

Passive and Active Colloidal Chemotaxis in a Microfluidic Channel

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KU LEUVEN



<http://pdebuyl.be/>

Outline

1 Introduction

2 Mesoscopic & stochastic simulations

- Mesoscopic simulation
- Stochastic simulation
- Chemical concentration
- Surface interaction

3 Results

- Passive sphere
- Active sphere
- Nanomotor

4 Conclusions

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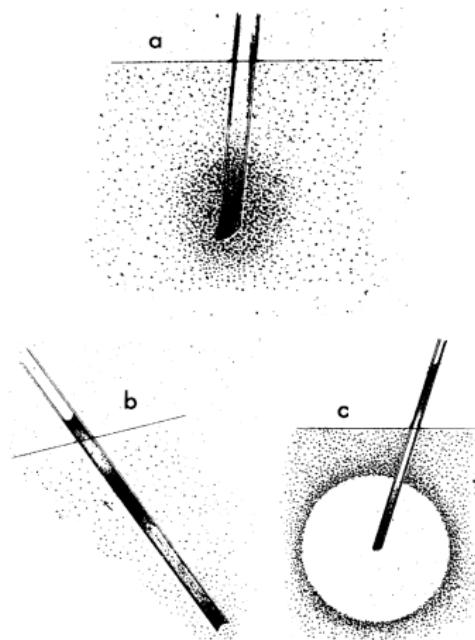
4 Conclusions

Bacterial chemotaxis - *Chromatium okenii*

- Miyoshi (1898) J. Coll. Sci. Imp. Univ. Jap. **10**, 143 (taken from Berg, *E. Coli in Motion*, Springer, 2004)

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Synthetic chemotaxis

Experiments

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Simulations

- Chen *et al* Soft Matter **12**, 1876 (2016)

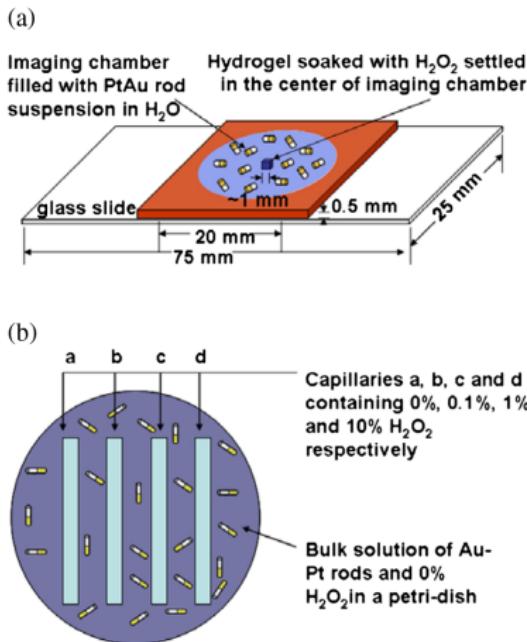
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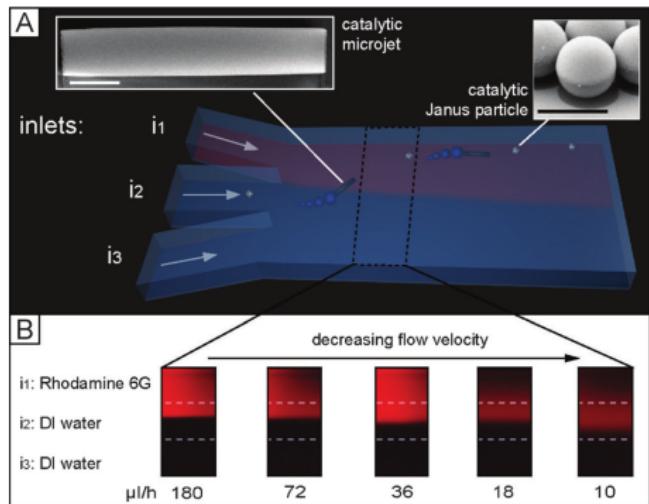
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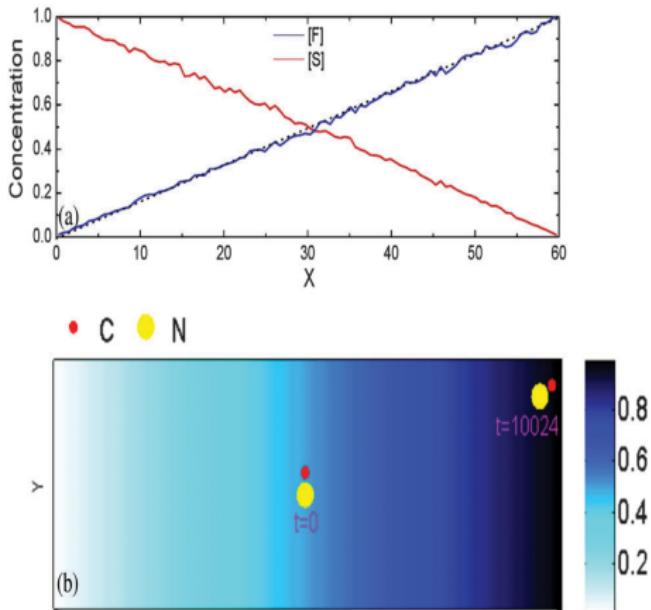
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Enzymatic chemotaxis

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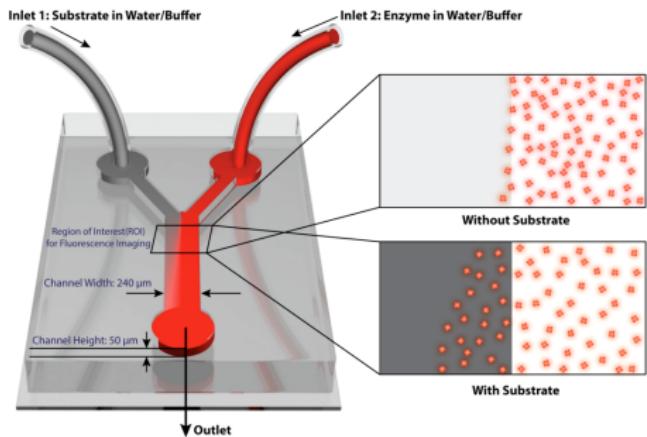
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Objectives

- Understand the mechanism for chemotaxis
 - ▶ For passive and active colloids
 - ▶ Under a flow
- Provide simulation models to explore chemotactic behavior
 - ▶ “Experimental setup”
 - ▶ Chemical activity
 - ▶ Surface interaction
- Lay the foundation for later work on enzyme chemotaxis

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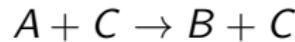
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Mesoscopic simulation

Microfluidic channel

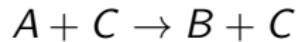
- MPCD fluid (here, close to 1M particles).
- Flow: constant acceleration for the solvent, bounce-back BC and ghost particles in z.
- Gradient device: two inlets for the different chemical species.
- For the colloids: Molecular Dynamics.
- Activity:



Mesoscopic simulation

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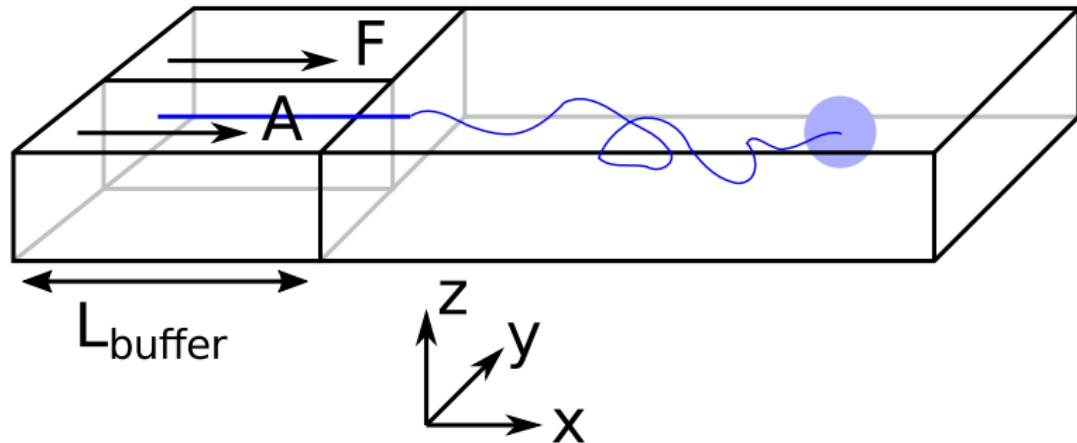
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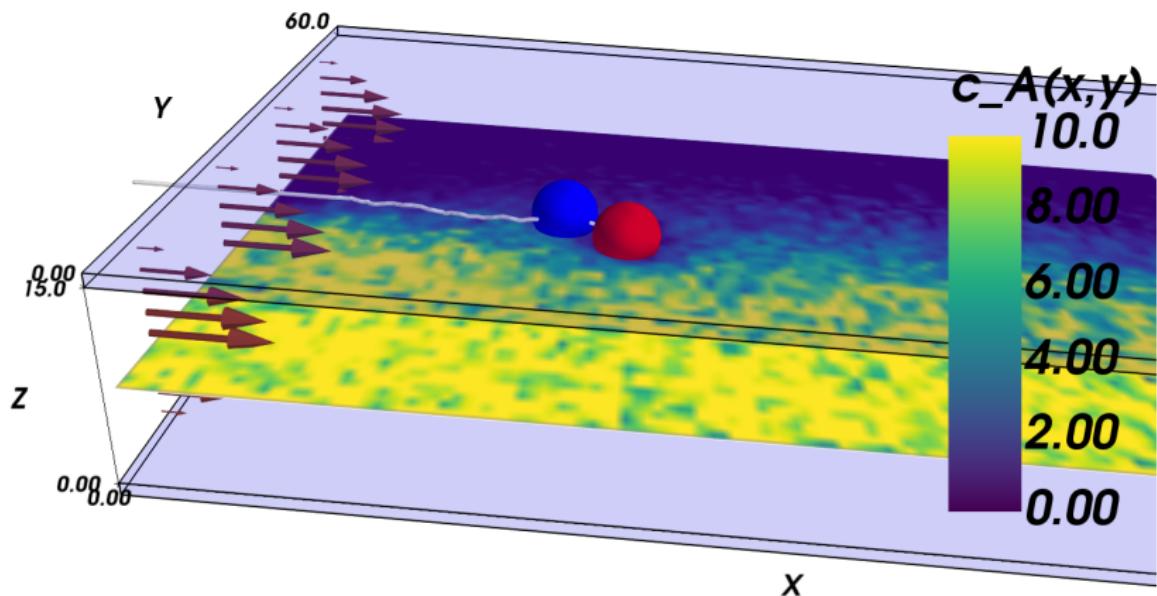
Software

- All simulations were performed with RMPCDMD
<http://lab.pdebuyl.be/rmpcdmd/>
de Buyl *et al*, J. Open Res. Software [5, 3](#) (2017)

Mesoscopic simulation



Mesoscopic simulation



Stochastic simulation

Langevin equation for the sphere

$$\begin{aligned}\dot{x} &= v_{\text{flow}} + \sqrt{2D}\xi_x \\ \dot{y} &= \frac{F_y(x/v_{\text{flow}}, y)}{\gamma} + \sqrt{2D}\xi_y\end{aligned}$$

Langevin equation for the dimer nanomotor

$$\begin{pmatrix} \dot{x} - v_{\text{flow}} \\ \dot{y} \end{pmatrix} = \begin{pmatrix} \cos \phi & -\sin \phi \\ \sin \phi & \cos \phi \end{pmatrix} \begin{pmatrix} \frac{F_{||}}{\gamma_{||}} + \sqrt{2D_{||}}\xi_{||} \\ \frac{F_{\perp}}{\gamma_{\perp}} + \sqrt{2D_{\perp}}\xi_{\perp} \end{pmatrix}$$

$$\dot{\phi} = \frac{\mathcal{T}}{\gamma_r} + \sqrt{2D_r}\xi_{\phi}$$

$F_{||}$ & F_{\perp} : forces on the axes of the dimer. \mathcal{T} : torque on the dimer.

Chemical concentration

- For high Pe, at the center of the channel $z = L_z/2$
Ismagilov et al, Appl. Phys. Lett. [76, 2376 \(2000\)](#)

$$v_{\text{flow}} \partial_x c_\alpha(x, y) = D \partial_y^2 c_\alpha(x, y)$$

- Solution:

$$c_A(x, y) = c_0 \left(1 - \frac{1}{2} \operatorname{erfc} \left(\frac{L_y/2 - y}{\sqrt{4Dx/v_{\text{flow}}}} \right) \right)$$

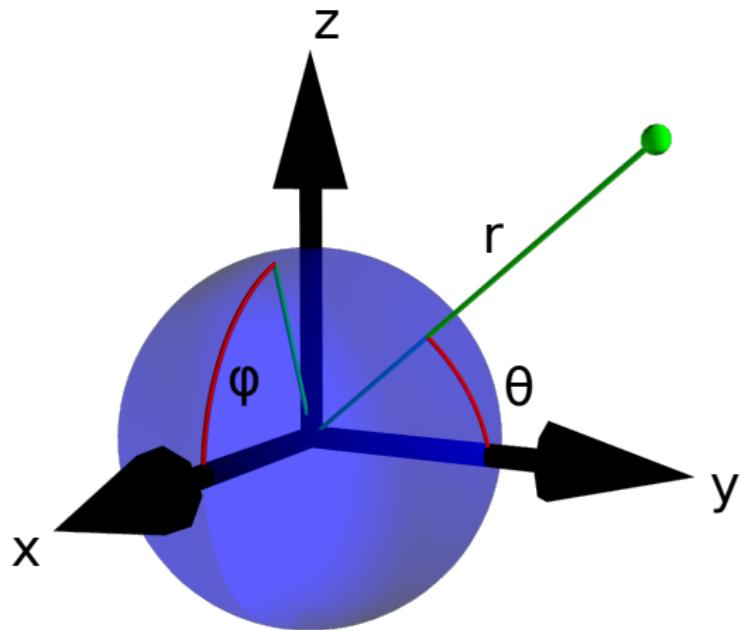
$$c_F(x, y) = c_0 - c_A(x, y)$$

$$c_B(x, y) = 0$$

Chemical concentration

Spherical coordinates

$$\begin{cases} x = r \cos \varphi \sin \theta \\ y = r \cos \theta \\ z = r \sin \varphi \sin \theta \end{cases}$$



Chemical concentration

Ansatz for $c_A(x, y)$

- $\lambda = \partial_y c_A(x, y)$
-

$$c_A = c_0 + c_1 \frac{R}{r} + c_2 \left(\frac{R}{r} \right)^2 \cos \theta + \lambda r \cos \theta$$

Chemical concentration

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$$\begin{cases} c_0 &= c_A(x, y) \\ c_1 &= -\frac{k_0}{k_0+k_D} c_0 \\ c_2 &= -\frac{k_0}{k_0+2k_D} \lambda R \end{cases}$$

Surface interaction

- Methodology used in Rückner and Kapral, Phys. Rev. Lett. [98, 150603 \(2007\)](#)
- Integrating the force over the surface gives

$$\vec{F} = \frac{2}{\beta} \sum_{\alpha} \Lambda_{\kappa,\alpha} \int_{r=R} d\mathbf{r} c_{\alpha}(R\hat{r}) \vec{1}_r ,$$

where we have defined

$$\Lambda_{\kappa,\alpha} = \int_0^R dr r \left(e^{-\beta V_{\kappa,\alpha}(r)} - 1 \right) .$$

- c_{α} is the concentration of chemical species α .
- $V_{\kappa,\alpha}$ is the interaction potential between colloid κ and fluid species α .
- $\beta = (k_B T)^{-1}$

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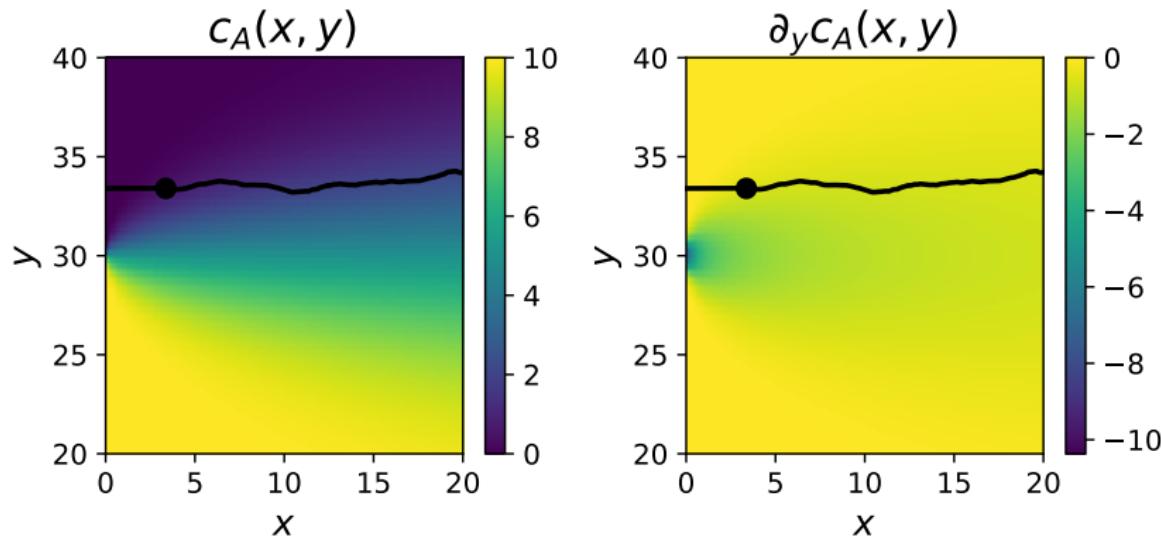
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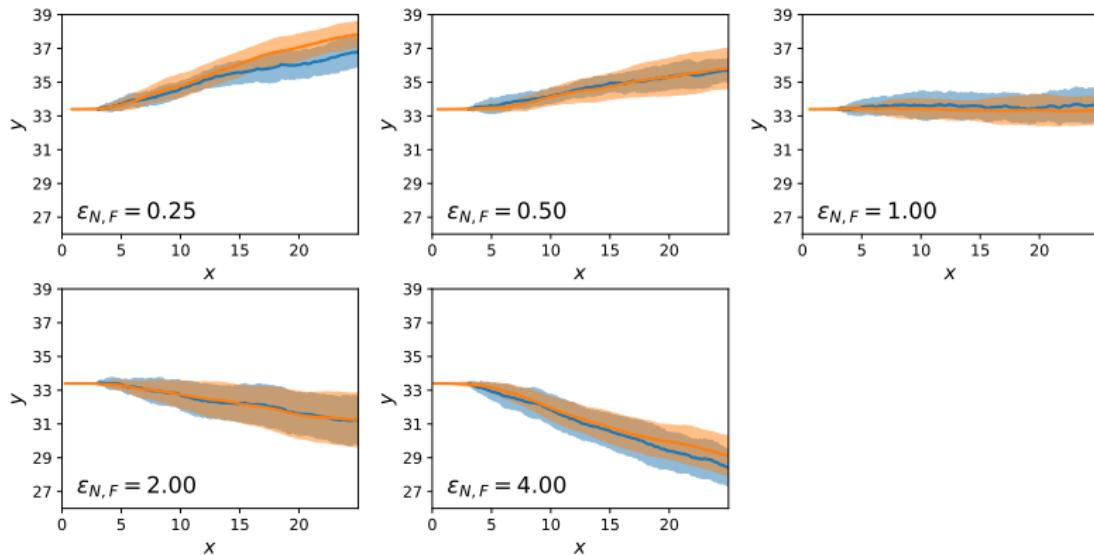
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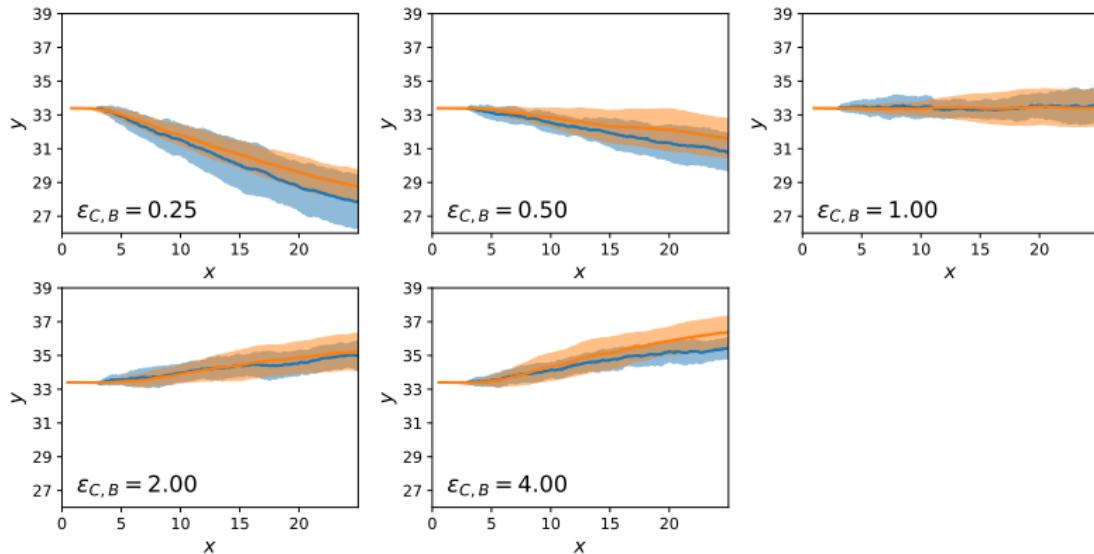
Pseudocolor represents the concentration (left) and the magnitude of the gradient (right).



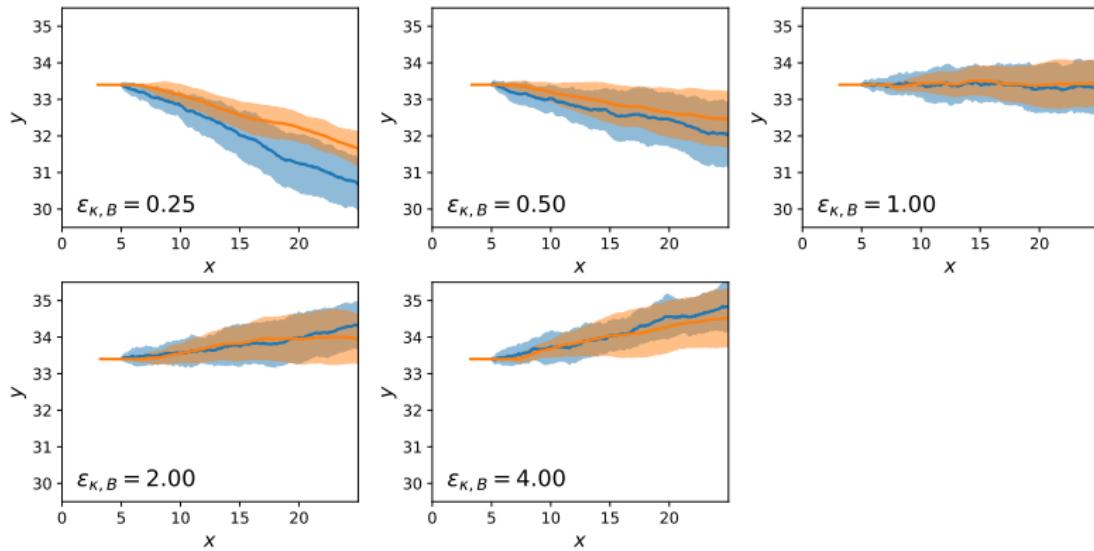
Passive sphere - only A and F, $\epsilon_A = 1$



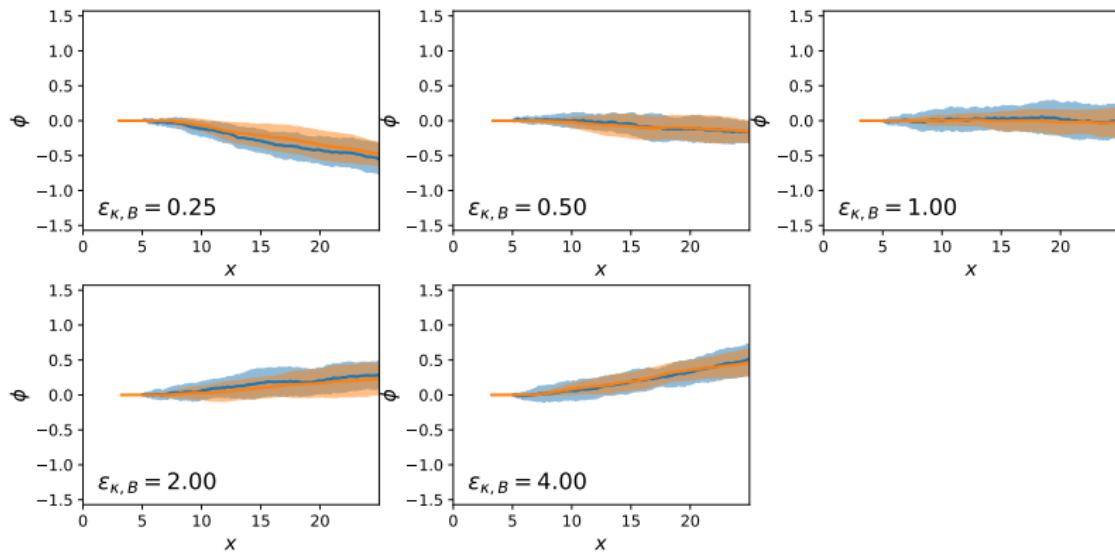
Active sphere - $A + C \rightarrow B + C, \epsilon_A = 1$



Nanomotor - $A + C \rightarrow B + C$, $\epsilon_A = 1$, $\epsilon_{\kappa,B}$ is varied



Nanomotor angle - $A + C \rightarrow B + C$, $\epsilon_A = 1$, $\epsilon_{\kappa,B}$ is varied



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- Perspectives:
 - ▶ Other motors
 - ▶ Integration with enzyme chemo-mechanical models
- Thank you!

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