

Reaction Kinetics

- Simple rate equations describe the progress of first-order and second-order reactions.
- The Michaelis-Menten equation relates the initial velocity of a reaction to the maximal reaction velocity and the Michaelis constant for a particular enzyme and substrate.
- An enzyme's overall catalytic efficiency is expressed as k_{cat}/K_M .
- A Lineweaver-Burk plot can be used to present kinetic data and to calculate values for K_M and V_{max} .
- Bisubstrate reactions can occur by an Ordered or Random sequential mechanism or by a Ping Pong mechanism.

Reaction Kinetics

- Enzyme kinetics: Study of the rates of enzyme- catalyzed reaction
- What are the uses of studying kinetics?

What we will cover:

1. Kinetics
2. Inhibition
3. Regulation

Plot of First-Order Rate Equation



The rate V is

1. quantities of A to disappear in a specified unit of time or
2. quantities of P to appear in a specified unit of time

$$V = -\Delta A / \Delta T = \Delta P / \Delta T$$

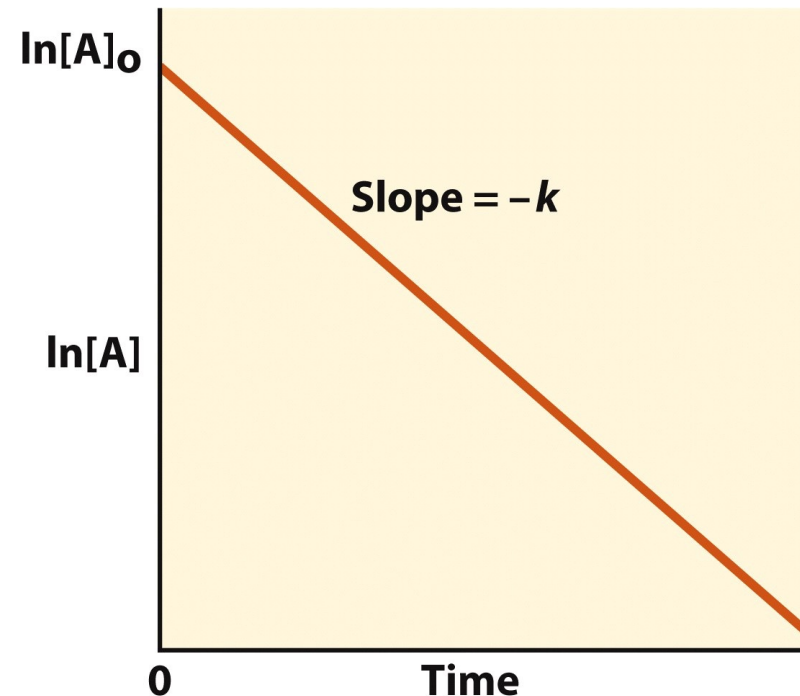
The rate of the reaction is directly related to the concentration of A by a proportionality constant, k .

k = rate constant

$$V = k[A]$$

Reactions that are directly proportional to the reactant concentration are ***first-order reactions***.

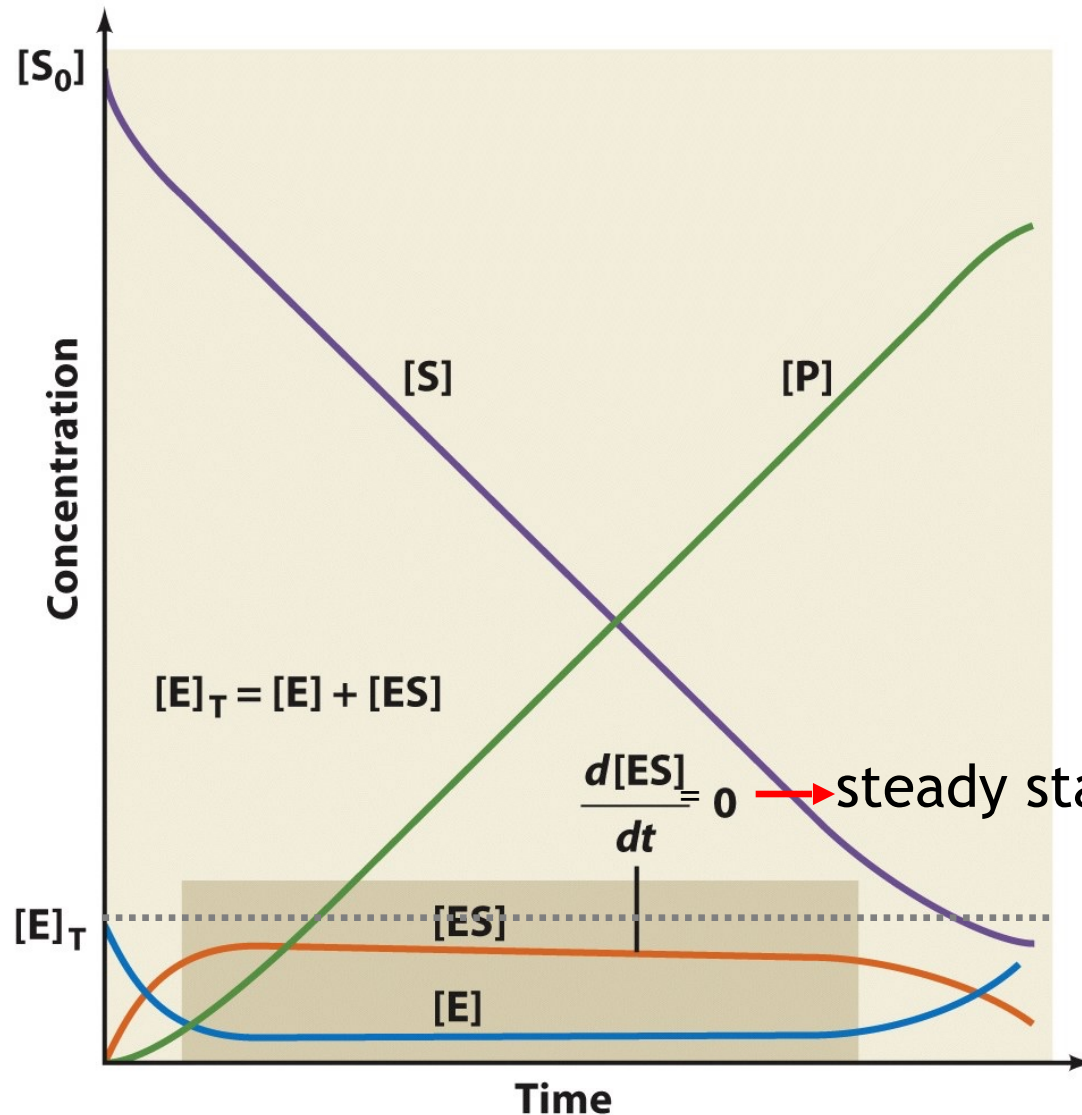
When $2A \longrightarrow P$ or $A + B \longrightarrow P$:
bimolecular reaction.



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- pseudo 1st order reaction
- zero order reaction

Progress Curve: Simple Enzyme-Catalyzed Reaction



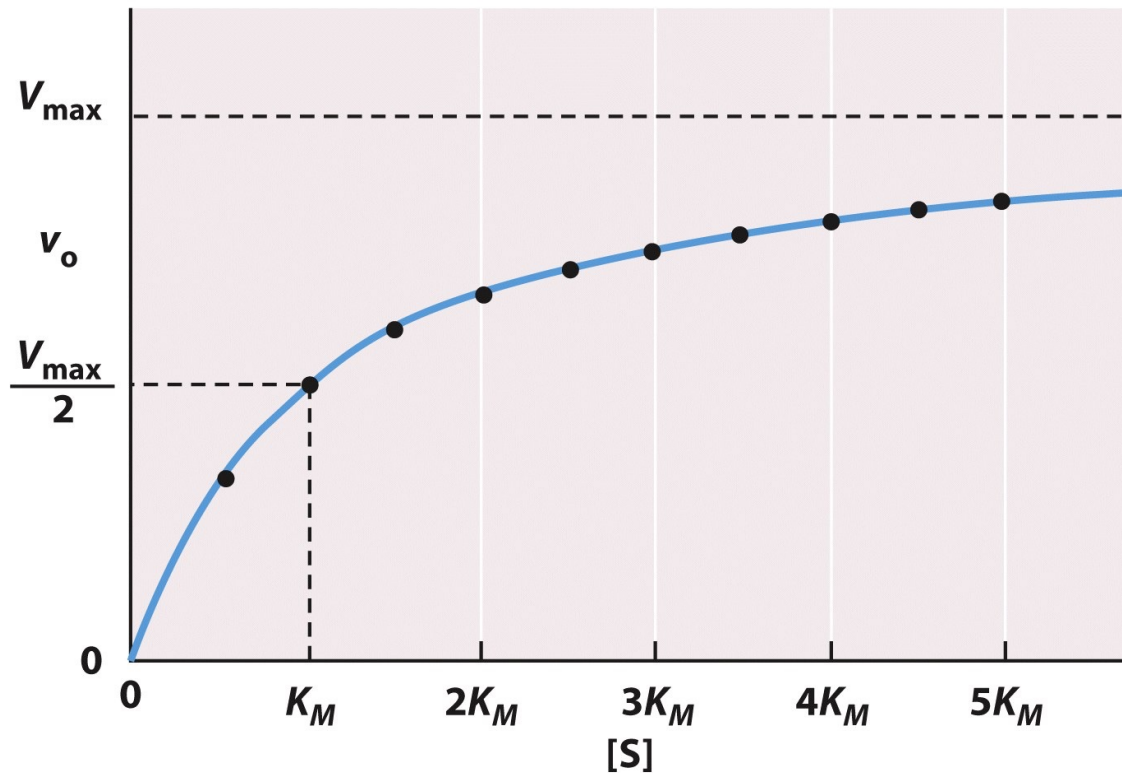
$[S]$: concentration of substrate
 $[P]$: concentration of product
 $[E]$: concentration of enzyme
 $[ES]$: enzyme bound to substrate
 $[E]_T$: concentration of total enzyme

Michaelis-Menten Kinetics



Three assumptions

1. ES is a necessary intermediate step
2. k_{-2} is negligible due to small [P]
3. Steady state: [ES] is a constant independent on [S] or [P]



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M-M equation

1. Finding K_M

$$[E][S]/[ES] = (k_{-1} + k_2)/k_1$$

2. Introduce $[E]_T$

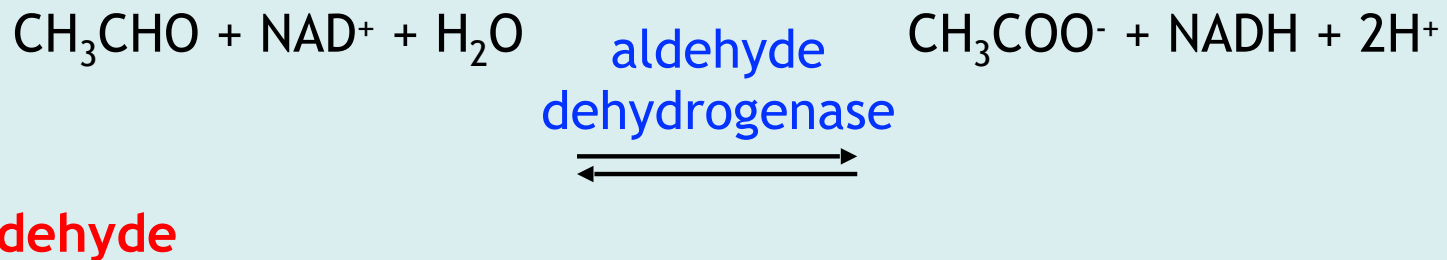
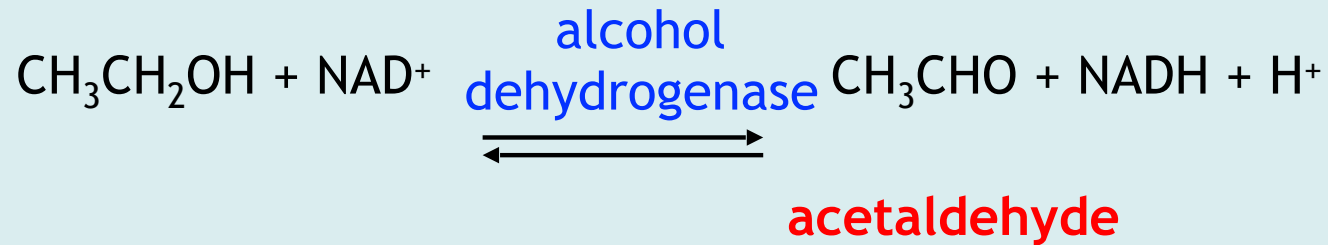
$$[E] = [E]_T - [ES]$$

3. Introduce $V_0 = k_2[ES]$

4. Find V_{max} when $[ES] = [E]_T$

Meaning of K_M

The physiological consequence of K_M in alcohol sensitivity



There are two forms of the AD: a low K_M mitochondrial form and a high K_M cytoplasmic form. What happen to the susceptible people?

Double-Reciprocal (Lineweaver-Burk) Plot

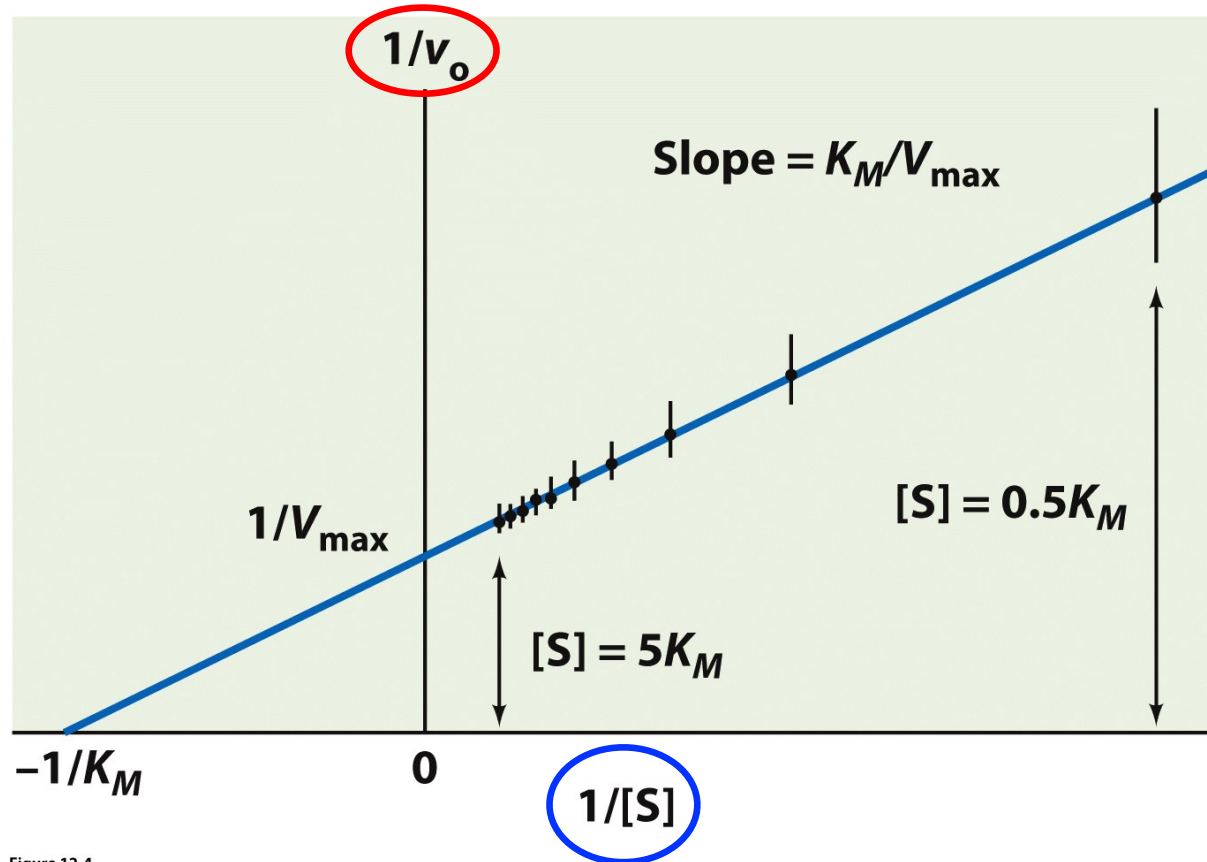


Figure 12-4
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double-reciprocal of MM equation

$$V_0 = V_{\max} \frac{[S]}{[S] + K_M}$$

$$\rightarrow \frac{1}{V_0} = \frac{K_M}{V_{\max}} \cdot \frac{1}{[S]} + \frac{1}{V_{\max}}$$

Enzyme Kinetic Parameters

a measure of catalytic efficiency

TABLE 12-1 The Values of K_M , k_{cat} , and k_{cat}/K_M for Some Enzymes and Substrates

Enzyme	Substrate	K_M (M)	k_{cat} (s^{-1})	k_{cat}/K_M ($M^{-1} \cdot s^{-1}$)
Acetylcholinesterase	Acetylcholine	9.5×10^{-5}	1.4×10^4	1.5×10^8
Carbonic anhydrase	CO ₂	1.2×10^{-2}	1.0×10^6	8.3×10^7
	HCO ₃ ⁻	2.6×10^{-2}	4.0×10^5	1.5×10^7
Catalase	H ₂ O ₂	2.5×10^{-2}	1.0×10^7	4.0×10^8
Chymotrypsin	<i>N</i> -Acetylglycine ethyl ester	4.4×10^{-1}	5.1×10^{-2}	1.2×10^{-1}
	<i>N</i> -Acetylvaline ethyl ester	8.8×10^{-2}	1.7×10^{-1}	1.9
	<i>N</i> -Acetyltyrosine ethyl ester	6.6×10^{-4}	1.9×10^2	2.9×10^5
Fumarase	Fumarate	5.0×10^{-6}	8.0×10^2	1.6×10^8
	Malate	2.5×10^{-5}	9.0×10^2	3.6×10^7
Urease	Urea	2.5×10^{-2}	1.0×10^4	4.0×10^5

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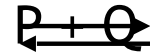
V_{max} : The maximal rate reveals the **turnover number** of an enzyme which is the number of substrate molecules converted into product by **an** enzyme molecule in a unit time when the enzyme is fully saturated with substrate = k_{cat} (= k_2 when the V is maximum)

$$V_{max} = k_{cat} [E]_T$$

Q: a 10^{-6} M solution of carbonic anhydrase catalyzes the formation of 0.6 M H₂CO₃ per second when the enzyme is fully saturated with substrate. What is the k_{cat} ?

Bisubstrate Reactions

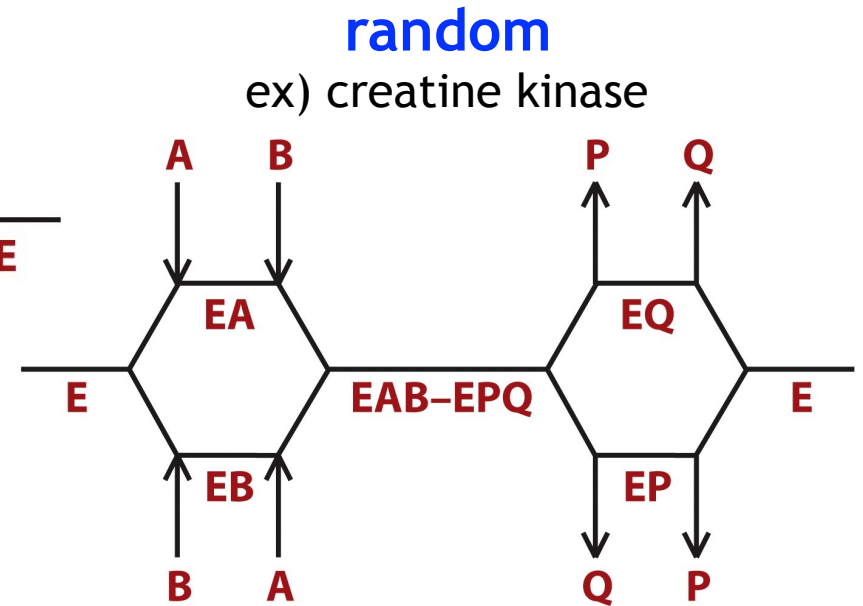
Most biological reactions: $A + B$



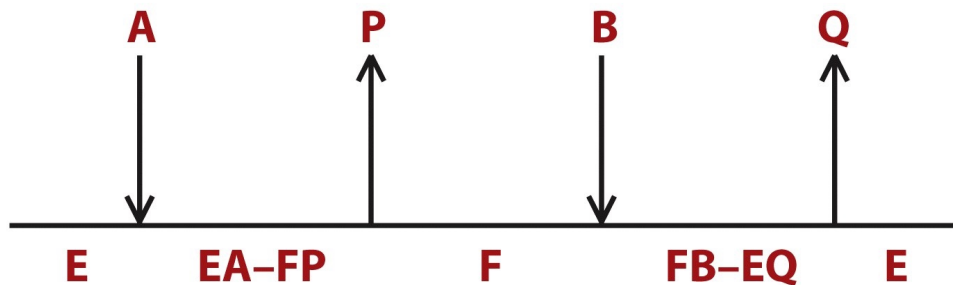
(1) sequential (ordered and random) and (2) ping-pong reaction



Many enzymes use NAD^+ or $NADH$
ex) lactate dehydrogenase



ping pong



: Group-transfer reactions in which one or more products are released before all substrates have been added.

F: enzyme temporarily modified with a functional group

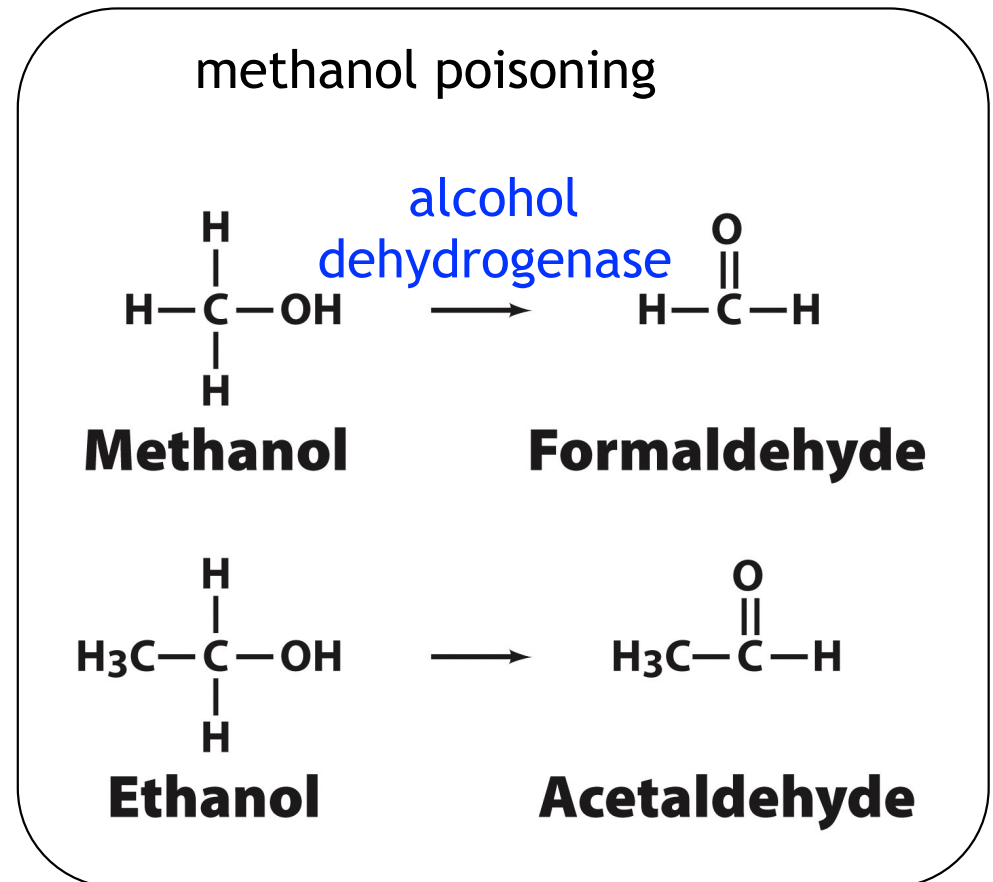
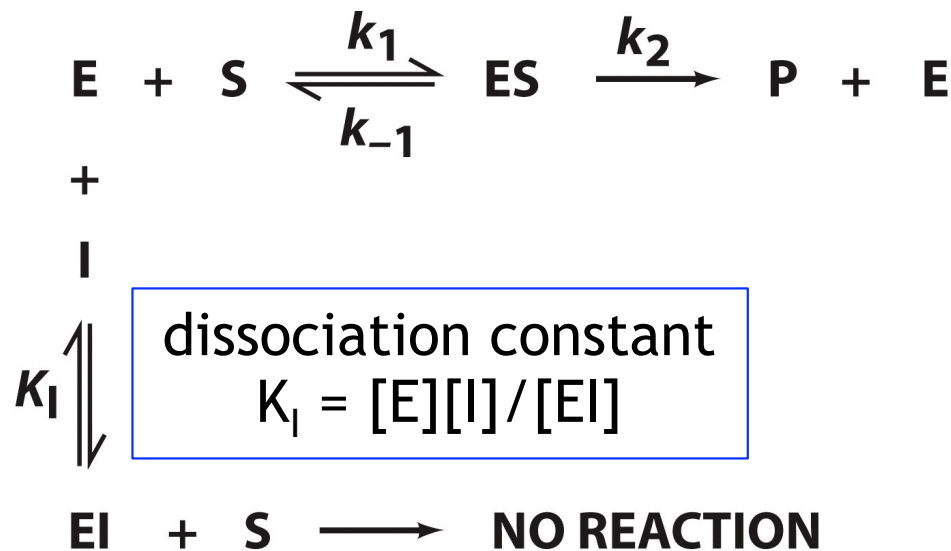
ex: aspartate aminotransferase

Enzyme Inhibition

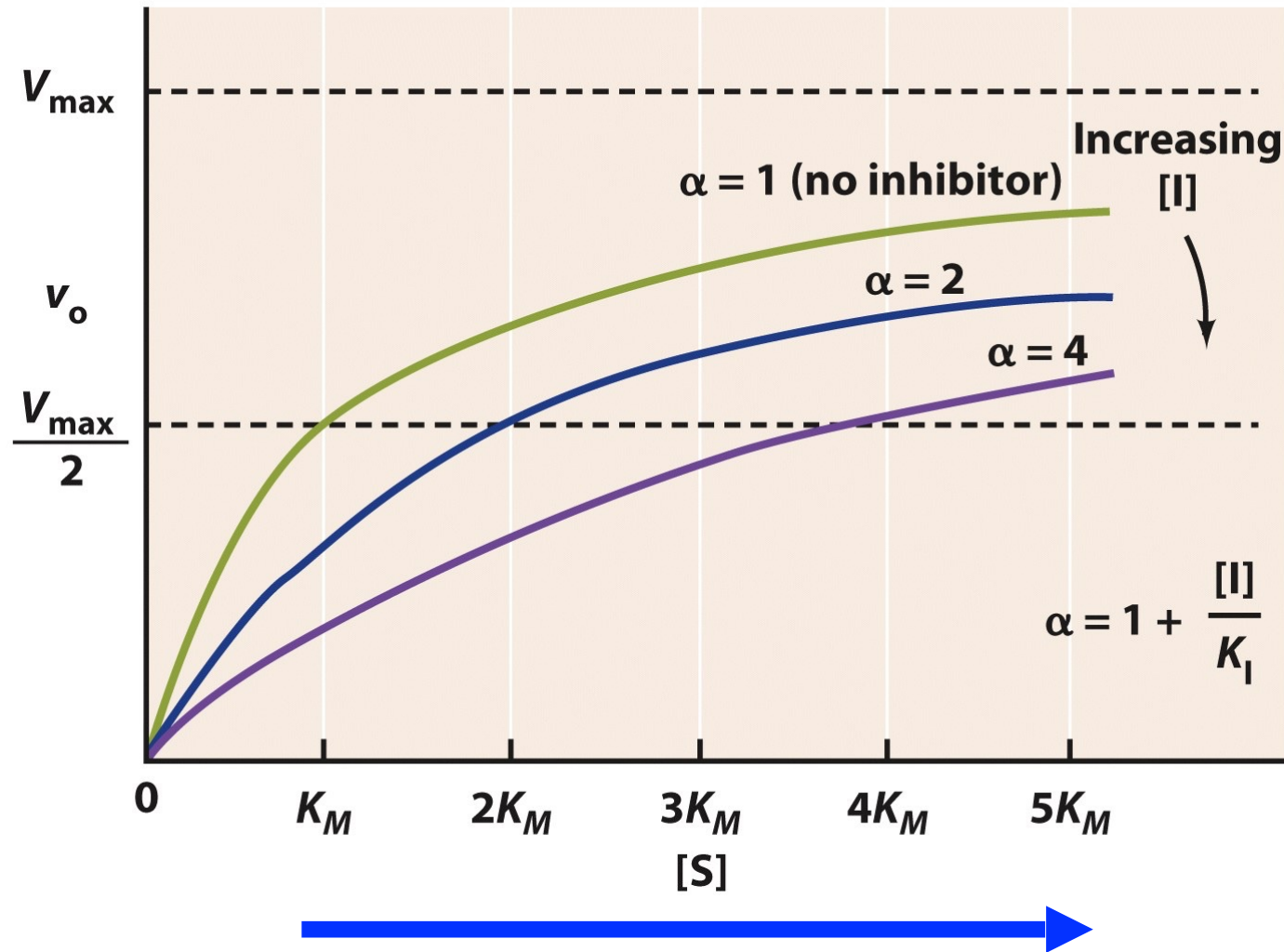
- Enzyme inhibition is a major control mechanism in biological systems.
- Many drugs and toxic agents act as inhibitors (transition state inhibitors!).
- Enzyme inhibitors interact **reversibly** or **irreversibly** (ex: Penicillin and Aspirin) with an enzyme to alter its K_M and/or V_{max} values. Among reversible inhibitors belong competitive and uncompetitive inhibitors.
- **A competitive inhibitor** binds to the enzyme's active site and increases the apparent K_M for the reaction.
- **An uncompetitive inhibitor** binds to ES complex and affects catalytic activity such that both the apparent K_M and the apparent V_{max} decrease.
- **A noncompetitive inhibitor or mixed inhibitor** can bind to free enzymes and reduce the number of available (functional) enzyme: decrease the turnover number. They alters both catalytic activity and substrate binding such that the apparent V_{max} decreases and

Competitive Enzyme Inhibition

- Inhibitors mimic the substrate: compete for the same site.
- The inhibition can be overcome by adding more substrate.
- Often act as drugs: e.g. ibuprofen, Statins



Competitive Enzyme Inhibition



more inhibitors increases K_M

More substrate is needed to obtain the same reaction rate.

The apparent $K_M = K_M(1 + [I]/K_I)$

Competitive Enzyme Inhibition

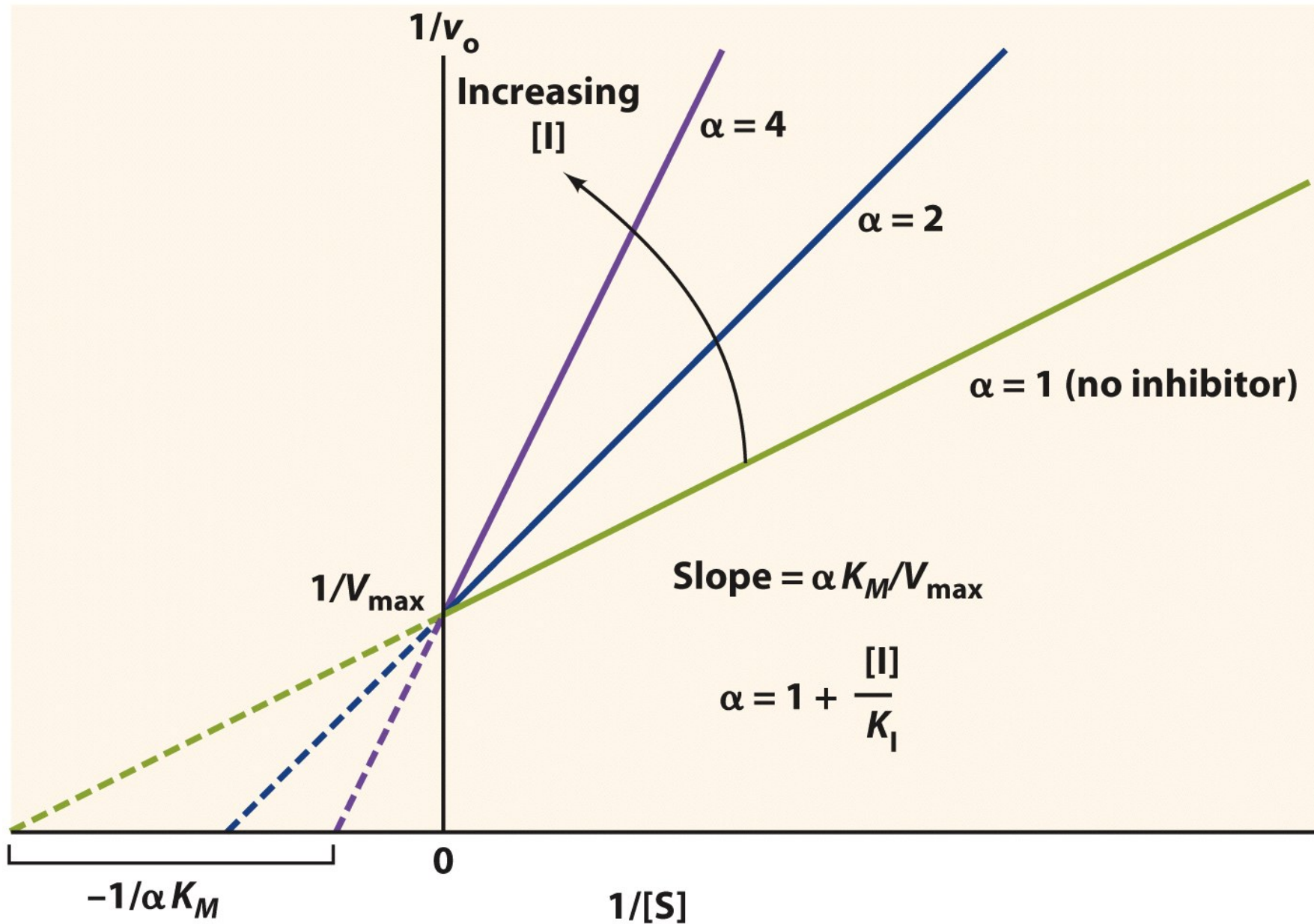
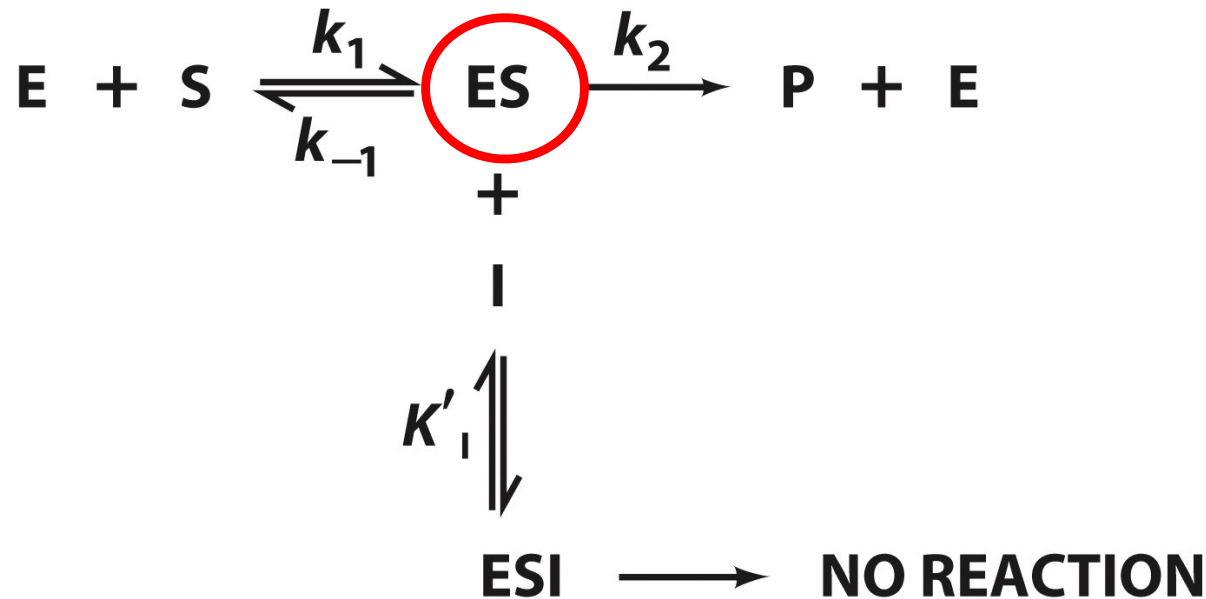


Figure 12-8

Uncompetitive Enzyme Inhibition



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ESI: Enzyme-Substrate-Inhibitor complex: a certain portion of ESI always exists, thus decreases V_{\max} - as if some enzymes are kidnapped in ES form!
 What would happen to K_M ?



$$K_M = (k_{-1} + k_2) / k_1$$

Uncompetitive Enzyme Inhibition

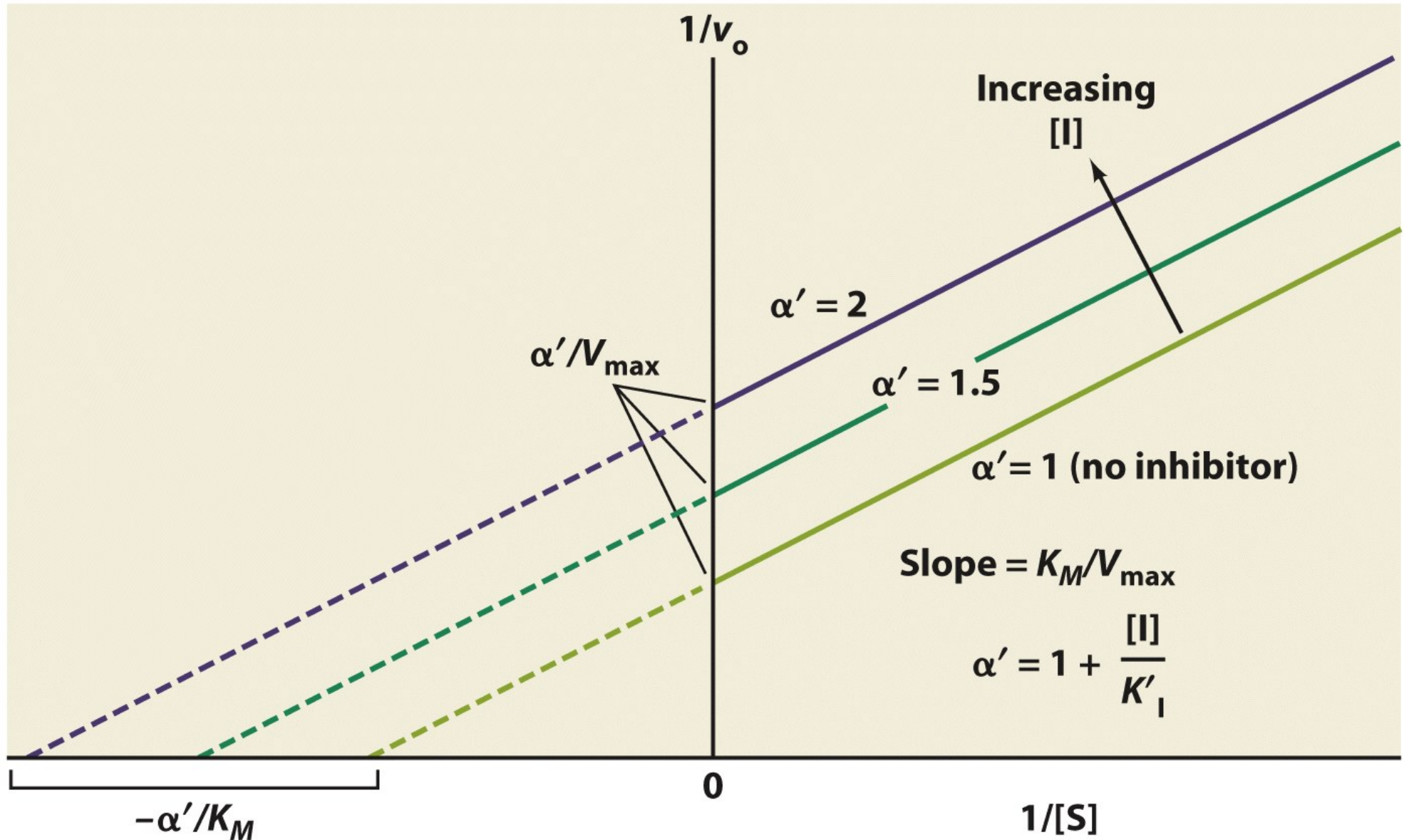
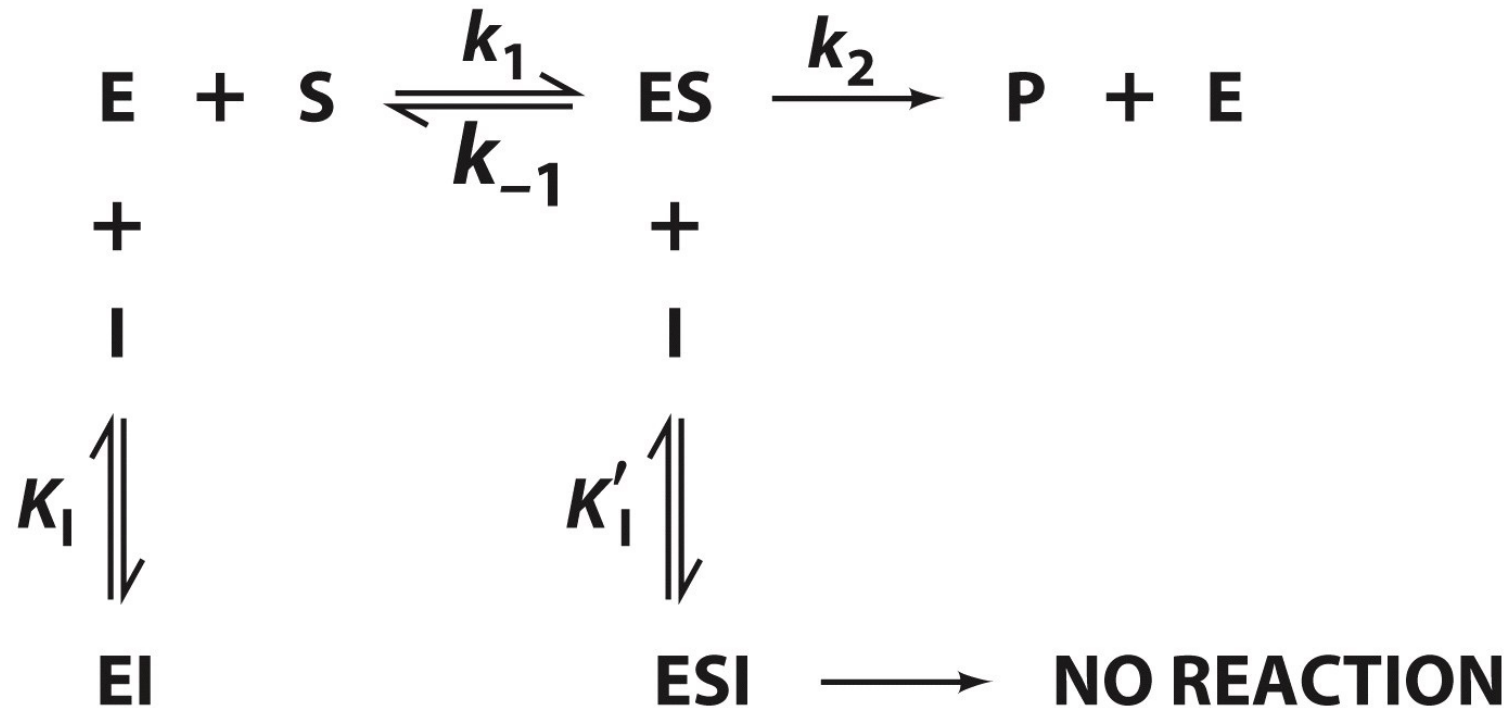
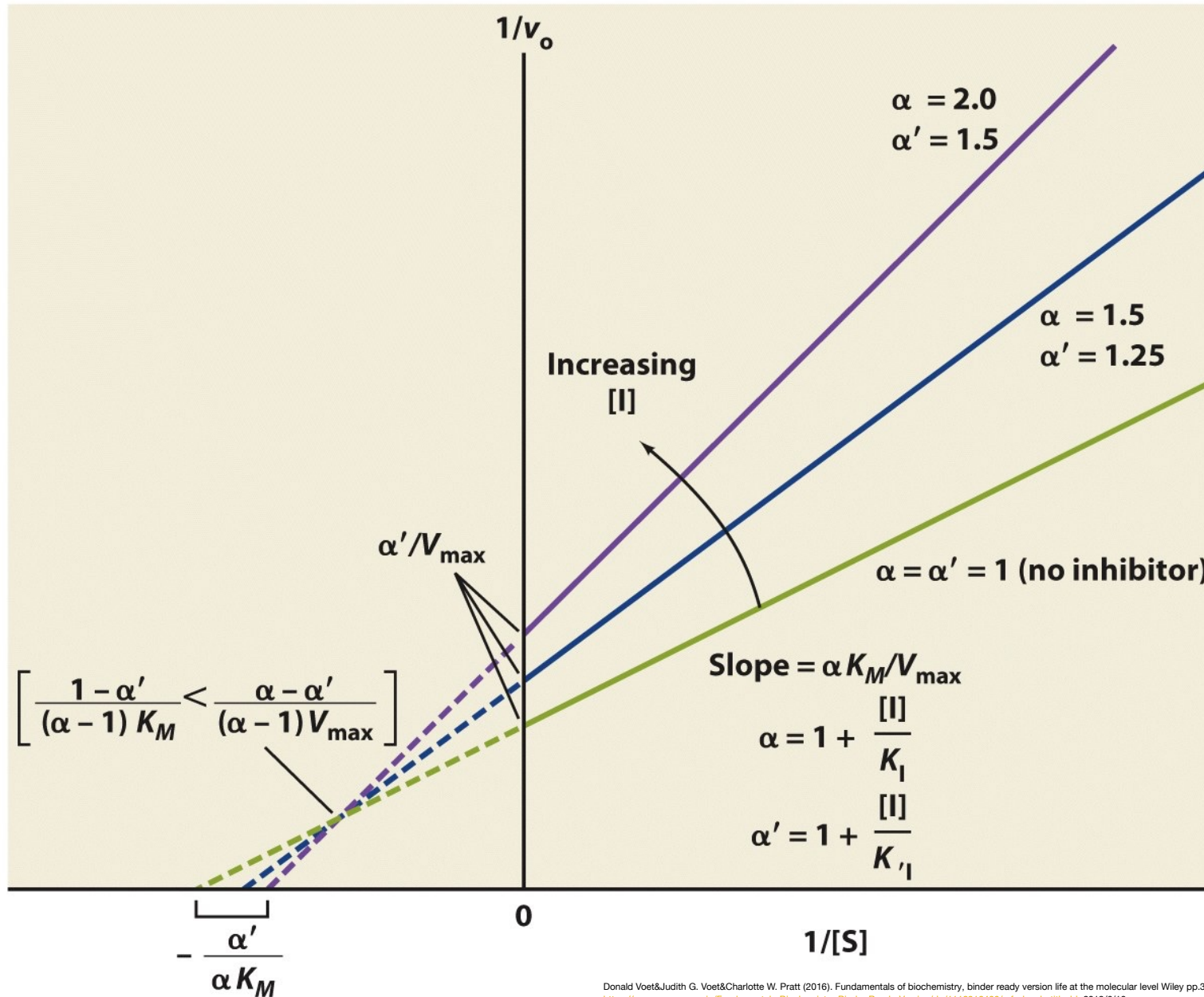


Figure 12-9

Mixed and Noncompetitive Enzyme Inhibition



Mixed and Noncompetitive Enzyme Inhibition



Enzyme Inhibitor Effects

TABLE 12-2 Effects of Inhibitors on Michaelis–Menten Reactions^a

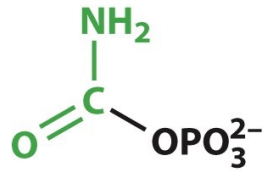
Type of Inhibition	Michaelis–Menten Equation	Lineweaver–Burk Equation	Effect of Inhibitor
None	$v_o = \frac{V_{\max}[S]}{K_M + [S]}$	$\frac{1}{v_o} = \frac{K_M}{V_{\max}} \frac{1}{[S]} + \frac{1}{V_{\max}}$	None
Competitive	$v_o = \frac{V_{\max}[S]}{\alpha K_M + [S]}$	$\frac{1}{v_o} = \frac{\alpha K_M}{V_{\max}} \frac{1}{[S]} + \frac{1}{V_{\max}}$	Increases K_M^{app}
Uncompetitive	$v_o = \frac{V_{\max}[S]}{K_M + \alpha'[S]} = \frac{(V_{\max}/\alpha')[S]}{K_M/\alpha' + [S]}$	$\frac{1}{v_o} = \frac{K_M}{V_{\max}} \frac{1}{[S]} + \frac{\alpha'}{V_{\max}}$	Decreases K_M^{app} and V_{\max}^{app}
Mixed (noncompetitive)	$v_o = \frac{V_{\max}[S]}{\alpha K_M + \alpha'[S]} = \frac{(V_{\max}/\alpha')[S]}{(\alpha/\alpha')K_M + [S]}$	$\frac{1}{v_o} = \frac{\alpha K_M}{V_{\max}} \frac{1}{[S]} + \frac{\alpha'}{V_{\max}}$	Decreases V_{\max}^{app} ; may increase or decrease K_M^{app}

$$^a\alpha = 1 + \frac{[I]}{K_I} \quad \text{and} \quad \alpha' = 1 + \frac{[I]}{K'_I}$$

Control of Enzyme Activity

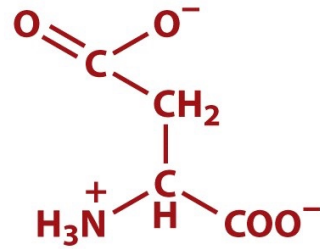
- Allosteric effectors bind to multisubunit enzymes such as aspartate transcarbamoylase, thereby inducing cooperative conformational changes that alter the enzyme's catalytic activity. (They don't follow MM)
- Phosphorylation and dephosphorylation of an enzyme such as glycogen phosphorylase can control its activity by shifting the equilibrium between more active and less active conformations.

Aspartate Transcarbamoylase Reaction



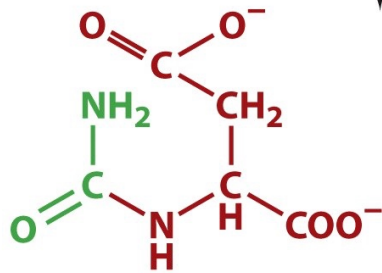
Carbamoyl phosphate

+



Aspartate

aspartate
transcarbamoylase

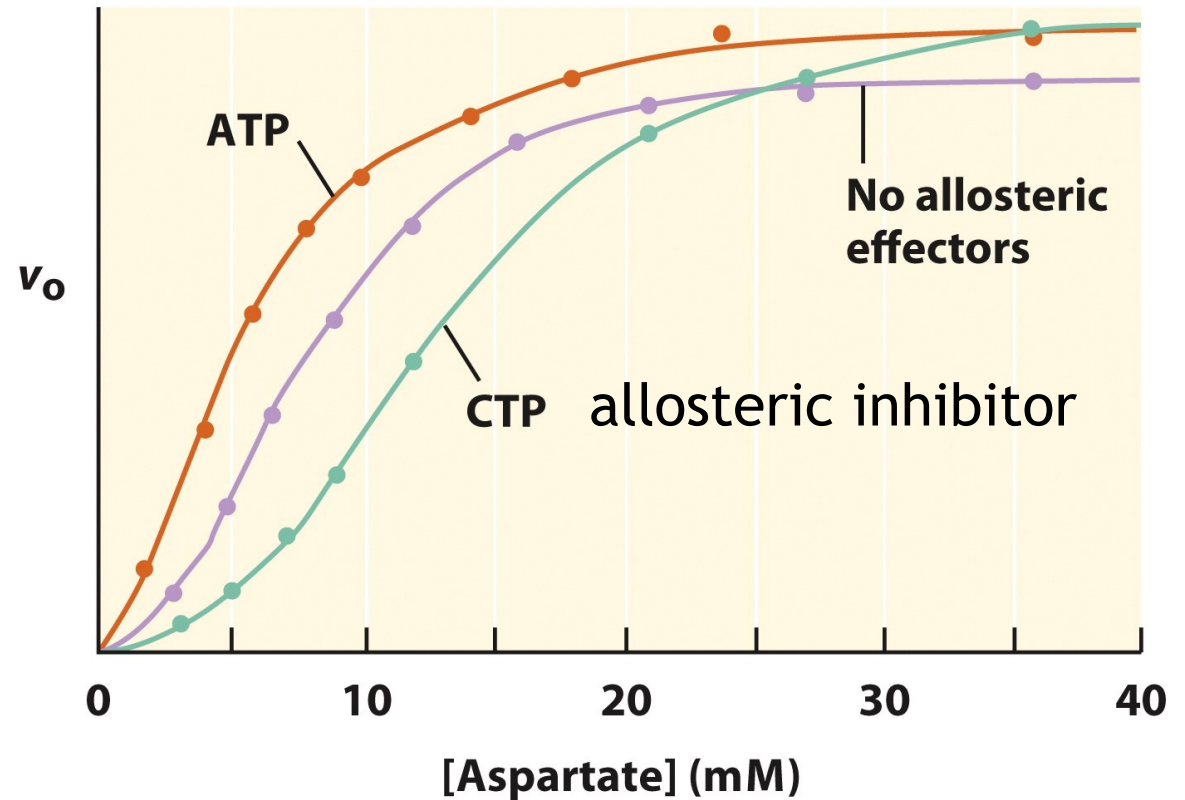


N-Carbamoylaspartate

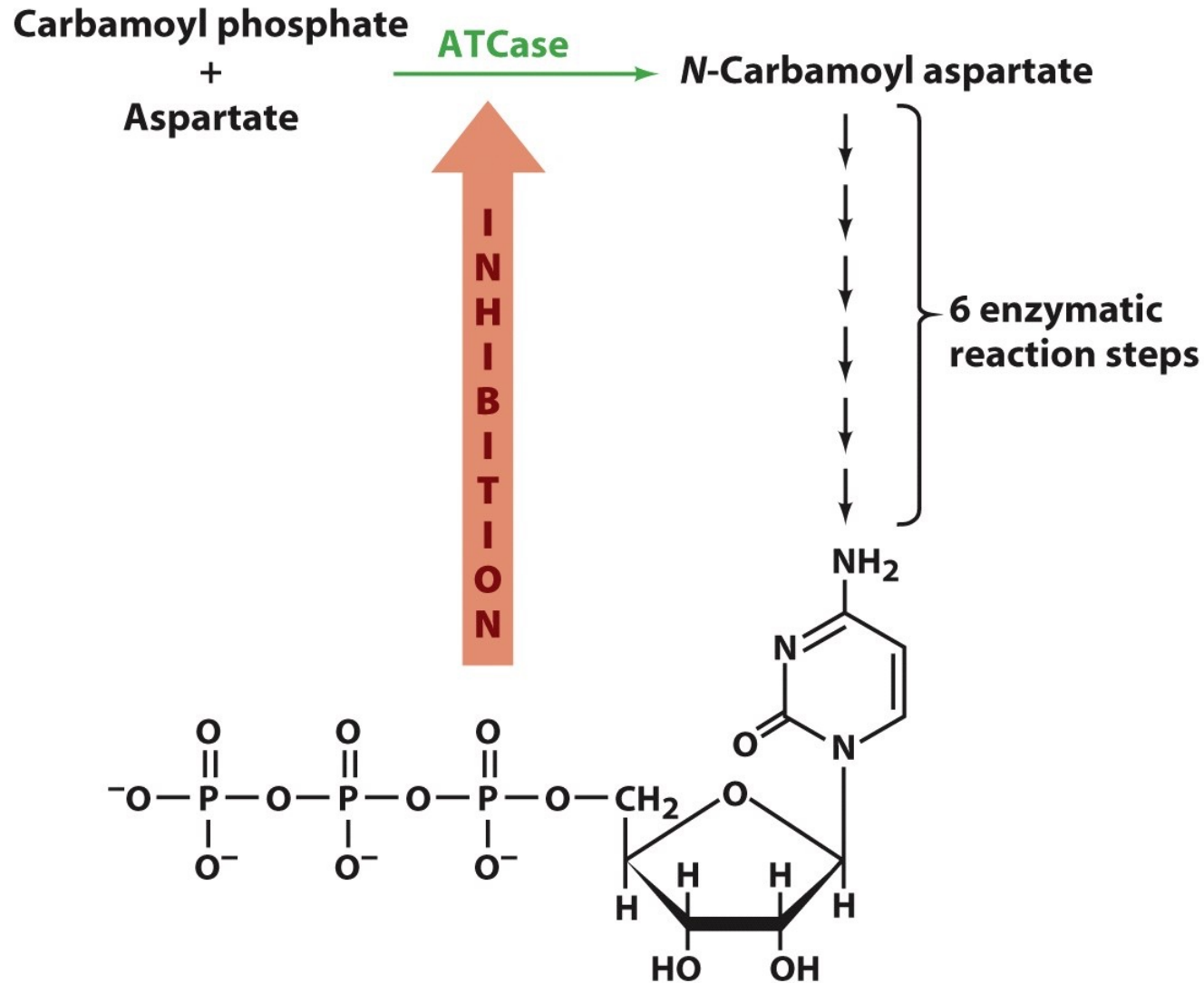
+ H_2PO_4^-

allosteric control:

- distinct regulatory site
- multisubunit
- cooperativity
- feedback inhibition

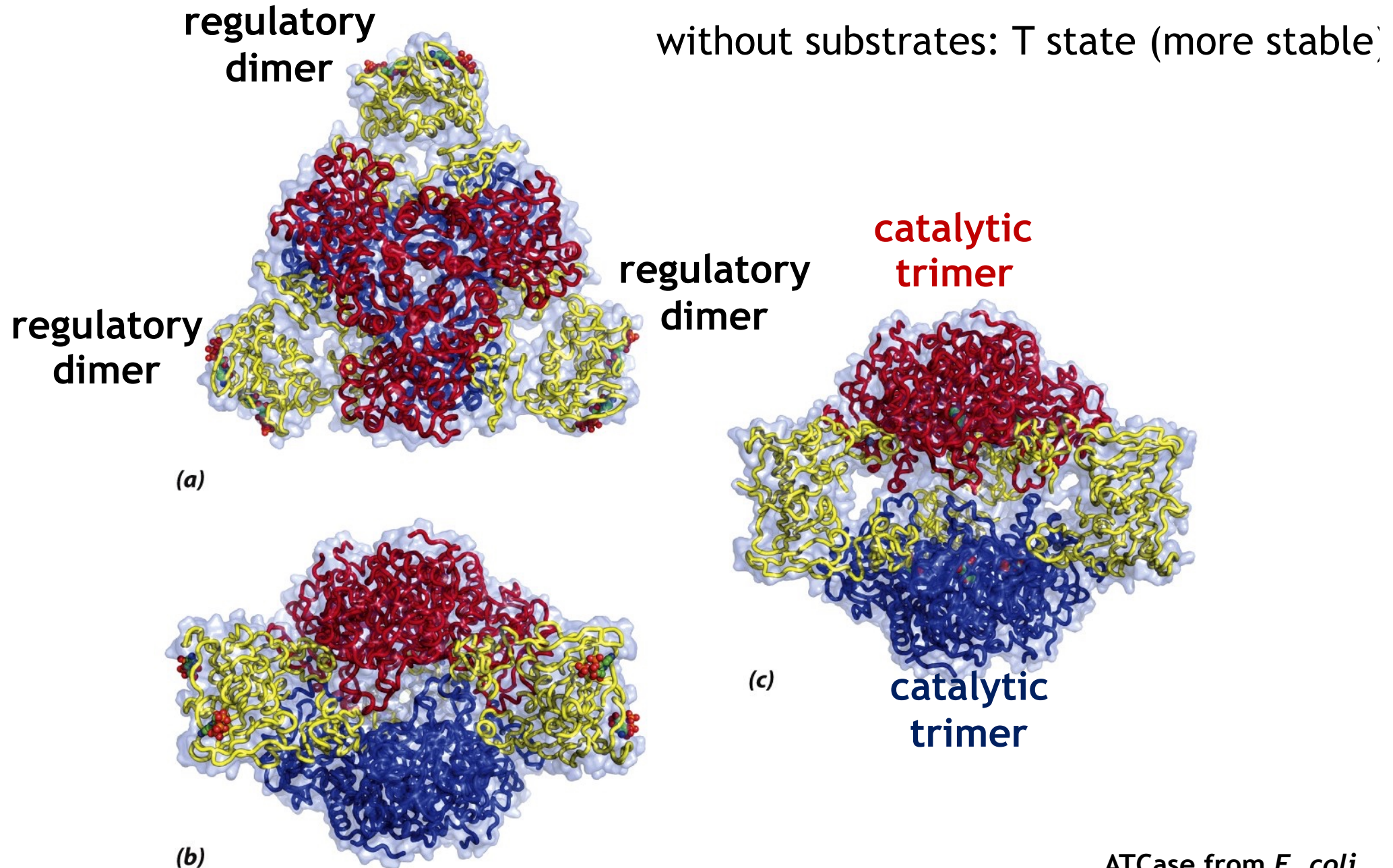


Pyrimidine Biosynthesis: ATCase Feedback Inhibition

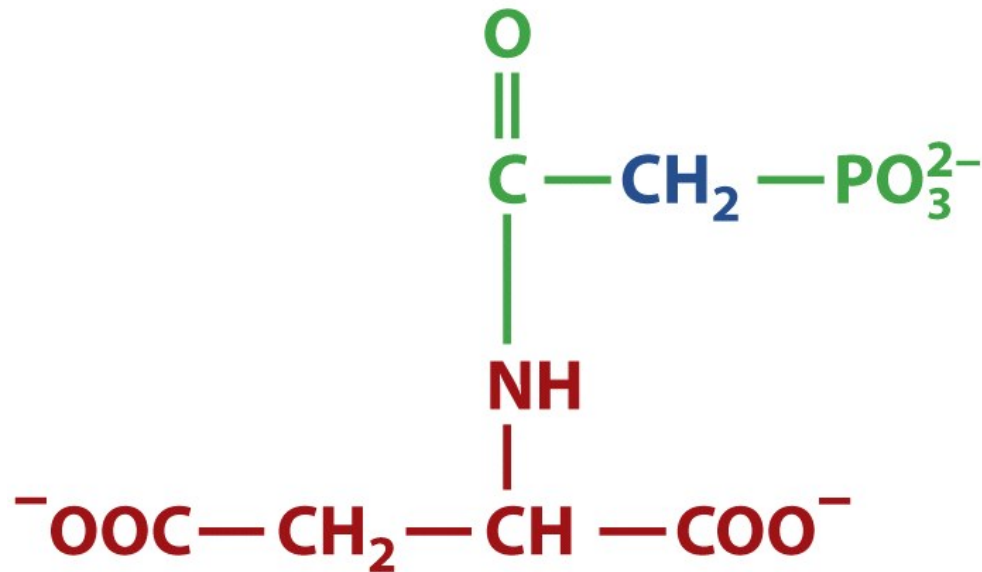


Cytidine triphosphate (CTP)

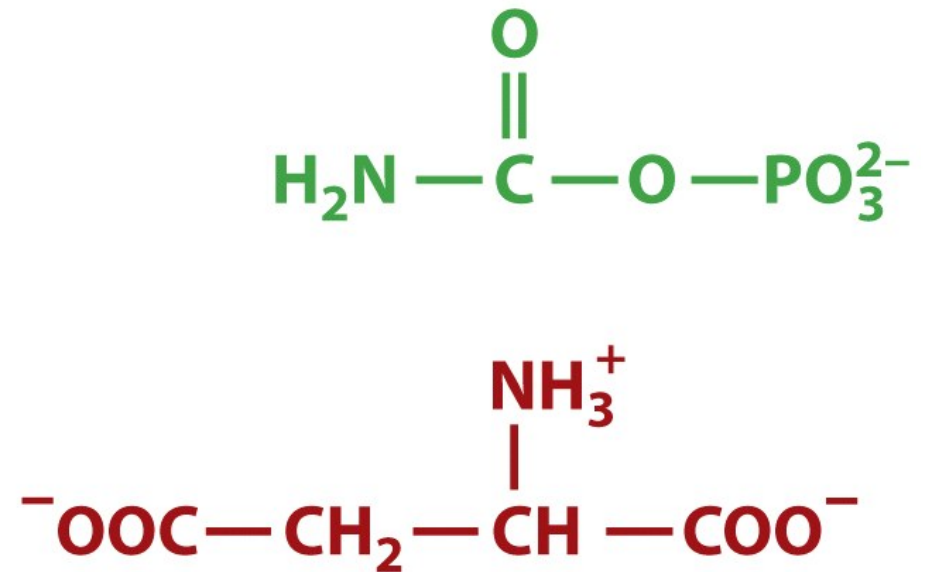
ATCase: T-State vs. R-State



Unreactive Bisubstrate Analog



***N*-(Phosphonacetyl)-
L-aspartate (PALA)**



**Carbamoyl phosphate
+
Aspartate**

ATCase: Conformational Changes

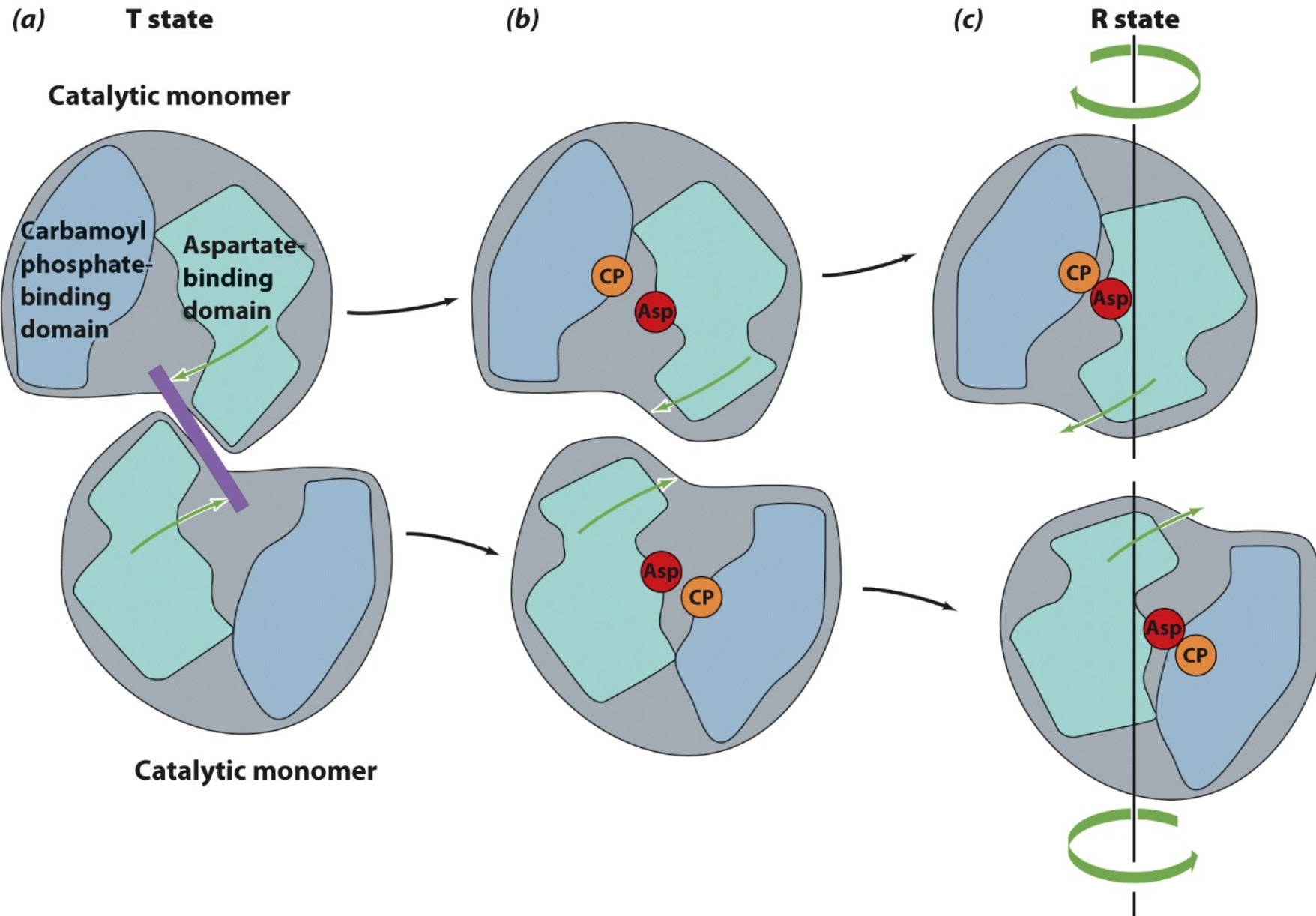
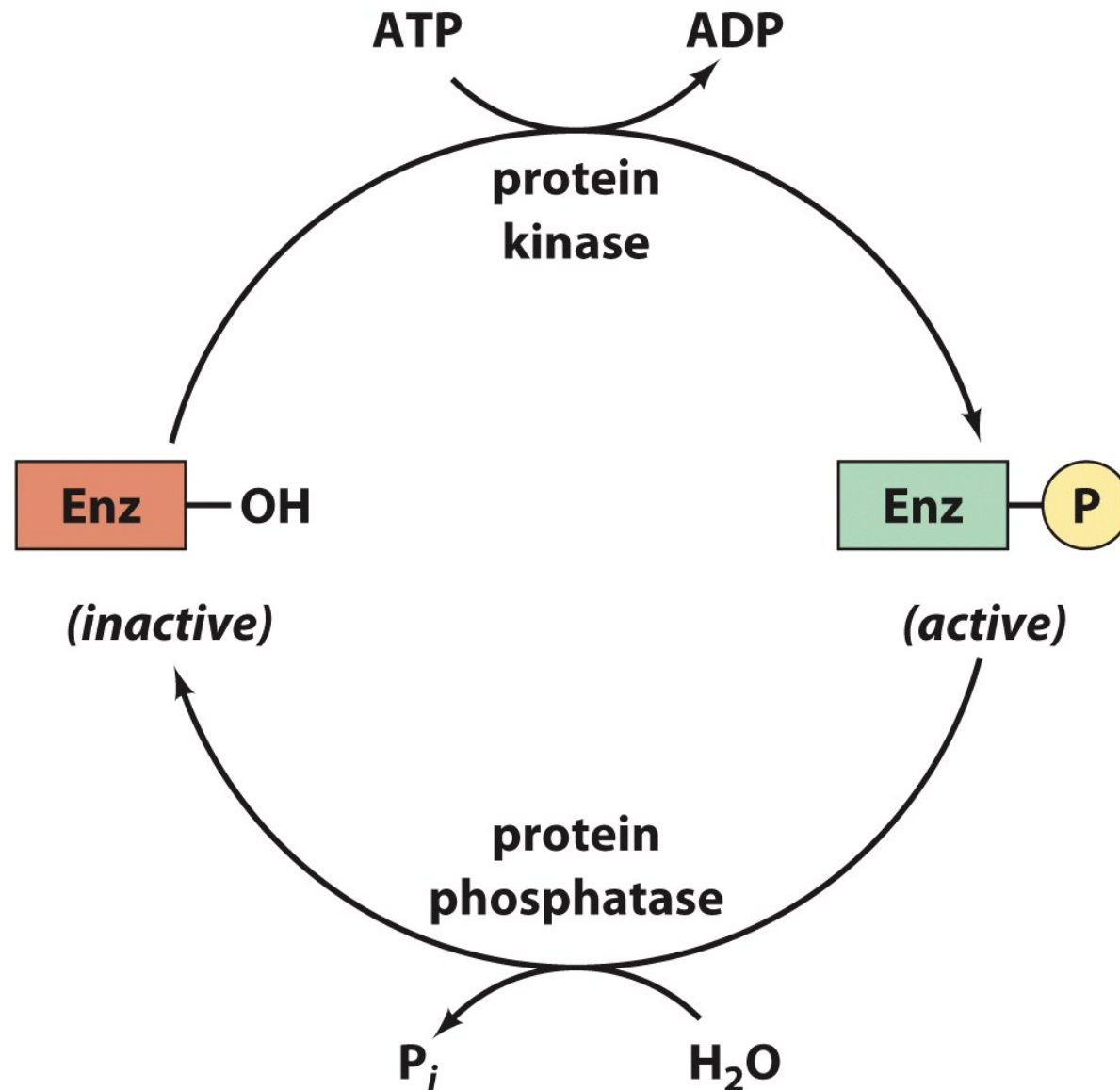


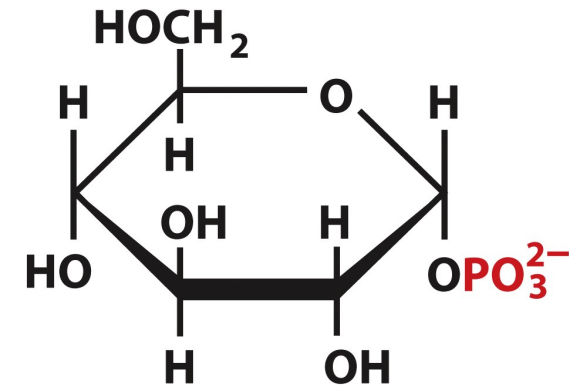
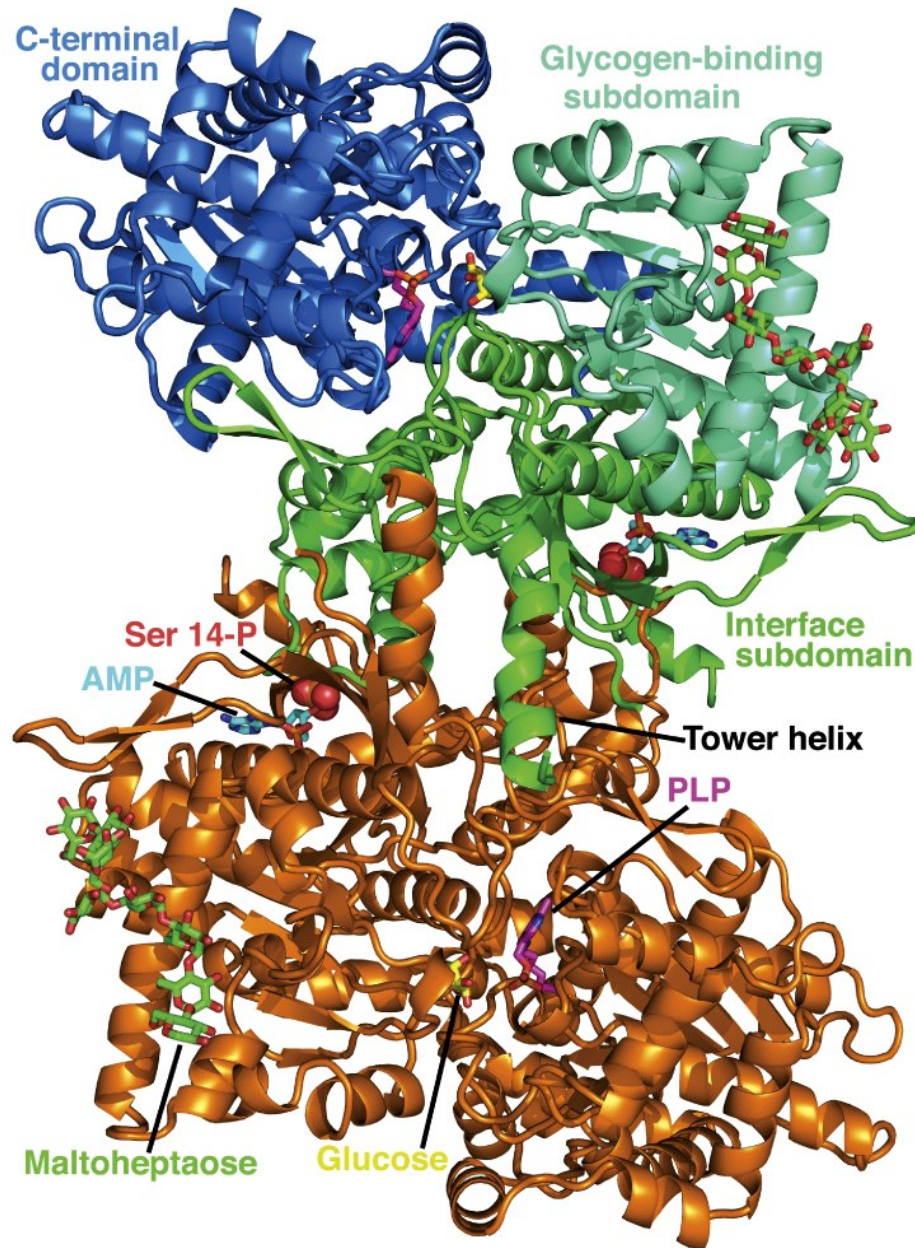
Figure 12-14

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Control by Covalent Modification: Phosphorylation



Rabbit Muscle Glycogen Phosphorylase



**Glucose-1-phosphate
(G1P)**

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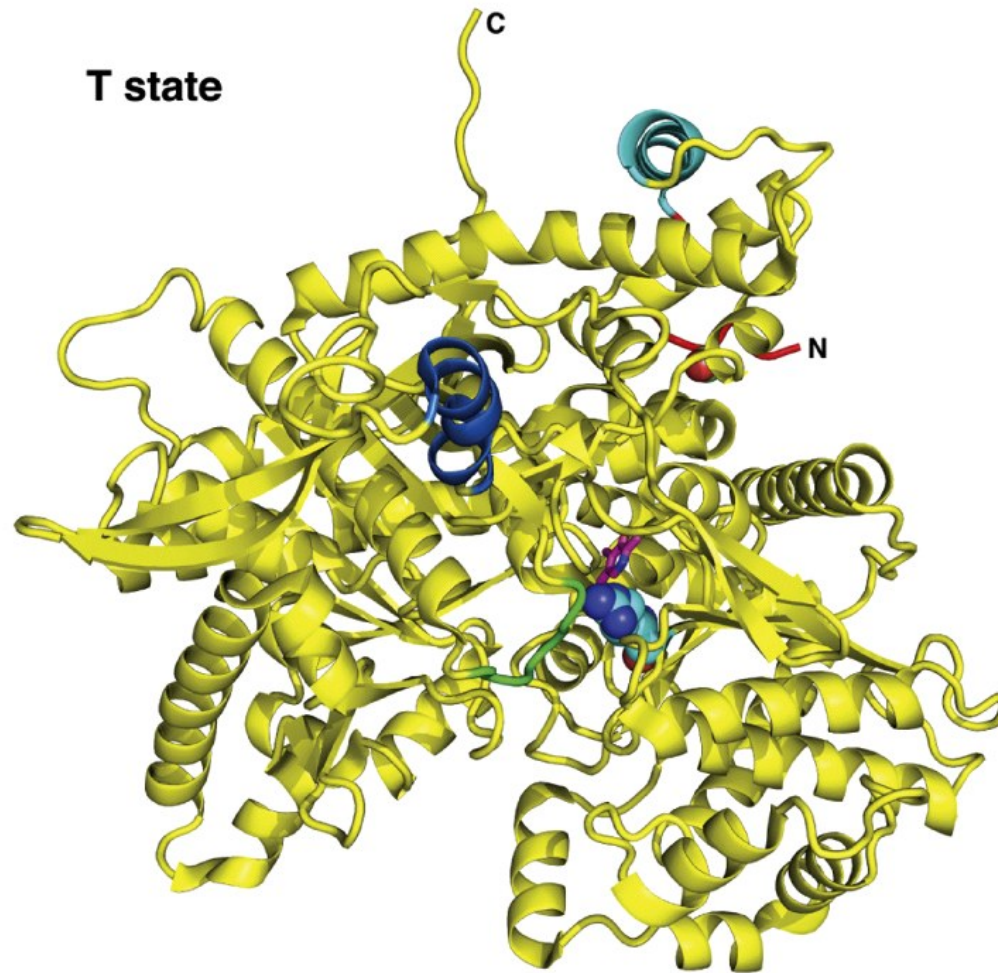
Figure 12-15

X-Ray structure coordinates courtesy of Stephen Sprang, University of Texas Southwest Medical Center

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Conformational Changes: Glycogen Phosphorylase

(a)



(b)

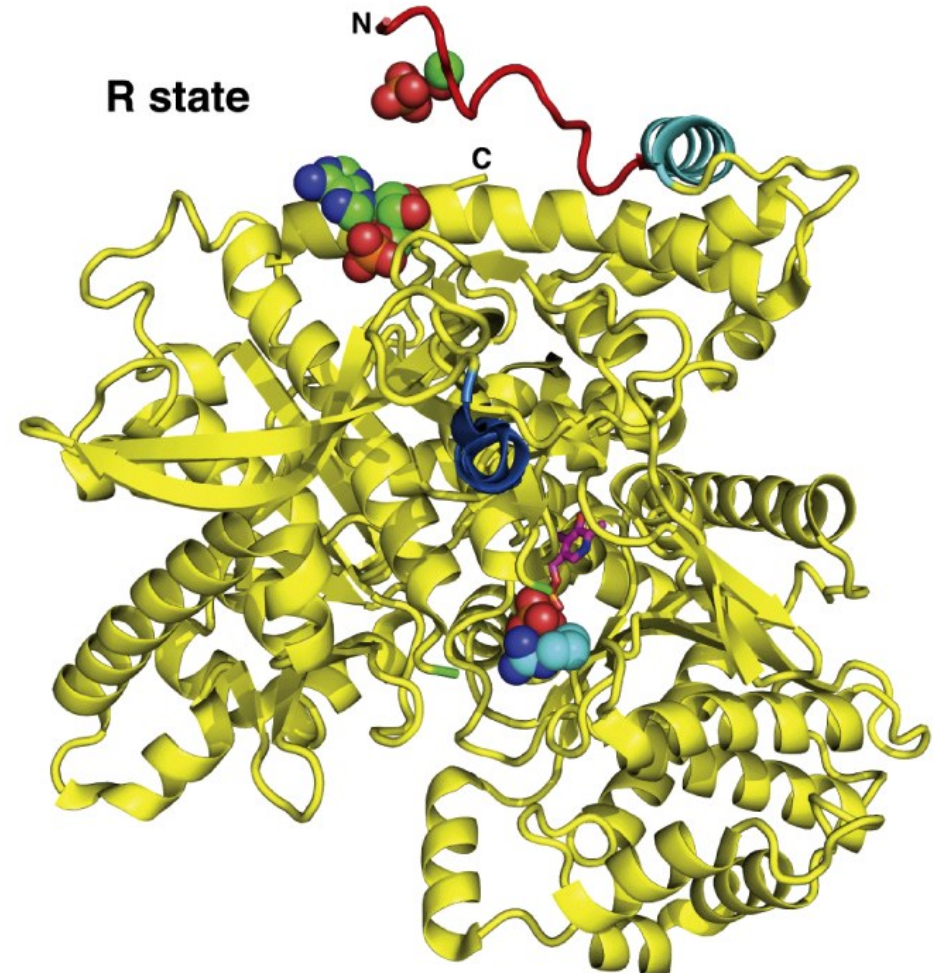


Figure 12-16

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Glycogen phosphorylase
PDBids [8GPB](#) and [7GPB](#)

Glycogen Phosphorylase: Control by Phosphorylation

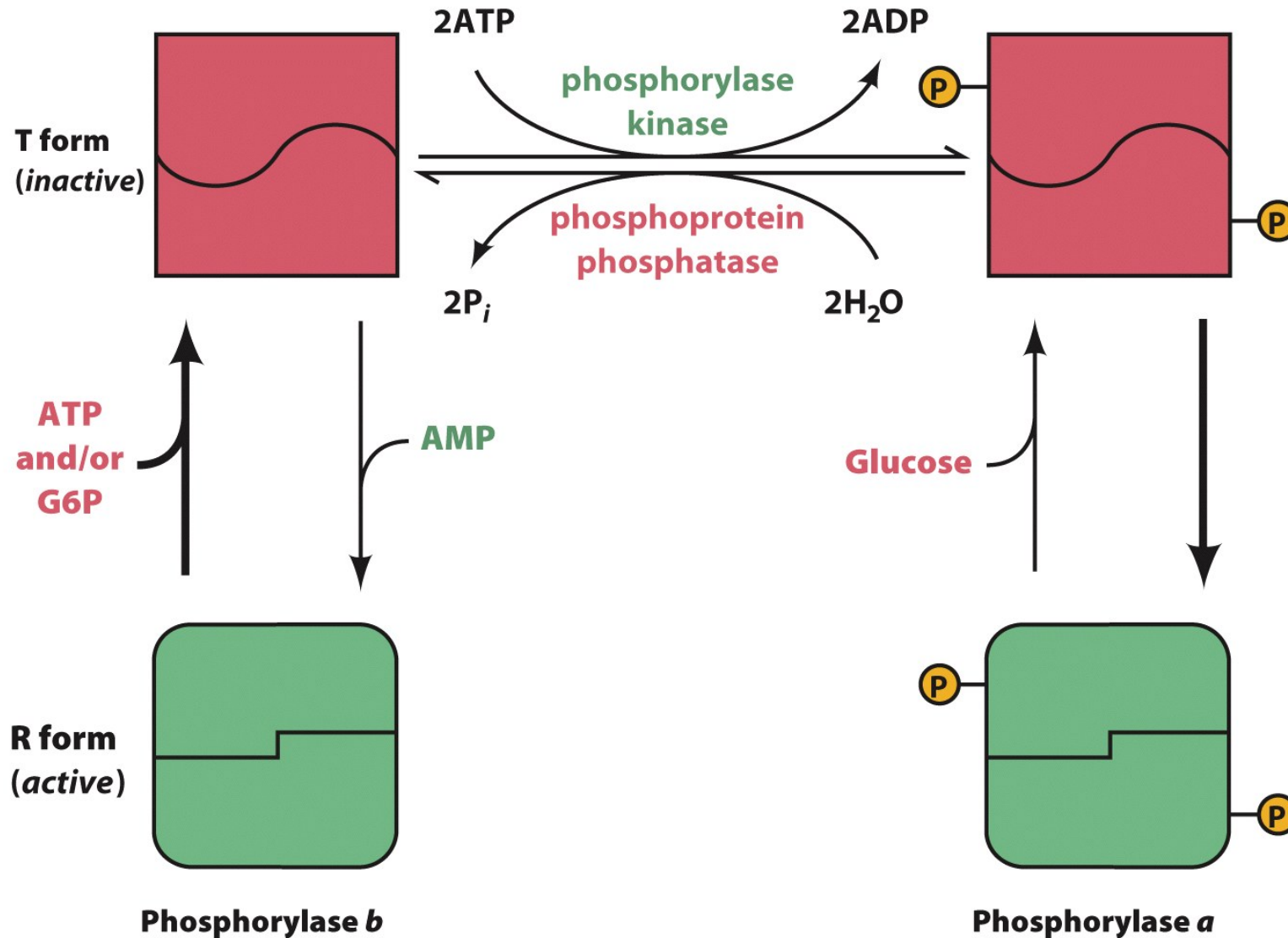


Figure 12-17

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