

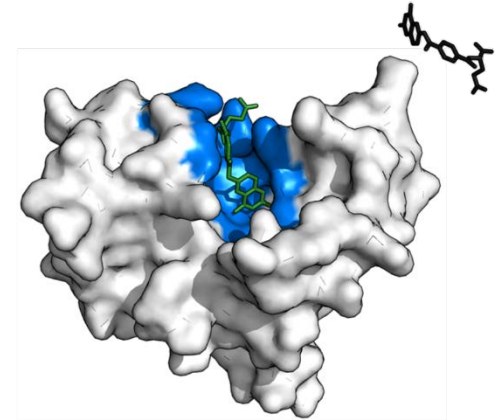
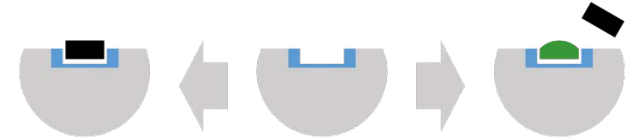
Lecture #6: Enzyme Inhibition and Kinetics

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A. Classification of Inhibitors

1. Irreversible

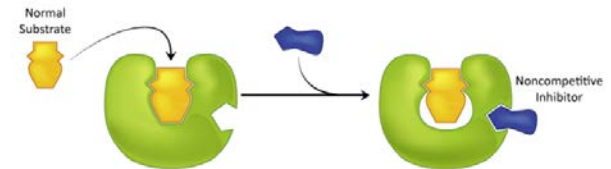
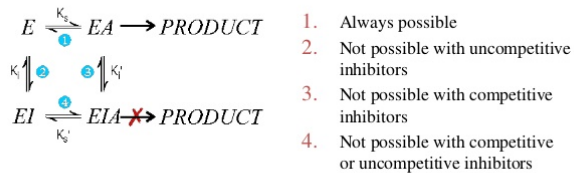
- Iodoacetamide
- DIFP
- Additional examples



2. Classification of Reversible Inhibitors

- Competitive
- Uncompetitive
- Mixed (special case =
- Substrate

Mixed Inhibition



A noncompetitive inhibitor is capable of all four reactions, but the classical noncompetitive inhibitor, as opposed to a mixed one, is a special case. With these inhibitors K_s and K_s' are equal to each other, as are K_i and K_i'

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Enzyme Inhibition

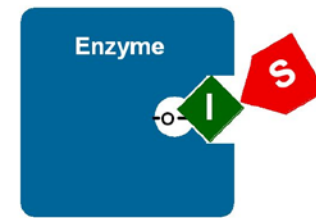
Inhibitors are compounds that decrease an enzyme's activity.

- **Irreversible inhibitors** (inactivators) react with the enzyme.
 - One inhibitor molecule can permanently shut off one enzyme molecule.
 - They are often **powerful toxins** but also may be used as drugs.
- **Reversible inhibitors** bind to and can dissociate from the enzyme.
 - They are often structural analogs of substrates or products.
 - They are often **used as drugs** to slow down a specific enzyme.
- **Reversible inhibitor** can bind to:
 - the free enzyme and prevent the binding of the substrate.
 - the enzyme-substrate complex and prevent the reaction.

1. Irreversible Inhibitor

- binds to the enzyme covalently
 - at the active site or elsewhere
- kinetic effect is like that of non-competitive inhibition
 - net effect is loss of active enzyme
 - time-dependent decrease in enzyme activity as $E + I \rightarrow EI$
- cannot be removed by physical means but may be removed by chemical means
 - breaks covalent bond
- depending upon the chemical treatment
 - enzyme may regain its full initial activity
 - have partial activity
 - or be completely inactive

Irreversible Inhibition



In irreversible inhibition, the inhibitor binds to the enzyme irreversibly through formation of a covalent bond with the enzyme, permanently inactivating the enzyme

Covalent Irreversible Drugs Can Silence Proteins

Reversible inhibitors

Traditional reversible drugs are in equilibrium with their target – continually binding, unbinding, & rebinding

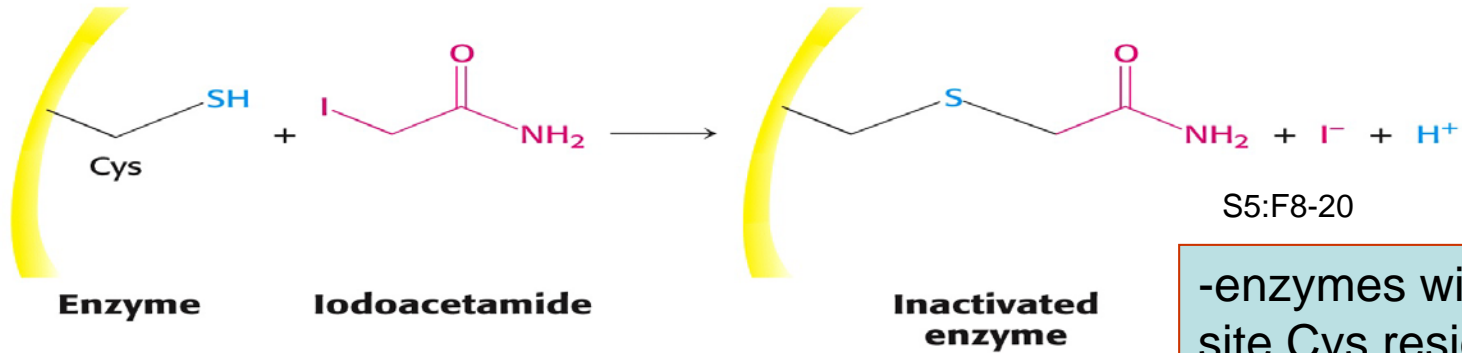
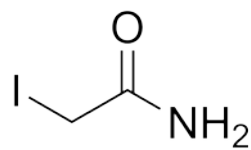


Covalent inhibitors

Covalent irreversible drugs bind specifically to a drug target and form a precisely directed, permanent bond with their target

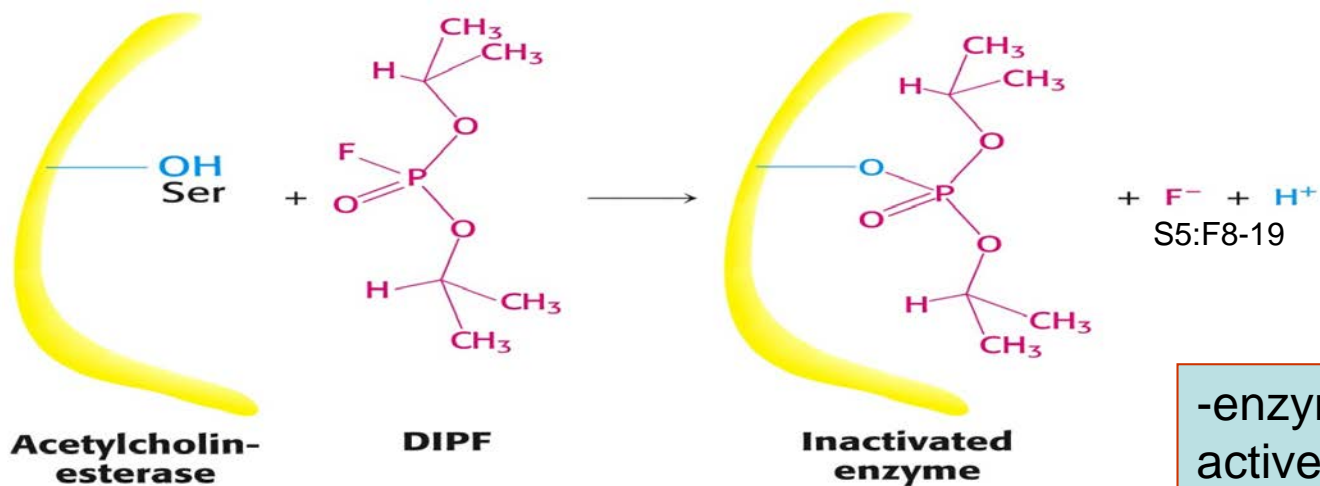
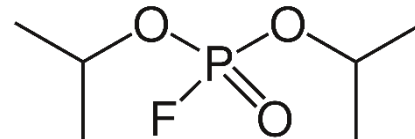


(a) Iodoacetamide



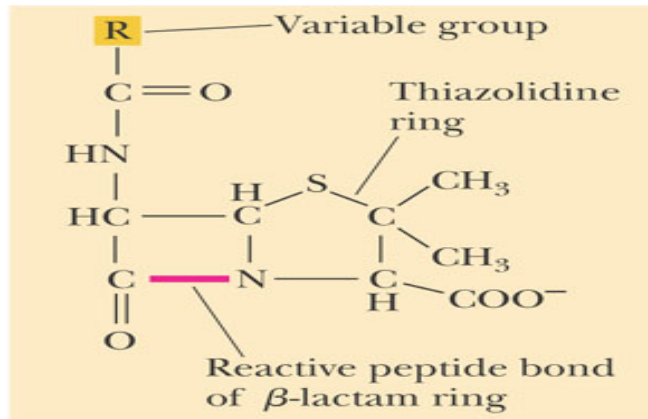
-enzymes with active site Cys residues

(b) DIPF (diisopropylfluorophosphate)

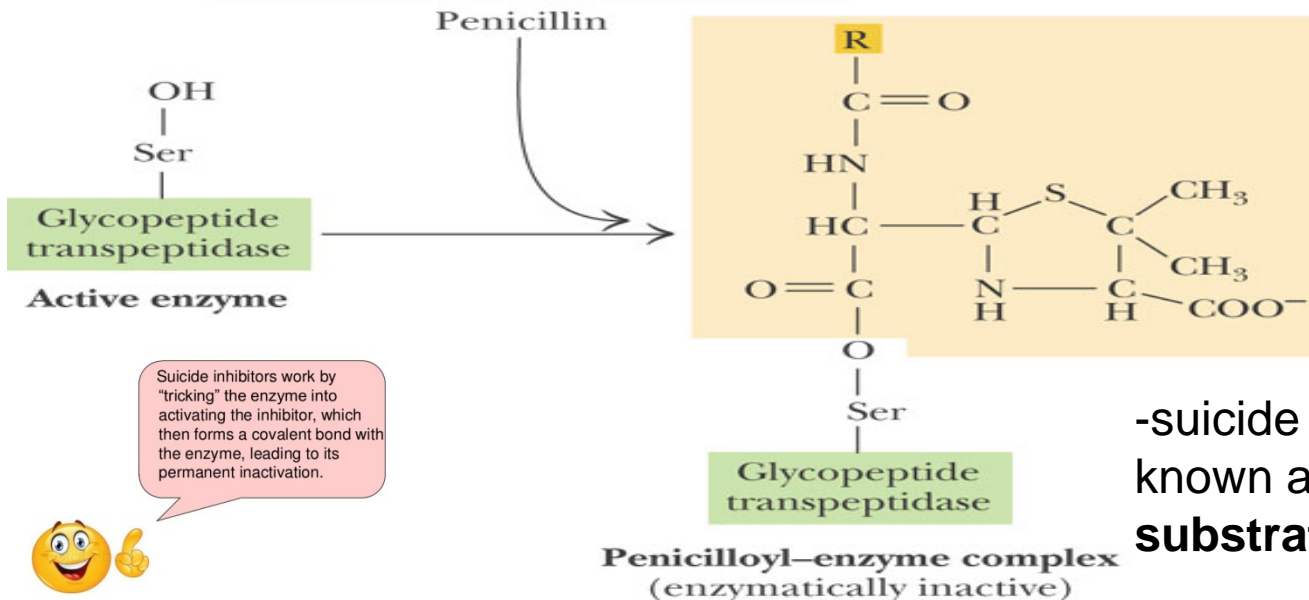


-enzymes with active site Ser

c. Suicide substrates--a substrate that when acted upon by an appropriate enzyme is **converted to product** that essentially **irreversibly inactivates** the enzyme, usually by covalent modification (normal catalytic reaction!)



- penicillin is an irreversible inhibitor of glycopeptide peptidase that is essential for bacterial cell wall synthesis
- conformation of penicillin around its reactive peptide bond resembles the TS of the normal substrate



Suicide inhibitors work by "tricking" the enzyme into activating the inhibitor, which then forms a covalent bond with the enzyme, leading to its permanent inactivation.

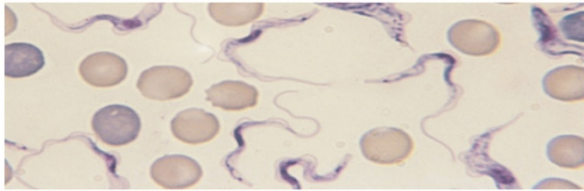


Trojan horse

-suicide substrates are also known as "**Trojan horse substrates**"

Treating African Sleeping Sickness with a Trojan Horse Inhibitor

- African sleeping sickness** is caused by protists called trypanosomes
- vaccines are ineffective because trypanosomes change surface coat proteins continuously
- vulnerable point is polyamine biosynthesis because ornithine decarboxylase (OC) catalyzes the 1st step in polyamine synthesis
- mammalian OC has short $t_{1/2}$ but trypanosome OC is a much more stable enzyme
- reagent that inactivates OC may be able to kill trypanosomes before host cells

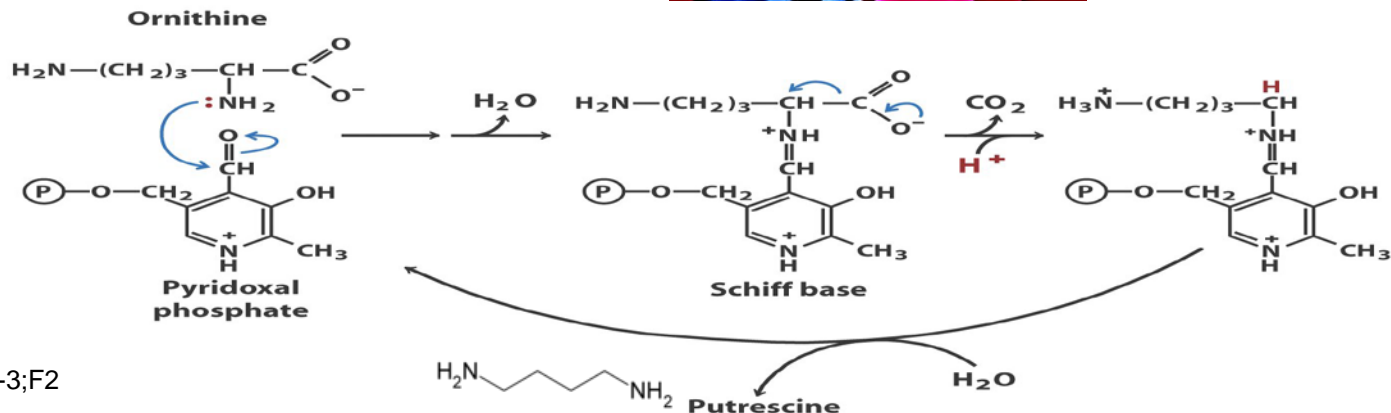


Trypanosoma brucei rhodesiense

L7:Bx-6-3;F1



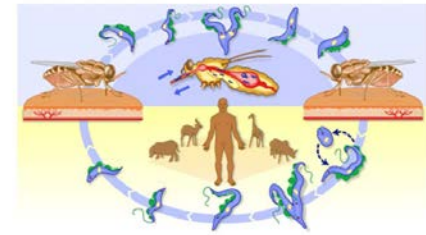
Tsetse fly



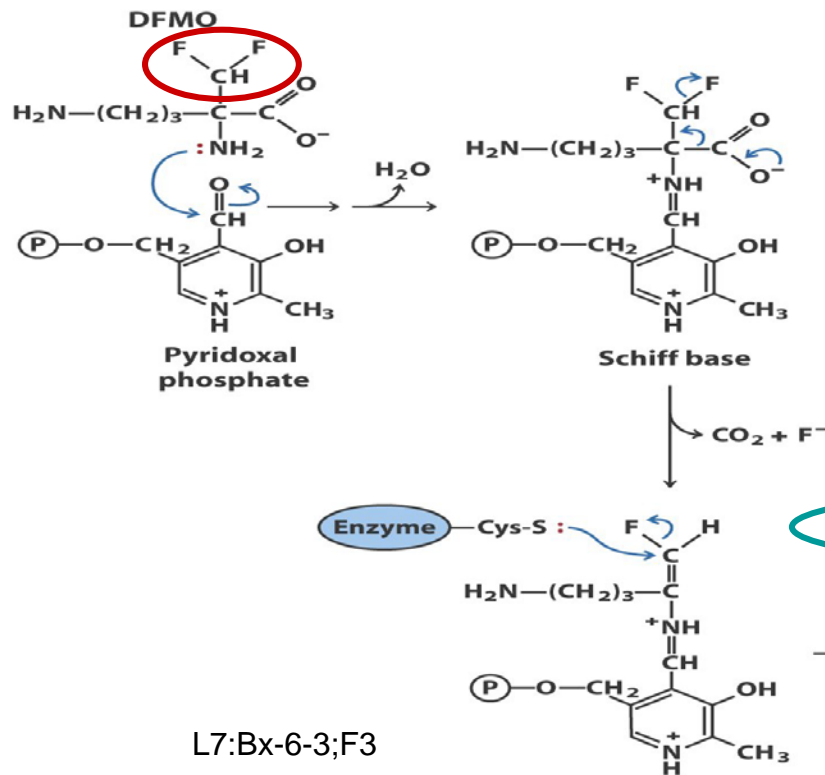
L7:Bx-6-3;F2

African Sleeping Sickness

Trypanosomiasis



A deadly disease spread by the tsetse fly.



- DFMO is a **suicide substrate** that is relatively inert in solution
- when DFMO binds to OC the enzyme is **quickly inactivated**
- inhibitor provides an alternative electron sink in 2 strategically placed **fluorine atoms**, which are excellent leaving groups
- DFMO is now used as a drug to treat African sleeping sickness

Competitive Inhibition

- Inhibitor competes with substrate for binding
 - binds active site
 - does not affect catalysis

Competitive inhibition

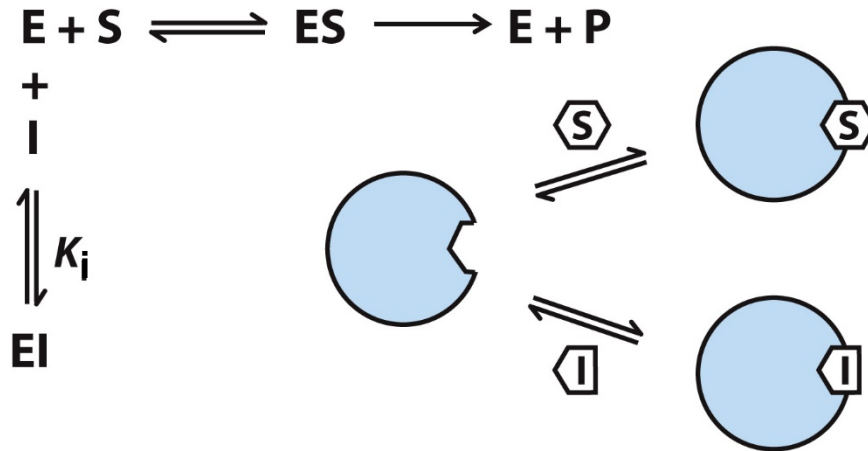


Figure 6-17a
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- No change in V_{\max} ; apparent increase in K_M
- Lineweaver-Burk: **lines intersect at the y-axis.**

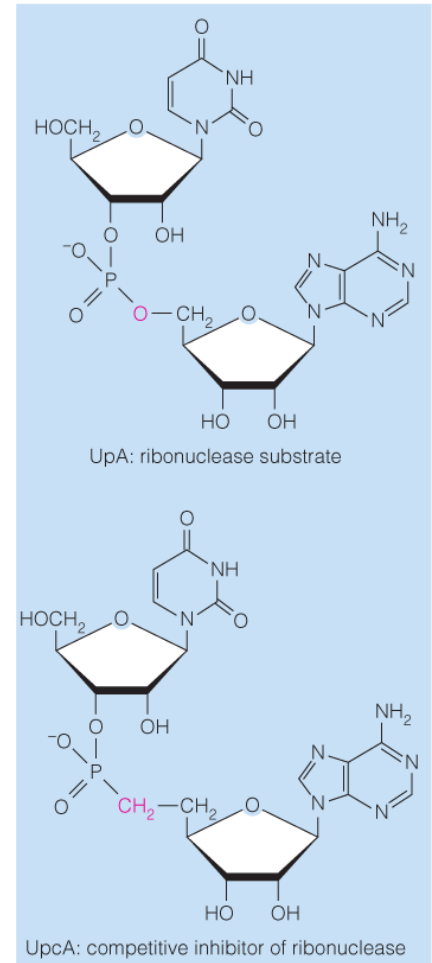


FIGURE 11.31

A substrate and its competitive inhibitor. The substrate UpA and the structurally similar molecule UpcA are competitors for the enzyme ribonuclease. The single difference between the substrate and the inhibitor is shown in magenta.

Competitive Inhibition (Fingerprint)

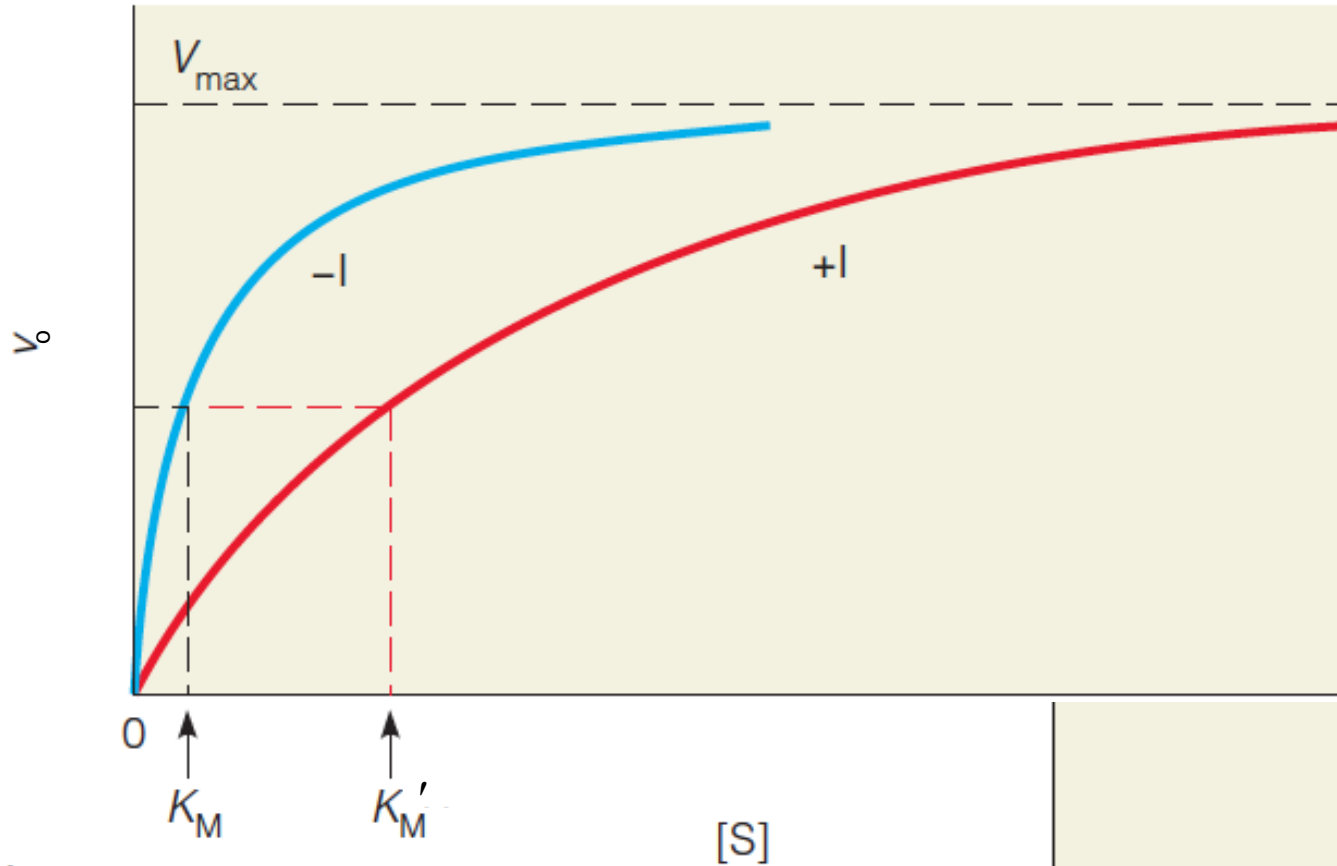
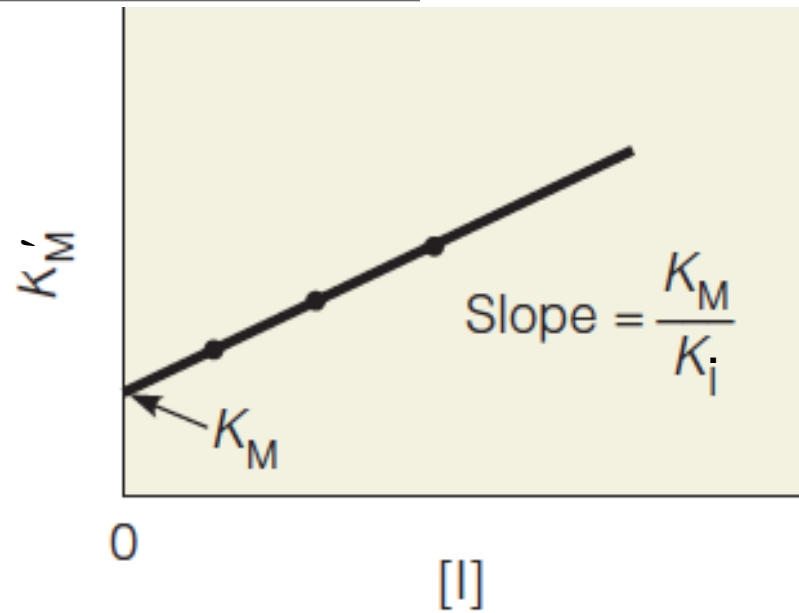


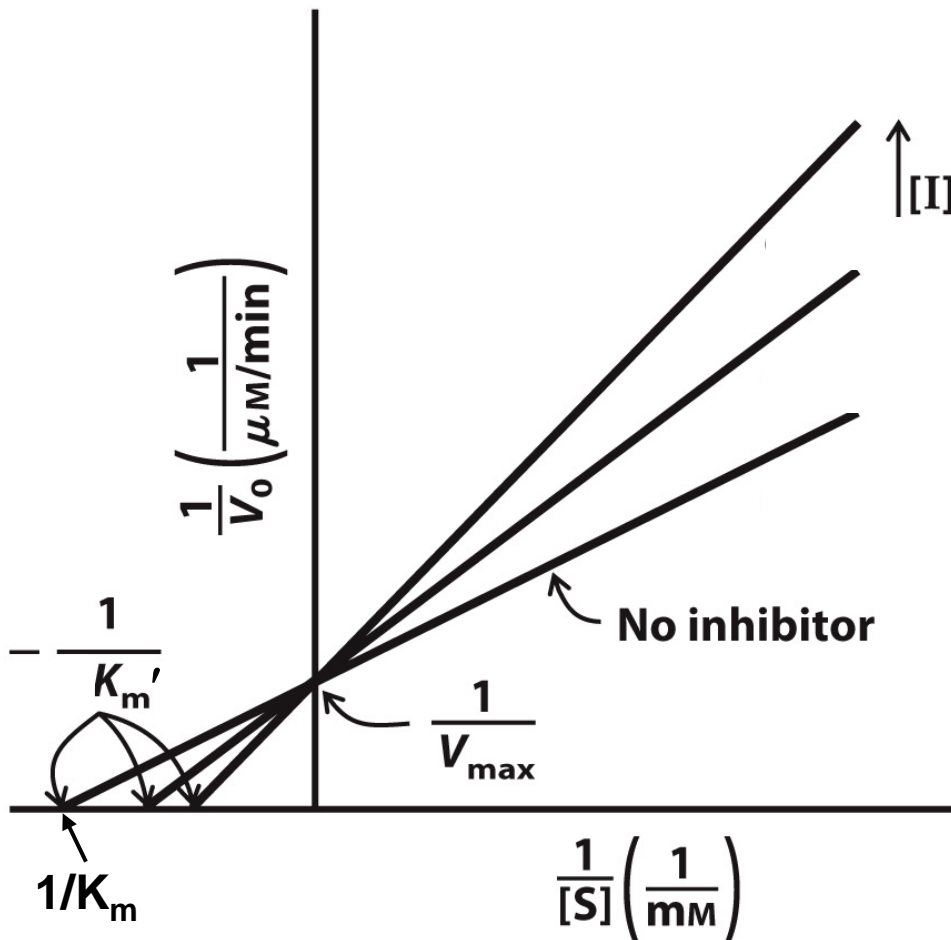
Fig. 11.30
Biochemistry
Matthews 2013

$$K'_M = K_M(1 + [I]/K_i) = K_M + (K_M/K_i)[I]$$



Competitive Inhibition

$$\text{LB: } \frac{1}{V_0} = \left(\frac{K_m'}{V_{\max}} \right) \frac{1}{[S]} + \frac{1}{V_{\max}}$$



Michaelis-Menten Equation:

$$v_0 = \frac{V_{\max} [S]}{K_M + [S]}$$

$$K_i = \frac{[E] + [I]}{[EI]}$$

$K_M' = \text{apparent } K_M$

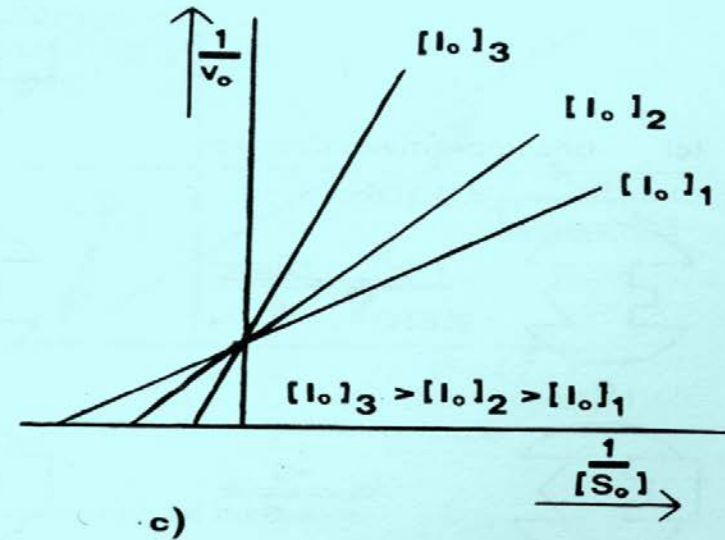
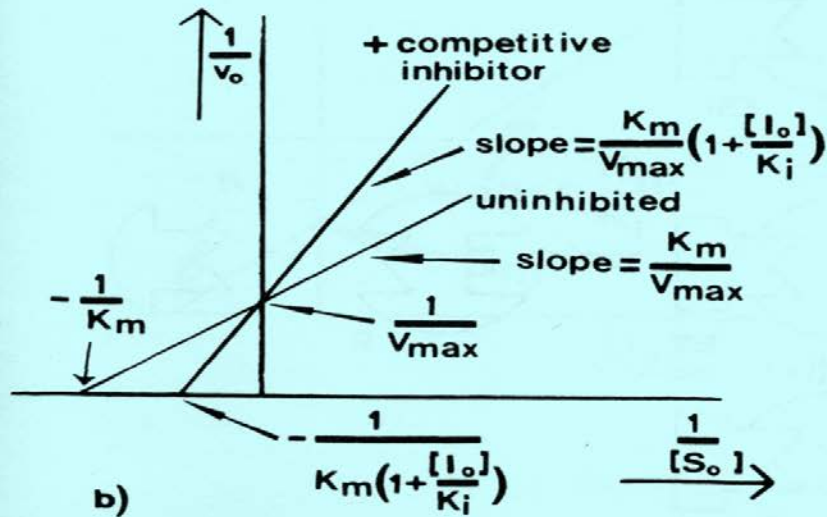
$$K_M' = K_M (1 + [I]/K_i)$$

K_M changes (increases)
but V_{\max} does not change

Box 6-2 Figure 1 modified

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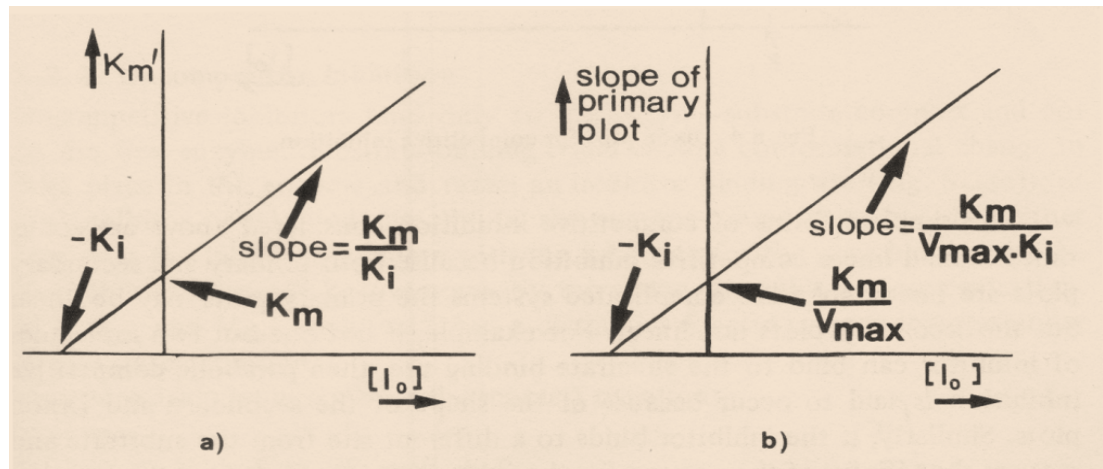
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-to study inhibition and to determine K_i , kinetic studies involve varying $[S_0]$ at several inhibitor concentrations ($[I_0]$)

Secondary Plots

- Graphs of the intercept or slope of the primary plot against $[I_0]$ will also be linear, the intercept on the $[I_0]$ axis giving $-K_i$



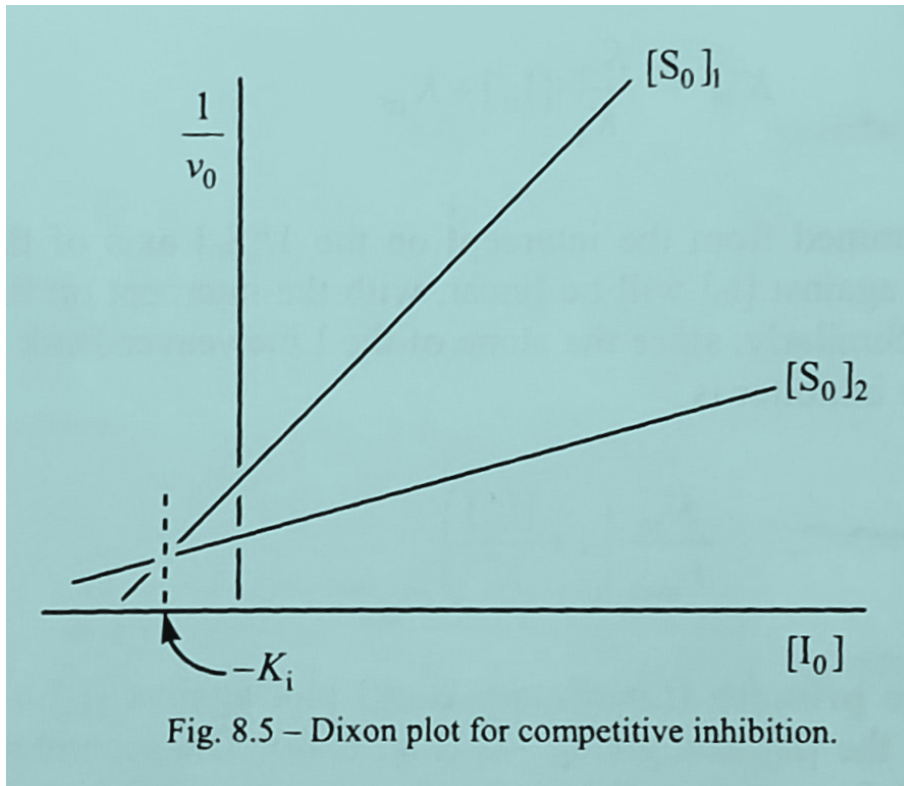
Secondary plots for competitive inhibition.

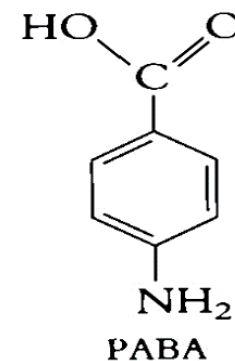
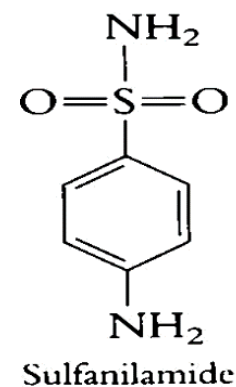
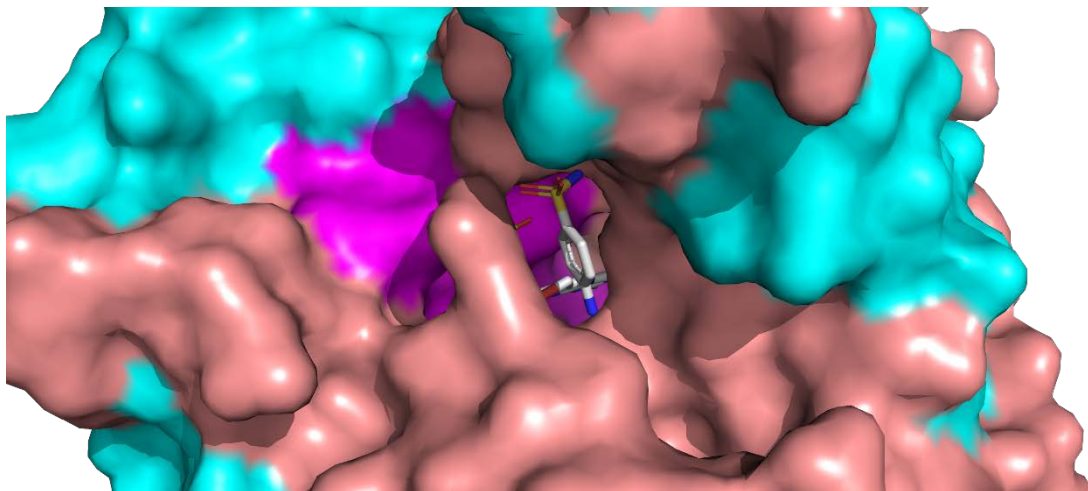
Dixon Plot

- Alternative graphical means of calculating K_i suggested by Dixon (1953)

$$\frac{1}{v_0} = \frac{K_M}{V_{\max} [S_0]} \cdot \frac{[I_0]}{K_i} + \frac{K_M}{V_{\max} [S_0]} + \frac{1}{V_{\max}}$$

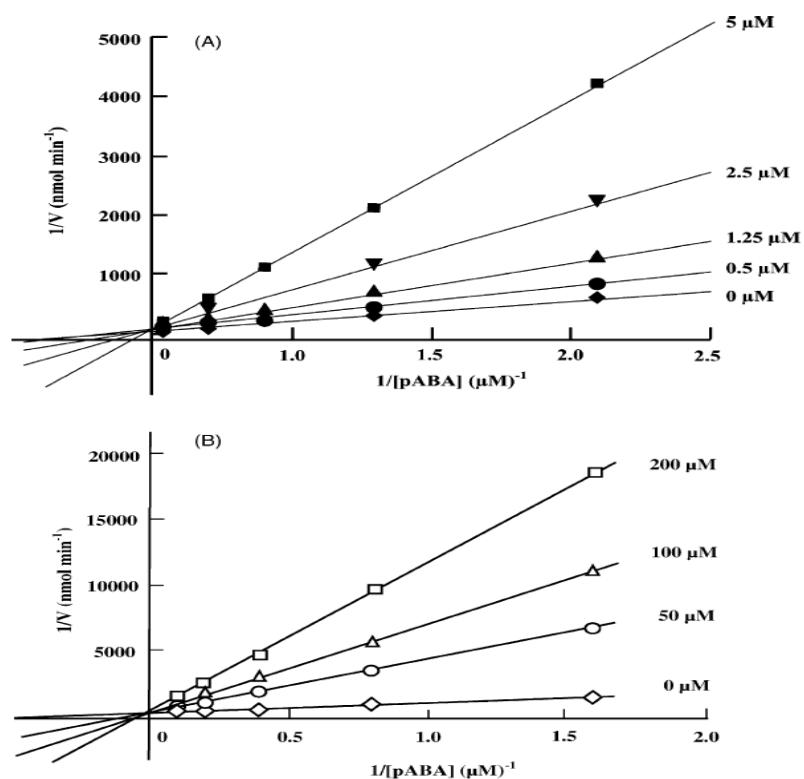
- At fixed $[S_0]$, a plot of $1/v_0$ against $[I_0]$ is linear
- When $[I_0] = -K_i$, then $1/v_0 = 1/V_{\max}$ and is independent of $[S_0]$
- Dixon plots for different $[S_0]$ values (fixed $[E_0]$) intersect where $[I_0] = -K_i$



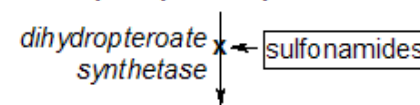


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W. Kasekarn et al. / Molecular & Biochemical Parasitology 137 (2004) 43–53



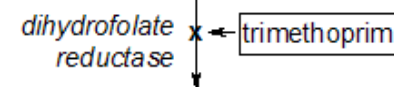
dihydropteroate diphosphate + p-aminobenzoic acid (PABA)



dihydropteroic acid



dihydrofolic acid



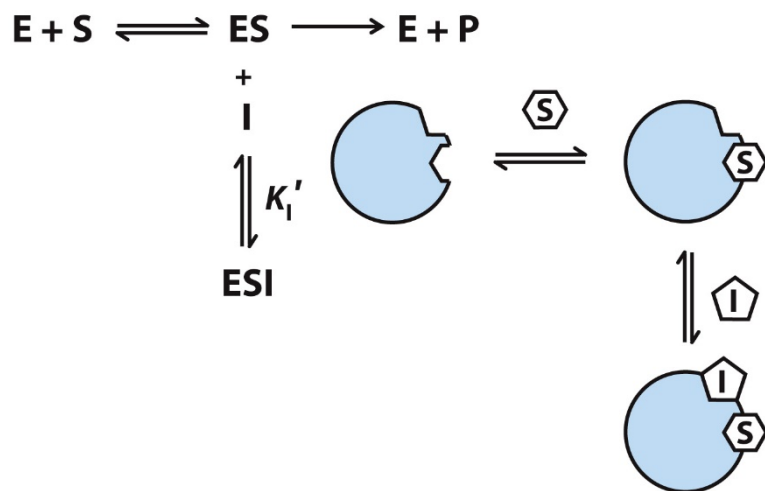
tetrahydrofolic acid

Fig. 5. Lineweaver–Burk plots of *p*/PPPK-DHPS inhibited by sulfa inhibitors. The assays were performed as described in the text in the presence of varying concentrations of *p*ABA while keeping the concentration of H₂PtCH₂OH constant at 20 μM: (A) sulfathiazole and (B) sulfadoxine. The concentrations of the inhibitors (μM) are as specified.

Uncompetitive Inhibition

- Inhibitor only binds to ES complex
 - does not affect substrate binding
 - inhibits catalytic function

Uncompetitive inhibition



Uncompetitive Inhibition Fingerprint

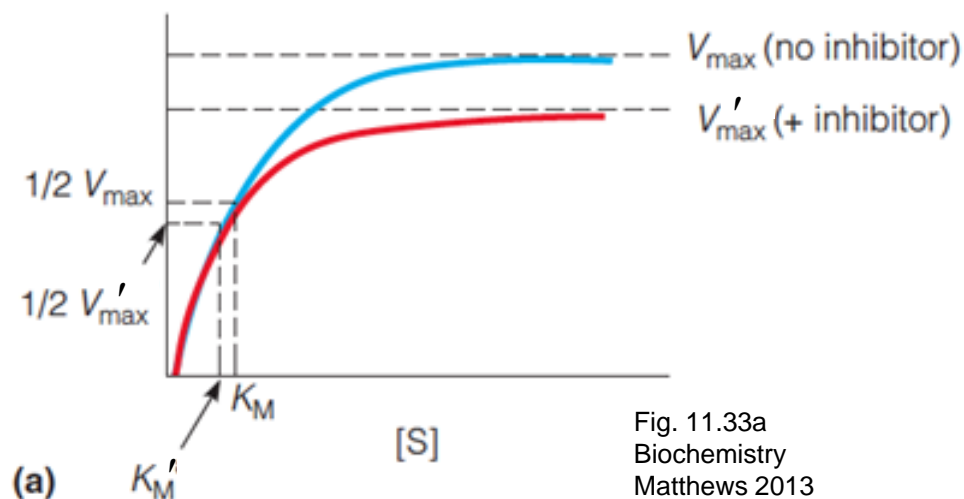


Fig. 11.33a
Biochemistry
Matthews 2013

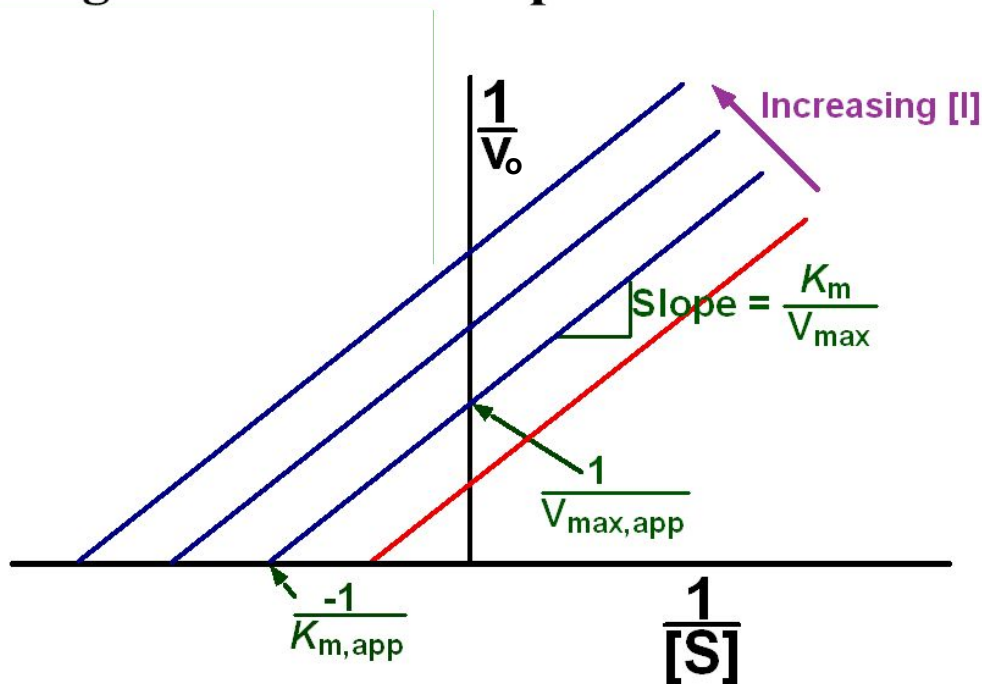
Figure 6-17b
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- Decrease in V_{\max} ; apparent decrease in K_M
- No change in K_M/V_{\max}
- Lineweaver-Burk: **lines are parallel.**

Uncompetitive Inhibition

$$\text{LB: } \frac{1}{v_0} = \frac{K_M'}{V_{\max}'} \left(\frac{1}{[S]} \right) + \frac{1}{V_{\max}'}$$

The Lineweaver-Burk plot is diagnostic for uncompetitive inhibition

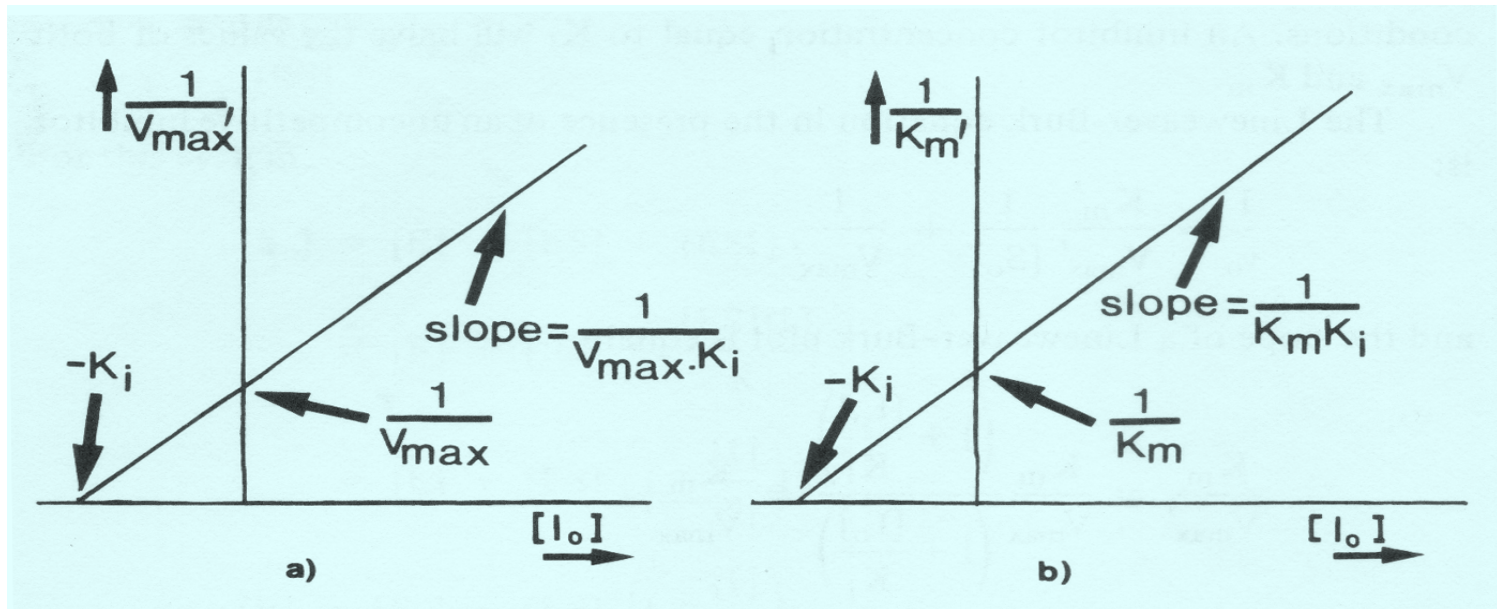


Michaelis-Menten Equation:

$$v_0 = \frac{V_{\max} [S]}{K_M' + [S]}$$

$$K_I = \frac{[ES] + [I]}{[ESI]}$$

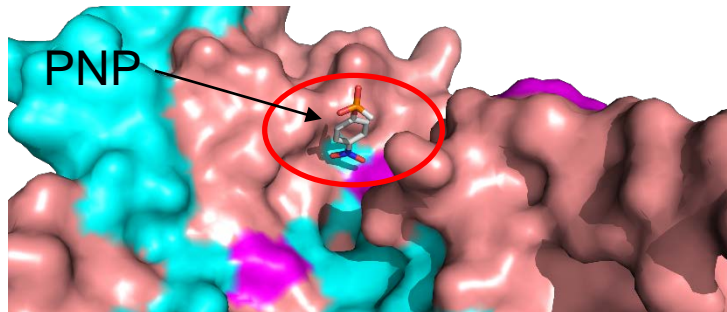
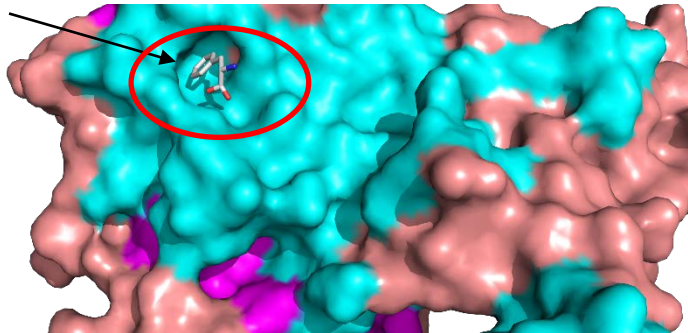
- both K_M and V_{\max} are apparent values (changed)
 - both are smaller than uninhibited values
- $V_{\max}' = V_{\max} / (1 + [I]/K_I)$
- $K_M' = K_M / (1 + [I]/K_I)$
- K_M/V_{\max} is unchanged so lines are parallel



Secondary plots for uncompetitive inhibition.

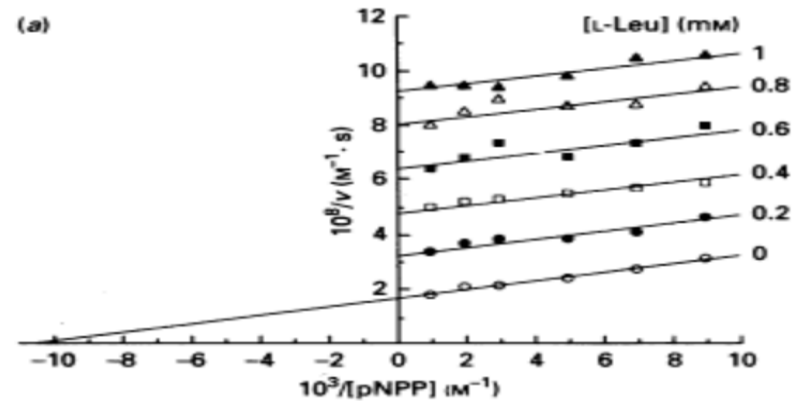
P:F8-7

L-Phe



Biochem. J. (1992) 286:23-30

M. F. Hoylaerts, T. Manes and J. L. Millán



Mixed Inhibition

Mixed inhibition

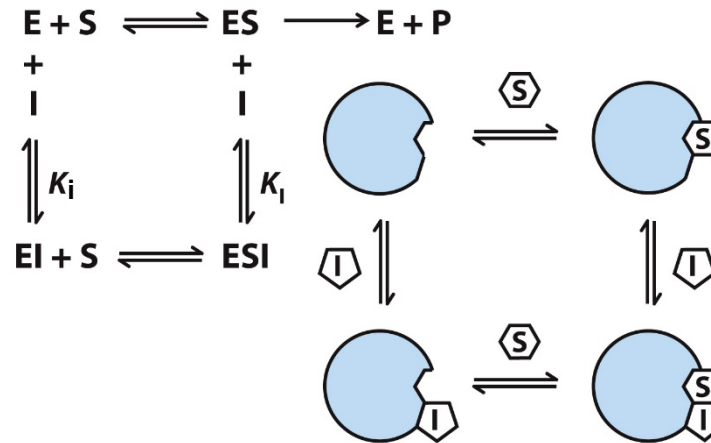


Figure 6-17c
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- Binds enzyme with or without substrate
 - binds to regulatory site
 - inhibits both substrate binding and catalysis
- Decrease in V_{\max} ; apparent change in K_M
- Lineweaver-Burk: **lines intersect left from the y-axis.**
- **Special case:** Noncompetitive inhibitors are mixed inhibitors such that there is no change in K_M

Non-competitive Inhibition

Non-competitive Inhibition:

- special case of mixed inhibition where $K_i = K_i$
- V_{\max} changes but not K_M
- Enzyme, S and E-S are in rapid equilibrium
- inhibitor binds to E and to ES but does **not affect** substrate binding

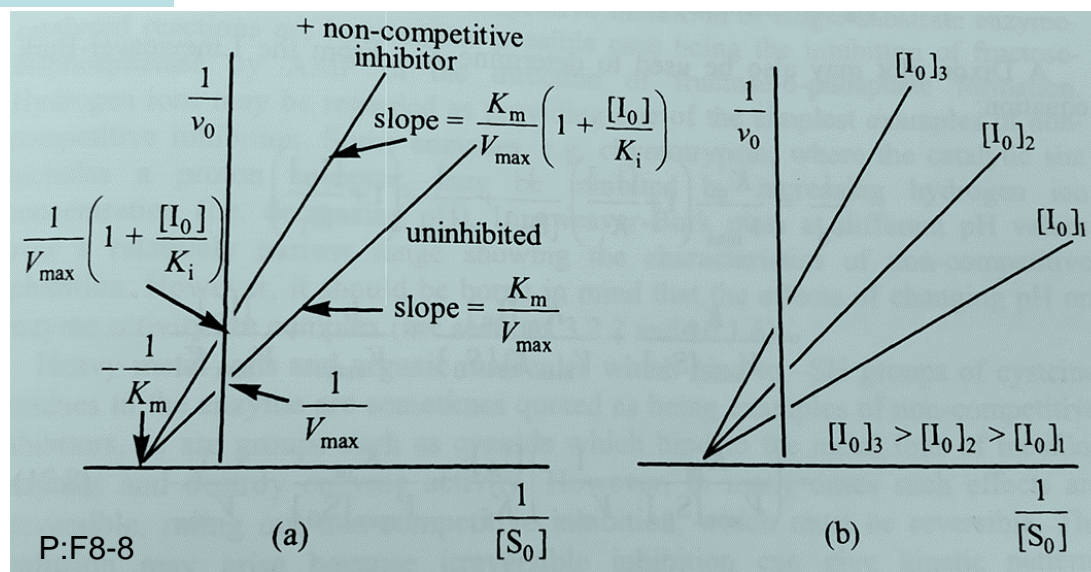
LB:
$$\frac{1}{v_0} = \frac{K_M}{V_{\max}} \left(\frac{1}{[S]} \right) + \frac{1}{V_{\max}}$$

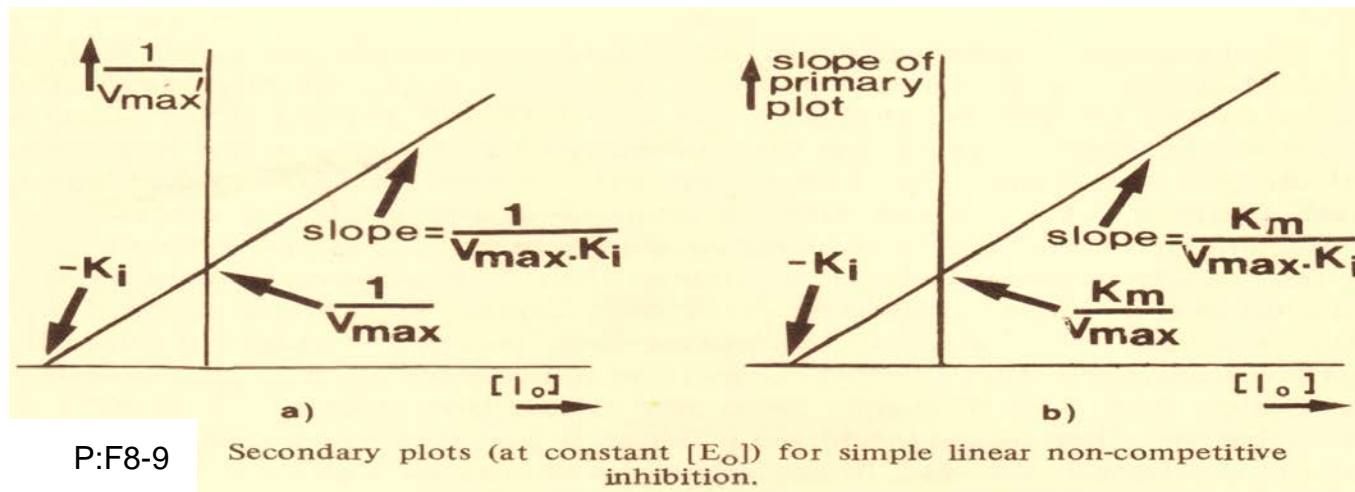
Michaelis-Menten Equation:

$$v_0 = \frac{V_{\max} [S]}{K_M + [S]}$$

$$K_i = \frac{[E][I]}{[EI]} \quad \text{and} \quad K_i = \frac{[ES] + [I]}{[ESI]}$$

$$V_{\max}' = V_{\max} / (1 + [I]/K_i)$$





P:F8-9

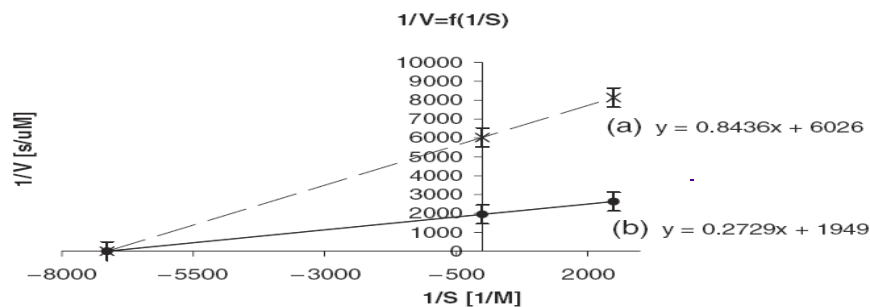
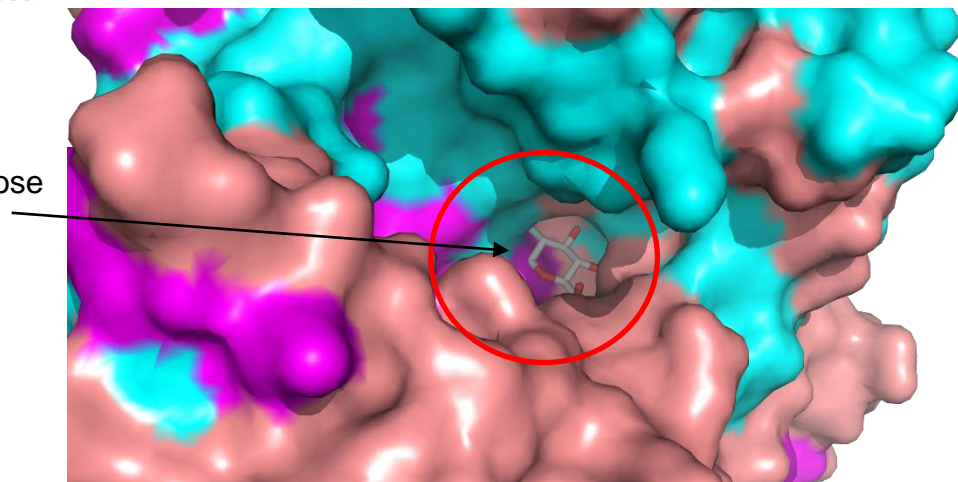


Fig. 6. Kinetics of ONPG hydrolysis by soluble Antarctic β -galactosidase in the presence (a) and absence (b) of glucose.

Makowski et al. (2006)

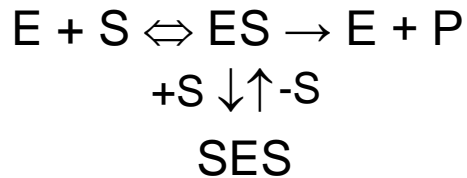
FEMS Microbiol. **59**, 535-542.

2-deoxy-2-fluoro-D-glucose



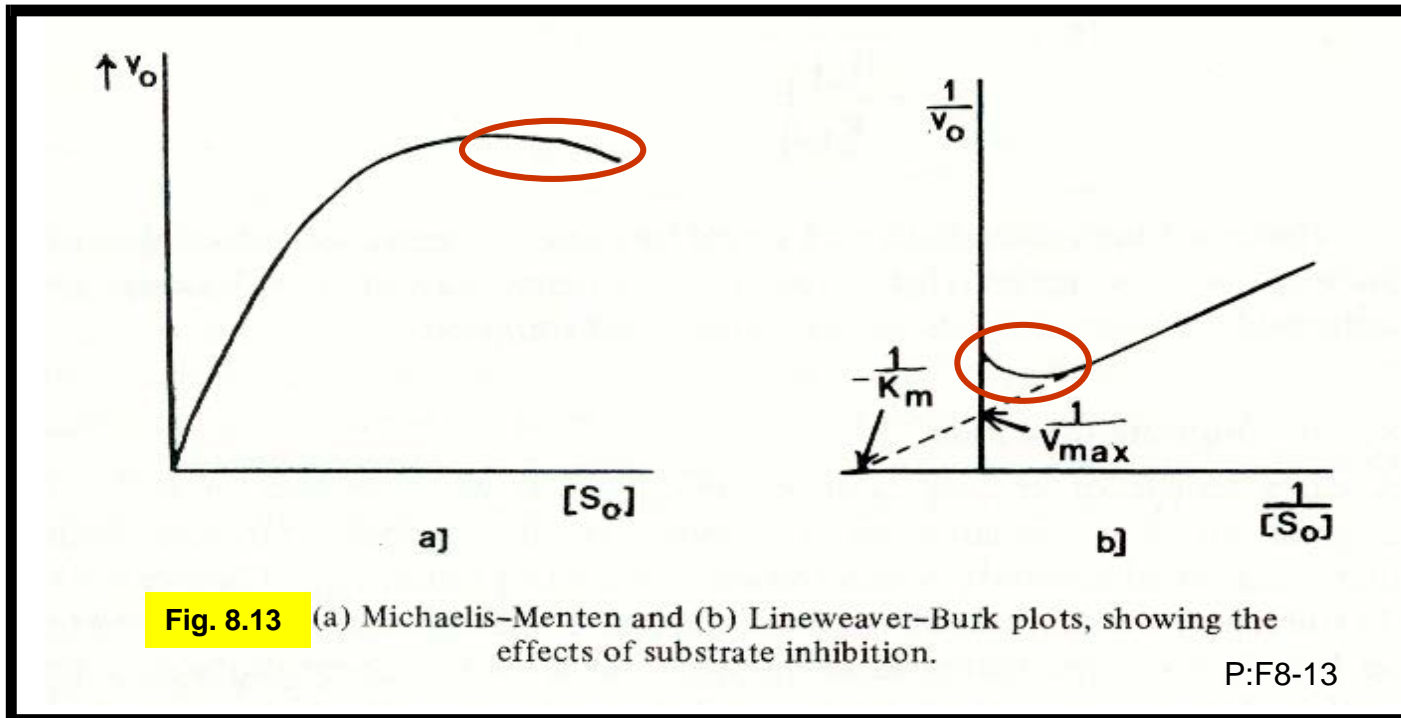
Substrate inhibition

- binding of second substrate molecule to the E results in formation of an inactive complex.



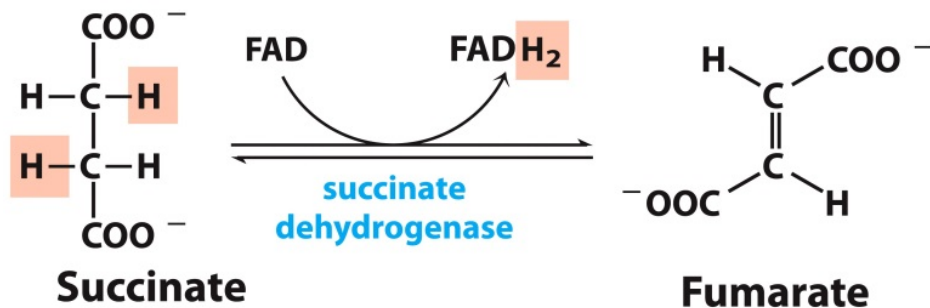
-an $[\text{S}]^2$ term in the rate equation means that it isn't Michaelis-Menten

-can be detected by non-linearity on standard plots





- succinate dehydrogenase shows substrate inhibition with its **succinate** substrate
- at high succinate concentrations above the K_M , the v_o starts to decrease with time
- often caused by active-site crowding



$$\Delta G'^{\circ} = 0 \text{ kJ/mol}$$

Unnumbered 16 p646
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-common occurrence so maximum **amount of substrate** that can be added to assay must be determined