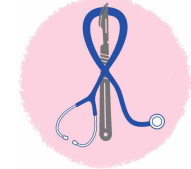


Enzymes & Coenzymes (1)



MED441
KING SAUD UNIVERSITY

Revised & Reviewed
by
Abdulaziz & Bahammam
Faye Weel Sendi



7

V1

Foundation
Block - KSU

Color Index:

- Main text
- Important
- Notes
- Boys slides'
- Girls slides'
- Extra

Editing File



Objectives

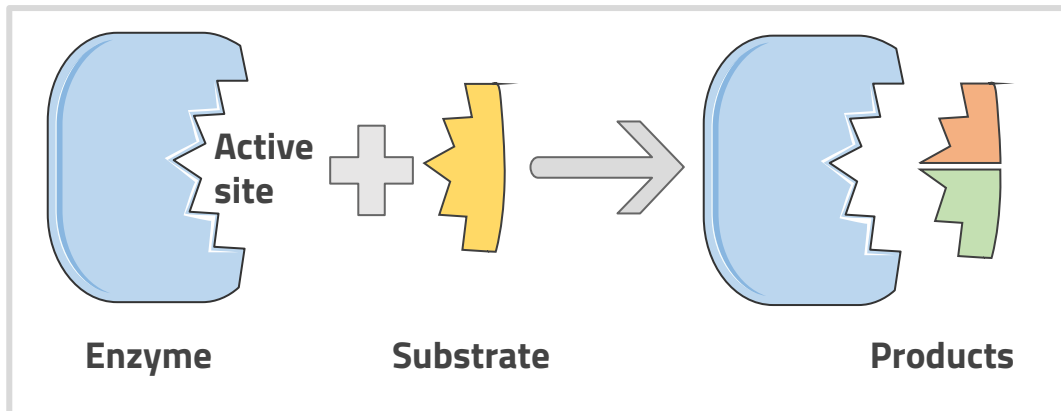
- Understand how enzymes are able to speed up the rate of biochemical reactions in the body.
- Identify classes of enzymes based on the type of reactions they catalyze.
- Comprehend basic terms of: coenzymes, isoenzymes, enzyme activity & specificity along with factors affecting their activity.
- Understand the enzyme kinetics.

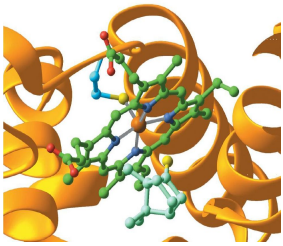
Enzymes



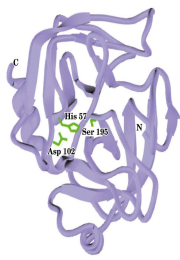
“**Biological catalysts** (محفزات) that speed up the rate of a reaction **without being consumed**/changed in the reaction”

Structure	all enzymes are protein in nature, but NOT all proteins are enzymes (others are structural, transporters, etc.) <u>Exception</u> : ribozymes are RNA (ribonucleic acid) molecules with enzymatic activity	
Function	bind to specific substrate “substance upon which enzymes act” (like a reactant) to convert them to product(s)	
Properties	1) Active site	▪ all enzymes have ≥ 1 active sites : “regions where substrate binds” (catalysis occurs once they bind)
	2) Specificity	enzymes are highly specific <ul style="list-style-type: none"> ▪ they interact with only 1 or few substrates if they have very similar structures ▪ they catalyze only 1 type of reaction even if it’s the same substrate (437)
	3) Regulation	enzymes can be activated or inhibited so that the rate of product formation responds to the need of the cell <ul style="list-style-type: none"> ▪ some enzymes also have regulatory sites that control the active site in different ways (next lecture’s topic)

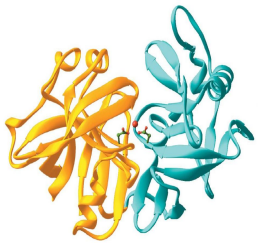




an **enzyme**
with its active site



Trypsin (a digestive
enzyme in small intestine)



Pepsin (a digestive
enzyme in stomach)

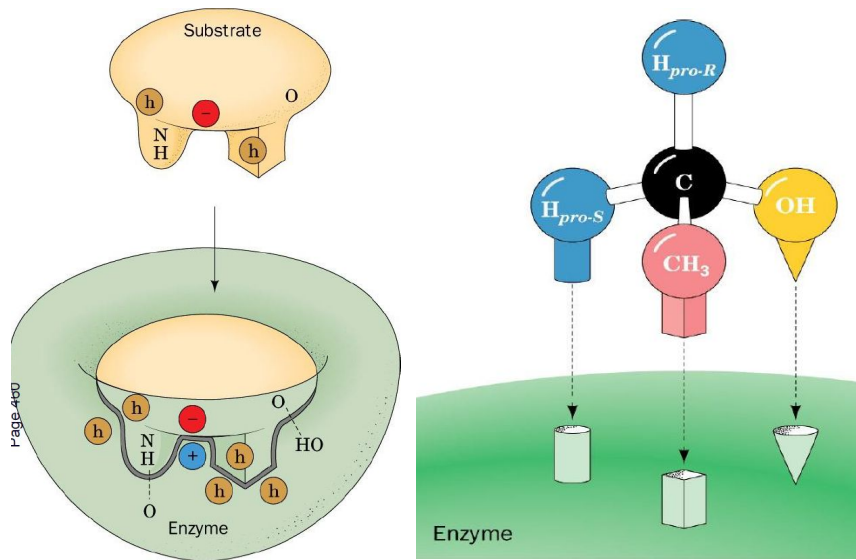


Models of enzyme-substrate binding

Lock & key binding

Active site fits the **exact dimensions** of substrate
[**exactly complementary**]

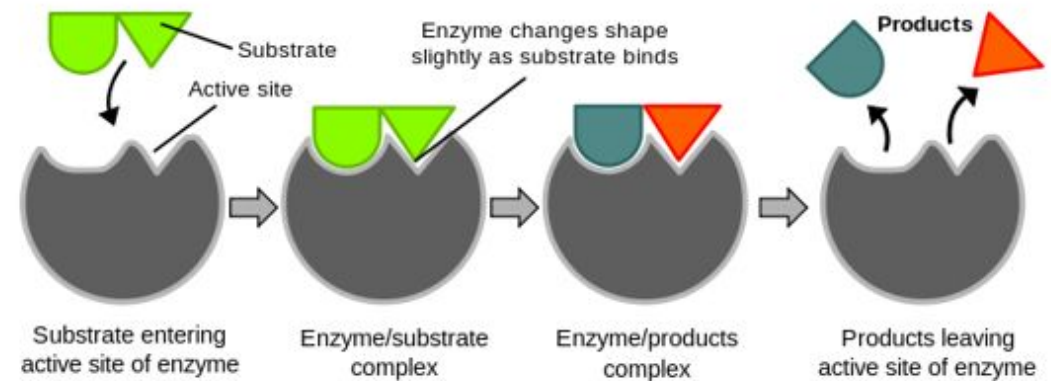
زي القفل ما يفتحه إلا مفتاح واحد نفس شكله بالضبط



Induced-fit binding

After the binding of substrate, the **enzyme changes its shape** (conformation) to fit more perfectly with substrate
[**not fully complementary**, after the reaction is done, it goes back to its original shape to be able to function again]

زي القفاز ياخذ شكل اليد بعد ما ينلبس





Classification of enzymes



[helpful video](#)

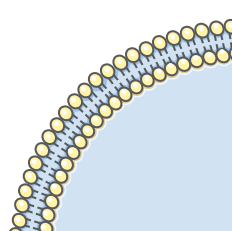
Classified into 6 types according to the type of reaction catalyzed:

(436)	Class	Type of reaction (439)
Omar	1. Oxidoreductases	Oxidation-reduction reaction
Tried	2. Transferases	Transfer of functional groups
Hard	3. Hydrolases	Hydrolysis (breaking bonds by adding water)
Learning	4. Lyases	Group elimination to form double/triple bonds
International	5. Isomerases	Isomerization (form isomers)
Languages	6. Ligases	Bond formation coupled with ATP hydrolysis



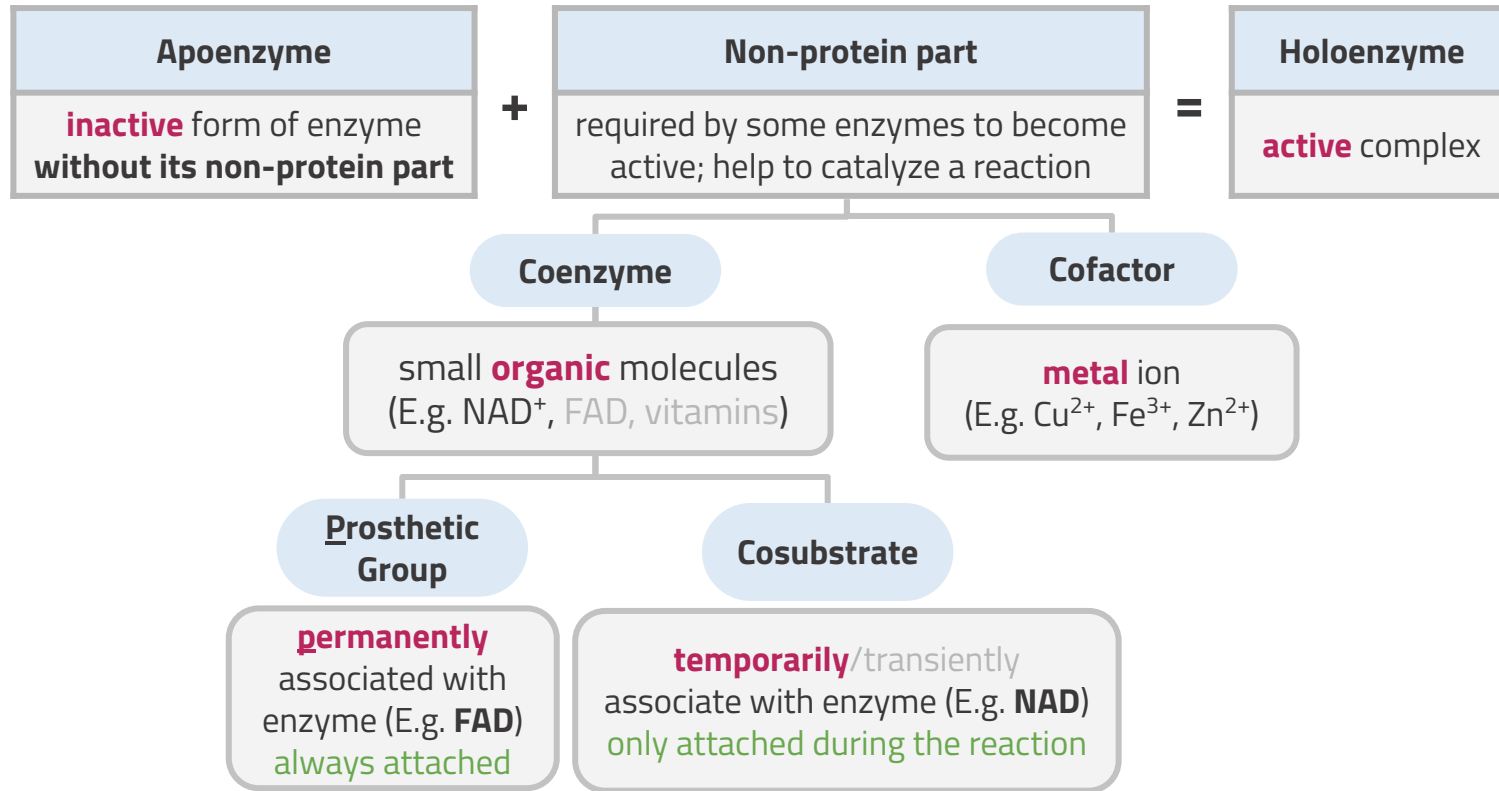
Nomenclature (naming)

	Common name	Systematic name
Rule	Suffix "-ase"	based on the rules given by IUBMB (International Union of Biochemistry & Molecular Biology): EC Class . Subclass . Subsubclass . Enzyme number (EC = Enzyme Commission)
Example	(E.g. amylase) <u>Exceptions:</u> pepsin, trypsin	<p><u>Example: EC 3.4.17.1</u> (carboxypeptidase A)</p> <p>Hydrolase, breaks a bond ← (439) → Which enzyme exactly? hypothetically #1 Which bond? peptide ← → Which peptide bond? alanine & serine for ex.</p> <p>*This is just an example, don't memorize it</p>

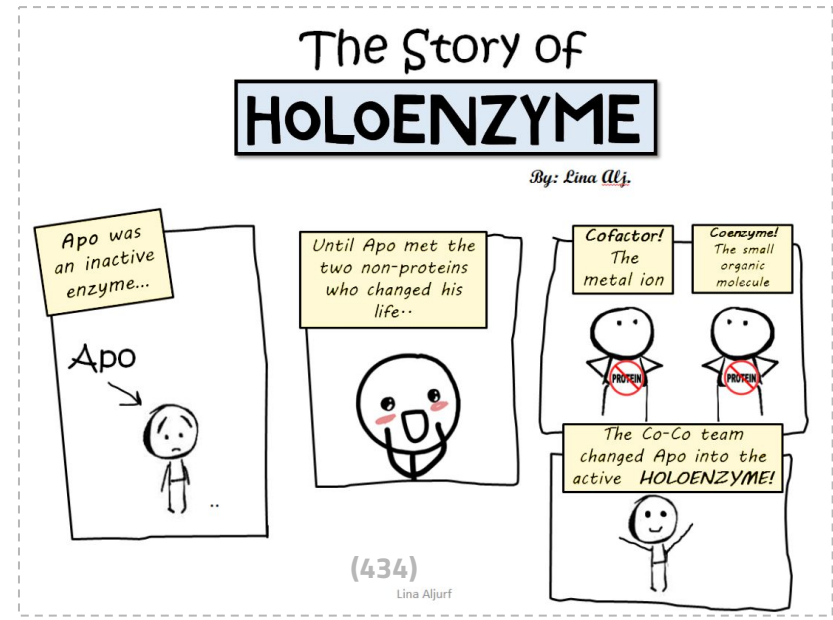


Basic terms

Some enzymes require **non-protein** groups to catalyze a reaction (become active):



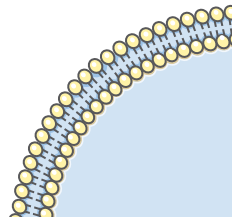
- The majority of enzyme require these molecules
- They are present in the active site as an additional molecule



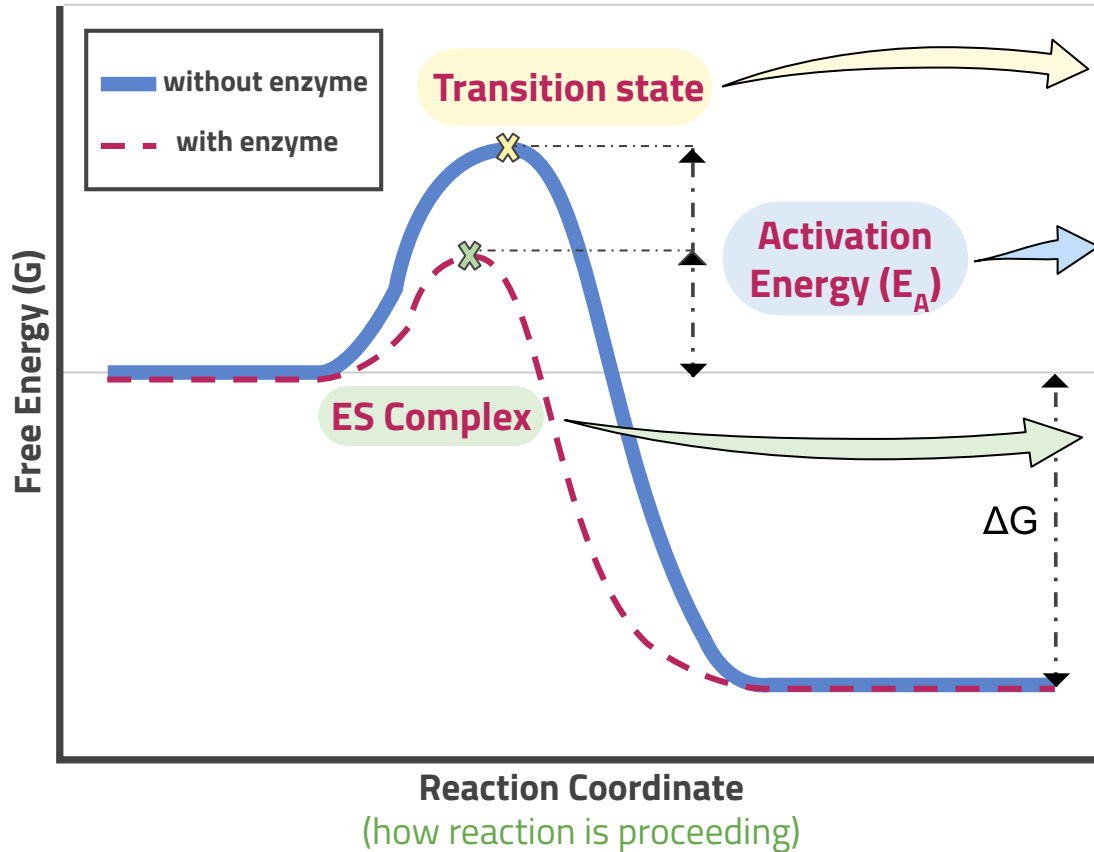
- Ribozymes** → **RNAs** (ribonucleic acid) with enzymatic activity (they're the only non-protein enzymes)
- Isoenzymes** → enzymes that catalyze the **same chemical reaction** but have slightly **different structures**
They're usually found in different locations [Ex. blood, tissue]
- Zymogens** → **inactive** enzyme precursors that require a biochemical change to become active
(E.g. cleavage of a peptide blocking the active site, phosphorylation). They're activated when needed
This provides a way of regulation. For example, [pepsinogen → pepsin] only when we're eating

(434): notice that zymogens & apoenzymes need different things to become active:
Apoenzymes require a coenzyme/cofactor
Zymogens require a biochemical change

(Aka proenzymes)



How enzymes work



It is a **high-energy** intermediate that reactants pass through in every chemical reaction; it has **greater energy** than that of reactants or products alone

It is the **difference in energy** between **reactants** & **transition state**

- If activation energy is available, the reaction can proceed forming products
- Uncatalyzed reactions are often slow due to high E_A

In order for enzymes to speed up reactions, they must:

- 1) **decrease the activation energy** required for a reaction, providing an **alternative transition state** of **lower energy** called **enzyme-substrate complex** and thus speed up the reaction.
- 2) **NOT alter the change in the free energy (ΔG)** "energy that can do work"
Why? because we don't want them to change the equilibrium of the reaction; rather, we need them to accelerate the rate by which equilibrium is reached, speeding up both forward & reverse rates

Summary of how enzymes work:

- 1) Activation energy is reduced
- 2) Free energy remains the same

(436)

Enzyme induction = increasing activity
Enzyme inhibition = decreasing activity

ΔG of products must be less than ΔG of reactants

Enzyme activity or velocity

Velocity: the rate of a reaction catalyzed by an enzyme

- **Enzyme activity** is expressed as μ **moles** of product formed/**min/mg** enzyme

Enzyme velocity can be measured by either:

- how much product is increased / formed (more common)
- how much substrate conc is decreased

Factors that affect enzyme activity

1) Temperature

Every enzyme has an **optimal temp** for catalyzing a reaction → in **humans** most enzymes have an optimal temp of **37°C**

- The **rate** of an enzyme reaction **initially increases** with rise in temperature (increase in velocity → until it reaches **peak** velocity (very active))
- But at **high** temp [E.g. above 40°C] enzymes are **denatured** → become **inactive**

2) pH

Every enzyme has an **optimal pH** for catalyzing a reaction, which is the peak of a bell-shaped curve

↳ most have highest activity between **pH 6-8** [**Exception:** pepsin has highest activity at **pH 2** in the stomach]

- pH affects **catalysis** through either the **substrate** or **ionizable groups** in the active site of enzyme

Extra: Ionizable groups are those that are able to gain/lose H^+ , such as $COOH$ & NH_2 in amino acids. Catalysis requires both substrate & active site to have a specific chemical group in either ionized or unionized states, which is directly affected by pH

3) [S] & [E] concentration

*Square brackets [] depict concentration

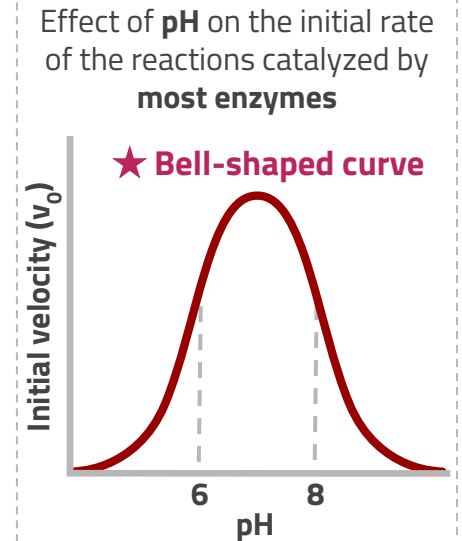
At low [S]

- the reaction velocity/rate (v) is **proportional to [S]** (increases initially with increasing [S])

at high [S]

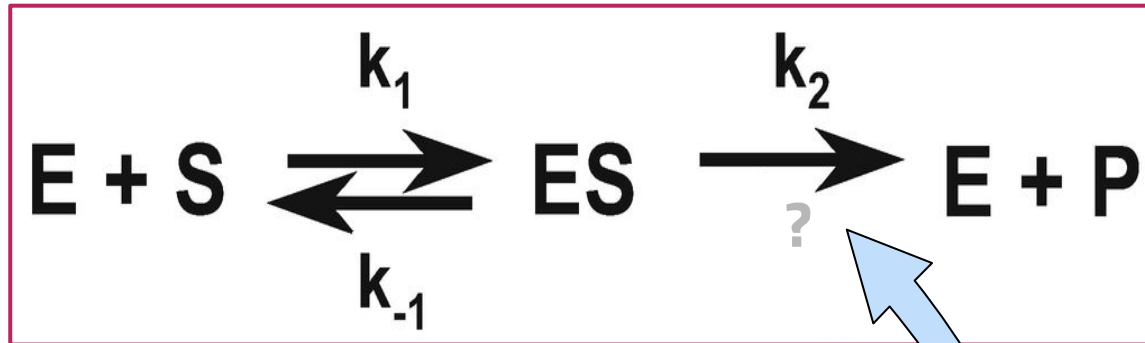
- further addition of **substrate** has **no effect** on enzyme velocity because enzyme is saturated & all active sites are engaged
- the rate of an enzyme reaction is directly **proportional to [E]** if [S] is higher than [E]

For example, if there are 50 enzyme molecules & 70 substrates, only 50 substrates will be catalyzed as we have a limited number of active sites



Enzyme Kinetics

Basic reaction model



In this reaction model, ES complex has 2 fates:
1) Continue to form product [right]
2) Go backwards [left]

S = Substrate
E = Enzyme
ES = Enzyme-Substrate complex
P = Product
 k_1 , k_{-1} , k_2 = Rate constants

Extra:

Why is k_{-2} missing? Although step 2 is reversible, we do not consider its reverse reaction because our study of kinetics is during the early phase, where there is a very low concentration of product, so k_{-2} is negligible



Initial rate of enzyme reaction



[helpful video](#)

1) Pre-steady state kinetics

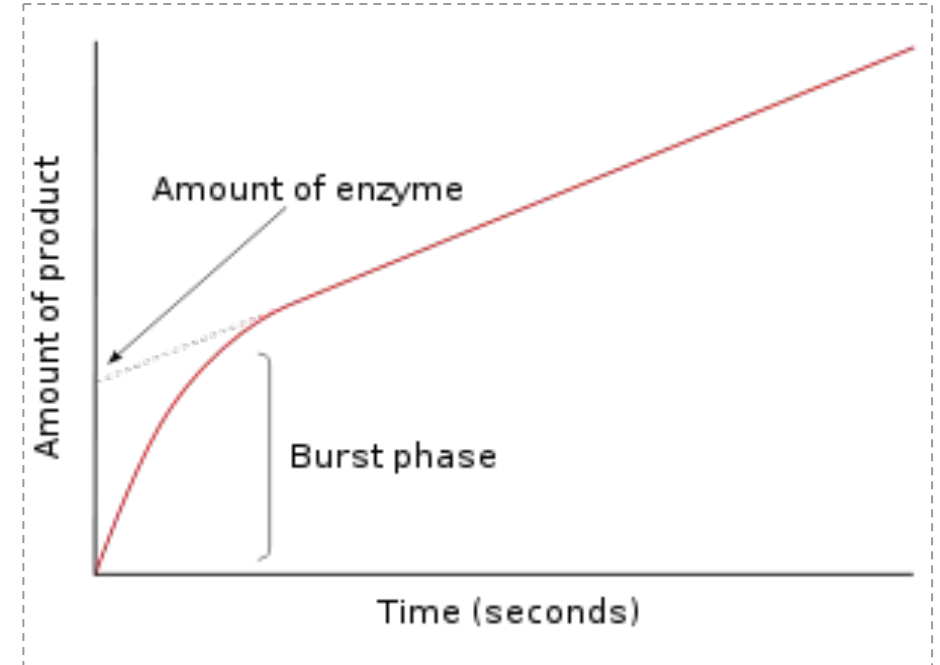
When an enzyme is mixed with high $[S]$, there is an **initial** short period of time (a few 100 microseconds) during which **intermediates** leading to the formation of product (**ES**) **gradually build up** (no product has formed yet)

أول ما نلاحظ الإنزيم مع substrate يبدأون يرتبطون ببعض ويكونون ال ES، والتي وهو لسا ما يعتبر ناتج تفاعل (product)

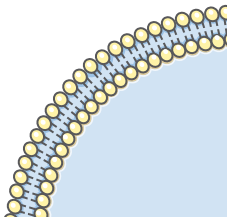
2) Steady state kinetics

After initial state, the **reaction rate** and **concentration** of intermediates change **slowly** with time, and **the intermediate is said to be in steady state** because its **rate of synthesis is equal to its rate of degradation**
rate of formation of ES = rate of breakdown of ES to $(E+S)$ & $(E+P)$

بعدها يثبت تركيز ال ES complex لأنه يبني ويتحطم بنفس المعدل (نلاحظ ثبات الميل)، وهنا يتكون ال product



Summary	Pre-steady state	<ul style="list-style-type: none"> ES builds up No product yet
	Steady state	<ul style="list-style-type: none"> ES is constant Product forms



Michaelis-Menten

- The model of enzyme kinetics was first proposed by **Michaelis & Menten** in 1913 and **later modified** by Briggs & Haldane
- The **Michaelis-Menten equation** describes the relationship of **initial rate** of an enzyme reaction to the **[S]**
- It measures the **initial velocity (v_o)** of an enzyme reaction

❖ Equation

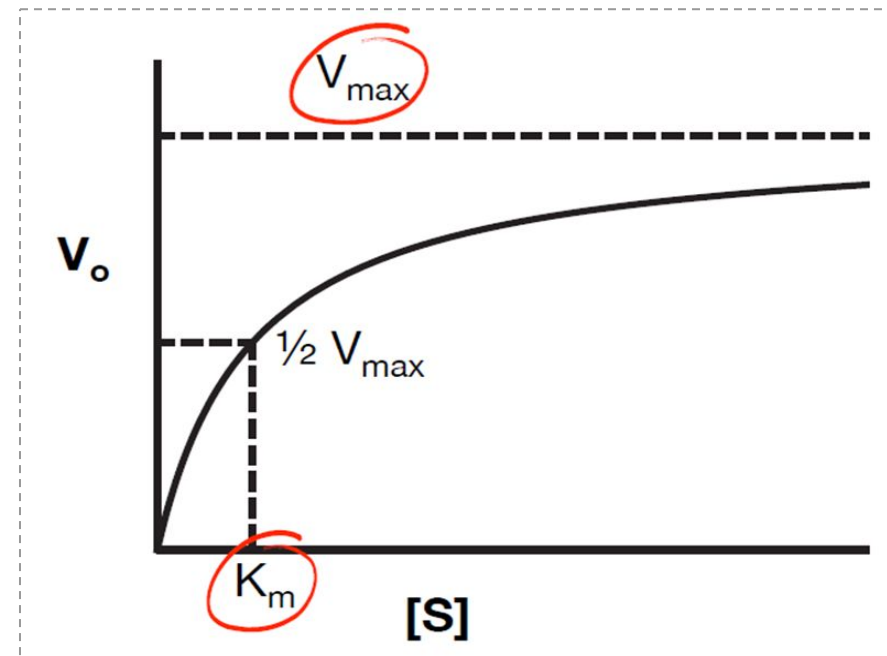
maximum velocity

$$v_o = \frac{V_{\max} \times [S]}{K_m + [S]}$$

substrate concentration
(at which v_o is calculated)

Michaelis constant
(explained in next slide)

❖ Graph



K_m (Michaelis Constant)

Quantitatively (numerically):

- It is the substrate concentration at which the **initial rate is one-half of the maximum rate ($\frac{1}{2} V_{\max}$)**
- It is the substrate concentration **required to saturate half of all of the active sites of an enzyme**

Qualitatively (how it benefits us or what it indicates):

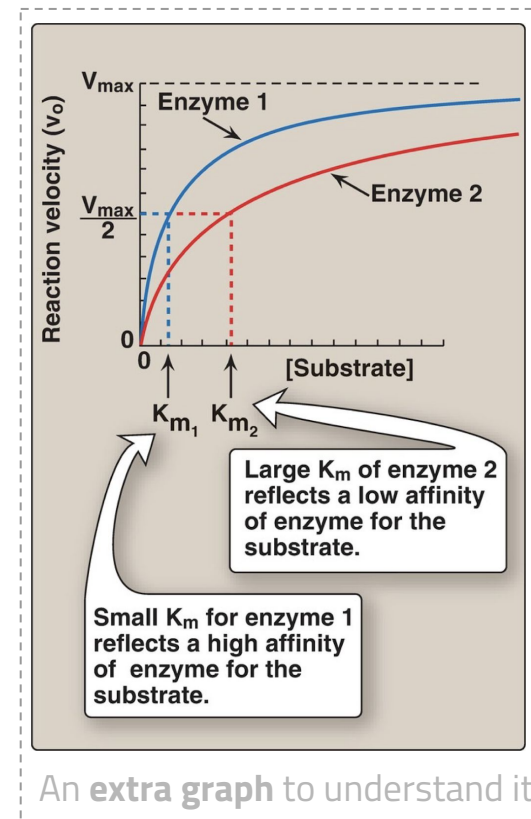
The K_m value of a substrate depends on its **affinity** with an enzyme

High K_m

means **low affinity** with enzyme
(**more substrate** is needed to saturate the enzyme)

Low K_m

means **high affinity** with enzyme
(**less substrate** is needed to saturate the enzyme)

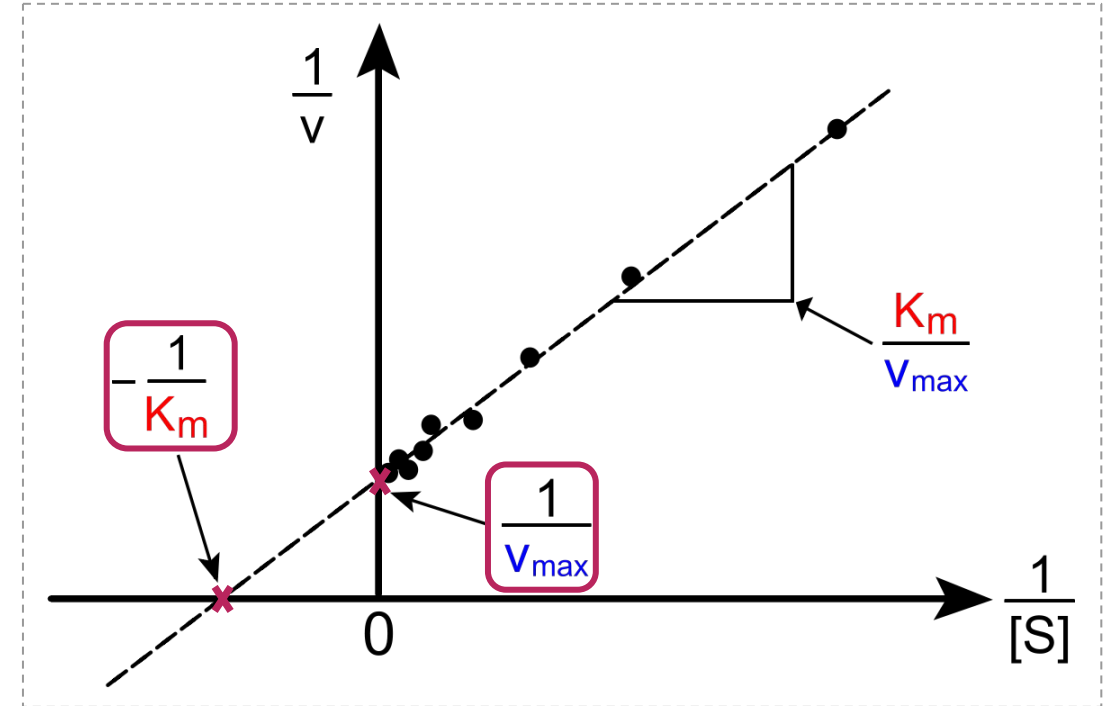


Lineweaver-Burk

- ❖ **Also called:** double-reciprocal plot
- ❖ **It's obtained from:** taking reciprocals (مقلوب) of the **Michaelis Menten equation**
- ❖ **It's plotted to:**
 - calculate the K_m and V_{max} values more accurately
 - determine the mechanism of action of enzyme inhibitors

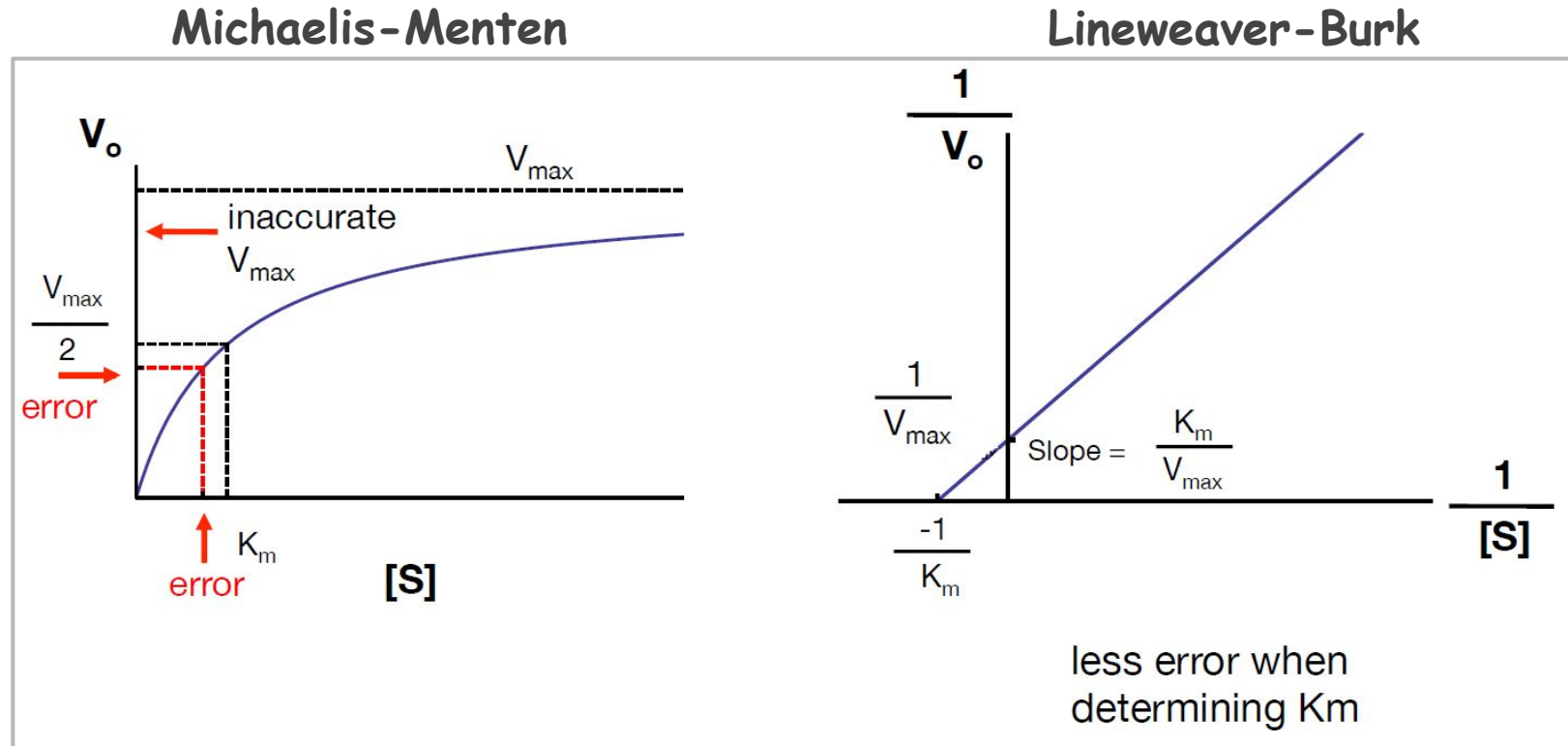
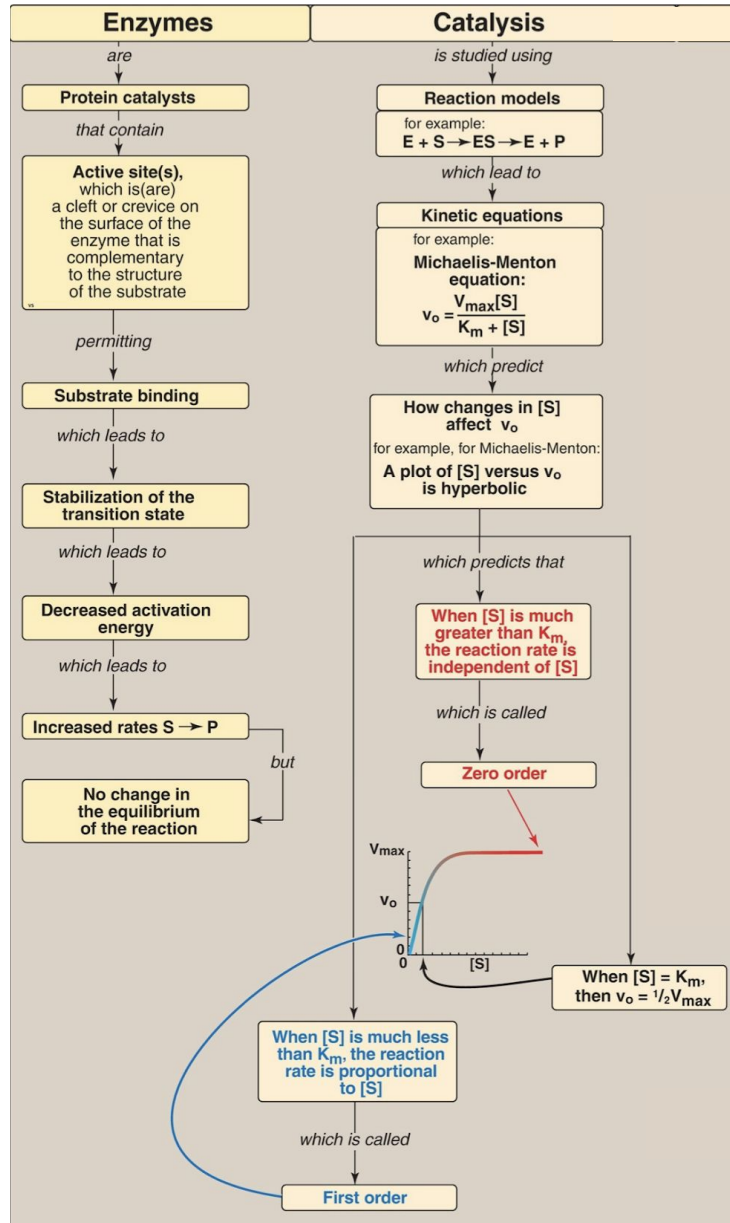
How is it different than the Michaelis-Menten plot?

The Michaelis-Menten plot is hyperbolic (curved), which makes it inaccurate as it gives multiple values for v_{max} . Thus, Lineweaver & Burk suggested to linearize it (graph the same data as a straight line) to improve accuracy.



Summary

Side-by side comparison



Quiz

Q1: Which class of enzymes requires ATP hydrolysis to function?

- A Hydrolases B Ligases C Lyases D Transferases

Q2: Which is true about isoenzymes?

- A They catalyze the same reaction B They have the same structure C They react with the same substrate D A & C

Q3: When enzymes decrease the activation energy, the free energy (ΔG) is:

- A Increased B Decreased C Unchanged D It depends

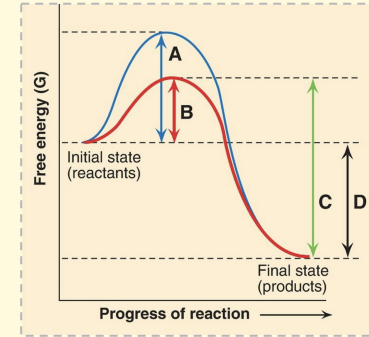
Q4: Which of the factors affecting enzyme activity follows a bell-shaped curve?

- A pH B temperature C [E] D [S]

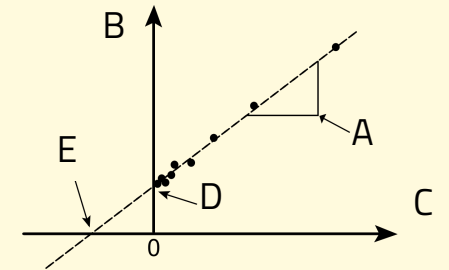
Q5: High K_m value indicates:

- A Low affinity with enzymes B High affinity with enzymes C Initial velocity equals zero D Steady state reaction

Answer Key: A (5) A (4) C (3) D (2) B (1)



Q6: Which letter represents the free energy of the reaction?

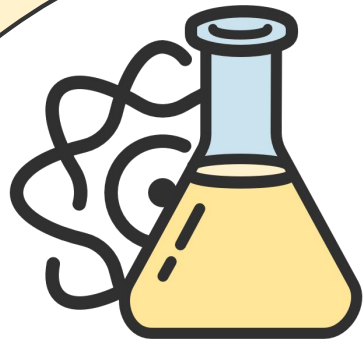


Q7: What does the letter D represent in the Lineweaver-Burk plot above?

Answers

Q6: D

Q7: V_{max} inverse ($1/V_{max}$)



Biochemistry 441

Girls



★ **Ghadah Alarify - Leader**

Yara Almufleh
Reema Alrashedi
Wareef Almousa
Joud Alangari
Fay Alluhaidan
Sarah Alhamlan
Arwa Almobeirek
Jumana AL-qahtani

Latifa Alkhdiri
Alanoud Alhaider
Futoon Almotairi
Manal Aldhirgham
Raaoum Jabor
Norah alawlah
Shahad Helmi
Rand Aldajani

Boys



★ **Khalid Alhamdi - Leader**

Ahmed Alayban
Sultan Alosaimi
Abdullah Alomran
Bassam Alghizzi
Ibrahim Aljurayyan
Mohammed Almutairi
Turki Alkhalifa
Malik Alshaya

Faisal Alhmoud
Abdulrahman Alnoshan
Ahmed Alqahtani
Hamad Alshaalan
Anas Alharbi
Mohammed Alwahibi
Saad Alghadir
Firas Alqahtani