

Basic Biochemistry

ABE 580

Classes of Biomolecules

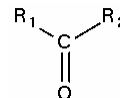
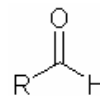
- Carbohydrates
- Lipids
- Amino Acids
- Nucleic Acids
- Other

Carbohydrates

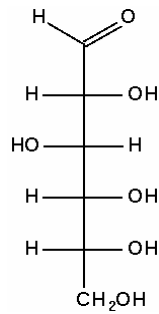
- Sugars
- Composed of C, H, O ($C_nH_{2n}O_n$)
- Biological Uses
 - Energy source/storage
 - Structural Strength or Support
 - Tag for IDing proteins, etc.
- Monosaccharides
- Oligo- and Polysaccharides

Monosaccharides

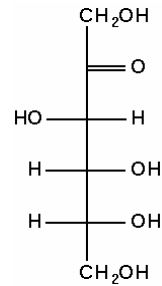
- Aldose
 - Sugars having an aldehyde moiety
- Ketose
 - Sugars having a ketone moiety
- Hexoses
 - Sugars having 6 carbons: $C_6H_{12}O_6$
- Pentoses
 - Sugars having 5 carbons: $C_5H_{10}O_5$
 - Ribose, Xylose



Hexose Isomers Fischer Projections



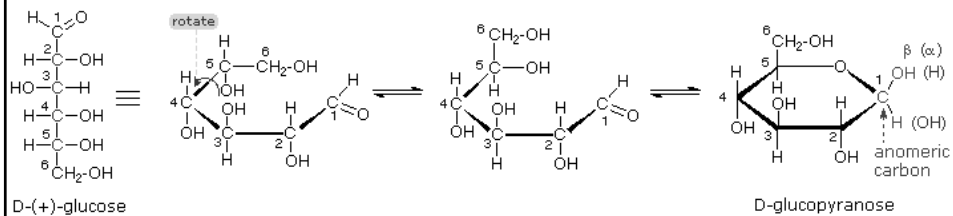
Glucose
Aldose



Fructose
Ketose

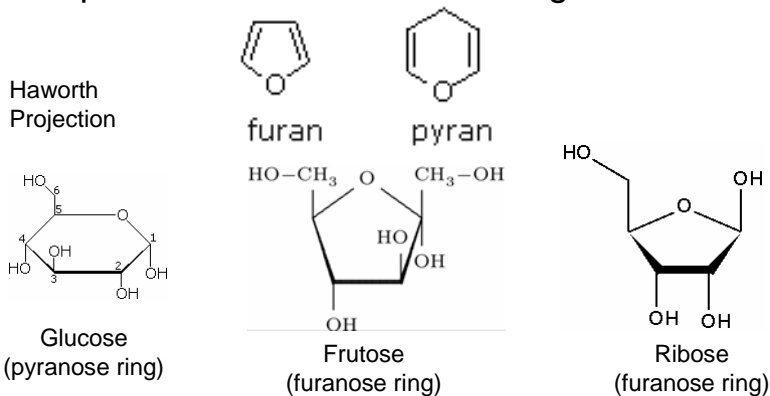
Chain vs Ring Structure

- Aldehyde (aldoses) and ketone (ketose) can form hemiacetal with hydroxyl of carbon on opposite end of chain

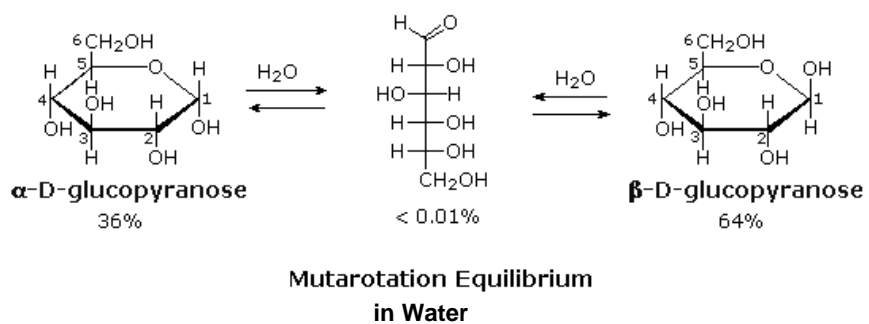


Ring Structures

- Aldohexoses form 6 member rings – pyranose
- Ketohexoses form 5 member rings - pyranose
- Aldopentoses form 5 member rings - furanose

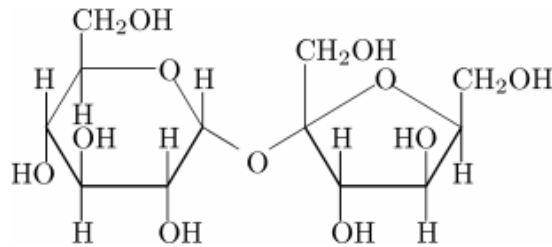


Mutarotation Equilibrium



Carbohydrate Bonds

- Glycosidic Bonds: C – O – C
- Between hydroxyls
 - Ring structures form these bonds!



Glucose α 1-2 fructose
Sucrose

Two Important Polysaccharides

- Polysaccharide – “many saccharides”
- Starch = Glucose α 1-4 Glucose
 - Two forms
 - Amylose (straight chain)
 - Amylopectin (branched chain)
 - Important energy storage (plants) and food
- Cellulose = Glucose β 1-4 Glucose
 - Important structural sugar (wood)
 - Straight chain

Difference in Glycoside Bonds Makes All the Difference

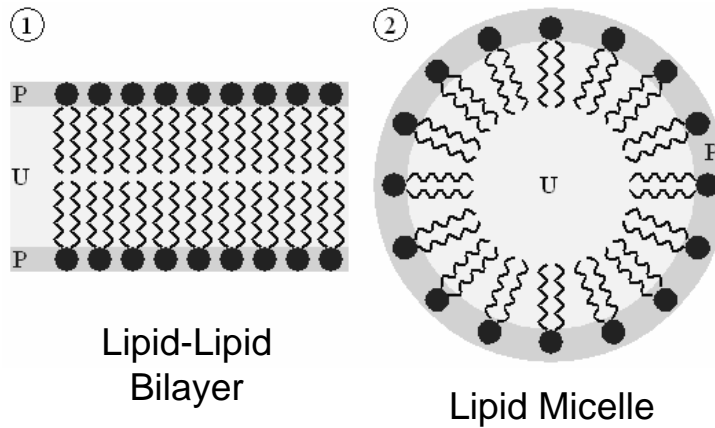
- Starch
 - Amylose chains form helix by hydrogen bonds between the hydroxyl groups of glucose
 - Allows it to be compacted for storage
 - Easy to break hydrogen bonds with water to “uncoil” and hydrolyze into glucose
- Cellulose
 - Straight, ribbon-like chains that stack into crystals through hydrogen bonds
 - Very strongly bound (high number of H-bonds)
 - Gives cellulose strength and resistance to hydrolysis
- 3-D Models

Lipids

- Hydrocarbons (largely C & H)
- Important Functions
 - Membranes! Cells, organelles
 - Energy storage (higher energy density than carbohydrates)
- Fatty acids – carbon chains with acetyl on one end
 - Saturated: no C=C (double bonds)
 - Unsaturated: Some C=C (double bonds)
- Phospholipids – hydrophobic carbon chains with hydrophilic head

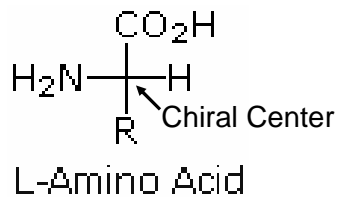
Lipids are Amphiphilic

- Hydrophilic Head – Hydrophobic Tail



Amino Acids

- 20+ Amino acids
- Amino acid polymers = proteins
- Amino acids = amine + carboxylic acid



R = side chain

R = acidic, basic, hydrophilic,
hydrophobic, charged,
uncharged

Proteins

- Proteins are linear polymers of amino acids
- Peptide bonds = amine to carboxylic acid
- Structure and function of protein determined by 3-D structure that polymer assumes in water
- 3-D structure a function of lowest energy confirmation
 - Hydrophobic groups associate away from water
 - Hydrophilic groups associate toward water
- Protein Function
 - Enzymes catalyze biochemical reactions
 - Structure and support
 - Communication between cells (hormones)

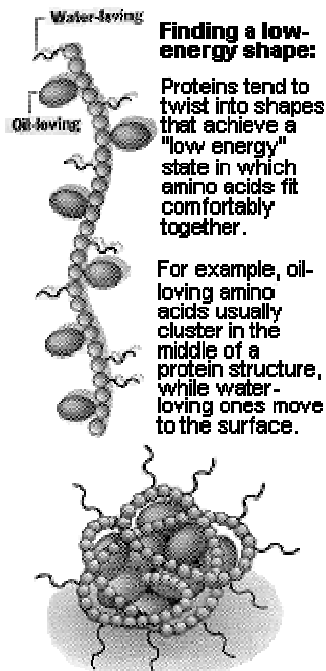
Protein Folding

Folding@Home

Distributed computing project studying protein folding using novel computational methods and large scale distributed computing, to simulate the thermodynamics of protein folding.

<http://folding.stanford.edu/>

3-D Models

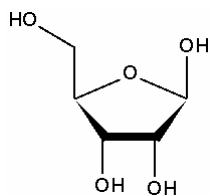


Nucleic Acids

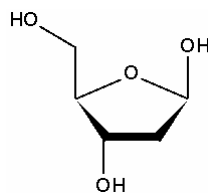
- Composed of 3 parts
 - Pentose (sugar)
 - Heterocyclic Base
 - Phosphate
- A unit of all 3 (sugar+base+phosphate) called a **nucleotide**

Nucleic Acids

- Pentose
 - Ribose (RNA)
 - Deoxyribose (DNA) = lacking 1 hydroxyl group on sugar C2



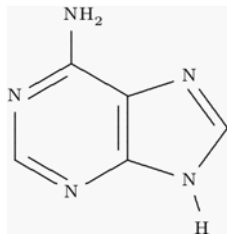
Ribose



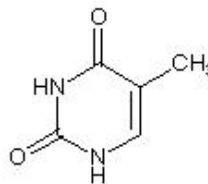
Deoxyribose

Nucleic Acids

- Bases
 - 5 bases
 - 4 used in RNA (U, A, C, G)
 - 4 used in DNA (T, A, C, G)
 - 2 Types
 - Purine (A, G)
 - Pyrimidine (T, U, C)



Adenine



Thymine

Polynucleotides

- DNA and RNA polymers store and convey genetic information
- Polymer consists of sugar backbone linked by -C-PO₄-C-
- Bases (esp. DNA) can associate through complementary pairing (A-T, C-G)
- 3-D structure is double helix
- 3-D Model

Genetic Code

- Sets of 3 bases (codon)
- $4^3 = 64$ codons
- Each codon specifies a unique amino acid
- Each amino acid specified by at least 1 codon
- Start codon (methionine)
- Stop codons (UAG, UAA and UGA)

Central Dogma of Biology

- DNA is transcribed to mRNA
- <http://www.johnkyrk.com/DNAtranscription.html>
- mRNA is translated into protein
- <http://www.johnkyrk.com/DNAtranslation.html>

- DNA makes RNA
- RNA makes Proteins