

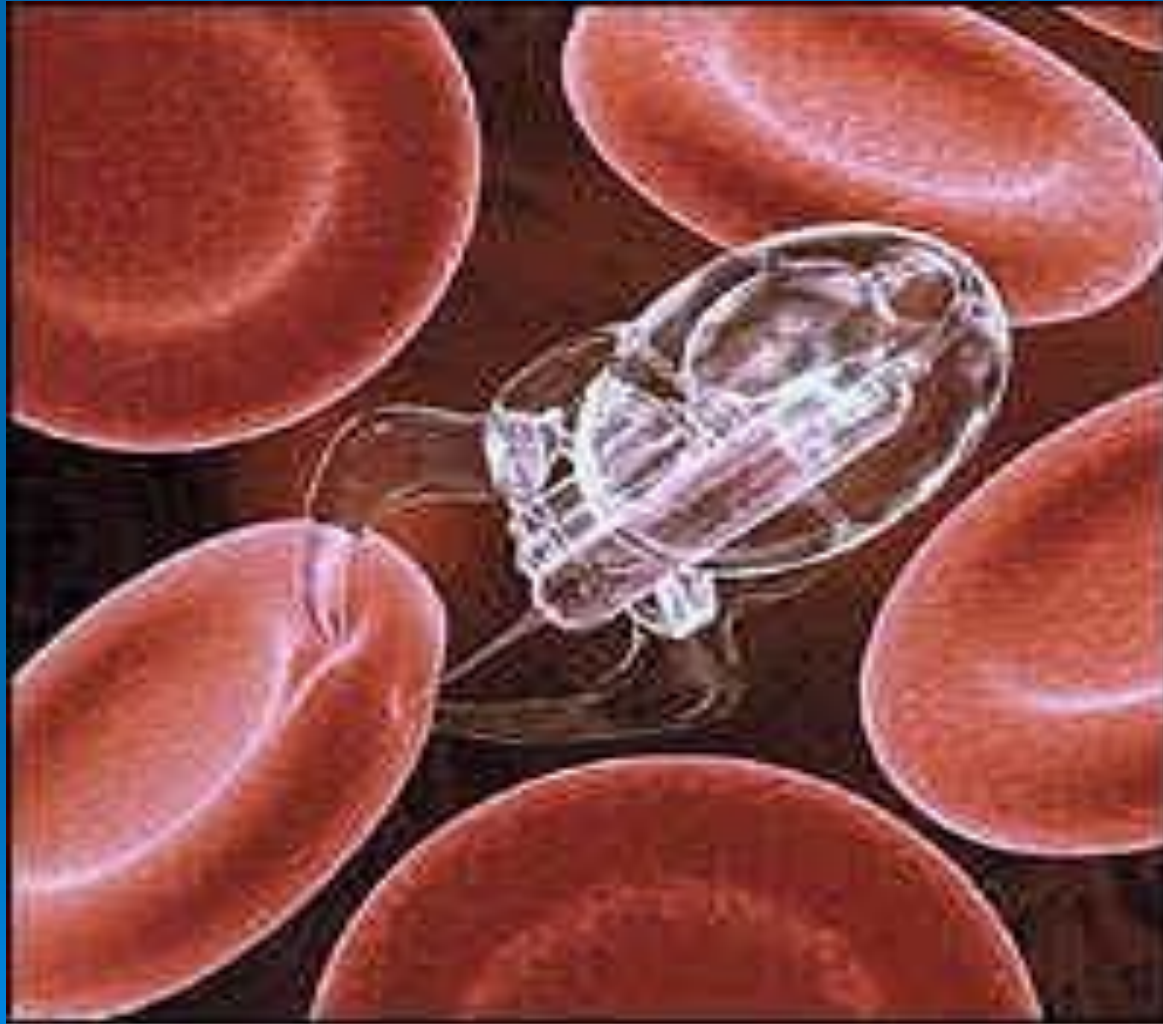
生物奈米馬達

中原大學 生物科技學系

吳宗遠

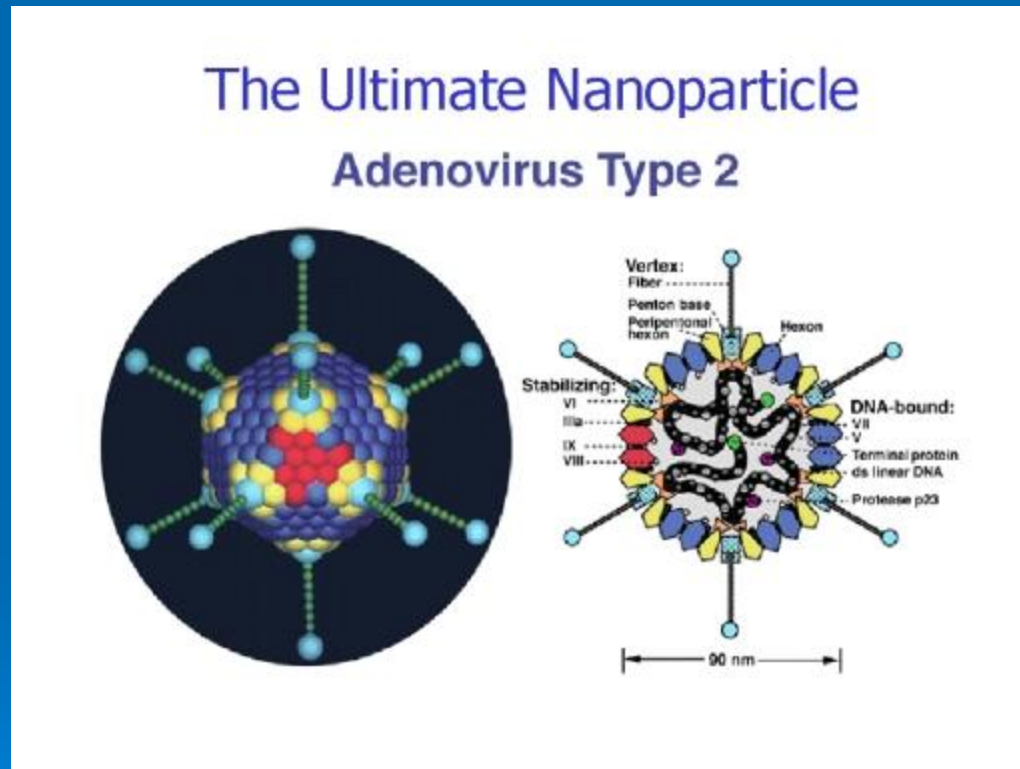


微小機械與細胞



Virus---self replicated particles (**machine**)

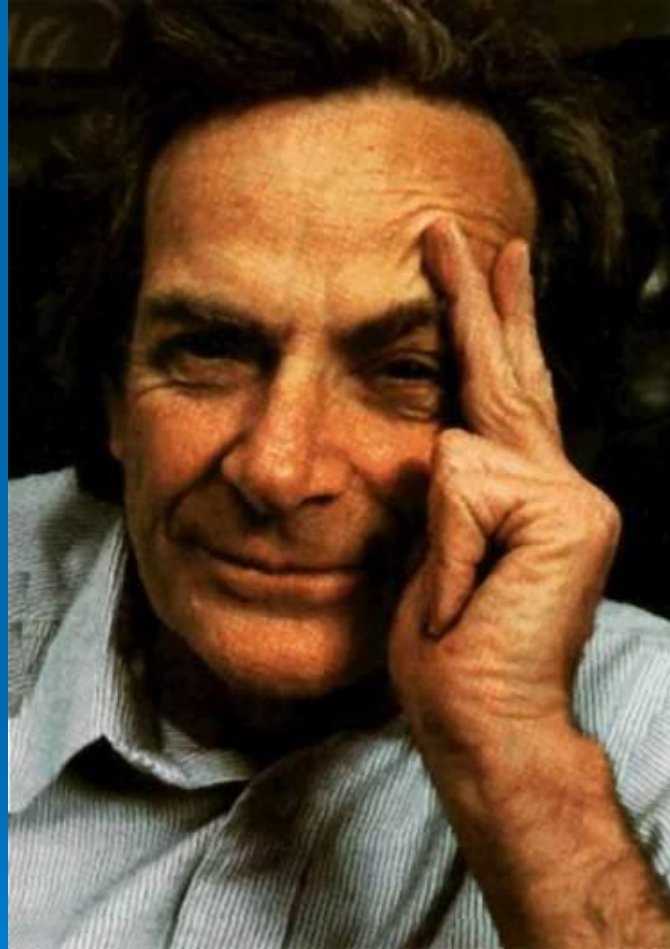
(奈米獵殺)



Virus: **proteins** plus **nucleic acids** (DNA or RNA)

The self replicated **nanomachine**

The man who dare to think **small**



Small is beautiful!

徐志摩 (1897-1931)



Large is beautiful !

The New Biochemistry: Macromolecular Machines

1998 Welch Conference, Houston, TX, October 26 & 27, 1998

Joseph L. Goldstein, Program Chairman

Machines That Control Cell Division

Ira Herskowitz – Discussion Leader

- Robert Weinberg – Oncogenes and Tumor Suppressors
- Steve Elledge – Cyclin Kinase Complexes
- Nikola Pavletich – Cell Cycle Regulatory Proteins
- Robert Huber – Proteasome

Machines That Produce DNA, RNA, and Protein

Steve McKnight – Discussion Leader

- Mike O'Donnell – DNA Replicase
- Tom Cech – Telomerase
- Robert Tjian – Basal Transcription Complex
- Cynthia Wolberger – Multimeric DNA-Binding Proteins

Machines That Transduce Signals

Susan Taylor – Discussion Leader

- Tony Pawson – Protein Modules
- Al Gilman – Heterotrimeric G Proteins
- Pierre Chambon – Nuclear Receptor Superfamily
- Michael Brown – SREBP Pathway

Machines That Move Molecules

Hans Deisenhofer – Discussion Leader

- Ulrich Hartl – Molecular Chaperones
- Peter Walter – Unfolded Protein Response Pathway
- Rod McKinnon – Ion Channels
- Thomas Südhof – Synaptic Vesicle Cycle

奈米馬達的最適材料可能
是蛋白質 (protein)

WHY?



Molecular Motors

linear motion motor

Rotatory motion motor

All this molecular motor in the cell are **protein molecules**

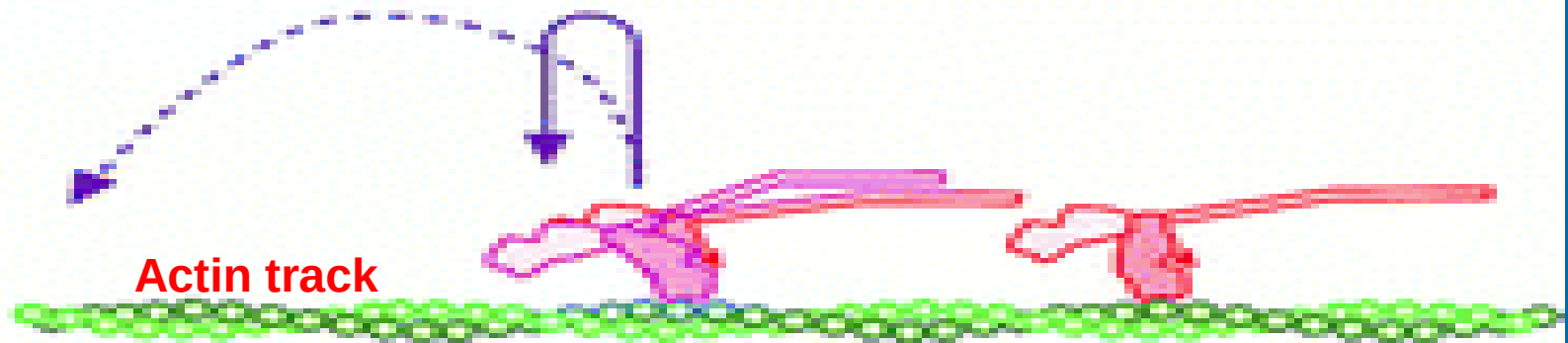
What is the energy source?

Chemical energy convert to mechanical energy

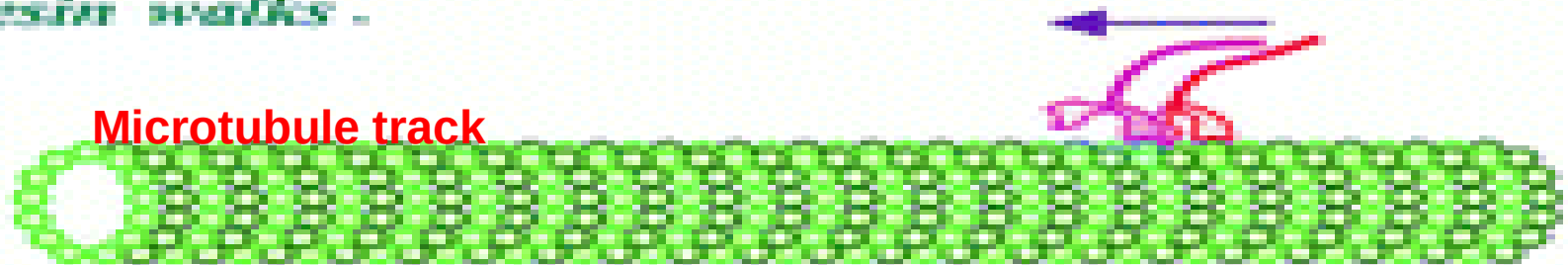


Linear molecular motors: **track** dependent

Myosin runs (hops) -



Kinesin walks -



Three β subunits (of F_1) crawl -



(a)

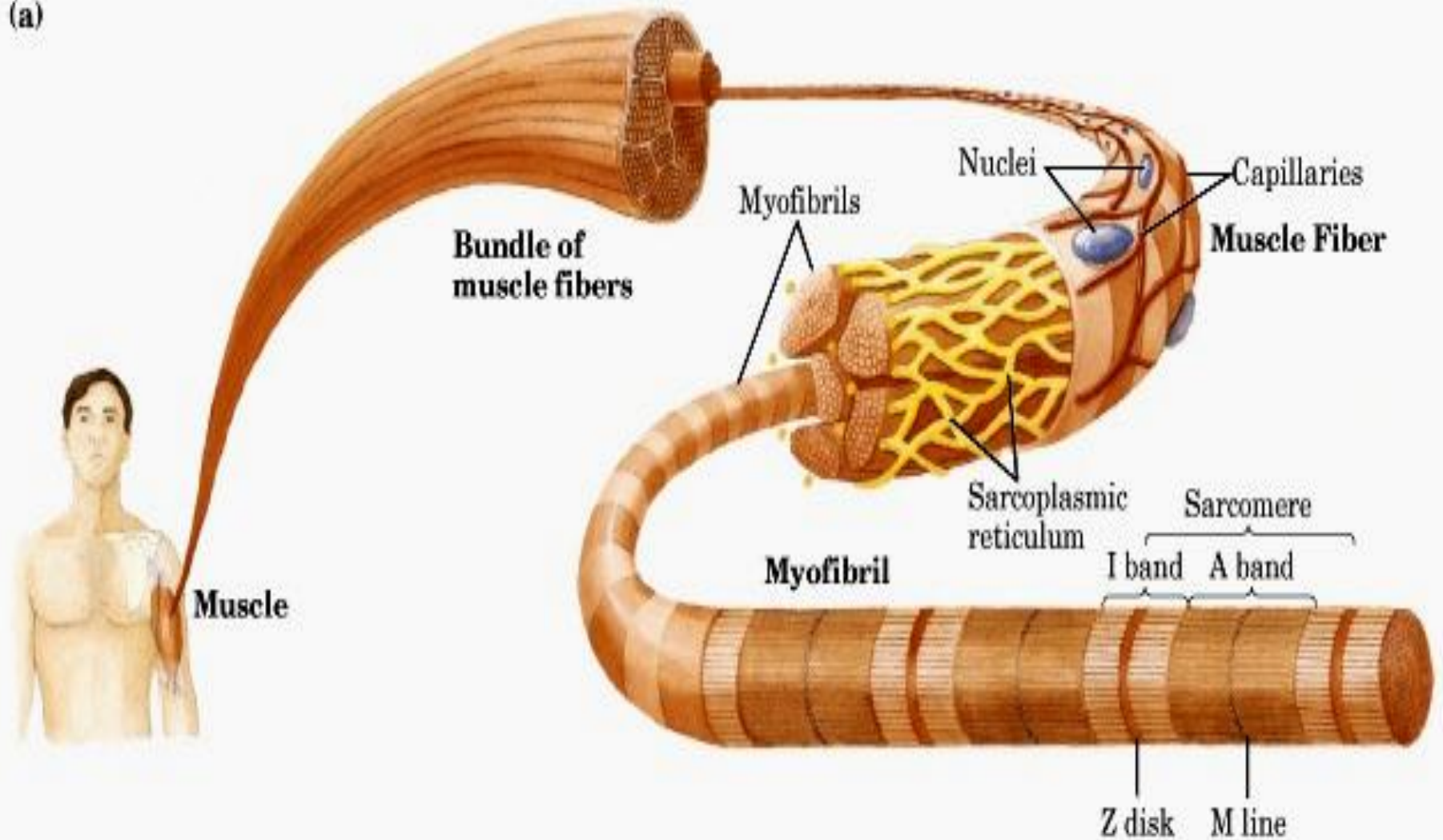


Figure (a) Structure of **skeletal muscle**. Muscle fibers consist of single, elongated, multinucleated cells that arise from the fusion of many precursor cells.

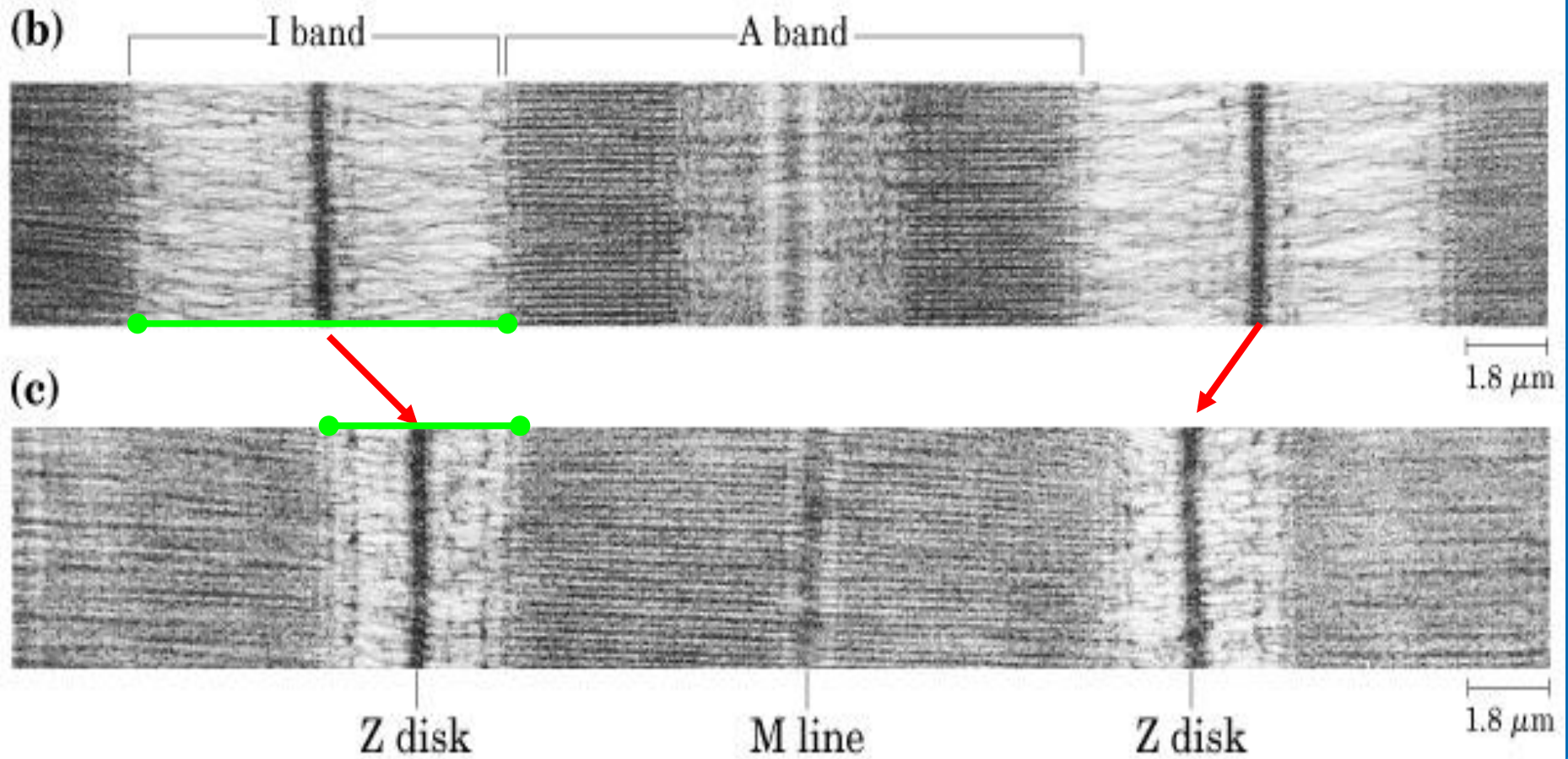


Figure (b) 、 (c) Structure of skeletal muscle.

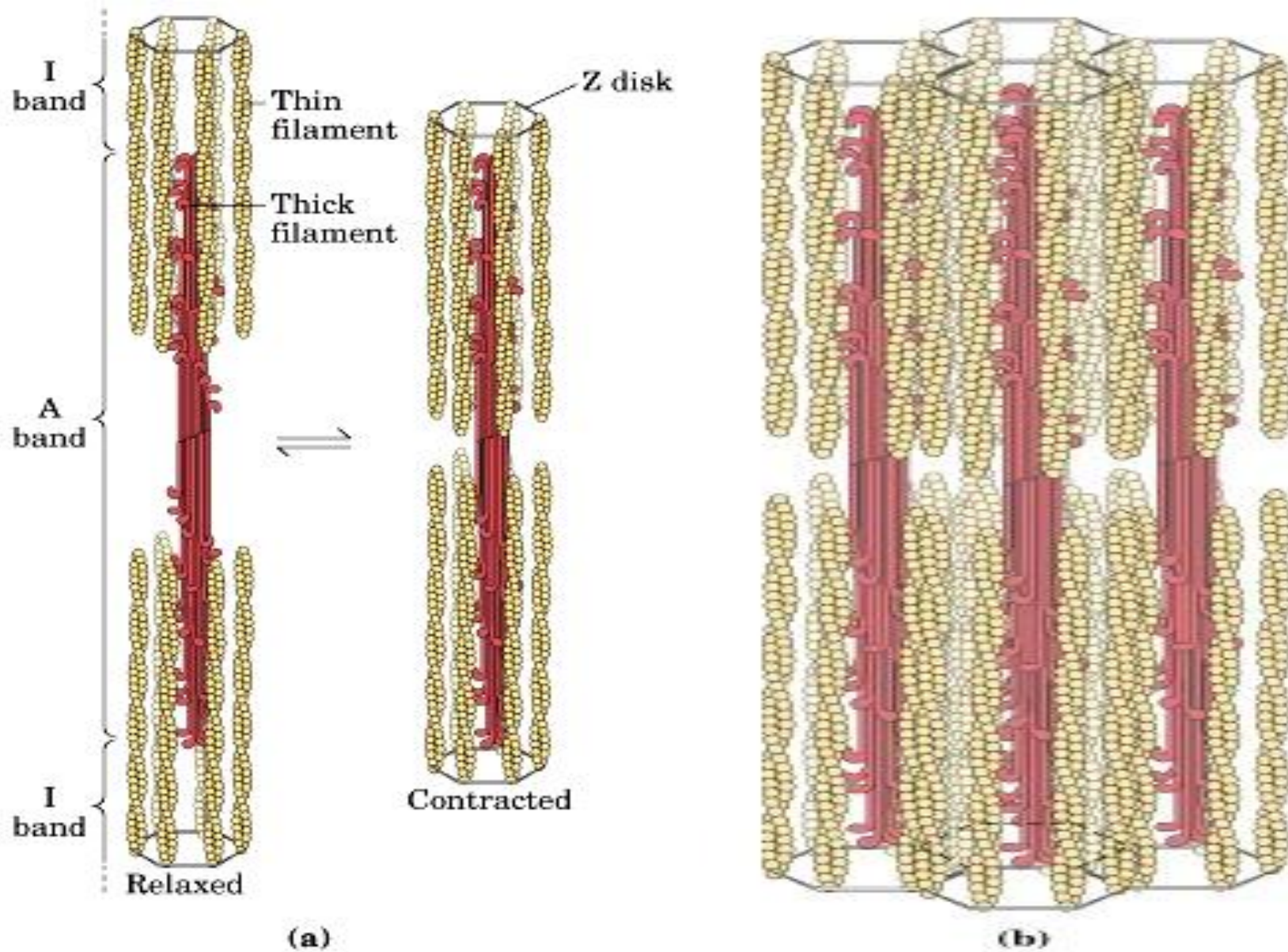


Figure 7-32 (a) Muscle contraction.

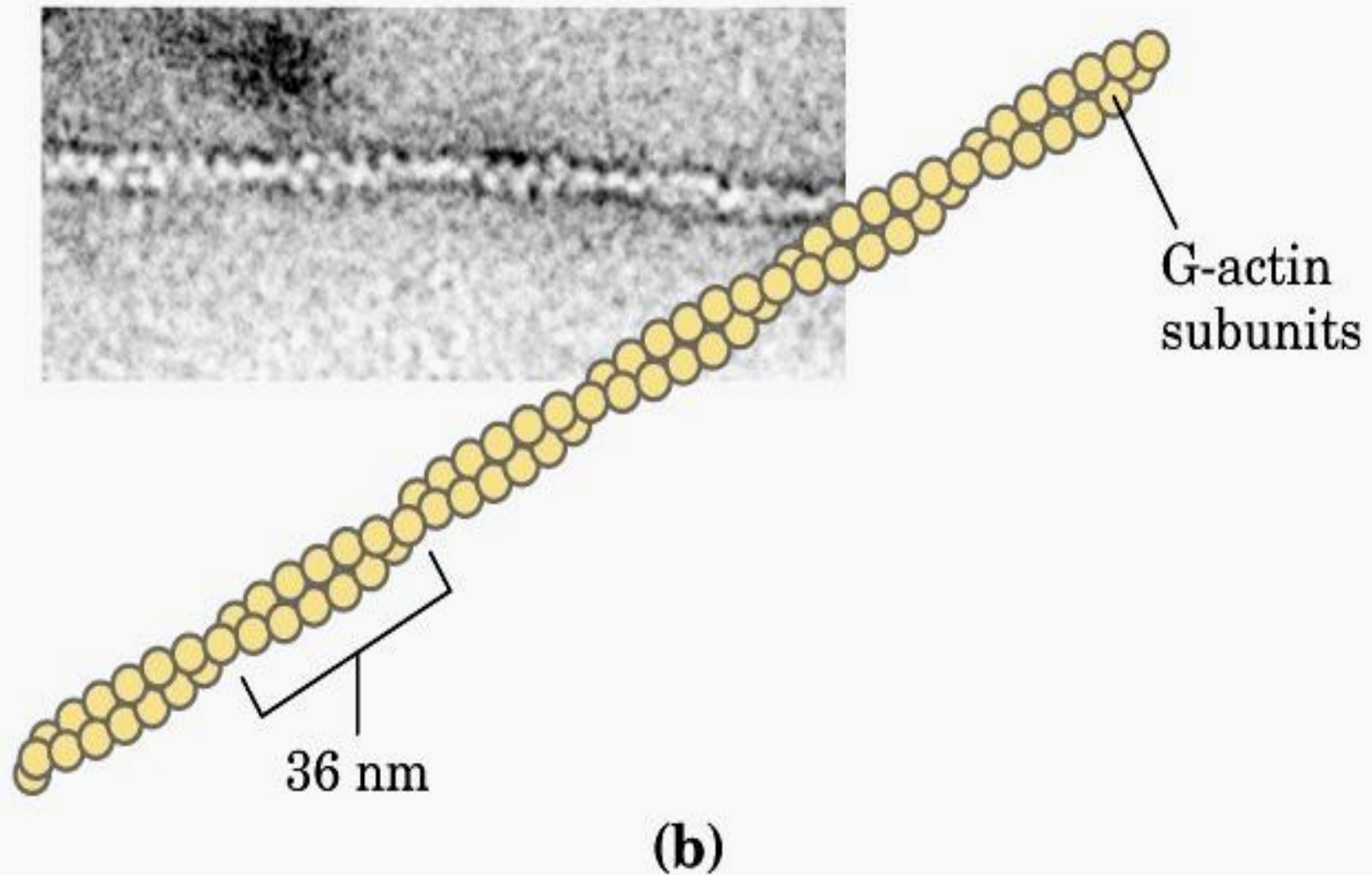


Figure (b) The major components of muscle. **F- actin**. The art of protein biochemistry is to isolate the proteins from “observation” tissues or cells.

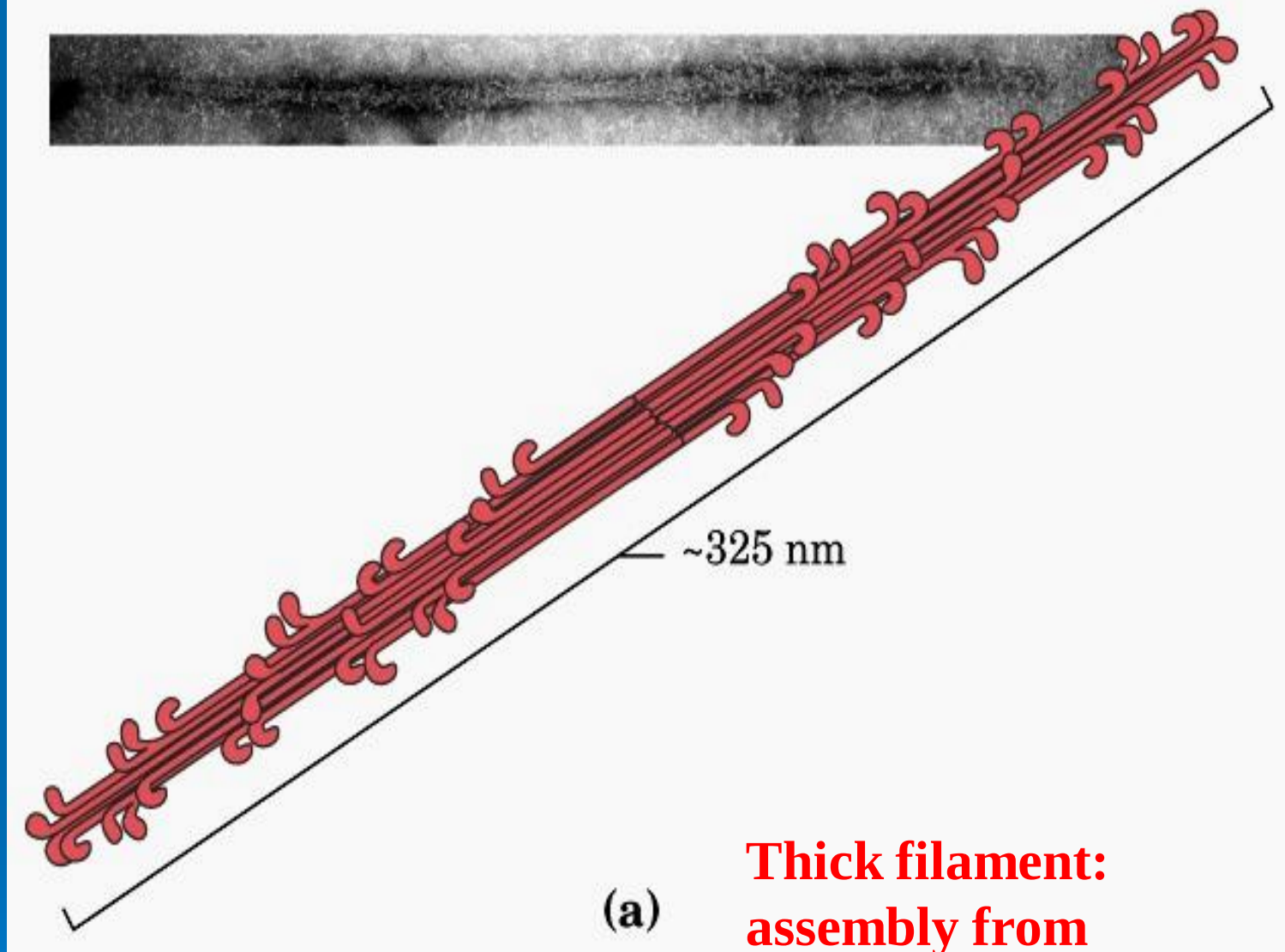


Figure (a) The major components of muscle. thick filament

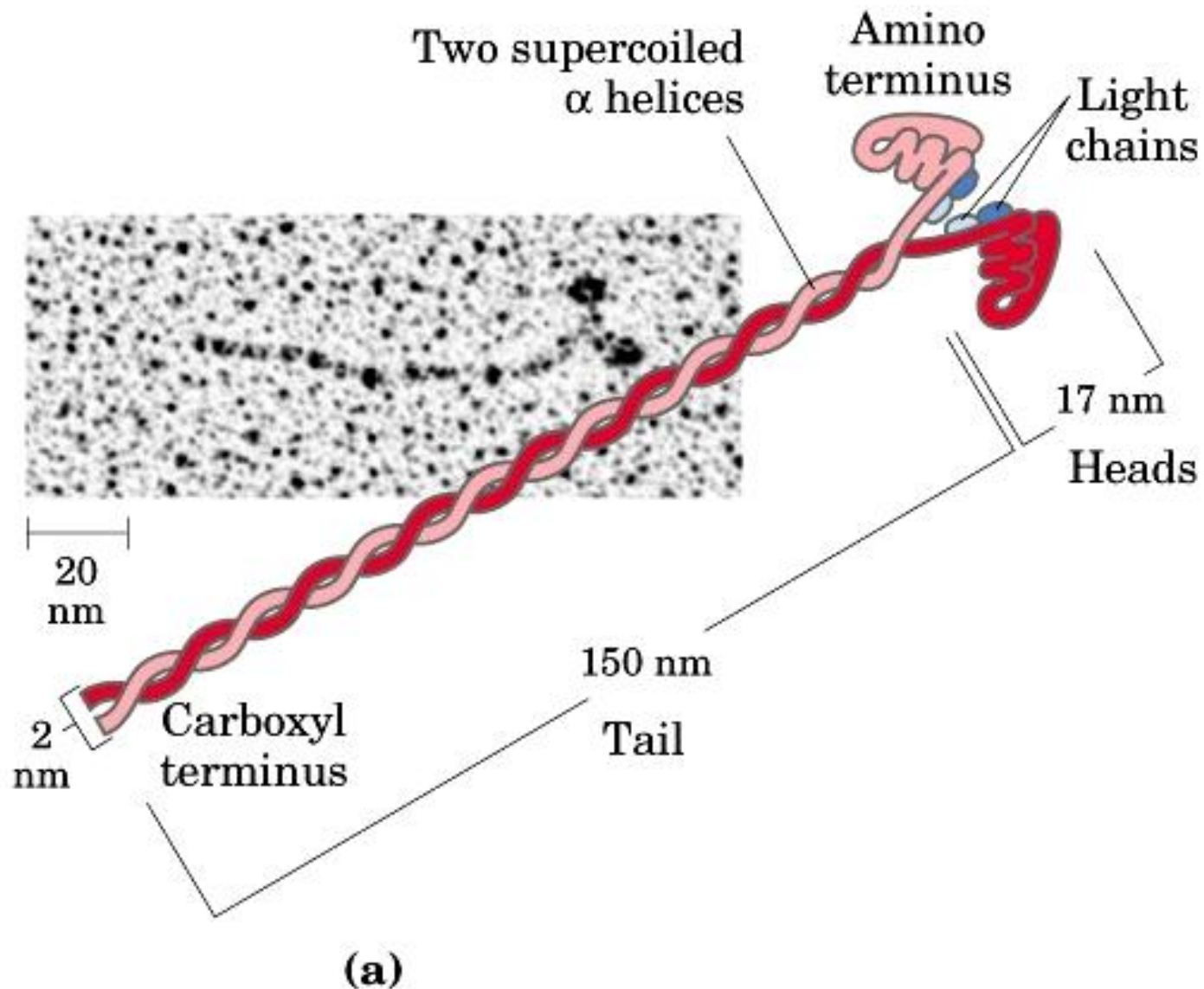


Figure (a) Myosin. Two heavy chain.

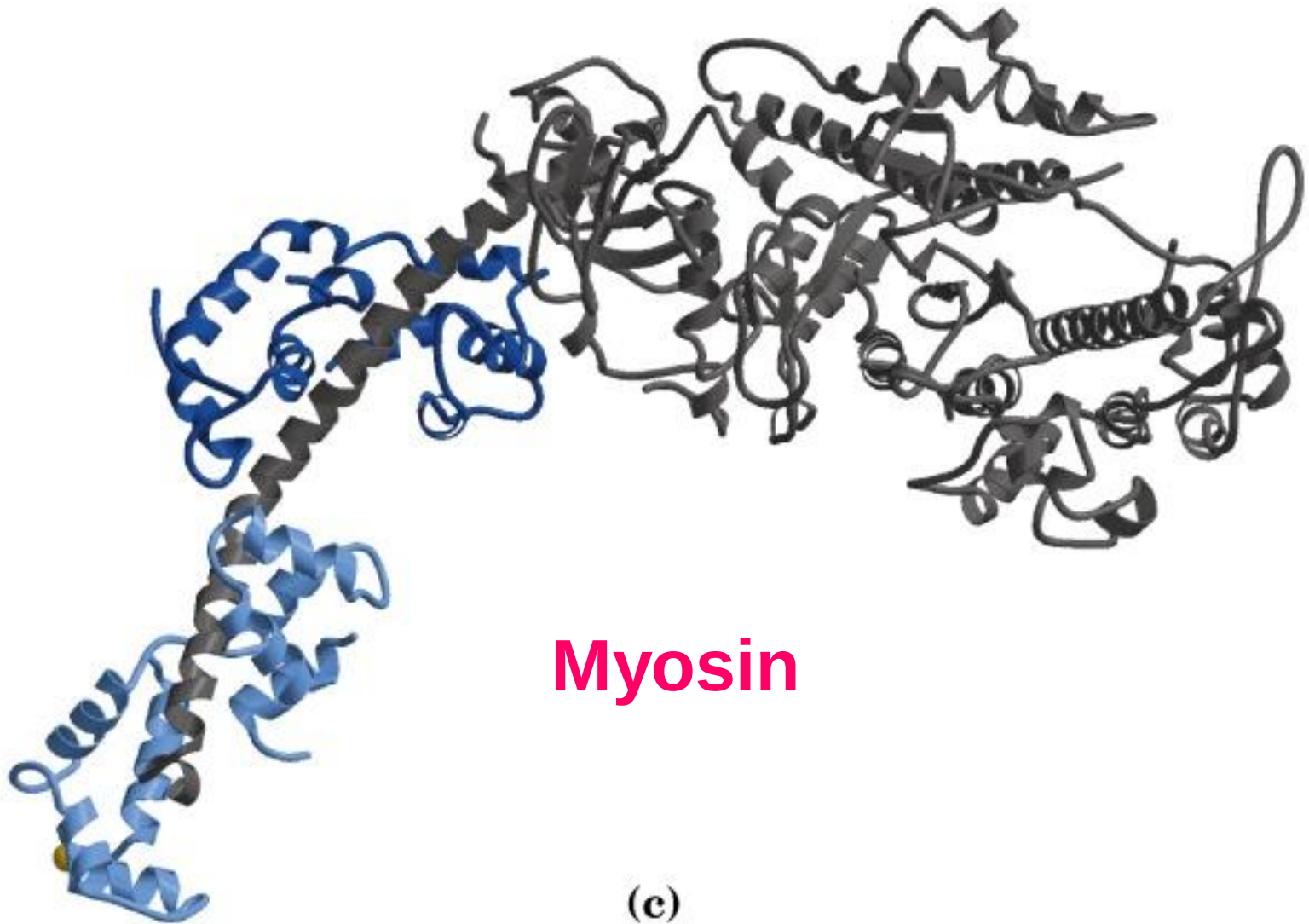
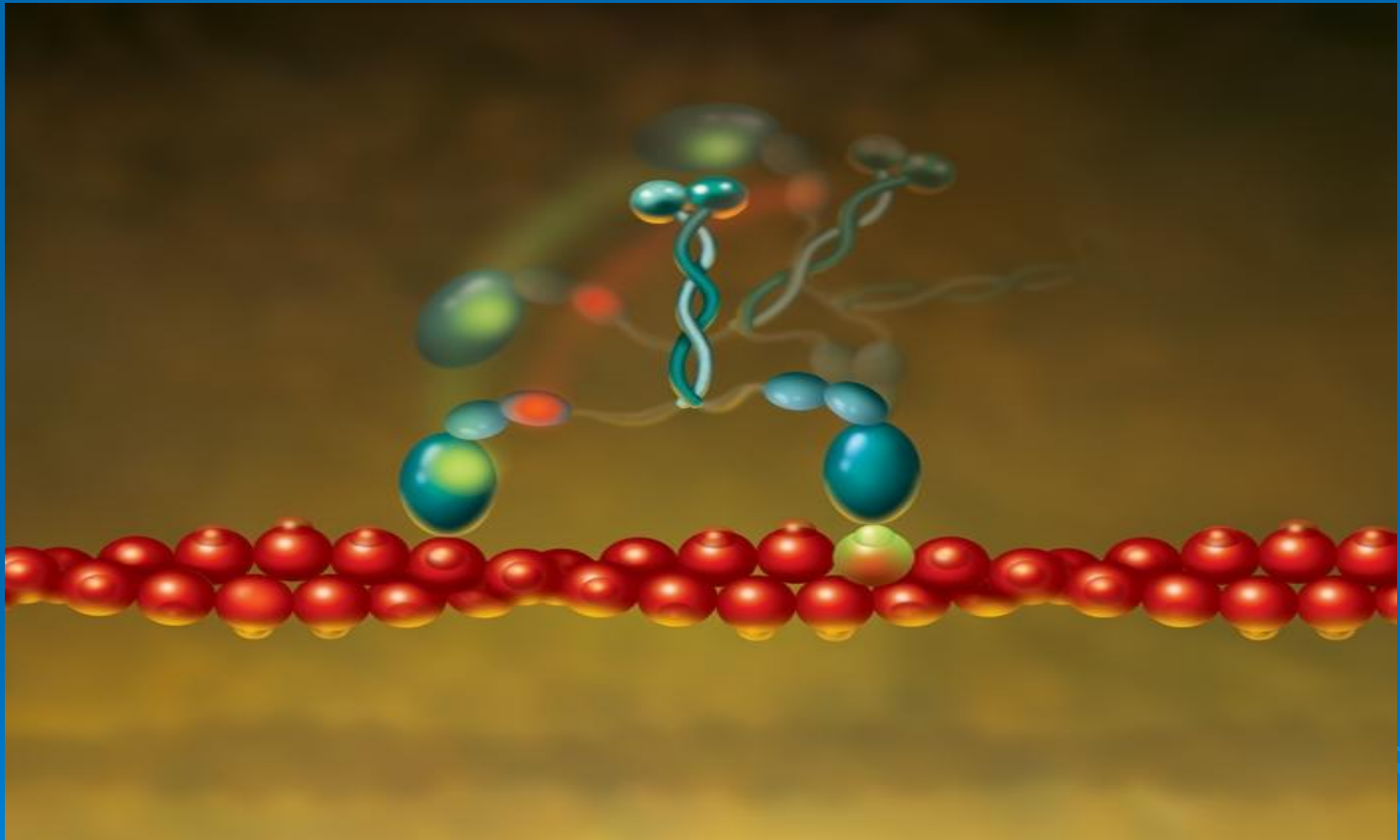


Figure (c) Myosin. Ribbon representation of the myosin S1 fragment.

Kinesin/dynein



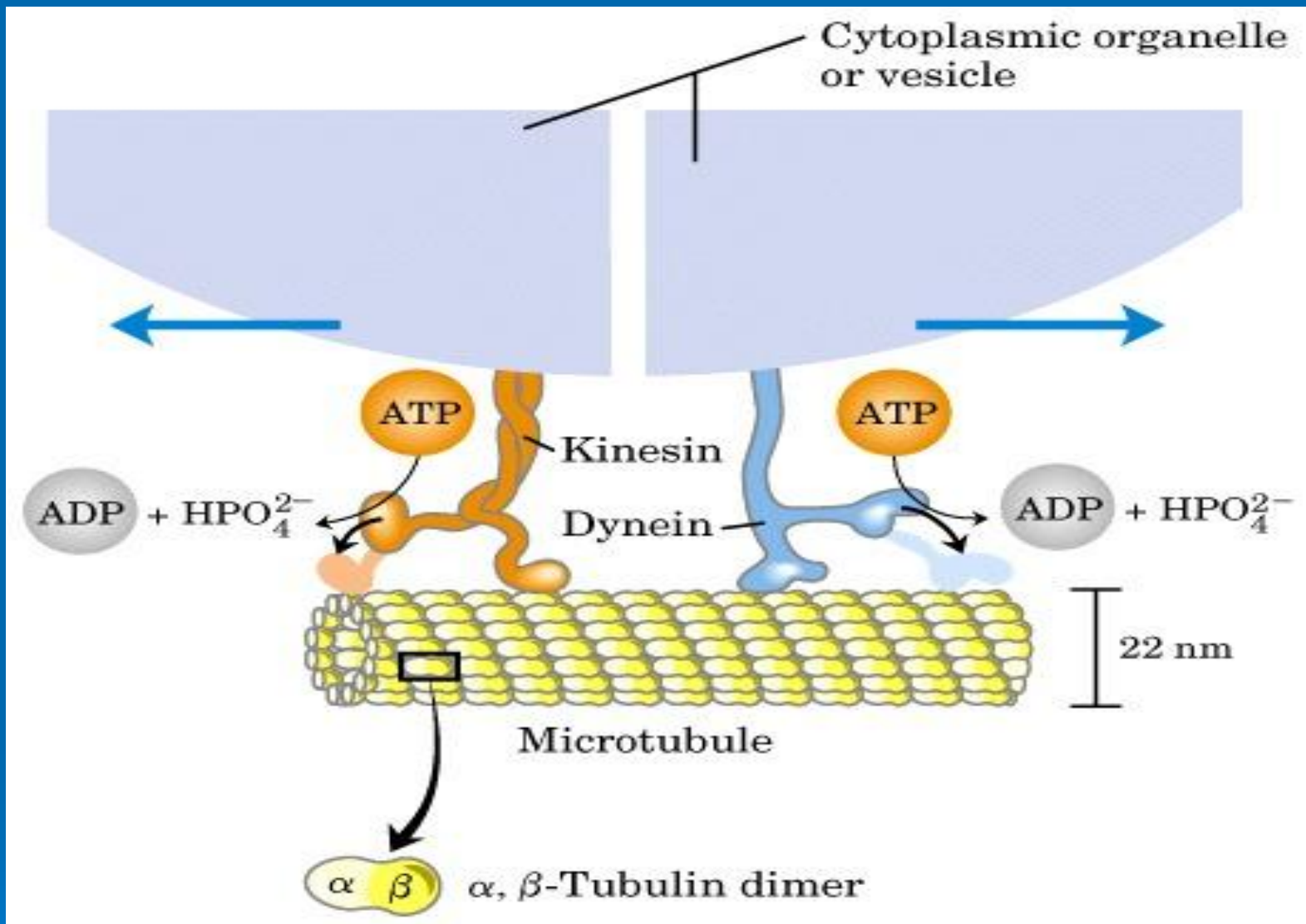
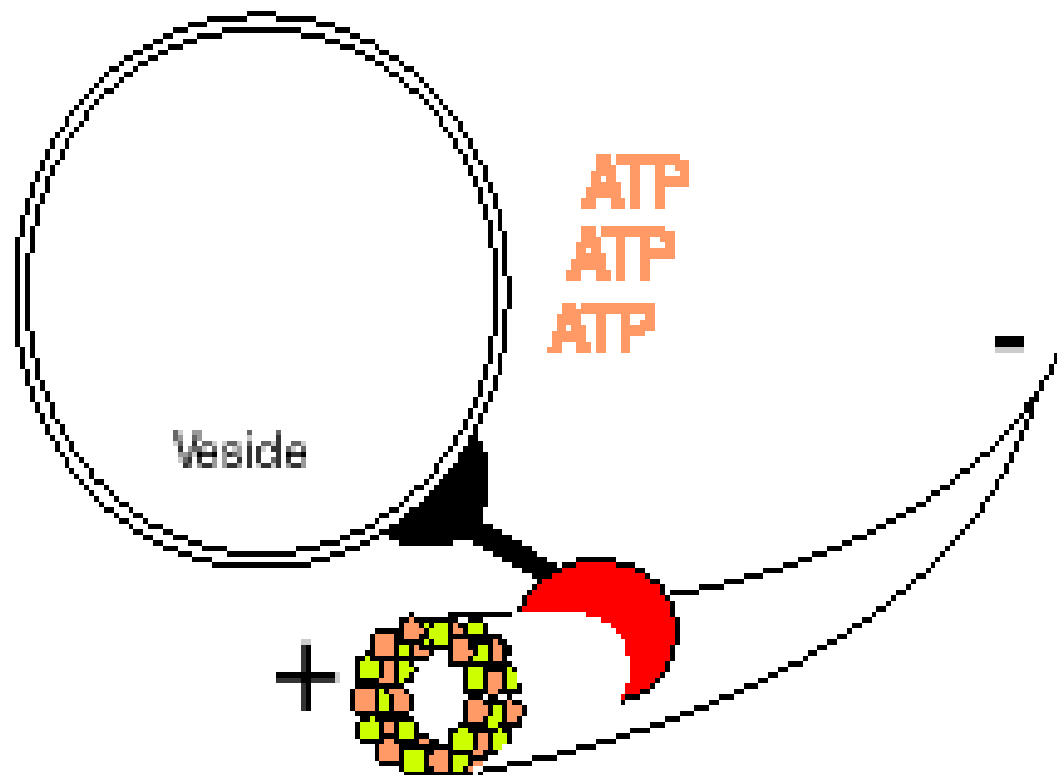
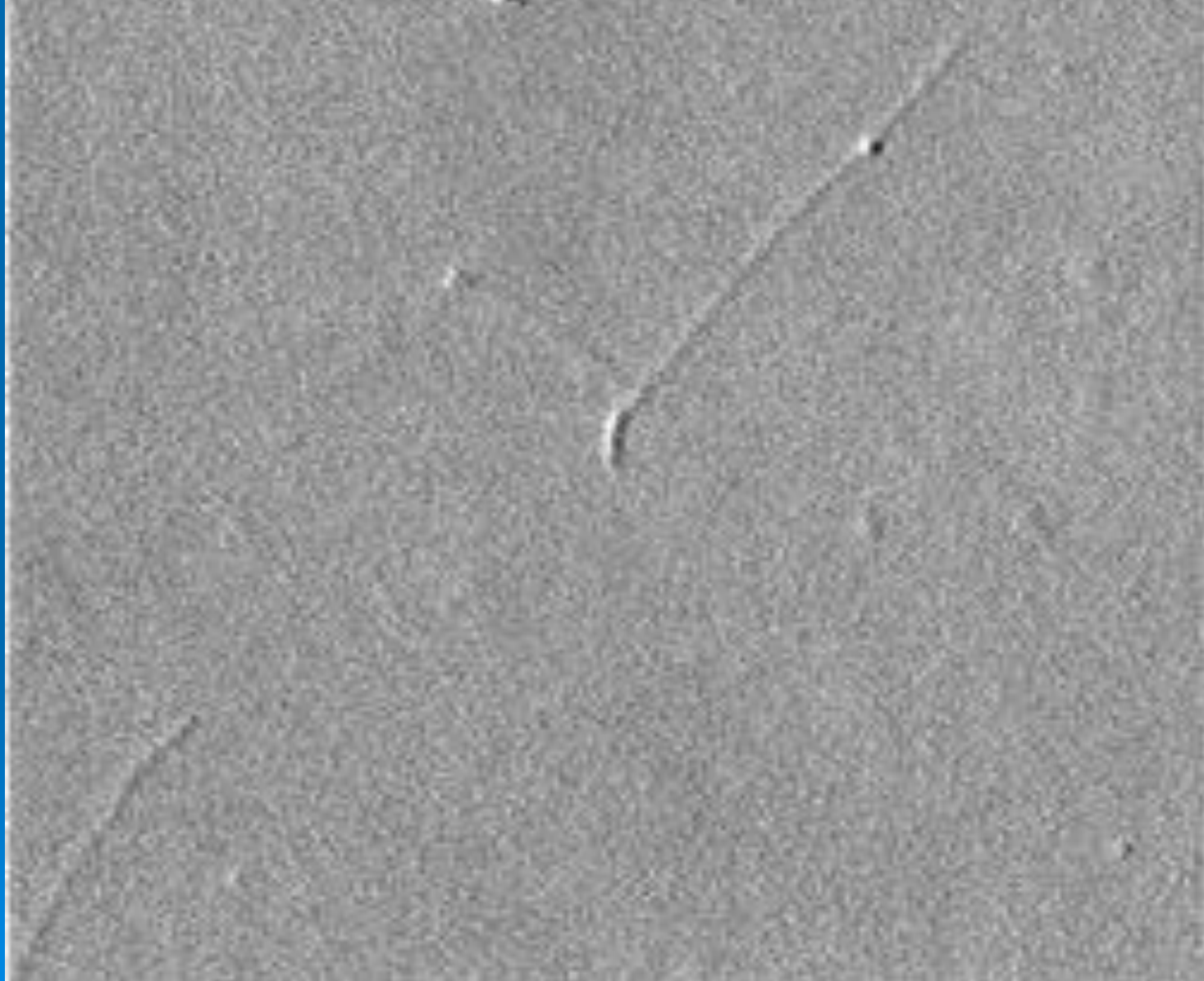


Figure : **Kinesin** and cytoplasmic **dynein** are ATP-driven **molecular motors** that can attach to cytoplasmic organelles or vesicles and drag along microtubular “rails” at a rate of about **1 μm/s**.



KH02/HW0 6/3/36

01143141



Toward Devices Powered by Biomolecular Motors

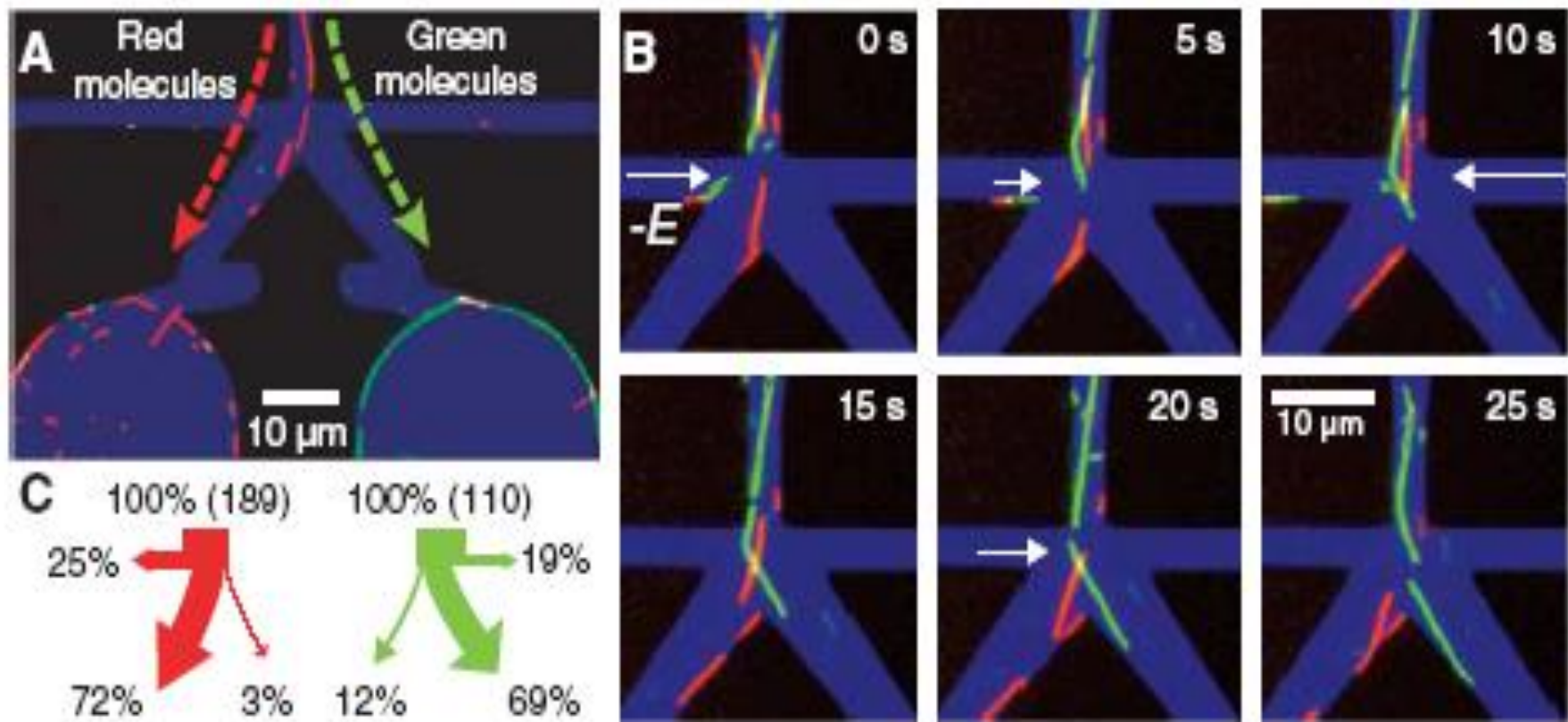
Biomolecular motors can be used in nanometer-scale devices to perform mechanical work. This approach will assist the development of active nanostructures.



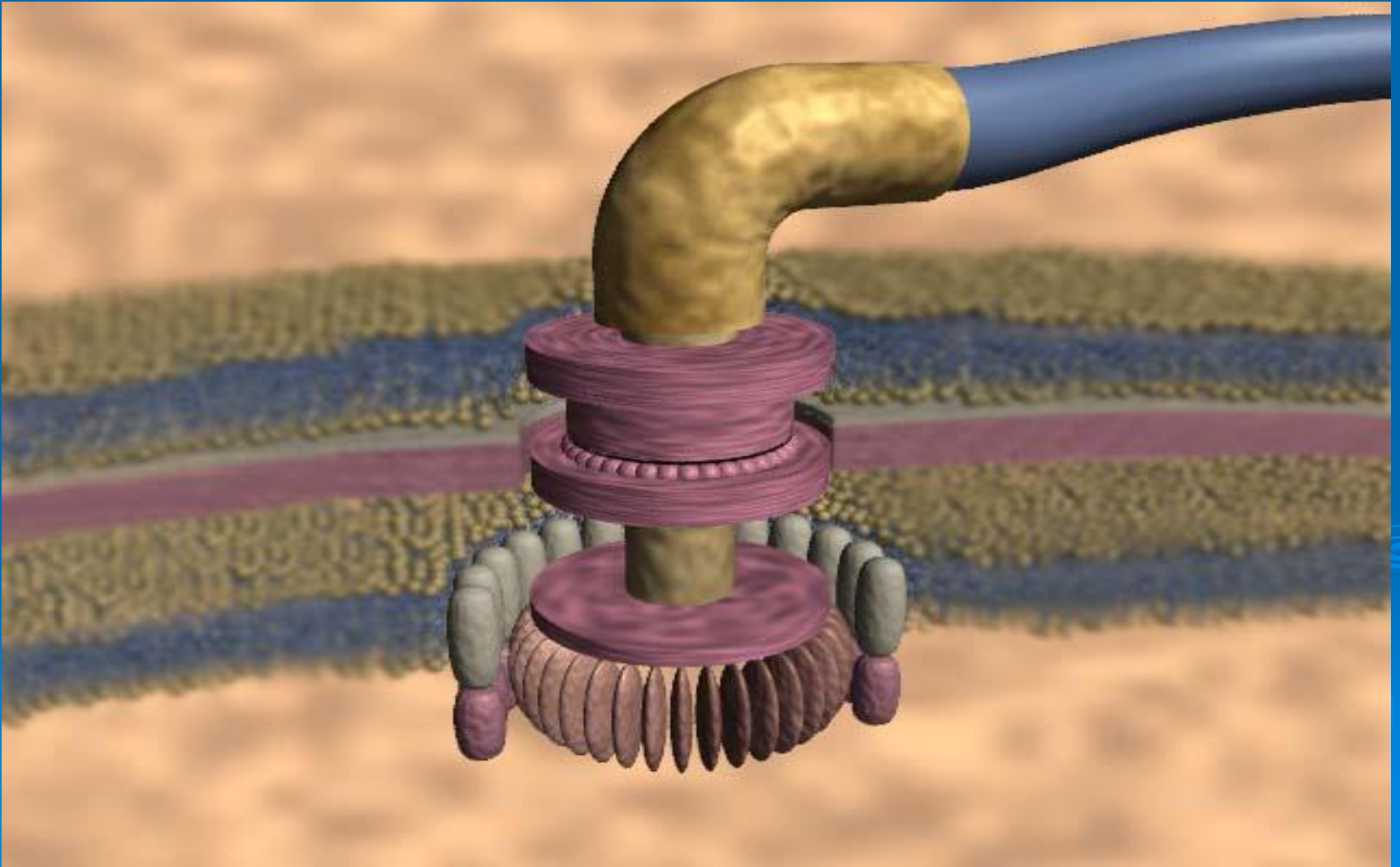
Nanofluidics with molecular motors. In van den Heuvel *et al.*'s work (2), an electric field is used to steer the microtubules into one of two arms of a Y junction; the microtubules move perpendicular to the field. The microtubules are transported by kinesin motor proteins.

Molecular Sorting by Electrical Steering of Microtubules in Kinesin-Coated Channels

Martin G. L. van den Heuvel, Martijn P. de Graaff, Cees Dekker*



Bacterial flagella: a rotatory motor



The ATP synthase



The Eukaryotic cell

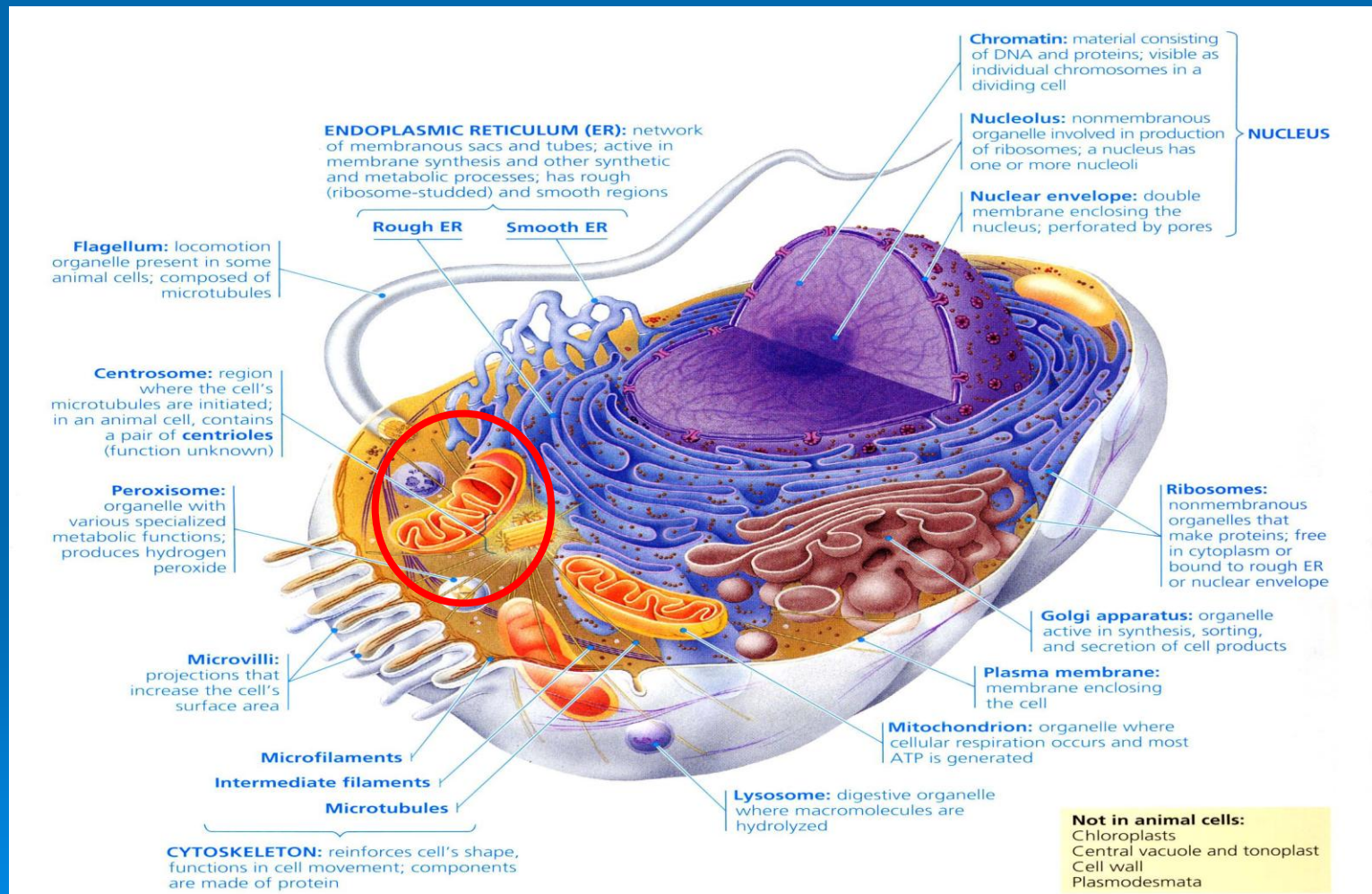
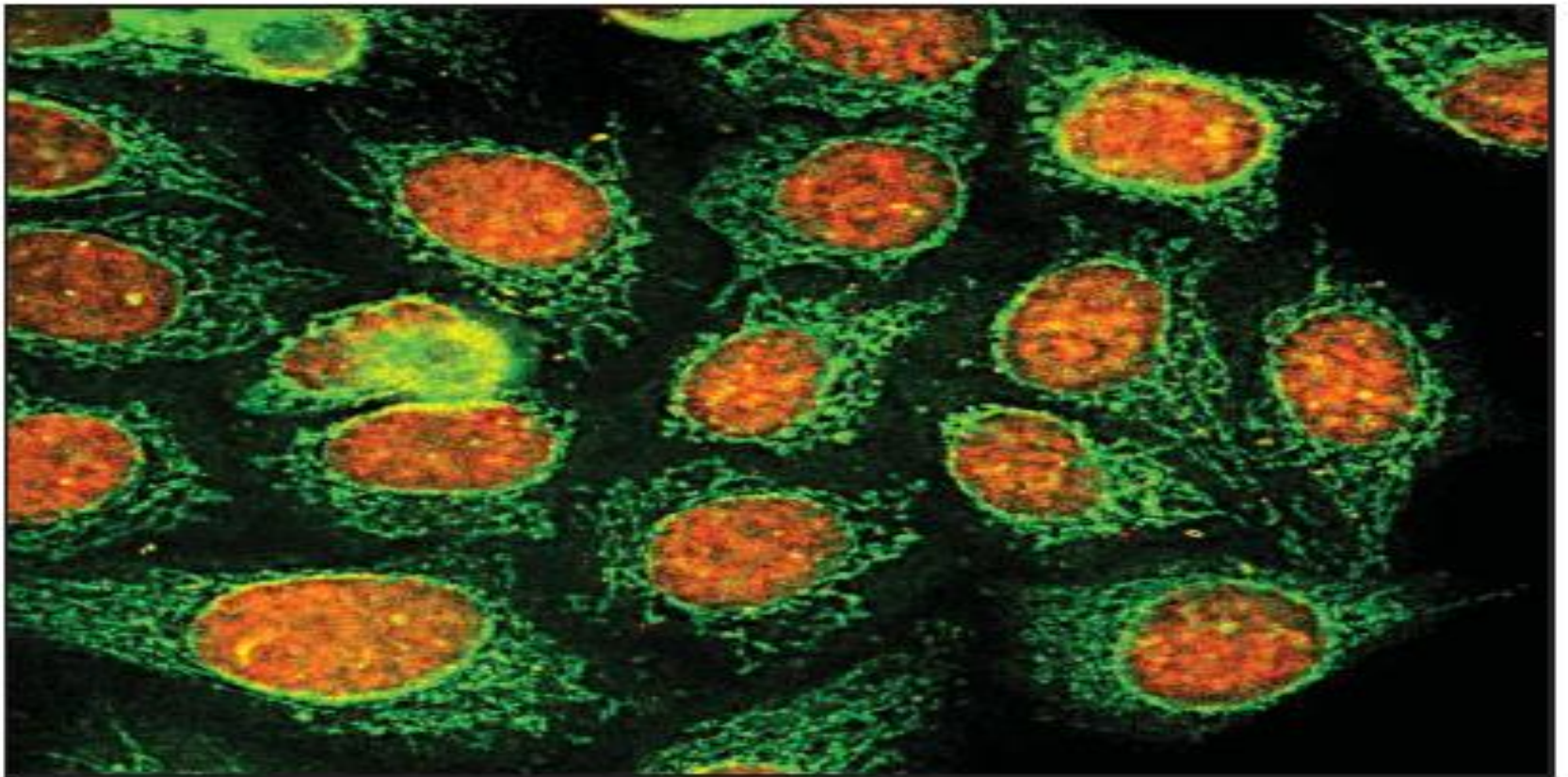


FIGURE 7.7 Overview of an animal cell.

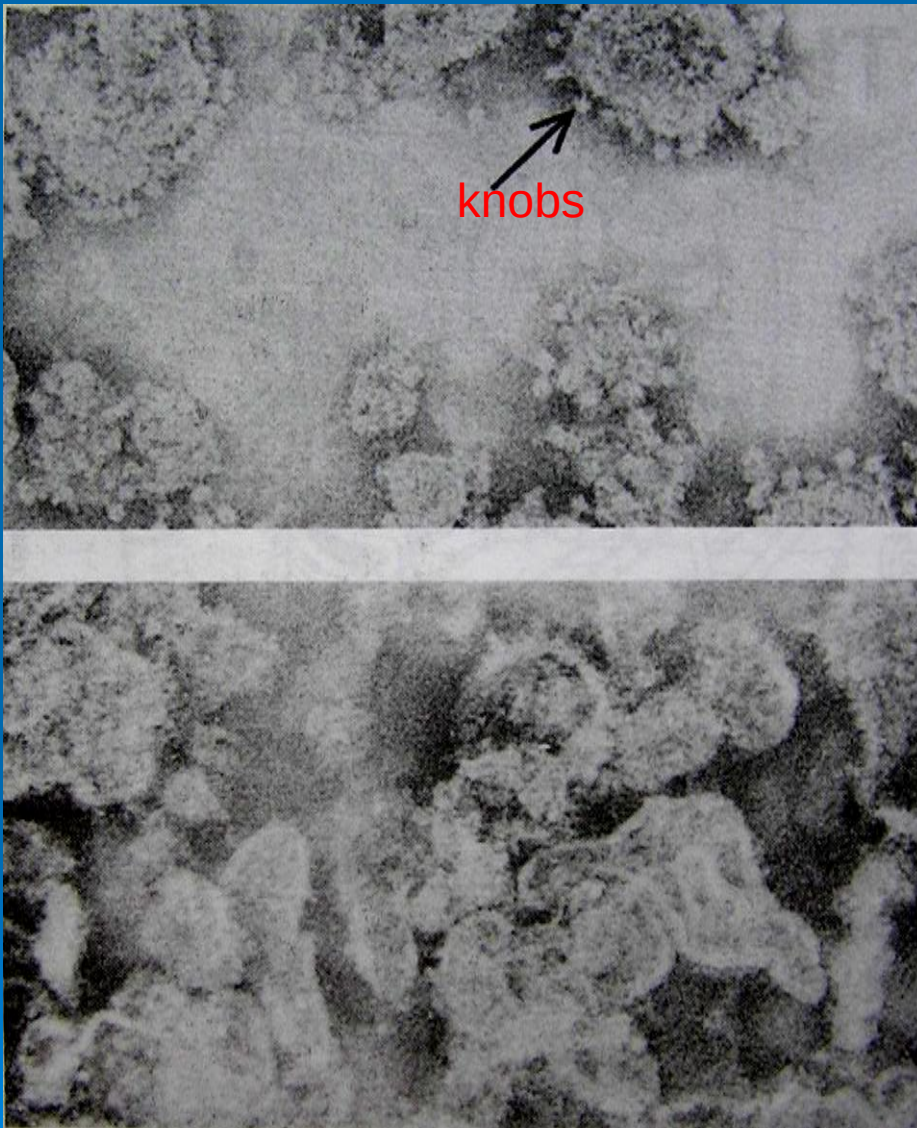
(Campbell Biology, 6th edition)



Double duty. Green quantum dots cling to mitochondria in the cytoplasm; orange ones label proteins in the same cells' nuclei.

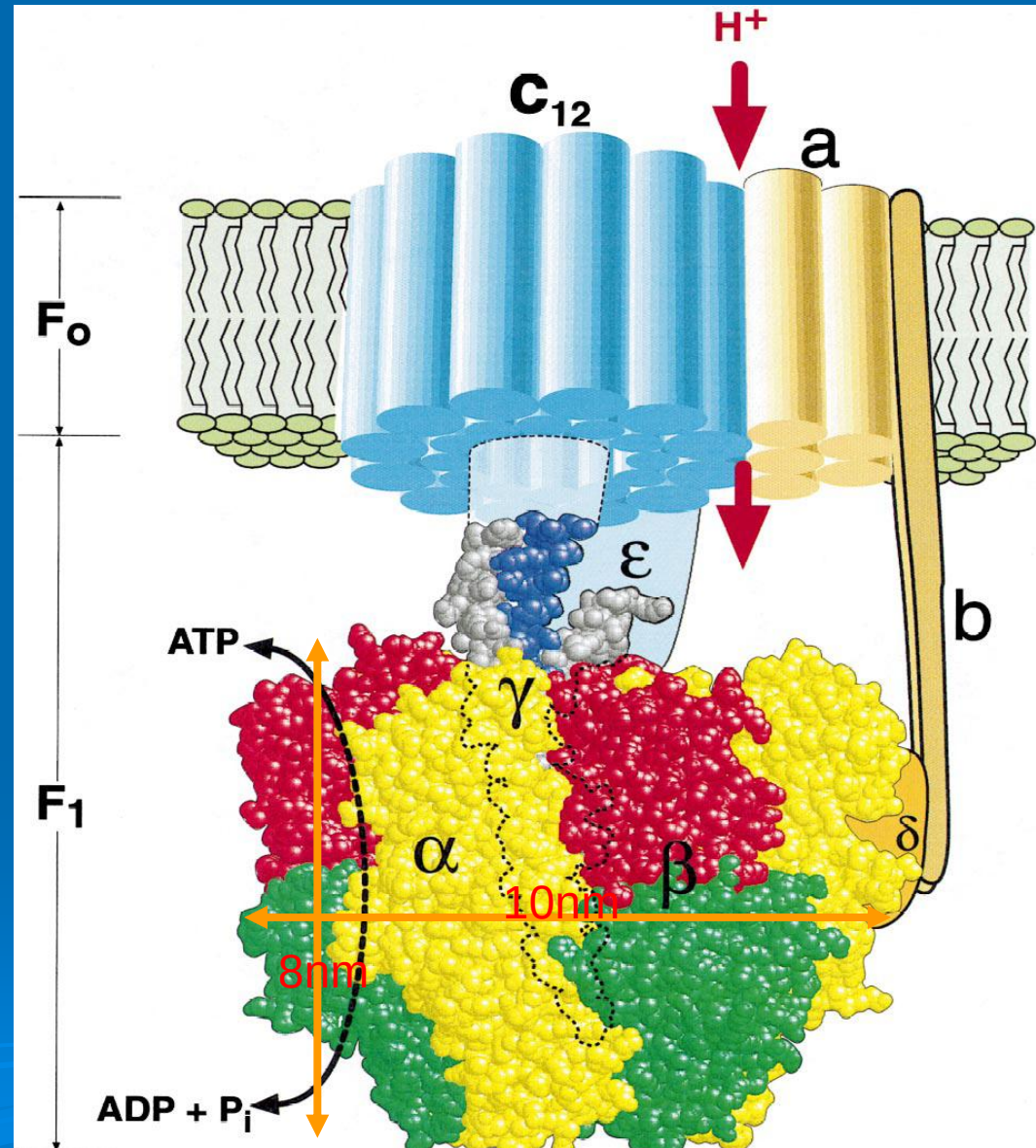
F₁ and F₀ Proteins in Mitochondria

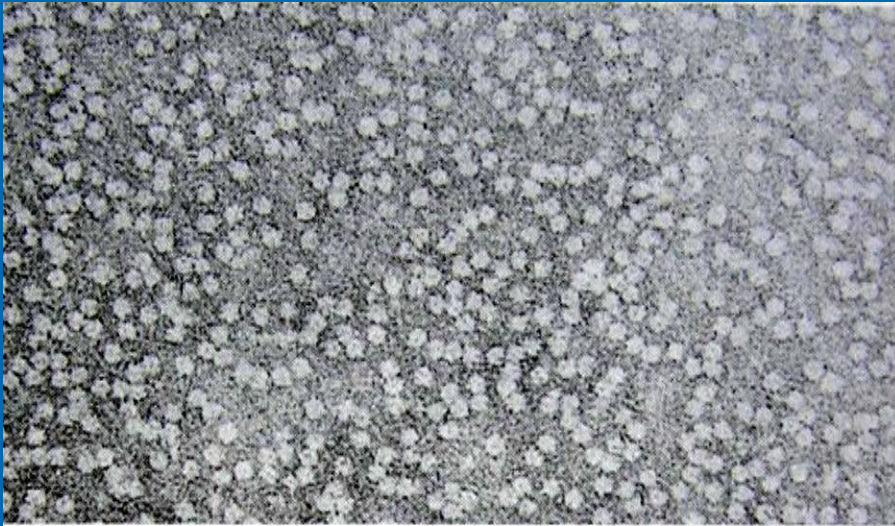
- Early studies were done with submitochondrial particles (SMP, inside-out vesicles)
- They synthesize ATP in the presence of respiratory substrates, or hydrolyze ATP in the absence of substrates
- Both respiration and ATP hydrolysis generate a Δp in the lumen $\Rightarrow$ ATP synthesis/ hydrolysis is tightly coupled with Δp
- ATP synthase can be visualized under EM in negatively stained SMP (with phosphotungstate)



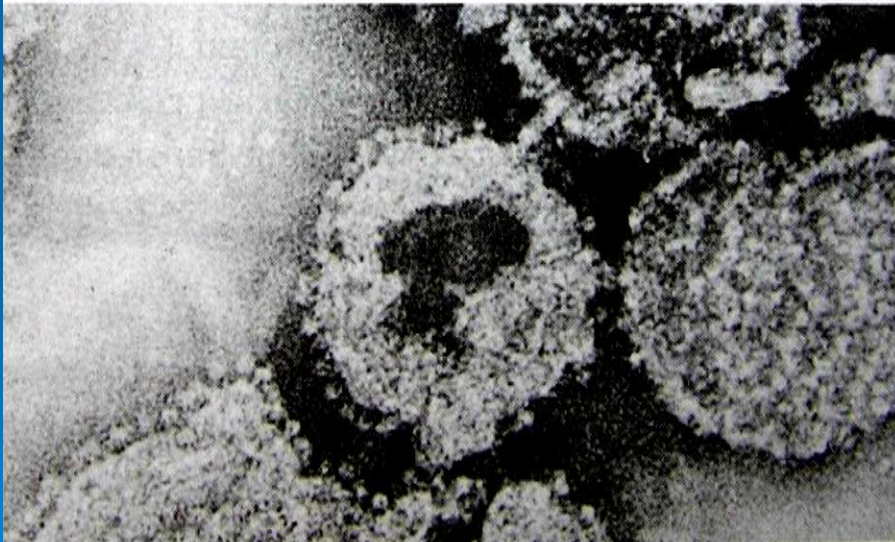
- Negatively stained SMP, with knobs facing outside
- Tightly **coupled** and sensitive to **oligomycin**
- Wash with urea/chelating agents, knobs were lost
- SMP became uncoupled, that is inhibited by oligomycin
- The **H⁺ channel** was termed F_o (fraction oligomycin)

- F_o – **hydrophobic**, forming H^+ channel and coupling H^+ to ATP synthesis
- F_1 – **hydrophilic**, site of ATP synthesis



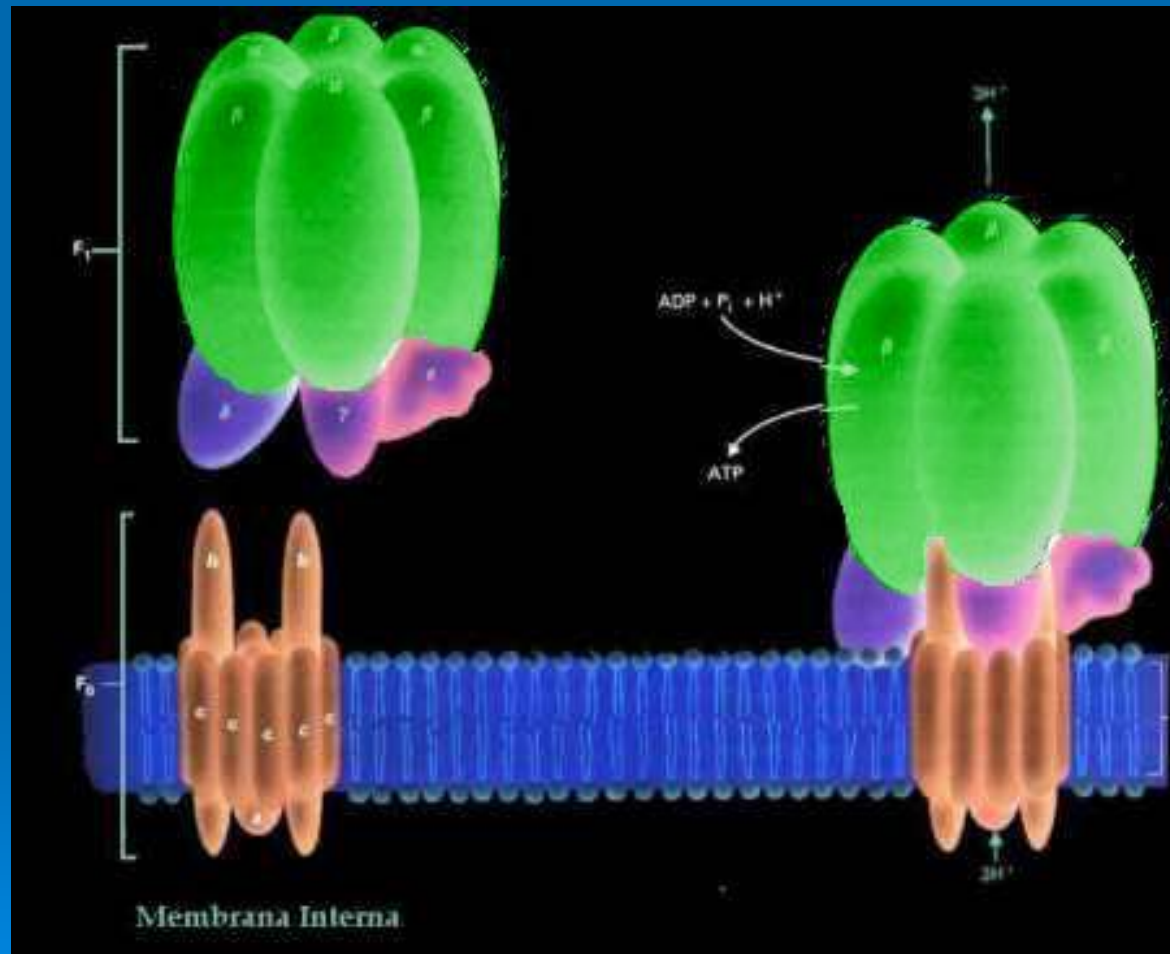


- ATPase activity was isolated, and **not** inhibited by oligomycin (termed F_1 , fraction 1)

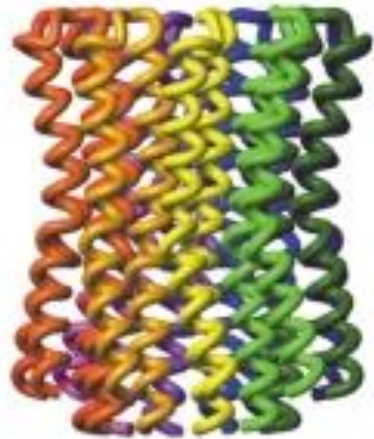


- After **reconstitution**, coupling was back
- **F_1 acts as a plug on F_0**

The assembly of Fo and F1

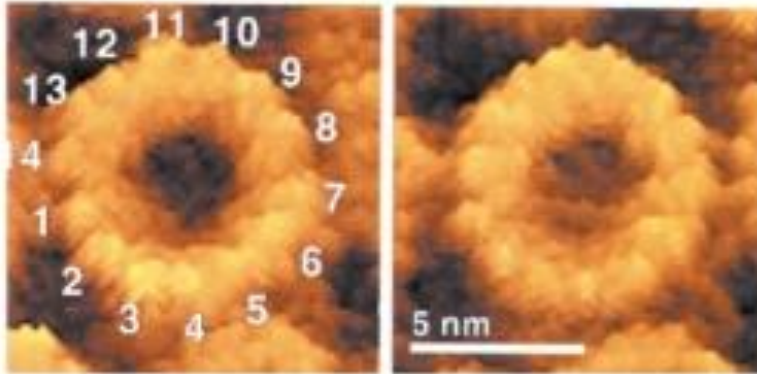
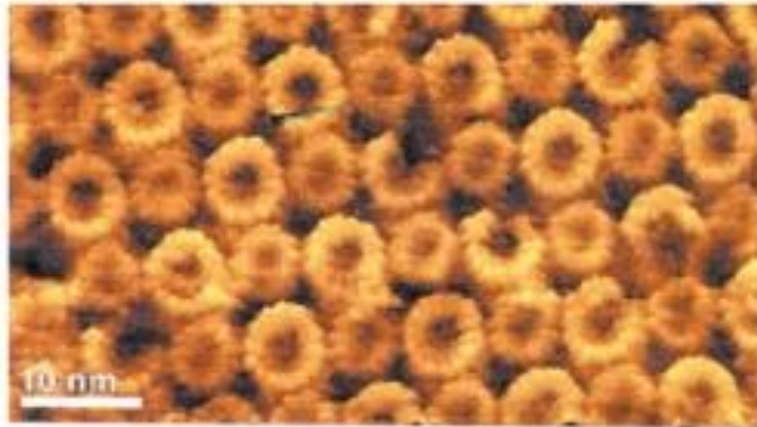


yeast



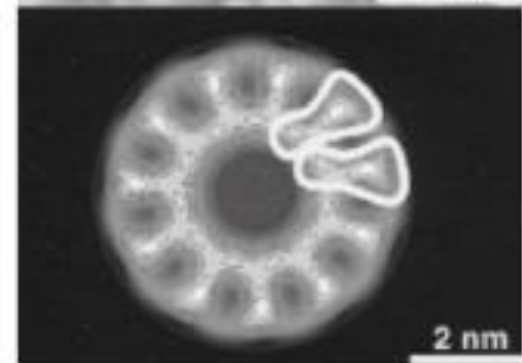
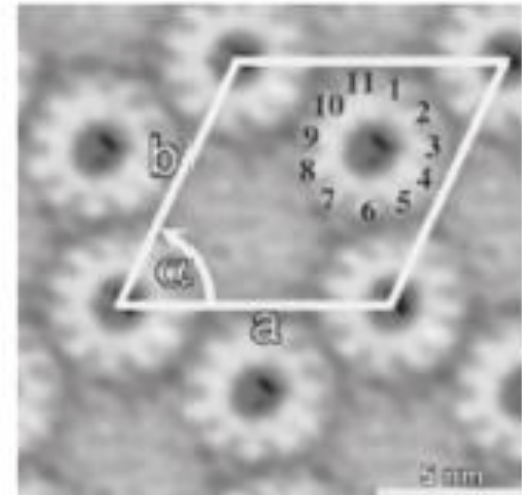
10 copies
in yeast

spinach



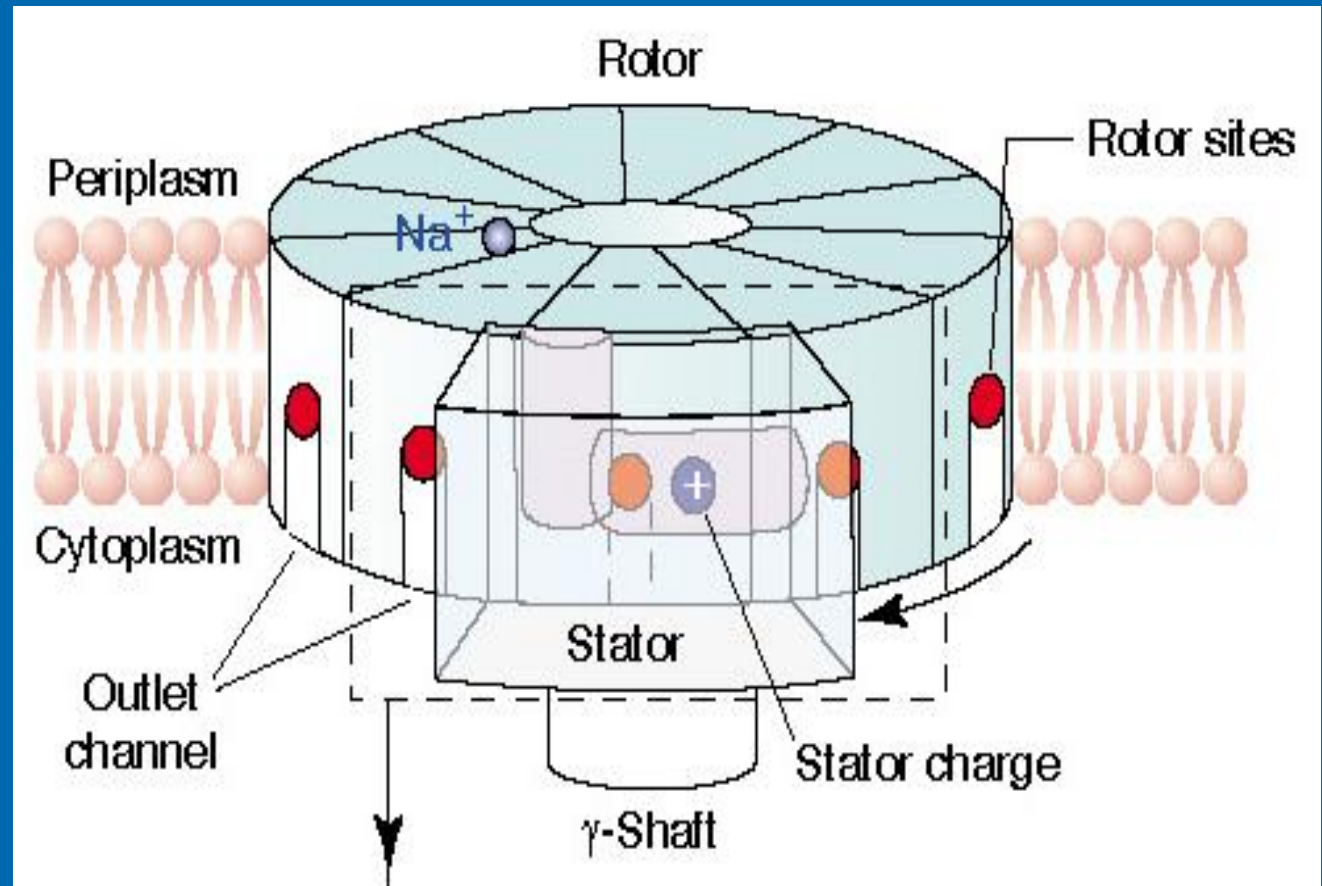
14 copies
in spinach
chloroplasts

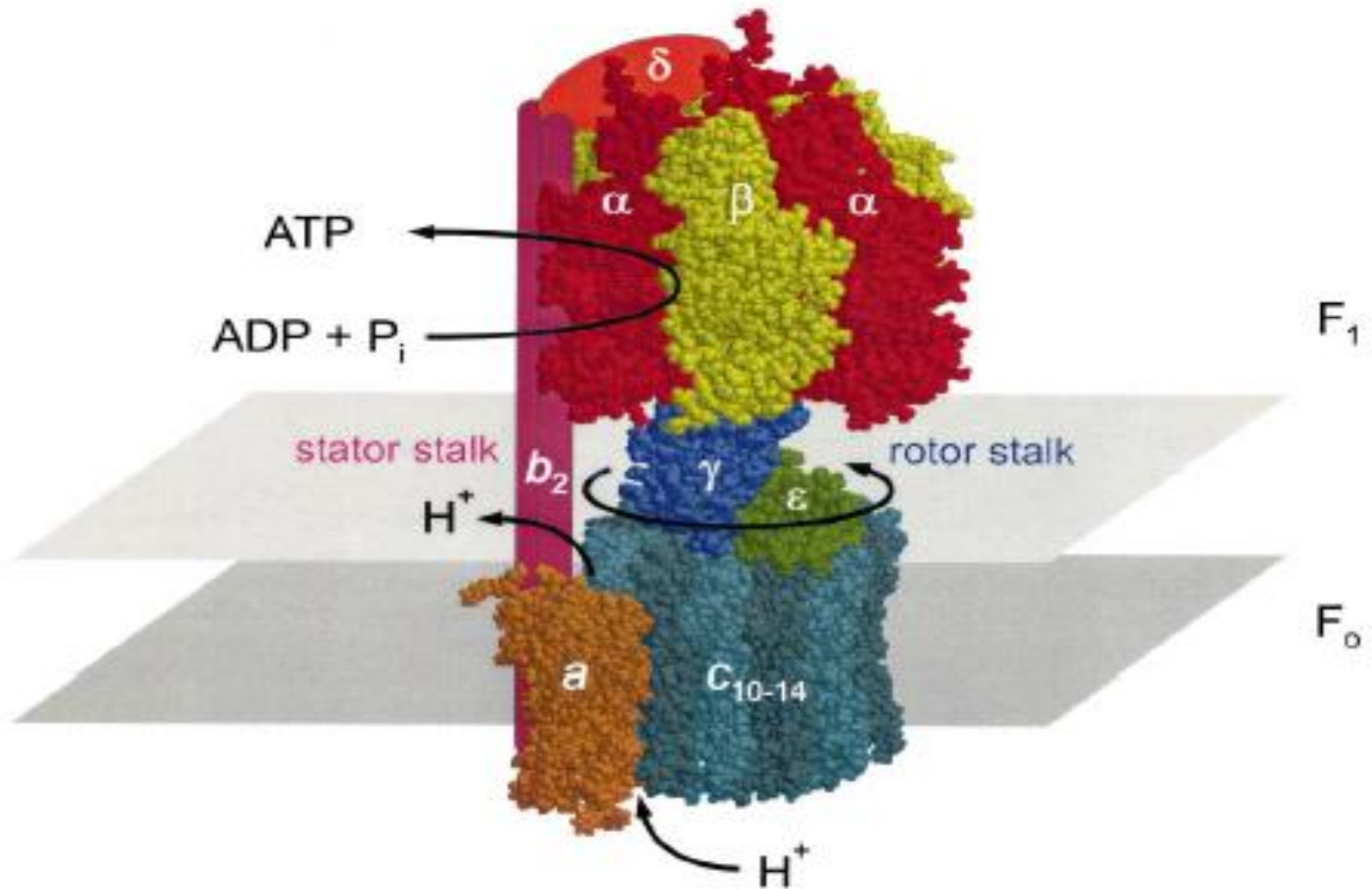
lyobacter



11 copies
in *lyobacter*

- The F_0 motor is made up of a transmembrane cylinder of 10–14 c subunits, depending on the species



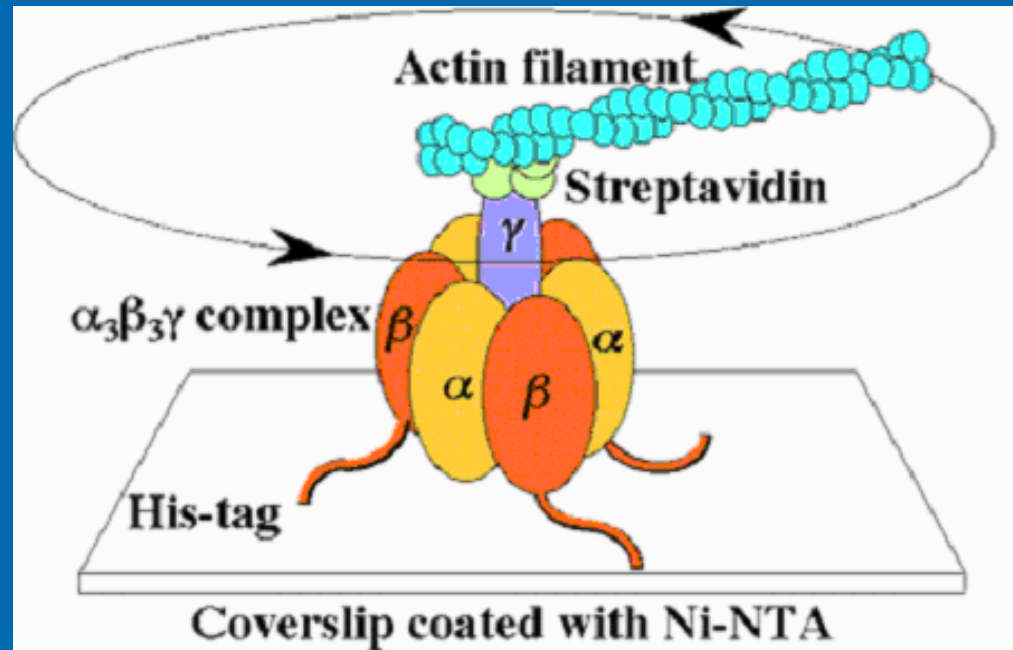


E.Coli ATP synthase

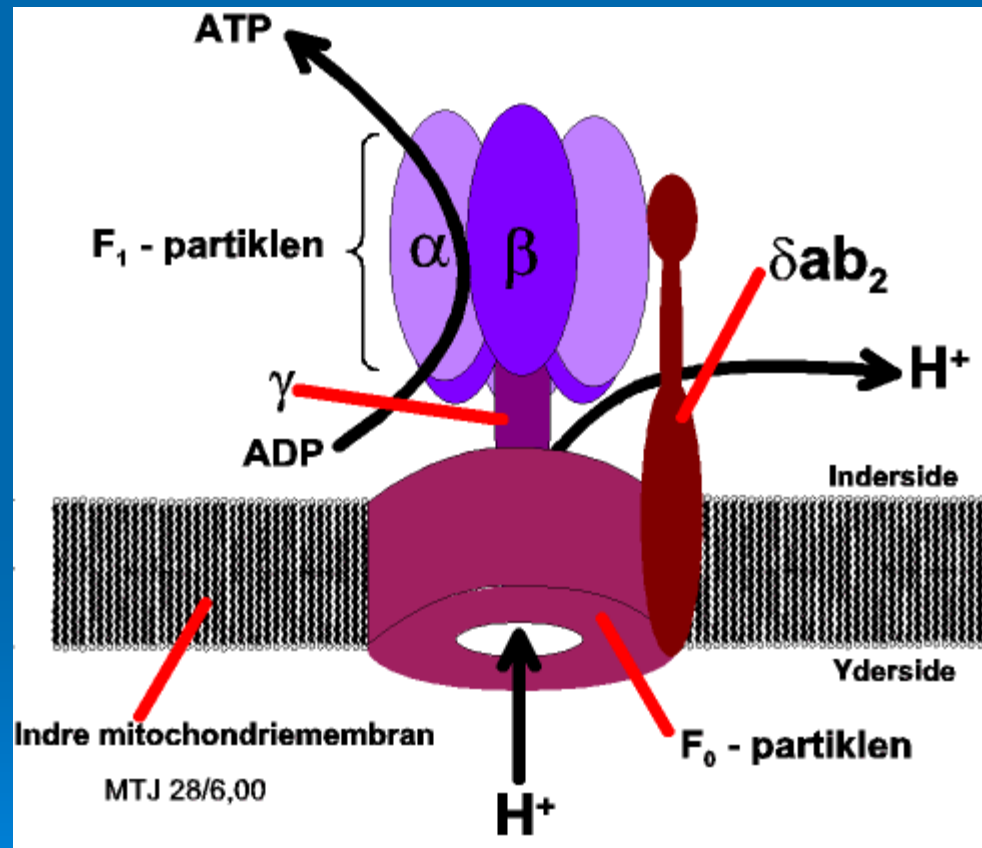
c, γ and ϵ are rotor, a, b, α , β and δ are stator

Visualize the rotation

- A $\alpha_3\beta_3\gamma$ complex from bacteria
- N terminus of β had polyhistidine tag attached, which enabled the anchoring of the complex to microscopic slide coated with **Ni ions**
- All the naturally occurring cysteine residues were removed, and insert one in γ at a position distant from the $\alpha\beta$ core
- A long and **fluorescent labeled actin filament** was attached to the **single introduced cysteine**







How to prepare your nano-materials?

1. **Protein molecules** by
genetic engineering

2. **Nano propeller** by
nanofabricated technology

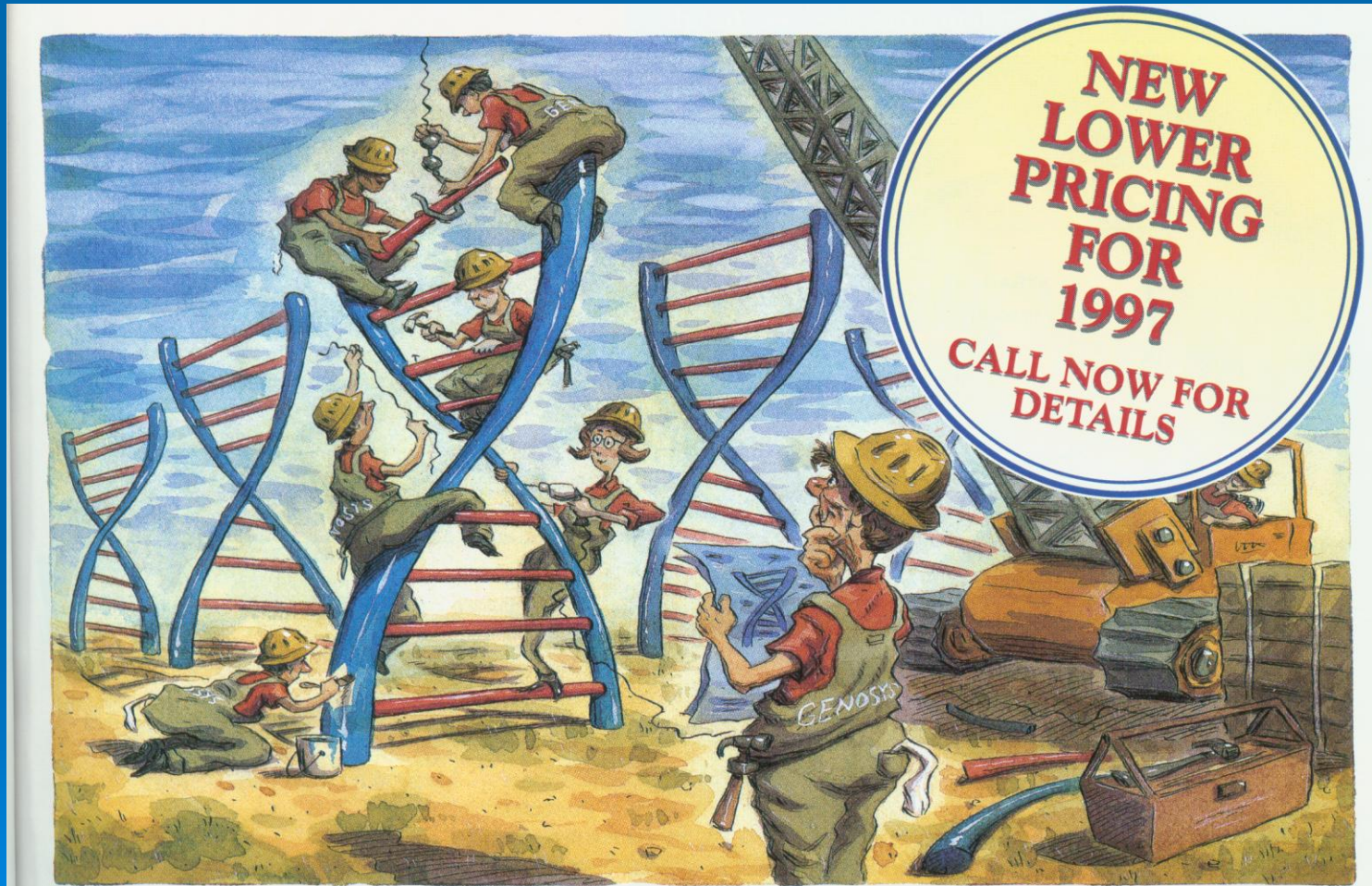
How to **assemble** these materials?

Recombinant protein production

Genetic engineering



基因工程(Genetic Engineering)

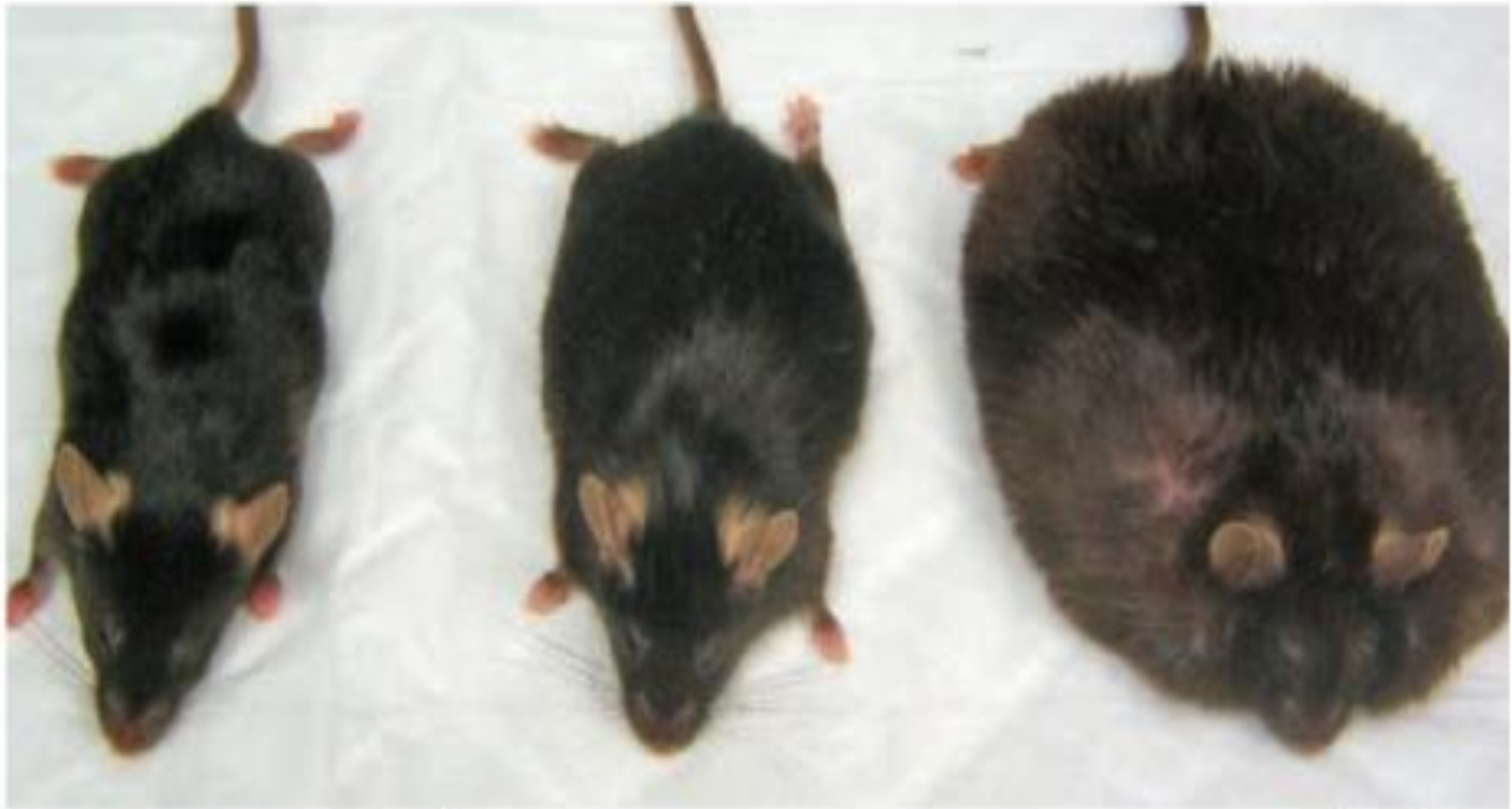




The Obesity Gene

The mouse on the left has a defective gene that causes obesity. Researchers have found a human relative of this gene.





Trimming down. Mice that can't respond to the appetite-regulating hormone leptin grow obese (*right*). Mice lacking the perilipin protein that coats lipid droplets burn off the excess fat and become almost as slender (*middle*) as normal mice (*left*).

Science, VOL.311, 2006, P. 1232-1234

比利時藍牛 (Belgian blue)



Mutation in Myostatin



Neonate

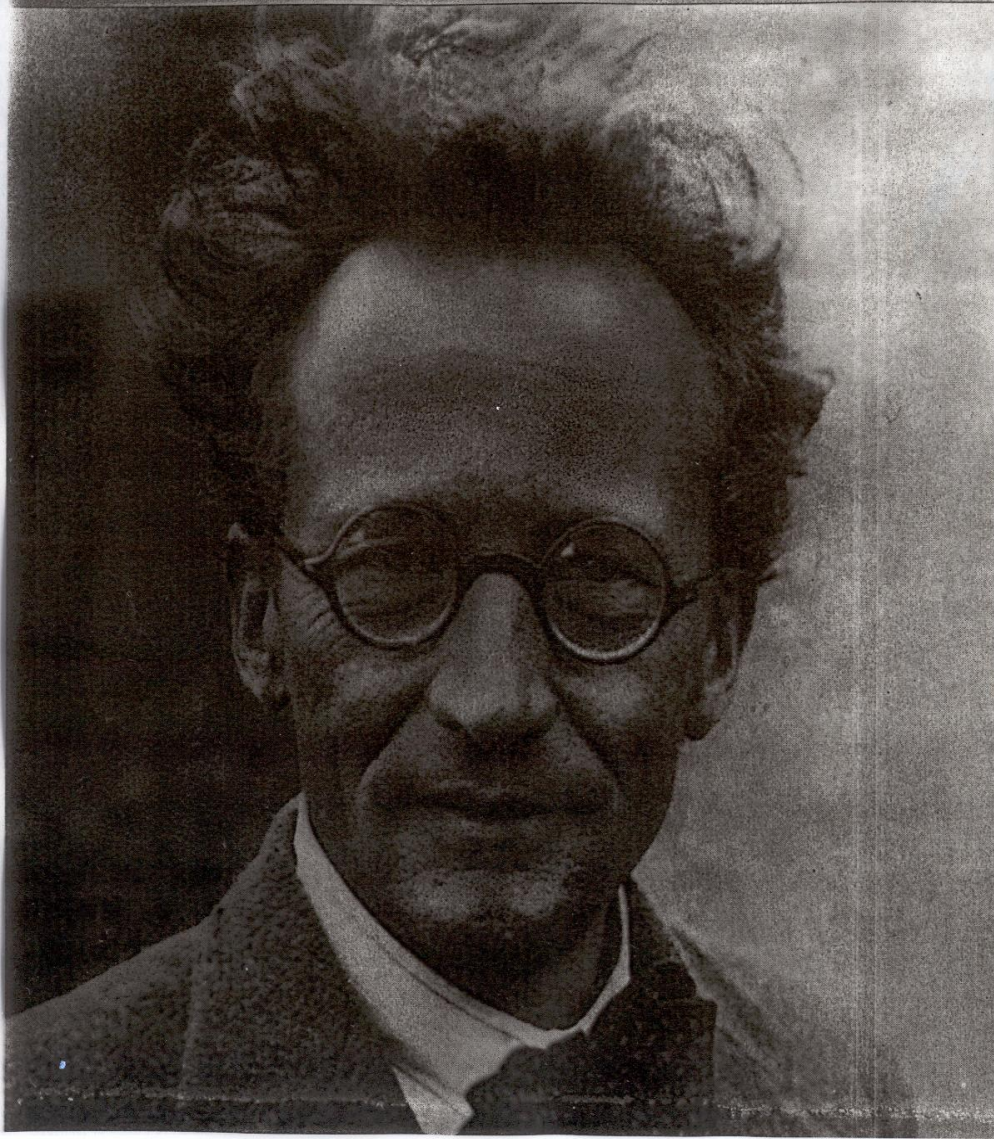
7 Months

Natural boost. A Berlin child who carries a mutation in the myostatin gene has had bodybuilder muscles since birth.



Schrödinger

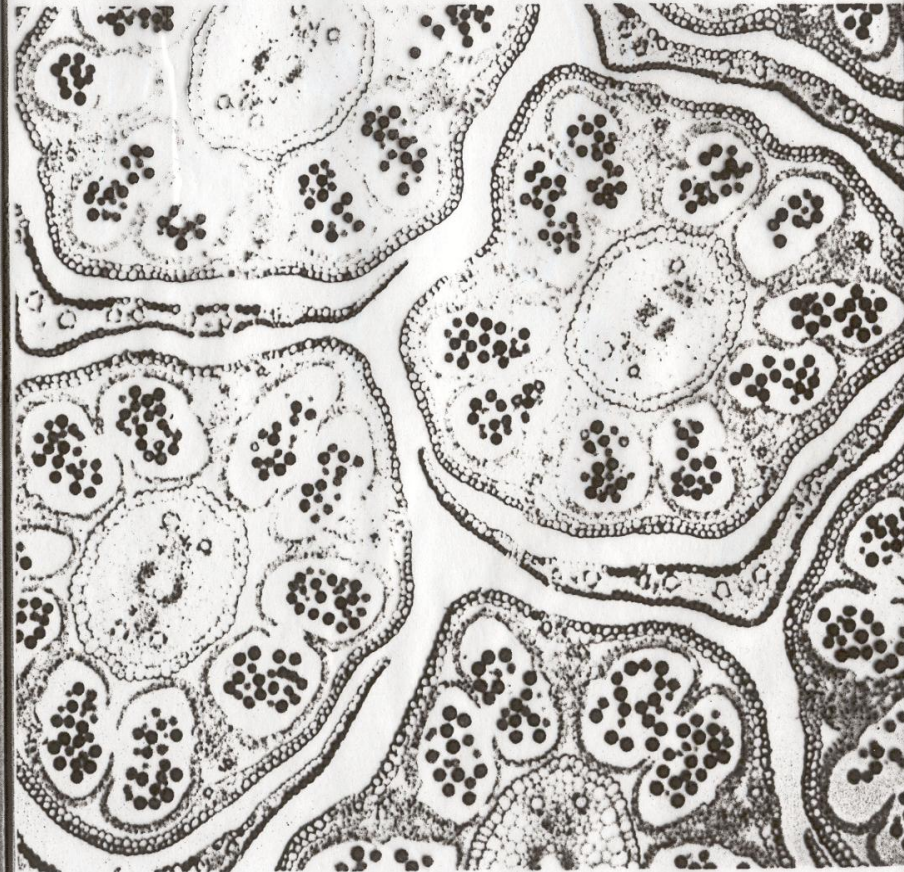
LIFE AND THOUGHT



WHAT IS LIFE?

*with Mind and Matter and
Autobiographical Sketches*

ERWIN SCHRÖDINGER



Canto

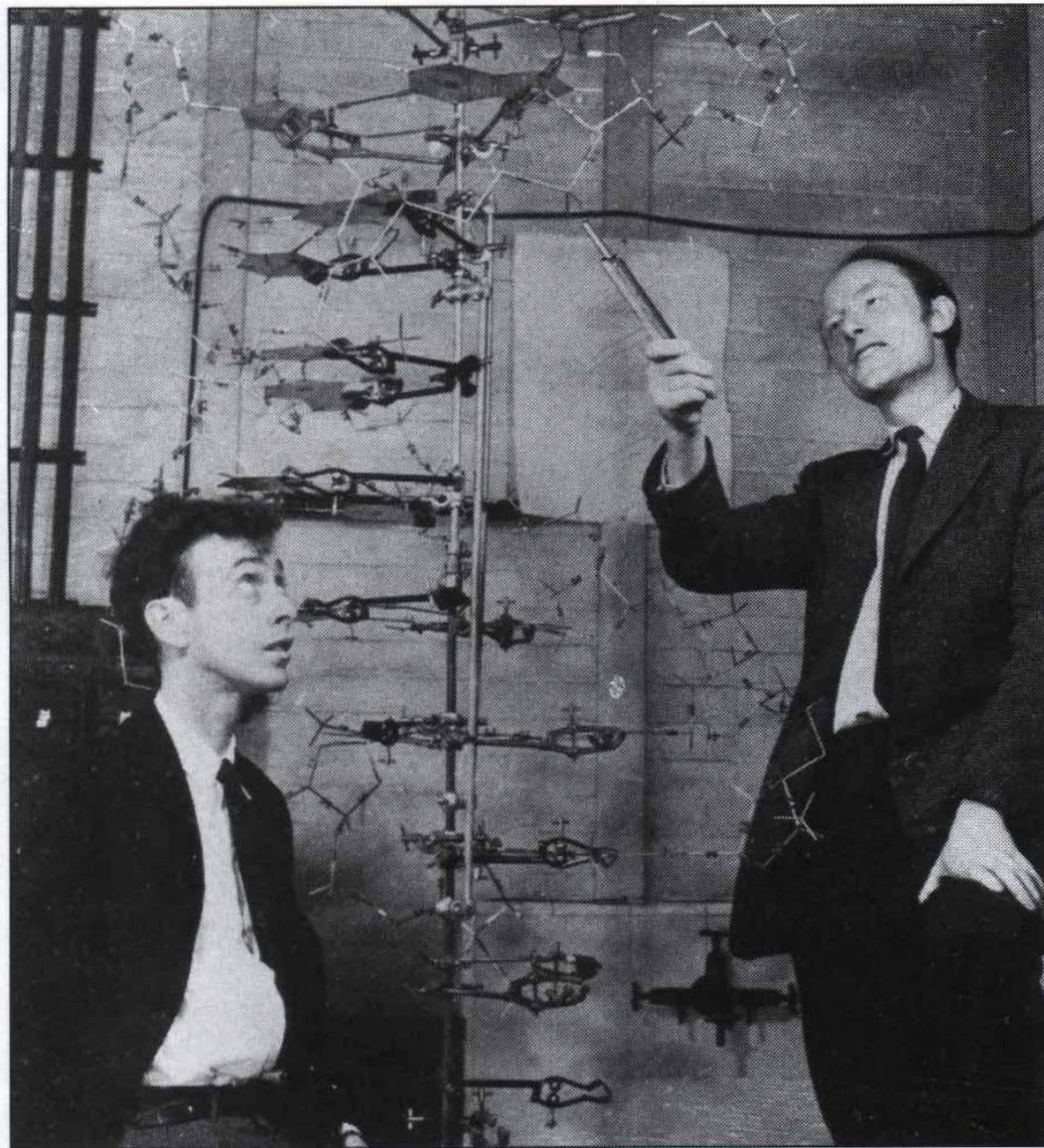
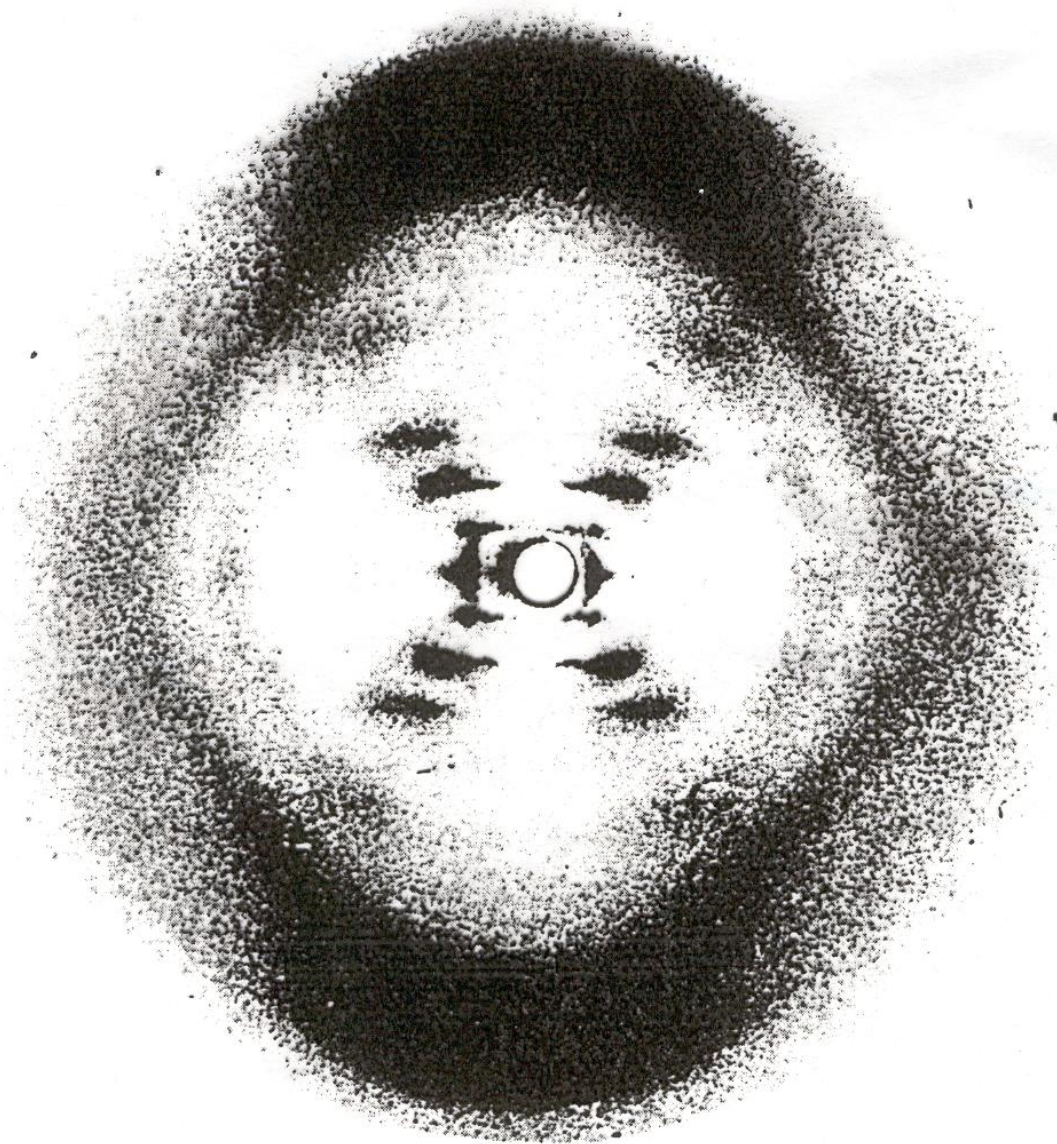
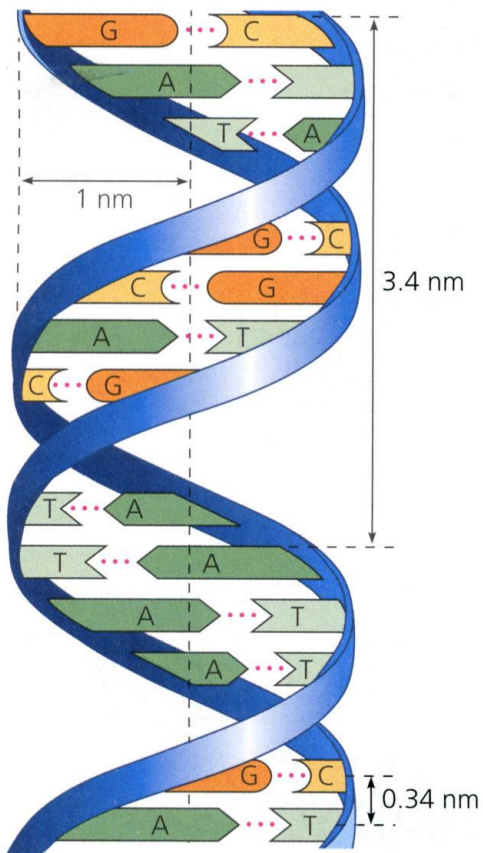


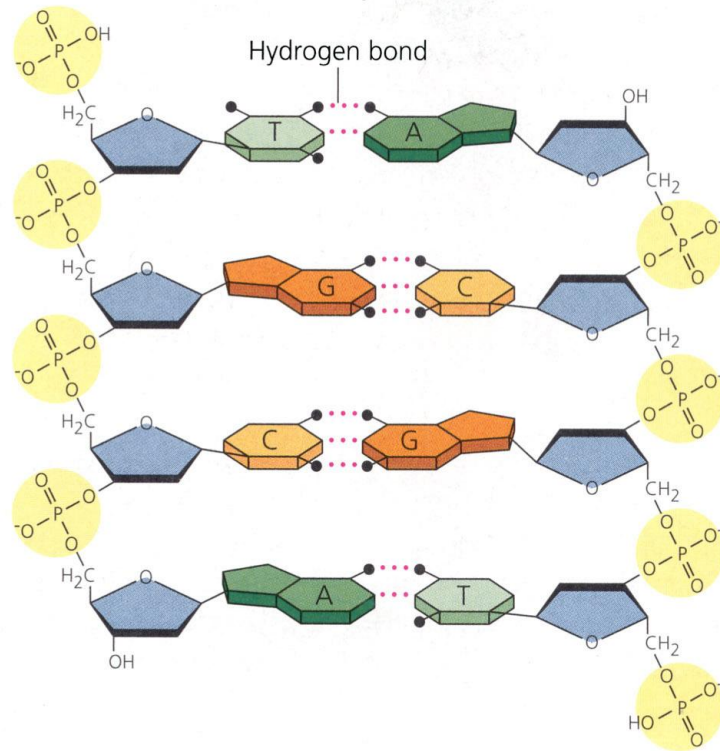
Figure 1.9 James Watson (left) and Francis Crick.



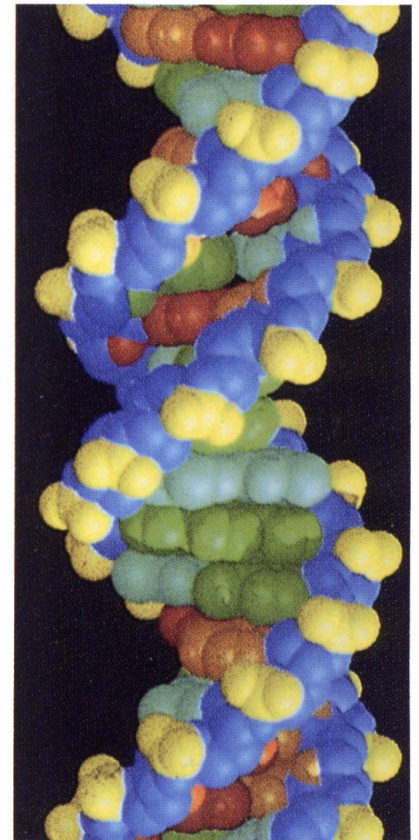
*An X-ray photograph of DNA in the B form,
taken by Rosalind Franklin late in 1952.*



(a) Key features of DNA structure

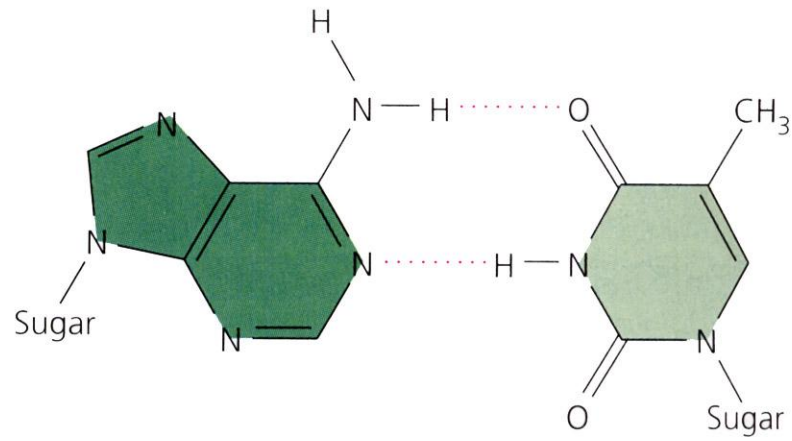


(b) Partial chemical structure



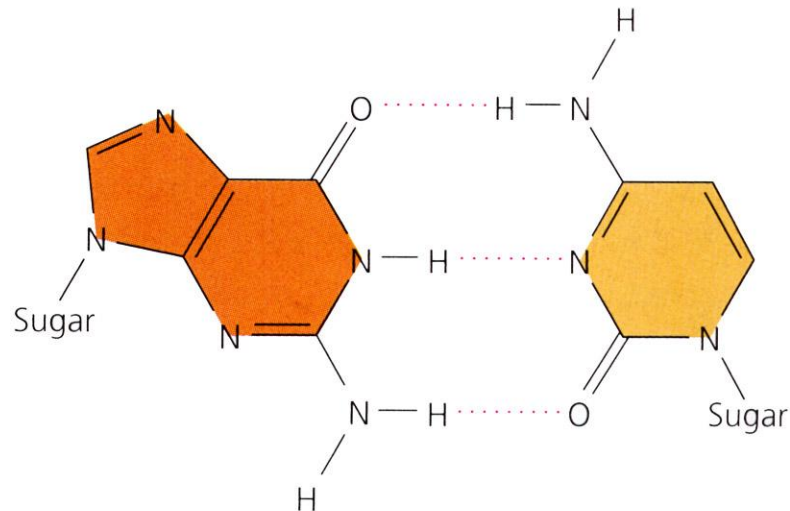
(c) Space-filling model

FIGURE 16.5 The double helix.



Adenine (A)

Thymine (T)



Guanine (G)

Cytosine (C)

FIGURE 16.6 Base pairing in DNA.

基因: 染色體上的雙股DNA 的排列順序



.....ATGCCGTA.....

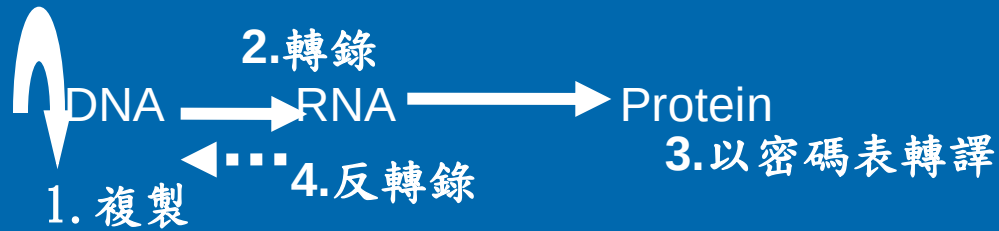
.....TACGGCAT.....

BED

BAD

BAT

← 單字



		Second base					
		U	C	A	G		
First base (5' end)	U	UUU } Phe	UCU } Ser	UAU } Tyr	UGU } Cys	U	Third base (3' end)
		UUC }	UCC }	UAC }	UGC }	C	
		UUA } Leu	UCA }	UAA Stop	UGA Stop	A	
		UUG }	UCG }	UAG Stop	UGG Trp	G	
	C	CUU } Leu	CCU } Pro	CAU } His	CGU } Arg	U	
		CUC }	CCC }	CAC }	CGC }	C	
		CUA }	CCA }	CAA } Gln	CGA }	A	
		CUG }	CCG }	CAG }	CGG }	G	
	A	AUU } Ile	ACU } Thr	AAU } Asn	AGU } Ser	U	
		AUC }	ACC }	AAC }	AGC }	C	
		AUA }	ACA }	AAA } Lys	AGA } Arg	A	
		AUG Met or start	ACG }	AAG }	AGG }	G	
	G	GUU } Val	GCU } Ala	GAU } Asp	GGU } Gly	U	
		GUC }	GCC }	GAC }	GGC }	C	
		GUA }	GCA }	GAA } Glu	GGA }	A	
		GUG }	GCG }	GAG }	GGG }	G	

圖二 生命科學中定理與密碼表

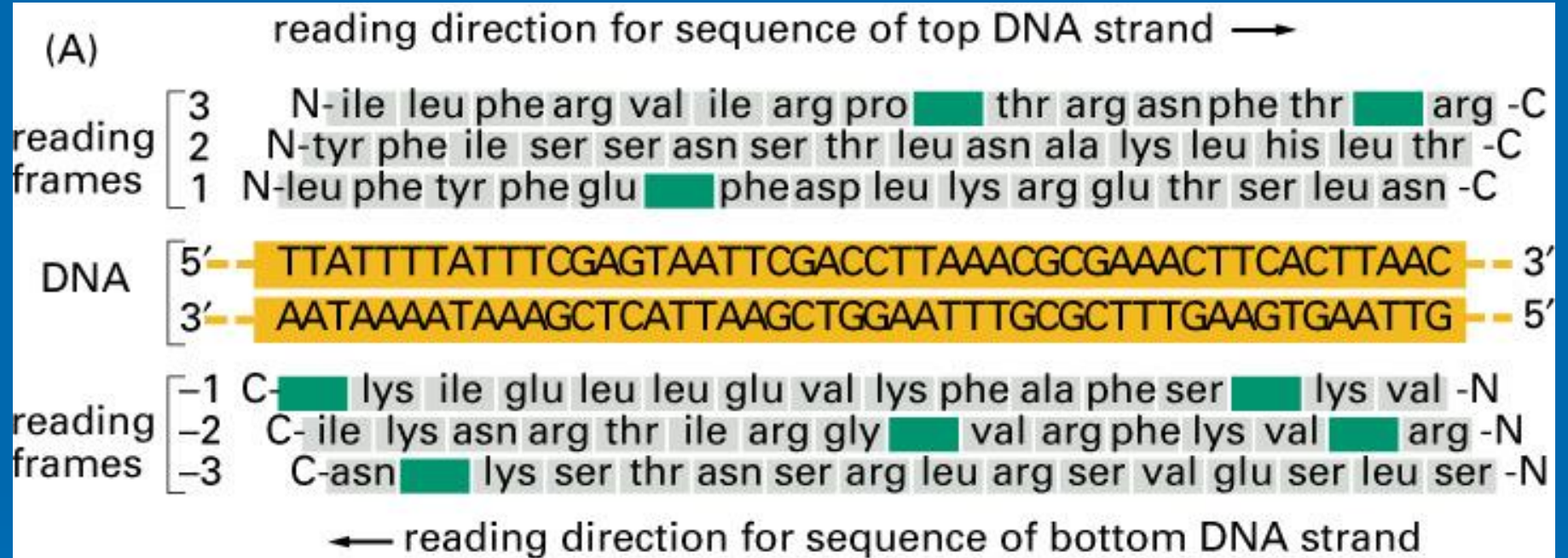


Figure 8-38 part 1 of 2. Molecular Biology of the Cell, 4th Edition.

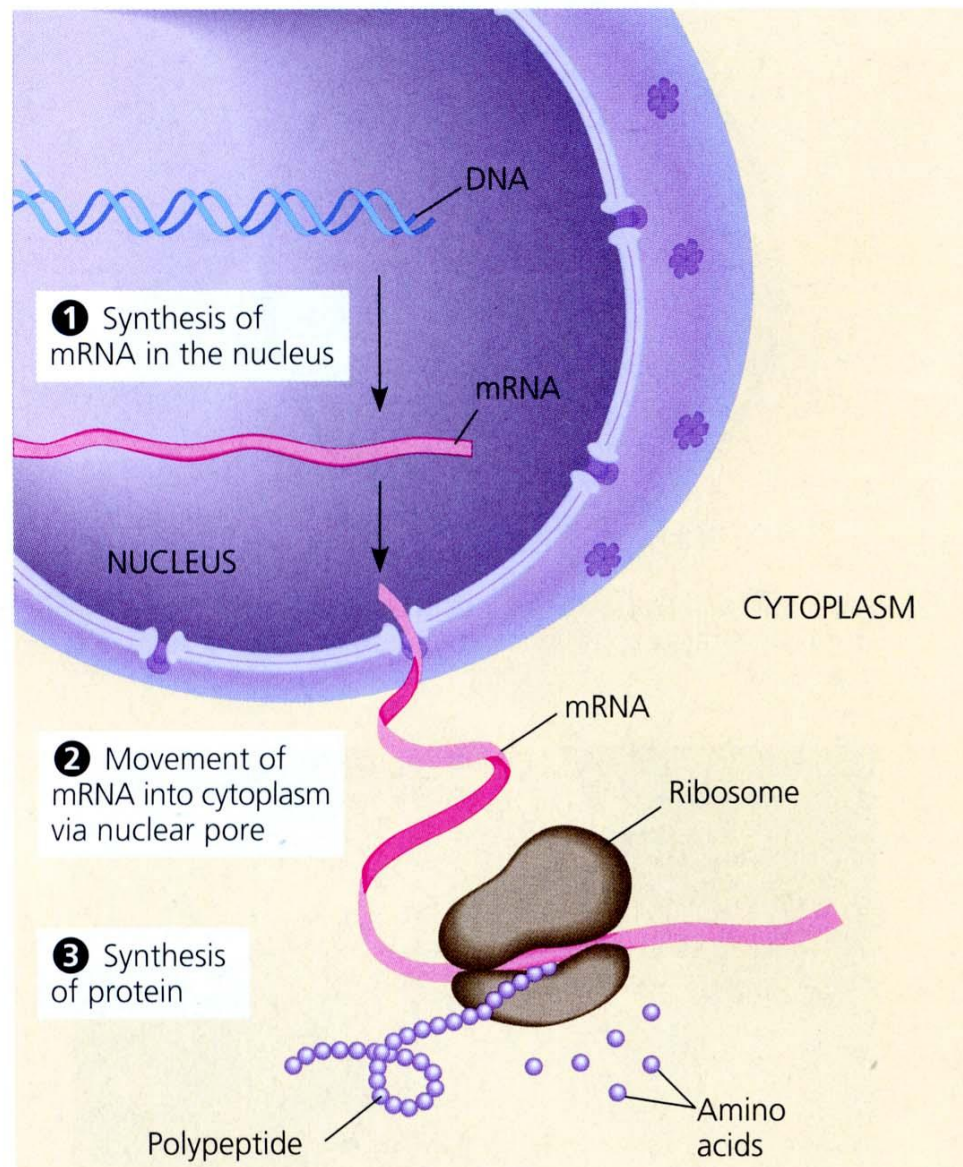


FIGURE 5.28 DNA → RNA → protein: a diagrammatic overview of information flow in a cell.

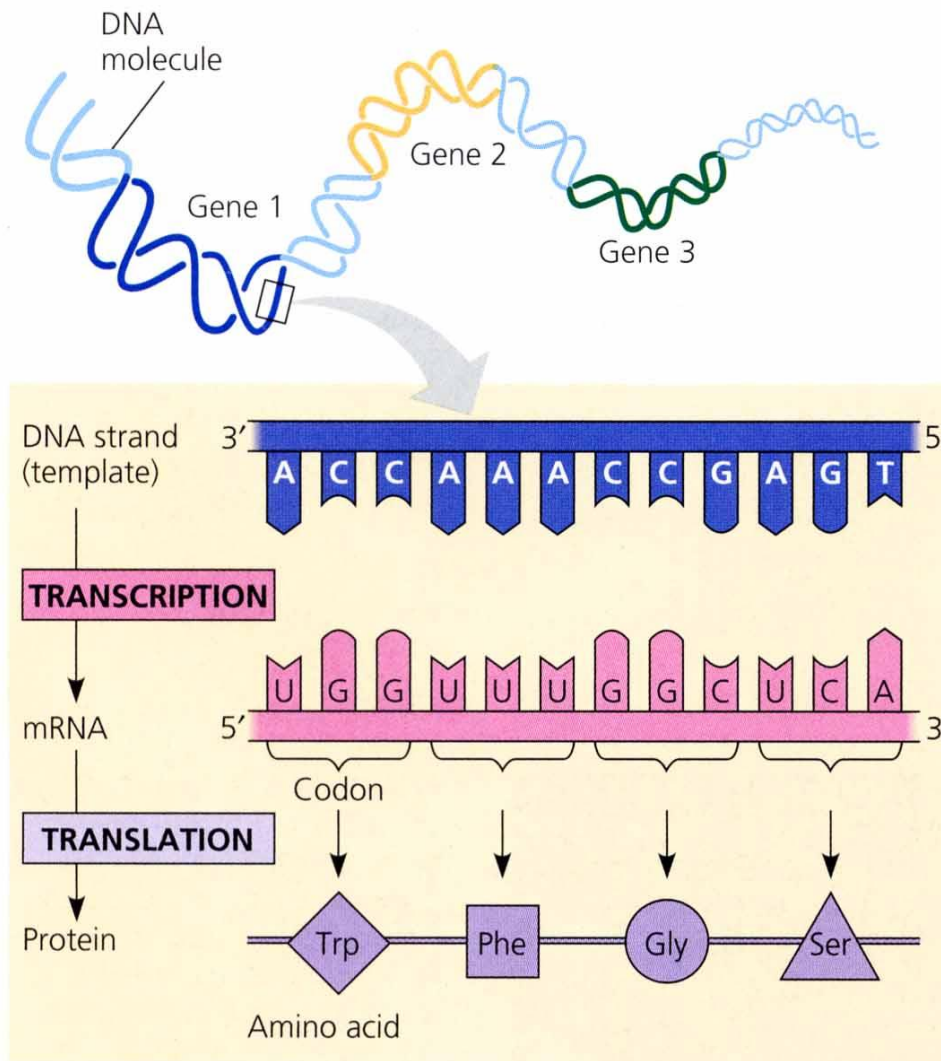


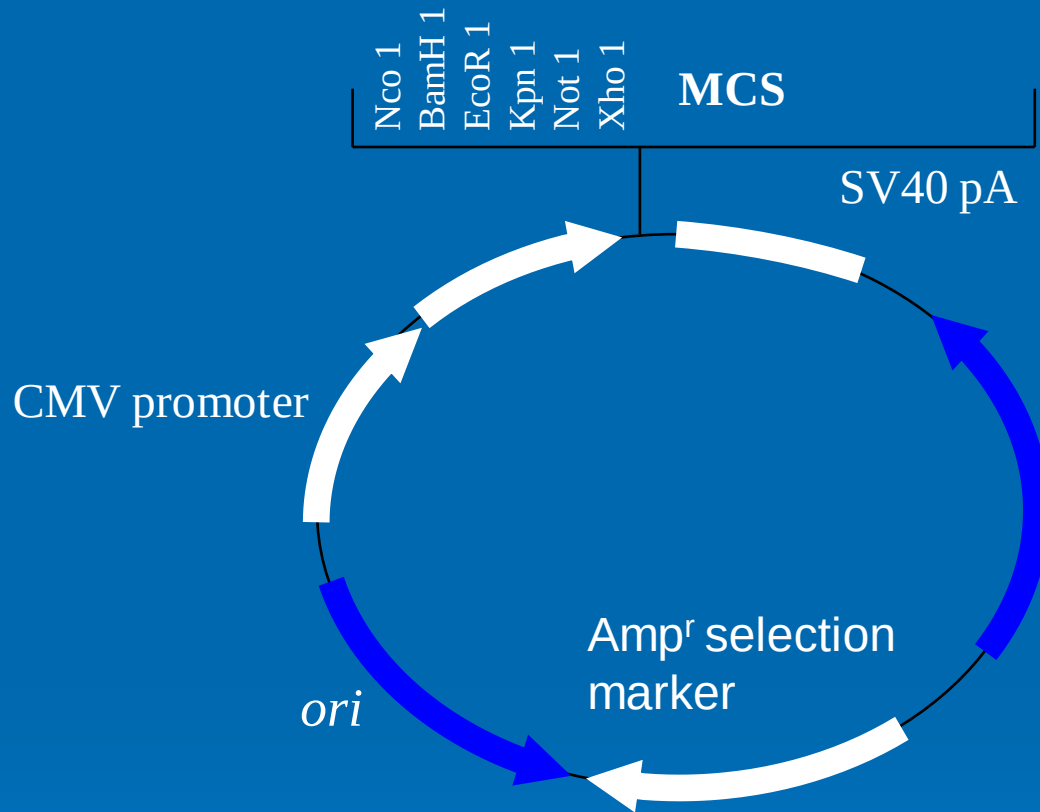
FIGURE 17.3 The triplet code.



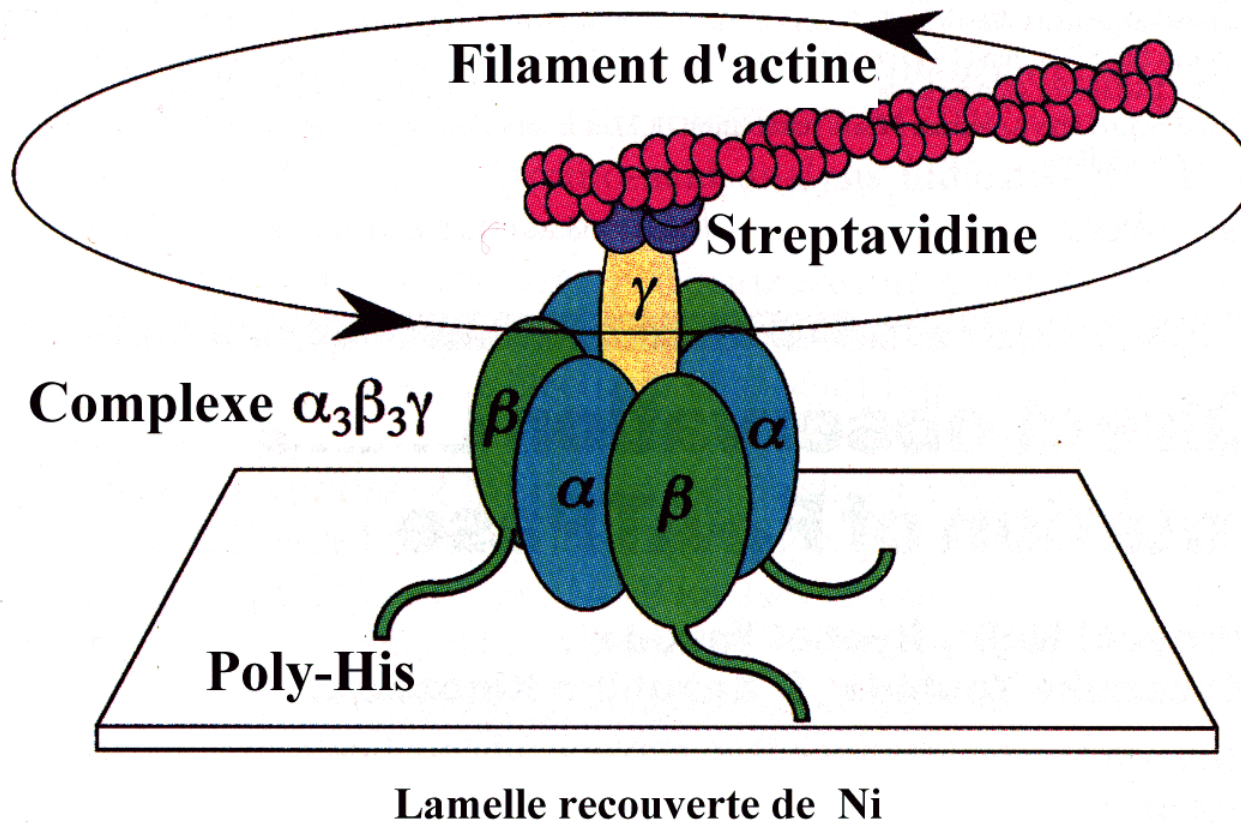
FIGURE 17.5 A tobacco plant expressing a firefly gene.

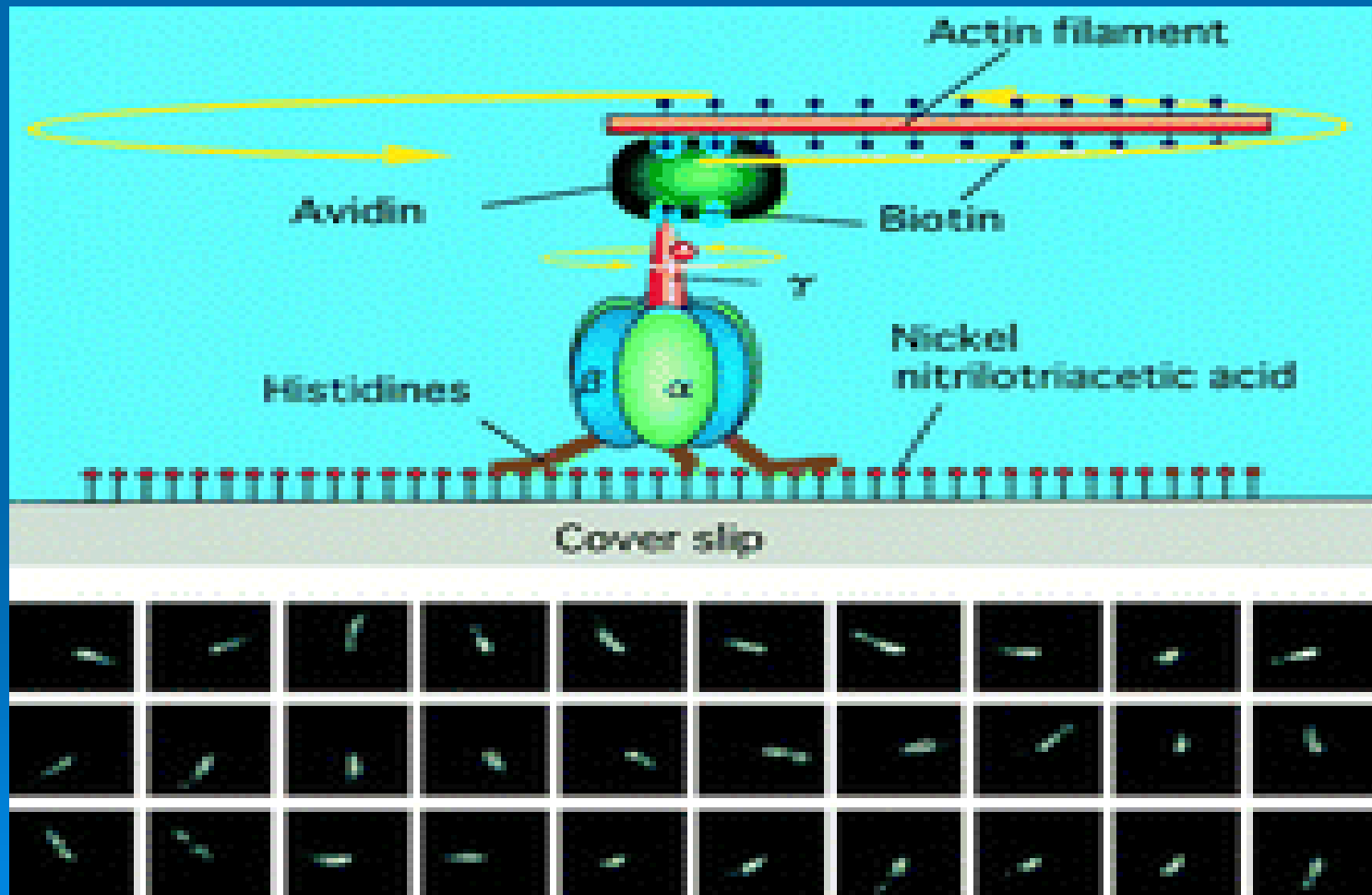
1ml IRES
6'th days





圖三:表現載體 (expression vector)



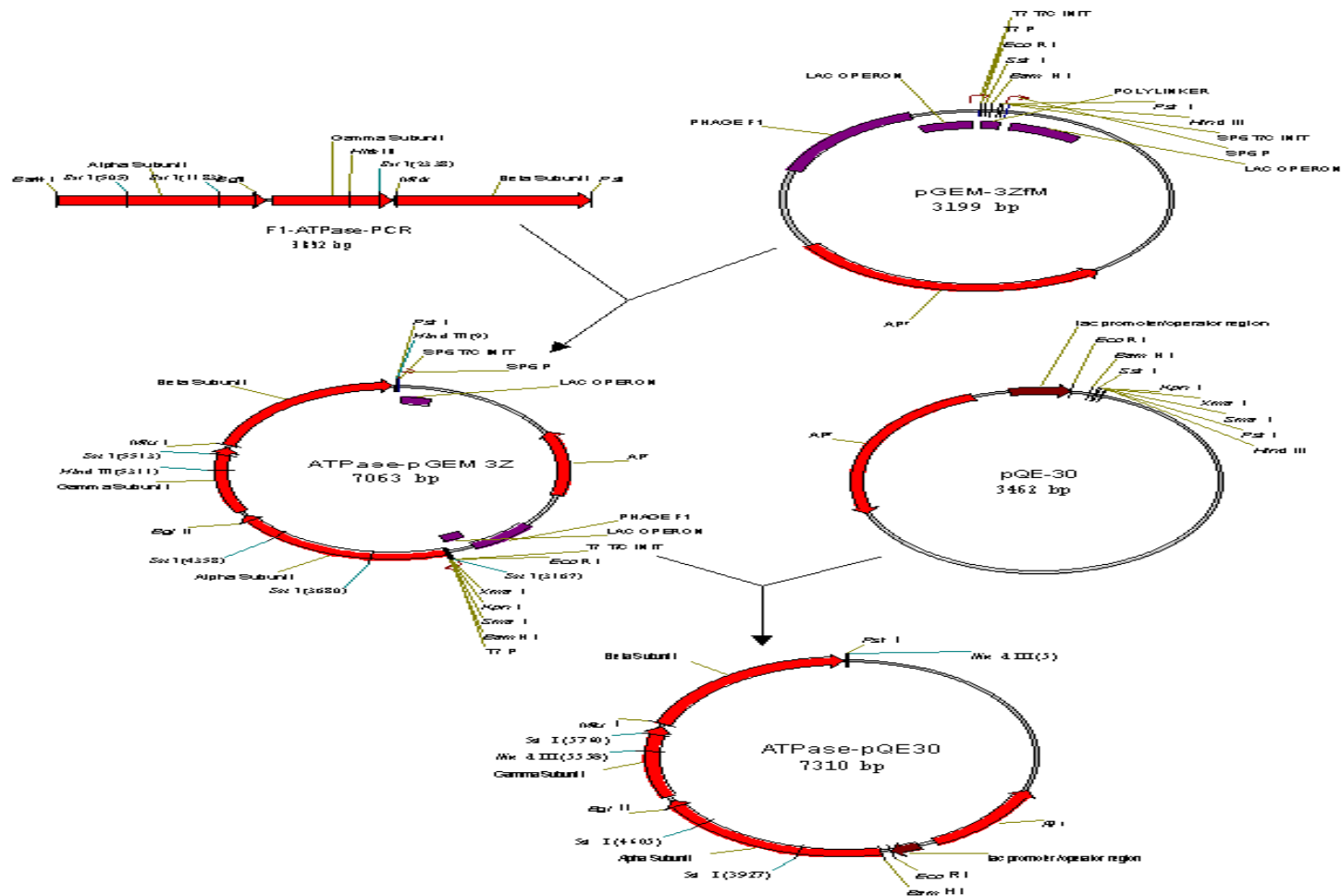


Powering an Inorganic Nanodevice with a Biomolecular Motor

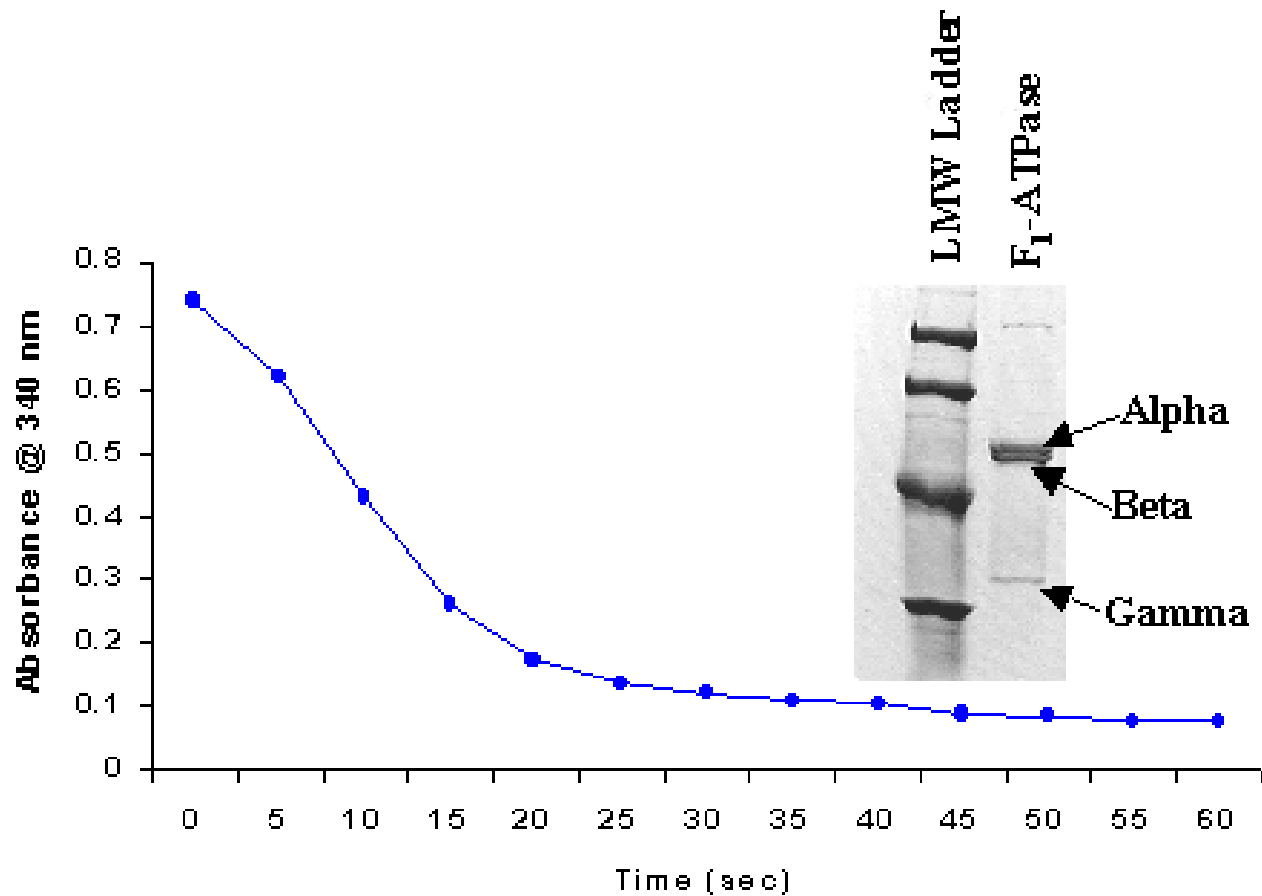
**Ricky K. Soong,^{1,2} George D. Bachand,^{1,2} Hercules P. Neves,^{1,2}
Anatoli G. Olkhovets,^{1,3} Harold G. Craighead,^{1,3}
Carlo D. Montemagno^{1,2*}**

Biomolecular motors such as F_1 -adenosine triphosphate synthase (F_1 -ATPase) and myosin are similar in size, and they generate forces compatible with currently producible nanoengineered structures. We have engineered individual biomolecular motors and nanoscale inorganic systems, and we describe their integration in a hybrid nanomechanical device powered by a biomolecular motor. The device consisted of three components: an engineered substrate, an F_1 -ATPase biomolecular motor, and fabricated nanopropellers. Rotation of the nanopropeller was initiated with 2 mM adenosine triphosphate and inhibited by sodium azide.

Cloning the ATPase from thermophilic *Bacillus*



Protein analysis by SDS-PAGE



Nanofabricated technology

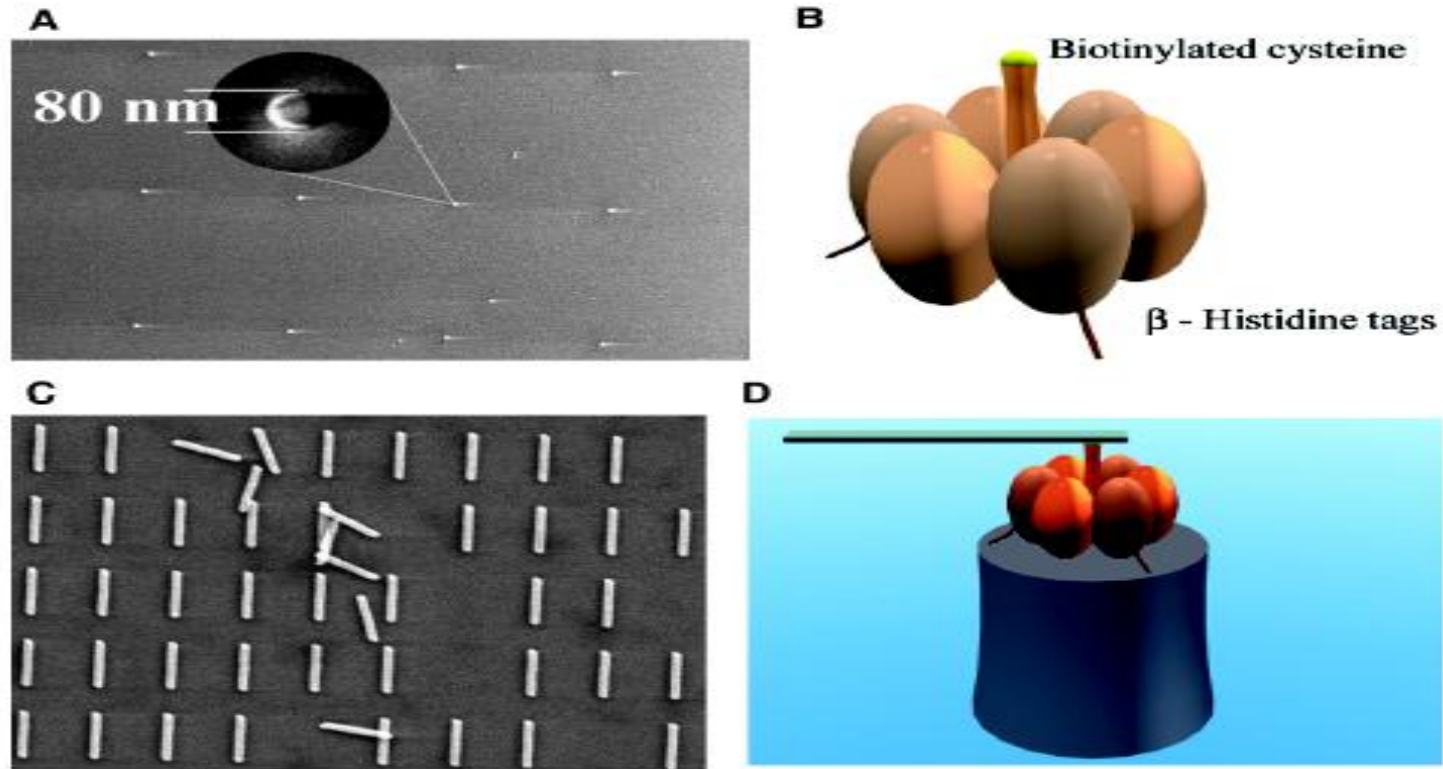
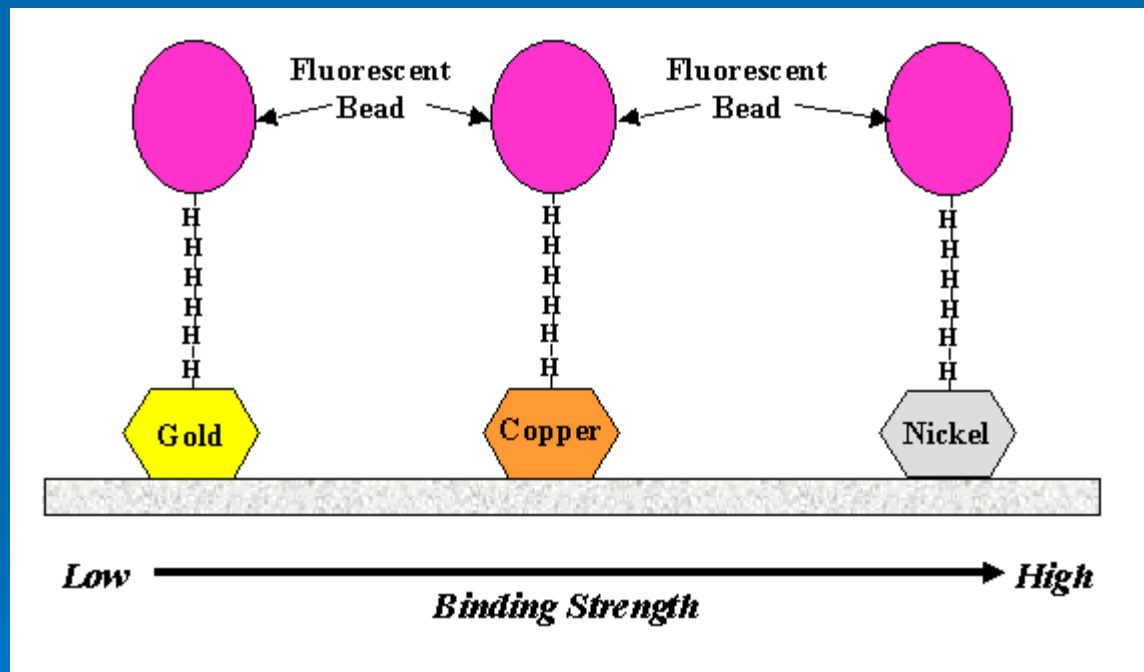


Fig. 1. Schematic diagram of the F₁-ATPase biomolecular motor-powered nanomechanical device. The device consisted of (A) a Ni post (height 200 nm, diameter 80 nm), (B) the F₁-ATPase biomolecular motor, and (C) a nanopropeller (length 750 to 1400 nm, diameter 150 nm). The device (D) was assembled using sequential additions of individual components and differential attachment chemistries.

The assembly of protein and nano-inorganic element



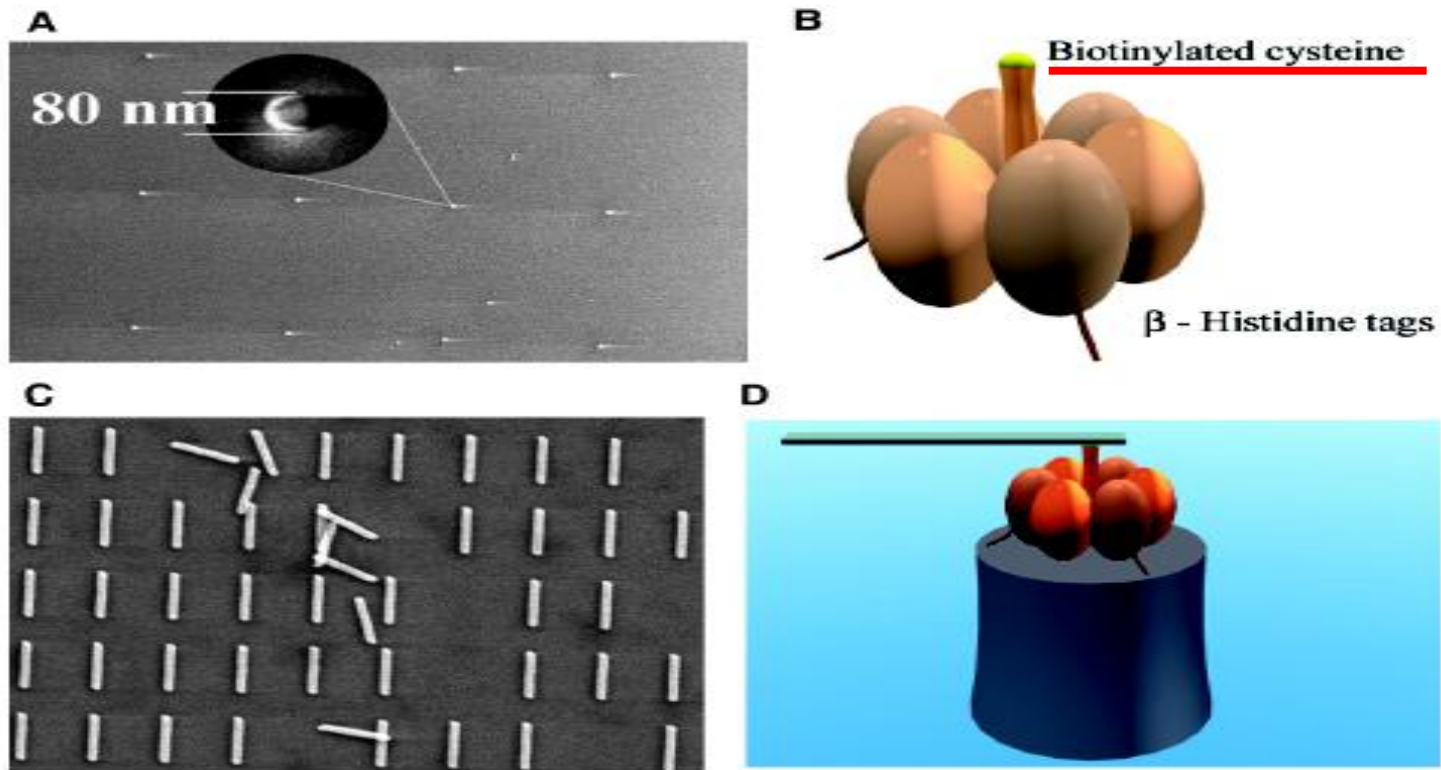


Fig. 1. Schematic diagram of the F_1 -ATPase biomolecular motor-powered nanomechanical device. The device consisted of (A) a Ni post (height 200 nm, diameter 80 nm), (B) the F_1 -ATPase biomolecular motor, and (C) a nanopropeller (length 750 to 1400 nm, diameter 150 nm). The device (D) was assembled using sequential additions of individual components and differential attachment chemistries.

table 7-1

Some Protein Dissociation Constants

Protein	Ligand	K_d (M)*
Avidin (egg white) [†]	Biotin	1×10^{-15}
Insulin receptor (human)	Insulin	1×10^{-10}
Anti-HIV immunoglobulin (human) [‡]	gp41 (HIV-1 surface protein)	4×10^{-10}
Nickel-binding protein (<i>E. coli</i>)	Ni ²⁺	1×10^{-7}
Calmodulin (rat) [§]	Ca ²⁺	3×10^{-6}
		2×10^{-5}

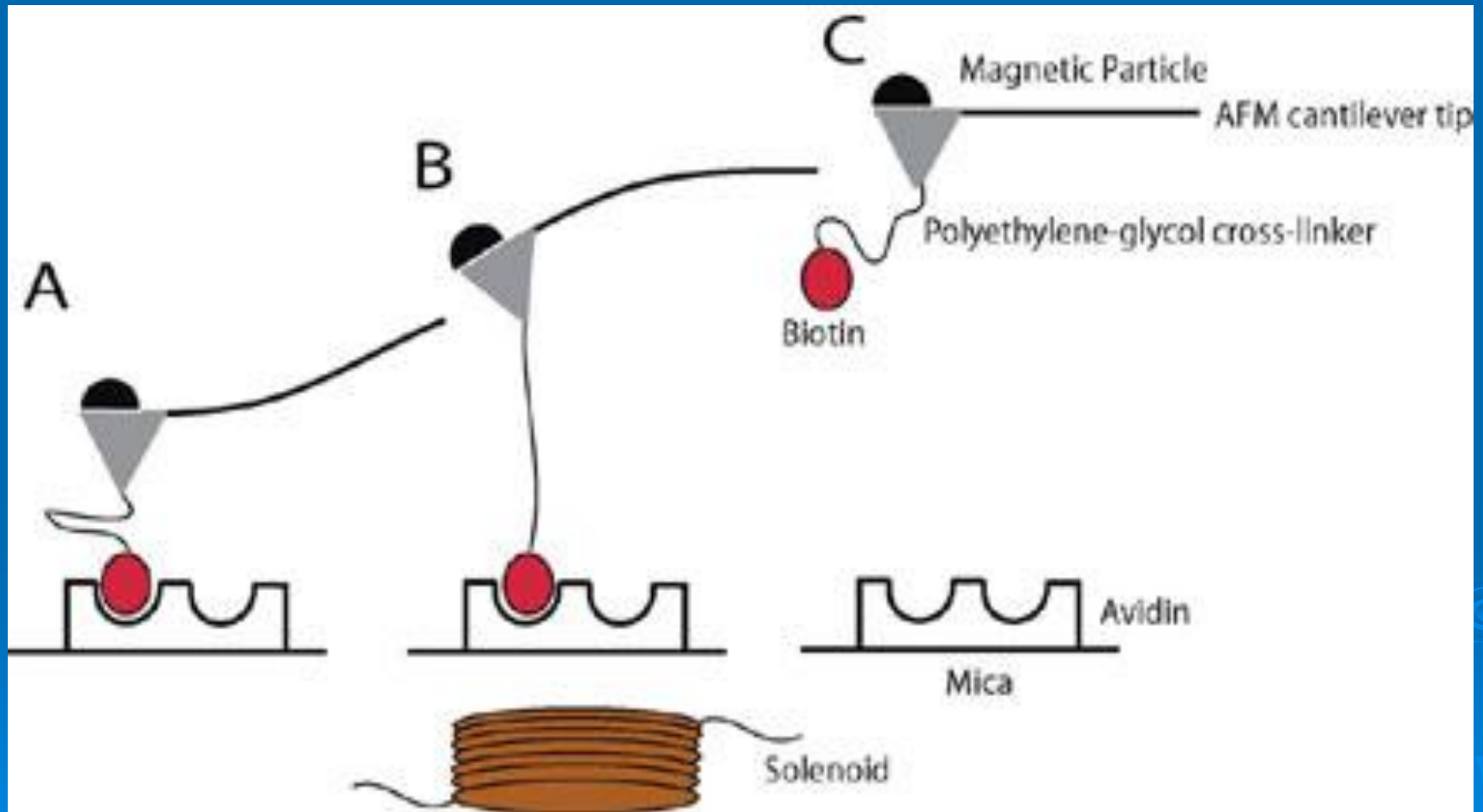
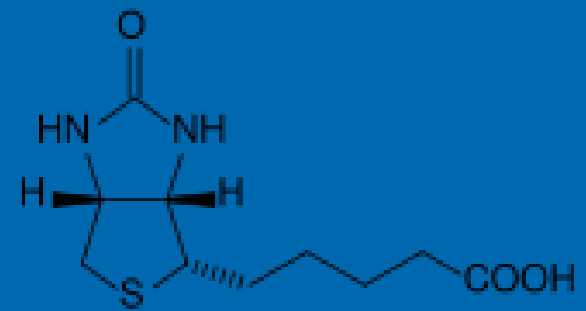
*A reported dissociation constant is valid only for the particular solution conditions under which it was measured. K_d values for a protein-ligand interaction can be altered, sometimes by several orders of magnitude, by changes in solution salt concentration, pH, or other variables.

[†]Interaction of avidin with the enzymatic cofactor biotin is among the strongest noncovalent biochemical interactions known.

[‡]This immunoglobulin was isolated as part of an effort to develop a vaccine against HIV. Immunoglobulins (described later in the chapter) are highly variable, and the K_d reported here should not be considered characteristic of all immunoglobulins.

[§]Calmodulin has four binding sites for calcium. The values shown reflect the highest- and lowest-affinity binding sites observed in one set of measurements.

Biotin



The observation of nanomotor work

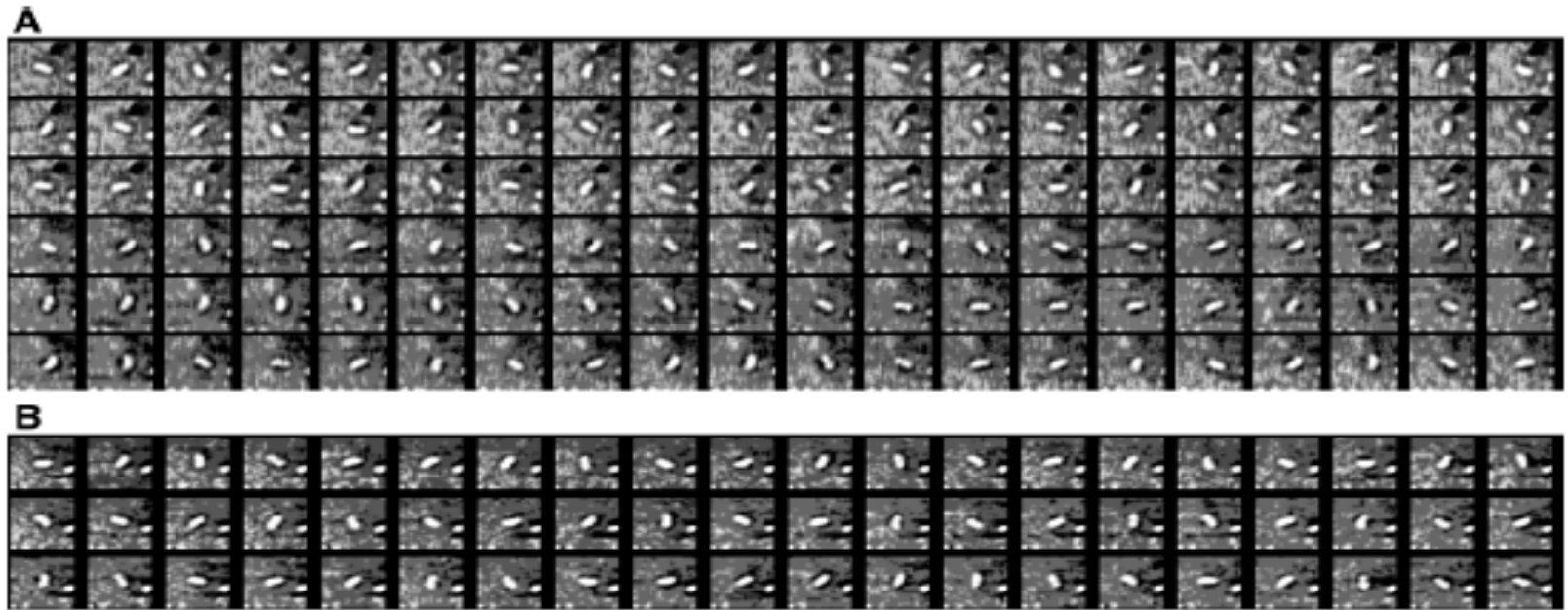
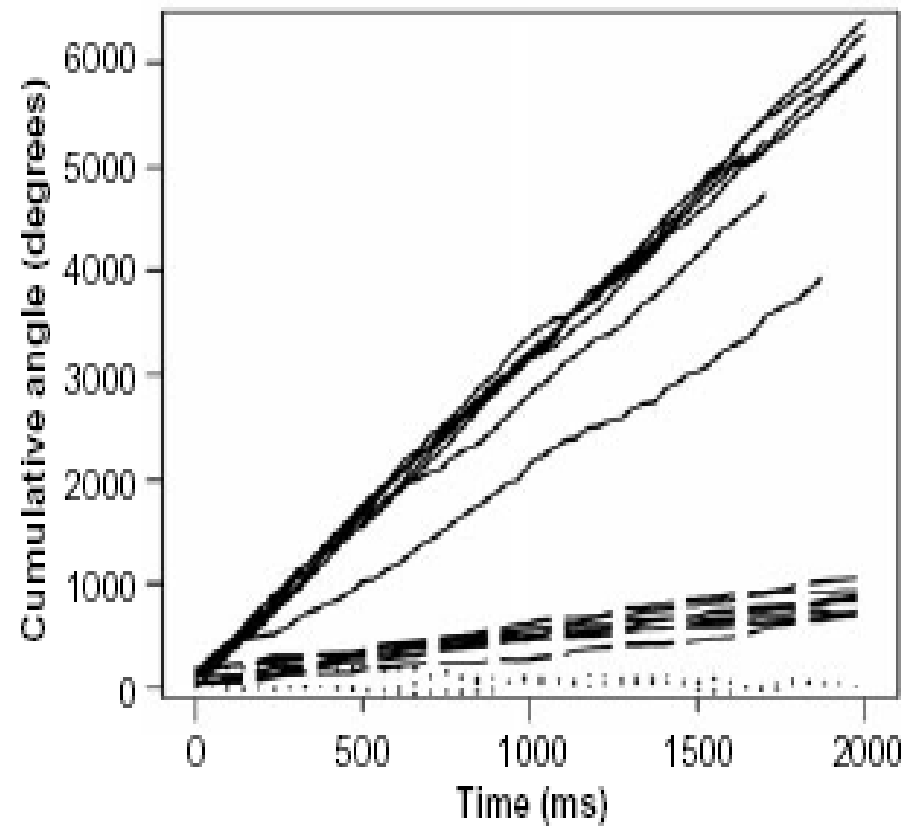


Fig. 2. Image sequence (viewed left to right) of nanopropellers being rotated anticlockwise at 8.3 rps (A) and 7.7 rps (B) by the F_1 -ATPase biomolecular motor. Observations were made using 100 $\times$ oil immersion or 60 $\times$ water immersion and were captured with a CCD video camera (frame rate 30 Hz). The rotational velocity ranged from ~ 0.8 to 8.3 rps, depending on propeller length. Data were recorded for up to 30 min; however, propellers rotated for almost 2.5 hours while ATP was maintained in the flow cell. These sequences can be viewed as movies at the Nanoscale Biological Engineering and Transport Group Web site (<http://falcon.aben.cornell.edu/News2.htm>).

Fig. 3. Time course of F_1 -ATPase γ sub-unit rotation. Each line represents data from a rotating nanopropeller. Solid lines, propellers 750 nm long; dashed lines, propellers 1400 nm long; dotted lines, propellers 1400 nm long in the presence of NaN_3 .





Thanks for your patient



謝謝指教

