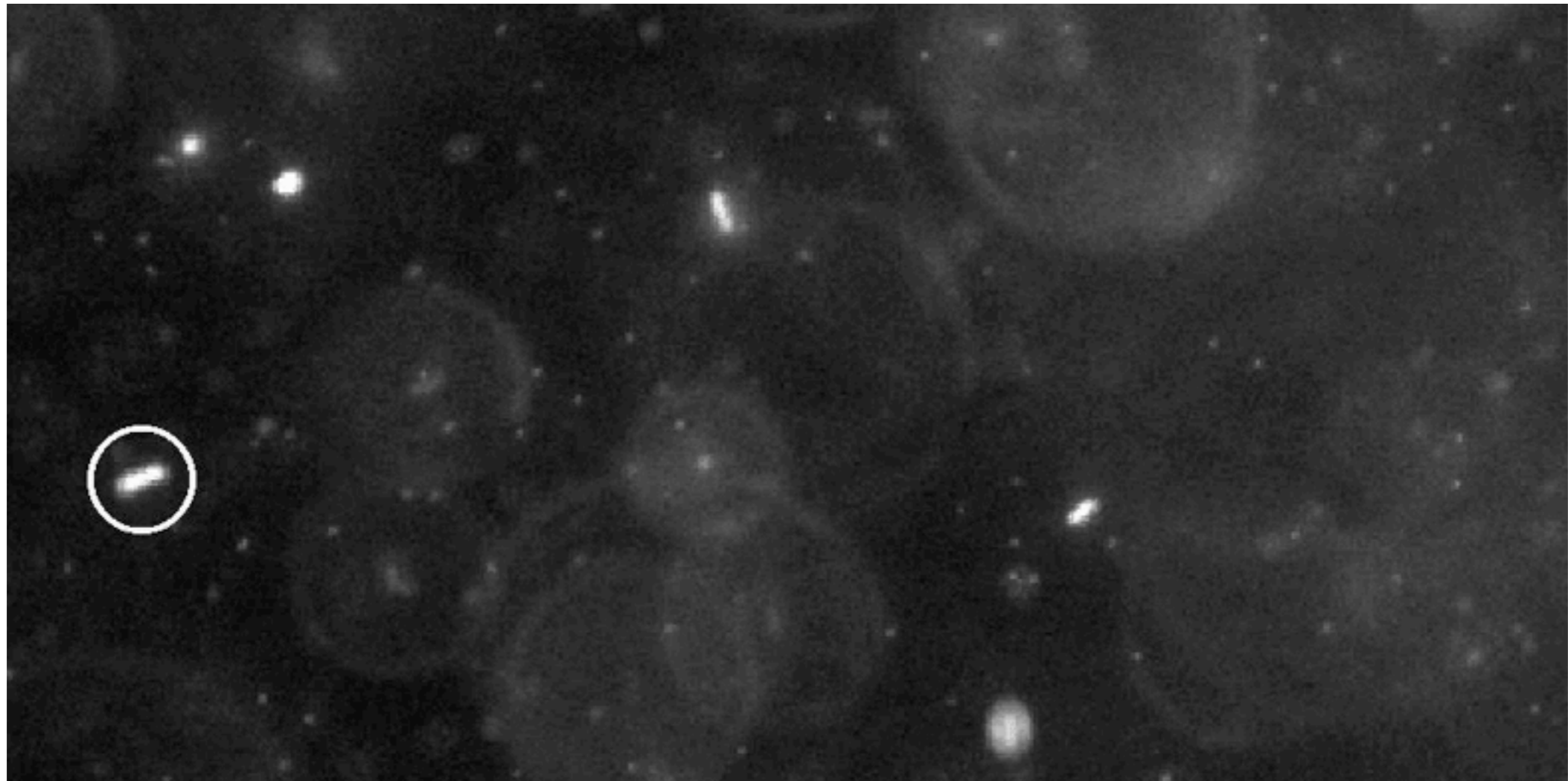


# Dilute bacterial suspensions

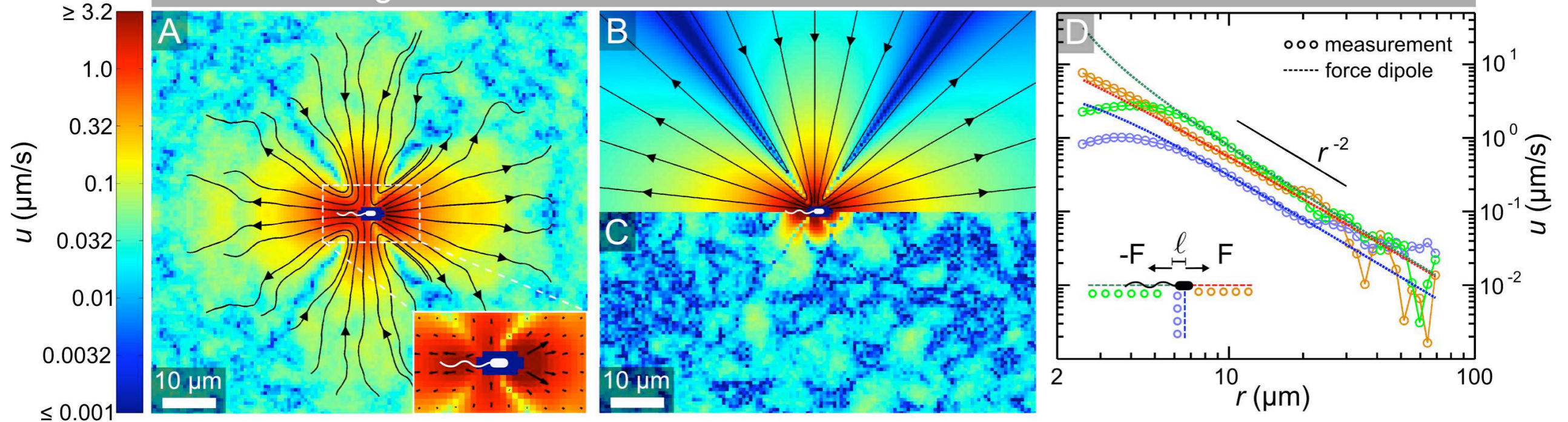
18.S995 - L06 & 07

# *E.coli* (non-tumbling HCB 437)



# *E. coli* (non-tumbling HCB 437)

Free swimming



$$\mathbf{u}(\mathbf{r}) = \frac{A}{|\mathbf{r}|^2} \left[ 3(\hat{\mathbf{r}} \cdot \hat{\mathbf{d}})^2 - 1 \right] \hat{\mathbf{r}}, \quad A = \frac{\ell F}{8\pi\eta}, \quad \hat{\mathbf{r}} = \frac{\mathbf{r}}{|\mathbf{r}|}$$

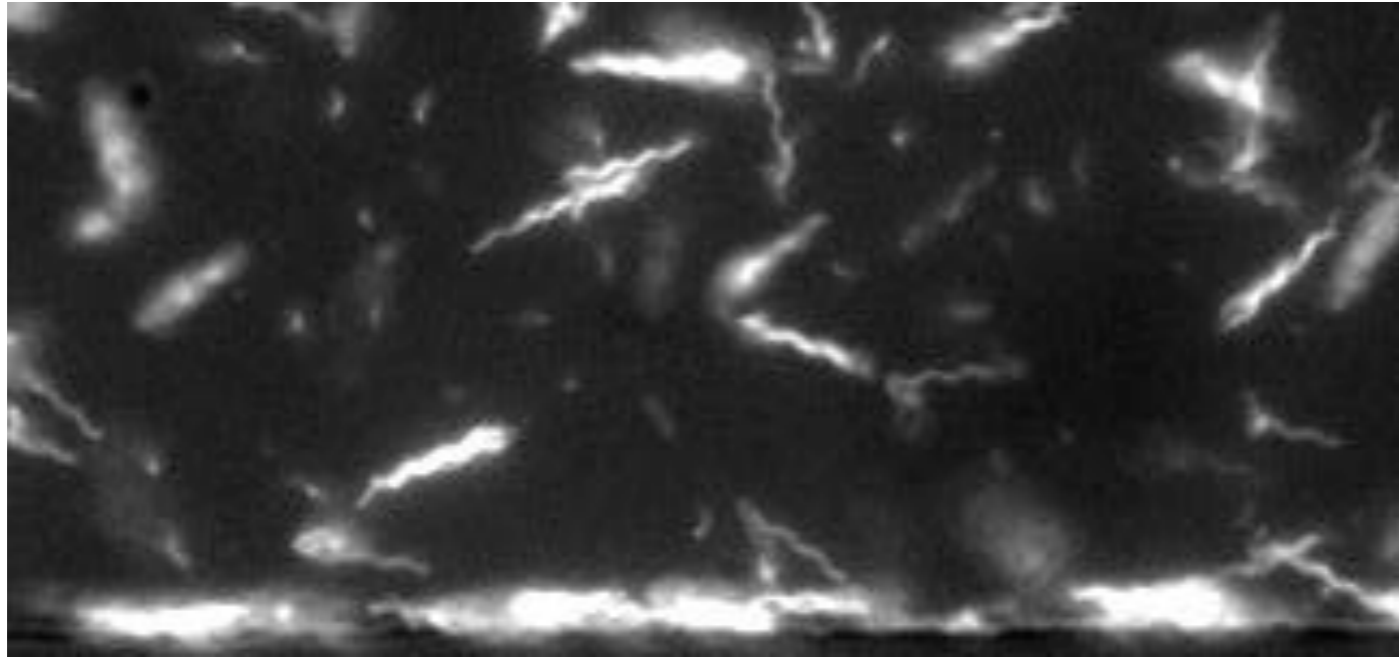
$$V_0 = 22 \pm 5 \mu\text{m/s}$$

$$\ell = 1.9 \mu\text{m}$$

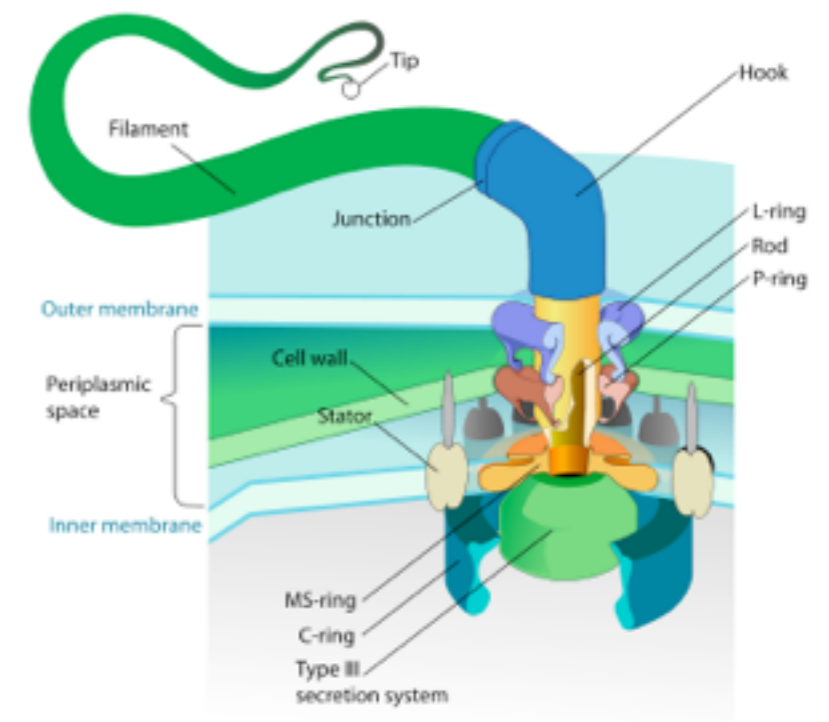
$$F = 0.42 \text{ pN}$$

# Bacterial run & tumble motion

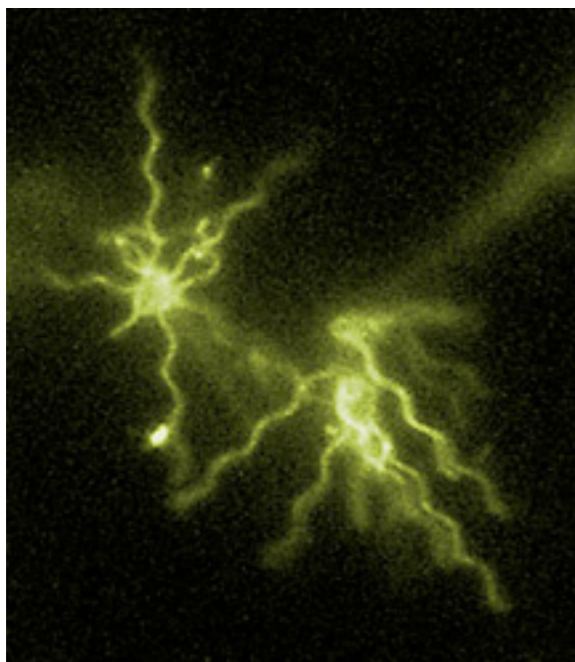
movie: V. Kantsler



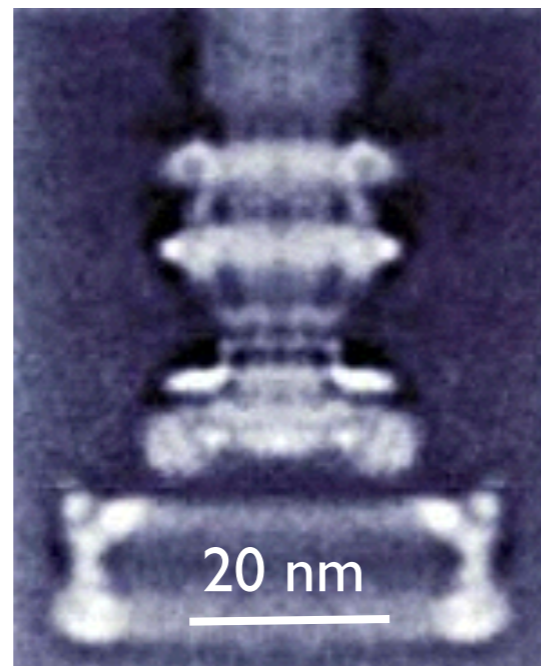
~20 parts



source: wiki



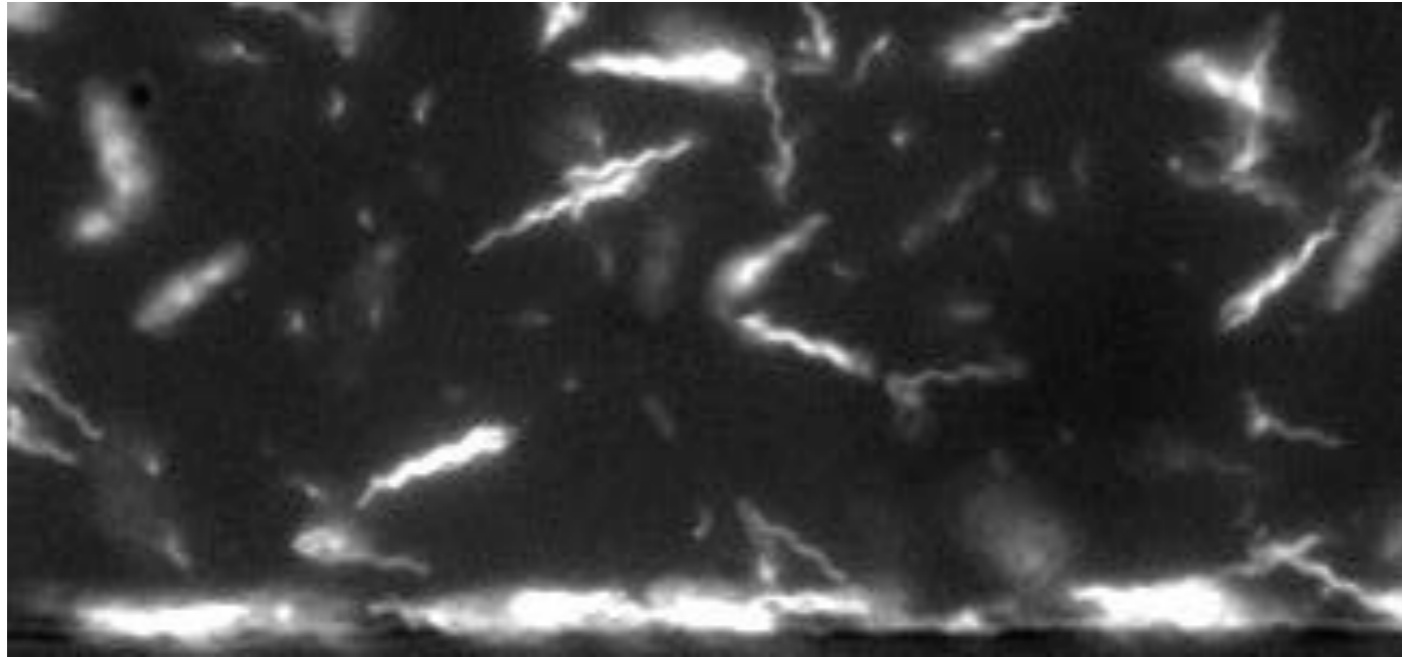
Berg (1999) Physics Today



Chen et al (2011) EMBO Journal

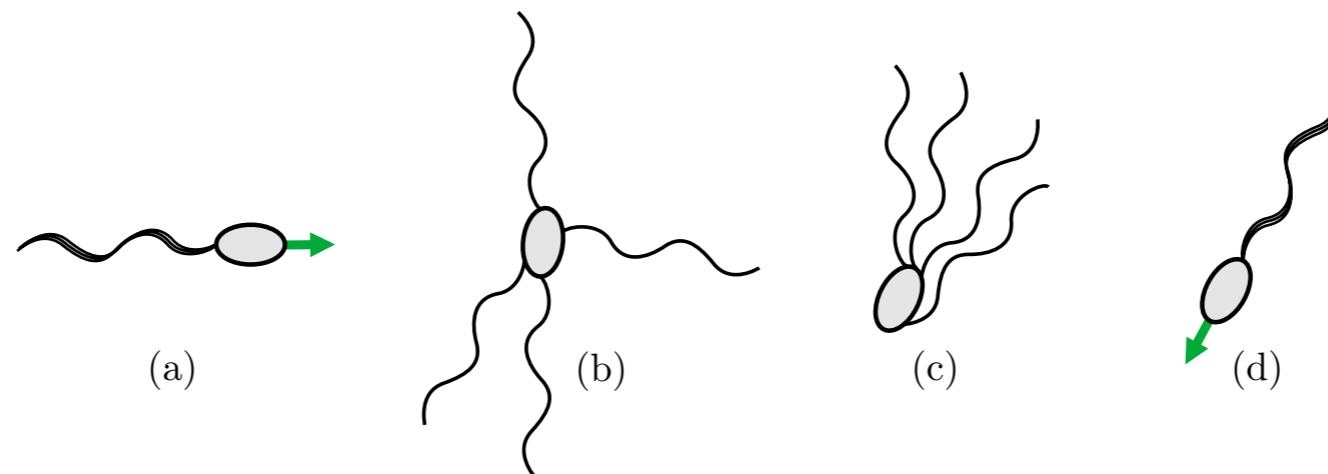
# Bacterial run & tumble motion

movie: V. Kantsler



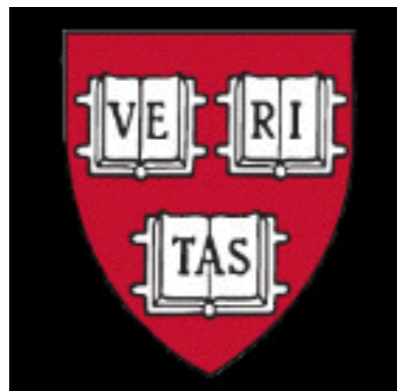
Rep. Prog. Phys. **72** (2009) 096601

E Lauga and T R Powers



**Figure 15.** Bundling of bacterial flagella. During swimming, the bacterial flagella are gathered in a tight bundle behind the cell as it moves through the fluid ((*a*) and (*d*)). During a tumbling event, the flagella come out the bundle (*b*), resulting in a random reorientation of the cell before the next swimming event. At the conclusion of the tumbling event, hydrodynamic interactions lead to the relative attraction of the flagella (*c*), and their synchronization to form a perfect bundle (*d*).

for more movies, see also

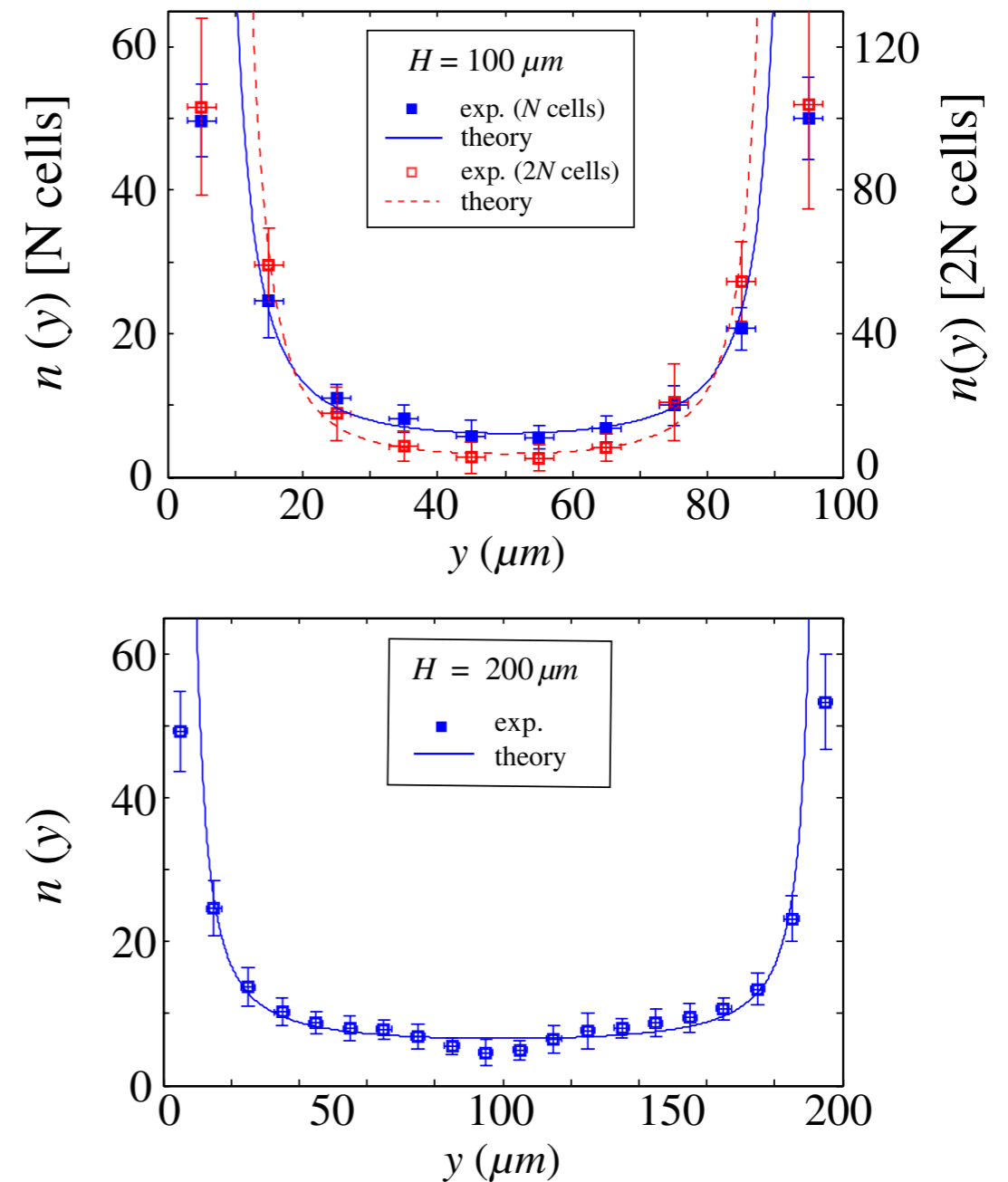
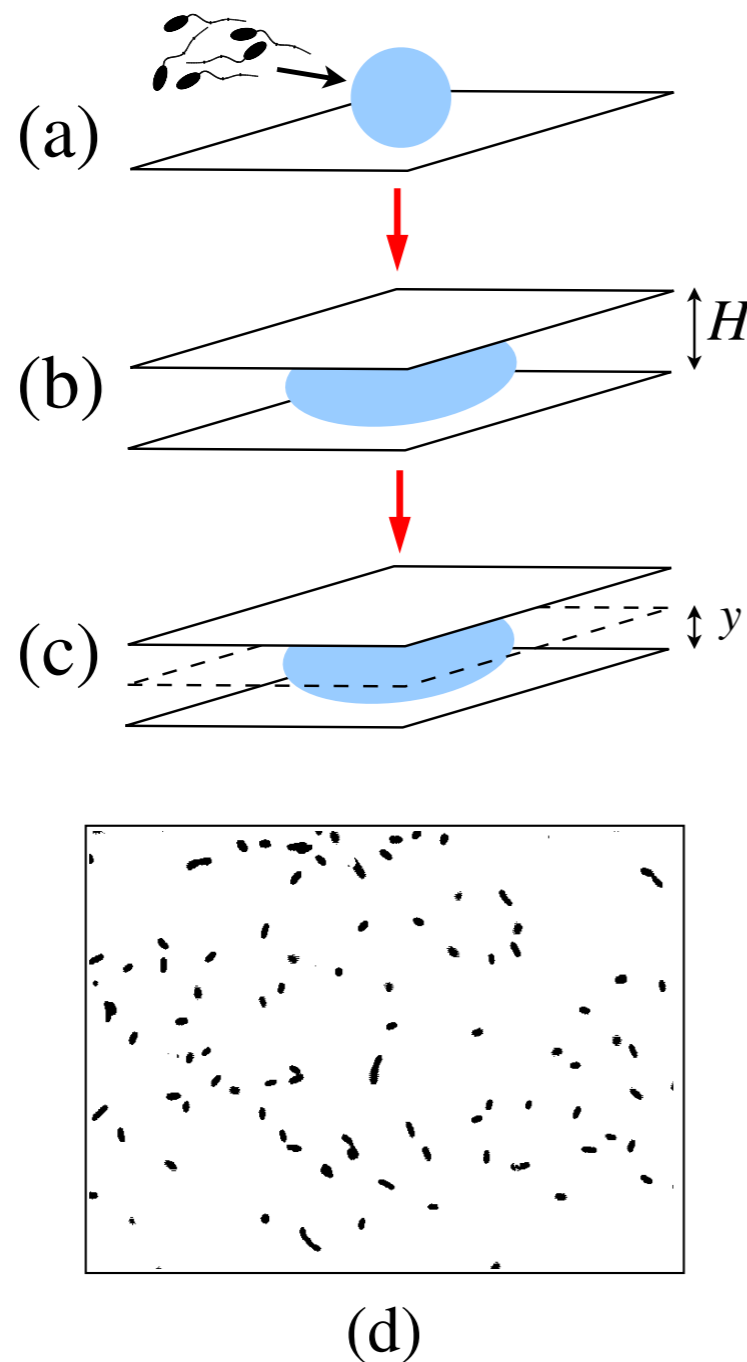


Howard C. Berg, PI  
Bacterial motility and behavior

<http://www.rowland.harvard.edu/labs/bacteria/movies/index.php>

# Hydrodynamic Attraction of Swimming Microorganisms by Surfaces

Allison P. Berke,<sup>1</sup> Linda Turner,<sup>2</sup> Howard C. Berg,<sup>2,3</sup> and Eric Lauga<sup>4,\*</sup>



# Goals

- minimal SDE model for microbial swimming
- wall accumulation & density profile

## 1.3 Dilute microbial suspensions

A minimalist model for the locomotion of an isolated microorganism (e.g., alga or bacterium) with position  $\mathbf{X}(t)$  and orientation unit vector  $\mathbf{N}(t)$  is given by the coupled system of Ito SDEs

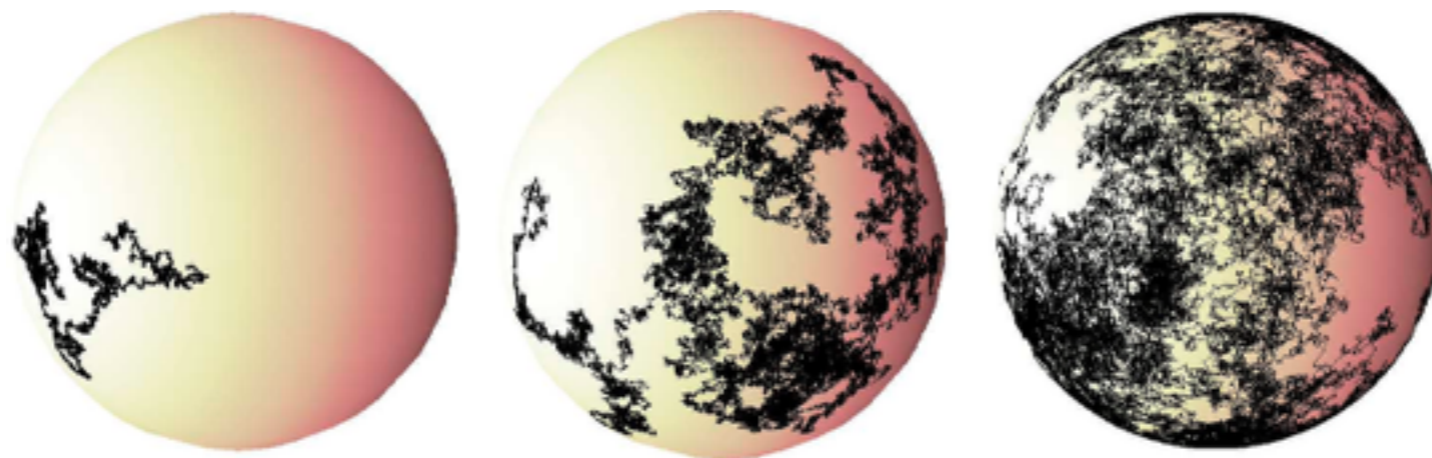
$$d\mathbf{X} = V\mathbf{N}dt + \sqrt{2D_T} * d\mathbf{B}(t), \quad (1.45a)$$

$$d\mathbf{N} = (1 - d)D_R\mathbf{N}dt + \sqrt{2D_R}(\mathbf{I} - \mathbf{N}\mathbf{N}) * d\mathbf{W}(t). \quad (1.45b)$$

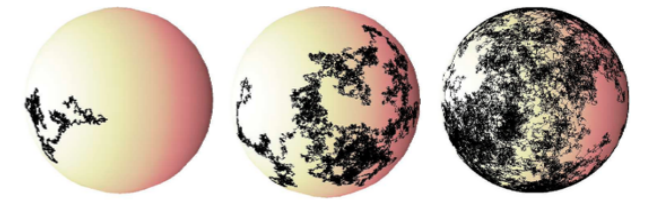
To confirm that Eq. (1.45b) conserves the unit length of the orientation vector,  $|\mathbf{N}|^2 = 1$  for all  $t$ , it is convenient to rewrite Eqs. (1.45) in component form:

$$dX_i = VN_idt + \sqrt{2D_T} * dB_i(t), \quad (1.46a)$$

$$dN_j = (1 - d)D_R N_jdt + \sqrt{2D_R}(\delta_{jk} - N_jN_k) * dW_k(t). \quad (1.46b)$$



# BM on the unit sphere



$$dN_j = (1 - d)D_R N_j dt + \sqrt{2D_R} (\delta_{jk} - N_j N_k) * dW_k(t).$$

For the constraint  $|\mathbf{N}|^2 = 1$  to be satisfied, we must have  $d|\mathbf{N}|^2 = 0$ . Applying the  $d$ -dimensional version of Ito's formula, see Eq. (A.12), to  $F(\mathbf{N}) = |\mathbf{N}|^2$ , one finds indeed that

$$\begin{aligned} d|\mathbf{N}|^2 &= 2N_j * dN_j + (\partial_{N_i} \partial_{N_j} N_k N_k) D_R (\delta_{ij} - N_i N_j) dt \\ &= 2N_j * \left[ (1 - d)D_R N_j dt + \sqrt{2D_R} (\delta_{jk} - N_j N_k) * dW_k(t) \right] + \\ &\quad \partial_{N_i} (\delta_{jk} N_k + N_k \delta_{jk}) D_R (\delta_{ij} - N_i N_j) dt \\ &= 2(1 - d)D_R dt + \\ &\quad (\delta_{jk} \delta_{ik} + \delta_{ik} \delta_{jk}) D_R (\delta_{ij} - N_i N_j) dt \\ &= 0. \end{aligned} \tag{1.47}$$

# Orientation correlations

To understand the dynamics (1.46), it is useful to compute the orientation correlation,

$$\langle \mathbf{N}(t) \cdot \mathbf{N}(0) \rangle = \mathbb{E}[\mathbf{N}(t) \cdot \mathbf{N}(0)] = \mathbb{E}[N_z(t)], \quad (1.48)$$

where we have assumed (w.l.o.g.) that  $\mathbf{N}(0) = \mathbf{e}_z$ . Averaging Eq. (1.46b), we find that

$$\frac{d}{dt} \mathbb{E}[N_z(t)] = (1 - d) D_R \mathbb{E}[N_z(t)], \quad (1.49)$$

implying that, in this model, the memory loss about the orientation is exponential

$$\langle \mathbf{N}(t) \cdot \mathbf{N}(0) \rangle = e^{(1-d)D_R t}, \quad (1.50)$$

# Mean square displacement

$$\begin{aligned}
 d|\mathbf{X}|^2 &= 2X_j * dX_j + (\partial_{X_i} \partial_{X_j} X_k X_k) D_T \delta_{ij} dt \\
 &= 2X_j * dX_j + (\delta_{jk} \delta_{ik} + \delta_{ik} \delta_{jk}) D_T \delta_{ij} dt \\
 &= 2X_j * dX_j + 2d D_T dt \\
 &= 2X_j [V N_j dt + \sqrt{2D_T} * dB_j(t)] + 2d D_T dt,
 \end{aligned} \tag{1.51}$$

averaging and dividing by  $dt$ , gives

$$\frac{d}{dt} \mathbb{E}[\mathbf{X}^2] = 2V \mathbb{E}[\mathbf{X}(t) \mathbf{N}(t)] + 2d D_T. \tag{1.52}$$

The expectation value on the rhs. can be evaluated by making use of Eq. (1.50):

$$\begin{aligned}
 \mathbb{E}[\mathbf{X}(t) \cdot \mathbf{N}(t)] &= \mathbb{E} \left[ \int_0^t d\mathbf{X}(s) \cdot \mathbf{N}(t) \right] \\
 &= V \mathbb{E} \left[ \int_0^t ds \mathbf{N}(s) \cdot \mathbf{N}(t) \right] \\
 &= V \int_0^t ds \langle \mathbf{N}(t) \cdot \mathbf{N}(s) \rangle \\
 &= V \int_0^t ds e^{(1-d)D_R(t-s)} \\
 &= \frac{V}{(d-1)D_R} [1 - e^{(1-d)D_R t}].
 \end{aligned}$$

# Mean square displacement

By inserting this expression into Eq. (1.52) and integrating over  $t$ , we find

$$\mathbb{E}[\mathbf{X}^2] = \frac{2V^2}{(d-1)^2 D_R^2} [(d-1)D_R t + e^{(1-d)D_R t} - 1] + 2dD_T t. \quad (1.53)$$

If  $D_T$  is small, then at short times  $t \ll D_R^{-1}$  the motion is ballistic

$$\mathbb{E}[\mathbf{X}^2] \simeq V^2 t^2 + 2dD_T t, \quad (1.54)$$

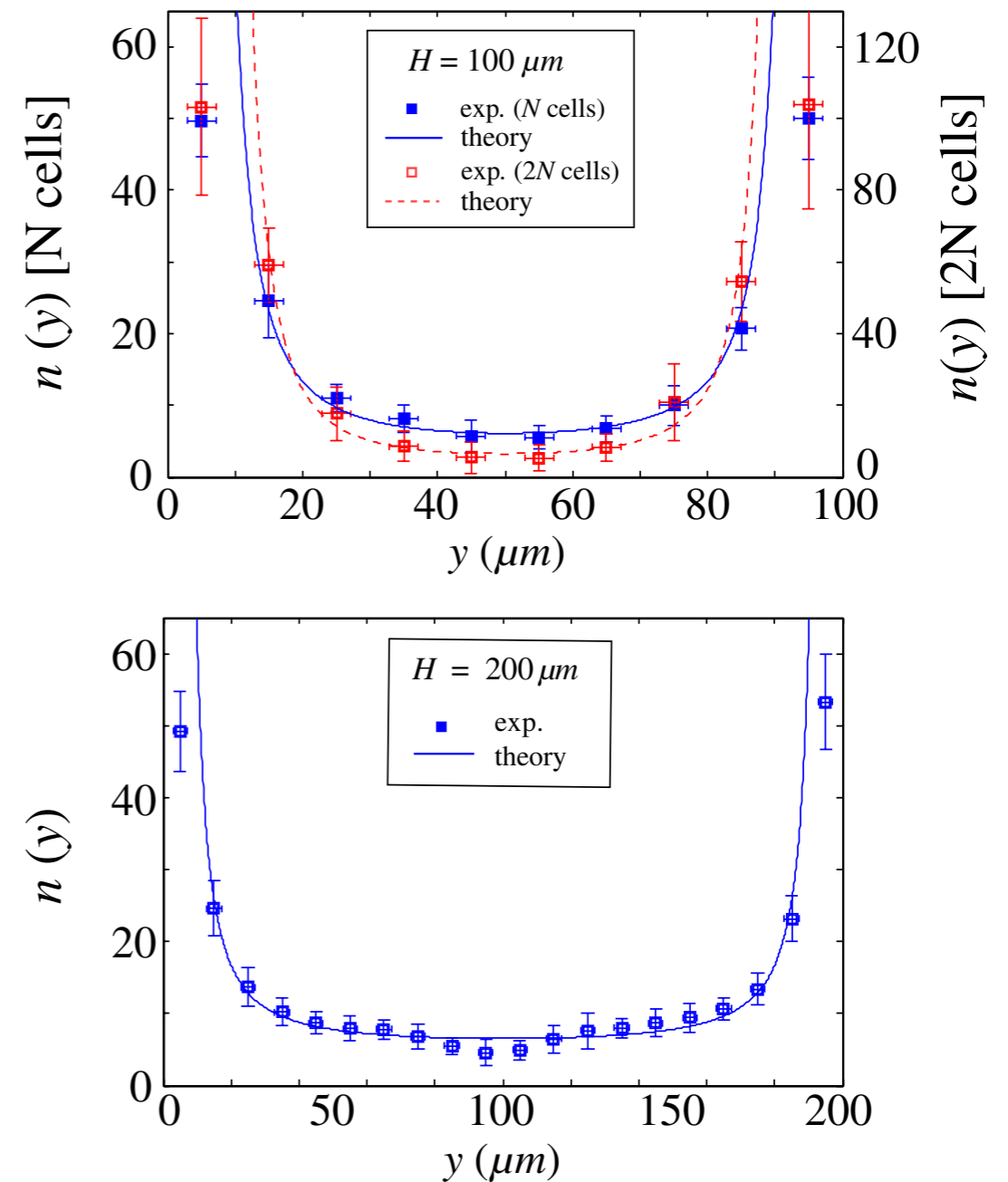
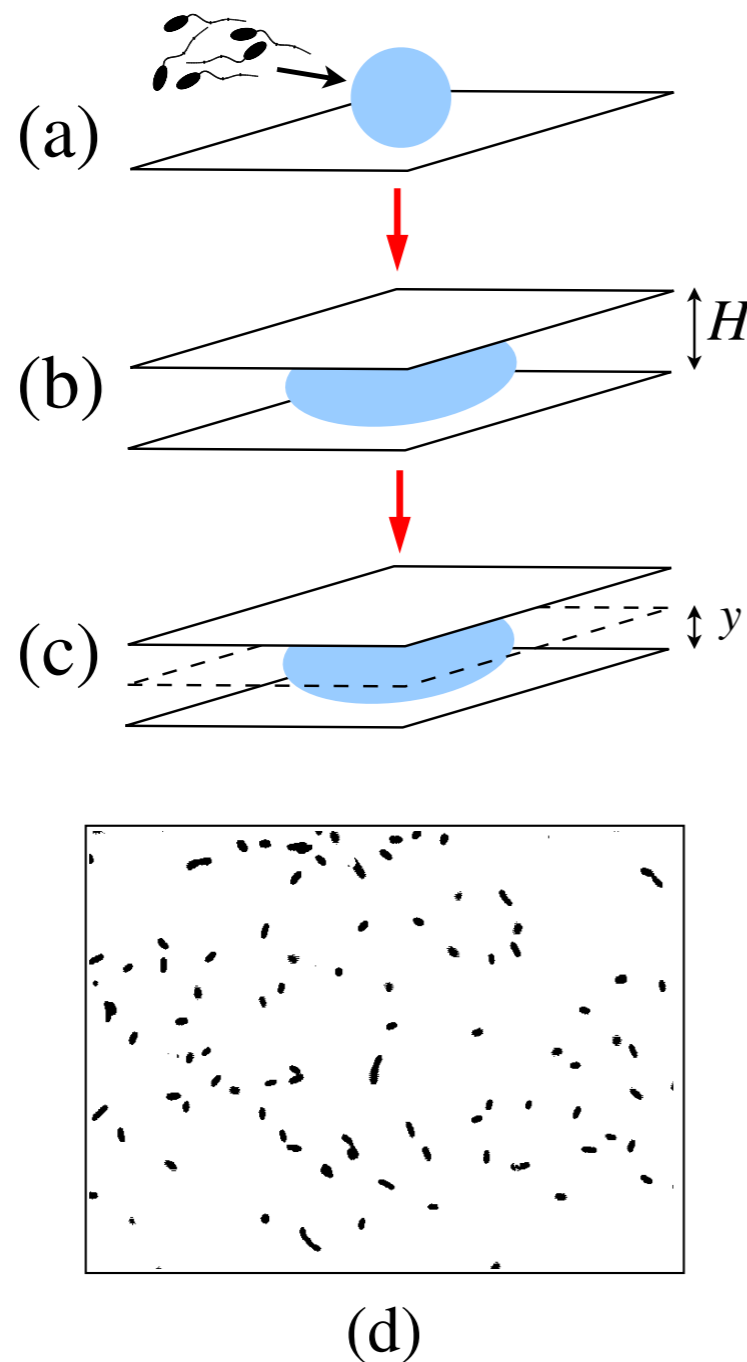
At large times, the motion becomes diffusive, with asymptotic diffusion constant

$$\lim_{t \rightarrow \infty} \frac{\mathbb{E}[\mathbf{X}^2]}{t} = \frac{2V^2}{(d-1)D_R} + 2dD_T. \quad (1.55)$$

Inserting typical values for bacteria,  $V \sim 10\mu\text{m/s}$  and  $D_R \sim 0.1/\text{s}$ , and comparing with  $D_T \sim 0.2\mu\text{m}^2/\text{s}$  for a micron-sized colloids at room temperature, we see that active swimming and orientational diffusion dominate the diffusive dynamics of microorganisms at long times.

# Hydrodynamic Attraction of Swimming Microorganisms by Surfaces

Allison P. Berke,<sup>1</sup> Linda Turner,<sup>2</sup> Howard C. Berg,<sup>2,3</sup> and Eric Lauga<sup>4,\*</sup>



# ‘Hydrodynamic’ fields

**Concentration profile between two walls** An interesting question that is relevant from a medical perspective concerns the spatial distribution of bacteria and other swimming microbes in the presence of confinement. Restricting ourselves to dilute suspensions<sup>10</sup>, we may obtain a simple prediction from the model (1.45) by considering the FPE for the associated PDF  $p(t, \mathbf{x}, \mathbf{n})$ . Given  $p$  and the total number of bacteria  $N_b$  in the solutions, we obtain the spatial concentration profile by integrating over all possible orientations

$$c(t, \mathbf{x}) = N_b \int_{\mathbb{S}_d} d\mathbf{n} p(t, \mathbf{n}, \mathbf{x}). \quad (1.56a)$$

The associated mean orientation field reads

$$\mathbf{u}(t, \mathbf{x}) = N_b \int_{\mathbb{S}_d} d\mathbf{n} p(t, \mathbf{n}, \mathbf{x}) \mathbf{n}. \quad (1.56b)$$

The FPE for the Ito-SDE (1.45) can be written as a conservation law

$$\partial_t p = -(\partial_{x_i} J_i + \partial_{n_i} \Omega_i), \quad (1.57a)$$

where

$$J_i = (V n_i - D_T \partial_{x_i}) p \quad (1.57b)$$

$$\Omega_i = D_R \left\{ (1 - d) n_i p - \partial_{n_j} [(\delta_{ij} - n_i n_j) p] \right\}. \quad (1.57c)$$

# Concentration field

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$$\Omega_i = D_R \left\{ (1 - d) n_i p - \partial_{n_j} [(\delta_{ij} - n_i n_j) p] \right\}. \quad (1.57c)$$

Focusing on the three-dimensional case,  $d = 3$ , we are interested in deriving from Eq. (1.57) the stationary concentration profile  $c$  of a suspension that is confined by two quasi-infinite parallel walls, which are located  $z = \pm H$ . That is, we assume that the distance between the walls is much smaller than their spatial extent in the  $(x, y)$ -directions,  $2H \ll L_x, L_y$ . To obtain an evolution equation for  $c$ , we multiply Eq. (1.57a) by  $N_b$  and integrate over  $\mathbf{n}$  with

$$\int_{\mathbb{S}_d} d\mathbf{n} \partial_{n_i} \Omega_i = 0. \quad (1.58)$$

This yields the mass conservation law

$$\partial_t c = -\nabla \cdot (V \mathbf{u} - D_T \nabla c). \quad (1.59)$$

# Orientation (velocity) field

The FPE for the Ito-SDE (1.45) can be written as a conservation law

$$\partial_t p = -(\partial_{x_i} J_i + \partial_{n_i} \Omega_i), \quad (1.57a)$$

where

$$J_i = (V n_i - D_T \partial_{x_i}) p \quad (1.57b)$$

$$\Omega_i = D_R \left\{ (1 - d) n_i p - \partial_{n_j} [(\delta_{ij} - n_i n_j) p] \right\}. \quad (1.57c)$$

To obtain also an evolution equation for  $\mathbf{u}$ , we multiply Eq. (1.57a) by  $n_k$ ,

$$\partial_t (n_k p) = -\partial_{x_i} (n_k J_i) - n_k \partial_{n_i} \Omega_i. \quad (1.60)$$

and note that

$$n_k \partial_{n_i} \Omega_i = \partial_{n_i} (n_k \Omega_i) - (\partial_{n_i} n_k) \Omega_i = \partial_{n_i} (n_k \Omega_i) - \delta_{ik} \Omega_i. \quad (1.61)$$

# Orientation (velocity) field

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This allows us to rewrite (1.60) as

$$\begin{aligned} \partial_t(n_k p) &= -\partial_{x_i}(n_k J_i) + \Omega_k - \partial_{n_i}(n_k \Omega_i) \\ &= -\partial_{x_i}[V n_k n_i p - D_T \partial_{x_i}(n_k p)] + \\ &\quad D_R \{-2n_k p - \partial_{n_j}[(\delta_{kj} - n_k n_j)p]\} - \partial_{n_i}(n_k \Omega_i) \\ &= -\partial_{x_i}[V n_k n_i p - D_T \partial_{x_i}(n_k p)] - 2D_R n_k p - \\ &\quad \partial_{n_j}(n_k \Omega_j + (\delta_{kj} - n_k n_j)p). \end{aligned} \quad (1.62)$$

# Orientation (velocity) field

To obtain also an evolution equation for  $\mathbf{u}$ , we multiply Eq. (1.57a) by  $n_k$ ,

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Multiplying by  $N_b$  and integrating over  $\mathbf{n}$  with appropriate boundary conditions gives

$$\partial_t u_k = -\partial_{x_i}[V N_b \langle n_k n_i \rangle p - D_T \partial_{x_i} u_k] - 2D_R u_k,$$

where we have abbreviated

$$\langle n_i n_k \cdots \rangle = \int_{\mathbb{S}_d} d\mathbf{n} p(t, \mathbf{n}, \mathbf{x}) n_i n_k \cdots. \quad (1.63)$$

To obtain a closed linear system of equations for the fields  $(c, \mathbf{u})$ , we neglect<sup>11</sup> the higher-order moments  $N_b \langle n_k n_i \rangle$  in (1.63) and find

$$\partial_t \mathbf{u} \simeq -2D_R \mathbf{u} + D_T \nabla^2 \mathbf{u}. \quad (1.64)$$

# Stationary profiles

$$\partial_t \mathbf{u} \simeq -2D_R \mathbf{u} + D_T \nabla^2 \mathbf{u}. \quad (1.64)$$

To find the stationary density and orientation profiles, we look for solutions of the form  $c = \rho(z)$  and  $u_x = u_y = 0, u_z = u(z)$ . According to Eqs. (1.59) to (1.63), the functions  $\rho$  and  $u_z$  must satisfy

$$0 = Vu - D_T c', \quad (1.65)$$

$$0 = -2D_R u + D_T u'', \quad (1.66)$$

and it is physically plausible that they also fulfill the symmetry<sup>12</sup> requirements  $\rho(z) = \rho(-z)$  and  $u(z) = -u(-z)$ . Hence, solution takes the form

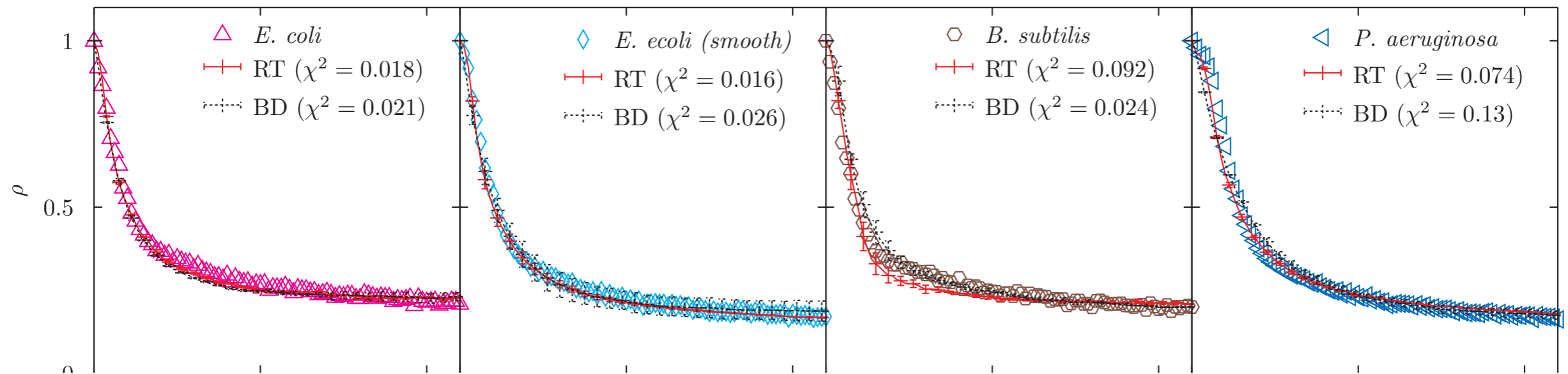
$$u(z) = A \sinh(z/\Lambda), \quad (1.67a)$$

$$\rho(z) = A \frac{V\Lambda}{D_T} [\cosh(z/\Lambda) - 1] + \rho_0, \quad (1.67b)$$

where  $\Lambda = \sqrt{D_\perp/(2D_R)}$ .

The cosh-profile (1.67b) agrees qualitatively with experimental measurements for dilute bacterial suspensions [BTBL08, LT09].

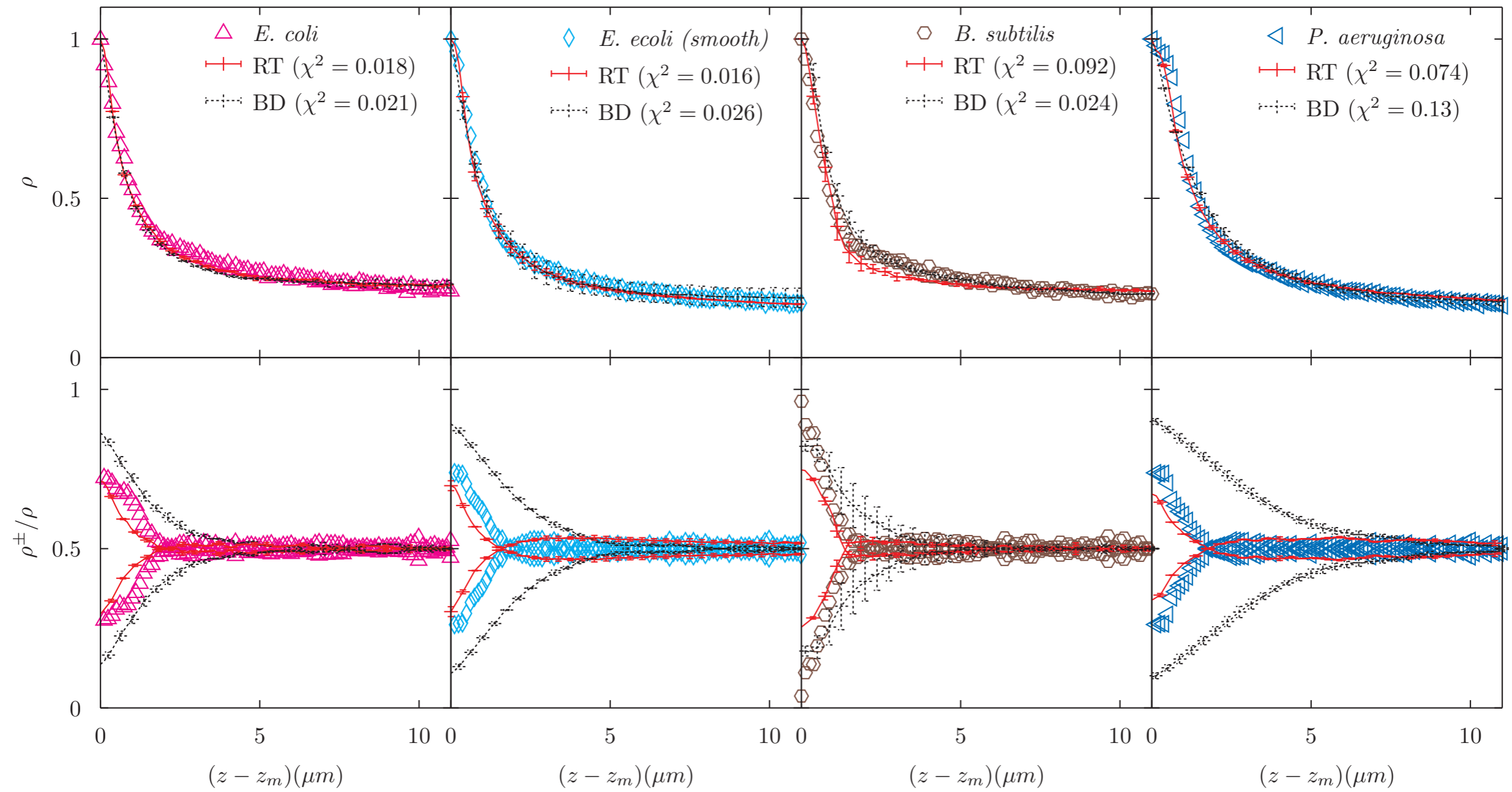
But ...



Density profiles seem ok ... what about fluxes?

joint work with Peter Lu, Rik Wensink & Jeff Guasto

# But ...



# Need to include run & tumbling

**BD:**

$$\begin{aligned}\delta \mathbf{r} &= \mathbf{D}_T \cdot \mathbf{F}_{bw} \delta t + v_0 \hat{\mathbf{u}} \delta t + \sqrt{2\delta t D_T} \delta \mathbf{r}_G \\ \delta \hat{\mathbf{u}} &= D_R (\mathbf{T}_{bw} \times \hat{\mathbf{u}}) \delta t + \sqrt{2\delta t D_R} \delta \hat{\mathbf{u}}_G\end{aligned}\quad (\text{SF-SDE})$$

$$P_T^{-1} = D_T / v_0 \ell \text{ and } P_R^{-1} = D_R \ell / v_0$$

**RT:**

$$\begin{aligned}\ell^{-1} \delta \mathbf{r} &= \frac{\mathbf{F}_{bw}}{F_a} \delta \tau \\ \delta \hat{\mathbf{u}} &= \xi \left( \frac{\mathbf{T}_{bw}}{F_a \ell} \times \hat{\mathbf{u}} \right) \delta \tau + \left[ \left( \hat{\mathbf{u}} \times \frac{\Delta \hat{\mathbf{u}}}{\Delta \tau_{\text{tumble}}} \right) \times \hat{\mathbf{u}} \right] \delta \tau\end{aligned}$$

TABLE I: Main bacterial parameters used for the fit.

culture	$\ell$ ( $\mu m$ )	$a$ ( $a_{\text{eff}}$ )	$v_0$ ( $\mu m/s$ )	$\theta_T$	$\Delta t_{\text{tumble}}$ (s)	$\Delta t_{\text{run}}$ (s)
<i>E. coli</i>	$\sim 3$	$\sim 2(5.6)$	$\sim 20$	$68^\circ$	$\sim 0.1$	$\sim 1$
<i>E. coli</i> (smooth)	$\sim 3$	$\sim 2(5.6)$	$\sim 20$	$0^\circ$	0	$\infty$
<i>B. subtilis</i>	$\sim 5$	$\sim 6(11.5)$	$\sim 50$	$\sim 40^\circ$	$\sim 0.1$	$\sim 0.5$
<i>P. aeruginosa</i>	$\sim 2$	$\sim 4(9.8)$	$\sim 40$	$\sim 110^\circ$	$\sim 0.1$	$\sim 0.5$

	RT		BD	
	rotation $P_R^{-1}$	translation $P_T^{-1}$	rotation $P_R^{-1}$	translation $P_T^{-1}$
<i>E. coli</i>	$0.04 \pm 0.005$	$0.1 \pm 0.01$	$0.11 \pm 0.005$	$0.31 \pm 0.005$
<i>E. coli</i> (smooth)	$0.02 \pm 0.005$	$0.07 \pm 0.005$	$0.07 \pm 0.001$	$0.34 \pm 0.001$
<i>B. subtilis</i>	$0.14 \pm 0.05$	$0.02 \pm 0.005$	$0.10 \pm 0.001$	$0.09 \pm 0.001$
<i>P. aeruginosa</i>	$0.02 \pm 0.005$	$0.08 \pm 0.005$	$0.04 \pm 0.001$	$0.49 \pm 0.005$

# Need to include run & tumbling

