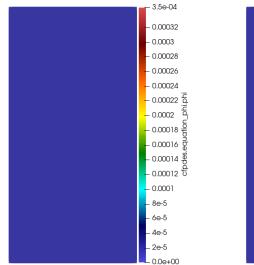


Figure: Tumor v at t = 1000

Chemoattractant φ Evolution (I)



- 3.5e-04 - 0.00032 - 0.0003 - 0.00028 - 0.00026 - 0.00024 _ 0.00022 년 - 0.0002 0.00018 - 0.00016 _ 0.00014 မြို့ _ 0.00012 ပိ - 0.0001 - 8e-5 _ 2e-5 - 0.0e+00

Figure: φ at t = 0

Figure: φ at t=10

Chemoattractant φ Evolution (II)

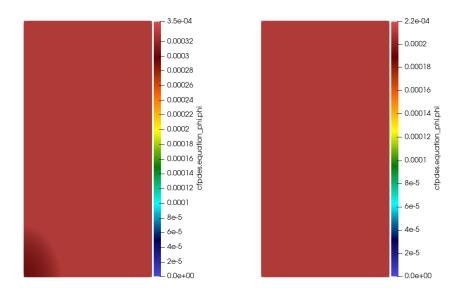


Figure: φ at t = 100

Figure: φ at t = 500

Chemoattractant ψ Evolution (I)

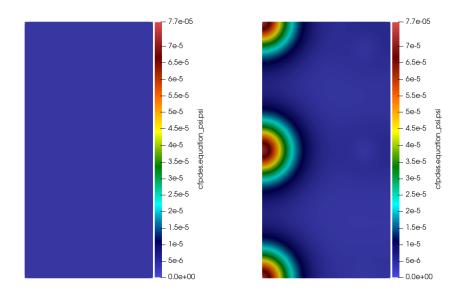


Figure: ψ at t=0

Figure: ψ at t=10

Chemoattractant ψ Evolution (II)

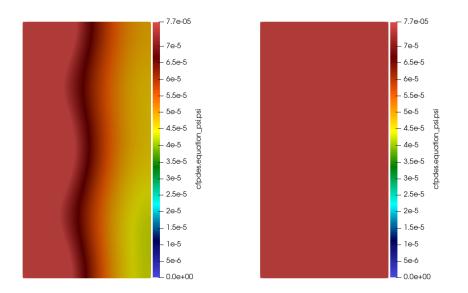


Figure: ψ at t=100 Figure: ψ at t=500

Conclusion and Future Work

Summary:

- Developed a one-chamber model based on diffusion-reaction-chemotaxis equations.
- Simulated cellular and chemical interactions on a rectangular domain (one chamber case), using feelpp toolboxes.

Challenges:

- Extend the model to a two-chamber geometry with HDG coupling.
- Validate results with experimental Organ-on-Chip data.
- Solve the inverse problem for parameter calibration.

Thank you all for your attention. Any questions?