Chapter 1

Topic: Overview of basic principles

Four major themes of biochemistry

- I. What are living organism made from?
- II. <u>How do organism acquire and use energy?</u>
- III. How does an organism maintain its identity across generation?
- IV. How organism communicate with its environment?

Central dogma of molecular biology

1866 Gregor Mendel: Plants contains a pair of factors, one inherited from each parents.

1869 Friedrich Mieschet isolated nucleic acids

1944 Avery, Mac Leod, Mc Carty – DNA carry genetic information.

1958 F. Crieck formulated central dogma of molecular biology



Biochemistry uses tools of chemistry to explain life on molecular level

Approach : reductionism

Living organism \Rightarrow cells \Rightarrow organelles/membranes \Rightarrow <u>molecules</u> Molecules \Rightarrow composition??? function ??? structure (3D) ??? reaction mechanism



Holistic approach – even complete set of information about all molecules in organism does not reveal how the organism live. We need to study intact cells and organisms to understand how they live.

Processes of current interest:

Genomics: study of genetic code and its regulation

Proteomics: study or protein function as determined by genetic code

<u>Cloning</u>: insertion of foreign DNA into cell or duplication of an organism through a genetic modification

Epigenetics: the study of heritable changes in gene expression that occur without a change in DNA sequence.

Biochemically important molecules:

Bio-macromolecules

proteins (various structures and functions)nucleic acidsmembranes

small molecules called metabolites.



Cellular structure

- Viruses smallest organized particle (nucleic acid and protein)
- Prokaryotic cells- smallest self-replicating organisms
 - Simple structure (bacterial cell, no defined nucleus)
- Eukaryotes complex organism
 - Plasma membrane, cell wall, cytoplasm, cytosol, organells

All living organism single cell to complex organisms:

viruses (20 – 200nm), *eukaryotic cells* (10 – 100 µm)

composed of same building blocks

Similar principles in –

replication metabolism ability to organize at the cell level



Key metabolic processes are common to many organisms.

Different organisms have macromolecules of similar structure and common biochemical processes suggest that all organisms evolved from a common ancestor.



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TATA box binding protein



Sulfolobus archaea





Arabidopsis thaliana









Figure 1.1 Biochemistry, Eighth Edition © 2015 Macmillan Education [Photos: (Left) Eye of Science/Science Source; (middle) Holt Studios/Photo Researchers; (right) Time Life Pictures/Getty Images] On the basis of biochemical characteristics, all organisms can be placed in one of three domains:



Prokaryotes:

Bacteria and Archaea Single cell organism Missing organelles

Archaea: metabolic pathways similar to those of eukaryotes (enzymes involved in transcription and translation.) Often organism that lives in harsh condiditions

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Electron micrographs of *E. coli* cells. Condensed DNA (**nucleoid** is shown in red)



Small molecules

- ATP, AMP. NADH --coenzymes
- sacharides
- lons
- lipids

Covalent and non-covalent interactions are important for the structure and stability of biological molecules

Covalent bonds, formed by electron sharing between two adjacent atoms, are the strongest bonds.

A typical C-C covalent bond has a distance of 1.54 Å and a bond energy of 355 kJ mol⁻¹ (85 kcal mol⁻¹).

Energy units: Joules (J) 1J is the energy that is required to move 1 meter against a force of 1 N. Distances: 1 Angstrom is 0.1 nm or 10⁻¹⁰ m.

I. Disulfide bond:

I. Peptide bond:





Resonance structures.



Resonance structures of adenine

Noncovalent bonds Ionic interactions.

Ionic interactions, also called electrostatic interactions, occur between oppositely charged molecules.

Electrostatic interactions in water have a bond distance of ≈ 3 Å and a bond energy, given by Coulomb energy, of 5.86 kJ mol⁻¹ (1.4 kcal mol⁻¹)



Noncovalent bonds Hydrogen bonds.

Hydrogen bonds occur between an electronegative atom and a hydrogen covalently bonded to another electronegative atom.

Hydrogen bonds vary in bond distance from 1.5 Å to 2.6 Å with bond energies from 4-20 kJ mol⁻¹ (1-5 kcal mol⁻¹)



Hydrogen- Hydrogenbond donor bond acceptor





О—Н----N



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van der Waal interactions.

Van der Waals interactions occur between two molecules that sufficiently close (the van der Walls contact distance) such that transient electronic asymmetries in one molecule induce complementary interaction in the partner molecule.

Van der Waals interactions have bond energies from 2-4 kJ mol⁻¹ (0.5-1.0 kcal mol⁻¹)



Water is a polar molecule. The charges on the molecule are not evenly distributed.



Water molecules solvate proteins and ions in the solutions Water molecules also form a channels through protein structure necessary for proton transport

The hydrophobic effect.

Nonpolar molecules in water can be driven together by the hydrophobic effect which is powered by the increase in entropy of water. The associated interactions are called hydrophobic interaction.



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Noncovalent forces and DNA molecule





Stability of DNA double helix

When a double helix forms, charge repulsion occurs between the negatively charged phosphates of the backbone.

These repulsive forces are reduced by the high dielectric constant of water and interaction of positively charged ions with the phosphate groups.



Figure 1.13 *Biochemistry*, Eighth Edition © 2015 Macmillan Education Hydrogen bonds form between bases, which are in the hydrophobic interior away from the disrupting effects of water.



Unnumbered 1 p10 *Biochemistry*, Eighth Edition © 2015 Macmillan Education In the interior of the helix, bases are stacked and interact with one another through van der Waals interactions.



Biochemistry, Eighth Edition © 2015 Macmillan Education The First Law of Thermodynamics states the total energy of a system and its environment is constant.

The Second Law of Thermodynamics states the total entropy of a system plus that of its surroundings <u>always</u> <u>increases</u>.

Entropy can decrease locally in the system if there is a corresponding increase in entropy in the surroundings.

Thermodynamic of chemical reactions:



 ΔG – change in the free energy

 ΔH – change in the enthalpy,

 ΔS – change in the entropy of the system

Spontaneous reaction, nonspontaneous reaction

Comparing to simple chemical reaction – entropy contribution is significant in biological reactions.

Chemical equilibrium and standard state

 $\Delta G = \Delta G^{\circ} + RTInK_{eq}$ K_{eq} equilibrium constant

equilibrium

 $\Delta G = 0 => \Delta G'^\circ = - RTInK_{eq}$

ΔG " free energy change at standard conditions

Equilibrium constant of reaction can be calculated from the standard free energy and vice versa

Living organism – high organization of structures => low entropy Organism maintain this thermodynamically unfavorable state as long As they are obtaining free energy from food. Acid-base reactions are central in many biochemical processes

Acid-base reactions involve the addition or removal of a H⁺ ion. Such reactions are important in biochemistry.

pH is a measure of the H⁺ concentration and is defined by

 $pH = -log[H^+]$

 H^+ and OH^- ions are formed upon the dissociation of H_2O .

 $H_2O \Longrightarrow H^+ + OH^-$

The equilibrium constant K_{eq} for the dissociation of water is given by

 $K_{eq} = [H^+][OH^-]/[H_2O]$ K_w, the ion constant of water is given by $K_w = K_{eq} \times [H_2O]$ This can be simplified to $K_w = [H^+][OH^-]$

At pH 7, $[H^+] = [OH^-] = 10^{-7}$

From this, we can $[OH^-] = 10^{-14}/[H^+]$ and $[H^+] = 10^{-14}/[OH^-]$

$$K_{\rm w} = [{\rm H^+}][{\rm OH^-}] = 1.0 \times 10^{-14} \,{\rm M^2}$$

 $K_{\rm w} = [{\rm H^+}][{\rm OH^-}]$

Acid-base reactions can disrupt the double helix

As base is added to a solution of double helical DNA, the helix is disrupted or denatured.



Figure 1.16 Biochemistry, Eighth Edition © 2015 Macmillan Education The chemical basis of the denaturation is the disruption of base-pairing. For example, the loss of a proton by the base guanine prevents base pairing with cytosine.



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More generally, proton removal from any molecule HA is described by

 $K_{\rm a} = [\rm H^+][\rm A^-]/[\rm HA]$

The pK_a value indicates the susceptibility of proton removal.

 $pK_{a} = -\log(K_{a})$ When pH = pK_a $-\log[H^{+}] = -\log([H^{+}][A^{-}]/[HA])$ and $[H^{+}] = [H^{+}][A^{-}]/[HA]$

 $1 = [A^{-}]/[HA]$ or $[A^{-}] = [HA]$

Thus, when the pH is equal to the pK_a , the concentration of the protonated form of HA (the acid) is equal to the deprotonated form A⁻ (the base).

The N-1 proton of guanine has a pK_a of 9.7. When the pH is near or exceeds this value, the proton is lost, base pairing is disrupted and the helix is denatured.

Buffers regulate pH in organisms and in the laboratory

An acid-base conjugate pair resists changes in the pH of a solution.

In other words, it acts as a buffer. A buffer is most effective at a pH near its pK_a .



Figure 1.17 Biochemistry, Eighth Edition © 2015 Macmillan Education The ionization reaction of a weak acid is given by

$$HA \Longrightarrow H^+ + A^-$$

The equilibrium constant for this reaction is

 $K_{\rm a} = [{\rm H}^+][{\rm A}^-]/[{\rm HA}]$

Taking the logarithms of both sides yields

 $\log(K_{a}) = \log([H^{+}]) + \log([A^{-}]/[HA])$

Substituting pK_a and pH and rearranging yield the Henderson-Hasselbalch equation

 $pH = pK_a + \log([A^-]/[HA])$

Consideration of this equation reveals that weak acids are most effective as buffers at pH near their pK_a .