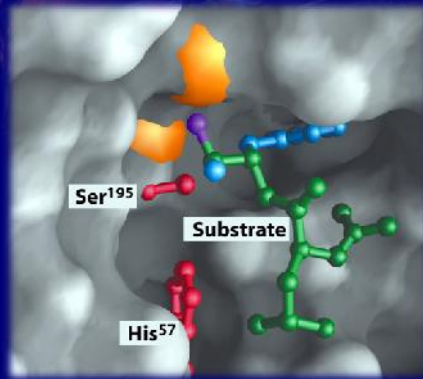
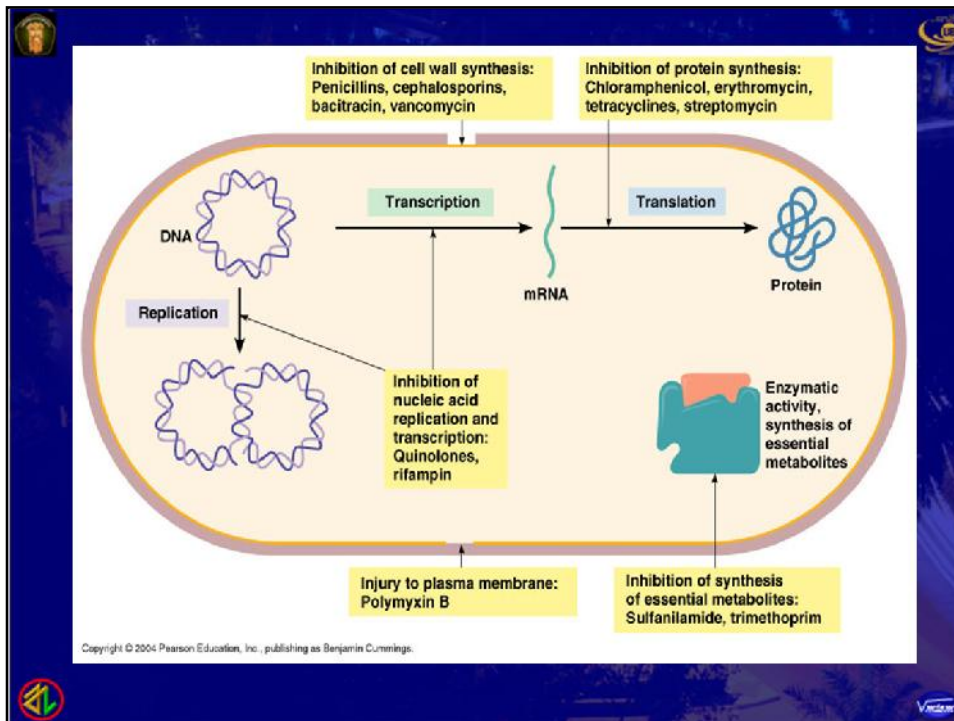
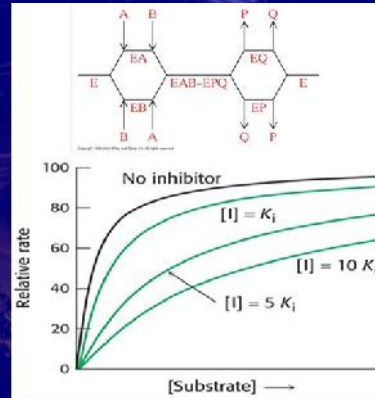


LECTURE 4: REACTION MECHANISM & INHIBITORS



Chymotrypsin



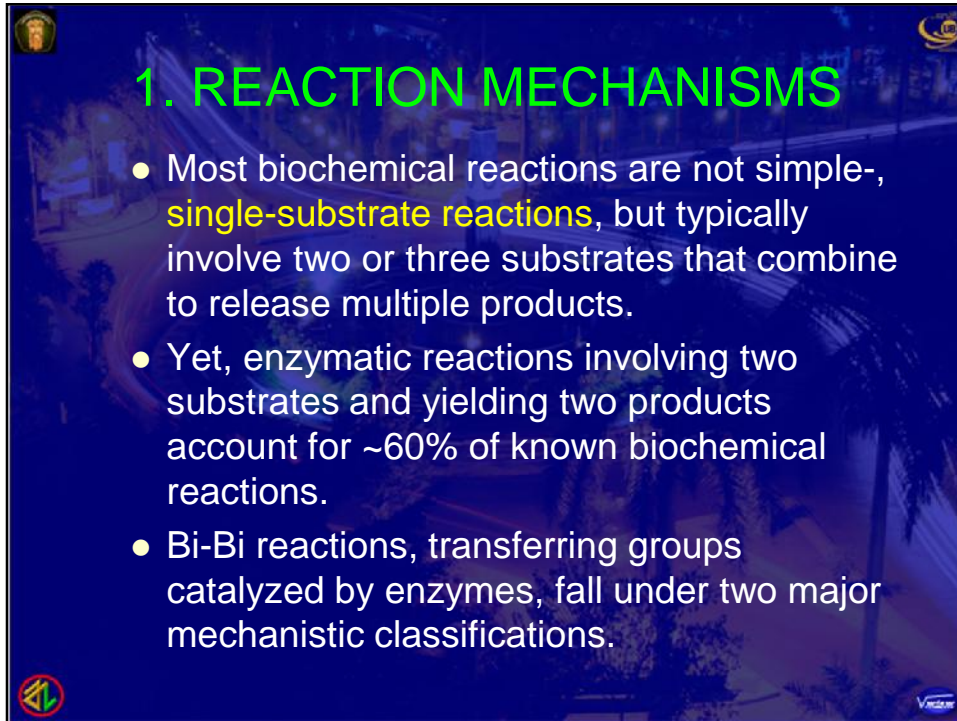
LECTURE OUTCOMES

After mastering the present lecture materials, students will be able to

1. to explain reaction mechanisms of between enzyme and substrate
2. to explain the influence of irreversible inhibitors on enzymatic reactions
3. to explain the influence of reversible inhibitors on enzymatic reactions
4. to explain competitive, uncompetitive, noncompetitiive inhibition of enzymatic reactions
5. to calculate K_M and V_{max} of reactions catalyzed by enzymes with the presence of inhibitors

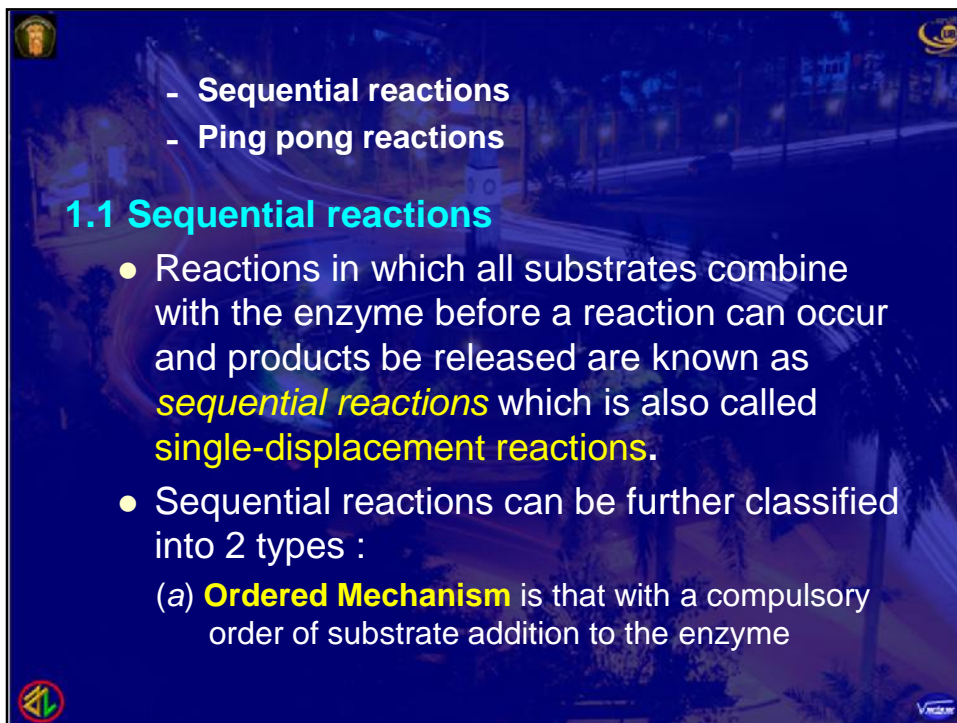
LECTURE LAYOUT

1. REACTION MECHANISMS
 - 1.1 Sequential Reactions
 - 1.2 Ping-Pong Reactions
 - 1.3 Kinetics of Bi-Bi Reactions
2. REACTON INHIBITION
 - 2.1 Irreversible Inhibition
 - 2.2 Reversible Inhibition
 - 2.2.1 Competitive inhibition,
 - 2.2.2 Uncompetitive inhibition.
 - 2.2.3 Noncompetitive inhibition
 - 2.3 Feedback Inhibition



1. REACTION MECHANISMS

- Most biochemical reactions are not simple-, **single-substrate reactions**, but typically involve two or three substrates that combine to release multiple products.
- Yet, enzymatic reactions involving two substrates and yielding two products account for ~60% of known biochemical reactions.
- Bi-Bi reactions, transferring groups catalyzed by enzymes, fall under two major mechanistic classifications.



- Sequential reactions
- Ping pong reactions

1.1 Sequential reactions

- Reactions in which all substrates combine with the enzyme before a reaction can occur and products be released are known as **sequential reactions** which is also called **single-displacement reactions**.
- Sequential reactions can be further classified into 2 types :
 - (a) **Ordered Mechanism** is that with a compulsory order of substrate addition to the enzyme

(b) **Random mechanism** is that with no preference for the order of substrate addition

A B P Q

E EA EAB̃ EPQ EQ E

Sequential Reactions: Ordered Mechanism

All substrates must combine with enzyme before reaction can occur

Bisubstrate reactions

(a)

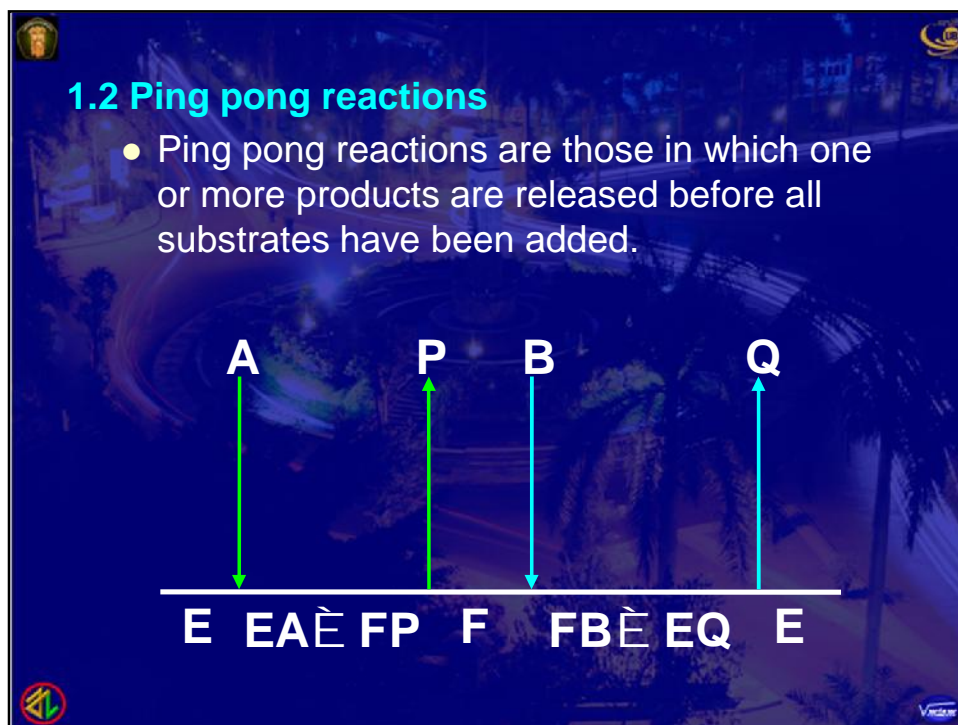
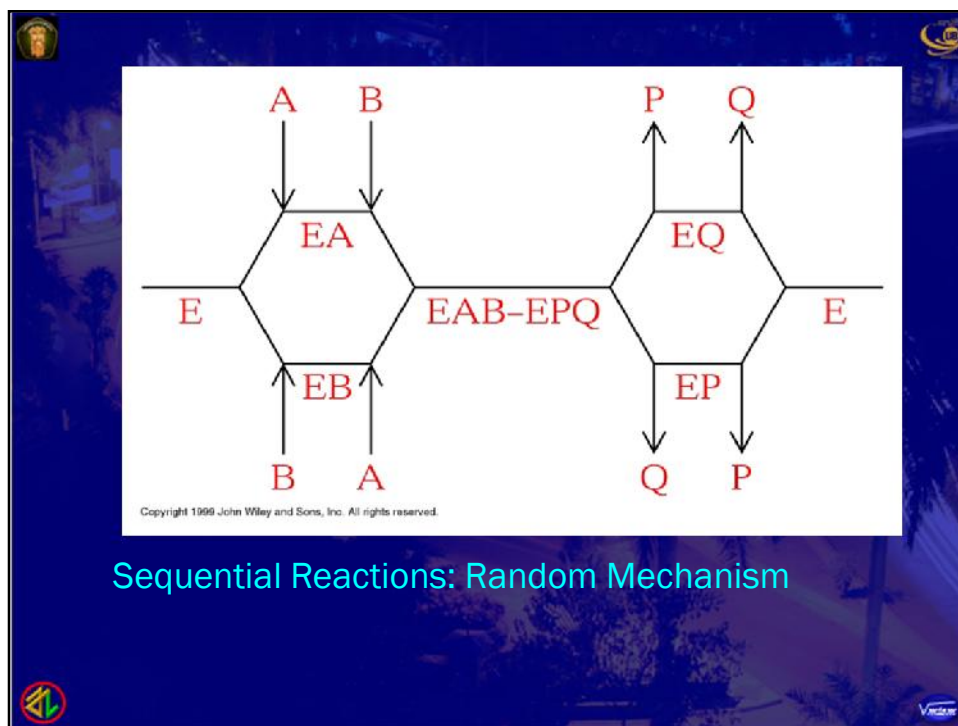
$$\text{R}_1-\overset{\text{O}}{\parallel}{\text{C}}-\text{NH}-\text{R}_2 + \text{H}_2\text{O} \xrightarrow{\text{trypsin}} \text{R}_1-\overset{\text{O}}{\parallel}{\text{C}}-\text{O}^- + \text{H}_3\text{N}^+-\text{R}_2$$

Polypeptide

(b)

$$\text{CH}_3-\overset{\text{H}}{\underset{\text{H}}{\text{C}}}-\text{OH} + \text{NAD}^+ \xrightarrow[\text{H}^+]{\text{alcohol dehydrogenase}} \text{CH}_3-\overset{\text{O}}{\parallel}{\text{C}}-\text{H} + \text{NADH}$$

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1.3 Kinetics of Bi-Bi Reactions

- Steady state kinetic measurements can be utilized to distinguish among the foregoing bisubstrate mechanisms. For doing so, one must first derive their *rate equations*.
- This can be done in much the same way as for single-substrate-enzymes, *i.e.*, solving a set of simultaneous linear equations consisting of an equation expressing the steady state condition for each kinetically distinct enzyme complex plus one equation representing the conservation condition for the enzyme.

- The rate equations for the above-described bisubstrate mechanisms in the absence of products are given below in double reciprocal form.

For ordered Bi-Bi reactions:

$$\frac{1}{V} = \frac{1}{V_m} + \frac{K_M^A}{V_m[A]} + \frac{K_M^B}{V_m[B]} + \frac{K_S^A K_M^B}{V_m[A][B]}$$

$$Y = a + bX_1 + cX_2 + dX_3$$

$$Y = 1/V, a = 1/V_m, b = K_M^A/V_m, c = K_M^B/V_m, d = K_S^A K_M^B/V_m, X_1 = 1/[A], X_2 = 1/[B], \text{ and } X_3 = 1/[A][B]$$

For random Bi Bi reactions :

$$\frac{1}{V} = \frac{1}{V_m} + \frac{K_S^A K_M^B}{V_m K_S^B [A]} + \frac{K_M^B}{V_m [B]} + \frac{K_S^A K_M^B}{V_m [A][B]}$$

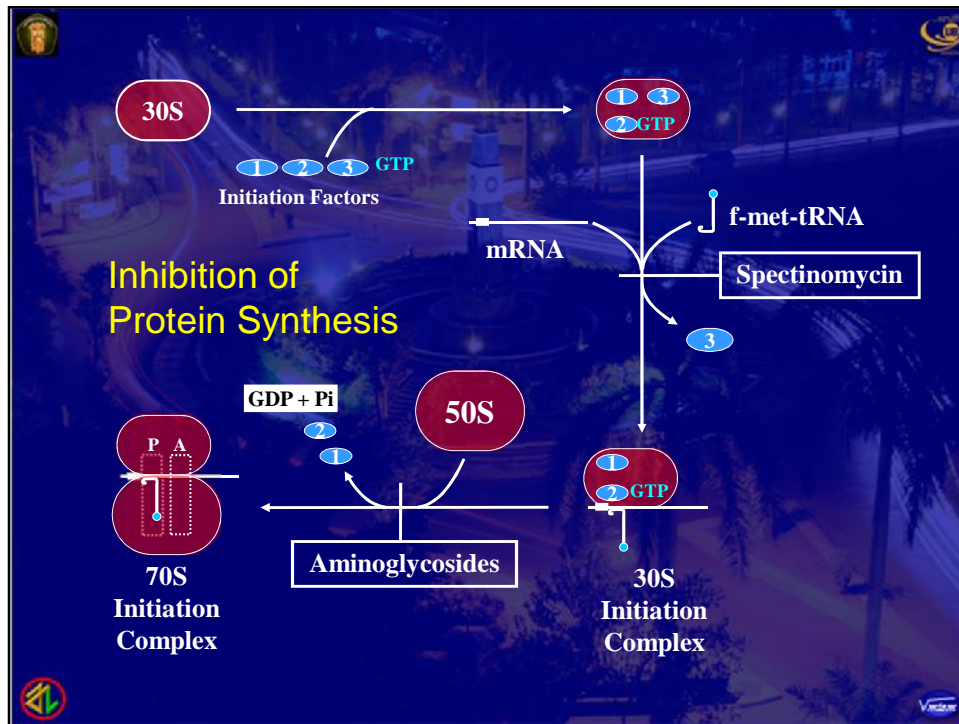
$Y = a + bX_1 + cX_2 + dX_3$
 $Y = 1/V$, $a = 1/V_m$, $b = K_S^A K_M^B / (V_m K_S^B)$, $c = K_M^B / V_m$, $d = K_S^A K_M^B / V_m$, $X_1 = 1/[A]$, $X_2 = 1/[B]$, and $X_3 = 1/([A][B])$

For ping pong Bi Bi reactions :

$$\frac{1}{V} = \frac{1}{V_m} + \frac{K_M^A}{V_m [A]} + \frac{K_M^B}{V_m [B]}$$

2. REACTION INHIBITION

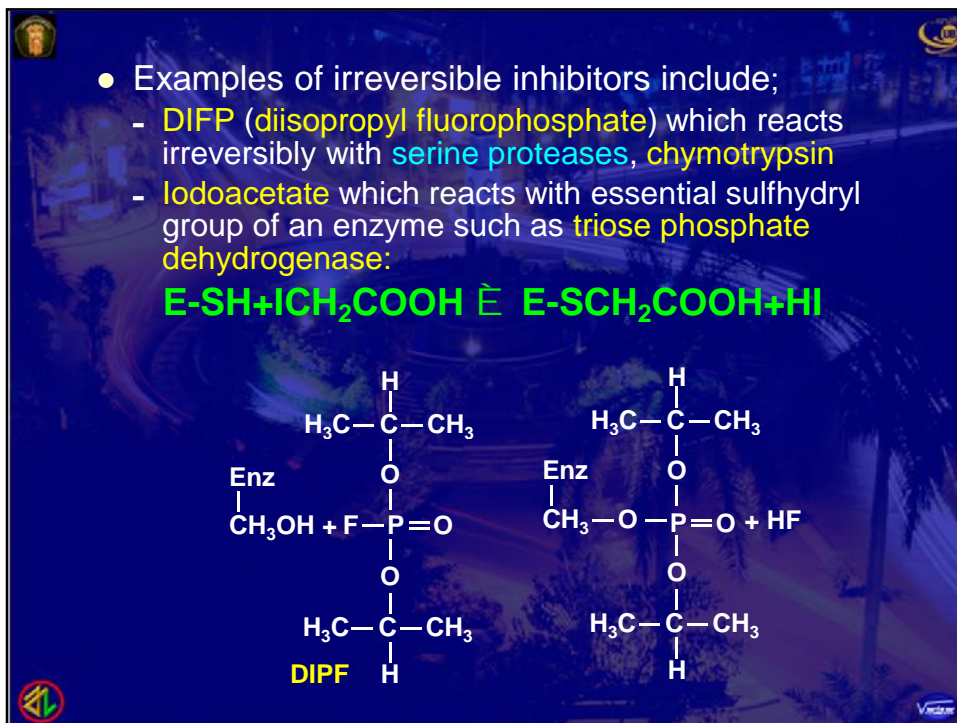
- An important number of compounds have the ability to combine with certain enzymes in either a reversible or irreversible manner, and thereby
 1. block the enzyme, but do not usually destroy it
 2. reduce the rate of enzymic reactions
 3. work specifically in general, and at low concentrations
- Such compounds are called **INHIBITORS** and include;
 - drugs,
 - antibiotics,
 - poisons,
 - anti metabolites
 - products of enzymic reactions.



- Two general classes of inhibitors are recognized ;
 1. **Irreversible**
 2. **Reversible**

2.1 Irreversible

- An irreversible inhibitor forms a **covalent bond** with a specific function, usually an amino acid residue, which may, in some manner, be associated with the catalytic activity of the enzyme.
 - Many examples of enzyme inhibitors covalently bind not at the active site, but physically block the active site

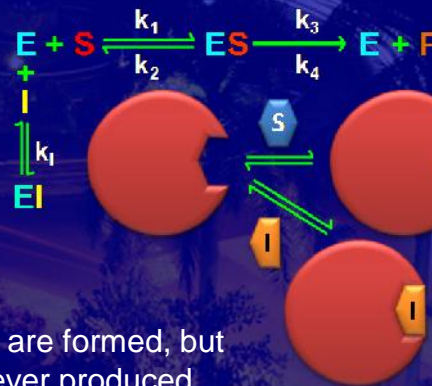


2.2 Reversible Inhibition

- As the term implies, this type of inhibition involves equilibrium between the enzyme and the inhibitor, the equilibrium constant (K_i) being a measure of the affinity of the inhibitor for the enzyme.
- Three distinct types of reversible inhibition are known;
 - Competitive inhibition,
 - Uncompetitive inhibition
 - Noncompetitive inhibition

2.2.1 Competitive Inhibition

- Compounds that may or may not be structurally related to the natural substrate combine reversibly with the enzyme at or near the active site
- The inhibitor and the substrate therefore compete for the same site according to the reaction as shown on the right side.
- ES and EI complexes are formed, but EIS complexes are never produced.



One can conclude that high concentrations of substrate will overcome the inhibition by causing the reaction sequence to swing to the right. The velocity of reaction can be calculated by the following equation

- The rate of reaction can be calculated from the following equation.

$$V = \frac{V_{max} [S]}{K_M (1 + \frac{[I]}{K_I}) + [S]}$$

Competitive

- Among other enzymes that may undergo competitive inhibition is succinic dehydrogenase, which readily oxidizes succinic acid to fumaric acid.
- If increasing concentrations of malonic acid, which closely resembles succinic acid in structure, are added, however, succinic dehydrogenase activity falls markedly
- This inhibition can now be reversed by increasing in turn the concentration of the substrate succinic acid.

Product	Substrate		Competitive Inhibitor		
	Succinate	Glutarate	Malonate	Oxalate	
<chem>C(=O)C(O)C(=O)[O-]</chem>	<chem>C(=O)C(O)C(=O)[O-]</chem>	<chem>C(=O)C(O)C(O)C(=O)[O-]</chem>	<chem>C(=O)C(O)C(=O)[O-]</chem>	<chem>C(=O)C(O)C(=O)[O-]</chem>	<chem>C(=O)C(=O)C(=O)[O-]</chem>
<chem>C(=O)C(O)C(=O)[O-]</chem>	<chem>C(=O)C(O)C(=O)[O-]</chem>	<chem>C(=O)C(O)C(O)C(=O)[O-]</chem>	<chem>C(=O)C(O)C(=O)[O-]</chem>	<chem>C(=O)C(O)C(=O)[O-]</chem>	<chem>C(=O)C(=O)C(=O)[O-]</chem>
<chem>C(=O)C(O)C(=O)[O-]</chem>	<chem>C(=O)C(O)C(=O)[O-]</chem>	<chem>C(=O)C(O)C(O)C(=O)[O-]</chem>	<chem>C(=O)C(O)C(=O)[O-]</chem>	<chem>C(=O)C(O)C(=O)[O-]</chem>	<chem>C(=O)C(=O)C(=O)[O-]</chem>
<chem>C(=O)C(O)C(=O)[O-]</chem>	<chem>C(=O)C(O)C(=O)[O-]</chem>	<chem>C(=O)C(O)C(O)C(=O)[O-]</chem>	<chem>C(=O)C(O)C(=O)[O-]</chem>	<chem>C(=O)C(O)C(=O)[O-]</chem>	<chem>C(=O)C(=O)C(=O)[O-]</chem>

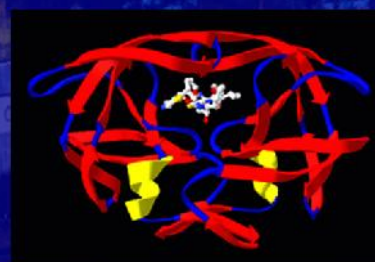
Succinate Dehydrogenase
Competitive Inhibition

2.2.2 Uncompetitive Inhibition

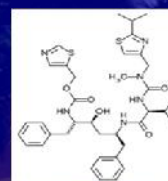
- Compounds that combine only with the ES complex, not with the free enzyme, are called uncompetitive inhibitors. The inhibition is not overcome by high substrate concentrations.



- HIV protease* in a complex with the protease inhibitor *ritonavir*
- The structure of the protease is shown by the red, blue and yellow ribbons. The inhibitor is shown as the smaller ball-and-stick structure near the centre. Created from PDB



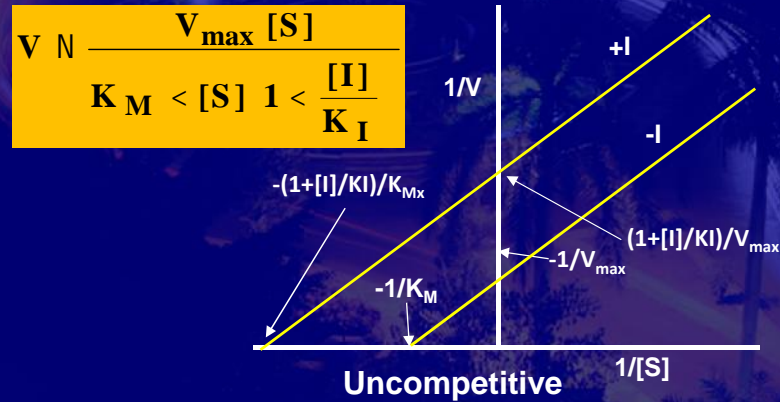
Peptide-based protease inhibitor *ritonavir*



Human immunodeficiency virus

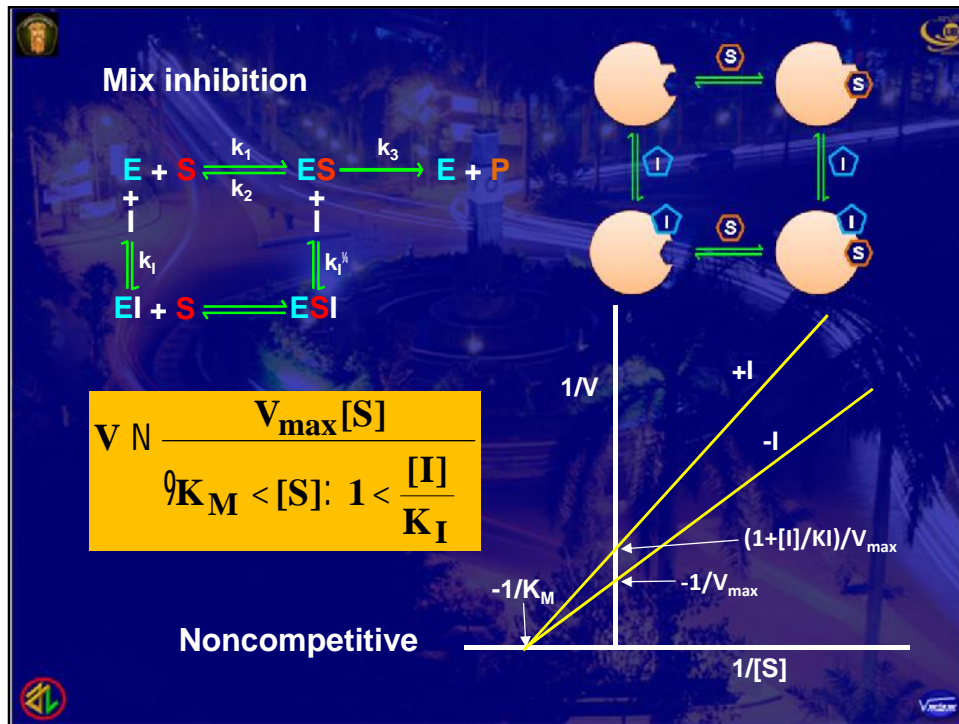
Scanning electron micrograph of HIV-1 (in green) budding from cultured lymphocyte. Multiple round bumps on cell surface represent sites of assembly and budding of virions

- K_M value is consistently smaller than the K_M value of the uninhibited reaction which implies that S is more effectively bound to the enzyme in the presence of the inhibitor.
- The equation used to calculate the velocity of the noncompetitive inhibition is as follows



2.2.3 Noncompetitive Inhibition

- Compounds that reversibly bind with either the enzyme or the enzyme substrate complex are designated as noncompetitive inhibitors
- Noncompetitive inhibition therefore differs from competitive inhibition in that the inhibitor can combine with ES, and S can combine with EI to form in both instances EIS.
- This type of inhibition is not completely reversed by high substrate concentration since the closed sequence will occur regardless of the substrate concentration
- Since the inhibitor binding site is not identical to nor does it modify the active site directly, the K_M is not altered.




2.3 Feedback Inhibition (Allosteric Effectors)

- The activity of some enzymes is controlled by certain molecules binding to a specific regulatory (or **allosteric**) site on the enzyme, distinct from the active site.
- Different molecules can either inhibit or activate the enzyme, allowing sophisticated control of the rate. Only a few enzymes can do this, and they are often at the start of a long biochemical pathway.
- They are generally **activated by the substrate** of the pathway and **inhibited by the product** of the pathway, thus only turning the pathway on when it is needed.

The switch: Allosteric inhibition

Allosteric means “other site”




The diagram shows a yellow enzyme labeled 'E'. It has a pink, irregularly shaped region on its left side labeled 'Active site'. On its right side, there is a red, irregularly shaped region labeled 'Allosteric site'. Red arrows point from the text labels to their respective parts of the enzyme.

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Switching off

- These enzymes have two receptor sites
- One site fits the substrate like other enzymes
- The other site fits an inhibitor molecule



The diagram shows the same yellow enzyme 'E' as in the previous slide. A black, irregularly shaped molecule labeled 'Substrate' is positioned near the pink 'Active site', but it is not fitting. An orange, cross-shaped molecule labeled 'Inhibitor molecule' is fitting into the red 'Allosteric site'. Text labels with arrows point to the substrate and inhibitor.

Substrate cannot fit into the **active site**

Inhibitor fits into **allosteric site**

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Feedback inhibition occurs when isoleucine binds to an allosteric site on threonine deaminase

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- This process is known as **feedback inhibition**.

Feedback Inhibition

HOW TO SOLVE THE EQUATIONS

Type of Inhibition	Equation	V_{max}	K_m
None	$v = \frac{V_{max}[S]}{K_m + [S]}$	—	—
Competitive	$v = \frac{V_{max}[S]}{K_m \left(1 + \frac{I}{K_i}\right) + [S]}$	No change	Increased
Noncompetitive	$v = \frac{V_{max}[S]}{(K_m + S) \left(1 + \frac{I}{K_i}\right)}$	Decreased	No change
Uncompetitive	$v = \frac{V_{max}[S]}{K_m + [S] \left(1 + \frac{I}{K_i}\right)}$	Decreased	Decreased

1. Competitive inhibitor

$$V = \frac{V_{max}[S]}{K_M \left(1 + \frac{[I]}{K_I}\right) + [S]}$$

$$\frac{1}{V} = \frac{K_M \left(1 + \frac{[I]}{K_I}\right)}{V_{max}} \frac{1}{[S]} + \frac{1}{V_{max}}$$

- $y = 1/V$; $x = 1/[S]$
- $a = 1/V_{max}$
- $b = K_M(1 + [I]/K_I)/V_{max}$

2. Uncompetitive

$$v = \frac{v_{\max}[S]}{K_M + [S] \left(1 + \frac{[I]}{K_I}\right)}$$

$$\frac{1}{v} = \frac{K_M}{v_{\max}} \frac{1}{[S]} + \frac{1 + \frac{[I]}{K_I}}{v_{\max}}$$

- $y = 1/v$; $x = 1/[s]$
- $a = (1 + [I]/K_I)/v_{\max}$
- $b = K_M/v_{\max}$

3. Noncompetitive Inhibition

$$v = \frac{v_{\max}[S]}{K_M + [S] \left(1 + \frac{[I]}{K_I}\right)}$$

$$\frac{1}{v} = \frac{K_M}{v_{\max}} \frac{1}{[S]} + \frac{1 + \frac{[I]}{K_I}}{v_{\max}}$$

- $y = 1/v$; $x = 1/[s]$
- $a = (1 + [I]/K_I)/v_{\max}$
- $b = K_M(1 + [I]/K_I)/v_{\max}$



QUIZ (10 min)

- How is enzyme specificity achieved ?
- Calculate V_{max} & K_M from the following data, and does the reaction obey Michaelis-Menten kinetics ?

[DNA] mol total nucleotides/L	Free nucleotides in solution, V (pmol/L)	
	0 min	10 min
1.0×10^{-5}	0.05	5.1
1.0×10^{-6}	0.04	4.5
1.0×10^{-7}	0.06	3.2
1.0×10^{-8}	0.04	1.4
1.0×10^{-9}	0.04	0.23

ANSWERS

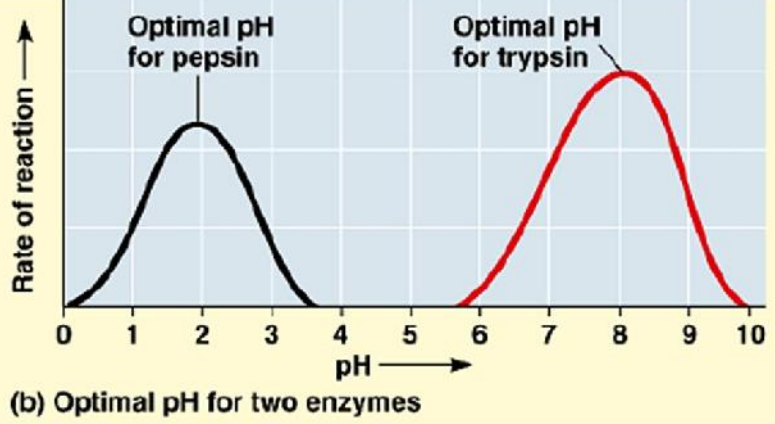
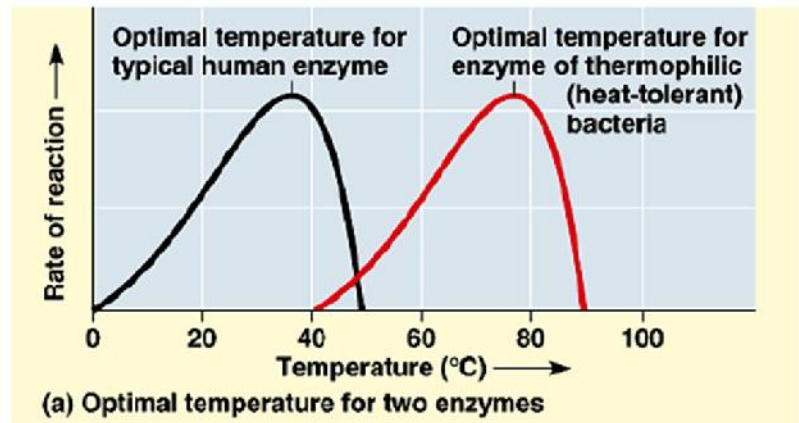
1. The enzyme specificity is achieved through the characteristic of active site
2. $V_{max} = 4.36695$
 $K_M = 2.2E-08$
 $R^2 = 0.999864$, so the reaction obeys Michaelis-Menten kinetics

SOAL

- Diketahui suatu reaksi enzimatik tanpa dan dengan inhibitor dengan $[I] = 2,2 \cdot 10^4 M$.
- Hitunglah K_M dan V_{max} tanpa dan dengan I serta K_I

[S]	V(-I)	V(+I)
$1 \cdot 10^{-4}$	28	17
$1.5 \cdot 10^{-4}$	36	23
$2.0 \cdot 10^{-4}$	43	29
$5 \cdot 10^{-4}$	65	50
$7.5 \cdot 10^{-4}$	74	61

Figure 6.16 Environmental factors affecting enzyme activity



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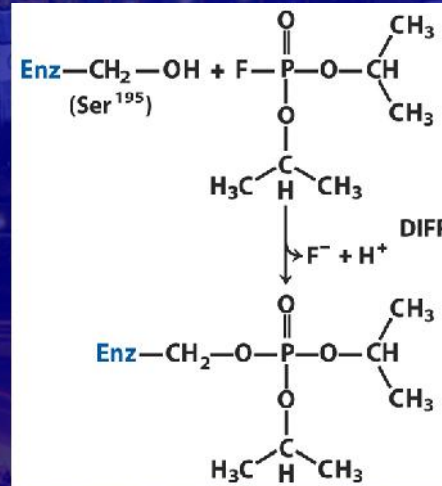
	▶ Competitive	■ Non-competitive	◀ Uncompetitive
Cartoon Guide	<p>Substrate Inhibitor Compete for active site</p>	<p>Different site</p>	
Equation and Description	$E + S \rightleftharpoons ES \rightarrow E + P$ $+ I \rightleftharpoons EI$ $\downarrow \uparrow$ E/I <p>[I] binds to free [E] only, and competes with [S]; increasing [S] overcomes inhibition by [I].</p>	$E + S \rightleftharpoons ES \rightarrow E + P$ $+ I \rightleftharpoons EI$ $\downarrow \uparrow$ $E/I + S \rightleftharpoons EIS$ <p>[I] binds to free [E] or [ES] complex; Increasing [S] can not overcome [I] inhibition.</p>	$E + S \rightleftharpoons ES \rightarrow E + P$ $+ I \rightleftharpoons ESI$ $\downarrow \uparrow$ E/S <p>[I] binds to [ES] complex only, increasing [S] favors the inhibition by [I].</p>

	▶ Competitive	■ Non-competitive	◀ Uncompetitive
Direct Plots	<p>V_{max} v_o K_m K_m' [S], mM</p>	<p>V_{max} v_o V_{max}' [S], mM $K_m = K_m'$</p>	<p>V_{max} v_o V_{max}' [S], mM K_m K_m'</p>
Double Reciprocal	<p>V_{max} unchanged K_m increased</p> <p>Intersect at Y axis $1/v_o$ I $1/V_{max}$ $1/K_m$ $1/[S]$</p>	<p>V_{max} decreased K_m unchanged</p> <p>Intersect at X axis $1/v_o$ I $1/V_{max}$ $1/K_m$ $1/[S]$</p>	<p>Both V_{max} & K_m decreased</p> <p>Two parallel lines $1/v_o$ I $1/V_{max}$ $1/K_m$ $1/[S]$</p>

Juang RH (2004) BCbasics

Example of a suicide inhibitor

- Diisopropylfluorophosphate (DIFP) forms a covalent bond with an active -site residue (Ser) of the enzyme chymotrypsin.
- Every molecule that reacts is inactivated irreversibly.
- Here, a key active site Ser is bound irreversibly to the inhibitor, preventing it from doing its "normal" job.



Irreversible inhibitor

12

9/28/05

The effect of enzyme inhibition

- **Irreversible inhibitors:** Combine with the functional groups of the amino acids in the active site, irreversibly

Examples: nerve gases and pesticides, containing organophosphorus, combine with serine residues in the enzyme acetylcholine esterase