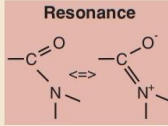


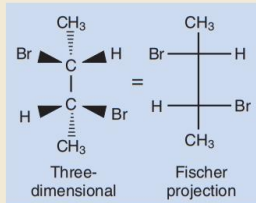
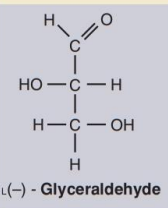
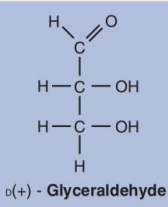
**BONDS & STRUCTURE IN ORGANIC COMPOUNDS****A. Bonding Principles**

- Most bonds are **polar covalent**; the more electronegative atom is the “-” end of the bond
Example: For $>C=O$, O is negative, C is positive
- Simplest Model: Lewis Structure:** Assign valence electrons as bonding electrons and non-bonding lone-pairs; more accurate bonding models include **valence-bonds, molecular orbitals** and **molecular modeling**
- Resonance:** The average of several Lewis structures describes the bonding
Example: The peptide bond has some $>C=N<$ character

**B. Molecular Structure****Typical Behavior of C, N & O**

Atom	sp ³	sp ²	sp
C 4 e ⁻ 4 bonds	-C-C-	>C=C<	-C≡C-
N 5 e ⁻ 3 bonds, 1 lone pair	>N-	R=N-	-C≡N
O 6 e ⁻ 2 bonds, 2 lone pairs	-O-	>R=O	

- Geometries of valence electron hybrids:** sp² - planar, sp³ - tetrahedral, sp - linear
- Isomers and structure**
 - Isomers:** same formula, **different** bonds
 - Stereoisomers:** same formula and bonds, different **spatial** arrangement
 - Chiral = optically active:** Produces + or - rotation of plane-polarized light
 - D:** Denotes dextrorotary based on clockwise rotation for glyceraldehyde
 - L:** Denotes levorotary based on counter-clockwise rotation for glyceraldehyde; **insert (-) or (+) to denote actual polarimeter results**
 - D/L** denotes **structural** similarity with **D** or **L** glyceraldehyde
 - Chiral:** Not identical with mirror image
 - Achiral:** Has a plane of symmetry
 - Racemic:** 50/50 mixture of stereoisomers is optically inactive; + and - effects cancel
 - R/S notation:** The four groups attached to the chiral atom are ranked a,b,c,d by molar mass
 - The lowest (**d**) is directed away from the viewer and the sequence of a-b-c produces clockwise (**R**) or counter-clockwise (**S**) configurations
 - This notation is less ambiguous than **D/L**; works for molecules with >1 chiral centers
- Nomenclature:** Use **D/L** (or **R/S**) and +/- in the compound name:
Example: **D** (-) lactic acid
- Fisher-projection:** Diagram for chiral compound
- Molecular conformation:** All molecules exhibit structural variation due to free rotation about **C-C** single bond; depict using a **Newman-diagram**
- Alkene:** **cis** and **trans** isomers; $>C=C<$ does not rotate; common in fatty acid side chains

**C. Common Organic Terminology**

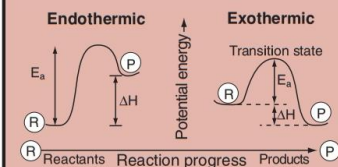
- Saturated:** Maximum # of Hs (all C-C)
- Unsaturated:** At least one $>C=C<$
- Nucleophile:** Lewis base; attracted to the + charge of a nucleus or cation
- Electrophile:** Lewis acid; attracted to the electrons in a bond or lone pair

Carbon-chain Prefixes

1 meth-	7 hept-	13 tridec-	19 nonadec-
2 eth-	8 oct-	14 tetradec-	20 eicos-
3 prop-	9 non-	15 pentadec-	22 docos-
4 but-	10 dec-	16 hexadec-	24 tetracos-
5 pent-	11 undec-	17 heptadec-	26 hexacos-
6 hex-	12 dodec-	18 octadec-	28 octacos-

REACTIONS, ENERGY & EQUILIBRIUM**A. Mechanisms**

- Biochemical reactions involve a number of simple steps that together form a **mechanism**
- Some steps may establish **equilibria**, since reactions can go forward, as well as backward; the slowest step in the mechanism, the **rate-determining step**, limits the overall reaction rate and product formation
- Each step passes through an energy barrier, the **free energy of activation** (E_a), characterized by an unstable configuration termed the **transition state** (**TS**); E_a has an **enthalpy** and **entropy** component

**B. Key Thermodynamic Variables**

- Standard conditions: 25°C, 1 atm, solutions = 1 M
- Enthalpy (H):** ΔH = heat-absorbed or produced
 $\Delta H < 0$ exothermic
 $\Delta H > 0$ endothermic

C. Standard Enthalpy of Formation, ΔH_f^0

- $\Delta H = \sum \text{prod } \Delta H_f^0 - \sum \text{react } \Delta H_f^0$
- Entropy (S):** ΔS = change in disorder
- Standard Entropy, S^0 :**
 $\Delta S = \sum \text{prod } S^0 - \sum \text{react } S^0$

4. Gibbs-Free Energy (G):

- $\Delta G = \Delta H - T\Delta S$; the capacity to complete a reaction
 $\Delta G = 0$ at equilibrium steady state $K_{eq} = 1$
 $\Delta G < 0$ **exergonic** spontaneous large K_{eq}

 $\Delta G > 0$ **endergonic**not spontaneous small K_{eq}
 $\Delta G = -RT \ln(K_{eq})$ - connection with equilibrium**D. Standard-Free Energy of Formation, ΔG_f^0 :**

- $\Delta G = \sum \text{prod } \Delta G_f^0 - \sum \text{react } \Delta G_f^0$
- For coupled reactions: Hess's Law:**
- Combine reactions, add ΔG , ΔH , ΔS
- An exergonic step can overcome an endergonic step
Example: ATP/ADP/AMP reactions are exothermic and exergonic; these provide the energy and driving force to complete less spontaneous biochemical reactions; **Example:** $ATP + H_2O \Rightarrow ADP + \text{energy}$

E. Equilibrium**1. LeChatlier's Principle**

- Equilibrium shifts to relieve the stress due to changes in reaction conditions
- K_{eq} increases: Shift equilibrium to the product side
- K_{eq} decreases: Shift equilibrium to the reactant side

2. Equilibrium and temperature changes

- For an exothermic process, heat is a product; a decrease in temperature increases K_{eq}
- For an endothermic process, heat is a reactant; an increase in temperature increases K_{eq}

3. Entropy and Enthalpy factors

- $\Delta G = \Delta H - T\Delta S$
- $\Delta H < 0$ promotes spontaneity
 - $\Delta S > 0$ promotes spontaneity
 - If $\Delta S > 0$, increasing T promotes spontaneity
 - If $\Delta S < 0$, decreasing T lessens spontaneity
Note: T is always in Kelvin;
 $K = ^\circ C + 273.15$

KINETICS: RATES OF REACTIONS**A. Determination of Rate**For a generic reaction, $A + B \Rightarrow C$:

- Reaction rate:** The rate of producing C (or consuming A or B)
- Rate-law:** The mathematical dependence of the rate on [A], [B] and [C]
- Multiple-step reaction:** Focus on **rate-determining step** - the slowest step in the mechanism controls the overall rate

B. Simple Kinetics

- First-order: Rate = $k_1[A]$
Examples: SN1, E1, aldose rearrangements
- Second order: Rate = $k_2[A]^2$ or $k_2[A][B]$
Examples: SN2, E2, acid-base, hydrolysis, condensation

C. Enzyme Kinetics

- An enzyme catalyzes the reaction of a substrate to a product by forming a

stabilized complex; the enzyme reaction may be 10^3 - 10^{15} times faster than the uncatalyzed process**2. Mechanism:**

- Step 1.** $E + S = k_1 \Rightarrow ES$
- Step 2.** $ES = k_2 \Rightarrow E + S$
- Step 3.** $ES = k_3 \Rightarrow \text{products} + E$
[E] = total enzyme concentration,
[S] = total substrate concentration,
[ES] = enzyme-substrate complex concentration, k_1 - rate ES formation, k_2 - reverse of step 1, k_3 - rate of product formation

3. Data analysis:

- Examine steady state of [ES]; rate of ES formation equal rate of disappearance
 $K_m = (k_2 + k_3)/k_1$ (Michaelis constant)
 v - reaction speed = $k_3[ES]$
 $V_{max} = k_3[E]$

Michaelis-Menten Equation:

$$v = \frac{V_{max}[S]}{K_m + [S]}$$

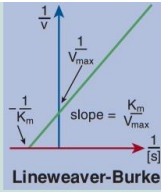


4. Practical solution:

Lineweaver-Burk approach:
 $1/v = K_m/V_{max}(1/[S]) + 1/V_{max}$
 The plot "1/v vs. 1/[S]" is linear

Slope = K_m/V_{max}
 y - intercept = $1/V_{max}$
 x - intercept = $-1/K_m$

Calculate K_m from the data



Lineweaver-Burke

D. Changing Rate Constant (k)

1. Temperature increases the rate constant:

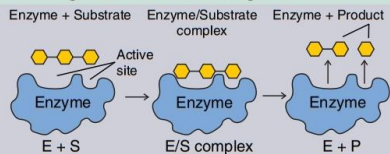
Arrhenius Law: $k = Ae^{-E_a/RT}$

• Determining E_a : Graph "ln(k) vs. 1/T"; calculate E_a from the slope

2. **Catalyst:** Lowers the activation energy; reaction occurs at a lower temperature

3. Enzymes

- Natural protein catalysts; form substrate-enzyme complex that creates a lower energy path to the product
- In addition, the enzyme decreases the **Free Energy of Activation**, allowing the product to more easily form
- Enzyme mechanism is very specific and selective; the ES complex is viewed as an "induced fit" lock-key model since the formation of the complex modifies each component



E. Energetic Features of Cellular Processes

1. **Metabolism:** The cellular processes that use nutrients to produce energy and chemicals needed by the organism

- Catabolism:** Reactions which break molecules apart; these processes tend to be **exergonic** and **oxidative**
- Anabolism:** Reactions which assemble larger molecules; biosynthesis; these processes tend to be **endergonic** and **reductive**

2. Anabolism is coupled with catabolism by ATP, NADPH and related high-energy chemicals

3. Limitations on biochemical reactions

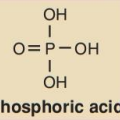
- All required chemicals must either be in the diet or be made by the body from chemicals in the diet; harmful waste products must be detoxified or excreted
- Cyclic** processes are common, since all reagents must be made from chemicals in the body
- Temperature** is fixed; activation energy and enthalpy changes cannot be too large; enzyme catalysts play key roles

ORGANIC ACIDS & BASES

	Acid	Base
Arrhenius	aqueous H ₃ O ⁺	aqueous OH ⁻
Bronsted-Lowry	proton donor	proton acceptor
Lewis	electron-pr acceptor electrophile	electron-pr donor nucleophile

A. Amphoteric

- A substance that can react as an **acid or a base**
- The molecule has acid and base functional groups; **Example:** amino acids
- This characteristic also allows amphoteric compounds to function as single-component buffers for biological studies



Phosphoric acid

B. Acids

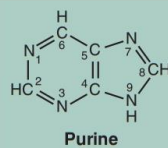
- $K_a = [A^-][H^+]/[HA]$
 $pK_a = -\log_{10}(K_a)$
- Strong acid:** Full dissociation: HCl, H₂SO₄ and HNO₃; Phosphoric acid
- Weak acid:** $K_a \ll 1$, large pK_a
- Key organic acid: **RCOOH**
Examples: Fatty acid: R group is a long hydrocarbon chain; Vitamin C is ascorbic acid; nucleic acids contain acid phosphate groups

Common Acids & pK_a

Acid	pK _a	Acid	pK _a
Acetic	4.75	Formic	3.75
Carbonic	6.35	Bicarbonate	10.33
H ₂ PO ₄ ⁻	7.21	HPO ₄ ²⁻	12.32
H ₃ PO ₄	2.16	NH ₄ ⁺	9.25

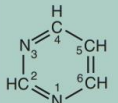
C. Organic Bases

- $K_b = [OH^-][B^+]/[BOH]$
 $pK_b = -\log_{10}(K_b)$
- Strong base:** Full dissociation: NaOH, KOH
- Weak base:** $K_b \ll 1$, large pK_b
- Organic: Amines & derivatives**
Examples: NH₃ ($pK_b = 4.74$), hydroxylamine ($pK_b = 7.97$) and pyridine ($pK_b = 5.25$)
- Purine:** Nucleic acid component: adenine (6-aminopurine) & guanine (2-amino-6-hydroxypurine)



Purine

6. **Pyrimidine:** Nucleic acid component: cytosine (4-amino-2-hydroxypyrimidine), uracil (2,4-dihydroxypyrimidine) & thymine (5-methyluracil)



Pyrimidine

D. Buffers

- A combination of a weak acid and salt of a weak acid; equilibrium between an acid and a base that can shift to **consume excess acid or base**
- Buffer can also be made from a weak base and salt of weak base
- The **pH of a buffer** is roughly equal to the pK_a of the acid, or pK_b of the base, for comparable amounts of acid/salt or base/salt
- Buffer pH is approximated by the **Henderson Hasselbalch equation**
 Note: This is for an acid/salt buffer

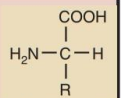
Henderson Hasselbalch Equation:
 $pH = pK_a + \log(\text{salt/acid})$

Common Buffers

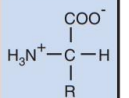
Buffer composition	approx. pH
acetic acid + acetate salt	4.8
ammonia + ammonium salt	9.3
carbonate + bicarbonate	6.3
diacid phosphate + monoacid phosphate	7.2

E. Amino Acids

- Amino acids** have amine (base) and carboxylic acid functionality; the varied chemistry arises from the chemical nature of the R- group
 • **Essential amino acids:** Must be provided to mammals in the diet
- Polymers of amino acids** form proteins and peptides
 • Natural amino acids adopt the L configuration
- Zwitterion;** self-ionization; the "acid" donates a proton to the "base"
 • **Isoelectric point, pI:** pH that produces balanced charges in the **Zwitterion**



L Amino acid



Zwitterion

MAJOR TYPES OF BIOCHEMICAL REACTIONS

Reaction Type	Reactants/Conditions	Products/Notes
Addition Nucleophilic: Electrophilic:	Add to a >C=C< Nucleophile attacks >C=O	Hydrogenate Hydrate Hydroxylate
Substitution Nucleophilic:	Replace a group on alkane (OH, NH ₂) SN1 or SN2	Amination of R-OH deamination
Elimination: E1 and E2	Reverse of addition, produce >C=C<	Dehydrogenate Dehydrate
Isomerization	Change in bond connectivity	aldose => pyranose
Oxidation- loss of e- Reduction- gain of e- Coupled Processes	Biochemical: Oxidize: Add O or remove H Reduce: Reverse of oxidize Metals: Change valence	ROH to >C=O Hydrogenate fatty acid
Hydrolysis	Water breaks a bond, add -H and -OH to form new molecules	Hydrolyze peptide, sucrose triglyceride
Condensation	R-NH or R-OH combine via bridging O or N	Form peptide or amylose

TYPES OF ORGANIC COMPOUNDS

Type of Compound	Examples	Notes
Alkane	$\begin{array}{c} & \\ -\text{C}-\text{C}- \\ & \end{array}$	ethane C ₂ H ₆ , methyl (Me) -CH ₃ , ethyl (Et) -C ₂ H ₅
Alkene	>C=C<	ethene C ₂ H ₄ , unsaturated fatty acids
Aromatic ring	-C ₆ H ₅	benzene - C ₆ H ₆ , phenylalanine
Alcohol	R-OH	methanol Me-OH, diol = glycol (2 -OH), glycerol (3 -OH)
Ether	R''-O-R'	ethoxyethane Et-O-Et, or diethyl ether
Aldehyde	$\begin{array}{c} \text{O} \\ \\ \text{R}-\text{C}-\text{H} \end{array}$	methanal H ₂ CO or formaldehyde, aldose sugars
Ketone	$\begin{array}{c} \text{O} \\ \\ \text{R}-\text{C}-\text{R}' \end{array}$	Me-CO-Me 2-propanone or acetone ketose sugars
Carboxylic acid	$\begin{array}{c} \text{O} \\ \\ \text{RC}-\text{OH} \end{array}$	Me-COOH ethanoic acid or acetic acid Me-COO ⁻ Acetate ion
Ester	$\begin{array}{c} \text{O} \\ \\ \text{RC}-\text{OR}' \end{array}$	Me-CO-OEt, ethyl acetate, Lactone: cyclic ester, Triglycerides
Amine	N-RR'R''	H ₃ C-NH ₂ , methyl amine, R-NH ₂ (1°) - primary, RR'NH (2°) - secondary, RR'R''N (3°) - tertiary
Amide	$\begin{array}{c} \text{O} \\ \\ \text{R}-\text{C}-\text{NRR}' \end{array}$	H ₃ C-CO-NH ₂ , acetamide Peptide bonds
Cyclic Ethers:	<p>Pyran Furan</p>	

BIOCHEMICAL COMPOUNDS

A. Carbohydrates: Polymers of Monosaccharides

1. **Carbohydrates** have the general formula $(CH_2O)_n$
2. **Monosaccharides**: Simple sugars; building blocks for polysaccharides

Common Sugars

Triose	3 carbon	glyceraldehyde
Pentose	5 carbon	ribose, deoxyribose
Hexose	6 carbon	glucose, galactose, fructose

a. **Aldose**: Aldehyde type structure:

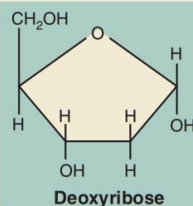
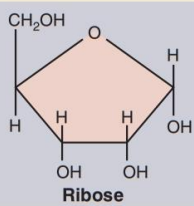
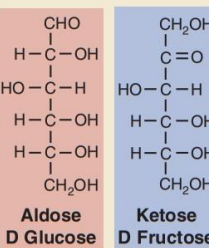


b. **Ketose**: Ketone type structure:



c. **Ribose and deoxyribose**:

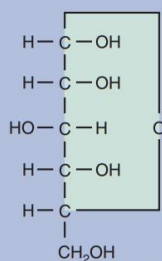
Key component in nucleic acids and ATP



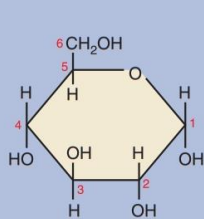
d. Monosaccharides cyclize to ring structures in water

- 5-member ring: **Furanose** (ala furan)
- 6-member ring: **Pyranose** (ala pyran)
- The ring closing creates two possible structures: α and β forms
- The carbonyl carbon becomes another chiral center (termed **anomeric**)
- α : -OH on #1 below the ring; β : OH on #1 above the ring
- Haworth figures and Fischer projections are used to depict these structures (see figure for glucose below)

Fischer Projection



Haworth Figure



α -D-Glucopyranose

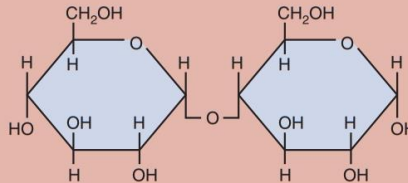
2. Polysaccharides

- a. Glucose and fructose form polysaccharides
- b. Monosaccharides in the pyranose and furanose forms are linked to form polysaccharides; dehydration reaction creates a bridging oxygen
- c. Free anomeric carbon reacts with -OH on opposite side of the ring
- d. Notation specifies form of monosaccharide and the location of the linkage; termed a **glycosidic bond**

e. Disaccharides

Disaccharide

- 2 units
- Lactose (β -galactose + β -glucose) β (1,4) link
- Sucrose (α -glucose + β -fructose) α , β (1,2) link
- Maltose (α -glucose + α -glucose) α (1,4) link



Maltose - Linked α D Glucopyranose

f. Oligosaccharides

- 2-10 units
- May be linked to proteins (glycoproteins) or fats (glycolipids)
- **Examples of functions**: cellular structure, enzymes, hormones

g. Polysaccharides

- >10 units
- Examples**:
- **Starch**: Produced by plants for storage
- **Amylose**: Unbranched polymer of α (1,4) linked glucose; forms compact helices
- **Amylopectin**: Branched amylose using α (1,6) linkage
- **Glycogen**: Used by animals for storage; highly branched polymer of α (1,4) linked glucose; branches use α (1,6) linkage
- **Cellulose**: Structural role in plant cell wall; polymer of β (1,4) linked glucose
- **Chitin**: Structural role in animals; polymer of β (1,4) linked N-acetylglucosamine

3. Carbohydrate Reactions

- a. Form polysaccharide via condensation
- b. Form glycoside: Pyranose or furanose + alcohol
- c. Hydrolysis of polysaccharide
- d. Linear forms are reducing agents; the aldehyde can be oxidized
- e. Terminal -CH₂-OH can be oxidized to carboxylic acid (uronic acid)
- f. Cyclize acidic sugar to a lactone (cyclic ester)
- g. Phosphorylation: Phosphate ester of ribose in nucleotides
- h. Amination: Amino replaces hydroxyl to form amino sugars
- i. Replace hydroxyl with hydrogen to form deoxy sugars (deoxyribose)

B. Fats and Lipids

1. **Lipid**: Non-polar compound, insoluble in water

Examples: steroids, fatty acids, triglycerides

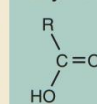
2. **Fatty acid: R-COOH**

Essential fatty acids cannot be synthesized by the body: linoleic, linolenic and arachidonic

3. Properties and structure of fatty acids:

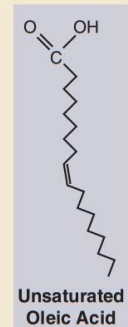
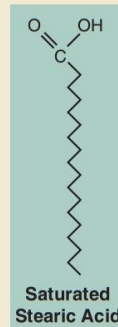
- a. **Saturated**: Side chain is an **alkane**
- b. **Unsaturated**: Side chain has at least one $>C=C<$; the name must include the position # and denote cis or trans isomer
- c. **Solubility in water**: <6 C soluble, >7 insoluble; form micelles
- d. **Melting points**: Saturated fats have higher melting points; cis-unsaturated have lower melting points

Fatty Acid



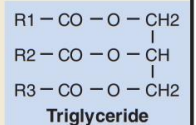
Common Fatty Acids

Common Name	Systematic	Formula
Acetic acid	ethanoic	CH ₃ COOH
Butyric	butanoic	C ₃ H ₇ COOH
Valeric	pentanoic	C ₄ H ₉ COOH
Myristic	tetradecanoic	C ₁₃ H ₂₇ COOH
Palmitic	hexadecanoic	C ₁₅ H ₃₁ COOH
Stearic	octadecanoic	C ₁₇ H ₃₅ COOH
Oleic	cis-9-octadecenoic	C ₁₇ H ₃₃ COOH
Linoleic	cis, cis-9, 12 octadecadienoic	C ₁₇ H ₃₁ COOH
Linolenic	9, 12, 15-octadecatrienoic (all cis)	C ₁₇ H ₂₉ COOH
Arachidonic	5, 8, 11, 14-eicosatetraenoic (all trans)	C ₁₉ H ₃₁ COOH



4. Common fatty acid compounds

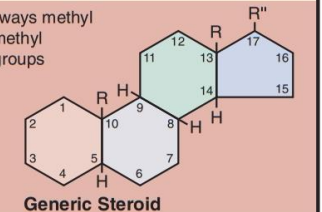
- a. **Triglyceride** or **triacylglycerol**: Three fatty acids bond via ester linkage to glycerol
- b. **Phospholipids**: A phosphate group bonds to one of three positions of fatty acid/glycerol; R-PO₄⁻ or HPO₄⁻ group



5. Examples of other lipids

- a. **Steroids**: Cholesterol and hormones
- Examples**: testosterone, estrogen

R = Nearly always methyl
R' = Usually methyl
R'' = Various groups



b. Fat-soluble vitamins:

- Vitamin A: polyunsaturated hydrocarbon, all trans
- Vitamins D, E, K

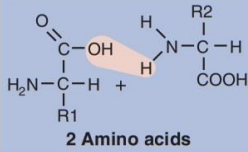
6. Lipid reactions

- a. **Triglyceride**: Three-step process: dehydration reaction of fatty acid and glycerol
- b. The reverse of this reaction is hydrolysis of the triglyceride
- c. **Phosphorylation**: Fatty acid + acid phosphate produces phospholipid
- d. **Lipase** (enzyme) breaks the ester linkage of triglyceride

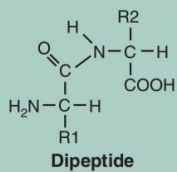


C. Proteins and Peptides - Amino Acid Polymers

1. Peptides are formed by linking amino acids; all natural peptides contain L-amino acids



a. **Dipeptide:** Two linked amino acids
 b. **Polypeptide:** Numerous linked amino acids
 c. The **peptide bond** is the linkage that connects a pair of amino acids using a dehydration reaction; the N-H of one amino acid reacting with the -OH of another => -N- bridge
 d. The dehydration reaction links the two units; each amino acid retains a reactive site



2. The nature of the peptide varies with amino acids since each R- group has a distinct chemical character
 a. R- groups end up on alternating sides of the polymer chain
 b. Of the **20 common amino acids:** 15 have neutral side chains (7 polar, 8 hydrophobic), 2 acidic and 3 basic; the variation in R- explains the diversity of peptide chemistry (see table, pg. 6)

3. **Proteins** are polypeptides made up of hundreds of amino acids
 a. Each serves a specific function in the organism
 b. The structure is determined by the interactions of various amino acids with water, other molecules in the cell and other amino acids in the protein

4. **Types of proteins:**
 a. **Fibrous:** Composed of regular, repeating helices or sheets; typically serve a structural function
Examples: keratin, collagen, silk
 b. **Globular:** Tend to be compact, roughly spherical; participates in a specific process:
Examples: enzyme, globin
 c. **Oligomer:** Protein containing several subunit proteins

Common Protein

Examples	Mol Wt	Function
fibrinogen	450,000	Physical structures
hemoglobin	68,000	Binds O ₂
insulin	5,500	Glucose metabolism
ribonuclease	13,700	Hydrolysis of RNA
trypsin	23,800	Protein digestion

5. Peptide Structure:

a. **Primary structure:** **Primary Structure**
 The linear sequence of Ala-Ala-Cys-Leu
 • Ala-Ala-Cys-Leu or A-A-C-L denotes a peptide formed from 2 alanines, a cysteine and 1 leucine
 • The order is important since this denotes the connectivity of the amino acids in the protein
 b. **Secondary structure:** Describes how the polymer takes shape
Example: Helix or pleated sheet
Factors: H-bonding, hydrophobic interactions, disulfide bridges (cysteine), ionic interactions
 c. **Tertiary structure:** The overall 3-dimensional conformation

d. **Quaternary structure:** The conformation of protein subunits in an oligomer

6. Chemical reactions of proteins:

a. Synthesis of proteins by DNA and RNA
 b. Peptides are dismantled by a hydrolysis reaction breaking the peptide bond
 c. **Denaturation:** The protein structure is disrupted, destroying the unique chemical features of the material
 d. **Agents of denaturation:** Temperature, acid, base, chemical reaction, physical disturbance

7. Enzymes

a. Enzymes are proteins that function as biological catalysts
 b. **Nomenclature:** *Substrate* + -ase
Example: The enzyme that acts on phosphoryl groups (R-PO₄) is called *phosphatase*
 8. Enzymes are highly selective for specific reactions and substrates

Six Classes of Enzymes (Enzyme Commission)

Type	Reaction
1. Oxidoreductase	Oxidation-reduction Examples: oxidize CH-OH, >C=O or CH-CH; Oxygen acceptors: NAD, NADP
2. Transferase	Functional group transfer Examples: transfer methyl, acyl- or amine group
3. Hydrolase	Hydrolysis reaction Examples: cleave carboxylic or phosphoric ester
4. Lyase	Addition reaction Examples: add to >C=C<, >C=O, aldehyde
5. Isomerase	Isomerization Example: modify carbohydrate, cis-trans fat
6. Ligase	Bond formation, via ATP Examples: form C-O, C-S or C-C

9. An enzyme may require a **cofactor**
Examples: Metal cations (Mg²⁺, Zn²⁺ or Cu²⁺); vitamins (called **coenzymes**)
 10. **Inhibition:** An interference with the enzyme structure or ES formation will *inhibit* or block the reaction
 11. **Holoenzyme:** Fully functional enzyme plus the cofactors
 12. **Apoenzyme:** The polypeptide component

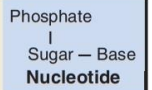
D. Nucleic Acids: Polymers of Nucleotides

1. **Nucleotide:** A phosphate group and organic base (pyrimidine or purine) attached to a sugar (ribose or deoxyribose)
 • Name derived from the base name
 • **Example:** Adenylic acid = adenosine-5'-monophosphate = 5' AMP or AMP
 2. **Nucleoside:** The base attached to the sugar
 • Nomenclature: Base name + idine (pyrimidine) or + osine (purine)
 • **Example:** adenine riboside = adenosine; adenine deoxyriboside = deoxyadenosine

Nucleic Acid Components

Base	Nucleoside	Nucleotide
adenine	Adenosine Deoxyadenosine	Adenylic acid, AMP dAMP
guanine	Guanosine Deoxyguanosine	Guanilylic acid, GMP dGMP
cytosine	Cytidine Deoxycytidine	Cytidylic acid, CMP dCMP
uracil	Uridine	Uridylic acid, UMP
thymine	Thymidine	Thymidylic acid, dTMP

3. **Cyclic nucleotides:** The phosphate group attached to the 3' position bonds to the 5' carbon 3', 5' cyclic AMP = cAMP and cGMP

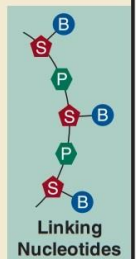


4. Additional Phosphates

a. A nucleotide can bond to 1 or 2 additional phosphate groups
 b. AMP + P => ADP - Adenosine diphosphate
 ADP + P => ATP - Adenosine triphosphate
 c. ADP and ATP function as key biochemical energy-storage compounds

5. **Glycosidic bond:** Linkage between the sugar and base involve the anomeric carbon (carbon #1)
 >C-OH (sugar) + >NH (base) => linked sugar - base

6. **Linking Nucleotides:** The polymer forms as each phosphate links two sugars; #5 position of first sugar and #3 position of neighboring sugar



7. **Types of nucleic acids:**
 Double - stranded **DNA** (deoxyribonucleic acid) and single - stranded **RNA** (ribonucleic acid)

8. **Components of a nucleotide:** sugar, base and phosphate
 a. Sugar: **ribose** (RNA) or **deoxyribose** (DNA)
 b. **Bases:** **purine** (adenine and guanine) and **pyrimidine** (cytosine, uracil (RNA) and **thymine** (DNA))

9. In DNA, the polymer strands pair to form a **double helix**; this process is tied to base pairing

10. Chargaff's Rule for DNA:

a. Adenine pairs with thymine (A: T) and guanine pairs with cytosine (C: G)
 b. Hydrogen bonds connect the base pairs and supports the helix
 c. The sequence of base pairs along the DNA strands serves as genetic information for reproduction and cellular control

P	P
S-T...A-S	
P	P
S-C...G-S	
P	P
S-G...C-S	
P	P

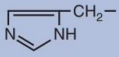

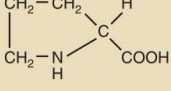
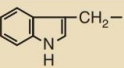
Chargaff's Rule

11. **DNA vs RNA:** DNA uses deoxyribose, RNA uses ribose; DNA uses the pyrimidine thymine, RNA uses uracil

12. Role of DNA & RNA in protein synthesis

a. DNA remains in the nucleus
 b. **Messenger-RNA (m-RNA):** Enters the nucleus and copies a three-base sequence from DNA, termed a **codon**. **m-RNA** then passes from the nucleus into the cell and directs the synthesis of a required protein on a ribosome
 c. **Transfer-RNA (t-RNA):** Carries a specific amino acid to the **ribosomal-RNA (r-RNA)** and aligns with the m-RNA codon
 d. Each codon specifies an amino acid, STOP or START; a protein is synthesized as different amino-acids are delivered to the ribosome by t-RNA, oriented by m-RNA and r-RNA, then chemically connected by enzymes



COMMON AMINO ACIDS					
hydrophobic = yellow, basic = blue, acidic = red, polar = green					
Amino acid MW essential - e		pK _a pK _b	pI R-pK _a		-R
Alanine 89.09	Ala A	2.33 9.71	6.00	hydrophobic	H ₃ C-
Arginine e 174.20	Arg R	2.03 9.00	10.76 12.10	basic	$\text{H}_2\text{N}-\overset{\text{NH}}{\parallel}{\text{C}}-\text{NH}-\text{CH}_2-\text{CH}_2-\text{CH}_2-$
Asparagine 132.12	Asn N	2.16 8.73	5.41	polar	$\text{H}_2\text{N}-\overset{\text{O}}{\parallel}{\text{C}}-\text{CH}_2-$
Aspartate 133.10	Asp D	1.95 9.66	2.77 3.71	acidic	HOOC-CH ₂ -
Cysteine 121.16	Cys C	1.91 10.28	5.07 8.14	polar	HS-CH ₂ -
Glutamate 147.13	Glu E	2.16 9.58	3.22 4.15	acidic	HOOC-CH ₂ -CH ₂ -
Glutamine 146.15	Gln Q	2.18 9.00	5.65	polar	$\text{H}_2\text{N}-\overset{\text{O}}{\parallel}{\text{C}}-\text{CH}_2-\text{CH}_2-$
Glycine 75.07	Gly G	2.34 9.58	5.97	polar	-H
Histidine e 155.16	His H	1.70 9.09	7.59 6.04	basic	
Isoleucine e 131.18	Ile I	2.26 9.60	6.02	hydrophobic	$\text{CH}_3-\text{CH}_2-\overset{\text{HC}}{\underset{\text{CH}_3}{\diagdown}}-$
Leucine e 131.18	Leu L	2.32 9.58	5.98	hydrophobic	$\text{CH}_3-\text{CH}_2-\overset{\text{HC}}{\underset{\text{CH}_3}{\diagdown}}-\text{CH}_2-$
Lysine e 146.19	Lys K	2.15 9.16	9.74 10.67	basic	$\text{H}_2\text{N}-\text{CH}_2-\text{CH}_2-\text{CH}_2-\text{CH}_2-$
Methionine e 149.21	Met M	2.16 9.08	5.74	hydrophobic	CH ₃ -S-CH ₂ -CH ₂ -
Phenylalanine e 165.19	Phe F	2.18 9.09	5.48	hydrophobic	
Proline 115.13	Pro P	1.95 10.47	6.30	hydrophobic	
Serine 105.09	Ser S	2.13 9.05	5.68	polar	HO-CH ₂ -
Threonine e 119.12	Thr T	2.20 8.96	5.60	polar	$\text{CH}_3-\overset{\text{CH}}{\underset{\text{OH}}{\diagdown}}-$
Tryptophan e 204.23	Trp W	2.38 9.34	5.89	hydrophobic	
Tyrosine 181.19	Tyr Y	2.24 9.04	5.66 10.10	polar	HO-C ₆ H ₄ -CH ₂ -
Valine - e 117.15	Val V	2.27 9.52	5.96	hydrophobic	$\text{CH}_3-\overset{\text{HC}}{\underset{\text{CH}_3}{\diagdown}}-$

AMINO ACID RNA CODONS
• Phe UUU UUC
• Thr ACU ACC ACA ACG
• Lys AAA AAG
• Leu UUA UUG CUU CUC CUA CUG
• Ala GCU GCC GCA GCG
• Asp GAU GAC
• Glu GAA GAG
• Ile AUU AUC AUA
• Tyr UAU UAC
• Cys UGU UGC
• Met START AUG
• STOP UAA UAG UGA
• Trp UGG
• Val GUU GUC GUA GUG
• His CAU CAC
• Arg CGU CGC CGA CGG AGA AGG
• Ser UCU UCC UCA UCG
• Gln CAA CAG
• Ser AGU AGC
• Pro CCU CCC CCA CCG
• Asn AAU AAC
• Gly GGU GGC GGA GGG

ABBREVIATIONS USED IN BIOLOGY & BIOCHEMISTRY			
aa	amino acid	Lys	aa lysine
A	aa alanine adenine - purine base	M	aa methionine Molar (moles/L)
Ala	aa alanine	m	milli (10 ⁻³)
ADP	adenosine diphosphate	Man	mannose sugar
AMP	adenosine monophosphate	Met	aa methionine
Arg	aa arginine	mL	milliliter
Asn	aa asparagine	mm	millimeter
Asp	aa aspartate	N	aa asparagine
atm	atmosphere (pressure unit)		Avogadro's number elemental nitrogen
ATP	adenosine triphosphate	n	nano (10 ⁻⁹)
C	aa cysteine cytosine - pyrimidine elemental carbon	O	orotidine elemental oxygen
cal	calorie	P	aa proline phosphate group elemental phosphorous
Cys	aa cysteine	p	pico (10 ⁻¹²)
D	aa aspartate Dalton	Phe	aa phenylalanine
DNA	deoxyribonucleic acid	Pro	aa proline
dRib	2-deoxyribose sugar	Q	aa glutamine coenzyme Q, ubiquinone
E	aa glutamate	R	aa arginine gas constant
F	aa phenylalanine	Rib	ribose sugar
Fru	fructose sugar	RNA	ribonucleic acid
G	aa glycine guanine - purine base	S	aa serine Svedberg unit
Gal	galactose sugar	s	second (unit)
Glc	glucose sugar	Ser	aa serine
Glu	aa glutamate	T	aa threonine thymine - pyrimidine absolute temperature
H	aa histidine	Thr	aa threonine
h	hour Planck's constant	Trp	aa tryptophan
His	aa histidine	Tyr	aa tyrosine
I	aa isoleucine inosine elemental iodine	U	uracil - pyrimidine
Ile	aa isoleucine	V	aa valine volt (electrical potential)
J	Joule (energy unit)	Val	aa valine
K	aa lysine Kelvin - absolute T elemental potassium	W	aa tryptophan elemental tungsten
k	kilo (10 ³)	X	xanthine
L	aa leucine liter (volume)	Y	aa tyrosine
Lac	lactose sugar	yr	year
Leu	aa leucine		