

# Special Properties of Water

class – 22 (16.11.24)

LS2103 (Autumn 2024)

IISER Kolkata

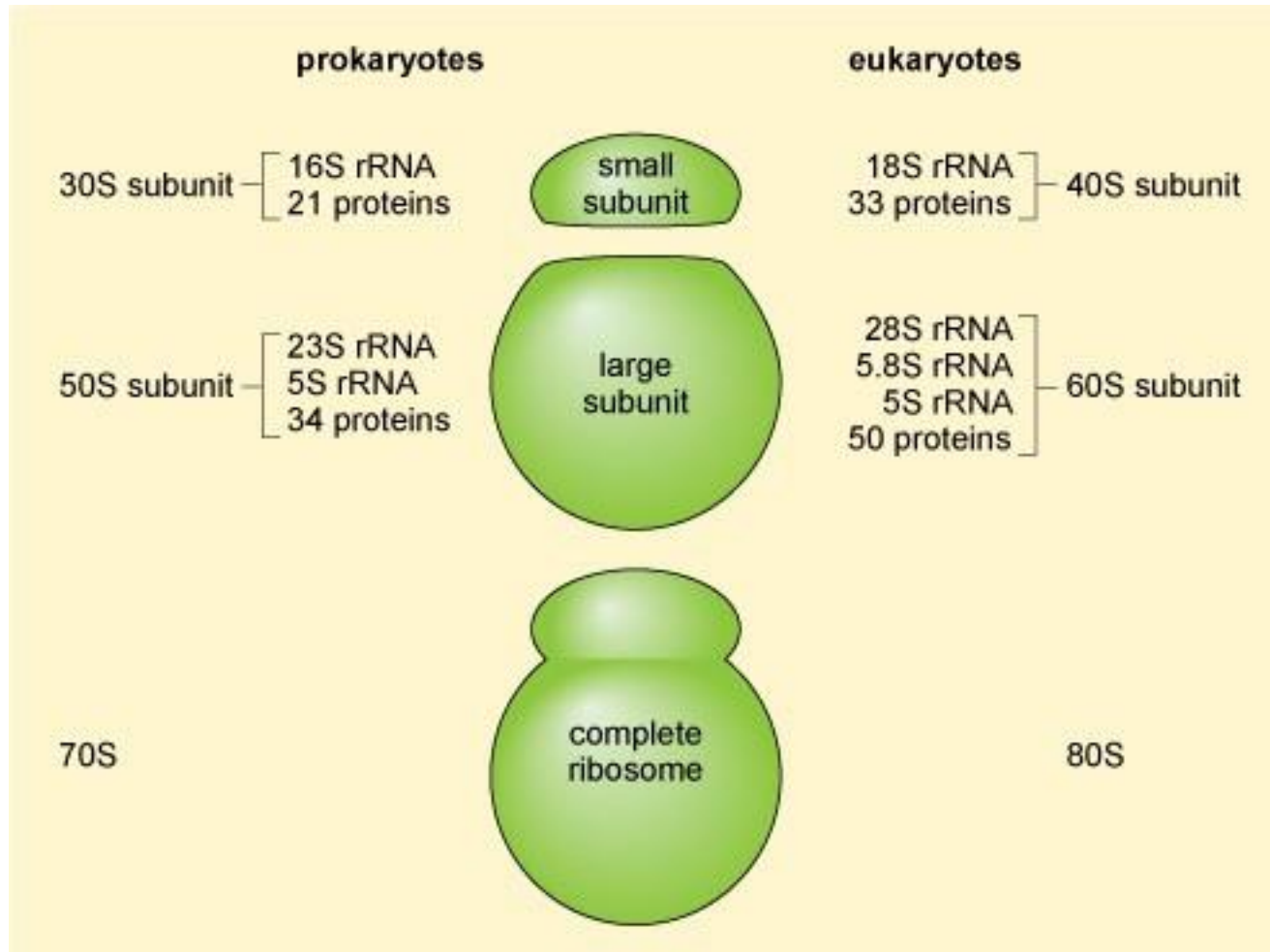
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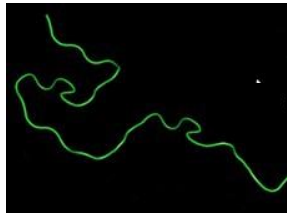
<https://www.iiserkol.ac.in/~n.sengupta/>

# Ribosome subunits

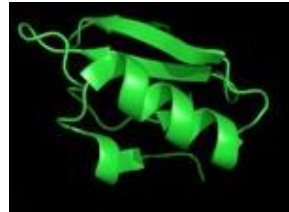


Svedberg units are non additive – why?

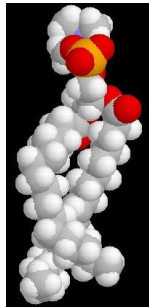
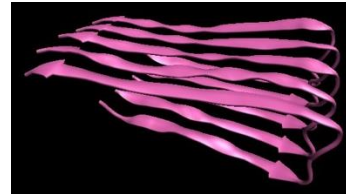
# Biological “solvent” is predominantly water



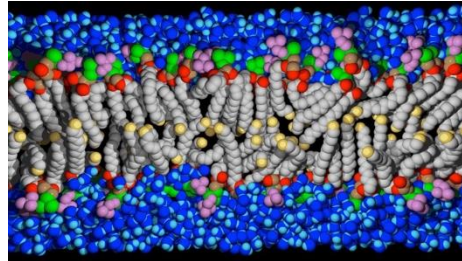
polypeptides



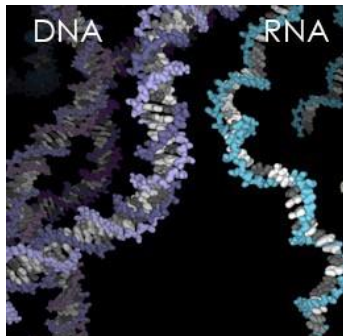
protein structures



phospholipids



membranes



nucleic acids

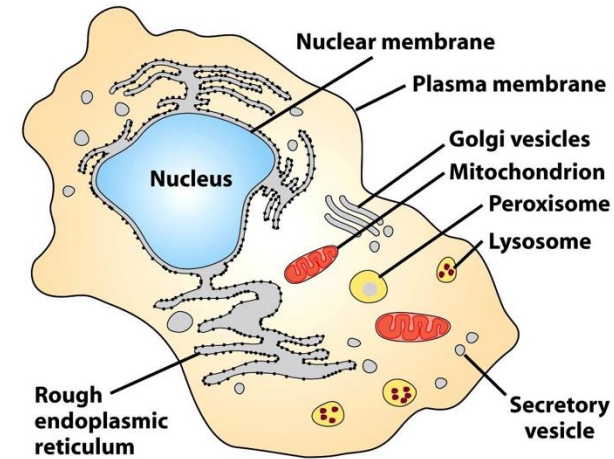
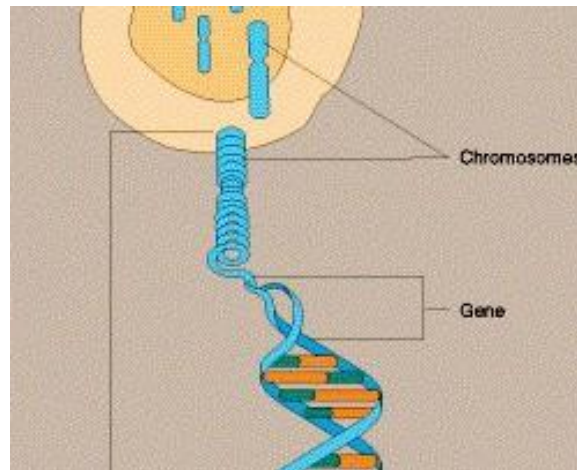
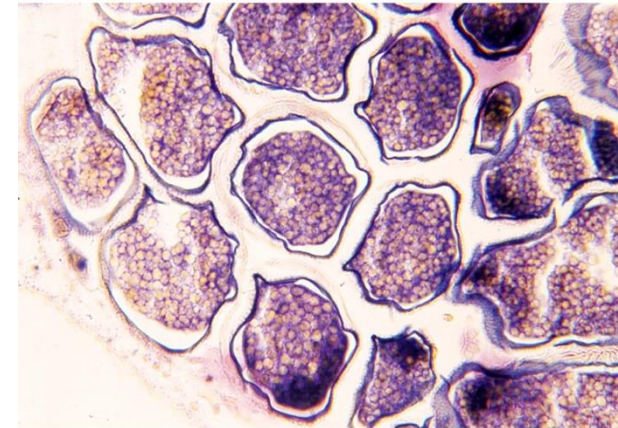


Figure 1-2b part 2  
Molecular Cell Biology, Sixth Edition  
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# Biological “solvent” is predominantly water

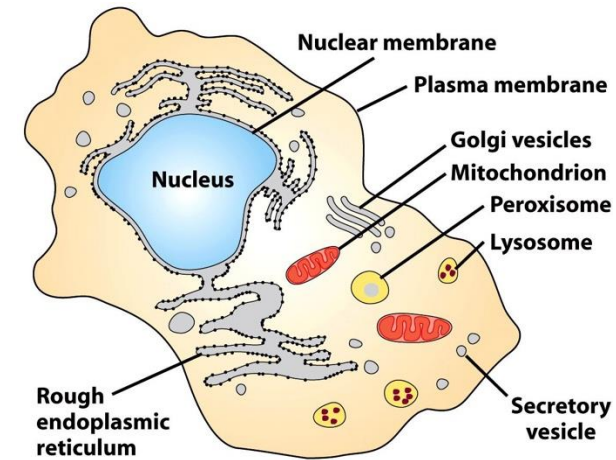
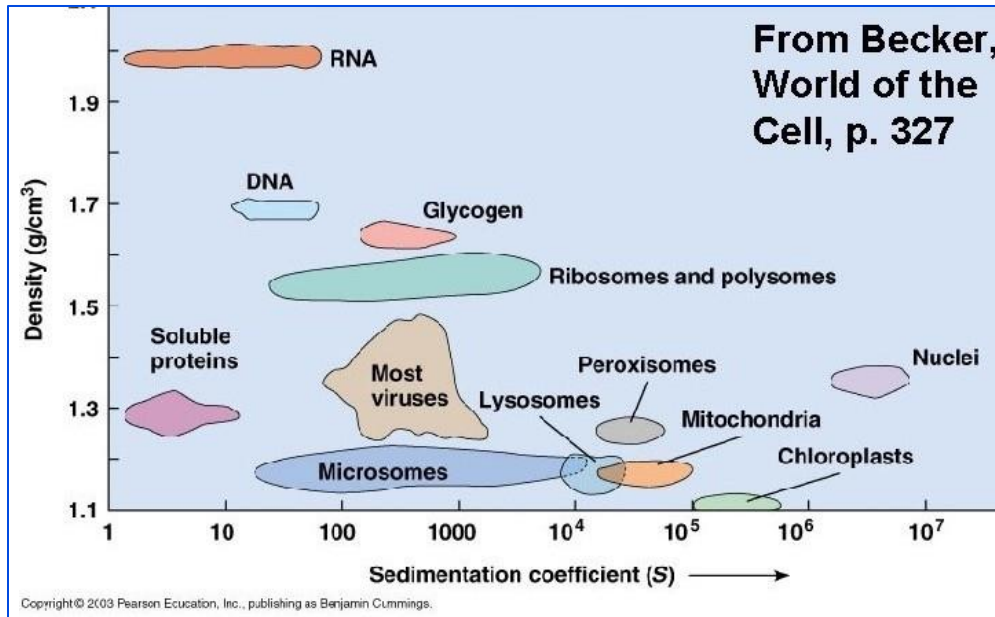
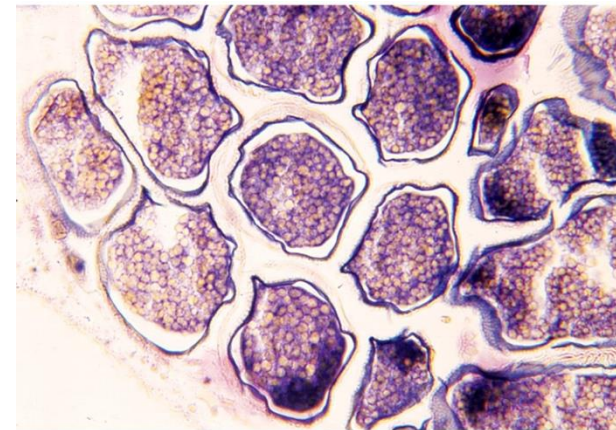


Figure 1-2b part 2  
Molecular Cell Biology, Sixth Edition  
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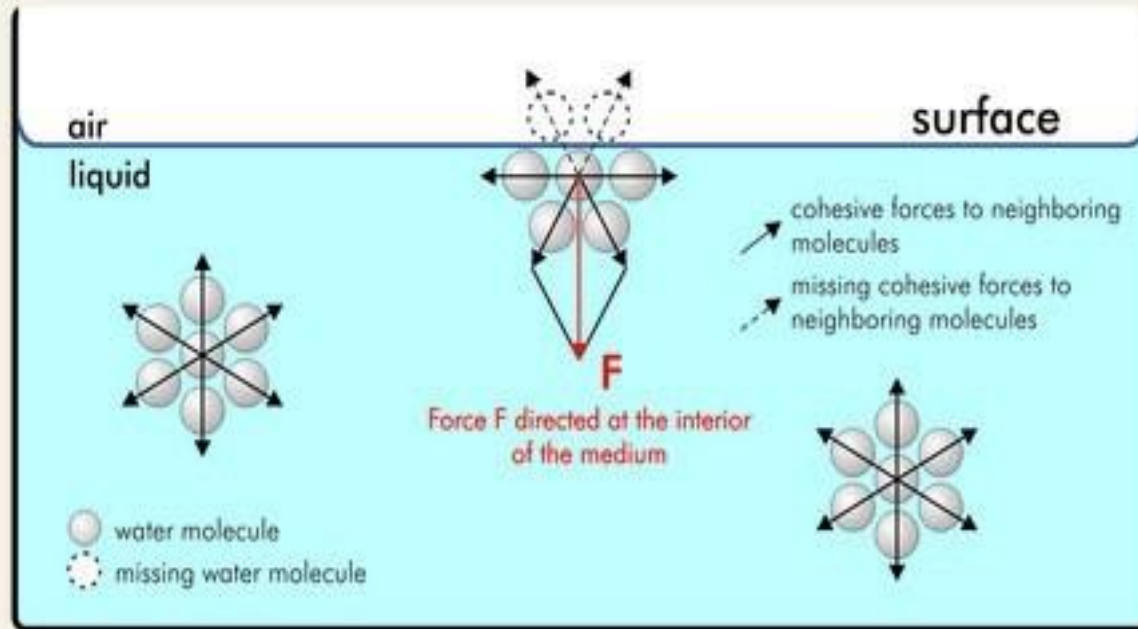




# Biological “solvent” is predominantly water

**Surface Tension:** Attraction of surface particles to the (inner) bulk that resists external force

**Units / Dimensions:** Energy per unit area, or, Force per unit length



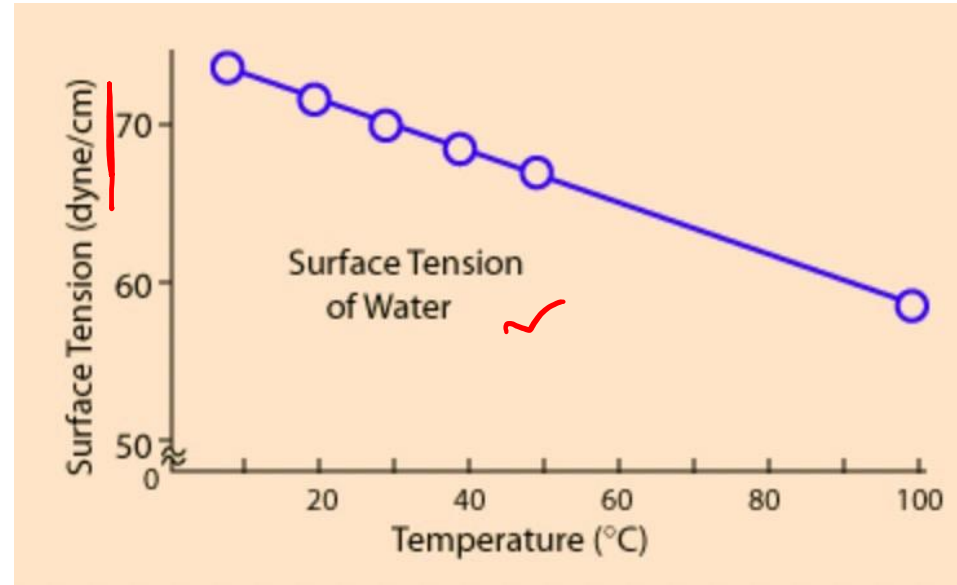
<https://www.sita-process.com/information-service/process-parameter-surface-tension/overview/>

# Biological “solvent” is predominantly water

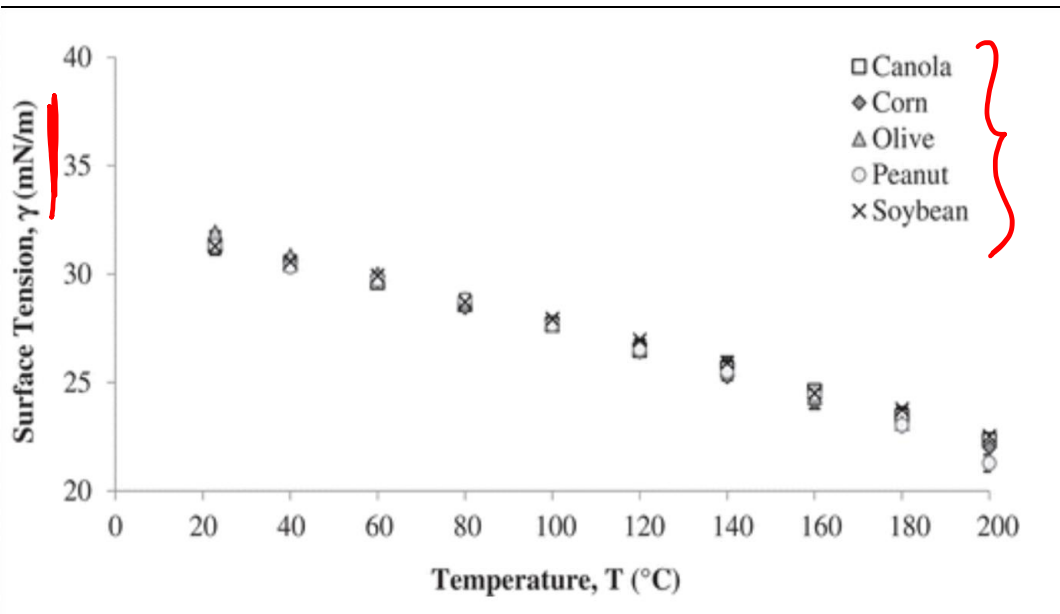
## Comparison with other liquids

- Water has higher surface tension than other common solvents

Sahasrabudhe et al., Int. J. Food. Prop., 2017



<http://hyperphysics.phy-astr.gsu.edu/hbase/surten.html>

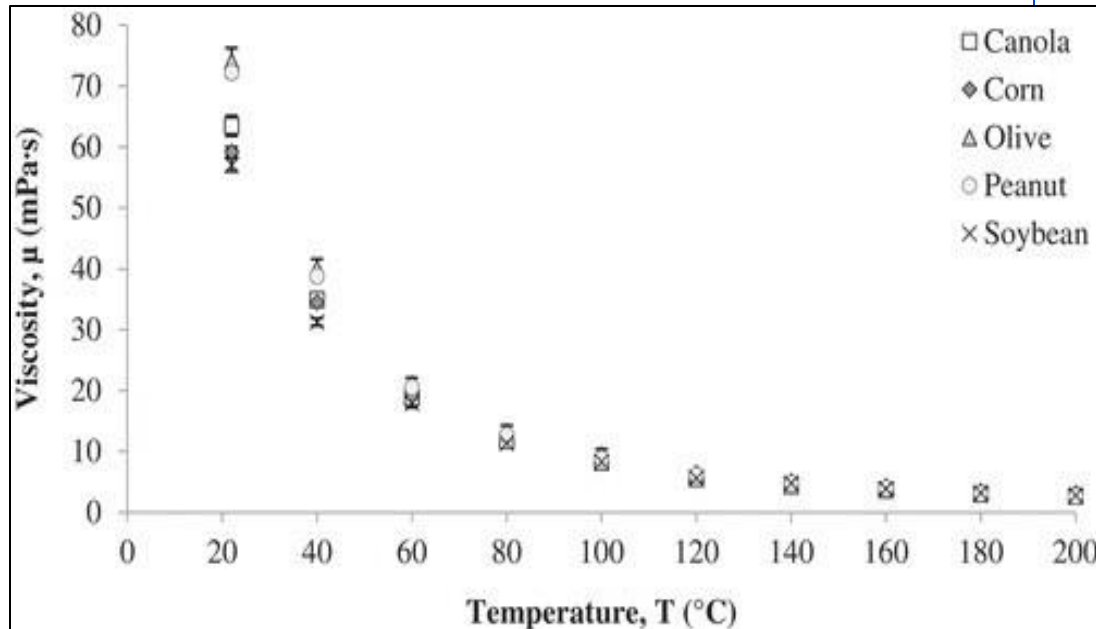


# Biological “solvent” is predominantly water

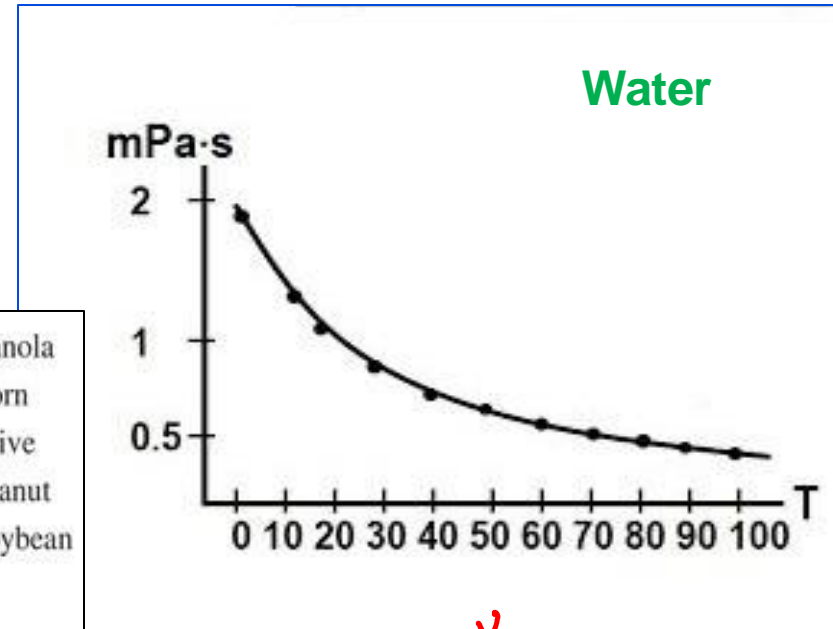
## Comparison with other liquids

### Dynamic viscosity ( $\eta$ ):

Sahasrabudhe et al., Int. J. Food. Prop., 2017



- Water has **higher surface tension** than other common solvents
- Water has **lower viscosity** than other common solvents

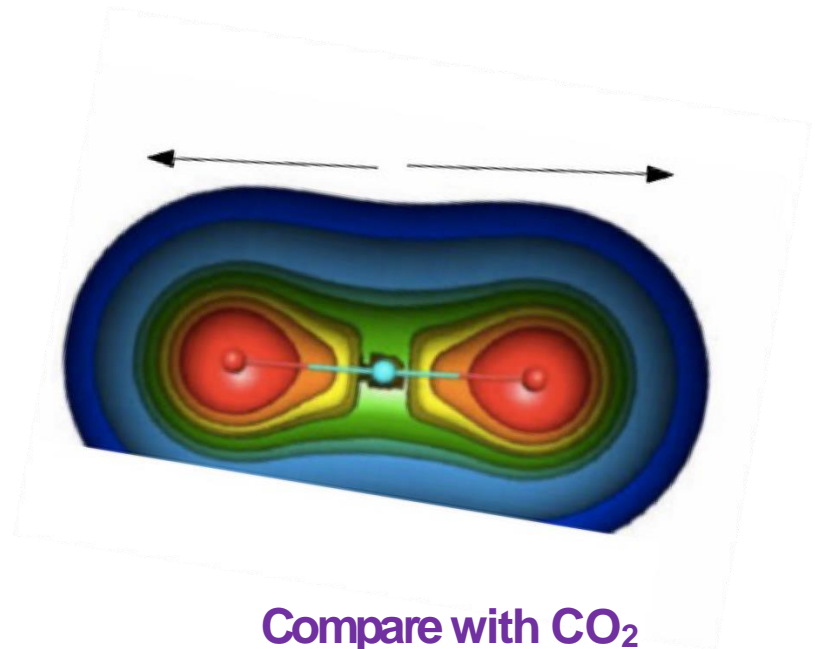
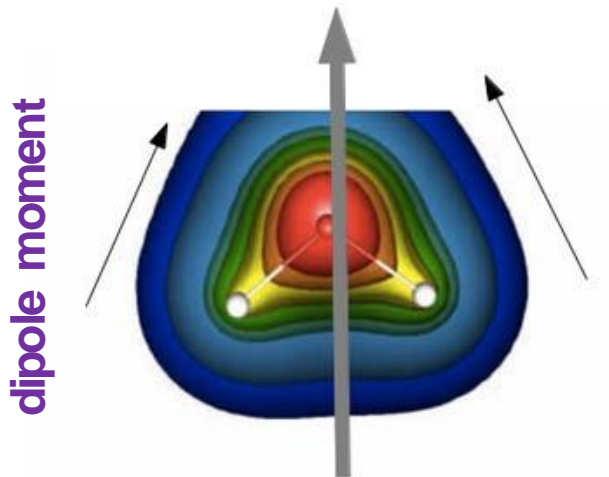
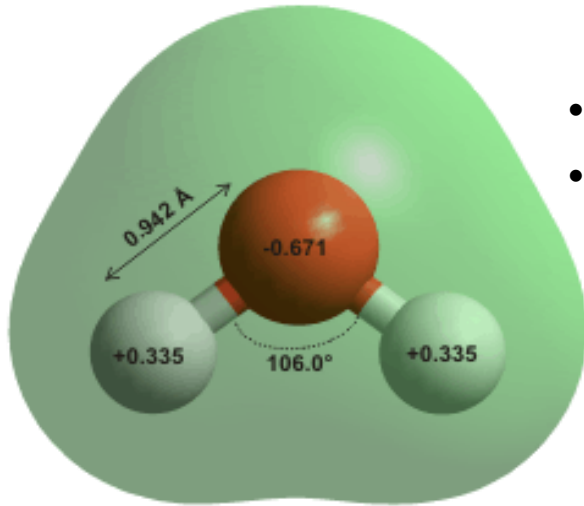


$T(^{\circ}\text{C})$	$\mu(\text{mPa}\cdot\text{s})$
0	~1.8
10	1.308
20	1.002
30	0.7978
40	0.6531
50	0.5471
60	0.4658
70	0.4044
80	0.3550
90	0.3150
100	0.2822

# The molecular picture of water

## What is 'special'?

- Two lone pairs of electrons at the oxygen atom
- Resulting shape: Bent structure and **net dipole moment** (~1.84 Debye)



Compare with  $\text{CO}_2$   
(no net dipole moment)



# Results in hydrogen bonded and tetrahedral water network

**Hydrogen bond (HB):** A mainly electrostatic attraction between a H-atom covalently bonded to one electronegative atom ('HB donor'), and another electronegative atom ('HB acceptor').

- HBs are strongly "directional"
- They can "co-operate"

$$q = 1$$

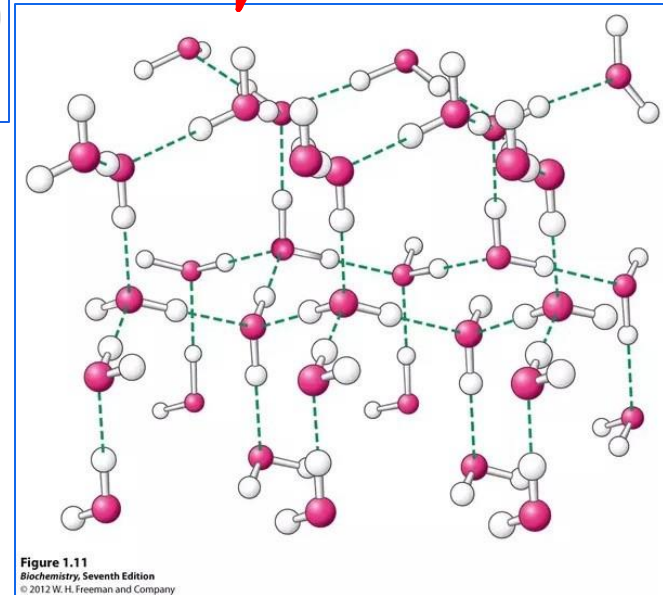
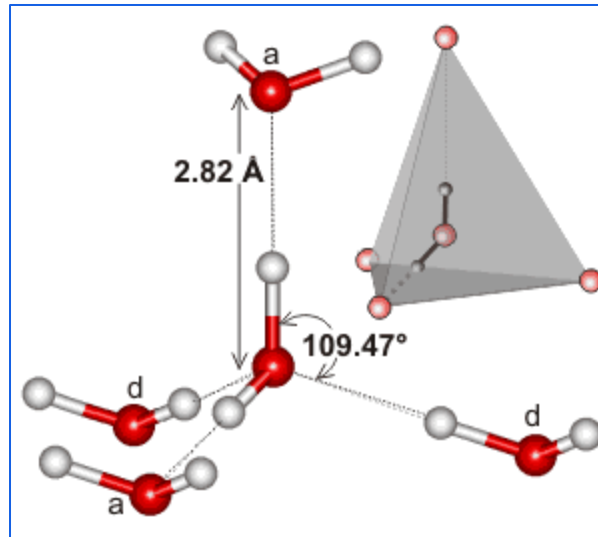
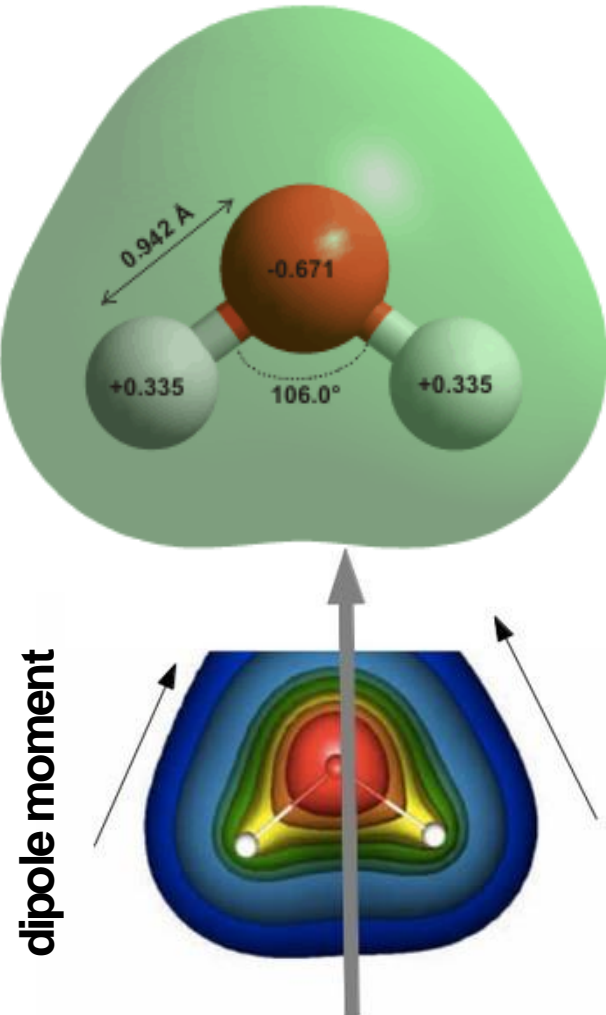


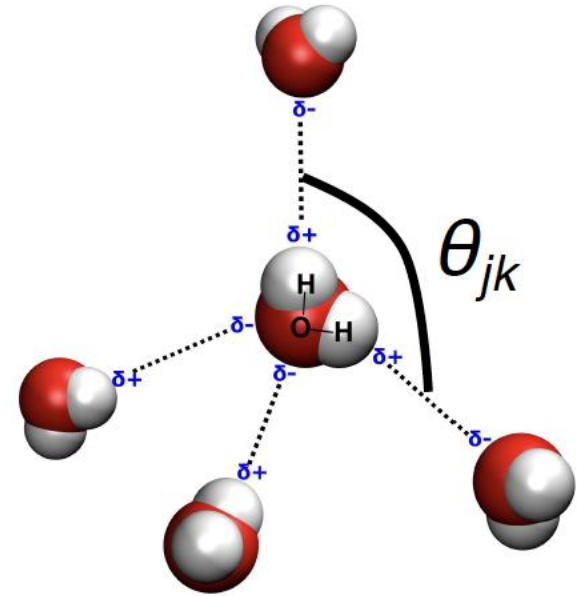
Figure 1.11  
Biochemistry, Seventh Edition  
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# Water tetrahedrality

**Orientational Order Parameter:**

$$q = 1 - \frac{3}{8} \sum_{j=1}^3 \sum_{k=j+1}^4 \left( \cos \theta_{jk} + \frac{1}{3} \right)^2$$

$q = 1$  for perfect ice



Errington and Debenedetti, Nature, 2001

<https://www.nature.com/articles/35053024>

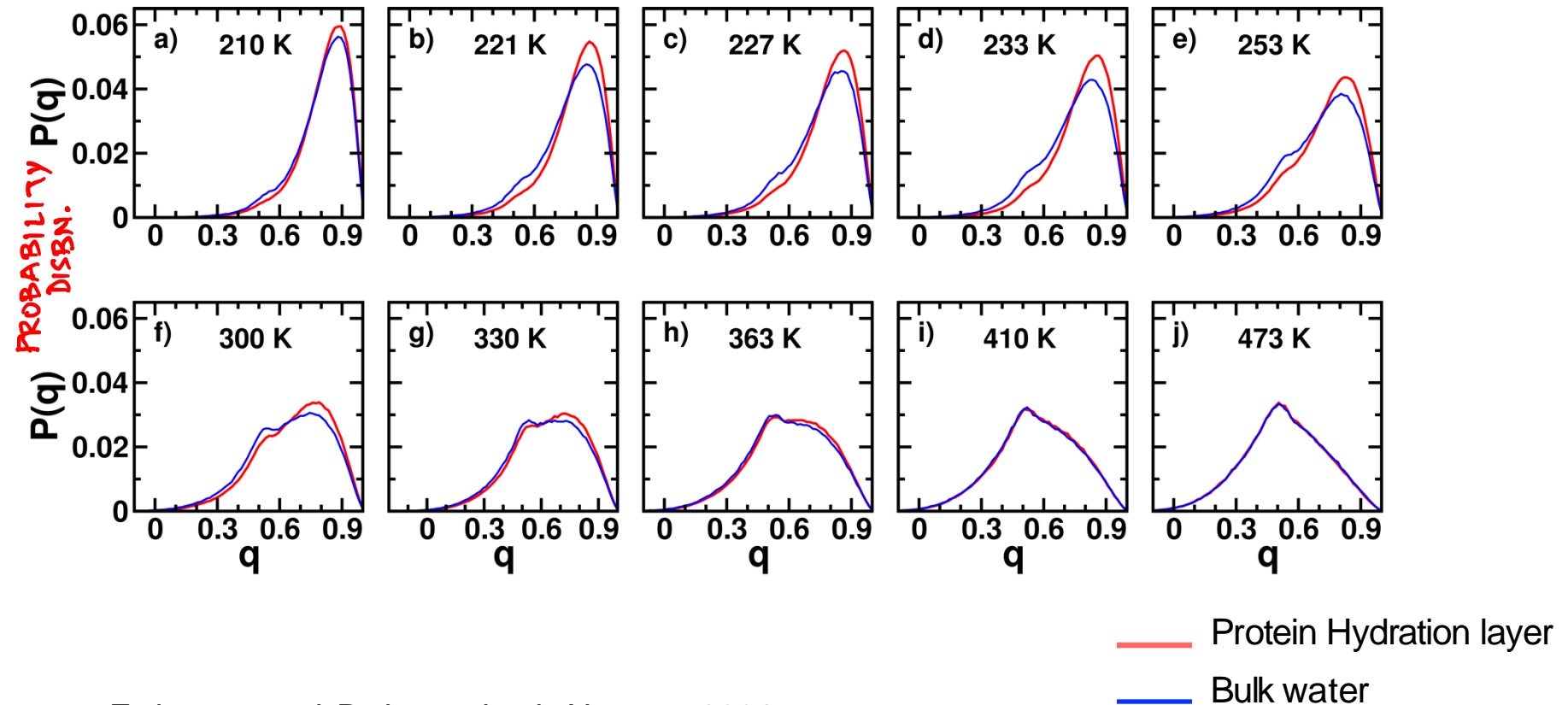
# Hydration layer ordering (\*)

(results from computer simulations)

$q = 1$  for perfect ice



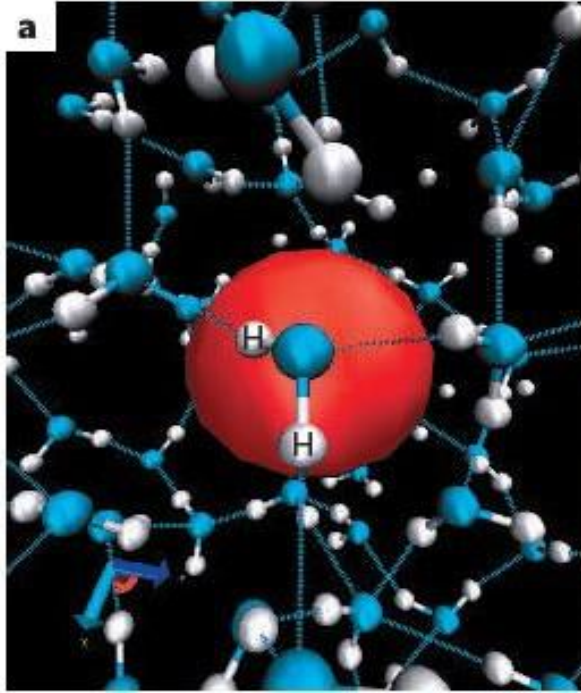
Ordering modified with temperature, and in presence of solute (biomolecule)



Errington and Debenedetti, Nature, 2001

<https://www.nature.com/articles/35053024>

Under "no perturbations", every  $\text{H}_2\text{O}$  will "make & break" H-Bonds  
**Molecular scale inspection** *with other  $\text{H}_2\text{O}$ s*

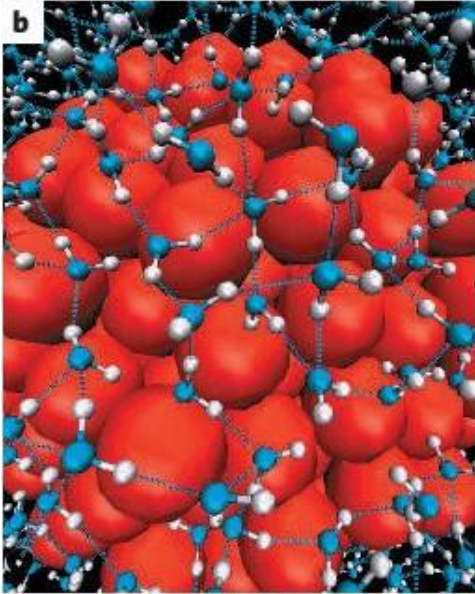


## Solute radius comparable to $\text{H}_2\text{O}$

- Insignificant loss in H-bonds of nearby waters
- Marginal overall re-ordering of H-bonding network

Chandler, Nature Reviews, **2005**, 437, 640

# Molecular scale inspection



## Large solute (eg. protein)

- No. of water H-bonds reduced significantly
- **Interface formed to minimize loss in H-bonds**
- Interface tends to move away from solute
- **Energetic cost of solvating the solute** (ie. forming the interface):

*Assuming a spherical solute of radius  $R'$ ,*

*Free  
Energy  
cost*

$$\underline{\underline{\Delta G}} \approx (4\pi R'^2)\gamma$$

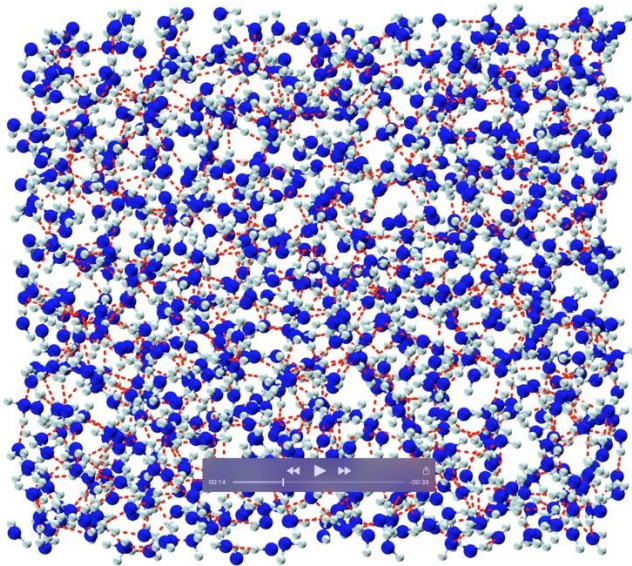
$\gamma$ : Surface tension

Chandler, Nature Reviews, 2005, 437, 640

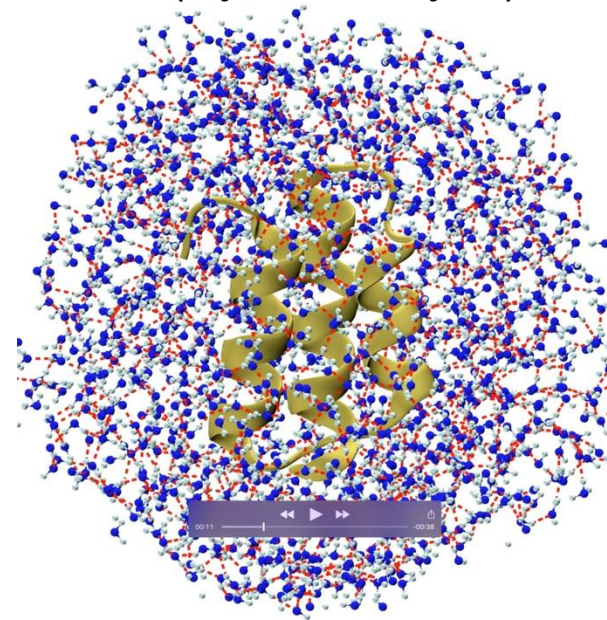


# Visualizing the H-bonded network (computer simulations)

Bulk water



Water close to a biomolecule  
(‘hydration layer’)

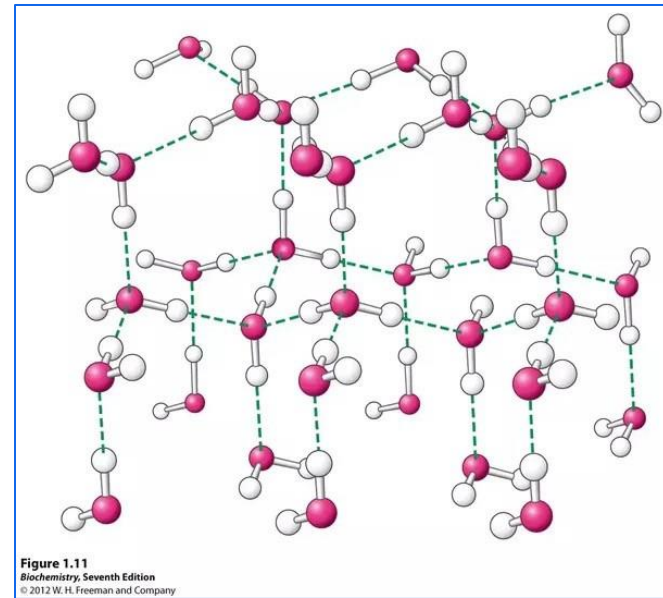
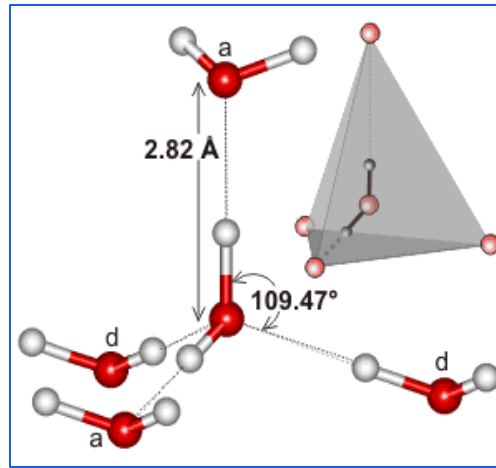


- Protein-water (P-W) H-bonds **compete with** water-water (W-W) H-bonds
- **Lifetimes and energetics** of P-W and W-W H-bonds may differ
- Water helps the protein achieve **functional flexibility**
- Water plays **thermodynamic role** in folding, enzyme-ligand binding, etc

**Hydrogen bond (HB):** A mainly electrostatic attraction between a H-atom covalently bonded to an electronegative atom ('HB donor'), and another electronegative atom ('HB acceptor').

**Pure water:**  $E_{\text{Hbond}} \sim 5.6 \text{ kCal mol}^{-1}$ , ie.  $\sim 9.7 \text{ RT}$  at room temperature

**Prob.** A Raman spectroscopy study showed that breakage of a single H-bond within pure water at room temperature (300 K) is commensurate with an **enthalpy increase of 1.9 kcal mol<sup>-1</sup>**, and an **entropy increase of 2.4 k<sub>B</sub>**.



- What is the **free energy cost** of breaking a single H-bond at room temperature?