

Biochemistry

Metabolism

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Glycolysis

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FMN and FAD are able to sequentially transfer one e⁻

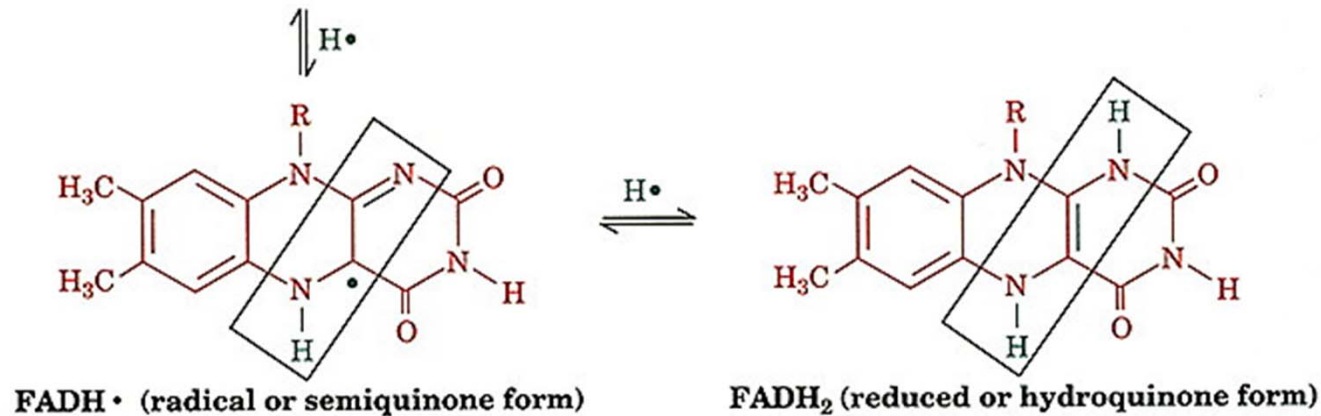
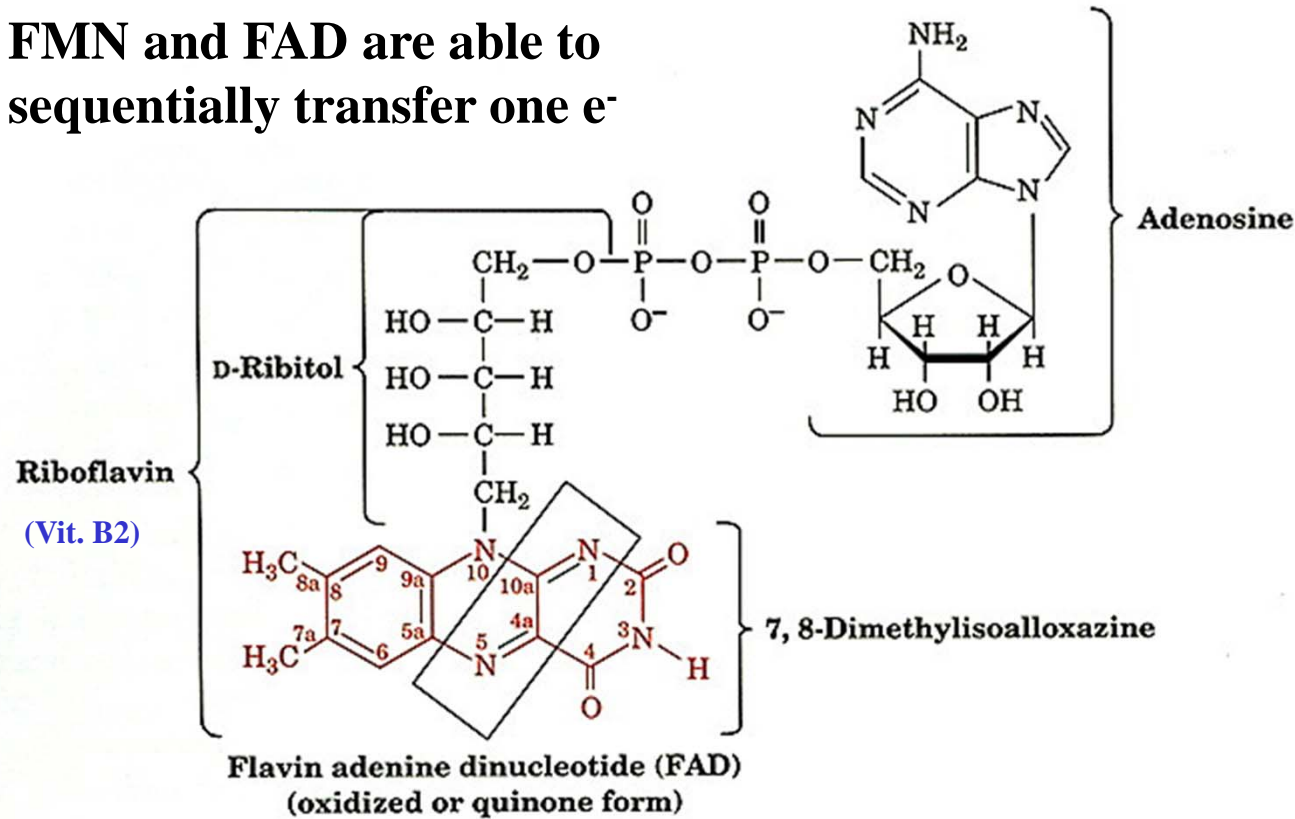
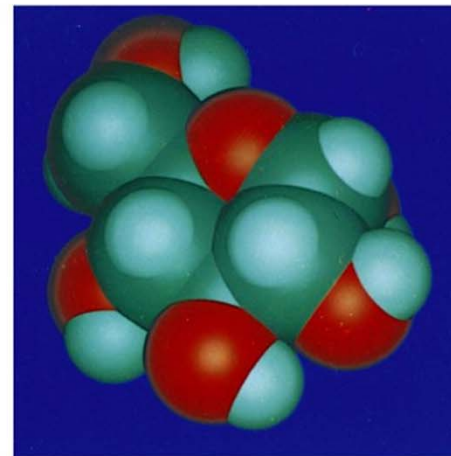
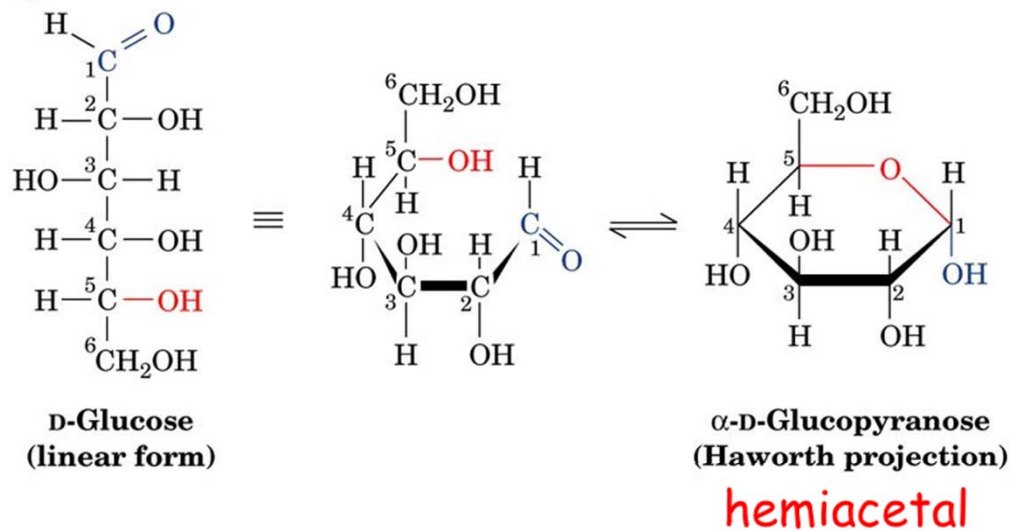


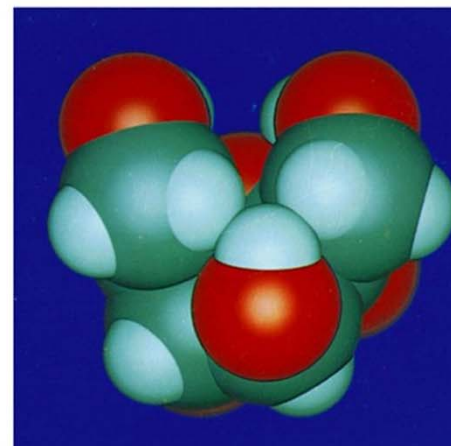
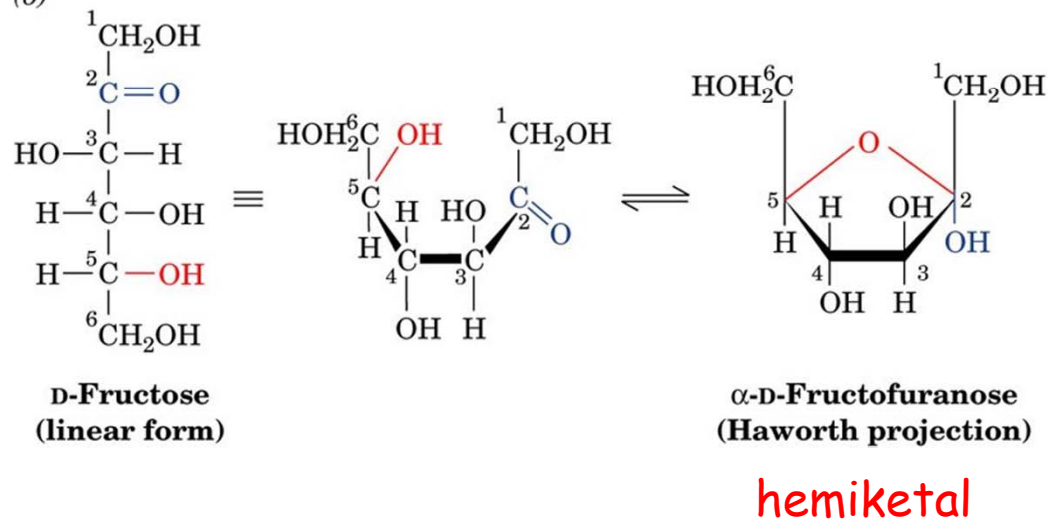
TABLE 13–9 Some Enzymes (Flavoproteins)
That Employ Flavin Nucleotide Coenzymes

<i>Enzyme</i>	<i>Flavin nucleotide</i>	<i>Text page(s)</i>
Acyl–CoA dehydrogenase	FAD	638
Dihydrolipoyl dehydrogenase	FAD	605
Succinate dehydrogenase	FAD	612
Glycerol 3-phosphate dehydrogenase	FAD	714–715
Thioredoxin reductase	FAD	869
NADH dehydrogenase (Complex I)	FMN	696–697
Glycolate oxidase	FMN	767

(a)



(b)



Space-filling models courtesy of Robert Stodola, Fox Chase Cancer Center

Figure 11-4 Cyclization reactions for hexoses.

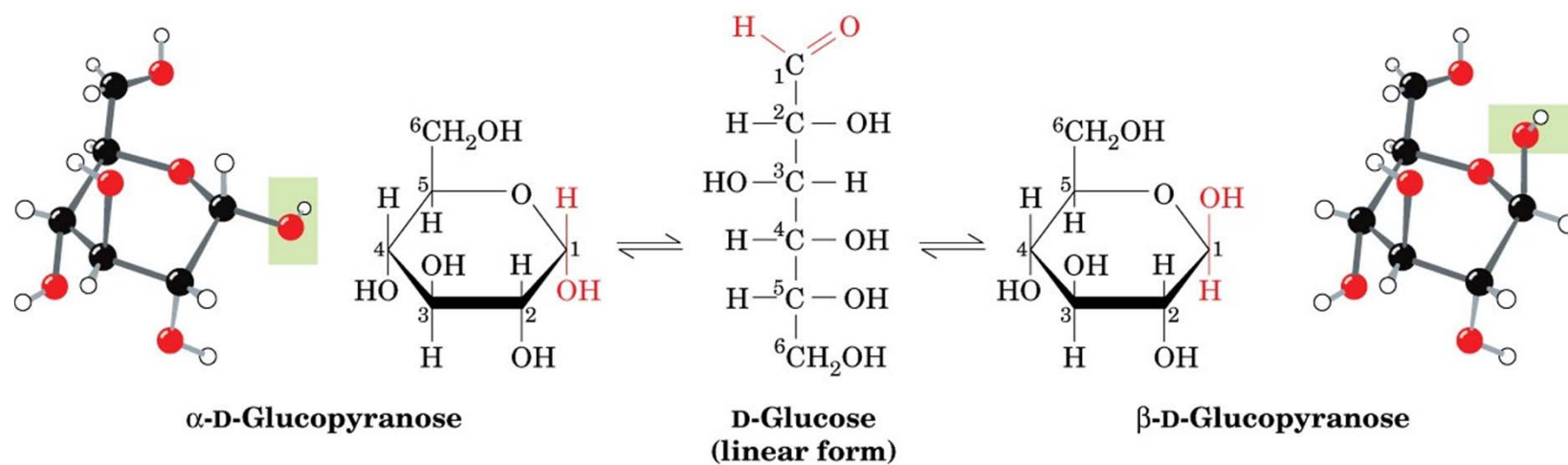
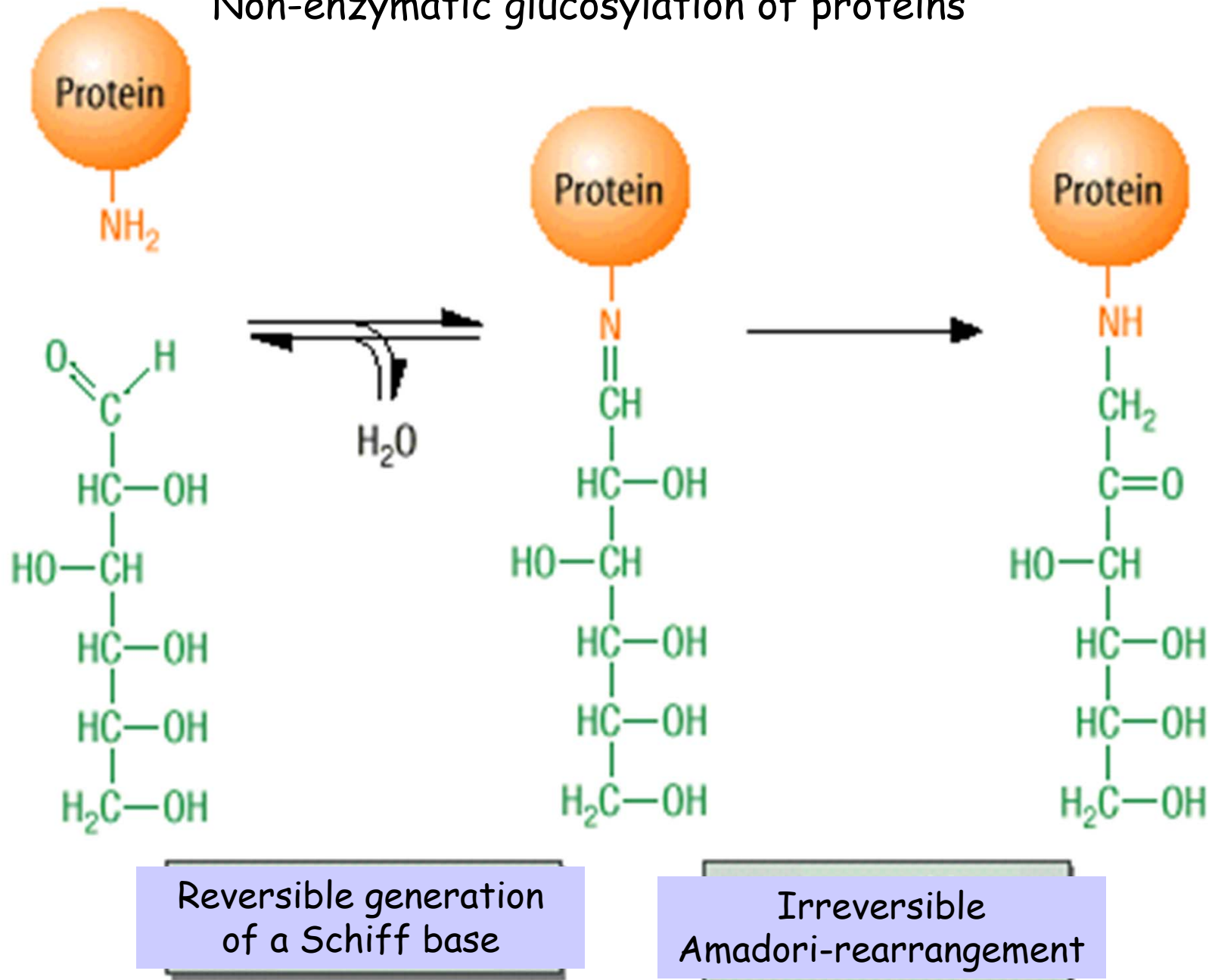


Figure 11-5 The anomeric monosaccharides α -D-glucopyranose and β -D-glucopyranose, drawn as both Haworth projections and ball-and-stick models.

Non-enzymatic glucosylation of proteins

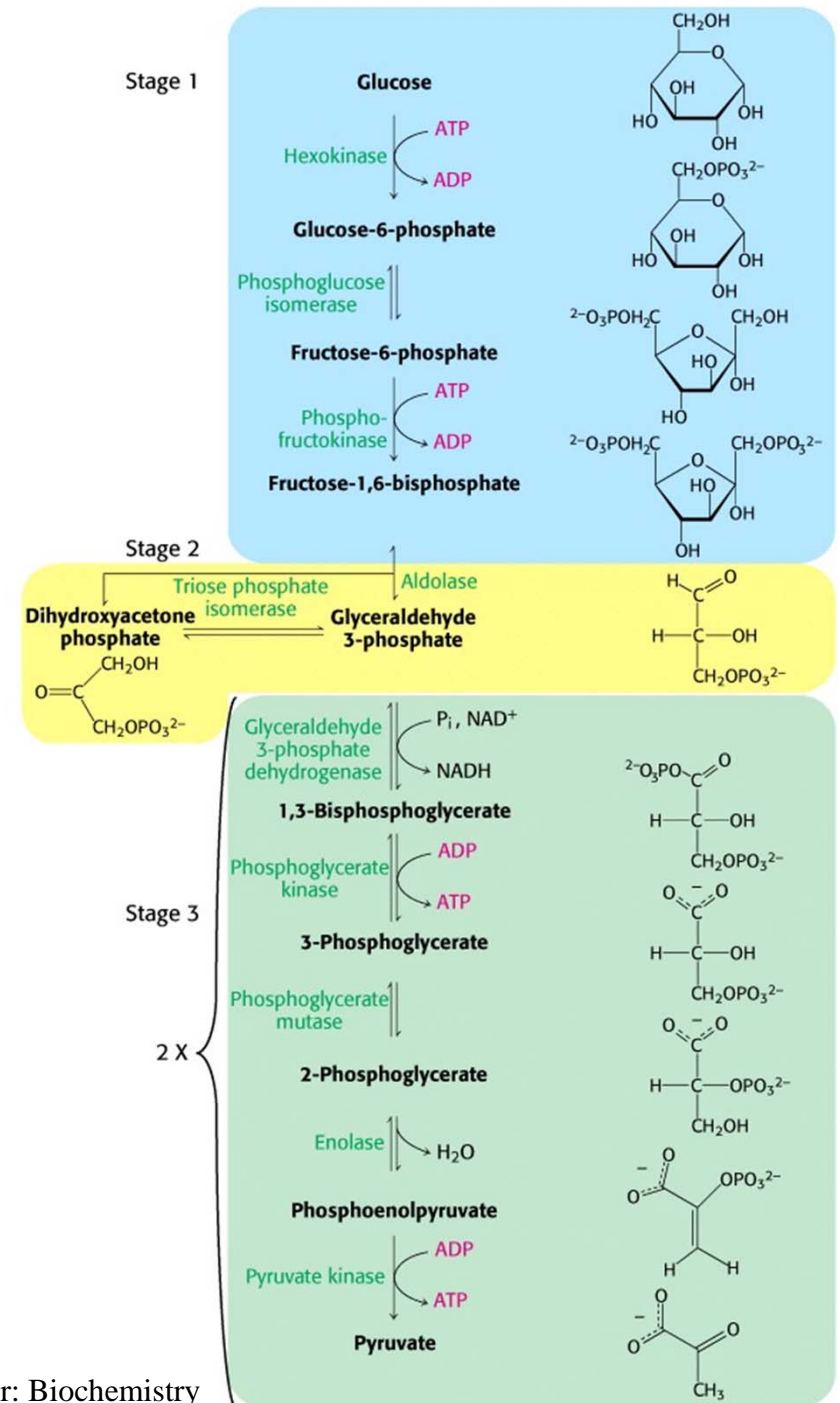
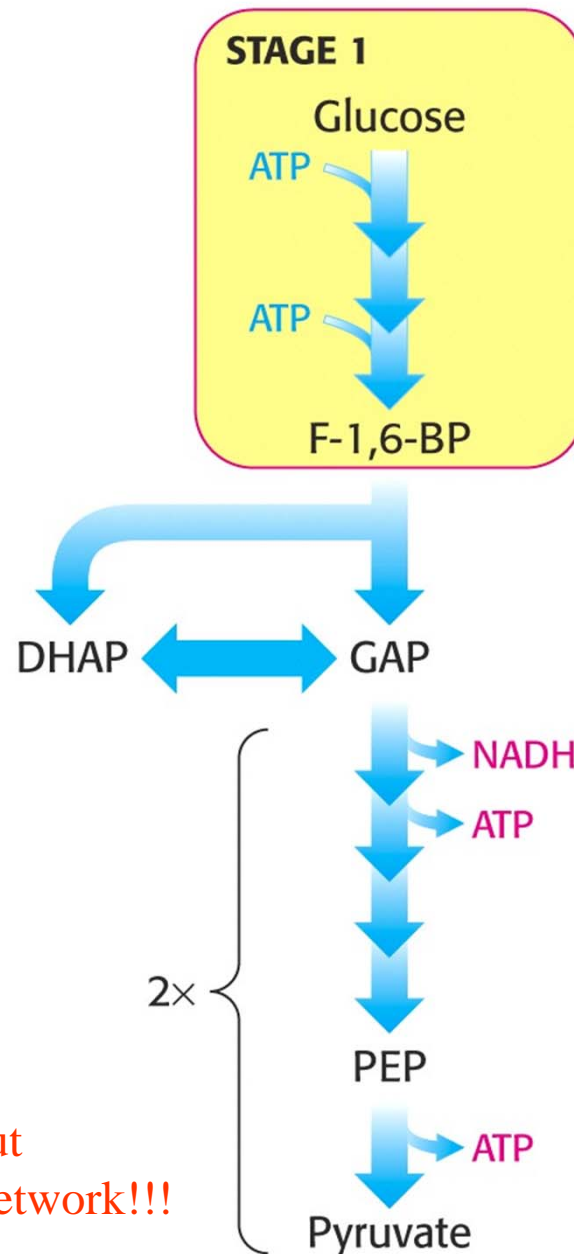


Glycolysis - overview

Preparatory phase:

Payoff phase:

Not an one-way road but
a branched metabolic network!!!



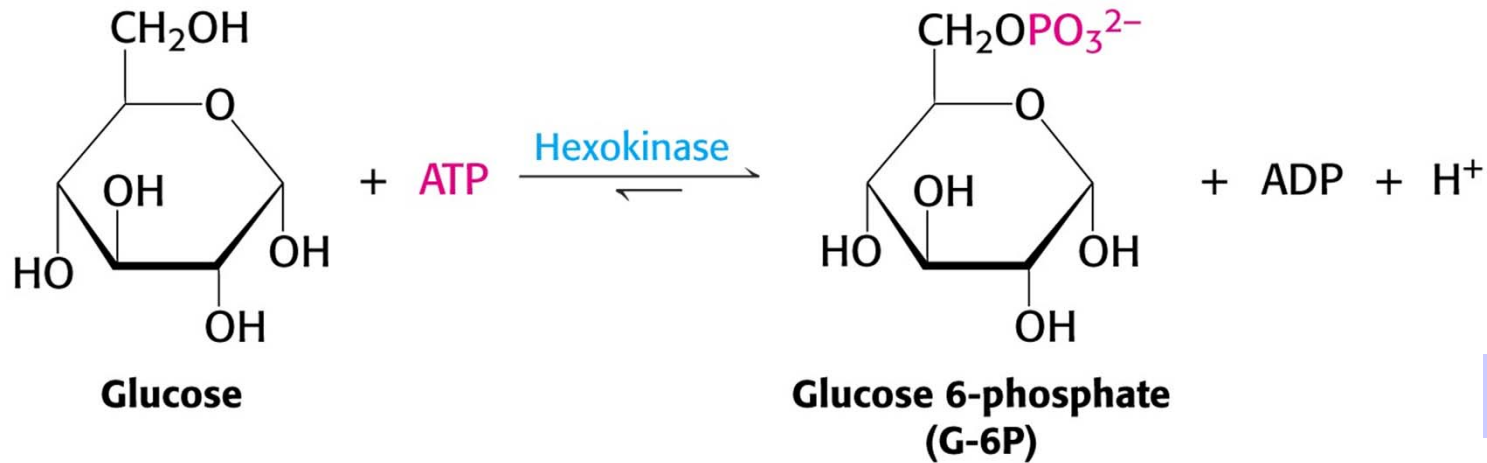
Energetics of glycolysis

TABLE 16.3 Reactions of glycolysis

Step	Reaction	Enzyme	Reaction type	$\Delta G^{\circ'}$ in kcal mol ⁻¹ (kJ mol ⁻¹)	ΔG in kcal mol ⁻¹ (kJ mol ⁻¹)
1	Glucose + ATP \longrightarrow glucose 6-phosphate + ADP + H ⁺	Hexokinase	Phosphoryl transfer	-4.0 (-16.7)	-8.0 (-33.5)
2	Glucose 6-phosphate \rightleftharpoons fructose 6-phosphate	Phosphoglucose isomerase	Isomerization	+0.4 (+1.7)	-0.6 (-2.5)
3	Fructose 6-phosphate + ATP \longrightarrow fructose 1,6-bisphosphate + ADP + H ⁺	Phosphofructokinase	Phosphoryl transfer	-3.4 (-14.2)	-5.3 (-22.2)
4	Fructose 1,6-bisphosphate \rightleftharpoons dihydroxyacetonephosphate + glyceraldehyde 3-phosphate	Aldolase	Aldol cleavage	+5.7 (+23.8)	-0.3 (-1.3)
5	Dihydroxyacetone phosphate \rightleftharpoons glyceraldehyde 3-phosphate	Triose phosphate isomerase	Isomerization	+1.8 (+7.5)	+0.6 (+2.5)
6	Glyceraldehyde 3-phosphate + P _i + NAD ⁺ \rightleftharpoons 1,3-bisphosphoglycerate + NADH + H ⁺	Glyceraldehyde 3-phosphate dehydrogenase	Phosphorylation coupled to oxidation	+1.5 (+6.3)	+0.6 (+2.5)
7	1,3-Bisphosphoglycerate + ADP \rightleftharpoons 3-phosphoglycerate + ATP	Phosphoglycerate kinase	Phosphoryl transfer	-4.5 (-18.8)	+0.3 (+1.3)
8	3-Phosphoglycerate \rightleftharpoons 2-phosphoglycerate	Phosphoglycerate mutase	Phosphoryl shift	+1.1 (+4.6)	+0.2 (+0.8)
9	2-Phosphoglycerate \rightleftharpoons phosphoenolpyruvate + H ₂ O	Enolase	Dehydration	+0.4 (+1.7)	-0.8 (-3.3)
10	Phosphoenolpyruvate + ADP + H ⁺ \longrightarrow pyruvate + ATP	Pyruvate kinase	Phosphoryl transfer	-7.5 (-31.4)	-4.0 (-16.7)

Note: ΔG , the actual free-energy change, has been calculated from $\Delta G^{\circ'}$ and known concentrations of reactants under typical physiologic conditions. Glycolysis can proceed only if the ΔG values of all reactions are negative. The small positive ΔG values of three of the above reactions indicate that the concentrations of metabolites in vivo in cells undergoing glycolysis are not precisely known.

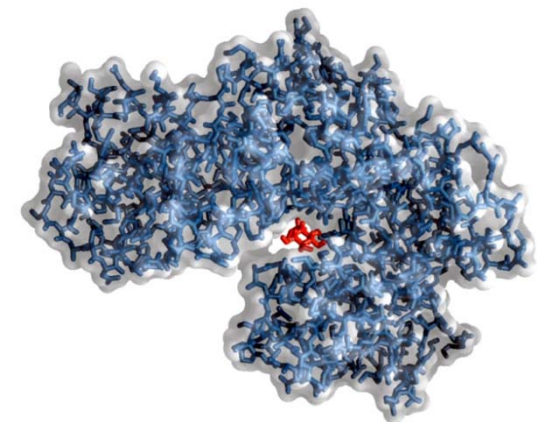
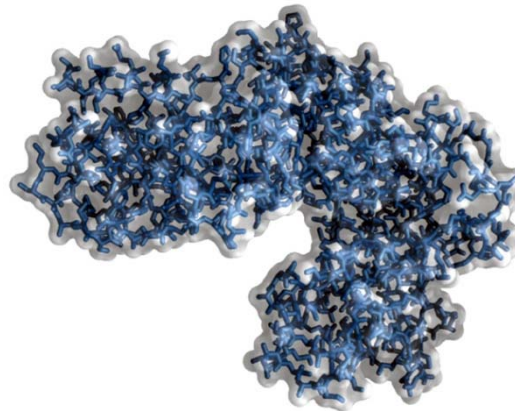
Glycolytic steps: Phosphorylation catalysed by **hexokinase**



„Isoenzyme“

- ubiquitous
- unspecific (catalyses phosphorylation of several hexoses)
- co-substrate Mg-ATP-complex
- $K_{\text{MGlc}} < 100 \mu\text{M}$!!!

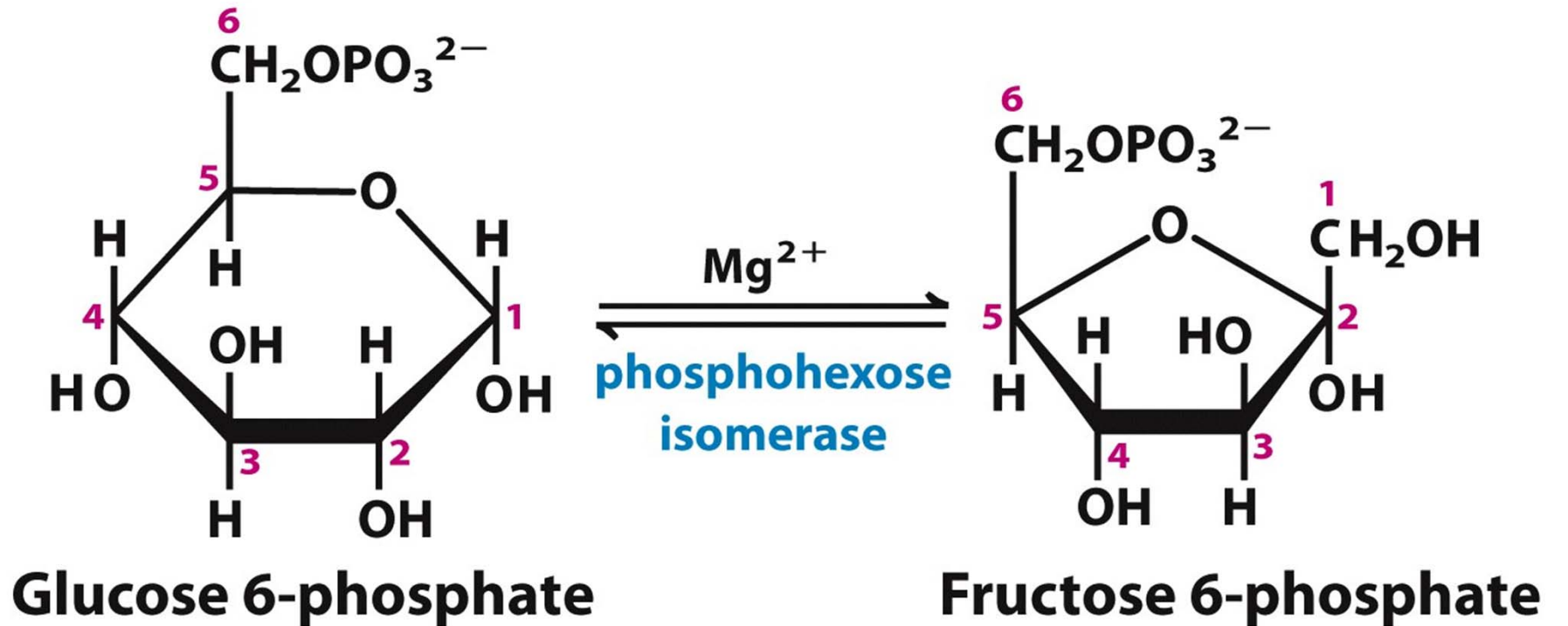
• Note:
in the liver **Glucokinase**:
 $K_{\text{MGlc}} = 10 \text{ mM}$



Induced fit

Berg, Tymoczko, Stryer: Biochemistry

Glycolytic steps: **Isomerisation** catalysed by **glucose phosphate isomerase**

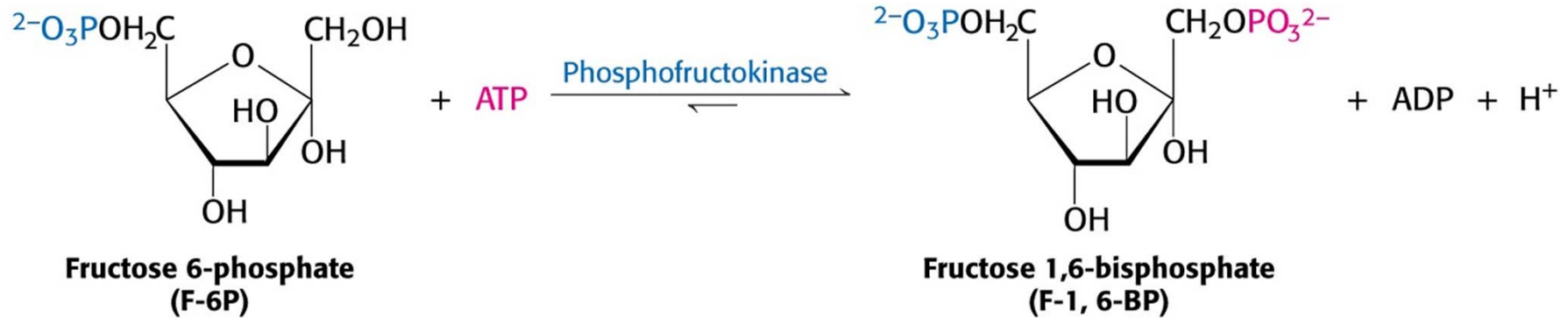


$$\Delta G'^{\circ} = 1.7 \text{ kJ/mol}$$

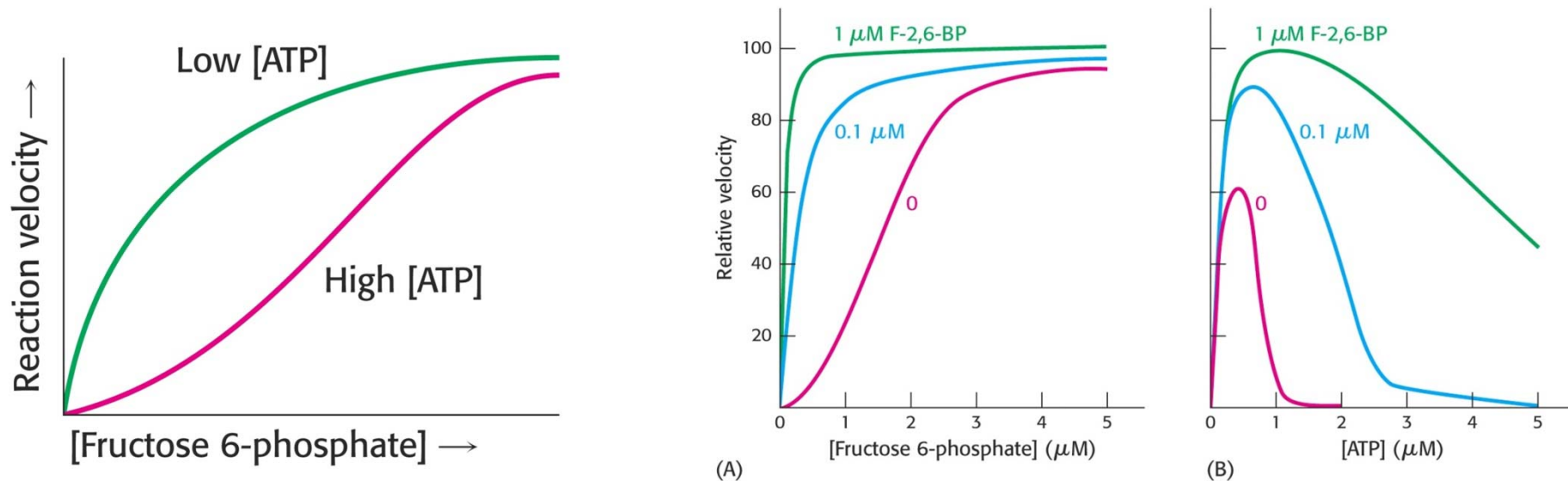
Reaction mechanism:

General acid/base catalysis

Glycolytic steps: **Phosphorylation** catalysed by **phosphofructokinase (PFK)**
(rate-limiting step)

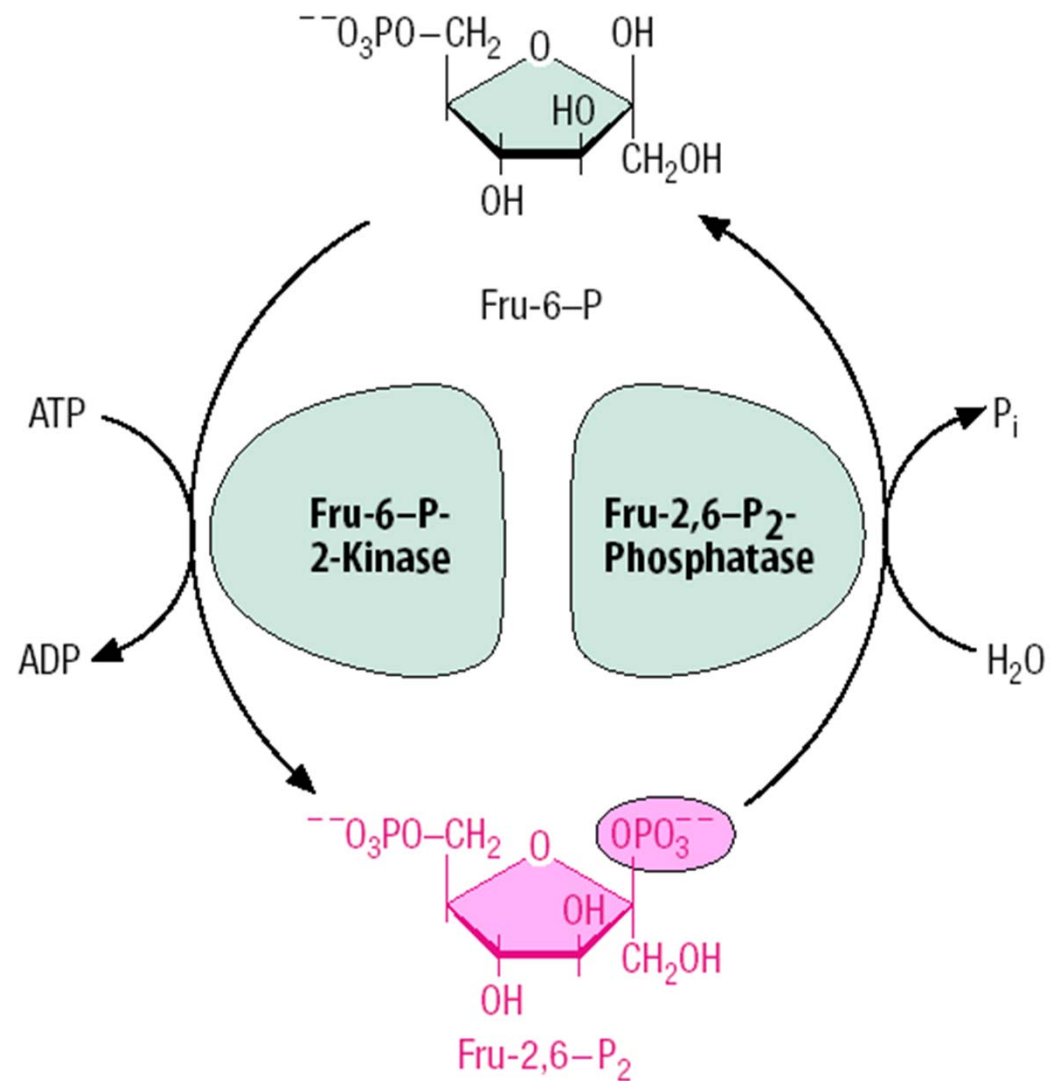
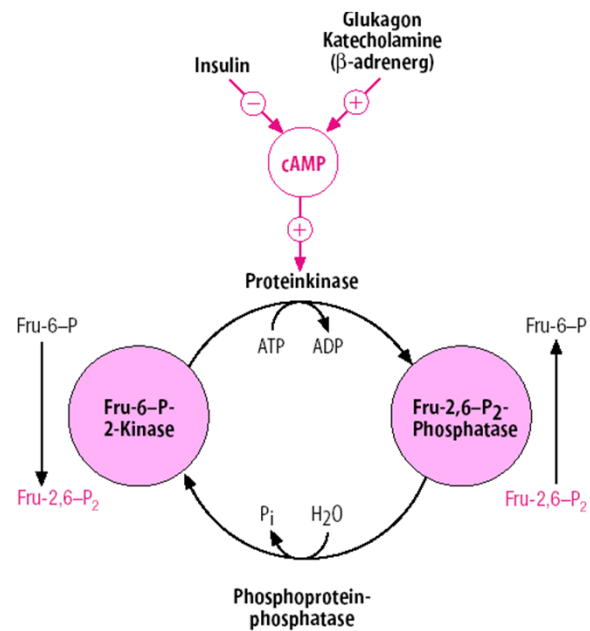
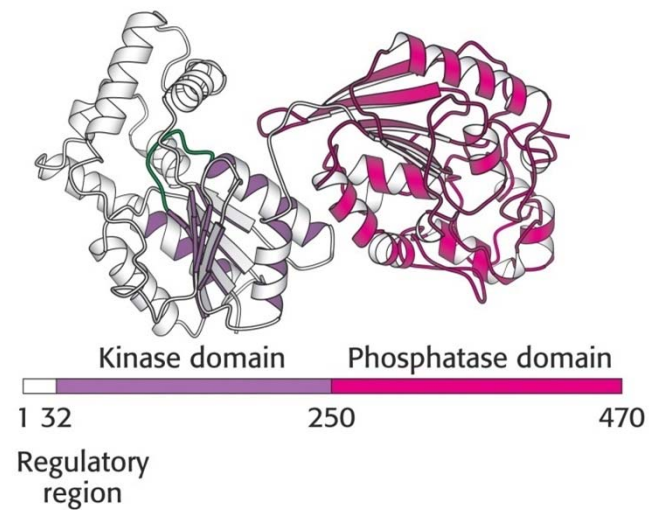


Allosteric modulators of PFK: ATP, AMP, Citrate, H⁺, F-2,6-BP (Activators, Inhibitors)



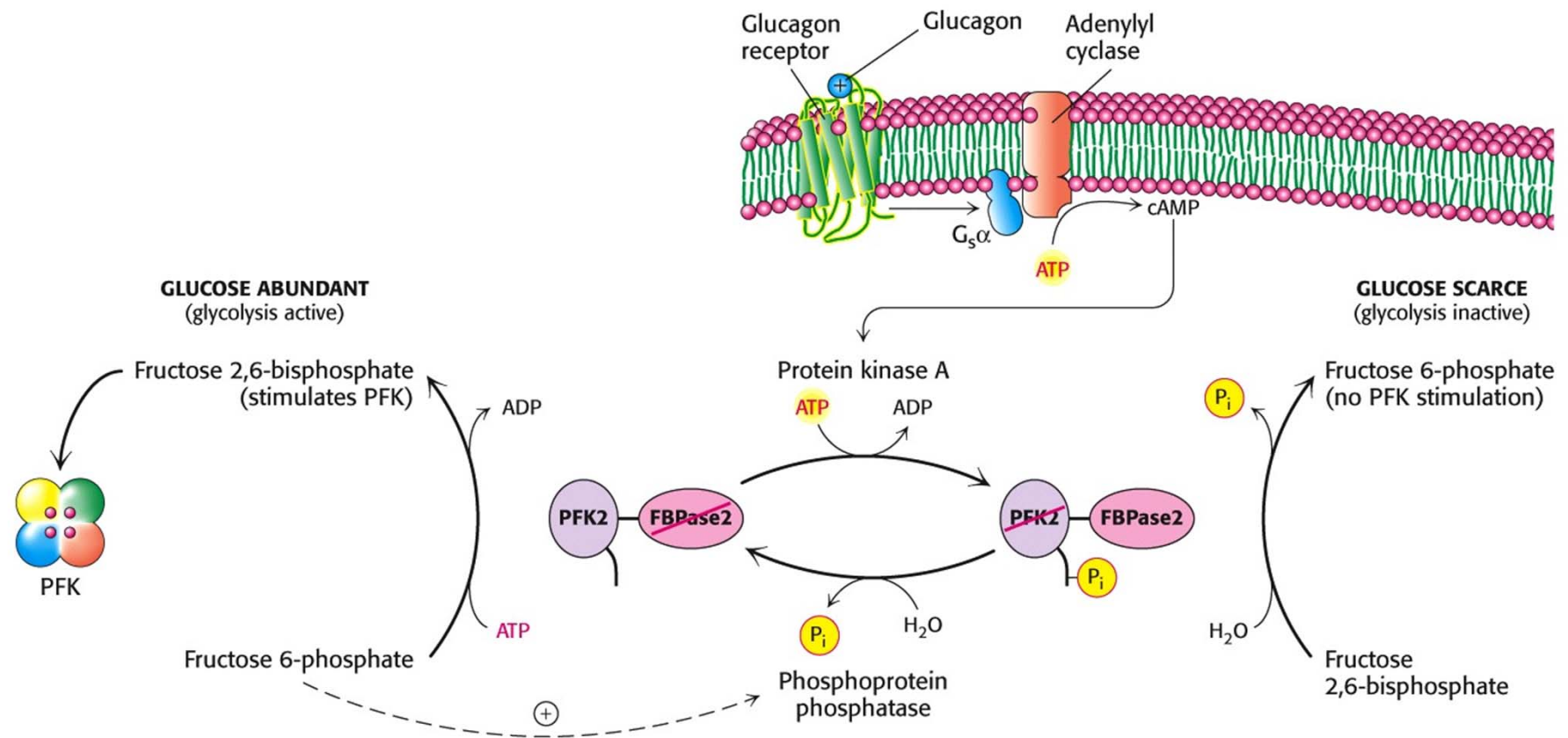
Regulation of **F1,6BPase** is mirror-inverted but opposite !!!

PFK2: a bifunctional (tandem)-enzyme



Hormone regulation of the enzyme is tissue-specific!!!

cAMP-mediated hormonal regulation of PFK via F2,6BP



General mechanism of the activation of effector proteins associated with GPCRs

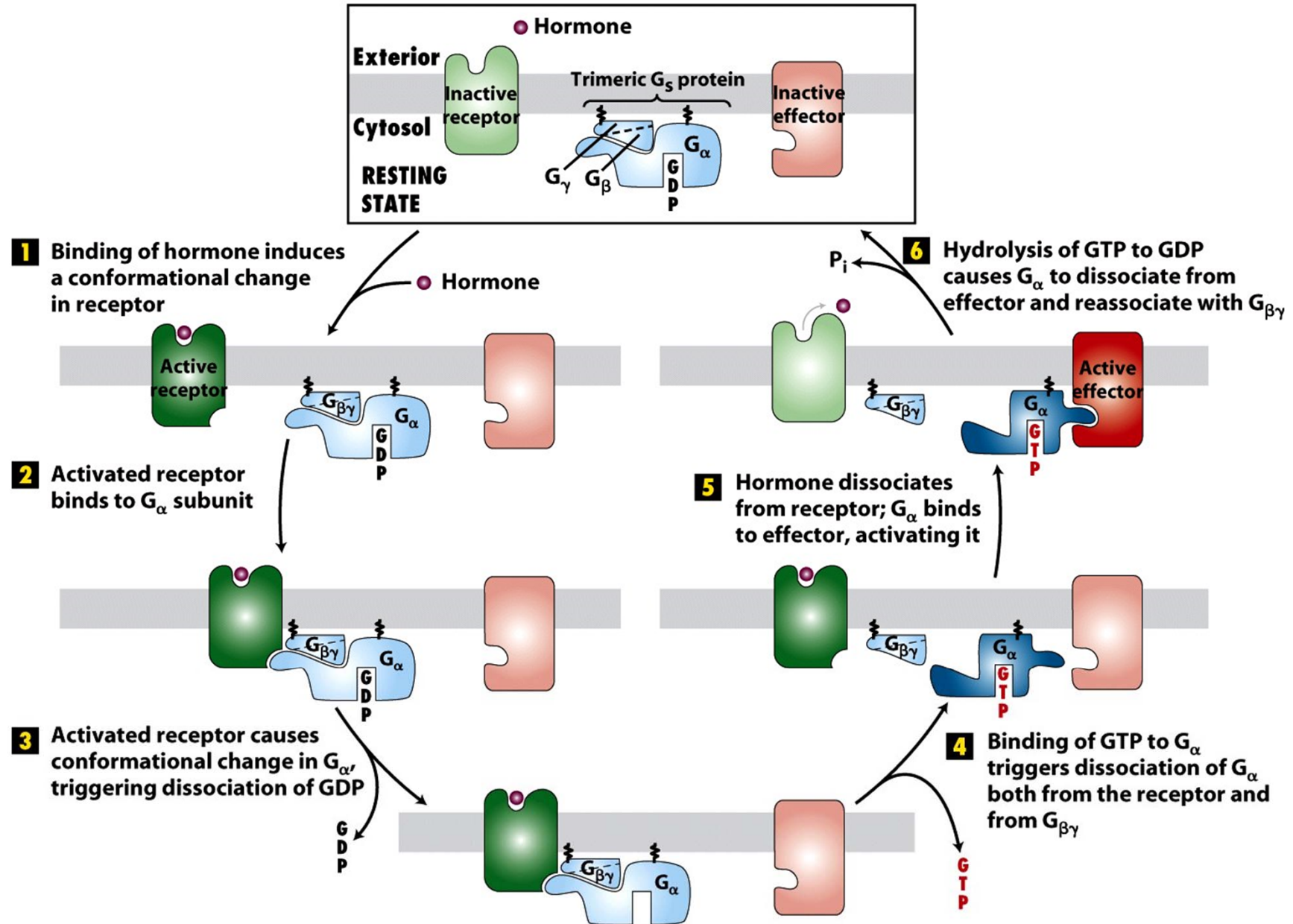


Figure 15-13
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Hormone-induced activation and inhibition of adenylyl cyclase in adipocytes

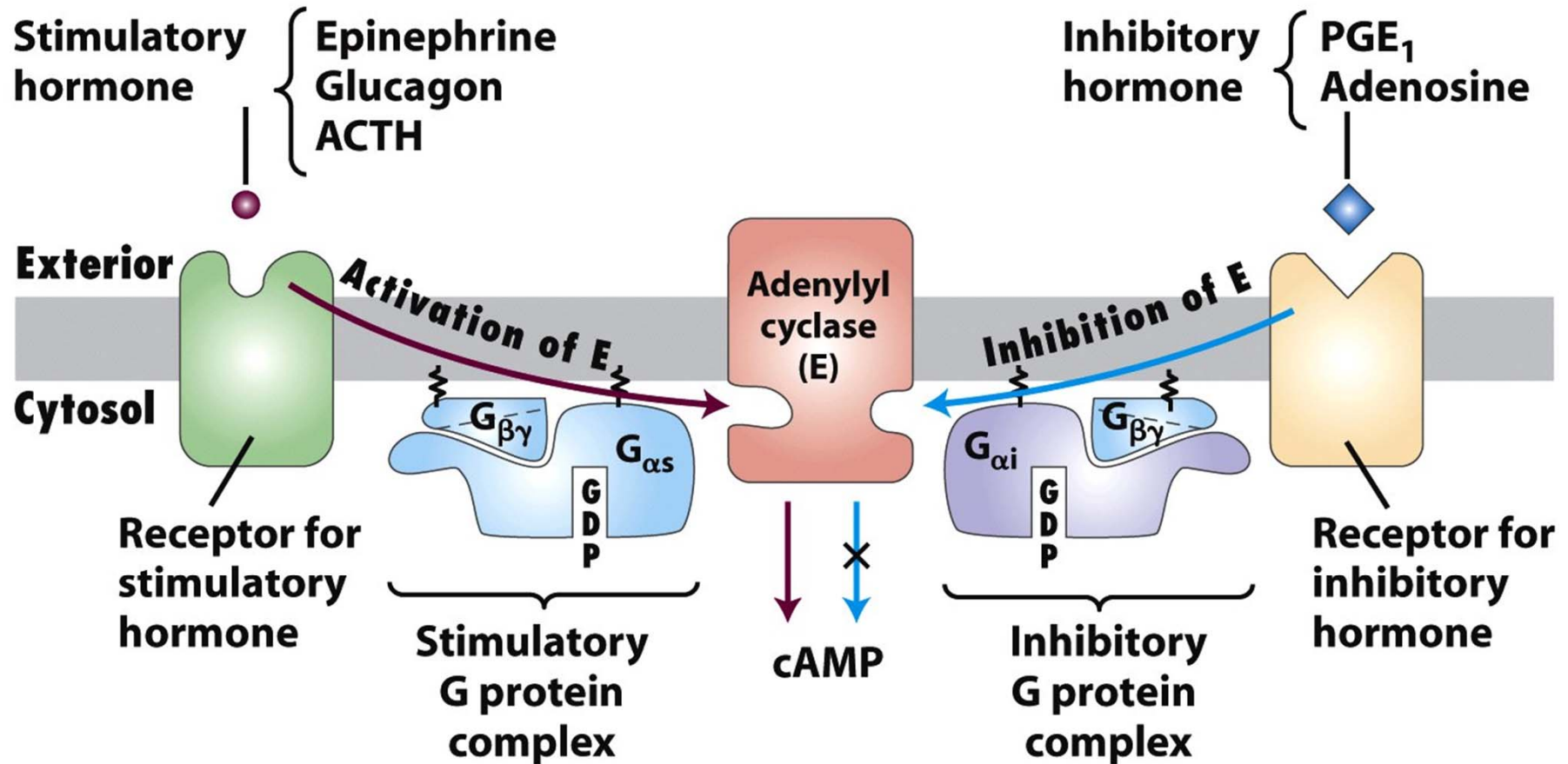


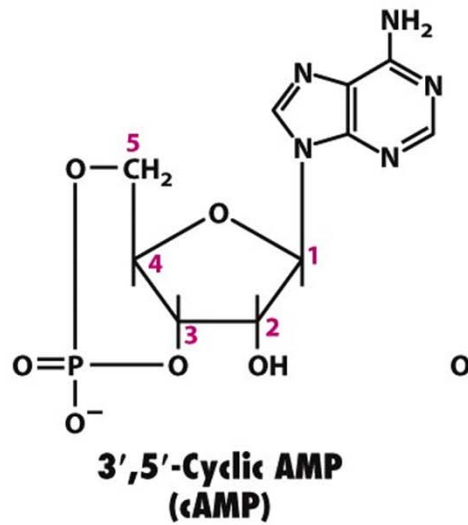
Figure 15-21
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TABLE 15.2 G-protein families and their functions

G_{α} class	Initiating signal	Downstream signal
$G_{\alpha s}$	β -Adrenergic amines, glucagon, parathyroid hormone, many others	Stimulates adenylate cyclase
$G_{\alpha i}$	Acetylcholine, α -adrenergic amines, many neurotransmitters	Inhibits adenylate cyclase
$G_{\alpha t}$	Photons	Stimulates cGMP phosphodiesterase
$G_{\alpha q}$	Acetylcholine, α -adrenergic amines, many neurotransmitters	Increases IP_3 and intracellular calcium
$G_{\alpha 13}$	Thrombin, other agonists	Stimulates Na^+ and H^+ exchange

PLC- β

Source: Z. Farfel, H. R. Bourne, and T. Iiri. *N. Engl. J. Med.* 340(1999):1012.



**Activates protein kinase A
(PKA)**

Adenylyl cyclase generates
the second messenger cAMP
using ATP as a substrate

Figure 15-9
Molecular Cell Biology, Sixth Edition
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Structure and activation of Protein kinase A by cAMP

Protein kinase A (PKA)

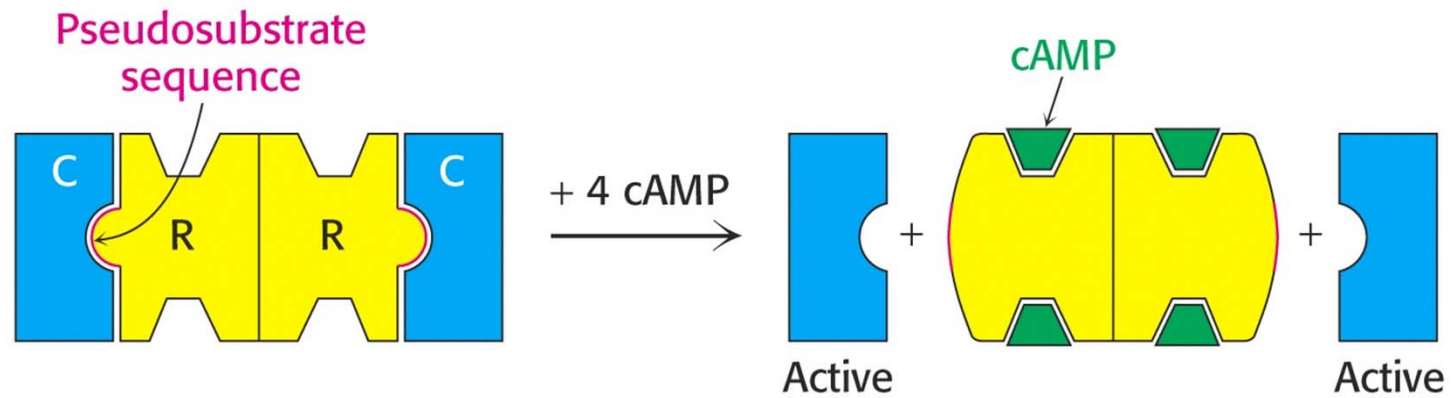


TABLE 15-2 Cellular Responses to Hormone-Induced Rise in cAMP in Various Tissues*

TISSUE	HORMONE INDUCING RISE IN cAMP	CELLULAR RESPONSE
Adipose	Epinephrine; ACTH; glucagon	Increase in hydrolysis of triglyceride; decrease in amino acid uptake
Liver	Epinephrine; norepinephrine; glucagon	Increase in conversion of glycogen to glucose; inhibition of glycogen synthesis; increase in amino acid uptake; increase in gluconeogenesis (synthesis of glucose from amino acids)
Ovarian follicle	FSH; LH	Increase in synthesis of estrogen, progesterone
Adrenal cortex	ACTH	Increase in synthesis of aldosterone, cortisol
Cardiac muscle	Epinephrine	Increase in contraction rate
Thyroid gland	TSH	Secretion of thyroxine
Bone	Parathyroid hormone	Increase in resorption of calcium from bone
Skeletal muscle	Epinephrine	Conversion of glycogen to glucose
Intestine	Epinephrine	Fluid secretion
Kidney	Vasopressin	Resorption of water
Blood platelets	Prostaglandin I	Inhibition of aggregation and secretion

*Nearly all the effects of cAMP are mediated through protein kinase A (PKA), which is activated by binding of cAMP.

SOURCE: E. W. Sutherland, 1972, *Science* **177**:401.

Table 15-2

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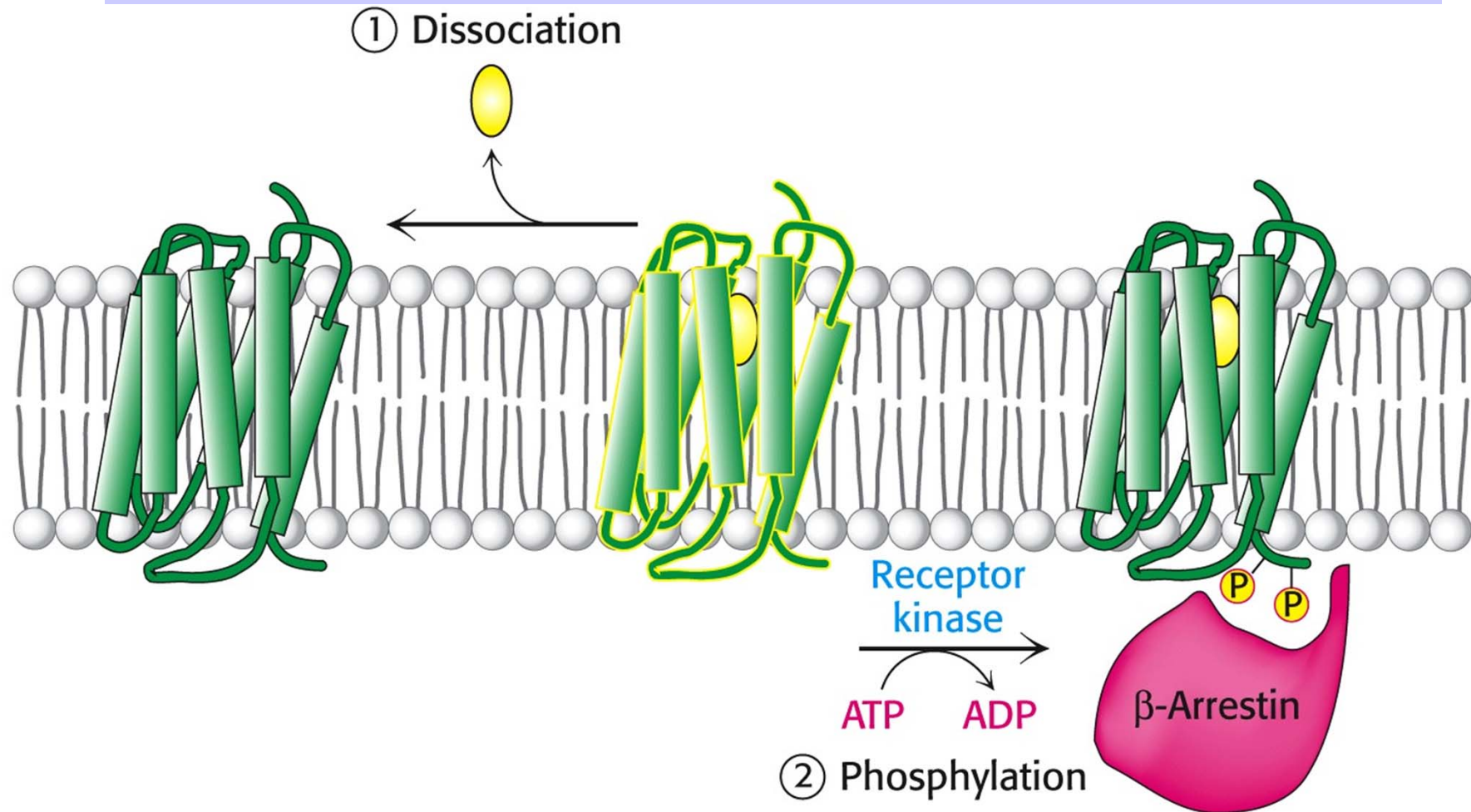
TABLE 12-3 Some Enzymes and Other Proteins Regulated by cAMP-Dependent Phosphorylation (by PKA)

<i>Enzyme/protein</i>	<i>Sequence phosphorylated*</i>	<i>Pathway/process regulated</i>
Glycogen synthase	RASCTSSS	Glycogen synthesis
Phosphorylase <i>b</i> kinase		
α subunit	VEFRRLSI	Glycogen breakdown
β subunit	RTKRSGSV	
Pyruvate kinase (rat liver)	GVLRRASVAZL	Glycolysis
Pyruvate dehydrogenase complex (type L)	GYLRRASV	Pyruvate to acetyl-CoA
Hormone-sensitive lipase	PMRRSV	Triacylglycerol mobilization and fatty acid oxidation
Phosphofructokinase-2/fructose 2,6-bisphosphatase	LQRRRGSSIPQ	Glycolysis/gluconeogenesis
Tyrosine hydroxylase	FIGRRQSL	Synthesis of L-DOPA, dopamine, norepinephrine, and epinephrine
Histone H1	AKRKASGPPVS	DNA condensation
Histone H2B	KKAKASRKESYSVYVK	DNA condensation
Cardiac phospholamban (cardiac pump regulator)	AIRRAST	Intracellular $[Ca^{2+}]$
Protein phosphatase-1 inhibitor-1	IRRRRPTP	Protein dephosphorylation
PKA consensus sequence [†]	XR(R/K)X(S/T)B	Many

*The phosphorylated S or T residue is shown in red. All residues are given as their one-letter abbreviations (see Table 3-1).

[†]X is any amino acid; B is any hydrophobic amino acid.

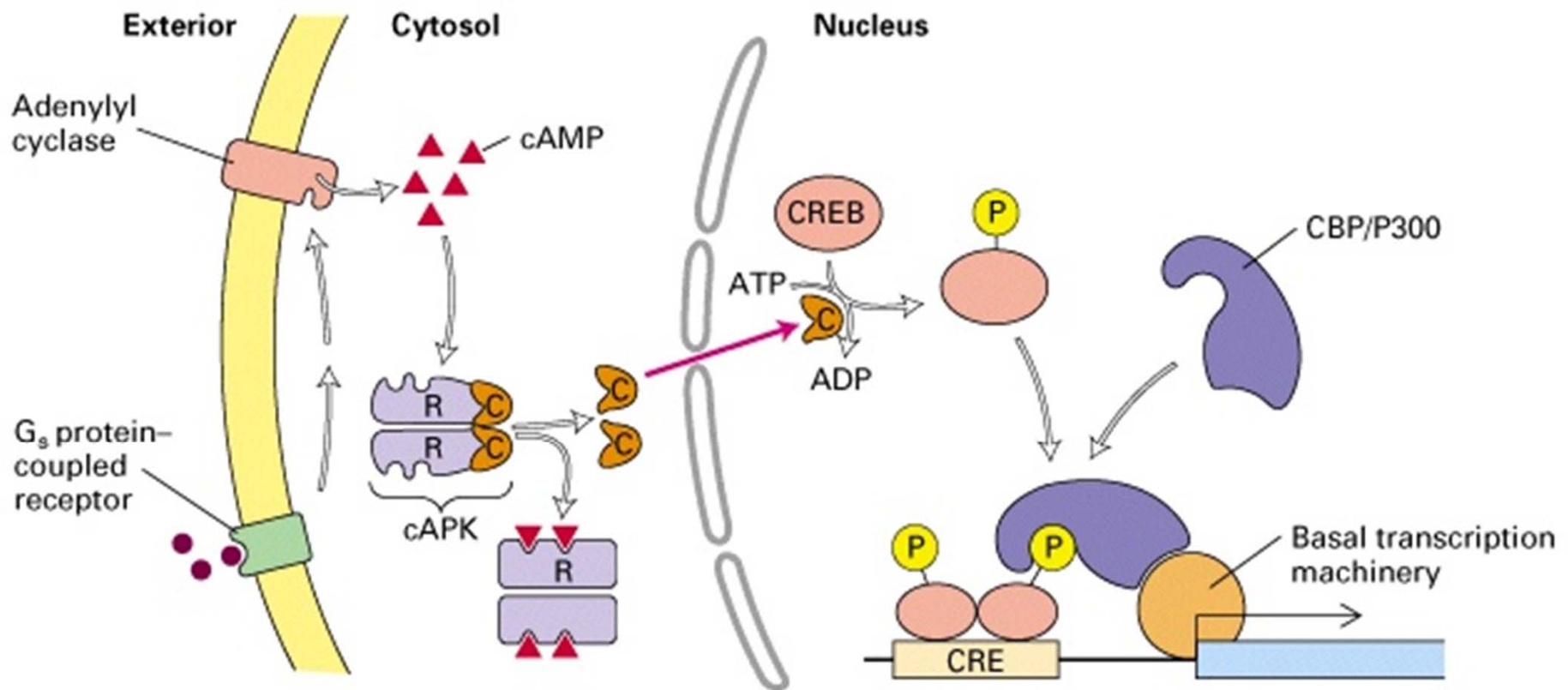
Termination/desensitization of the signal transduction process



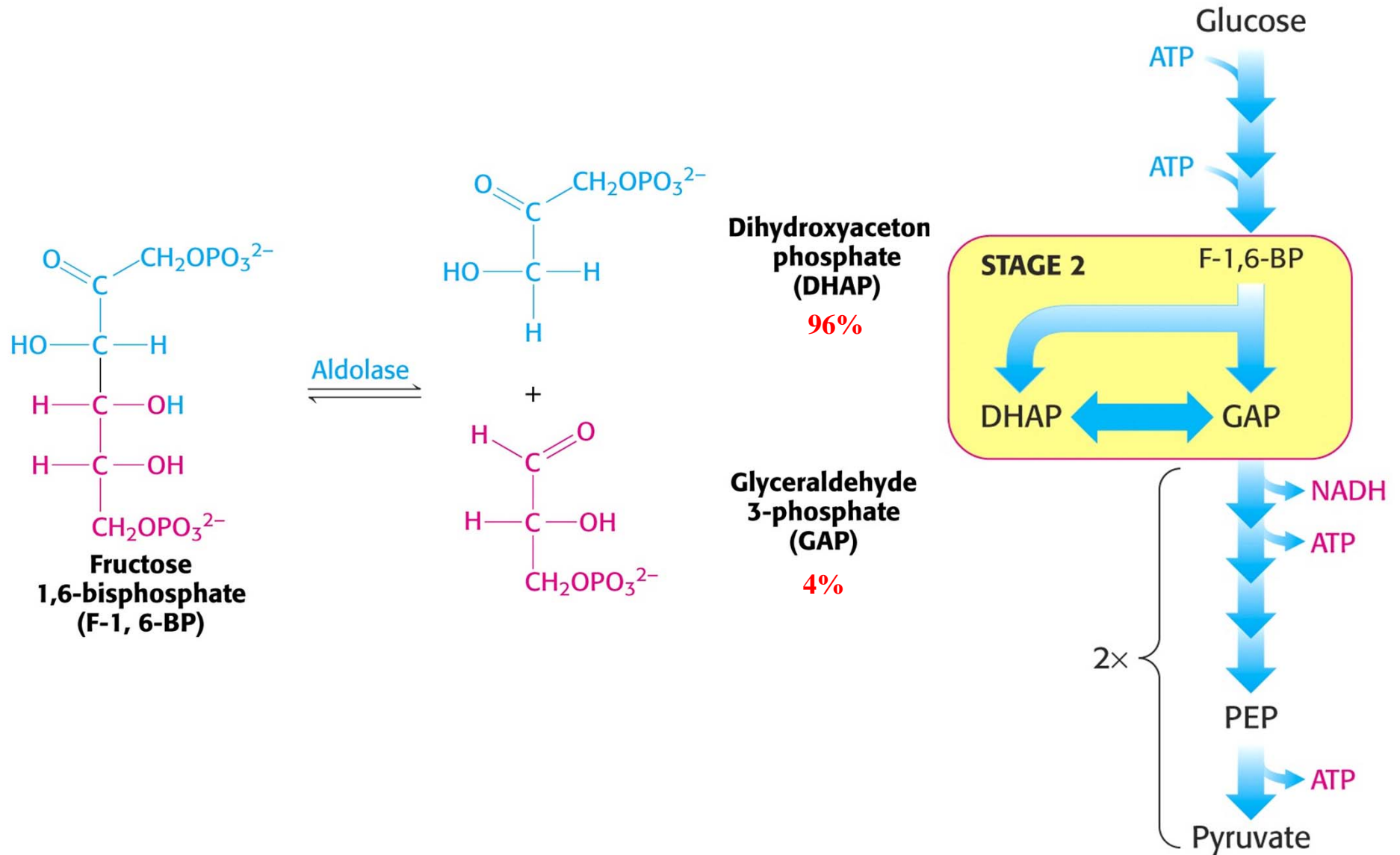
3. Phosphodiesterase (PDE) catalyses hydrolysis of cAMP (calcium-dependent)
4. GTP-hydrolysis

CREB links cAMP signals to transcription

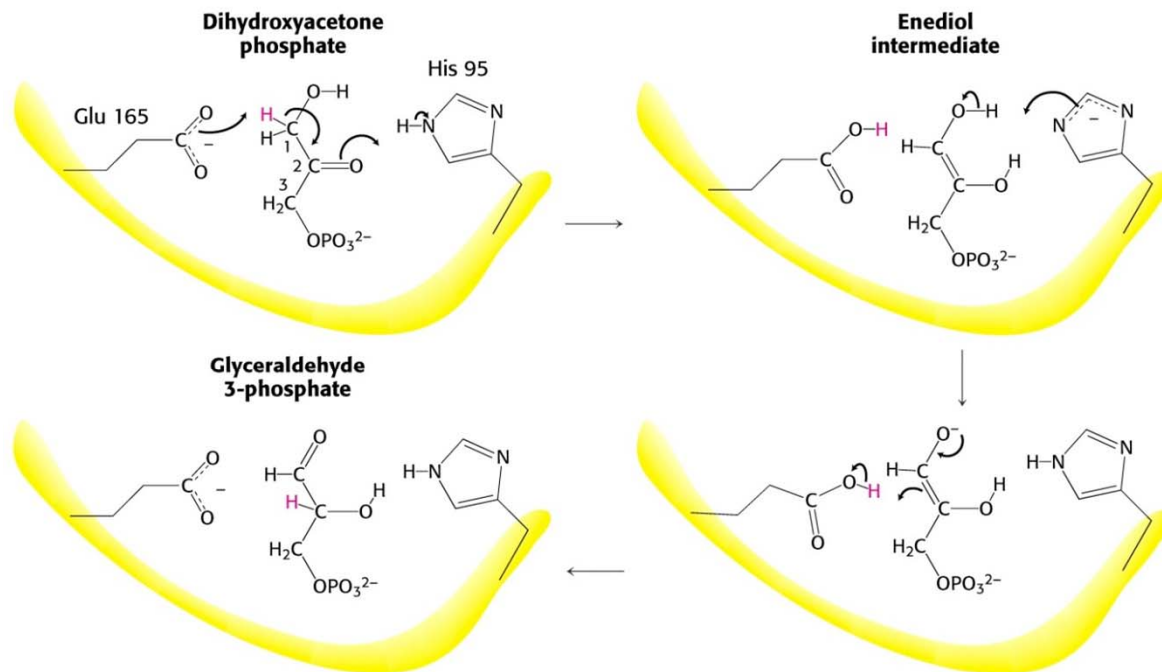
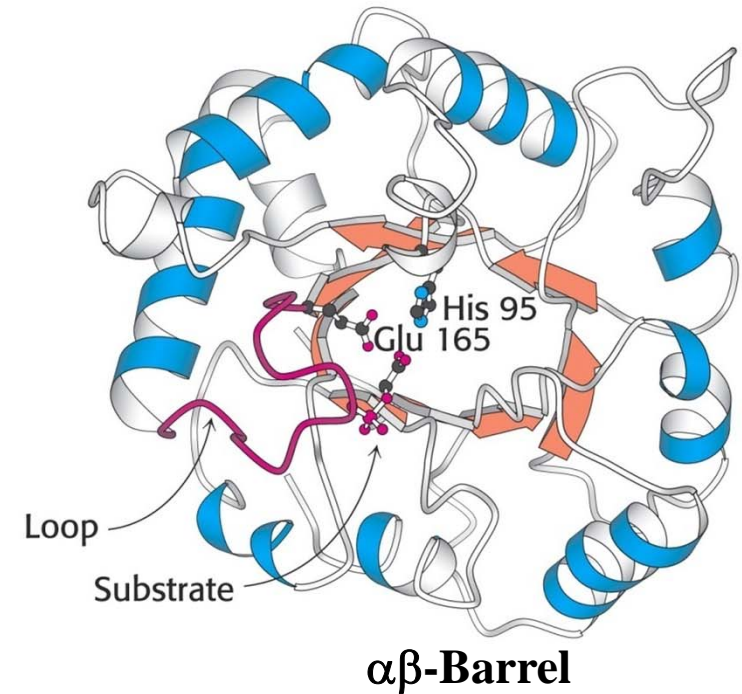
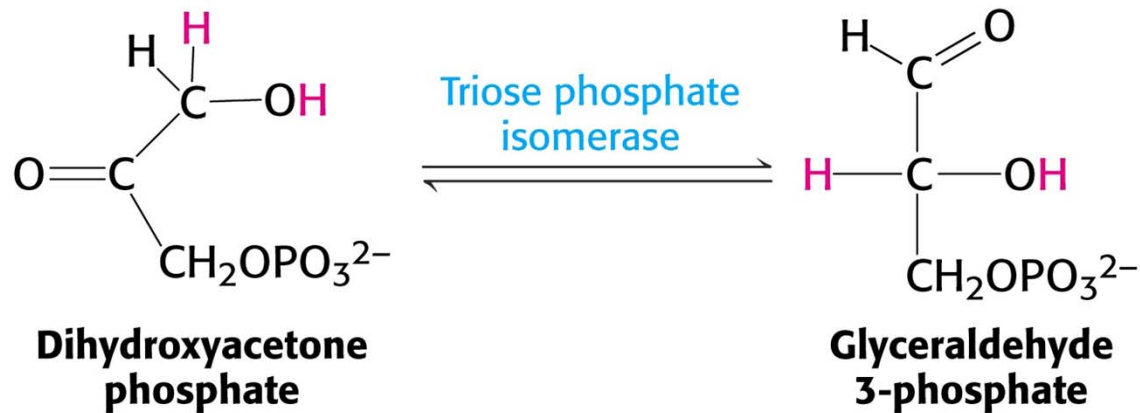
(a) G protein – cAMP pathway



Glycolytic steps: Aldol cleavage catalysed by **aldolase**



Glycolytic steps: Isomerisation catalysed by **triose phosphate isomerase (TIM)**



- catalysis via an endiol-intermediate
- Mech: general acid/base catalysis
- $K = [\text{GAP}]/[\text{DHAP}] = 4,7 \times 10^{-2}$
 $[\text{DHAP}] \gg [\text{GAP}]$
- Enzyme works diffusion controlled
- $k_{\text{cat}}/K_M = 2 \times 10^8 \text{ M}^{-1}\text{s}^{-1}$

Glycolysis

Stage 3

2 X

Glyceraldehyde
3-phosphate
dehydrogenase

P_i, NAD^+

NADH

1,3-Bisphosphoglycerate

Phosphoglycerate
kinase

ADP

ATP

3-Phosphoglycerate

Phosphoglycerate
mutase

2-Phosphoglycerate

Enolase

H_2O

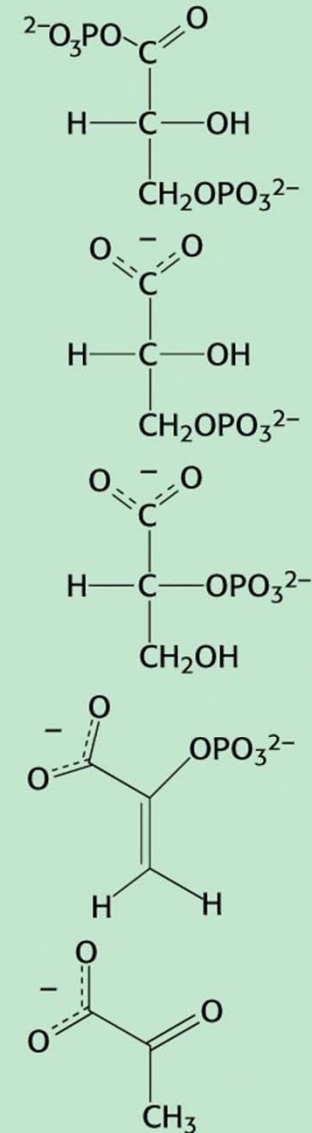
Phosphoenolpyruvate

Pyruvate kinase

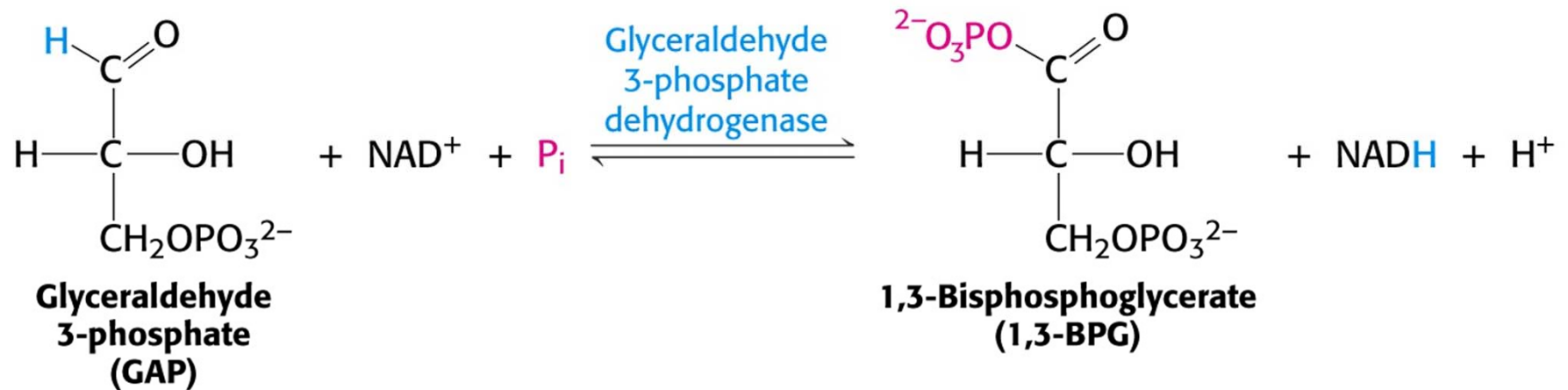
ADP

ATP

Pyruvate



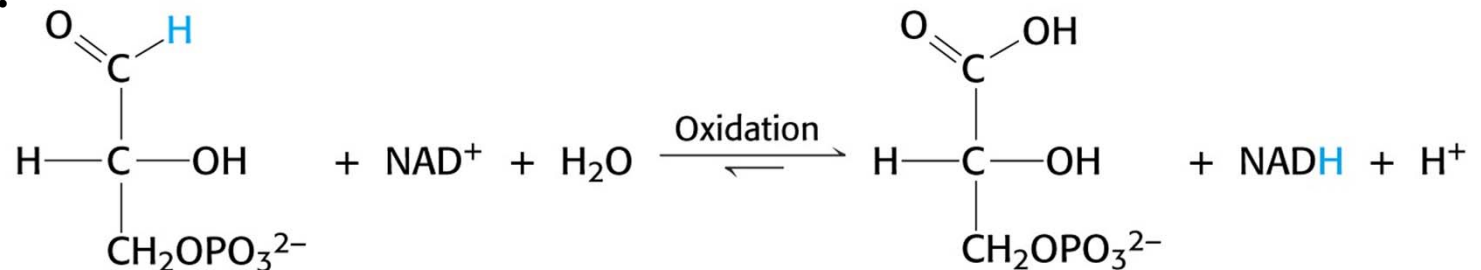
Glycolytic steps: Oxidative phosphorylation catalysed by **GAP-DH**



Two-step-reaction:

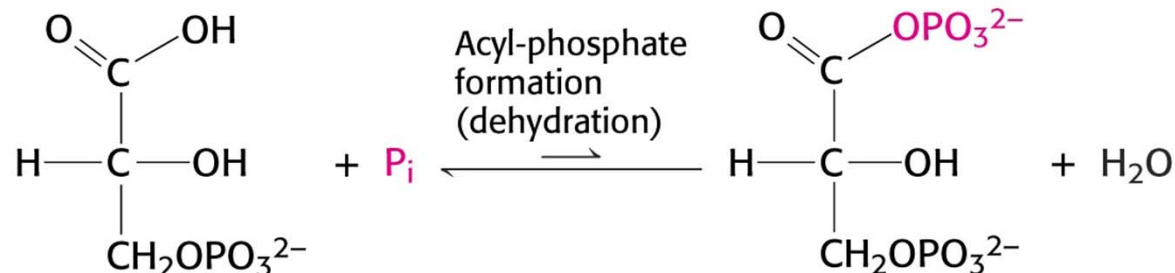
exergonic

$$\Delta G^{0'} = -43,1 \text{ kJ/mol}$$

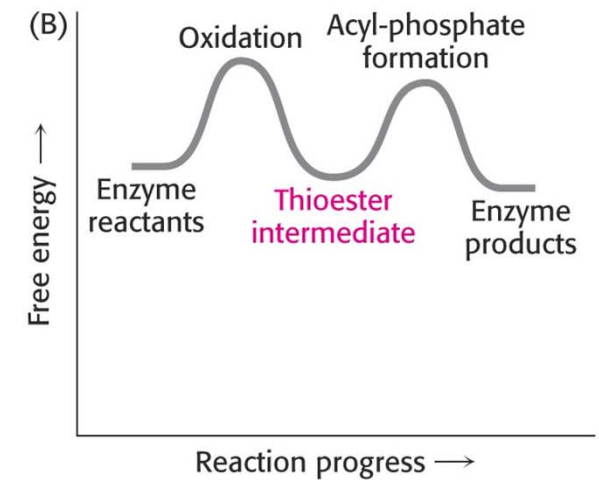
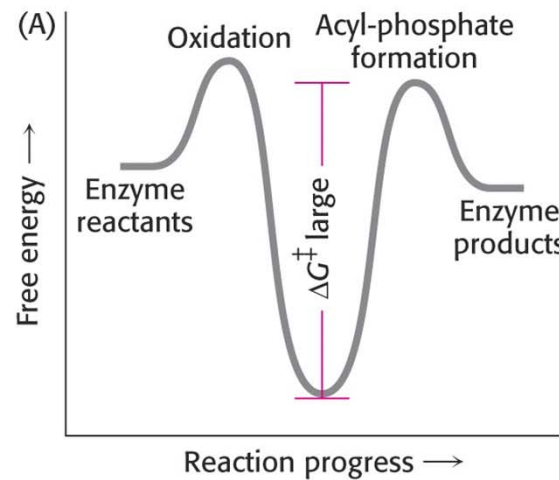
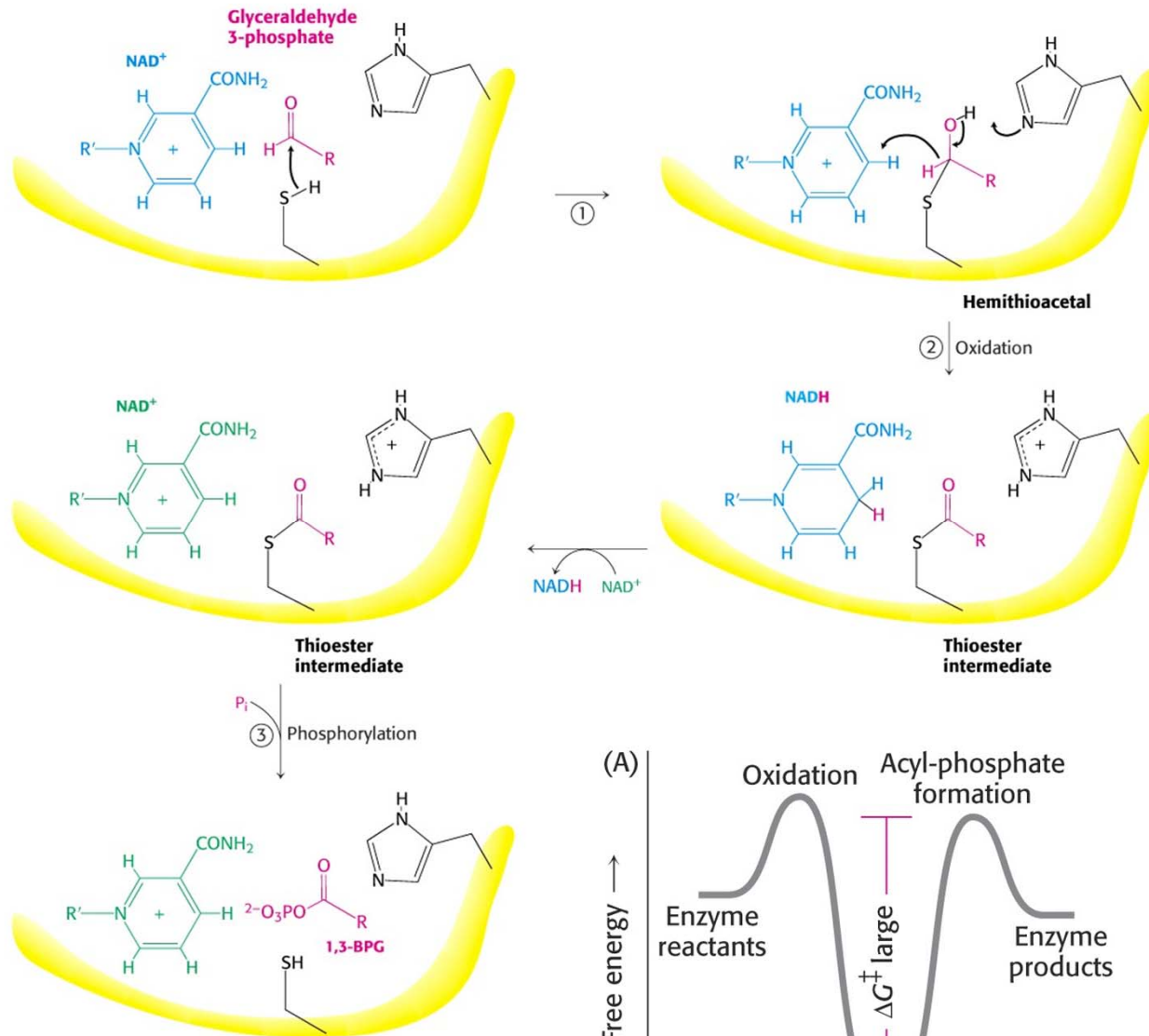


endergonic

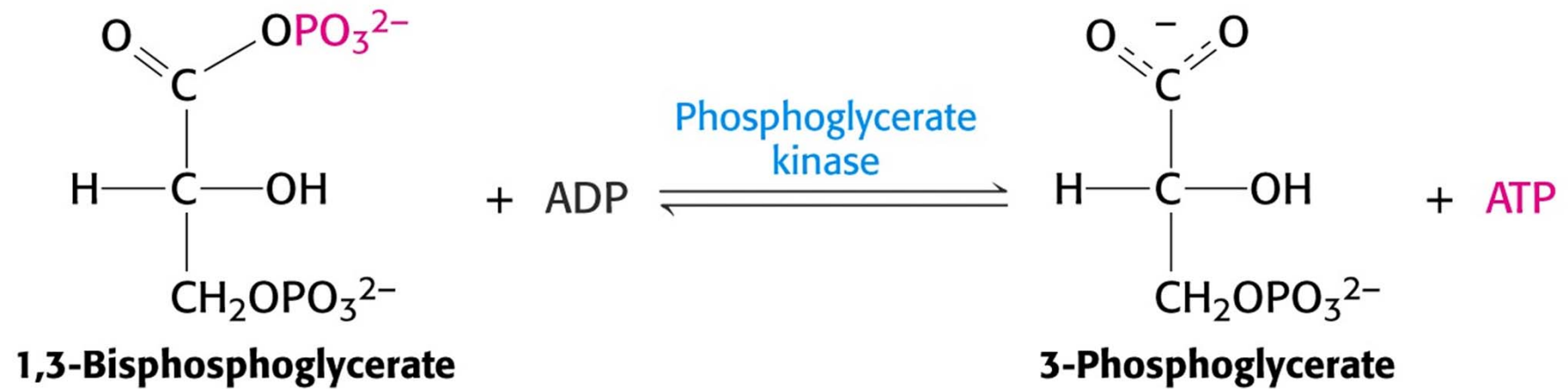
$$\Delta G^{0'} = +49,4 \text{ kJ/mol}$$



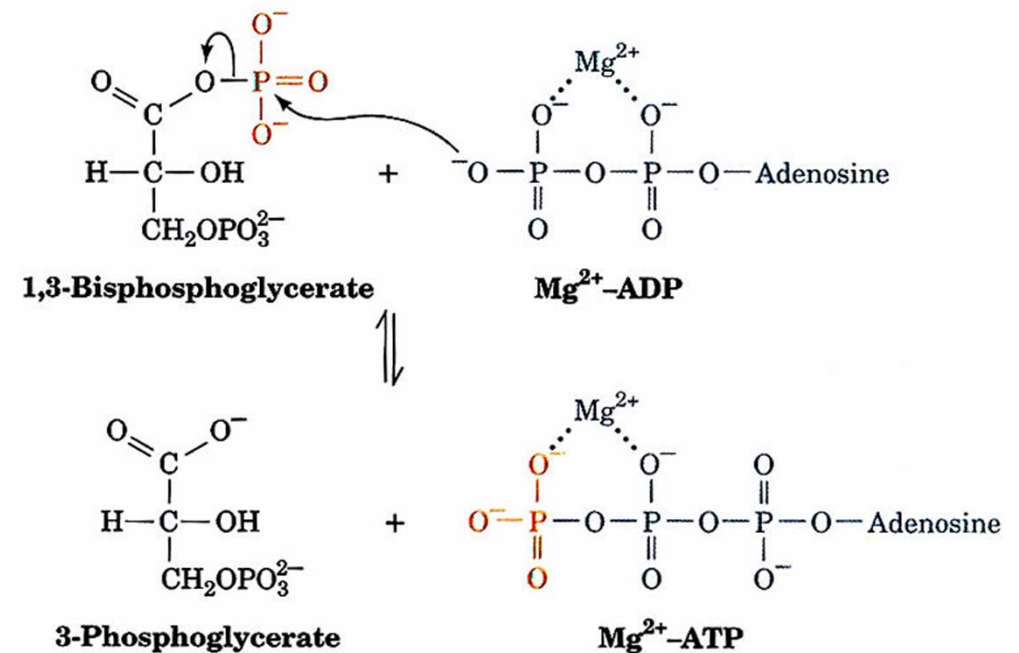
Mechanism of the GAP-DH catalysed reaction



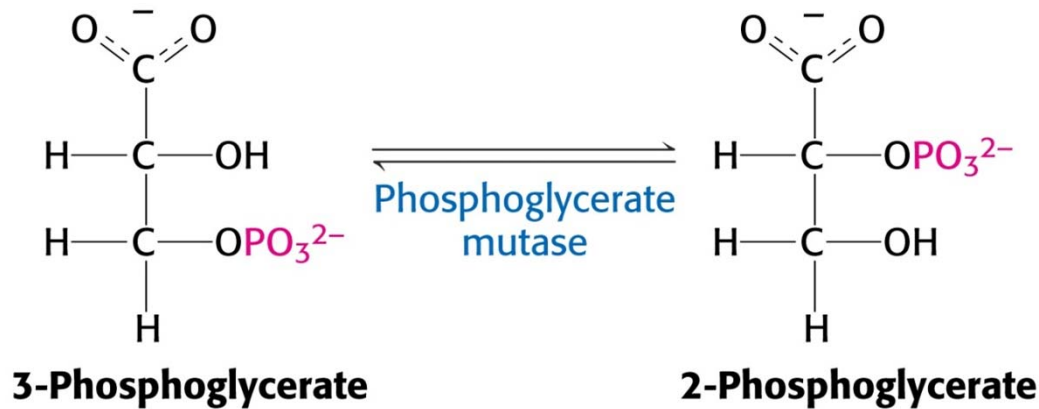
Glycolytic steps: Phosphorylation catalysed by **phosphoglycerate kinase**



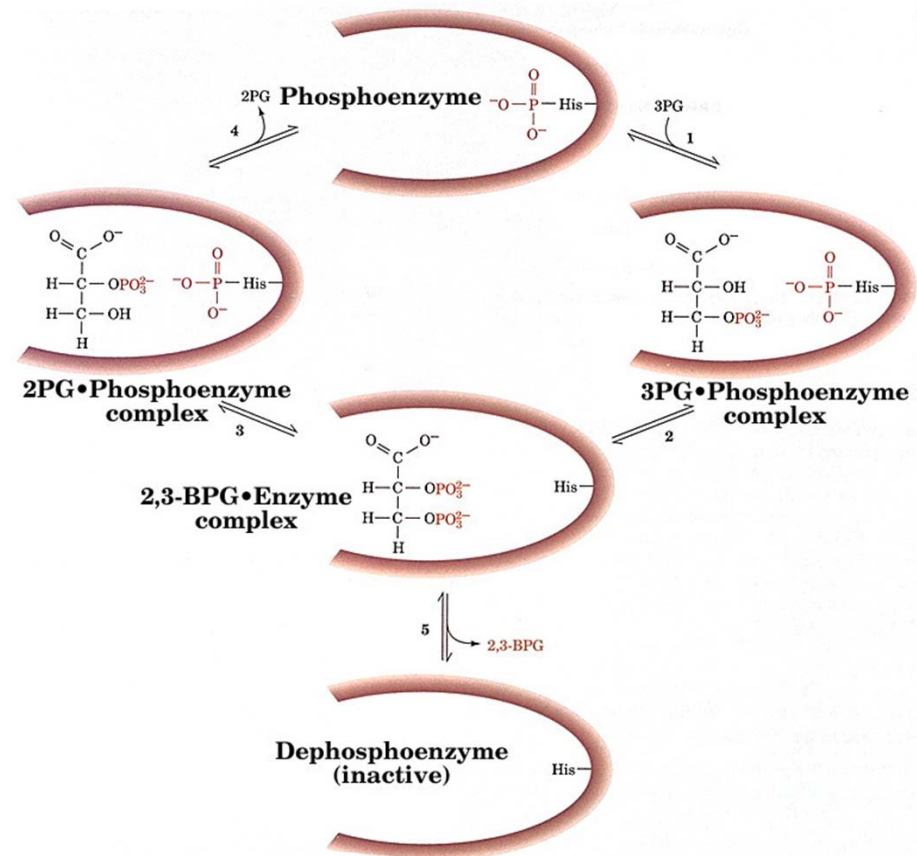
The phosphoglycerate kinase reaction generates the first ATP molecules:



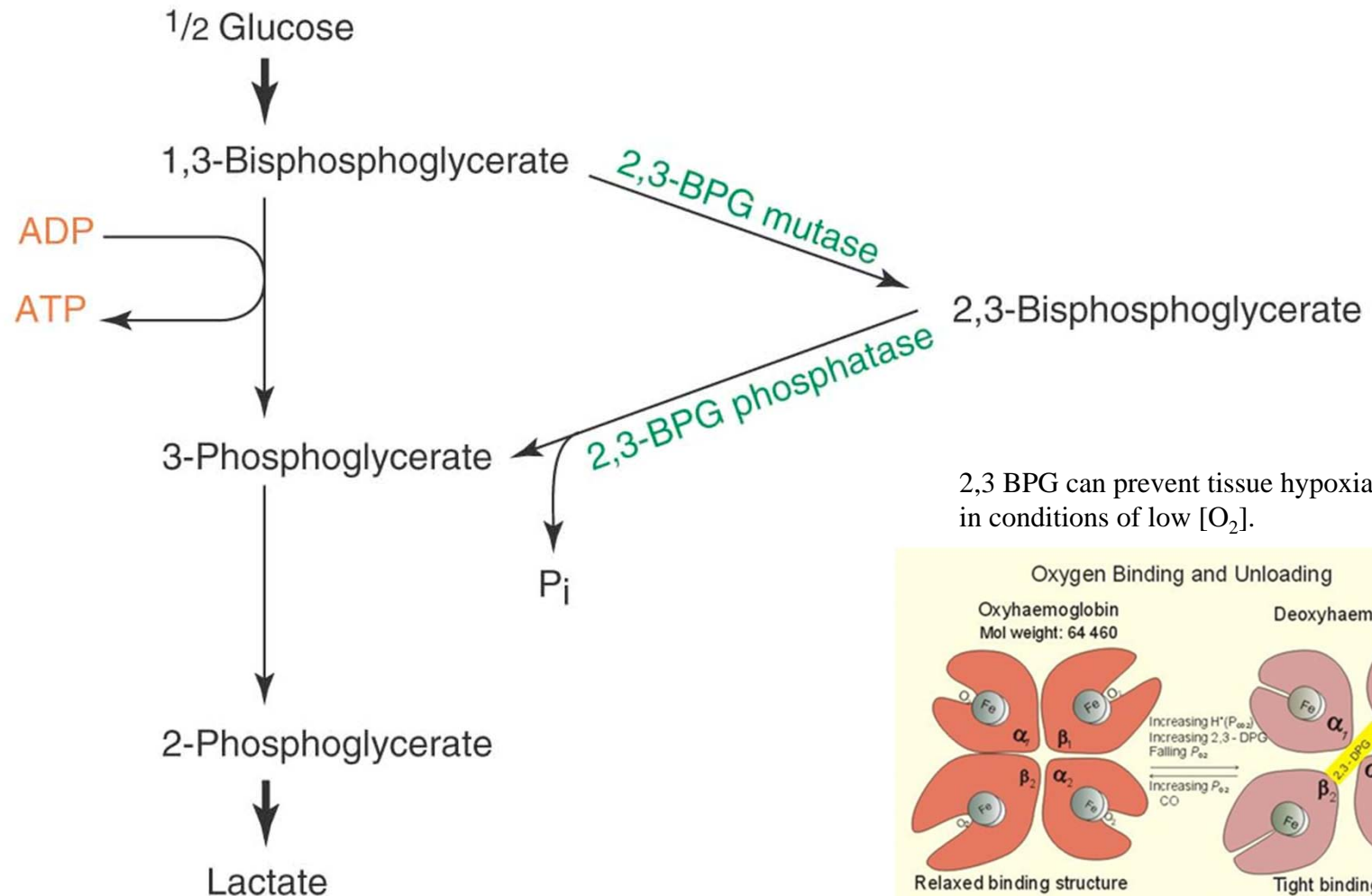
Glycolytic steps: **Phosphoglycerate mutase** catalyzes a phosphoryl group transfer from C3 to C2



- **Rationale** for this reaction in glycolysis: It repositions the phosphate to make PEP (high energy molecule) in the following reaction (enolase)
- Note the **phospho-histidine intermediates** - a bit of **2,3-BPG** (Hb) is required to phosphorylate His (discovered by Zelda Rose)
- **Nomenclature note:** a “mutase” catalyzes migration of a functional group within a substrate

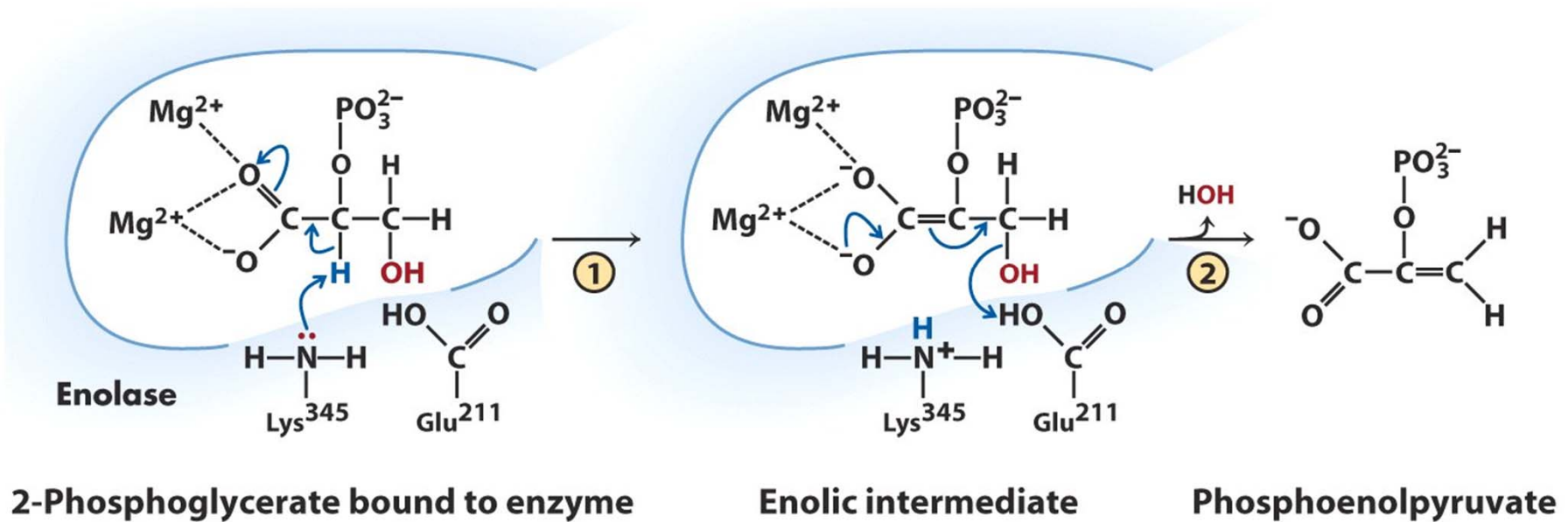
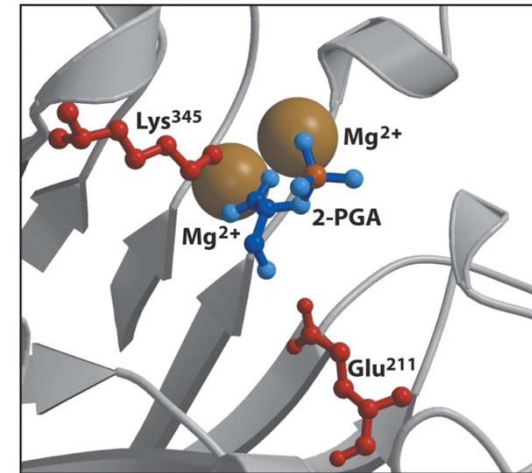
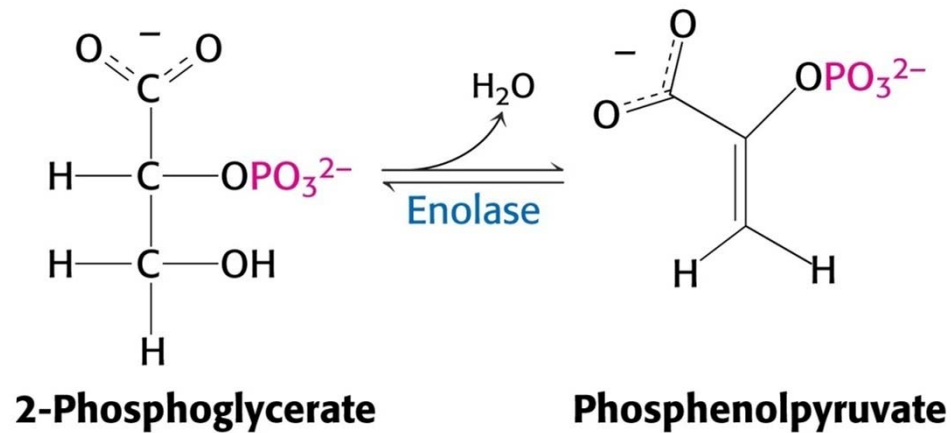


The 2,3-BPG Shunt

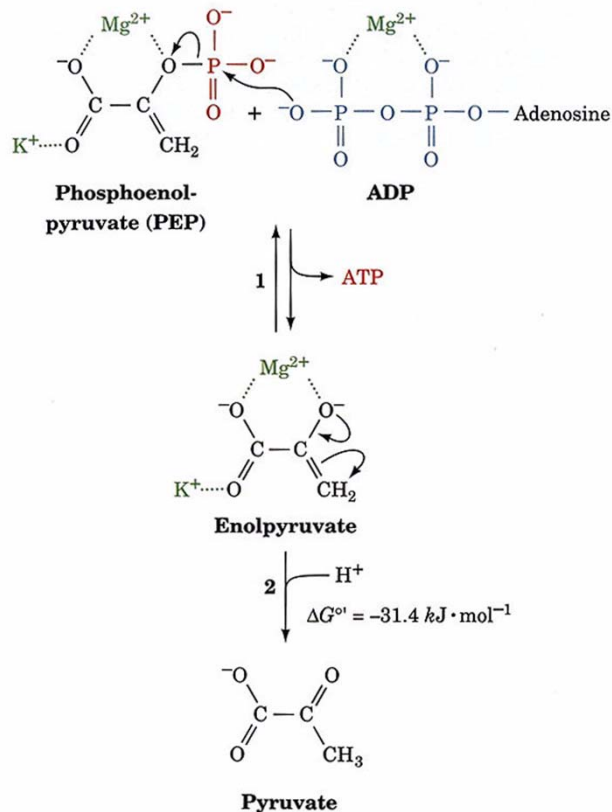
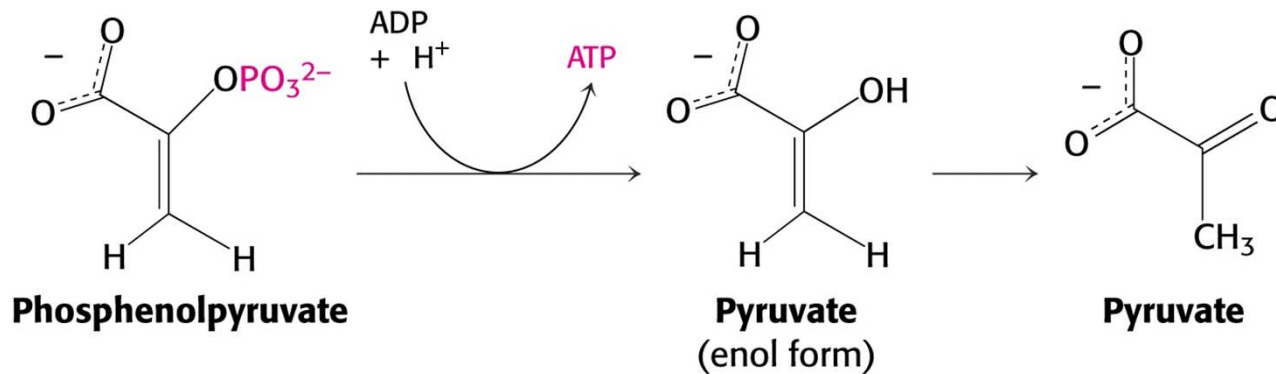


The reactions of 2,3-bisphosphoglycerate (2,3-BPG) shunt are catalyzed by the bifunctional enzyme, 2,3-BPG mutase/phosphatase.

Glycolytic steps: **Dehydration** catalysed by **enolase**



Glycolytic steps: phosphorylation catalysed by **pyruvate kinase**



The pyruvate kinase reaction converts PEP to pyruvate, driving synthesis of ATP. Pyruvate kinase generates 2 ATP (debt was paid in the phosphoglycerate kinase reaction so both ATPs are a gain of energy).

- enzyme requires **Mg²⁺** or **Mn²⁺**
- in mammals allosterically regulated:
 - Activators: F-1,6-BP, PEP
 - Inhibitors: ATP, Citrate, Ala