

Run-and-tumble motion and differential dynamic microscopy

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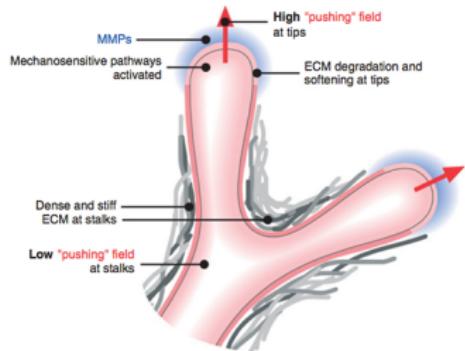
Outline

- 1 Pattern formation in biology
- 2 Patterns from interactions of two species
- 3 Differential dynamic microscopy
- 4 Run-and-tumble motion with obstacles
- 5 Summary and future work

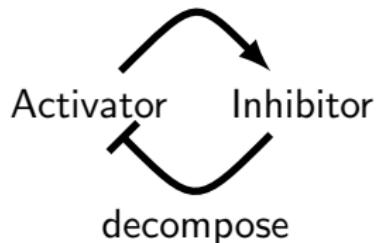
Patterns



Patterns

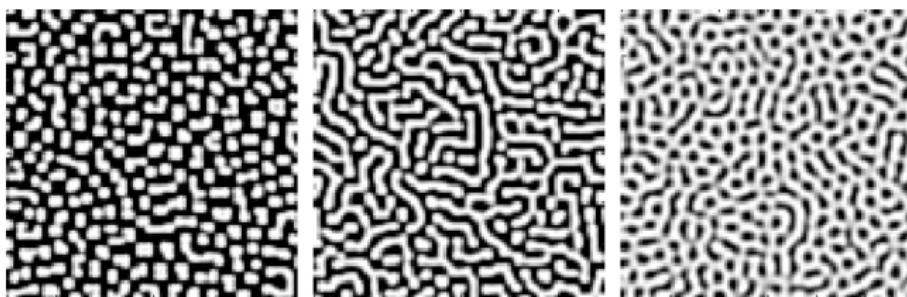


Turing pattern (1952)



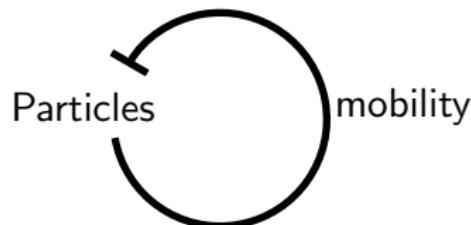
$$\frac{\partial A}{\partial t} = D_A \nabla^2 A + f(A, I)$$

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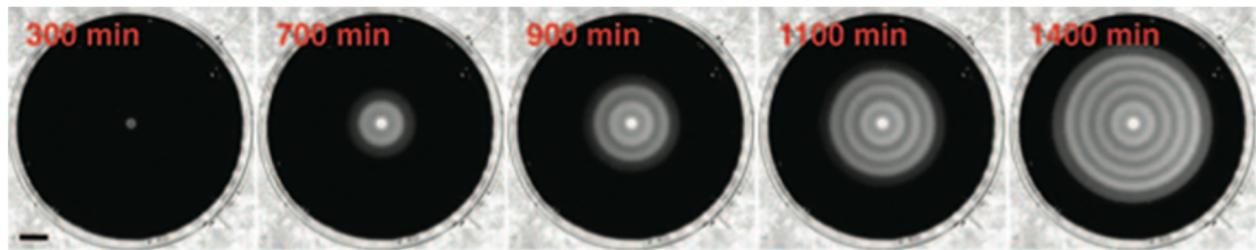


(Wikipedia.org)

Density-dependent mobility induced pattern



$$\frac{\partial \rho}{\partial t} = \nabla [D(\rho) \nabla \rho]$$

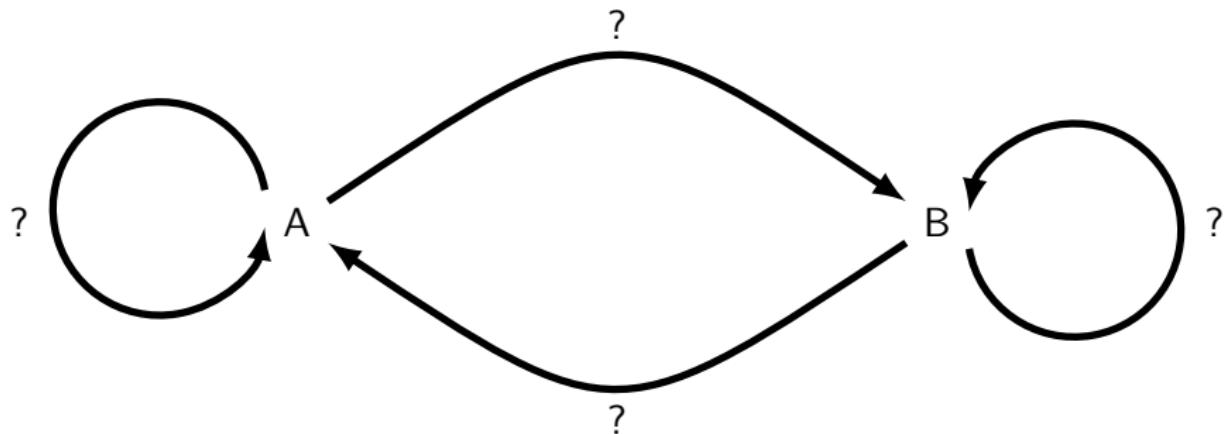


(2011, Chenli Liu, Xiongfei Fu, et al, Science)

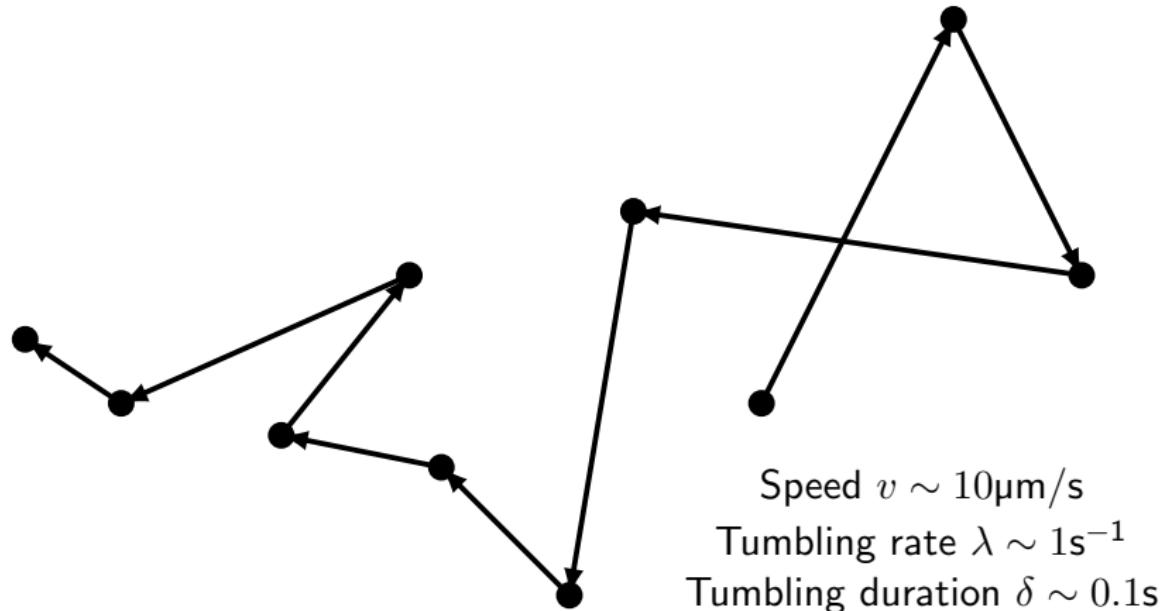
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Interactions among two species



Run-and-tumble motion of *E. coli*



$$\frac{\partial \rho}{\partial t} = \nabla \cdot (D \nabla \rho) + \nabla \left[\frac{v}{d\lambda} \nabla \left(\frac{v}{1 + \lambda \delta} \rho \right) \right].$$

Linear stability analysis

$$\frac{\partial \rho_a}{\partial t} = \nabla \left[\frac{v_a^2}{d\lambda_a(\rho_b)(1 + \lambda_a(\rho_b)\delta_a)} \nabla \rho_a - \frac{v_a^2 \delta_a \lambda'_a(\rho_b) \rho_a}{d\lambda_a(\rho_b)(1 + \lambda_a(\rho_b)\delta_a)^2} \nabla \rho_b \right],$$

$$\frac{\partial \rho_b}{\partial t} = \nabla \left[\frac{v_b^2}{d\lambda_b(\rho_a)(1 + \lambda_b(\rho_a)\delta_b)} \nabla \rho_b - \frac{v_b^2 \delta_b \lambda'_b(\rho_a) \rho_b}{d\lambda_b(\rho_a)(1 + \lambda_b(\rho_a)\delta_b)^2} \nabla \rho_a \right].$$

Bifurcation condition

$$\lambda'_a(\rho_{b0}) \lambda'_b(\rho_{a0}) > \frac{(1 + \lambda_a(\rho_{a0})\delta_a)(1 + \lambda_b(\rho_{b0})\delta_b)}{\delta_a \delta_b \rho_{a0} \rho_{b0}}.$$

Eigenvector

$$\left(\frac{v_a^2}{d\lambda_a(\rho_{b0})(1 + \lambda_a(\rho_{b0})\delta_a)} + \eta \right) \delta \rho_a = \frac{v_a^2 \delta_a \lambda'_a(\rho_{b0}) \rho_{a0}}{d\lambda_a(\rho_{b0})(1 + \lambda_a(\rho_{b0})\delta_a)^2} \delta \rho_b.$$

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Two general principles

$$\lambda'_a(\rho_{b0})\lambda'_b(\rho_{a0}) > \frac{(1 + \lambda_a(\rho_{a0})\delta_a)(1 + \lambda_b(\rho_{b0})\delta_b)}{\delta_a\delta_b\rho_{a0}\rho_{b0}} \implies$$

$$\lambda'_a(\rho_{b0}) > 0, \lambda'_b(\rho_{a0}) > 0 \text{ or } \lambda'_a(\rho_{b0}) < 0, \lambda'_b(\rho_{a0}) < 0.$$

- $\lambda'_a(\rho_{b0}) > 0, \lambda'_b(\rho_{a0}) > 0$: **mutual inhibition**
 $\delta\rho_a \propto \delta\rho_b$: co-migrating pattern.
- $\lambda'_a(\rho_{b0}) < 0, \lambda'_b(\rho_{a0}) < 0$: **mutual activation**
 $\delta\rho_a \propto -\delta\rho_b$: segregating pattern.

The similar conclusions hold for density dependent speed (from Curatolo) or tumbling duration.

Two general principles

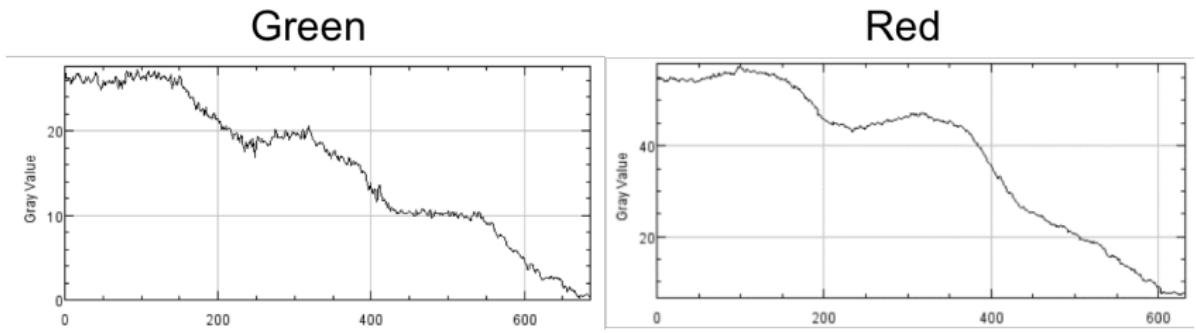
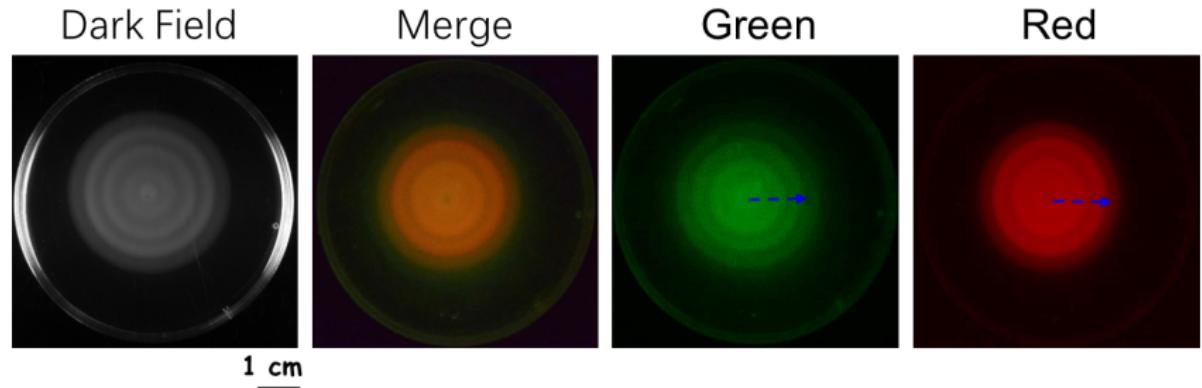
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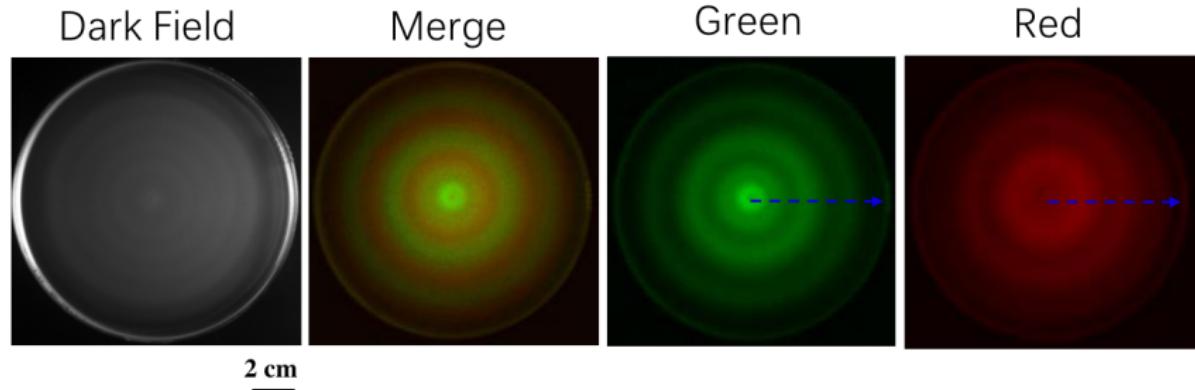
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Experiments - mutual inhibition



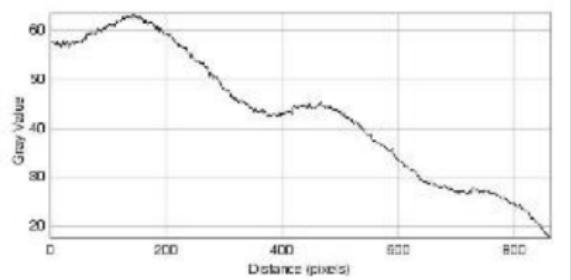
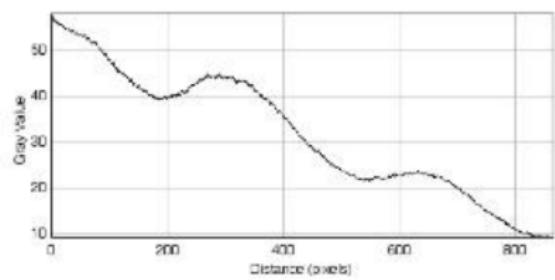
Experiments - mutual activation



2 cm

Green

Red



A question

How do we know the designed system works as we expected?

⇒ How can we measure the speed/tumbling rate/tumbling duration of *E. coli* ?

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Why not tracking?

- Special equipment for 3D measurement.
- Difficult to measure tumbling duration.
- Valid only for low density.
- Laborious.
- Poor statistics ($\sim 10^2$ cells).

If I am telling you there is a method:

- Usual equipment automatically doing 3D measurement.
- Easier to measure tumbling duration.
- Valid for high density.
- Easy.
- Good statistics ($\gg 10^4$ cells).

Why not tracking?

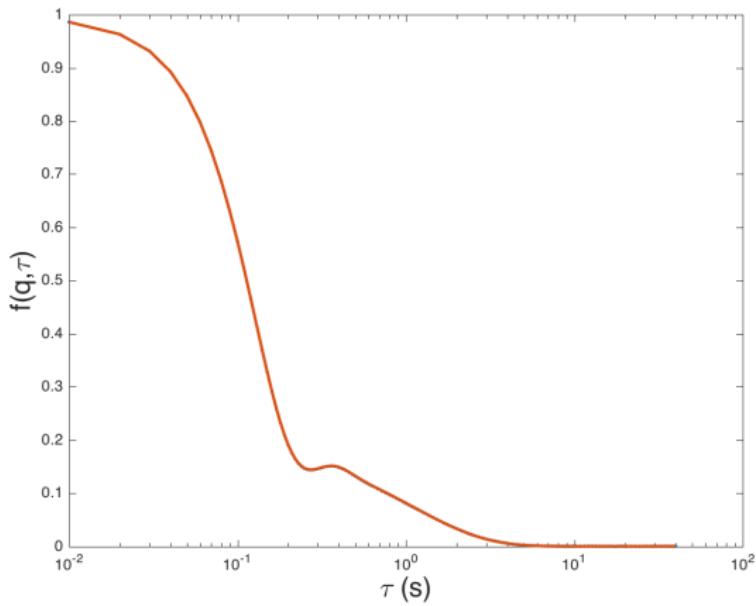
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Intermediate scattering function (ISF)

$$f(q, \tau) = \frac{\langle \Delta\rho(q, t)\Delta\rho^*(q, t + \tau) \rangle_e}{\langle \Delta\rho(q, t)\Delta\rho^*(q, t) \rangle_e}.$$



Differential dynamic microscopy

The principle of differential dynamic microscopy

It can be proven for ergodic point particles that

$$f(q, \tau) = \frac{\langle \Delta\rho(q, t)\Delta\rho^*(q, t + \tau) \rangle_t}{\langle \Delta\rho(q, t)\Delta\rho^*(q, t) \rangle_t} = p(q, \tau),$$

which is the solution of corresponding master equation with initial condition to be $p(x, t) = \delta(x)$.

- $f(q, \tau)$ can be calculated from image intensity $I(q, t)$, via

$$g(q, \tau) = \langle |I(q, t + \tau) - I(q, t)|^2 \rangle_t = A(q)(1 - f(q, \tau)) + B(q).$$

- $p(q, \tau)$ can be obtained from solving the master equation.

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- $p(q, \tau)$ can be obtained from solving the master equation.

Master equation of run-and-tumble particles

$$\frac{\partial p(\mathbf{x}, \mathbf{v}, t)}{\partial t} = D\nabla^2 p - \mathbf{v} \cdot \nabla p - \lambda p + \frac{\lambda}{\Omega} \int p(\mathbf{x}, \mathbf{v}', t) d\Omega'.$$

$$p(q, v, s) = \frac{\arctan(qv/(s + Dq^2 + \lambda))}{qv - \lambda \arctan(qv/(s + Dq^2 + \lambda))}.$$

Adding a distribution of v and a contribution of dead cells,

$$f(q, \tau) = (1 - \alpha)e^{-Dq^2\tau} + \alpha \int_0^\infty p(q, v, t) P(v) dv,$$

where

$$P(v) = \frac{v^Z}{\Gamma(Z+1)} \left(\frac{Z+1}{\bar{v}} \right)^{Z+1} e^{-(Z+1)v/\bar{v}}.$$

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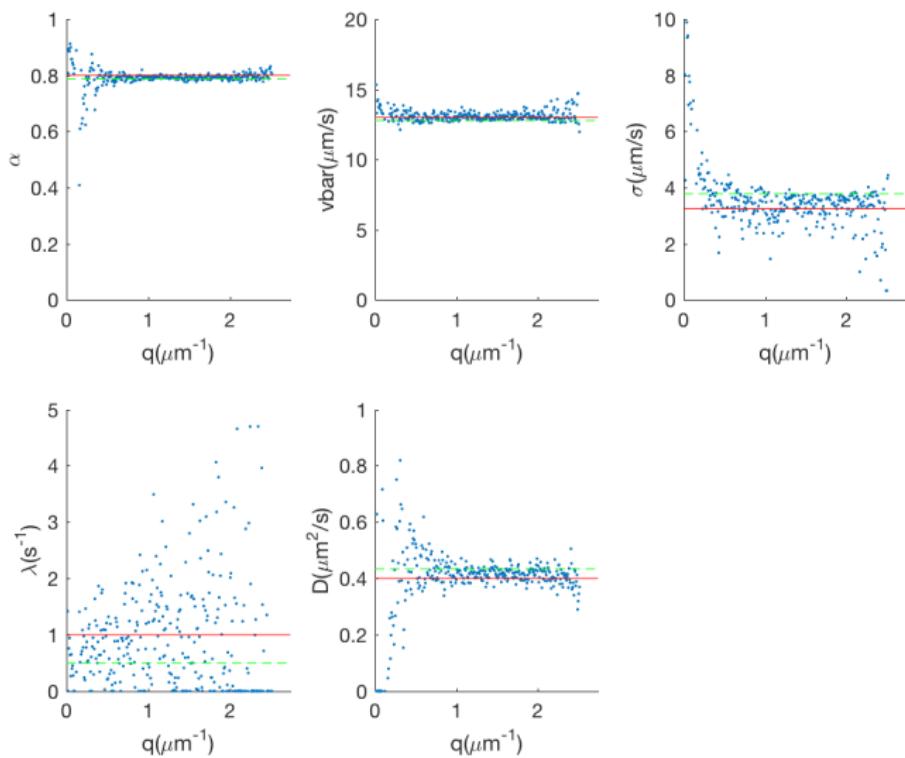
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Simulation

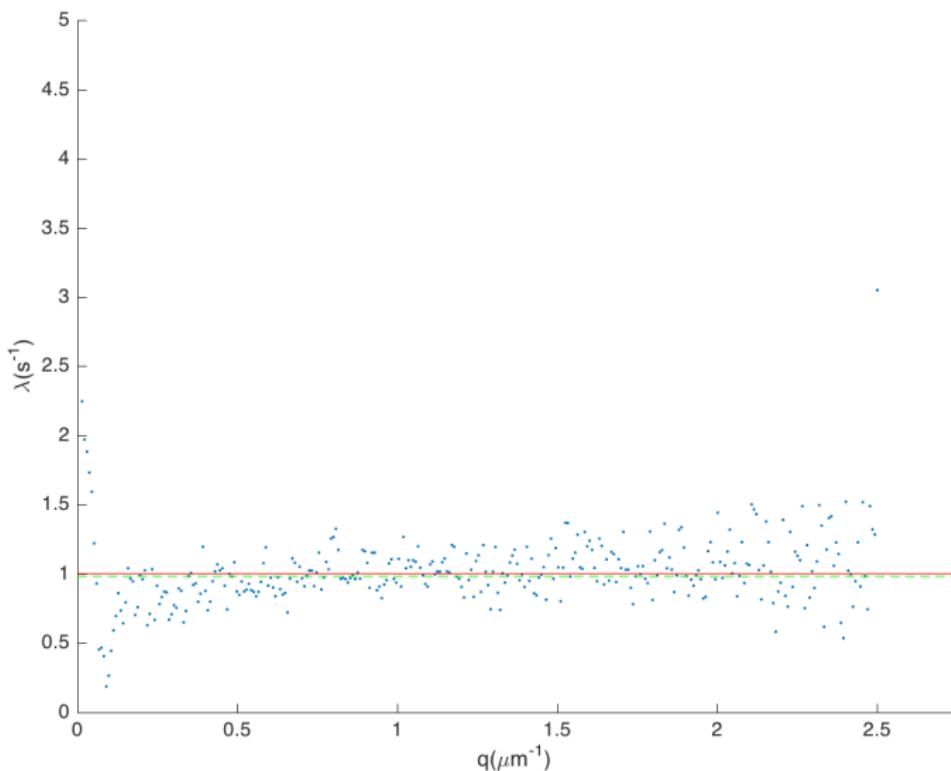


- 512×512 pixels.
- Pixel size corresponds to real microscopy systems.
- Depth of fields $\sim 40 \mu\text{m}$.
- Particle number $\sim 10^4$.

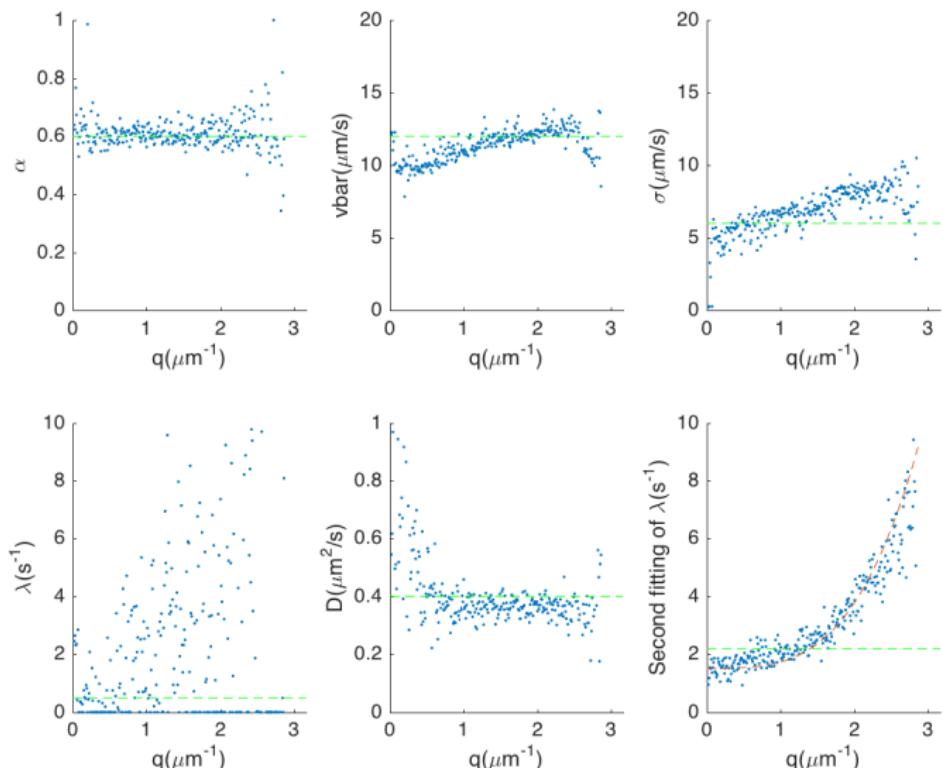
Fitting result of simulation



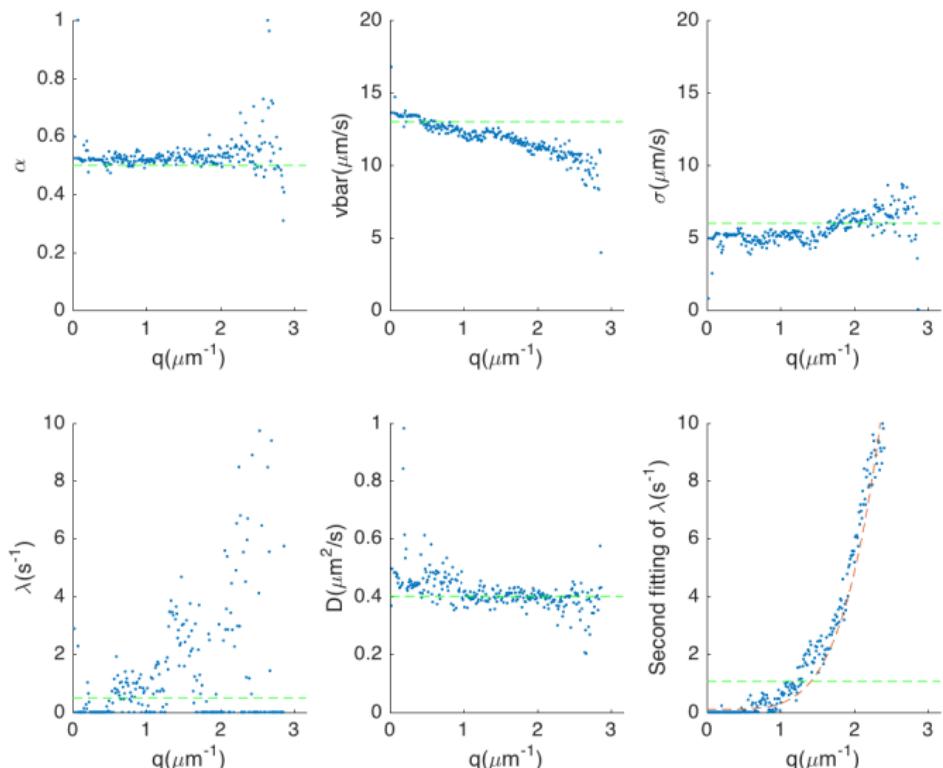
A second fitting with fixed parameters



Fitting result of experiment of wild type *E. coli*



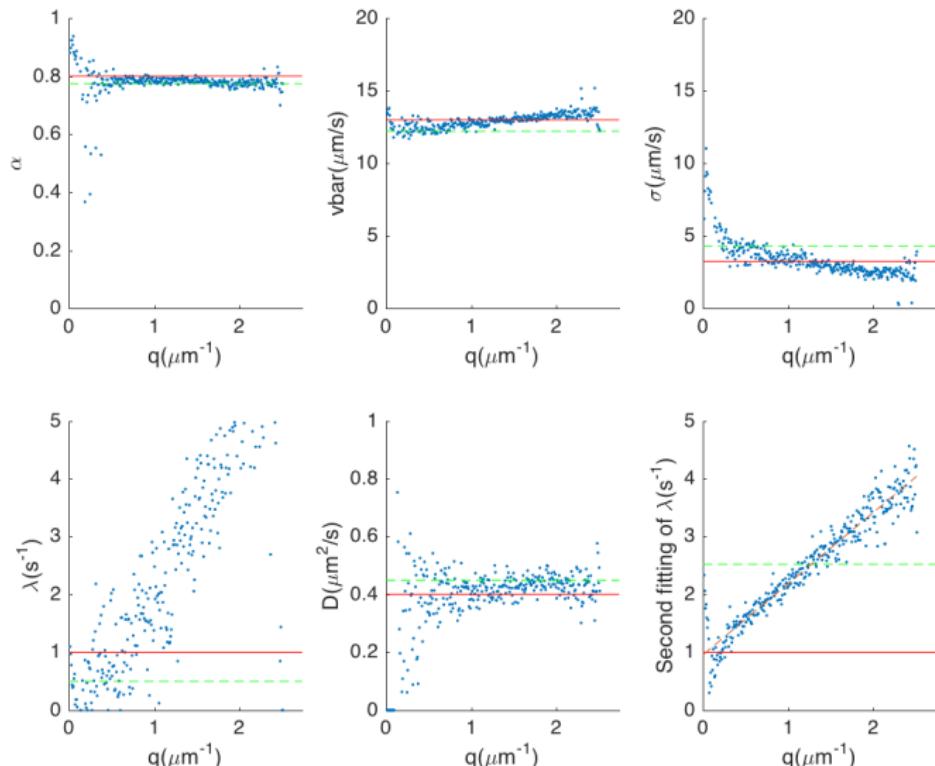
Fitting result of experiment of Δ CheY *E. coli*



What is missing?

- Finite tumbling duration.
- Rotational diffusion.
- Active rotational diffusion causing bias in tumbling.
- Lévy's walk (power law distribution in running time).

Simulation with finite tumbling duration



Finite tumbling duration

Master equation

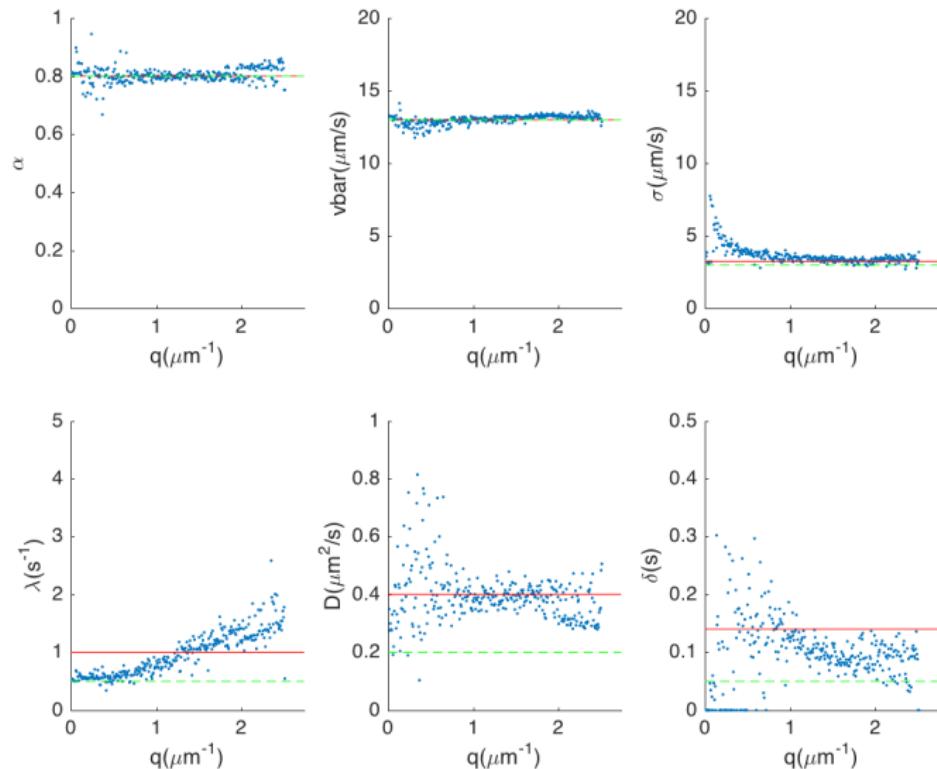
$$\begin{aligned}\frac{\partial p_r(\mathbf{x}, \mathbf{v}, \tau)}{\partial \tau} &= D \nabla^2 p_r - \mathbf{v} \cdot \nabla p_r - \lambda p_r + \frac{p_t}{\delta \Omega}, \\ \frac{\partial p_t(\mathbf{x}, \tau)}{\partial \tau} &= D \nabla^2 p_t + \lambda \int p_r(\mathbf{x}, \mathbf{v}', \tau) d\Omega' - \frac{p_t}{\delta}.\end{aligned}$$

Intermediate scattering function

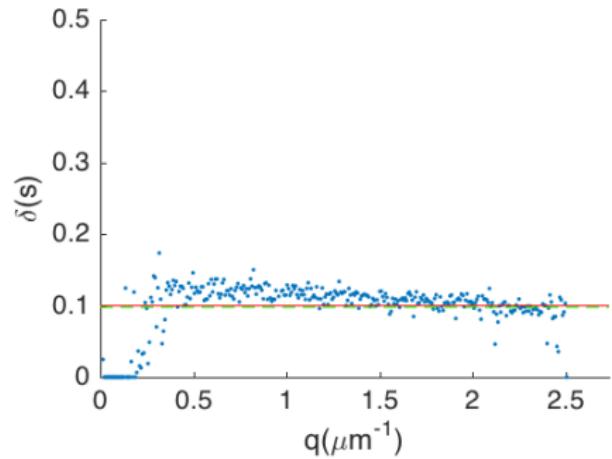
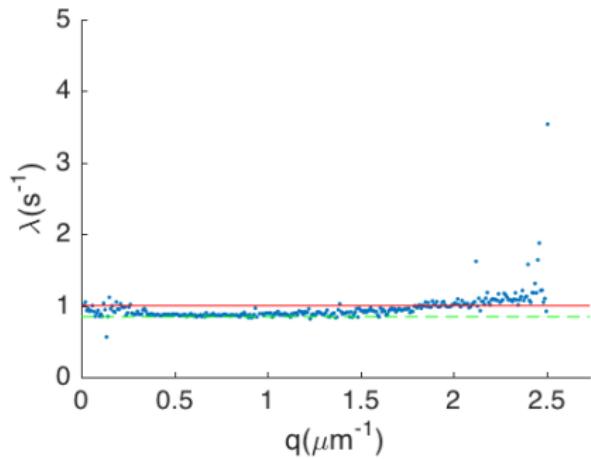
$$p(q, v, s) = \frac{1}{(\delta(s + Dq^2) + 1)(\delta\lambda + 1)} \left(\frac{(\delta(s + \lambda + Dq^2) + 1)^2 \arctan(qv/(s + \lambda + Dq^2))}{qv(\delta(s + Dq^2) + 1) - \lambda \arctan(qv/(s + \lambda + Dq^2))} + \delta^2 \lambda \right).$$

$$f(q, \tau) = (1 - \alpha)e^{-Dq^2\tau} + \alpha \int_0^\infty p(q, v, t) P(v) dv.$$

Fitting with tumbling duration



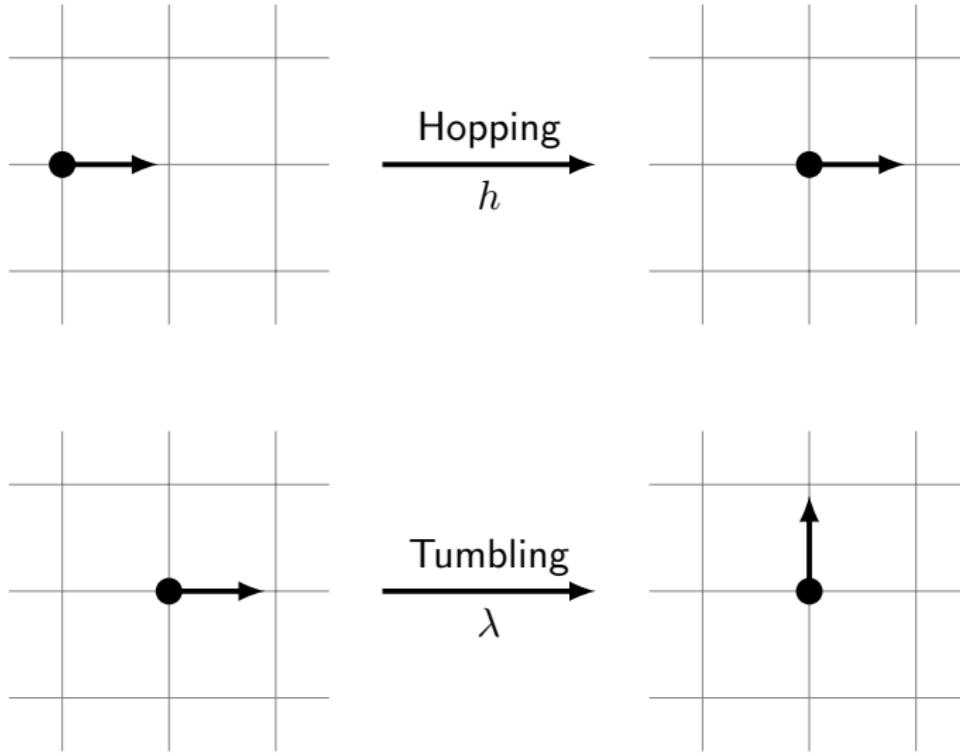
Fitting with tumbling duration



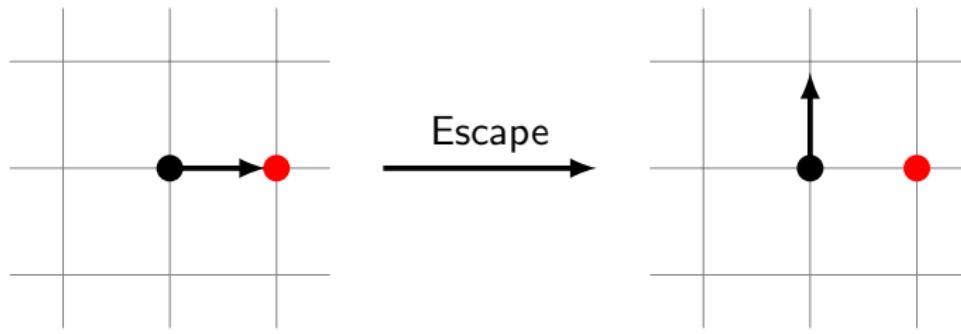
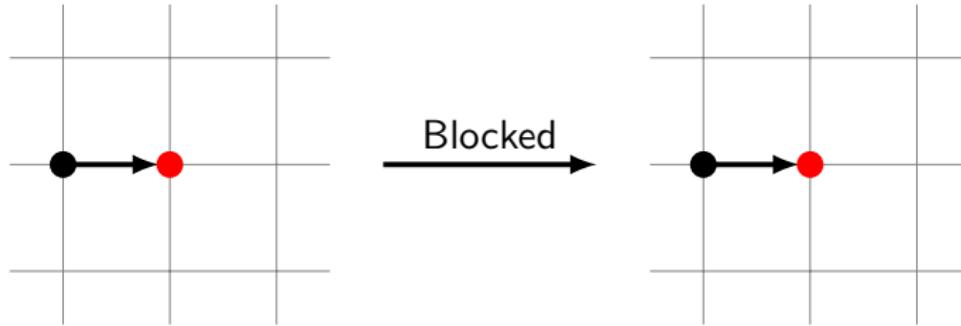
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Lattice model



Lattice model



Mean field approximation

- c : Concentration of obstacles.

Master equation

$$\begin{aligned}\frac{\partial p_r(\mathbf{x}, \hat{\mathbf{v}}, t)}{\partial t} &= -\nabla \cdot (\hat{\mathbf{v}} h p_r) - \lambda p_r - \gamma_d(c) h c p_r + \frac{\lambda(1-c)}{\Omega} \rho_M, \\ \frac{\partial p_b(\mathbf{x}, \hat{\mathbf{v}}, t)}{\partial t} &= -\lambda p_b + \frac{\lambda c}{\Omega} \rho_M + \gamma_d(c) h c p_r, \\ \rho_M &= \int (p_r + p_b) d\Omega.\end{aligned}$$

Effective diffusion coefficient

$$D_{\text{eff}} = \frac{h^2 \lambda (1-c)}{d [h c \gamma_d(c) + \lambda]^2}.$$

Mean field approximation

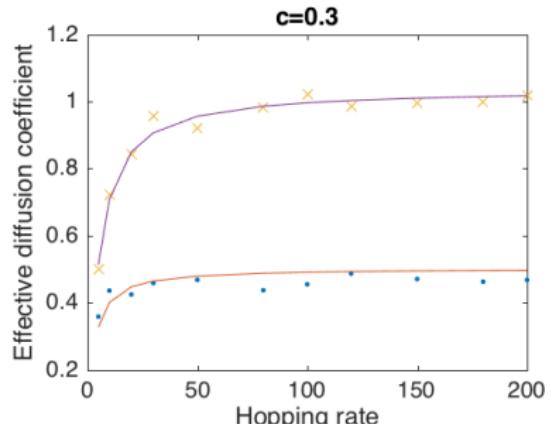
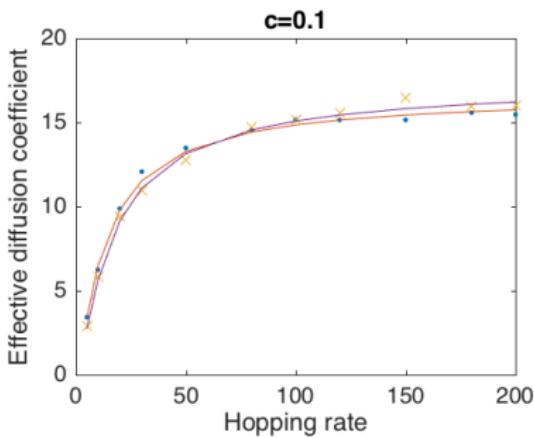
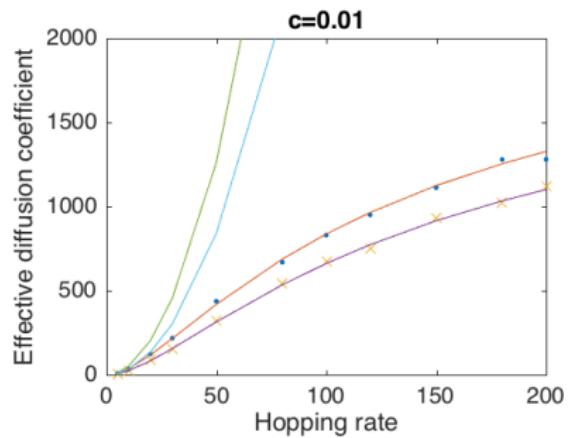
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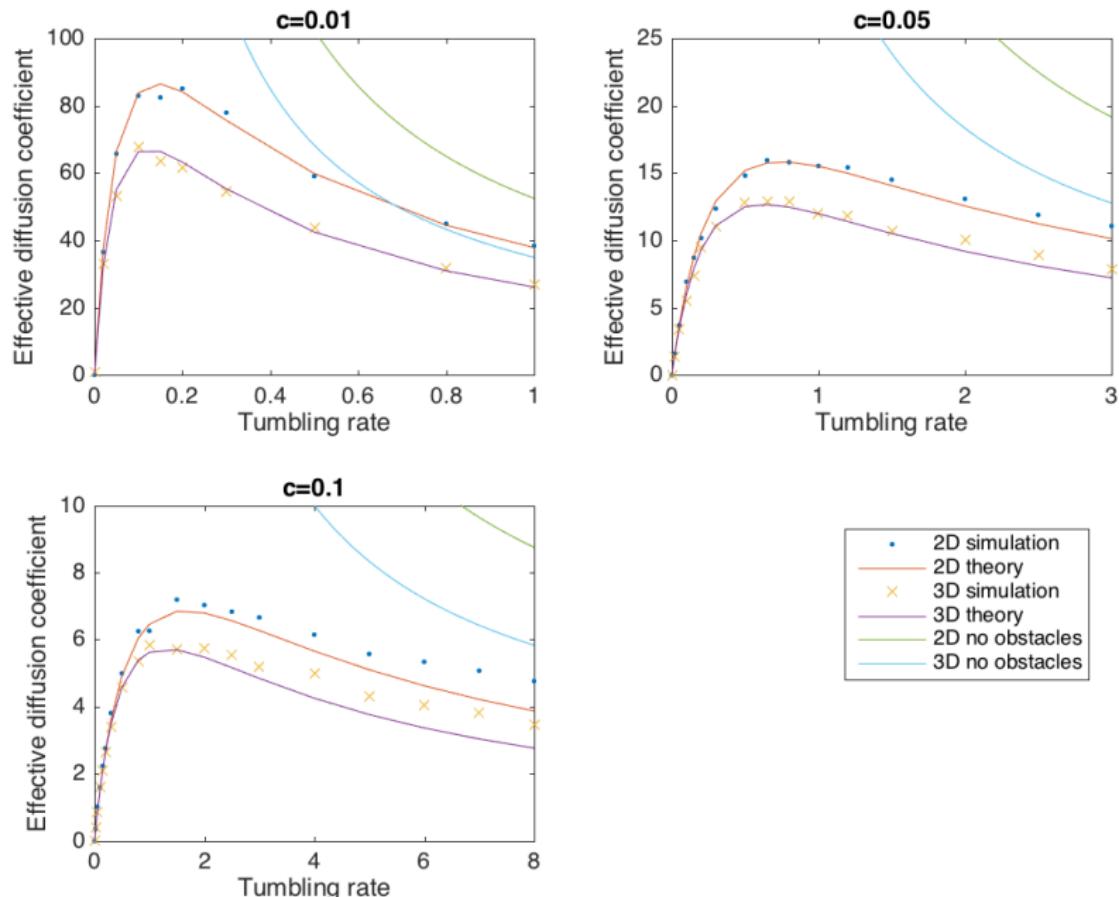
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- 2D simulation
- 2D theory
- ✖ 3D simulation
- 3D theory
- 2D no obstacles
- 3D no obstacles



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Summary

New mechanism for pattern formation

- Mutual inhibition of mobility: co-migrating pattern.
- Mutual activation of mobility: segregating pattern.

New development of differential dynamic microscopy

- Tumbling rate can be measured for run-and-tumble particles with instantaneous tumbling.
- Tumbling duration can be measured for run-and-tumble particles if the tumbling rate is known.
- The motion of *E. coli* may have some ingredients we haven't known.

New attempt for *E. coli* motion with obstacles

- A valid lattice model.
- A mean field approximation.

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Future work

Pattern formation

- Experiments: tuning the pattern.
- Three-species, four-species, ...: more complex network motif.
- Growth, death, ...: more kinds of interactions.

Differential dynamic microscopy

- Way to specify the tumbling rate of run-and-tumble particles with finite tumbling duration: multi-scale imaging?
- Measure rotational diffusivity.
- Effect of Levy's walk.

E. coli moving in agar

- Existence of better continuous approximation: calculation of $\gamma_d(c)$.
- Possibility of DDM.

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What's more?

- Applying DDM to study the motion of other "particles": mammalian cells, fishes, birds, sheep, humans, ...
- Studying their collective behaviour.
- Finding principles in coordination and self-organization.
- Answering questions in organ development, morphogenesis, microbe infection, ...

Acknowledgement

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- Dr. Adrian Daerr
(University Paris Diderot)
- Dr. Huang, Wei
- Prof. Peter Lenz
- Mr. Zhou, Nan
- Ms. Agnese Curatolo
(University Paris Diderot)
- Dr. Alexandre Solon
(MIT)
- Dr. Lina Hamouche
(University Paris Diderot)

Case no obstacle: compare with continuous model

Master equation

$$\frac{\partial P(\mathbf{x}, \mathbf{e}_i, t)}{\partial t} = hP(\mathbf{x} - \mathbf{e}_i, \mathbf{e}_i, t) - hP - \lambda P + \frac{\lambda}{2d} \sum_i P(\mathbf{x}, \mathbf{e}_i, t).$$

Mean square displacement - lattice model

$$\langle \Delta x^2(t) \rangle = \left(\frac{2h^2}{d\lambda} + \frac{h}{d} \right) t + \frac{2h^2}{d\lambda^2} (e^{-\lambda t} - 1).$$

Mean square displacement - continuous model

$$\langle \Delta x^2(t) \rangle = \frac{2v^2}{d\lambda} t + \frac{2v^2}{d\lambda^2} (e^{-\lambda t} - 1).$$

- Necessary condition of lattice model to be valid: $h \gg \lambda/2$.

Numerical inverse Laplace transformation - Week's method

$$\mathcal{L}\{f(t)\} = F(s) .$$

- Möbius transformation: $s = \sigma - b \frac{z+1}{z-1}$.
- Expand the function:

$$(s - \sigma + b)F(s) = \frac{2b}{1-z} F\left(\sigma - b \frac{z+1}{z-1}\right) = \sum_{n=0}^{\infty} a_n z^n .$$

- Laplace transformation of Laguerre polynomial: $\mathcal{L}[L_n(2bt)] = \frac{(s-2b)^n}{s^{n+1}}$.

- $\mathcal{L}^{-1}[F(s)] = f(t) = \sum_{n=0}^{\infty} a_n e^{(\sigma-b)t} L_n(2bt)$.

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$$a_n = \frac{1}{2\pi} \int_0^{2\pi} e^{-in\theta} \frac{2b}{1-e^{i\theta}} F\left(\sigma - b \frac{e^{i\theta}+1}{e^{i\theta}-1}\right) d\theta .$$