

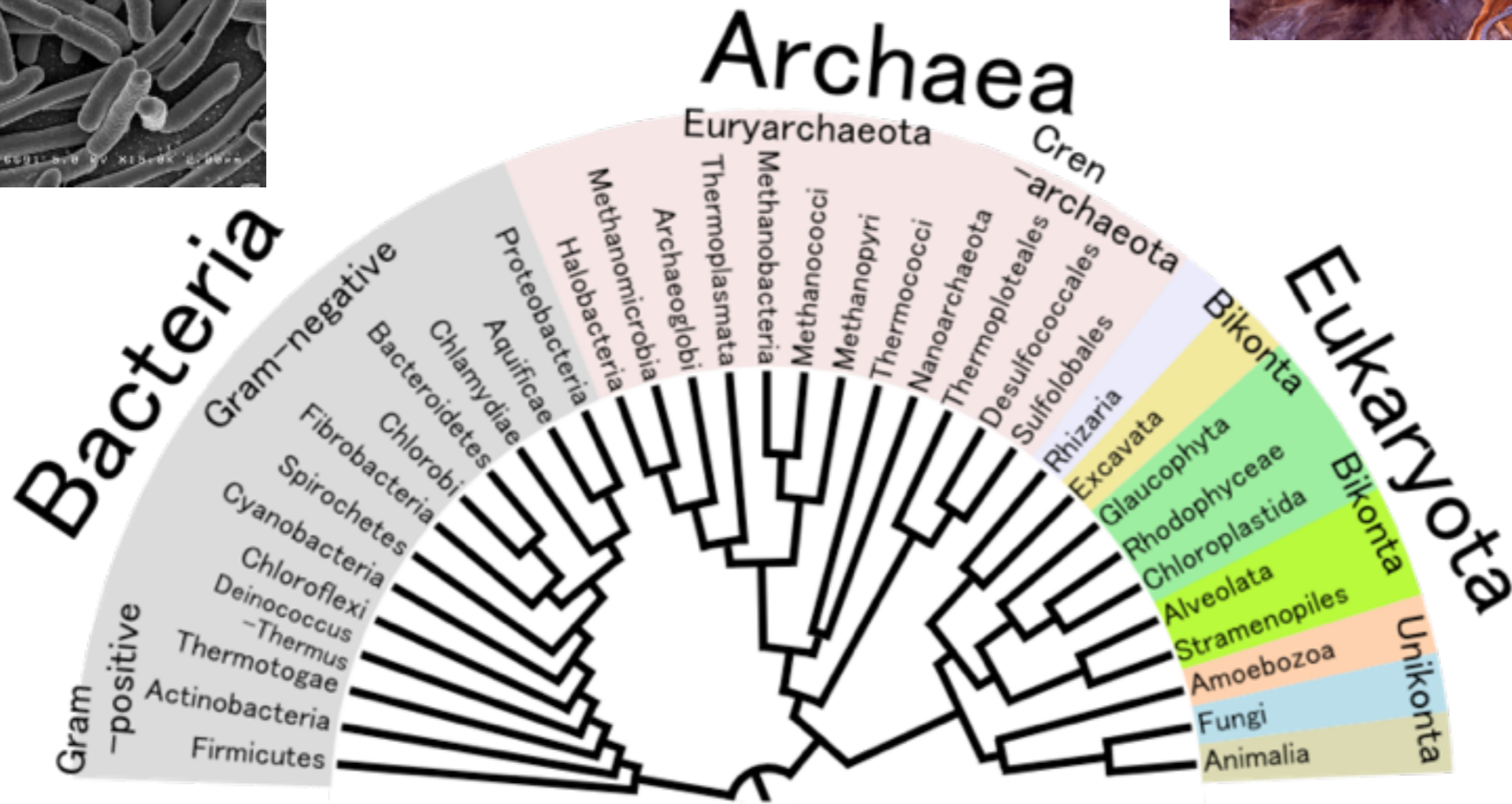
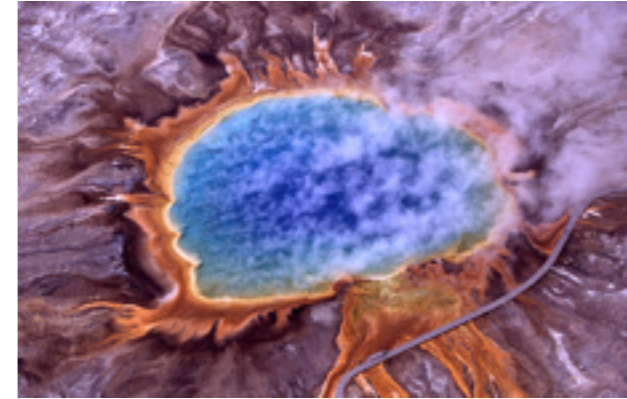
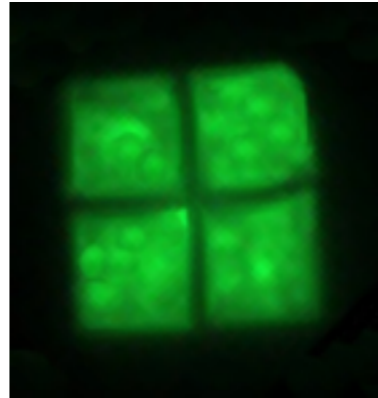
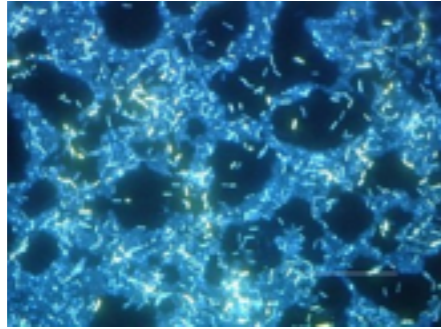
# (Some) Numbers and Maths in Biology

Jörn Dunkel  
E17-412  
[dunkel@math.mit.edu](mailto:dunkel@math.mit.edu)

<http://bionumbers.hms.harvard.edu/>

**B10NUMB3R5**  
THE DATABASE OF USEFUL BIOLOGICAL NUMBERS

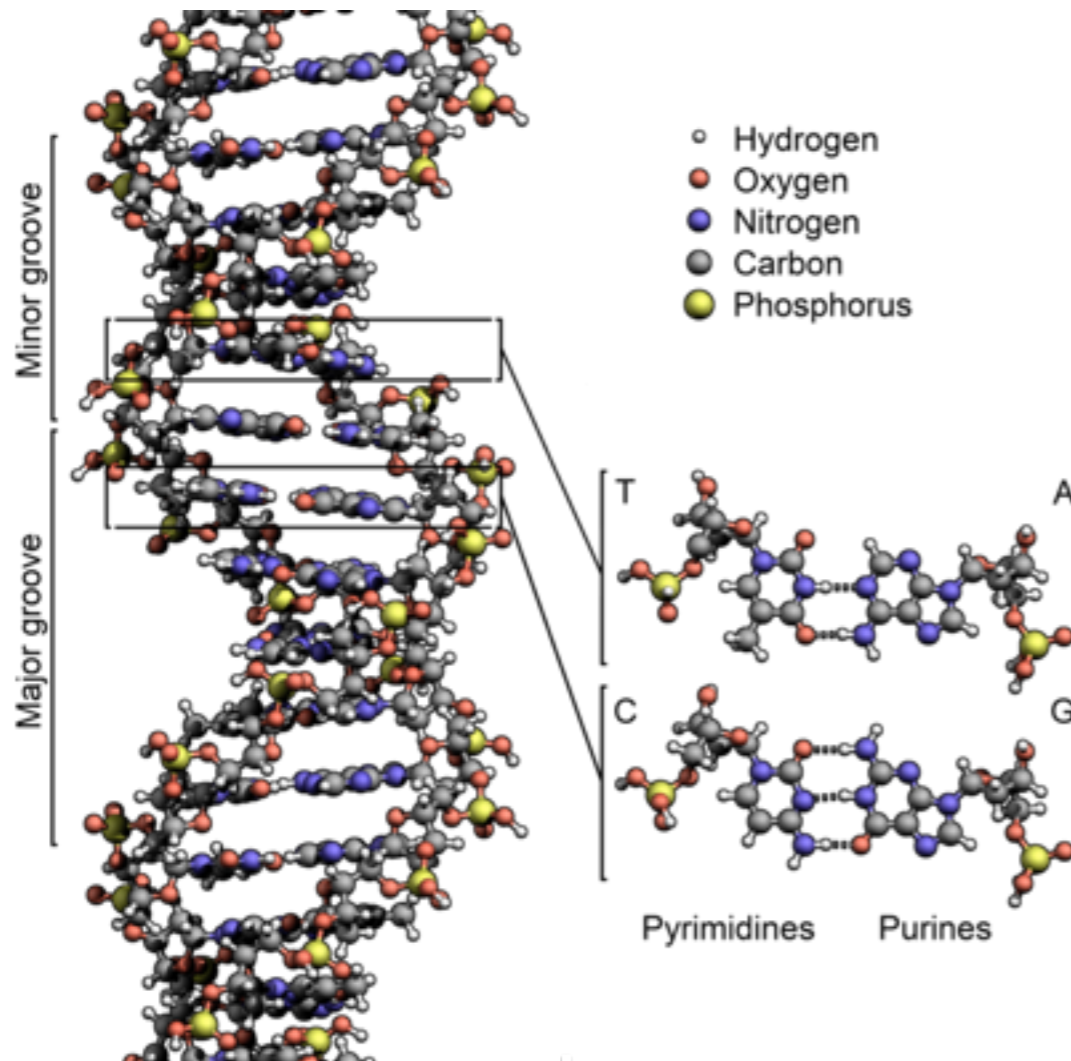
# Phylogenetic tree



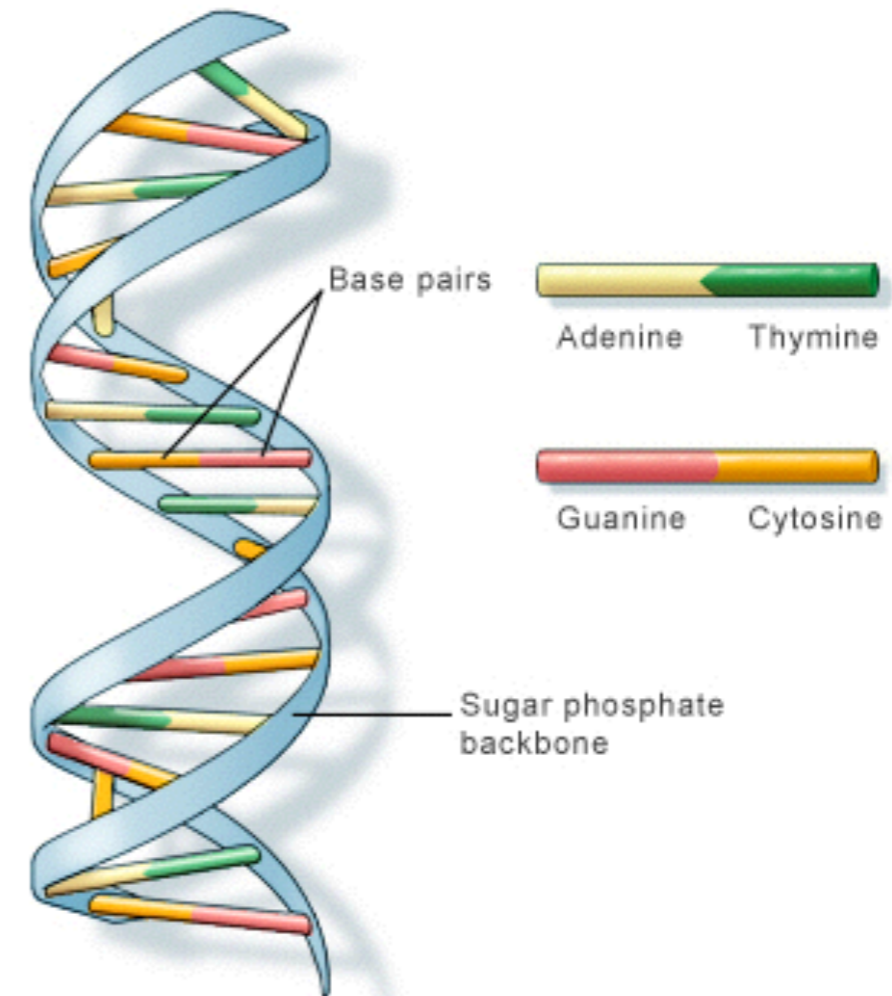
source: wiki



# DNA



source: wiki



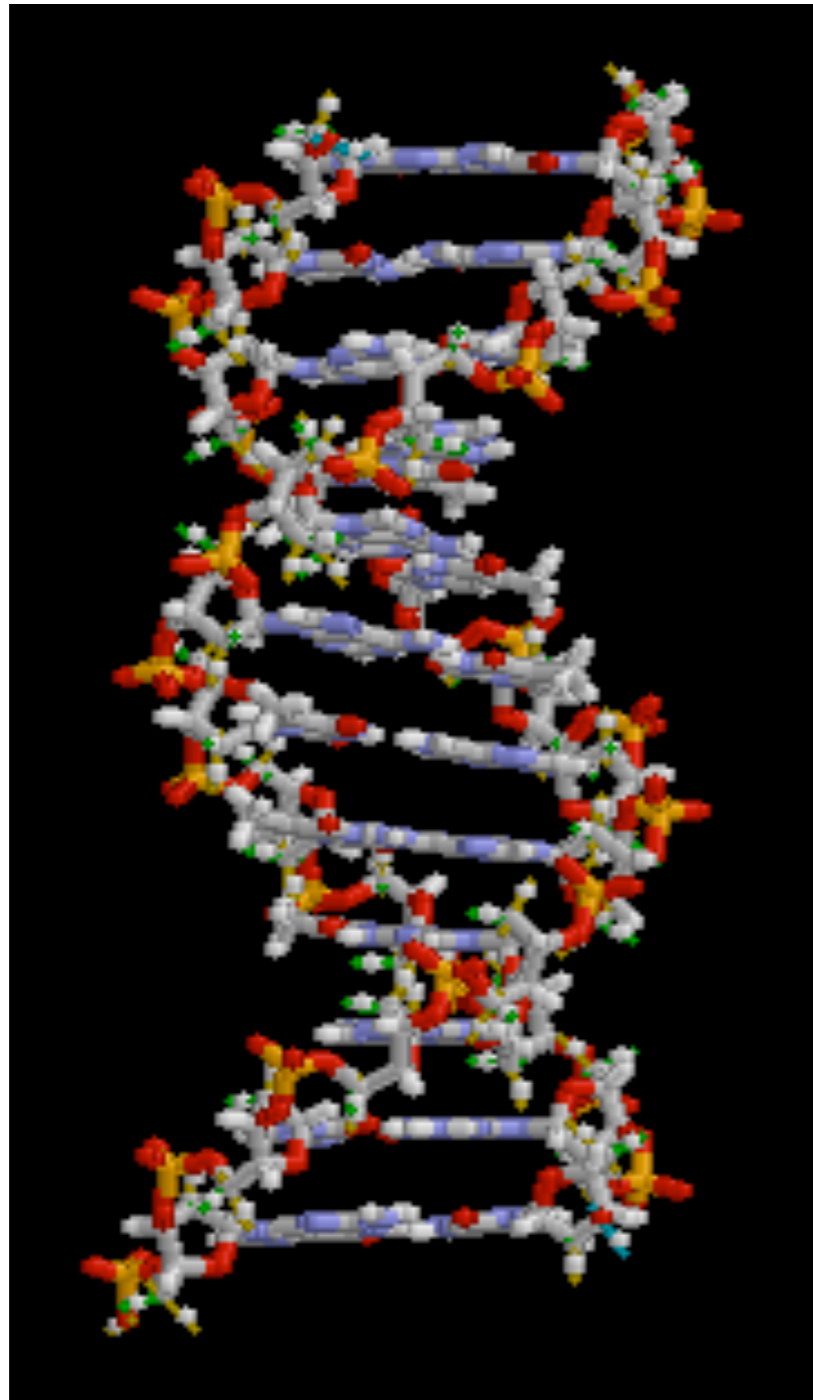
U.S. National Library of Medicine

<http://ghr.nlm.nih.gov/handbook/basics/dna>

- DNA contour length in bacteria:  $\sim 1.5\text{mm}$
- Length of DNA in nucleus of mammals:  $\sim 2\text{m}$

dunkel@math.mit.edu

# DNA = biopolymer pair

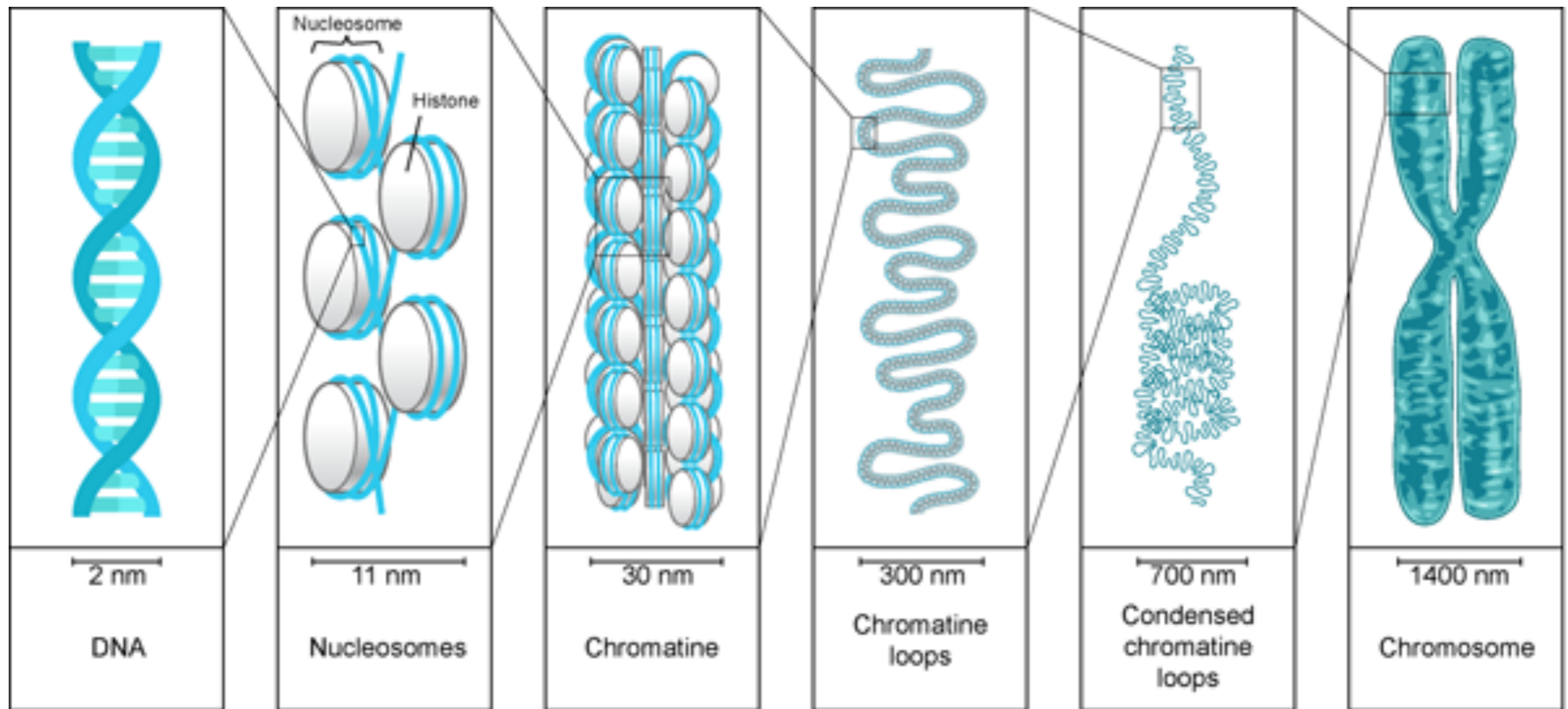


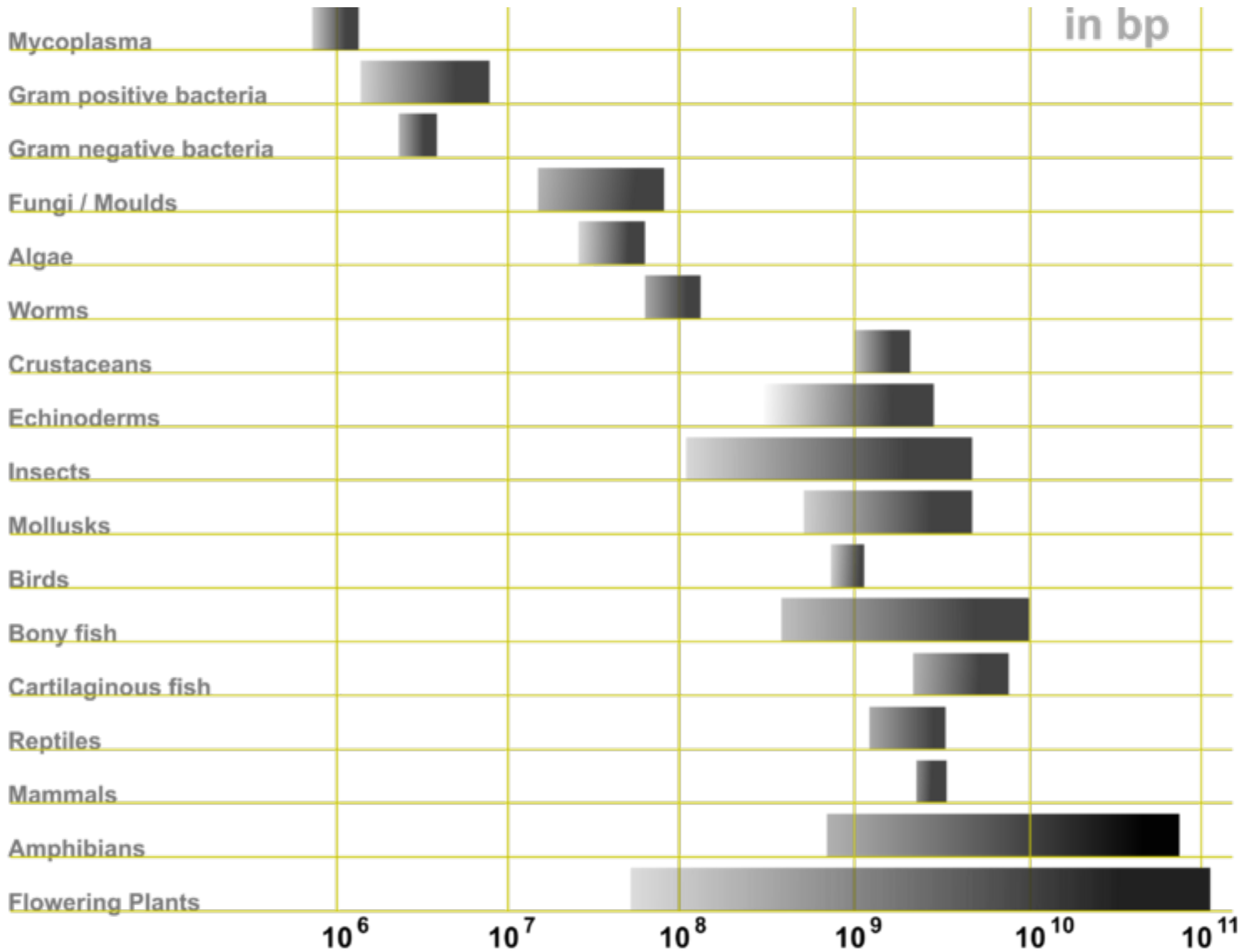
~ 3m per cell

~  $10^{14}$  cells/human

> max. distance between  
Earth and Pluto  
(~50 AU =  $7.5 \times 10^{12}$  m)

# DNA packaging in eukaryotes

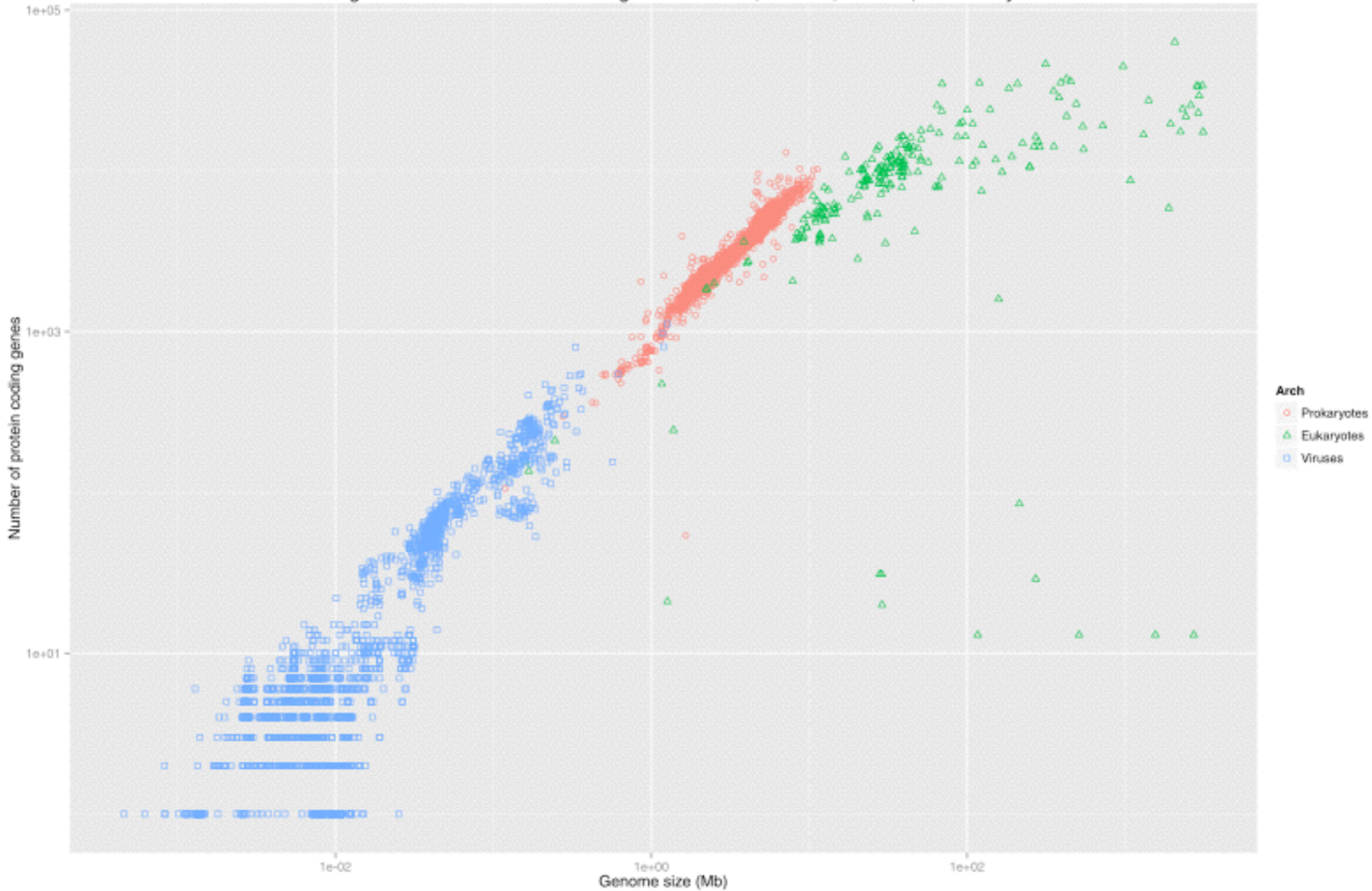




source: wiki

mass 1pg = 978Mb

The total genome size and the number of genes in viruses, bacteria, archaea, and eukaryotes.



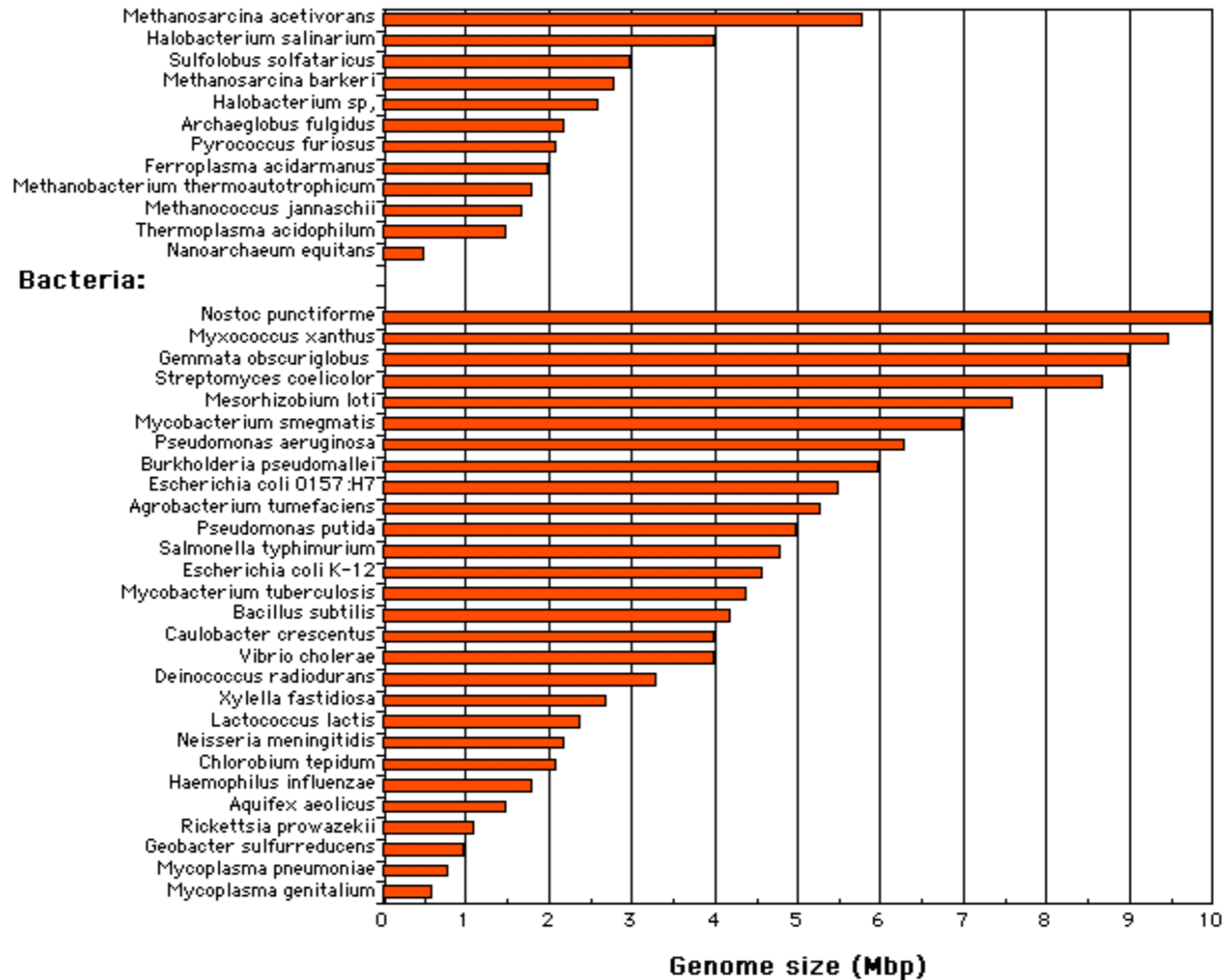
source: wiki

dunkel@math.mit.edu

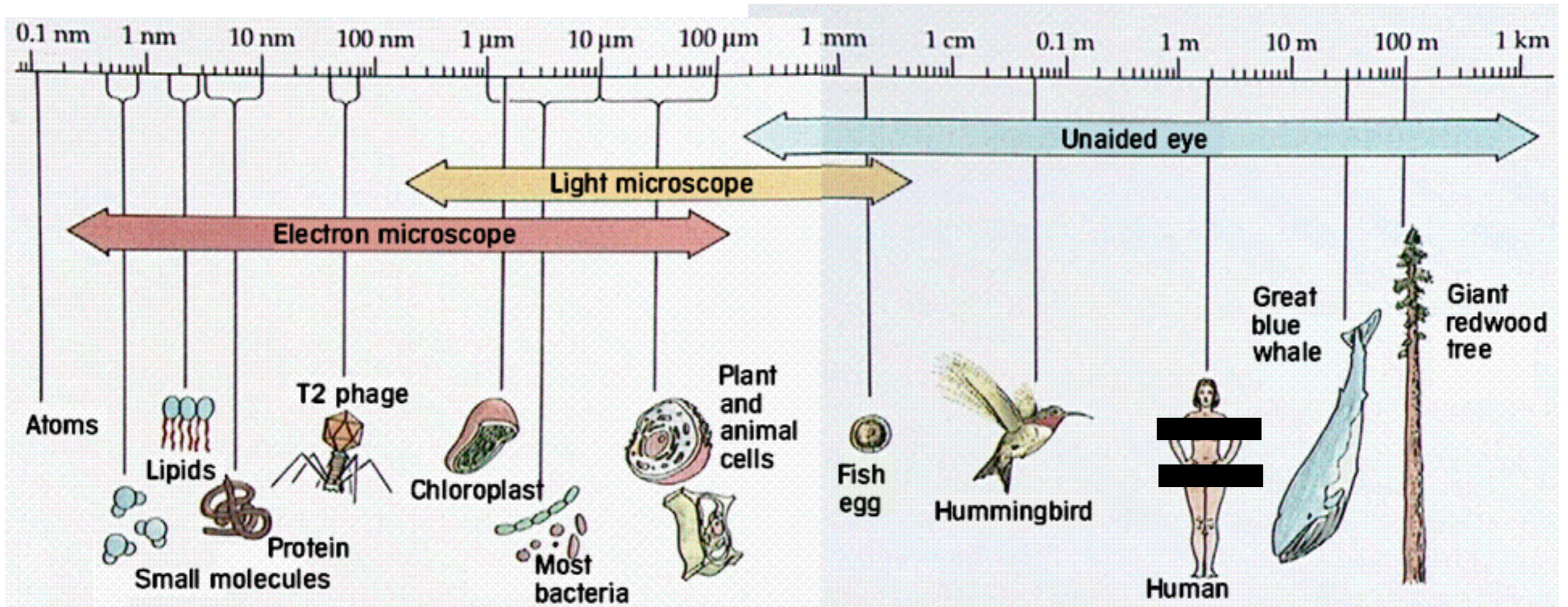


# Prokaryotes

## Archaea:



# Typical length scales

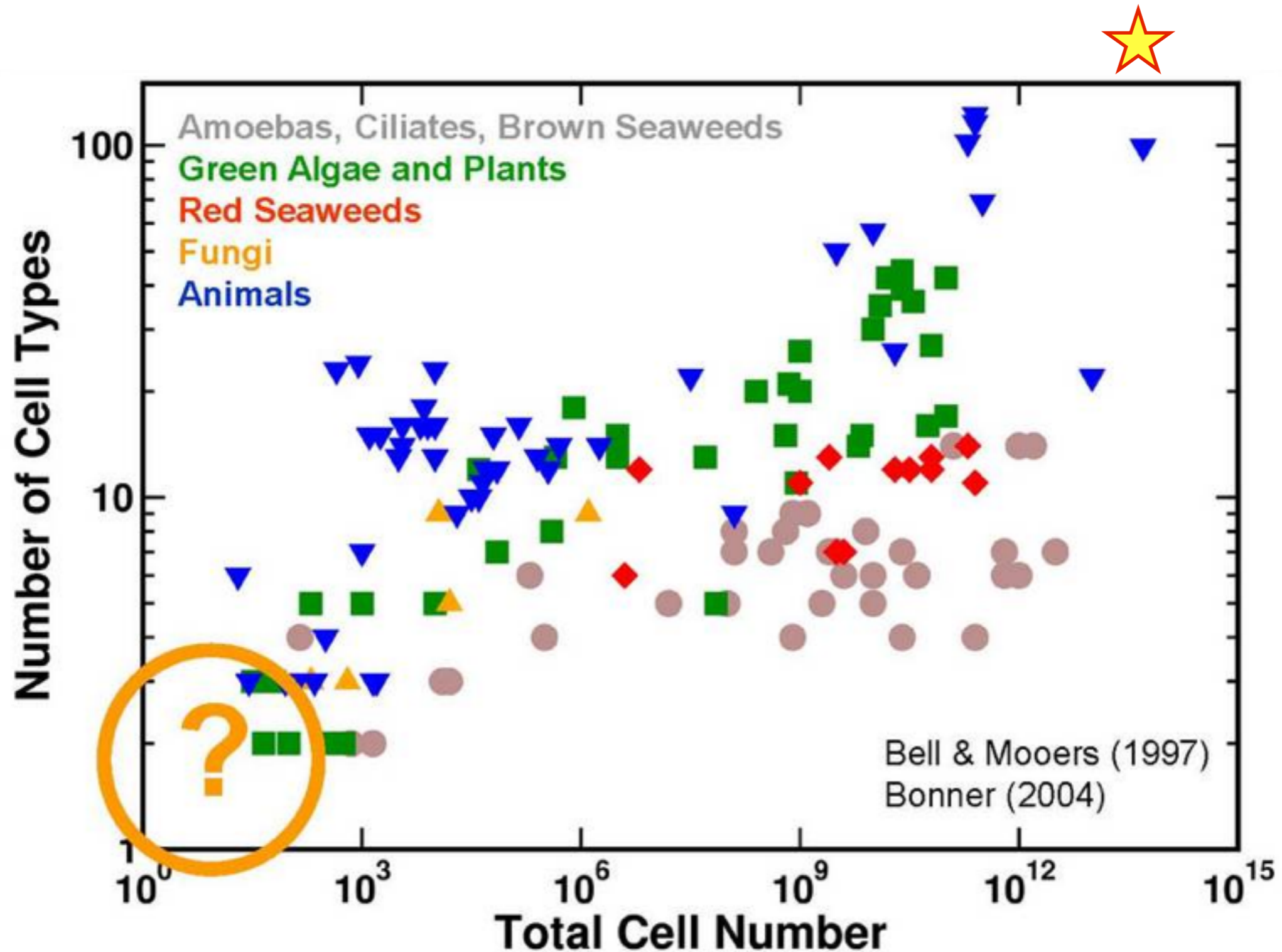


<http://www2.estrellamountain.edu/faculty/farabee/BIOBK/biobookcell2.html>

# Species estimates

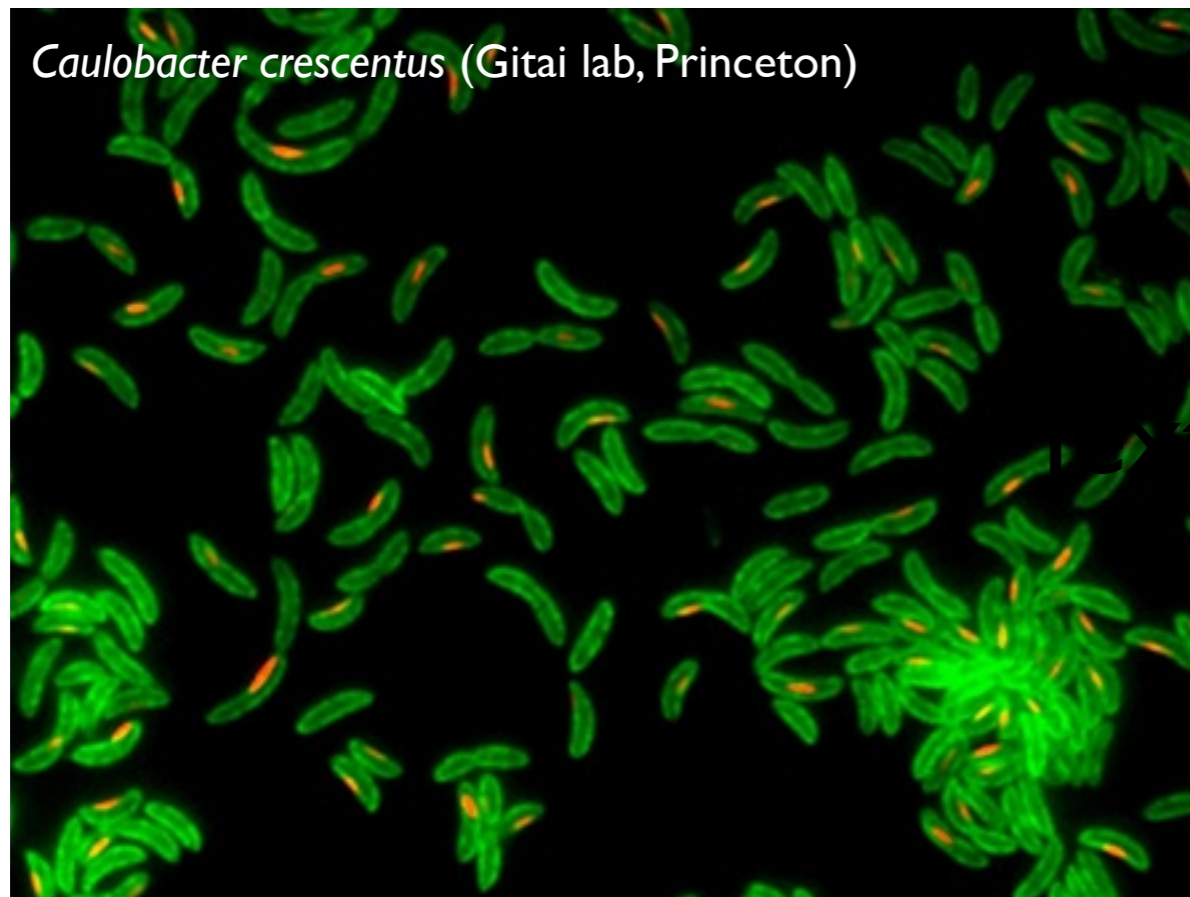
- estimated number of eukaryotic species on Earth: 8.7 million (Nature, 2011)
- undiscovered: 86% land spec. & 91% marine spec
- ~ 300,000 plant species
- prokaryotic biomass ~ eukaryotic biomass
- oldest known fossilized prokaryotes from 3.5 billion years ago

# Size-Complexity relation



# Unicellular organisms

## Bacteria



size  $\sim 1\mu\text{m}$   
doubling time  $\sim 2\text{h}$

## Algae



size  $\sim 10\mu\text{m}$   
doubling time  $\sim 5\text{-}8\text{h}$

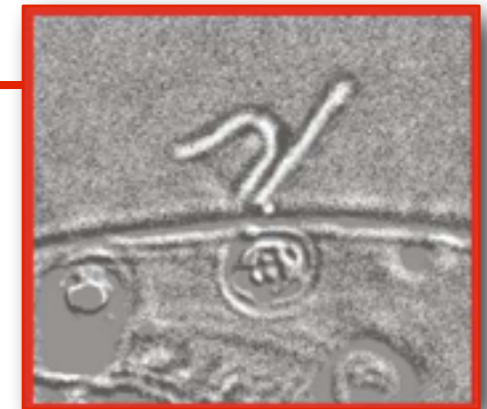
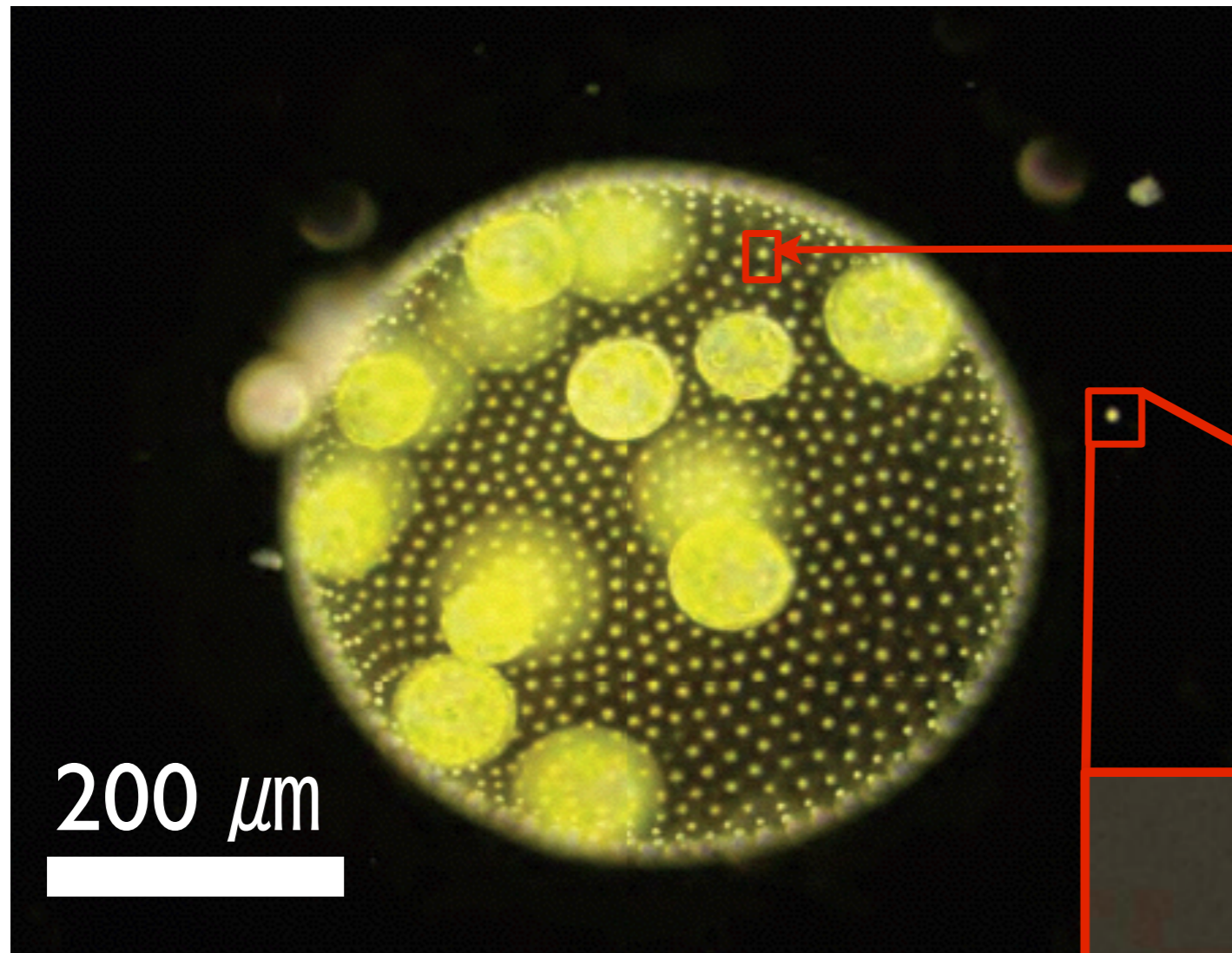
## Amoeba



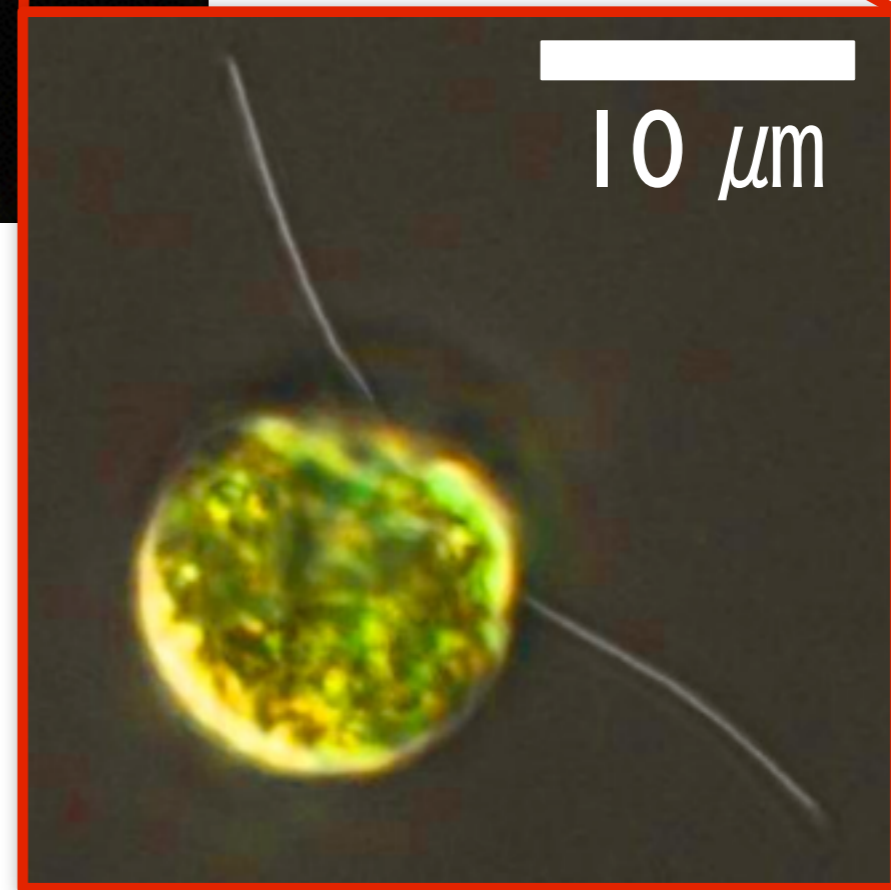
size  $\sim 1\text{mm}$   
doubling time  $\sim 1\text{d}$

evolution from  
unicellular to multicellular ?

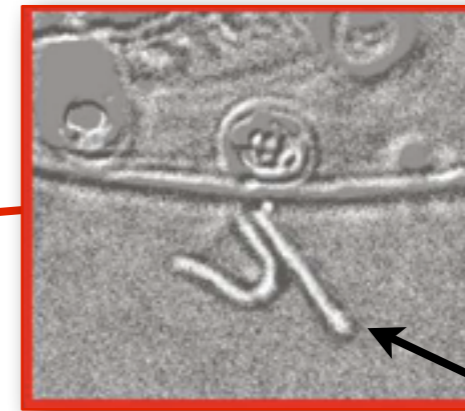
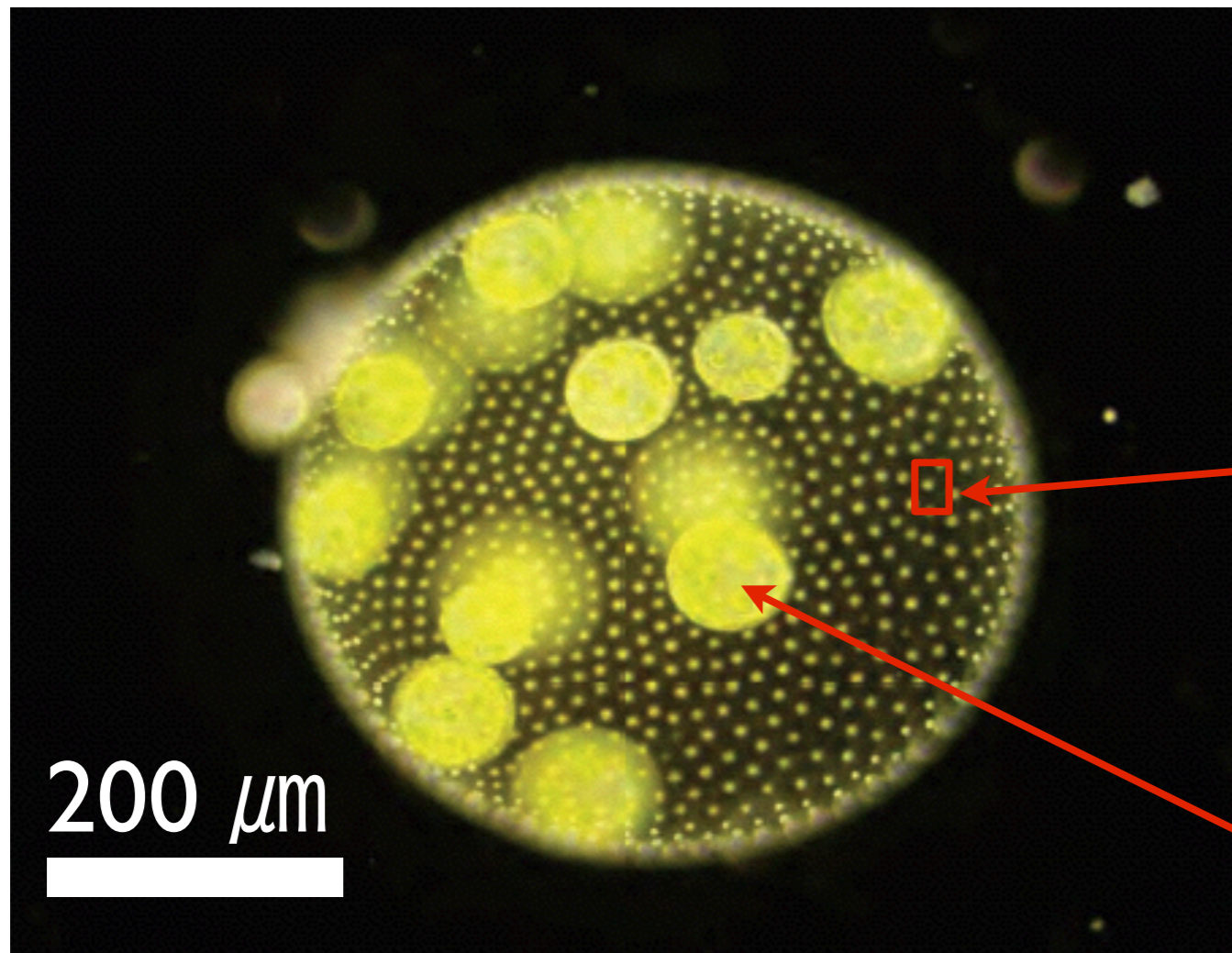
# Volvox carteri



# Chlamydomonas reinhardtii



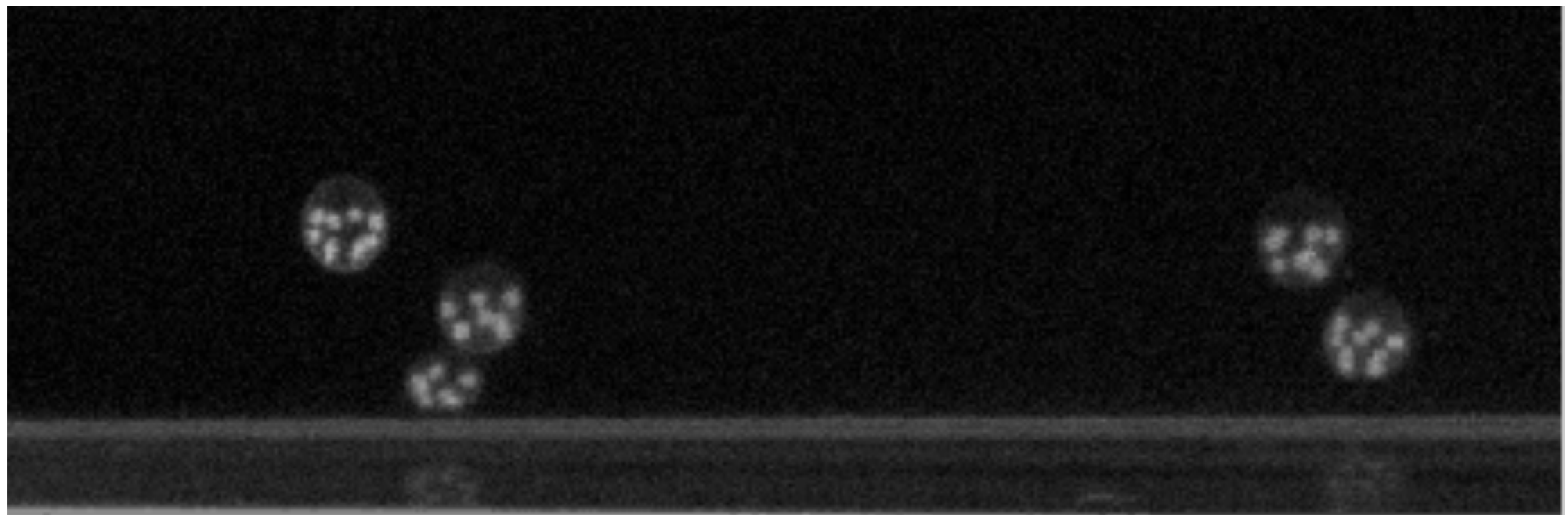
# Volvox carteri



somatic  
cell

cilia

daughter colony

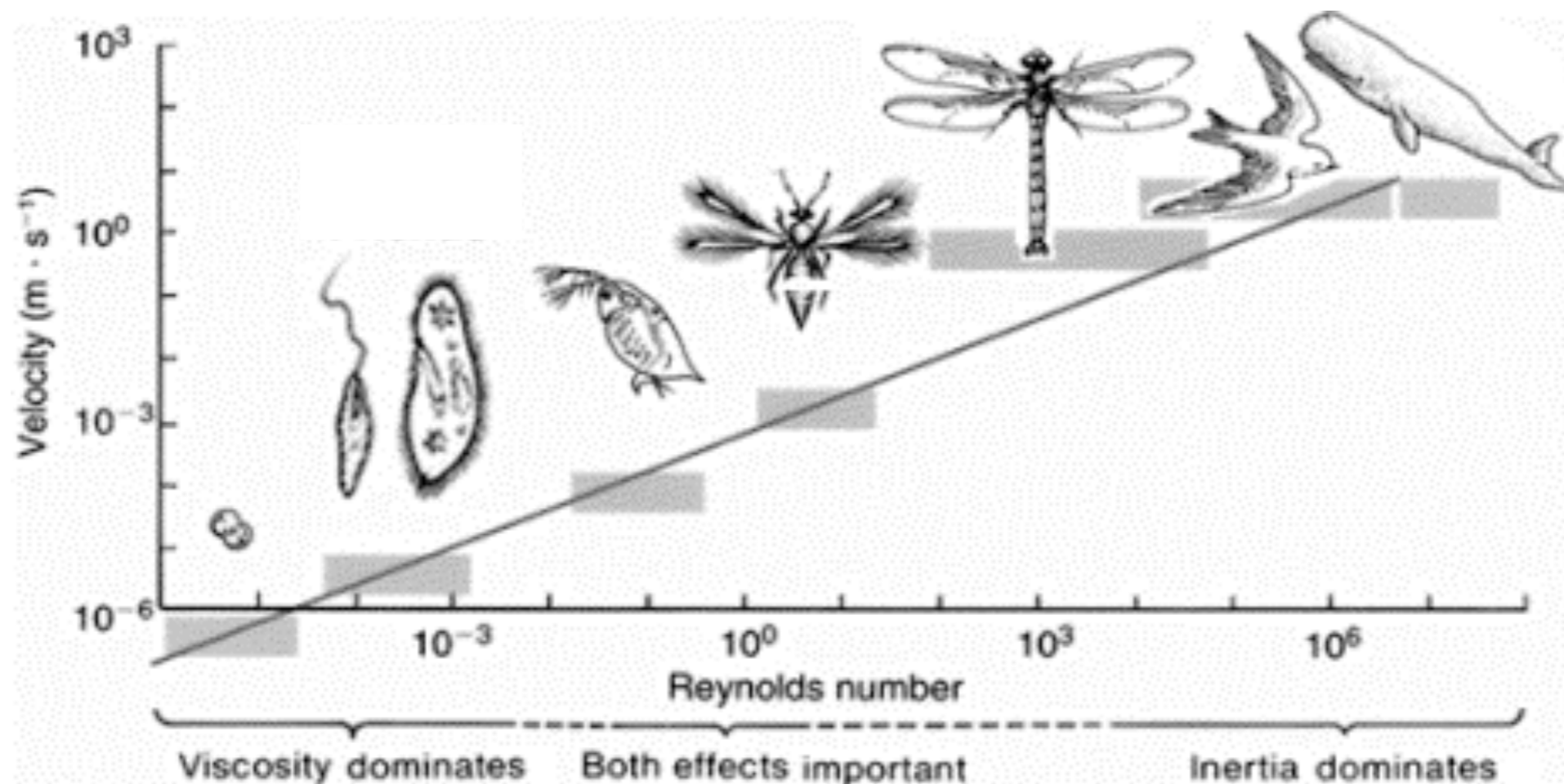




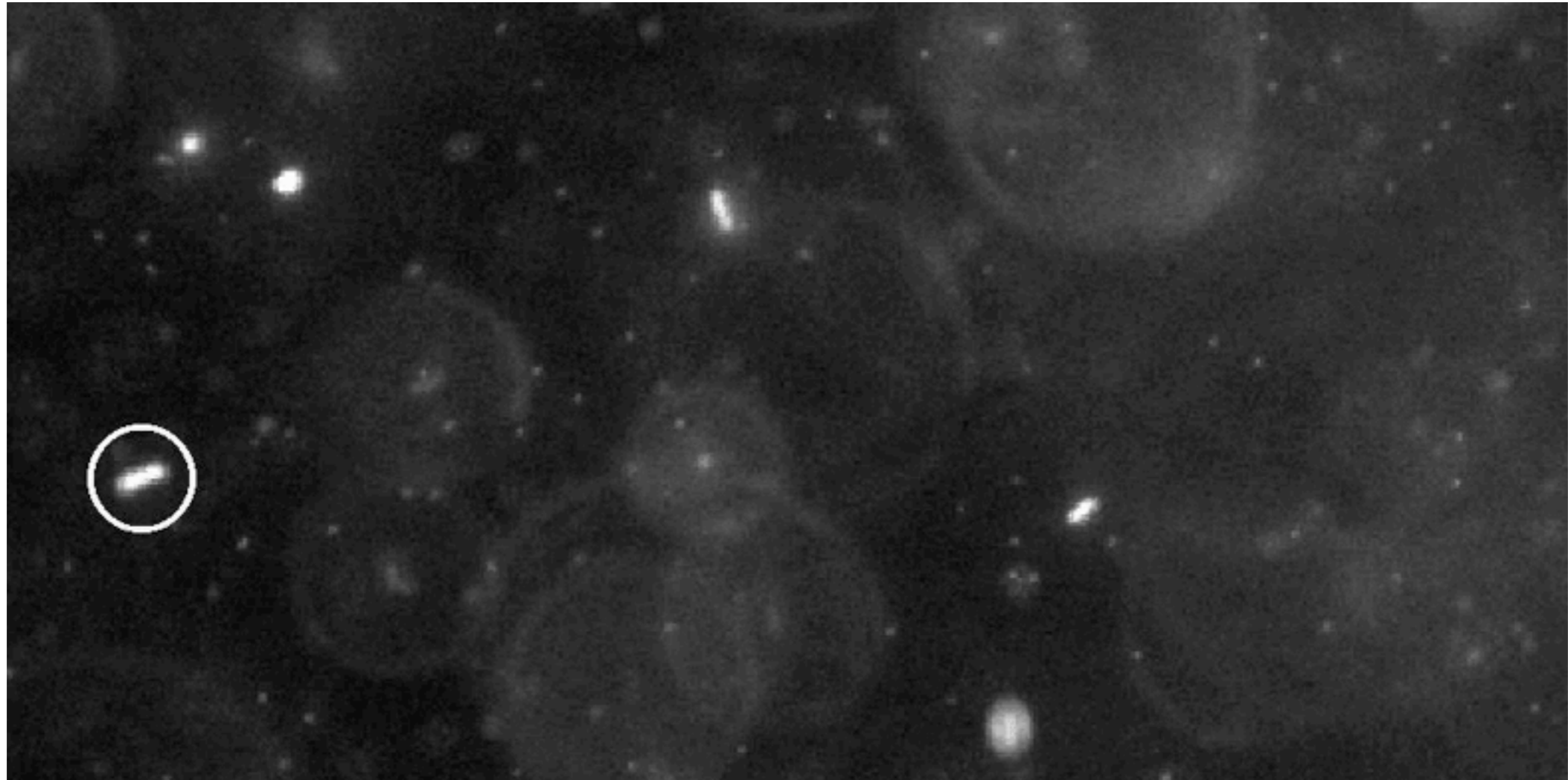
how do organisms  
achieve **locomotion** ?

# Reynolds numbers

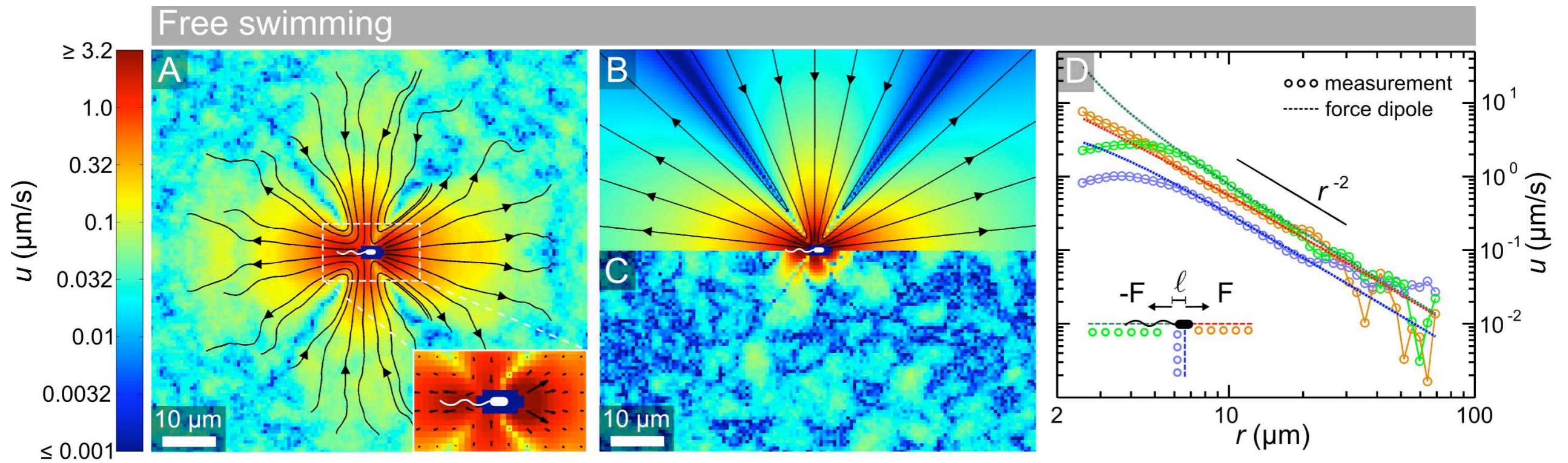
$$Re = \frac{\rho U L}{\mu} = \frac{U L}{\nu}$$



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



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



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

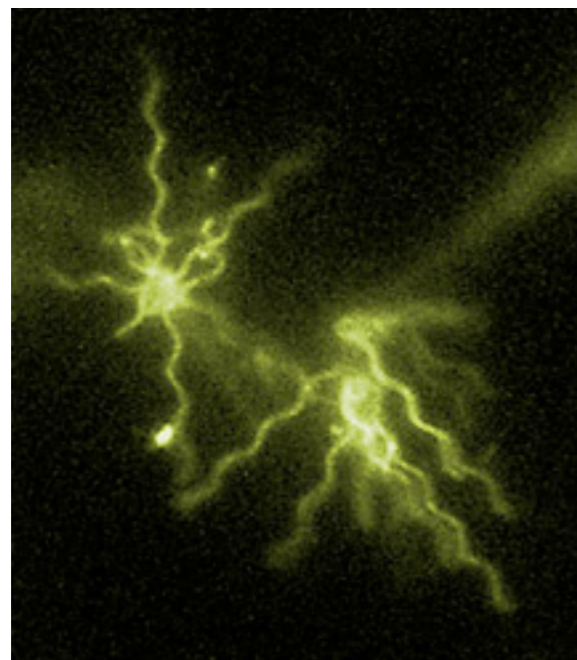
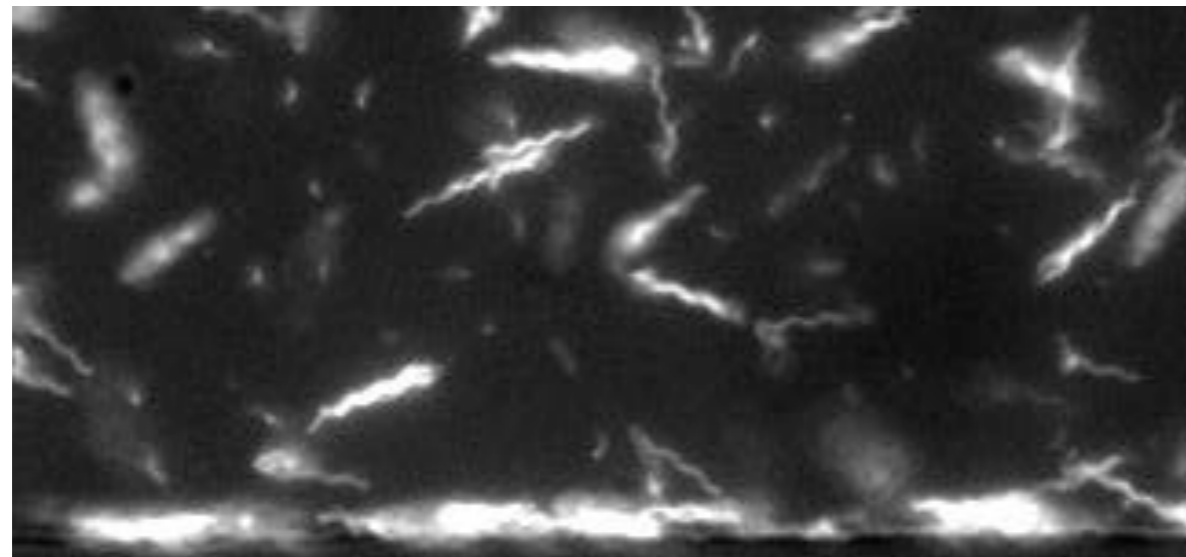
$$V_0 = 22 \pm 5 \mu\text{m/s}$$

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

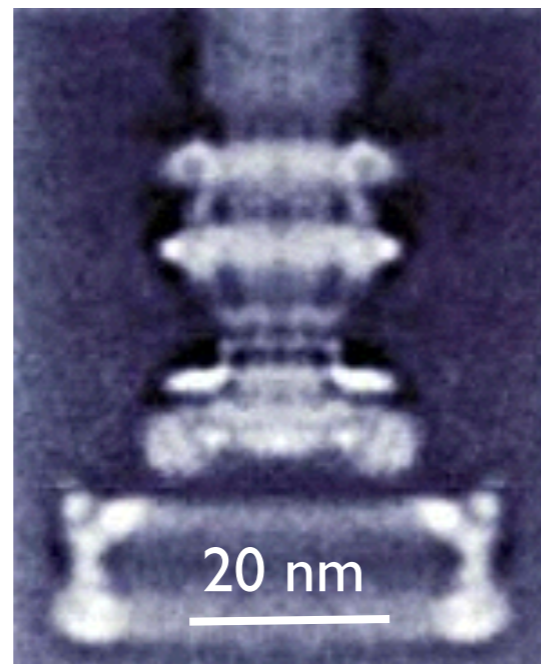
$$F = 0.42 \text{ pN}$$

# Bacterial motors

movie: V. Kantsler

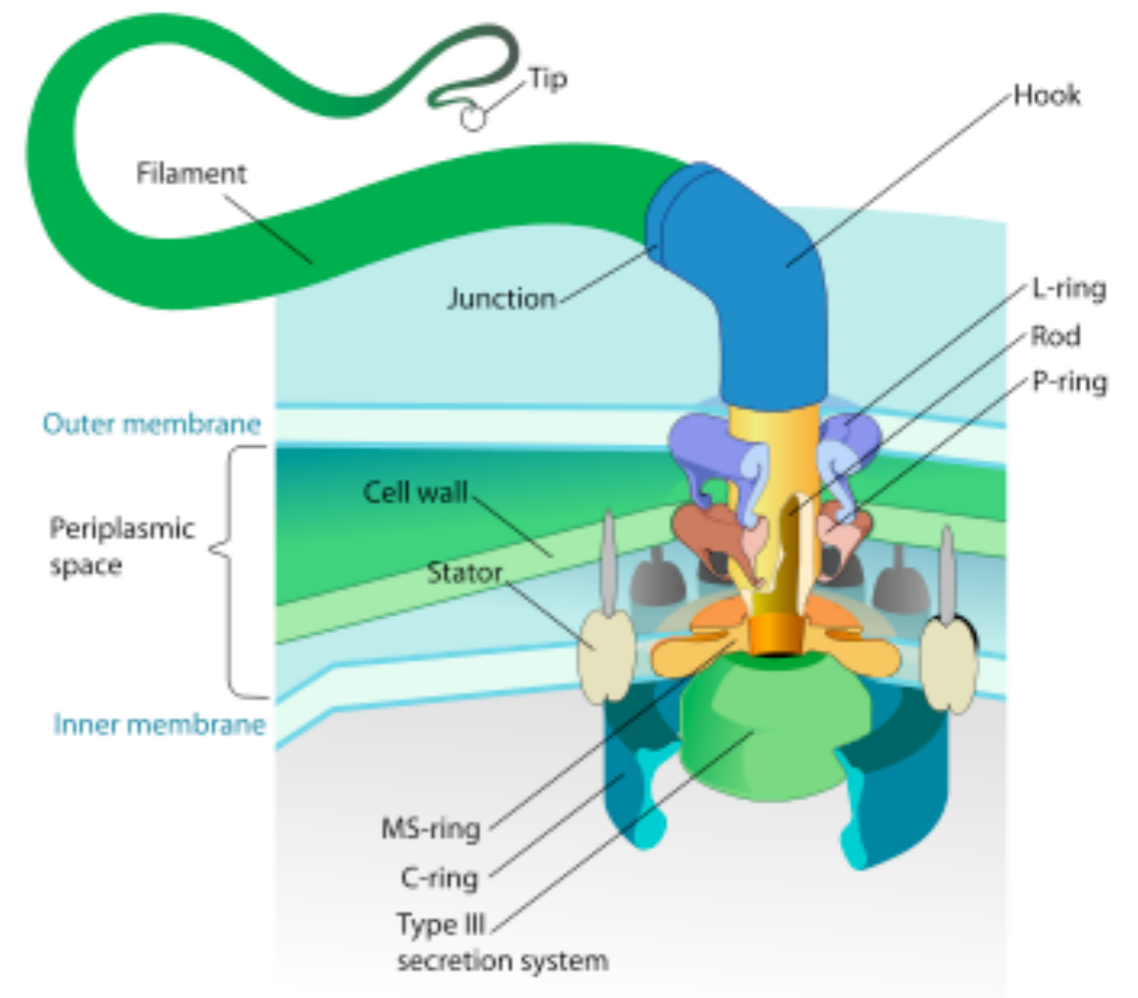


Berg (1999) Physics Today



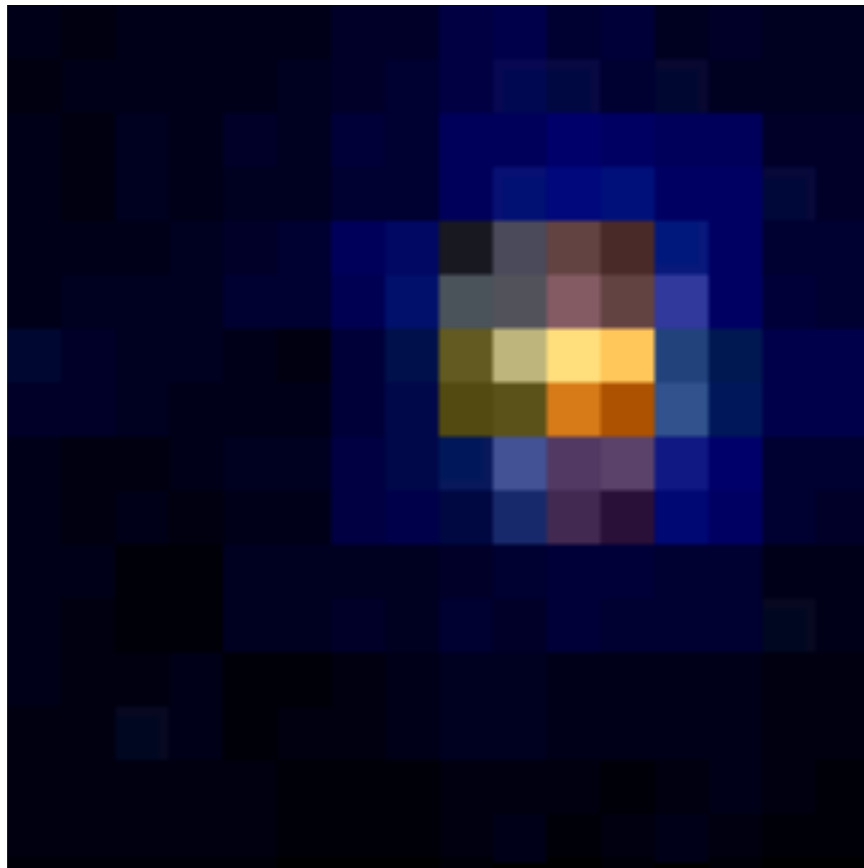
Chen et al (2011) EMBO Journal

~20 parts

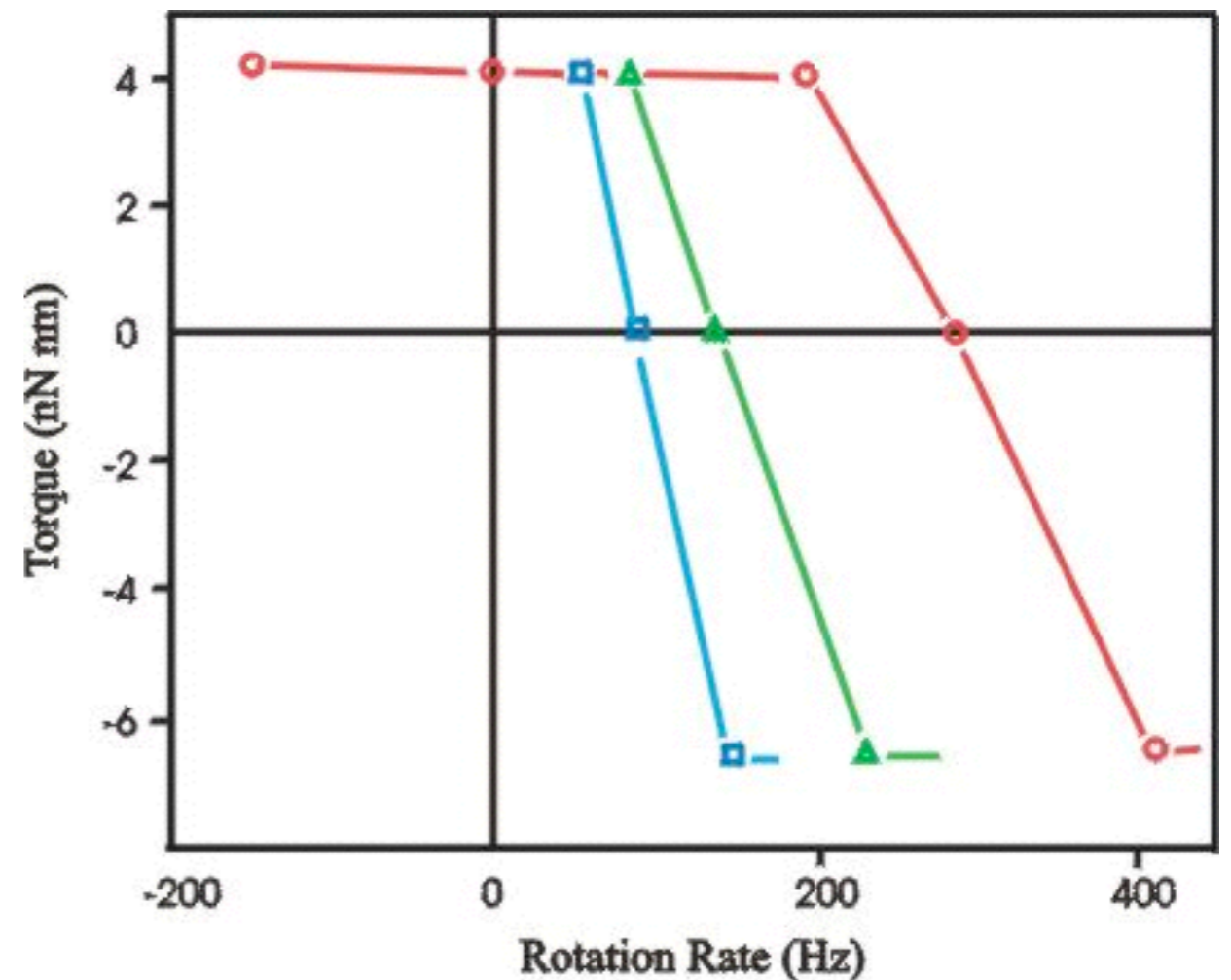


source: wiki

# Torque-speed relation

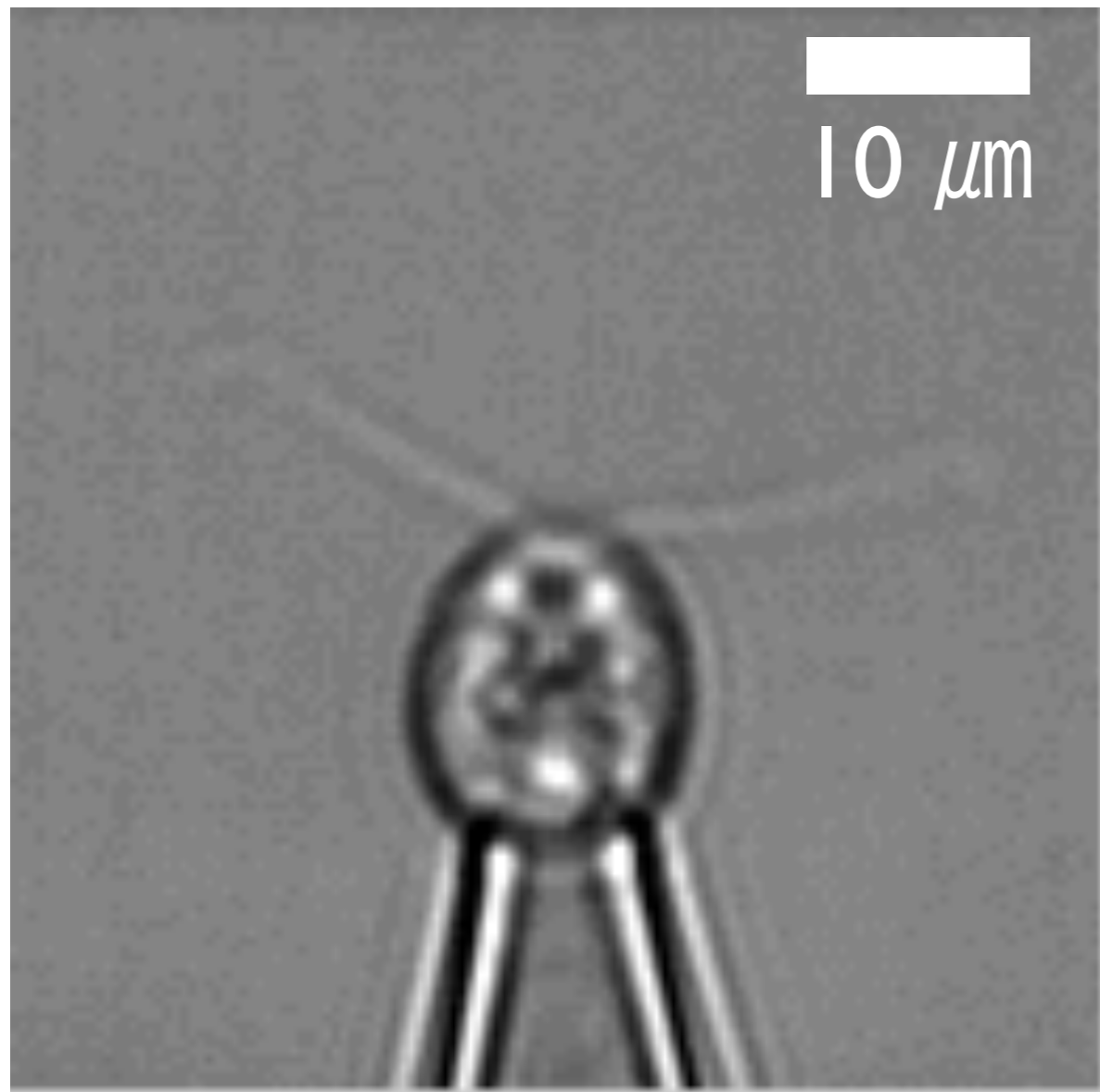
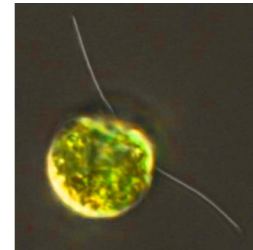


*200 nm fluorescent bead attached to a flagellar motor  
26 steps per revolution  
30x slower than real time  
2400 frames per second  
position resolution ~5 nm*

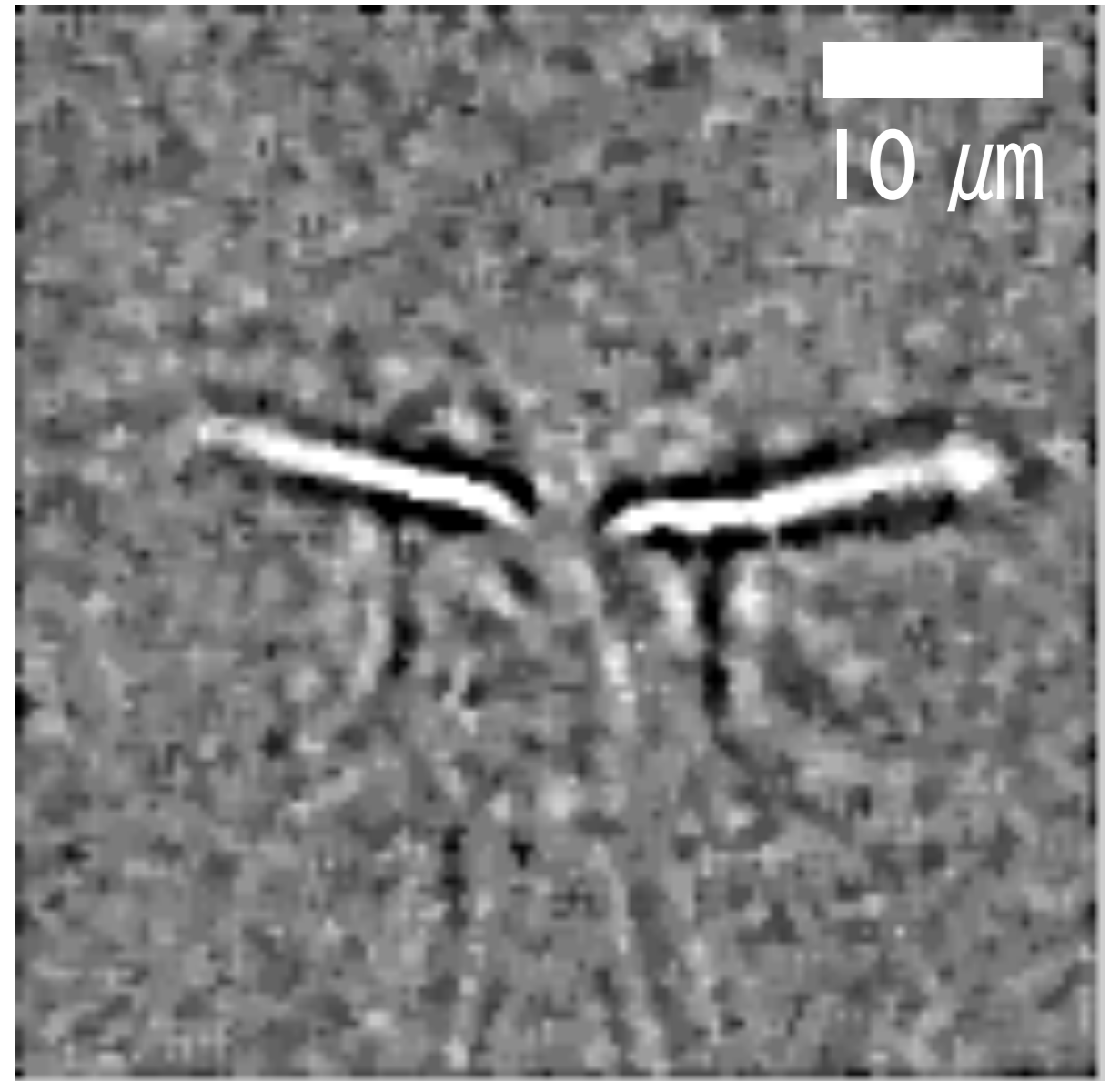


Berry group, Oxford

# *Chlamydomonas* alga

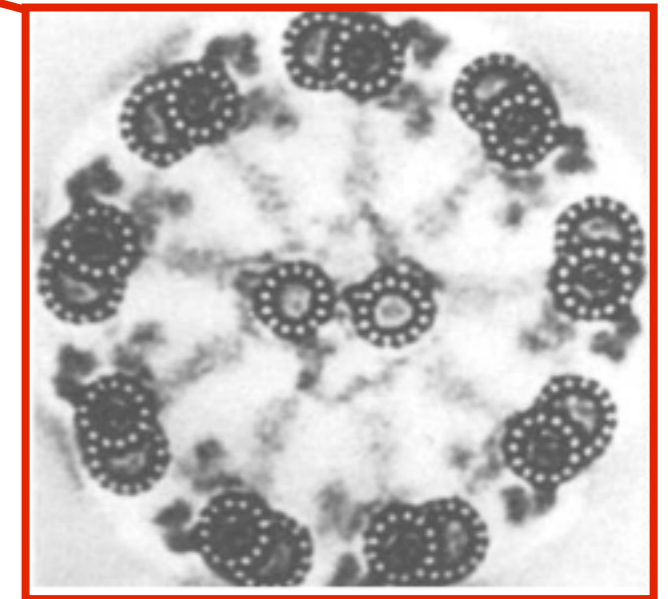
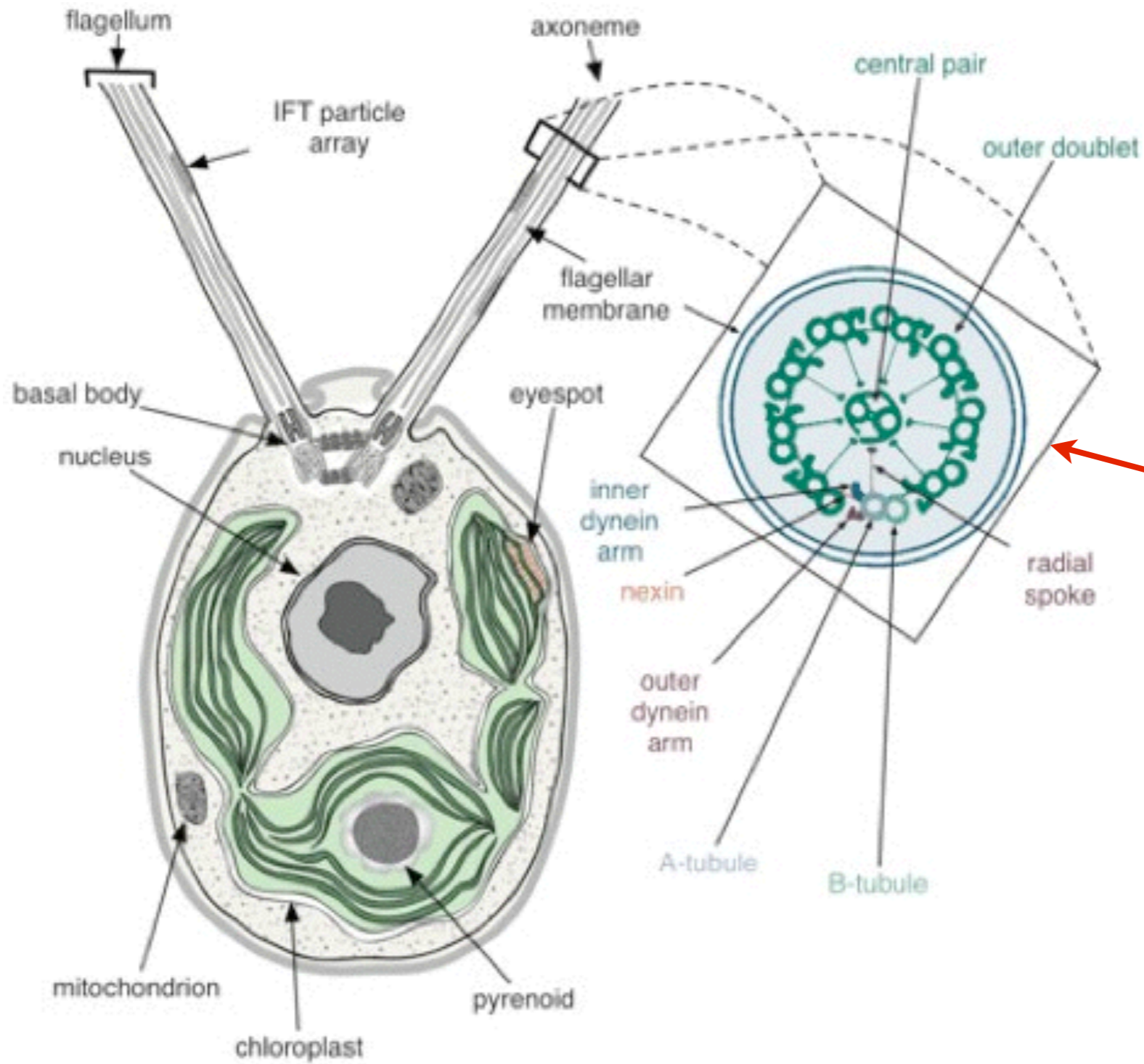
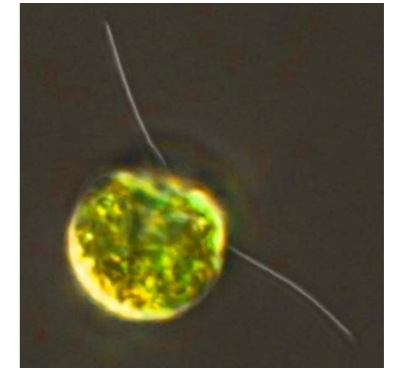


~ 50 beats / sec



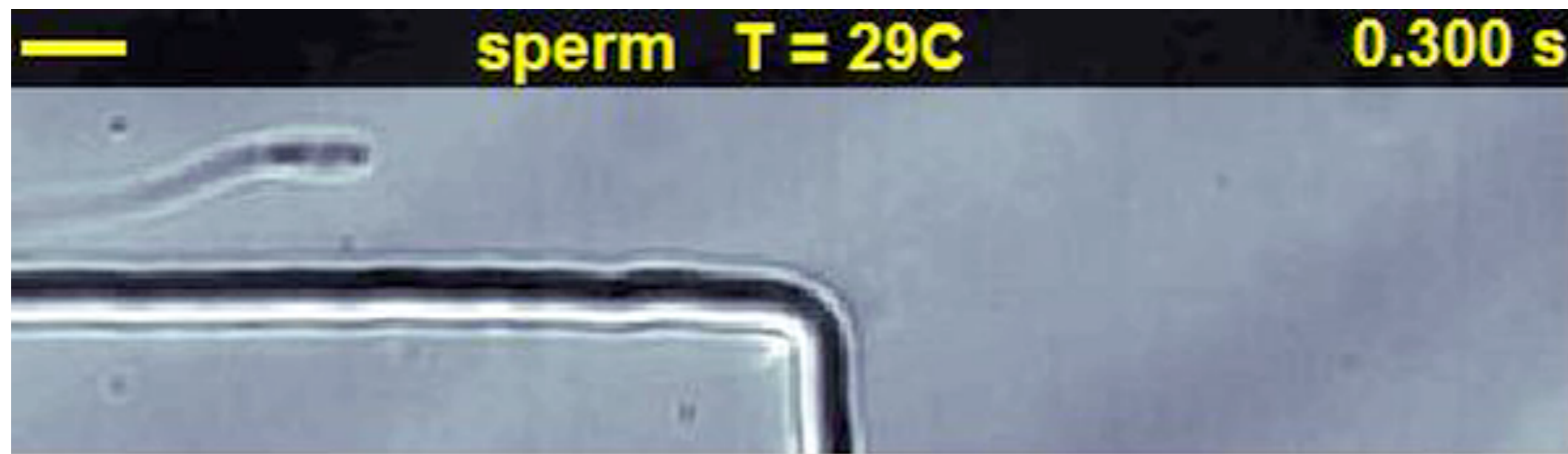
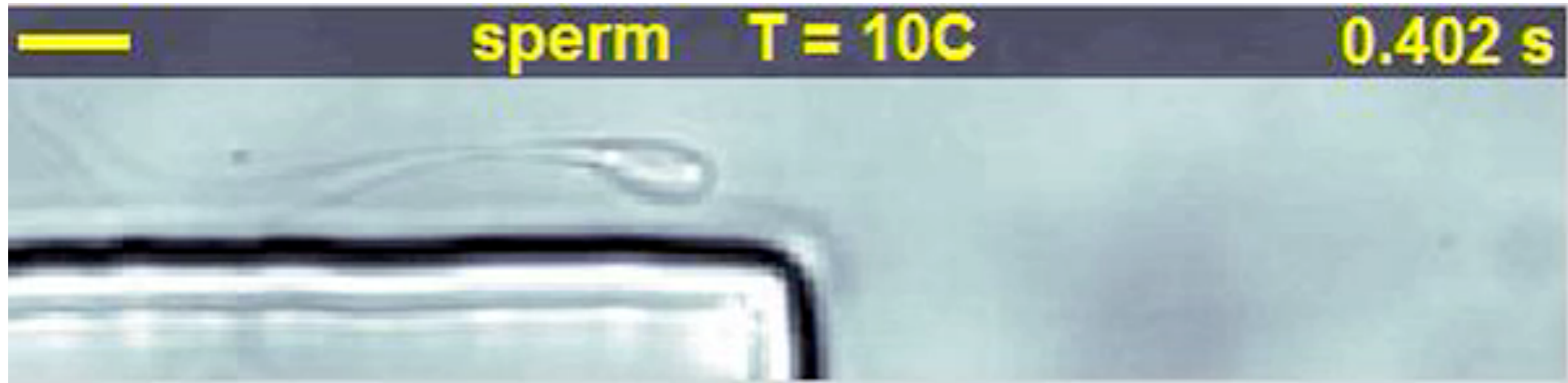
speed ~100 μm/s

# Chlamy

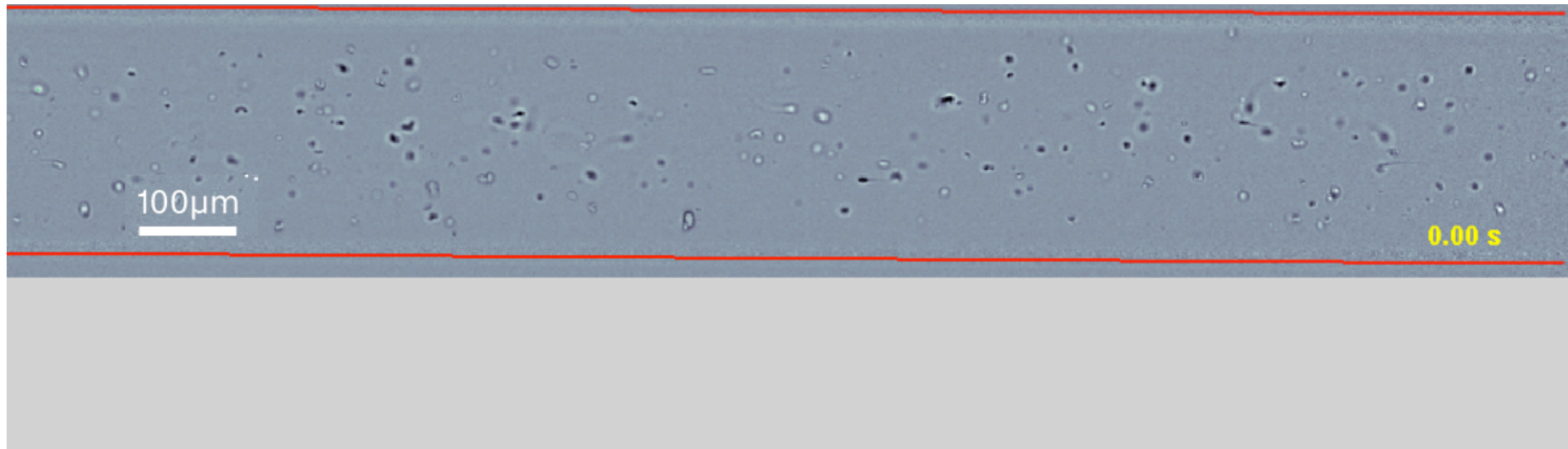




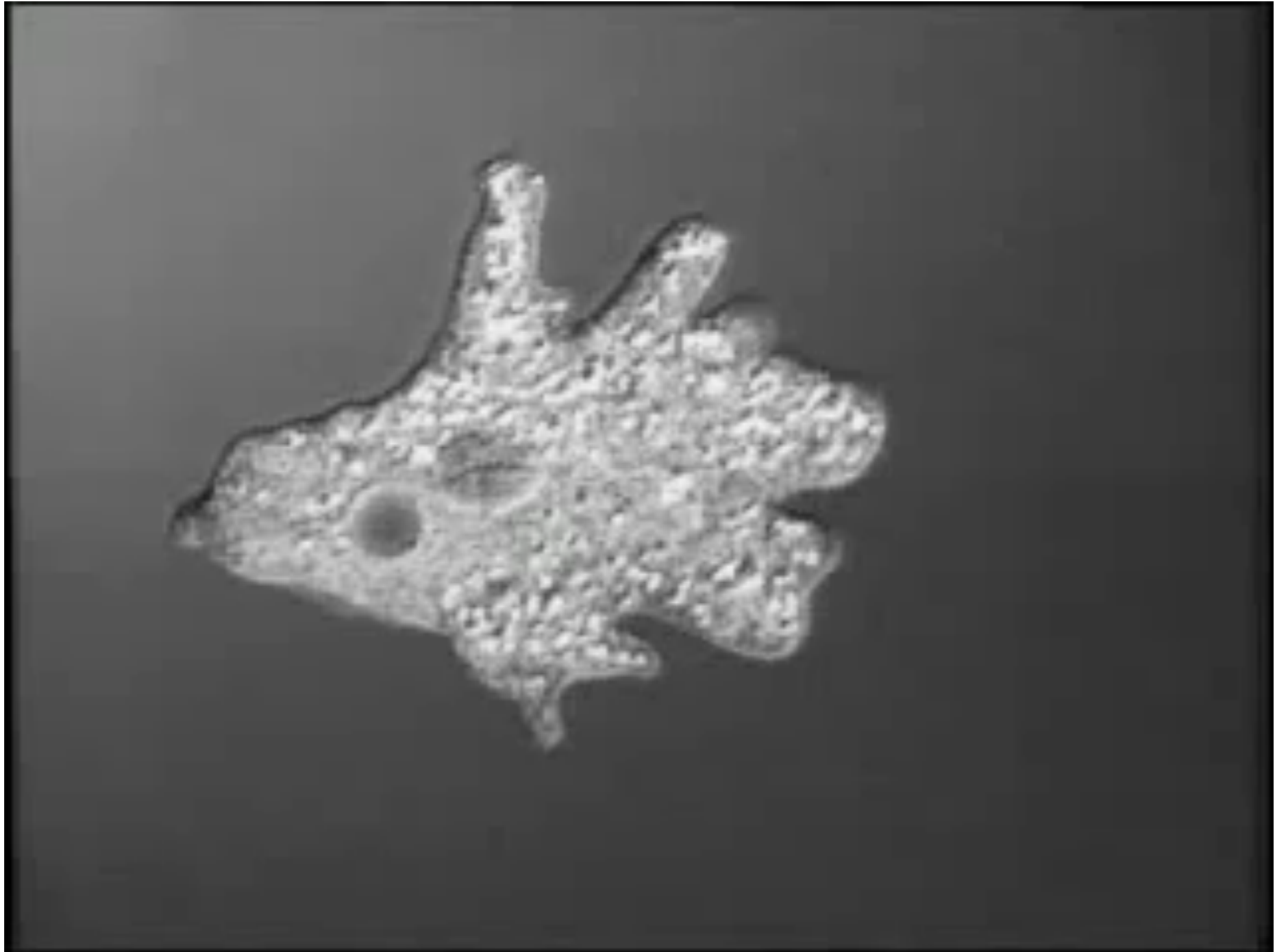
# Sperm near surfaces



# Surface + shear flow

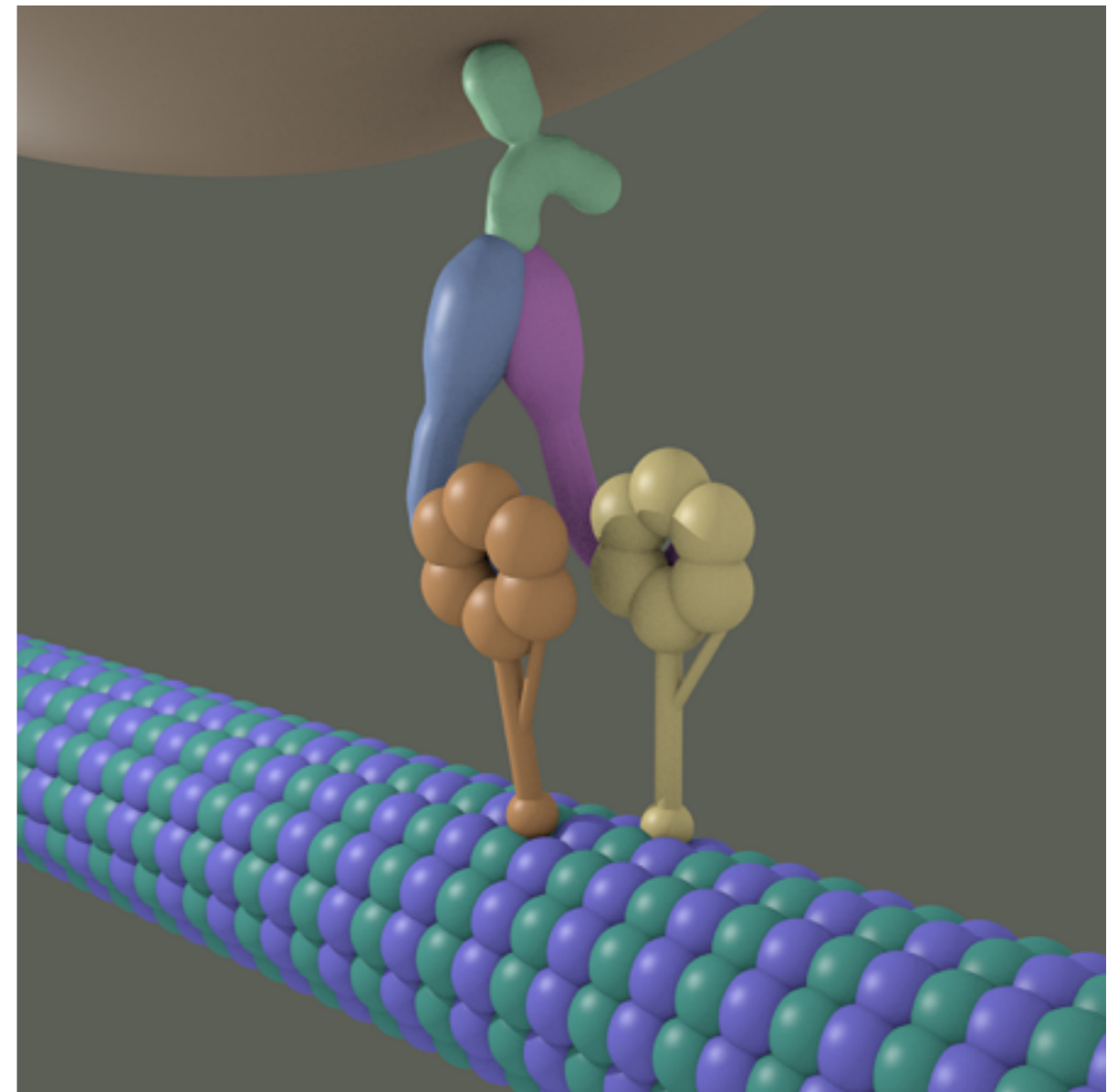
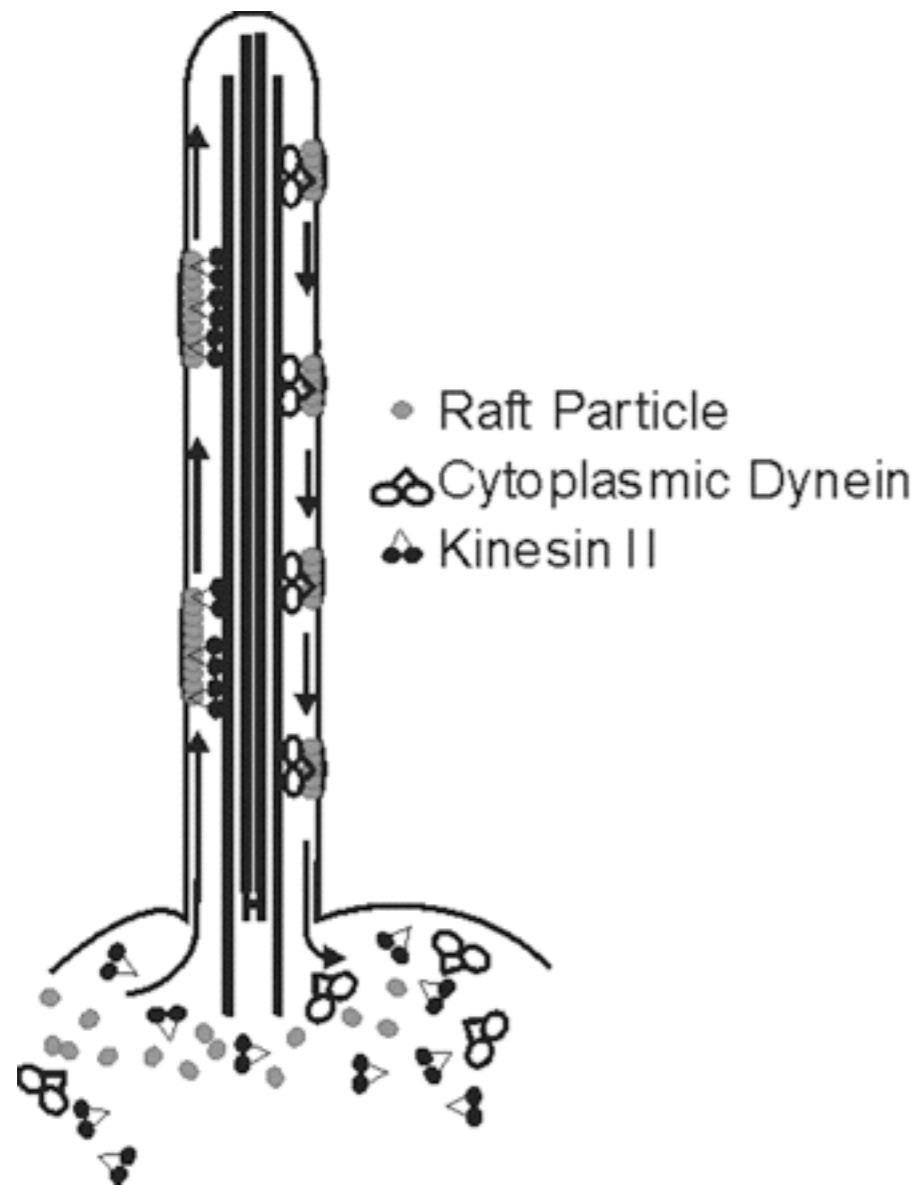


# Amoeba



# Eukaryotic motors

Sketch: dynein molecule carrying cargo down a microtubule

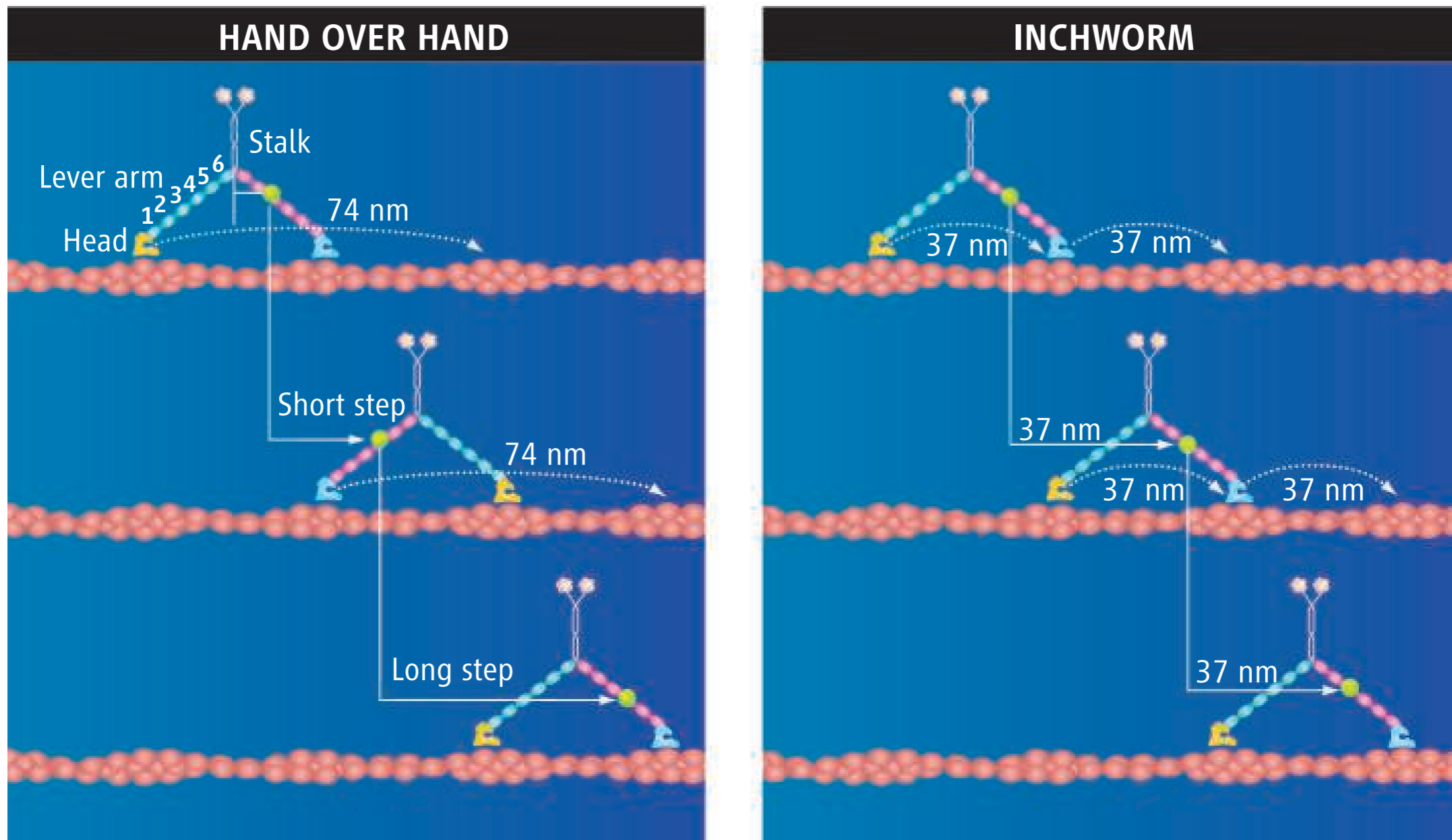


<http://www.plantphysiol.org/content/127/4/1500/F4.expansion.html>

Yildiz lab, Berkeley

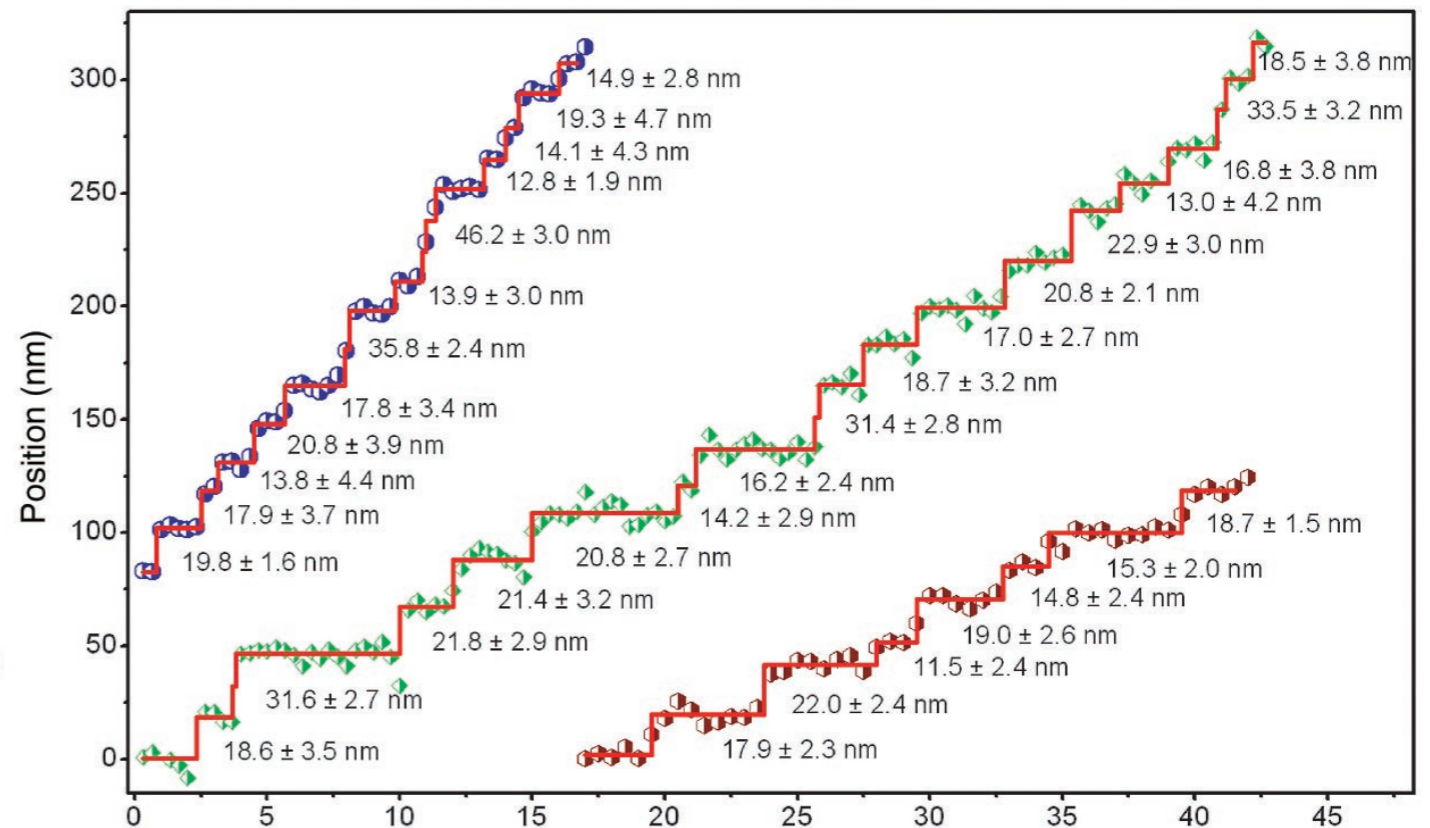
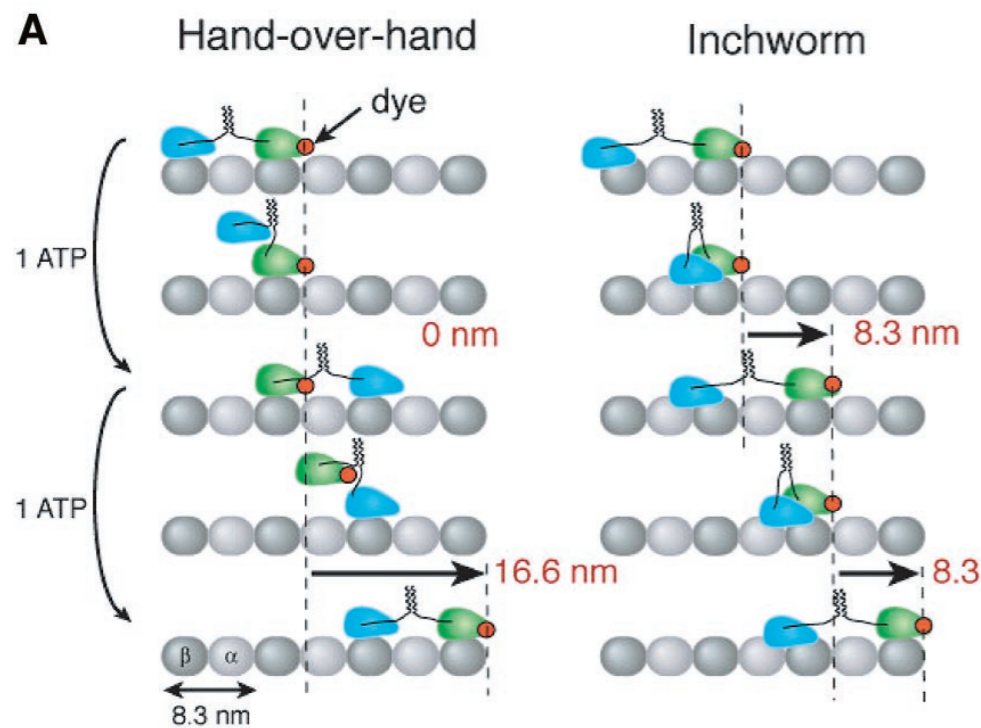
dunkel@math.mit.edu

# Walking modes



**Myosin V: Walking or inchworming?** Predicted movement for the heads and a dye molecule label (green dot) on the lever arm in the hand-over-hand model (**left**) and the inchworm model (**right**). The FIONA assay has revealed that myosin V, along with kinesin and myosin VI, walks hand-over-hand.

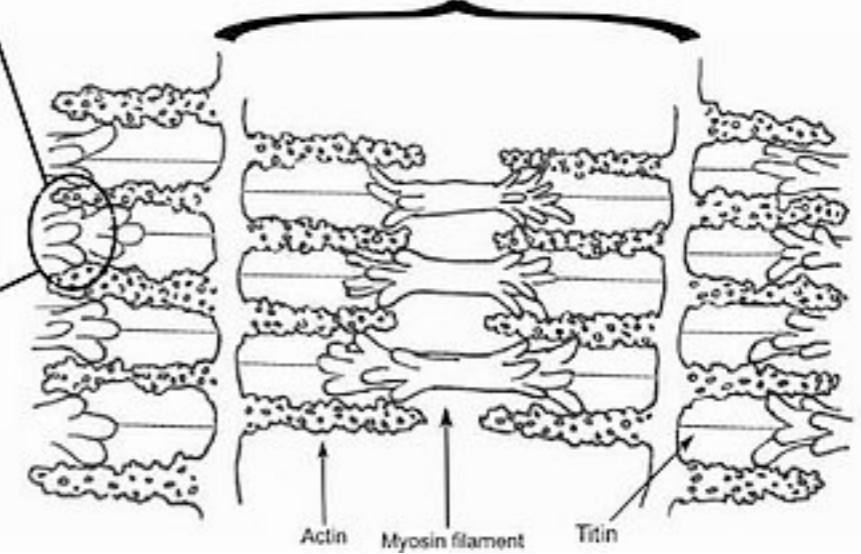
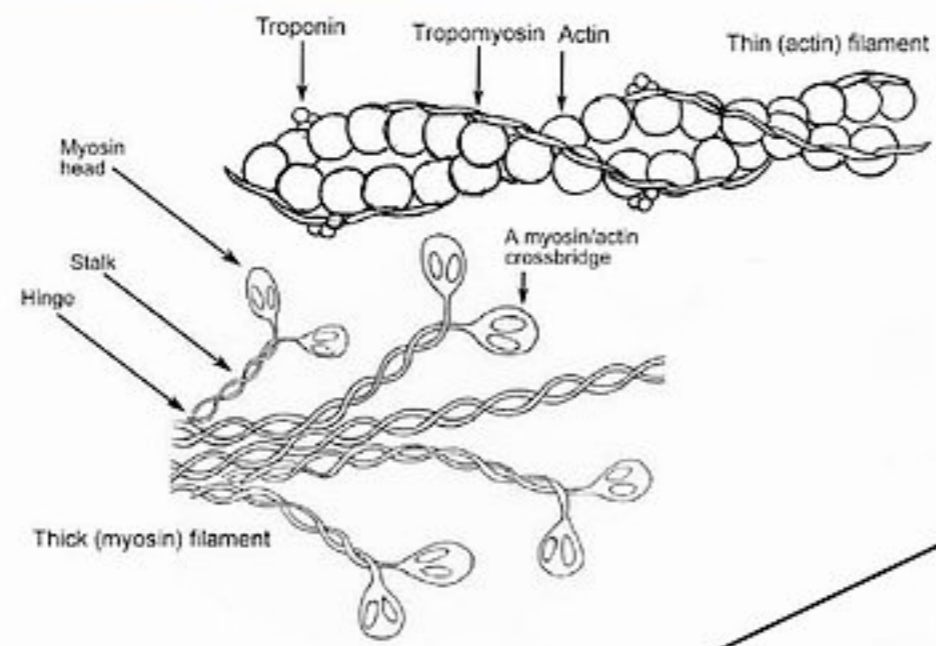
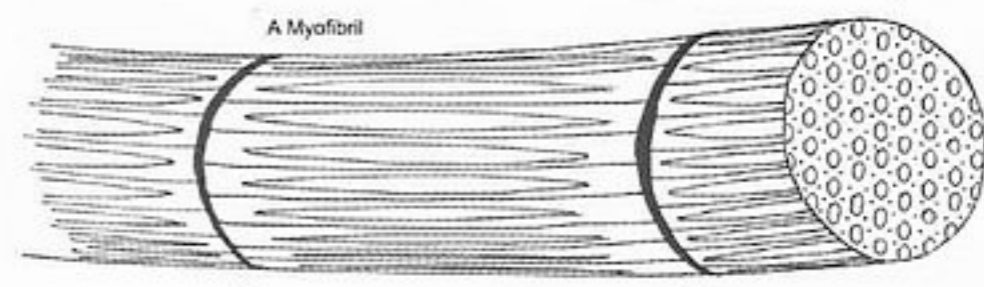
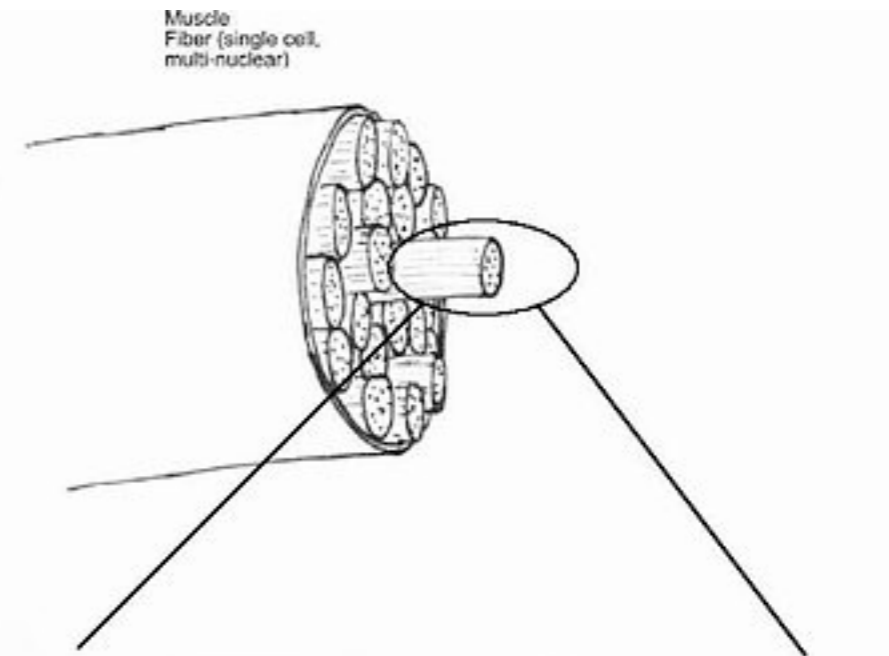
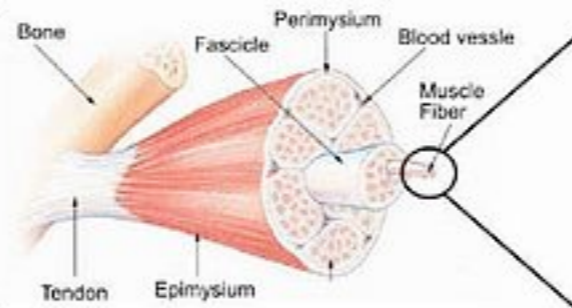
# Kinesin walks hand-over-hand



Yildiz et al (2005) Science

# Intracellular transport



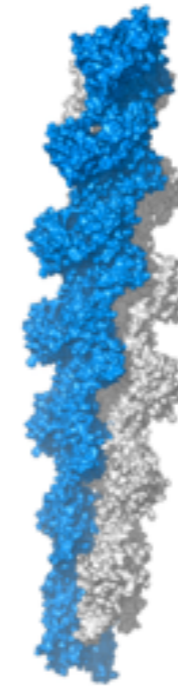
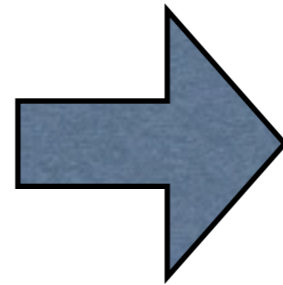
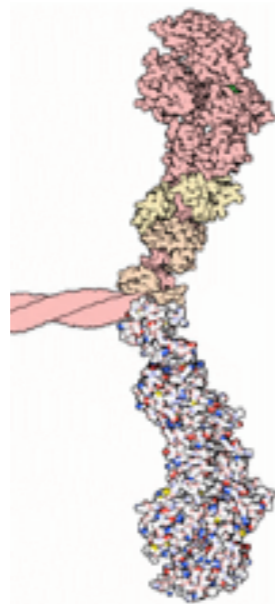


wiki

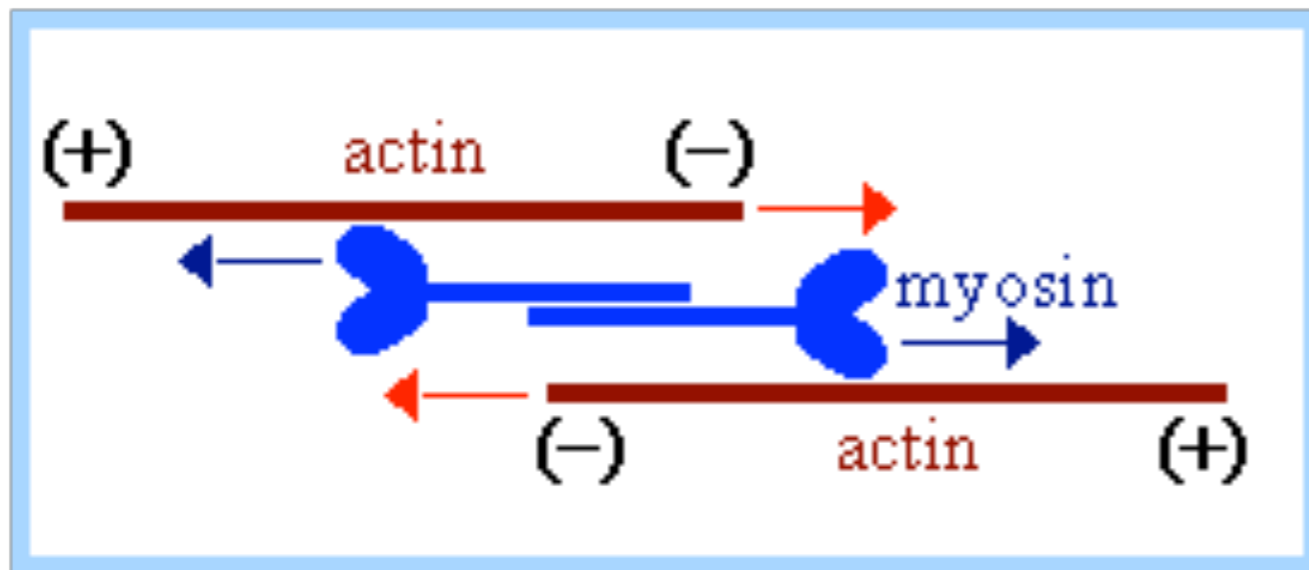


# Actin-Myosin

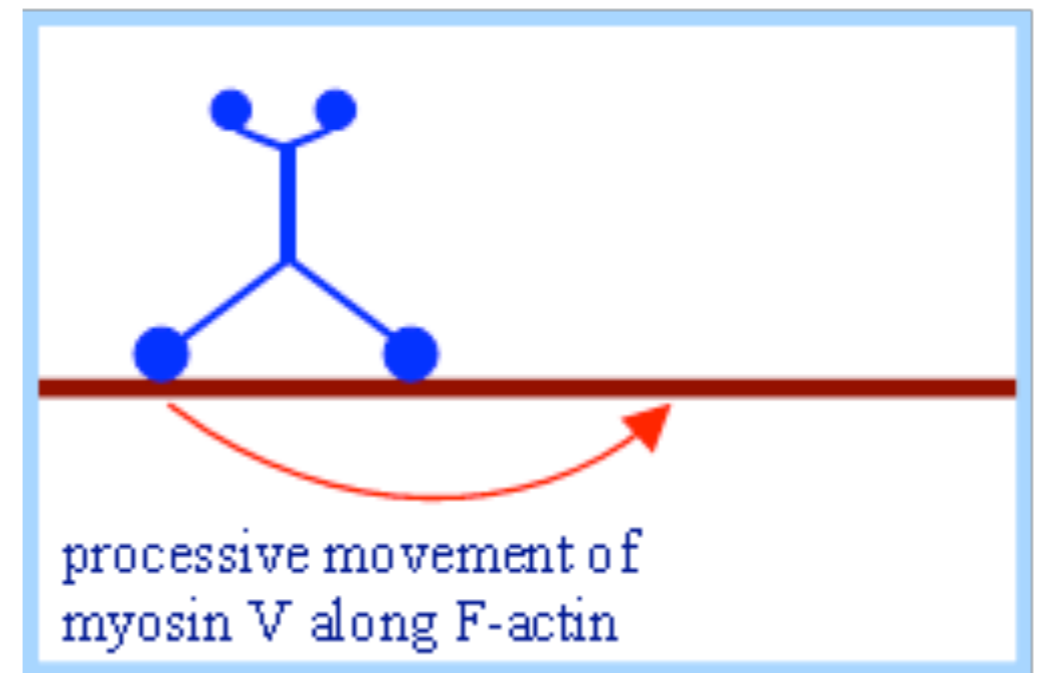
Myosin



F-Actin  
helical filament



myosin-II



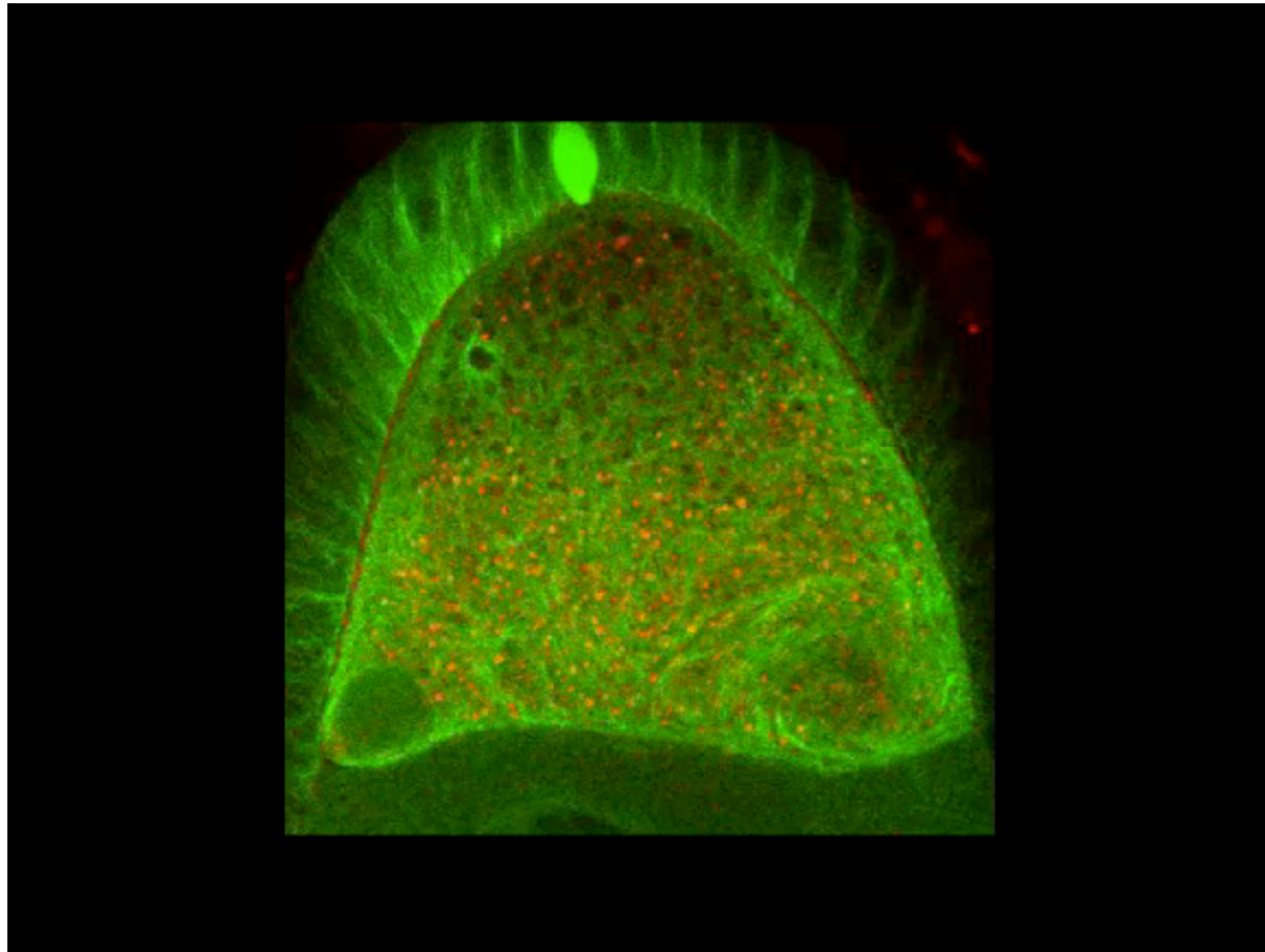
processive movement of  
myosin V along F-actin

myosin-V

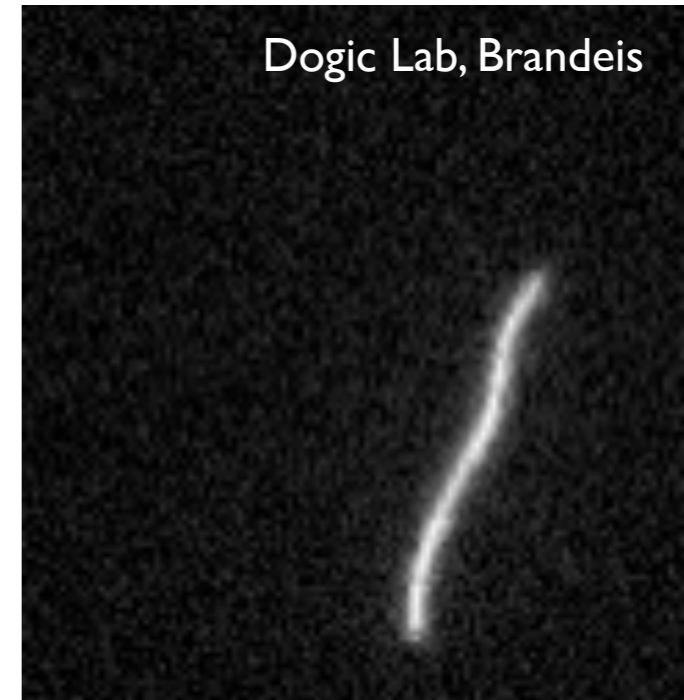
**our lecture course:**

**generic models of  
micro-motors**

# Polymers & filaments ( $D=1$ )



*Drosophila* oocyte

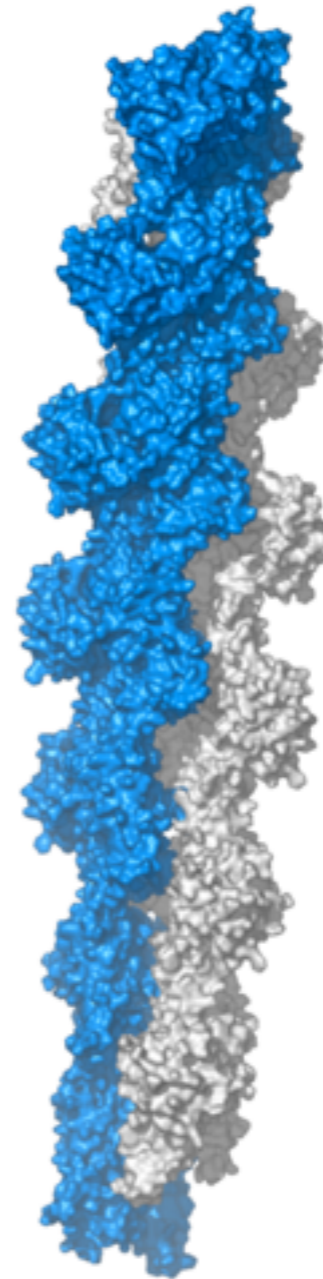


Physical parameters  
(e.g. bending rigidity)  
from fluctuation  
analysis

# Actin in 2D



Dogic Lab (Brandeis)

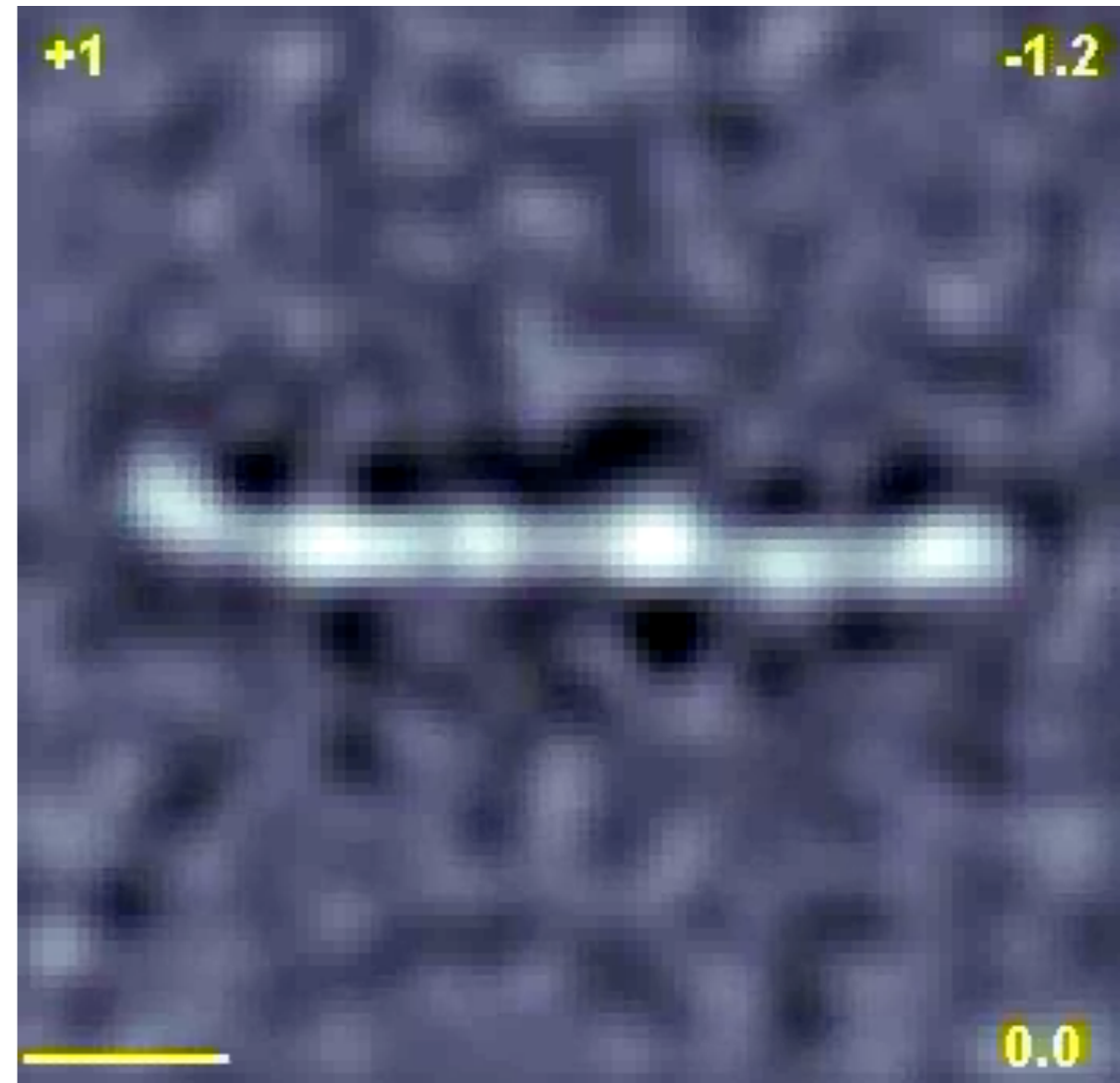
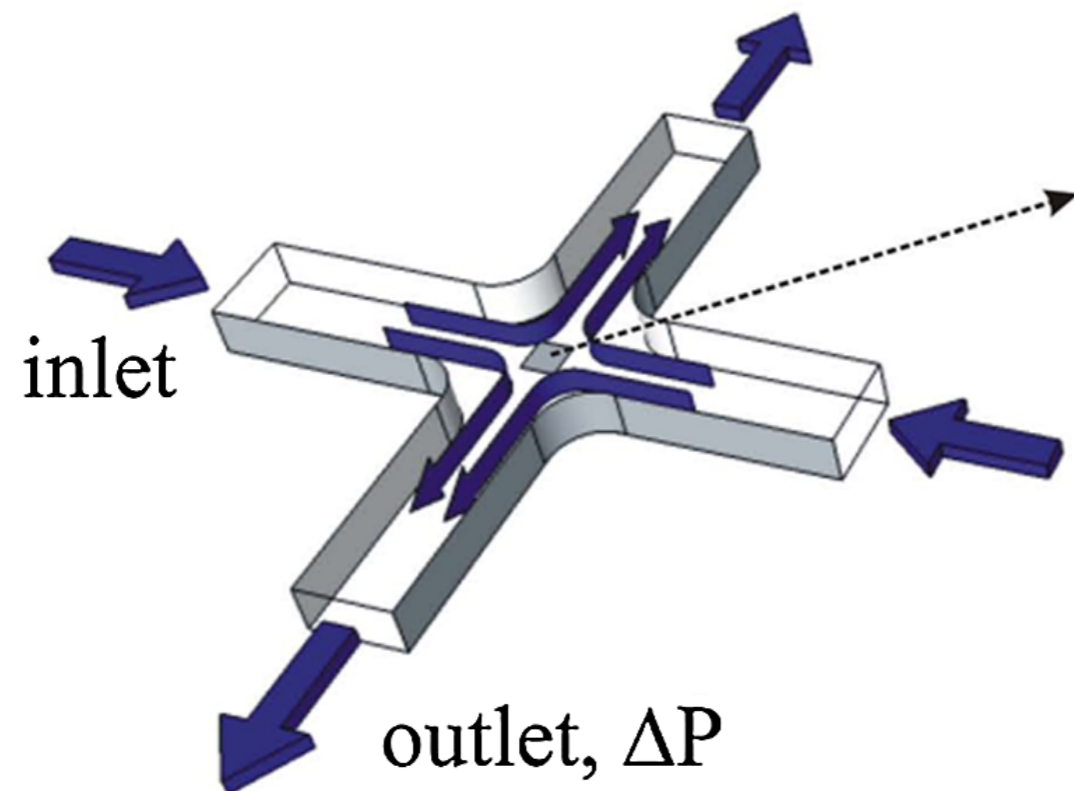


F-Actin

helical  
filament

# Actin in flow

(a)

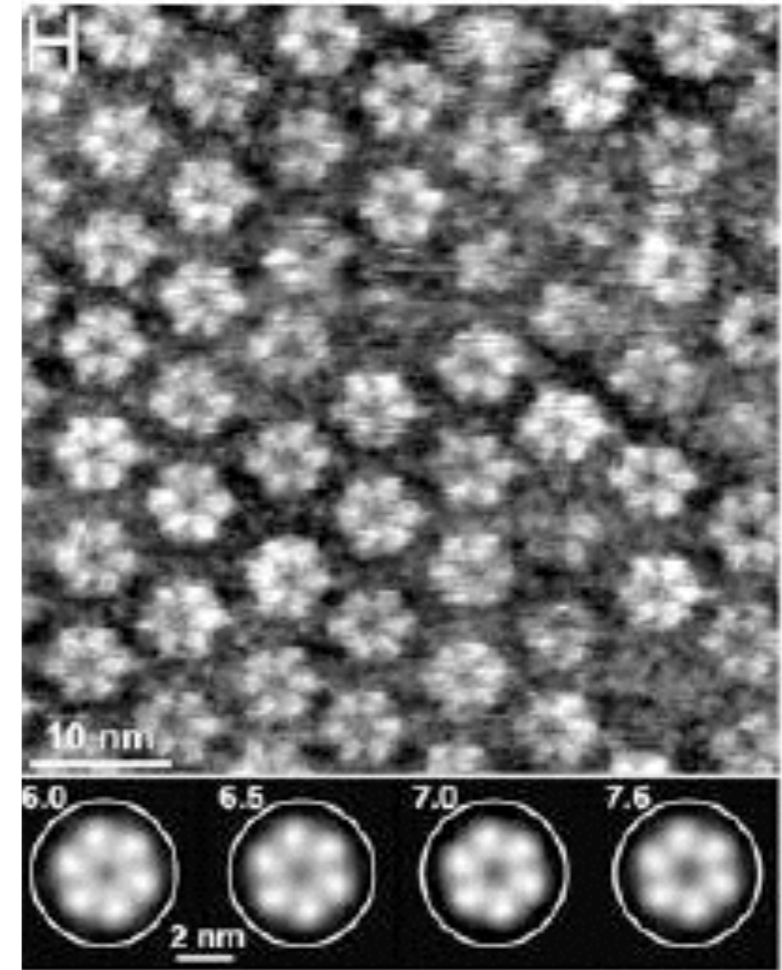
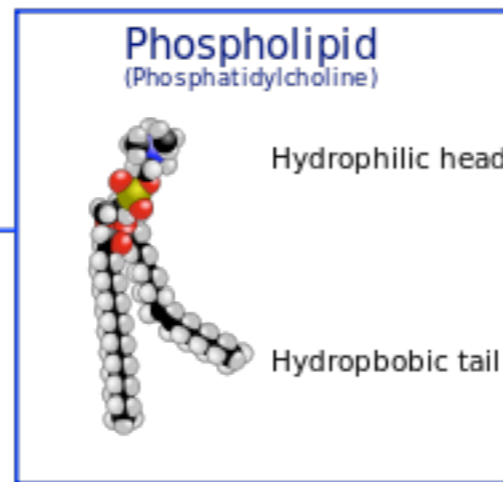
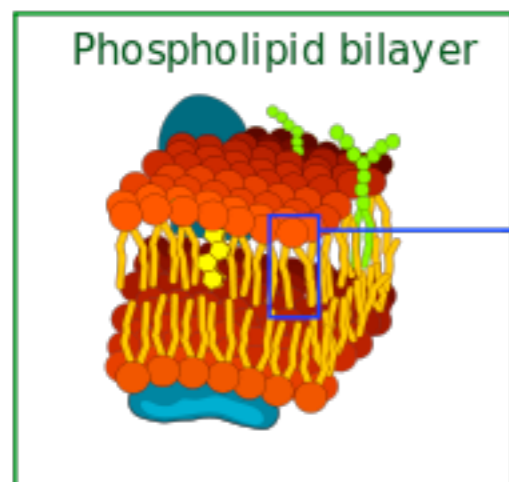
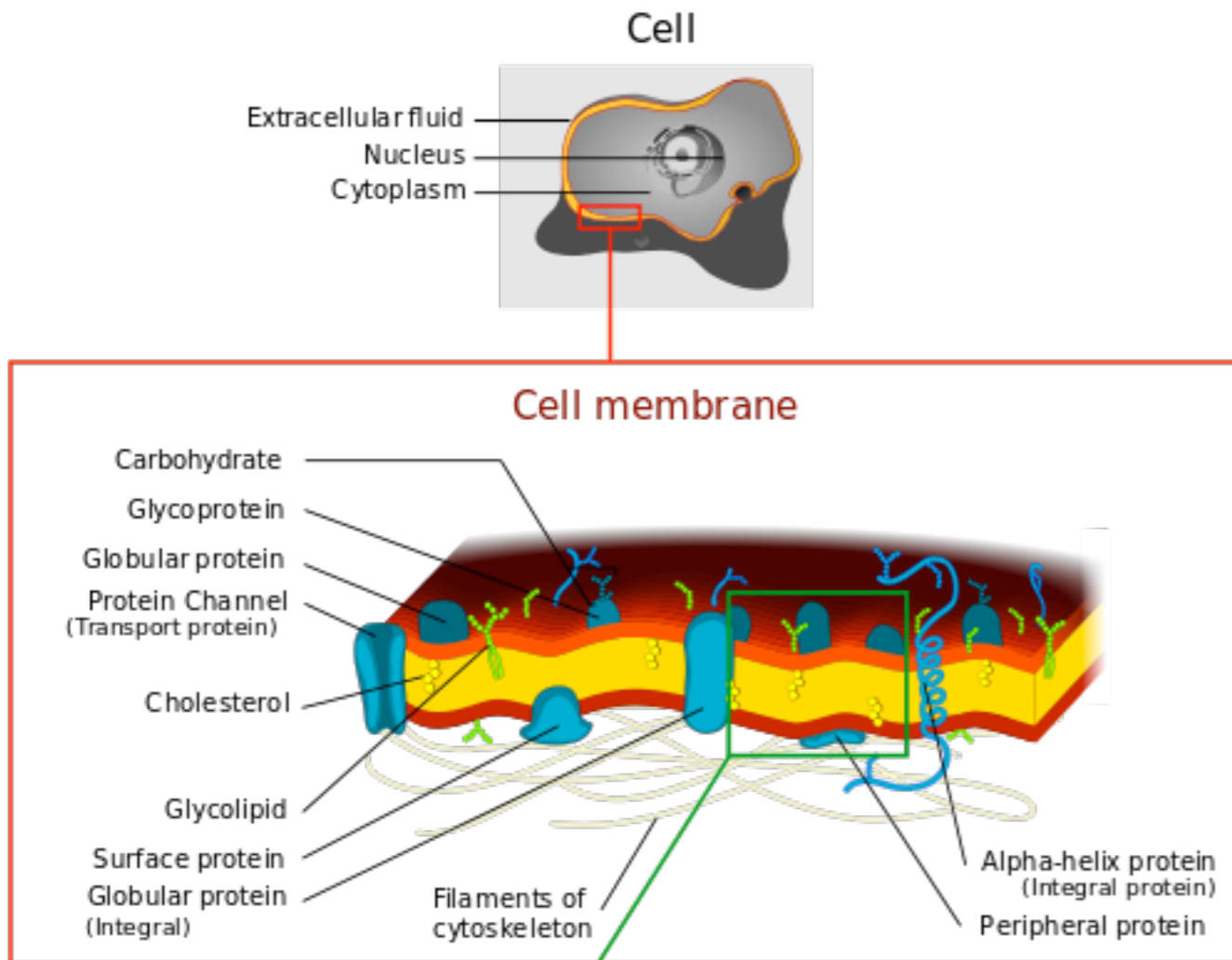


Kantsler & Goldstein (2012) PRL

**our lecture course:**

- **polymer models**
- **how to relate fluctuations to mechanical properties**

# Cell membranes (D=2)

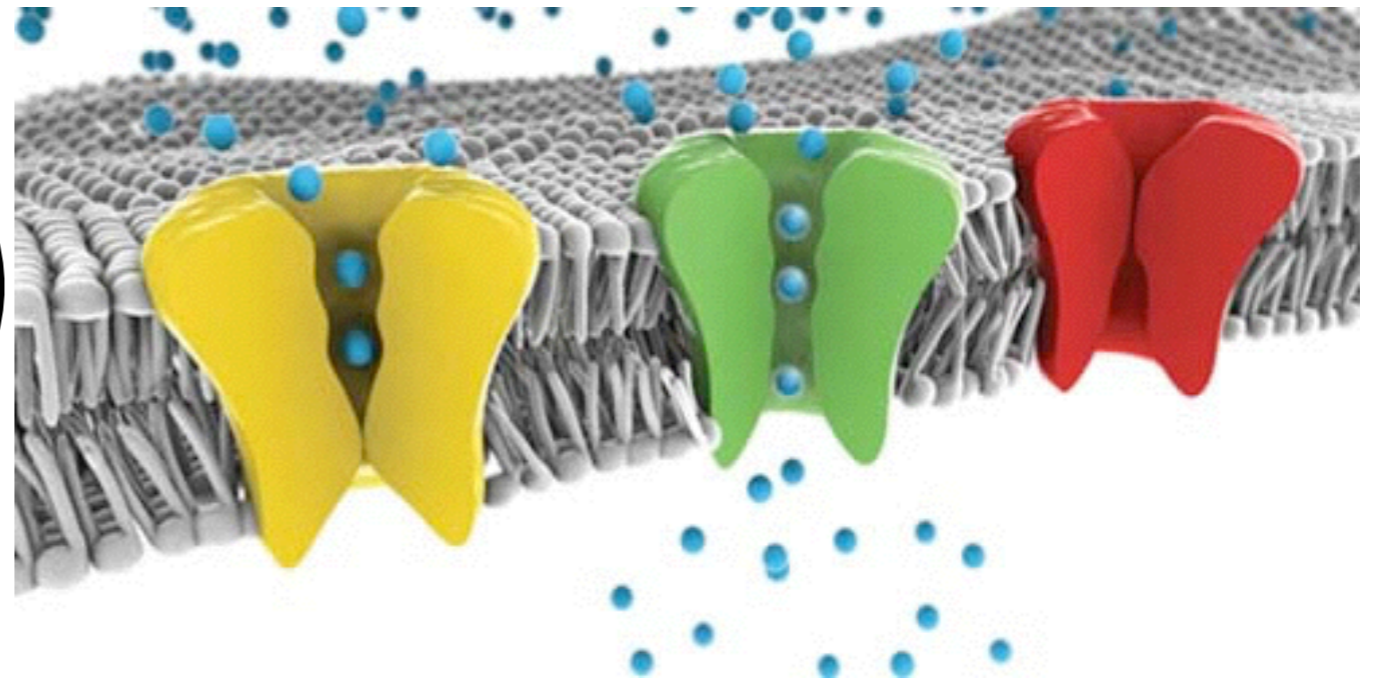


[http://www.sbmp-itn.eu/sbmpr/research\\_method/](http://www.sbmp-itn.eu/sbmpr/research_method/)

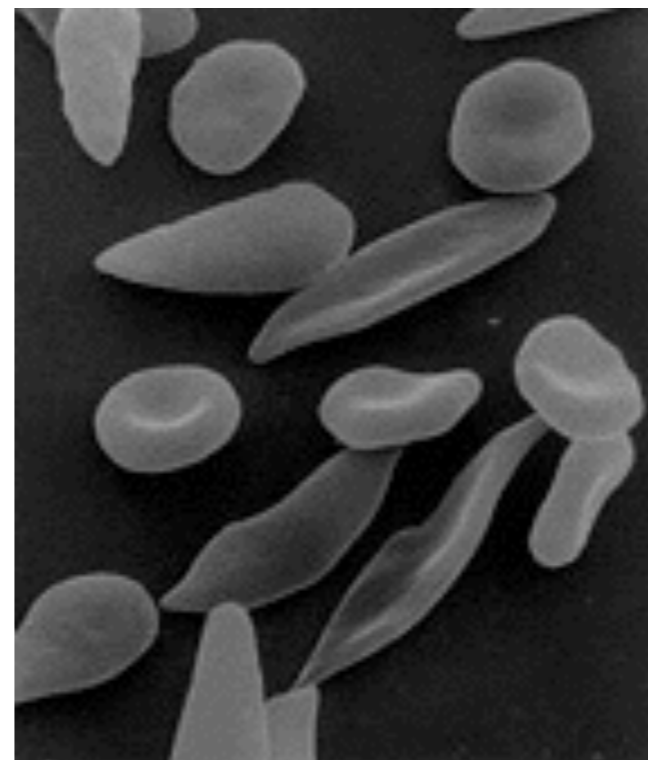
# Cell membranes ( $D=2$ )

Illustration by J.P. Cartailier. Copyright 2007, Symmation LLC.

**transport:**  
stochastic  
escape problems



**shape:**  
differential  
geometry



red blood cells  
affected by  
sickle-cell disease

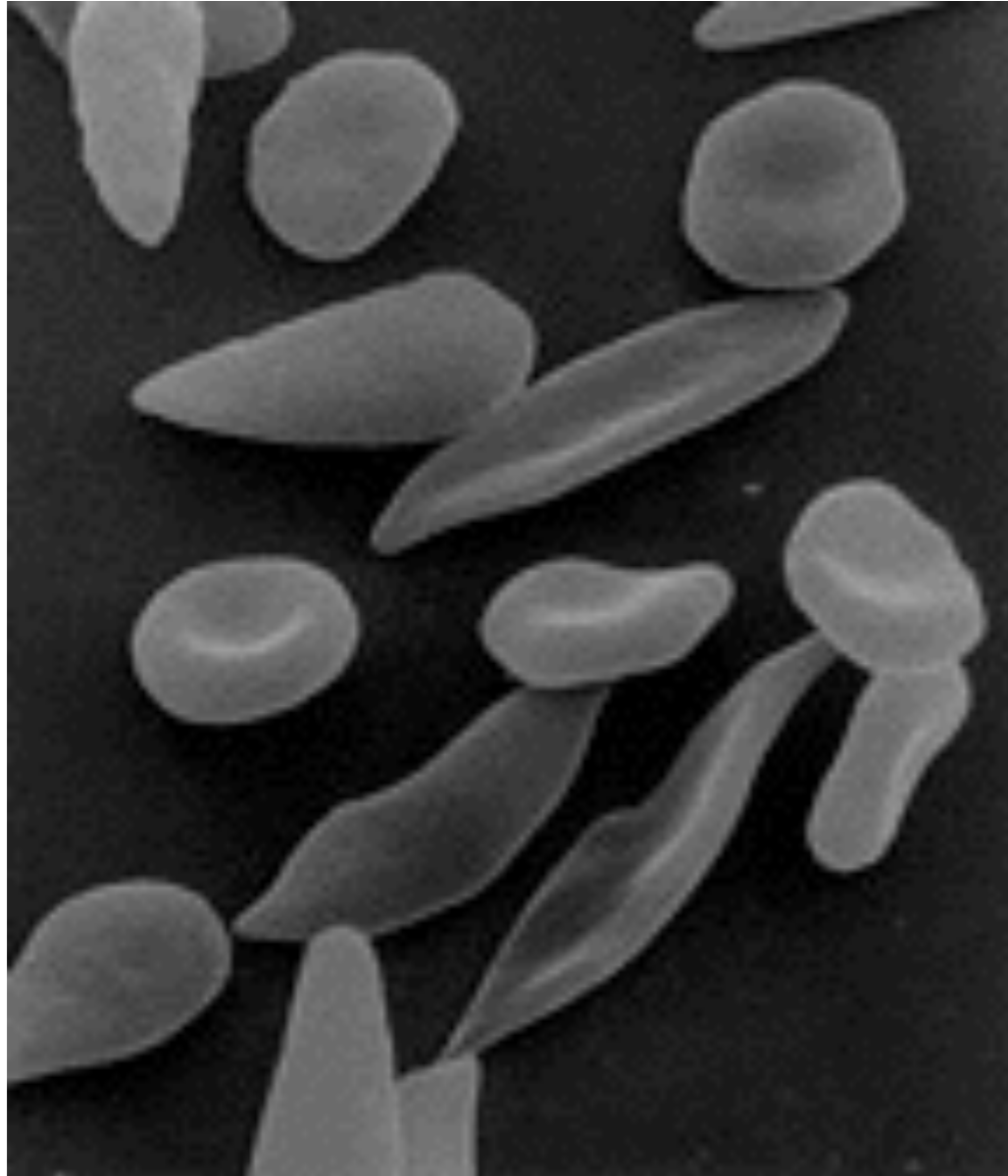
source: wiki

dunkel@math.mit.edu

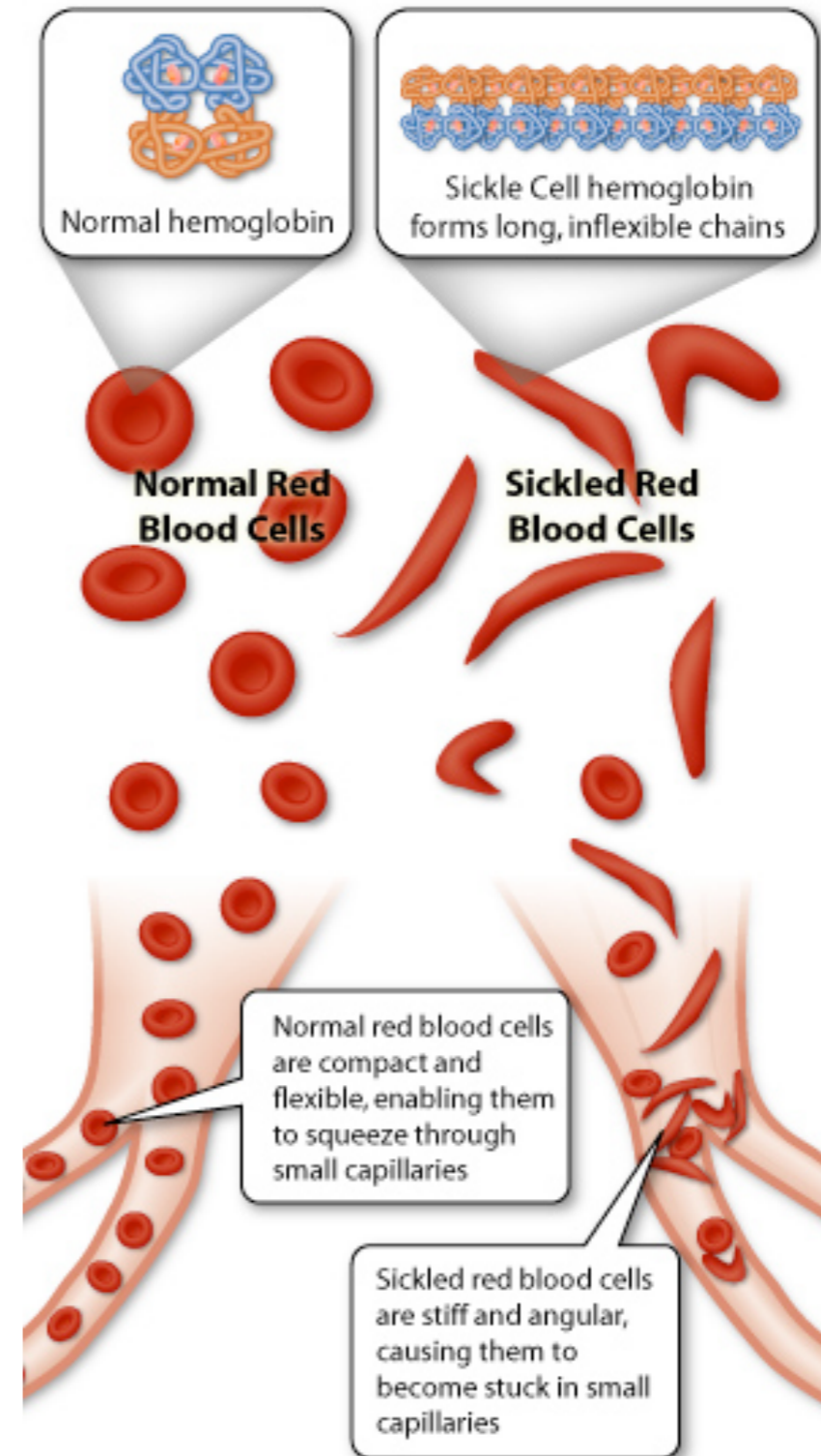


# Blood cells: shape & function

source: wiki

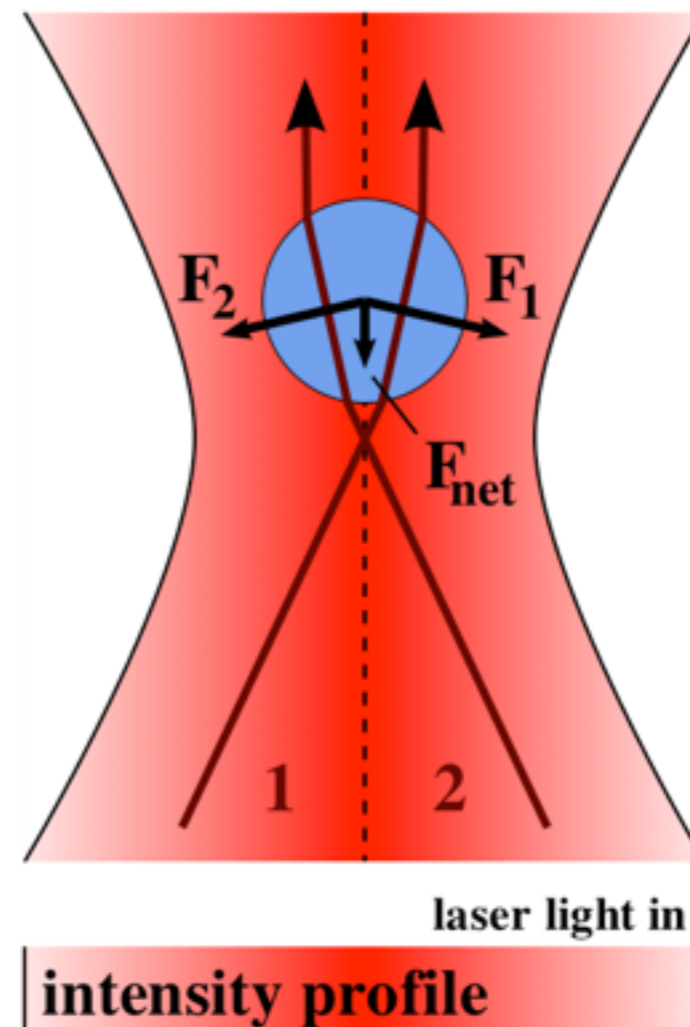
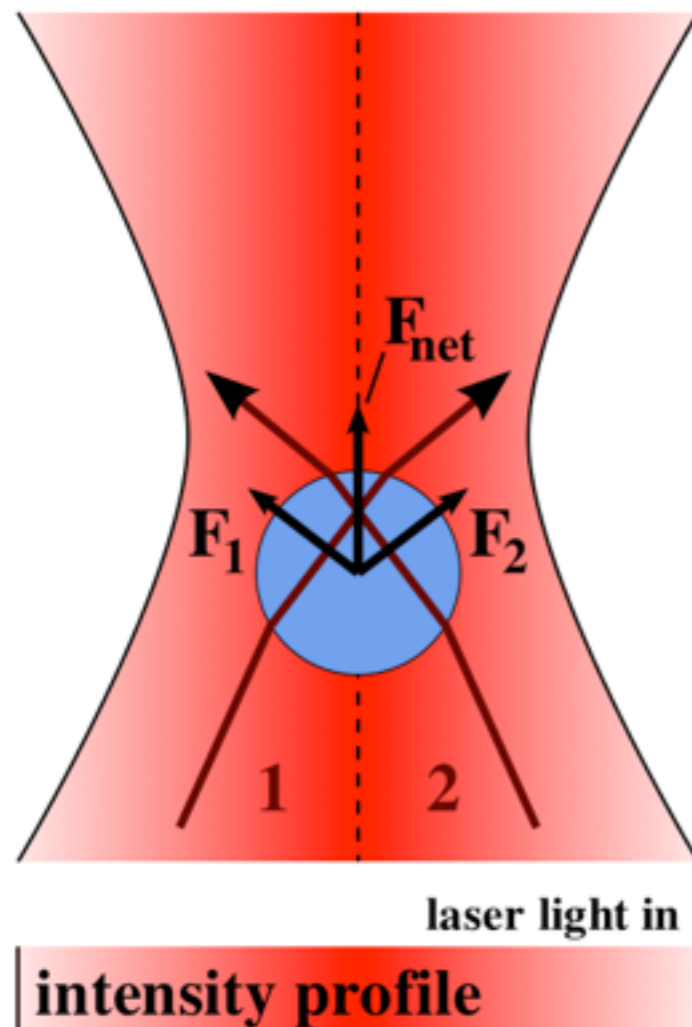


red blood cells  
affected by sickle-  
cell disease



# Optical tweezer

source: wiki



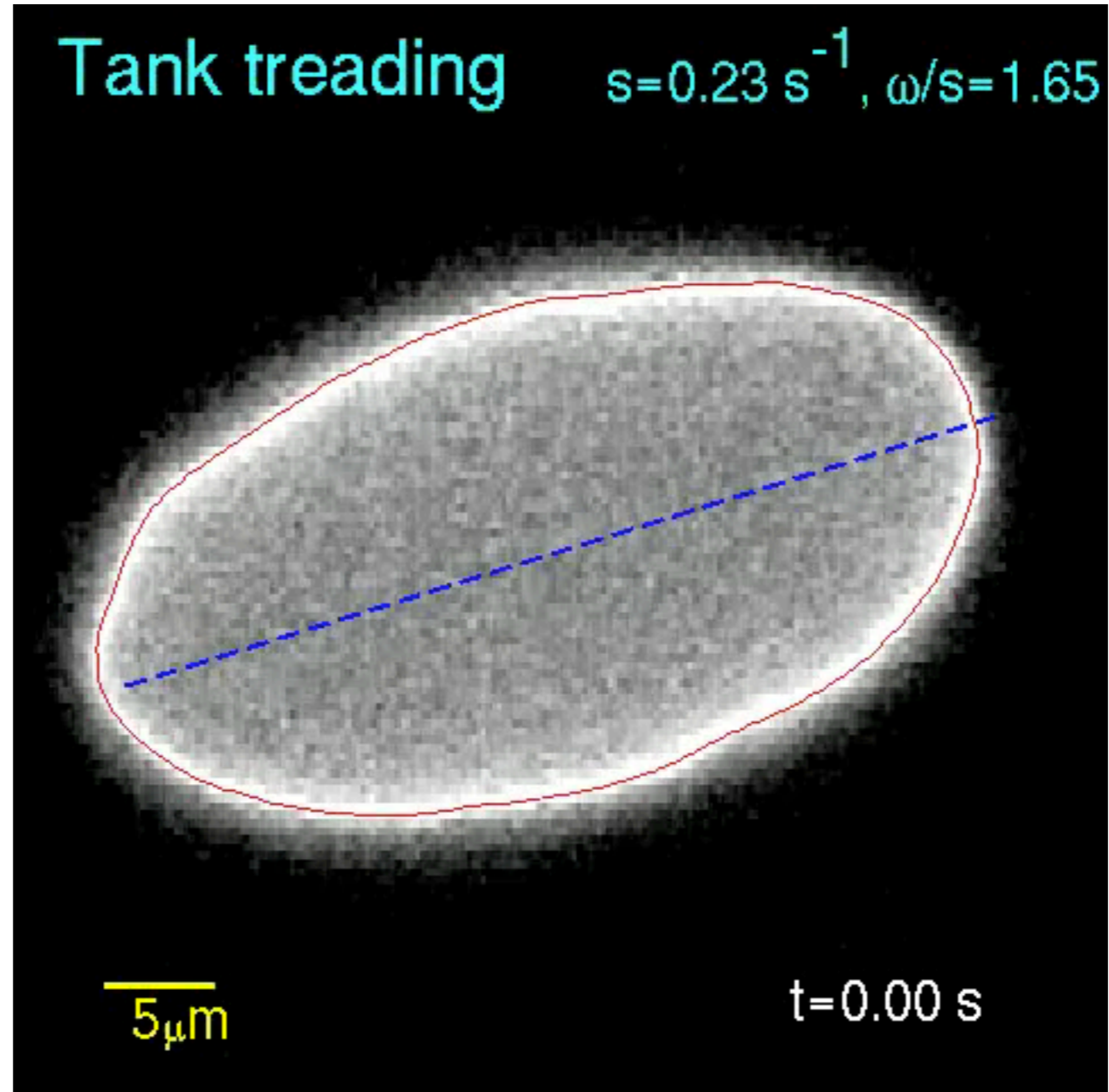
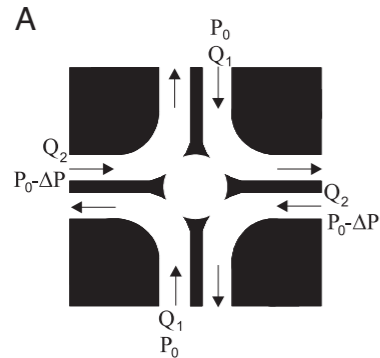
<http://www.nature.com/ncomms/journal/v4/n4/extref/ncomms2786-s1.swf>

# Dynamics of a vesicle in general flow

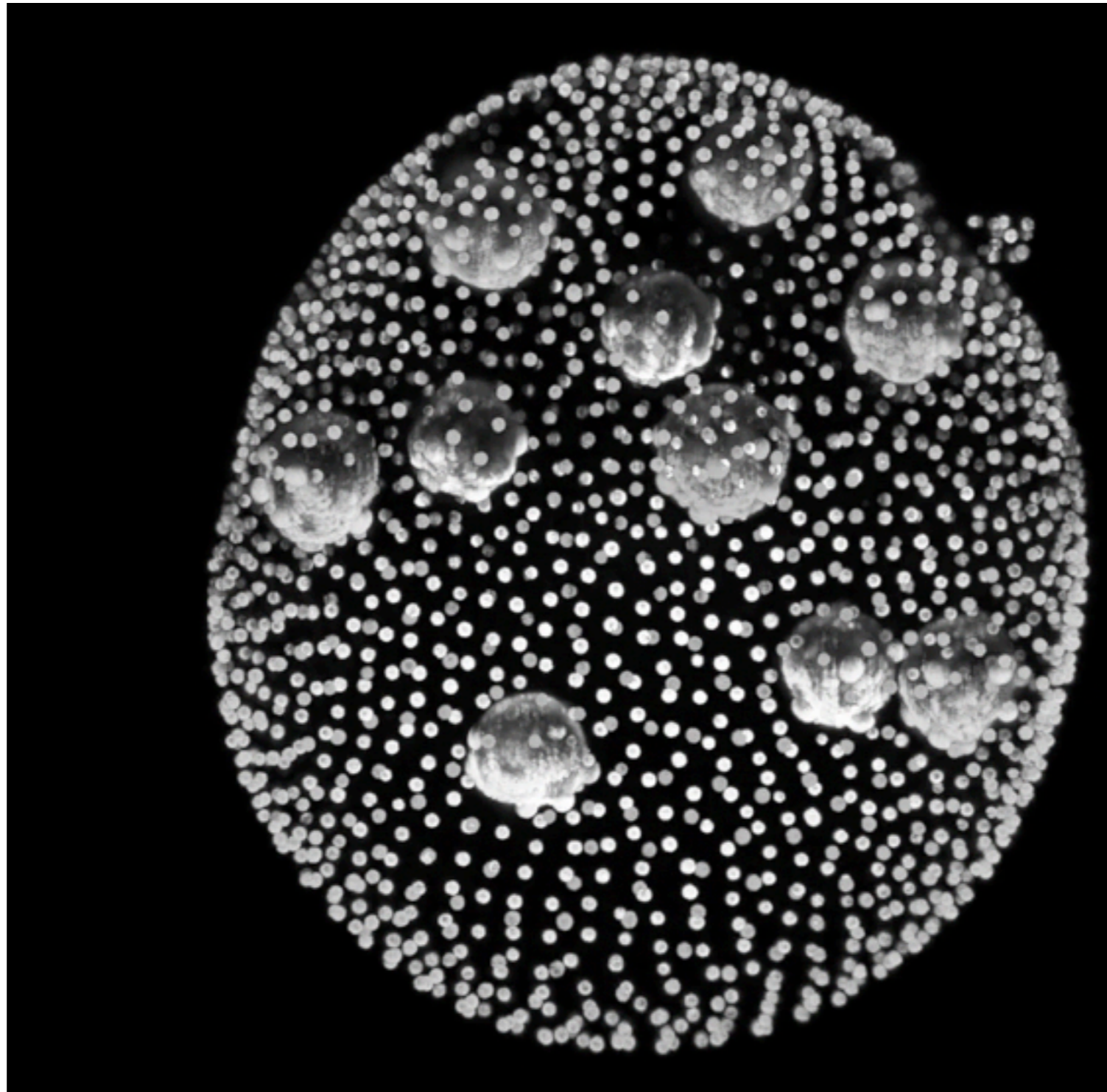
J. Deschamps, V. Kantsler, E. Segre, and V. Steinberg<sup>1</sup>

Department of Physics of Complex Systems, Weizmann Institute of Science, Rehovot, 76100 Israel

11444–11447 | PNAS | July 14, 2009 | vol. 106 | no. 28



# Volvox inversion



**our lecture course:**

- **‘differential geometry’ of membranes**

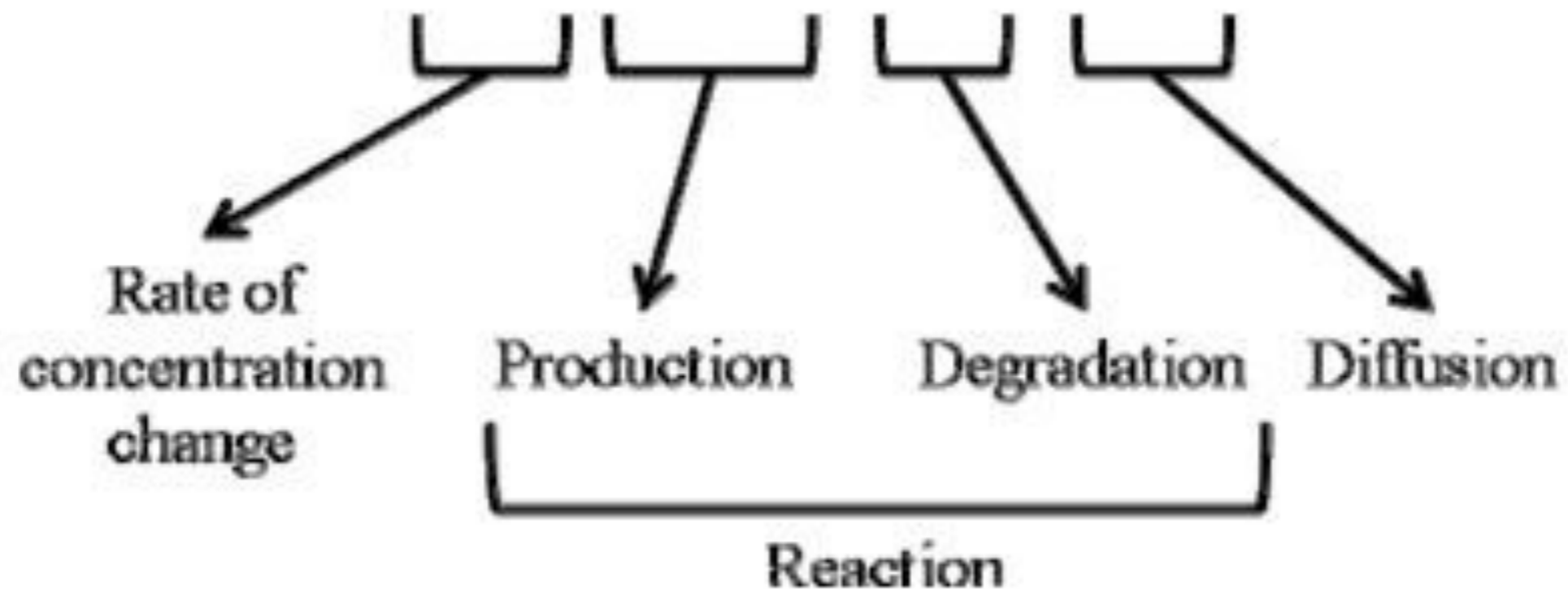
# Stationary patterns



# Turing model

$$\frac{\partial u}{\partial t} = F(u, v) - d_u v + D_u \Delta u$$

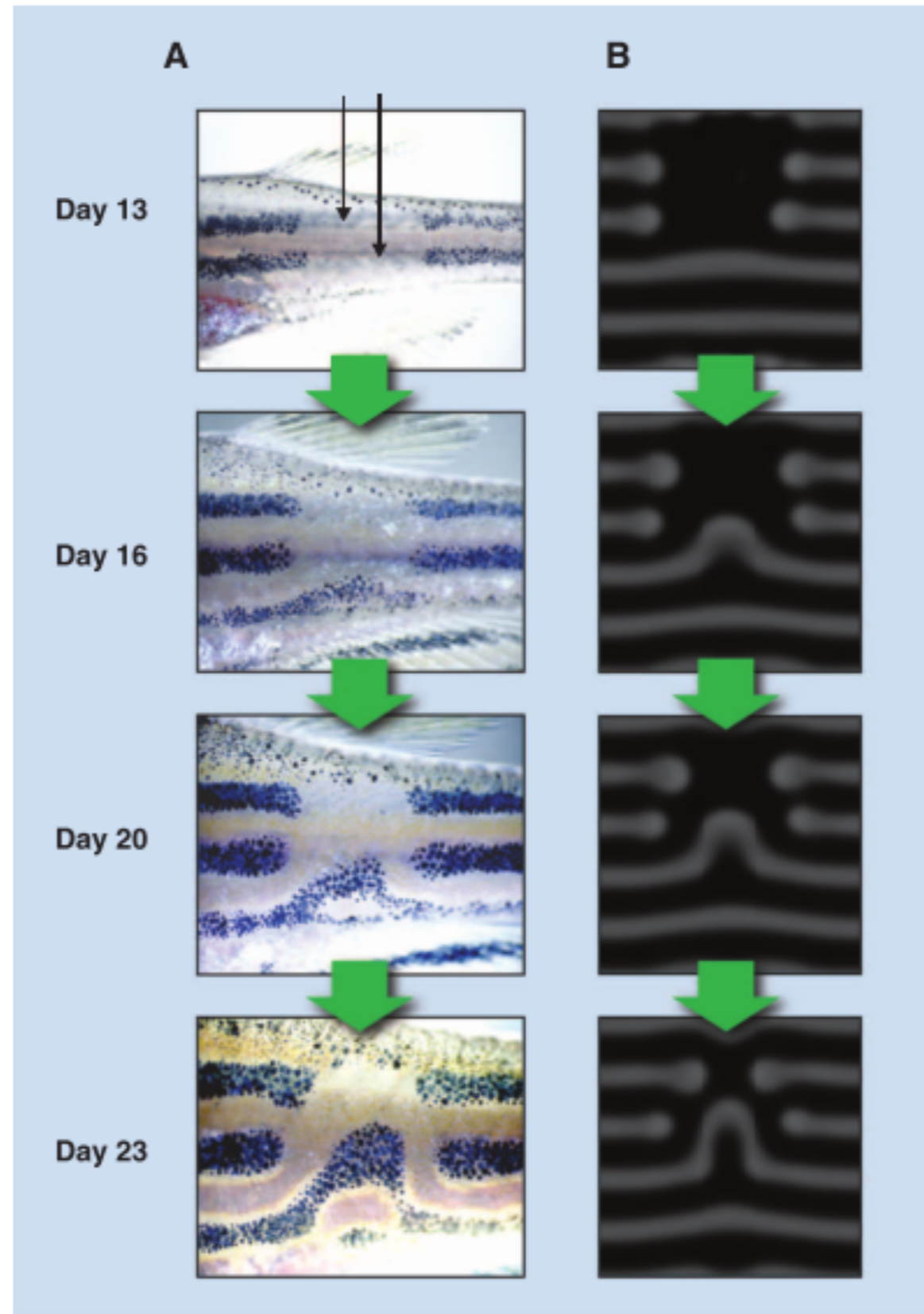
$$\frac{\partial v}{\partial t} = G(u, v) - d_v v + D_v \Delta v$$



wiki



# The matching of zebrafish stripe formation and a Turing model



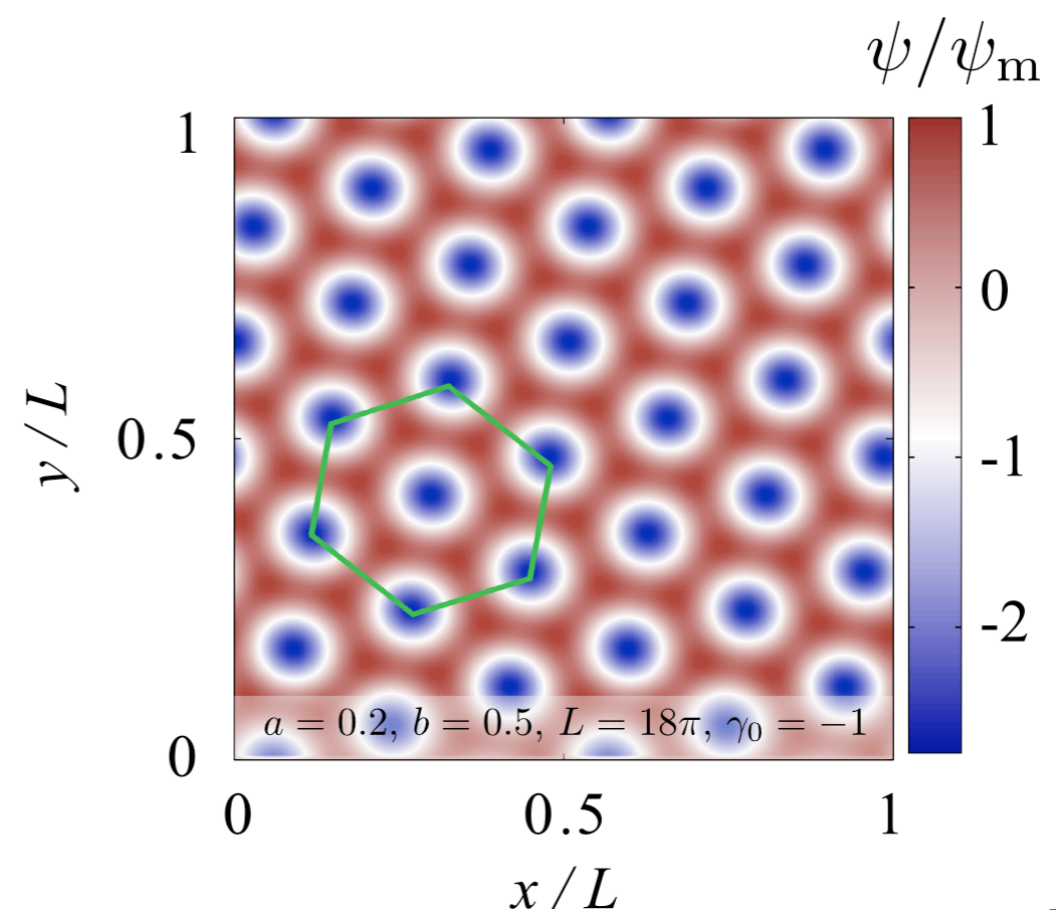
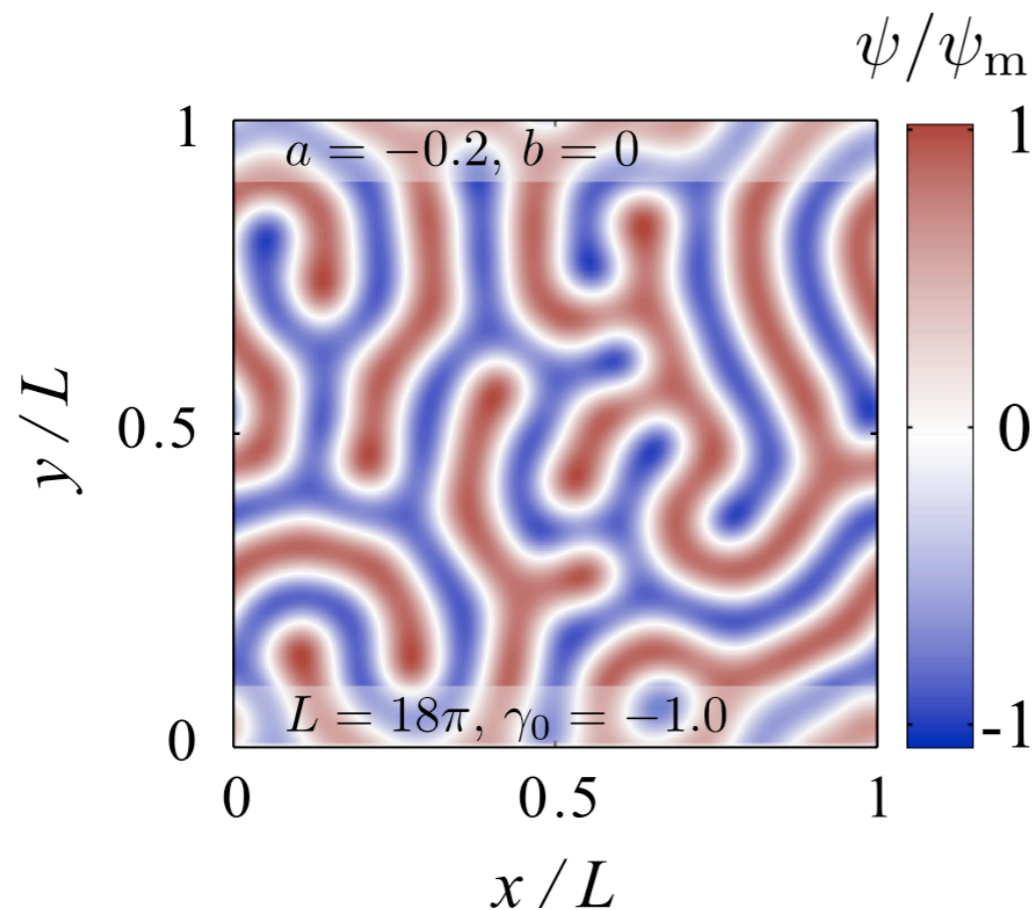


# Scalar field theory

2d Swift-Hohenberg model

$$\partial_t \psi = -U'(\psi) + \gamma_0 \nabla^2 \psi - \gamma_2 (\nabla^2)^2 \psi$$

$$U(\psi) = \frac{a}{2} \psi^2 + \frac{b}{3} \psi^3 + \frac{c}{4} \psi^4$$



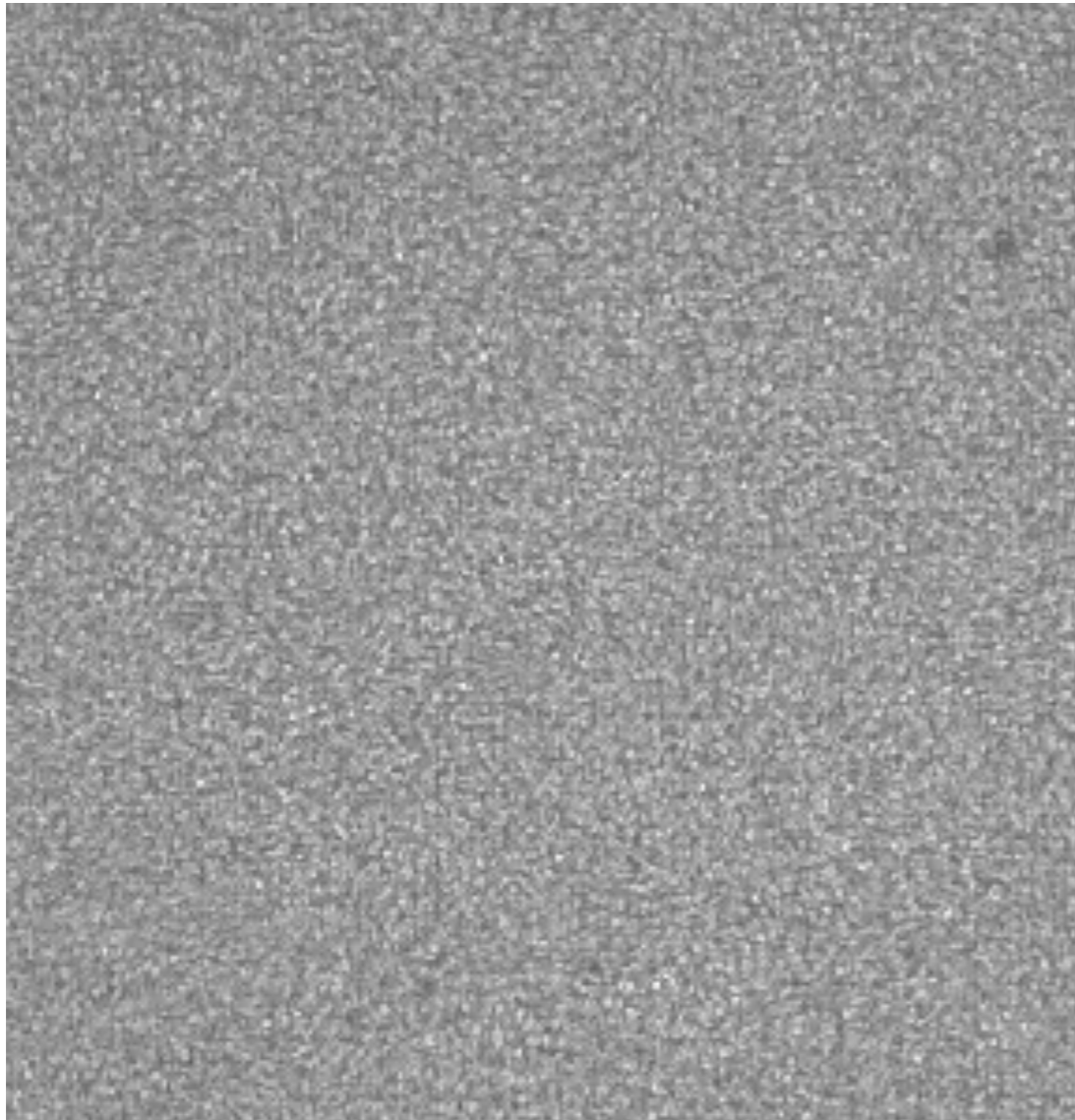


# Active patterns

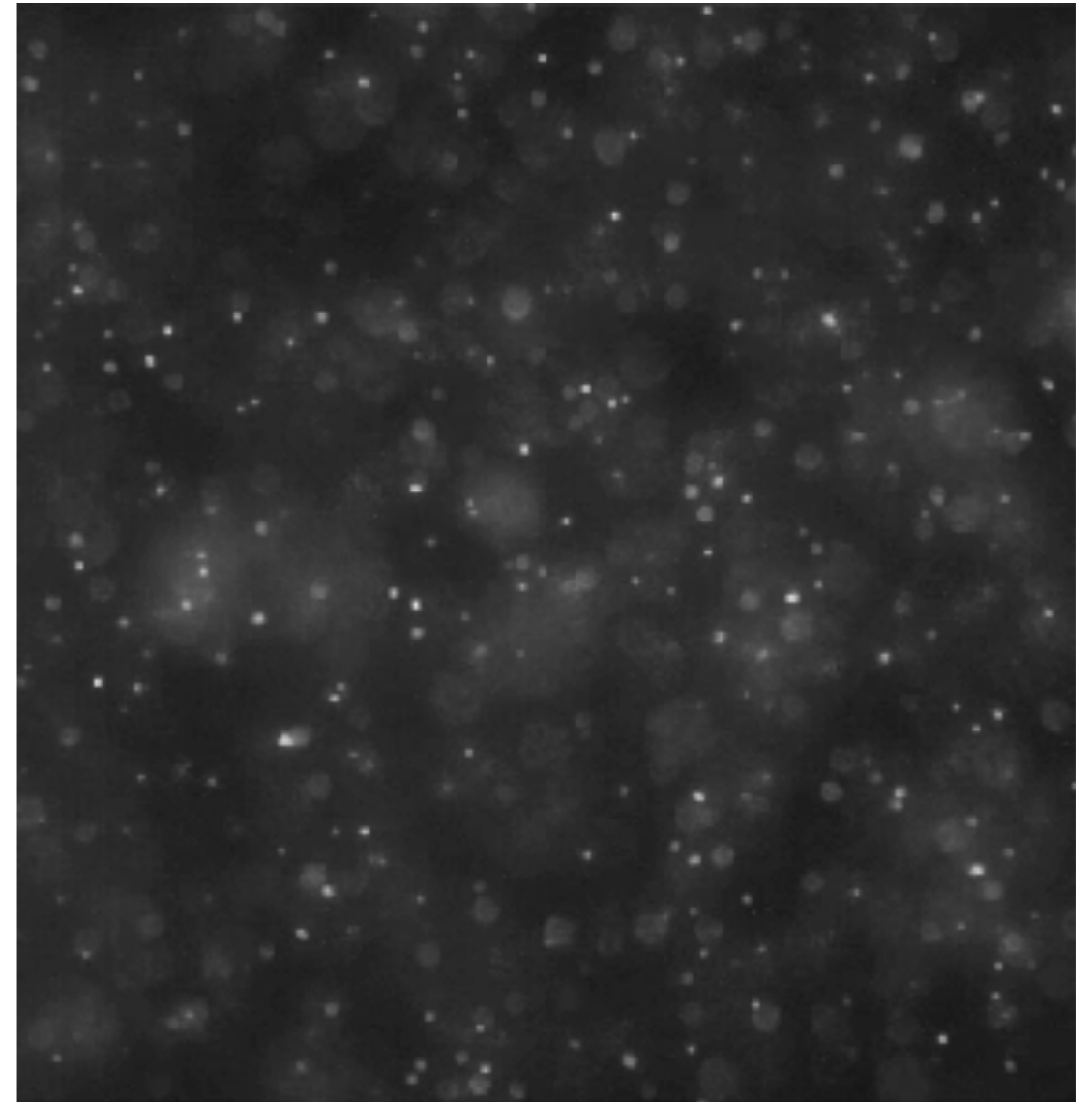
PRL (2013)

*B. subtilis*

tracer



bright field

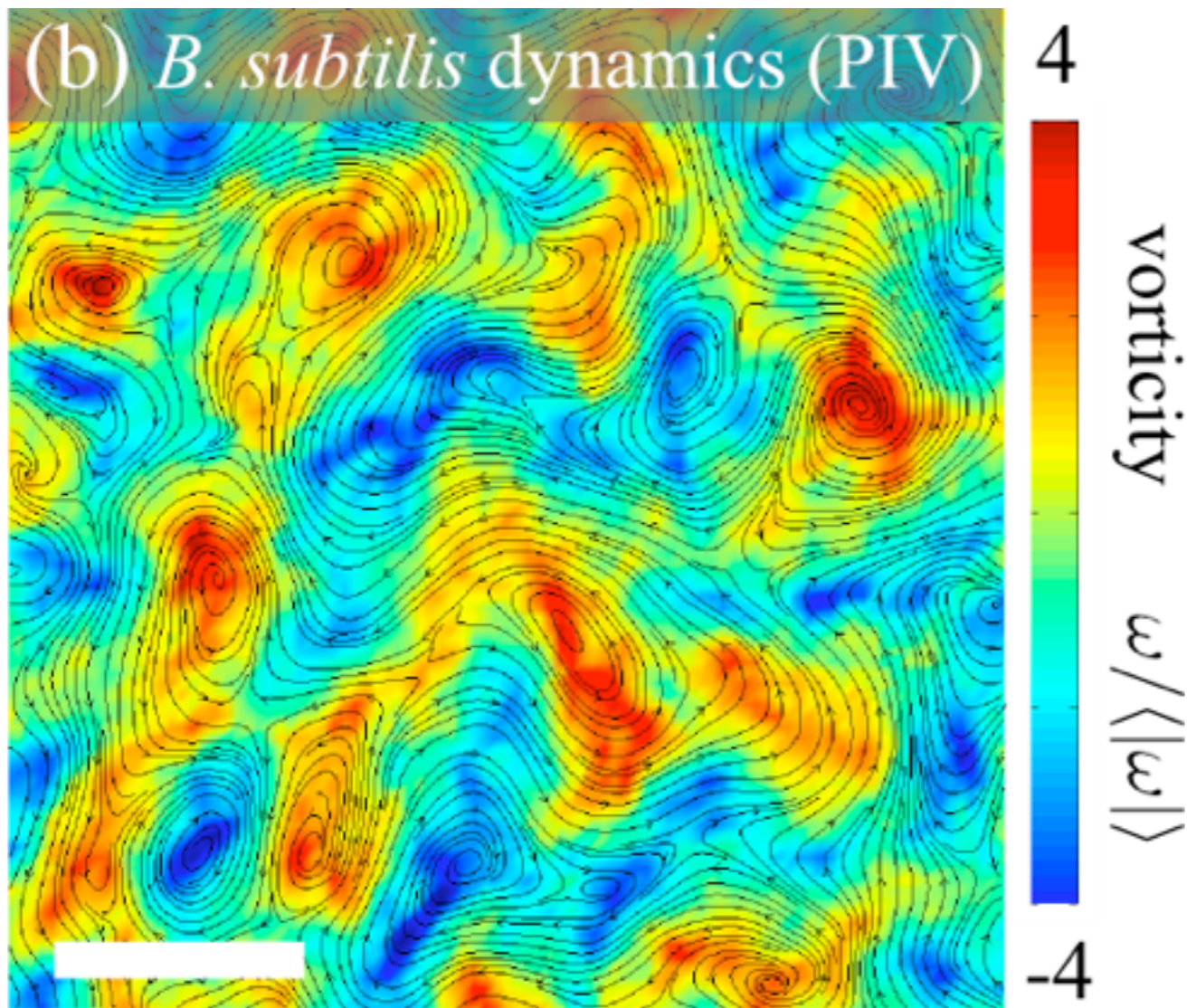


fluorescence

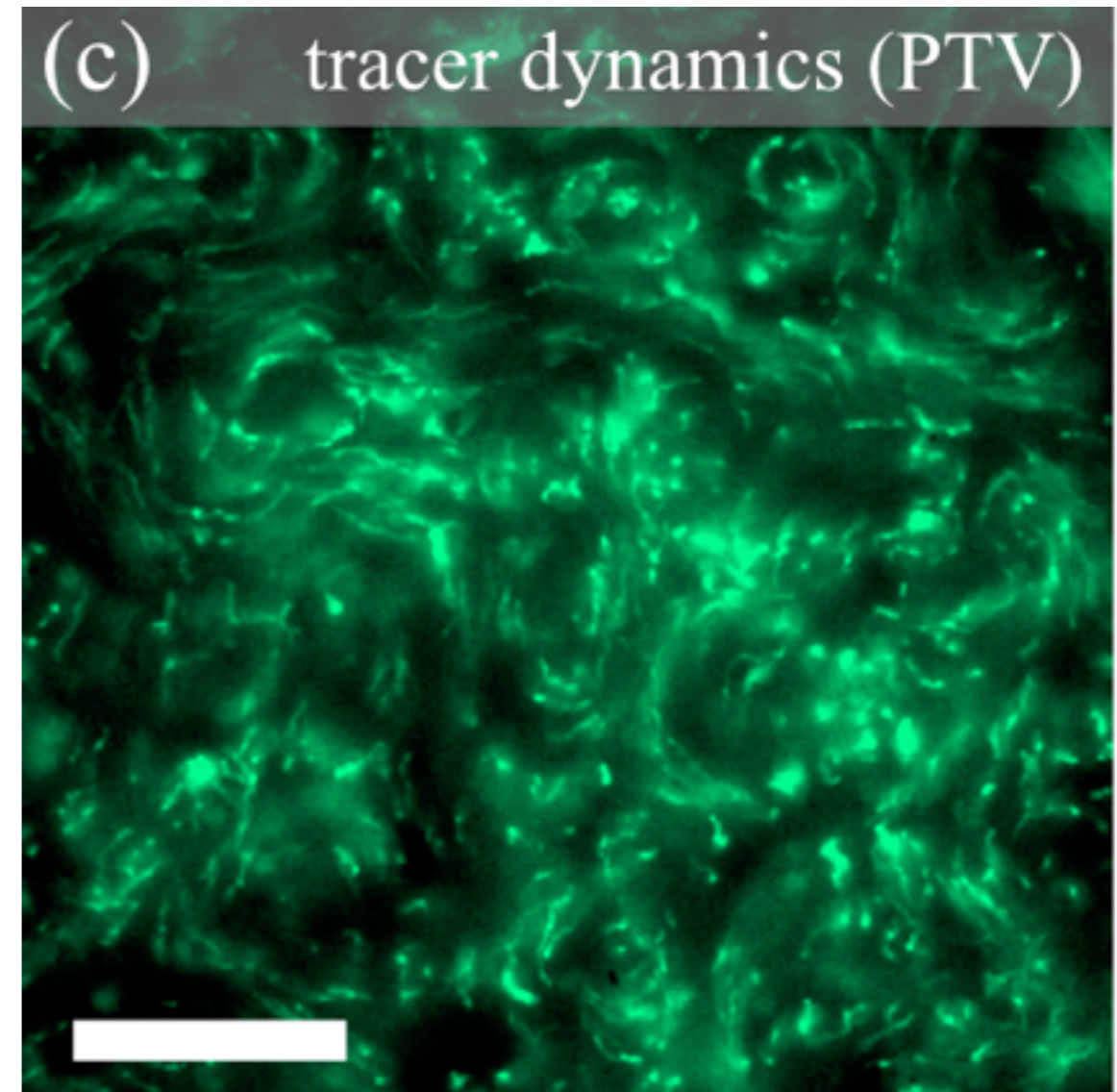


# 3D bacterial suspension

PRL (2013)



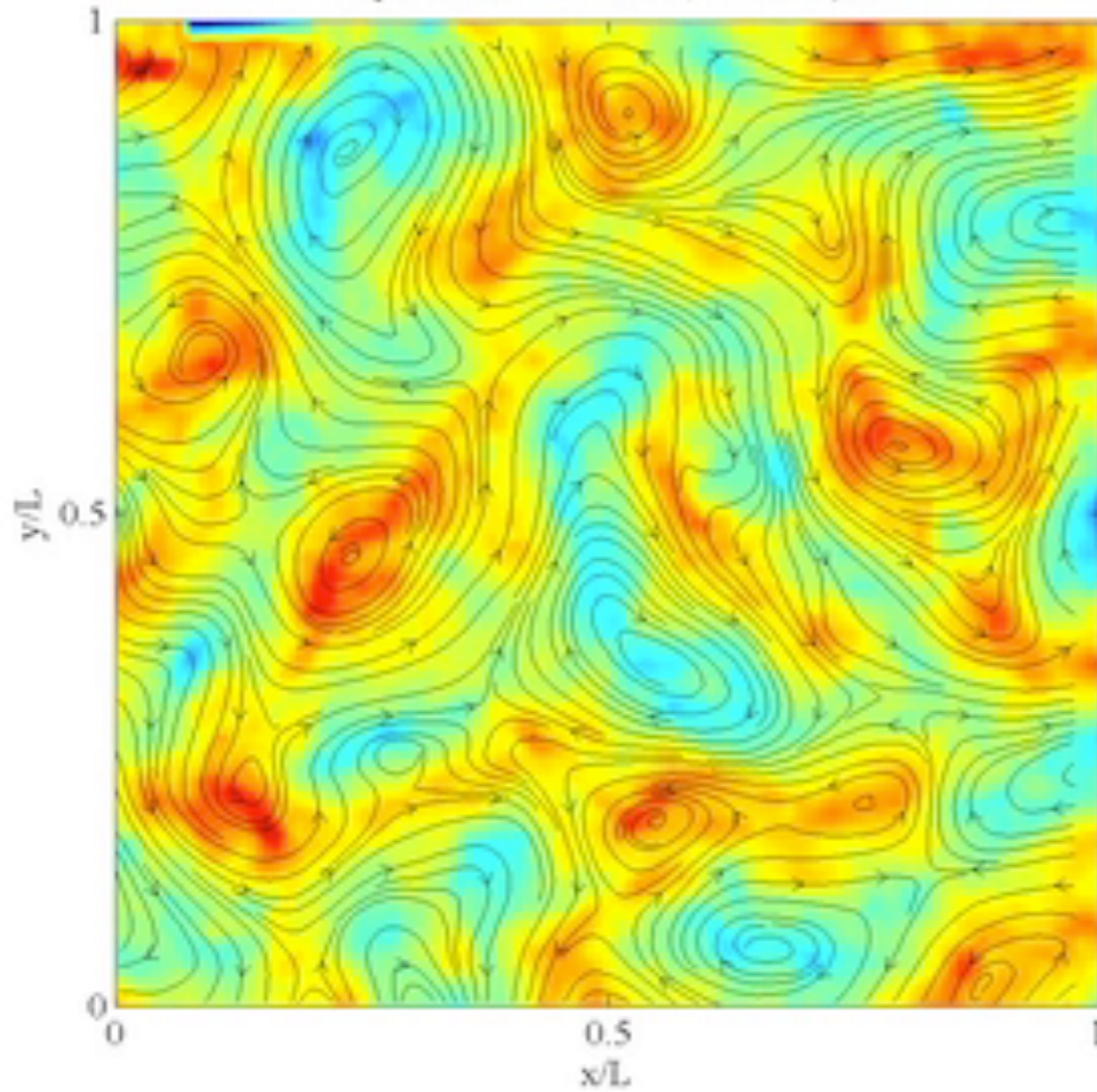
bright field



# 3D suspension

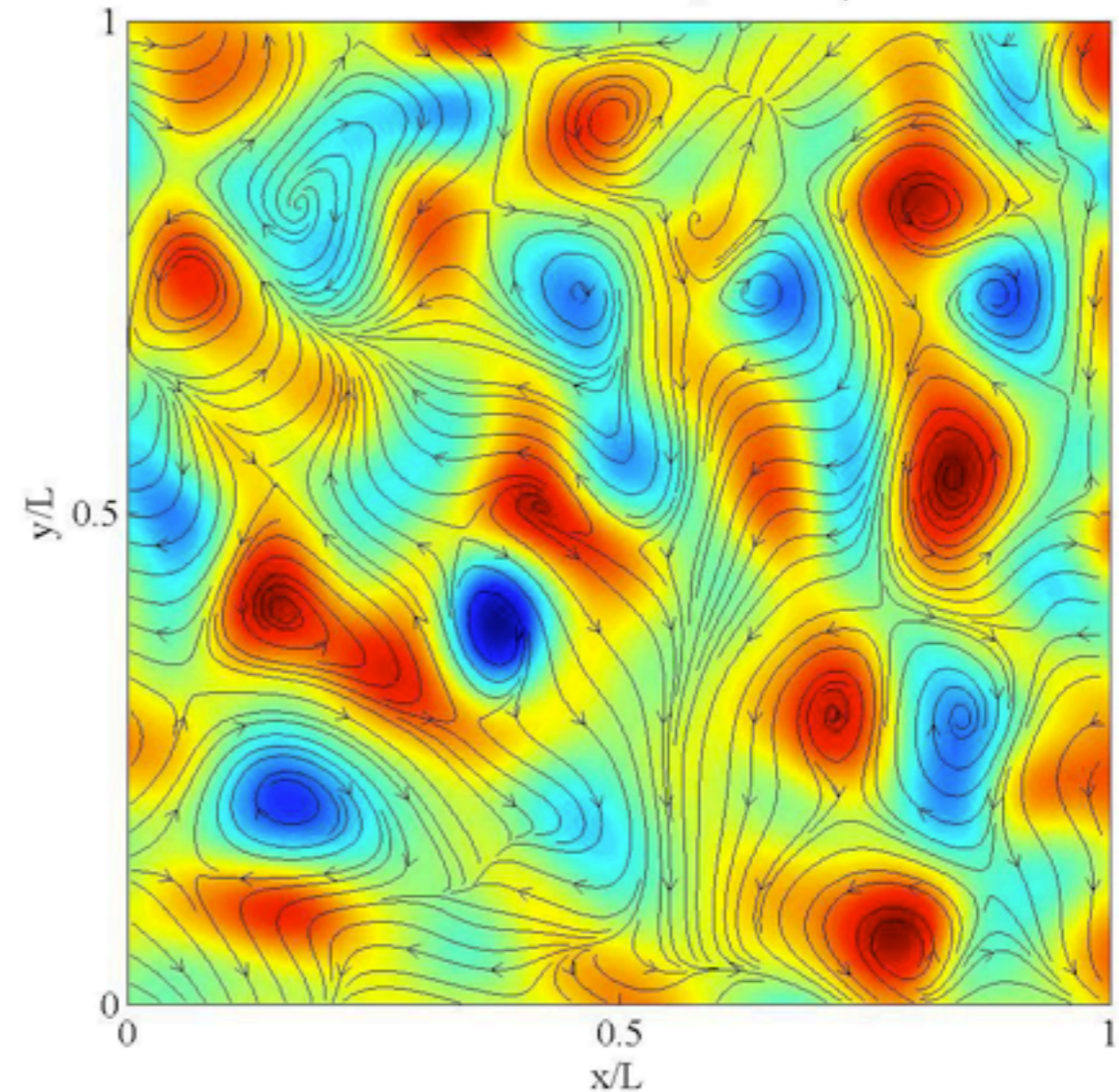
PRL (2013)

Experiment:  $t = 0.1 \text{ s}$ ,  $L = 276 \mu\text{m}$



Experiment:  
quasi-2D slice

Simulation:  $t = 8.7 \text{ s}$ ,  $L = 300 \mu\text{m}$



Theory:  
2D slice

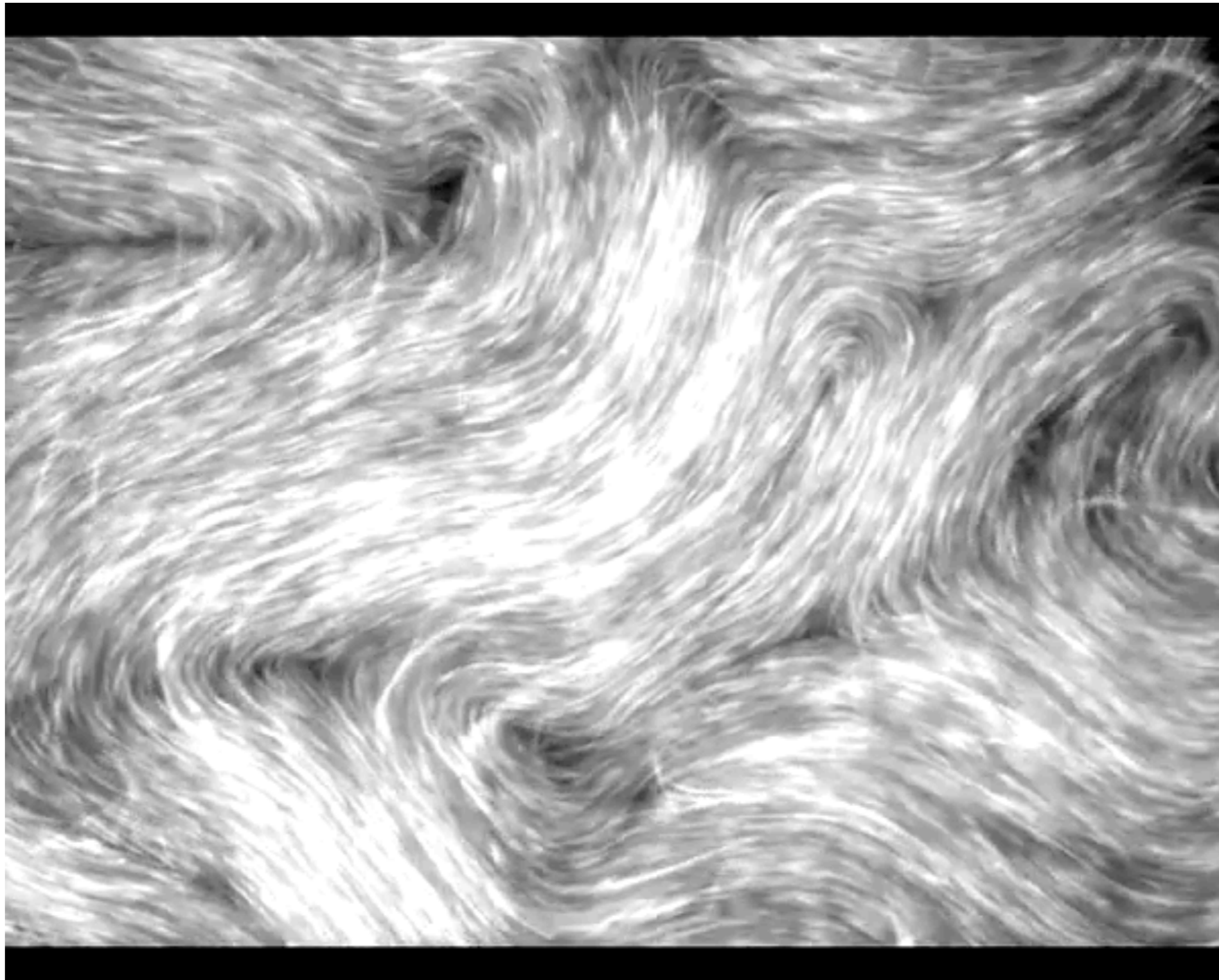
# Vector field theory

incompressibility

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

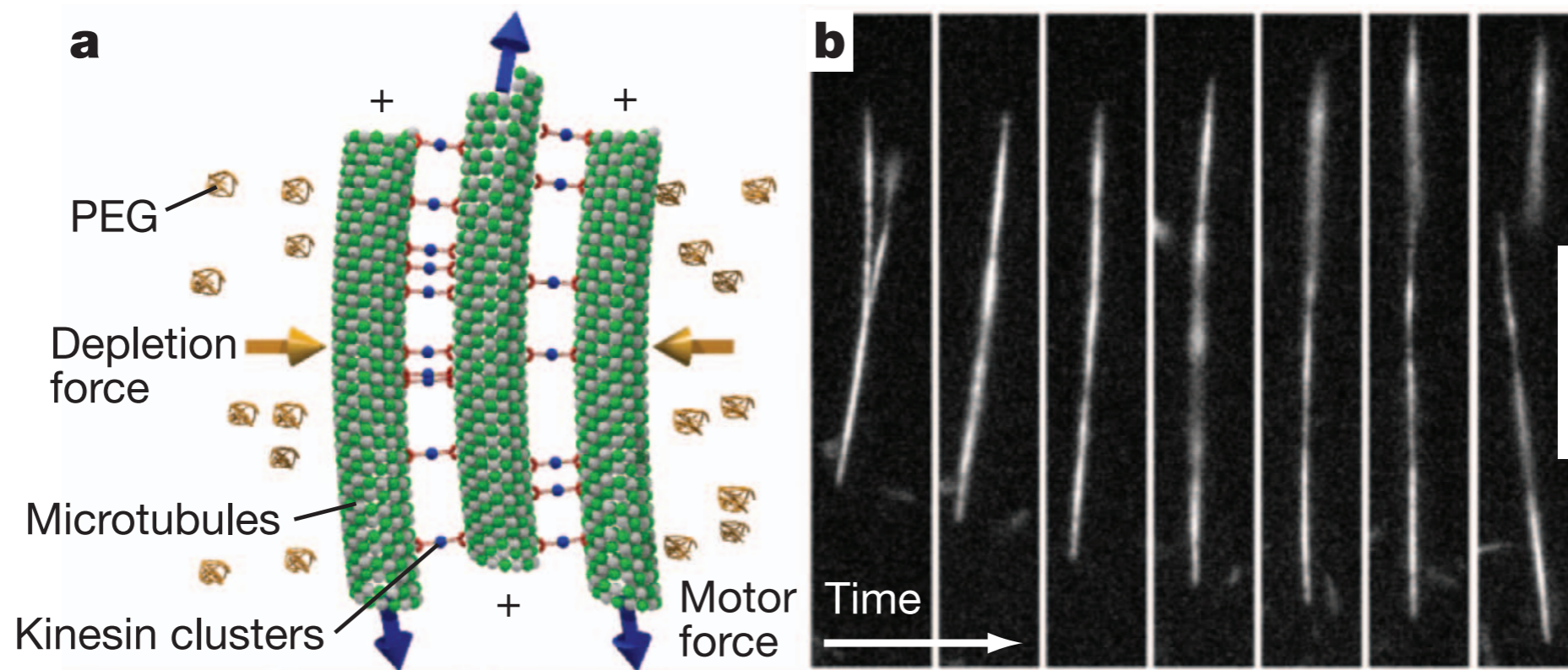
$$\begin{aligned} (\partial_t + \lambda_0 \mathbf{v} \cdot \nabla) \mathbf{v} = & - \nabla(p + \lambda_1 \mathbf{v}^2) - (\beta \mathbf{v}^2 + \alpha) \mathbf{v} + \\ & + \Gamma_0 \nabla^2 \mathbf{v} - \Gamma_2 (\nabla^2)^2 \mathbf{v} \end{aligned}$$

# Active nematics



Dogic lab (Brandeis) Nature 2012

# Active nematics



Dogic lab (Brandeis) Nature 2012

no head or tail  $\Rightarrow$  Q-tensor order-parameter

$$Q_{ij} = Q_{ji}, \quad \text{Tr } Q = 0, \quad Q = \begin{pmatrix} \lambda & \mu \\ \mu & -\lambda \end{pmatrix}.$$

$$\Delta = \sqrt{\lambda^2 + \mu^2}, \quad \Lambda^{\pm} = \pm \Delta$$

# Matrix field theory

$$\partial_t Q_{ij} + v_k \partial_k Q_{ij} = - \frac{\delta \mathcal{F}}{\delta Q_{ij}}$$

$$v_k = D \partial_n Q_{nk}$$



# biological networks

# Tokyo rail network by *Physarum plasmodium*



Tero et al (2010) Science

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