

ENZYMODOLOGY

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- Unit-2 Enzyme Inhibition
- Lecture-2
- Content- Reversible and Irreversible enzyme Inhibition

- Inhibitor : An Enzyme inhibitor is a Enzyme compound that decreases or diminish the rate or velocity of an enzyme-catalyzed reaction by influencing the binding of S and /or its turnover number.
- The inhibitor may be organic or inorganic in nature Inhibitors - drugs, antibiotics ,toxins and antimetabolite or natural products of enzyme reaction.

Inhibition

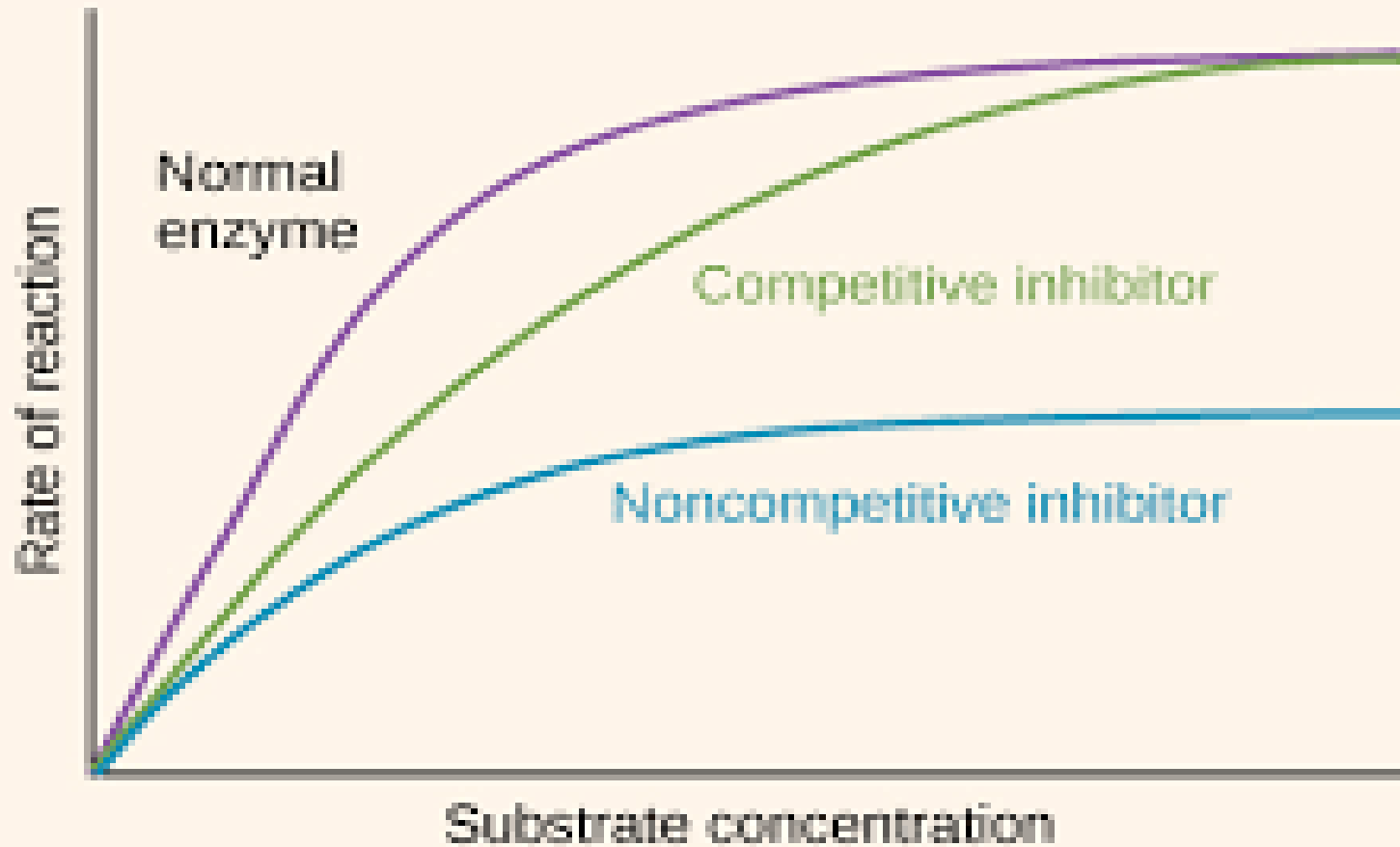
- Type of enzyme Inhibitors
- Competitive Allosteric Suicide Inhibitors
- Non- Competitive
- Uncompetitive
- Reversible Inhibition Inhibitor binds non-covalently (weak interaction) with Enzyme If inhibitor is removed – action of E fully restored - Reversible
- An Equilibrium is established between the free inhibitor & EI Complex and is defined by an equilibrium constant (K_i) $E + I \rightleftharpoons EI$ The activity of Enzyme is fully restored on removing the Inhibitor by dialysis

Reversible Irreversible

- Inhibitor binds reversibly to the same site that the substrate binds - competes with the S for binding. □ Substrate analogue – I closely resembles the S
- I can be reversed by increasing the conc. of S – reversible
- Degree of inhibition - depend on the conc. of S & I and on the relative affinities of the enzyme for S & I

Competitive Inhibition

Enzyme Inhibition



- Competitive Inhibition Apo enzyme Velocity is decreased - effective concentration of enzyme is reduced K_m is increased -affinity of the enzyme towards substrate is apparently decreased in presence of the inhibitor
- V_{max} is not changed No inhibitor Inhibitor V_{max}
 $\frac{1}{2} V_{max}$ K_m New K_m [s] v $1/V_m$ $-1/K_m$ $1/S$ $1/V$
 No Inhibitor $-1/K_m$ Inhibitor
- Malonate is a competitive inhibitor of SDH COOH
 CH_2 CH_2 COOH Succinate dehydrogenase FAD
 FADH₂ Succinate COOH H -C C-H COOH COOH
 CH_2 COOH Malonate Fumarate Similarity in three
 dimensional structure b/w S and I

- Antibacterial action of sulpha drugs (sulfonamide) - structural analog of PABA
NC(=O)c1ccc(N)cc1 PABA - para amino benzoic acid
NC(=O)c1ccc(N)cc1 NH₂
- Bacteria Pteroid synthetase Sulfonamide inhibits the bacterial enzyme
NC(=O)c1ccc(N)cc1 + NC(=O)c1ccc(N)cc1 → NC(=O)c1ccc(N)cc1 + 7,8-dihydropteroic acid
 Folic acid Non toxic to human -human cannot synthesize Folic acid
- Clinically useful Competitive Inhibition Drugs Target Enzyme Therapeutic Use
 STATINS - Atorvastatin, simvastatin HMG CoA reductase Decrease plasma Cholesterol level
- Antihyperlipidemic agents Allopurinol Xanthine oxidase Gout Methotrexate Dihydrofolate reductase Cancer Captopril & Enalapril Angiotensin converting enzyme High blood pressure Dicoumarol Vit.K-epoxide-reductase Anti-coagulan

- Non-competitive Inhibition Inhibitor binds at a site other than the active site of the enzyme □ I has no structural resemblance to the S – No competition for binding □ Increase in the S conc. does not relieve this I □ I & S binds at different site – formation of both EI and EIS complexes is possible. □ EIS – forms product at a slower rate than ES □ Reaction is slowed down but not halted.
- Non-competitive Inhibition
- Product is formed at slower rate but not E halted I
- K_m value is unchanged - I do not interfere with the binding of S to E V_{max} decreases - I cannot be overcome by increasing the conc. of S
 No inhibitor Inhibitor V_{max} V_{max} i 1/2 V_{max} K_m [s] □ v
 1/2 V_{max} i Inhibitor No Inhibitor 1 V_{max} 1 K_m 1/s 1/v -
- Non competitive inhibitor Inhibitor Enzyme inhibited Heavy metals – Ag²⁺, Hg²⁺, Pb²⁺ Binding with cysteinyl SH gr of E
 Pepstatin Pepsin Soyabean trypsin inhibitor Trypsin Ethanol or narcotic drugs Acid phosphatase

- Uncompetitive Inhibition : binds only to the ES complex , not to free E cause structural distortion of the active site - E catalytically inactive can't be reversed by increasing the [S] since I doesn't compete with S for the same binding site □ Inhibition of placental alkaline phosphatase (Regan iso-enzyme) by phenylalanine . Enzyme Enzyme S Enzyme I S I
- $E + S \rightleftharpoons ES \xrightarrow{k_2} E + P$ $E + S + I \rightleftharpoons ESI$ Inhibitor V_{max} V_{max} i
 $\frac{1}{2} V_{max}$ K_m [s] □ $\frac{1}{2} V_{max}$ i $V_{max} =$ Decreases
 $K_m =$ Decreases K_{mi} v I has no affinity for free E
 No Inhibitor I No I $-1/K_m$ $1/v$ $-1/K_m$ $1/V_{max}$
 $1/V_{maxi}$ $1/s$

- Irreversible Inhibition Inhibitor binds covalently (strong) with the enzyme irreversibly so it can't dissociate from the enzyme Inhibitor cause conformation change at active site of the E-destroying their capacity to function as catalysts. Enzyme activity is not regained on dialysis / by increasing the conc. of S \square A variety of poisons, such as iodoacetate, OP poisoning and oxidizing agents act as irreversible inhibition.
- Irreversible Inhibition In terms of kinetics – irreversible is similar to non competitive inhibition
 V_{max} – Decreased K_m – No change V_{max}
 $V_{max} \ i \ \frac{1}{2}$
 $V_{max} \ K_m \ [s] \ \square \ v \ \frac{1}{2} \ V_{max} \ i$ No inhibitor Inhibitor

Thanks