

Chapter 13

Speeds of motors

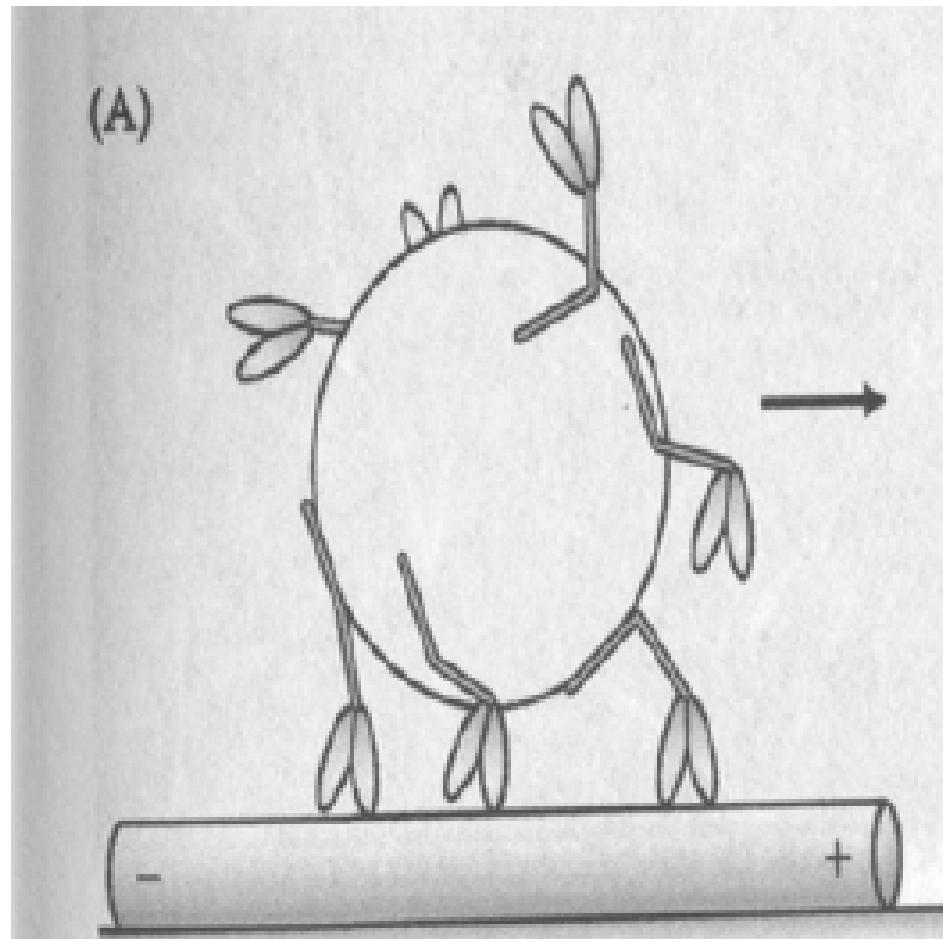
Speeds of motors *in vivo*

- Myosin IB – 200nm/s
- Cytoplasmic dynein – 1000nm/s
- Conventional kinesin – 1800nm/s
- Rowers vs. porters (myosin II vs. kinesin)

In vitro motility assays

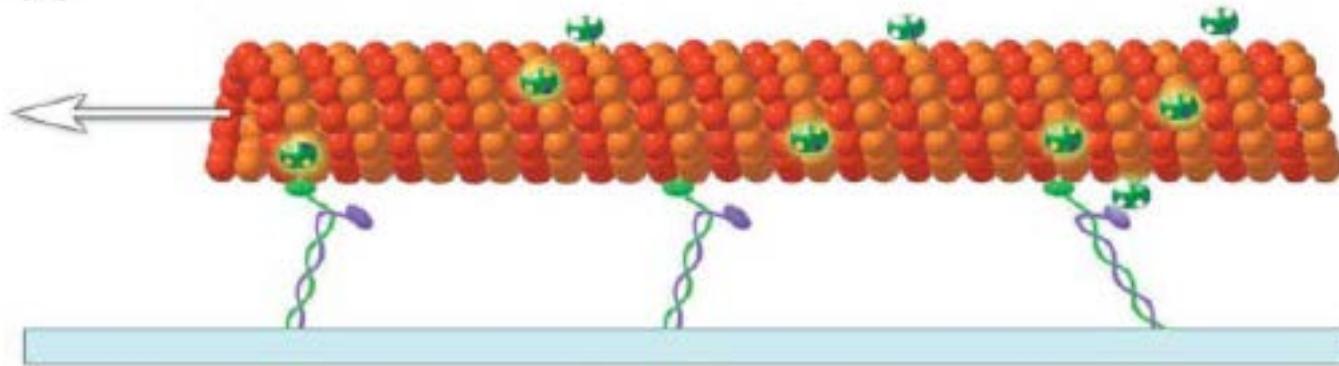
- Sheetz and Spudich first observed the transport of fluorescent beads coated with purified myosin moving along actin.
- Bead Assay, Stepping Assay, Gliding assay

Bead assay

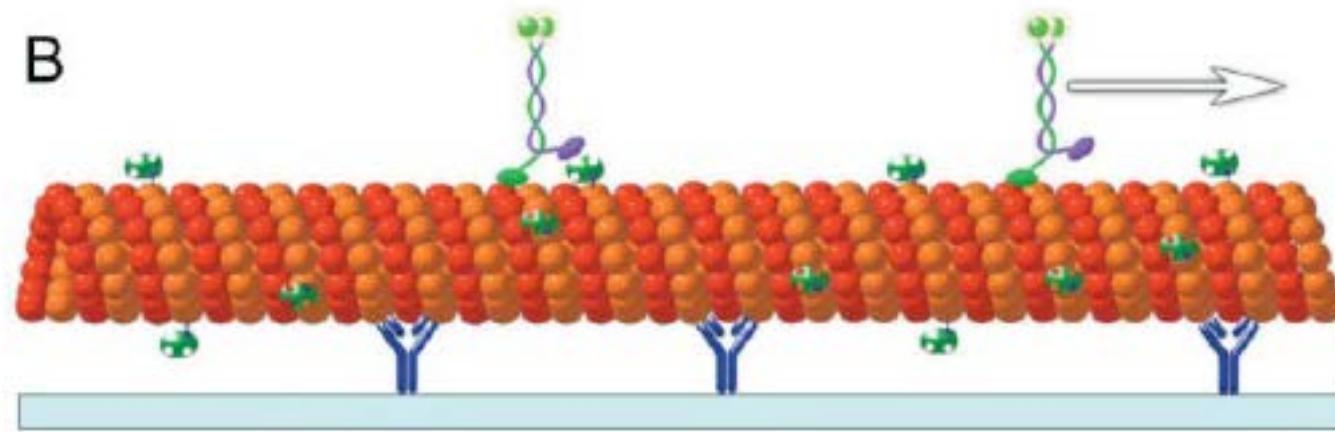


Gliding and stepping assays

A



B

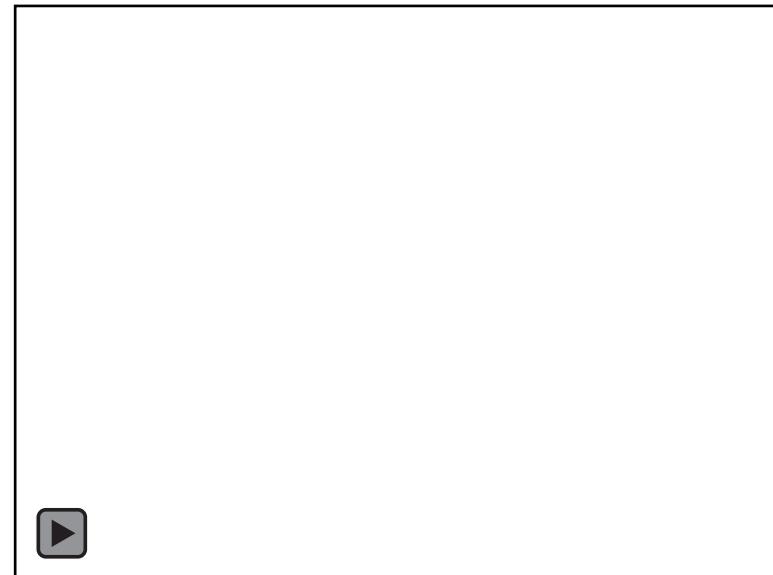


 kinesin-1	 streptavidin	 biotin
 microtubule	 α -tubulin antibody	

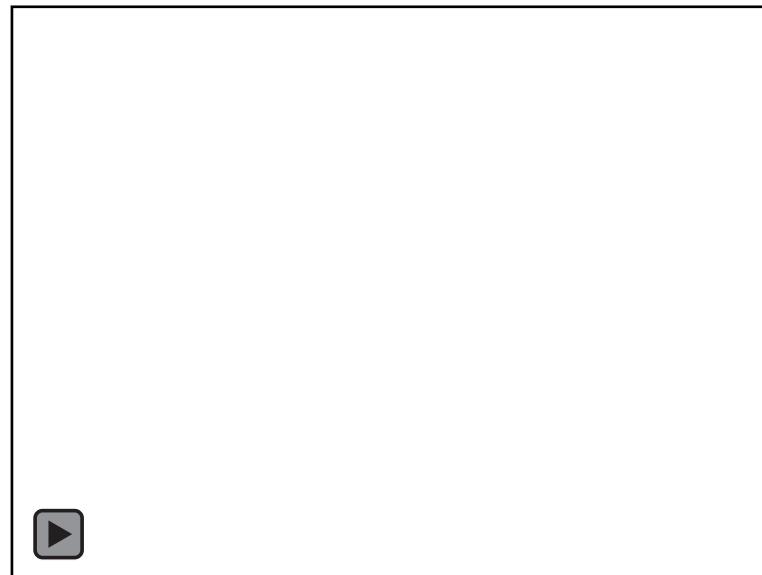
<https://www.youtube.com/watch?v=y0QCkObJIto>



<https://www.youtube.com/watch?v=2UdPsVSWwrE>



<https://www.youtube.com/watch?v=FzY86nkgumU>



In vitro motility assays

- Polarity-marked microtubules
- Mechanical loads very small

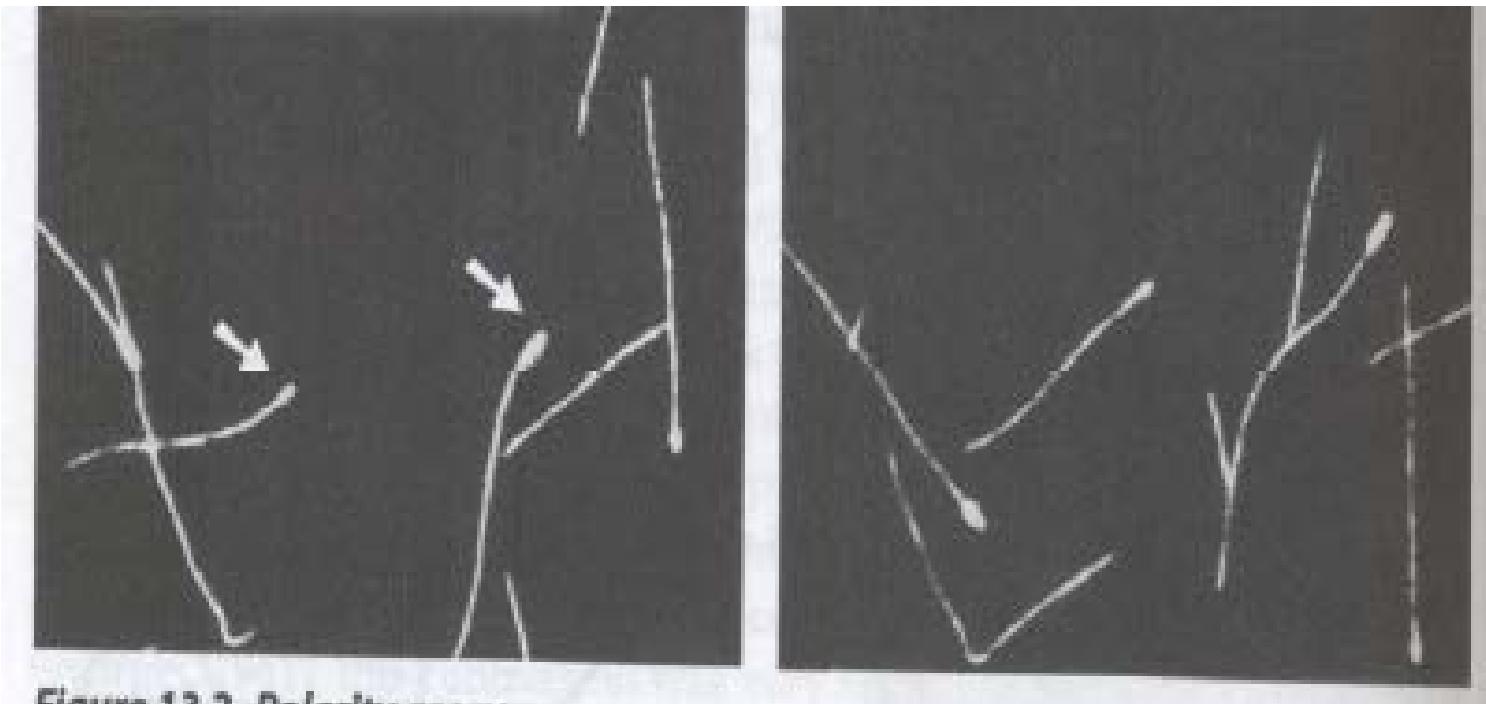


Figure 13.9. Polarized microtubules.

Processive and non-processive motors

- Conventional kinesin is a processive motor, 100s of 8-nm steps
- Kinesin hydrolyzes about 125 molecules of ATP following initial binding to MT (biochemically verified) – 1 step per ATP
- Ncd is a non-processive, minus end directed motor protein

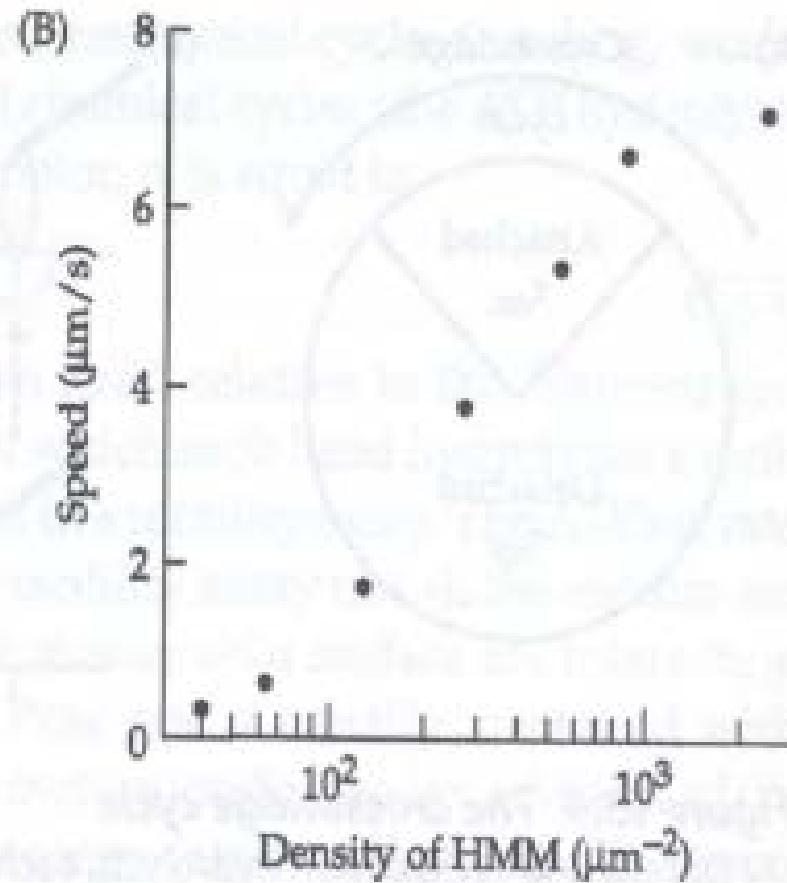
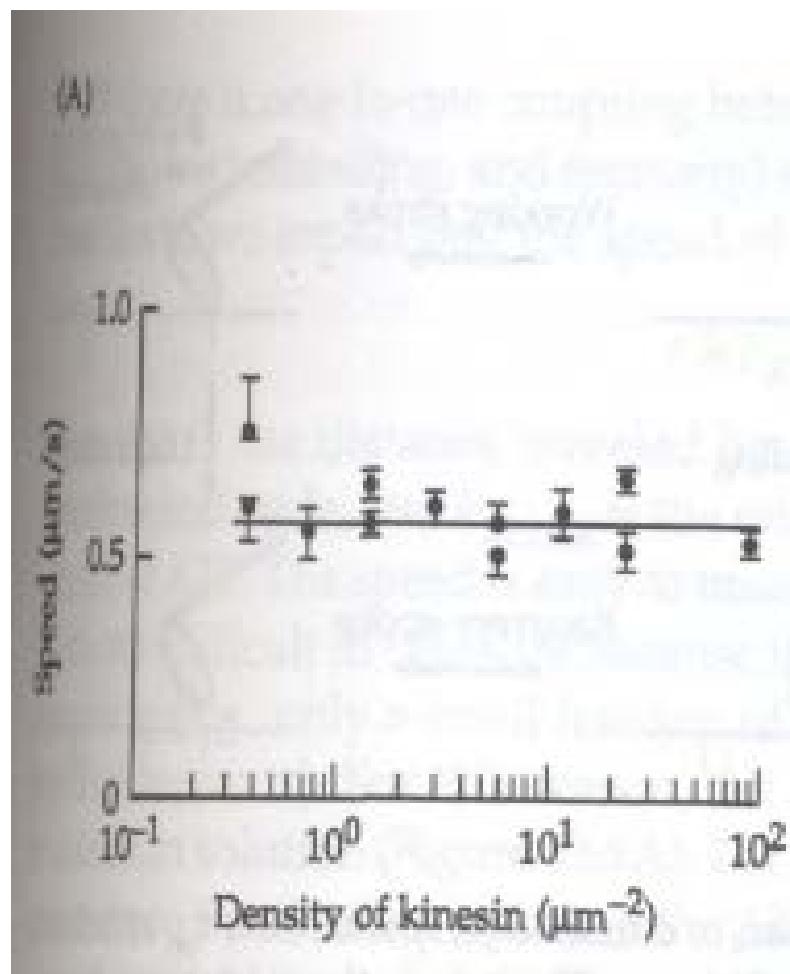
Processive and non-processive motors

- Muscle myosin II is non-processive
- Threshold density of myosin II is required for continuous motility of actin filaments → 4000 molecules / μm^2
- Myosin V and VI are processive
- Cytoplasmic dynein is processive, but outer-arm axonemal dynein is not.

Processive and non-processive motors

- Processivity/non-processivity -- dependence of speed of movement on the number of participating molecules
- Speed of kinesin – independent of no. of motors/filament length → co-ordination required
- Myosin – actin gliding occurs below threshold value when methyl cellulose added (prevents diffusion of filaments perpendicular to their long axes)

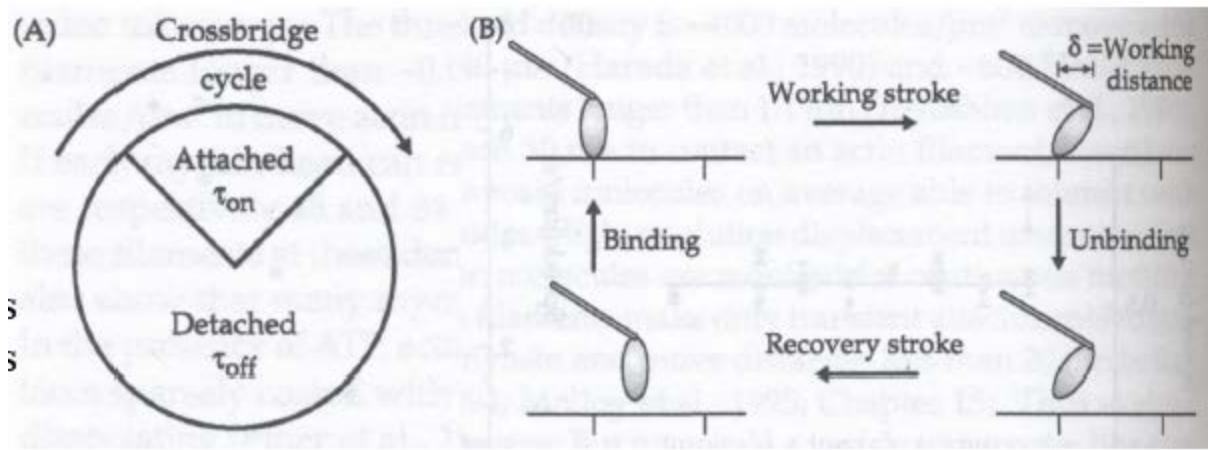
Processive and non-processive motors



Hydrolysis cycle and duty ratio

- Motors move along filaments through distances that are large compared to molecular dimensions → the motor reaction must be cyclic in which motor repeatedly bind and unbinds the filament

Hydrolysis cycle and duty ratio



- Average time motor domain spends attached to filament - τ_{on} (working stroke)
- Average time motor remains detached – τ_{off} (recovery stroke)
- Duty ratio – fraction of the time that each head spends in its attached phase:

$$r = \frac{\tau_{on}}{\tau_{on} + \tau_{off}} = \frac{\tau_{on}}{\tau_{total}}$$

Hydrolysis cycle and duty ratio

- Minimum no. Of heads required for continuous movement (N_{min})

$$r \cong \frac{1}{N_{min}}$$

- Guarantees that there will usually be at least one head bound to the filament
- Eg. Conventional 2-headed kinesin, dynein, myosin V - r must be at least 0.5 for each head.
- But for skeletal muscle myosin and outer arm dynein, large assemblies of 50-100 crossbridges, r is small ~0.01 to 0.02

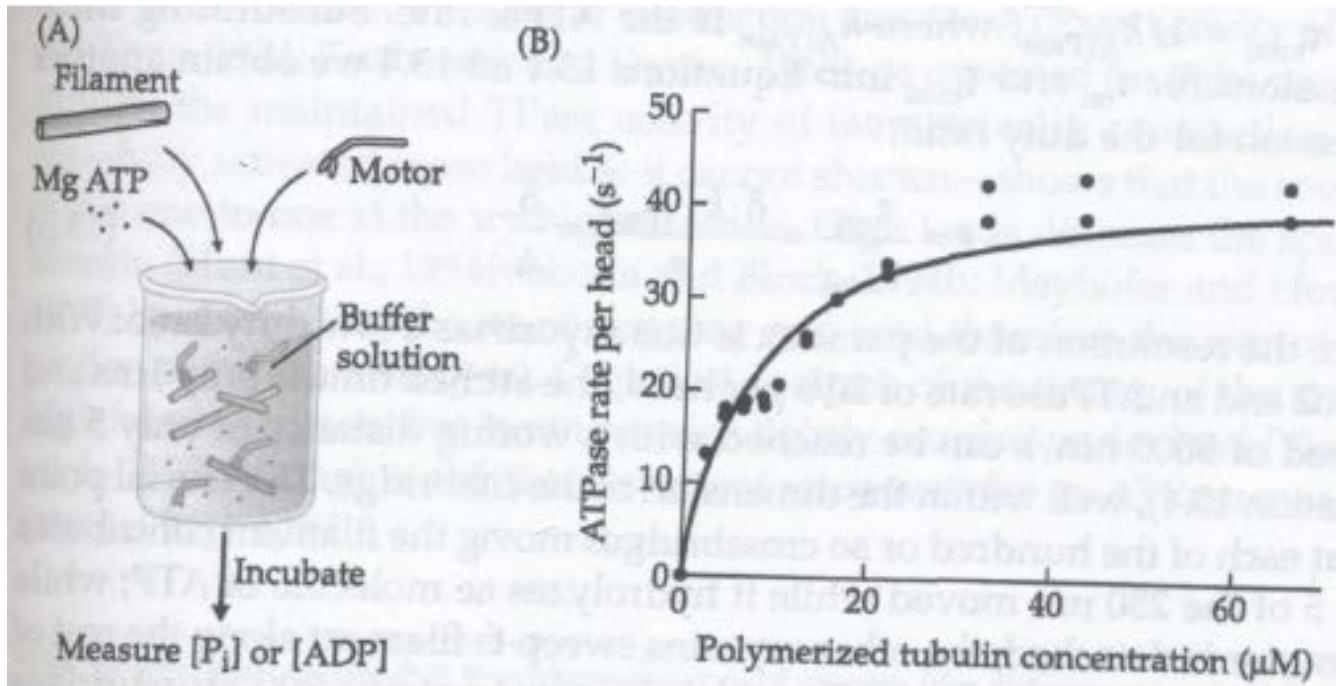
Hydrolysis cycle and duty ratio

- If there is one-to-one coupling between mechanical cycles and chemical cycles, then we expect speed of the motor:

$$v = k_{ATPase} \Delta$$

- k_{ATPase} – rate at which each head hydrolyzes a molecule of ATP, Δ – distance moved by each head per mechanical cycle.

Measurement of k_{ATPase}



- Maximum ATPase rate measured in solution (k_{cat}) is assumed to correspond to the ATPase during motility

Speed of motors and k_{cat}

Motor	Speed ^a in vivo (nm/s)	Speed ^b in vitro (nm/s)	In vitro ATPase ^c (s ⁻¹)	Function
Myosins				
1. Myosin IB	ND ^d	200	6	Amoeboid motility, hair cell adaptation
2. Myosin II	6000	8000	20	Fast skeletal muscle
3. Myosin II	200	250	1.2	Smooth muscle contraction
4. Myosin V	200	350	5	Vesicle transport
5. Myosin VI	ND	-58	0.8	Vesicle transport?
6. Myosin XI	60,000	60,000	ND	Cytoplasmic streaming in algae
Dyneins				
7. Axonemal	-7000	-4500	10	Sperm and ciliary motility
8. Cytoplasmic	-1100	-1250	2	Retrograde axonal transport, mitosis, transport in flagella
Kinesins				
9. Conventional	1800	840	44	Anterograde axonal transport

No direct relation!

Step size vs. k_{cat}

$$\Delta = \frac{v}{k_{ATPase}} = \begin{cases} 400 \text{ nm/ATP} & \text{For skeletal muscle myosin} \\ 450 \text{ nm/ATP} & \text{For outer-arm dynein} \end{cases}$$

- For HMM, 190nm/ATP , order of magnitude more than the crossbridge → ‘step-size paradox’ leading to the suggestion that myosin might take multiple steps for each ATP hydrolyzed.
- Duty ratio provides a simple explanation for the step-size paradox.

Hydrolysis cycle and duty ratio

- Filament moving with constant speed v , over array of fixed motor proteins.
- Assuming enough motors to ensure continuous motility (1 kinesin/50-100 myosin heads)

$$v = \frac{\delta}{\tau_{on}}$$

δ – working distance

τ_{on} – time for which head is attached

Hydrolysis cycle and duty ratio

- Total cycle time $\tau_{total} = \frac{1}{k_{ATPase}}$

$$\left. \begin{array}{l} r = \frac{\tau_{on}}{\tau_{total}} \\ \tau_{on} = \delta/v \\ \tau_{total} = 1/k_{ATPase} \end{array} \right\} \Rightarrow r = \frac{\delta k_{ATPase}}{v} \xrightarrow{v=k_{ATPase}\Delta} r = \frac{\delta}{\Delta}$$

(Cycle driven by ATP hydrolysis)

Myosin: $\left[\begin{array}{l} r = 0.02 \\ k_{ATPase} = 20/\text{s per head} \\ \tau_{on} = 1 \text{ ms} \\ v = 5000 \text{ nm/s} \\ \delta = 5 \text{ nm} \end{array} \right] \longrightarrow \text{within the dimension of the crossbridge}$

Hydrolysis cycle and duty ratio

- Each of the 100 or so crossbridges that move the filament contribute only 5nm of the 250nm moved while it hydrolyzes one molecule of ATP
- Each thick filament contains 300 crossbridges, for duty ratio of 2% in a rapidly contracting muscle → at all times, 6 or so crossbridges maintain contact between thin and thick filaments

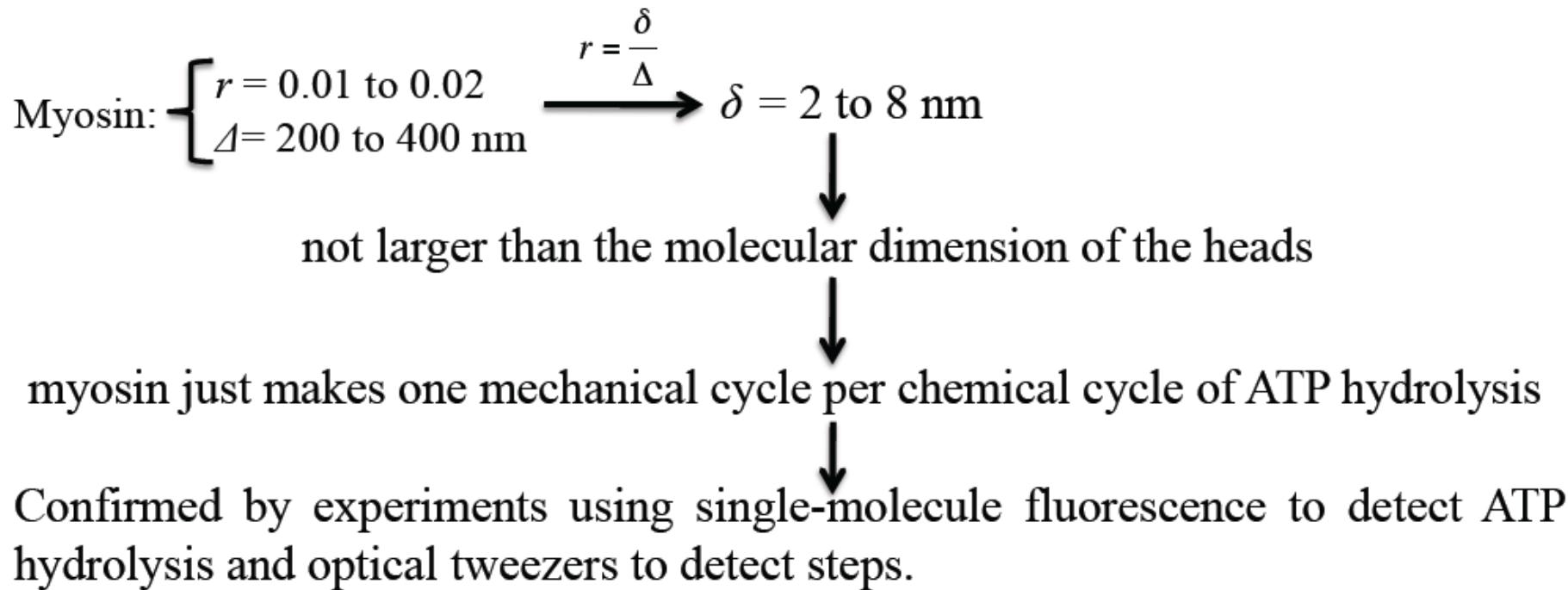
Processive motors

- Since kinesin has high r , high ATPase rate is needed to attain moderate speeds

kinesin:
$$\begin{cases} k_{ATPase} = 50/\text{s per head} \\ v = 800 \text{ nm/s} \\ \delta = 8 \text{ nm} \end{cases} \longrightarrow r = 0.5$$

Consistent with the duty ratio required to account for kinesin's processivity.

Estimation of working distance



Analogy to walking and running

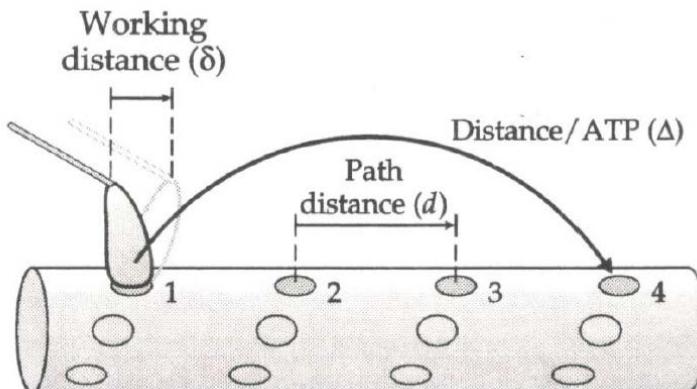
- Gait of a biped is walk if “the feet move alternatively with one foot not clear od the ground before the other touches”
- During run, times when both feet are off

Chapter 15

Steps and forces

Distances that characterize a motor reaction

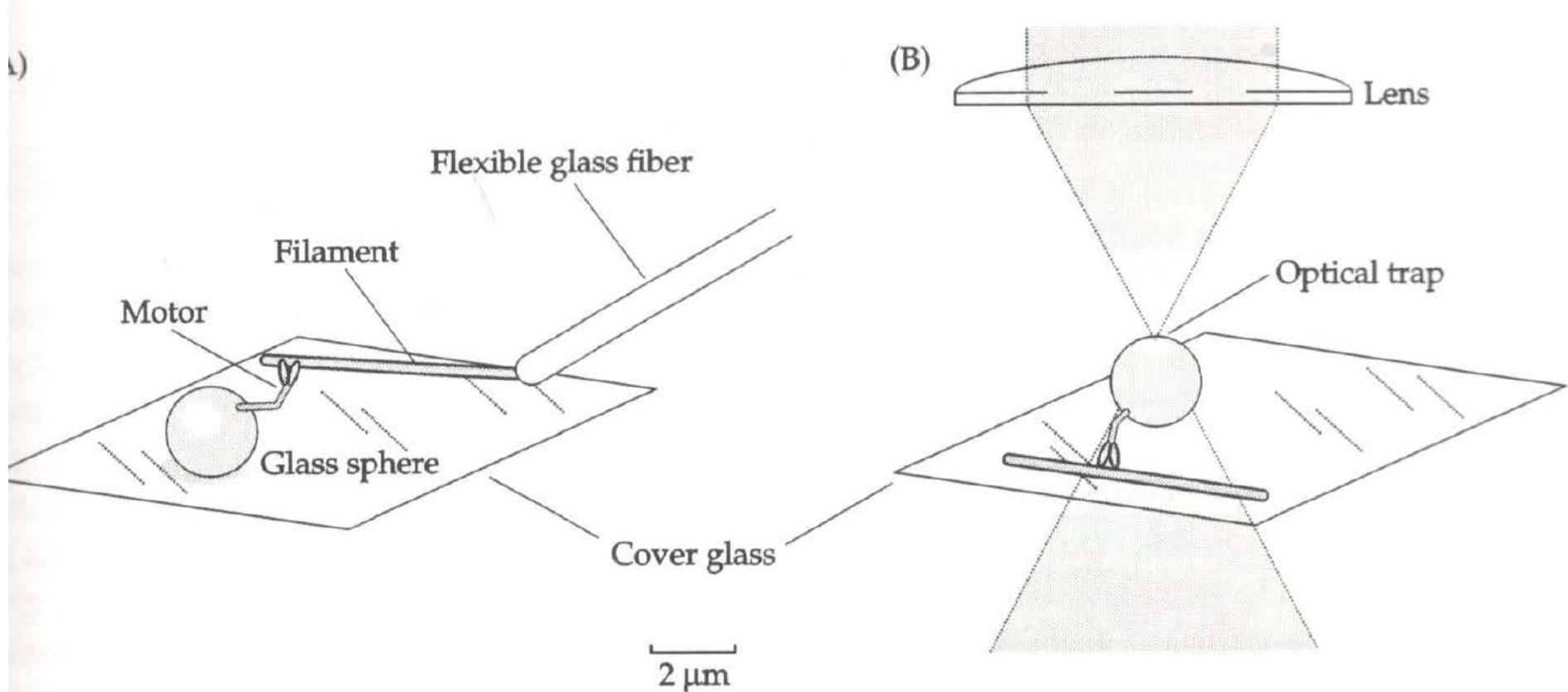
- Working distance δ (powerstroke distance)
- Distance per ATP for each motor domain Δ (Speed of movement/ATPase rate per head)
- Path distance d (Distance between the consecutive binding sites)



Single motor techniques

- Filament assay – MT held in force transducer and presented to a motor that is fixed to the surface
- Bead assay – Motor attached to a bead that is held in a force transducer and bead presented to a filament fixed on the surface
- Force transducers – cantilevered glass rods, AFMs or optical tweezers, must be able to produce and monitor forces in the piconewton range

Single motor techniques



Cantilevered springs

- Slender glass fiber whose base is held in a micromanipulator – spring exerting known forces on the motor (principle of AFM). Stiffness of cantilevered fiber:

$$\kappa_f = \frac{3\pi}{4} \frac{Er^4}{L^3} \text{ (Chapter 6)}$$

E- Young's modulus, L-length of fiber, r-radius

- Stiffness of 0.001 to 1 pN/nm with glass rods (E ~ 70 GPa) of 100 to 200 μm and radius 100-250 nm

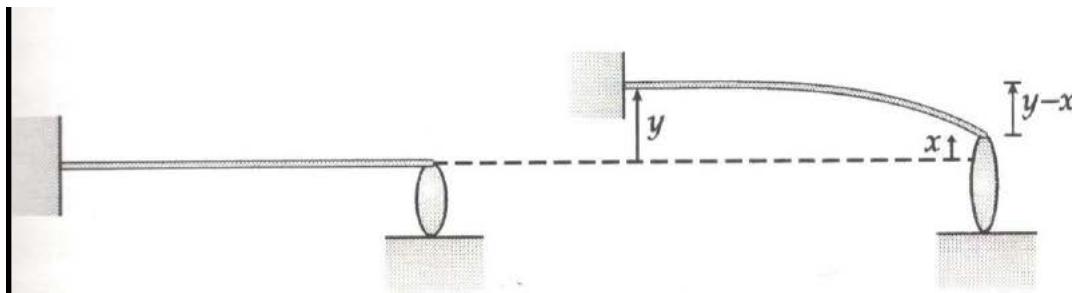


figure 15.3 Cantilever force transducer

The force equals $F = \kappa_f(y - x)$, where κ_f is the stiffness of the fiber, y is the displacement of the base of the fiber, and x is the displacement of the tip of the fiber, which is also equal to the displacement of the attached point of the protein. $y - x$ is the flexion in the fiber.

Cantilevered springs

- Time constant of a cantilevered spring

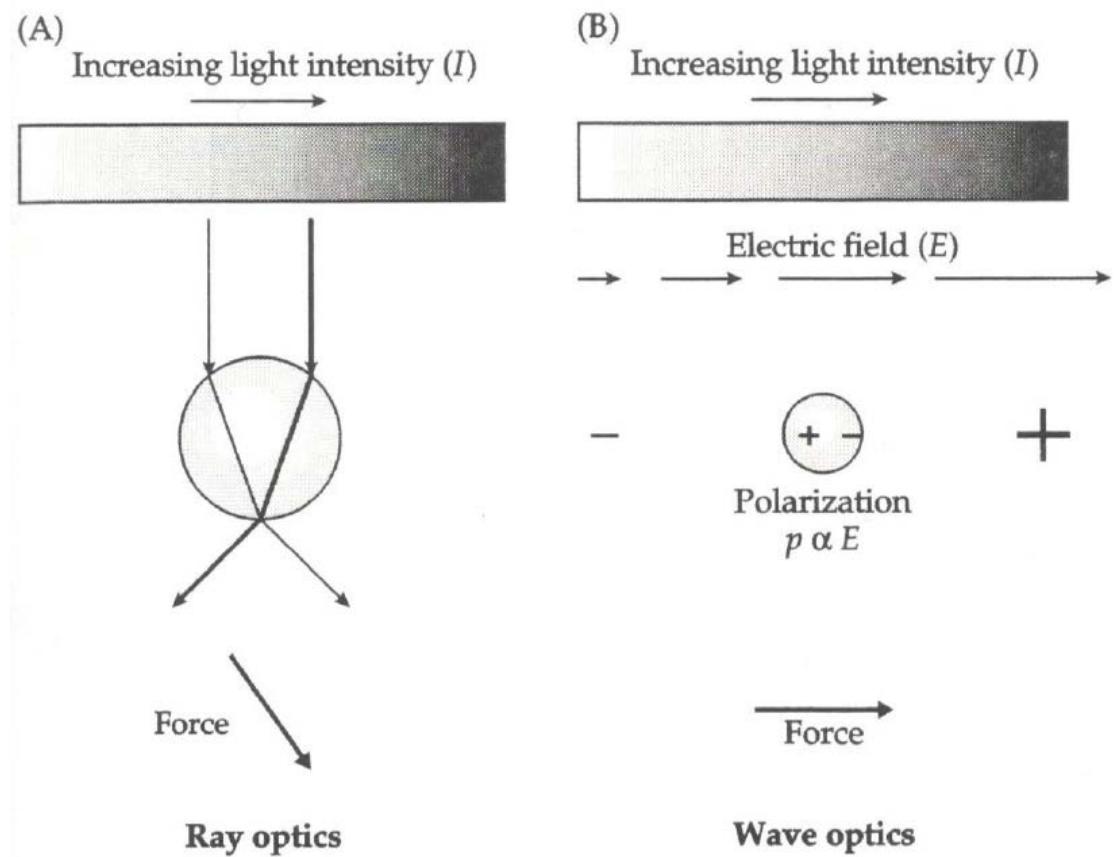
$$\tau \simeq 0.2 \frac{\eta \cdot L^4}{E \cdot r^4} \text{ (Chapter 6)}$$

η – Viscosity of the solution

- Time constant for glass fibres is $\sim 1\text{ms}$ \rightarrow small enough to resolve individual transitions in hydrolysis cycle
- Modify radius and length to get desired time constants and stiffnesses

Optical tweezers

- Laser beam focused down to a diffraction-limited spot using high NA lens
- Dielectric non-absorbing glass or plastic bead experiences a force that will tend to move it into the region of highest intensity (focus of the laser)
- If size of particle $\gg \lambda$, force due to change in momentum of the photons refracted by the particle.
- If size of particle $\ll \lambda$, force due to polarization of the particle induced by electric field component of the light wave



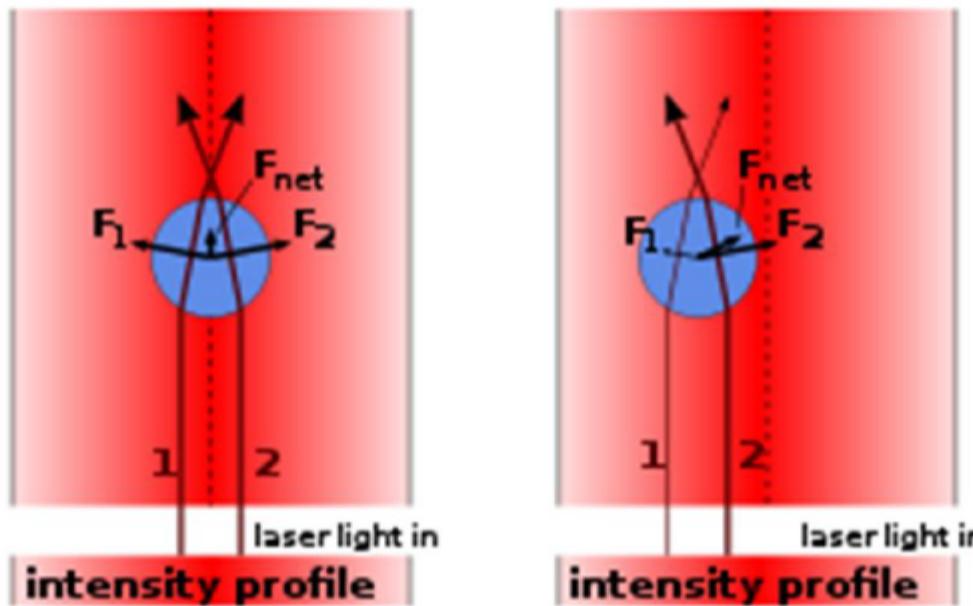


Figure 3: Lateral gradient forces on a particle in a Gaussian beam. The particle on the beam axis (left) has no net force in the lateral direction because the forces from Ray 1 and Ray 2 are equal from each side. The particle to the left of the beam axis (right) has a net force to the right since the higher intensity Ray 2 has a greater change in momentum than Ray 1.

Optical tweezers

- Force exerted by optical trap

$$F_{opt} = Q \frac{n P}{c}$$

- F_{opt} - force associated with the absorption of light (photon pressure)
- $n P / c$ – force exerted by light on a perfect absorber (c/n – speed of light in the medium, P -power)
- Q - dimensionless constant, trapping efficiency

Optical tweezers

- If the particle does not move very far from the centre of the trap, the trap will behave like a spring with spring constant:

$$\kappa_{opt} \sim Q \frac{n^2 P}{\lambda c}$$

- Spring constants in the range – 0.01 to 0.1 pN/nm using laser powers of $\sim 100\text{mW}$ on glass or plastic beads ($n=1.50$ or 1.57) of dia $1\mu\text{m}$

Optical tweezers

- Physical size of the trap is determined by λ and size of the object. Usually λ of around 1 μm used since less photodamage
- Traps much softer, produce less force than cantilever beams and AFMs
- But shorter time constants – better temporal resolution

Displacements

- Photodiode detector
- Accurate measurement of centre of mass of fiber or bead → mean of distribution is can be measured very precisely even if the distribution itself is quite broad.
- Precision of the displacement measurement is limited only by the number of photons that are counted (Appendix 15)

Calibration

- Thermal fluctuations of force transducers are picked up by the detectors
- This is used for the calculation of the stiffness of the force transducer in situ

Single molecule fluorescence

- TIRF and laser scanning confocal
- Limitations – limited lifetime of fluorophores
→ limited spatial and temporal resolution

Steps, Paths and Forces

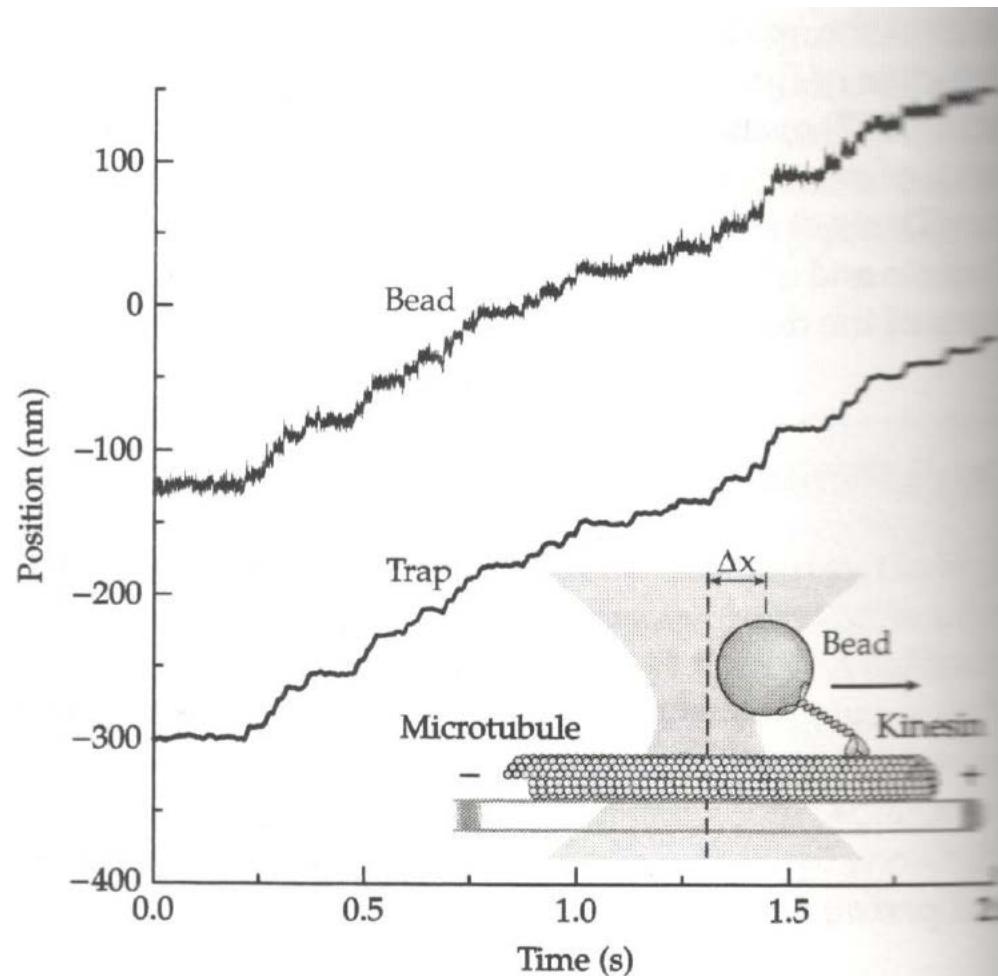
- Displacement sensitivity of optical tweezers, glass fibers and AFMs – 1nm sufficient for measuring step size of motor proteins
- Force sensitivity $\sim 1\text{pN}$, sufficient to resolve force of single motor.

Conventional kinesin

- Conventional kinesin is processive, pulling a $1\mu\text{m}$ glass bead 100s of nm on MT, even when under load of optical trap.
- At high force, motion is not smooth
- Steps have amplitude of 8 nm → kinesin moves from 1 tubulin dimer to the next along a protofilament (length of dimer – 8nm)
- No switching to neighbouring protofilament under low motor densities – no changing lanes by single motor

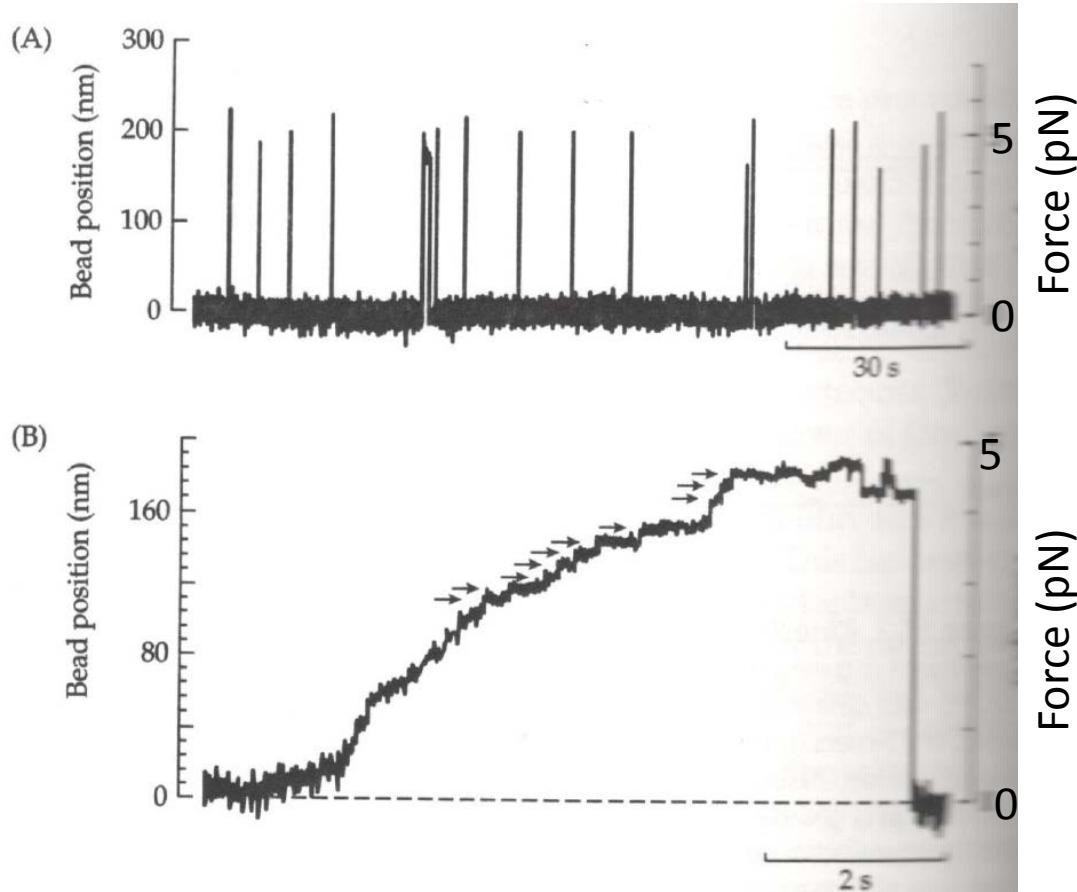
Conventional kinesin

- Maximum force that kinesin can work against $\sim 6\text{pN}$ → Measured by making kinesin walk away from the centre of the trap
- As distance increases, load increases and speed decreases, eventually leading to stalling of the motor. Stall force – 4 to 8 pN have been measured
- Speed of movement decreases \sim linearly with increase in opposing force and maximum force independent of ATP concentration and temperature



Conventional kinesin

- Maximum work done per step = $6\text{pN} \times 8\text{nm} = 48\text{pN}\cdot\text{nm}$
- Energy derived from hydrolysis of ATP = 100 pN.nm
- Therefore, maximum efficiency of kinesin $\sim 50\%$
- Maximum force is only 50% of the thermodynamic force (= chemical free energy/step size, for kinesin, thermodynamic force = $100\text{pN}\cdot\text{nm}/8\text{nm}=12.5\text{pN}$)
- Rate of forward stepping ~ 0 at high loads

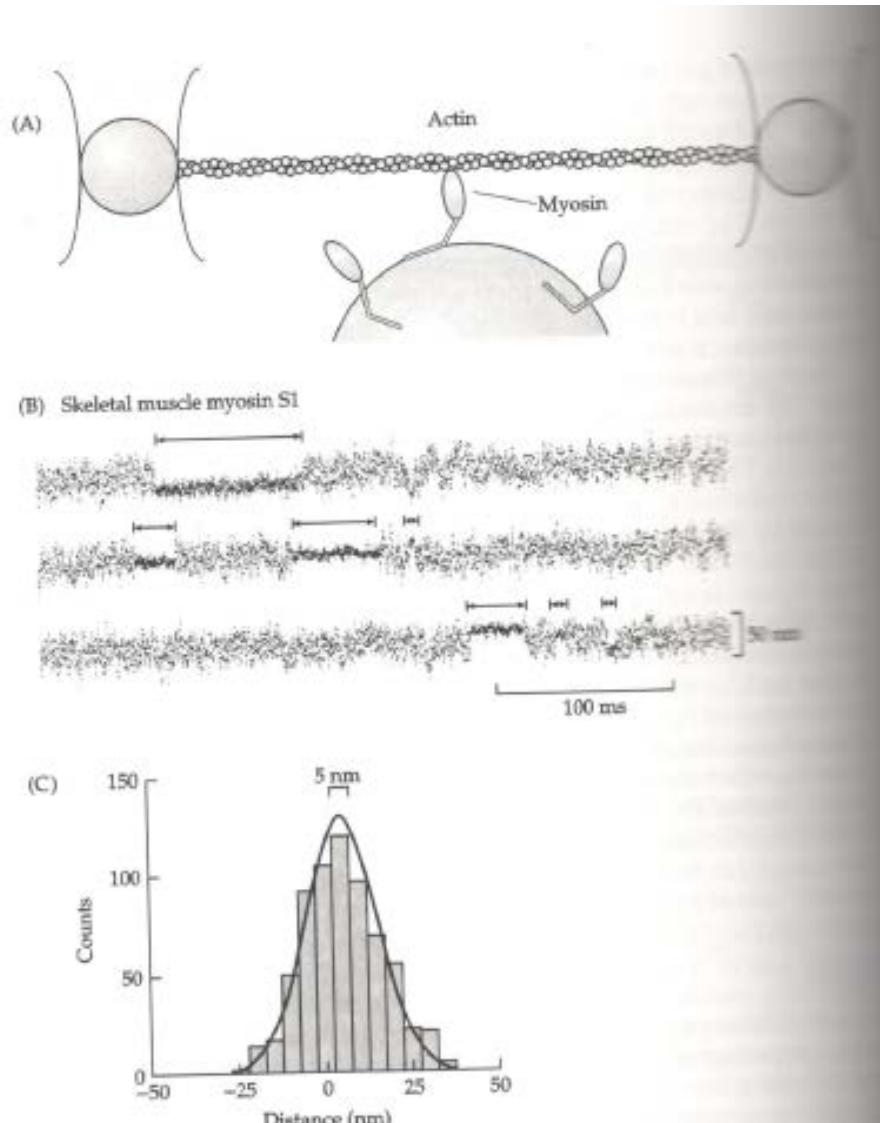


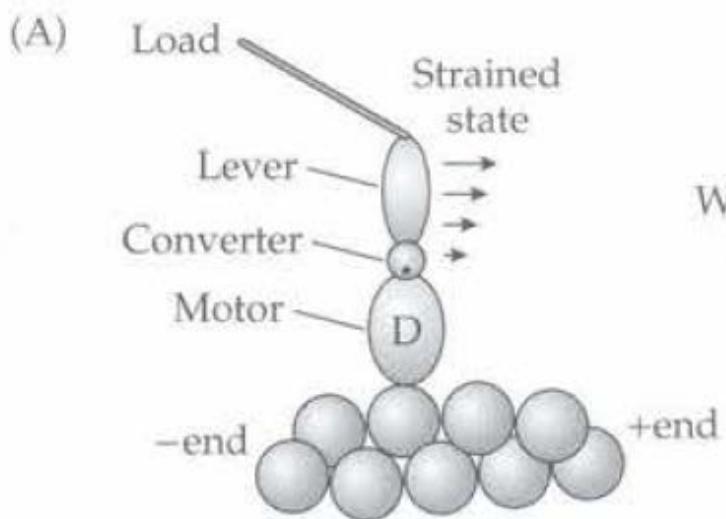
Myosin II

- Only interacts transiently with actin and does not step
- Discrete binding events observed → decrease in thermal fluctuations of the actin filament
- Not processive motor
- Binding of myosin displaces actin from average position, mean displacements + or –
→ Large thermal fluctuations of the actin filament sweep binding sites past the fixed motors; motor binds to filament and holds it in place, displacements can be + or – because fluctuations are symmetric

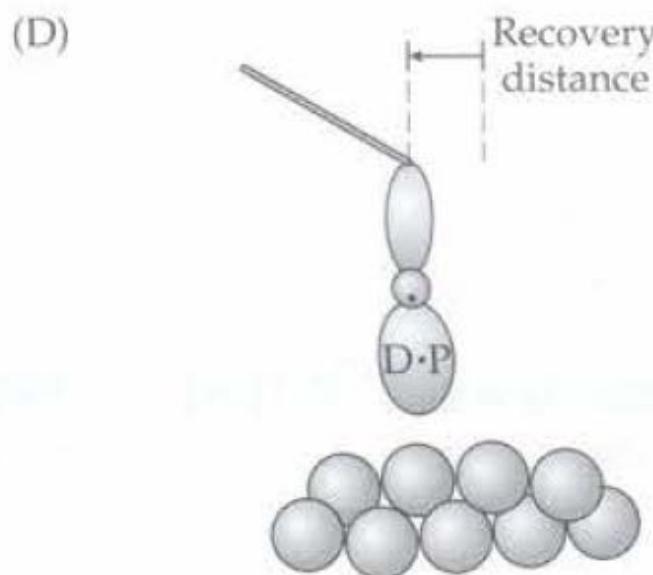
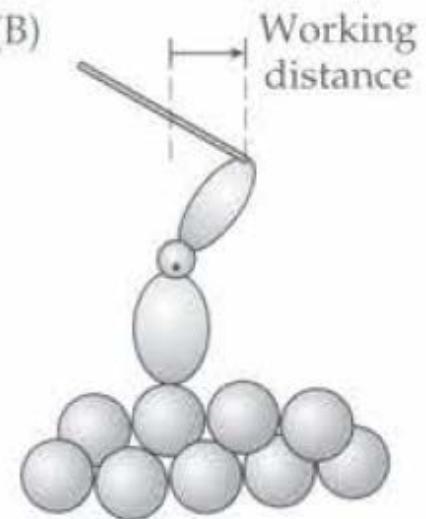
Myosin

- Despite variability in displacement, directional bias for myosin → myosin undergoes a directed conformational change of 5nm (working distance)
- Can be confirmed by rotating filament by 180 degrees
- Length of the rotating lever altered by replacing the light chain binding domain with artificial levers of various lengths → changes amplitude of the step
- Maximum force exerted by myosin crossbridge $\geq 10\text{pN}$





Working stroke



Recovery stroke
(progression to next binding site)



Structural basis for duty ratio

- Relation between the distance per ATP and the path distance

$$\Delta = n \cdot d$$

n- is an integer ≥ 0 .

- n=0 corresponds to futile ATP hydrolysis event, with no associated displacement (eg.: under high load)
- If $n > 1$, the motor jumps over one or more of the stepping stones (myosin at low load)

Structural basis for duty ratio

- Duty ratio is working distance divided by distance per ATP, $r = \delta/\Delta$
- Structural expression for duty ratio:

$$r = \frac{\delta}{n \cdot d}$$

- Steric constraint that is present in moving motor-filament system
- If working distance δ is smaller than path distance d , each individual crossbridge must spend a significant time detached while other crossbridges move the filament: $r < 1$ and motility will require an assembly of crossbridges

Table 15.1 Motor distances *in vitro*

Motor Distance	Parameter	Skeletal muscle myosin	Conventional kinesin
Working distance	δ	5 nm ^a	≥ 8 nm
Path distance	d	36 nm	8 nm
Distance/ATP per head ^a	Δ	200–400 nm	16 nm

^aLow load

