

Bacterial Microfluidics:

The physics and engineering of flagellated bacterial motility

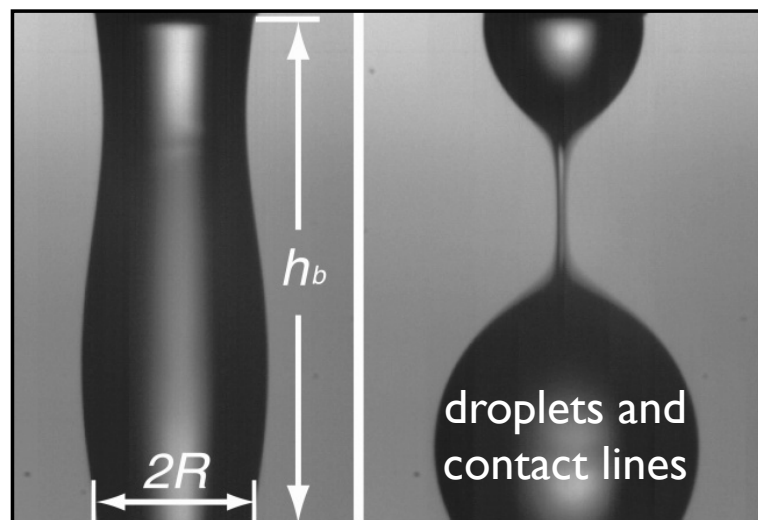
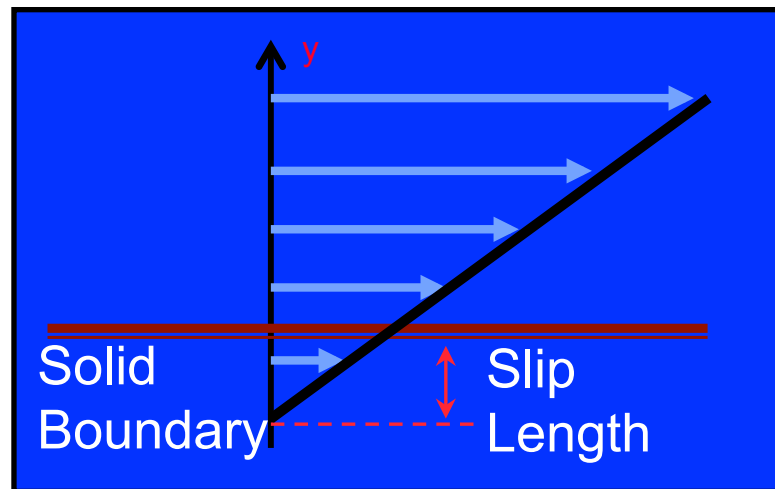
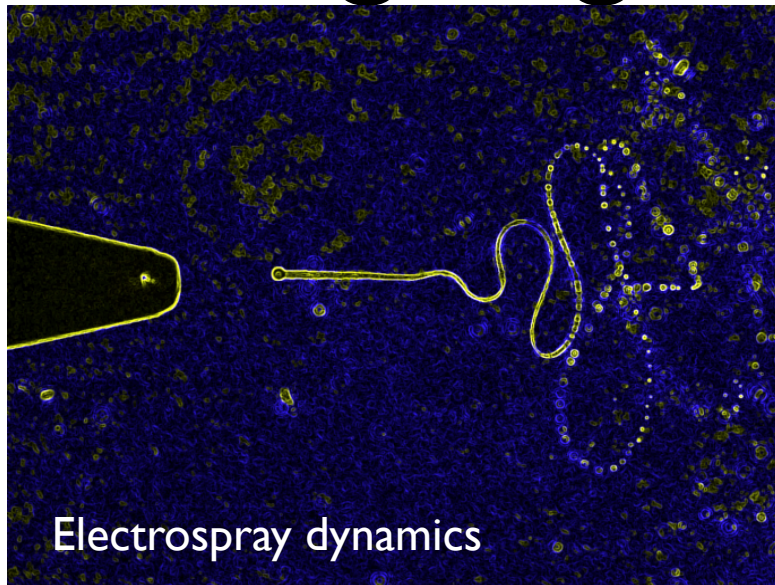
Kenny Breuer

School of Engineering, Brown University,
Providence, RI

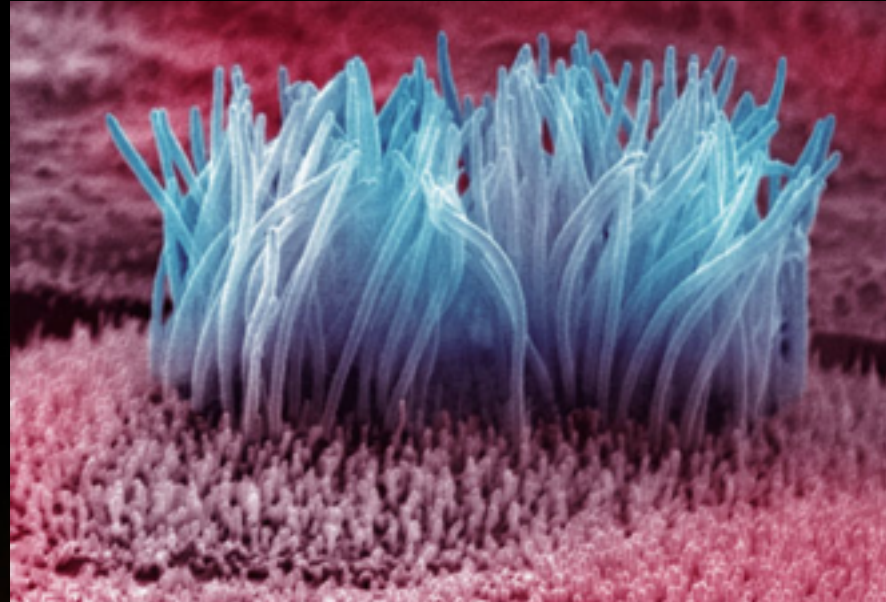
In collaboration with:

Tom Powers, Bin Liu, Qian Bian, MinJun Kim, Dave Gagnon

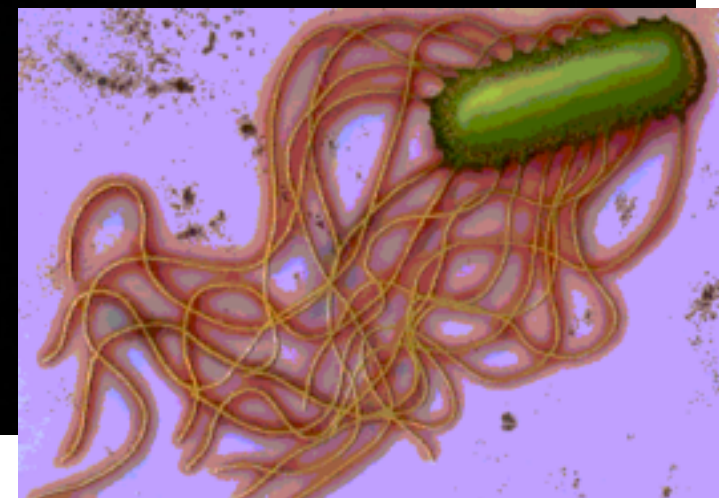
Stuff going on in my lab



Cilia and flagella in viscous fluids

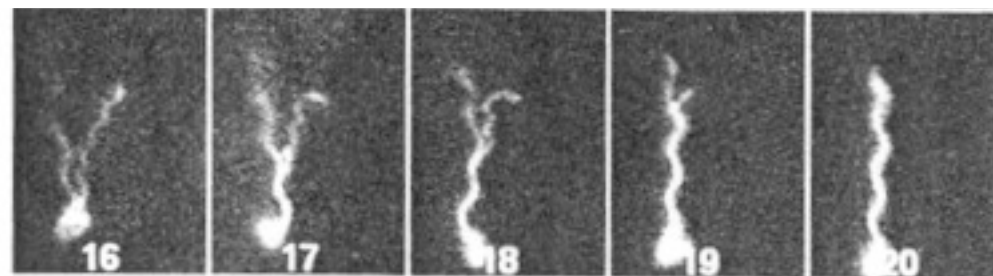
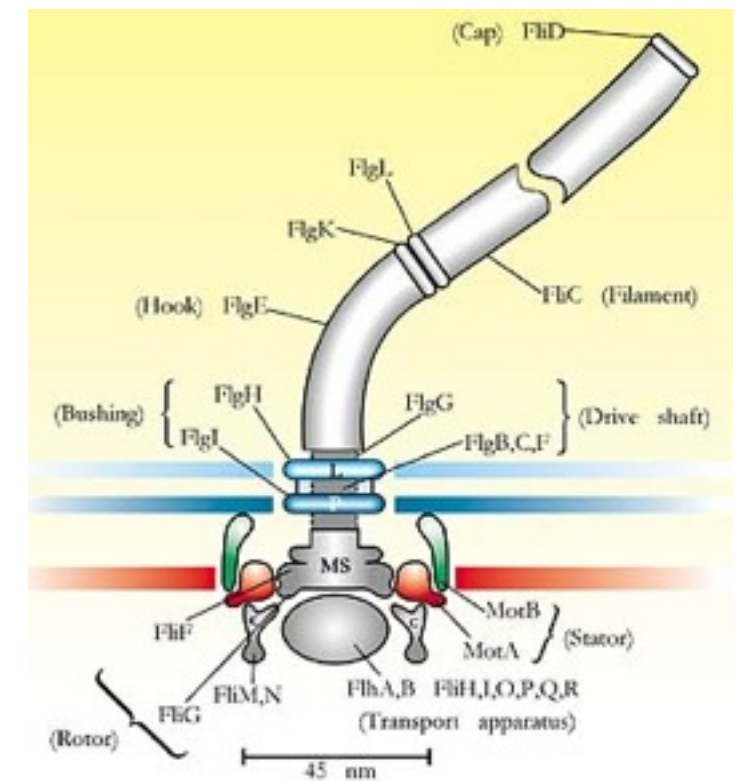
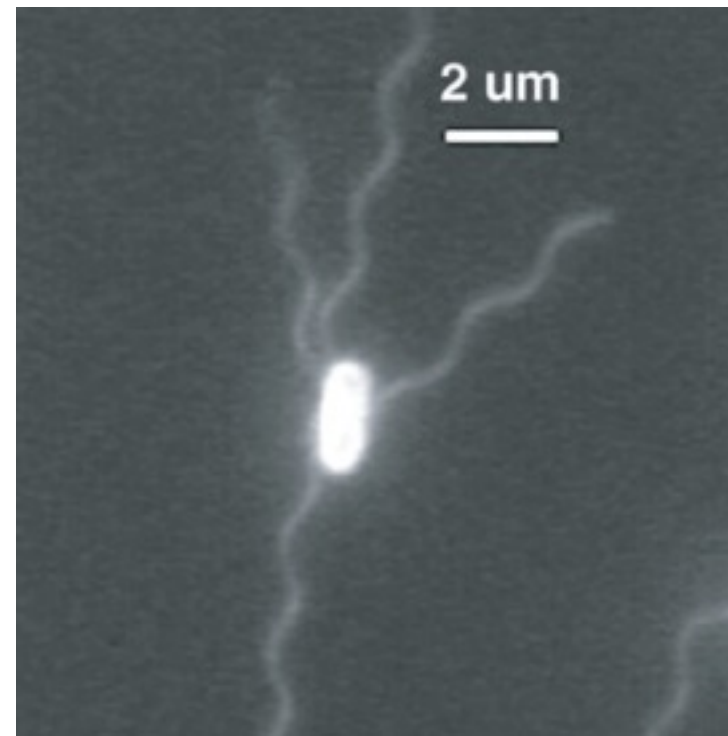


Bacterial motility



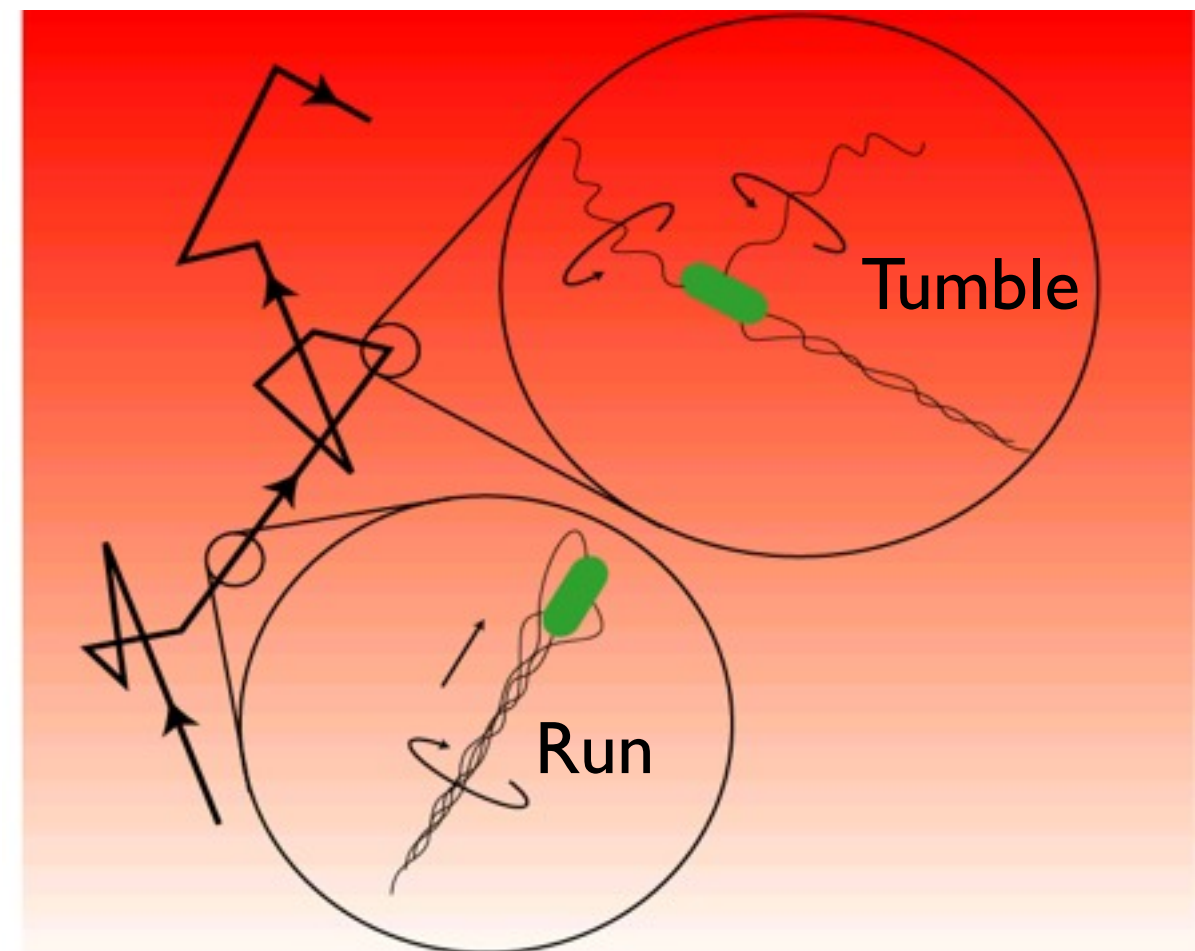
Motility of flagellated bacteria

H. Berg, Physics Today, Jan 2000;



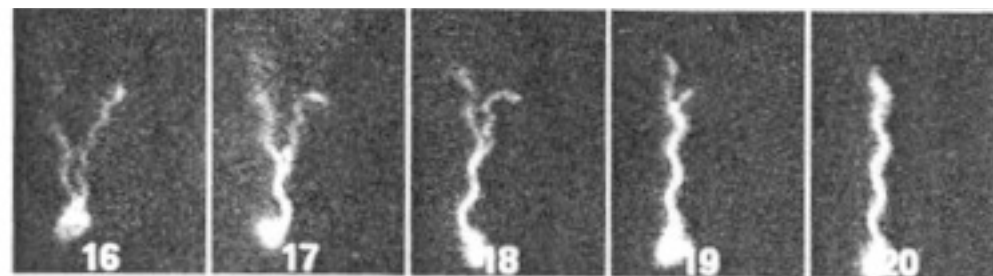
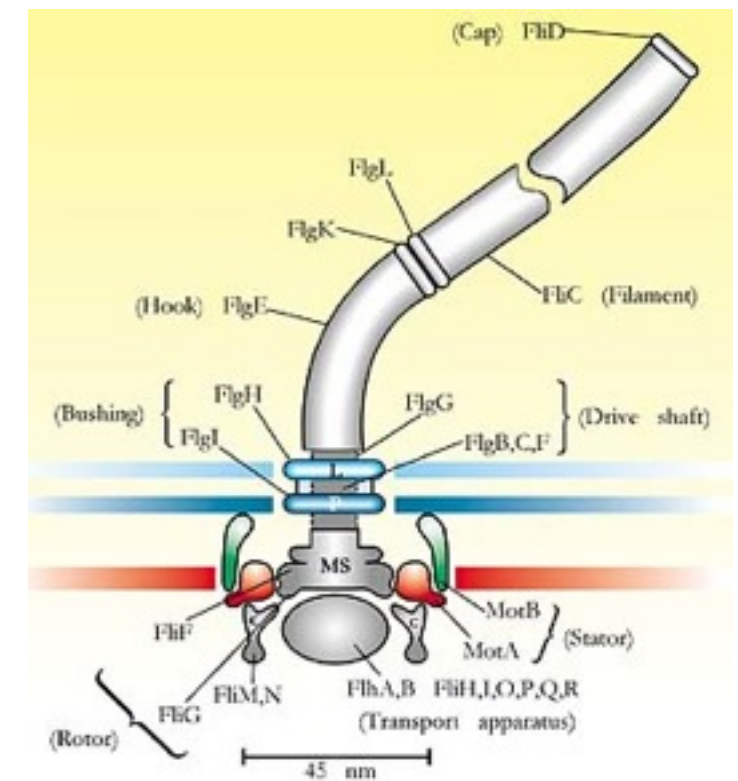
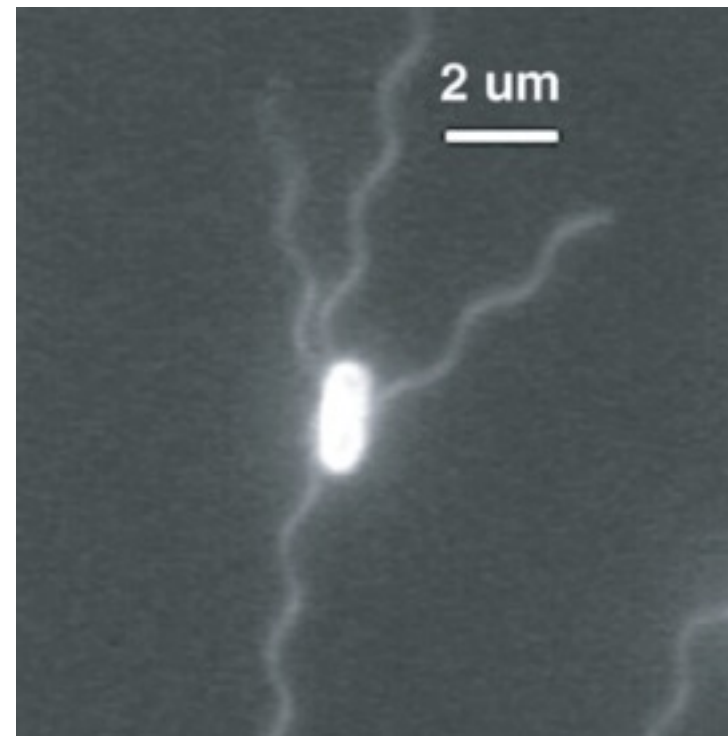
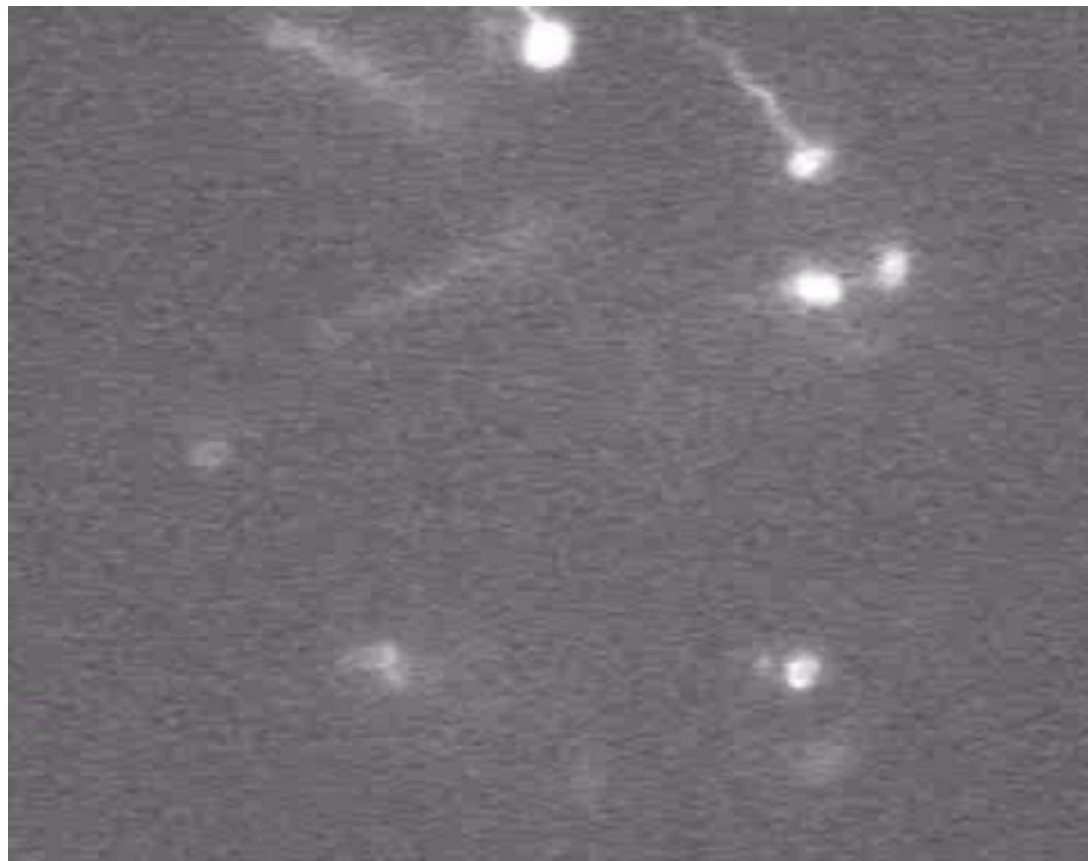
L. Turner, W. S. Ryu, and H. C. Berg (2000)

*In “real life”:
Newtonian and non-Newtonian media*



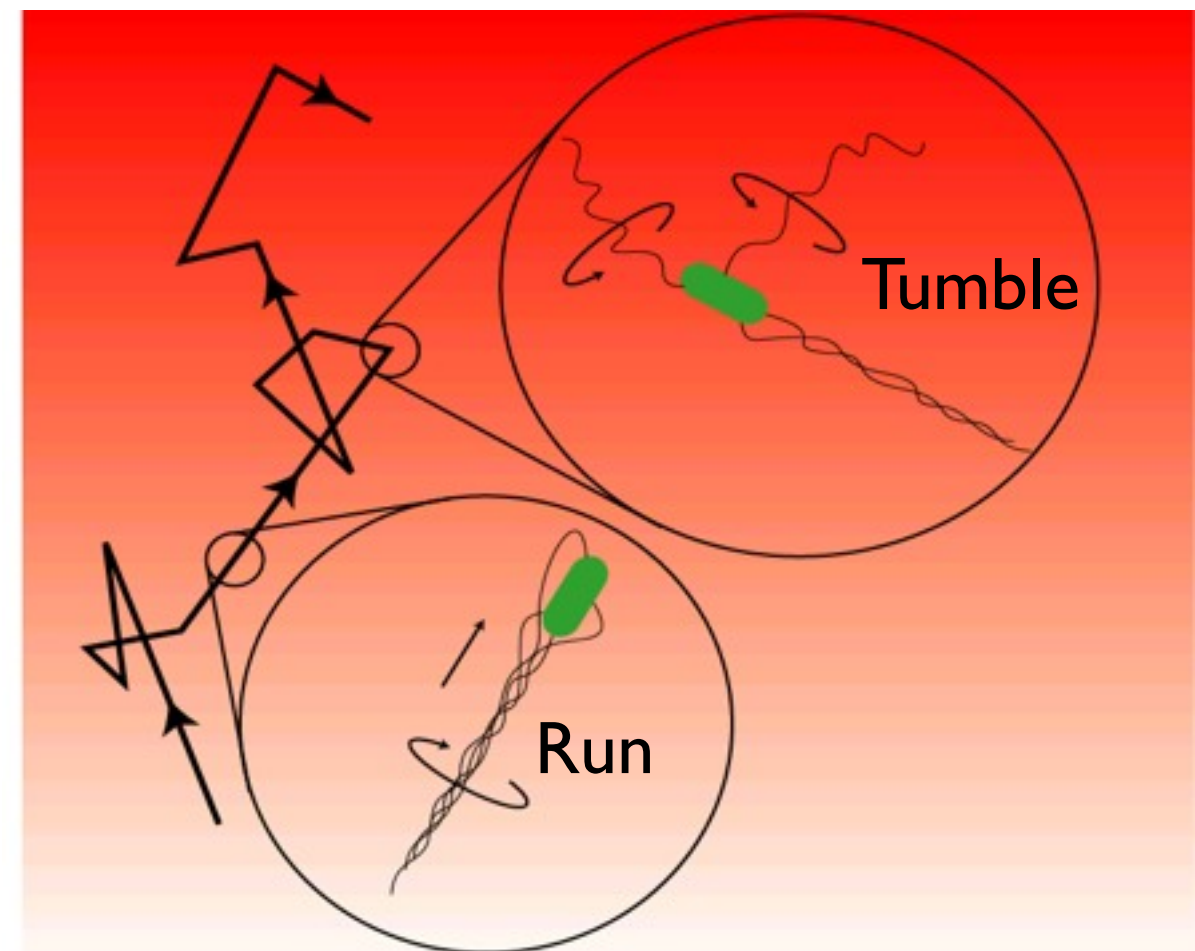
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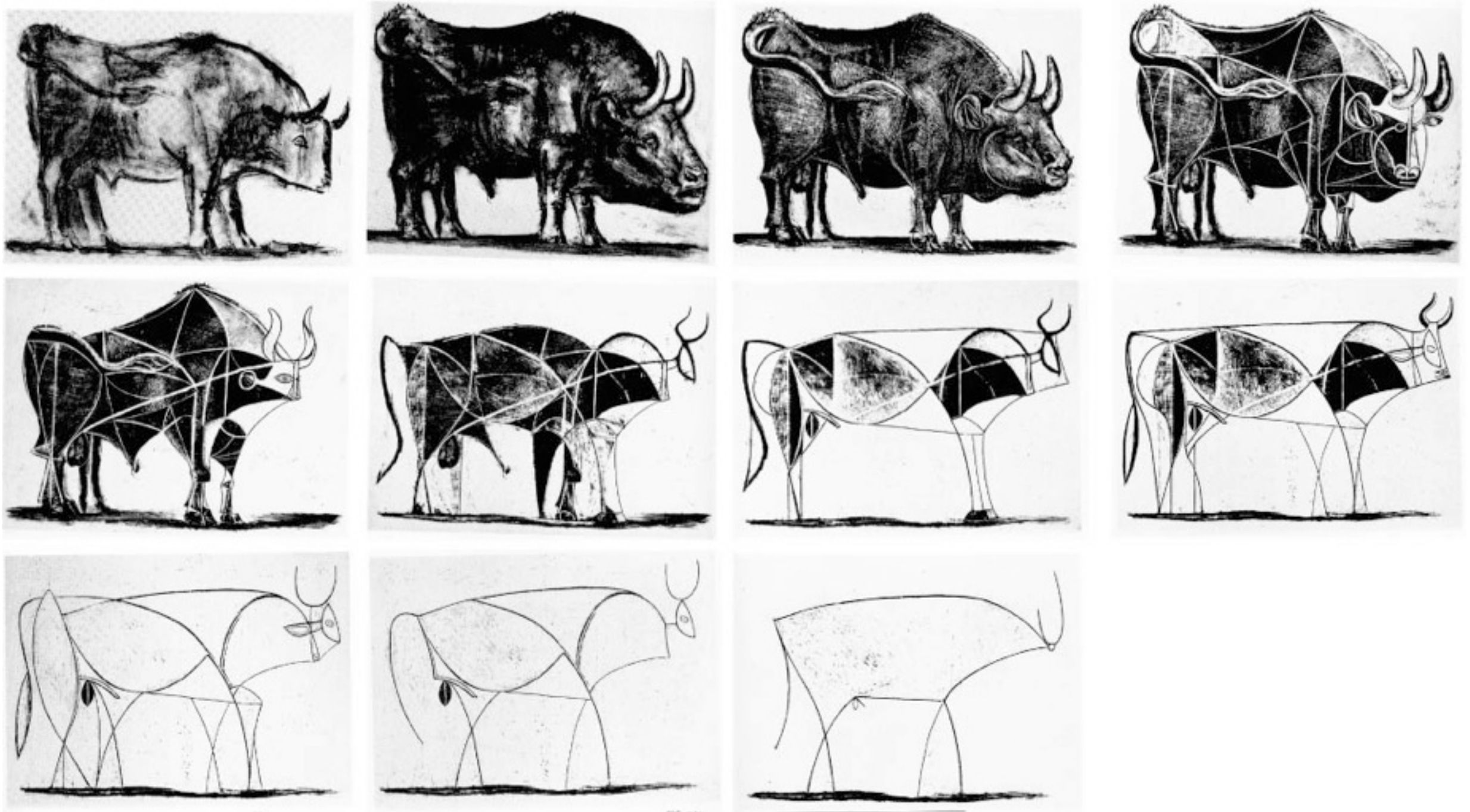
*In “real life”:
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Interesting questions

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 - Can bacteria affect the macroscopic world?
 - Can we harness their motion?
 - Can we control them?
 - etc.
- What next?

Bull #11

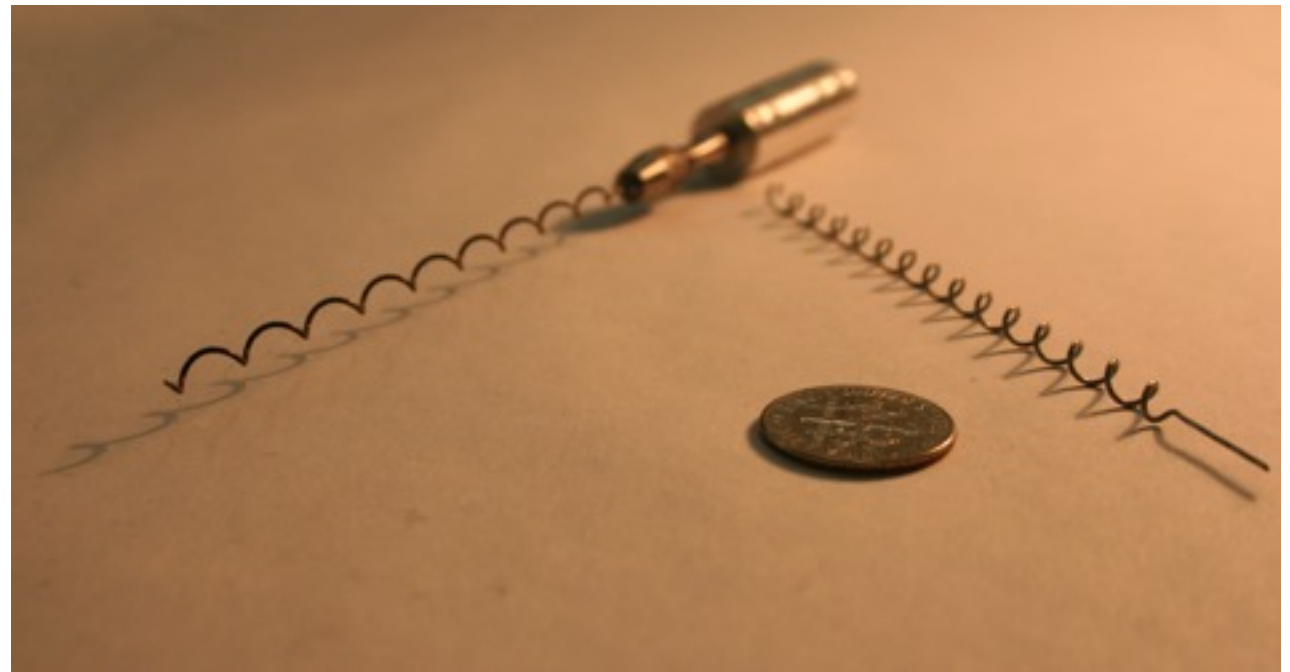
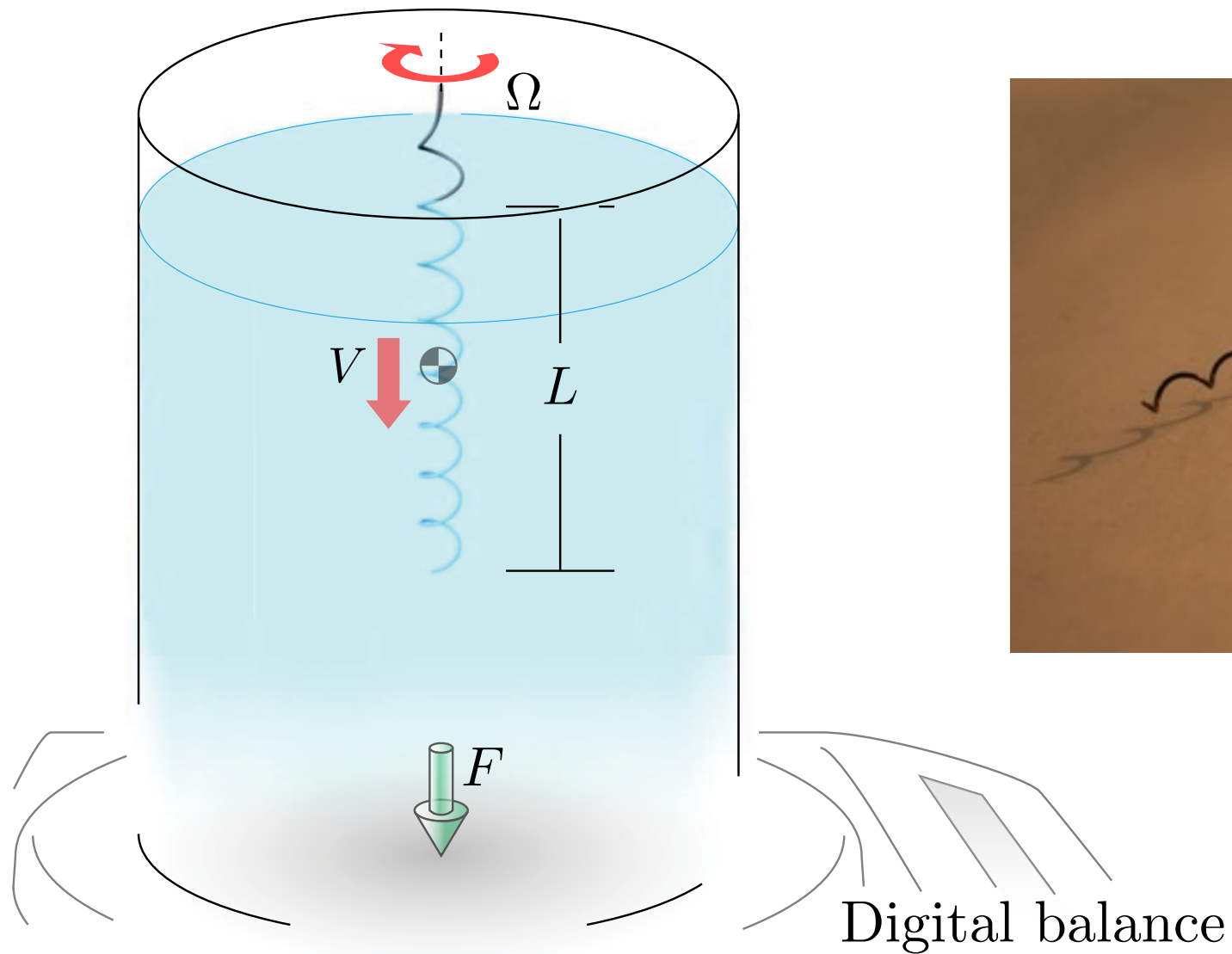


apologies to Picasso, thanks to Christophe Clanet

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Experimental setup (low-Re swimming helix)



fluids with high viscosity:

- silicone oil (Newtonian)
- Polybutene+ Polyisobutylene (viscoelastic)

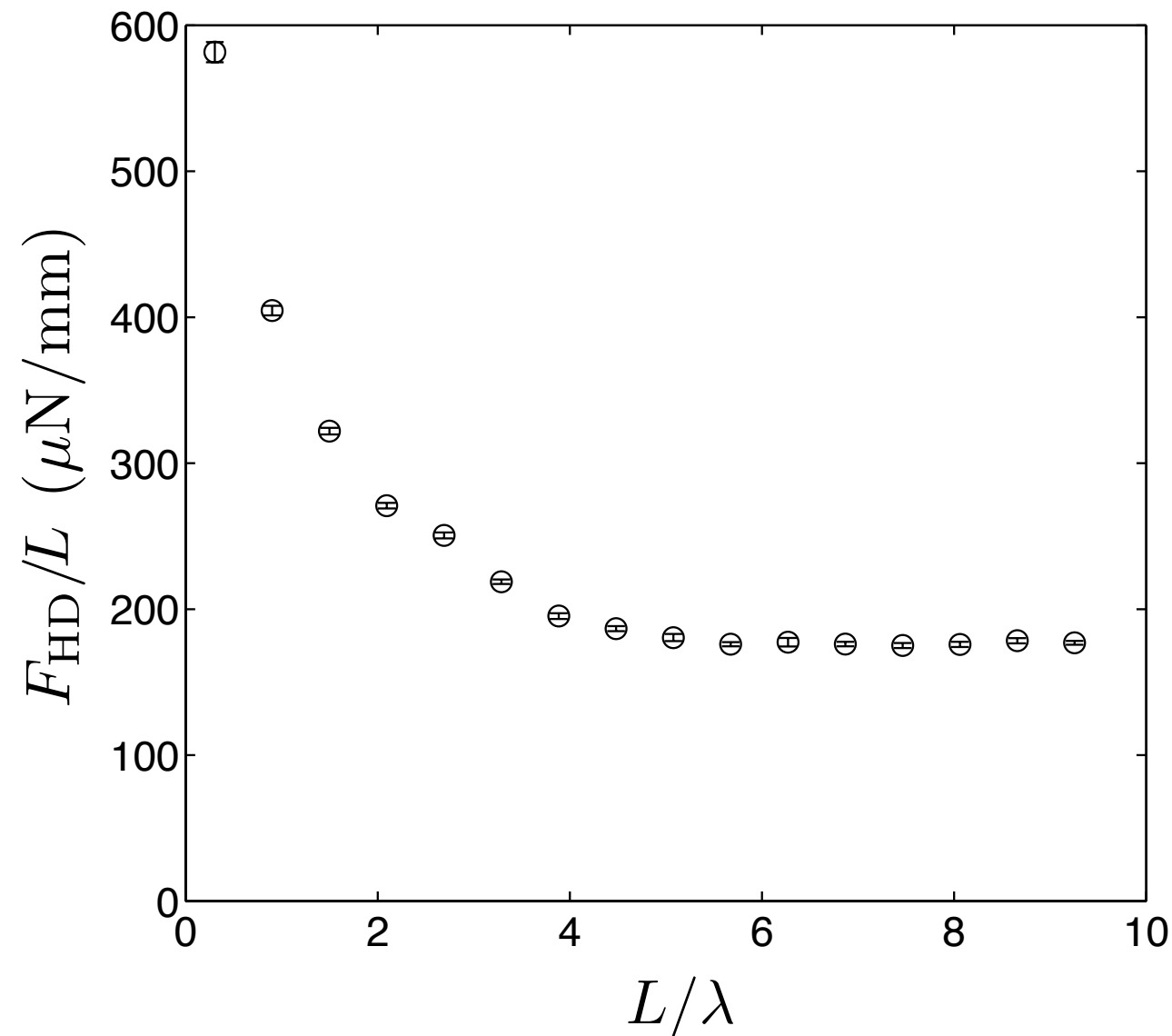
$$\text{Re} \sim 10^{-3}$$

Fluidic force on a rotating helix (Newtonian)

Fluidic force per unit length saturates with $L > 5 \lambda$

→ infinitely long helix

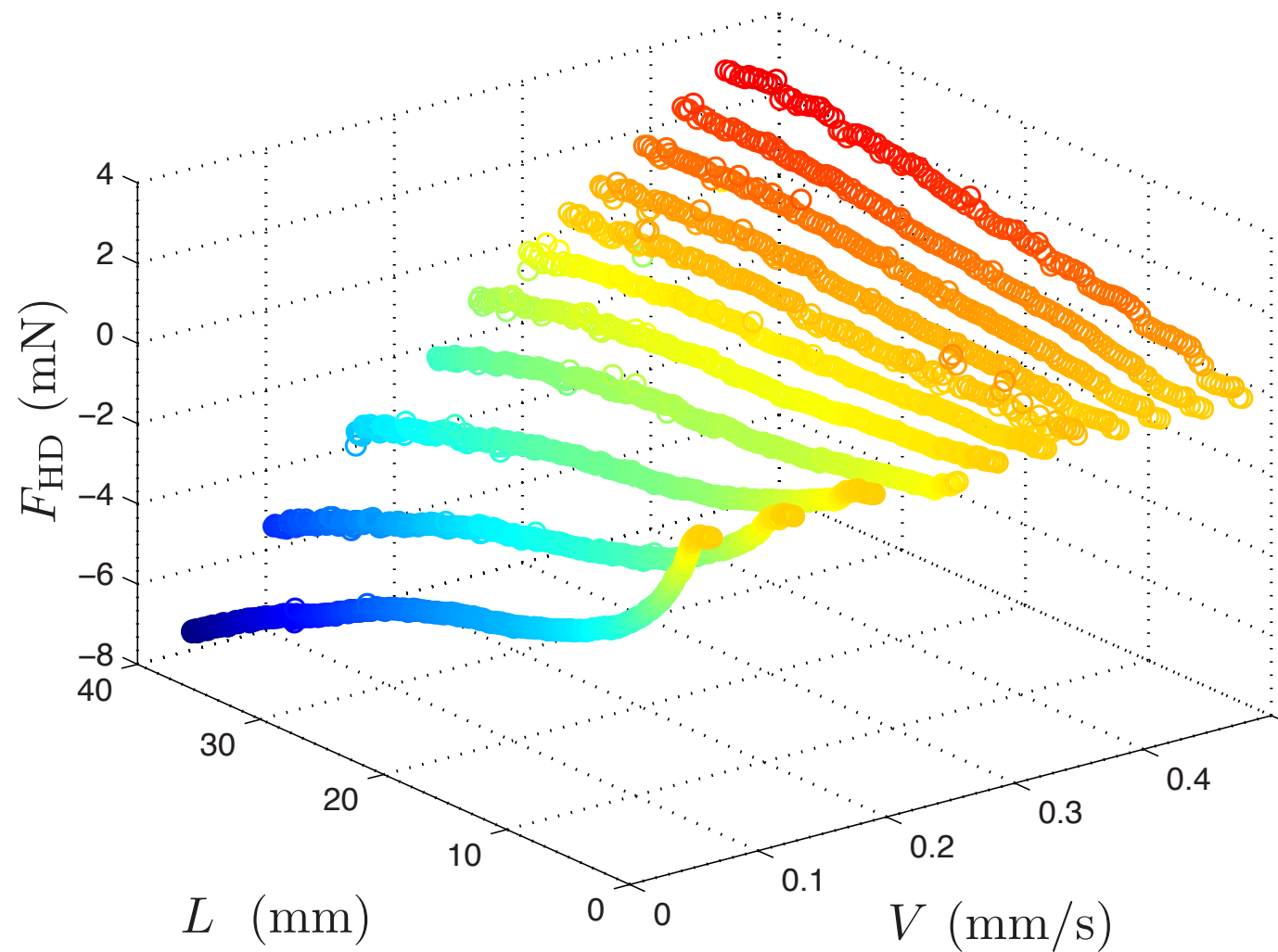
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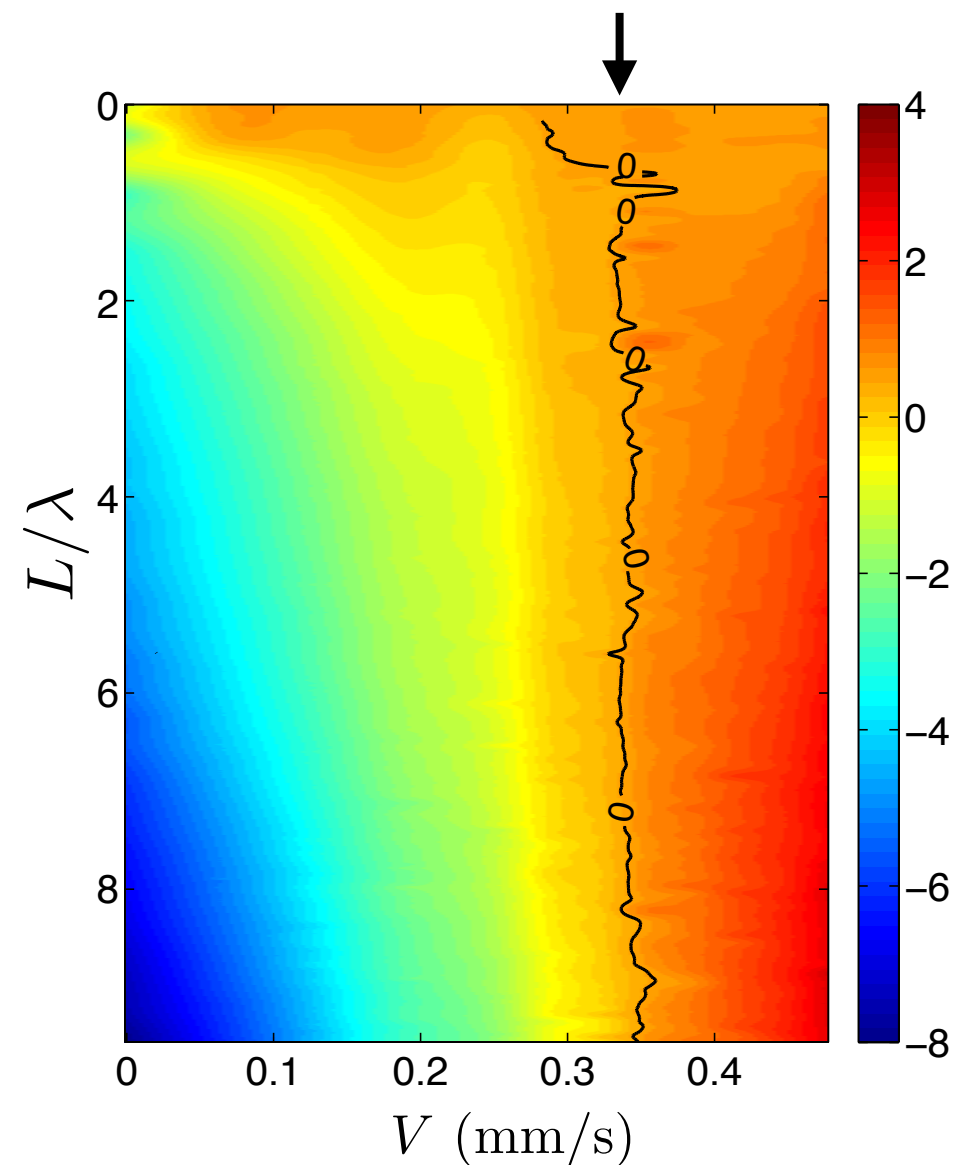
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Force-free swimming of a rotating helix (Newtonian)



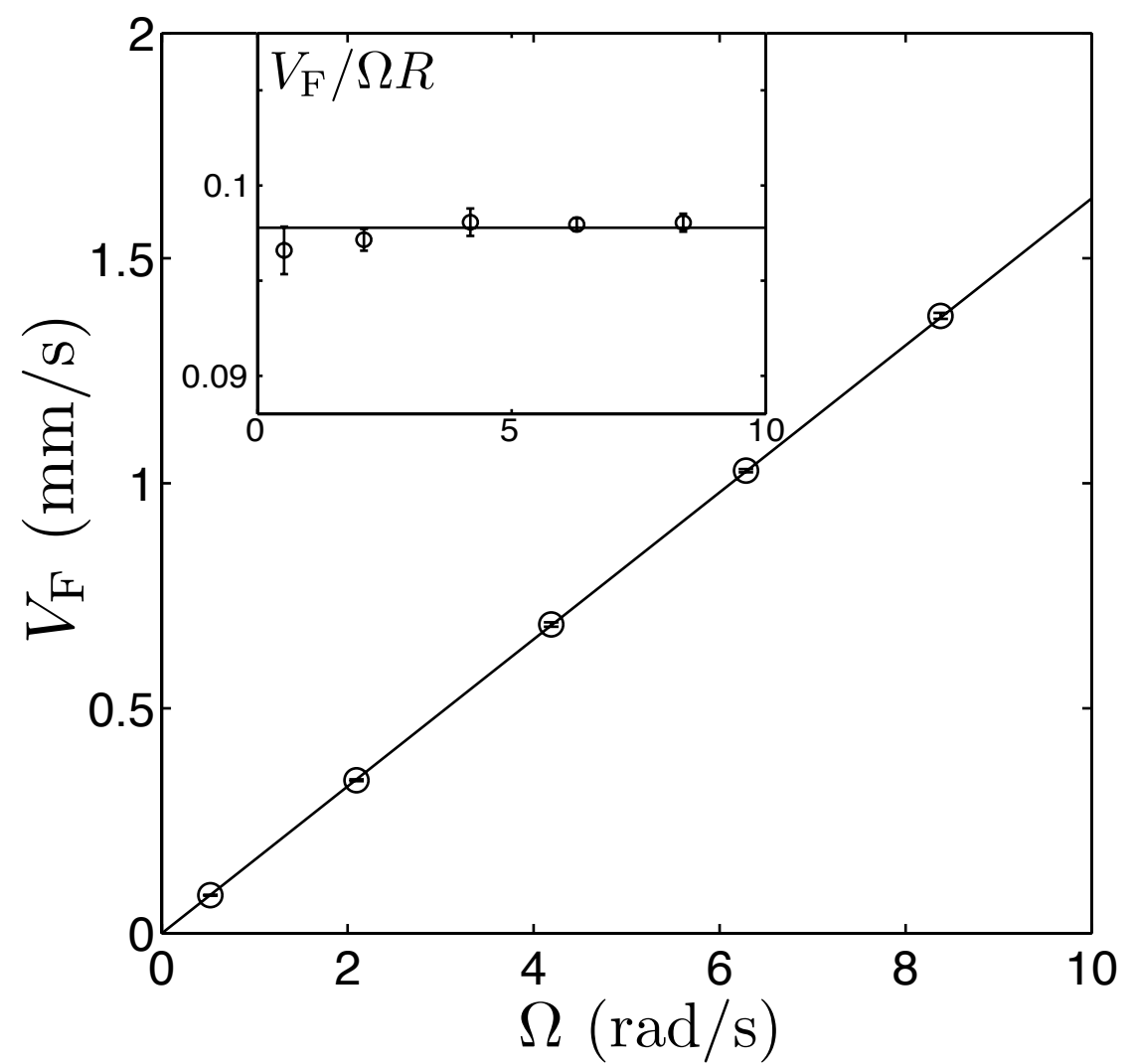
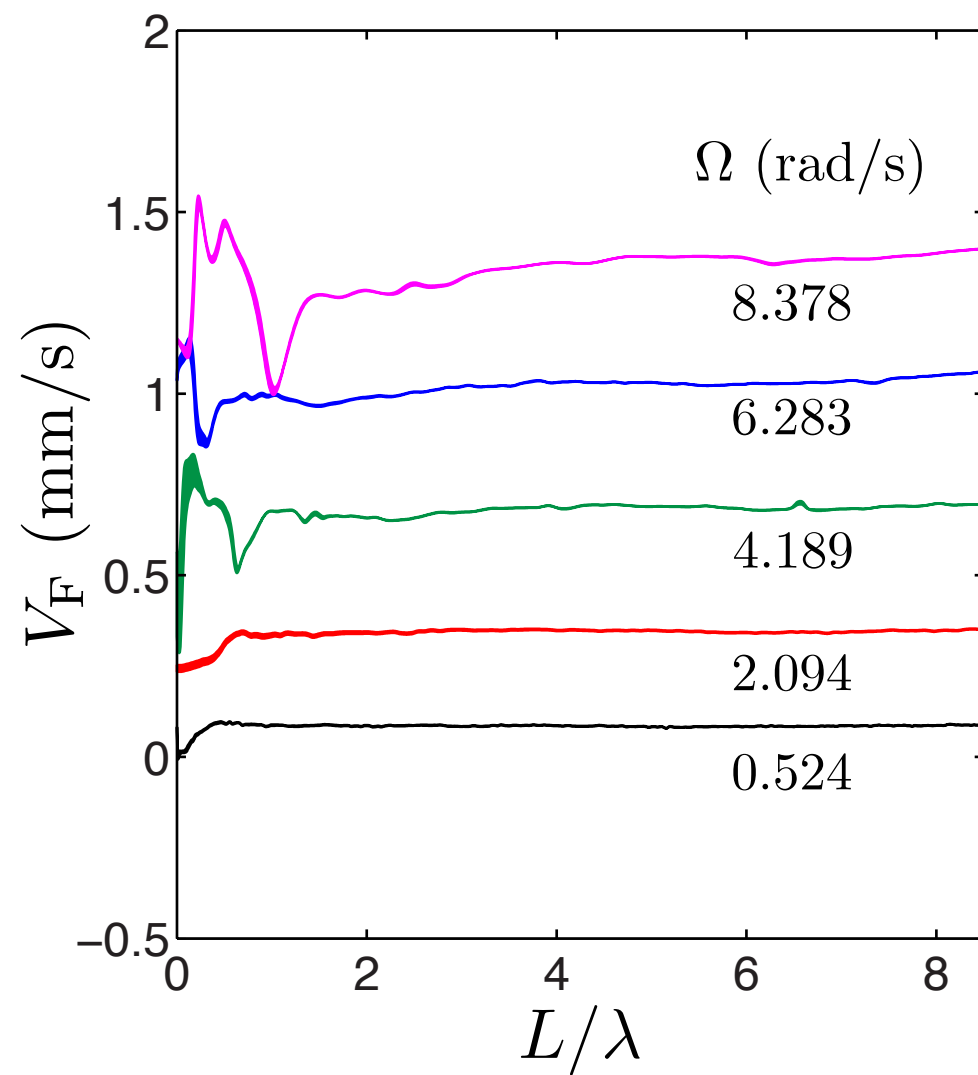
free swimming state ($F_{HD} = 0$)



sensitivity of force $\sim 10^{-5}$ N

sensitivity of swimming speed $\sim 0.3 \mu\text{m/s}$

Free swimming speed at different rotation rate



Computation of free swimming speed

1. Resistive force theory (no long-range interactions)

$$f_n = \mu C_n u_n, \quad f_s = \mu C_s u_s,$$

when swimming freely,

$$f_n \sin \theta + f_s \cos \theta = 0$$

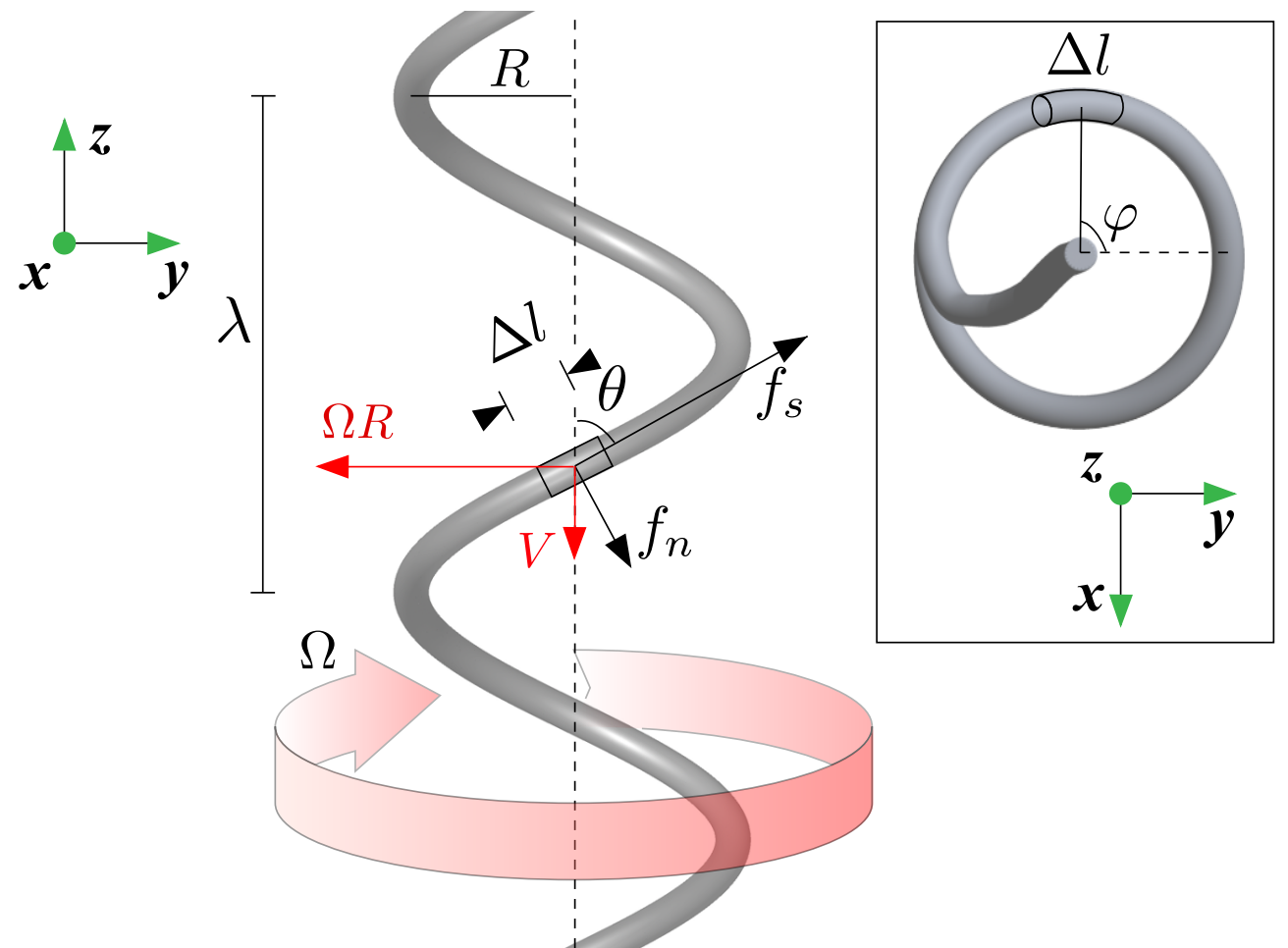
$$\downarrow$$

$$\frac{V_F}{\Omega R} = \frac{(C_n/C_s - 1) \sin \theta \cos \theta}{(C_n/C_s - 1) \sin^2 \theta + 1}$$

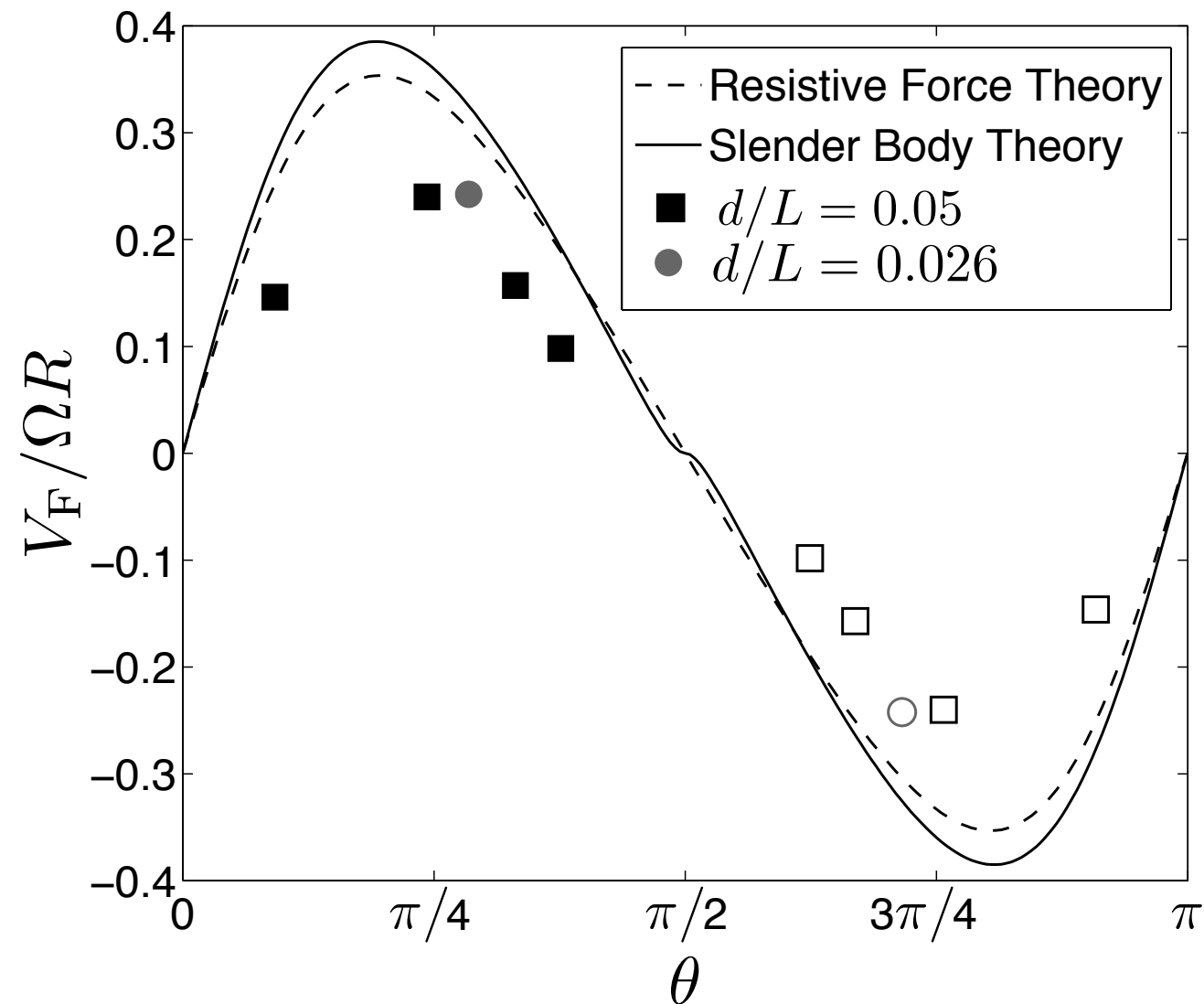
2. Slender body theory (incl. long range, no thickness effects)

$$V_F = \frac{f}{8\pi\mu} \int_{-\infty}^{\infty} \frac{\varphi \sin \varphi \cos \theta \csc^2 \theta}{(\xi(\varphi, \theta))^{3/2}} d\varphi,$$

$$\Omega R = \frac{f}{8\pi\mu \sin \theta} \int_{-\infty}^{\infty} \left(\frac{\cos \varphi}{(\xi(\varphi, \theta))^{1/2}} + \frac{\sin^2 \varphi}{(\xi(\varphi, \theta))^{3/2}} \right) d\varphi$$



Comparison between experiment and theory



d : diameter of the cross-section
of helical fiber

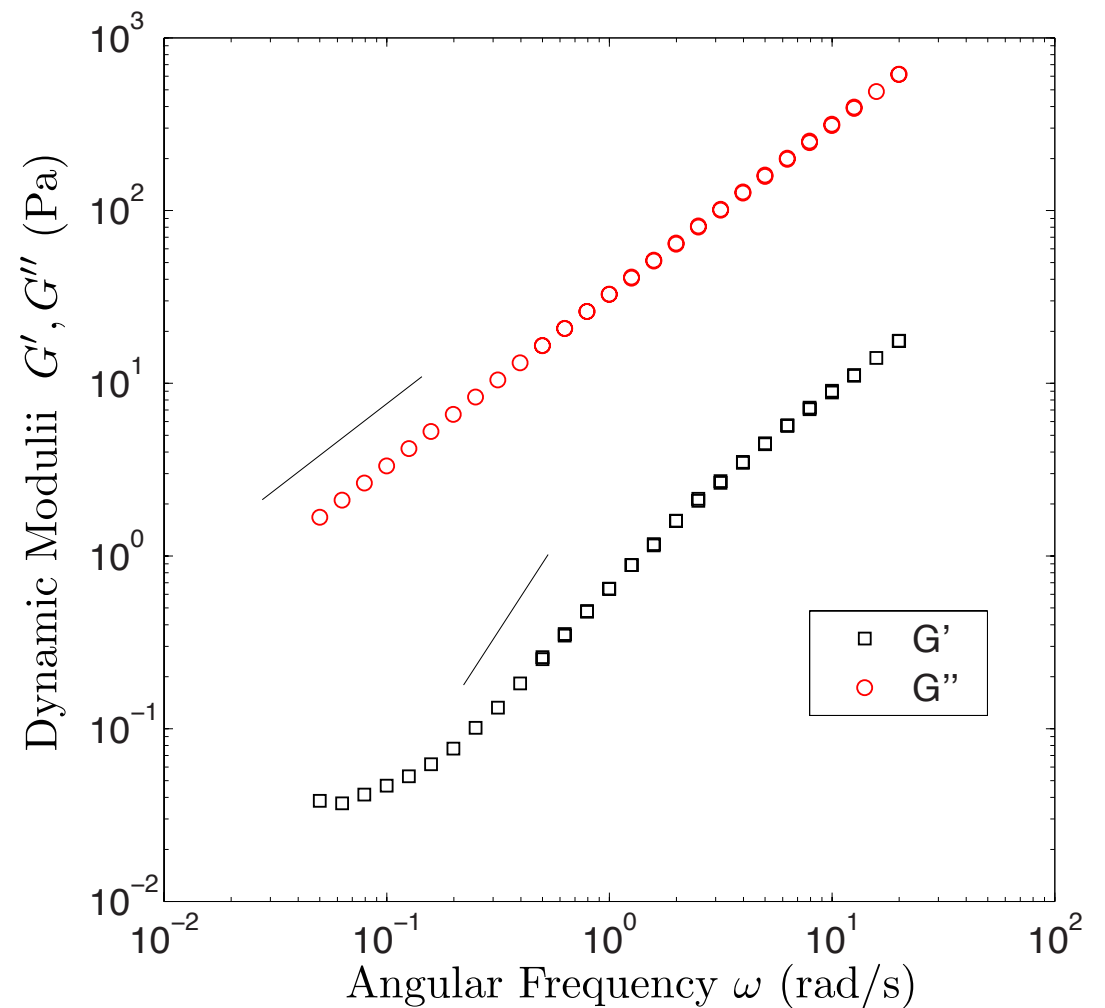
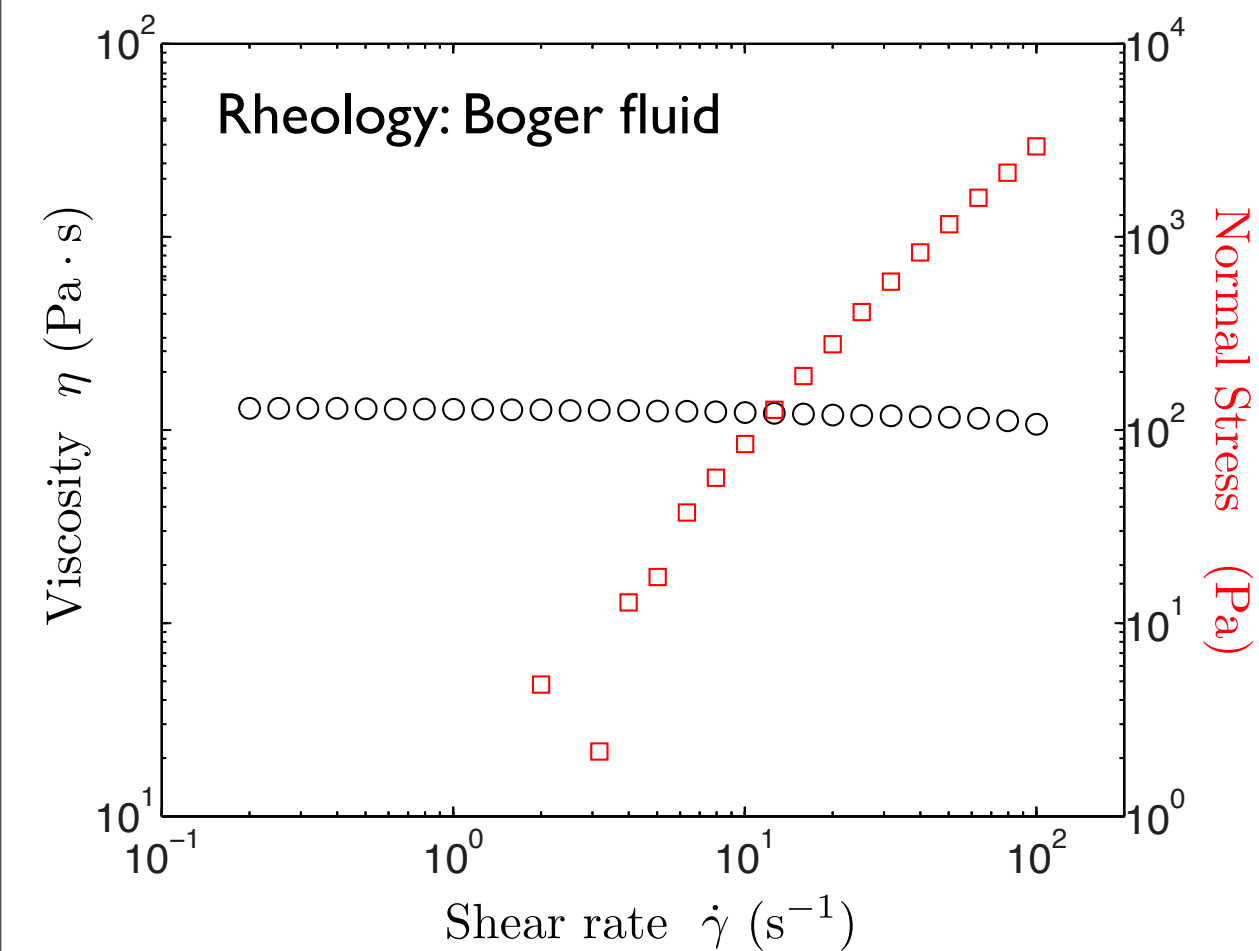
L : arc-length within a pitch

↑
Straight
filament

↑
Tightly
coiled

Rotating helix in Viscoelastic fluids

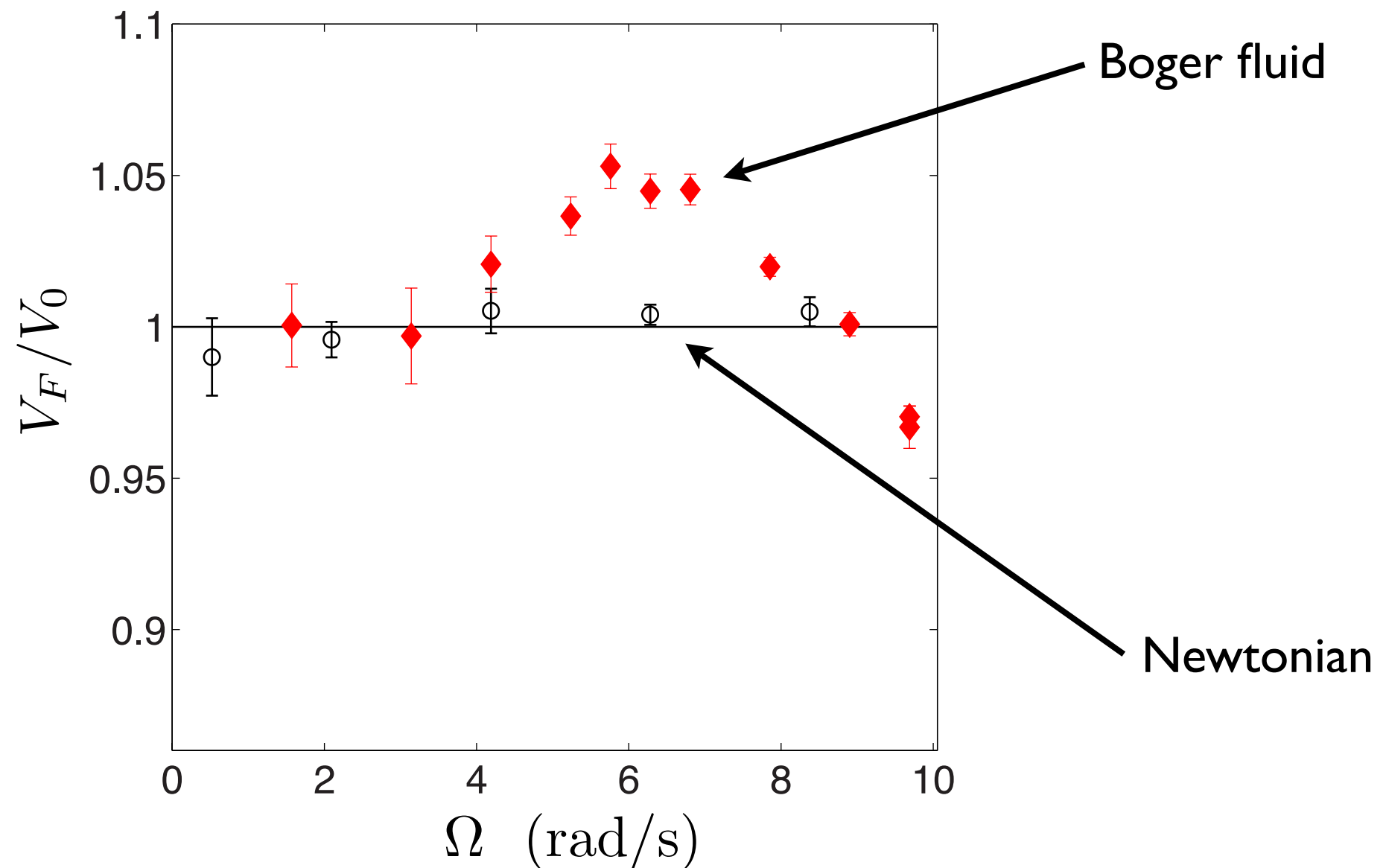
Polyisobutylene (PIB) suspended in Polybutene (PB)



Relaxation time: $\tau = 0.4 \text{ s}$

Rotating helix in Viscoelastic fluids

free swimming speed of rotating helix in viscoelastic fluids



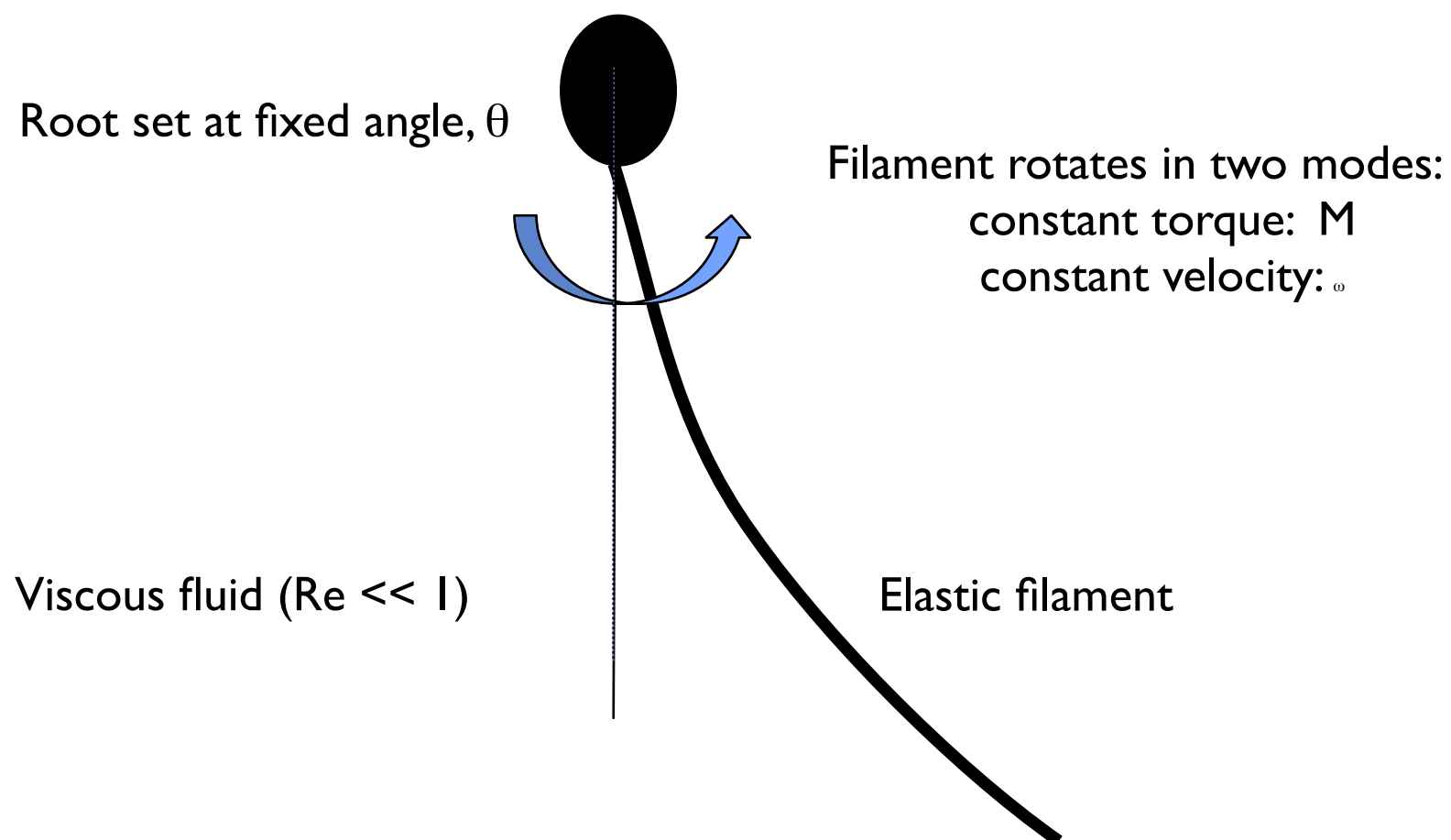
Next: fluid with stronger elasticity

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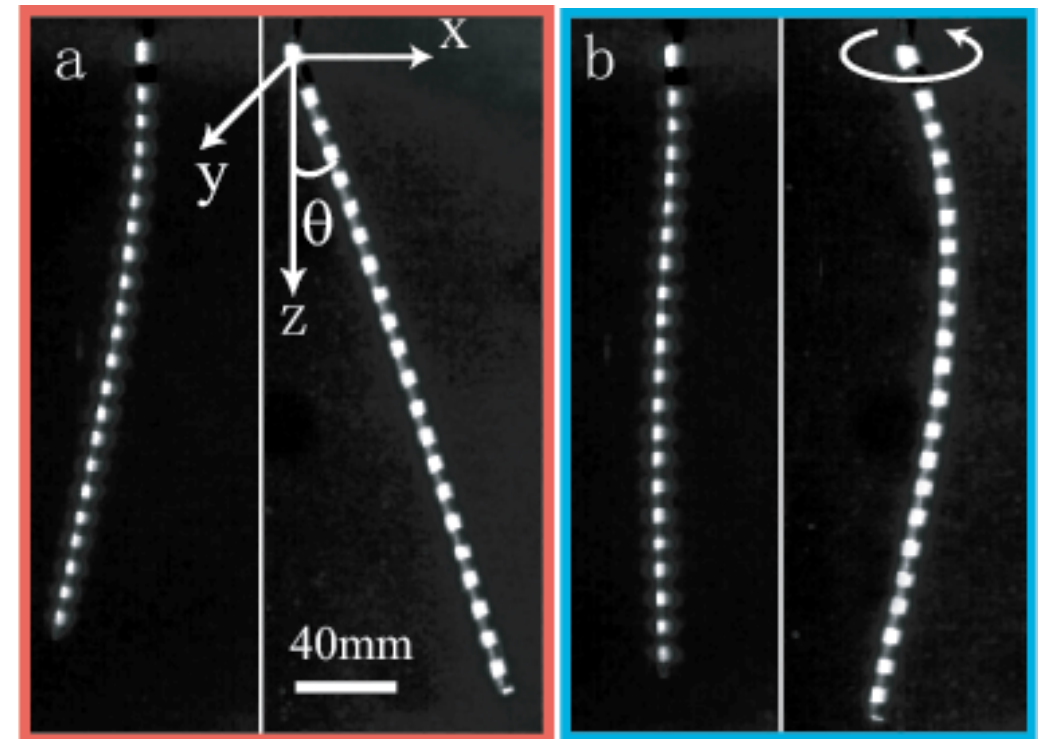
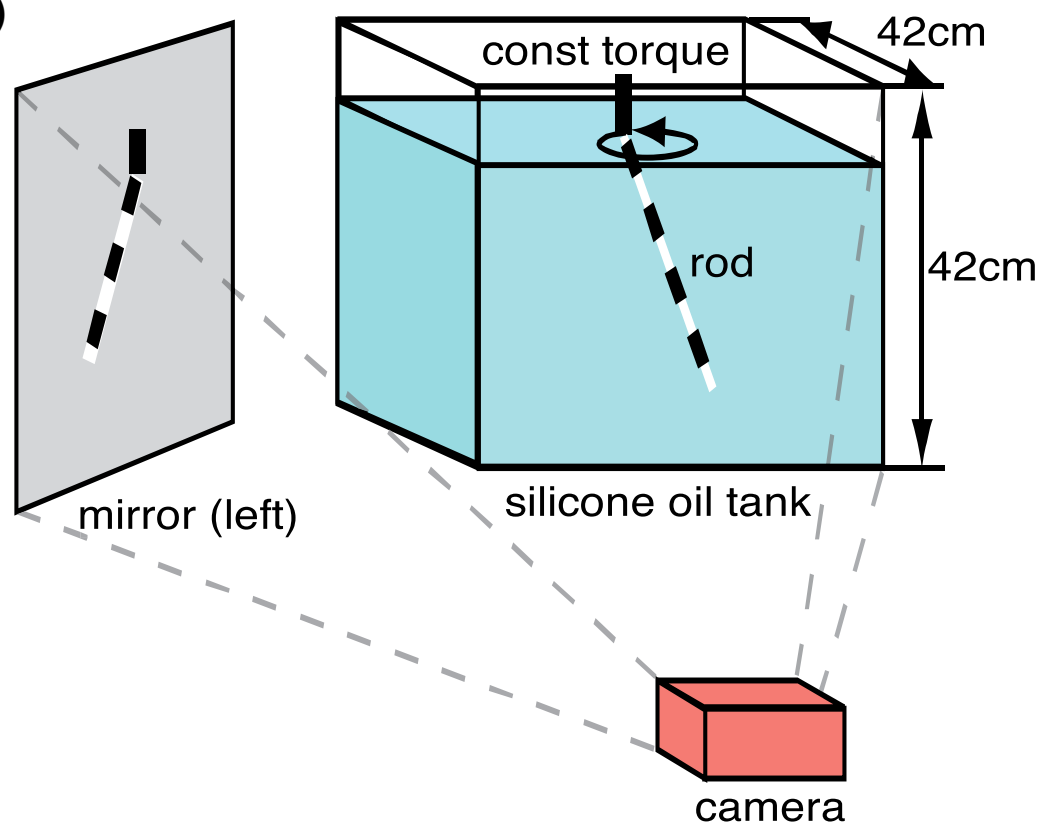
Elastic filament interacting with a viscous flow

- Motivation
 - Understand simplified behavior of flexible filaments in viscous flows
 - Look at a simpler model for microrobotic propulsion
 - Understand previous numerical experiments
 - (Manghi, Schlagberger & Netz, *PRL* 2006)



Shape Bifurcation of an Elastic Rotating Rod

(Video Sped up x10)



before bifurcation

after bifurcation

Experiment Parameters :

bending modulus $A \approx 3 \times 10^{-3} \text{ N/m}^2$

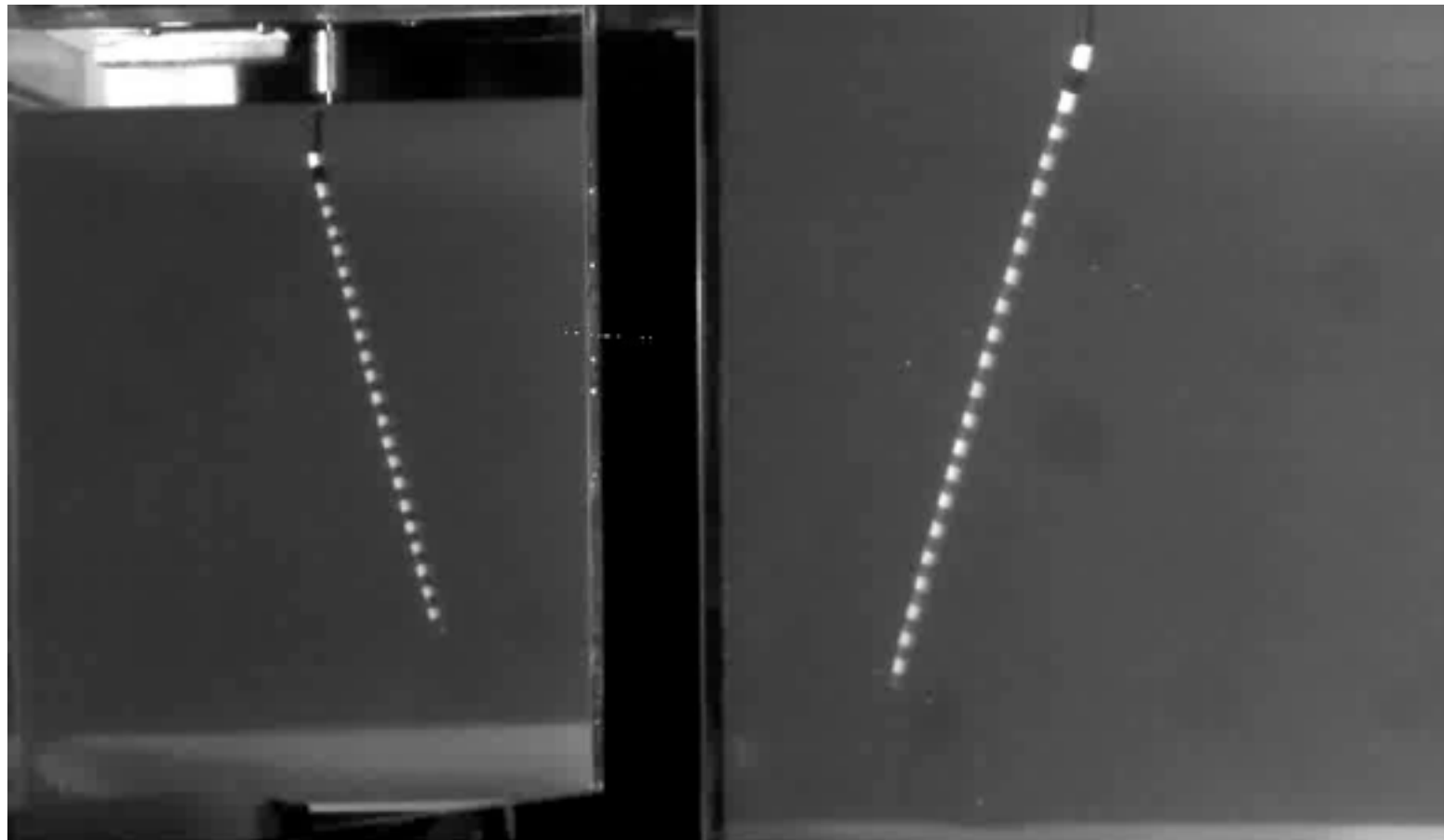
radius of rod $a = 2.5 \text{ mm}$

viscosity $\eta \approx 110 \text{ N}\cdot\text{s/m}^2$

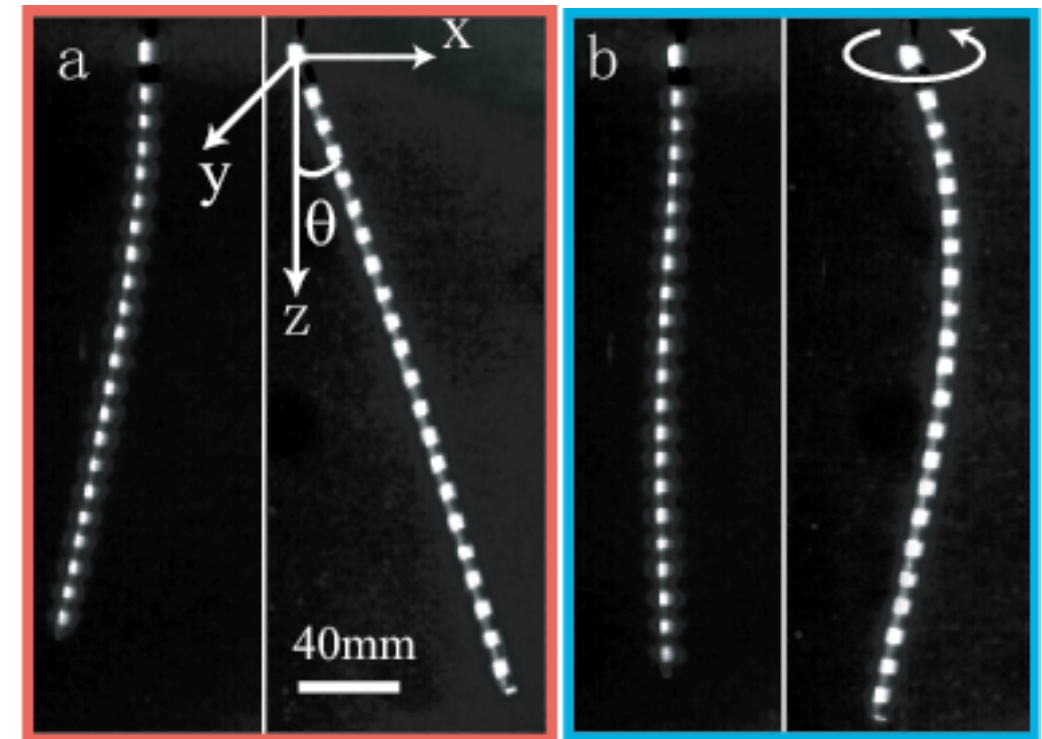
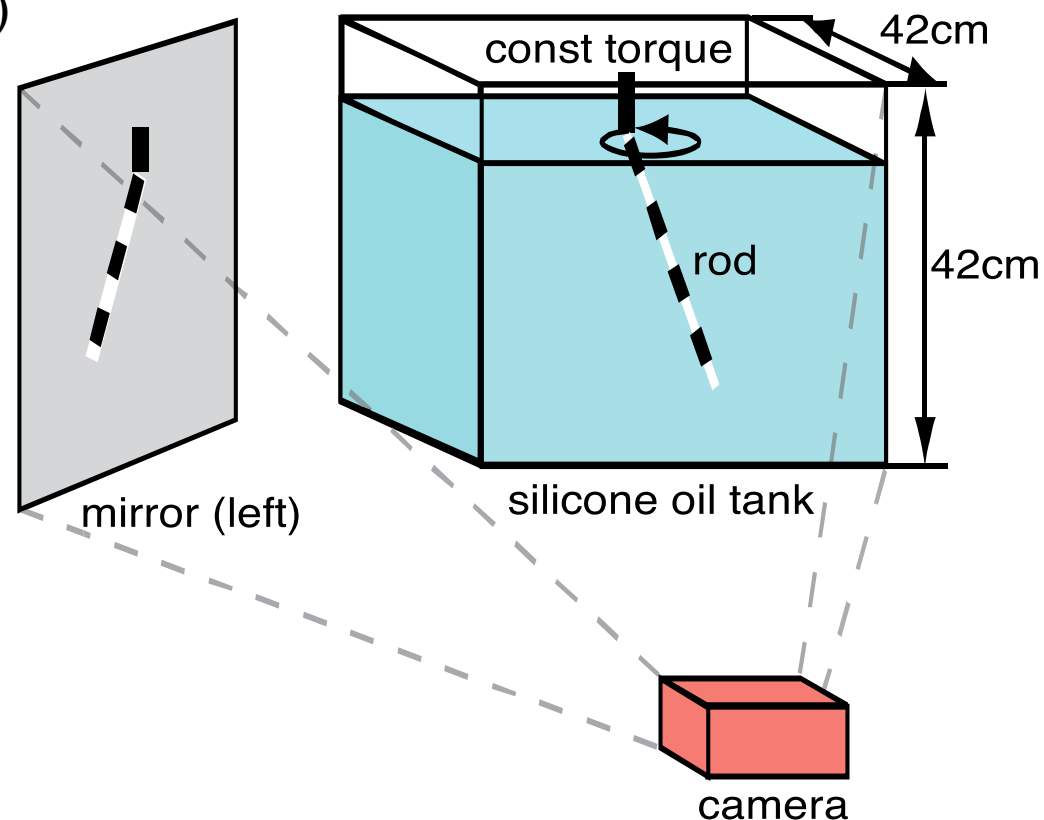
length of rod $L = 200 - 280 \text{ mm}$

tilted angle $\epsilon \approx 20^\circ - 30^\circ$

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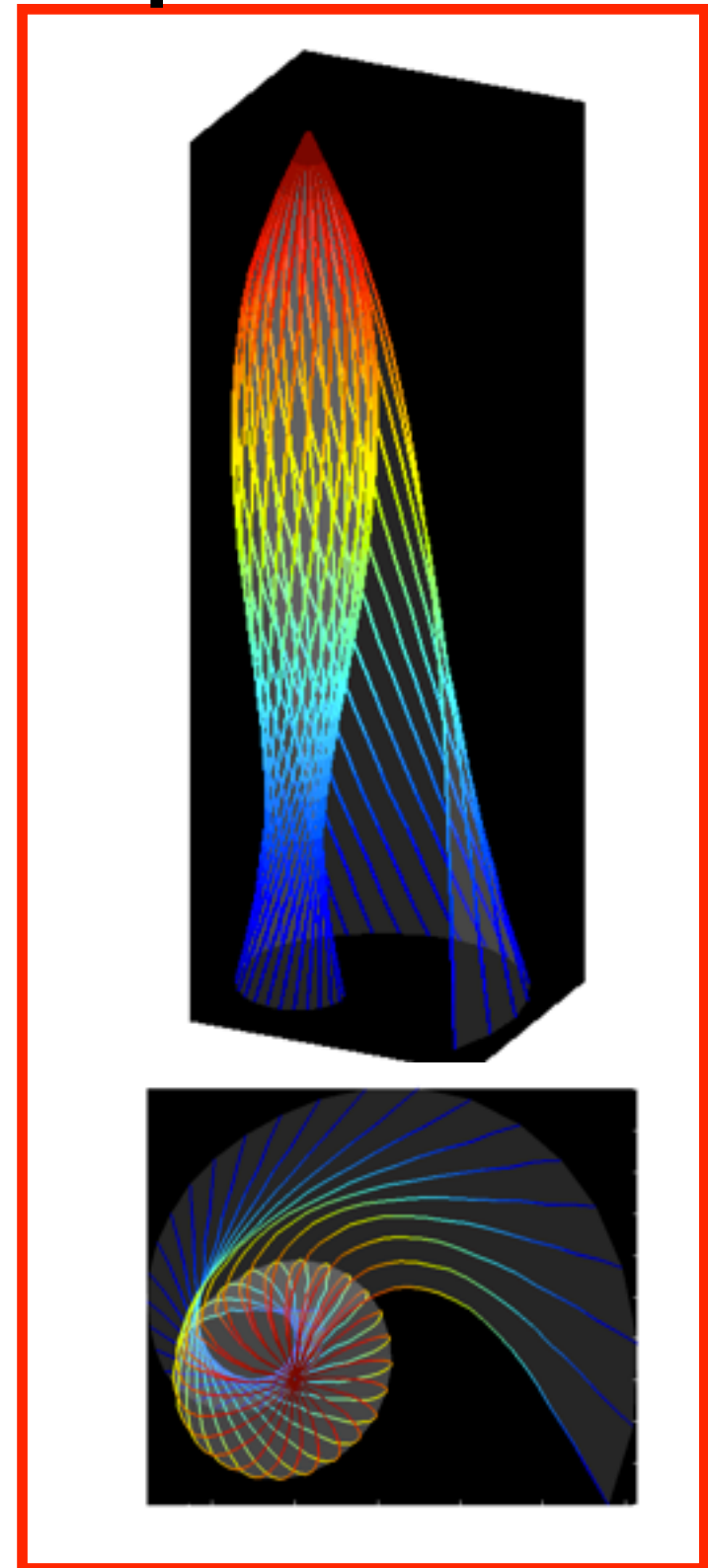
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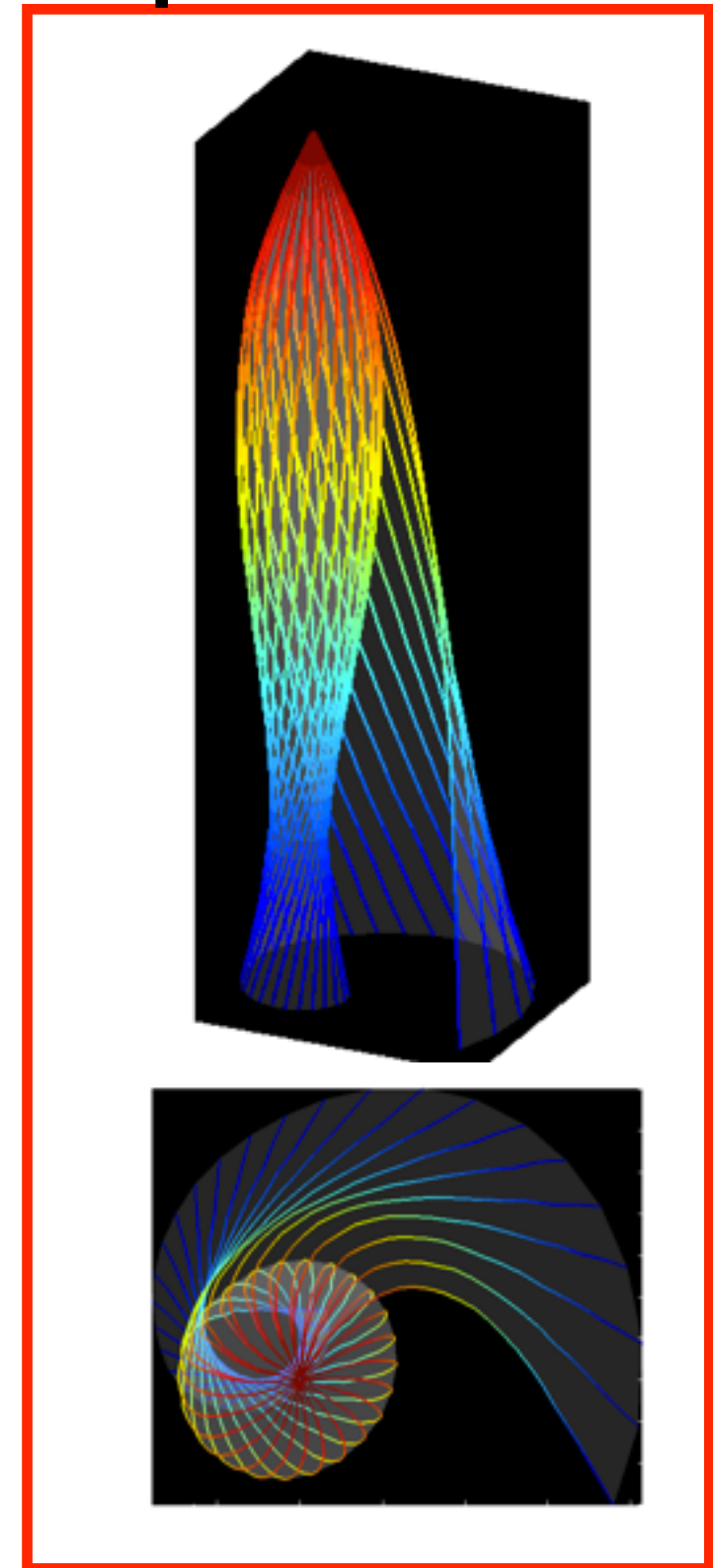
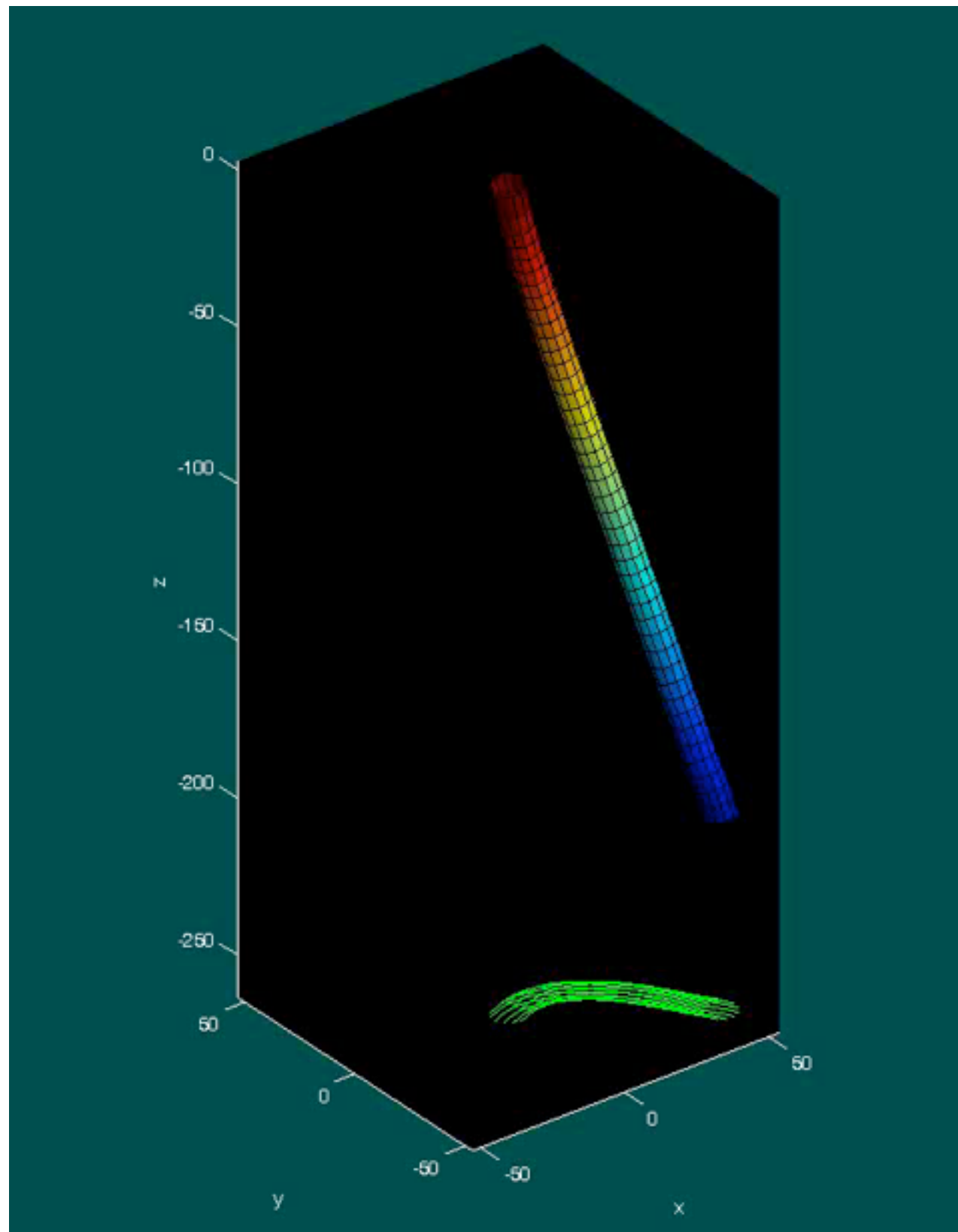
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Reconstructed motion from expt. data



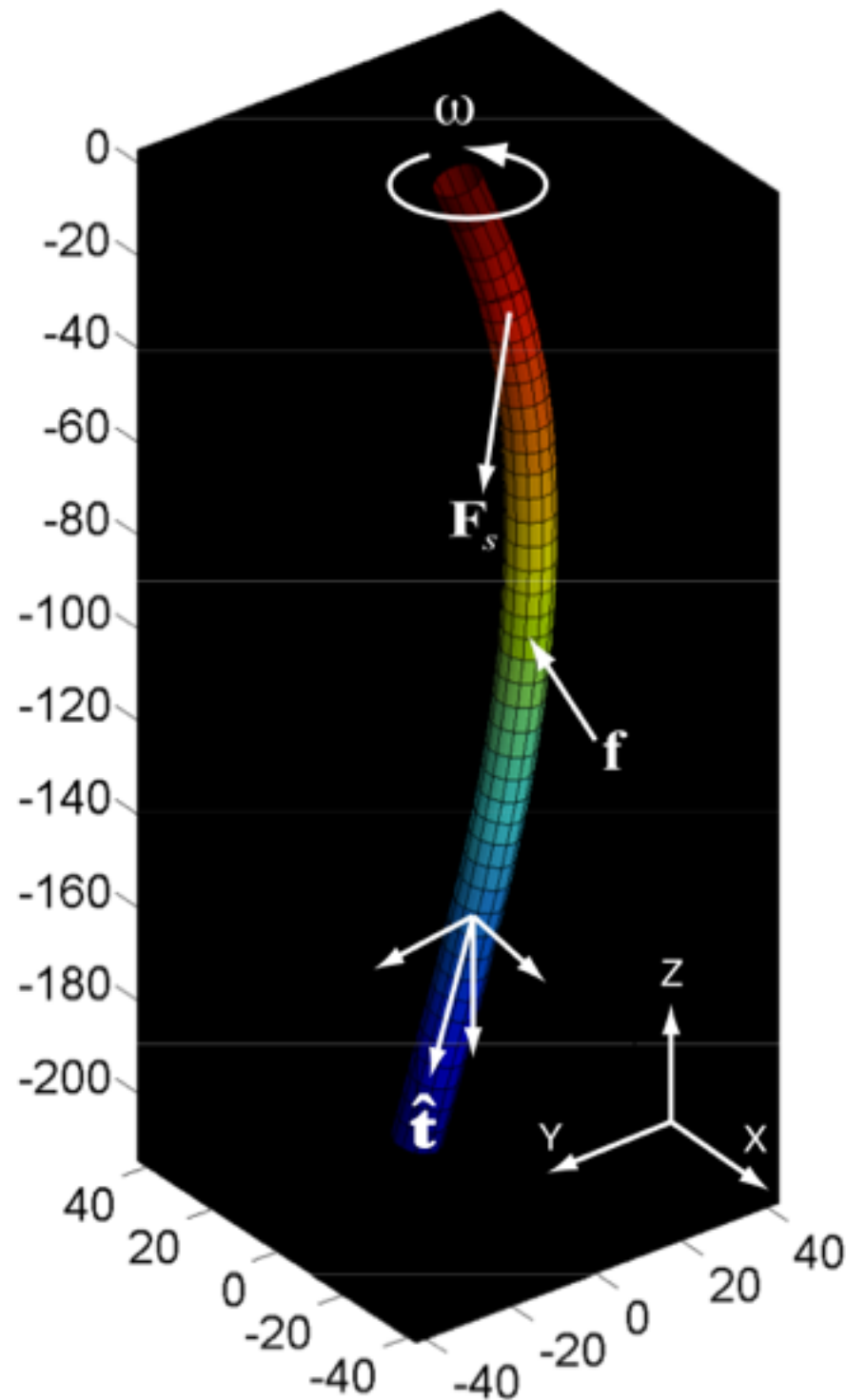
Motion around the bifurcation

Reconstructed motion from expt. data



Motion around the bifurcation

Theory



- Stokes flow (reversible flow)

$$\mu \nabla^2 \mathbf{v} = \nabla p$$

$$\nabla \cdot \mathbf{v} = 0$$

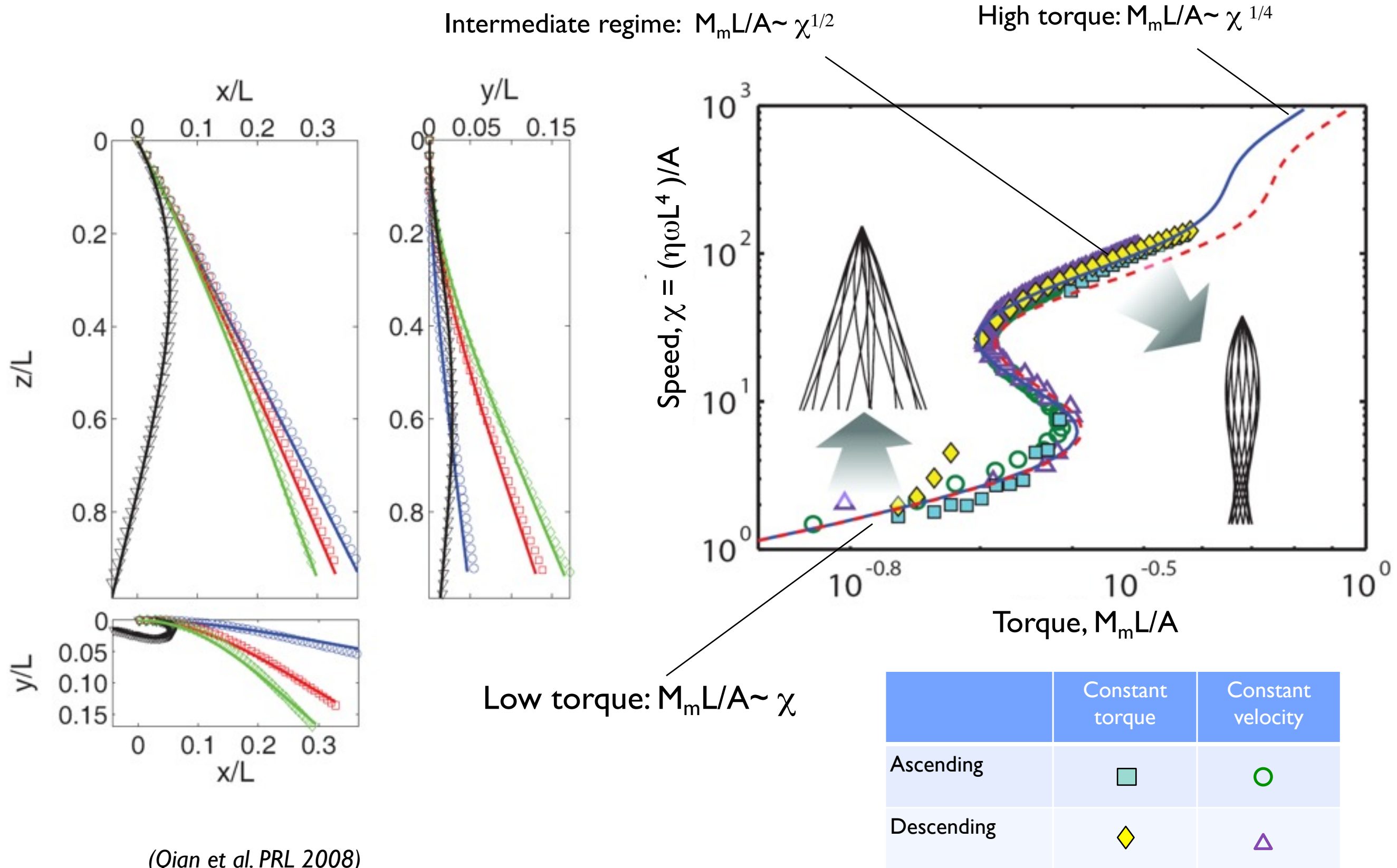
- Analyze in the rotating frame (steady)
- No forces due to non-inertial frame
- disregard twist
 - Only bending stiffness
- resistive force theory:

$$f_n = \mu C_n u_n, \quad f_s = \mu C_s u_s,$$

- Hydrodynamic forces are local
- balance of forces and moments
- Key non-dimensional parameters:

$$\theta, \quad \chi = \frac{\mu \omega L^4}{A} = (L/l_v)^4, \quad M = M_{mot} L/A$$

Comparison between theory and experiment

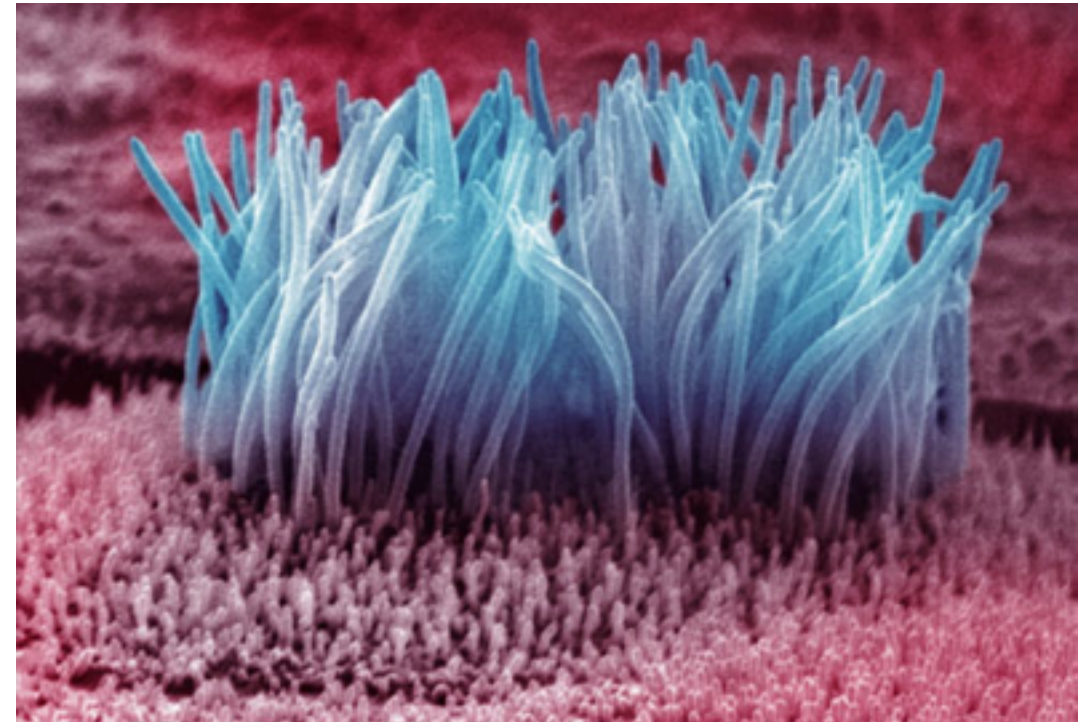


Interesting questions

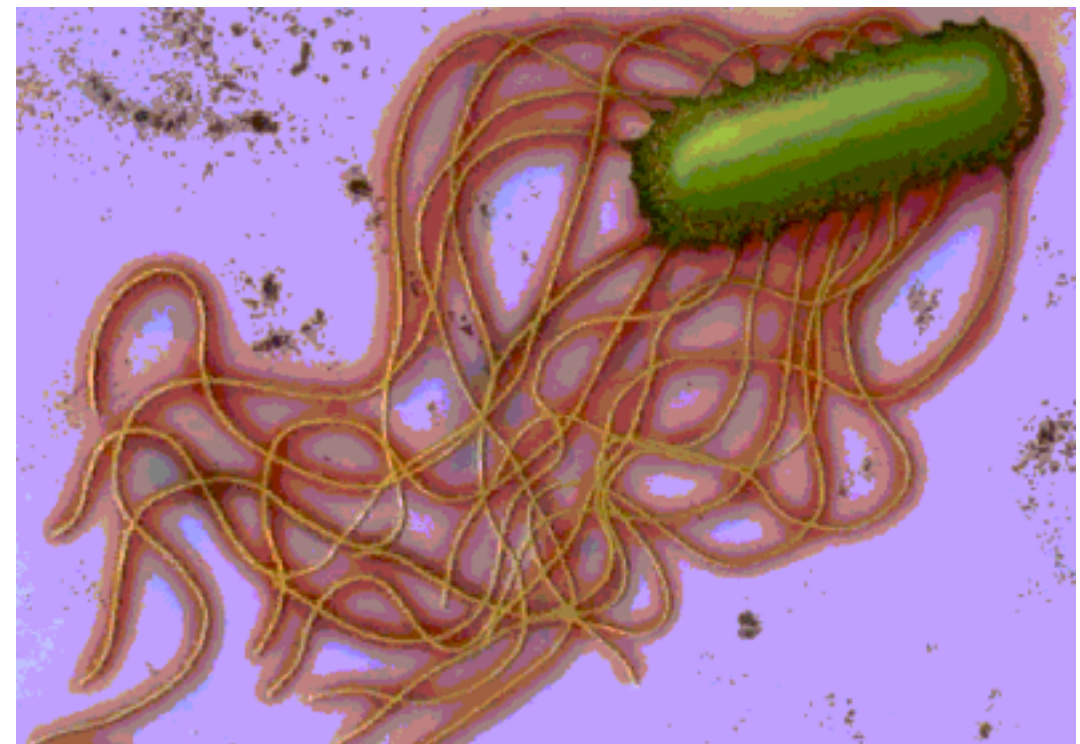
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Hydrodynamic interactions

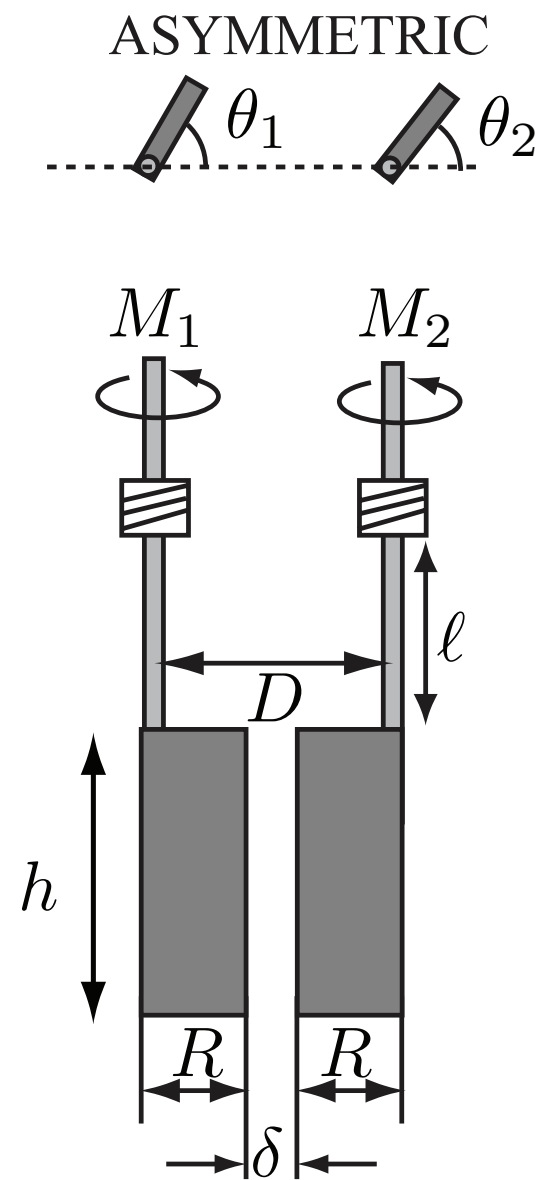
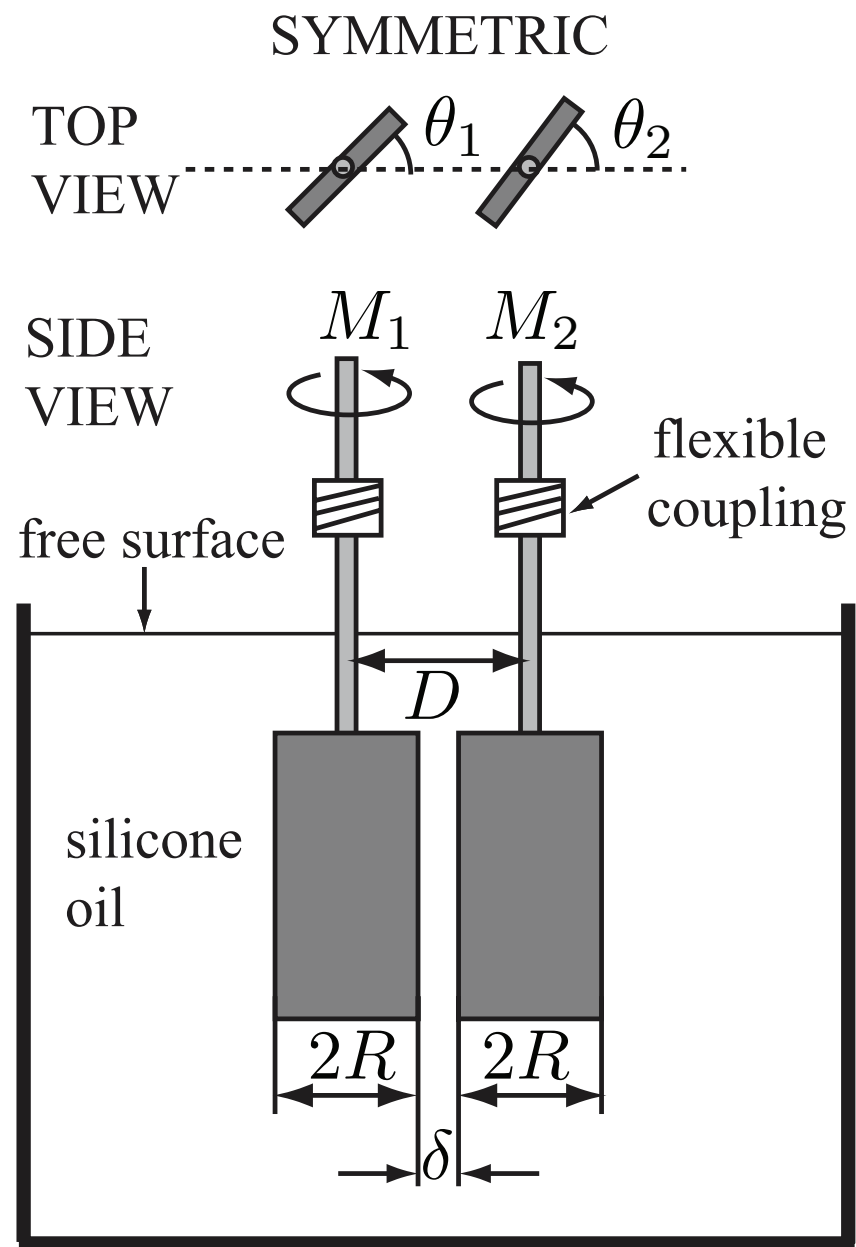
- Cilia and flagella occur in many biological (and engineering) systems.
- Collective motion important for transport and motility
- Interesting questions for flagella and cilia:
 - synchronization through hydrodynamic interactions
 - synchronization vs. chaotic motion
 - role of compliance
 - time-scales for interactions
 - effects of multiple filaments
 - effects of long range hydrodynamic forces



webmd.com



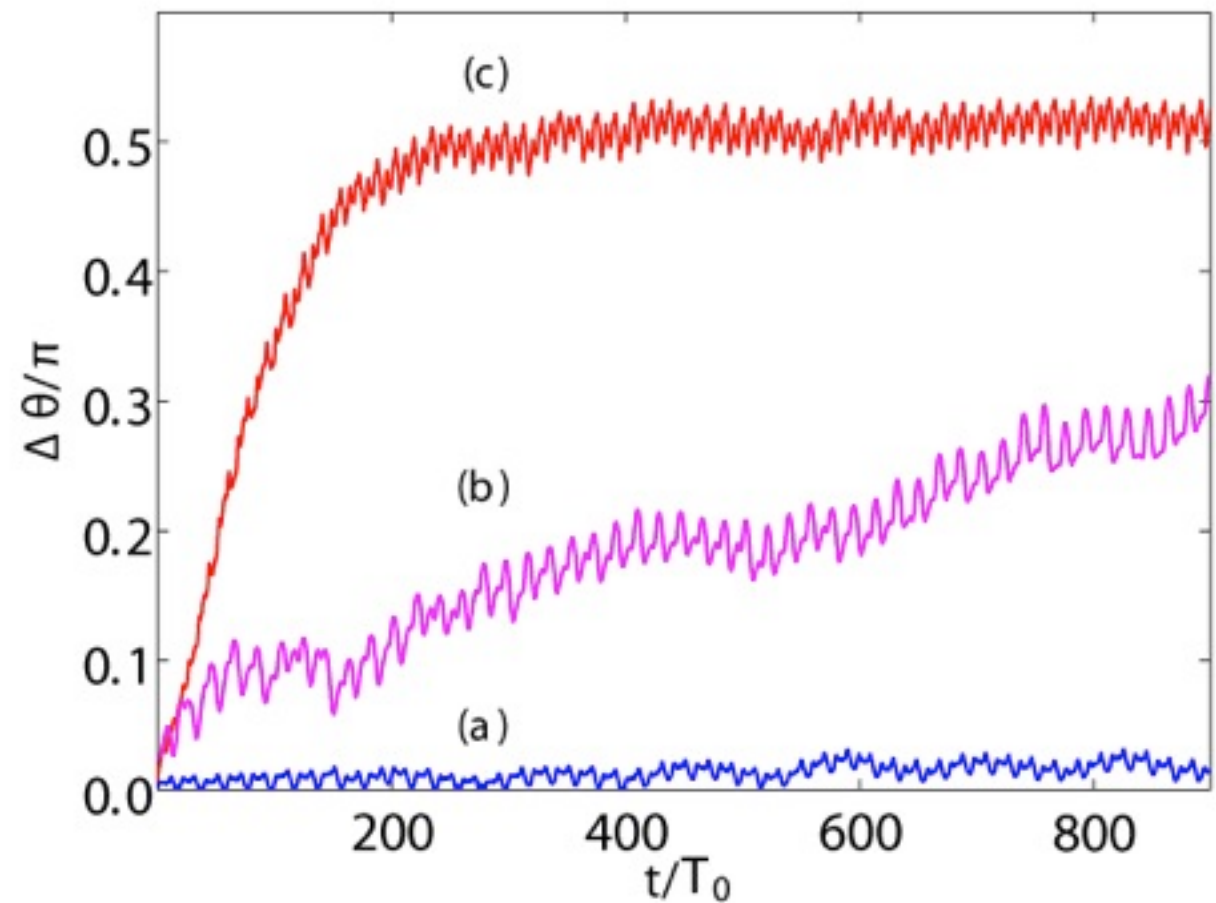
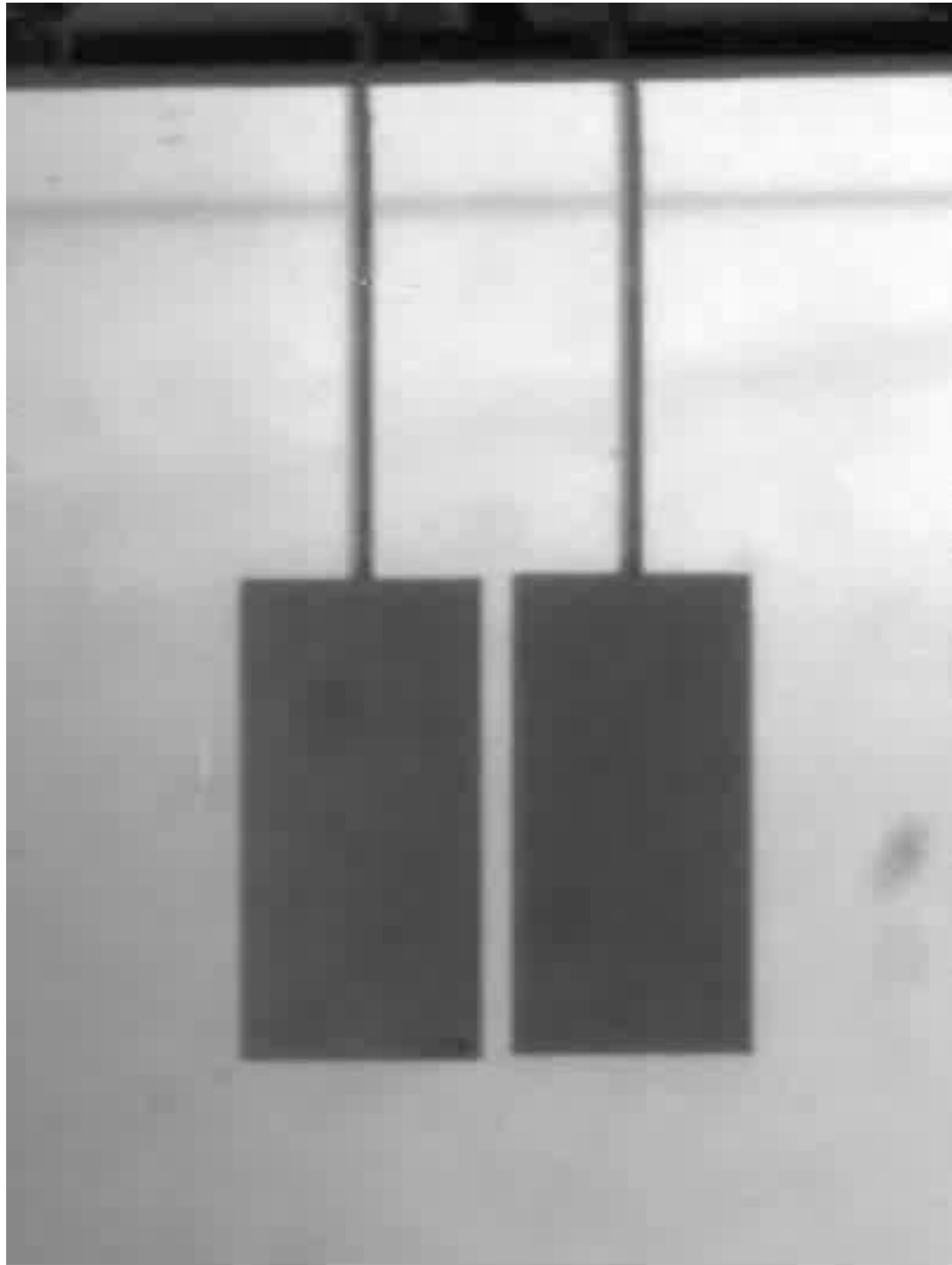
Two-paddle model system



$$Re = \mathcal{O}(10^{-3})$$



Two-paddle interactions



$$T_o = 6\pi\mu R^3/M$$

- a) Rigid shafts – no synchronization
- b) Rigid shafts – torque mismatch
– Phase wandering
- c) Flexible shafts – synchronization!

(Qian et al, PRE 2009)

Numerical simulation - Regularized

$$\mu \nabla^2 \mathbf{v} - \nabla p + \mathbf{F} = 0$$

$$\nabla \cdot \mathbf{v} = 0$$

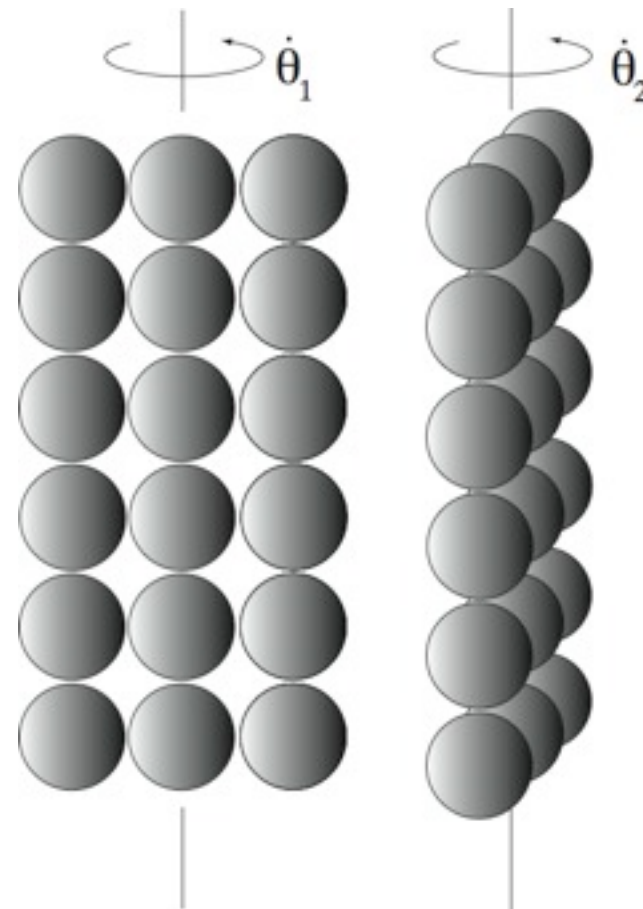
$$\mathbf{F} = \frac{15\epsilon^4}{8\pi} \frac{1}{(r^2 + \epsilon^2)^{7/2}}$$

(Cortez, 2001)

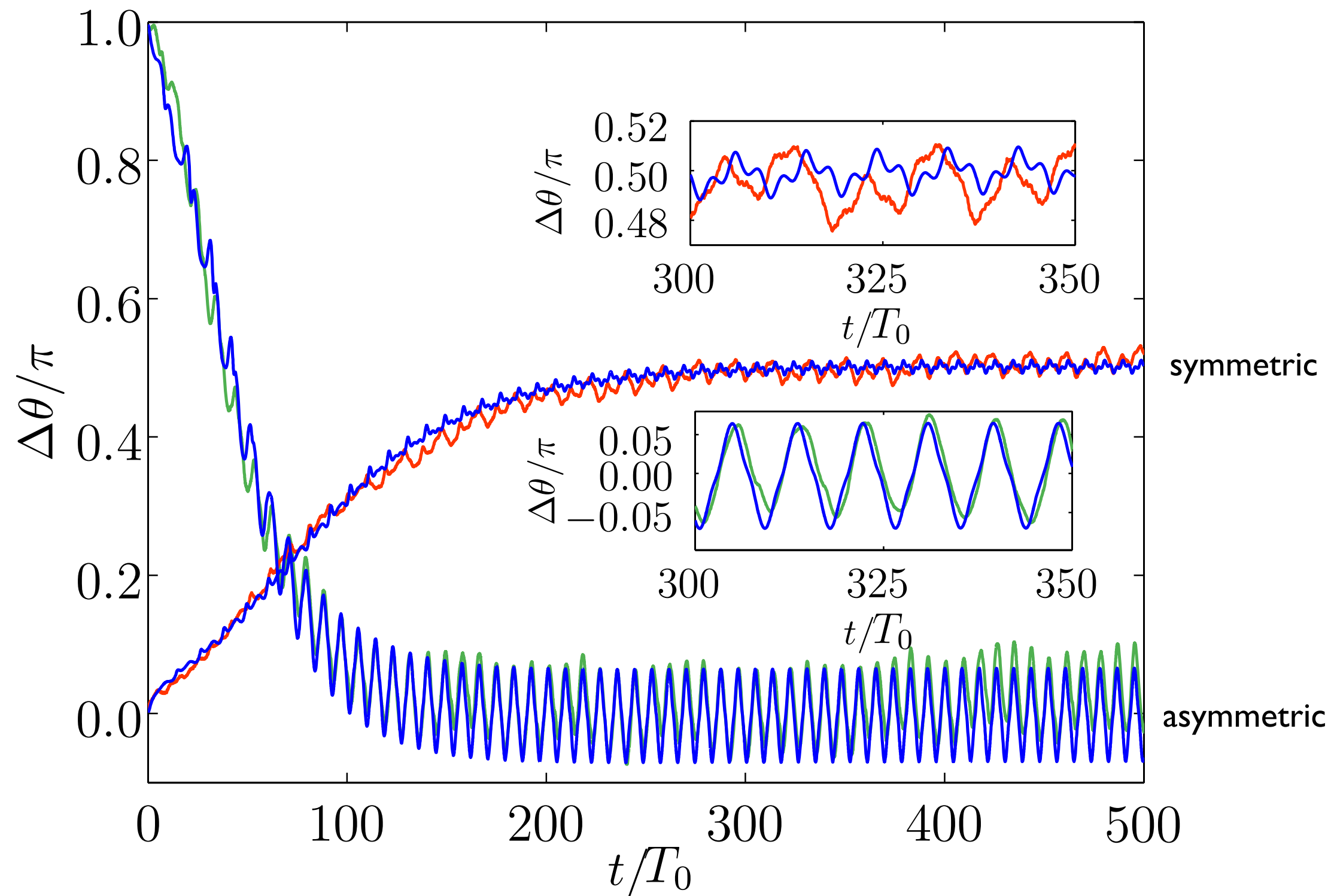
$$\begin{pmatrix} M_1 \\ M_2 \end{pmatrix} = \begin{pmatrix} A(\theta_1, \theta_2) & B(\theta_1, \theta_2) \\ B(\theta_1, \theta_2) & A(\theta_1, \theta_2) \end{pmatrix} \begin{pmatrix} \dot{\theta}_1 \\ \dot{\theta}_2 \end{pmatrix}$$

Kim & Powers (2004) - rigid helices – no synchronization

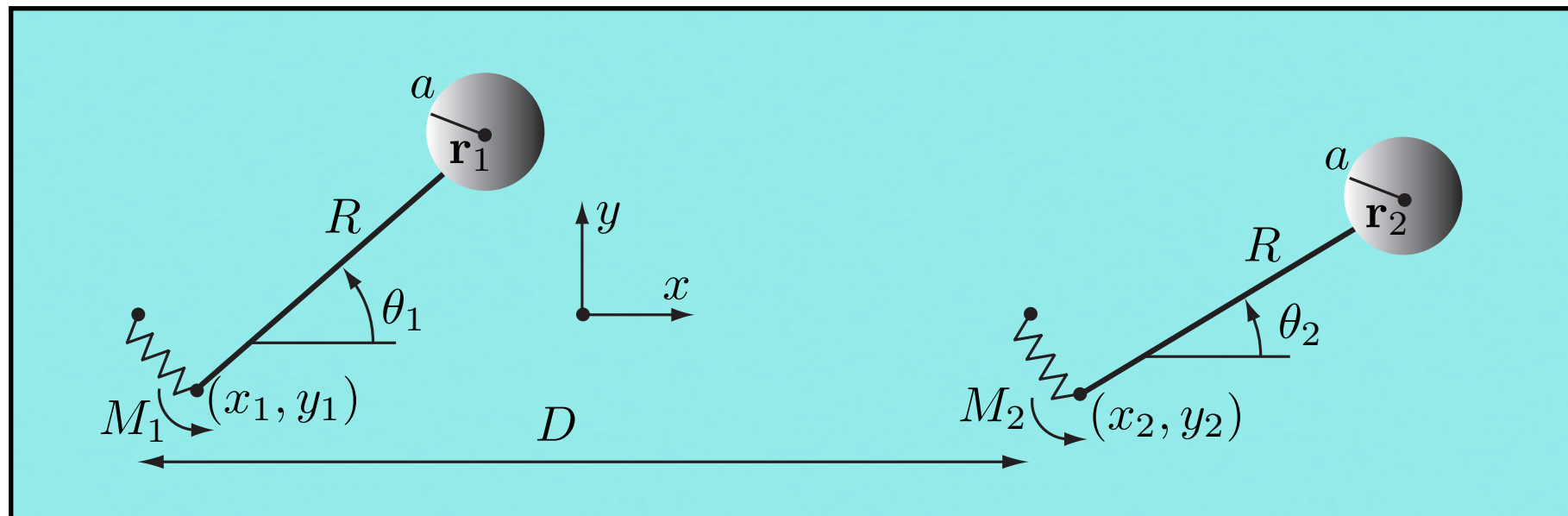
Reichert & Stark (2005) - flexible coupling leads to synchronization



Simulation and experiment



Analytical approach



velocity on ball i , induced by ball j

$$v_i = \frac{F_i}{6\pi\mu a} + \frac{1}{8\pi\mu} \sum_{i \neq j} \left[\frac{F_j}{|r_{ij}|} + \frac{(F_j \cdot r_{ij})r_{ij}}{|r_{ij}|^3} \right]$$

ignore “far field” term

$$a \ll R \ll D$$

Sum of forces and moments; for two balls:

$$-\frac{\Delta M}{M_1} + \Delta\dot{\theta} + \frac{9}{8} \frac{a}{D} \Delta\dot{\theta} \cos \Delta\theta + \frac{3}{8} \frac{a}{D} \cos(2\bar{\theta}) = 0$$

$$\frac{\Delta M}{M_1} + 2 - 2\dot{\bar{\theta}} - \frac{3}{4} \frac{a}{D} \dot{\bar{\theta}} [-3 \cos \Delta\theta + \cos(2\bar{\theta})] = 0$$

Niedermeyer et al.
(Chaos, 2008)

Qian et al.
(PRE, 2009)

multiple-scale (slow) evolution equation for phase difference: $(\mu\omega/k \ll 1)$

$$\frac{\langle \Delta\theta \rangle}{dt} = -\frac{9}{2} \frac{M_1}{kR^2} \frac{a}{D} \sin \langle \Delta\theta \rangle + \frac{\Delta M}{M_1}$$

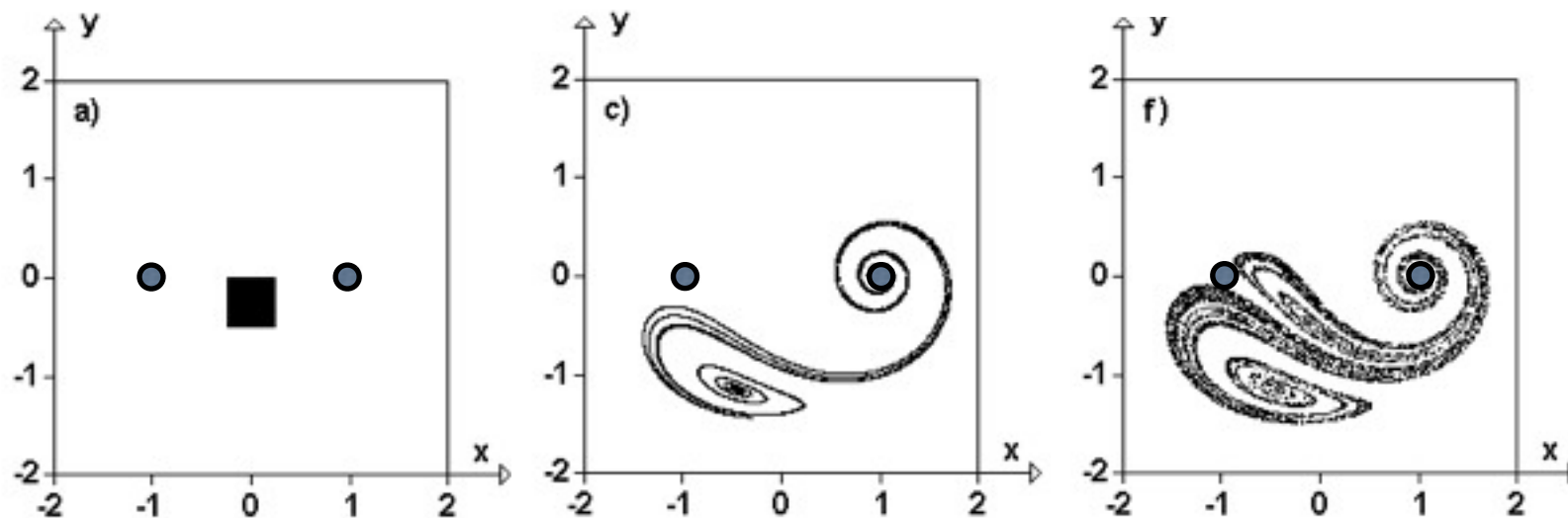
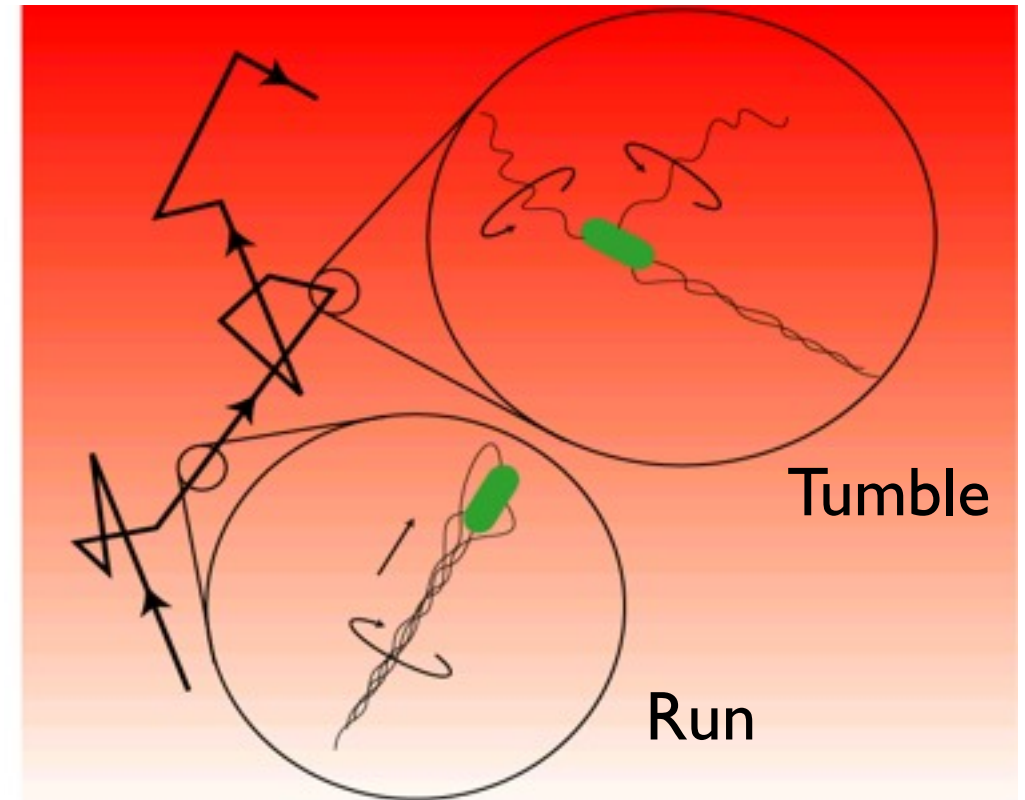
I/T time constant for synchronization

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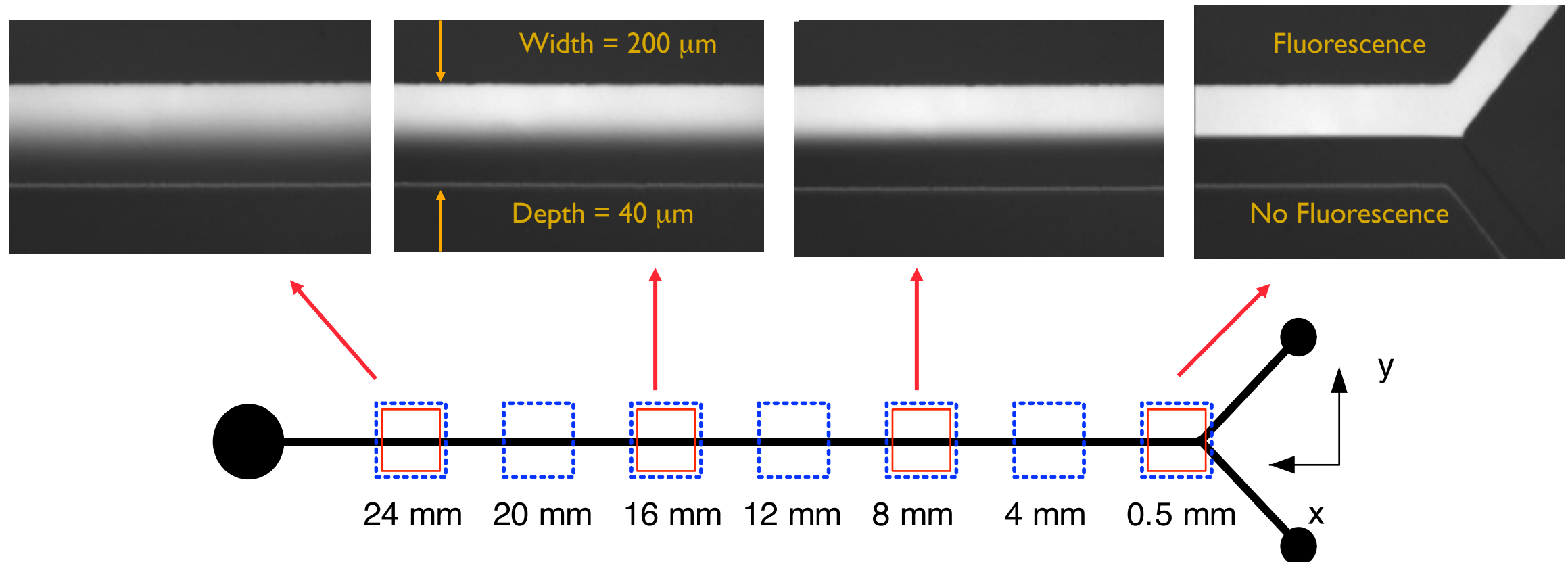
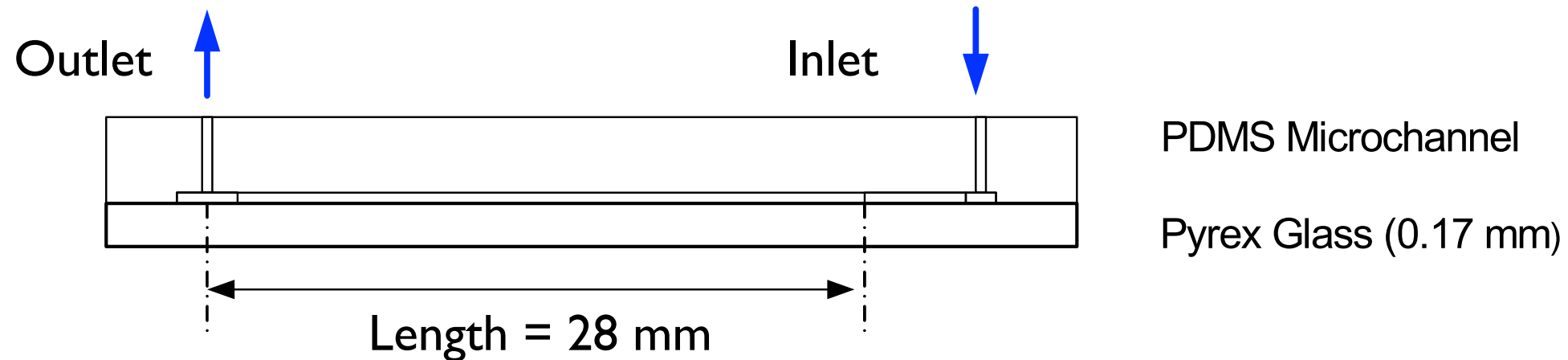
Motile bacteria as chaotic mixers

- Chaotic advection (Aref 1984)
- Viscous laminar flow
- Random switching of “blinking vortices”
- Bacteria have flagella that alternate rotation direction randomly
- Cell bodies execute random walk
 - Chaotic mixing in zero Re flow?

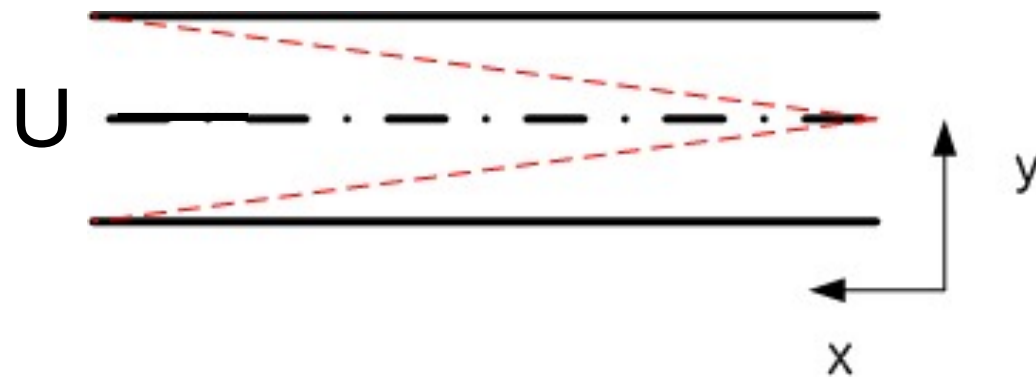


(Images courtesy of Károlyi György)

Mixing using freely swimming bacteria



Theory for standard diffusion.

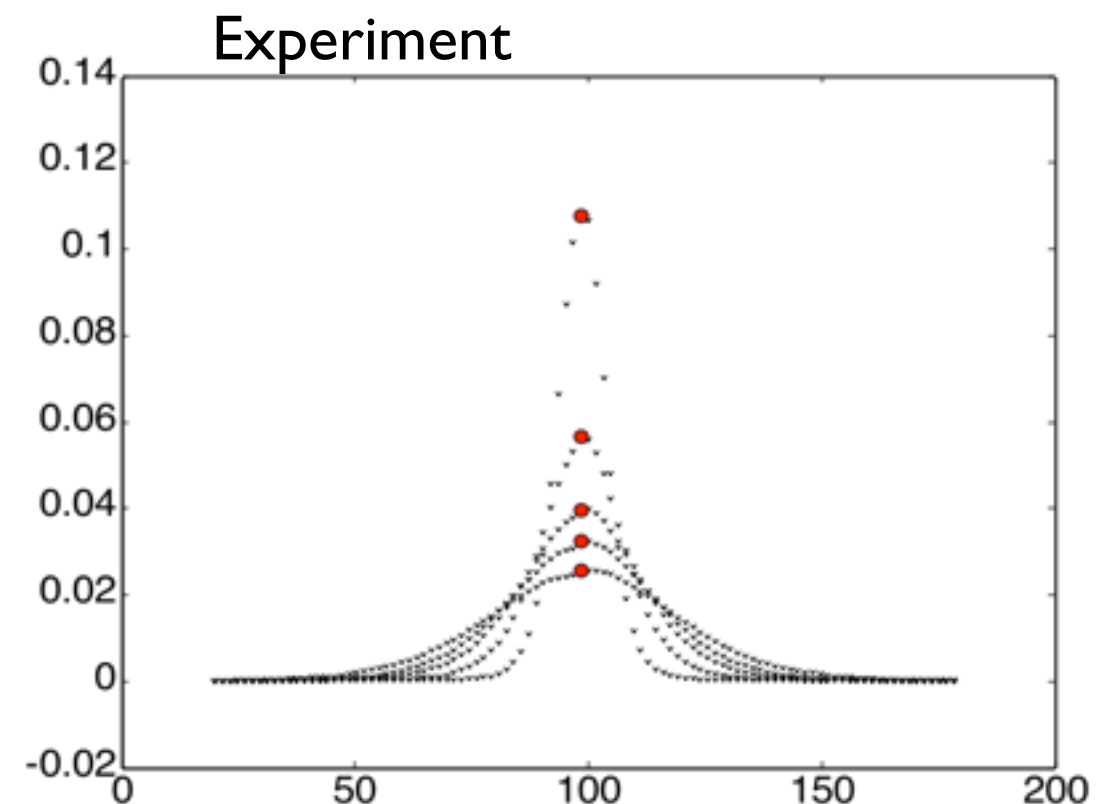
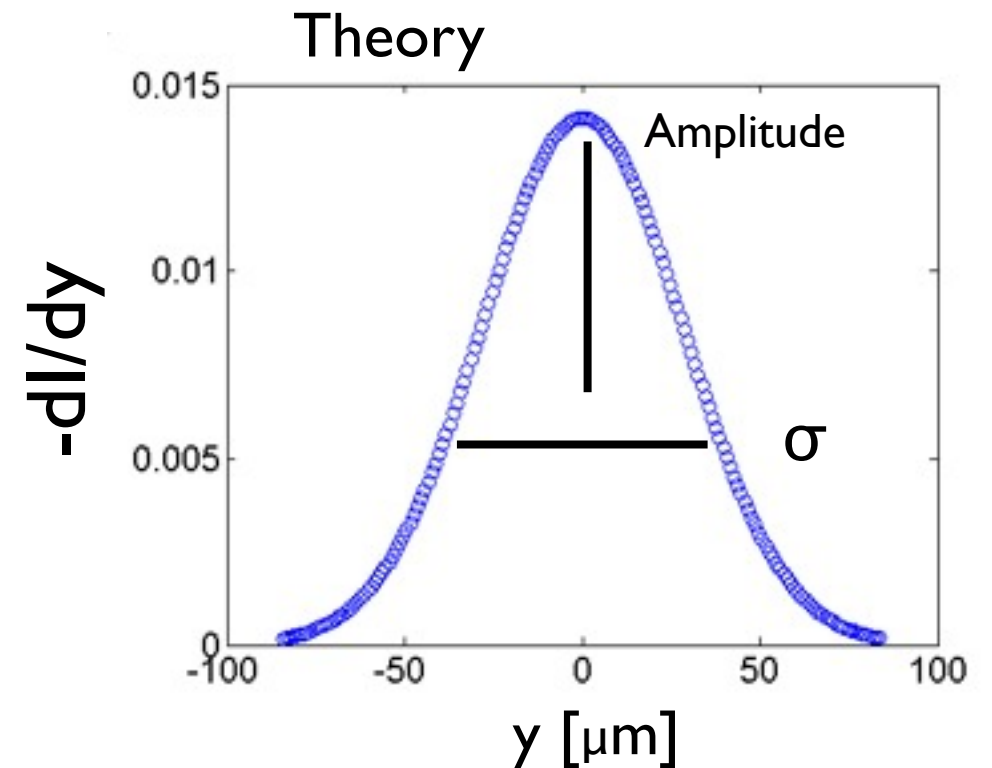


Diffusion Equation:
$$U \frac{\partial I}{\partial x} = D \frac{\partial^2 I}{\partial y^2}$$

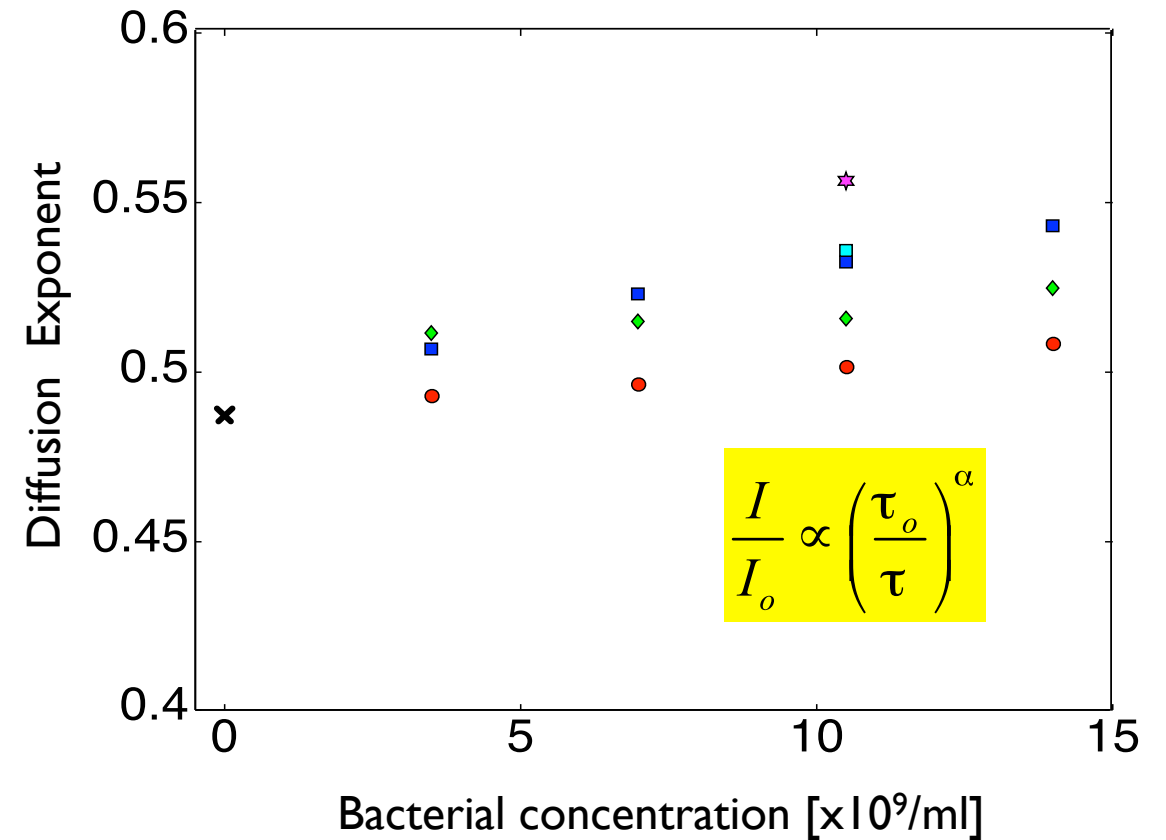
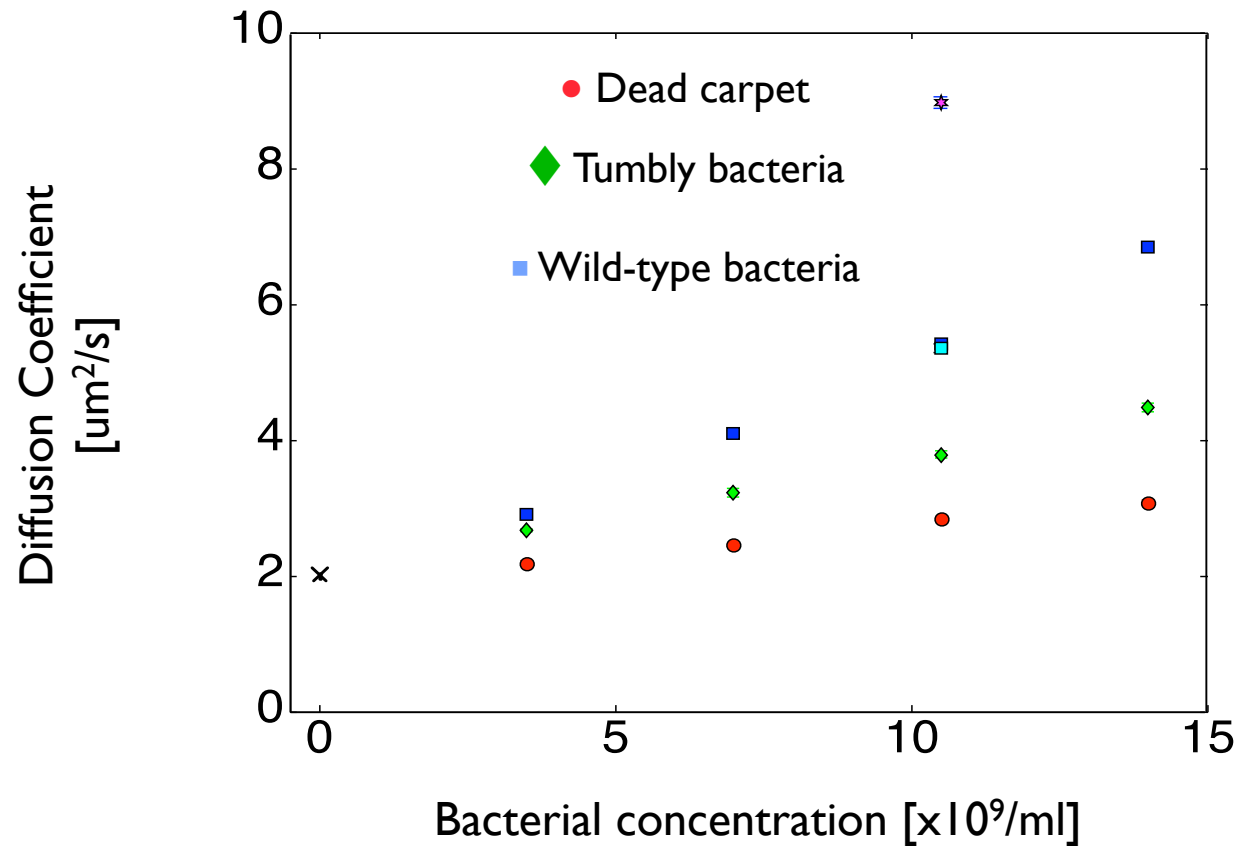
Intensity:
$$I(y, x) = \frac{1}{2} I_0 \operatorname{erfc} \frac{y}{2\sqrt{D\tau}}$$

Intensity Gradient:
$$-\frac{dI}{dy} = \frac{1}{2} \frac{I_0}{\sqrt{\pi D\tau}} \exp^{-\left(\frac{y}{2\sqrt{D\tau}}\right)^2}$$

Similarity Variable:
$$\tau = \frac{x}{U}$$



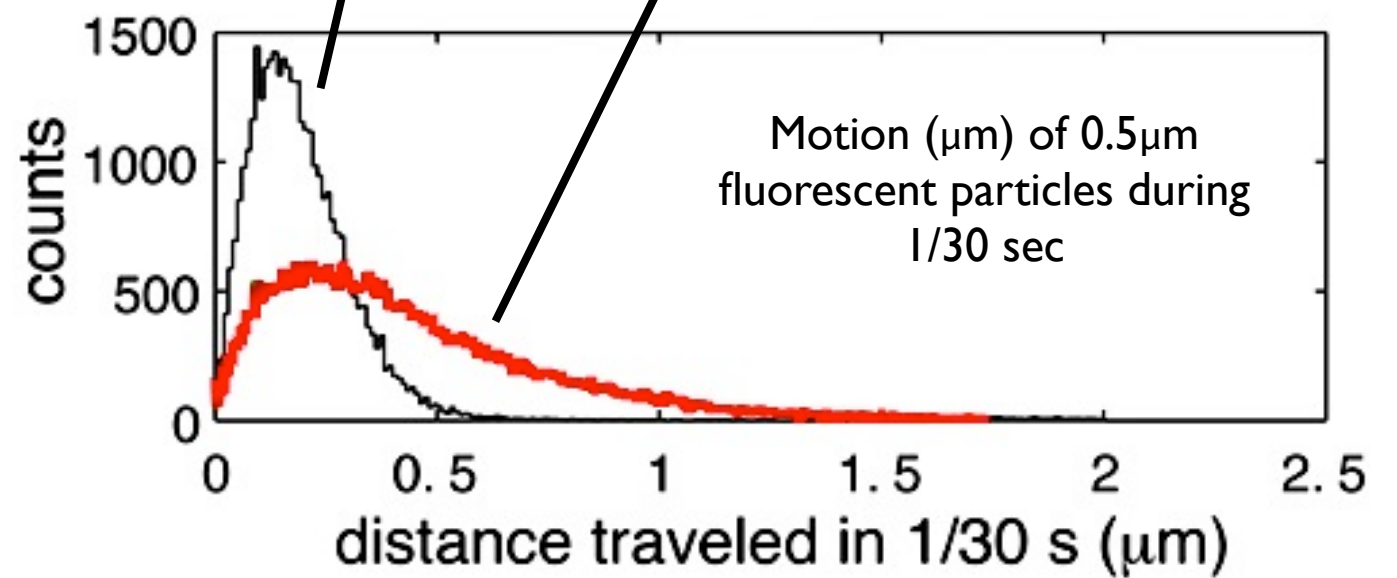
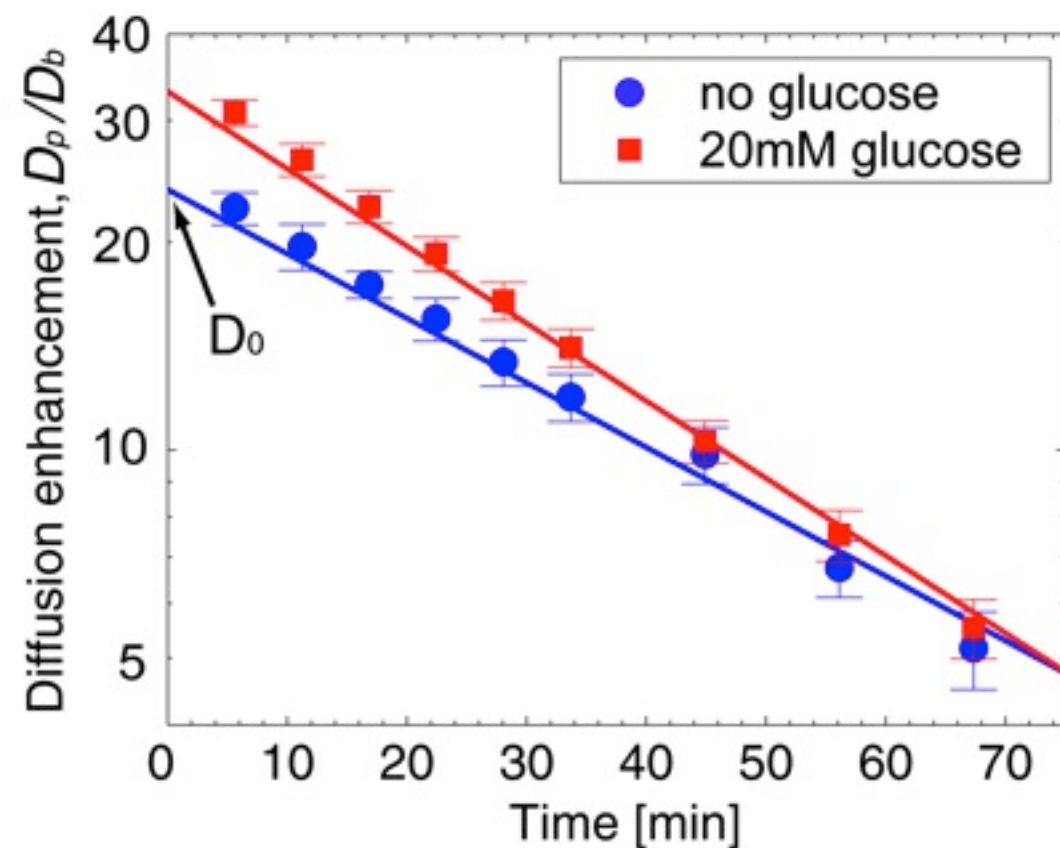
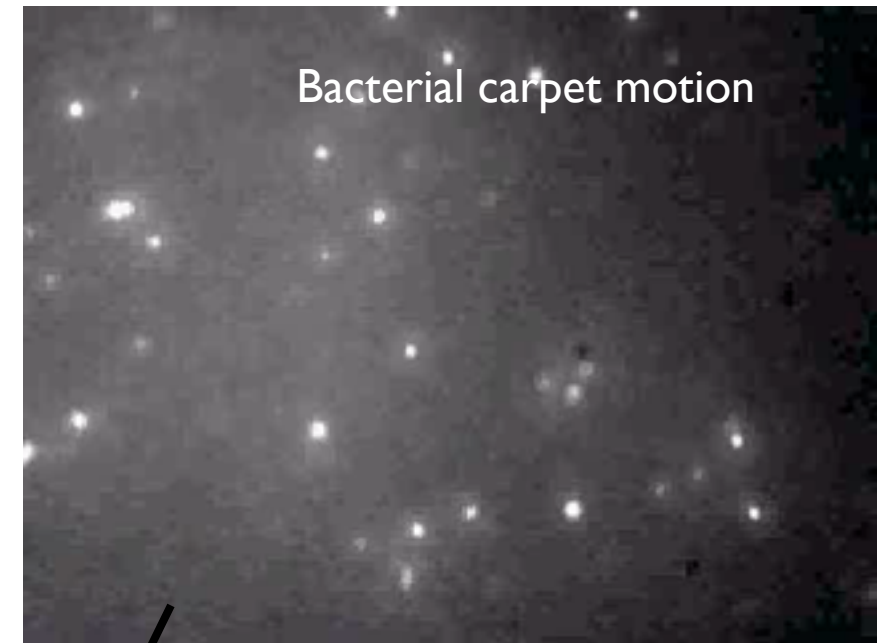
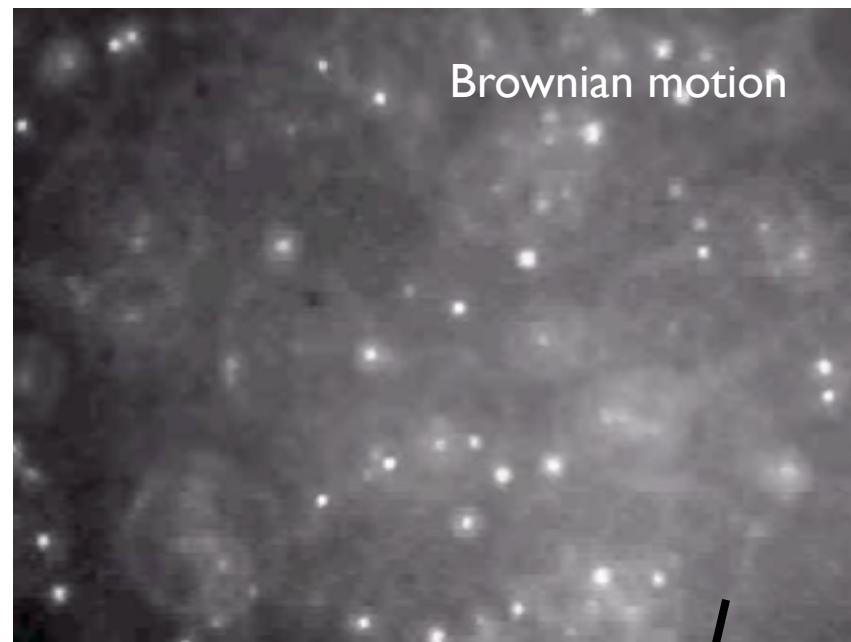
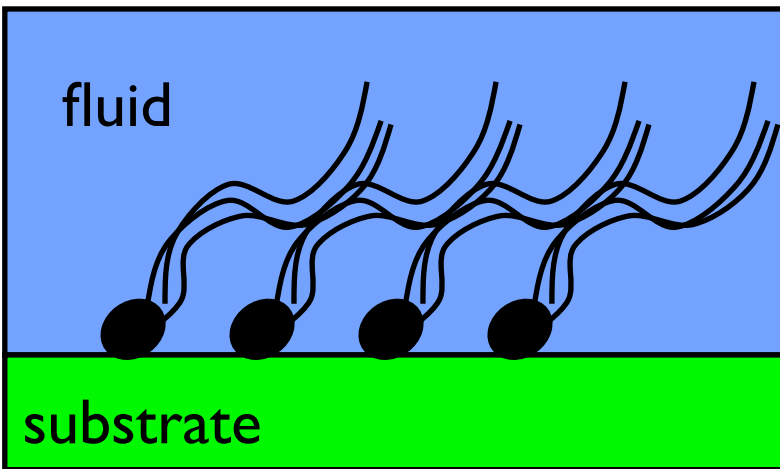
Enhanced diffusion due to bacteria



- Diffusion of small molecule rises from 20 – 80 $\mu\text{m}^2/\text{s}$
 - Large molecules and 1 μm beads show 50x enhancement
- Diffusion faster than standard “Fickian” diffusion
 - Introduction of new time scale

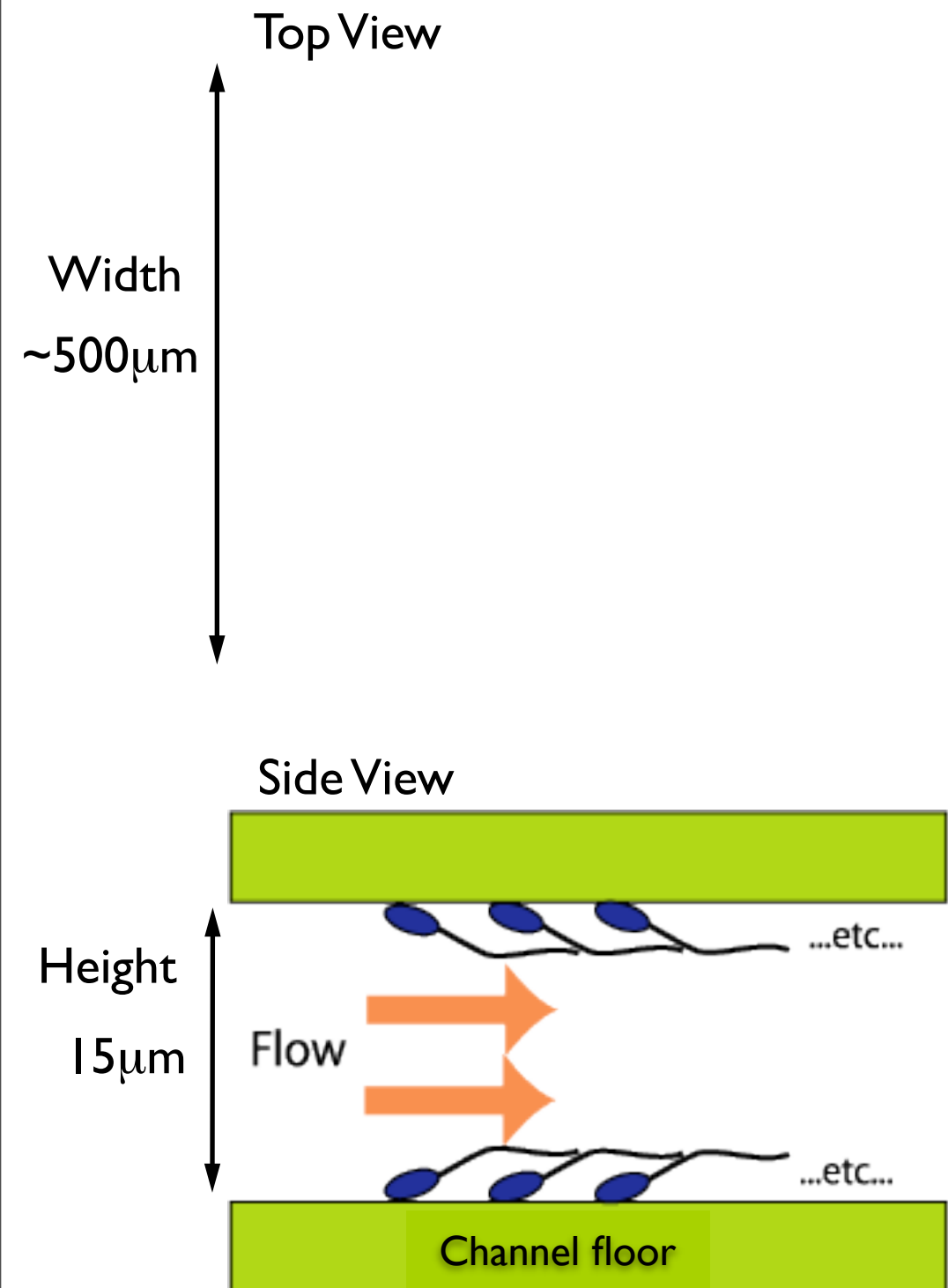
(Kim & Breuer, *Phys Fluids* 2004)

Bacterial carpets and enhanced mixing

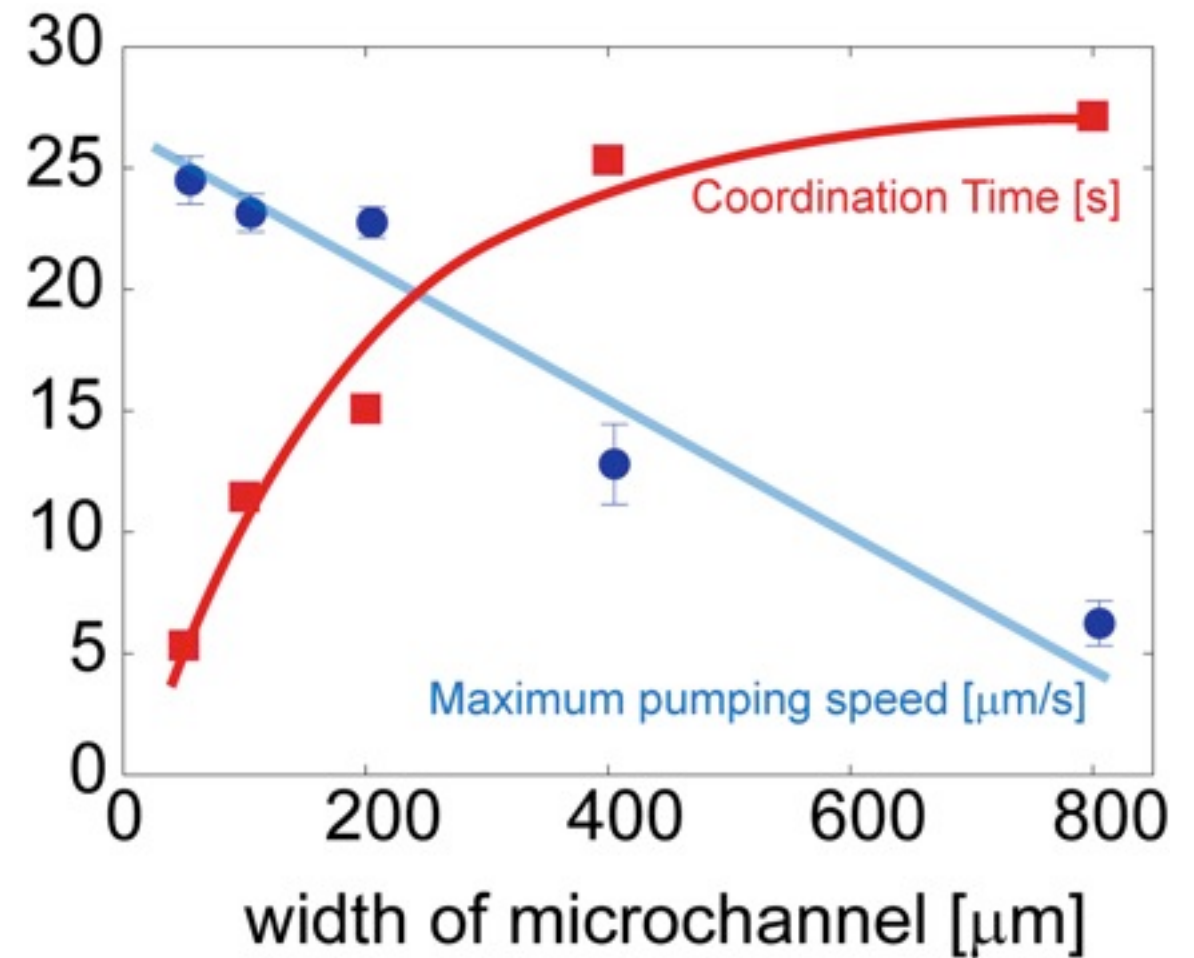


(Darnton, Turner, Breuer & Berg *Biophys J.* 2003)

Bacterial pumps

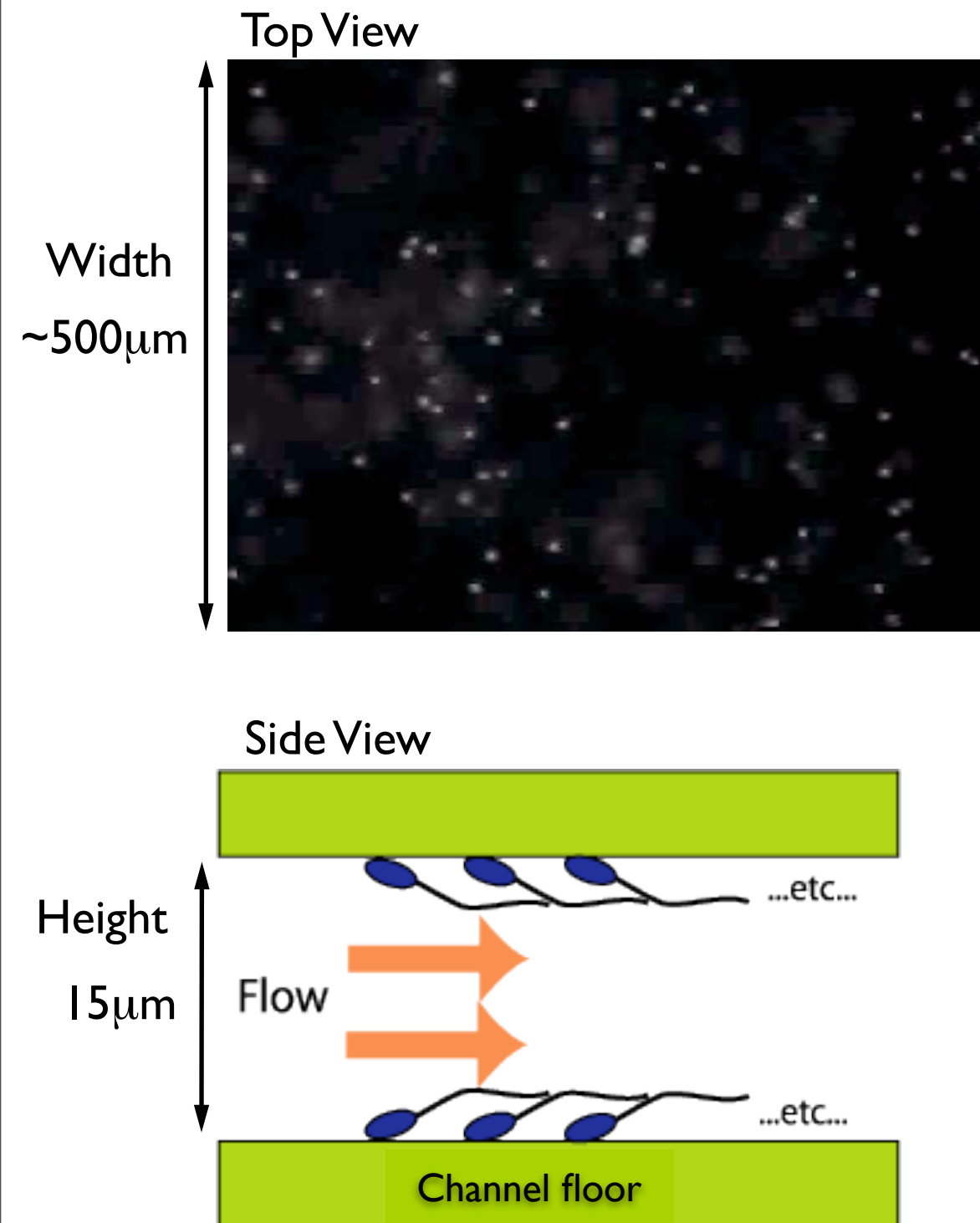


(Kim & Breuer, Small 2008)

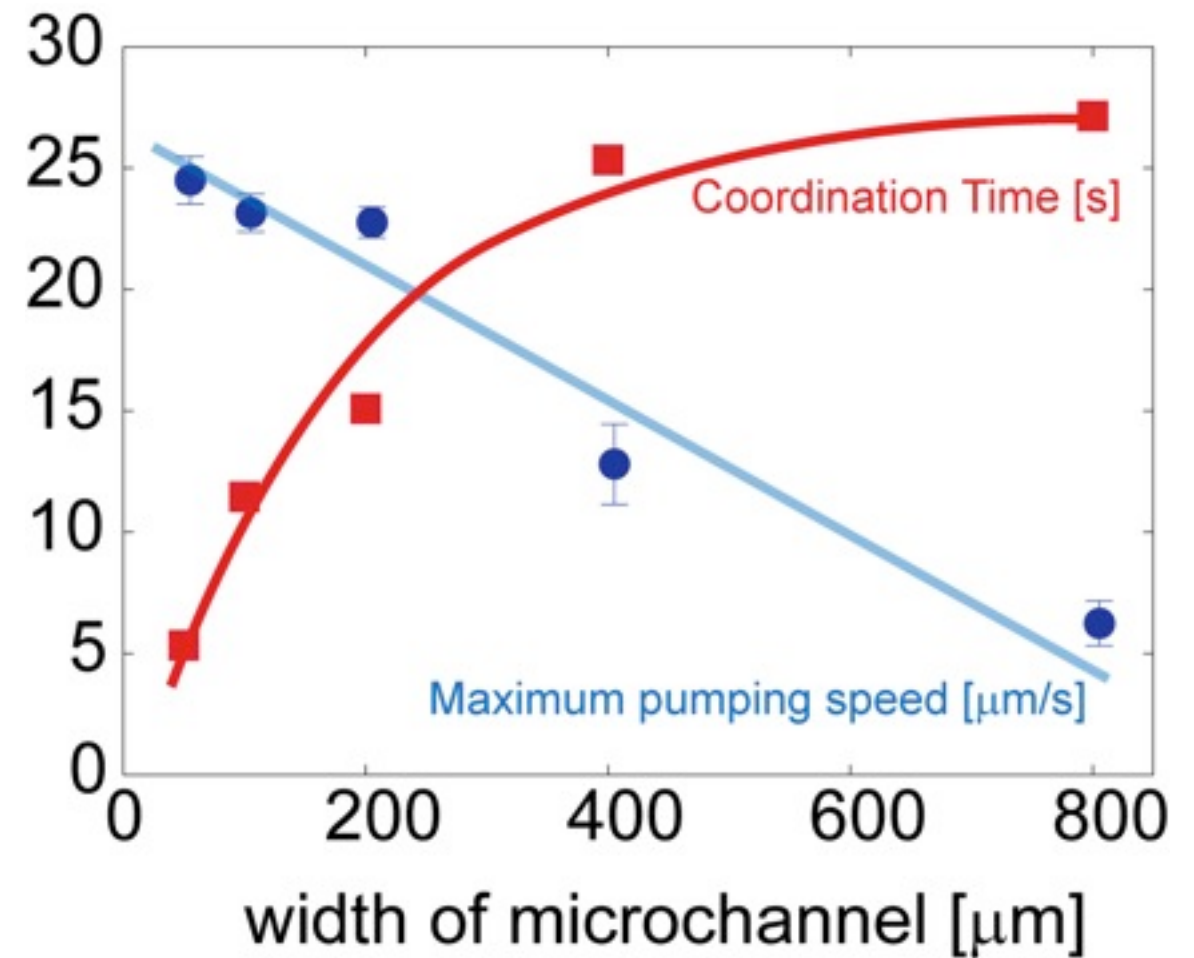


- Pumping arises spontaneously
- Direction determined by “last flush”
- “Crystallization” process

Bacterial pumps



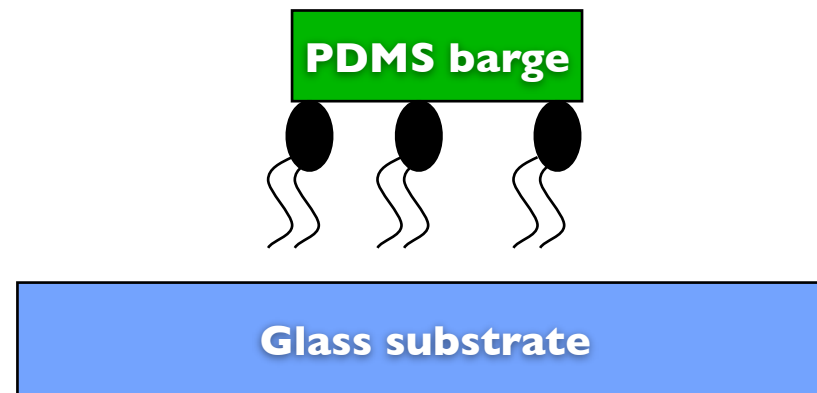
(Kim & Breuer, Small 2008)



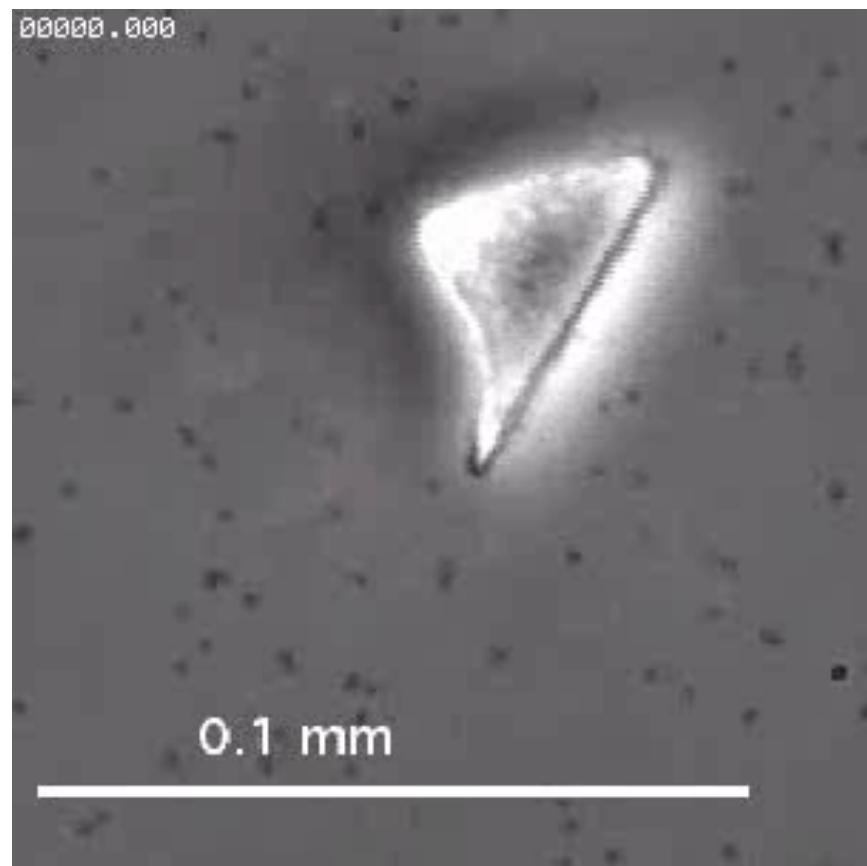
- Pumping arises spontaneously
- Direction determined by “last flush”
- “Crystallization” process

Bacterial barges

- Rotation 33 deg/s
- Surface area 1400 μm^2
- Long axis 65 μm
- Bacteria 300

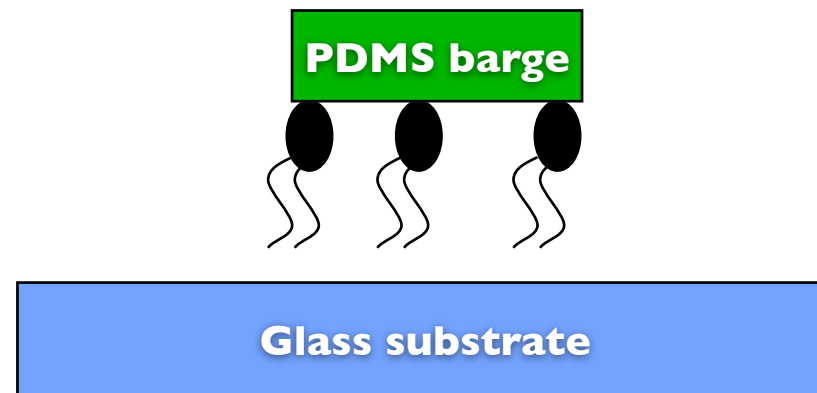


- Translates 5 $\mu\text{m/s}$
- Rotation 4.6 deg/s
- Surface area 4350 μm^2
- Long axis 92 μm
- Bacteria 1100

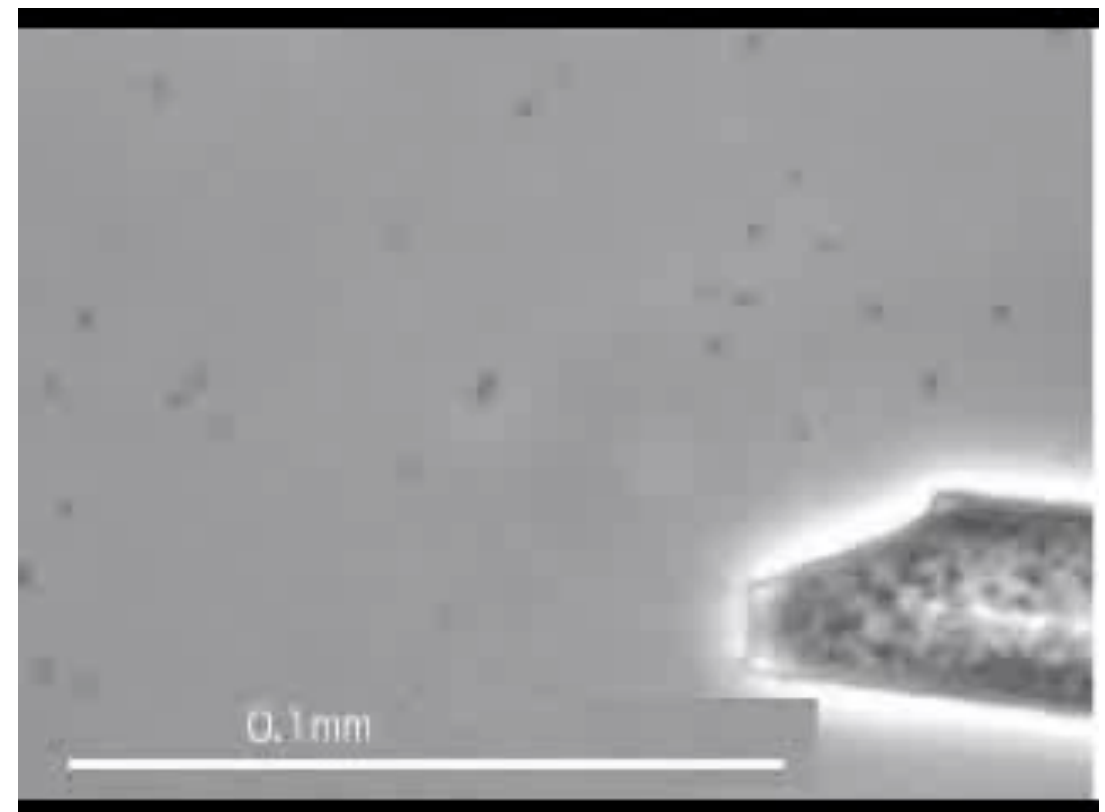
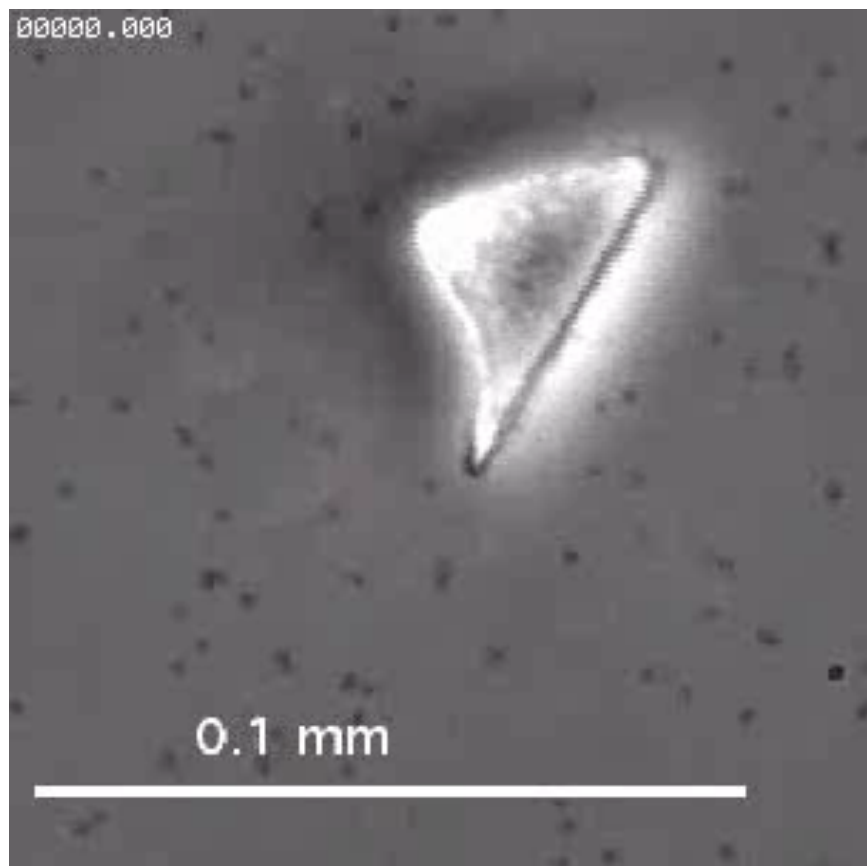


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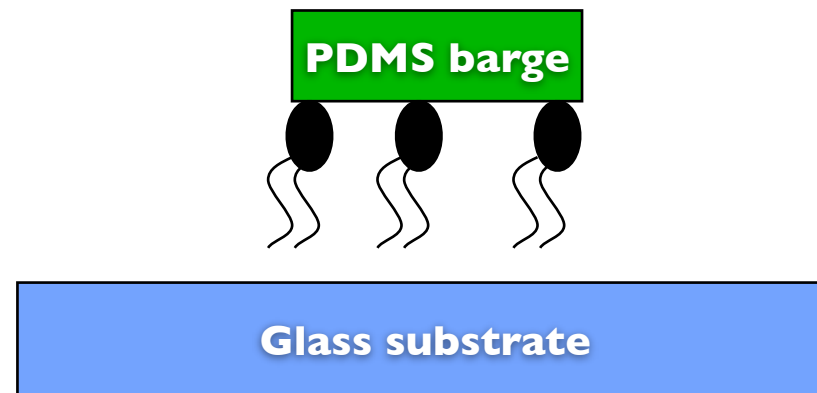


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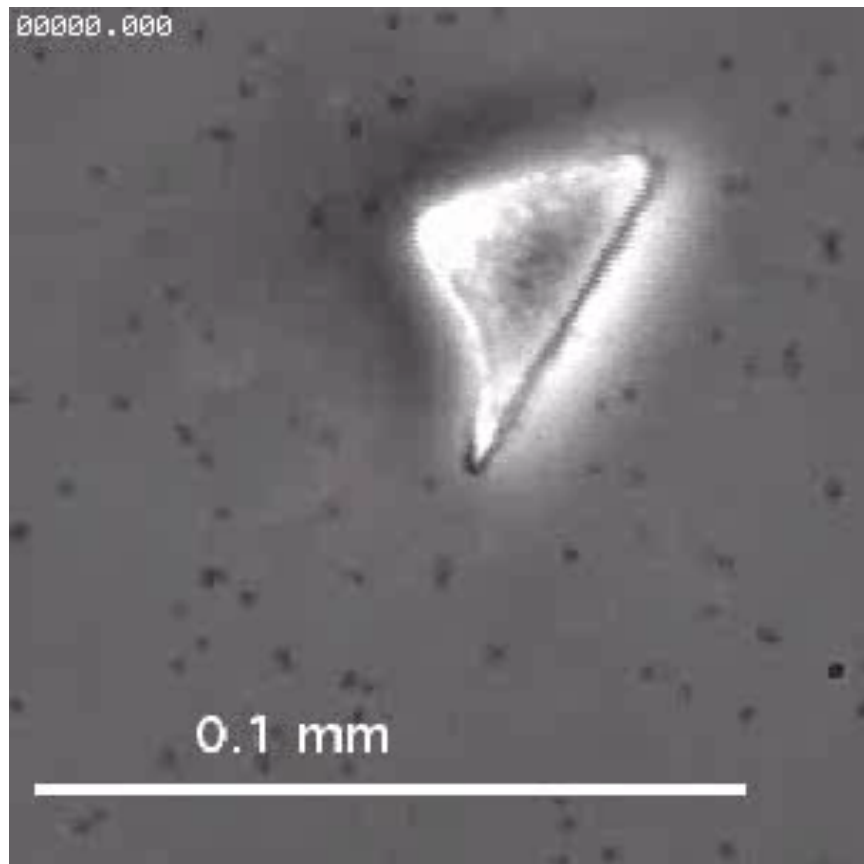


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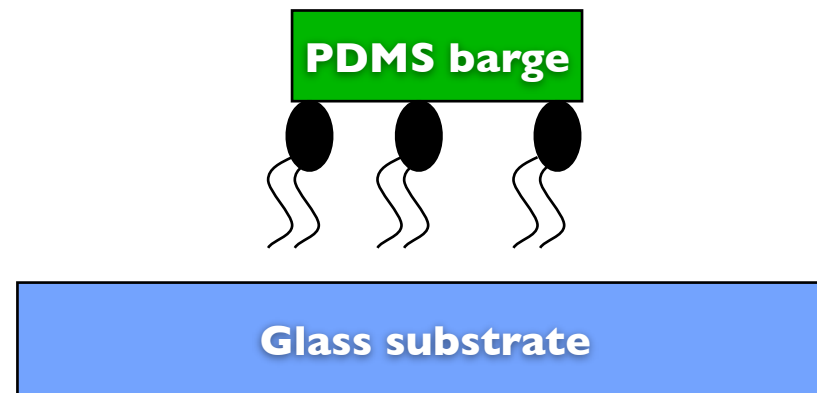


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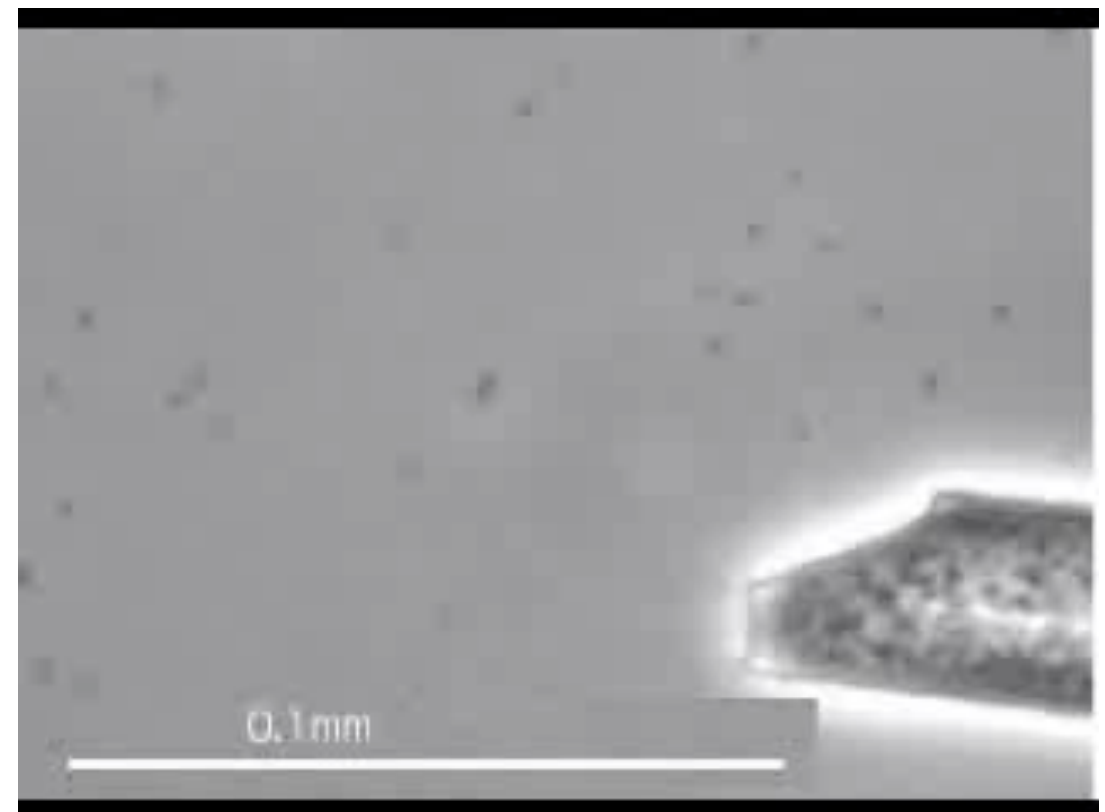


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- Bacteria 1100



Summary:

- Force-free swimming speed measured:
 - newtonian fluids
 - viscoelastic fluids underway
- Fluids-filament interactions:
 - coiling of an elastic filament due to viscous stresses
- Synchronization of adjacent filaments
 - hydrodynamic interactions
 - elastic compliance necessary
 - multiple paddles illustrate significantly more complex (& chaotic) behavior



- Acknowledgements: Tom Powers, Bin Liu, Min Jun Kim, Qian Bian, Dave Gagnon
- Support: NSF