

"Enzymes"

I. Overview

- Virtually all body reactions are mediated by enzymes.
- Enzymes are protein catalysts, usually intracellular.
- They increase reaction rates without being changed in the overall process.
- Among many energetically possible biologic reactions, enzymes selectively channel substrates into useful pathways.
- Enzymes direct all metabolic events.
- This chapter examines the nature and mechanisms of enzymes.

II. Nomenclature

A. Enzyme Names

- Each enzyme has two names:
 - a. Recommended name (short, everyday use)
 - b. Systematic name (complete, unambiguous identification)

B. Recommended Name

- Most enzyme names end with suffix “-ase”, attached to the substrate, e.g.,
 - Glucosidase
 - Urease
- Some names describe the action performed, e.g.,
 - Lactate dehydrogenase (LDH)
 - Adenylyl cyclase
- Some enzymes retain trivial names without indication of reaction, e.g.,
 - Trypsin
 - Pepsin

C. Systematic Name

- Systematic names reflect the chemical reaction catalyzed, including all substrate names.

- The suffix “-ase” is attached to a full description, e.g.,
 - Lactate:nicotinamide adenine dinucleotide (NAD⁺) oxidoreductase
- Each enzyme is assigned a classification number, e.g.,
 - Lactate:NAD⁺ oxidoreductase = 1.1.1.27
- Systematic names are unambiguous and informative but often cumbersome for general use.
- Enzymes are divided into six major classes, each with numerous subgroups.

II. Potentially Confusing Enzyme Nomenclature

- Enzymes with similar names but different functions/mechanisms:
 - Synthetases
 - Require ATP for their action.
 - Synthases
 - Do not require ATP.
 - Phosphatases
 - Use water to remove a phosphate group.

- Phosphorylases
 - Use inorganic phosphate to break a bond and generate a phosphorylated product.
- Dehydrogenases
 - Use NAD^+ or flavin adenine dinucleotide (FAD) as electron acceptors in redox reactions.
- Oxidases
 - Use oxygen as electron acceptor without incorporating oxygen atoms into the substrate.
- Oxygenases
 - Incorporate oxygen atoms into their substrates.

III. Properties

- An enzyme is an efficient, specific protein catalyst.
- It combines with a substrate at the enzyme active site.
- Performs chemistry on the substrate to convert it to product.

- Without enzymes, most biochemical reactions would be too slow to have physiologic importance in the human body.
- Enzymes increase the velocity of chemical reactions but are not consumed during the reaction.
- Some RNAs can catalyze reactions affecting phosphodiesterase and peptide bonds.
 - RNAs with catalytic activity are called ribozymes.
 - Ribozymes are much less common than protein catalysts.

A. Active Site

- Enzyme molecules contain a special pocket or cleft called the active site.
- The active site is formed by folding of the protein.
- Contains amino acid residues whose side chains participate in:
 - Substrate binding
 - Catalysis

- The substrate binds the enzyme forming an enzyme-substrate (ES) complex.
- Binding causes a conformational change in the enzyme (induced fit model).
- This allows rapid conversion of ES to enzyme-product (EP) complex.
- EP complex subsequently dissociates into free enzyme + product.

B. Efficiency

- Enzyme-catalyzed reactions proceed 10^3 to 10^8 times faster than uncatalyzed reactions.
- The number of substrate molecules converted to product per enzyme molecule per second is the turnover number (k_{cat}).
- Typical k_{cat} values are 10^2 to 10^4 s^{-1} .
- Note: k_{cat} is the rate constant for conversion of ES to E + P.

C. Specificity

- Enzymes are highly specific.
- Capable of interacting with one or very few substrates.
- Can catalyze only one type of chemical reaction.
- The set of enzymes synthesized within a cell determines which reactions occur in that cell.

D. Holoenzymes, Apoenzymes, Cofactors, and Coenzymes

- Some enzymes require nonprotein components for enzymatic activity.
- Holoenzyme = protein component + nonprotein component (active enzyme).
- Apoenzyme = enzyme without its nonprotein moiety (inactive enzyme).
- Nonprotein components must be present for catalytic function.

- Types of nonprotein components:
 - Cofactor: metal ion (e.g., zinc Zn^{2+} , iron Fe^{2+}).
 - Coenzyme: small organic molecule.
- Coenzymes (cosubstrates):
 - Bind transiently to the enzyme.
 - Dissociate in an altered state (example: NAD^+).
- Prosthetic group:
 - Coenzyme permanently bound to enzyme.
 - Returned to original form after reaction (example: FAD).
- Coenzymes are often vitamin-derived:
 - NAD^+ contains niacin.
 - FAD contains riboflavin.

E. Regulation

- Enzyme activity can be increased or decreased.
- Allows rate of product formation to match cellular needs.

F. Location within the Cell

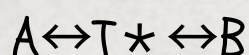
- Most enzymes act inside cells, confined by plasma membranes.
- Many enzymes are localized to specific organelles
- Compartmentalization benefits:
 - Isolates substrate/product from competing reactions.
 - Creates a favorable environment for reactions.
 - Organizes enzymes into specific metabolic pathways.

IV. Mechanism of Enzyme Action

- Enzyme action can be understood from two perspectives:
 - Energy changes during the reaction.
 - Chemical facilitation by the active site.

A. Energy Changes Occurring During the Reaction

- All chemical reactions have an energy barrier between reactants and products called the activation energy (E_a).
- Activation energy (E_a):
 - Energy difference between reactants and a high-energy intermediate called the transition state (T^*).
 - Transition state is a short-lived, high-energy intermediate formed during conversion from reactant to product.
- Reaction pathway:



where A = reactant, T^* = transition state, B = product.

- Activation energy significance:
 - E_a represents the peak of energy on the reaction coordinate.
 - High E_a causes slow rates for uncatalyzed reactions.

- Rate of reaction:
 - Molecules must have sufficient energy to overcome E_a to react.
 - Without enzymes, only a small fraction of molecules reach the transition state.
 - The rate depends on the number of molecules with enough energy to surpass E_a .
 - Lower $E_a \rightarrow$ more molecules can cross transition state \rightarrow faster reaction rate.

3. Alternate Reaction Pathway

- Enzymes provide an alternate reaction pathway with a lower activation energy (E_a).
- This allows reactions to proceed rapidly under cellular conditions.
- Enzymes do not change the free energy of reactants or products.
- Therefore, enzymes do not alter the equilibrium of the reaction.
- Enzymes accelerate the rate at which equilibrium is reached.

B. Active Site Chemistry

- The active site is a complex molecular machine, not just a substrate binding pocket.
- It uses diverse chemical mechanisms to facilitate substrate \rightarrow product conversion.
- Factors contributing to catalytic efficiency include:

1. Transition-State Stabilization

- Active site acts as a flexible molecular template binding substrate and initiating conversion to the transition state (T^*).
- Transition state has bond structures different from substrate or product.
- By stabilizing the transition state, the enzyme increases the concentration of the reactive intermediate.
- This stabilization accelerates the reaction.
- Note: The transition state cannot be isolated experimentally.

2. Catalysis

- The active site provides catalytic groups that increase the probability of transition state formation.
- Some enzymes use general acid-base catalysis: amino acid residues donate or accept protons.
- Other enzymes catalyze via transient covalent enzyme-substrate (ES) complexes.
- Example: Chymotrypsin (protein digestion enzyme in intestine) uses:
 - General base catalysis: Histidine gains a proton.
 - General acid catalysis: Histidine loses a proton.
 - Covalent catalysis: Serine forms a transient covalent bond with substrate.
- Histidine's pK near physiologic pH allows it to switch between protonated and deprotonated states.

3. Transition-State Visualization

- Enzyme-catalyzed substrate \rightarrow product conversion likened to removing a sweater from an uncooperative infant.

- High activation energy because both arms must be fully extended over the head, an unlikely substrate conformation without enzyme.
- Enzyme acts like a parent:
 - Binds substrate (forming ES complex).
 - Guides substrate to adopt extended arm posture (transition state).
- This conformational change facilitates reaction to product (disrobed baby).
- Note: ES complex is at a slightly lower energy than free substrate (explains small dip in energy curve at ES).

V. Factors Affecting Reaction Velocity

Enzymes show different responses to substrate concentration, temperature, and pH—both *in vitro* and *in vivo*.

A. Substrate Concentration

I. Maximal Velocity (V_{max})

- Reaction velocity (v) = number of substrate molecules converted to product per unit time.
 - Expressed as: μmol of product/second.
- As substrate concentration increases, the reaction rate increases — up to a point.
- At high $[S]$, all enzyme active sites are occupied, so the enzyme becomes saturated.
- Once saturation occurs, further increases in $[S]$ do not increase the rate \rightarrow velocity reaches V_{max} .
- V_{max} reflects the point where all enzyme molecules are bound to substrate.

2. Shape of the Enzyme Kinetics Curve

- Most enzymes follow Michaelis-Menten kinetics:
 - Plot of initial velocity (v_0) vs. substrate concentration = hyperbolic curve (like myoglobin's O_2 curve).
- Allosteric enzymes show a sigmoidal curve (like hemoglobin's O_2 curve):
 - Indicates cooperative binding and regulation.

B. Temperature

1. Velocity Increases with Temperature

- As temperature increases, more substrate molecules have sufficient energy to overcome activation energy (E_a).
- Velocity increases until it reaches a peak.

2. Velocity Decreases at High Temperature

- Higher temperatures denature enzymes, decreasing activity.
- Optimum for human enzymes: $\sim 35^{\circ}\text{C}$ – 40°C .
 - Start to denature $>40^{\circ}\text{C}$.
- Thermophilic bacterial enzymes: optimum $\approx 70^{\circ}\text{C}$.

C. pH

1. Effect on Active Site Ionization

- Proper ionization of functional groups (e.g. $-\text{NH}_3^+$) is critical.
- pH affects binding and catalysis.
 - E.g. deprotonation at high pH \rightarrow loss of activity.

2. Effect on Enzyme Denaturation

- Extremely low or high pH disrupts ionic interactions \rightarrow denatures protein structure.

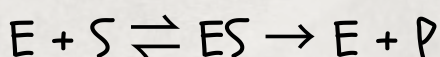
3. Variable pH Optimum

- Each enzyme has a specific pH for optimal activity, depending on its location:
 - Pepsin (stomach) → optimal at pH 2.
 - Neutral pH enzymes → inactive in acidic pH.

VI. Michaelis-Menten Kinetics:

- Proposed in 1913 by Leonor Michaelis and Maud Menten.
- Explains most enzyme-catalyzed reactions involving a single substrate.

Reaction Model



- E = Enzyme
- S = Substrate
- ES = Enzyme-substrate complex
- P = Product
- k_1 = rate constant for $E + S \rightarrow ES$
- k_{-1} = rate constant for $ES \rightarrow E + S$
- k_2 (or k_{cat}) = rate constant for $ES \rightarrow E + P$

A. Michaelis-Menten Equation

$$v_0 = (V_{\max} \times [S]) / (K_m + [S])$$

- v_0 = Initial velocity of the reaction
- V_{\max} = Maximum velocity = $k_{\text{cat}} \times [E]_{\text{total}}$
- K_m = Michaelis constant = $(k_{-1} + k_2) / k_1$
- $[S]$ = Substrate concentration

Assumptions of Michaelis-Menten Model

1. Enzyme and Substrate Relative Concentrations

- $[S] \gg [E]$, so only a small fraction of substrate is enzyme-bound.

2. Steady-State Assumption

- Concentration of ES complex remains constant during the reaction.
- Rate of ES formation = rate of ES breakdown.
- Applies to intermediates in any pathway where synthesis rate = degradation rate.

3. Initial Velocity (v_0)

- v_0 is measured immediately after mixing enzyme and substrate.
- At this early stage, product concentration is negligible, so reverse reaction ($P \rightarrow S$) can be ignored.
- Initial velocity allows clearer analysis of forward reaction kinetics.

B. Important Conclusions from Michaelis-Menten Kinetics

1. K_m Characteristics (Michaelis constant)

- K_m is unique for each enzyme-substrate pair.
- Reflects enzyme's affinity for the substrate.
- Defined as the substrate concentration at which $v = \frac{1}{2} V_{max}$.
- K_m is independent of enzyme concentration.

a. Low K_m

- Indicates high affinity \rightarrow enzyme binds substrate easily.
- Low $[S]$ is sufficient to reach half V_{max} .

b. High K_m

- Indicates low affinity \rightarrow more $[S]$ needed for half-saturation.

2. Velocity Relation to Enzyme Concentration

- When $[S]$ is not limiting, $v_0 \propto [E]$.
- Halving $[E] \rightarrow$ both v_0 and V_{max} are halved.

3. Reaction Order

a. First-Order Kinetics

- When $[S] \ll K_m$, velocity $\propto [S]$.
- Reaction is first-order with respect to substrate.

b. Zero-Order Kinetics

- When $[S] \gg K_m$, velocity = V_{max} .
- Reaction is zero-order \rightarrow independent of $[S]$ due to enzyme saturation.

c. Lineweaver-Burk Plot (Double-Reciprocal Plot)

- Developed in 1934 by Hans Lineweaver and Dean Burk.
- Overcomes limitations of hyperbolic curve by using reciprocals.

Lineweaver-Burk Equation:

$$1/v_0 = (K_m/V_{max}) \times (1/[S]) + 1/V_{max}$$

- Plot: $1/v_0$ vs $1/[S]$ \rightarrow yields a straight line.
- x-intercept = $-1/K_m$
- y-intercept = $1/V_{max}$
- Slope = K_m/V_{max}
- Useful for:
 - Calculating K_m and V_{max}
 - Identifying types of enzyme inhibition

VII. Enzyme Inhibition

Definition

- Any substance that decreases the velocity of an enzyme-catalyzed reaction.
- Two major types:
 - Irreversible inhibition
 - Reversible inhibition

A. Irreversible Inhibition

- Inhibitor forms covalent bonds with the enzyme.
- Permanent inactivation of enzyme function.
- Example: Lead (Pb^{2+})
 - Covalently binds to the sulfhydryl ($-\text{SH}$) group of cysteine residues.
 - Ferrochelatase, an enzyme in heme synthesis, is irreversibly inhibited by lead.

B. Reversible Inhibition

- Inhibitor binds via noncovalent interactions (e.g., hydrogen bonds, ionic bonds).
- Can form enzyme-inhibitor complex (EI).
- Enzyme activity can be restored by dilution or removal of inhibitor.
- Two major types:
 - Competitive inhibition
 - Noncompetitive inhibition

A. Competitive Inhibition

- Inhibitor competes with the substrate for the active site.
- Binds reversibly to the same site as the substrate.

I. Effect on V_{max}

- V_{max} remains unchanged.
- Because high $[S]$ can outcompete the inhibitor and restore full enzyme activity.

2. Effect on K_m

- Apparent K_m increases.
- More substrate is required to reach $\frac{1}{2} V_{max}$, indicating a decrease in enzyme affinity for the substrate in the presence of inhibitor.

3. Effect on Lineweaver-Burk Plot

- Inhibited and uninhibited plots intersect on the Y-axis at $1/V_{max} \rightarrow V_{max}$ is unchanged.
- The X-intercept changes:
 - Inhibited plot shifts leftward ($-1/K_m$ moves closer to zero).
 - Indicates an increase in apparent K_m .

- Competitive inhibitors increase slope of Lineweaver-Burk plot:

➤ Slope = K_m/V_{max} ↑

- Transition state analogs:
 - Stable molecules that mimic the transition state.
 - Bind more tightly than the actual substrate.
 - Act as potent competitive inhibitors.

4. Example: Statin Drugs

- Statins (e.g., atorvastatin, pravastatin) are competitive inhibitors.
- Inhibit HMG-CoA reductase, the rate-limiting enzyme in cholesterol synthesis.
- They are structural analogs of the natural substrate.
- Reduce de novo cholesterol synthesis → lower blood cholesterol levels.

B. Noncompetitive Inhibition

Mechanism

- Inhibitor binds to a site different from the substrate-binding site.
- Can bind:
 - To free enzyme (E)
 - Or to the enzyme-substrate complex (ES)
- Binding prevents the catalytic activity, regardless of whether the substrate is bound.

Effect on V_{max} and K_m

- V_{max} decreases → enzyme's overall capacity is reduced.
- K_m remains unchanged → substrate binding affinity is not affected.
- Lineweaver-Burk plot:
 - Same X-intercept ($-1/K_m$)
 - Higher Y-intercept ($1/V_{max}$)

1. Effect on V_{max}

- V_{max} decreases.
- Effect cannot be reversed by increasing substrate concentration.
- Indicates reduced overall enzyme catalytic capacity.

2. Effect on K_m

- K_m remains unchanged.
- Substrate binding is not affected, as inhibitor binds elsewhere (not at active site).
- Enzyme shows same affinity for substrate in presence of inhibitor.

3. Effect on Lineweaver-Burk Plot

- Plots show:
 - Same X-intercept ($-1/K_m$) → K_m unchanged
 - Higher Y-intercept ($1/V_{max}$) → V_{max} decreased

- Useful to distinguish noncompetitive inhibition from competitive inhibition.

C. Enzyme Inhibitors as Drugs

1. β -Lactam Antibiotics

- Include penicillin, amoxicillin.
- Inhibit bacterial enzymes involved in cell wall synthesis.

2. ACE Inhibitors

- Drugs like captopril, enalapril, lisinopril.
- Inhibit angiotensin-converting enzyme (ACE).
- Prevent conversion of angiotensin I \rightarrow angiotensin II.
- Result: Vasodilation and lower blood pressure.

3. Aspirin

- A nonprescription drug.
- Irreversibly inhibits cyclooxygenase (COX enzyme).
- Blocks prostaglandin and thromboxane synthesis.
- Results in anti-inflammatory and antiplatelet effects.

VIII. Enzyme Regulation

Importance of Regulation

- Enzyme regulation is essential for coordinating metabolic processes.
- Most enzymes respond to changes in substrate concentration.
 - Substrate levels often near K_m .
 - $\uparrow [\text{Substrate}] \rightarrow \uparrow \text{reaction rate} \rightarrow \text{helps normalize substrate levels.}$

Specialized Regulatory Mechanisms

- Some enzymes respond to:
 - Allosteric effectors
 - Covalent modification
 - Altered synthesis/degradation rates in response to physiological changes.

A. Allosteric Enzymes

Basic Characteristics

- Do not follow Michaelis-Menten kinetics.
- Regulated by effectors (bind noncovalently at sites other than active site).
- Usually multimeric (multiple subunits).
- Effector-binding site \neq catalytic site.

Types of Effectors

- Positive effectors: Increase enzyme activity.
- Negative effectors: Decrease enzyme activity.

Effects of Effectors

- Modify:
 - Affinity for substrate ($K_{0.5}$)
 - Catalytic activity (V_{max})
 - Or both

Role in Metabolic Pathways

- Often catalyze:
 - Committed step in a pathway.
 - Rate-limiting step.

1. Homotropic Effectors

- Substrate itself acts as the effector.
- Example: Cooperative binding seen in some enzymes (like hemoglobin* – although not an enzyme, it shows similar behavior).
- Term "homotropic" = same as substrate.

Cooperativity in Allosteric Enzymes

- Substrate often acts as a positive effector.
- Binding of substrate to one site enhances catalytic activity at other sites.
- This interaction between sites is called cooperativity.
- Leads to a sigmoidal (S-shaped) v_0 vs. $[S]$ curve.
 - In contrast to hyperbolic curve seen with Michaelis-Menten enzymes.
- Analogy: Similar to O_2 binding to hemoglobin.

2. Heterotropic Effectors

- Effector \neq substrate \rightarrow called heterotropic.

- Example: Feedback inhibition:
 - End product G binds to an allosteric site of an earlier enzyme (e.g., enzyme converting $D \rightarrow E$).
 - High $[G] \rightarrow$ inhibits the first irreversible step unique to the pathway.
 - Regulates product levels by controlling pathway flow.
- Common Example:
 - Phosphofructokinase-1 (PFK-1) in glycolysis:
 - Inhibited by citrate (not a substrate of PFK-1).
 - Citrate acts as a heterotropic inhibitor.

B. Covalent Modification

- Many enzymes are regulated by addition/removal of phosphate groups.
- Modification usually occurs on serine, threonine, or tyrosine residues.
- Protein phosphorylation is a key method of cellular regulation.

1. Phosphorylation and Dephosphorylation

- Phosphorylation catalyzed by protein kinases:
 - Add phosphate group to enzyme/protein.
 - Use ATP as phosphate donor.
- Dephosphorylation catalyzed by phosphoprotein phosphatases:
 - Remove phosphate groups.

2. Enzyme Response to Phosphorylation

- Effect of phosphorylation varies by enzyme:
 - Some become more active, others less active.
- Examples:
 - Glycogen phosphorylase (breaks down glycogen):
 - Activity increased by phosphorylation.
 - Glycogen synthase (synthesizes glycogen):
 - Activity decreased by phosphorylation.

C. Enzyme Synthesis

- Regulation can occur by changing enzyme quantity, not just activity.
- Achieved by altering the rate of enzyme synthesis or degradation.

Key Concepts:

- Induction: Increased enzyme synthesis.
- Repression: Decreased enzyme synthesis.
- Alters total number of active sites.

Example:

- High blood glucose → High insulin:
 - ↑ Synthesis of key glucose metabolism enzymes.

Additional Notes:

- Enzymes regulated by synthesis are typically:
 - Required only during certain stages or specific conditions.

- Housekeeping enzymes:
 - Used constantly → not typically regulated this way.
- Synthesis-based regulation is slow:
 - Takes hours to days.
 - In contrast, allosteric and covalent regulation acts in seconds to minutes.

IX. Enzymes In Human Blood

Overview:

- Most enzymes function intracellularly, but some are found in extracellular fluids like blood plasma.
- Plasma contains enzymes from two main sources:

A. Types of Enzymes Found in Blood Plasma

I. Actively Secreted Enzymes:

- A small group secreted into blood by specific cells.

- Example: Liver secretes zymogens (inactive precursors) of protease enzymes for blood coagulation.
 - These become activated in blood to perform enzymatic functions.

2. Enzymes Released from Cells During Normal Turnover:

- Not secreted purposefully — result of routine cell death.
- Are normally intracellular and nonfunctional in plasma.
- Their levels remain constant in healthy individuals:
 - Steady state: Rate of release = Rate of clearance.
- Elevated plasma enzyme levels suggest:
 - Tissue damage or increased cell death beyond normal turnover.

Additional Definitions:

- Blood plasma: Fluid, noncellular portion of blood.
- Serum:
 - Obtained after coagulation and centrifugation of whole blood.
 - Used in most laboratory enzyme activity assays.
- Plasma vs. Serum:
 - Plasma = Physiologic body fluid.
 - Serum = Laboratory-prepared fluid (no clotting factors).

A. Blood Plasma Enzyme Levels in Disease States

- Tissue damage → Cell lysis → Release of intracellular enzymes into plasma.
- These enzymes do not function outside the cell but are used for diagnosis.

- Clinical importance:
 - Enzyme levels correlate with:
 - Extent of tissue damage.
 - Therapy effectiveness.
 - Prognosis.

B. Plasma Enzymes as Diagnostic Tools

- Some enzymes are tissue-specific → Useful for localizing tissue damage.

Example:

- Alanine aminotransferase (ALT):
 - High in liver.
 - Elevated plasma ALT = Possible liver damage.
 - Included in liver function tests.
- Enzymes with wide tissue distribution:
 - Less specific for diagnosis.
 - Only indicate general cell injury, not the tissue source.

C. Isoenzymes

Definition:

- Isoenzymes (isozymes) = Variant forms of the same enzyme.
- Same catalytic activity, but differ in:
 - Amino acid sequence (genetically determined).
 - Physical properties (e.g., charge, structure).
- Due to differences in charged amino acids, they can be separated via:
 - Electrophoresis – movement in an electric field.

I. Clinical Importance of Isoenzymes:

- Different tissues have characteristic proportions of isoenzymes.
- Pattern of isoenzymes in blood plasma helps identify tissue damage.

Examples:

- LDH (Lactate Dehydrogenase):
 - Found in many tissues.
 - Has five isoenzymes: LD1-LD5
 - LD1: Myocardial muscle
 - LD2: Red blood cells
 - LD5: Liver, skeletal muscle
- CK (Creatine Kinase):
 - Isoenzymes: CK1 (BB), CK2 (MB), CK3 (MM)
 - Distribution:
 - CK1 (BB): Brain
 - CK2 (MB): Myocardium (cardiac-specific!)
 - CK3 (MM): Skeletal muscle

2. Isoenzyme Quaternary Structure

LDH Isoenzymes:

- All are tetramers (4 subunits).
- Two types of subunits: H (heart) and M (muscle).

- Combinations:

- LD1 = HHHH
- LD2 = HHHM
- LD3 = HHMM
- LD4 = HMMM
- LD5 = MMMM

CK Isoenzymes:

- All are dimers (2 subunits).
- Two types of subunits: B (brain) and M (muscle).
- Combinations:
 - CK1 = BB
 - CK2 = MB (unique to cardiac muscle)
 - CK3 = MM

Each isoenzyme has a characteristic electrophoretic mobility.

3. Historical Use in Diagnosis of Myocardial Infarction (MI)

- Before troponins, CK MB (CK2) was used as a biomarker for MI.
- CK MB is specific to myocardium:
 - Only tissue with >5% total CK activity as CK MB.
 - Its presence in blood = Myocardial damage.

CK MB Timeline Post-MI:

- Appears in plasma: 4-8 hours after chest pain.
- Peaks: ~24 hours.
- Used in early MI detection before troponin testing became standard.

Clinical Application: Diagnostic Use of Troponins

Troponins:

- Regulatory proteins in muscle contractility.
- Cardiac-specific isoforms: cTnT (Troponin T) and cTnI (Troponin I).

cTn Characteristics:

- Released into plasma after cardiac muscle damage.
- Highly sensitive and specific for myocardial injury.

Timeline of cTn Post-MI:

- Appear: 4-6 hours after onset.
- Peak: 24-36 hours.
- Remain elevated: 3-10 days.

Clinical Use:

- Considered the "gold standard" for MI diagnosis.
- Interpretation includes:
 - Elevated cTn
 - Clinical symptoms
 - ECG changes

Note: Though similar in timing to CK MB, cTn shows a greater change from baseline, making it more reliable.