

Enzyme Inhibitors

Molecules which bind and act to alter the reaction kinetics of an enzyme. By binding to enzyme reducing the affinity, velocity, catalytic turnover or all of the above. - examples include poisons and drugs

- Some inhibitors **reversibly** bind directly to active site competing with normal substrate (competitive inhibitor)
- Some inhibitors **reversibly** bind to the protein away from active site – altering structure of protein so the enzyme does not function (either won't bind or won't react substrate). Non and Un-competitive and allosteric...
- Other inhibitors bind covalently (**irreversibly**) to the active site, leaving the enzyme "dead". Uncompetitive – sometimes called suicide inhibitor

Competitive Inhibitors Compete With Substrate for the Same Site on the Enzyme

- Inhibitor is similar to substrate and both bind to or near active site. 'compete' for binding
- Inhibitor is unreactive - EI state
- Lineweaver Burke intersect at the Y axis

$$E + S \xrightleftharpoons[k_{-1}]{k_1} ES \xrightarrow{k_2} P + E$$

$$E + I \xrightleftharpoons{K_I} EI$$

$$EI + S \longrightarrow \text{NO REACTION}$$

Competitive Inhibitors Compete With Substrate for the Same Site on the Enzyme

Lineweaver-Burk plot of competitive inhibition, showing lines for no I, [I], and 2[I].

K_m is reduced but V_{max} is unchanged

Uncompetitive Inhibition, where I combines only with ES, but not with Free E

- binds covalently in the transition state
- Some are** suicide inhibitors
- binds to the ES complex
- lowers affinity and velocity
- lineweaver Burke plots are parallel

Note that both intercepts change but the slope (K_m/V_{max}) remains constant in the presence of I.

Uncompetitive Inhibition, where I combines only ES, but not with Free E

$$E + S \xrightleftharpoons[k_{-1}]{k_1} ES \xrightarrow{k_2} P + E$$

$$ES + I \xrightleftharpoons{K'_I} ESI$$

$$ESI \longrightarrow \text{NO REACTION}$$

Note that both intercepts change but the slope (K_m/V_{max}) remains constant in the presence of I.

Pure Noncompetitive Inhibition – where S and I bind to different sites on the enzyme

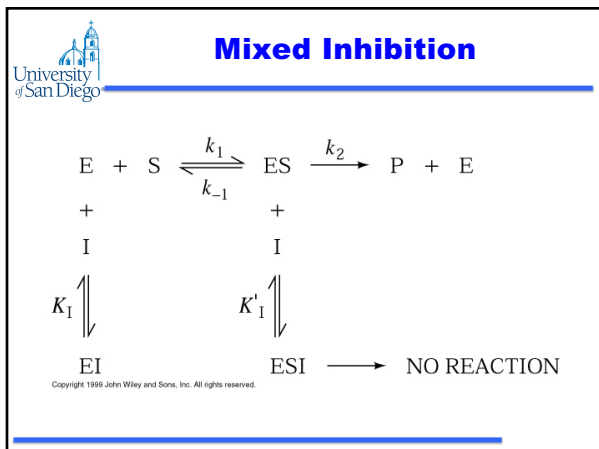
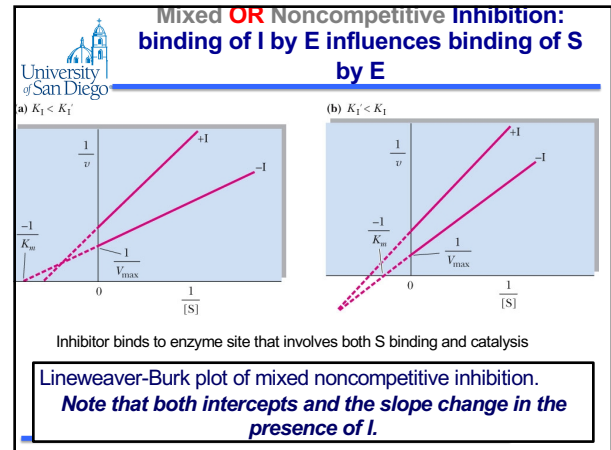
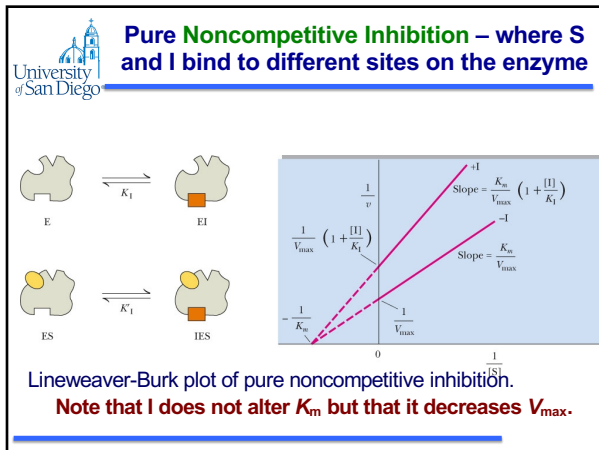
- inhibitor binds distal to active site
- affects enzyme rate not affinity
- binds E in ES or E
- Reversible
- Lineweaver Burke intersect at the Y axis

$$E + S \xrightleftharpoons[k_{-1}]{k_1} ES \xrightarrow{k_2} P + E$$

$$E + I \xrightleftharpoons{K_I} EI$$

$$EI + S \longrightarrow \text{NO REACTION}$$

Also Known as Mixed



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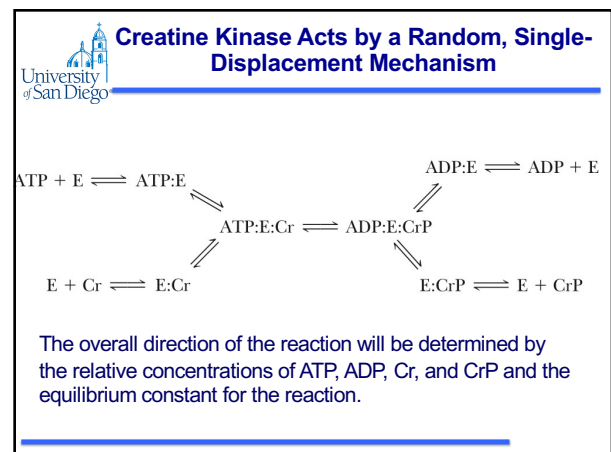
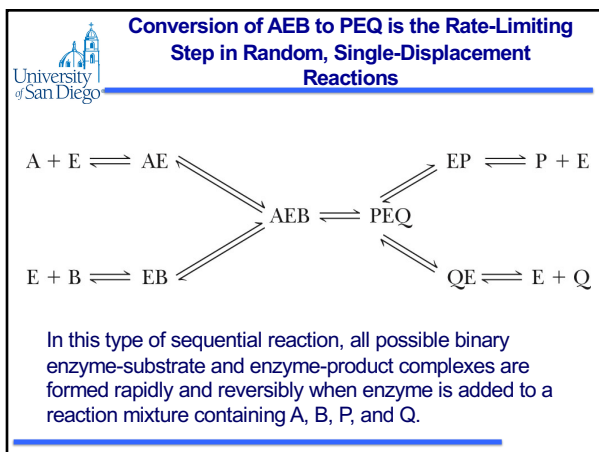
Kinetic Behavior of Bimolecular Reactions?

Enzymes often catalyze reactions involving two (or more) substrates

Bisubstrate reactions may be **sequential** or **single-displacement** reactions or **double-displacement (ping-pong)** reactions

Single-displacement reactions can be of two distinct classes:

1. **Random**, where either substrate may bind first, followed by the other substrate
2. **Ordered**, where a **leading substrate** binds first, followed by the other substrate



In an Ordered, Single-Displacement Reaction, the Leading Substrate Must Bind First

$$E \xrightarrow{A} AE \xrightarrow{B} AEB \rightleftharpoons PEQ \xrightarrow{} PE \xrightarrow{} E$$

The leading substrate (A) binds first, followed by B. Reaction between A and B occurs in the ternary complex and is usually followed by an ordered release of the products, P and Q.

The Double Displacement (Ping-Pong) Reaction

$$E \xrightarrow{A} AE \rightleftharpoons PE' \xrightarrow{} E' \xrightarrow{B} E'B \rightleftharpoons EQ \xrightarrow{} E$$

Double-Displacement (Ping-Pong) reactions proceed via formation of a covalently modified enzyme intermediate. Reactions conforming to this kinetic pattern are characterized by the fact that the product of the enzyme's reaction with A (called P in the above scheme) is released prior to reaction of the enzyme with the second substrate, B.

How do enzymes work?

It's the amino acids!

Stabilization of reactive intermediates

Enzyme-catalyzed reactions generate *reactive intermediates*, such as carbanions, carbocations, and radicals, that normally would require strong reaction conditions to generate in chemical reactions and would be very unstable in aqueous solution.

Its about transition state!

Enzymes stabilize the transition state and catalyze the reaction by providing appropriate functional groups of the protein or cofactor spatially and temporally appropriate manner.

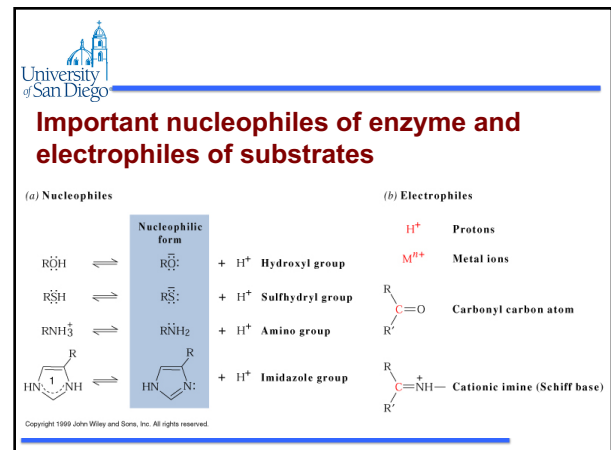
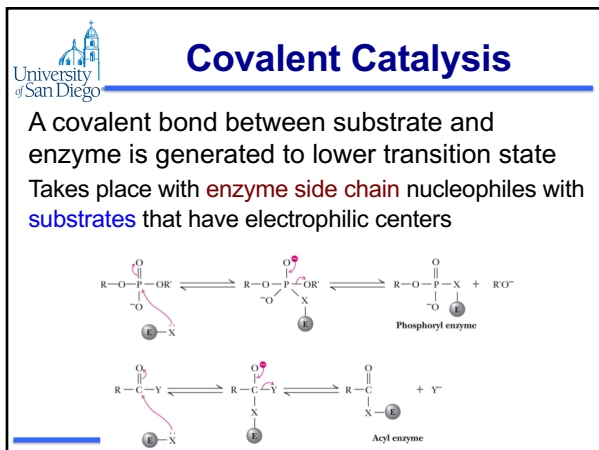
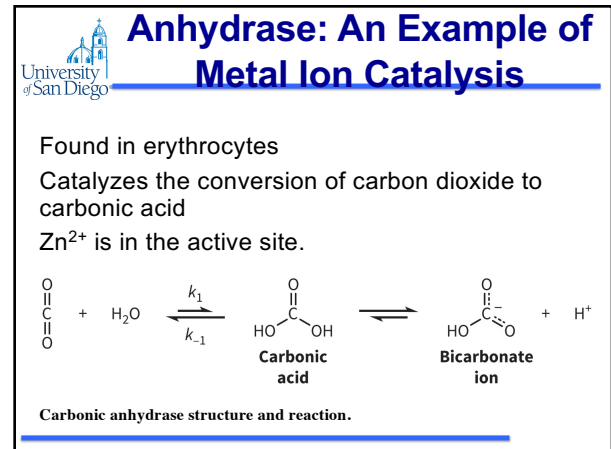
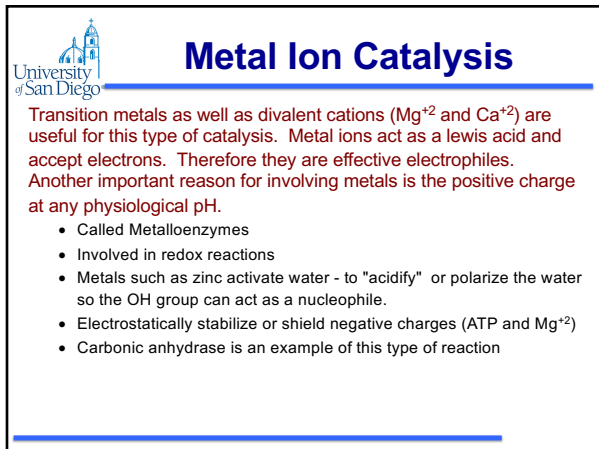
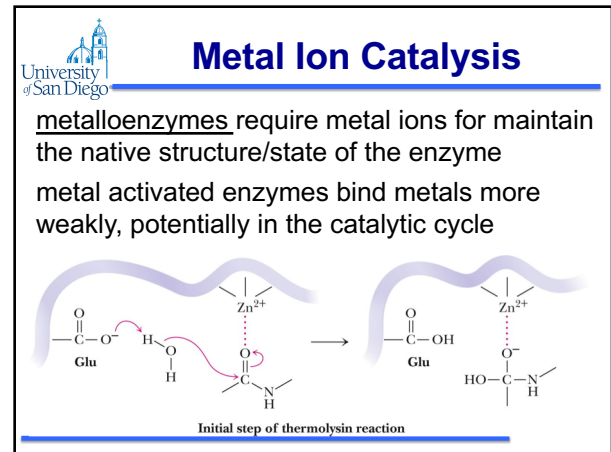
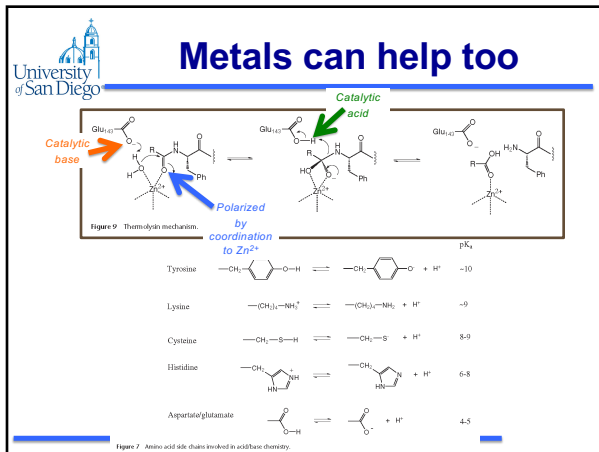
- There are five general mechanisms by which enzymes contribute to catalysis:

Acid/Base, covalent and metal ion catalysis. Proximity and orientation (strain) effects and preferential binding to transition state complex

Acid / Base catalysis

Chemical groups are made more reactive by adding or removing a proton from substrate to reduce (stabilizing) transition state free energy.

- Result of Acid/Base catalysis is making a reactive group more reactive by increasing its intrinsic electrophilic or nucleophilic character
- This can increase the rate 10-100 fold
- Microenvironment shifts in pKa allow for several amino acids to be involved: Asp, Glu, His, Cys, Tyr, and Lys often involved



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Covalent Catalysis

In some enzymes a nucleophilic side chain group forms an unstable covalent bond with the substrate. The enzyme-substrate complex then forms product. The pathway can require that the intermediate is more susceptible to nucleophilic attack by water than the original substrate.

Three stages of covalent catalysis:

- Formation of a bond between substrate and enzyme
- Removal of electrons to make a reactive center
- Elimination of the bond that was formed in step one

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Covalent Catalysis

1. Bond formation between deprotonated Cys nucleophile (STRONG nucleophilic thiolate anion) and electrophilic substrate
2. Lewis base transferring H^+ & e^- Intermediate
3. Elimination of (phosphate anion attack) bond forms new product

Figure 10: Mechanism of 8-oxotetrahydrofuran CYP4B5.

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Lysozyme: An Example of Acid-Base Catalysis

A natural antibiotic

An enzyme found in egg whites, tears, and mucus

The active site contains two negatively charged residues

Glutamic acid is protonated, whereas aspartate is not

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Role of Zn^{2+} in Carbonic Anhydrase

After Sheridan, R.P. and Allen, L.C., J. Am. Chem. Soc. 103, 1545 (1981). Copyright 1999 John Wiley & Sons, Inc. All rights reserved.

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Mechanism of Anhydrase

1. Water binds the active site zinc to generate the active enzyme.
2. Active site zinc polarizes water. Protons are shuttled out by histidine.
3. Carbon dioxide binds; the hydroxyl group from water attacks.
4. Bicarbonate is stabilized through the zinc ion prior to departure.

Mechanism of carbonic anhydrase.

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Chymotrypsin: A Covalent Catalysis

Serine protease

Cleaves dietary protein on the C-terminus of Tyr, Phe, and Trp

Secreted from the pancreas

Contains a catalytic triad containing Asp, His, and Ser

Inactive form = chymotrypsinogen

Chymotrypsin

Peptide bond cleaved by chymotrypsin

Polypeptide

Polypeptide fragments

$R = \text{Phe, Trp, Tyr, etc.}$