

Slide : Enzymes 5

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Sections : 4,5,6

■ Slide □ Sheet



# Biochemistry

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cybernetics  
ecology  
taxonomy  
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bacteriology  
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anatomy  
microbiology  
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cystology  
xenobiology  
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gnotobiotics  
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astrobiology  
biochemics  
physiology  
ethnobiology  
molecular  
bioecology  
virology  
zoology  
biometry  
enzymology  
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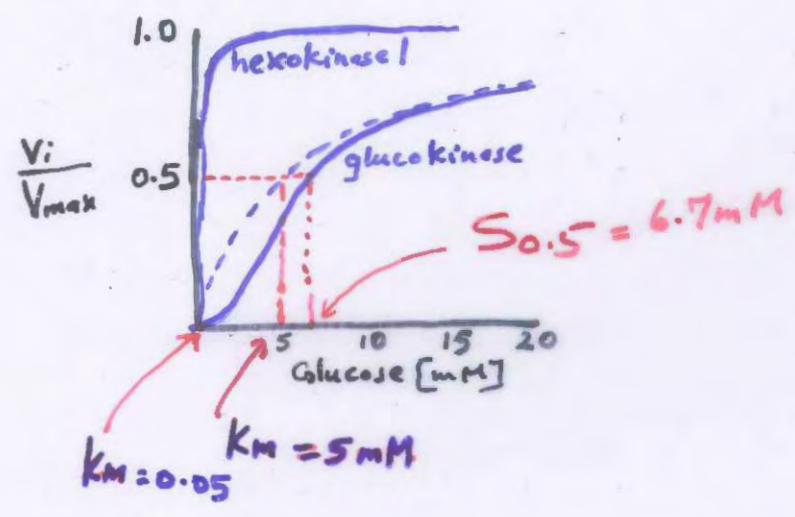
Mousa Suboh

# Hexokinase Isozymes have different $K_m$ values for glucose

HK-I in erythrocytes, muscle, brain & most other tissues  
(110 kD)  $K_m$  0.02 - 0.13 mM

GK isozyme in liver: 5 to 6 mM ( $K_m$ )  
(55 kD)

$K_m$  for GK as calculated from Michaelis-Menten Eq.



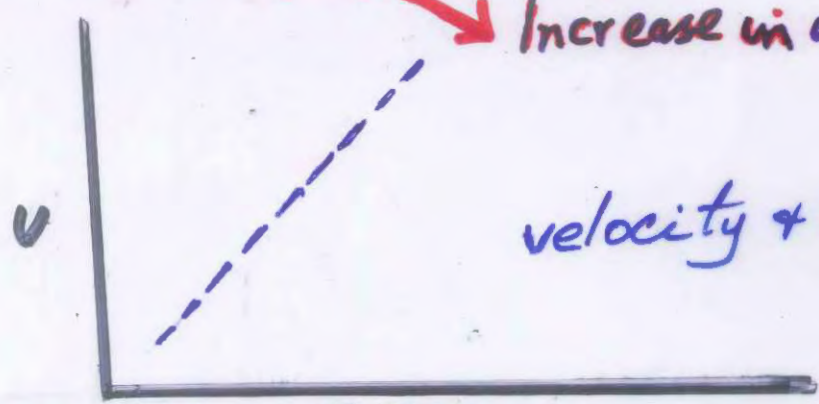
- Which enzyme activity increases after a high carbohydrate meal

-  $V_{max} = k_3 [E_T]$

Increase in  $V_{max}$

Increase in catalytic power  $k_{cat}$  and/or

Increase in amount of enzyme



velocity + Enz. conc.

[E]

# Enzyme Inhibitors

Reversible

Competitive  
(substrate analogues)

Non-Competitive

uncompetitive

Irreversible

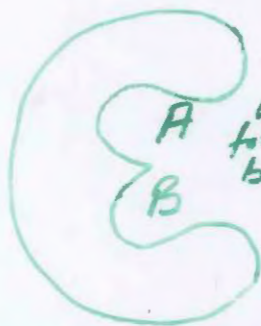
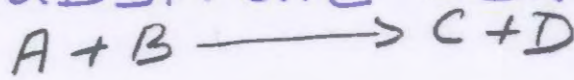
Group Specific

[e.g. DIPF,  
aspirin,  
Heavy metals  
...etc.]

Suicide

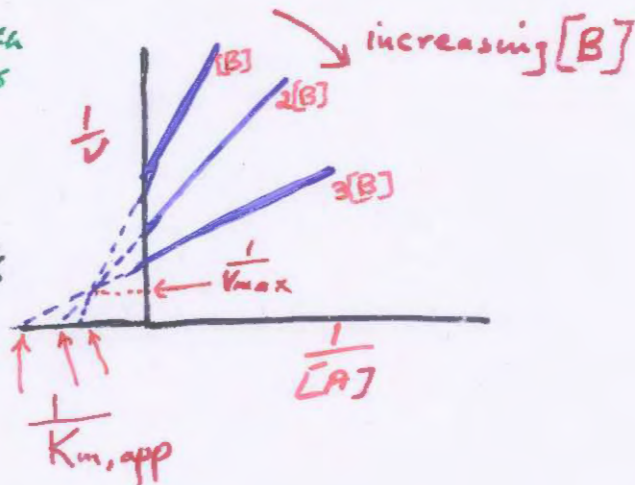
(by mechanism based inhibition)  
[e.g. Penicillins  
allopathic drugs  
other potential drugs]

# Multi Substrate Reactions:-



when the enzyme forms complex with both substrates

an apparent  $K_m$  depends on conc. of the other substrate



## II REG. by REV. INHIBITION

→ 1. - Competitive  
Increases  $K_m$

→ 2. - Noncompetitive  
I binds E in presence or absence of S  
decreases  $V_{max}$

X 3. - Uncompetitive

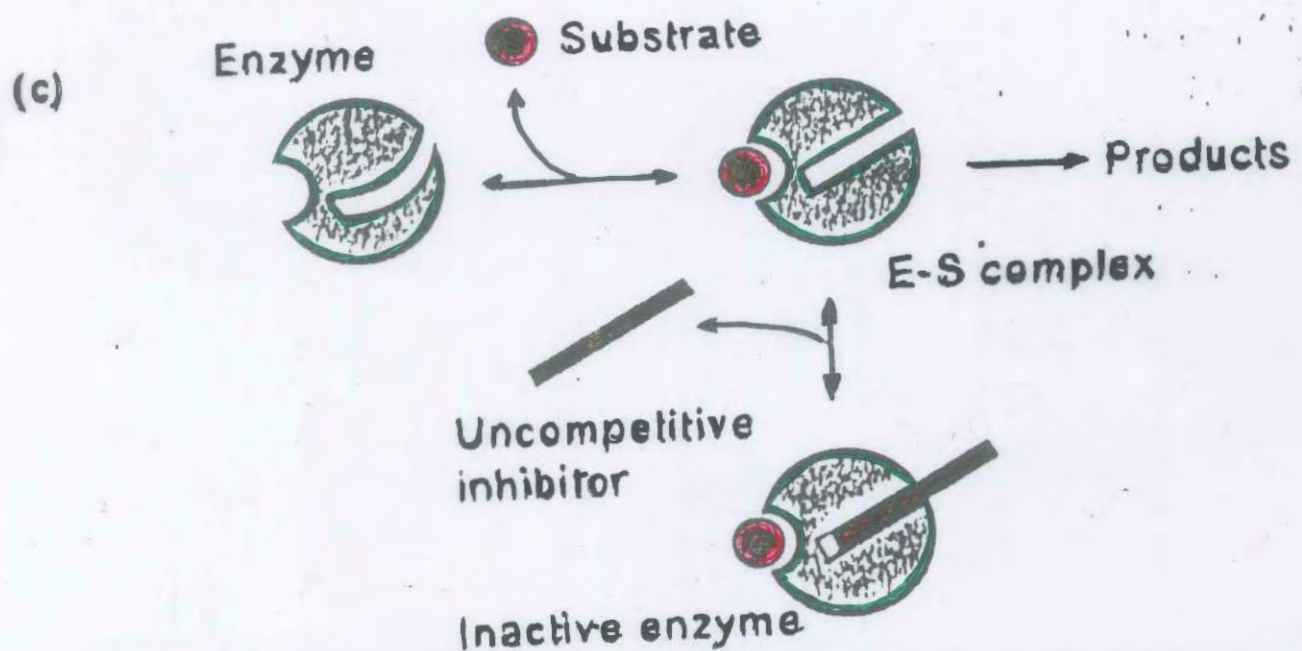
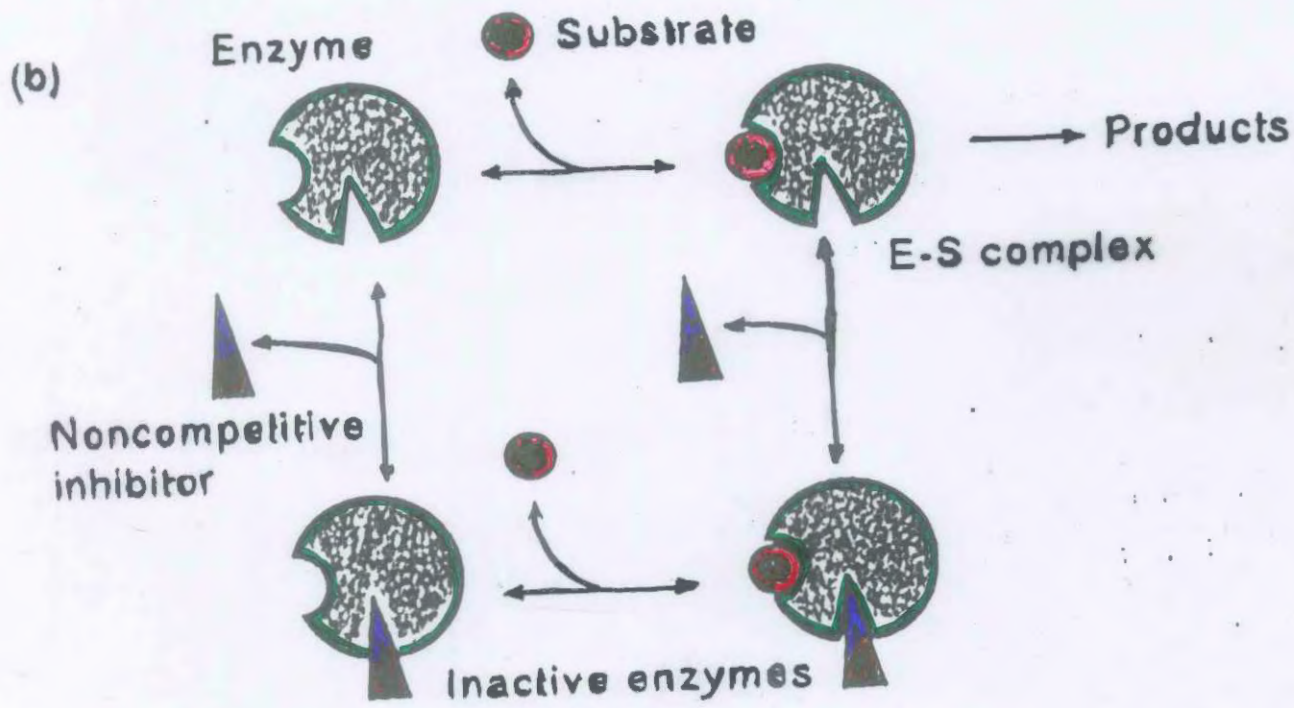
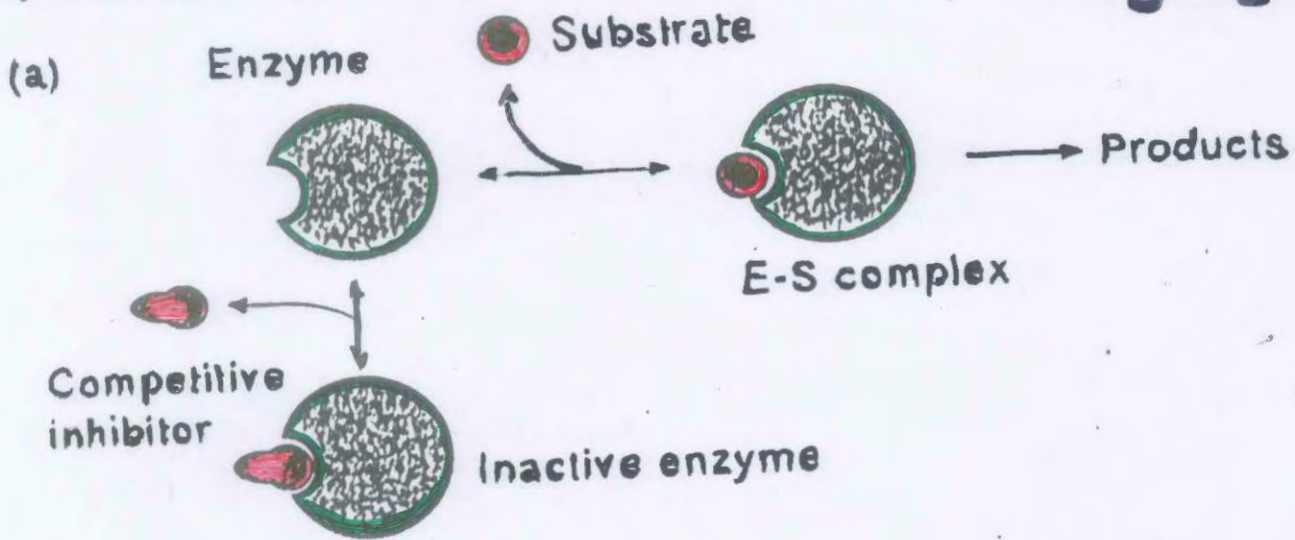
I binds only to ES complex. Occurs with enzyme of a multi substrate having an ordered sequence of substrate binding.

Decreases both  $V_{max}$  and  $K_m$  - Not encountered in medicine. Rare with single substrate reaction

### → Product inhibition

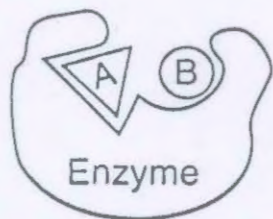
Most products are reversible inhibitor. For the enzymes that produce them - to prevent their accumulation  
e.g. Flu-b-P inhibition of hexokinase

# Reversible Inhibition - Single[S] 76

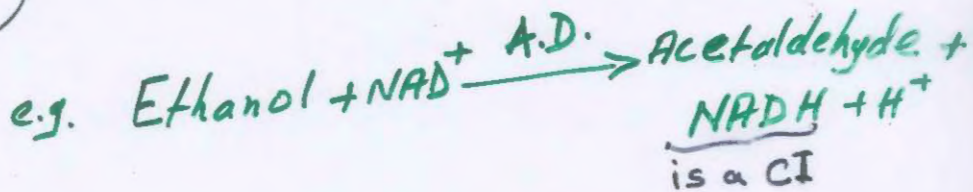


# - Reversible Inhibition of the Active Site

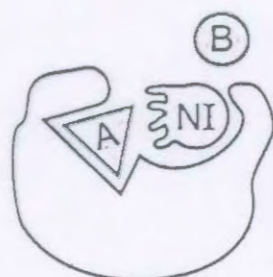
Reaction



Substrates both bind



CI is competitive with respect to A



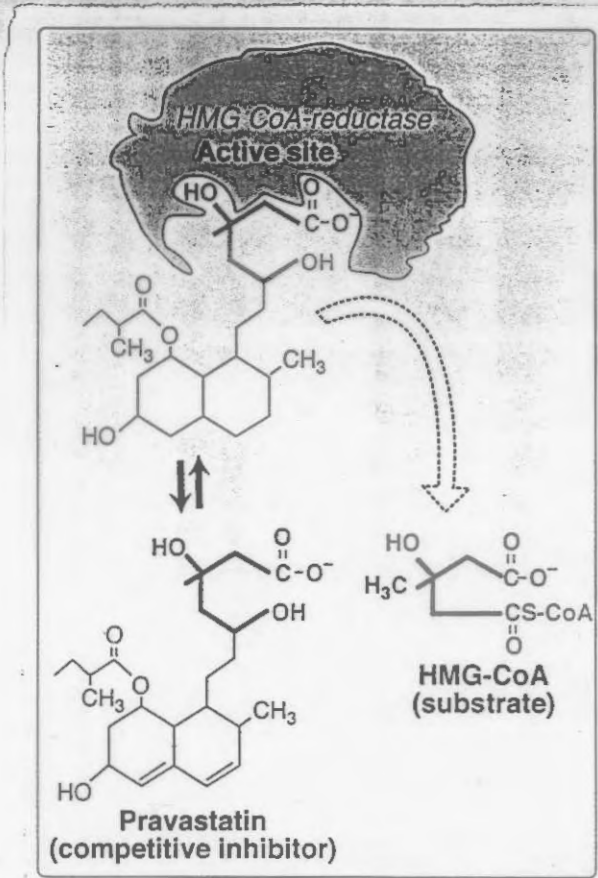
NI is noncompetitive with respect to A

Fig. 9.27. Competitive or noncompetitive inhibition with respect to a given substrate. A and B are substrates for the reaction. The enzyme has separate binding sites for each substrate, which overlap in the active site. The competitive inhibitor (CI) competes for the binding site of A, the substrate it most closely resembles. NI is a noncompetitive inhibitor with respect to the substrate A, and A can still bind to its binding site in the presence of NI. However, NI is competitive with respect to B because it binds to the B binding site. In contrast, an inhibitor that is uncompetitive with respect to A might resemble NI, but it could only bind to the B site after A is bound.

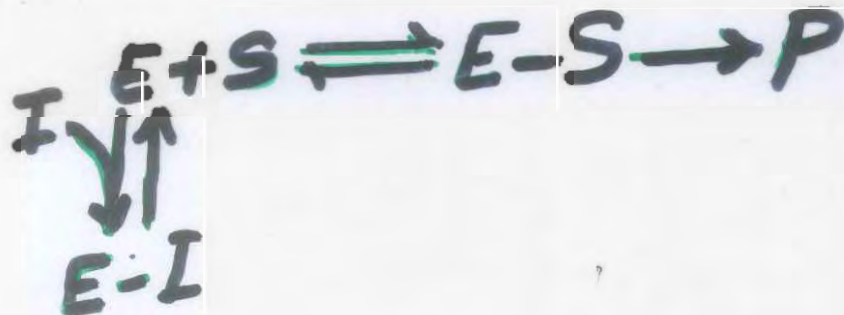
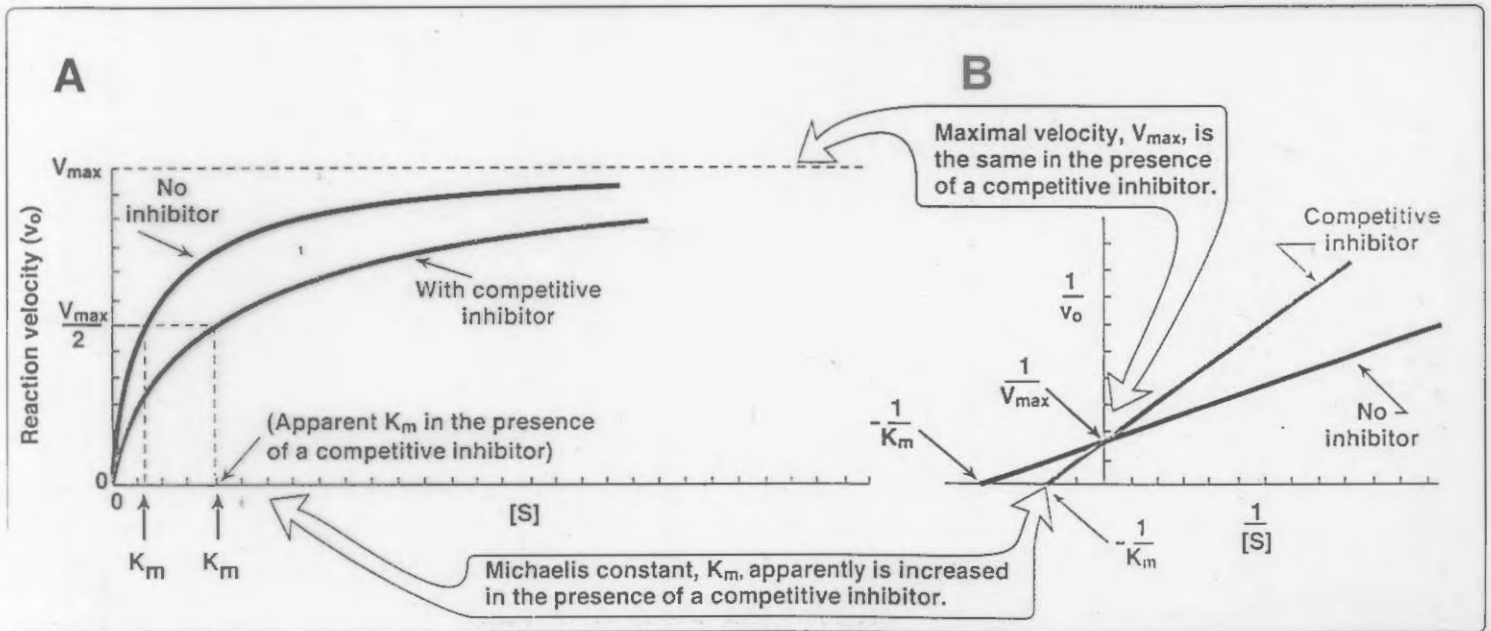
Competitive

non-Competitive

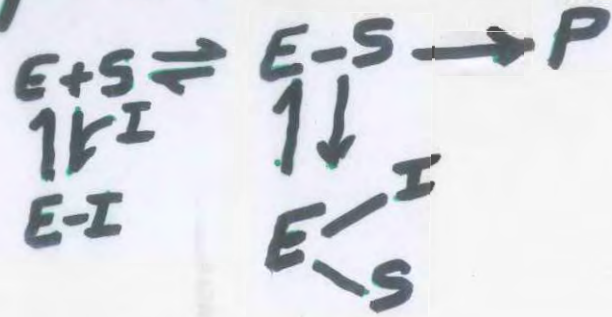
Uncompetitive



# Competitive Inhibitor

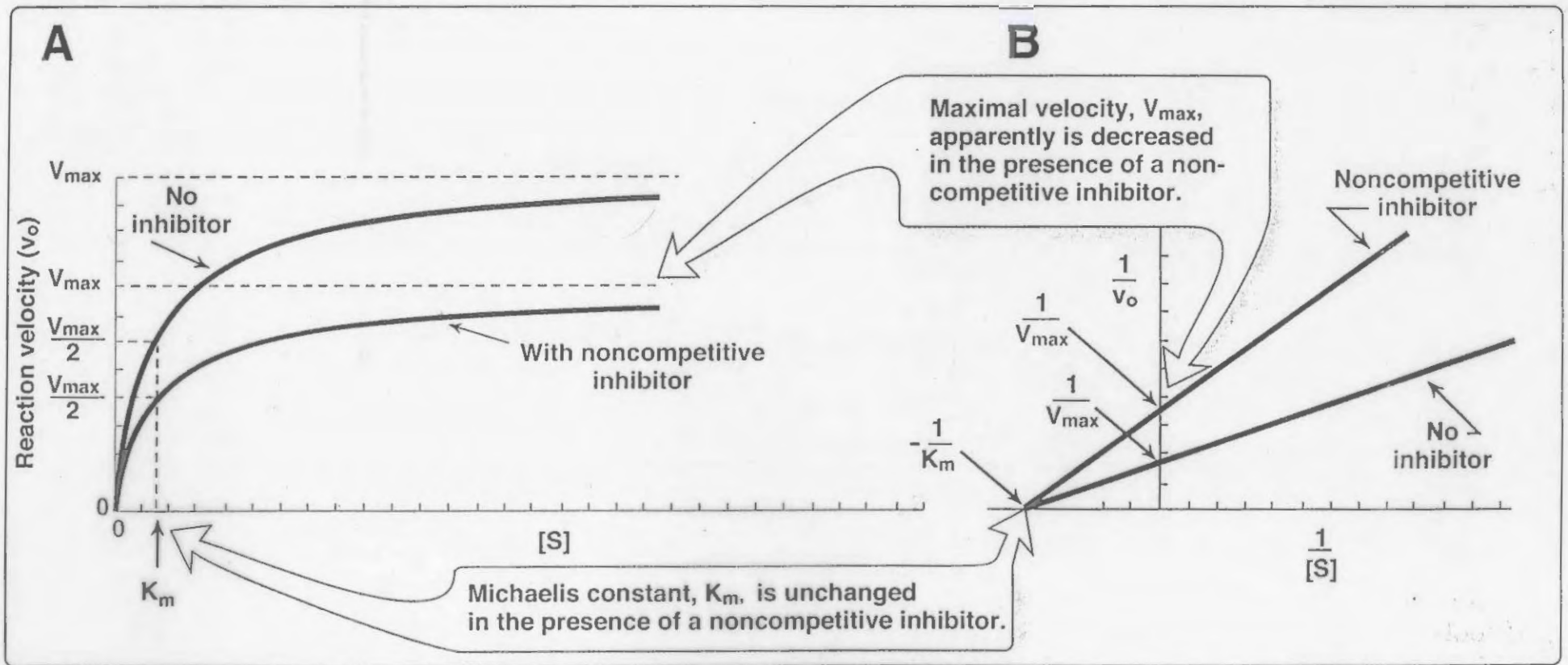


# Noncompetitive Inhibition



8C

*I has similar affinity to E + ES as below. If I has different affinity to E + ES, the lines will intersect above or below the abscissa — change both  $K_m$  +  $V_{max}$*





### III Regulation through Conformational change

#### A. Conformational changes in Allosteric Enzymes

- . Allosteric Enzymes
- . Allosteric sites
- . Allosteric Effectors
  - . Multi subunit
  - . Positive cooperativity
  - . Feedback Inhibition

#### B. Conformational changes from Covalent Modification

- Phosphorylation - dephosphorylation
- other covalent modification e.g. Acetyl ; ADP-ribose or lipid moieties

#### C. Modulator PROTEINS

e.g.  $Ca^{2+}$ -calmodulin  
G-Proteins

#### D. PRECURSOR CLEAVAGE :- Proteolytic cleavage

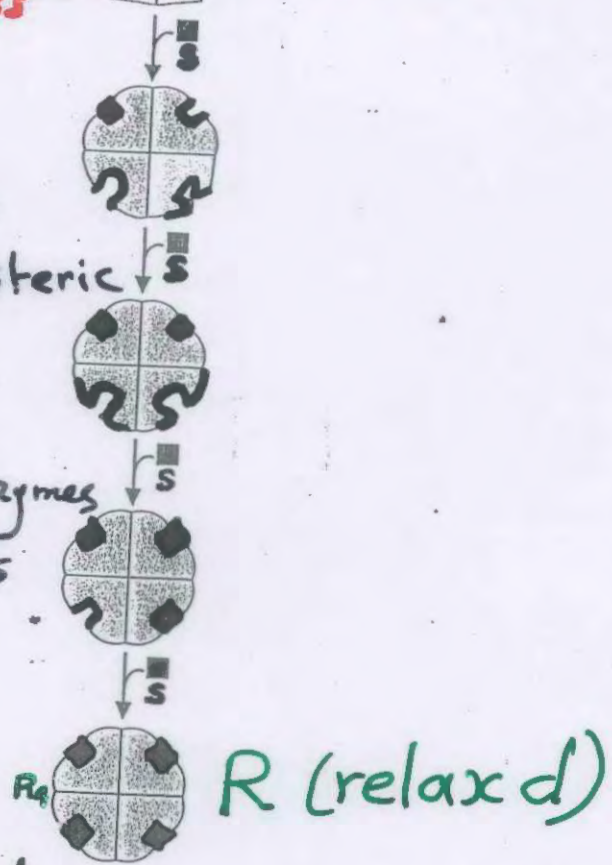
#### IV- AMOUNT OF ENZ. PRESENT

# A sequential model for Allosteric Enzymes

$T_0$   $T$  (taut)

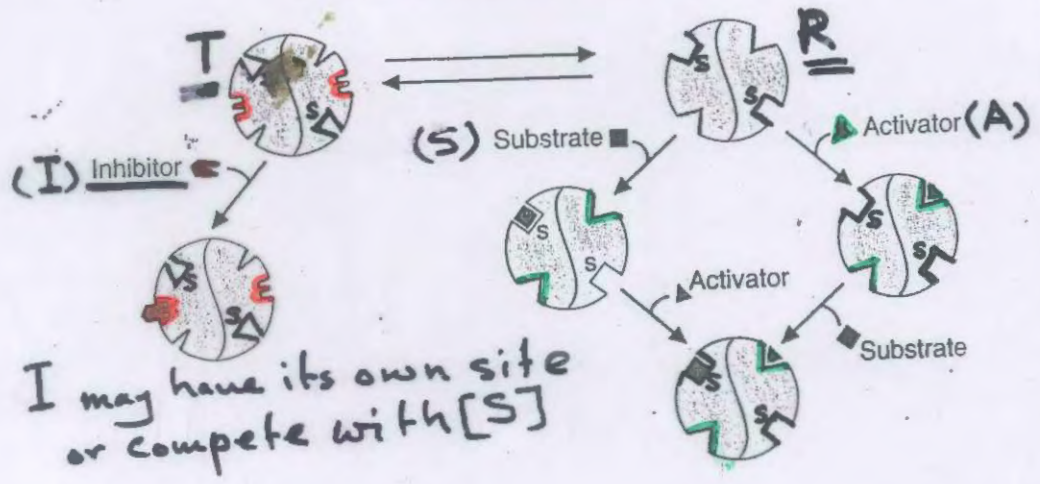
1- Cooperativity in substrate binding to allosteric enzymes

Most allosteric enzymes follow sequential models



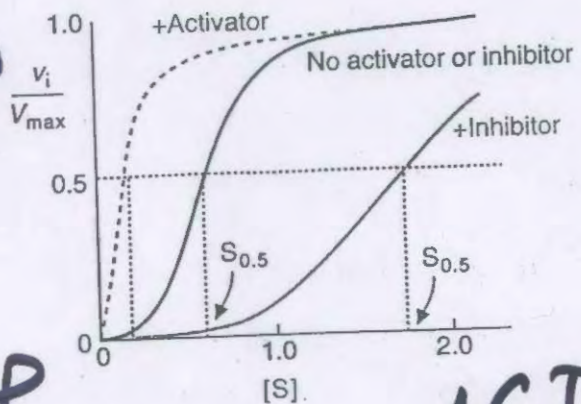
## 2- Allosteric Activators and Inhibitors

A model of an allosteric enzyme



I may have its own site or compete with  $[S]$

e.g. Activation of PFK and GP, ICD by ADP and AMP when ATP  $\downarrow$

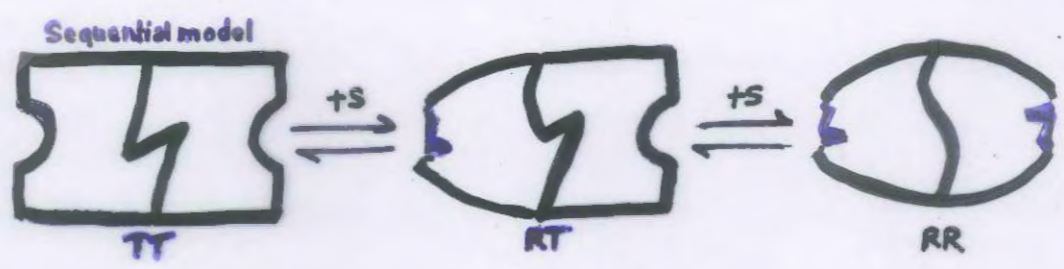


PFK + GP  $\rightarrow$   $\uparrow$  AMP

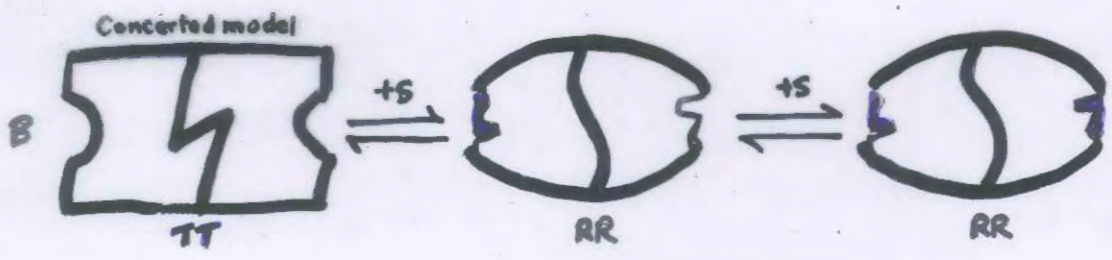
ICD  $\rightarrow$   $\uparrow$  ADP

# Allosteric Models:-

Most Enzymes → A



e.g. Hb → B



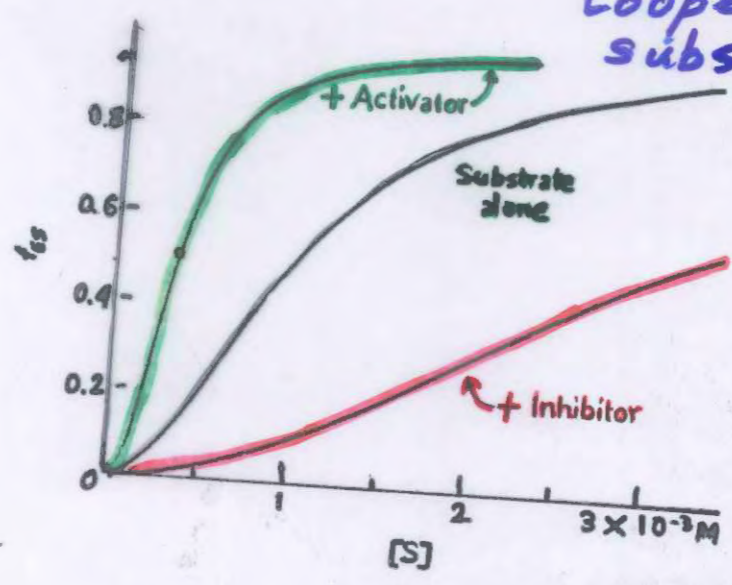
## Allosteric Enzymes

- multi subunits
- Allosteric site
- Reg. & catalytic subunits e.g. ACTase
- Cooperativity in substrate binding



REG. THROUGH CONFORMATIONAL CHANGES :-

Cooperativity in substrate Binding



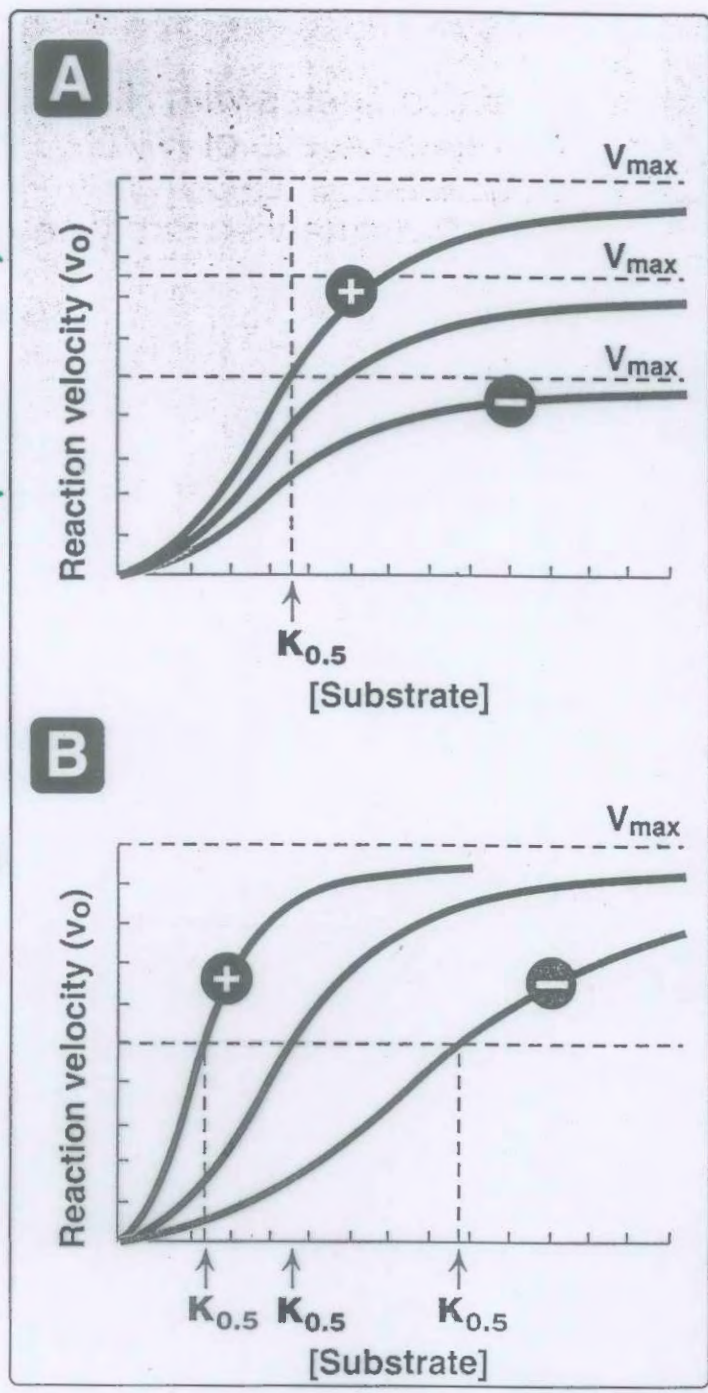
# Advantages of Allosteric Regulation

- Stronger effect than comp. & non-comp. I at the active site

- Allosteric effectors do not occupy active site can be activators

- Effector need not bear resemblance to [S] or [P]

- Rapid as conc. changes e.g. feed-back or signal molecules



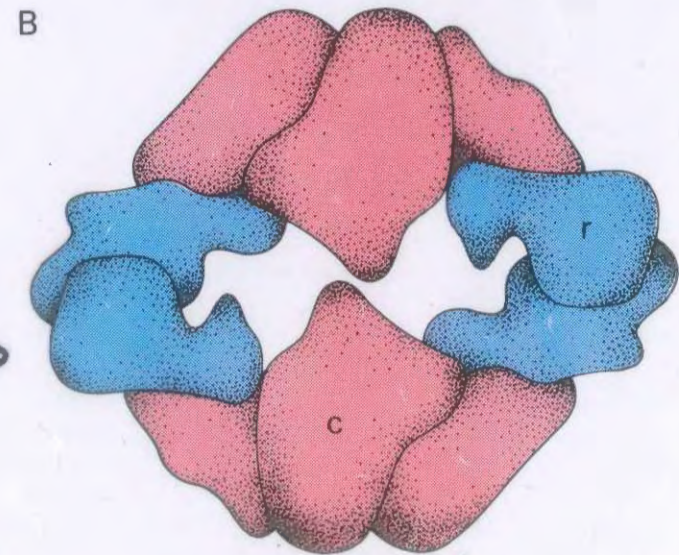
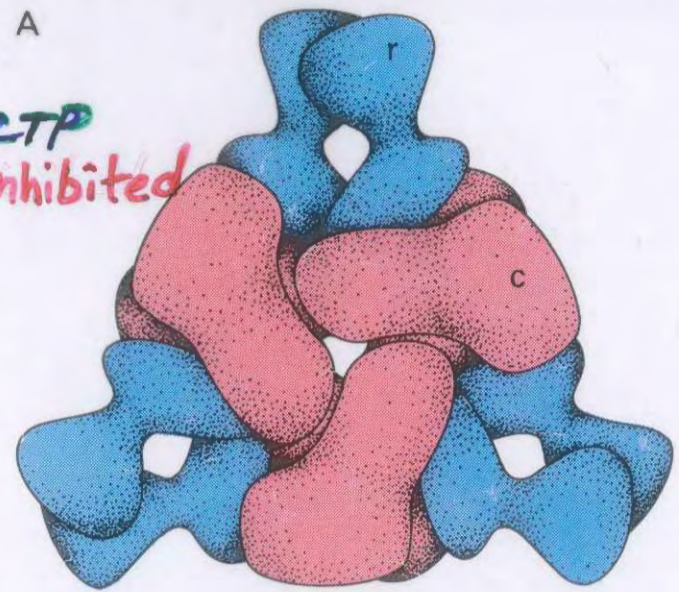
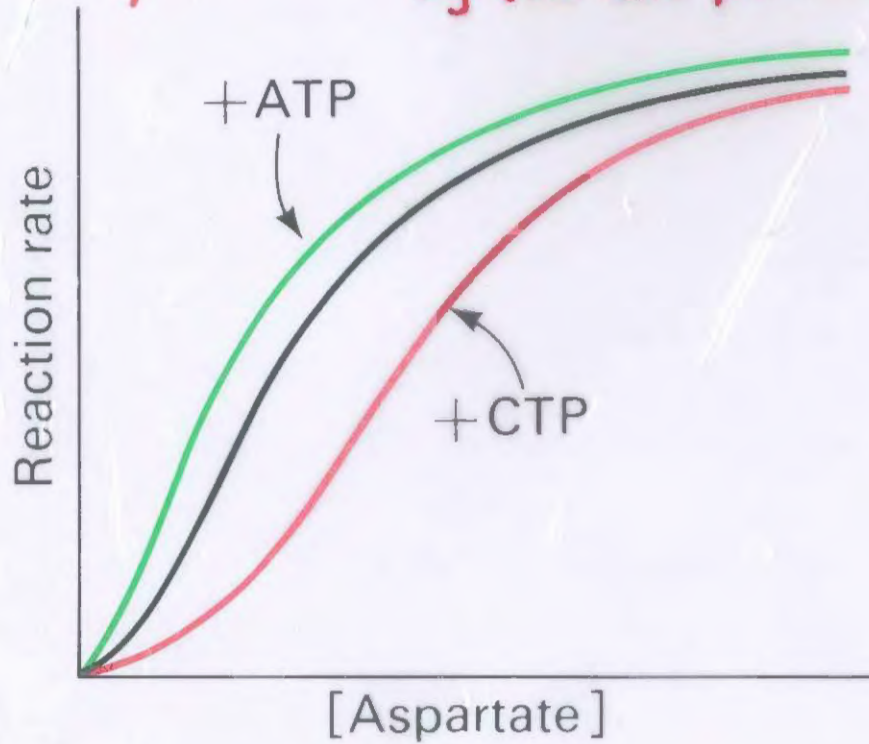
Affecting  $V_{max}$  or  $V_{max}$  &  $K_m$  ( $K_{0.5}$  or  $S_{0.5}$ )

K effectors

Asp + Carbamoyl phosphate  $\xrightarrow{\text{ATCase}}$

N-Carbamoyl-aspartate  $\rightarrow \rightarrow \rightarrow$  CTP

Aspartate Transcarbamoylase is Allosterically inhibited by the end product of its pathway



• Feed back regulation e.g. CTP

• Feed Forward Regulation

e.g. disposal of toxic compounds  
storage pathway e.g. glycogen synthesis

G6P is activator for glycogen synthase

Figures 10-2 and 10-5

Stryer: Biochemistry, Third Edition

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