

Lecture 15:

The Mathematics of Water

Lecturer:

Luis Cruz (for Brigita Urbanc)

Office: 12-909

(E-mail: *brigita@drexel.edu*)

Course website:

www.physics.drexel.edu/~brigita/COURSES/BIOPHYS_2011-2012/

Water is of key importance in the living world.

Continuous description versus motion of individual H₂O molecules

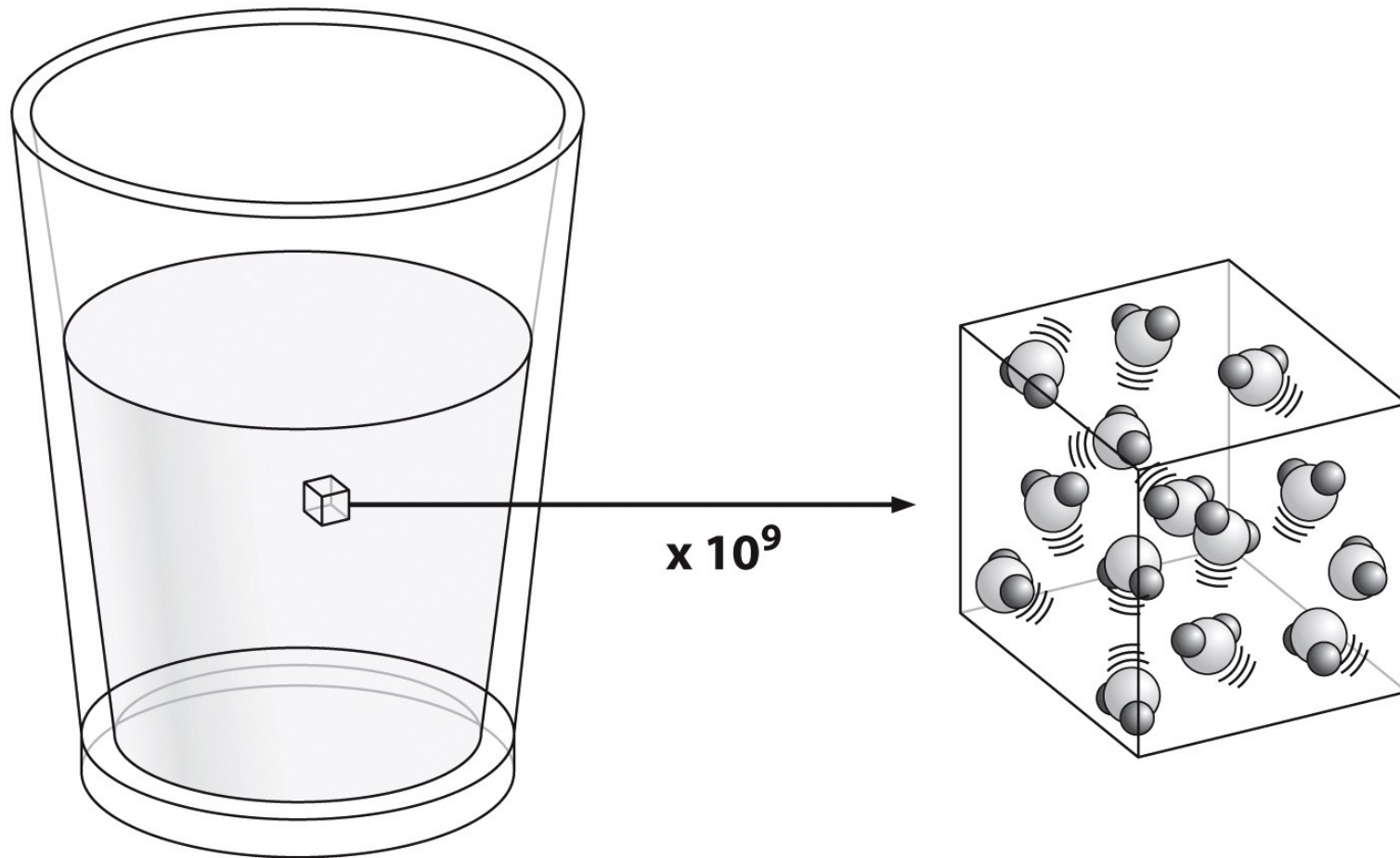


Figure 12.1 Physical Biology of the Cell (© Garland Science 2009)

Continuous description of water uses the velocity field, which in general can change with a spacial position (x,y,z) of a small volume of water and time:

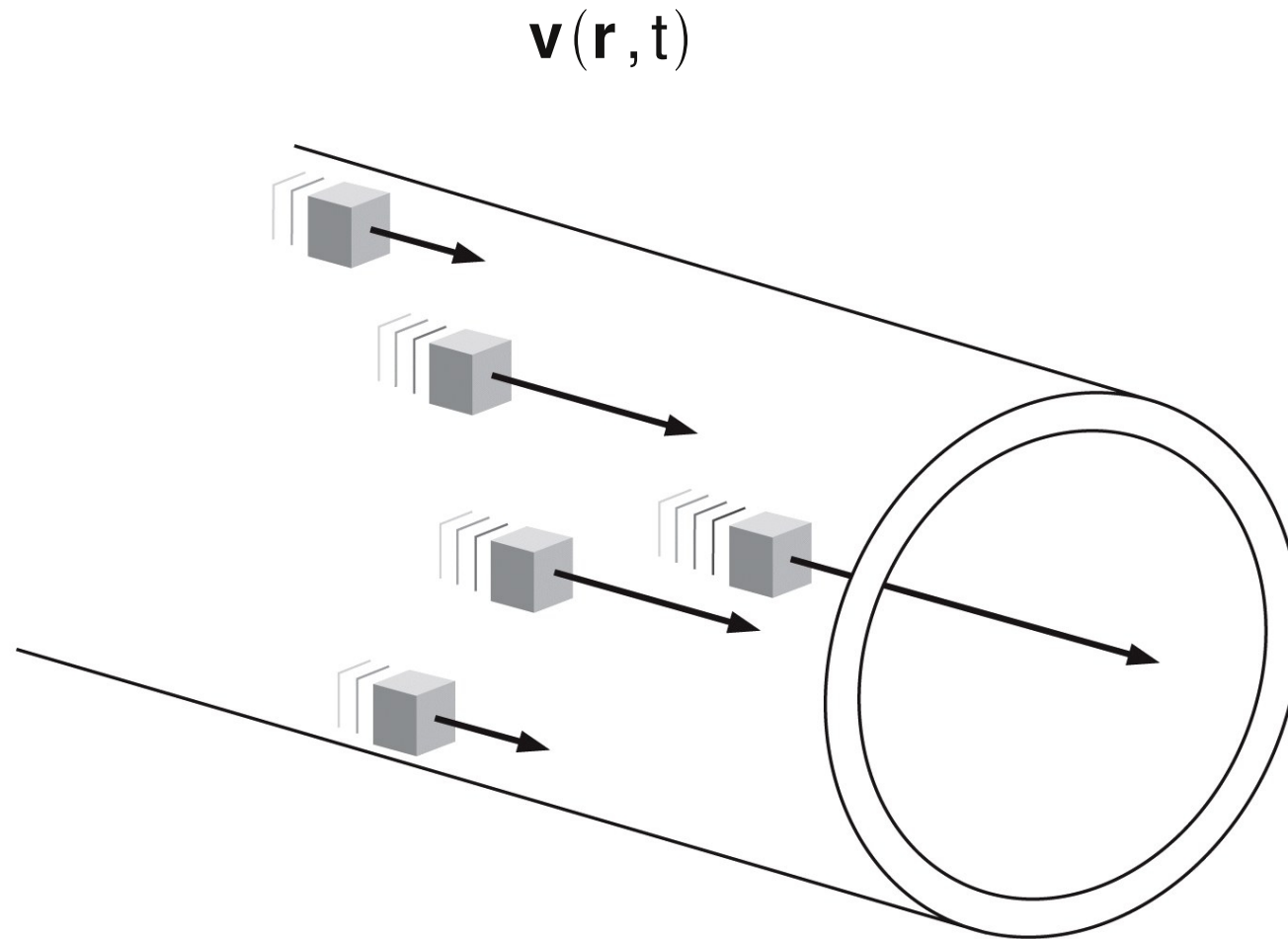


Figure 12.2 Physical Biology of the Cell (© Garland Science 2009)

Q: What distinguishes liquids from solids?

**A: Solids and liquids respond to applied forces differently:
Solids undergo deformation, liquids flow.**

**Mechanical properties
of liquids are described
by *viscosity* (in analogy
to Young modulus E
that describes solids).**

**For simple, Newtonian
fluids: pulling force
Results in a constant
velocity fluid motion:**

$$\frac{F}{A} = \eta \frac{v}{d}$$

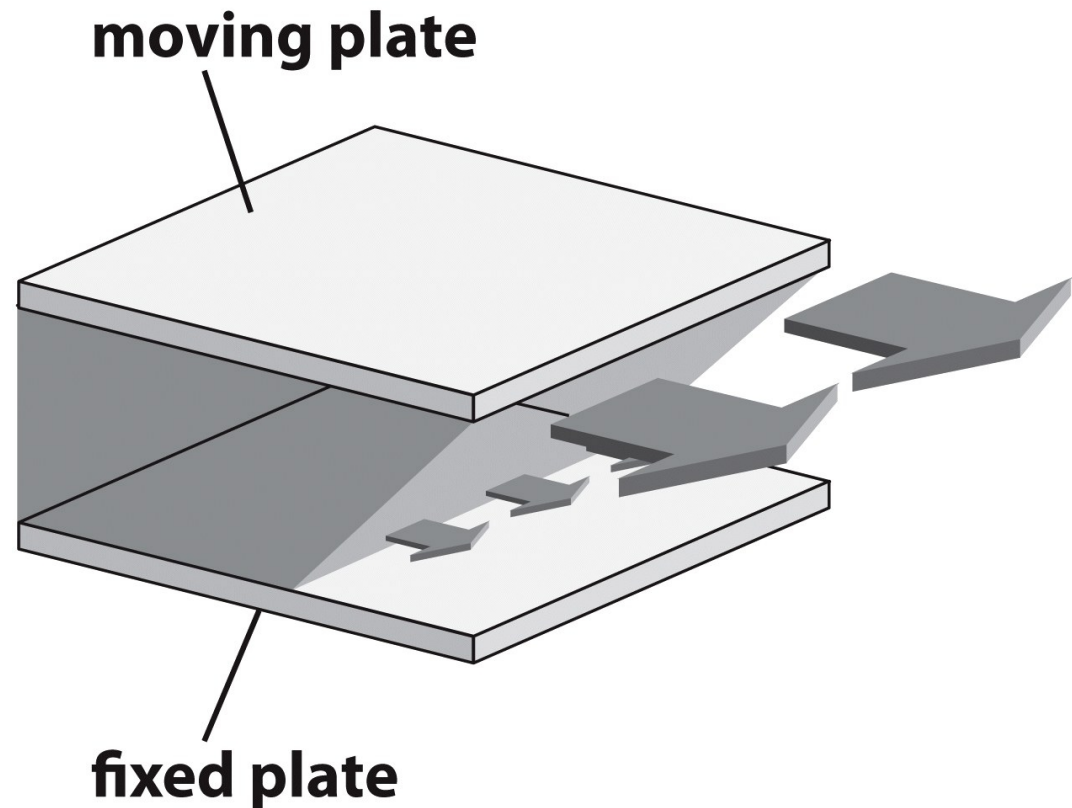


Figure 12.3 Physical Biology of the Cell (© Garland Science 2009)

Viscosity: $\eta [\text{Pa} \cdot \text{s}]$, where $1 \text{ Pa} = 10^{-5} \text{ N/m}^2 = 10^{-5} \text{ atm}$

Viscosity of water: $\eta = 1 \text{ mPa} \cdot \text{s} = 10^{-8} \text{ N/m}^2 \cdot \text{s} = 10^{-8} \text{ atm} \cdot \text{s}$

Equation of motion for a Newtonian fluid: $\mathbf{F} = m \mathbf{a}$

Consider a small fluid element and apply Newton's second law to its motion:

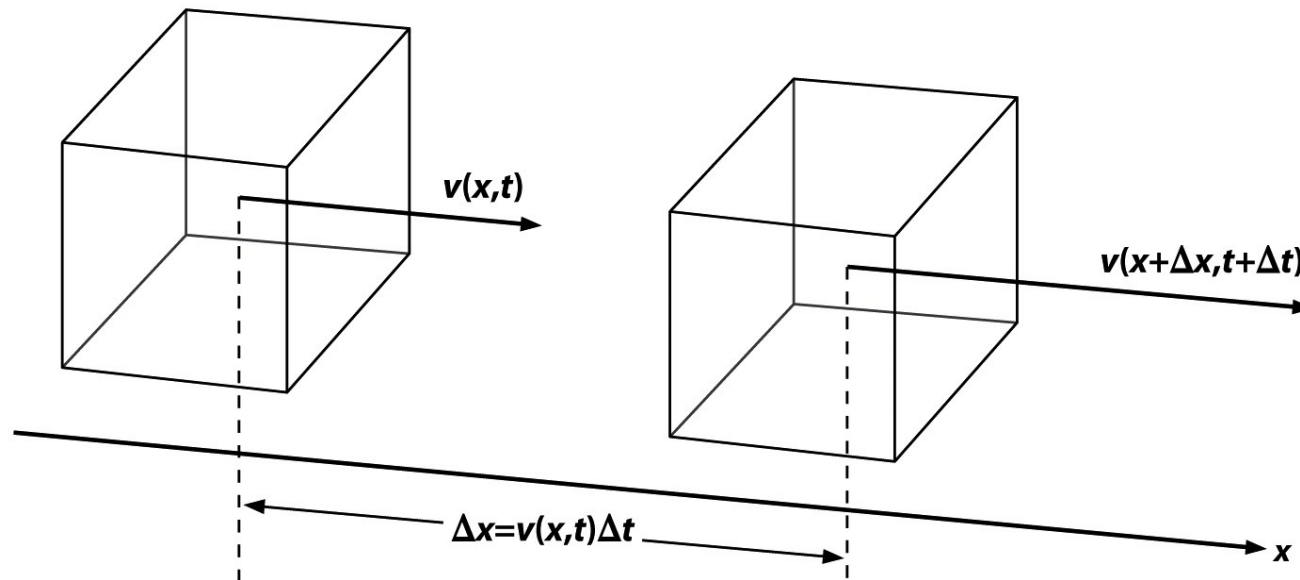


Figure 12.4 Physical Biology of the Cell (© Garland Science 2009)

In general, we need to consider: $\mathbf{r}=(x, y, z)$ $\mathbf{v}=(v_x, v_y, v_z)$

To derive the equation of motion, we will assume a one-dimensional Problem:

$v(x, t)=v(x, t)\mathbf{e}_x$ \mathbf{e}_x ...unit vector along x – direction

$$v(x+\Delta x, t+\Delta t) = v(x, t) + \frac{\partial v}{\partial x}\Delta x + \frac{\partial v}{\partial t}\Delta t$$

The element moves along the x-direction by Δx in time Δt :

$$v(x, t) = \frac{\Delta x}{\Delta t} \rightarrow \frac{\partial v}{\partial x}\Delta x = \frac{\partial v}{\partial x} \cdot \frac{\Delta x}{\Delta t} \cdot \Delta t = \frac{\partial v}{\partial x} \cdot v \cdot \Delta t$$

Thus we can calculate the change of velocity and thus acceleration along the x-direction:

$$\Delta v(x, t) = \left[\frac{\partial v}{\partial t} + v(x, t) \frac{\partial v}{\partial x} \right] \Delta t \rightarrow$$
$$a(x, t) = \frac{\Delta v(x, t)}{\Delta t} = \frac{\partial v(x, t)}{\partial t} + v(x, t) \frac{\partial v(x, t)}{\partial x}$$

We can express the acceleration of a fluid element at position \mathbf{x} and at time t as a *material derivative*:

$$\mathbf{a} = \frac{D \mathbf{v}}{Dt}$$

which is defined as: $\frac{D}{Dt} = \frac{\partial}{\partial t} + \mathbf{v} \cdot \nabla$

Material derivative consists of two parts:

- (1) conventional explicit time dependence reflecting the field variable \mathbf{v} that is changing with time
- (2) convective term: a material particle can be dragged into a region of space, where the field is different

Example of a temperature field $T(\mathbf{x}, t)$:

- (1) fluid is resting but T is changing with time

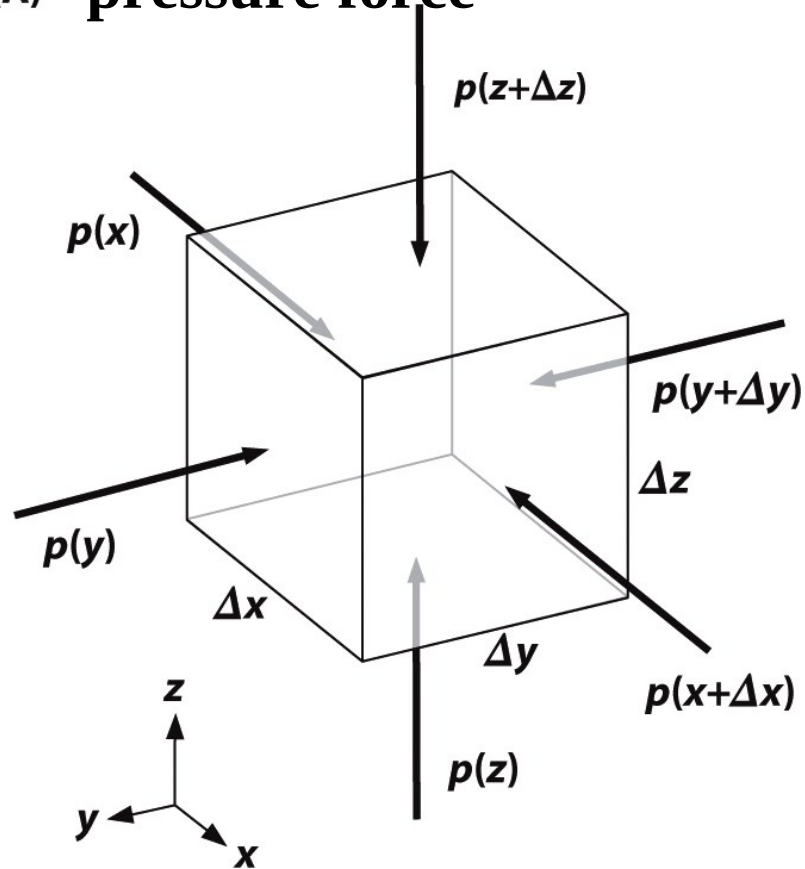
(e.g. heating): $\frac{D}{Dt} = \frac{\partial}{\partial t}$

- (2) fluid is moving, steady state T gradient: $\frac{D}{Dt} = \mathbf{v}_x \cdot \nabla$

Two different kinds of forces are acting on a fluid volume element

$$\Delta x \times \Delta y \times \Delta z$$

(A) pressure force



(B) viscous stress force

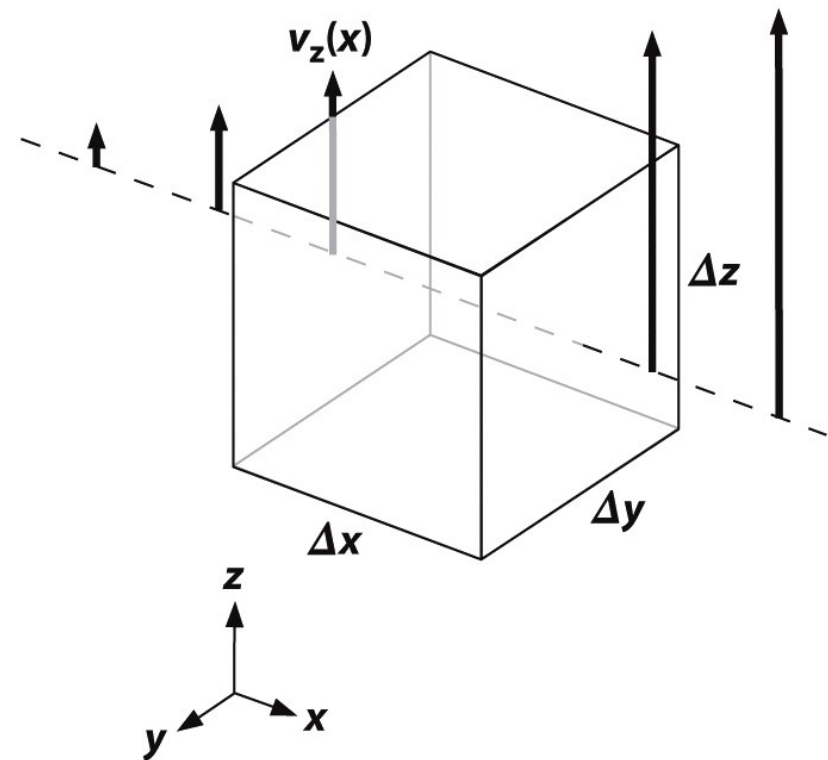


Figure 12.5 Physical Biology of the Cell (© Garland Science 2009)

Pressure Force: due to variations in the pressure along x, y, z

Viscous Stress Force: friction on the volume element due to a velocity gradient along x, y, z

The total pressure force along the x-direction:

$$\delta F_x^p = p(x, y, z, t) \Delta y \Delta z - p(x + \Delta x, y, z, t) \Delta y \Delta z$$

$$p(x + \Delta x, y, z, t) \approx p(x, y, z, t) + \frac{\partial p}{\partial x} \Delta x \rightarrow$$

$$\delta \mathbf{F}^p = -\nabla p \Delta x \Delta y \Delta z \quad \text{where} \quad \nabla p = \left(\frac{\partial p}{\partial x}, \frac{\partial p}{\partial y}, \frac{\partial p}{\partial z} \right)$$

The viscous force on the face at position x is:

$$-\eta \frac{\partial v_z(x)}{\partial x} \Delta y \Delta z \mathbf{e}_z$$

whereas on the face at position $x + \Delta x$ is:

$$+\eta \frac{\partial v_z(x + \Delta x)}{\partial x} \Delta y \Delta z \mathbf{e}_z$$

The total viscous stress force is then a sum of the two:

$$\begin{aligned}\delta \mathbf{F}_z^v &= -\eta \frac{\partial v_z(x)}{\partial x} \Delta y \Delta z \mathbf{e}_z + \eta \frac{\partial v_z(x + \Delta x)}{\partial x} \Delta y \Delta z \mathbf{e}_z \\ \frac{\partial v_z(x + \Delta x)}{\partial x} &\approx \frac{\partial v_z(x)}{\partial x} + \frac{\partial^2 v_z(x)}{\partial x^2} \Delta x \\ \delta \mathbf{F}_z^v &= \eta \frac{\partial^2 v_z(x)}{\partial x^2} \Delta x \Delta y \Delta z\end{aligned}$$

In general, the total viscous stress force can be written as:

$$\delta \mathbf{F}^v = (\eta \nabla^2 \mathbf{v} + \eta' \nabla (\nabla \cdot \mathbf{v})) \Delta x \Delta y \Delta z$$

If we only consider incompressible fluids, then $\nabla \cdot \mathbf{v} = 0$ and:

$$\delta \mathbf{F}^v = \eta \nabla^2 \mathbf{v} \Delta x \Delta y \Delta z$$

Incompressible fluids: the flow can distort and stretch the fluid element but preserves its volume.

Consider a volume element at a time t :

$$\Delta x \times \Delta y \times \Delta z$$

A moment later, at $t + \Delta t$:

$$\Delta x' = \Delta x \left(1 + \frac{\partial v_x}{\partial x} \Delta t \right)$$

$$\Delta y' = \Delta y \left(1 + \frac{\partial v_y}{\partial y} \Delta t \right)$$

$$\Delta z' = \Delta z \left(1 + \frac{\partial v_z}{\partial z} \Delta t \right)$$

Thus the volume change is:

$$\Delta x' \Delta y' \Delta z' = \Delta x \Delta y \Delta z \left[1 + \left(\frac{\partial v_x}{\partial x} + \frac{\partial v_y}{\partial y} + \frac{\partial v_z}{\partial z} \right) \Delta t \right]$$

$$\Delta x' \Delta y' \Delta z' = \Delta x \Delta y \Delta z (1 + \nabla \cdot \mathbf{v})$$

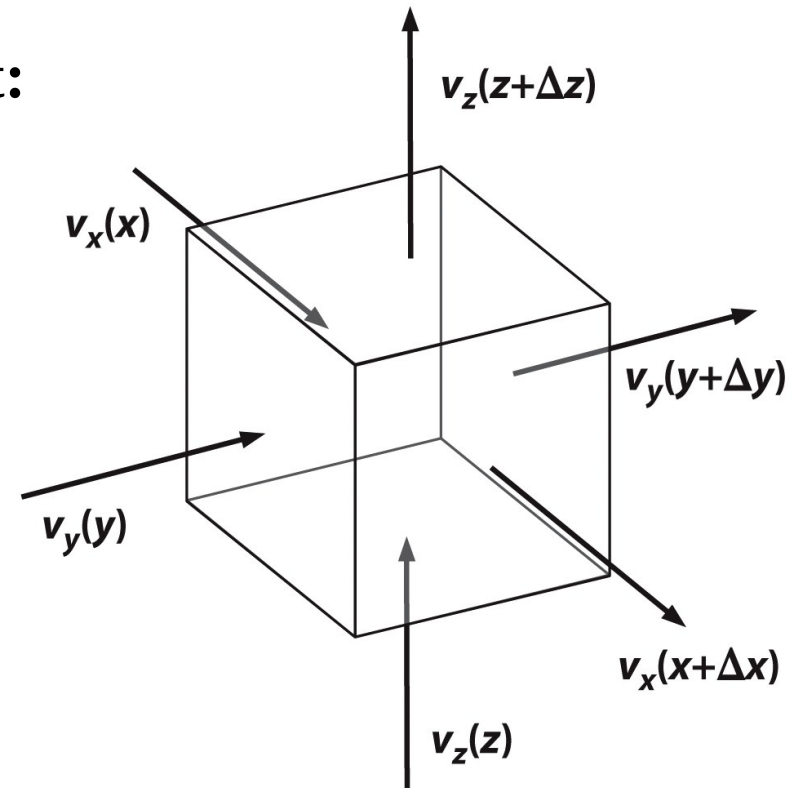


Figure 12.6 Physical Biology of the Cell (© Garland Science 2009)

The Newtonian Fluid and the Navier-Stokes Equation

$$\Delta m \mathbf{a} = \delta \mathbf{F}_p + \delta \mathbf{F}_v \quad \text{where} \quad \Delta m = \rho \Delta x \Delta y \Delta z$$

$$\frac{\partial \mathbf{v}}{\partial t} + (\mathbf{v} \cdot \nabla) \mathbf{v} = -\frac{1}{\rho} \nabla p + \nu \nabla^2 \mathbf{v} \quad \text{where} \quad \nu = \frac{\eta}{\rho}$$

kinematic viscosity

The Navier-Stokes equation is a second-order partial differential equation: need to specify the boundary conditions:

no-slip condition = at a solid boundary the fluid is at rest with respect to the solid

Fluid Dynamics of Blood

- a process of blood circulation
- blood vessels ranging in size from $2\text{ }\mu\text{m}$ to 1 cm
- use cylindrical symmetry to describe the blood flow

$$\mathbf{v} = v(r) \mathbf{e}_z$$

- consider only steady state:

$$\frac{\partial \mathbf{v}}{\partial t} = 0$$

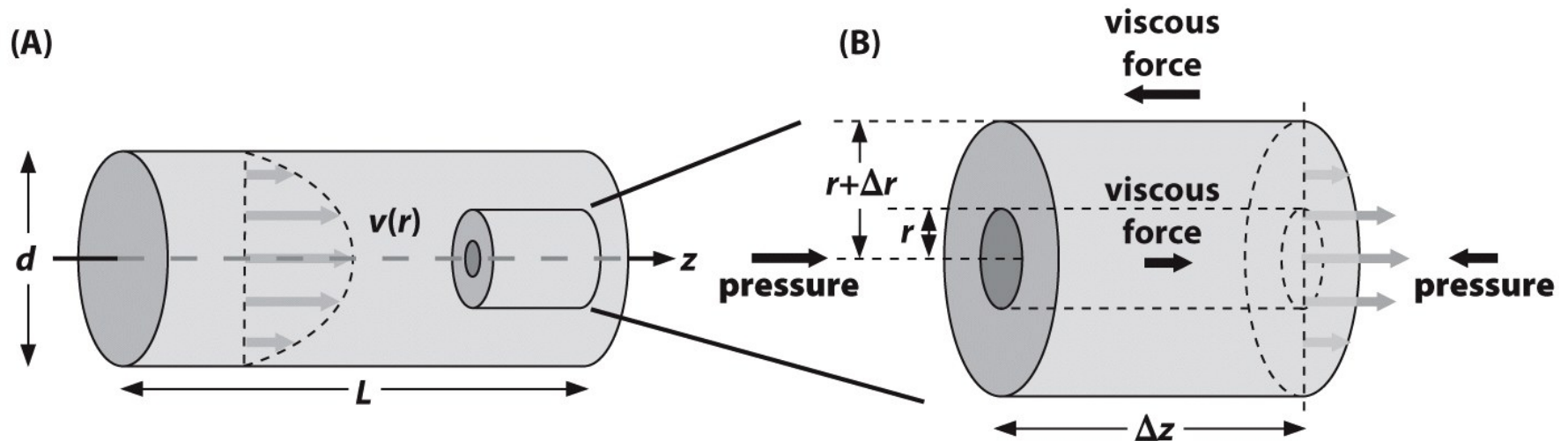


Figure 12.7 Physical Biology of the Cell (© Garland Science 2009)

Consider a small volume element in the shape of a hollow cylinder and calculate the opposing pressure forces:

$$\delta F_z^p = [p(z) - p(z + \Delta z)] 2\pi r \Delta r = -\frac{dp}{dz} 2\pi r \Delta r \Delta z$$

and the opposing viscous stress forces:

$$\delta F_z^v = \eta v'(r + \Delta r) 2\pi (r + \Delta r) \Delta z - \eta v'(r) 2\pi r \Delta z \quad \text{where} \quad v' = \frac{dv}{dr}$$

Use the Taylor expansion: $v'(r + \Delta r) \approx v'(r) + \frac{dv'}{dr} \Delta r$

To derive the expression for the viscous stress force:

$$\delta F_z^v = \eta v' 2\pi \Delta r \Delta z + \eta \frac{dv'}{dr} 2\pi r \Delta r \Delta z$$

In a steady state, a force balance on the hollow fluid cylinder is:

$$\delta F_z^p + \delta F_z^v = 0$$

This results in a differential equation relating the pressure and velocity fields:

$$\frac{1}{\eta} \frac{dp}{dz} = \frac{1}{r} \frac{dv}{dr} + \frac{d^2 v}{dr^2} = \frac{1}{r} \frac{d}{dr} \left(r \frac{dv}{dr} \right)$$

Laplacian of the velocity field in cylindrical coordinates

How do we solve this equation?

Integrate both sides along the z direction along the length L:

$$\frac{-1}{\eta} \Delta p = \frac{1}{r} \frac{d}{dr} \left(r \frac{dv}{dr} \right) L \quad \text{where} \quad \Delta p = p(0) - p(L)$$

pressure drop along the length of the pipe

Now integrate twice along r to get a solution:

$$v(r) = -\frac{\Delta p}{\eta L} \left(\frac{r^2}{4} - \frac{C_1}{r^2} + C_2 \right)$$

Take into account the boundary conditions:

$$v(r=0) < \infty \rightarrow C_1 = 0$$
$$v\left(r=\frac{d}{2}\right) = 0 \quad \text{no-slip} \rightarrow C_2 = \frac{-d^2}{16}$$

We derive the final expression for the fluid velocity in the pipe:

$$v(r) = \frac{\Delta p}{4\eta L} \left(\frac{d^2}{4} - r^2 \right)$$

The fluid moves the fastest in the center of the pipe and not at all at the boundaries. We need an average velocity $\langle v \rangle$ which can be related to the flow rate Q :

$$Q = \frac{\langle v \rangle \pi d^2}{4}$$

The average velocity and the resulting flow rate are:

$$\langle v \rangle = \frac{\int_0^{d/2} v(r) 2\pi r dr}{\frac{\pi d^2}{4}} = \frac{\Delta p d^4}{128\eta L} \rightarrow Q = \frac{\pi \Delta p d^4}{128\eta L}$$

Estimated blood flow through capillaries:

$$d \approx 5 \mu\text{m} \quad \text{for most animals}$$

A measured pressure difference across a capillary and its length:

$$\Delta p \approx 20 \text{ mmHg} \approx 3000 \text{ Pa}$$

$$L \approx 1 \text{ cm}$$

The average flow velocity through a capillary:

$$v = \frac{\Delta p d^2}{32 \eta L} \approx 0.02 \text{ cm/s} \quad (\eta \approx \eta_{\text{water}} = 10^{-3} \text{ Pa})$$

But the viscosity of blood is higher than that of water:

$$\eta \approx 3 \eta_{\text{water}} \rightarrow v \approx 0.007 \text{ cm/s} \quad \text{too small!}$$

$$v_{\text{measured}} \approx 0.05 \text{ cm/s}$$

Can blood be well represented as a Newtonian liquid? Not really.

Blood carries red blood cells and white blood cells (leukocytes)

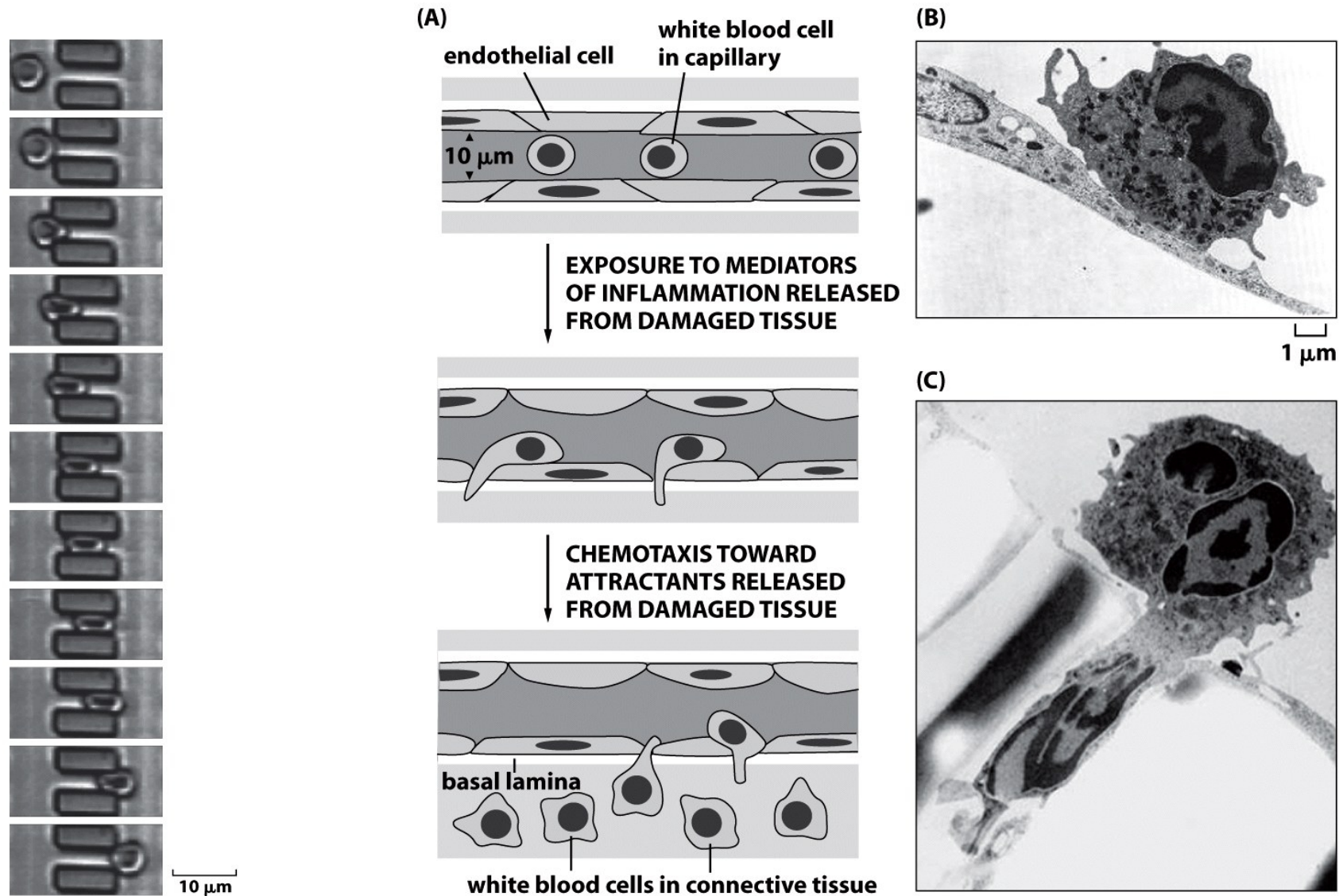


Figure 2.11 Physical Biology of the Cell (© Garland Sci

Figure 12.8 Physical Biology of the Cell (© Garland Science 2009)

The Low Reynolds number World

The 100 years experiment with a very viscous fluid: 1 drop every 10 years!

This system has the same viscous properties as water to the *E. coli* bacterium.

What is a Reynolds number Re ?
 $Re = \text{inertial term of N.-S. Eq.} / \text{viscous term of N.-S. Eq.}$

For a rigid body of length L moving with a speed U :

$$Re.N. = \frac{\rho U^2 / L}{\eta U / L^2} = \frac{\rho L U}{\eta} = \frac{L U}{\nu}$$



Figure 12.9 Physical Biology of the Cell (© Garland Science 2009)

For small Reynolds numbers, the Navier-Stokes equation simplifies into the Stokes equation:

$$\nabla p = \eta \nabla^2 \mathbf{v} \quad \text{incompressibility: } \nabla \cdot \mathbf{v} = 0$$

The meaning of Reynolds number can be found by comparing the kinetic energy to the work done by the viscous stress dissipation of an object of linear dimension a and speed u (where u/a is the rate of change of the fluid velocity):

$$KE \approx \rho a^3 u^2 \quad W \approx \eta \frac{u}{a} \times a^2 \times a \quad \rightarrow \quad \text{Re.N.} = \frac{KE}{W} = \frac{a u}{\nu}$$

Another way: compare the time needed to dissipate KE of the fluid element to the time needed to move a distance comparable to its size:

$$\tau_{\text{viscous}} = \frac{\rho a^2}{\eta} \ll \tau_{\text{inertial}} = \frac{a}{u} \quad \rightarrow \quad \text{Re.N.} = \frac{\tau_{\text{viscous}}}{\tau_{\text{inertial}}}$$

Flow around a sphere at different Reynolds numbers

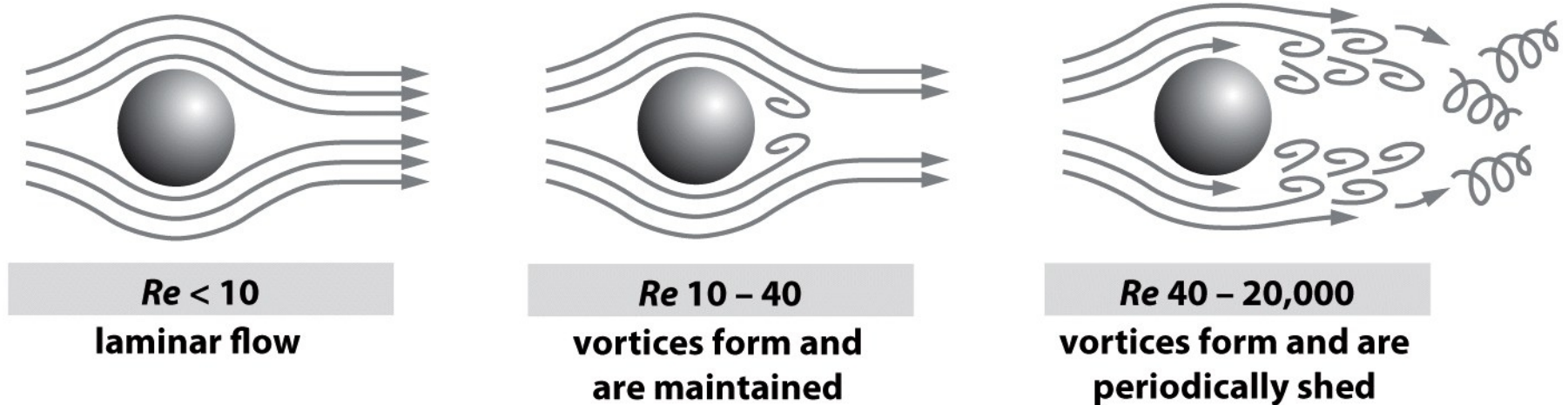


Figure 12.10 Physical Biology of the Cell (© Garland Science 2009)

The Stokes formula for a drag force on a sphere

- need to integrate the viscous stress and the pressure over the surface of a sphere
- viscous stress proportional to the spatial derivative of the speed $\frac{v}{R}$
- the area of the surface of the sphere: $4\pi R^2$
- the Stokes force/drag is then: $F_s = C\eta \frac{v}{R} \times 4\pi R^2 = 6\pi\eta Rv$

Stokes Drag in Single-Molecule Experiments

- ATP synthase is a rotary motor
- actin filament attached to this motor is tagged by a fluorescent molecule
- when the motor rotates, the filament spins around and the fluorescent tag allows us to measure the rotation
- existence of pauses in molecular motors

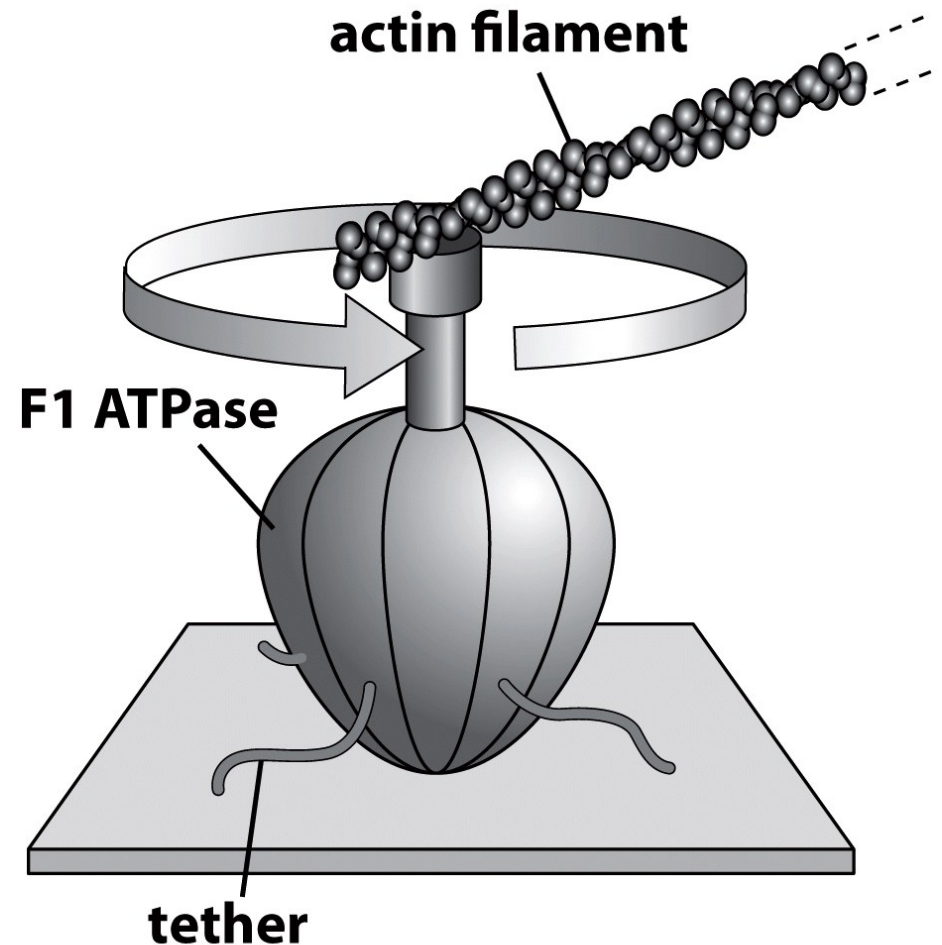


Figure 12.11 Physical Biology of the Cell (© Garland Science 2009)

Molecular motor: Motion of myosin on an actin filament

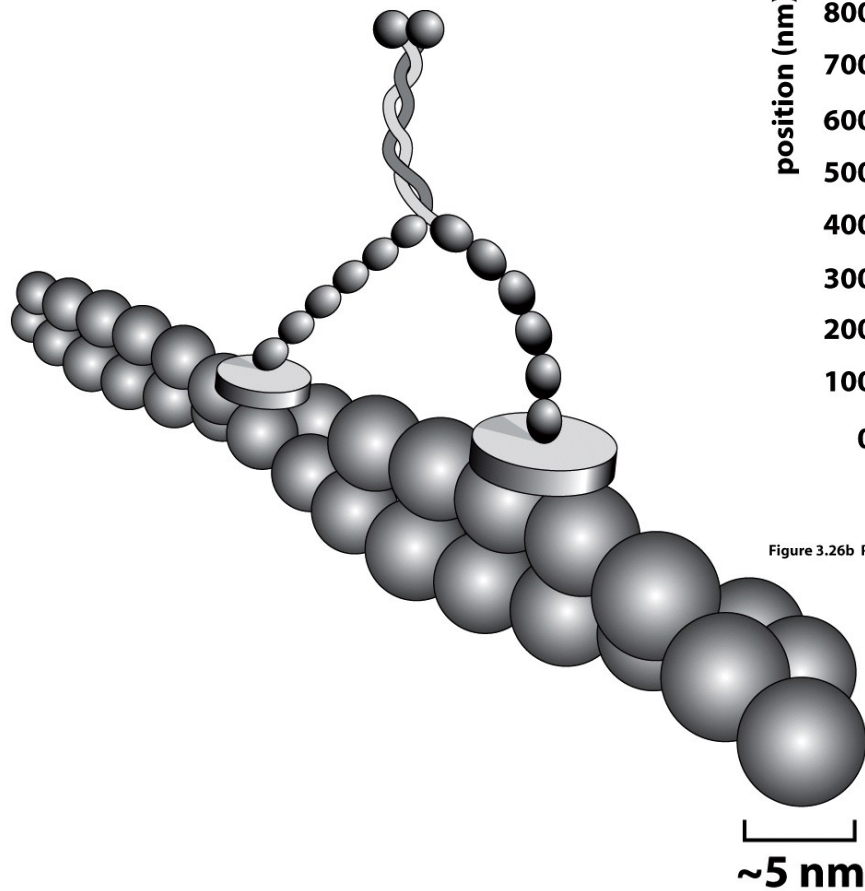


Figure 3.26a Physical Biology of the Cell (© Garland Science 2009)

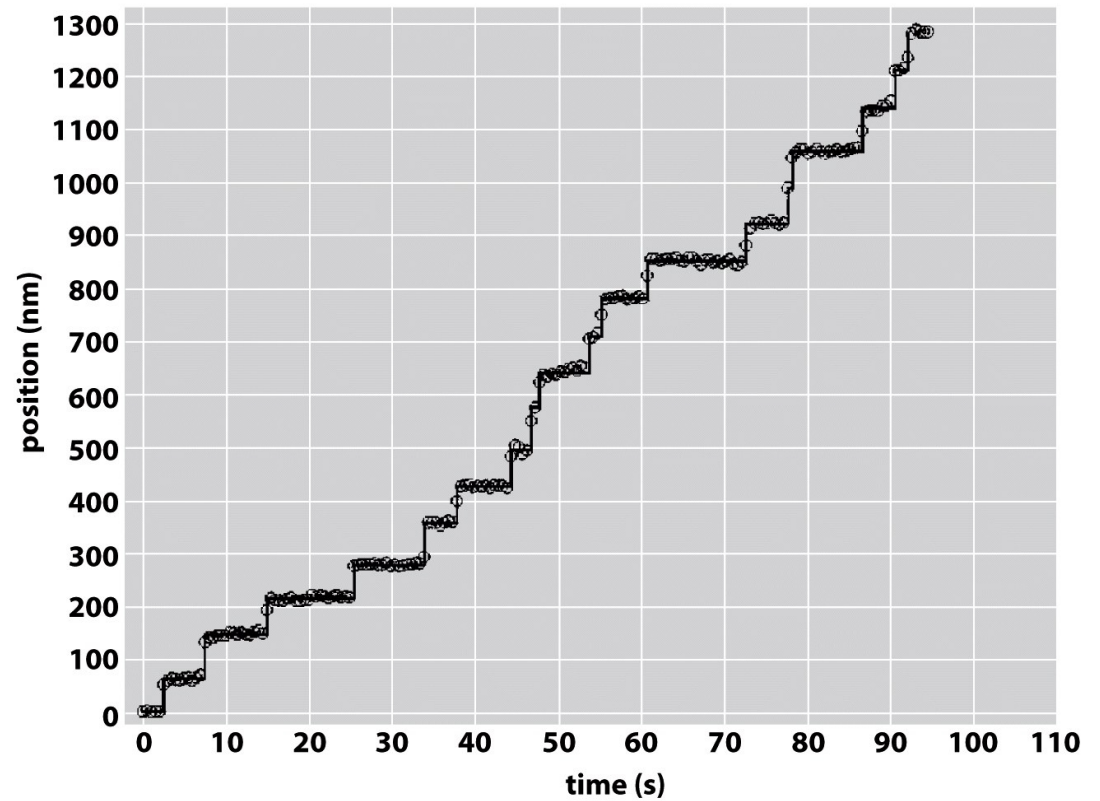


Figure 3.26b Physical Biology of the Cell (© Garland Science 2009)

Position as a function of time

Stokes Drag can be Neglected in Optical Tweezers Experiments

- can measure the speed of the motor as a function of the applied load
- How large is a drag force due to a bead tethered to the motor?
- the bead: 1 μm diameter

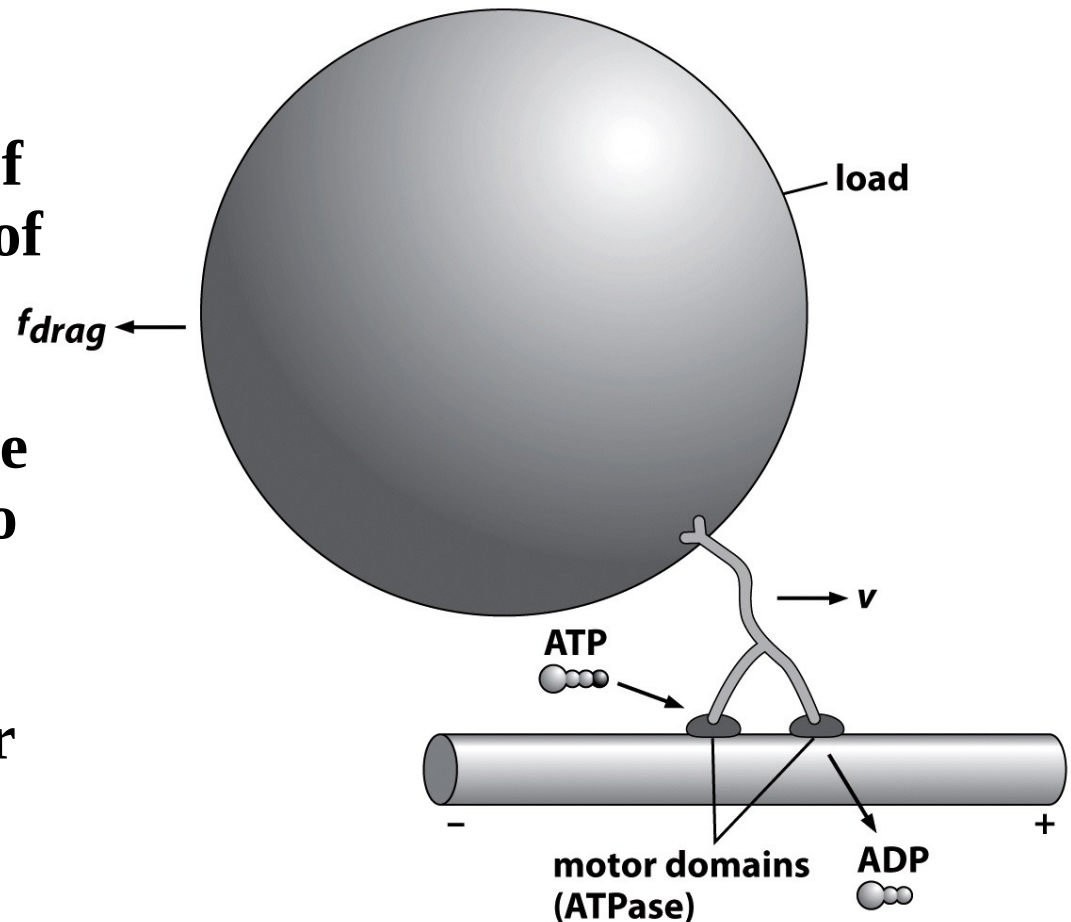


Figure 12.12 Physical Biology of the Cell (© Garland Science 2009)

$$F_S \approx 6\pi \times 10^{-3} \text{Ns/m}^2 \times 5 \times 10^{-7} \text{m} \times 10^{-6} \text{m/s} \approx 10^{-2} \text{pN} \ll 5 \text{pN}$$

Dissipative Time Scales and the Reynolds Number

Consider a damped harmonic oscillator and analyze its behavior at low Reynolds numbers:

$$m \frac{d^2 x}{dt^2} + \gamma \frac{dx}{dt} + kx = 0$$

A low Reynolds number means that the time scale over which the kinetic energy dissipates is small:

$$\frac{m (dx/dt)^2}{\tau_{\text{viscous}}} \approx \gamma (dx/dt)^2$$

**(rate of dissipation on the right is a viscous force times velocity)
Thus we get a simple expression:**

$$\tau_{\text{viscous}} = \frac{m}{\gamma}$$

which means that we can neglect the inertial term in the equation and get

$$\gamma \frac{dx}{dt} + kx = 0 \quad \text{with a solution:} \quad x(t) = x_0 e^{-(k/\gamma)t}$$

**Observation of stopping distances for objects of different sizes:
Consider a spherical object with initial velocity equal to a
diameter of a sphere per second**

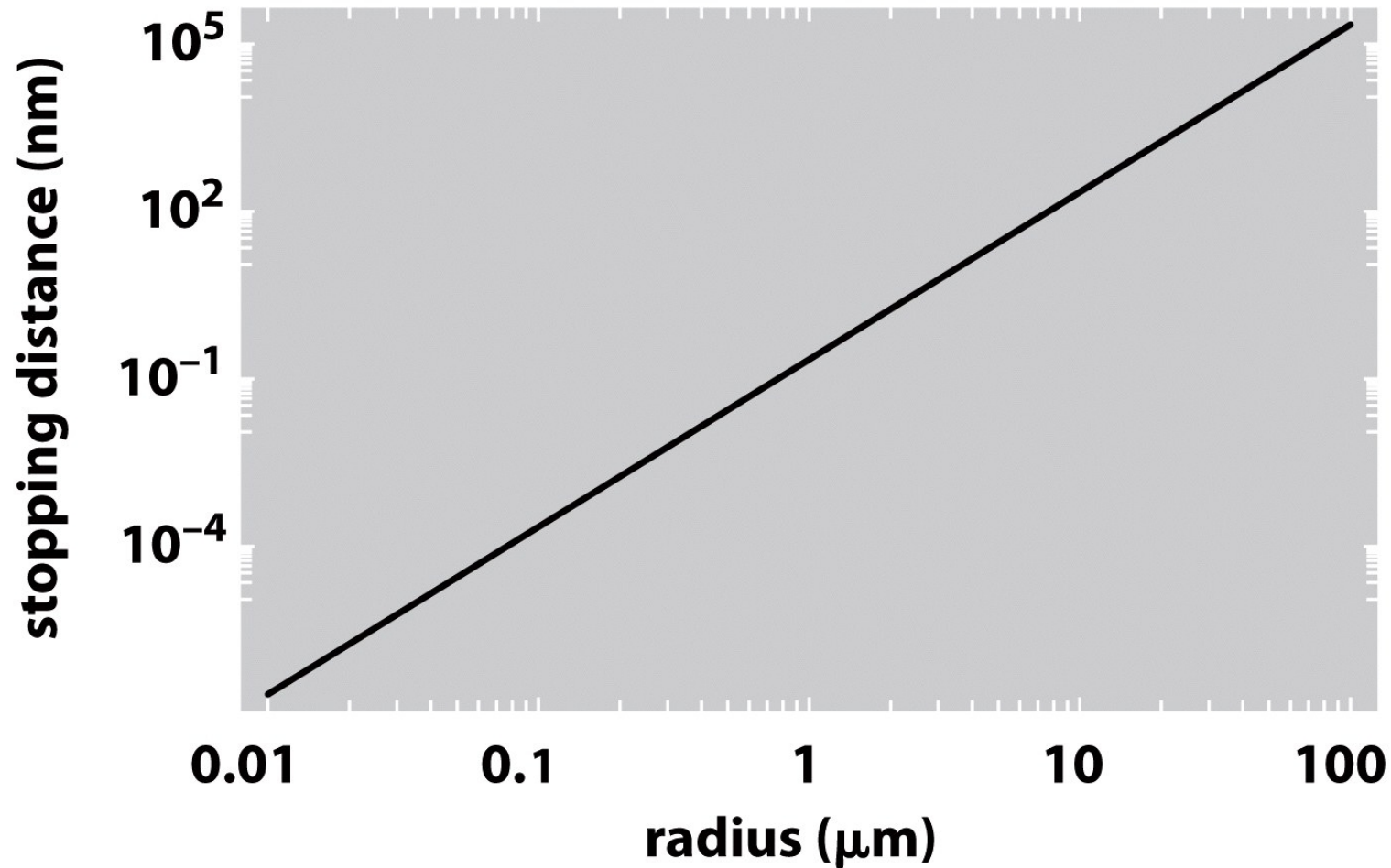


Figure 12.13 Physical Biology of the Cell (© Garland Science 2009)

E. coli moves by rotating its flagella

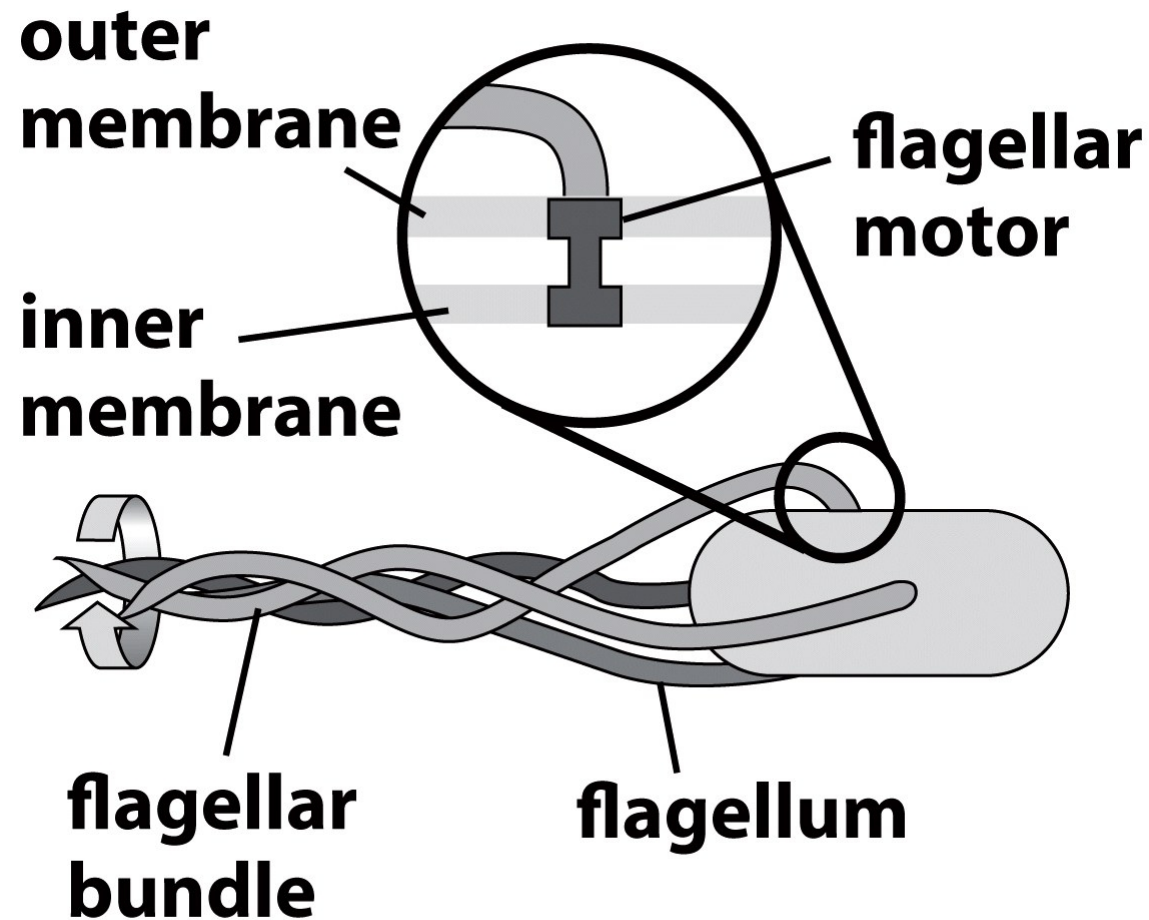


Figure 4.16a Physical Biology of the Cell (© Garland Science 2009)

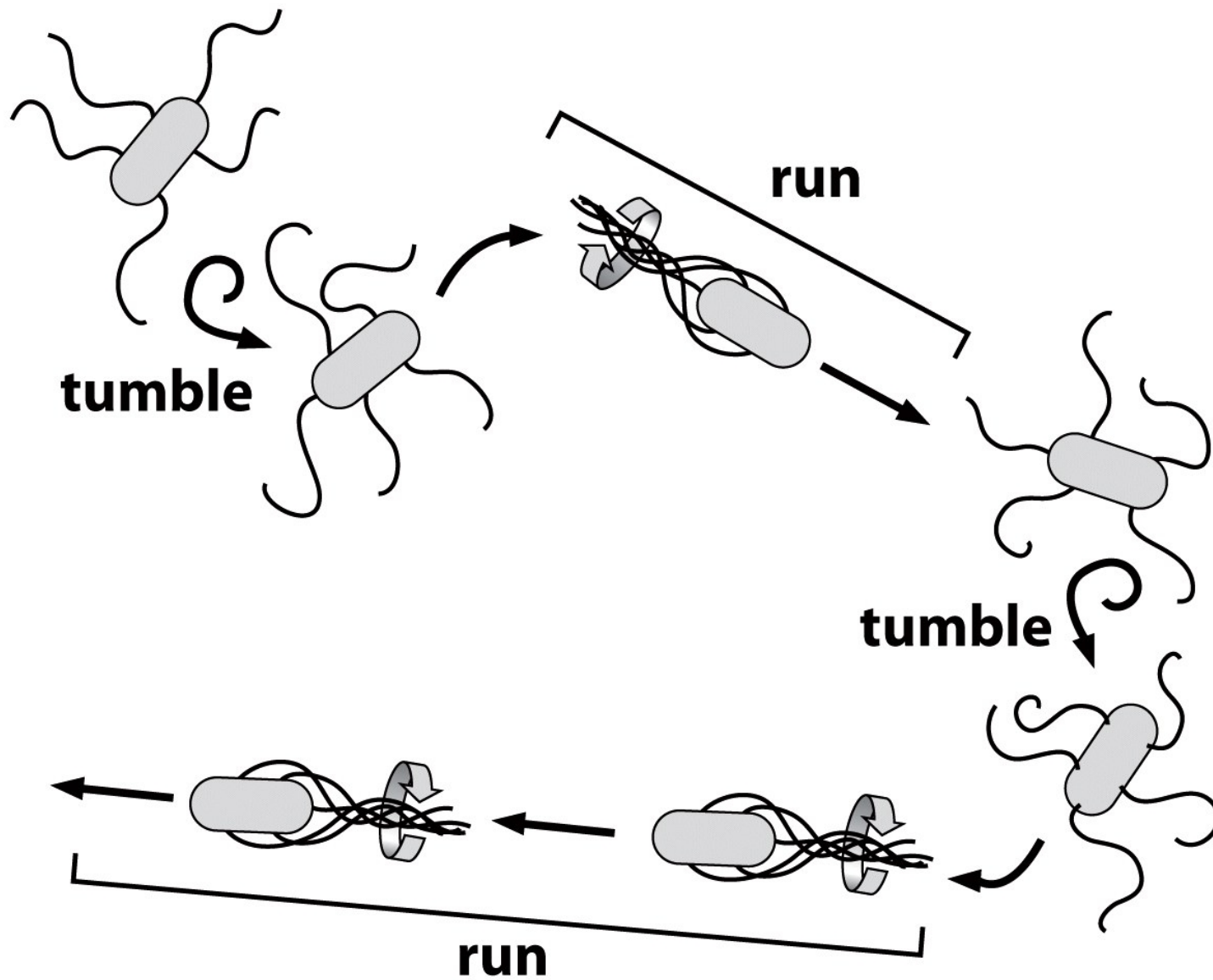


Figure 4.16b Physical Biology of the Cell (© Garland Science 2009)

Helically shaped flagella with a diameter D and pitch P :

$D \approx 0.5 \mu\text{m}$...diameter of the helically shaped flagellum

$P \approx 2 \mu\text{m}$...pitch of the helix (length of one helical turn)

f ...propulsion frequency $v = \pi D f$...linear velocity

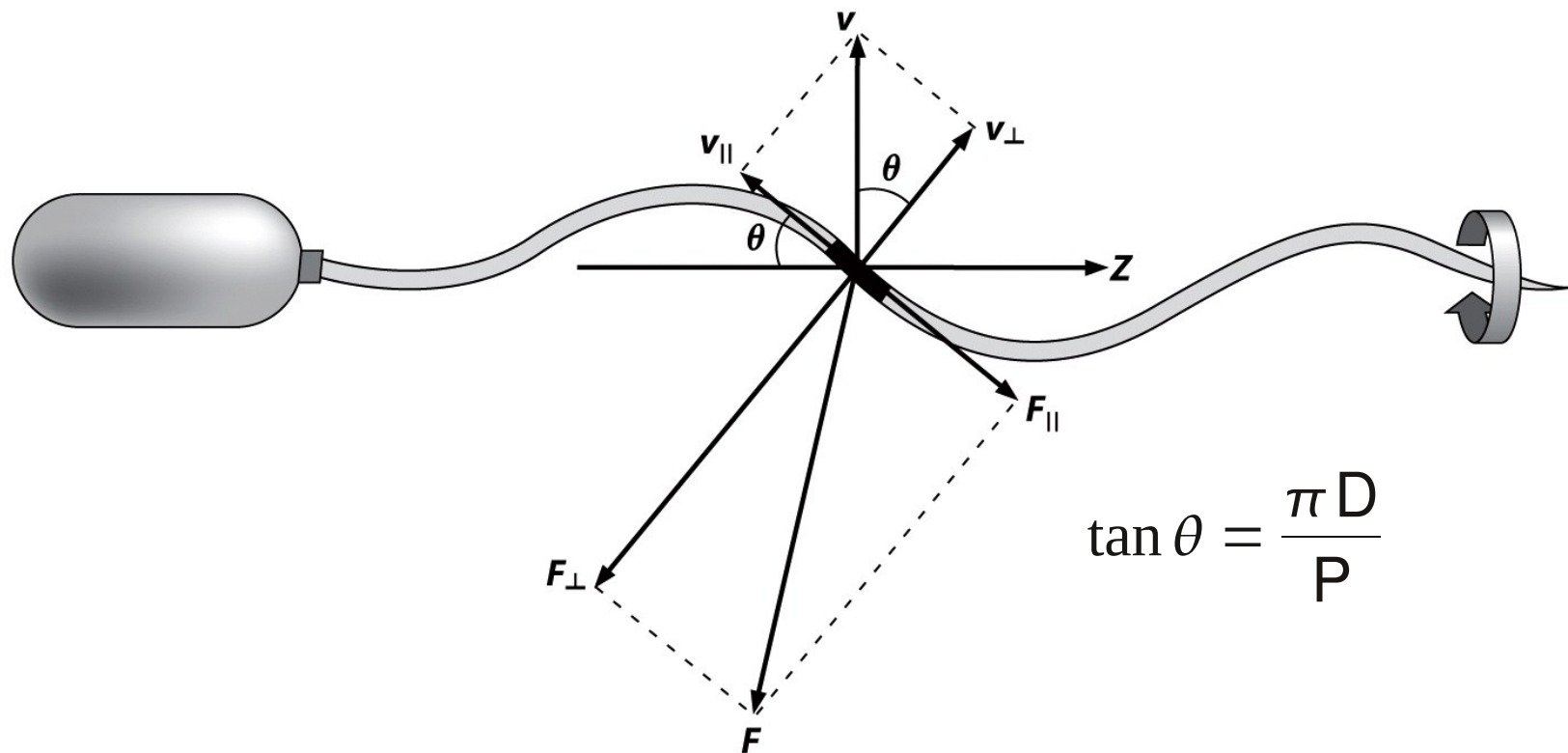


Figure 12.14 Physical Biology of the Cell (© Garland Science 2009)

The propulsive force relies on the difference in the drag coefficients between parallel and perpendicular motion of a rod:

→ **force parallel to the rod is:**

$$F_{\parallel} = \gamma_{\parallel} v \sin \theta \quad , \quad \text{where } \gamma_{\parallel} = 2 \pi \eta L$$

→ **force perpendicular to the rod is:**

$$F_{\perp} = \gamma_{\perp} v \sin \theta \quad , \quad \text{where } \gamma_{\perp} = 4 \pi \eta L$$

→ **the total force projected onto the negative z-direction (direction of motion) is then:** $F_p = -F_{\parallel} \cos \theta + F_{\perp} \sin \theta$

$$F_p = -2 \pi \eta L v \sin \theta \cos \theta + 4 \pi \eta L v \cos \theta \sin \theta = 2 \pi \eta L v \sin \theta \cos \theta$$

→ **the propulsion force is balanced by the drag force:**

$$F_D = 2 \pi \eta L V$$

→ **which results in the final speed of:**

$$V = \pi D f \sin \theta \cos \theta \approx 70 \mu\text{m/s} \quad (30 \mu\text{m/s} \text{ exp.}) \quad (v = \pi D f)$$

Centrifugation and Sedimentation

- **biochemical purification: to separate macromolecules or their complexes from the solution**
- ***centrifugation*: a simple example of diffusion in the presence of drift using the Stokes formula**
- **What is centrifugation? Spinning at up to 100,000 rpm $\sim 10^6$ g**
- **the centrifugal force that all molecules in the solution experience:**
$$m\omega^2 r$$
- **assume: the size of the sample much smaller than the distance from the rotation axis**
- **The mass of the molecule in the solvent needs to be corrected by the mass of displaced solvent: $m \rightarrow (\rho_p - \rho)V$**

The centrifugal force imparts a drift velocity to the biomolecules (low Reynolds numbers): centrifugal force equal to the frictional Force:

$$v_{\text{drift}} = \frac{m g_c}{\gamma} \quad \text{for a spherical molecule: } \gamma = 6 \pi \eta R$$

The quantity m/γ is quoted for different biomolecules in units of Svedberg:

$$1 \text{ svedberg} = 10^{-13} \text{ s}$$

which corresponds to a globular protein with a radius of 1 nm in water. Typical density of proteins: 1.35 g/cm^3

The drift velocity is then a quadratic function of the size of the Biomolecule:

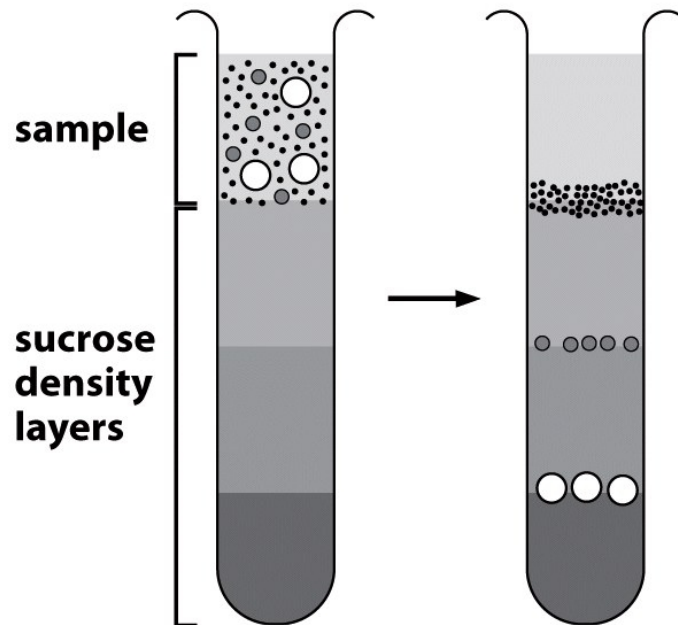
$$v_{\text{drift}} = \frac{(\rho_P - \rho) 4/3 \pi R^3}{6 \pi \eta R} = \frac{2(\rho_P - \rho)}{9 \eta} R^2 g_c$$

which allows for an effective separation of particles based on their respective sizes.

Rate Zonal Versus Isopycnic Centrifugation

(A)

RATE ZONAL CENTRIFUGATION



(B)

ISOPYCNIC CENTRIFUGATION

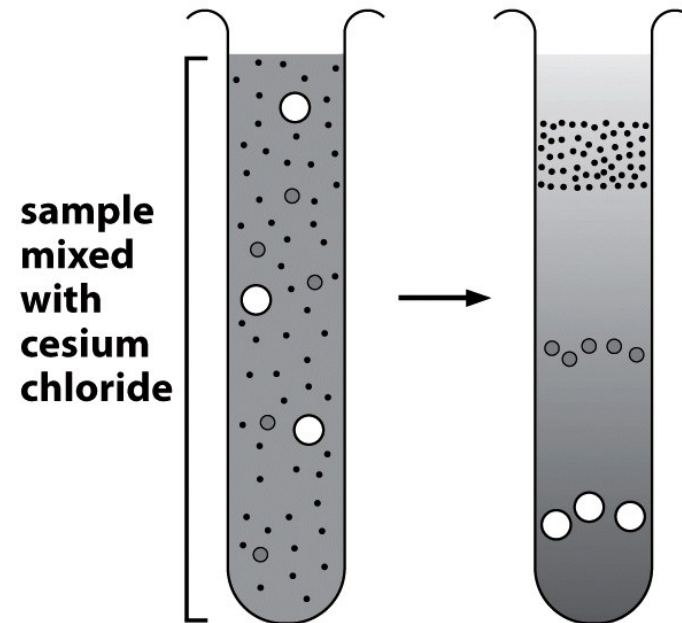


Figure 12.16 Physical Biology of the Cell (© Garland Science 2009)

$$x_c = v_{\text{drift}} t$$

$$\Delta x = \sqrt{2Dt}$$

Diffusion of biomolecules due to thermal fluctuations: a negative effect on the separation. A condition for separation:

$$|v_{\text{drift1}} - v_{\text{drift2}}|t > (\sqrt{2D_1} + \sqrt{2D_2})\sqrt{t}$$

which shows that we just need to wait long enough for two types of molecules to separate:

$$t_{\text{sep}} = \left(\frac{\sqrt{2D_1} + \sqrt{2D_2}}{|v_{\text{drift1}} - v_{\text{drift2}}|} \right)^2$$

However, a test tube has a finite length, so the separation needs to occur before the molecules reach the bottom of the tube, that is, the speed of spinning needs to be large enough:

$$v_{\text{drift}} t_{\text{sep}} < L \quad \rightarrow \quad g_c > \frac{1}{L} \frac{m_1}{\gamma_1} \left(\frac{\sqrt{2D_1} + \sqrt{2D_2}}{m_1/\gamma_1 - m_2/\gamma_2} \right)^2$$

The other separation method: isopycnic centrifugation relies on a density gradient of the solvent and different densities of the various macromolecules that needs to be separated.