

Slide : Enzymes 6

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Sections : 4,5,6

■ Slide □ Sheet



Biochemistry

biometrics
cybernetics
ecology
taxonomy
bionomics
biophysics
bacteriology
biological
radiobiology
anatomy
microbiology
science
life
cystology
xenobiology
embryology
exobiology
gnotobiotics
pharmacology
astrobiology
biochemistry
physiology
ethnobiology
molecular
bioecology
virology
zoology
biometry
enzymology
genetics
bionics



B Covalent Modification:-

1- Phosphorylation is a highly effective means of switching the activity of target proteins:-

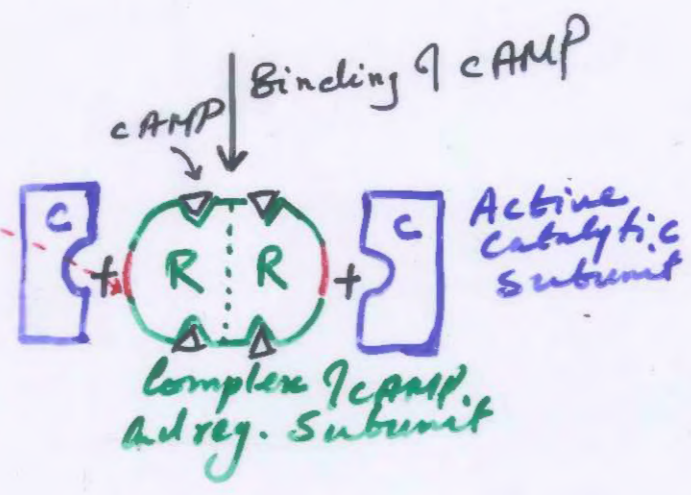
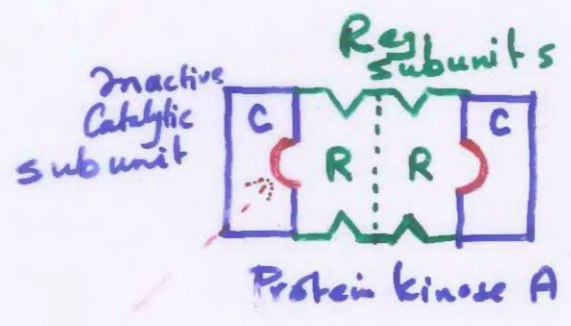
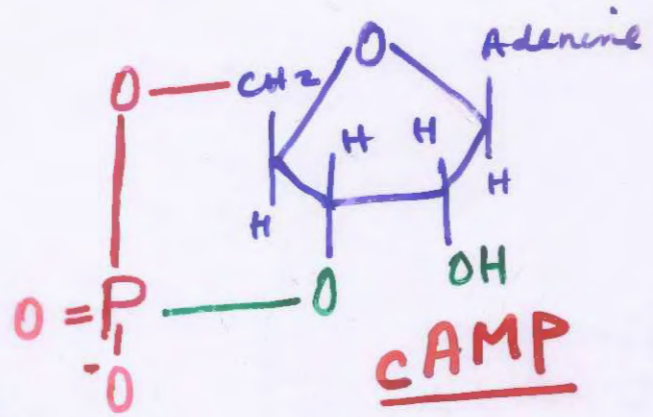
- Dedicated protein kinases - bind single protein
- Multifunctional protein kinases - phosphorylate many proteins
- Phosphorylation of Ser & Thr
- " " = Tyr. by tyr kinase
- Protein phosphatases :- reverse phosphorylation

Phosphorylation is a very effective process for controlling

- Phosphoryl gr. add two -ve charges
- Phosphoryl gr. can form three hydrogen bonds
- Free-energy of phosphorylation is large
~ - 12 Kcal/mole
- Phosphorylation-dephosphorylation can occur in less than a sec - min
- High amplification effect

Cyclic AMP activates Protein Kinase A by unleashing its catalytic subunits

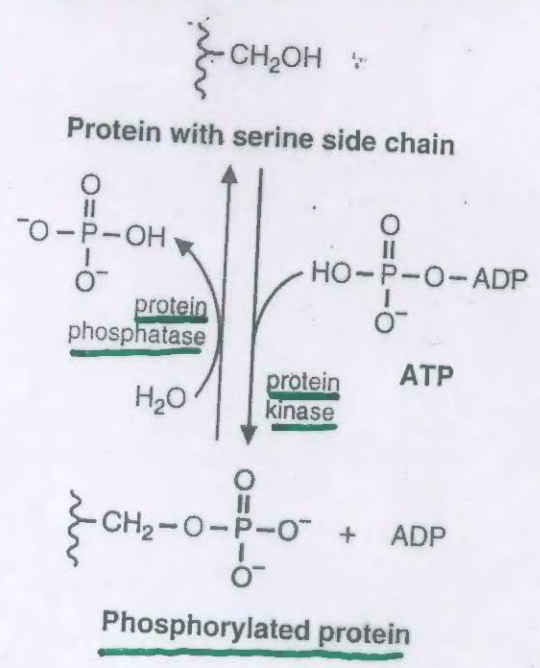
- PK-A
- cAMP



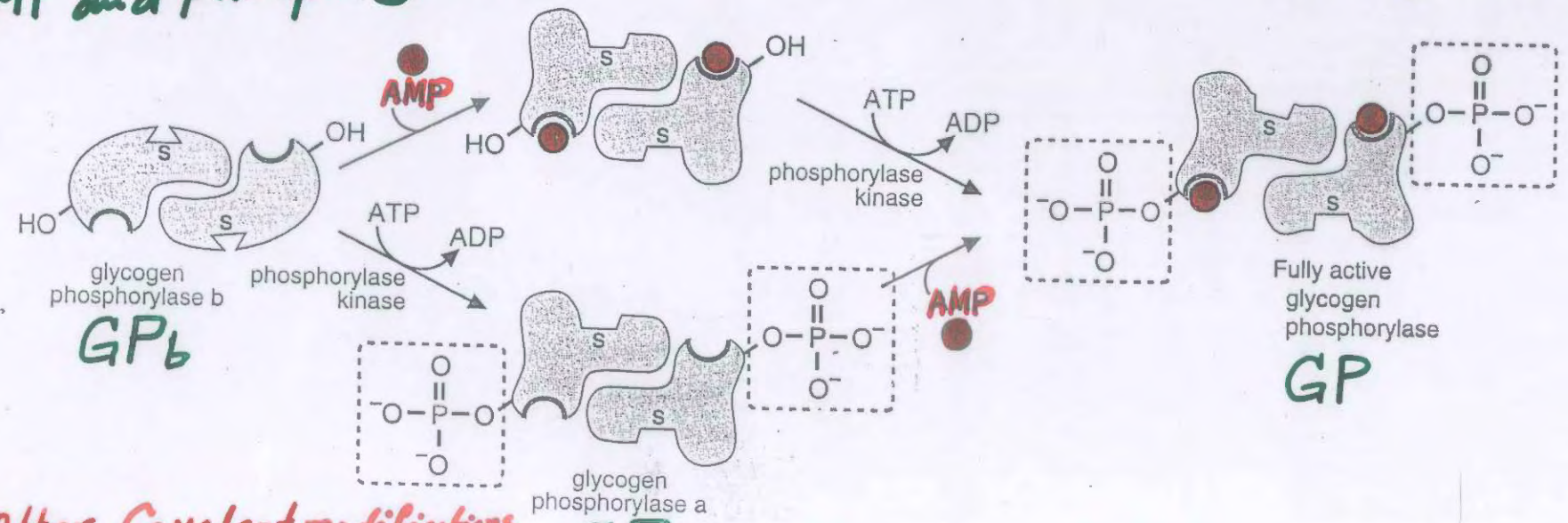
• Protein phosphatases

COVALENT MODIFICATION

(1) Phosphorylation — dephosphorylation of proteins



(2) Activation of GP by AMP and phosphorylation



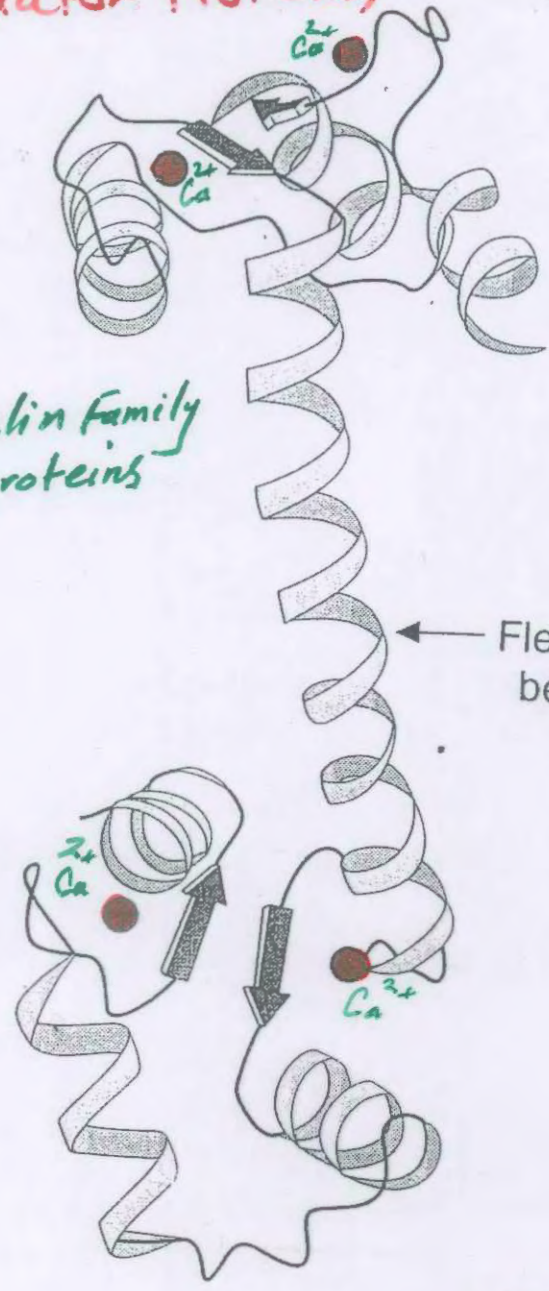
— Other Covalent modifications
 Acetyl-, ADP-ribose, lipid

GPa

C-Protein - Protein Interaction (Modulator Proteins)

(1)

Ca-Calmodulin Family
of modulator proteins



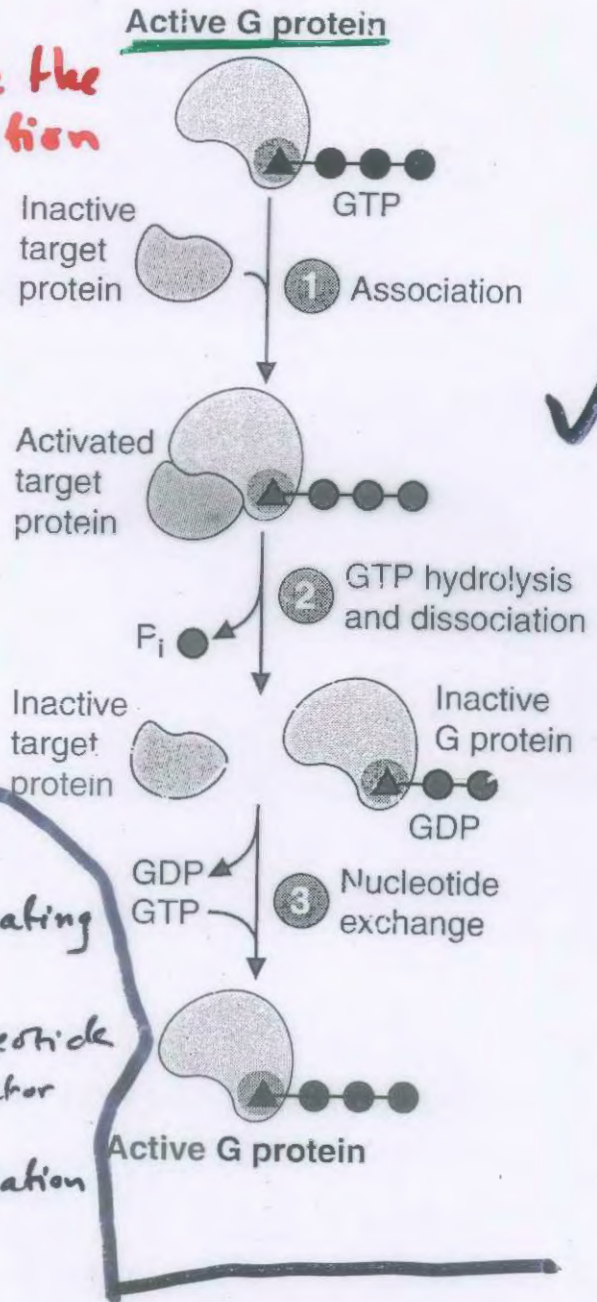
GP Kinase \uparrow by cAMP
also
GP Kinase \uparrow by Ca-Calmodulin

Flexible region
between domains

- **Tropinin C** binds Ca^{2+}
• regulatory subunit of tropinin - a regulator of
muscle contraction
• member of Ca^{2+} -Calmodulin superfamily

(2) Regulation by G proteins

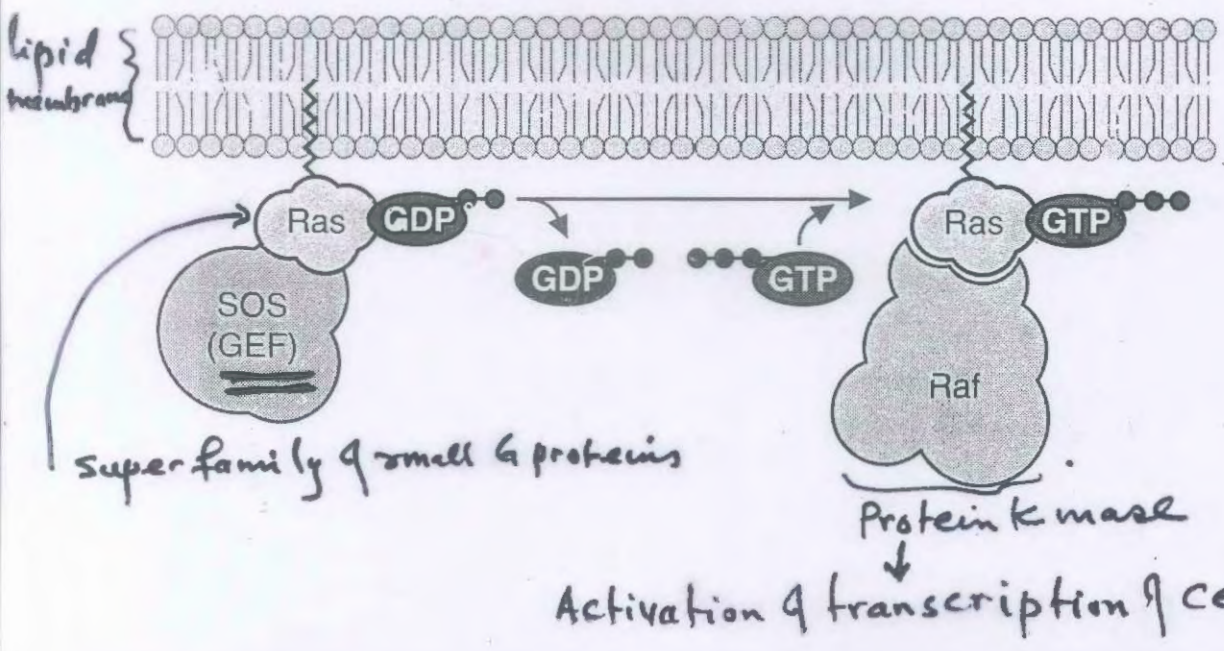
- G proteins are the master of regulation through reversible protein association



Reg. of G-proteins

- GAP - GTPase activating Proteins
- GEF - guanine nucleotide exchange factor
- GDI - GDP dissociation inhibitor

- ① Association of SOS and Ras
- ② Exchange of GTP for bound GDP
- ③ Ras-GTP binds Raf



Super family of small G proteins

Protein kinase

Activation of transcription of certain genes

D. PROTEOLYTIC CLEAVAGE

- Many Enzymes are Regulated by Specific Proteolytic Cleavage

- Digestive enzyme
- Blood clotting factors
- Pro enzyme or zymogen

- Isozymes Provide a Means of Regulation specific to Distinct tissues and Developmental stages

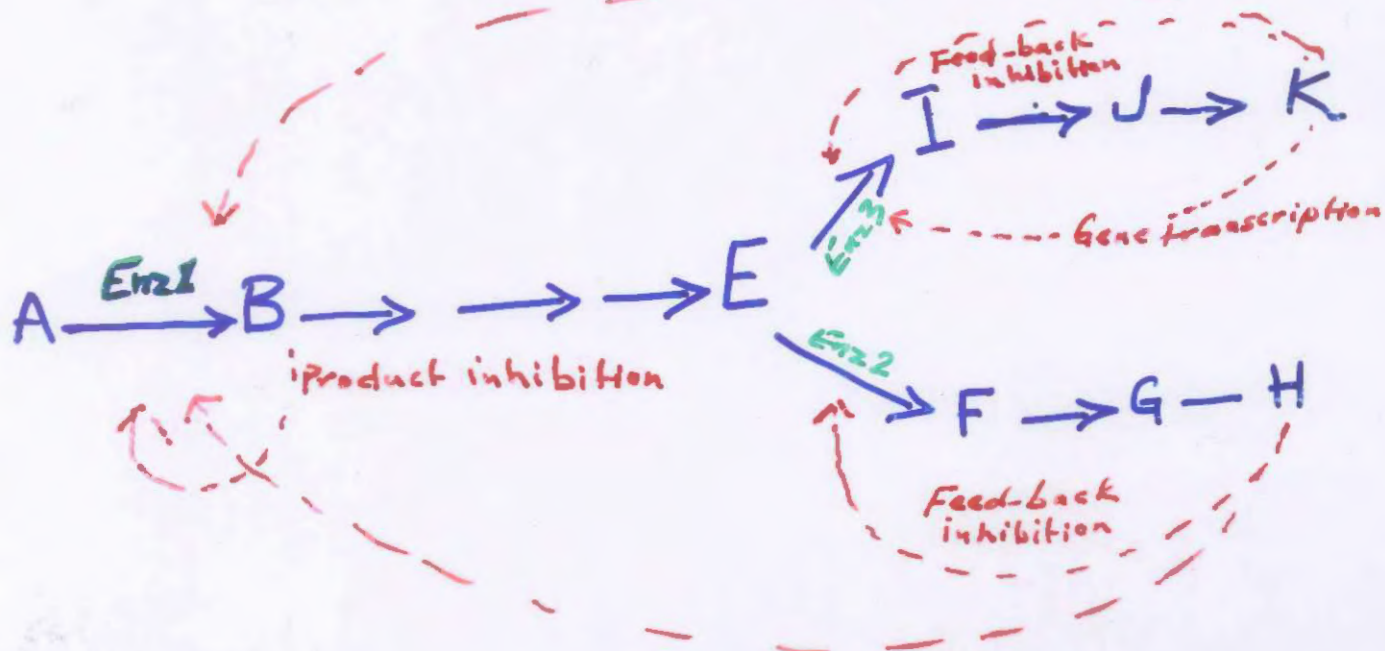
e.g. LDH

LDH-1 H ₄ Heart	LDH-2 H ₃ M ₁	LDH-3 H ₂ M ₂	LDH-4 H ₁ M ₃	LDH-5 M ₄ Liver, Muscle
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IV Regulation through Changes in Amount of Enzyme

- Enzyme synthesis
- Enzyme degradation

V Principles of Pathway Regulation ¹⁷



- Feedback regulation
- Feedforward regulation *as in enzymes involved in toxic compounds disposal*
- Product inhibition
- Synergistic effect

• Compartmentation

• Multi Enzyme Complexes

Mechanisms for regulating enzyme activity

REGULATOR EVENT

TYPICAL EFFECTOR

RESULTS TIME

REQ. FOR CHANGE

- Substrate availability

Substrate

change in velocity (v)

immediate

- Product inhibition

Reaction Product

changes in V_{max} and/or K_m immediate

- Allosteric control

Pathway end-product

changes in V_{max} and/or $K_{0.5}$ (S_{0.5}) immediate

- Covalent modification

another enzyme

changes in V_{max} and/or K_m immediate to minutes

- Synthesis or Degradation of Enzyme

has more or less metabolite

changing the amount of enzyme hours to days

Ribozymes (Catalytic RNAs)

Ribozymes are the RNA molecules that can act as catalysts.

The catalytic efficiency of catalytic RNAs is less than that of protein enzymes. The catalytic efficiency of known RNA is greatly enhanced by association with protein subunits.

Some Examples :-

- **Ribonuclease P (RNAase P)**
Catalyzes the maturation of tRNA by removing nucleotides from the 5'-end of the precursor mole.
- **Peptidyl transferase**
the RNA component of the ribosome that carries out protein synthesis
- **snRNA**
small nuclear RNA in association with proteins that catalyze splicing of exons to form final m-RNA