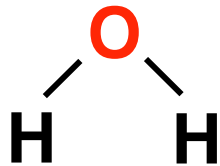


WATER, MACROMOLECULES

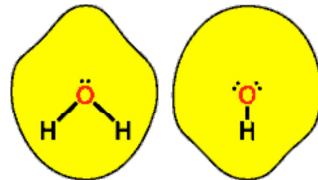
MIKLÓS KELLERMAYER

Structure of the water molecule

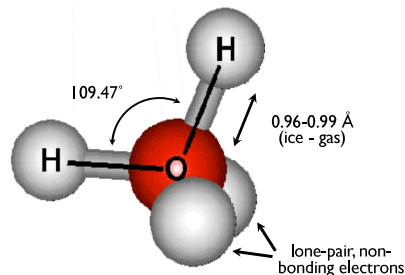
One of the smallest molecules
Barely larger than an atom



van der Waals radius: $\sim 3.2 \text{ \AA}$
Its shape is not spherical



Tetrahedral structure:



Water

- Source of *inspiration* (music, paintings).
- Thales (580, B.C.): "...water is source of all things..."
- Henry Cavendish (1783): water is H_2O .
- Only chemical that naturally exists in *all three states* (solid, liquid, gas).
- 71% of the Earth's surface is covered with water ("blue planet").
- Water is of utmost importance for life:
98% of jellyfish
94% of three-month human fetus
72% of newborn
60% of adult
- Average daily water intake: 2.4 liters.



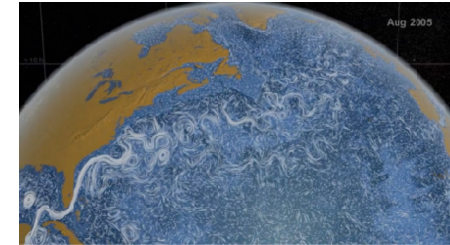
Georg Friedrich Händel (1685-1759): "**Water music**".



Georg Friedrich Händel (center) and King George I (right) on the Thames River, 17 July 1717.



Hokusai (1760-1849): Great wave off Kanagawa

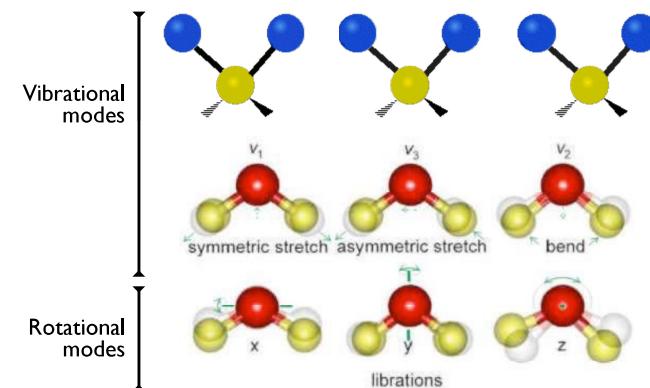


Perpetual motion of oceans on Earth's surface.



Dynamics of the water molecule

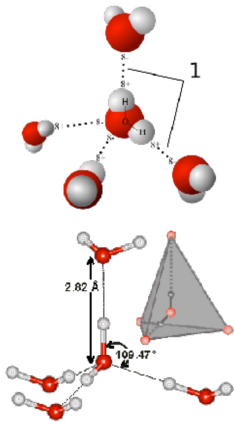
Rotational and vibrational motion



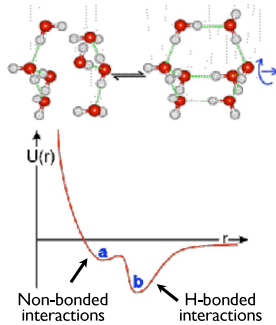
Absorption in the infrared and red spectral region →
"blue" color of natural waters: *blue planet*

Structure of liquid water

Hydrogen bonds in the vicinity of a water molecule: formation of the water pentamer

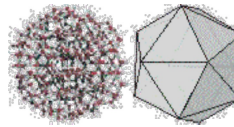


H-bridge: cohesion + repulsion
Cluster formation: bicyclo-octamer



From clusters to networks:
280 molecules form icosahedral structure

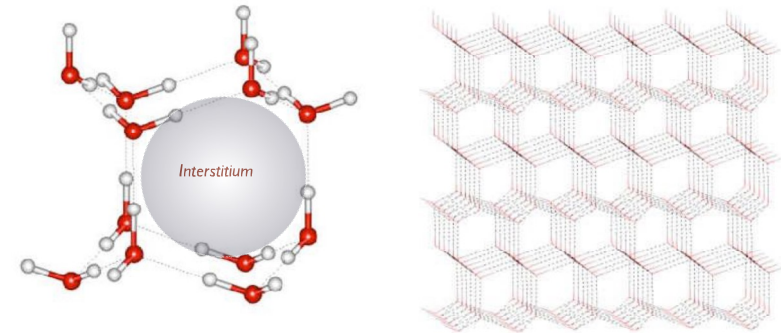
(icosahedron: regular polyhedron with 20 identical equilateral triangular faces)



Spatial networks:
May explain anomalous properties of water

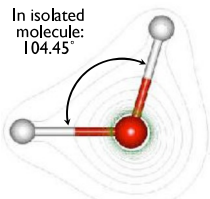
Structure of ice

- 9 different forms
- Conventional ice: hexagonal structure
- Coordination number: 4 (each molecule coordinates another four)
- Interstitium: could incorporate a water molecule - important in the diffusion of gases



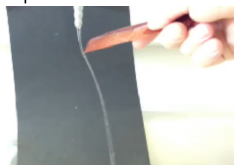
Physical properties of water I.

Large permanent dipole moment → Good solvent



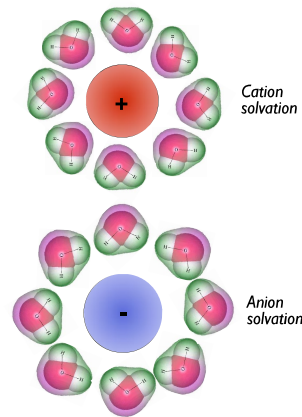
Chemical	Dipole moment
Polyethylene	2.25
Methanol	30
Ethylene glycol	37
Glycerol	47
Water	80
Titanium dioxide	86-173

Water stream bends in response to Coulombic forces



Dipole moment: amount of electrical energy stored in the material by an applied voltage, relative to vacuum. It shows how good an electrical insulator the material is. Consequence: water is good solvent.

Courtesy of Prof. Miklós Zrínyi

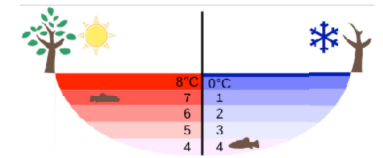
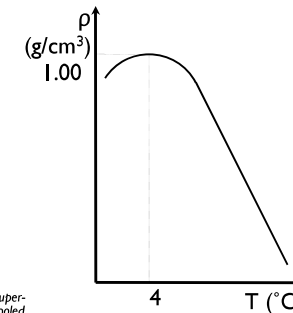


In the microwave oven: dipoles rotate according to the oscillating electromagnetic field. Water molecules acquire kinetic energy, which dissipates into the surroundings.

Physical properties of water II.

Anomalous density-temperature function

Temperature (°C)	Density (kg/m³)
+100	958.4
+80	971.8
+60	983.2
+40	992.2
+30	995.6502
+25	997.0479
+22	997.7735
+20	998.2071
+15	999.1026
+10	999.7026
+4	999.9720
0	999.8395
-10	998.117
-20	993.547
-30	983.854



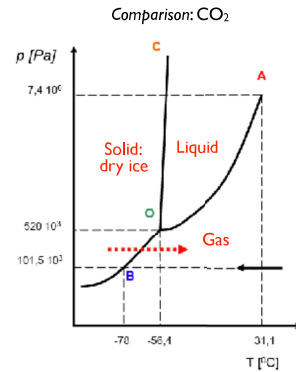
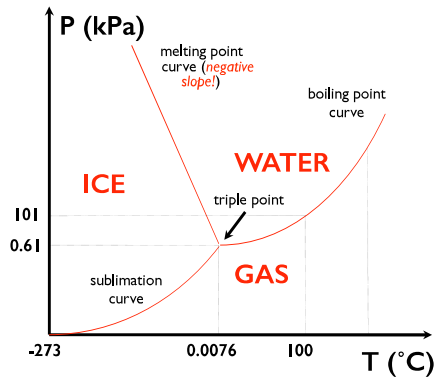
Consequences:

- 4 °C water is always at the bottom of the lake.
- Life persists under frozen lake.
- Creek runs under ice.

Physical properties of water III.

Anomalous phase diagram

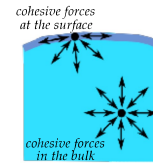
- Phase curve: two phases are in equilibrium
- Area between phase curves: a single phase is present
- Intersection of phase curves: triple point



Physical properties of water IV.

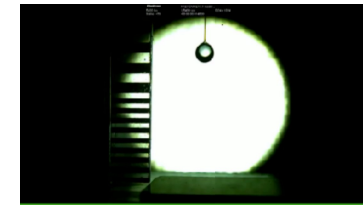
Large surface tension

Surface tension: contractive tendency of the liquid that resists external force.
Imbalance of cohesive forces in the **bulk** versus the **surface** of the liquid.



Chemical	Surface tension (mN/m)
Ethanol	24.4
Methanol	22.7
Acetone	23.7
Chloroform	27.1
Benzene	28.5
Water	72.9

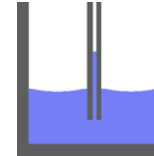
Consequences on **hydrophobic** surface



Consequences in macroscopic living systems

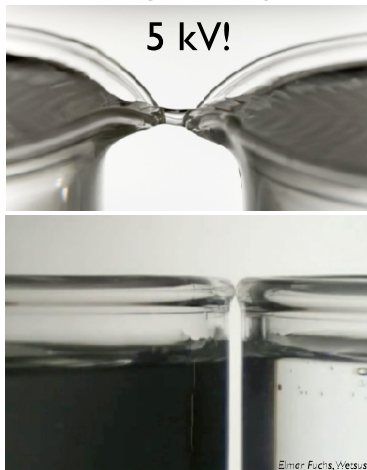


Consequences on **hydrophilic** surface

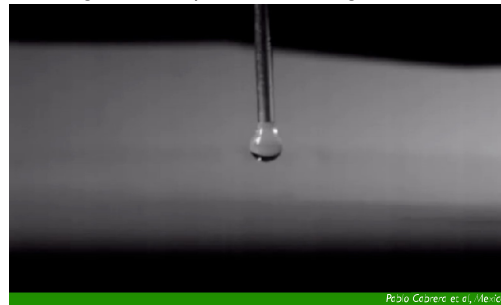


Further interesting features of water

Floatig water bridge

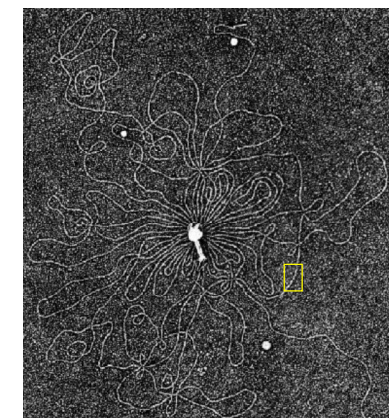
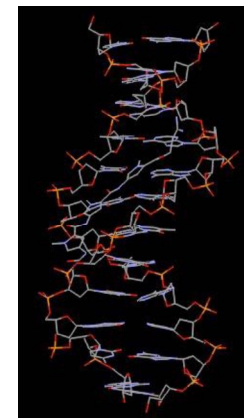


Persisting water droplets on vibrating water surface

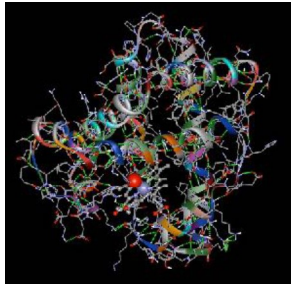


Number of water molecules in the cell	$\sim 1.6 \times 10^{14}$
Distance between water molecules	~ 0.4 nm

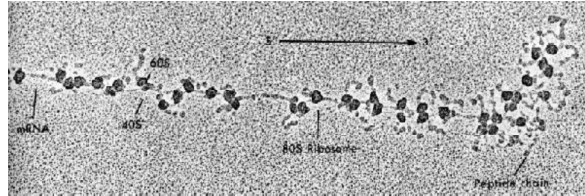
Biological macromolecules are **GIANT** molecules



Biological macromolecules are **EXCITING** molecules

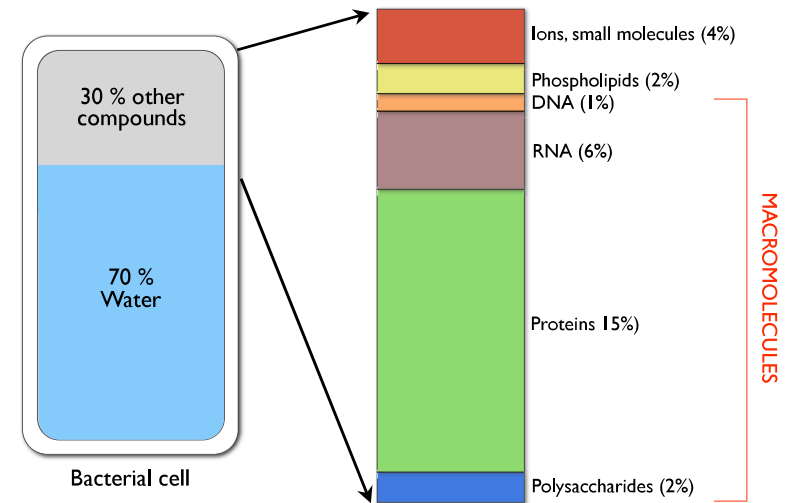


Structure of hemoglobin subunit



Newly synthesized protein (silk fibroin)

Proportion of macromolecules in the cell by mass is **LARGE**



Biological macromolecules: biopolymers

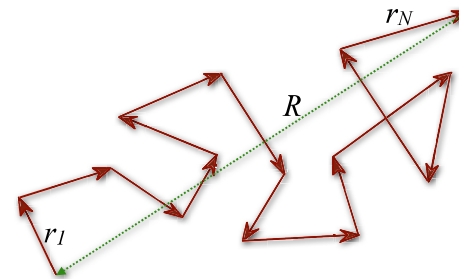
Polymers:
chains built up from monomers

Number of monomers: $N \gg 1$;
Typically, $N \sim 10^2 - 10^4$,
but, in DNA, e.g.: $N \sim 10^9 - 10^{10}$

Biopolymer	Monomer	Bond
Protein	Amino acid	Covalent (peptide bond)
Nucleic acid (RNA, DNA)	Nucleotide (CTUGA)	Covalent (phosphodiester)
Polysaccharide (e.g., glycogen)	Sugar (e.g., glucose)	Covalent (e.g., α -glycosidic)
Protein polymer (e.g., microtubule)	Protein (e.g., tubulin)	Secondary

Shape of the polymer chain resembles random walk

Brownian-movement -
“random walk”



“Square-root law”:

$$\langle R^2 \rangle = Nl^2 = Ll$$

R = end-to-end distance
 N = number of elementary vectors
 $l = |\vec{r}_i|$ = correlation length
 r_i = elementary vector
 $Nl = L$ = contour length
 l is related to **bending rigidity**.

In case of Brownian-movement R = displacement, N = number of elementary steps, L = total path length, and l = mean free path length.

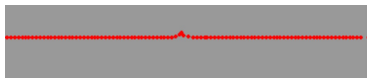
Biopolimer elasticity

Entropic (thermal) elasticity

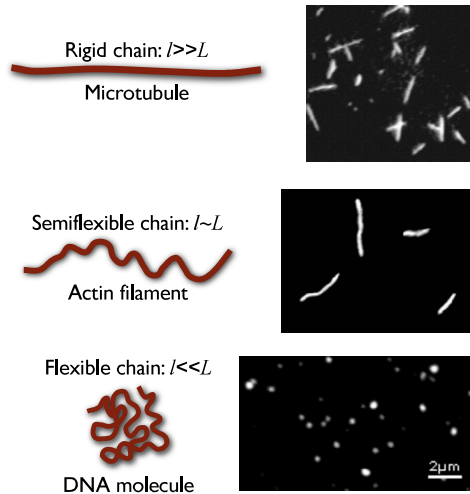
Polymer chain goes through thermal fluctuations of shape.

↓
Configurational entropy (orientational disorder of elementary vectors) increases.

↓
Due to the entropy maximization of the system the chain shortens (end-to-end distance falls below contour length).

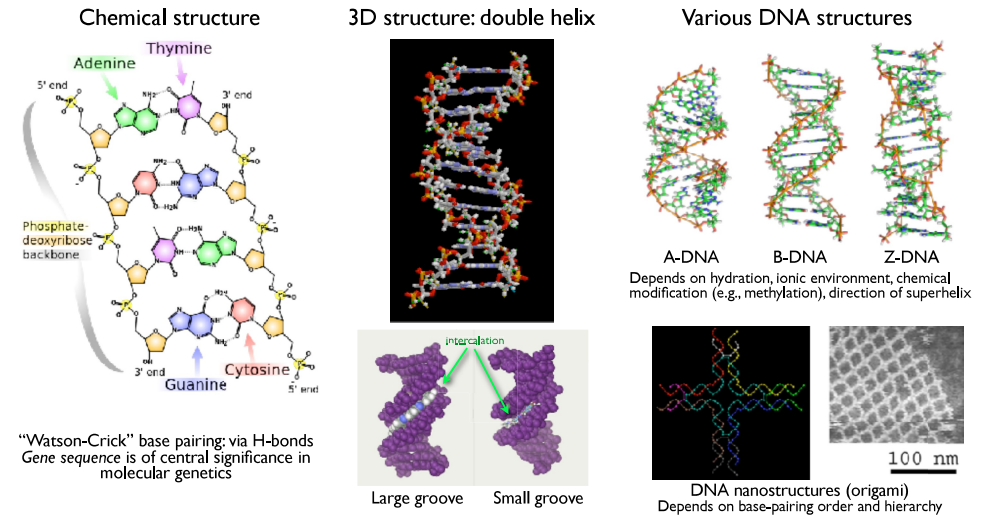


Relationship between persistence length (l) and contour length (L) in biopolymers



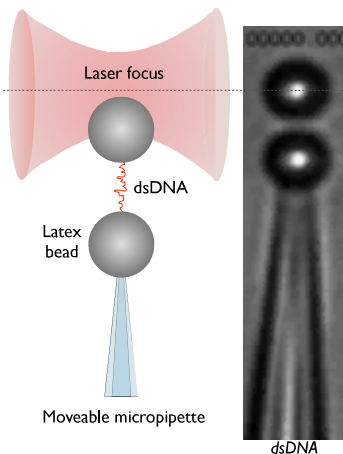
I. DNA: deoxyribonucleic acid

Function: molecule of biological information storage

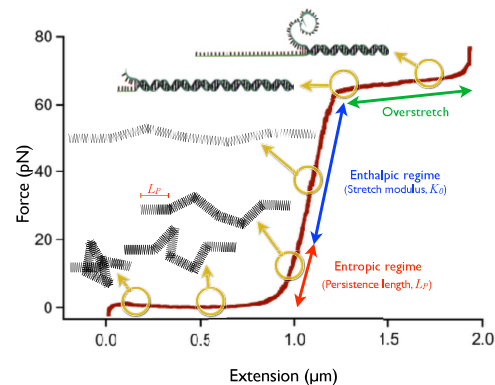


The DNA molecule is elastic!

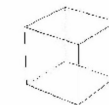
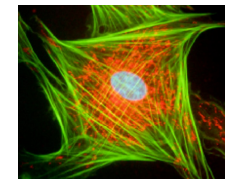
Force measurement: with optical tweezers



Force versus extension curve of a single dsDNA molecule



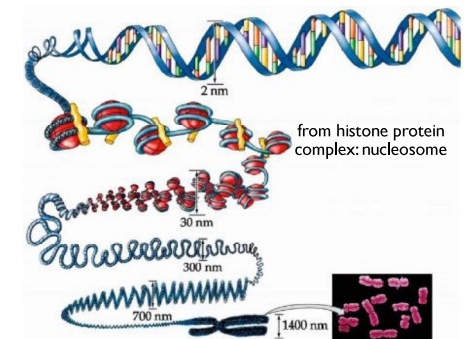
How much DNA is in a cell?



Simplified cell model: cube

Solution: DNA needs to be packed

Chromosome condensation

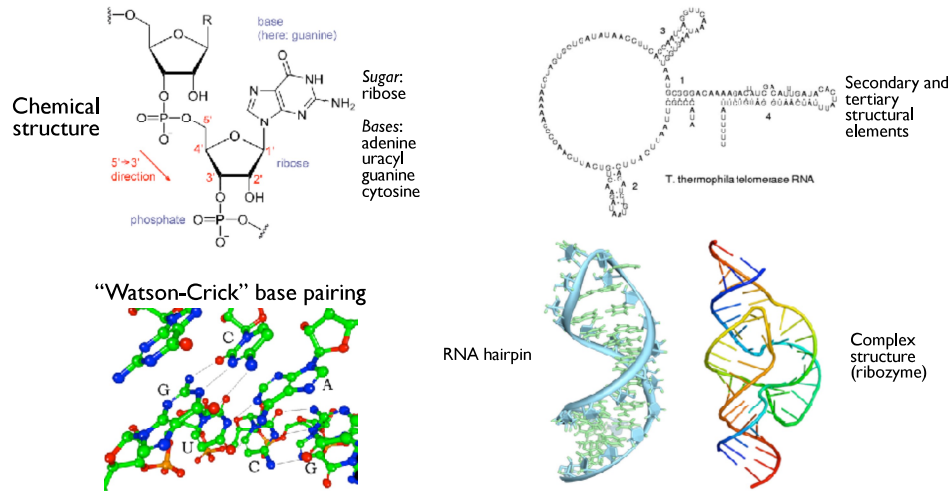


	Cell: 20 μm edge cube	Analog - Lecture hall: 20 m edge cube
DNA thickness	2 nm	2 mm
Full length of human DNA	~2 m	~2000 km (!!!) (Perimeter of Hungary: ~2200 km)
Persistence length of dsDNA	~50 nm	~50 cm
End-to-end distance (R)	~350 μm (!)	~350 m (!)
Volume of fully compacted DNA	~2 × 2 × 2 μm ³	~2 × 2 × 2 m ³ (= 8 m ³)

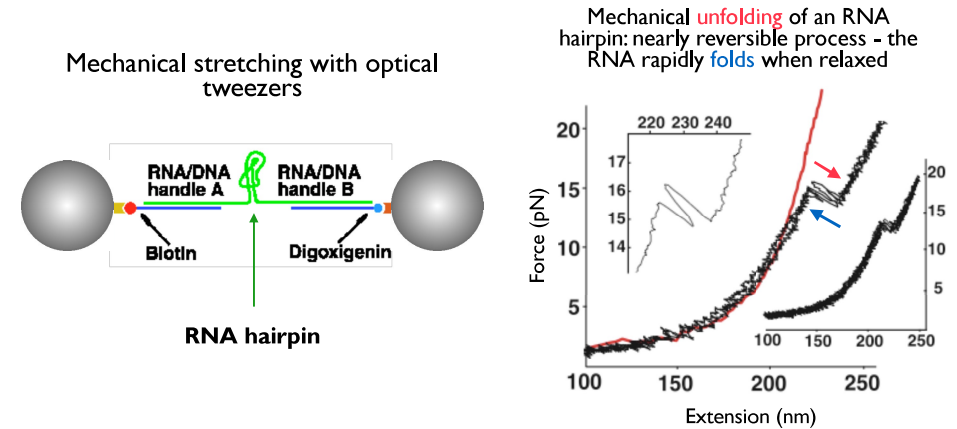
- **Condensins** play a role in high-order DNA packaging
- DNA chain: complex linear path with roadblocks!

2. RNA: Ribonucleic acid

Function: information transfer (transcription), structural element (e.g., ribosome), regulation (turning gene expression on and off)

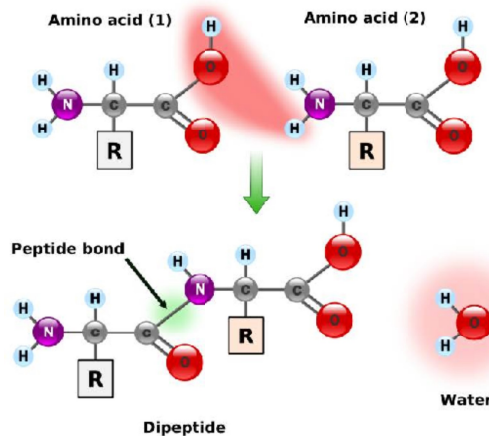


RNA structure can be perturbed with mechanical force



3. Proteins: polymers connected with peptide bonds

Function: most important molecules of the cell. Highly diverse functions - structure, chemical catalysis energy transduction, motoric functions, etc.



Formation of the peptide bond: condensation reaction followed by the release of water

Protein structure

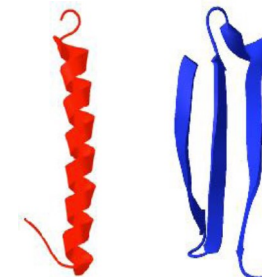
Primary

Amino acid sequence

Determines spatial structure as well.

Secondary

α-helix
β-sheet
β-turn (β-hairpin)

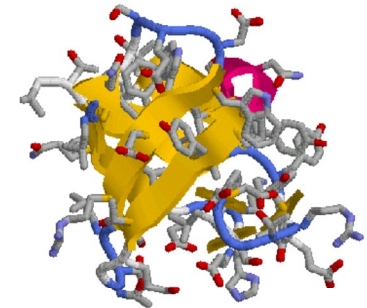


α-helix:
•right handed
•3.4 residue/turn
•H-bridges

β-sheet:
•parallel or
•antiparallel
•H-bridges between distant residues

Tertiary

3D structure of single-chain protein



*Quaternary structure: binding of independent subunits into a complex

Bonds holding protein structure together

- Weak (secondary) bonds
1. **Hydrogen bond**: proton sharing between proton-donor side chains.
 2. **Electrostatic interaction** (salt bridge): between oppositely charged residues.
 3. **van der Waals bond**: weak interaction between atoms (molecules) with closed electron shells.
 4. **Hydrophobe-hydrophobe interaction**: between hydrophobic residues (in the interior of the molecule).
- Covalent bond
5. **Disulfide bridge**: between cysteine side chains; connects distant parts of the protein chain.

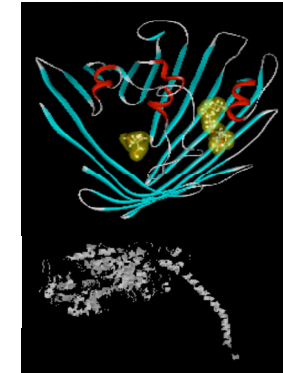
Protein structure classes

1. All alpha



calmodulin

2. All beta



porin

(3. Alpha-beta)

4. Multidomain

Domain:
folding subunit

myosin

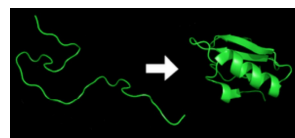
Although there are as many sequences as proteins, the spatial structures are classified into a surprisingly small number of classes!

How is the three-dimensional structure acquired?



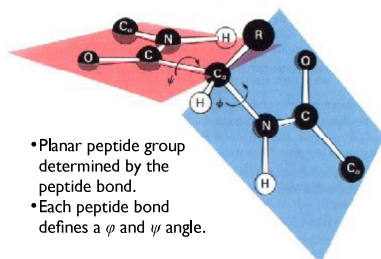
Christian Anfinsen
(1916-1995)

Anfinsen: proteins fold spontaneously (sequence determines structure)



Unfolded state
Native state (N)
Lowest energy

Levinthal's paradox (Cyrus Levinthal, 1969):
Are all available conformations explored?

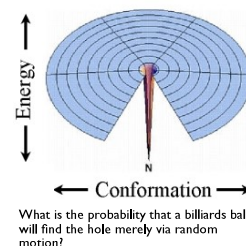


- Planar peptide group determined by the peptide bond.
- Each peptide bond defines a ϕ and ψ angle.

Number of possible conformations (degrees of freedom): i^n

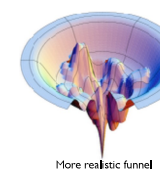
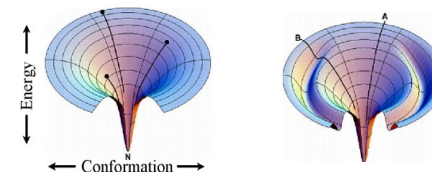
i = number of possible angular positions of a given ϕ or ψ angle
 n = total number of ϕ and ψ angles

Example: in a peptide composed of 100 residues the number of possible ϕ or ψ angles is 2. $n=198$. Number of possible conformations: 2^{198} (!!!)



Protein folding is guided by the shape of its conformational space

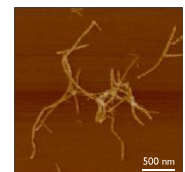
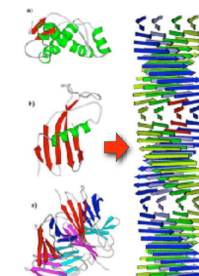
Shape of conformational space:
"Folding funnel"



- Proteins "slide down" the wall of the funnel.
- Folding funnel shape can be complex (determination of the shape is usually very difficult).
- A protein may get stuck at intermediate states (pathology).
- In the living cell chaperones assist folding.

Pathology

- Protein "folding diseases"
- Alzheimer's disease
- Parkinson's disease
- II-type diabetes
- Familial amyloidotic neuropathy

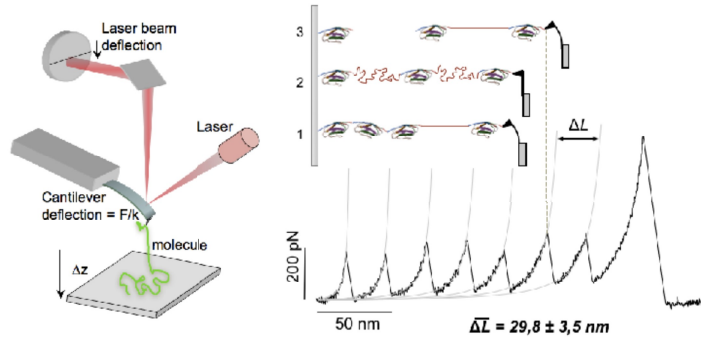


β -fibrils:
undissolved precipitate
cross- β structure

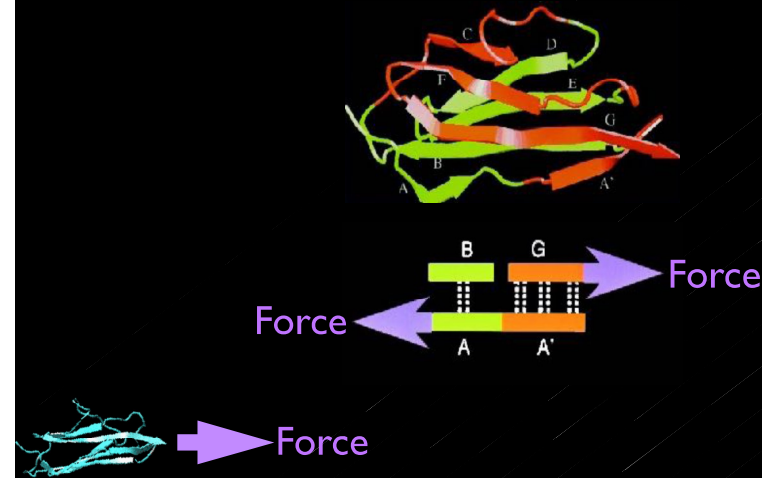
Methods of protein unfolding (denaturation)

- Heat
Break secondary chemical bonds
- Chemical agent
Disrupt secondary and tertiary structure
- Mechanical force

Mechanical unfolding of a single protein with atomic force microscope

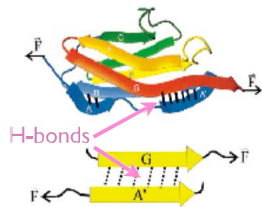


Structural basis of mechanical stability

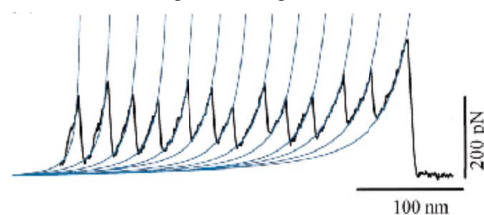


Biological logic of mechanical stability

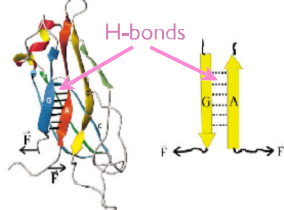
Parallel coupling of structure-stabilizing H-bonds



High unfolding forces



Serial coupling of structure-stabilizing H-bonds



Low unfolding forces

