

Introduction

class – 2 (07.08.24)

LS2103 (Autumn 2024)

IISER Kolkata

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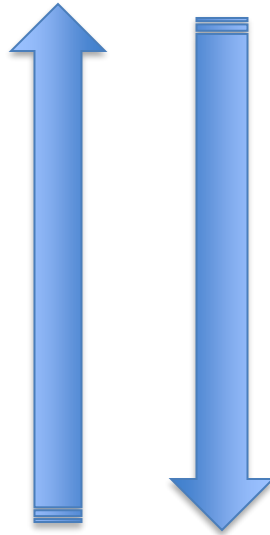
General Information:

1. **Approx. 28 classes, with 8 to 10 tutorial sessions**
2. **(-3) marks for not maintaining minimum 80% attendance**
3. **Evaluation:**
 - Internal assessment: Surprise class tests (best 4 out of 5): 20 marks
 - Mid-semester exam: 30 - 35 marks
 - End-semester: 45 - 50 marks
4. **Study Material:**
 - Lectures notes, extra material (to be shared)
 - Philip C. Nelson: *Biological Physics*
 - Phillips, Kondev, Theriot, Garcia: *Physical Biology of the Cell*

What is **Biophysics**?

Biology — The study of inception, sustenance and the evolution of life, in the context of the environment.

- Mechanistic understanding
- Bridge time, length scales
- Quantitative estimates
- Build falsifiable models, predictions

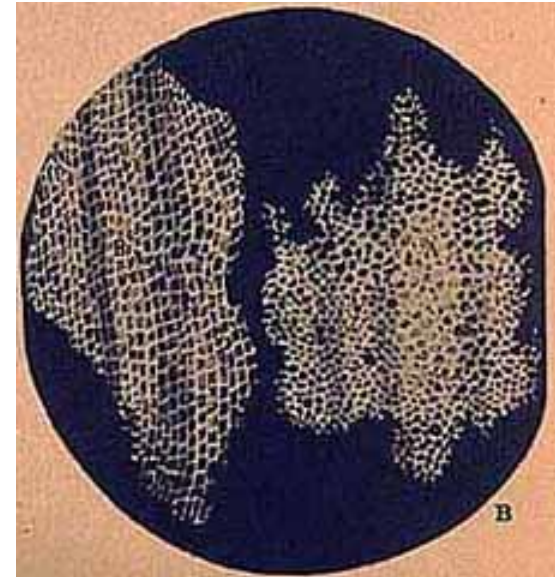


- Phenomenological problems
- Test physical hypotheses
- Make measurements
- Develop novel protocols

Physics — The study of matter and interactions: force, energy, charge, radiation, etc.

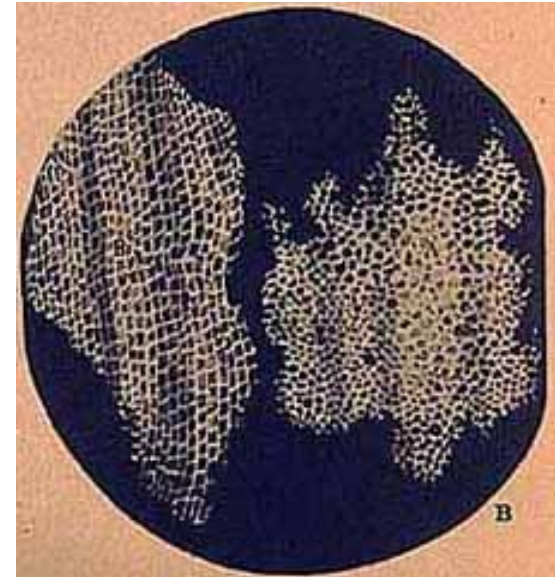
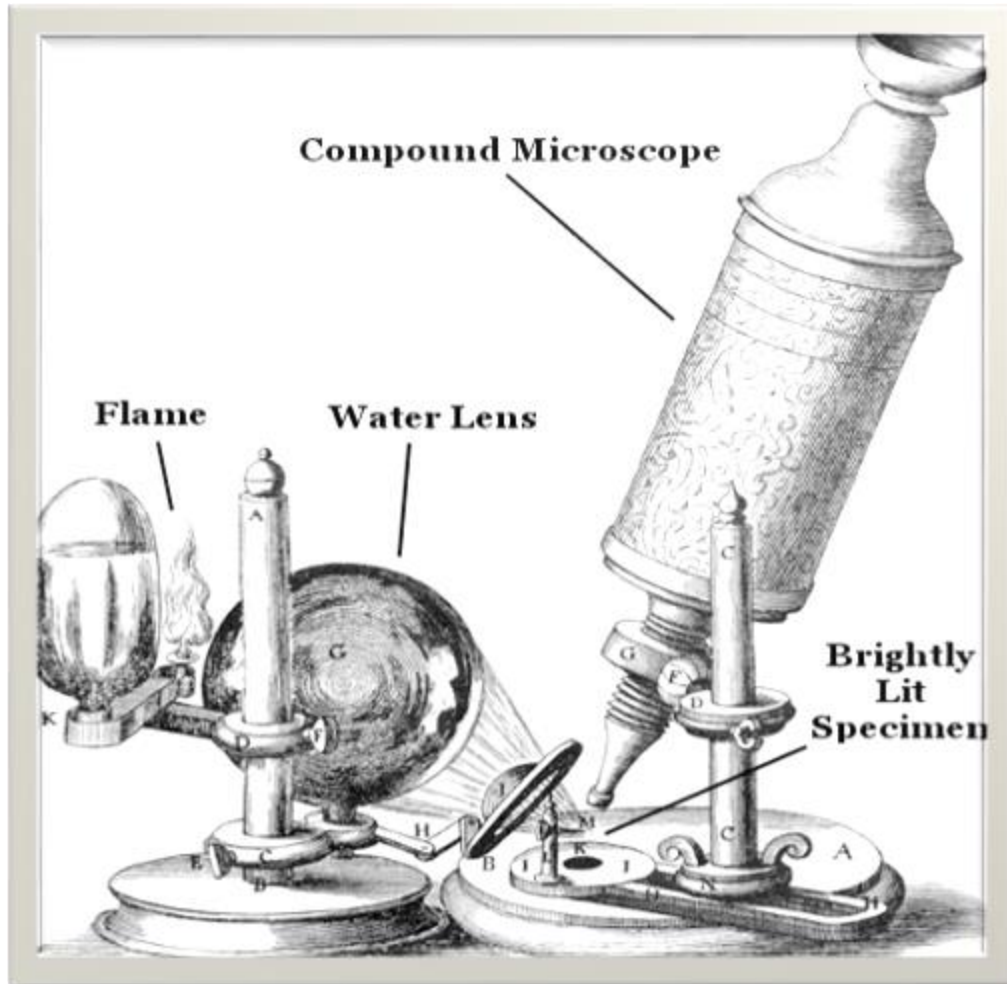
Early **Bi**ophysicists

Robert Hooke
(1635 - 1703)



- Discrete “empty” compartments in thin cork layer
- Named them “cells”
(analogous to dorm rooms in monasteries)
- No apparent connection to life
- Conclusion: spontaneous generation from ‘natural’ or ‘artificial’ heat!

Early **Bi**ophysicists



(in *Micrographia*, ca. 1665)
50x magnification

Early **Bi**ophysicists

Antonie van Leeuwenhoek
(1632 - 1723)



bacteria



Ciliophora



Foraminifera



- Teeming with movement, seemed “alive”!
- Named them “animalcules”
- Wrote series of letters to Royal Society, London
- Hooke asked to verify his observations
- Eventually sets **Cell Theory** in motion

Early **Bi**ophysicists



Simple microscope (single lens)
300X magnification

bacteria



Ciliophora



Foraminifera



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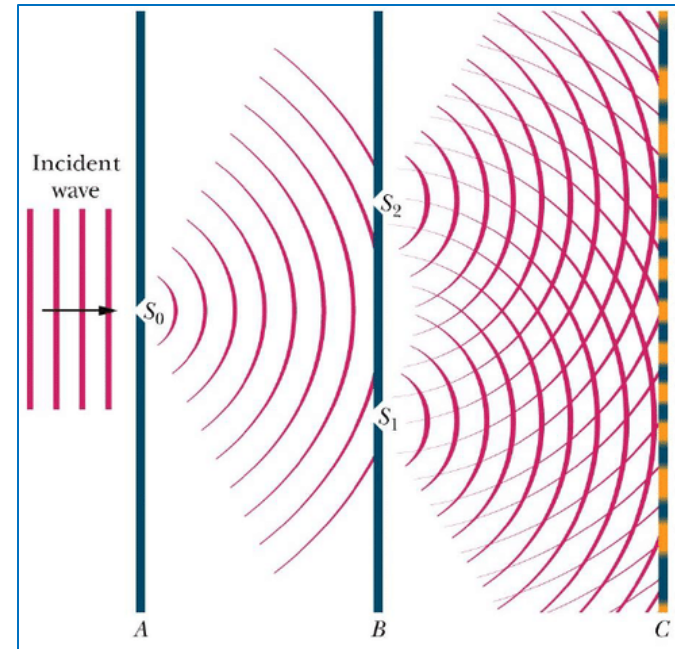
Early **Bio**physicists



Primary interest was in **sense perception**:

Discovered how the lens of the eye changes shape to focus on objects at differing distances.

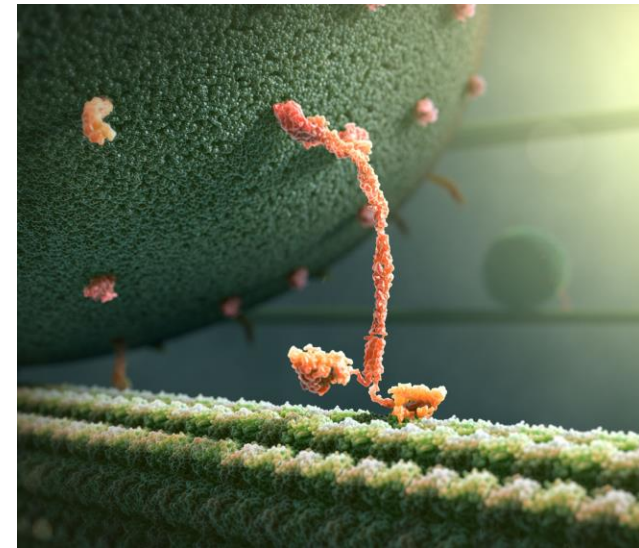
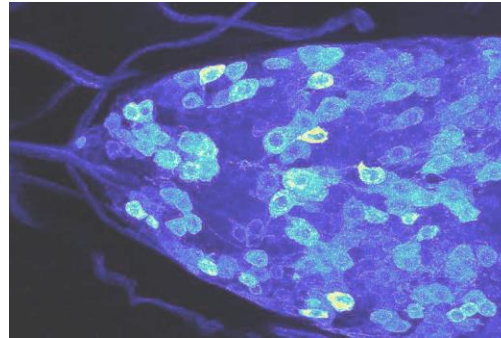
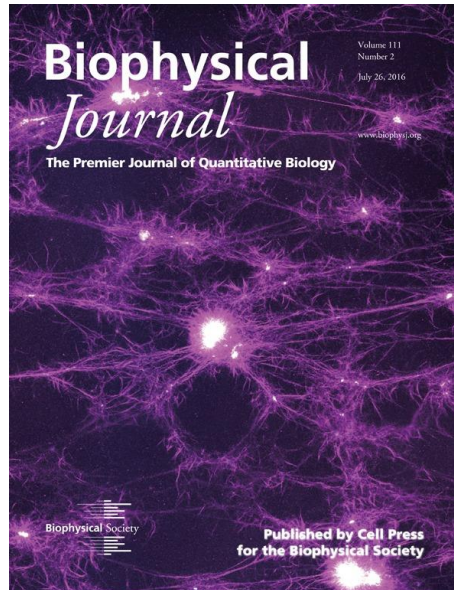
Double-slit experiment: Wave nature of light



Thomas Young (1773 - 1829)
physician and **physicist**

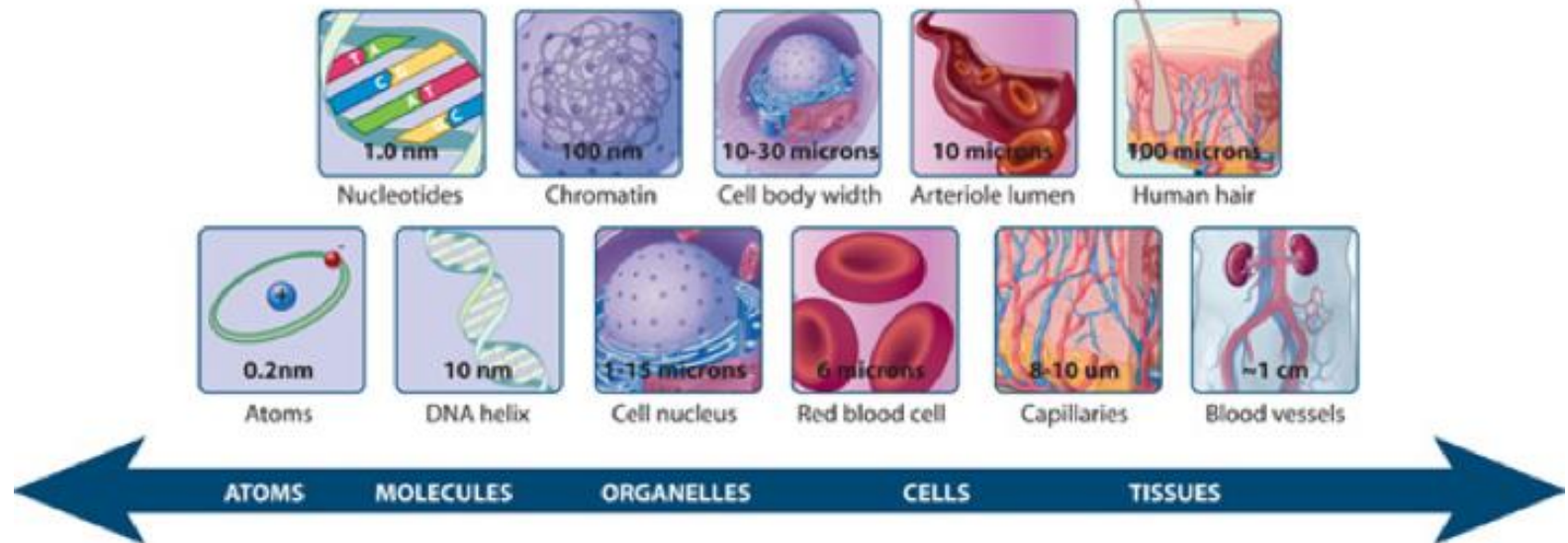
2016, volume 111(2)

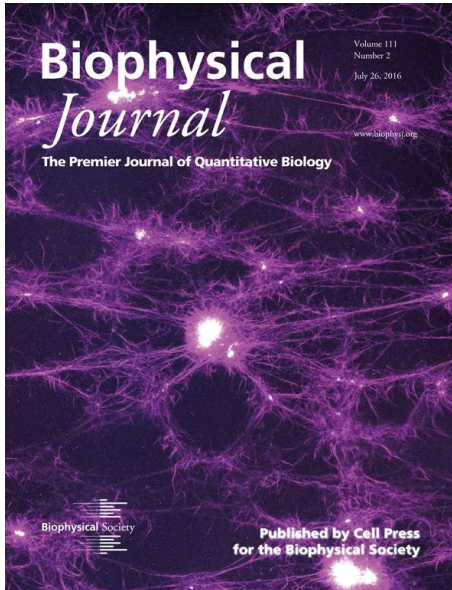
Drosophila-neuron image



Biophysics

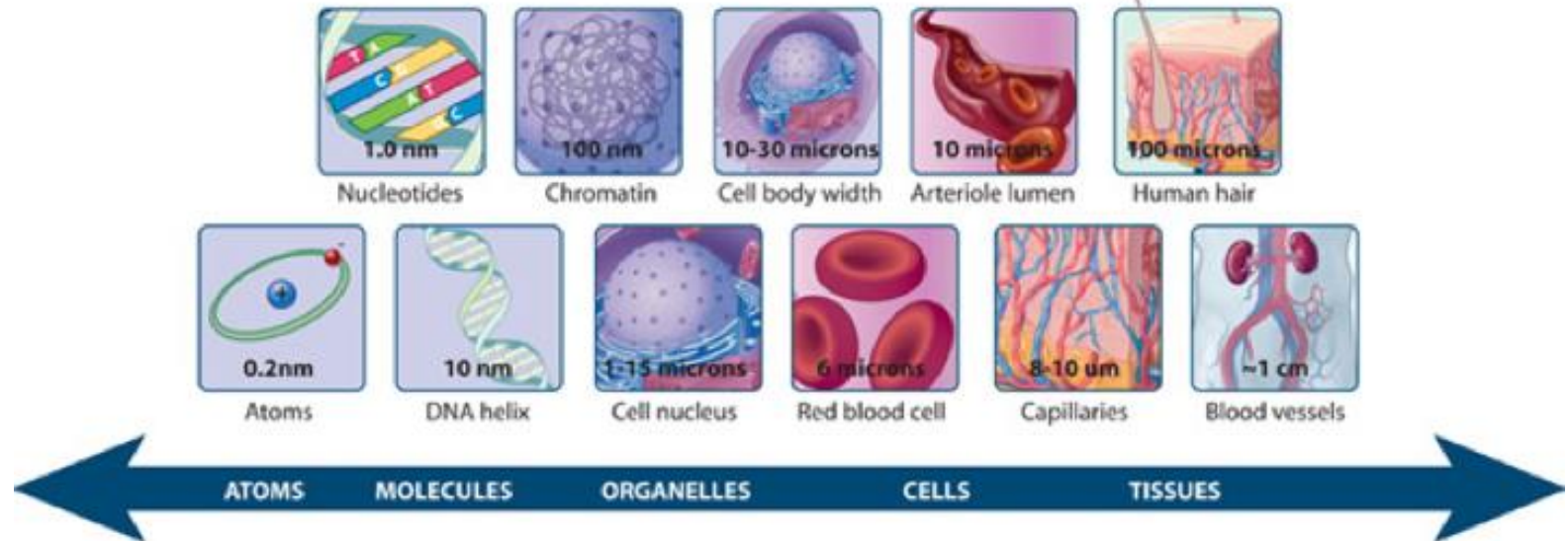
covers biology on wide *length* and *time* scales





- Contractile cytoskeletal network spontaneously formed in mixture of microtubule (*magenta*) and kinesin-5 (*cyan*).
- The network is composed of kinesin-accumulated nodes and bundled microtubule links.

covers biology on wide *length* and *time* scales



Bio**physics**

Energy processes underlie biology

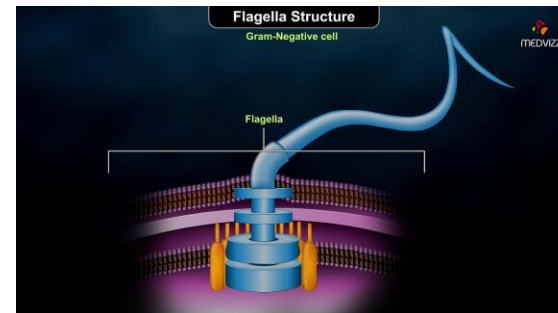
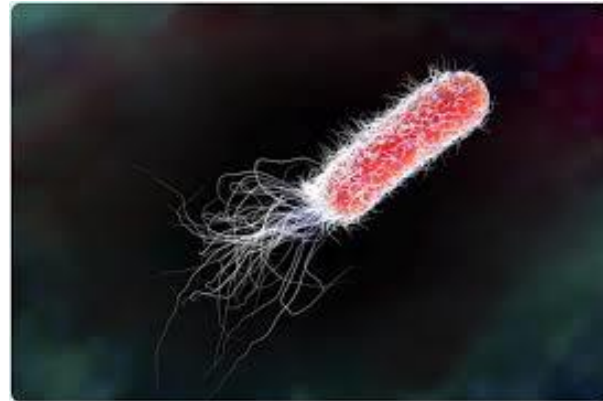


Life forms must continually transduce energy



Bio ↔ physics

Flow can be turbulent or laminar (in water) depending on the magnitude of forces



Inner structure converts chemical energy to mechanical energy

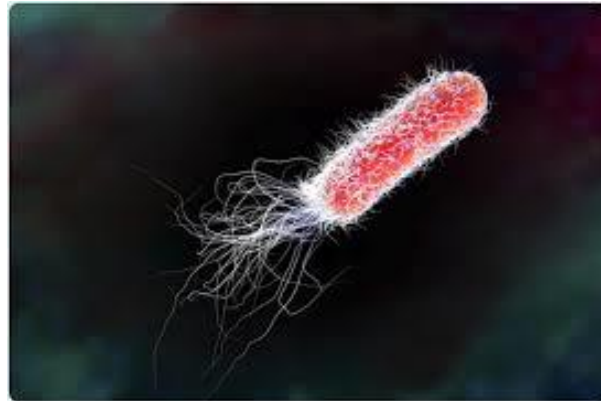
Biophysics

What are the scales of energy involved?

Force \sim Newtons



Force \sim picoNewton - nanoNewton



Biophysics

Neutrophil chasing bacteria in the midst of red blood cells (ca. 1950)

<https://www.youtube.com/watch?v=5yimbhkTqJo>



Can one make a (rough) estimate of the order of magnitudes of forces and energy?

Molecular Basis of Heredity

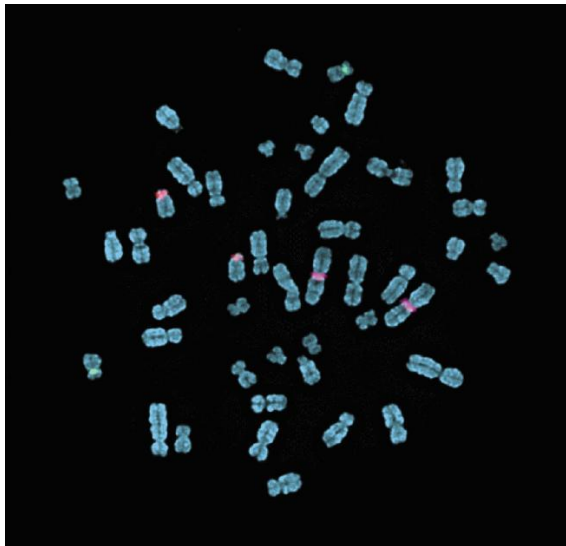
But what did DNA look like?

“**Genetics**” was coined in **1905** (by William Bateson), followed by the term “**gene**” in **1909** (by Wilhelm Johanssen)

1911:

Thomas Morgan, Alfred Sturtevant

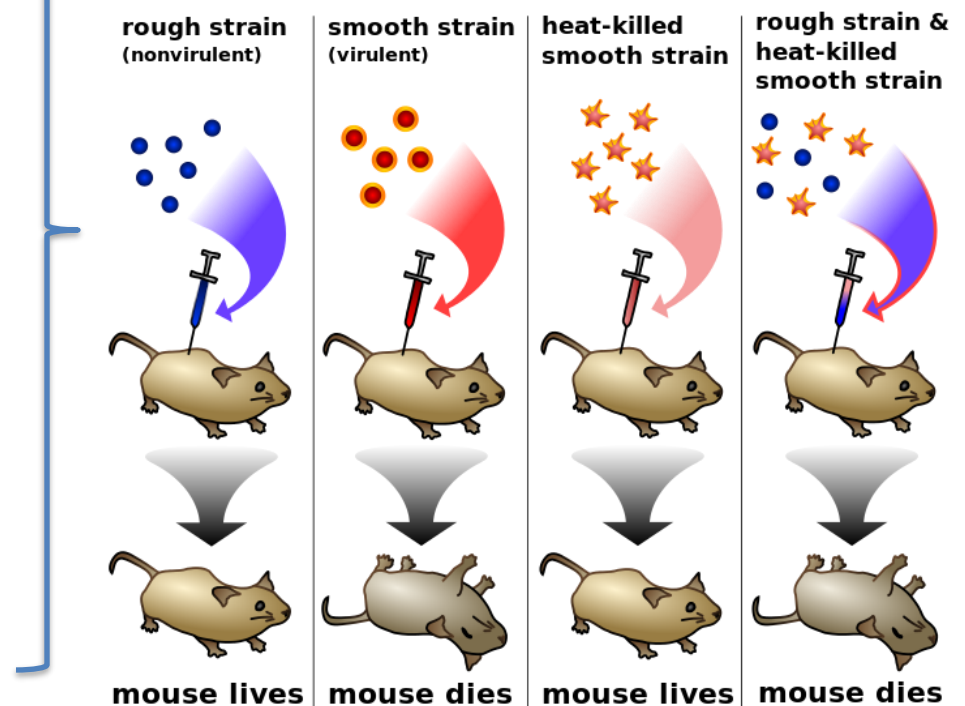
“Genes are arranged on **chromosomes**”
= protein + nucleic acids



1928:

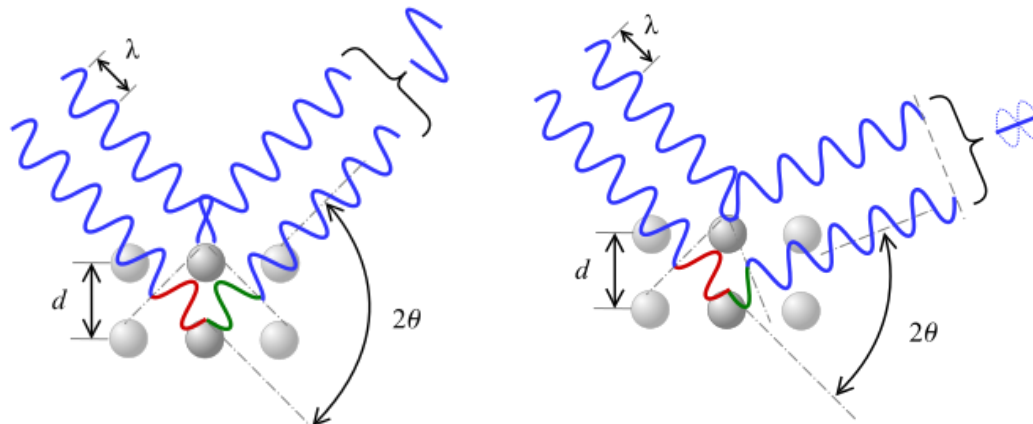
Frederick Griffith:

DNA could activate pneumococcus bacteria



Molecular Basis of Heredity

The double helix structure



Bragg's Law:

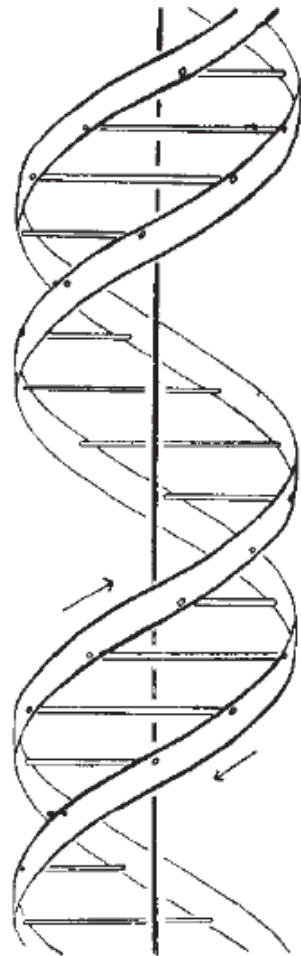
$$2d (\sin\theta) = n\lambda$$



Rosalind Franklin
(1920 - 1958)

Molecular Basis of Heredity

The double helix structure



This figure is purely diagrammatic. The two ribbons symbolize the two phosphate-sugar chains, and the horizontal rods the pairs of bases holding the chains together. The vertical line marks the fibre axis

No. 4356 April 25, 1953

NATURE

equipment, and to Dr. G. E. R. Deacon and the captain and officers of R.R.S. *Discovery II* for their part in making the observations.

¹ Young, F. B., Gerrard, H., and Jevons, W., *Phil. Mag.*, **40**, 149 (1920).

² Longuet-Higgins, M. S., *Mon. Not. Roy. Astro. Soc., Geophys. Supp.*, **5**, 285 (1949).

³ Von Arx, W. S., *Woods Hole Papers in Phys. Oceanog. Meteor.*, **11** (3) (1950).

⁴ Ekman, V. W., *Arkiv. Mat. Astron. Fysik. (Stockholm)*, **2** (11) (1905).

MOLECULAR STRUCTURE OF NUCLEIC ACIDS

A Structure for Deoxyribose Nucleic Acid

WE wish to suggest a structure for the salt of deoxyribose nucleic acid (D.N.A.). This structure has novel features which are of considerable biological interest.

A structure for nucleic acid has already been proposed by Pauling and Corey¹. They kindly made

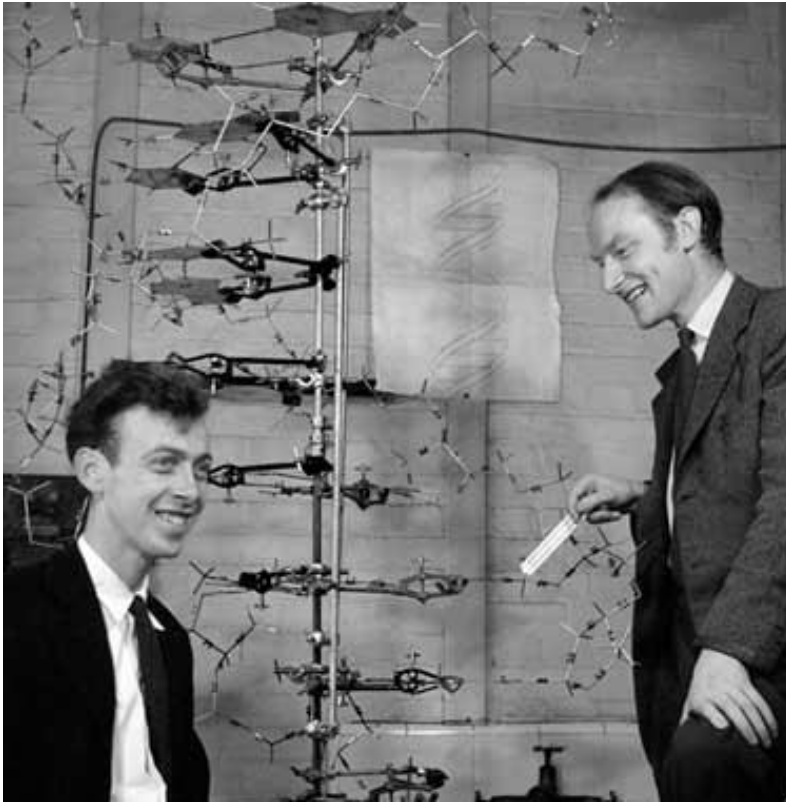
is a residue
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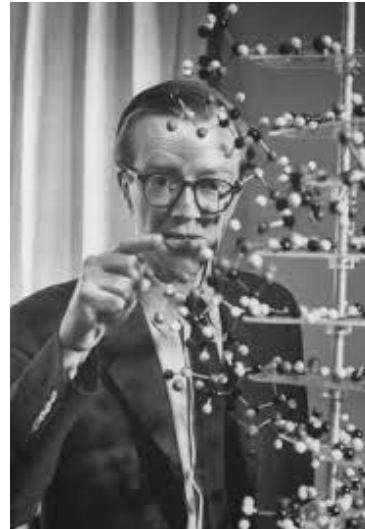
Molecular Basis of Heredity

The double helix structure



James Watson
(1928-)

Francis Crick
(1916–2004)



Maurice Wilkins
(1916 - 2004)



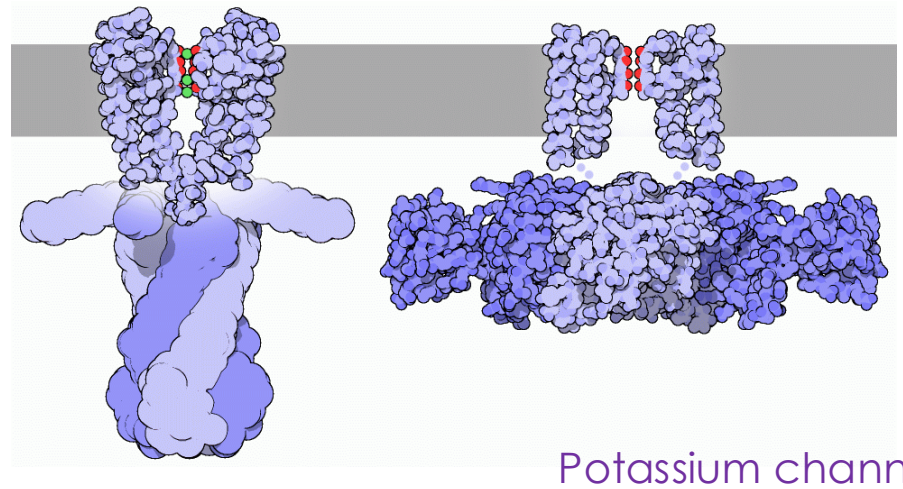
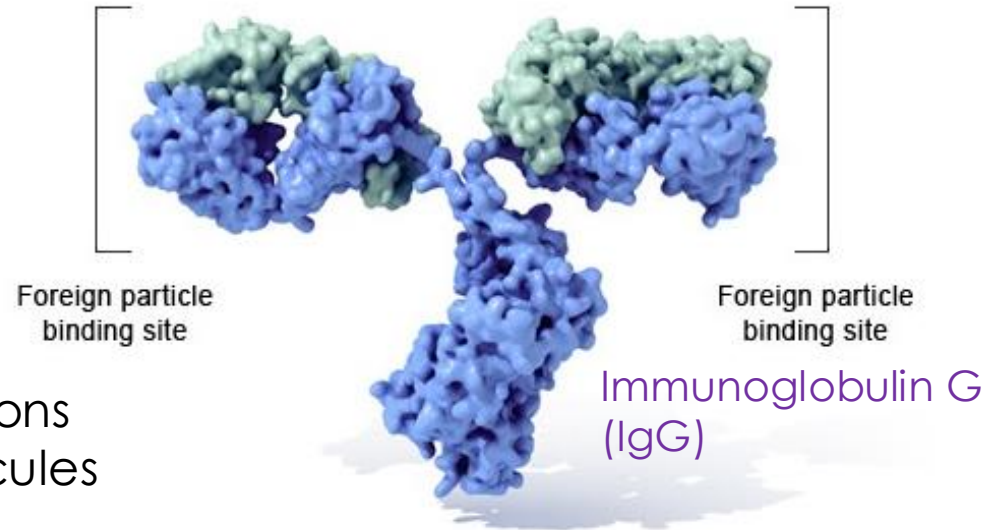
Rosalind Franklin
(1920 - 1958)

DNA as units of genetic information

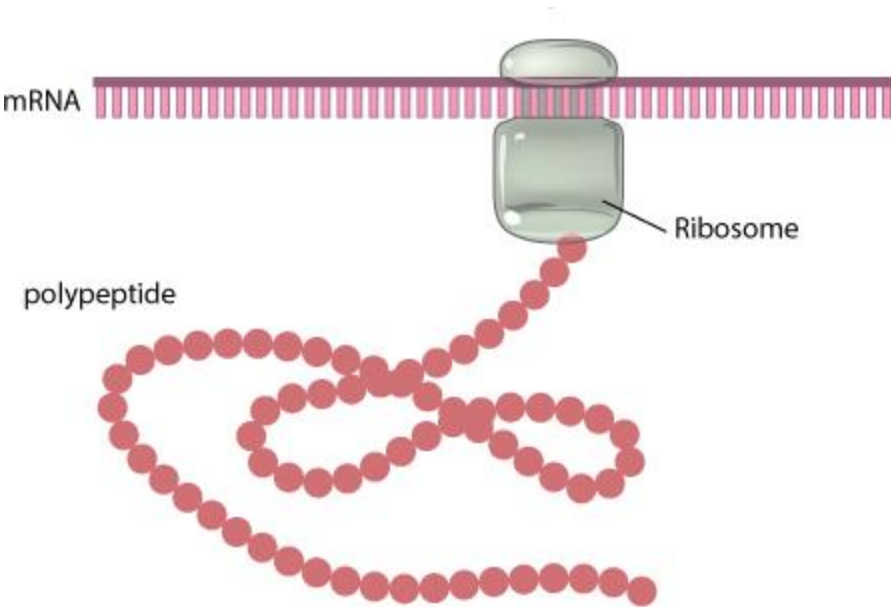
DNA encode PROTEINS, which are the “workhorses of life”, ie. they are the fundamental biological machinery

Roles include:

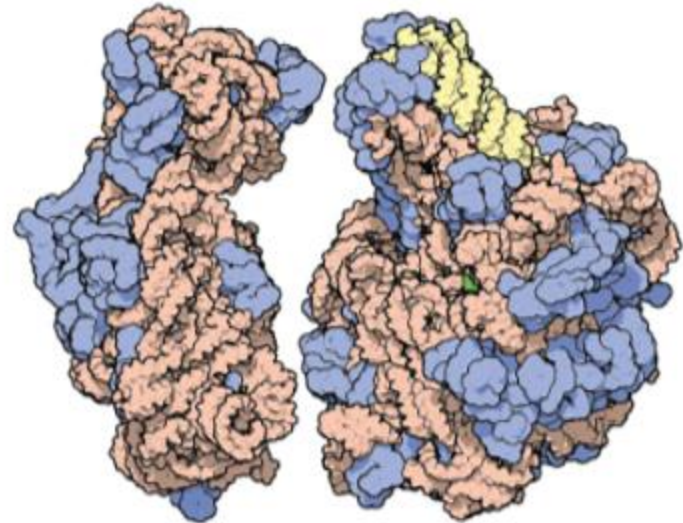
- Antibodies
- Enzymes
 - catalysts for chemical reactions
 - manufacturers of new molecules
- Messengers (eg. hormones)
- Structural units
- Channels and Transporters



How are proteins synthesized in the ribosome?



RIBOSOME the factory of protein synthesis



Ribosomes are composed of two subunits: a large subunit (PDB ID 1ffk), shown on the right, and a small subunit (PDB ID 1ffa), shown on the left. Of course, the term "small" is used in a relative sense here: both the large and the small subunits are huge compared to a typical protein. Both subunits are composed of long strands of RNA, shown here in orange and yellow, dotted with protein chains, shown in blue. When synthesizing a new protein, the two subunits lock together with a messenger RNA trapped in the space between. The ribosome then walks down the messenger RNA three nucleotides at a time, building a new protein piece-by-piece.

Image from the RCSB PDB's *Molecule of the Month* feature (www.pdb.org)



RESEARCH COLLABORATORY FOR STRUCTURAL BIOINFORMATICS
Belgium, The State University of New Jersey
San Diego Supercomputer Center & Skaggs School of Pharmacy &
Pharmaceutical Sciences, University of California, San Diego

Three structural biologists have won the 2009 Nobel Prize in Chemistry for studies of the structure and function of the ribosome: Venkatesan Ramakrishnan (MRC Laboratory of Molecular Biology), Thomas A. Steitz (Yale University), and Ada E. Yonath (Weizmann Institute of Science). The depositions of their first complete ribosome subunit structures (1ffg, 1ffk, and 1ffa) almost a decade ago ushered structural biology into a new era. Since that time, more than 120 ribosome structures consisting of 50S, 30S subunits and complete 70S ribosomes have been contributed by three Nobel scientists. The structures, complexed with and without antibiotics, tRNAs, and RNAs, initiation factors, and release factors, provide a basis for understanding how the ribosome works and are useful tools for drug development.

Ribosomal Structure

<https://www.nobelprize.org/prizes/chemistry/2009/illustrated-information/>



The Nobel Prize in Chemistry 2009

Venkatraman Ramakrishnan, Thomas A. Steitz, Ada E. Yonath

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The Nobel Prize in Chemistry 2009



Photo: U. Montan
Venkatraman Ramakrishnan
Prize share: 1/3



Photo: U. Montan
Thomas A. Steitz
Prize share: 1/3

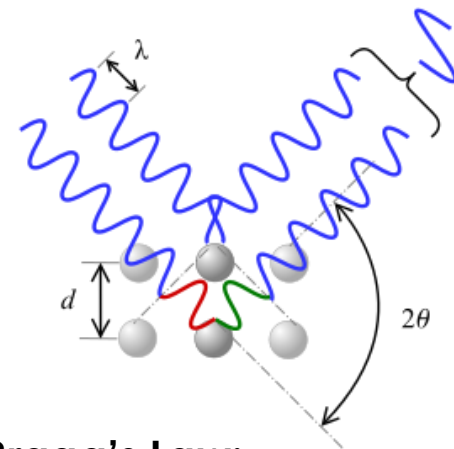


Photo: U. Montan
Ada E. Yonath
Prize share: 1/3

The Nobel Prize in Chemistry 2009 was awarded jointly to Venkatraman Ramakrishnan, Thomas A. Steitz and Ada E. Yonath *"for studies of the structure and function of the ribosome"*.

Photos: Copyright © The Nobel Foundation

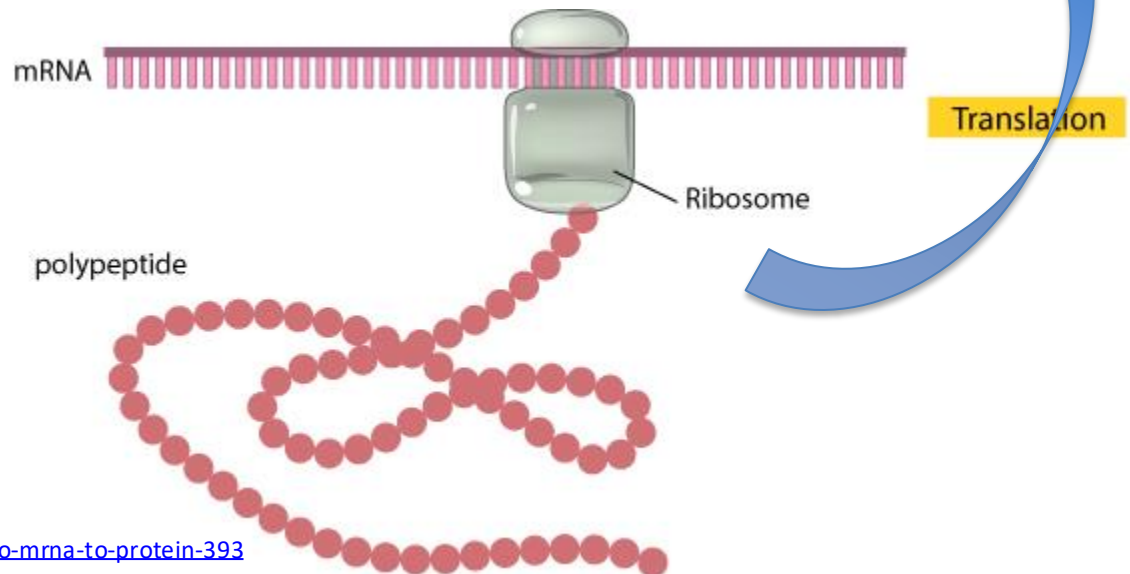
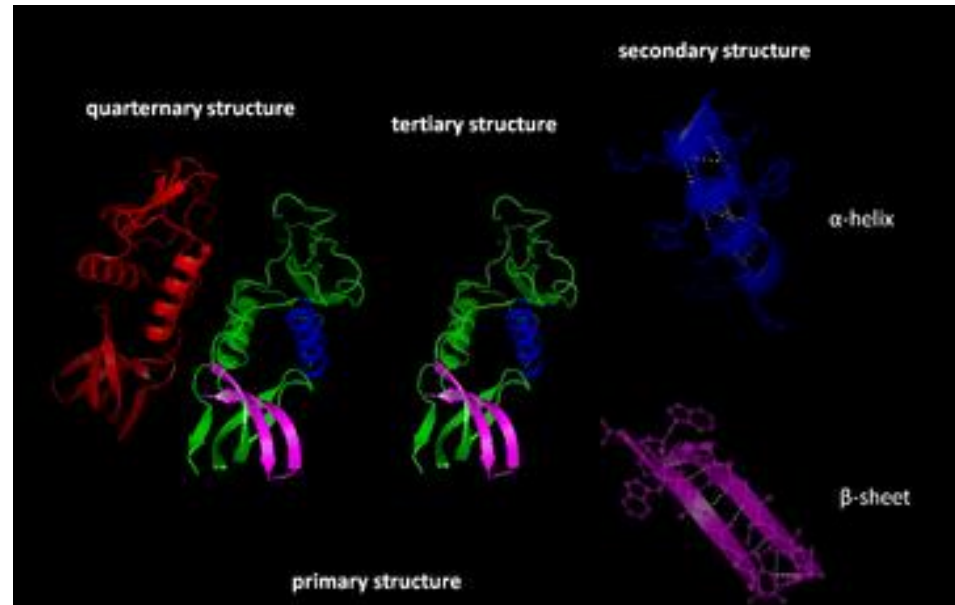
Their method was X-ray crystallography



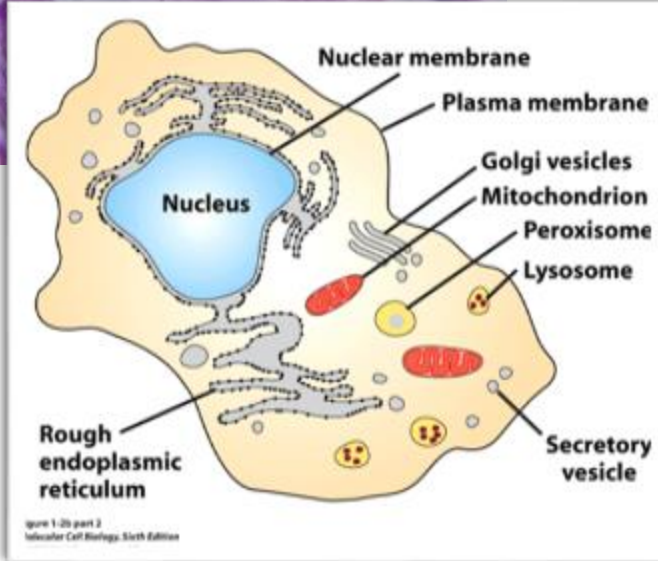
Bragg's Law:
$$2d (\sin \theta) = n\lambda$$

Nascently synthesized protein chain usually FOLDS into a functional form

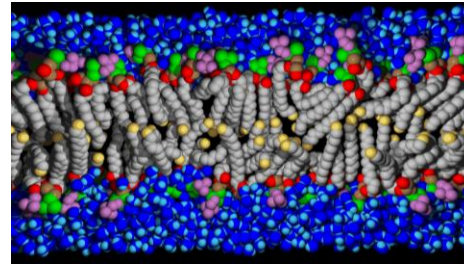
- Folding may be **spontaneous**
- Or it may be **assisted by** other proteins (**chaperones**)
- The environment (solvent, heat, viscosity, etc) strongly influences the folding *path* and stability of the *final folded structure*
- The **physics of protein folding is incompletely understood**



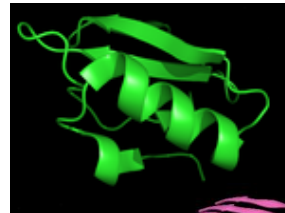
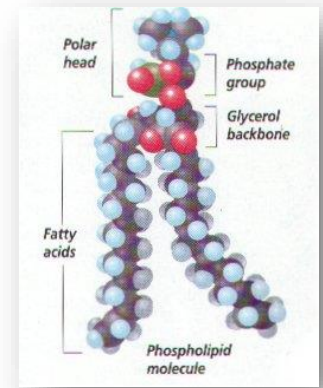
Biological Organization



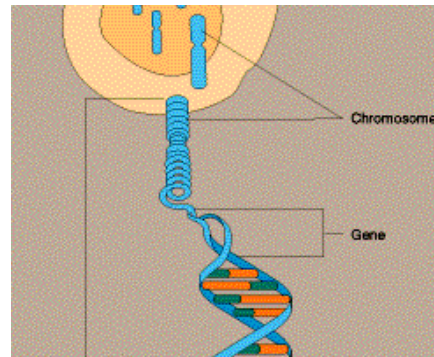
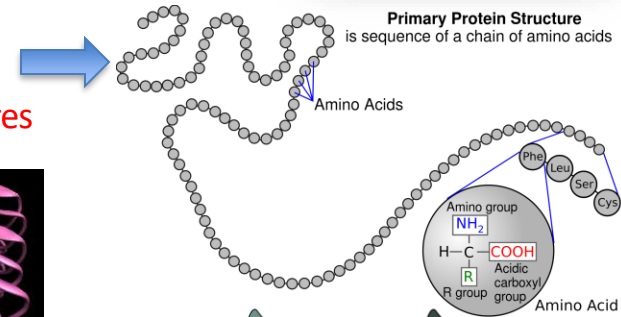
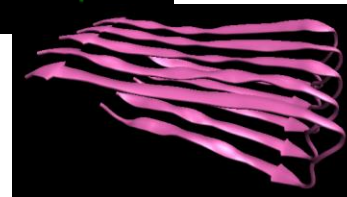
Molecular Assembly



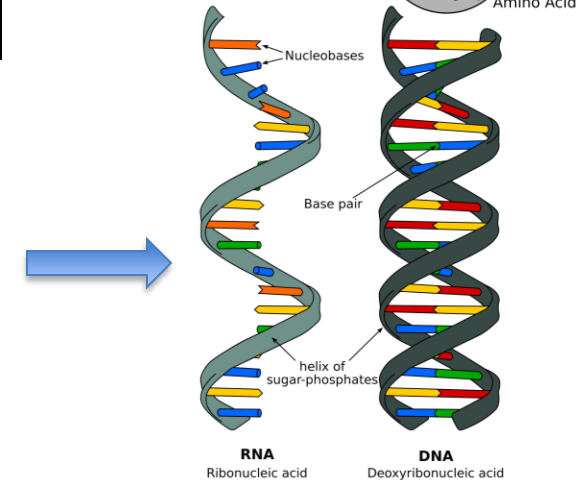
cell membranes



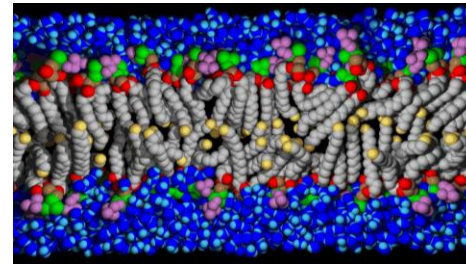
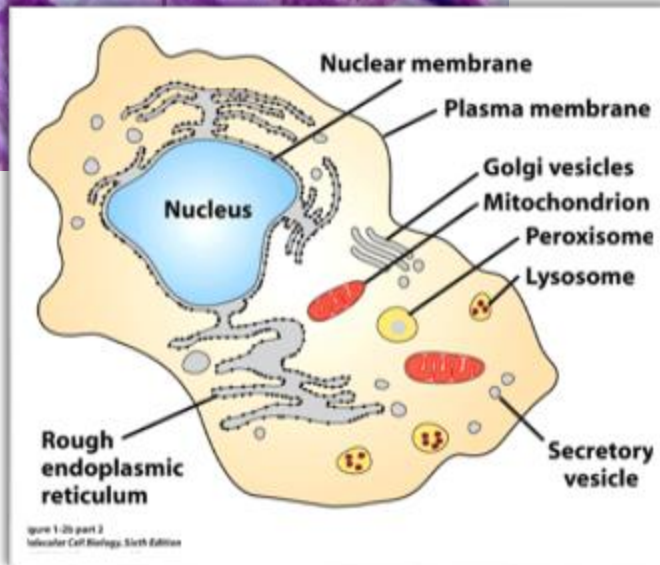
protein structures



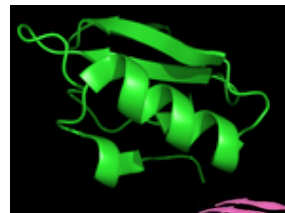
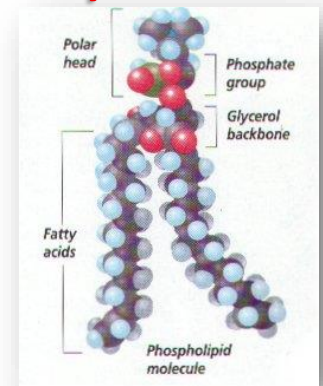
genetic material



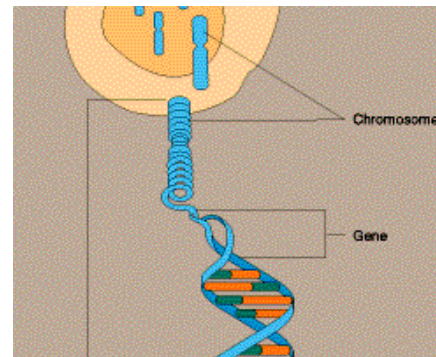
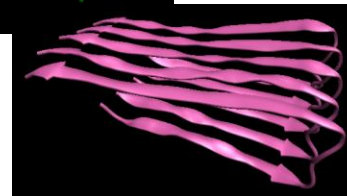
What Physical Principles Drive Molecular Assembly?



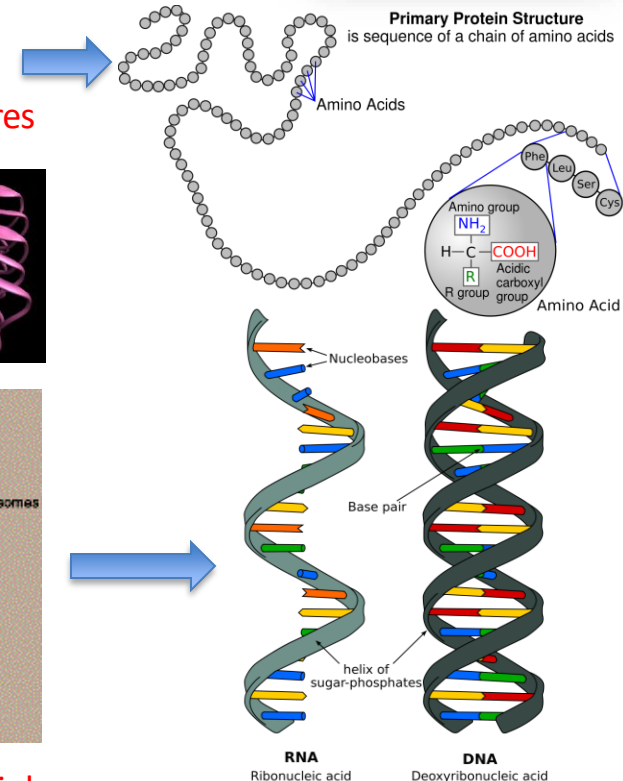
cell membranes



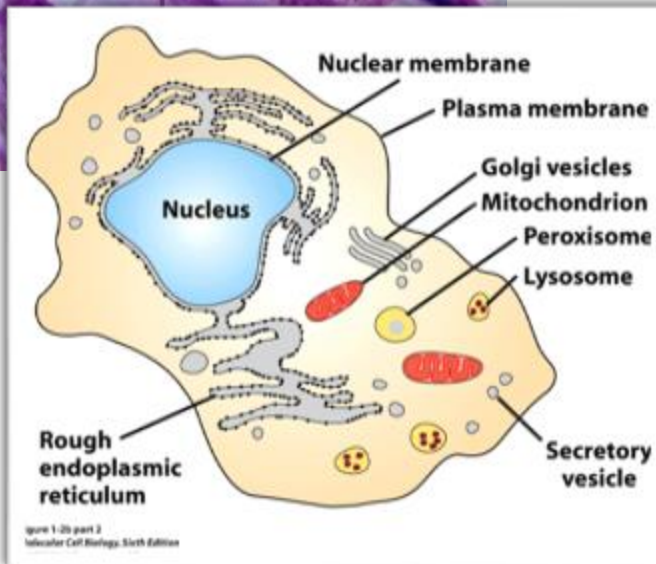
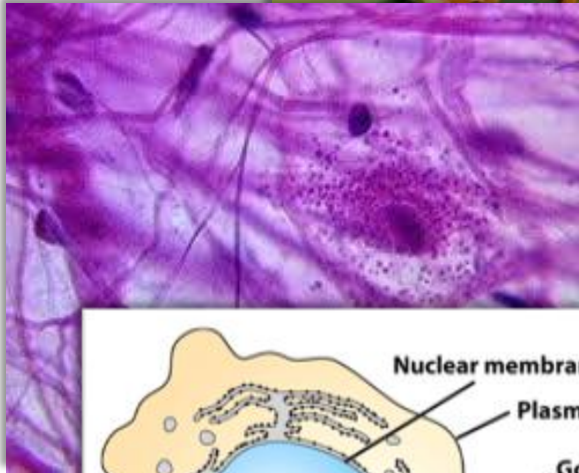
protein structures



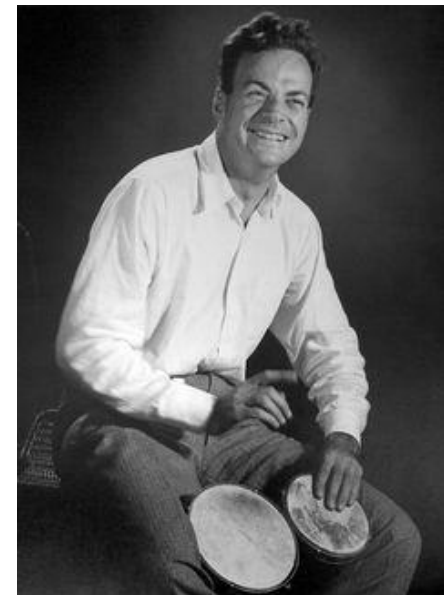
genetic material



What Physical Principles Drive Molecular Assembly?

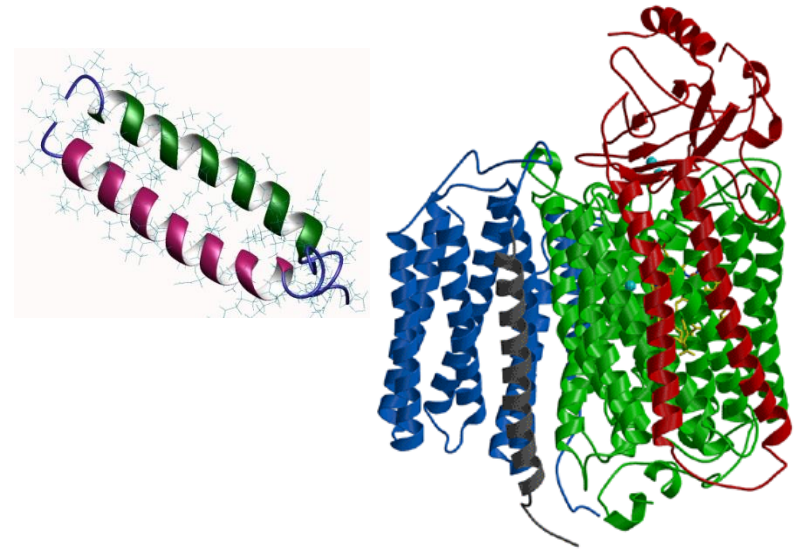


"Everything that *living things* do can be understood in terms of the *jiggling and wiggling of atoms.*"



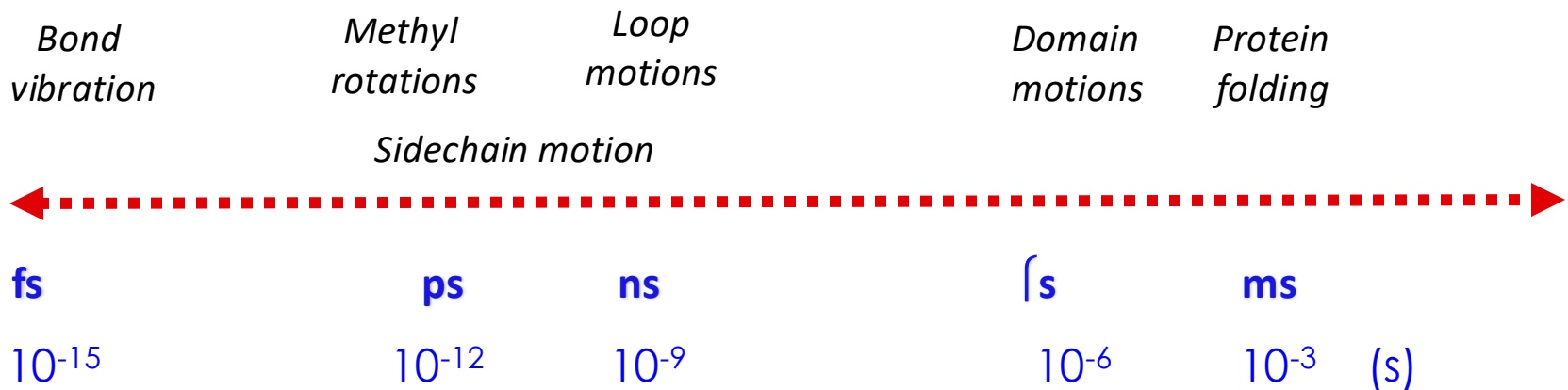
-- Richard Feynman
(1963, *Lectures in Physics*)

Biomolecules

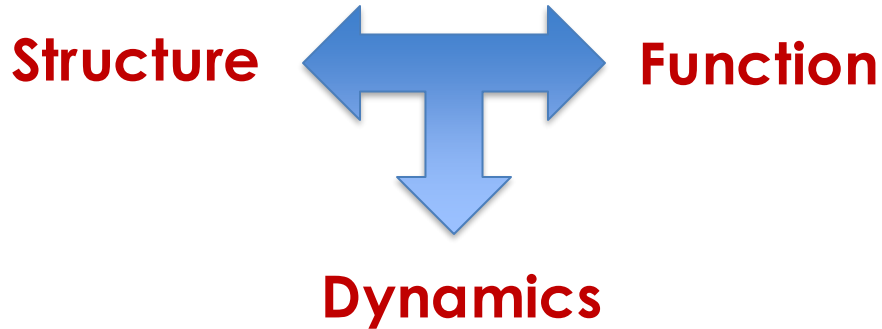


Dynamical Timescales

Henzler-Wildman & Kern, **Nature Rev.**, 2007



Biomolecules



Heat, Energy, Temperature, Work, Disorder

Dynamical Timescales

Henzler-Wildman & Kern, **Nature Rev.**, 2007

*Bond
vibration*

*Methyl
rotations*

*Loop
motions*

*Domain
motions*

*Protein
folding*

Sidechain motion



fs

10^{-15}

ps

10^{-12}

ns

10^{-9}

μs

10^{-6}

ms

10^{-3} (s)

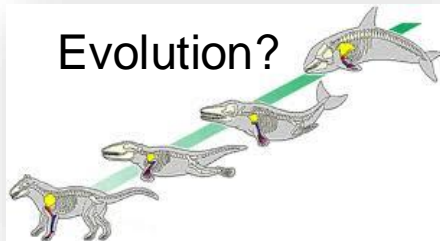


Biological Time, Length scales

1 year $\sim 3.2 \times 10^7$ s

Time Scales (s)

10^9
 10^8
 10^7
 10^5
 10^{-3}
 10^{-6}
 10^{-12}
 10^{-15}



atom,
chem. bond

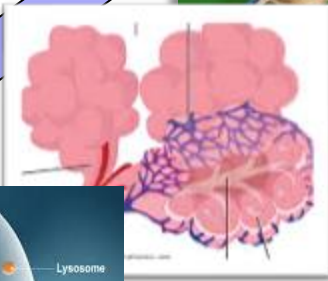
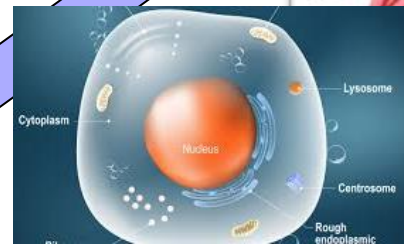
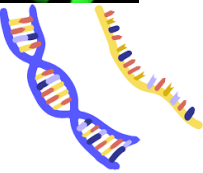
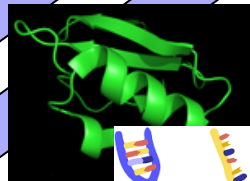
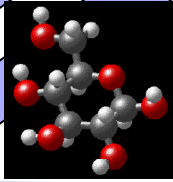
(Bio)
molecule

Cell
Organelle

Organ

Organism

Ecosystem



10^{-11}

10^{-9}

10^{-7}

10^{-5}

10^1

10^3

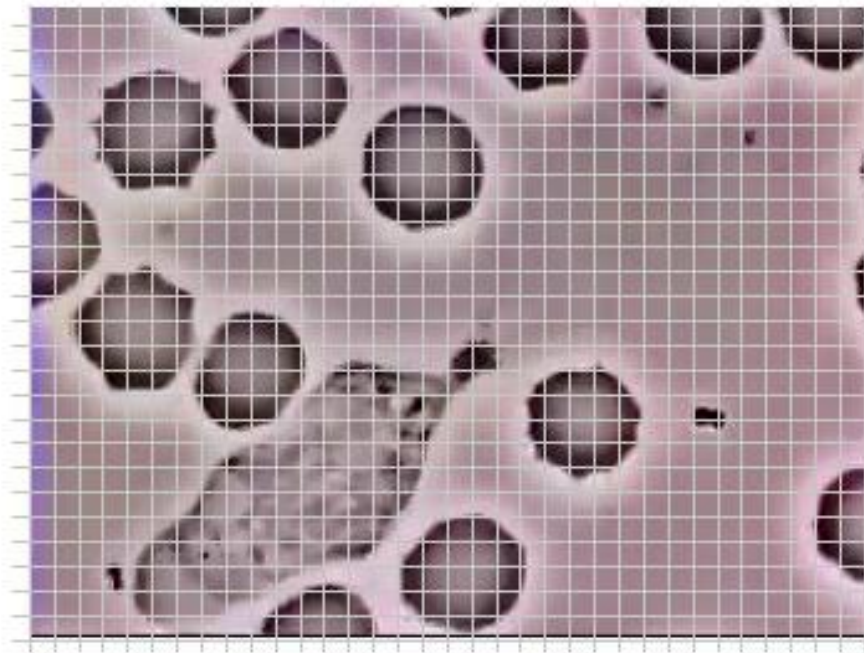
10^5

Length Scales (m)

Length, Time and Associated Scales

Neutrophil chasing bacteria in the midst of red blood cells (ca. 1950)

<https://www.youtube.com/watch?v=5yimbhkTqJo>



Red blood cell $\sim 8 \mu\text{m}$

What are the approx. sizes of:
a) the neutrophil, b) the bacteria it is chasing?

'Rule of Thumb': Length Scales

Table 1.1: Rules of thumb for biological estimates.

Quantity of interest	Symbol	Rule of thumb
<i>E. coli</i>		
Cell volume	$V_{E. coli}$	$\approx 1 \mu\text{m}^3$
Cell mass	$m_{E. coli}$	$\approx 1 \text{ pg}$
Cell cycle time	$t_{E. coli}$	$\approx 3000 \text{ s}$
Cell surface area	$A_{E. coli}$	$\approx 6 \mu\text{m}^2$
Macromolecule concentration in cytoplasm	$c_{E. coli}^{\text{macromol}}$	$\approx 300 \text{ mg/mL}$
Genome length	$N_{bp}^{E. coli}$	$\approx 5 \times 10^6 \text{ bp}$
Swimming speed	$v_{E. coli}$	$\approx 20 \mu\text{m/s}$
Yeast		
Volume of cell	V_{yeast}	$\approx 60 \mu\text{m}^3$
Mass of cell	m_{yeast}	$\approx 60 \text{ pg}$
Diameter of cell	d_{yeast}	$\approx 5 \mu\text{m}$
Cell cycle time	t_{yeast}	$\approx 200 \text{ min}$
Genome length	N_{bp}^{yeast}	$\approx 10^7 \text{ bp}$
Organelles		
Diameter of nucleus	d_{nucleus}	$\approx 5 \mu\text{m}$
Length of mitochondrion	l_{mito}	$\approx 2 \mu\text{m}$
Diameter of transport vesicles	d_{vesicle}	$\approx 50 \text{ nm}$

'Rule of Thumb': Length Scales

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Cell volume	$V_{E. coli}$	$\approx 1 \mu\text{m}^3$
Cell mass	$m_{E. coli}$	$\approx 1 \text{ pg}$
Cell cycle time	$t_{E. coli}$	$\approx 3000 \text{ s}$
Cell surface area	$A_{E. coli}$	$\approx 6 \mu\text{m}^2$
Macromolecule concentration in cytoplasm	$c_{E. coli}^{\text{macromol}}$	$\approx 300 \text{ mg/mL}$
Genome length	$N_{bp}^{E. coli}$	$\approx 5 \times 10^6 \text{ bp}$
Swimming speed	$v_{E. coli}$	$\approx 20 \mu\text{m/s}$

- I. What is the “surface-to-volume” ratio of *E. coli*?
 - Unit?
 - How is this expected to scale with size of organism?

'Rule of Thumb': Length Scales

Table 1.1: Rules of thumb for biological estimates.

Quantity of interest	Symbol	Rule of thumb
<i>E. coli</i>		
Cell volume	$V_{E. coli}$	$\approx 1 \mu\text{m}^3$
Cell mass	$m_{E. coli}$	$\approx 1 \text{ pg}$
Cell cycle time	$t_{E. coli}$	$\approx 3000 \text{ s}$
Cell surface area	$A_{E. coli}$	$\approx 6 \mu\text{m}^2$
Macromolecule concentration in cytoplasm	$c_{E. coli}^{\text{macromol}}$	$\approx 300 \text{ mg/mL}$
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Swimming speed	$v_{E. coli}$	$\approx 20 \mu\text{m/s}$

- III. Approx. how many times it's 'body size' does *E. coli* cover per second?
- How would you compare this movement with that of "very fast" humans?
(Typical Olympic Runner speed $\sim 30 \text{ km/h}$)