

Glucose Transport, Glycolysis and Hexose Metabolism

CHEM 420 – Principles of Biochemistry
Instructor – Anthony S. Serianni

Chapter 17: Voet/Voet, *Biochemistry*, 2011
Fall 2015

November 6 & 9

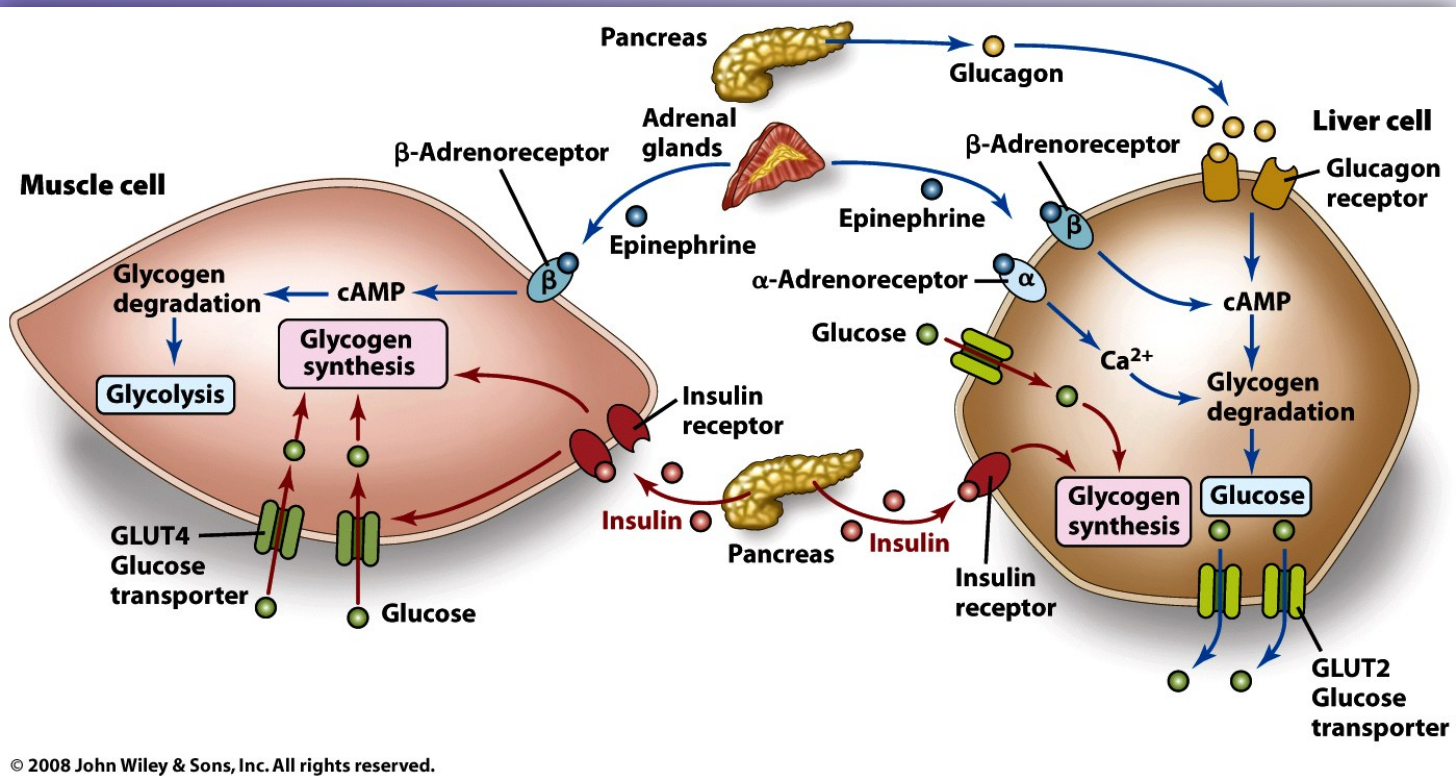
TABLE 12-1 Properties of Selected Members of Human Glucose Transporters (GLUT)

Transporters	Major Tissue Distribution	Properties
GLUT 1	Brain, microvessels, red blood cells, placenta, kidney, and many other cells	Low K_m (about 1 mM), ubiquitous basal transporter
GLUT 2	Liver, pancreatic β -cell, small intestine	High K_m (15–20 mM)
GLUT 3	Brain, placenta, fetal muscle	Low K_m , provide glucose for tissue cells metabolically dependent on glucose
GLUT 4	Skeletal and heart muscle, fat tissue (adipocytes)	K_m (5 mM), insulin responsive transporter
GLUT 5	Small intestine, testes	Exhibits high affinity for fructose
SGLT 1	Small intestine and renal tubules	Low K_m (0.1–1.0 mM)
SGLT 2	Renal tubules	Low K_m (1.6 mM)

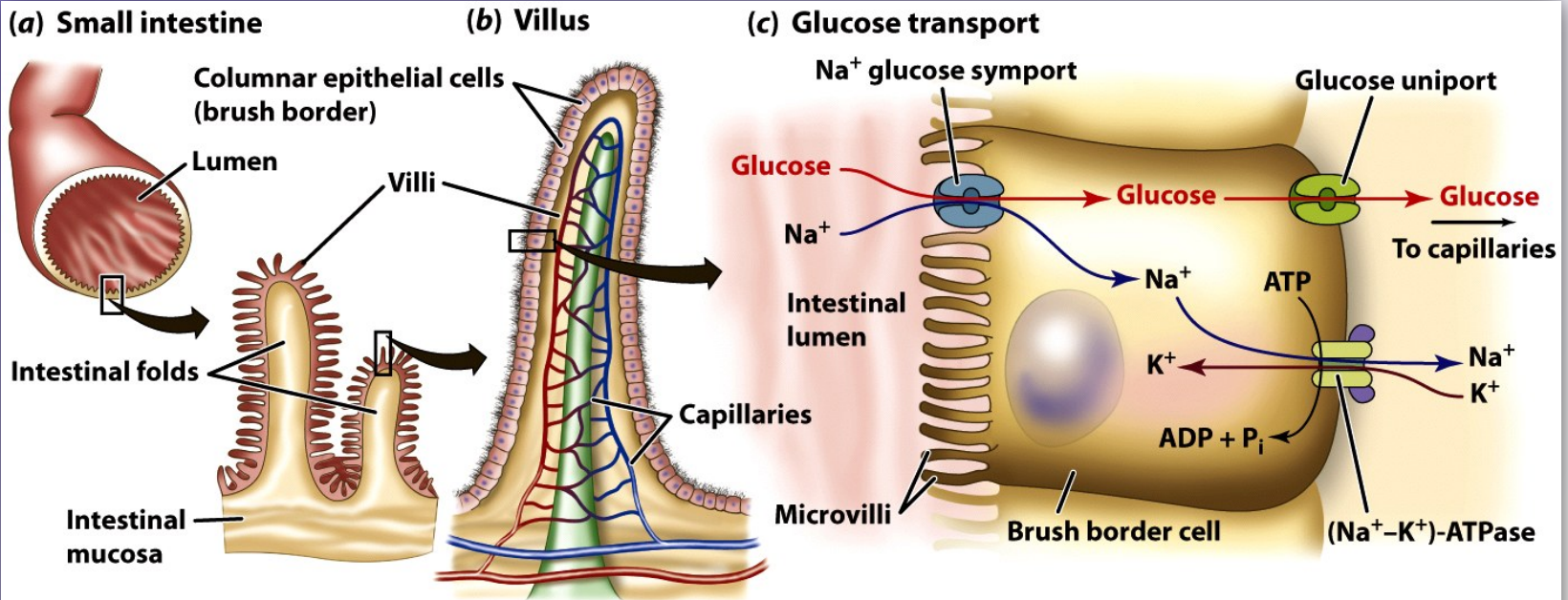
GLUT: concentration gradient-dependent facilitated transport with specific carrier; either insulin dependent or insulin independent

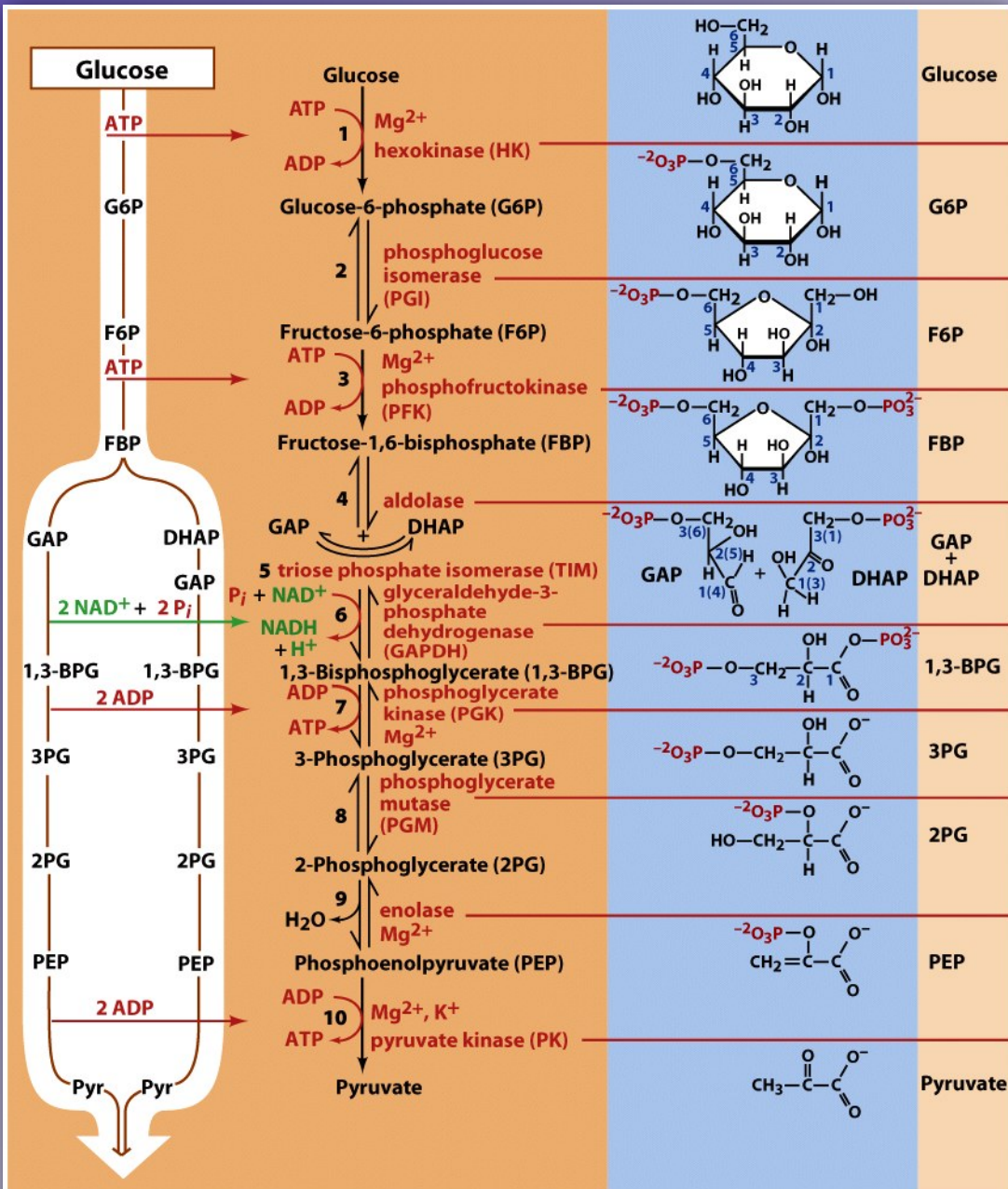
SGLT: active transport

Plasma membrane-bound glucose transporters



Active transport of glucose driven by an ion gradient





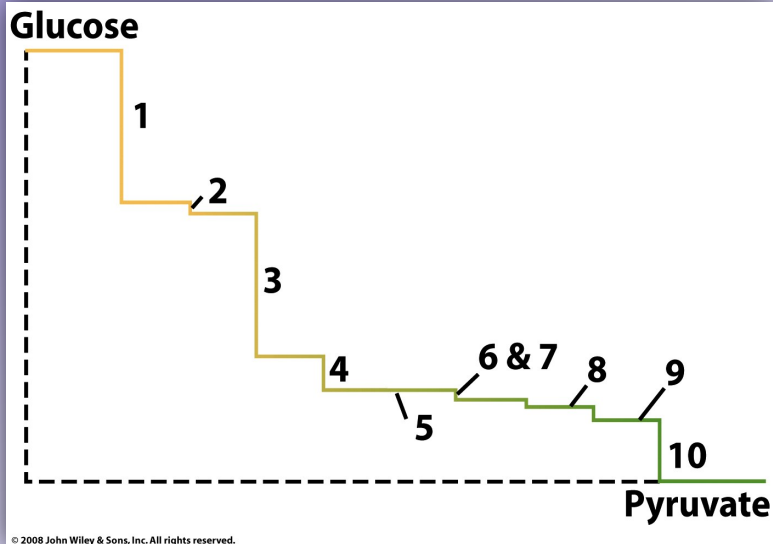
Overview of the ten reactions of glycolysis

Table 15-1 **$\Delta G^{\circ'}$ and ΔG for the Reactions of Glycolysis in Heart Muscle^a**

Reaction	Enzyme	$\Delta G^{\circ'}$ (kJ · mol ⁻¹)	ΔG (kJ · mol ⁻¹)
1	Hexokinase	-20.9	-27.2
2	PGI	+2.2	-1.4
3	PFK	-17.2	-25.9
4	Aldolase	+22.8	-5.9
5	TIM	+7.9	~0
6 + 7	GAPDH + PGK	-16.7	-1.1
8	PGM	+4.7	-0.6
9	Enolase	-3.2	-2.4
10	PK	-23.0	-13.9

^aCalculated from data in Newsholme, E.A. and Start, C., *Regulation in Metabolism*, p. 97, Wiley (1973).

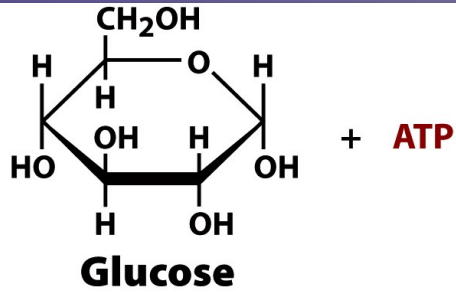
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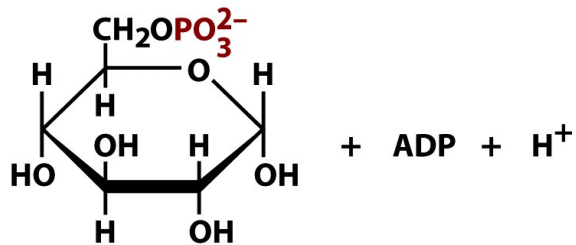
Free energy changes
in glycolysis

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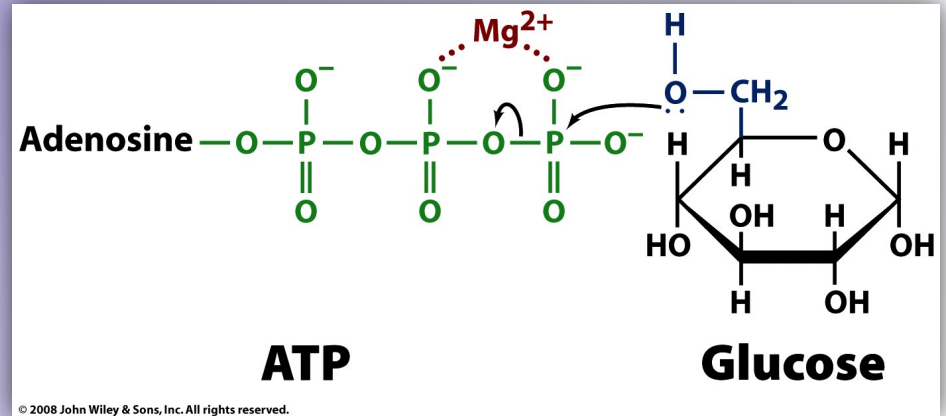
Hexokinase (muscle)



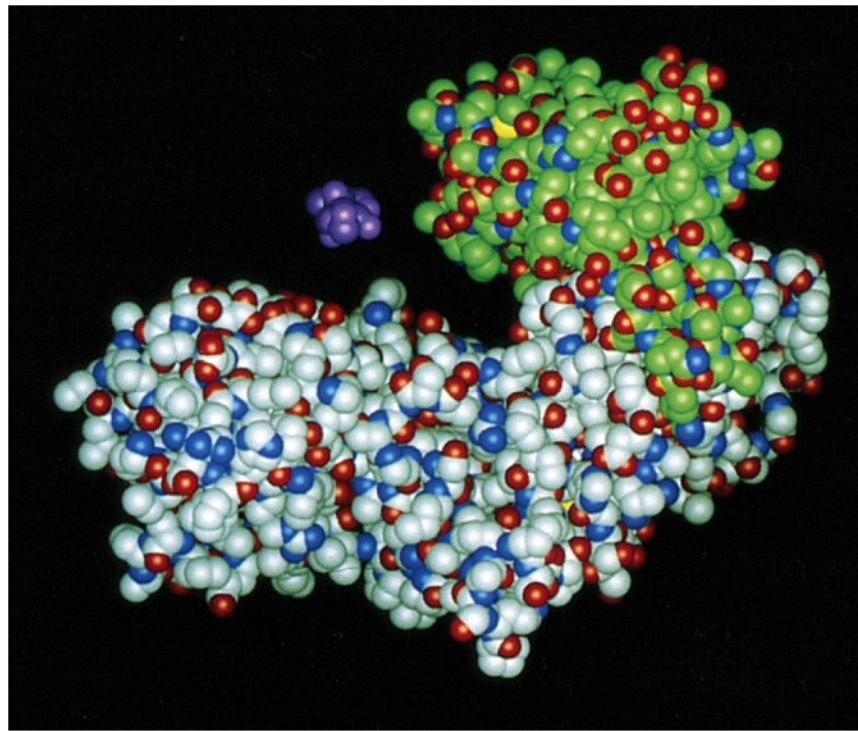
hexokinase
Mg²⁺



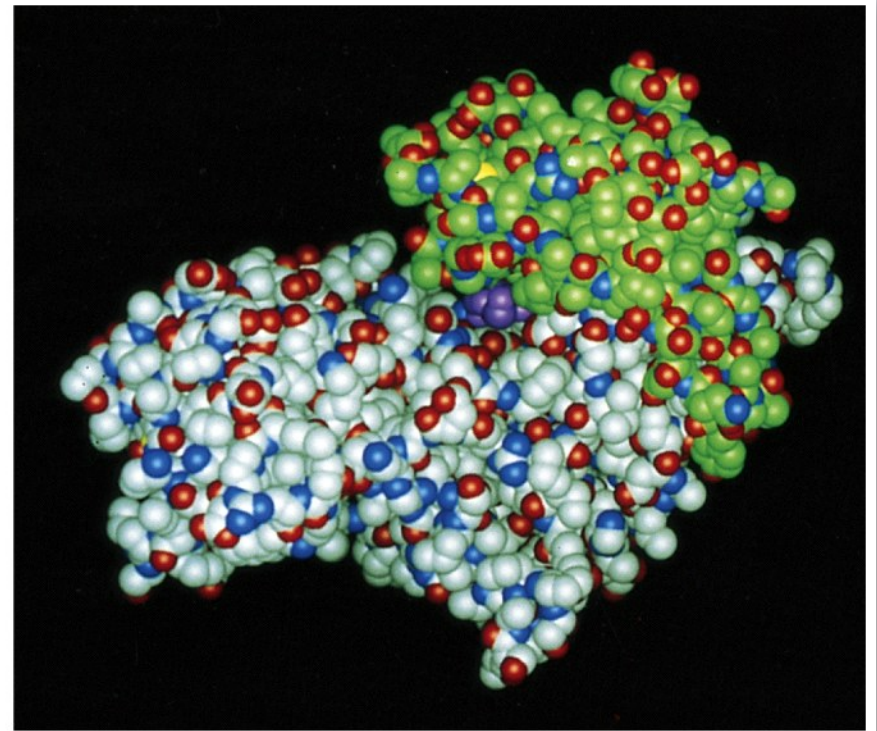
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Conformational change in hexokinase upon substrate binding

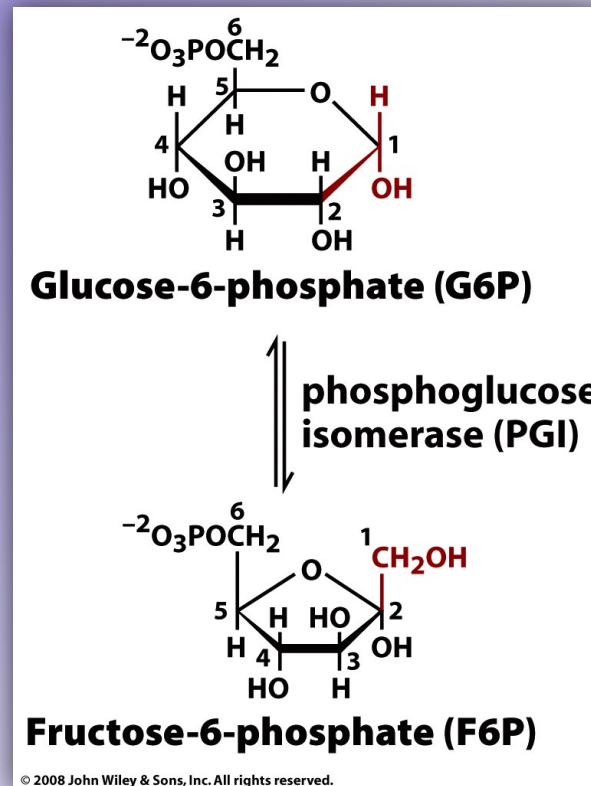


(a)

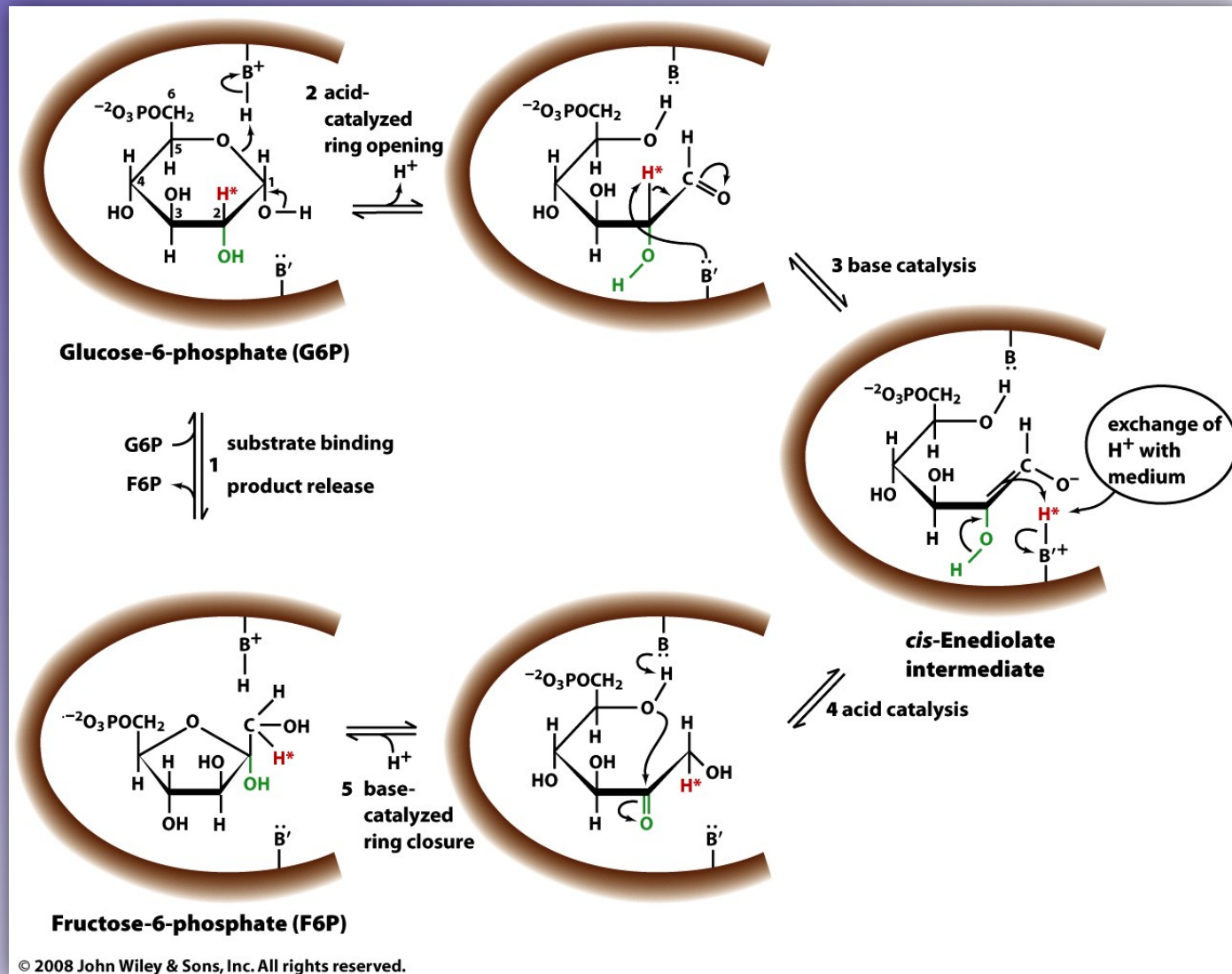


(b)

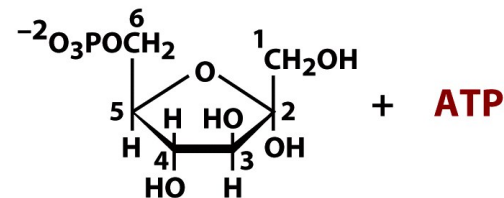
Phosphoglucoisomerase (PGI)



PGI mechanism

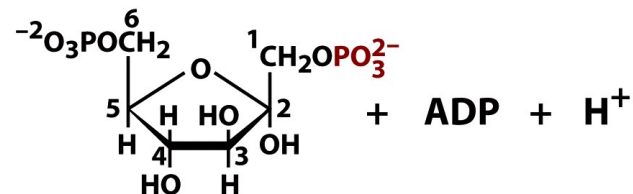


Phosphofructokinase (PFK)



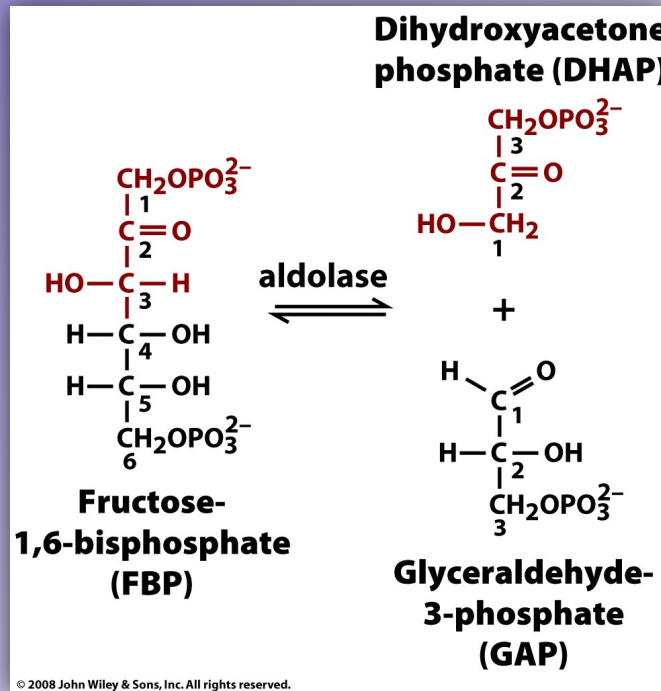
**Fructose-6-phosphate
(F6P)**

phosphofructokinase (PFK)
 Mg^{2+}

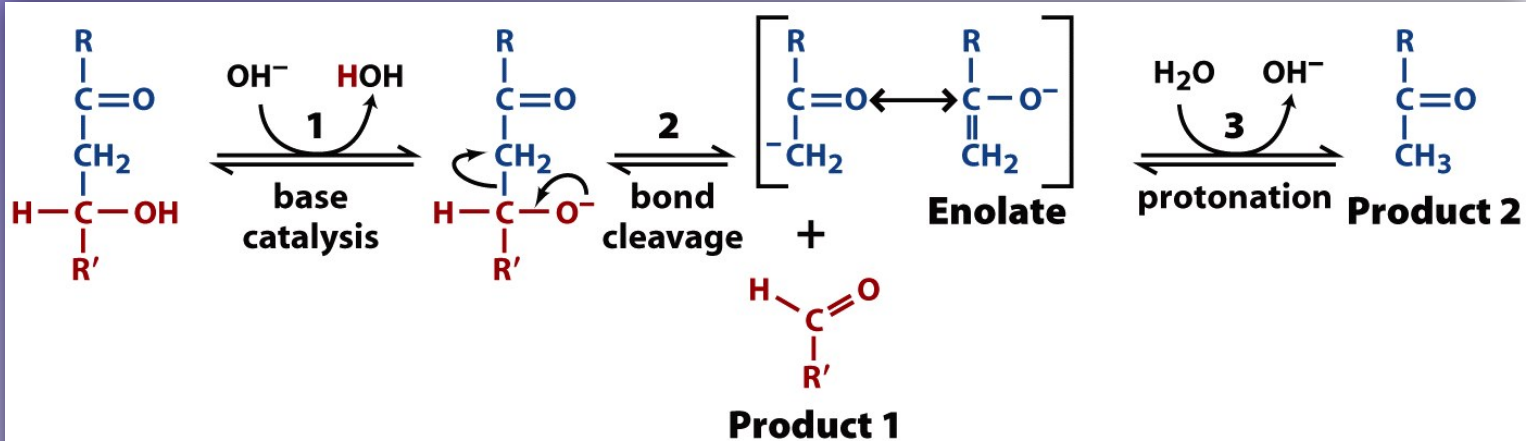


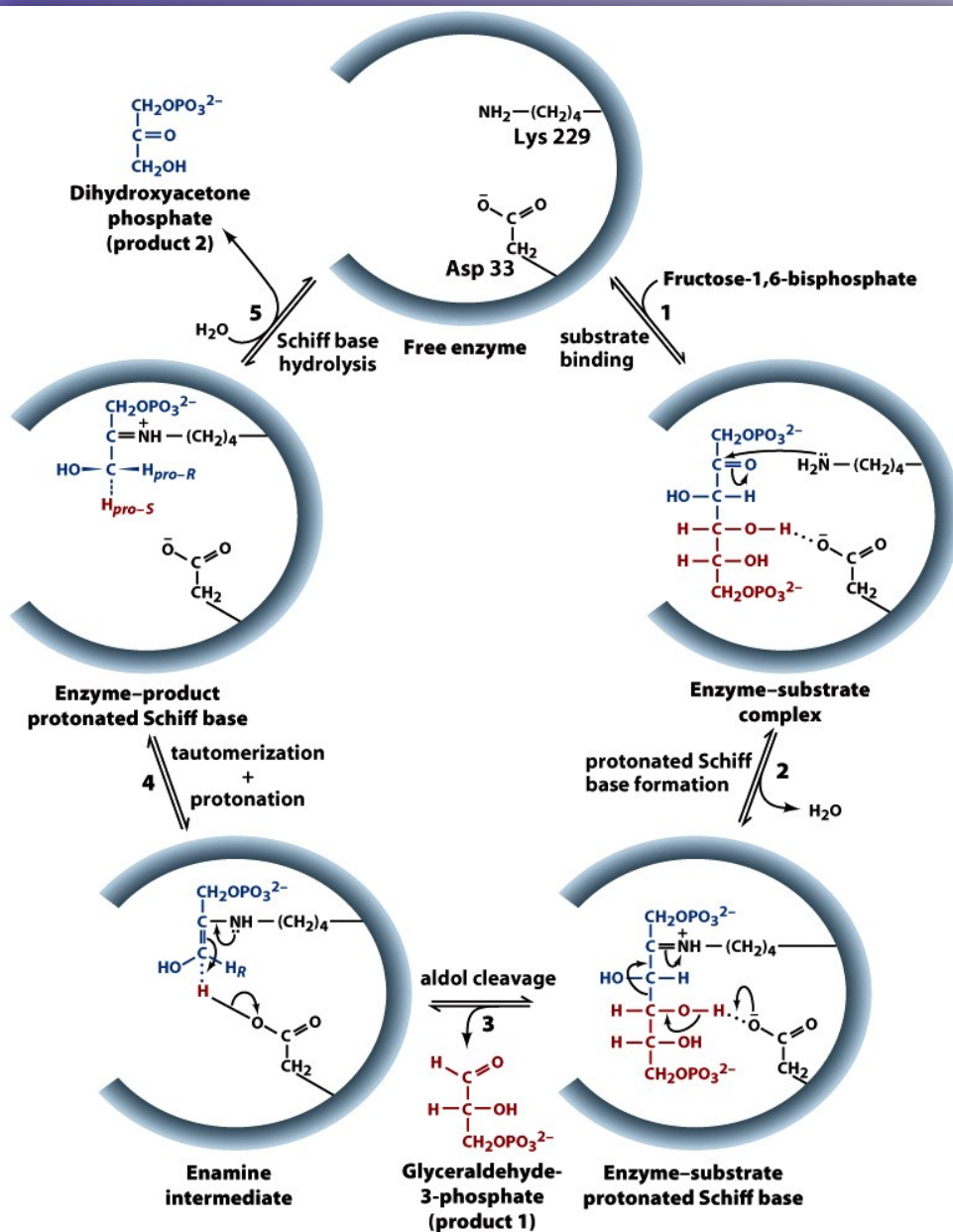
**Fructose-1,6-bisphosphate
(FBP)**

Aldolase



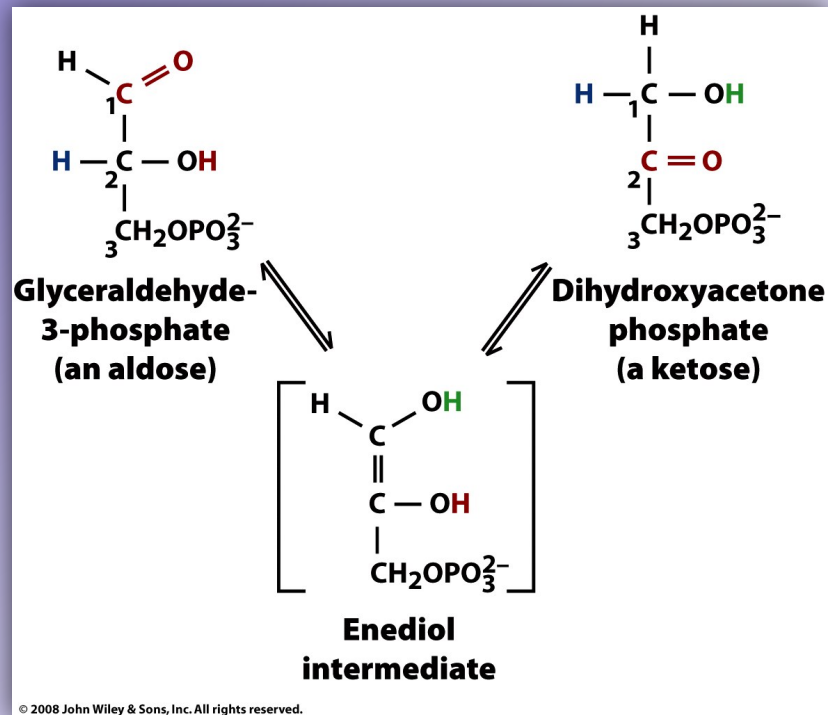
General mechanism of base-catalyzed aldol cleavage



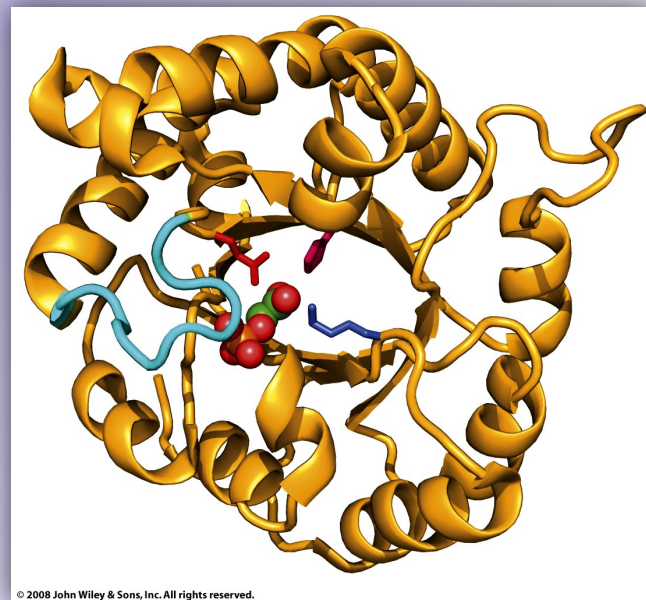
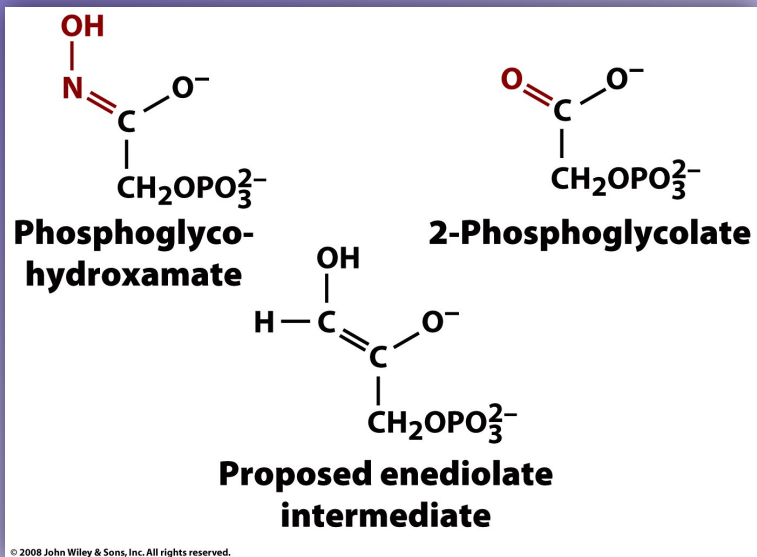


Proposed aldolase mechanism

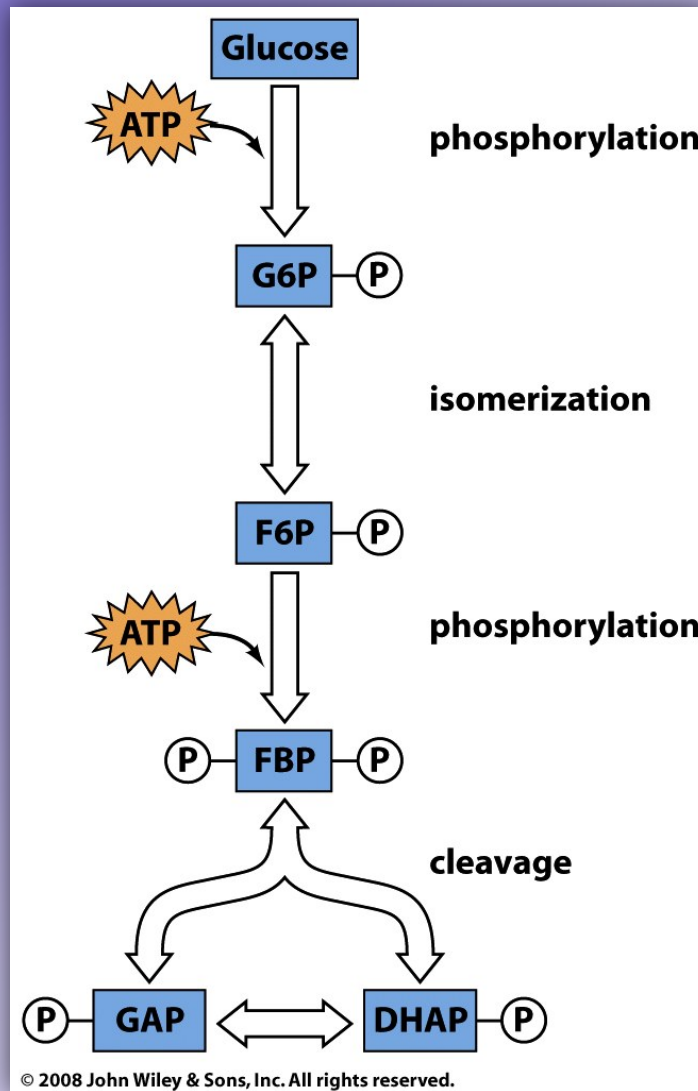
Triose-phosphate isomerase (TPI)



Transition-state analogs of TPI

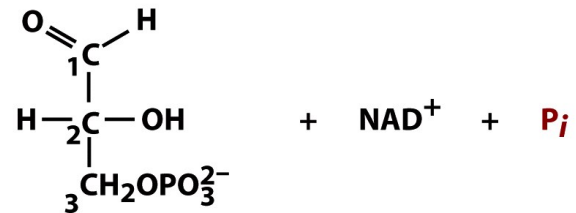


**TPI complexed
with 2-phosphoglycolate**

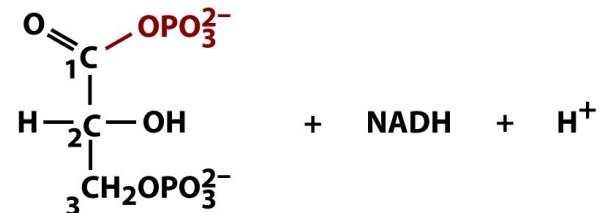
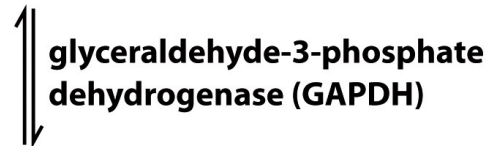


First half of glycolysis:
ATP investment phase

Glyceraldehyde 3-P dehydrogenase



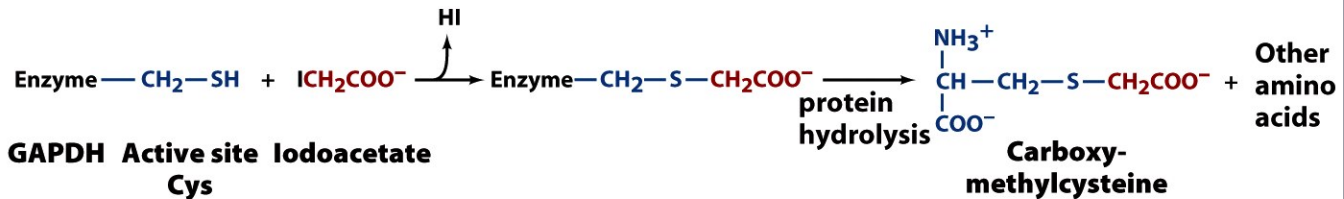
Glyceraldehyde-3-phosphate (GAP)



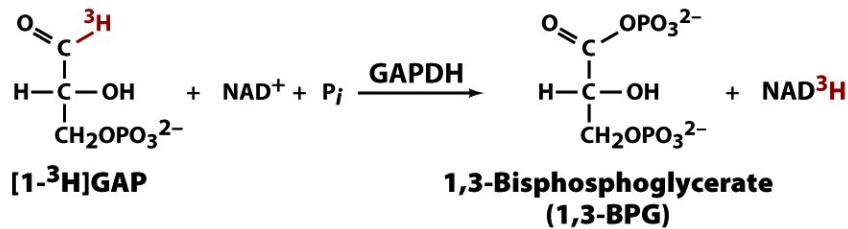
1,3-Bisphosphoglycerate (1,3-BPG)

Characterizing the G3P dehydrogenase reaction

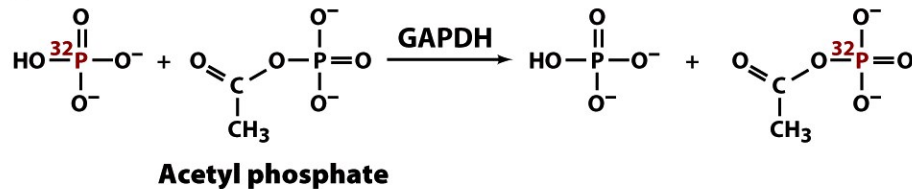
(a)

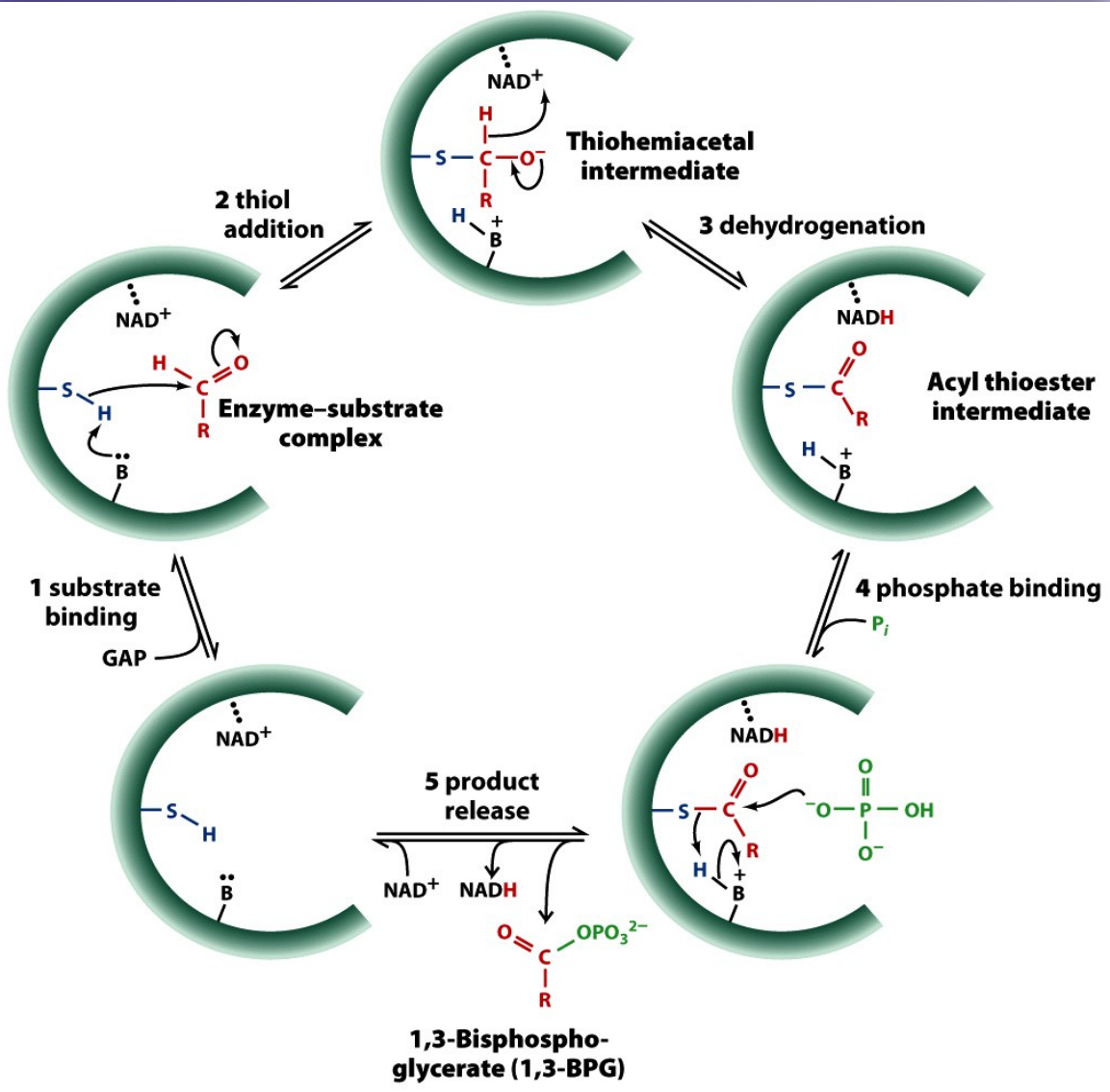


(b)



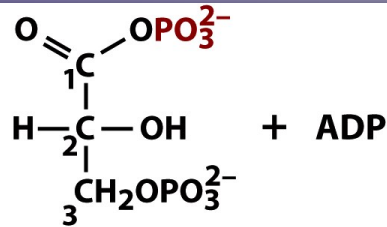
(c)



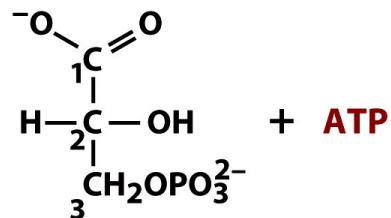


Proposed mechanism of GAP dehydrogenase

3-Phosphoglycerate kinase



1,3-Bisphosphoglycerate (1,3-BPG)



3-Phosphoglycerate (3PG)

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**First substrate-level
phosphorylation
reaction of glycolysis**

Mechanism of the PGK reaction

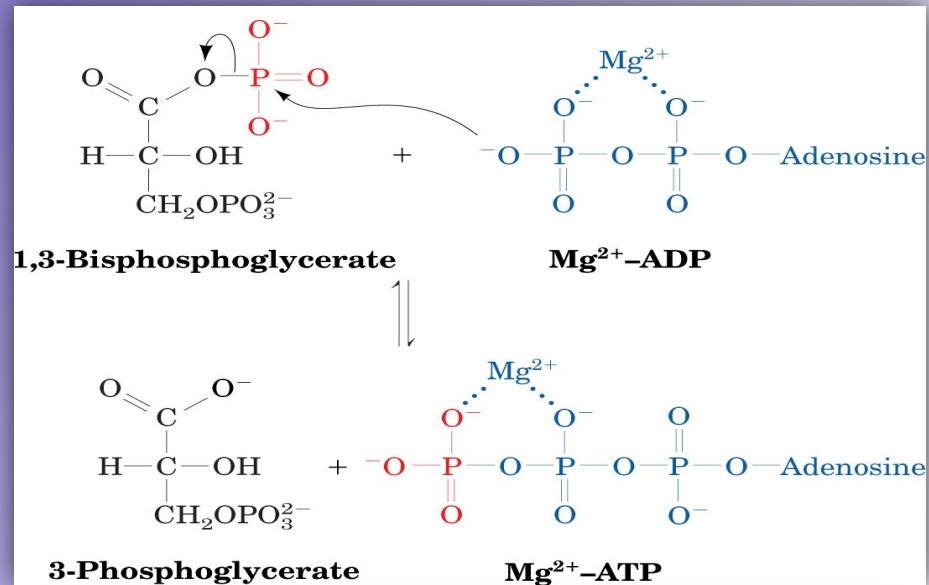


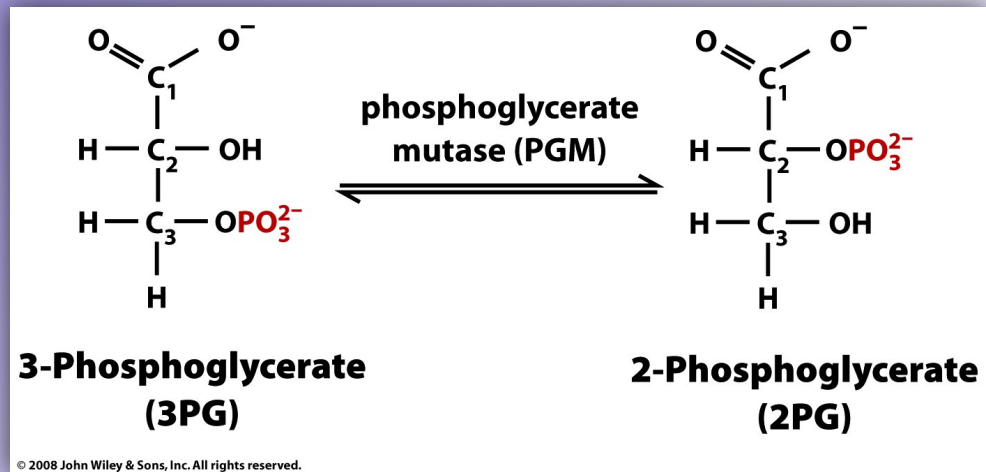
Table 15-1

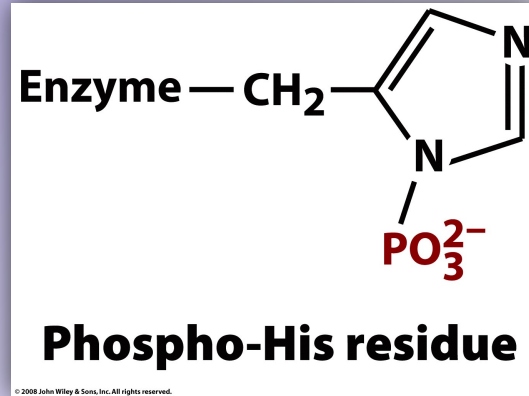
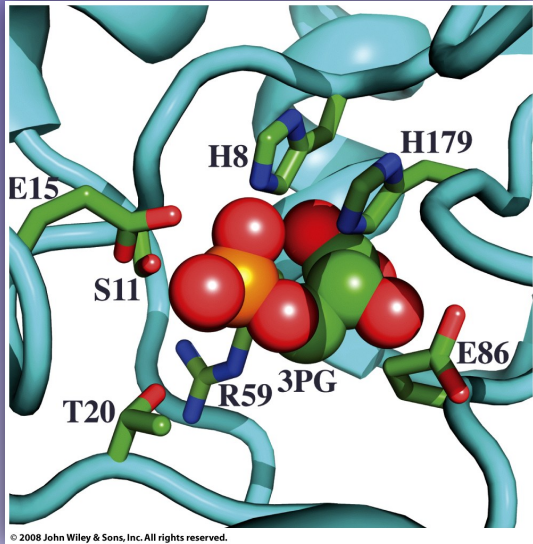
$\Delta G^{\circ'}$ and ΔG for the Reactions of Glycolysis in Heart Muscle^a

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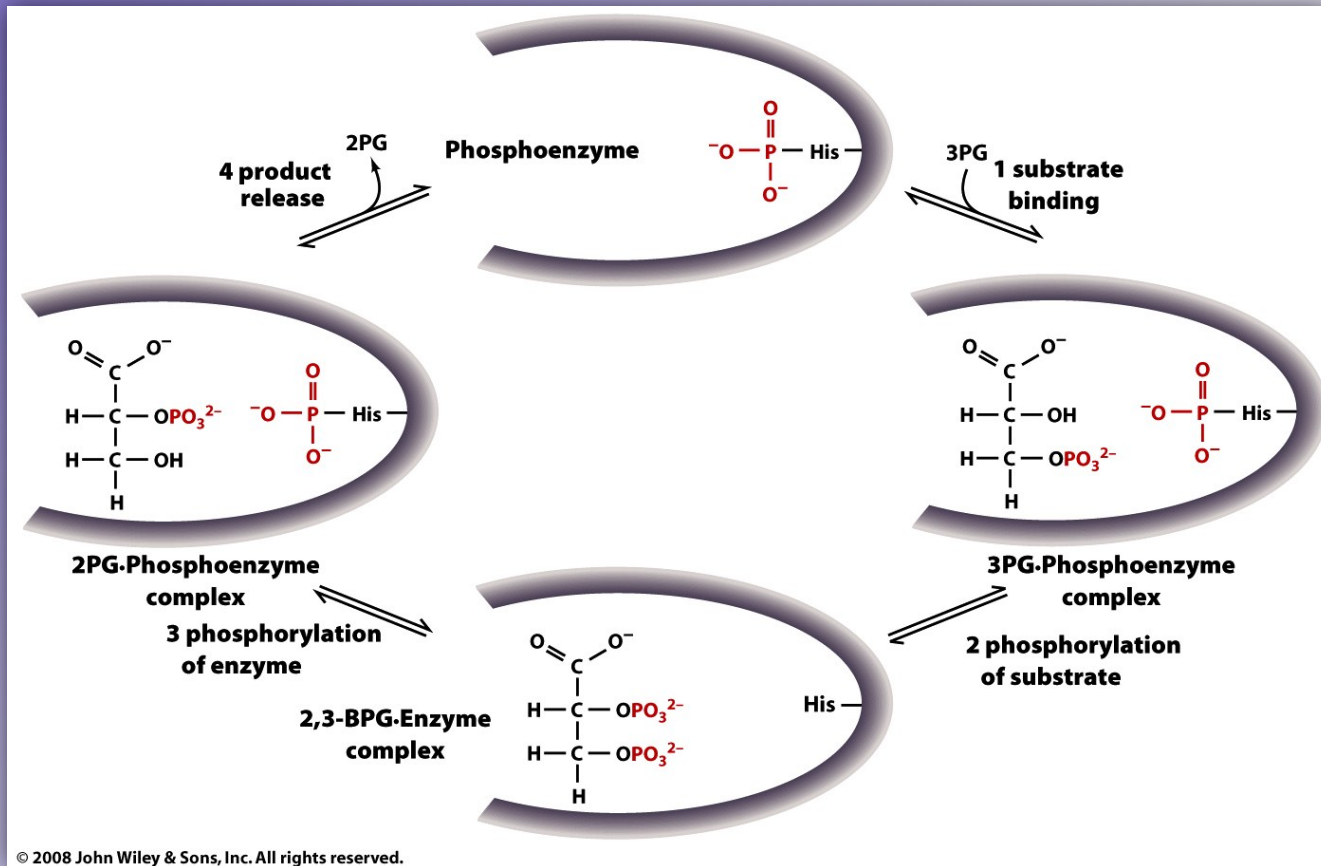
^aCalculated from data in Newsholme, E.A. and Start, C., *Regulation in Metabolism*, p. 97, Wiley (1973).

3-Phosphoglycerate mutase

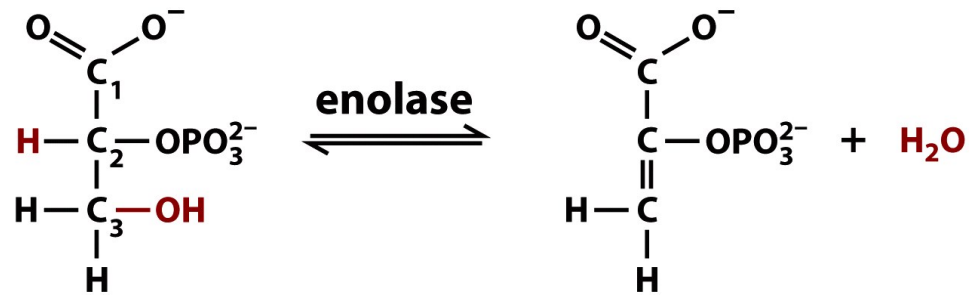




Proposed mechanism of 3-phosphoglycerate mutase



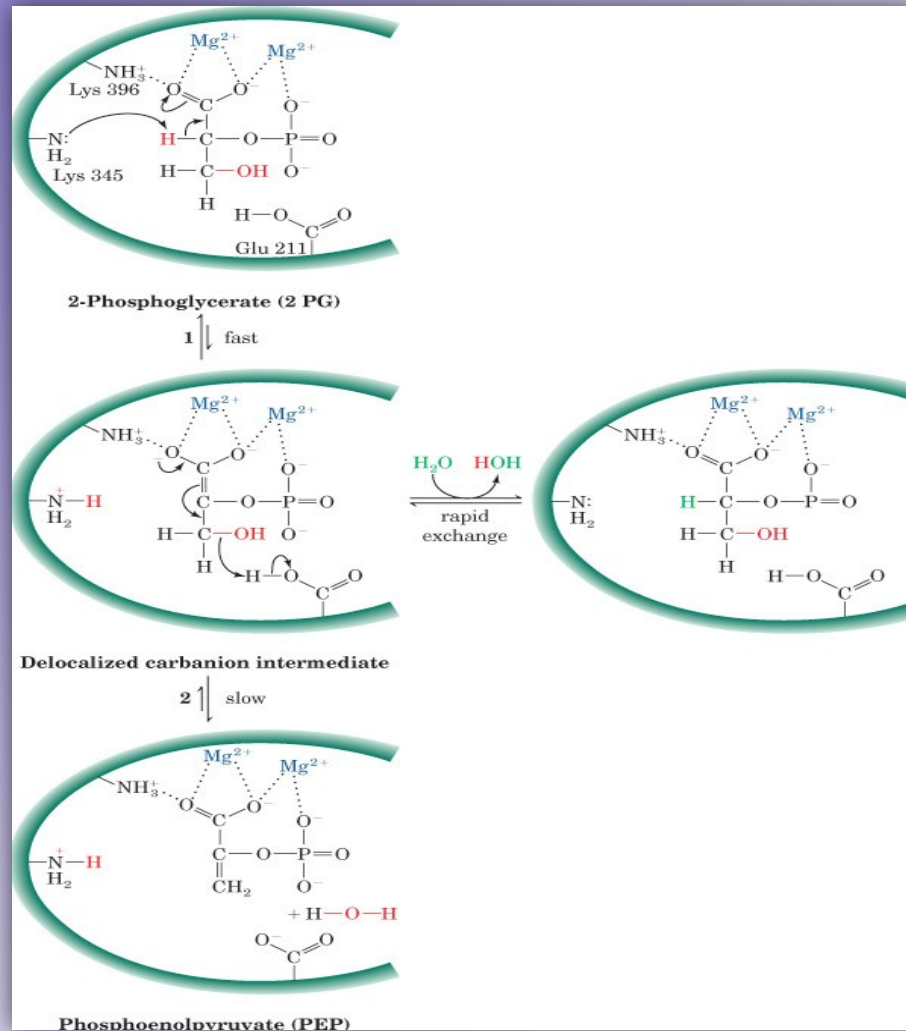
Enolase



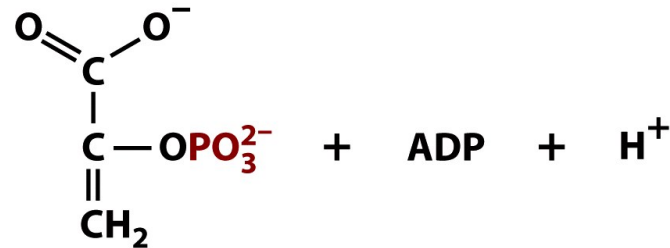
**2-Phosphoglycerate
(2PG)**

**Phosphoenolpyruvate
(PEP)**

Proposed reaction mechanism of enolase

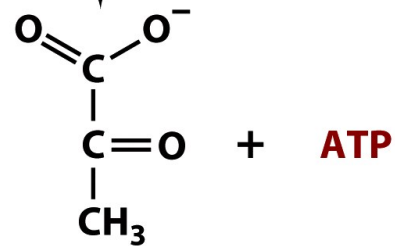


Pyruvate kinase



Phosphoenolpyruvate (PEP)

↓
pyruvate
kinase (PK)

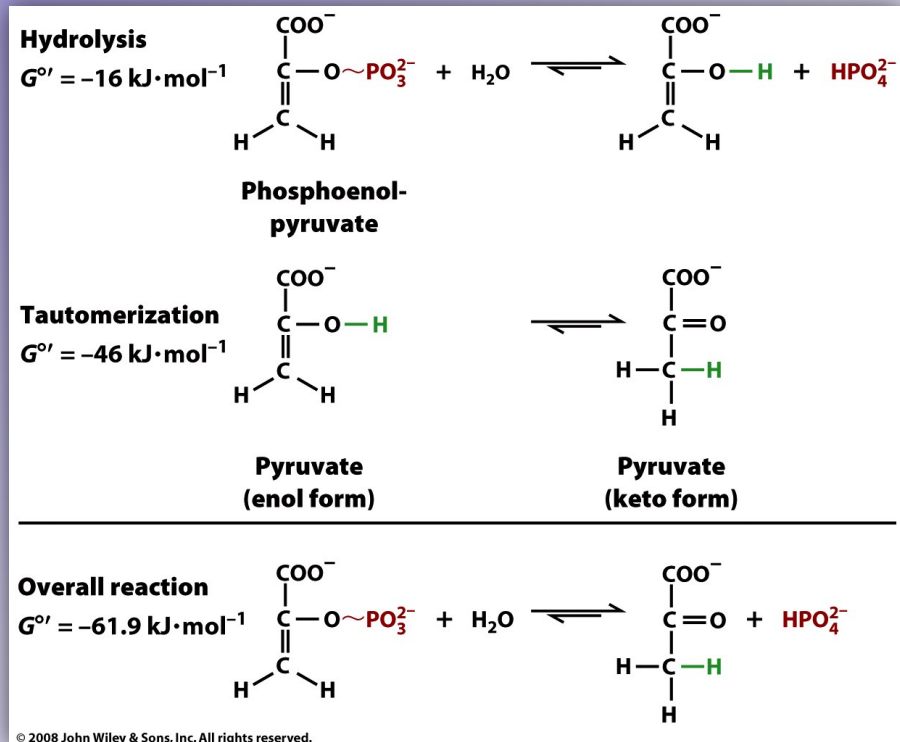
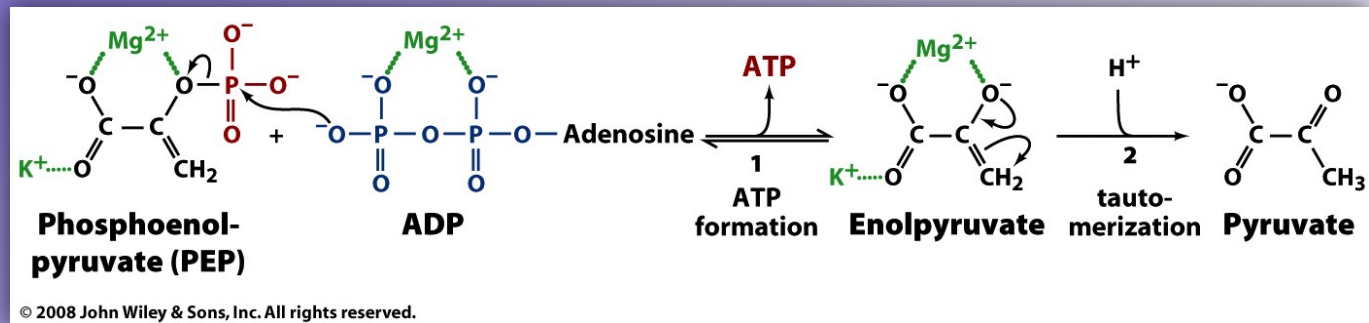


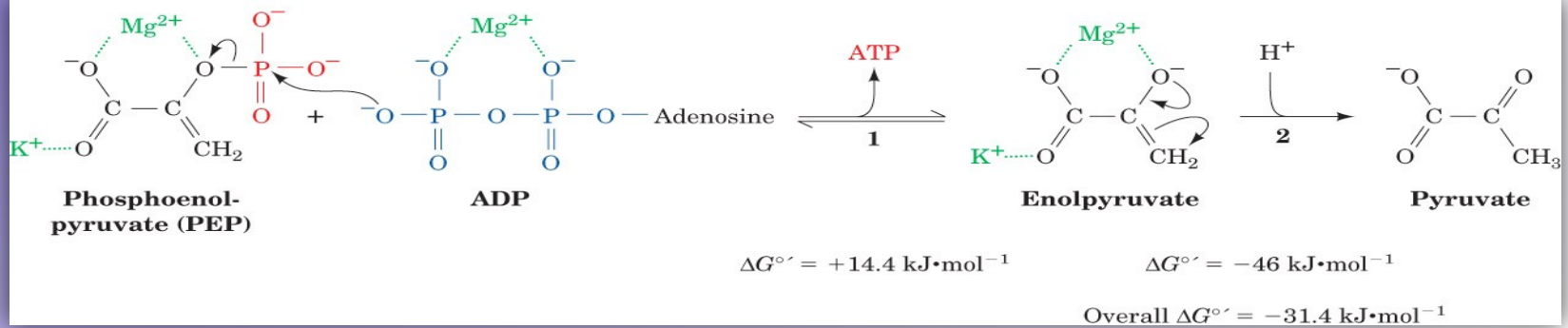
Pyruvate

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**Second substrate-level
phosphorylation
reaction of glycolysis**

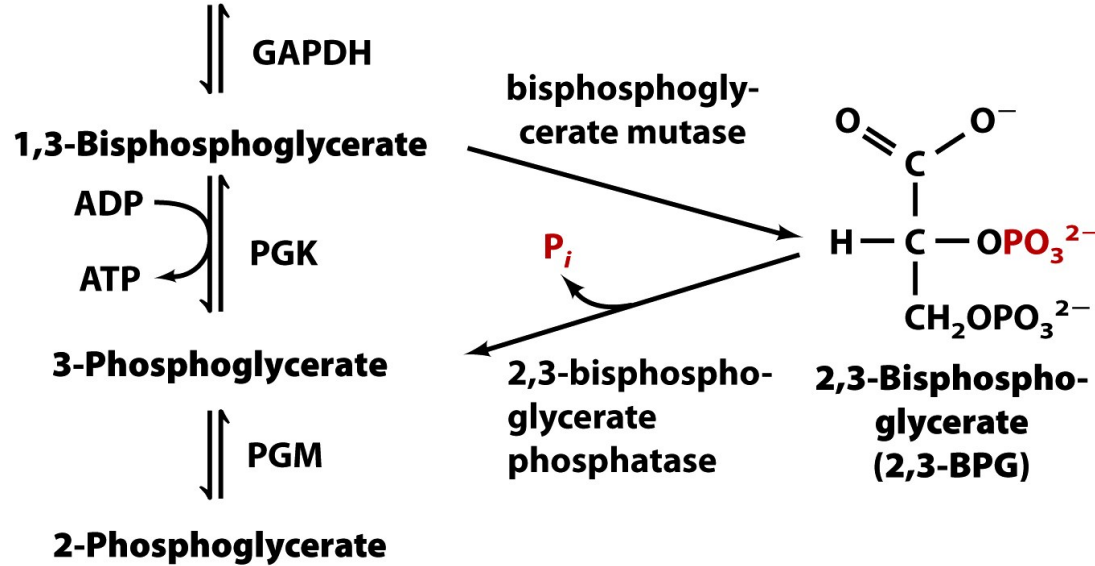
Explanation of the very (-) change in free energy associated with the PK reaction



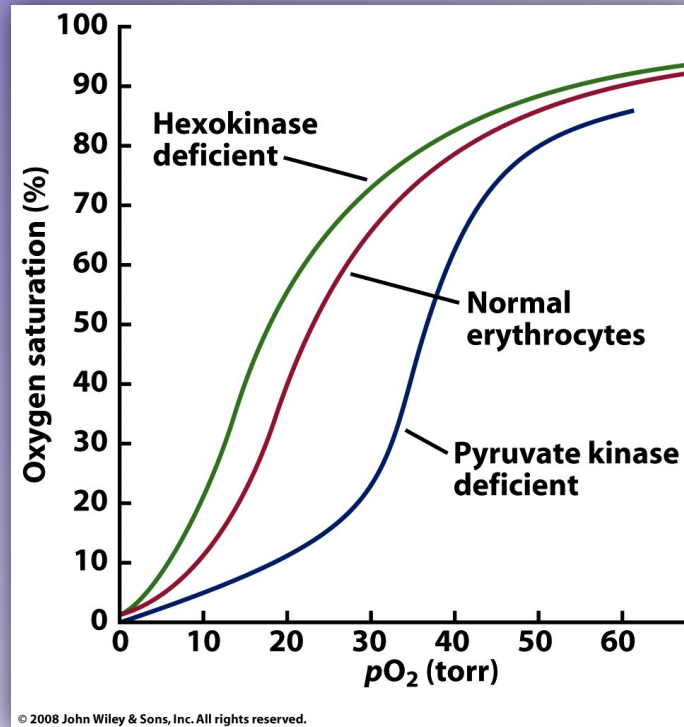


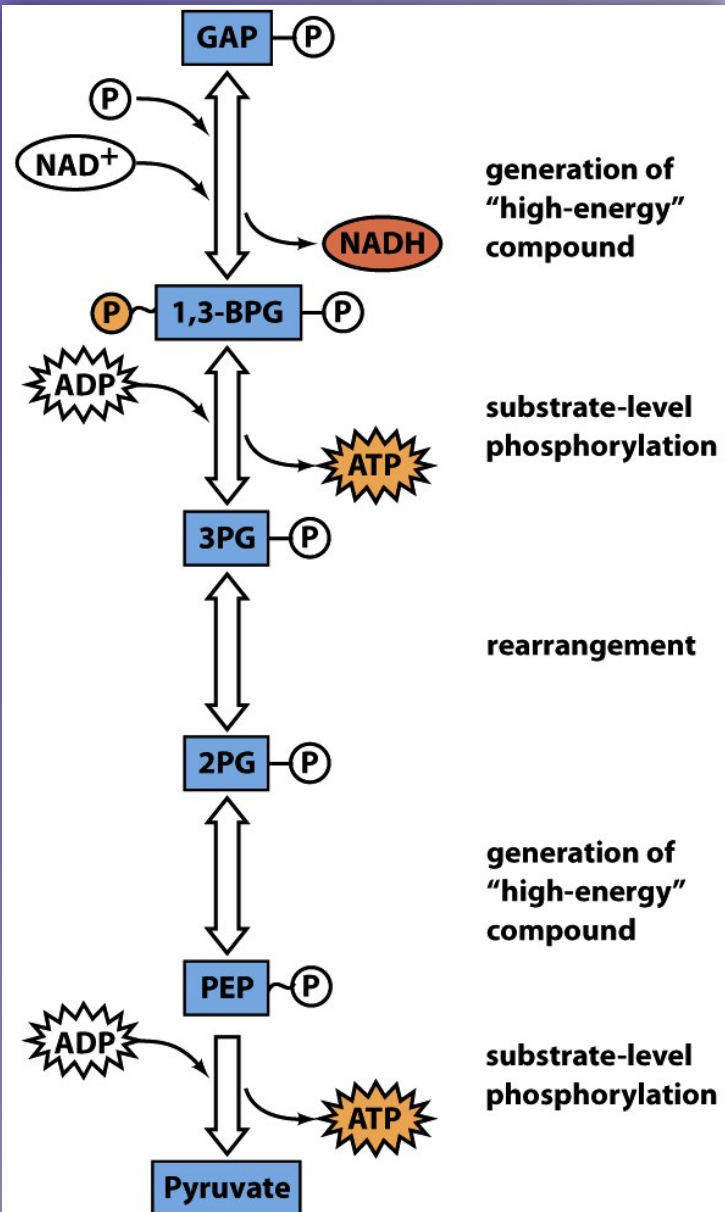
Glycolytic detour: 2,3-BPG biosynthesis in erythrocytes

Glyceraldehyde 3-phosphate



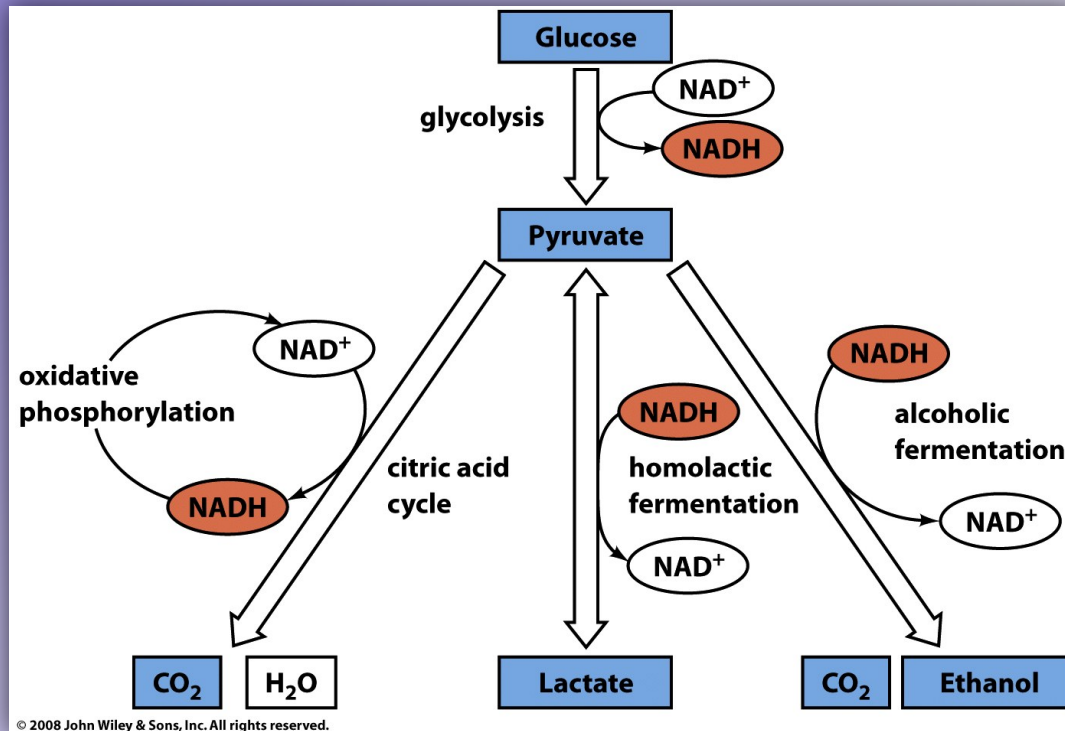
Impact of 2,3-BPG on O_2 -hemoglobin binding affinity: hexokinase and PK deficiencies



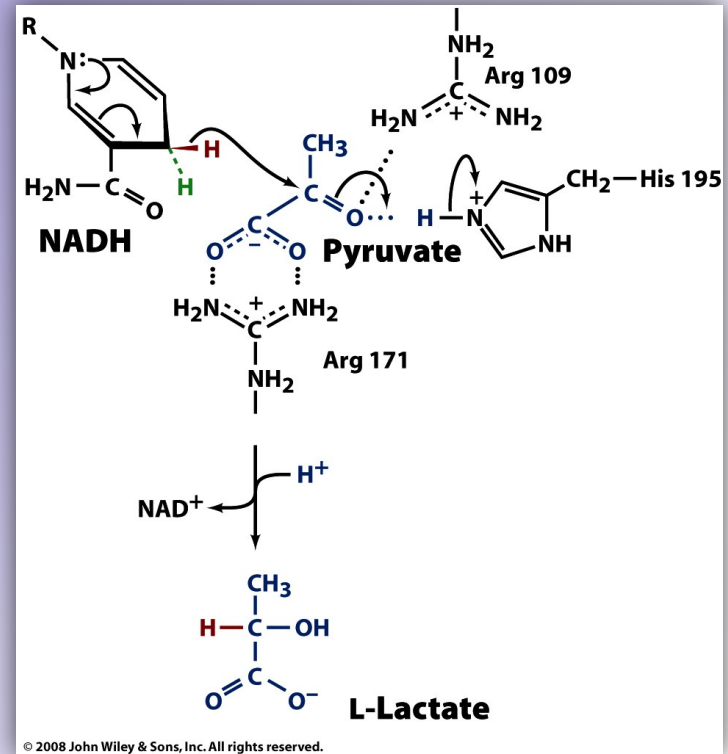
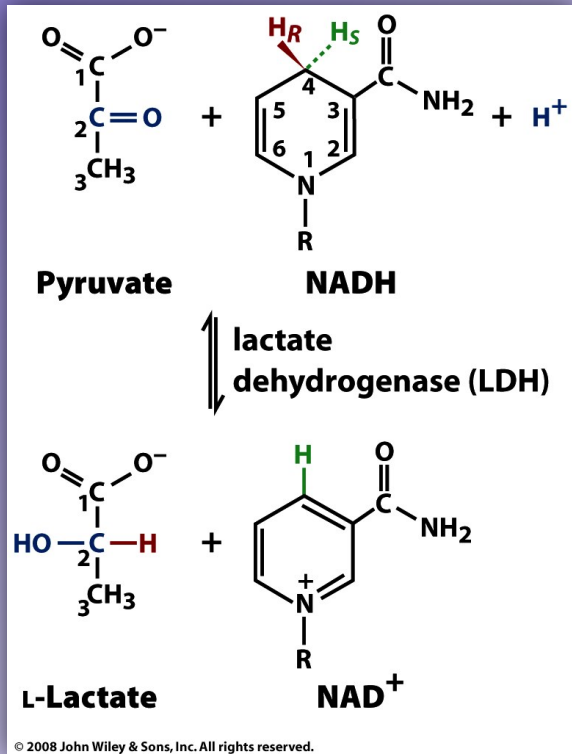


Second half of glycolysis: ATP-yielding phase

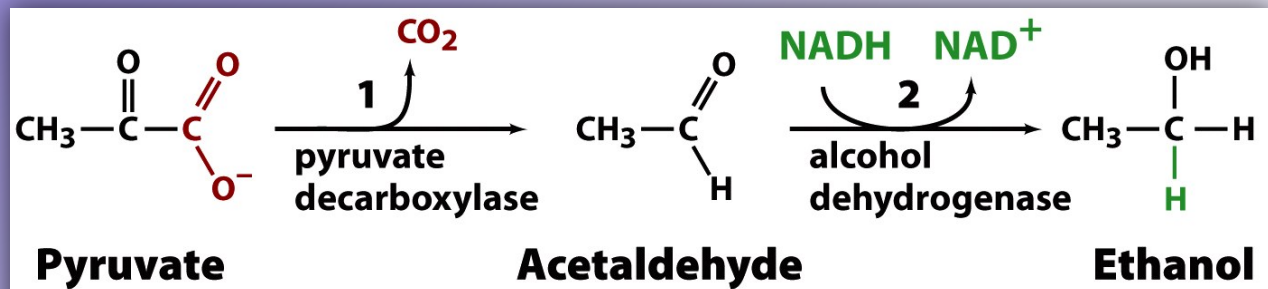
Metabolic fates of pyruvate



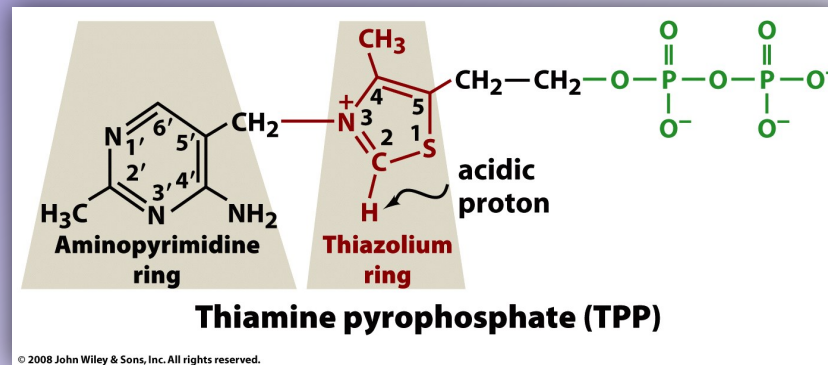
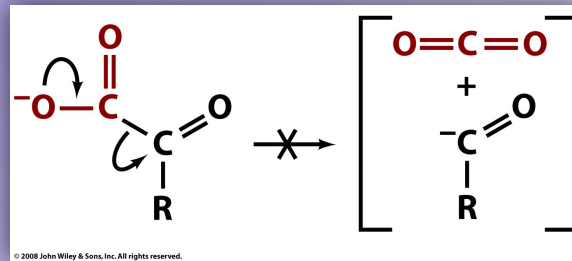
Conversion of pyruvate to lactate: LDH reaction

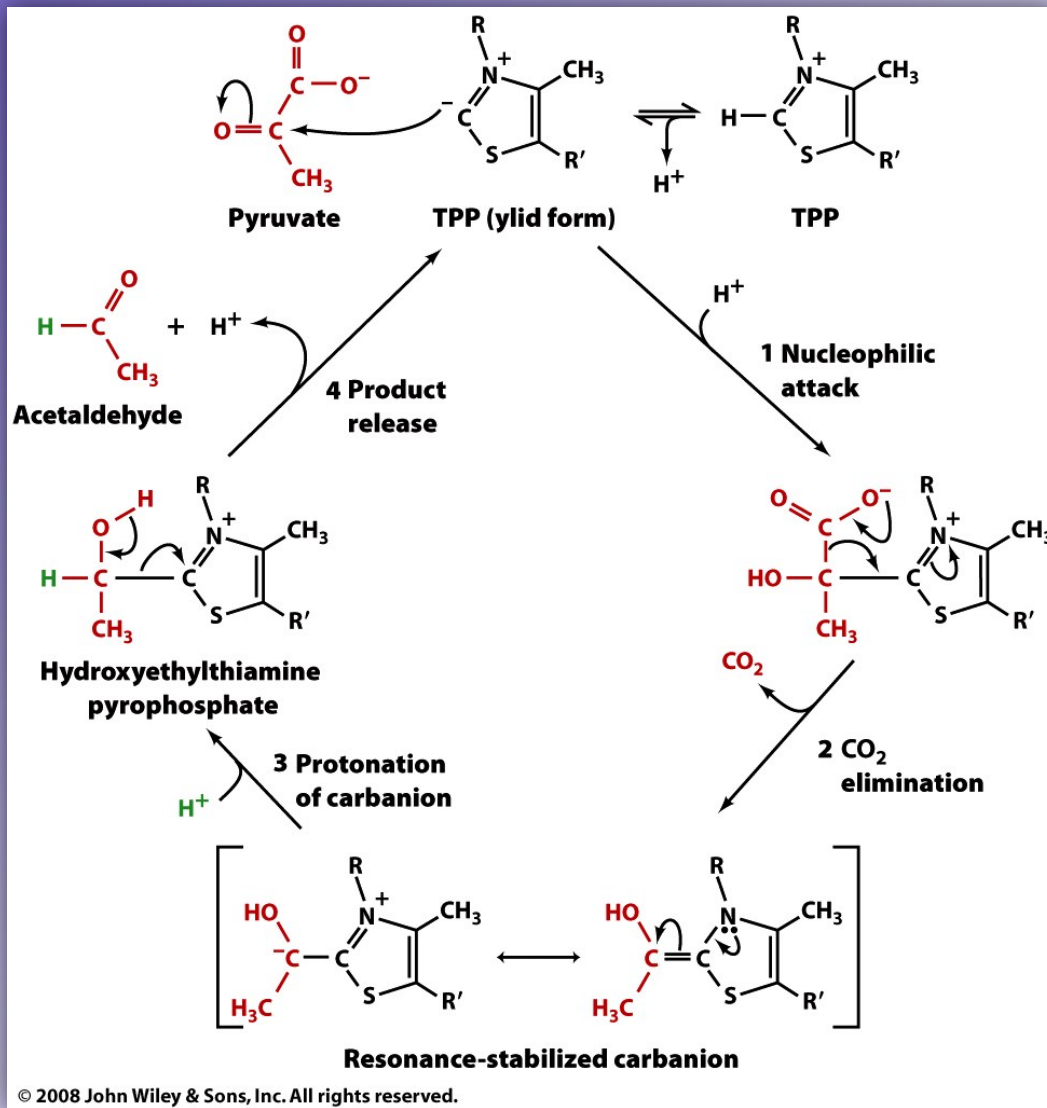


Conversion of pyruvate to ethanol via acetaldehyde



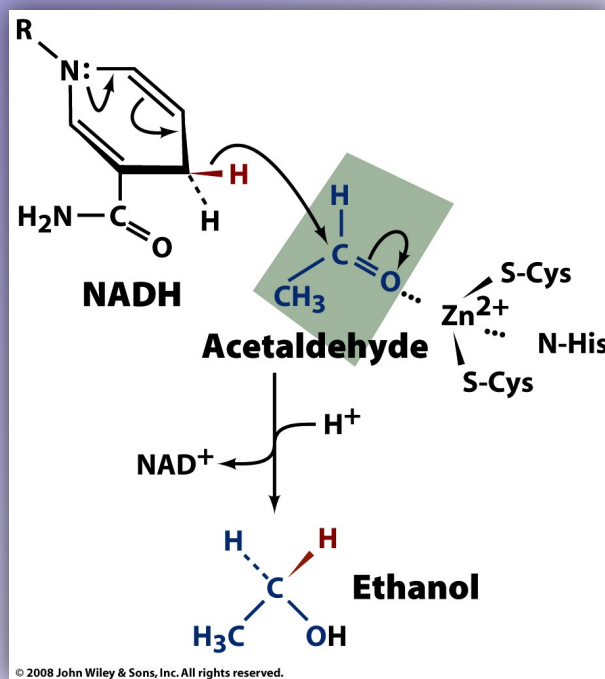
Enzyme-catalyzed decarboxylation of an α -ketoacid (pyruvate): TPP coenzyme is required for charge delocalization





Proposed mechanism of pyruvate decarboxylase

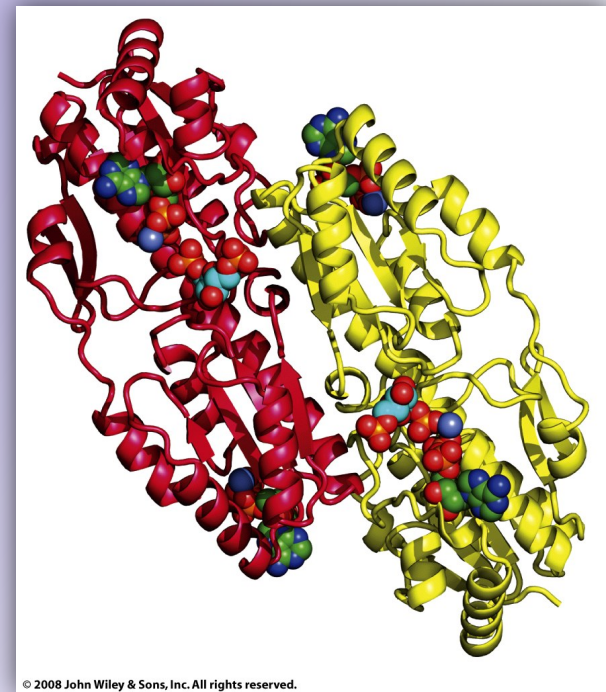
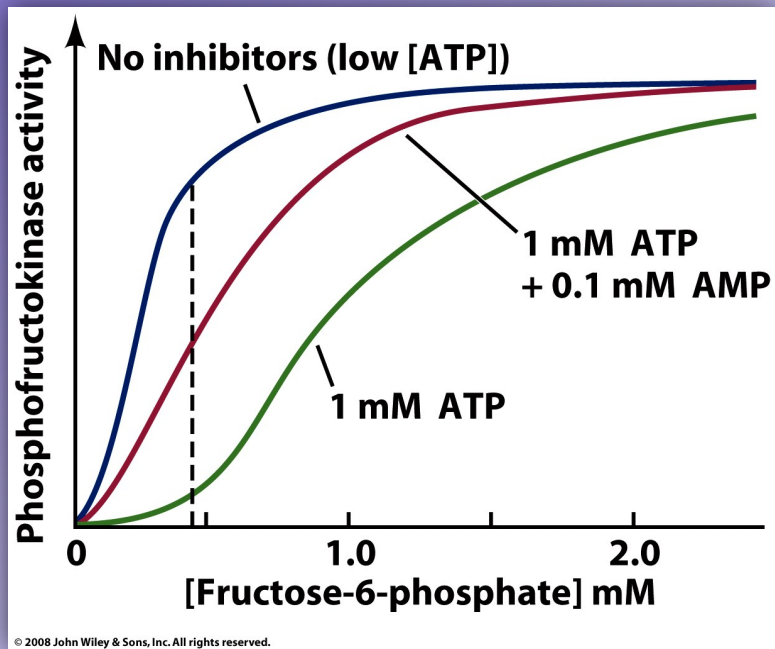
Alcohol dehydrogenase: Stereospecific transfer of hydride from NADH to acetaldehyde



Some effectors of the non-equilibrium enzymes of glycolysis

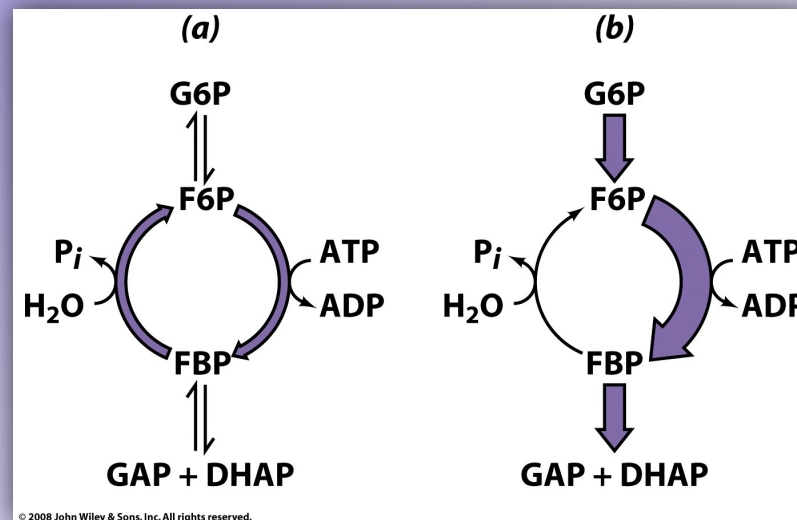
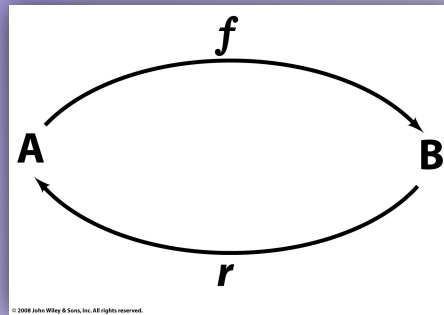
Enzyme	Inhibitors	Activators ^a
HK	G6P	–
PFK	ATP, citrate, PEP	ADP, AMP, cAMP, FBP, F2,6P, F6P, NH ₄ ⁺ , P _i
PK (muscle)	ATP	AMP, PEP, FBP

^aThe activators for PFK are better described as deinhibitors of ATP because they reverse the effect of inhibitory concentrations of ATP.



PFK is an allosteric enzyme (tetramer)

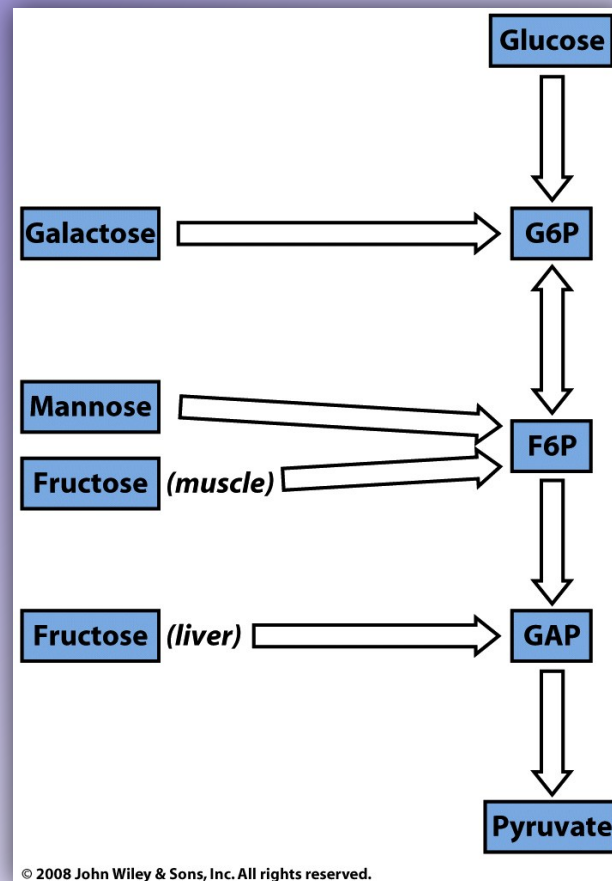
Control of glycolytic flux via substrate cycling



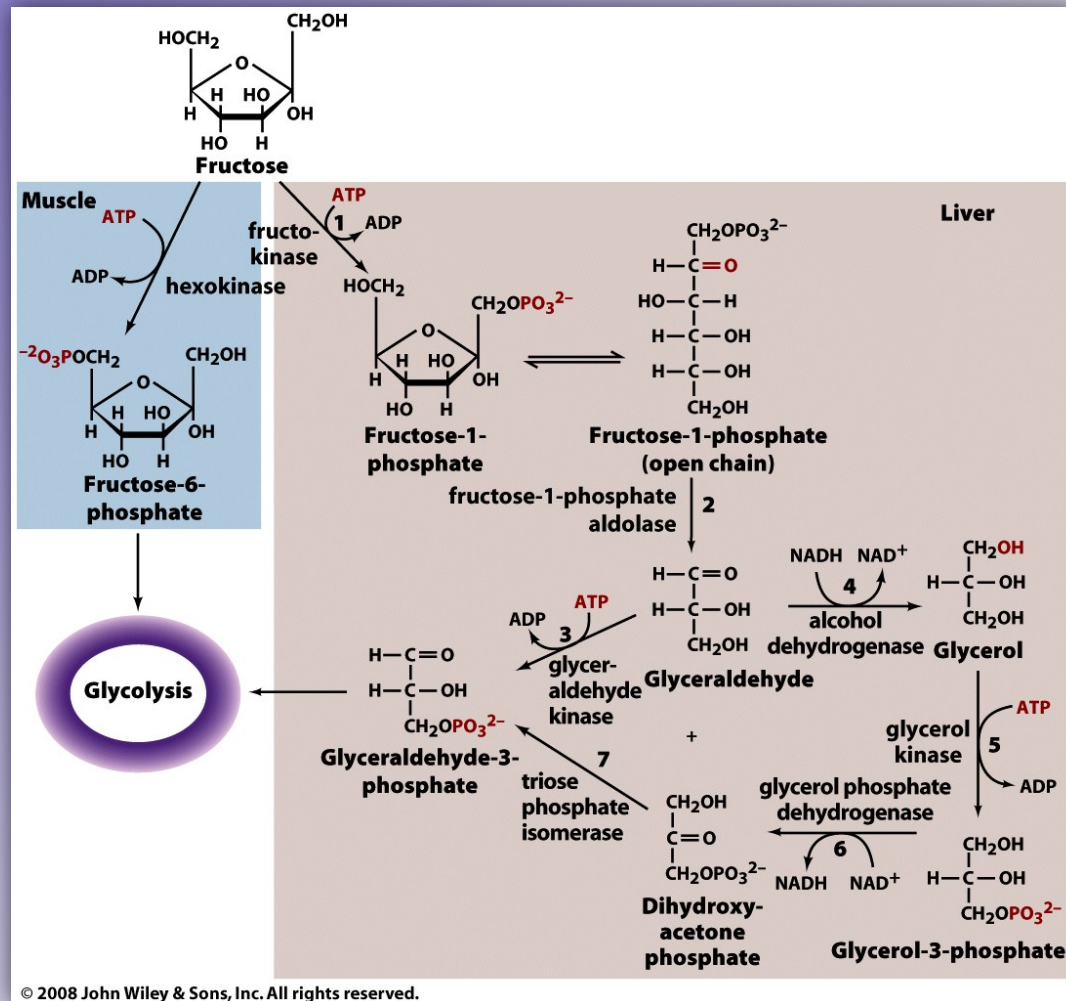
resting muscle

active muscle

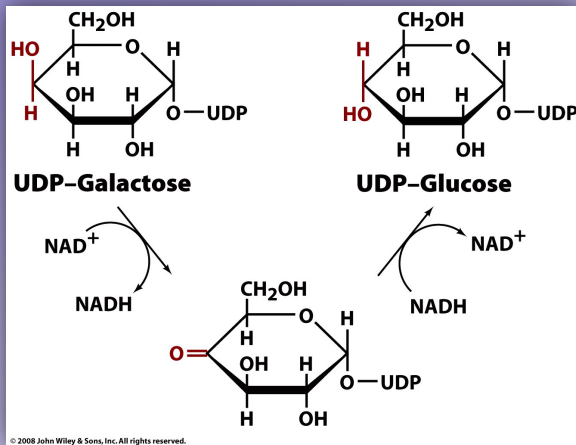
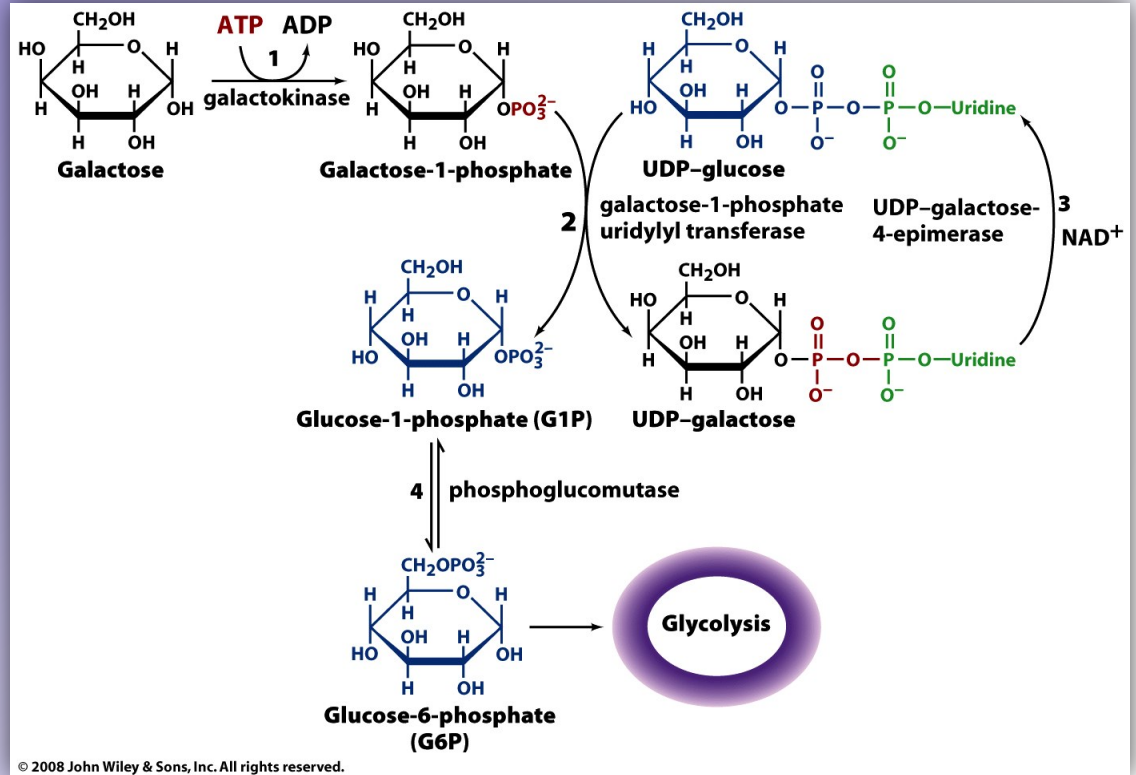
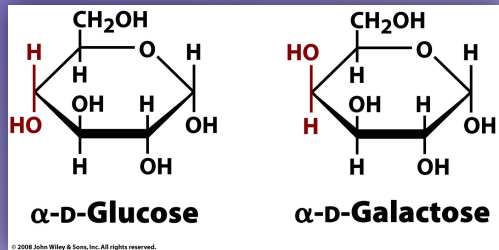
Entry routes of other monosaccharides into glycolysis



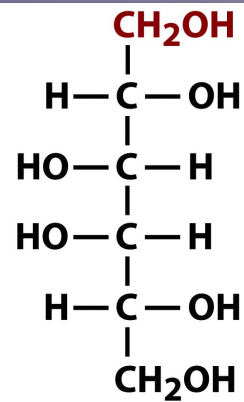
Metabolism of D-fructose is organ-dependent



Metabolism of D-galactose (human)



By-product of galactosemia (step 2 deficient)
(buildup in lens of the eye - cataracts)



Galactitol

Metabolism of D-mannose (human)

