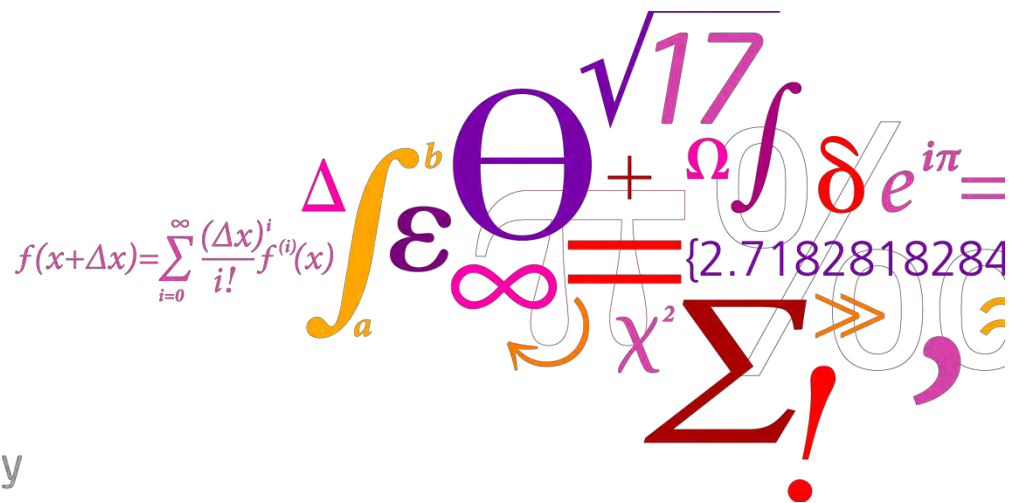


Microfluidic Systems for Cell Growth and Cell Migration Studies

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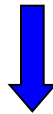
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Use of Comsol

- Expensive and/or time consuming fabrication processes → Need to minimize repeated fabrication runs and test cycles
- Competitive field → Design to product time should be minimized



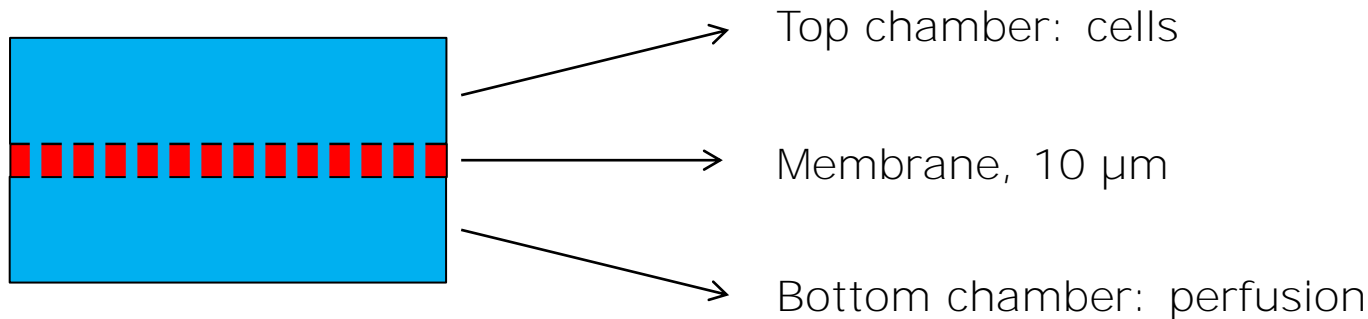
- Comsol contribution:
 - Optimise the performance of existing designs by calculating experimental parameters
 - Design and simulate the performance of new structures
 - Explain the experimental results

Overview

- Geometry and Physics
- Cell culture chambers
 - Design 1
 - Design 2
 - Results
- Cell migration chamber
 - Design
 - Results
- Conclusion and outlook

Geometry and Physics

- Common for all systems:

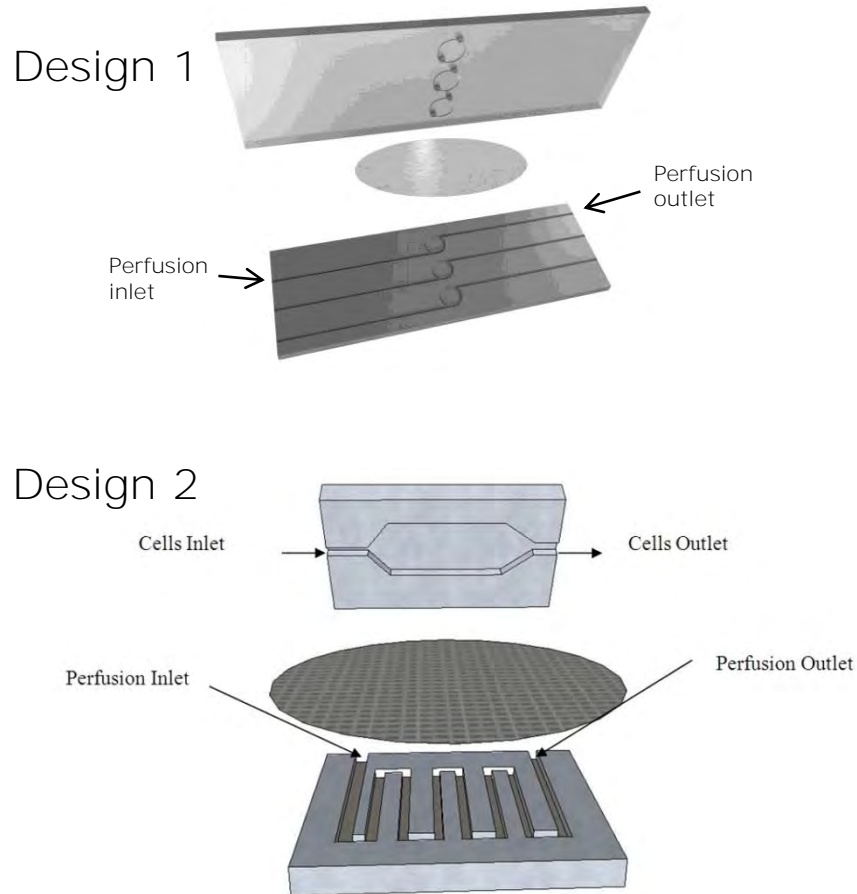


- **Incompressible Navier Stokes** coupled with **Convection and Diffusion**
- Flow solved in steady state and solution stored and used to solve for the concentration with time-dependent solver
- Membrane treated as a separate subdomain controlled by **Darcy's** law for porous flow

The membrane

- 10 μm thick, 5 μm pore size, porosity of 0.14
- In the subdomain settings for the Incompressible Navier Stokes a body force on the liquid was added, given by $\vec{F} = -\alpha \cdot \vec{U}$
- α is a constant calculated from the Darcy number (Da) as $\alpha = \frac{\eta}{\text{Da} \cdot L^2}$
- For the 10 μm thick porous membrane this gives $\alpha = 10^{11}$ kg/(m³s) for a Darcy number of 10^{-4} and a viscosity equal to that of water (0.001 Pa·s)
- Diffusion coefficient for various species was corrected with the membrane porosity
- The validity of the Darcy approximation was tested in a 2D simulation where a geometric approximation of the membrane was designed and simulated

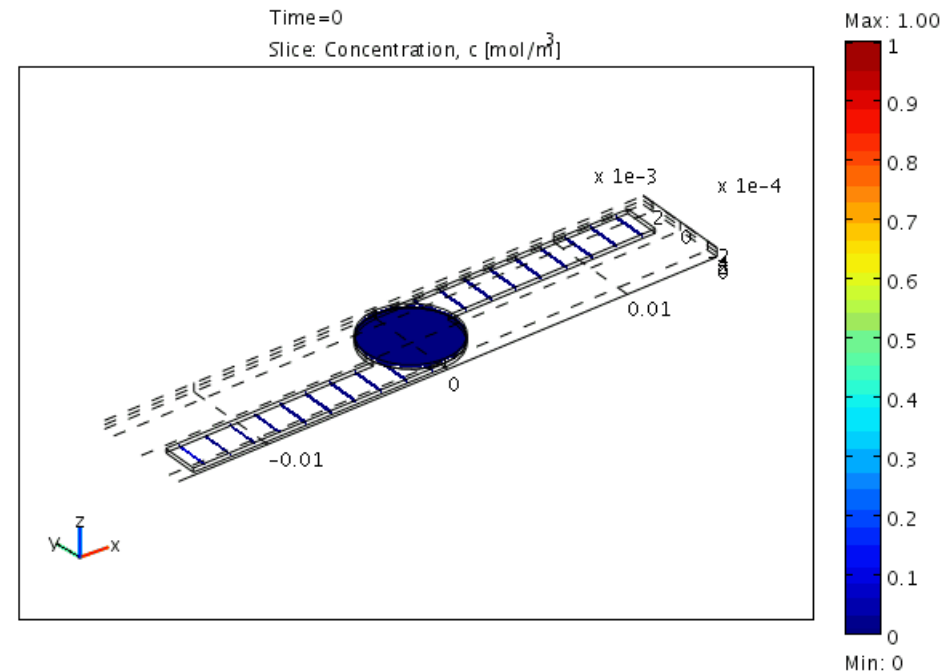
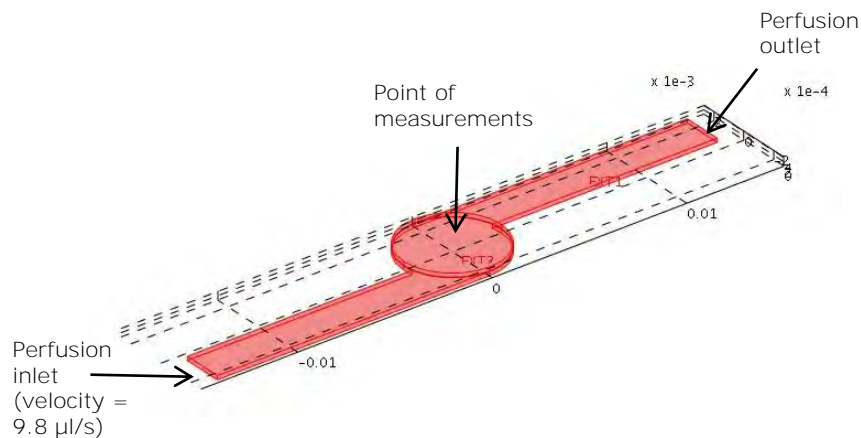
Cell culture chamber



- The two systems function essentially in the same way with some small differences:
 - Loading and unloading of cells by pipette (design 1) as opposed to flow (design 2)
 - Diffusion through large 5 mm circular opening as opposed to a meandering channel 500 μm wide
 - Depth of bottom chamber: 200 μm (design 1), 300 μm (design 2)
 - Depth of top chamber: 200 μm (design 1), 250 μm (design 2)
 - Perfusion inlet velocity: 9.8 $\mu\text{l}/\text{min}$ (design 1), 2.5 $\mu\text{l}/\text{min}$ (design 2)

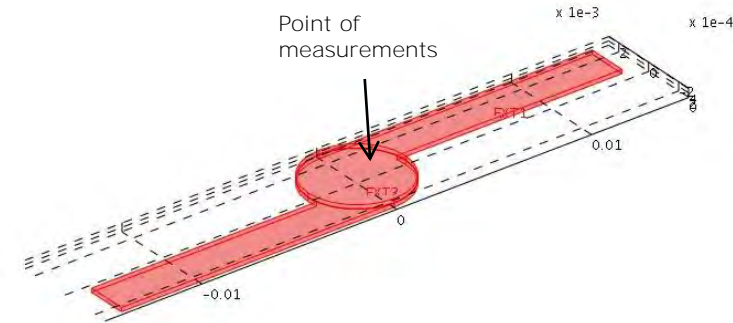
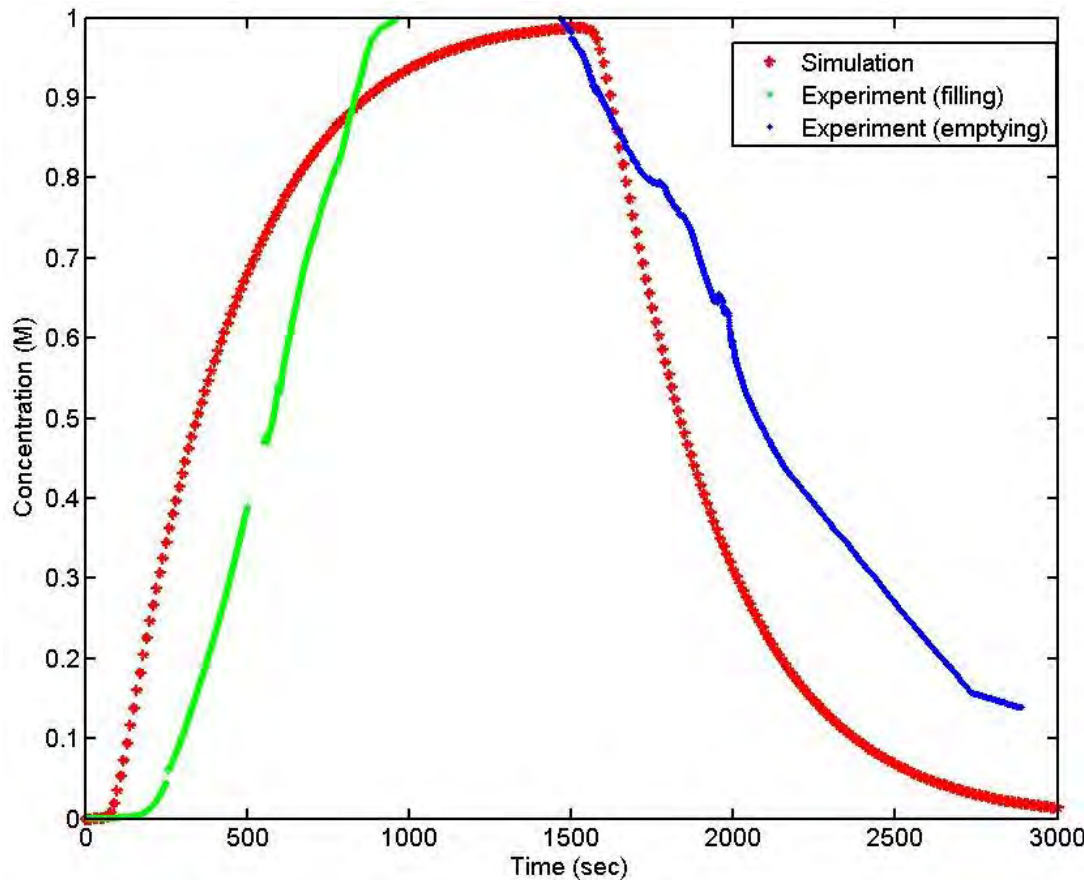
Design 1

- Simulating the filling and emptying of the top chamber with a solution of streptavidin ($D = 130 \mu\text{m}^2/\text{s}$). Boundary condition at perfusion inlet was set to $(t < 1500)$, i.e. 1 M for $t < 1500$ sec and 0 M for $t > 1500$ sec.
- Only part of the inlet and outlet channels were simulated
- Chamber practically at max concentration after 1000 sec and almost empty 1000 sec after solution switch



Results

- Concentration roughly at experimental measurement point (average of simulated concentrations in a volume of $25000 \mu\text{m}^3$ just below the top surface of the chamber towards the outlet)

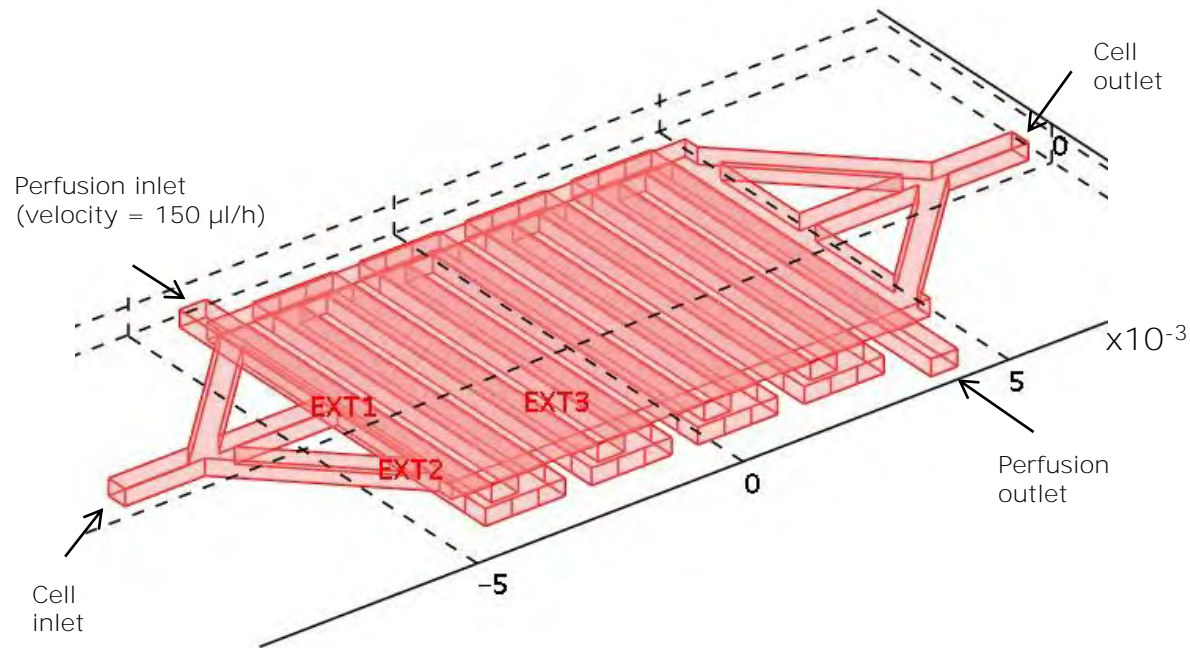


A few discrepancies

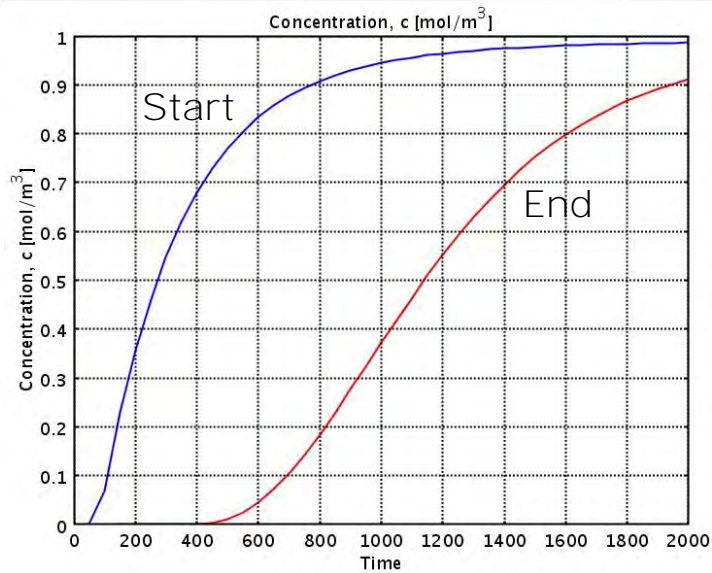
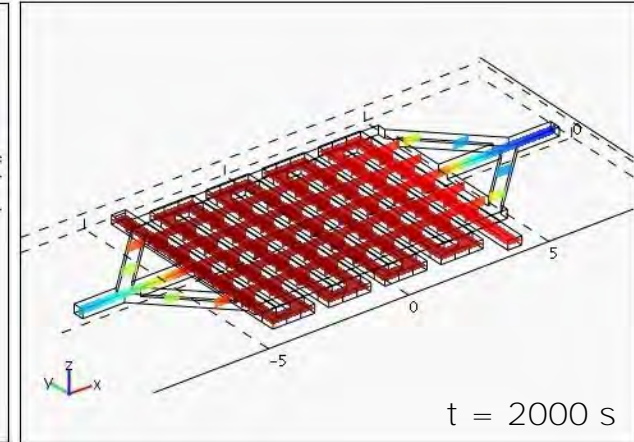
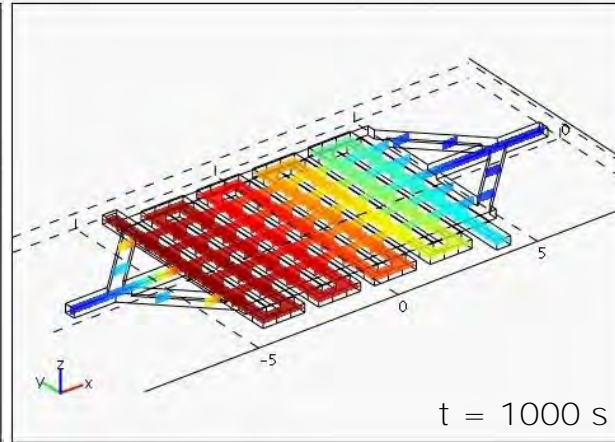
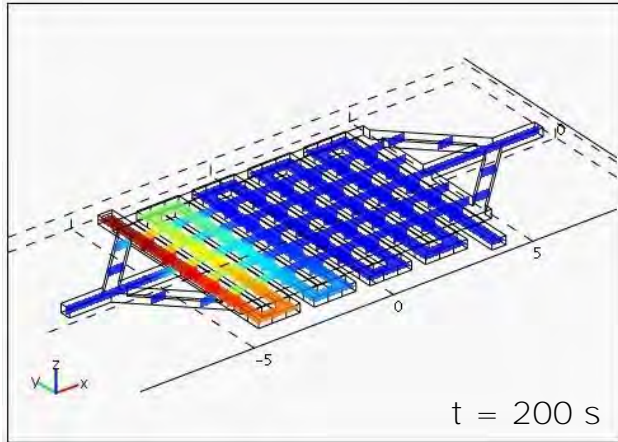
- Fluorescence signal first detectable after a certain concentration has been reached
- Unspecific binding on channel walls
- Uncertainty regarding exact measurement point

Design 2

- Simulating the filling of the top chamber with a solution of KCl ($D = 2000 \mu\text{m}^2/\text{s}$) as part of the cell treatment protocol for cytogenetic analysis



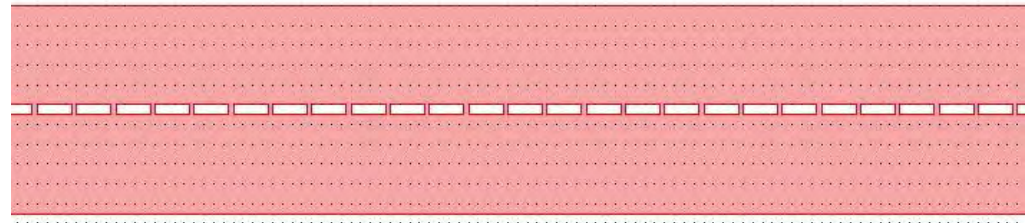
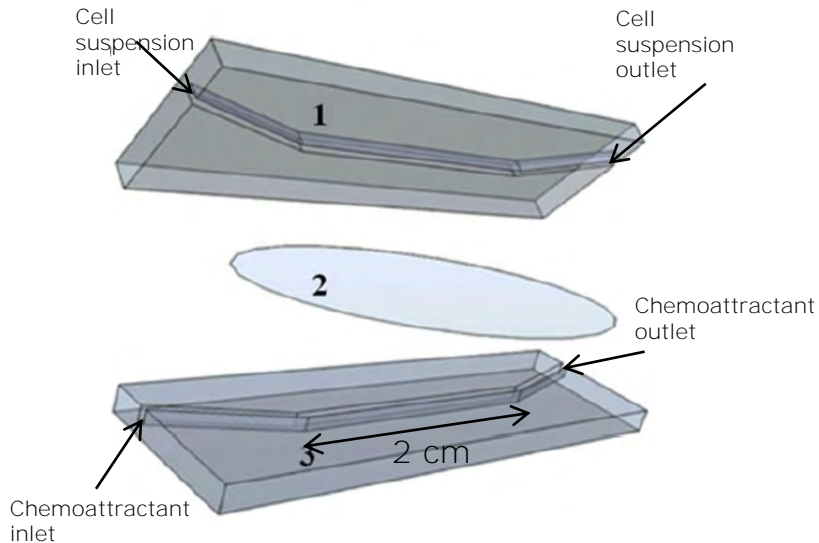
Results



Experimentally 25 min (1500 s) were used for filling the chamber with good results

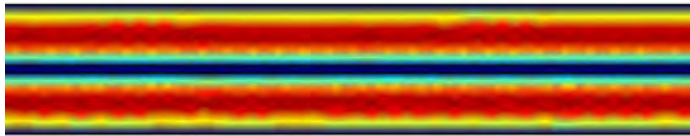
Cell migration chamber

- Parallel flow in the two channels. Inlet velocity at $1.67 \times 10^{-3} \text{ m/s}$
- Diffusion coefficient of chemoattractant calculated to be $1.76 \times 10^{-10} \text{ m}^2/\text{s}$
- Simulation conducted in 2D
 - By treating the membrane as a subdomain with a volume force on the liquid
 - By physically designing a $10 \mu\text{m}$ thick membrane with a large number of $5 \mu\text{m}$ wide holes to achieve a porosity of 0.14 over the 2 cm channel

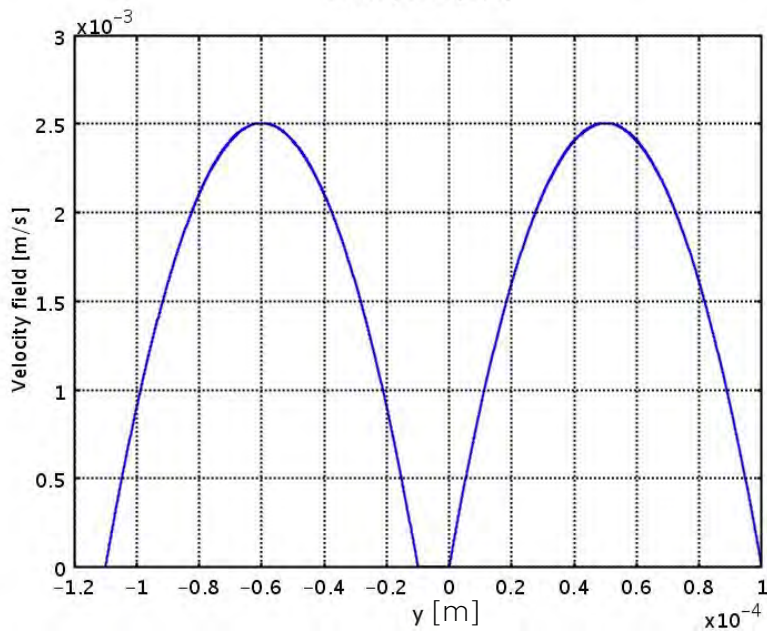


Results – velocity field

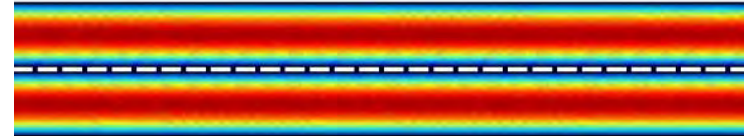
Darcy representation



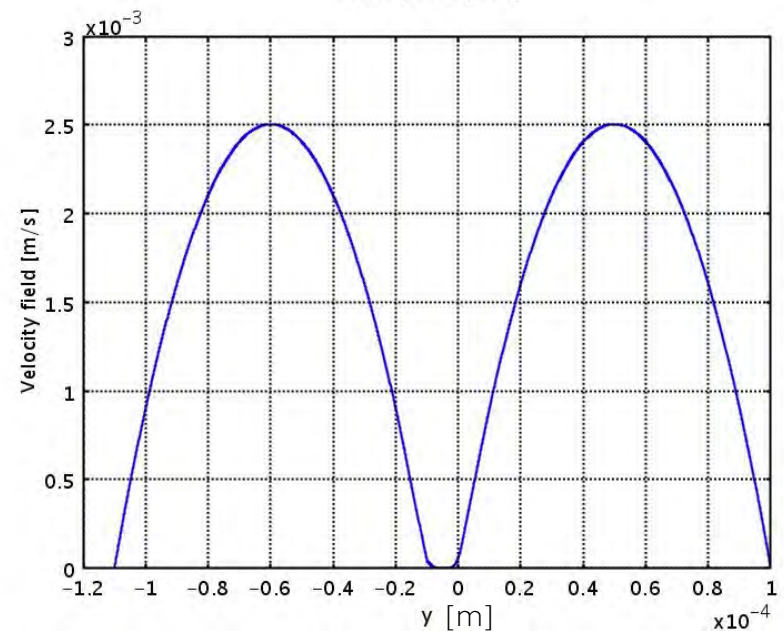
Velocity field [m/s]



Geometrical representation

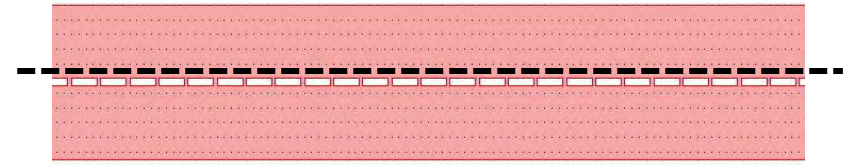


Velocity field [m/s]



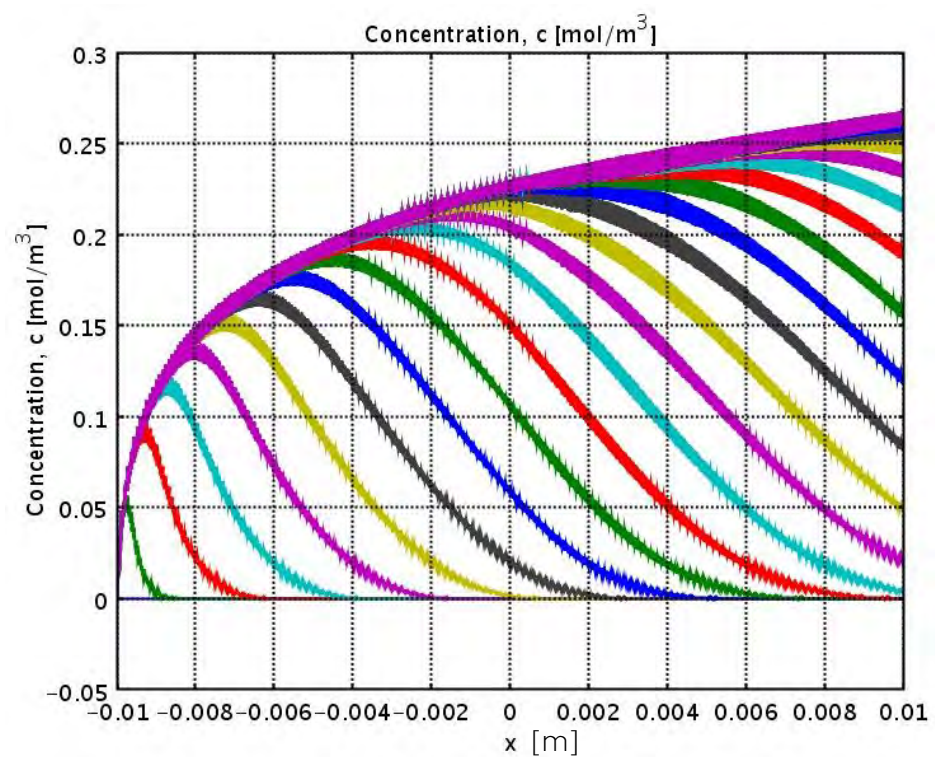
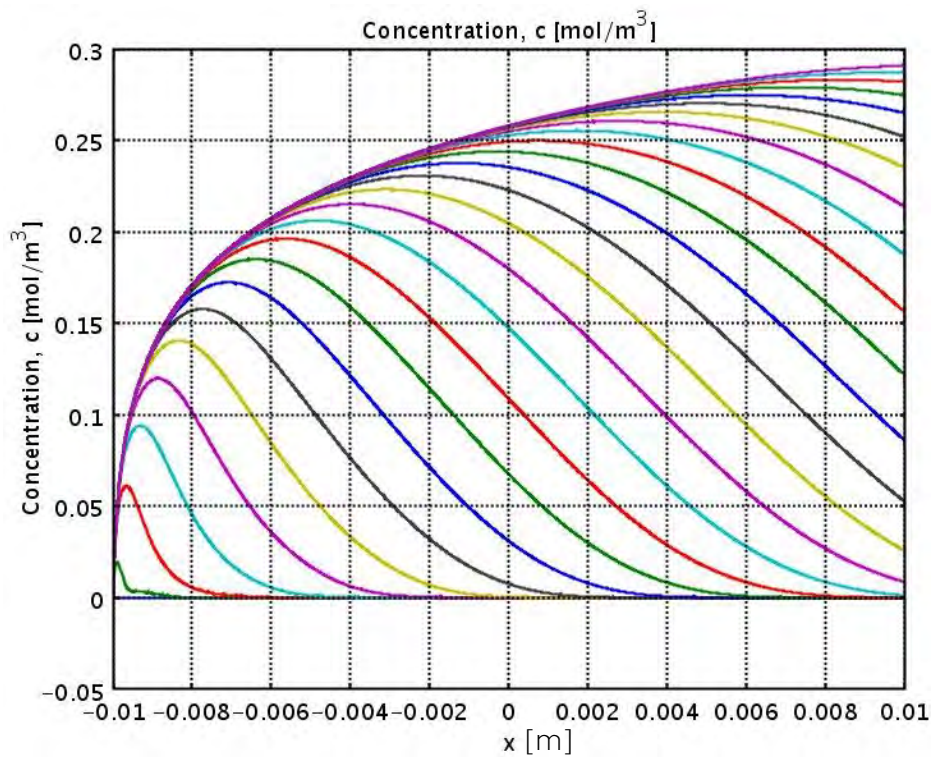
Results - concentration

5 μm above
membrane



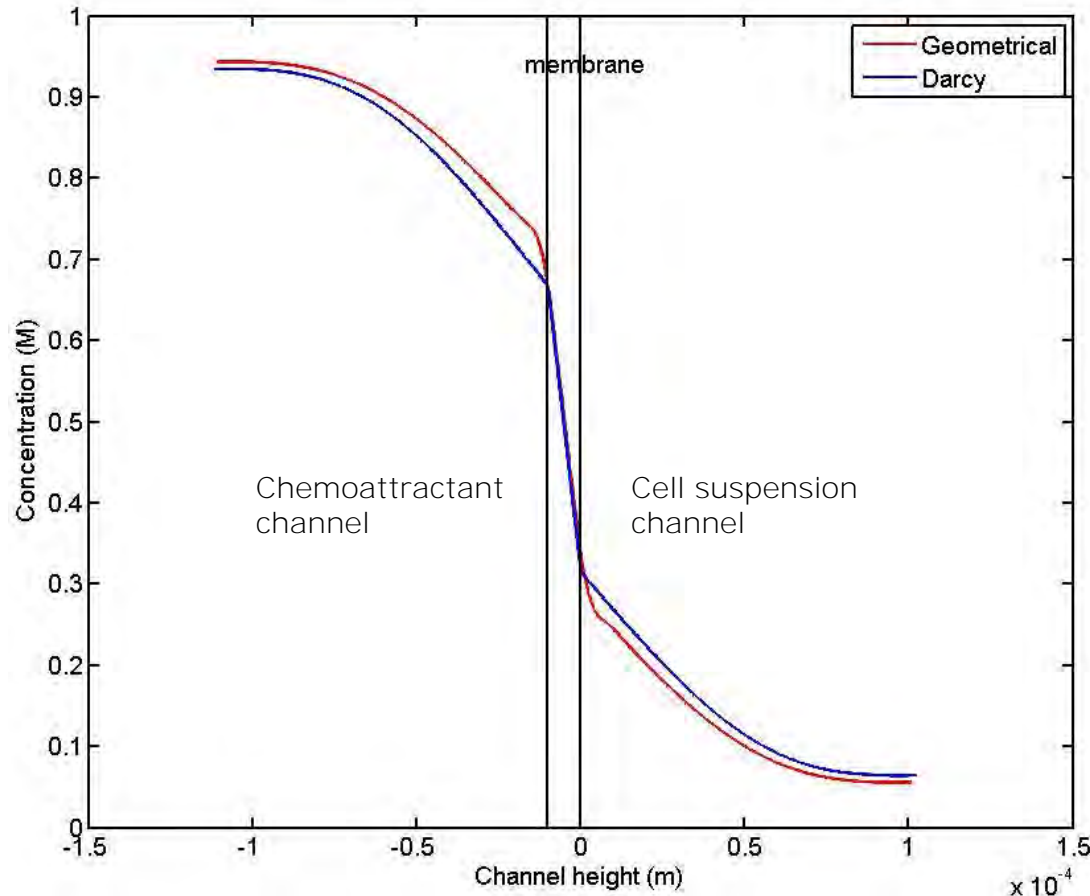
Darcy representation

Geometrical representation



Results - concentration

- Concentration gradient at the end of the channel at steady state ($t = 25$ s)



! The geometrical plot is taken at the middle of the last membrane opening (2 cm from inlet)

! Experimental data confirm that cells only migrate through the membrane close to the channel outlet, where the concentration is the highest

Conclusion and outlook

- Darcy modelling of the membrane gives similar results to those achieved with the geometrical representation of the membrane → Greatly reduces computational needs and geometric complexity
- Experimental results fit mostly with the theoretically predicted ones for all presented systems. Discrepancies can be due to:
 - Uncertainties for various simulation parameters such as diffusion coefficient and viscosity
 - Fabrication accuracy of the experimental device as opposed to the simulated one (micromilling error can be up to $\pm 5-10 \mu\text{m}$)
- The effect of the structural uncertainties needs to be quantified in the future

Thank you for your attention