

« وَلَا تَحْسَبَنَّ اللَّهُ عَاقِلًا عَمَّا يُعْمَلُ الظَّالِمُونَ »

Mays Aljundi ✓

Biochem ✓

رُبَمَا عَلَى حَاءِ الْحَيْرَةِ

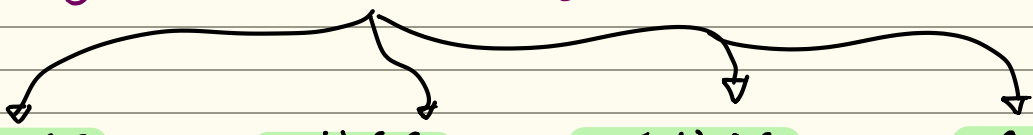
- نقطة لا تراها -

ENZYMES CLASSES

* Enzymes are classified into 6 groups:

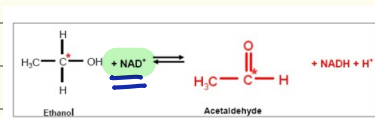
① Oxidoreductases (Largest group):

- Catalyze oxidation/reduction reactions involving the transfer of H atoms or e⁻
- They can be divided into 4 groups:



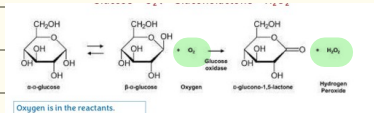
Dehydrogenases

- Removing H from substrate
- Need to have coenzymes:
 - ① NAD⁺ / NADH
 - ② FAD / FADH₂



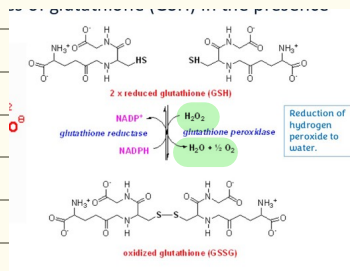
Oxidases

- O₂ considered as substrate
- H₂O₂ product



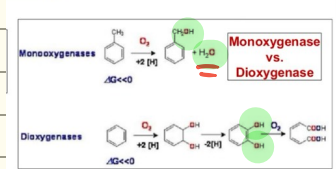
Peroxidases

- Oxidation by H₂O₂
- H₂O₂ serves as a substrate → H₂O₂ → H₂O + 1/2 O₂



Oxygenases

- Introducing O₂ into the substrate
- Reduced product is H₂O

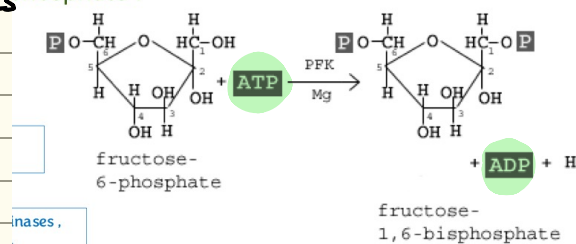


② Transferases:

- These enzymes transfer a functional group from one substrate to acceptor molecule
- Kinases are example (transferred group is phosphate)

Kinases phosphate:

((()))

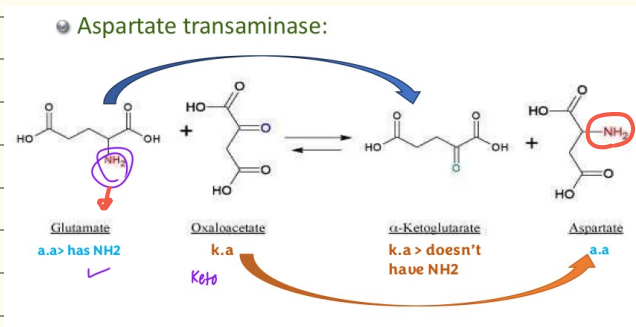


- ① you can see ATP
- ② Addition of phosphate group

هو اخذنا فوسفات من ATP يعني دخلت
جاء rxn

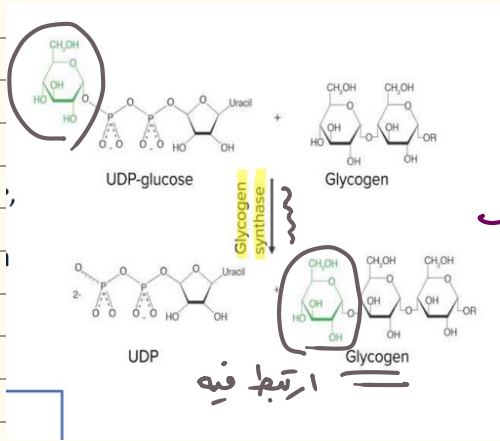
transaminases

((()))



- ① Transfer an amino functional group from amino acid to keto acid
- (((NH2), Keto acid)) amino acid (())

Synthases



⊙ The compound is physiologically IMP

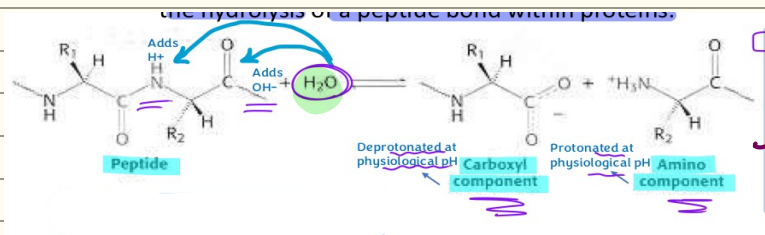


← تفرق بينه وبين الـ Synthase بقا الـ lyases
 «فانتزف double bond جديدة أو ما انتزف قول
 « single bond أو ما انتزف قول»

⊙ **Hydrolyases:**

- Catalyze cleavage reactions using water
- Such as: proteases, esterases, lipases, digestive enzymes

like: Trypsin, chymotrypsin, elastase

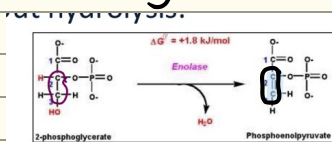


⊙ H₂O substrate

⊙ **Lyases:** (without H₂O, coenzymes, isomers etc.)

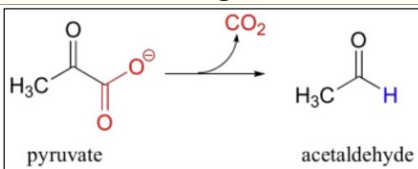
- cleave C-C, C-O, C-N and other bonds leaving double bonds or rings
- Such as:

▷ **Dehydrases**



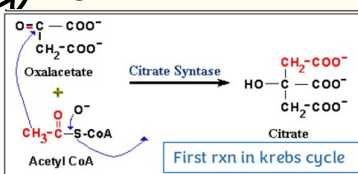
⇒ Removing H₂O from the substrate to give double bond

▷ **Decarboxylases**



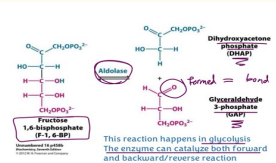
⇒ Replacement of carboxyl group by H

▷ **Synthases**



⇒ Addition of small molecules to a double bond

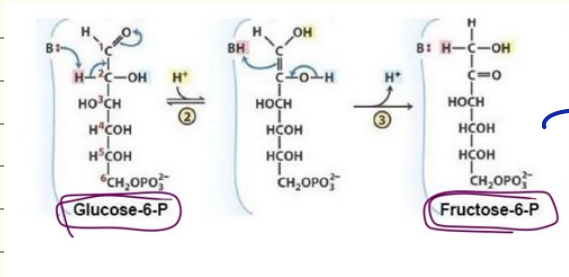
▷ **Special: Aldolase**



⇒ Breaks down the substrate into 2 products and forms double bond

⑤ Isomerases:

- Convert an isomer into another
- They don't require energy
- They rearrange the bond structure of a compound



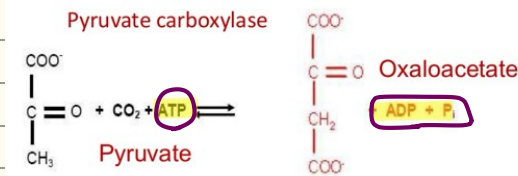
تخیرتے انکان اور bonds فتحول سے isomer
 ر isomer تانی

⑥ Ligases:

- join C-C, C-O, C-N bonds
- connect 2 molecules together
- use energy derived from ATP

ATP Substrate
 ADP product

ما باخذ functional group من
 ATP انتبھی



فانی phosphate transfer منو Ligase



اللہ یوفقم ویسرکم امرکم

"اللهم إني فوضت أمري إليك فأكفني،

واصرف عني ما يقلقني، وتولني بفضلك يا الله"

ENZYMES REGULATION

enzyme amount ↑

slow mechanisms

gene level
proteases
synthesis for isozymes

التقسيم (compartmentalization) بجزي

limit diffusion rate to produce higher collision rate
such as: lysosomal + fatty acid metabolism

Synthesis: cytosol degradation: Mitochondrial

هذو القتين ينذروا تحت
non-specific regulation
Temperature + pH مع

enzyme complexing نفس الي فوق بجزي limit

(oxidoreductases) ex: pyruvate dehydrogenase

60 subunit

decarboxylase (30) oxidation (20) Transfer (10)

2C, COA 3C, pyruvate

isoforms هذو نفس ال enzymes مختلف

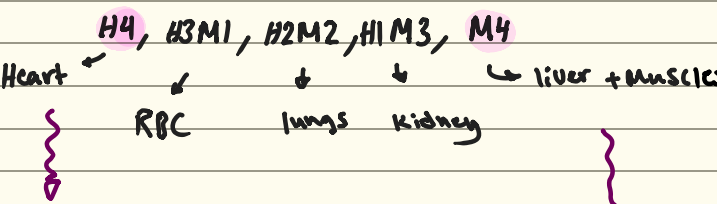
They catalyse same rxn

ال isozymes انما

diff genes
diff tissues
diff V_{max} & K_m

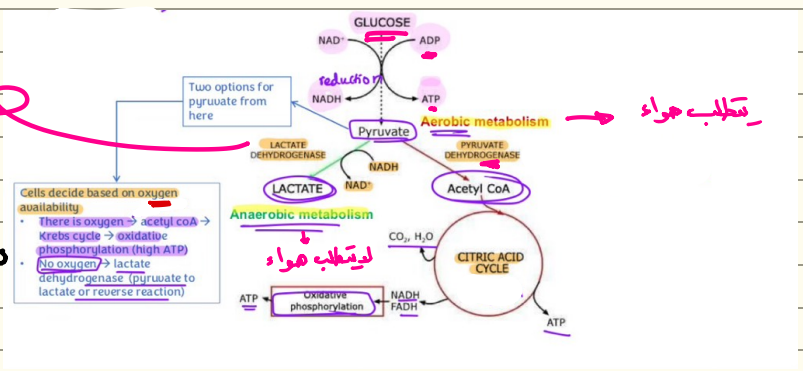
Aerobic vs Anaerobic metabolism

tetramer (reversible) rxn ال
عند 5 isozymes هم



LDH1 prefers to:
Lactate → pyruvate
High efficiency towards pyruvate

LDH5 prefers to:
pyruvate → Lactate
High efficiency towards lactate



* Differences between hexokinases + Glucokinases:

allosteric isozymes

NOTE: When glucose is phosphorylated it won't cross PM cuz of phosphate

① Hexokinase:

- A RBC + skeletal muscles enzyme } High efficiency to trap glucose
- Allosteric isozyme (phosphorylates glucose to glucose-6-phosphate)

1. Allosteric Enzymes:

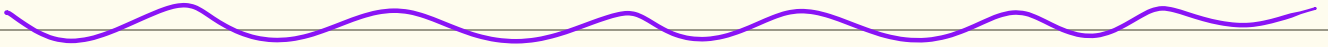
- Definition: Allosteric enzymes are enzymes that can be regulated by molecules binding to sites other than their active sites. These alternative binding sites are called allosteric sites.

favors T State

- It is inhibited by glucose-6-phosphate
- Low K_m , High affinity, low v_{max}

② Glucokinases:

- A Liver enzyme } low efficiency
- It is activated by insulin and inhibited by glucagon
- High K_m , low affinity



Regulation of enzymatic activity

INHIBITORS

Regulation of enzymatic activity

Inhibitors

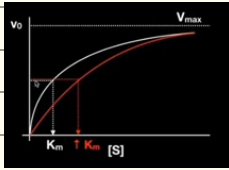
All physiological inhibitor

Competitive
non Competitive

Reversible
Irreversible

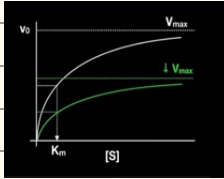
low concentration of active enzyme
(Synthetic inhibitors)

Covalent
Transition State analog
Heavy metals



Competitive inhibition

Substrate compete with enzyme.
Inhibition is reversible.
Vmax remains the same but Km increases.



non-competitive inhibition

Inhibitor binds to E or ES complex, not the catalytic site.
Km remains the same but Vmax decreases.

Irreversible

Covalent inhibitors

tight covalent bond

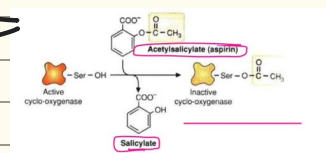
Example: diisopropyl fluorophosphate (DFP)

it inhibits acetylcholinesterase

acetylcholine degradation (paralysis) → acetylcholine → acetyl + choline

Aspirin (acetylsalicylic acid)

prostaglandin precursor → acetyl group



Transition State analogs

يربط مع Sub و يخلق
 يحد من ال enzyme من عمله اسprin
 يربط Tightly و يربط covalent
 ال drugs ما بقدر نفسها بحيث تكاثر ال transition State

Substrate analogs
Suicide inhibitors

هذه اهم

1 Methotrexate

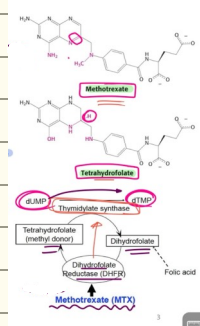
Synthetic inhibitor يعالج ال cancer , rheumatoid

يعتبر Structural analog ال folat

يعتبر Sub للانزيم dihydrofolate

يعتبر Coenzyme ال thymidylase

يتم تثبيط تكوين Nucleotide base من خلال تثبيط الانزيم ال
 يمنع deoxyTMP



2 Penicillin

The first antibiotic to be discovered.

بنفس Structure تجع alanyl-alanine

يعمل تثبيط ال glycopeptidyl transpeptidase

هو بيبي و بي attach مع penicillin و بيكلم الملقمة تبعت lactam β , و بيغير مفتوحة bacteria cell wall

Heavy metals

تعتبر non-specific على doses عالية

ال ال و ال يربط مع Sulfhydryl , بعدهم عن ال active site

ال ال Pb , Hg , Cu²⁺ , Fe²⁺ , Mn²⁺ عن طريق irreversible mechanism

toxic

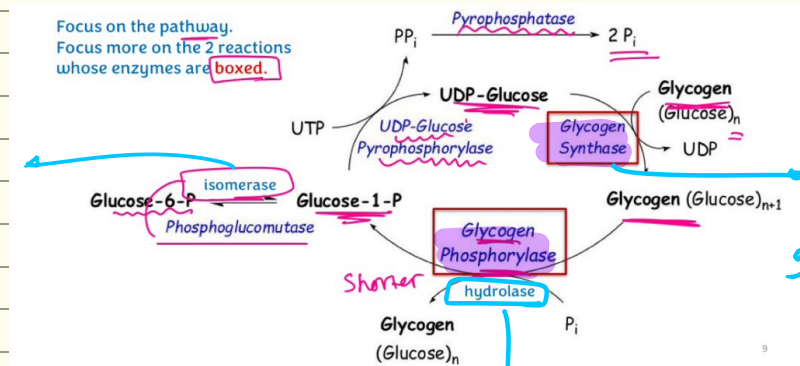
Regulation through conformational changes
through changing of the enzyme structure.



Regulation: conformational changes

- These regulatory mechanisms include
 - Allostery ✓
 - Covalent modulation ✓
 - Protein-protein interactions between regulatory & catalytic subunits or between two proteins; ✓
 - Proteolytic cleavage ✓

السكر (glycogen) و metabolism :
 - تحلل السكر (Synthesis) أو (degradation) في ال cytosol على عكس fatty acids metabolism



حول glucose 6P إلى glucose 1P

حول glucose 1P إلى glycogen

حول glycogen إلى glucose 1P

when inactive

2 catalytic → Quaternary ①
 2 Regulate → Serine / threonine ②

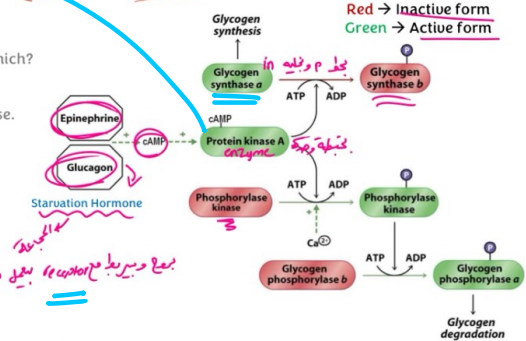
Regulation by phosphorylation

③ CAMP يرتبط فيه ليتغير (4 CAMP are needed)

adenyly cyclase هو

Focus on

Which enzyme acts on which? What is the effect? Active/Inactive forms. The physiological response.



يتم تفعيله بـ CAMP في ال activation

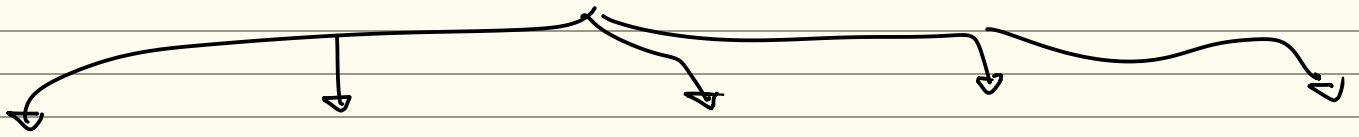
Reversible covalent modification

Reversible covalent modification

phosphorylation نزيد فوسفاتين phosphoryl donor ATP
 dephosphorylation نزيل فوسفاتين hydrolysis

phosphatases كالتالي Kinases
 amplification بتال

أشكاله :



Adenylation

Uridylation

ADP-ribosylation

Methylation

Acetylation

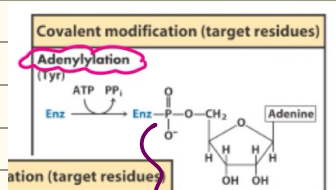
• Addition of adenylyl group

• Addition of uridylyl group

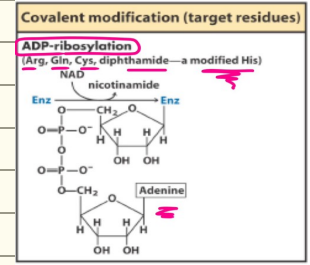
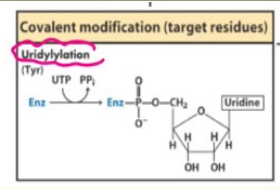
• Addition of adenosine diphosphate ribosyl group

• Addition of methyl group

• Addition of acetyl group
 $\text{CH}_3-\text{C}(=\text{O})-$

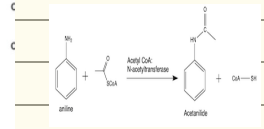
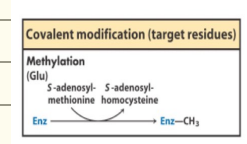


phosphodiester linkage



• Masking (-) charges

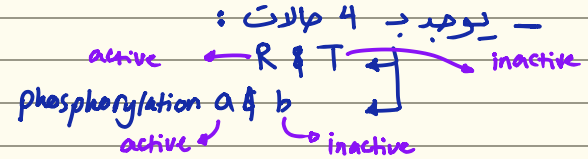
• MASKING (+) charges



allostery \leftarrow تنظيم

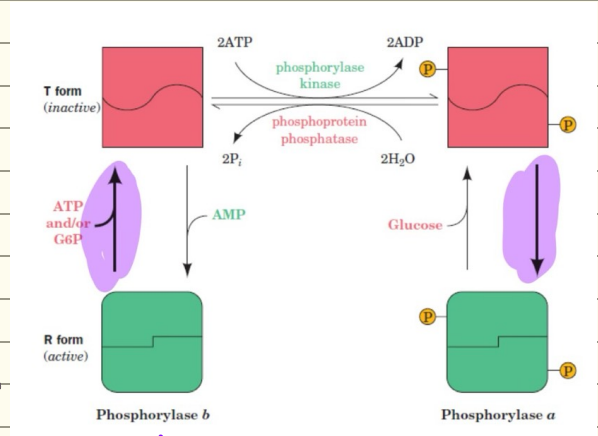
Glycogen phosphorylase *

allosteric



(Negative allosteric) ATP favors T State
 (positive allosteric) AMP favors R State

وفن الشيء \rightarrow phosphorylase a \leftarrow usually inactive cuz the equilibrium favors T state
 لا ينحاز يكون R State

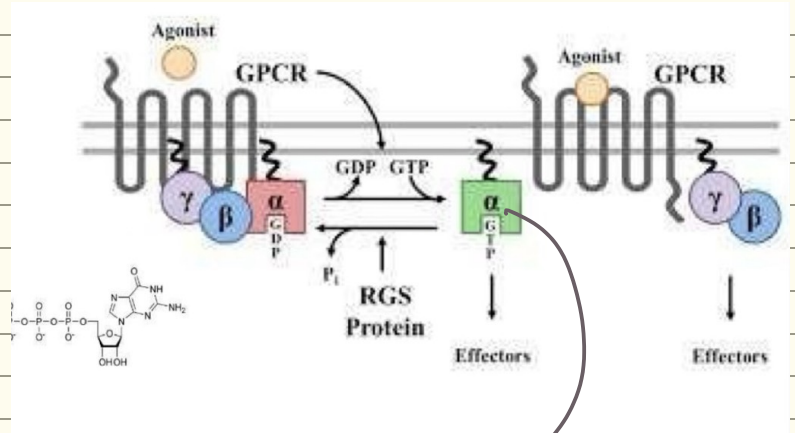


ندرج كنت

protein-protein interactions between
regulatory & Catalytic subunits or between
2 proteins

: G proteins ✖

- ↓, β, α ← Trimeric
- G protein-coupled receptors يوصلوا الرائل من Signals محمدك غير
- مهمين لـ Smell و taste و vision و hearing
- عندهم 7 transmembrane domains receptors
- لا يصير في replacement من GDP → GTP active
- لا يصير في hydrolysis من GTP → GDP inactive



يمكن تربط مع CAMP والي تربط بعدين مع PKA

تجزئة بروتين
proteolytic cleavage

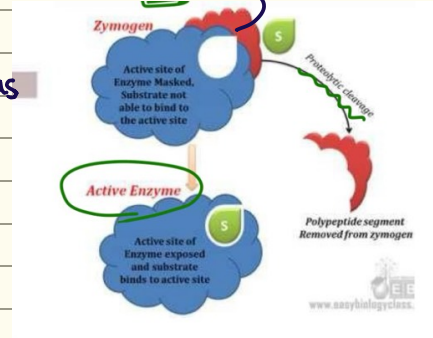
: (proteolytic activation) Irreversible covalent modification *

: (Trypsinogen) Zymogens

They are proenzymes that require proteolysis (lysis/cleavage of specific regions of their protein structure) in order to be active.

← 6 amino acids متعلق

pro region
present at N terminus



* موجود في بطن pancreas

Allosteric regulation

: Allosteric enzymes *

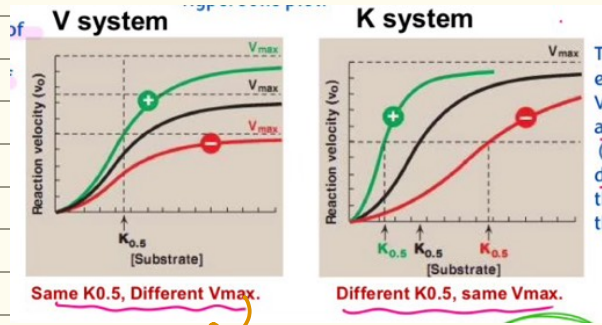
multi-subunit proteins

Quaternary

The binding of regulatory molecules triggers conformational changes in active site

allosteric site → *بيير بطرا*

Positive cooperativity means that the binding of one substrate to the active site of one subunit, makes it easier for another substrate to bind to the active site of another subunit.



بأثر على Vmax

بأثر على K0.5

: Notes

Allosteric enzymes and metabolism

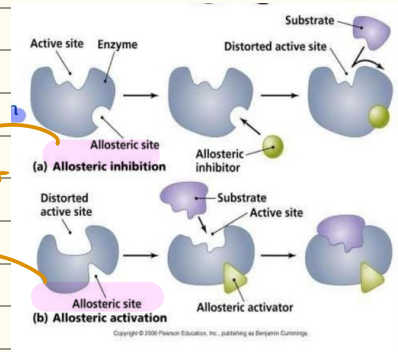
Recall that: the regulation of the enzyme is gradient. It changes its shape gradually, not in an on/off situation.

- Allosteric inhibitors usually have a much stronger effect on enzyme velocity than competitive and noncompetitive inhibitors.
- Allosteric enzymes are not limited to regulation through inhibition whereby allosteric effectors may function as activators.
- The allosteric effector needs not bear any resemblance to substrate or product of the enzyme.
- The effect of an allosteric effector is rapid occurring as soon as its concentration changes in the cell. It's also a huge effect on the enzyme's activity.
- Feedback regulation of metabolic pathways by end products or by signal molecules that coordinate multiple pathways.

Which is a difference between them and Michaelis-Menten's regulators that only function as inhibitors.

less active
Substrate wont fit

more active
Substrate will fit



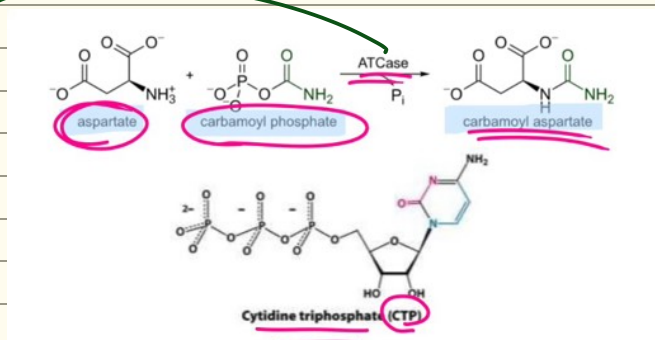
(ATPase) : Aspartate transcarbamoylase

هو مسؤول لادول خلو الكلى
12 polypeptide chains فيه

6 catalytic
6 regulatory

2 forms → موجود
R form
T form

inhibited by CTP



اليسر لله بكم كاف عباده

@ishaiokh