

David L. Nelson and Michael M. Cox

LEHNINGER

PRINCIPLES OF BIOCHEMISTRY

Fifth Edition

CHAPTER 14

**Glycolysis, Gluconeogenesis, and the
Pentose Phosphate Pathway**

Dr.Sulيمان Al-Khalil

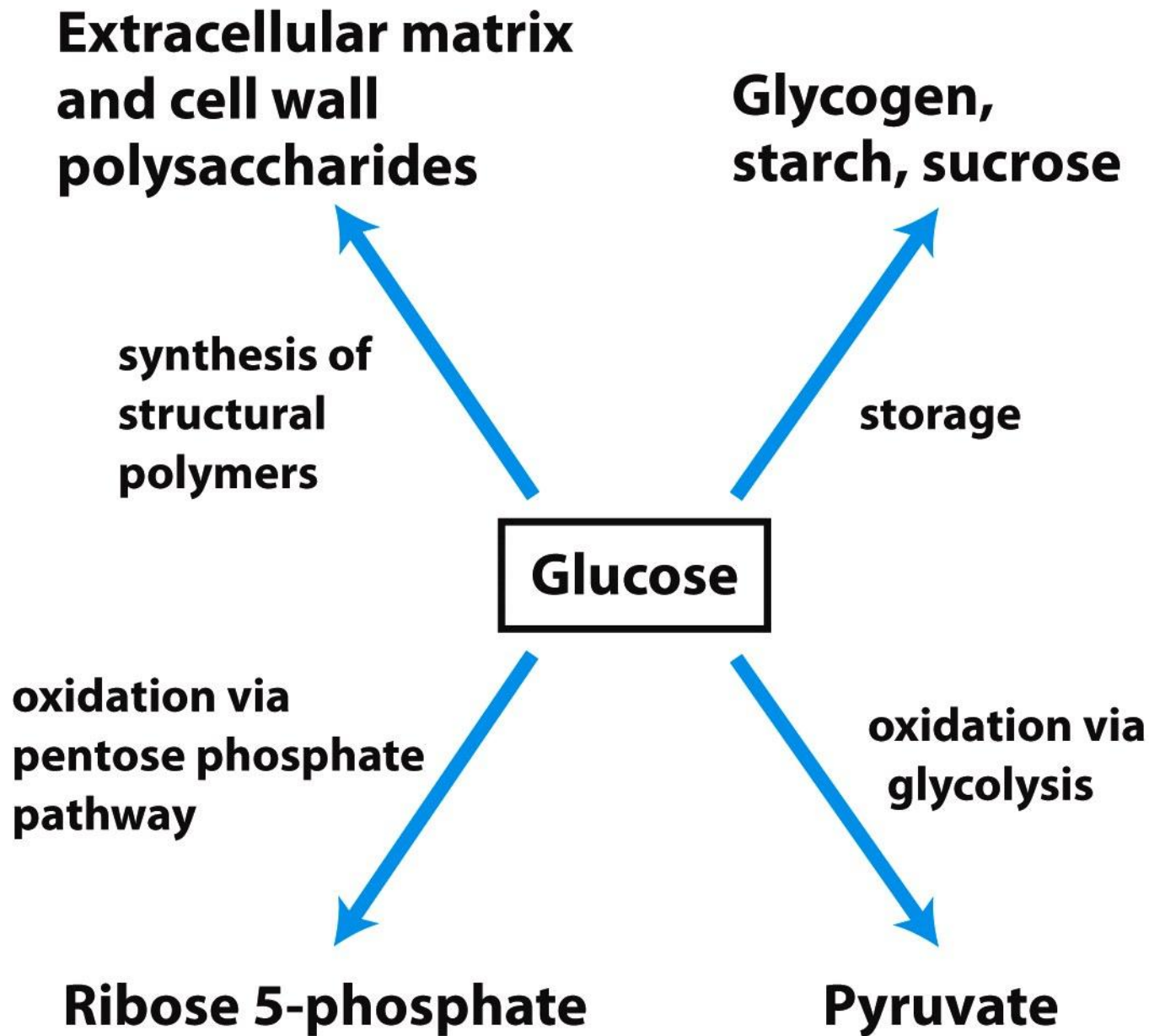


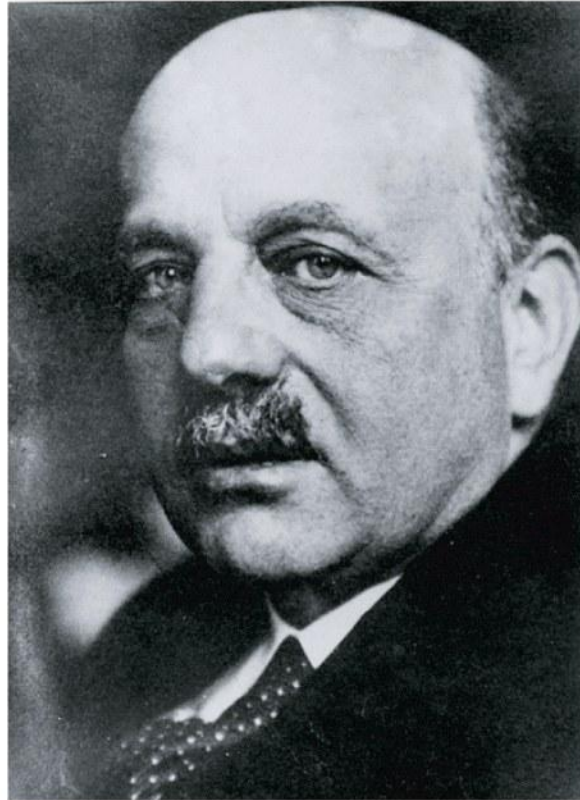
Figure 14-1

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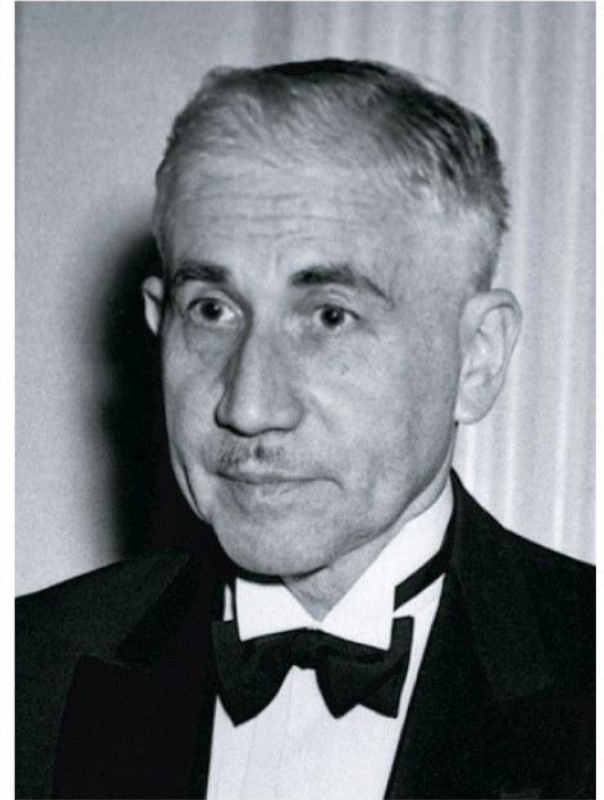
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Hans von Euler-Chelpin
1873–1964



Gustav Embden
1874–1933



Otto Meyerhof
1884–1951

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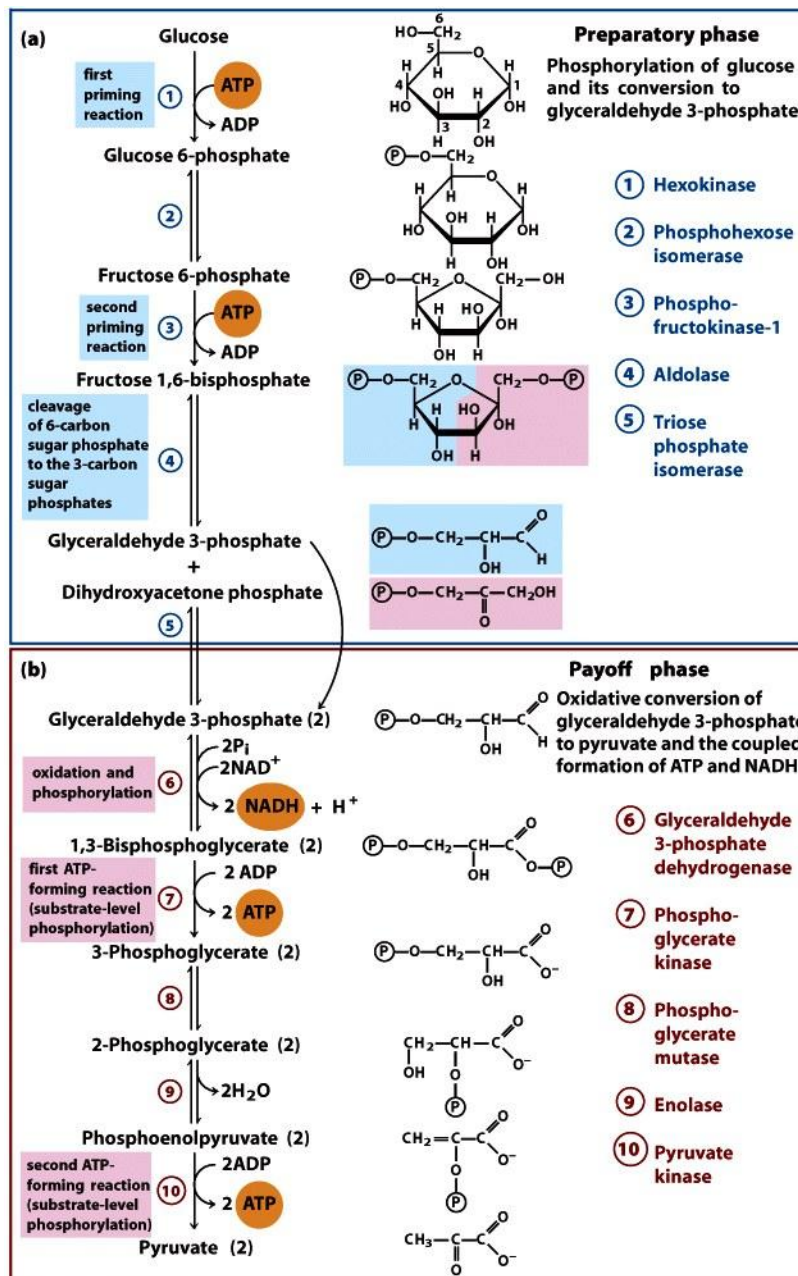


Figure 14-2

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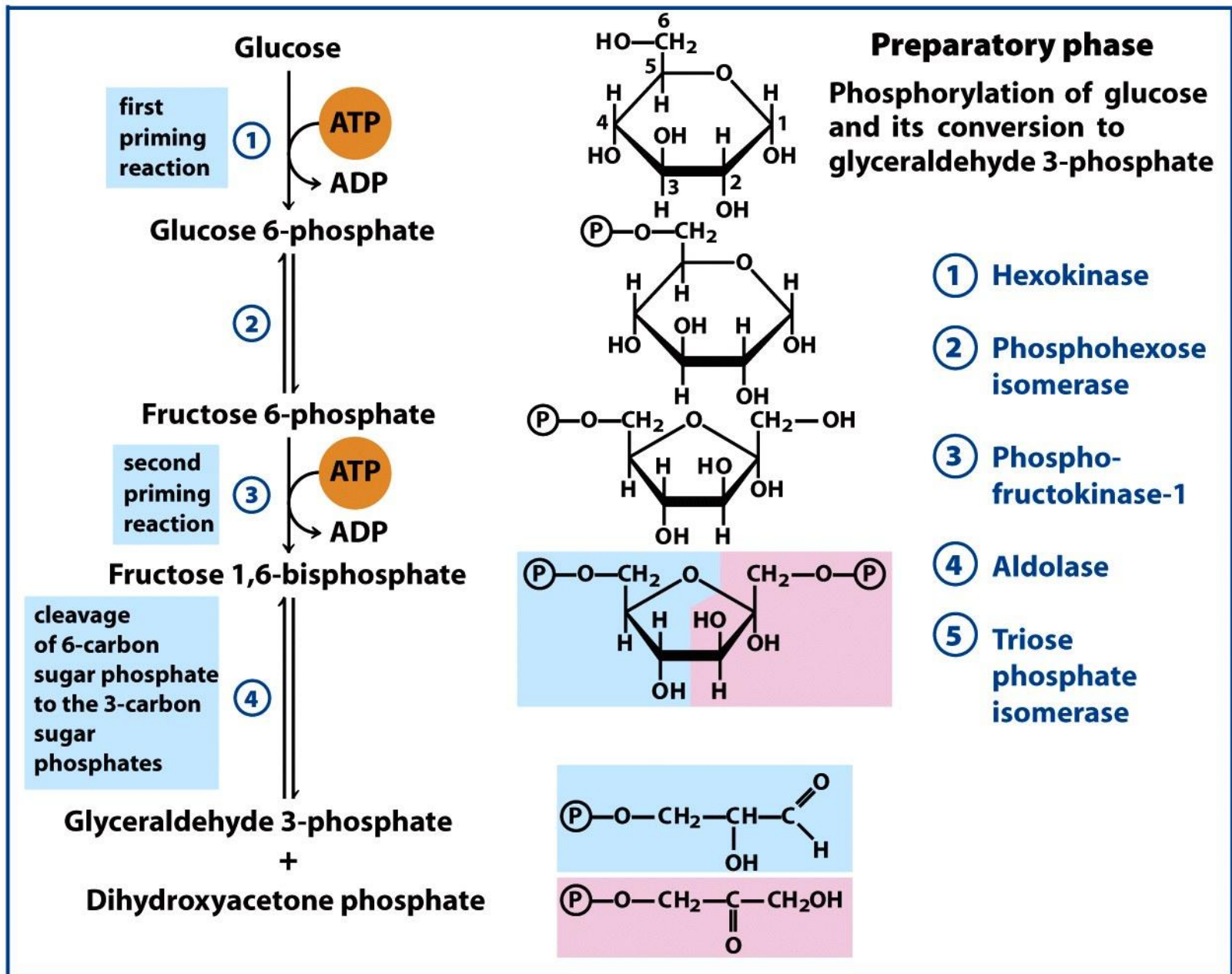


Figure 14-2a

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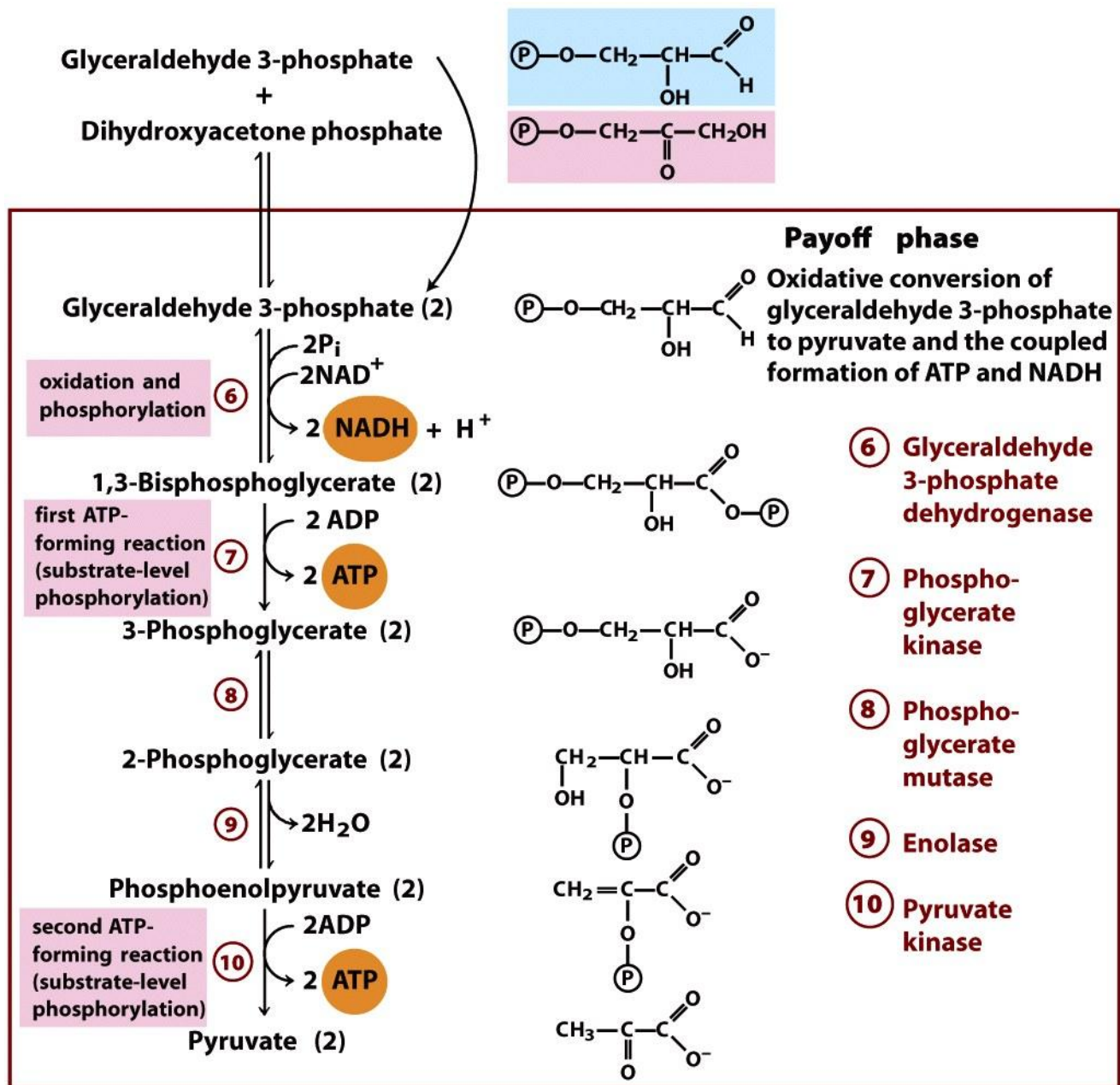


Figure 14-2b

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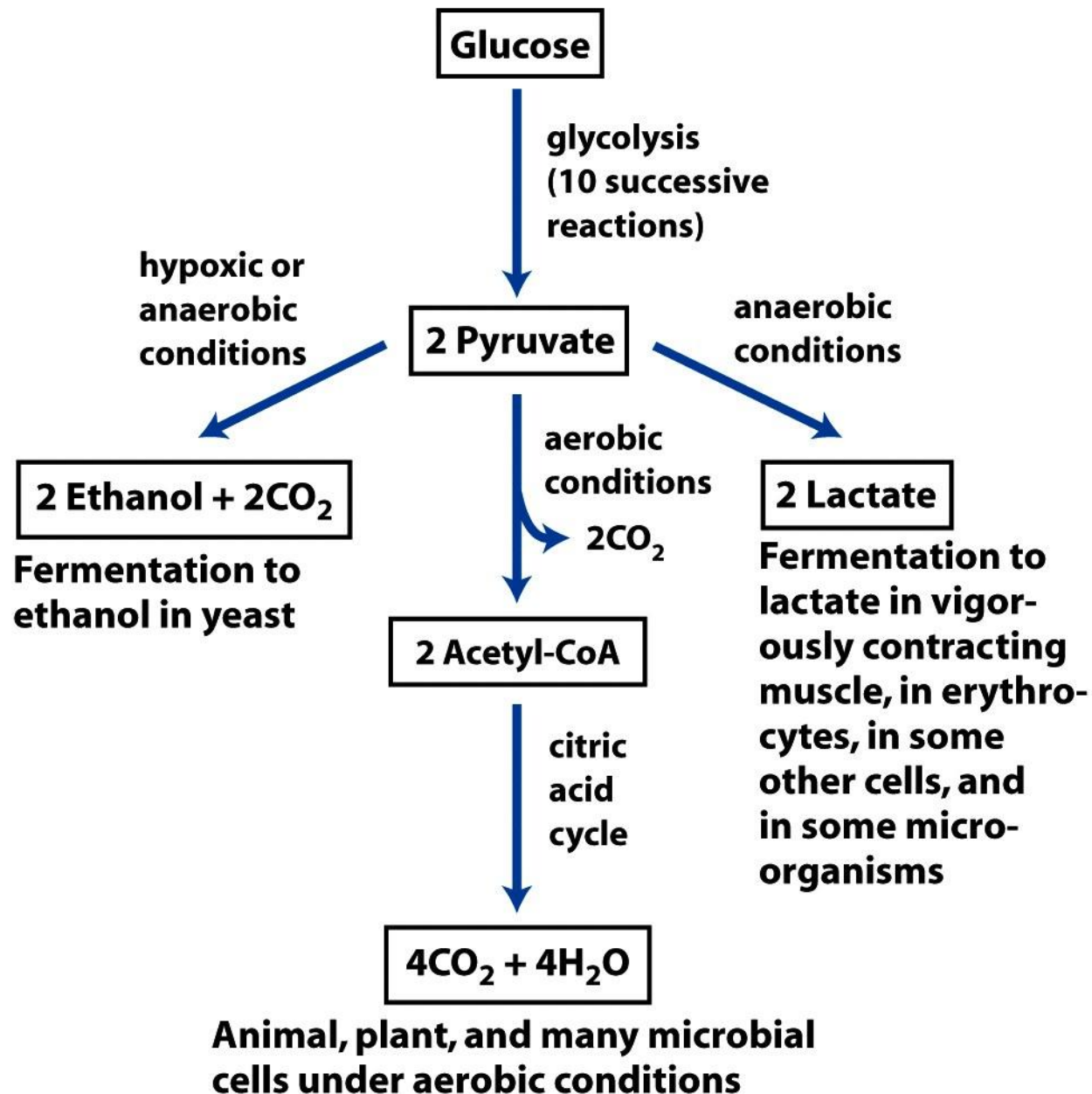
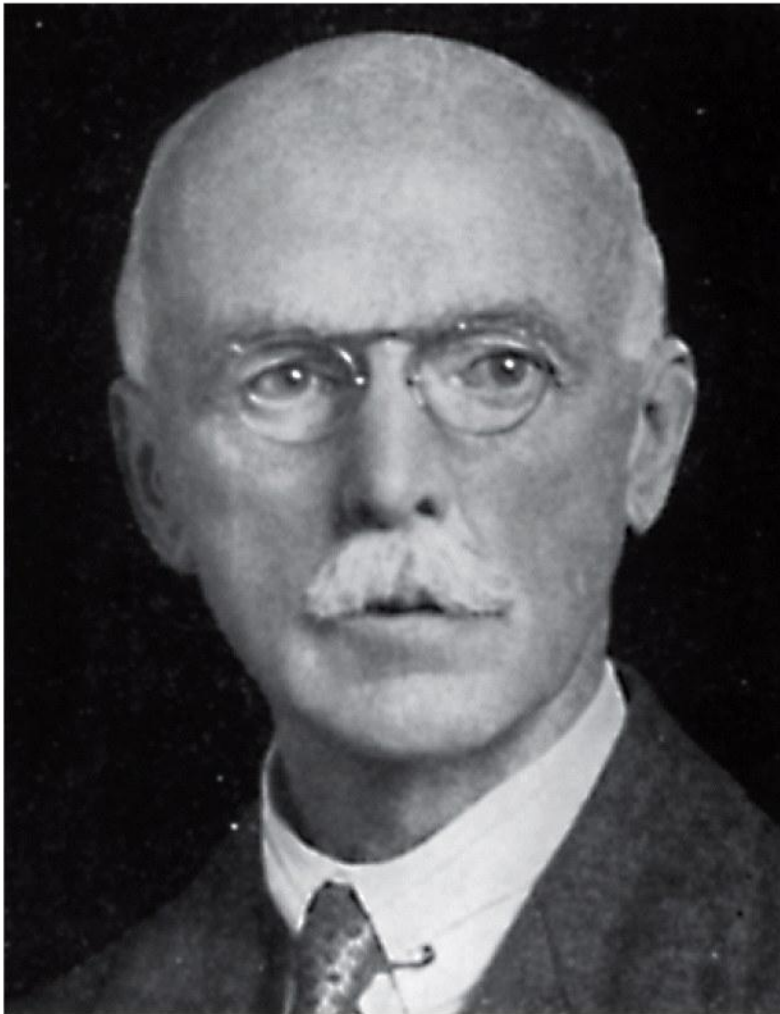


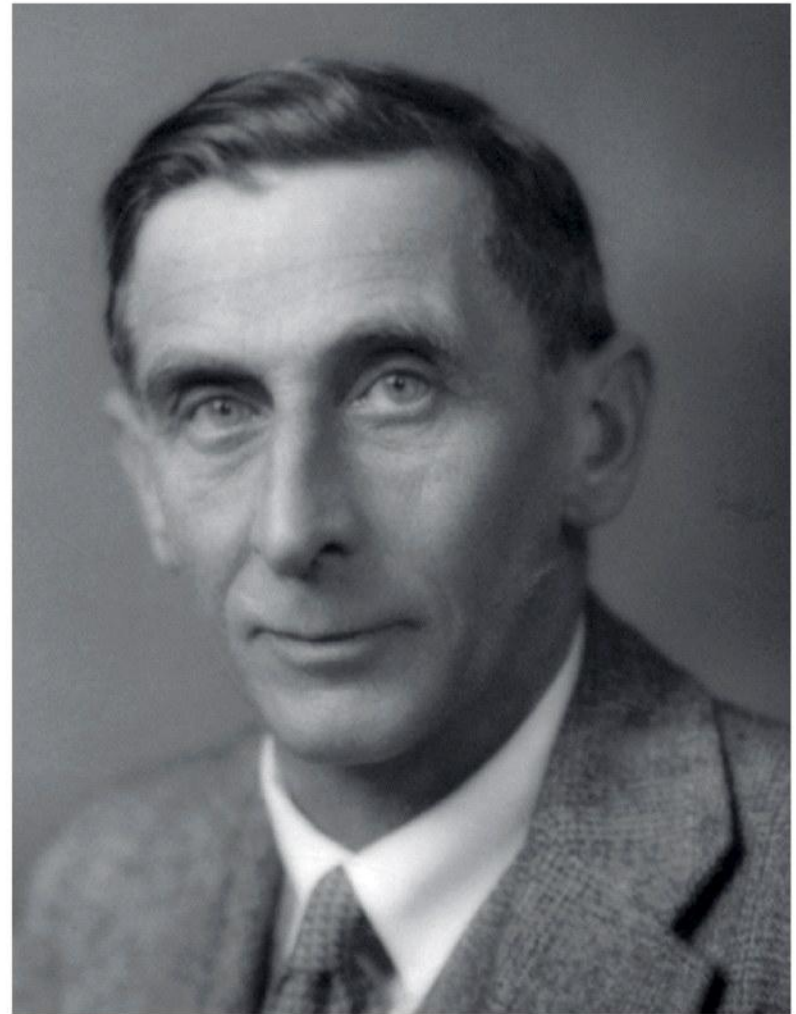
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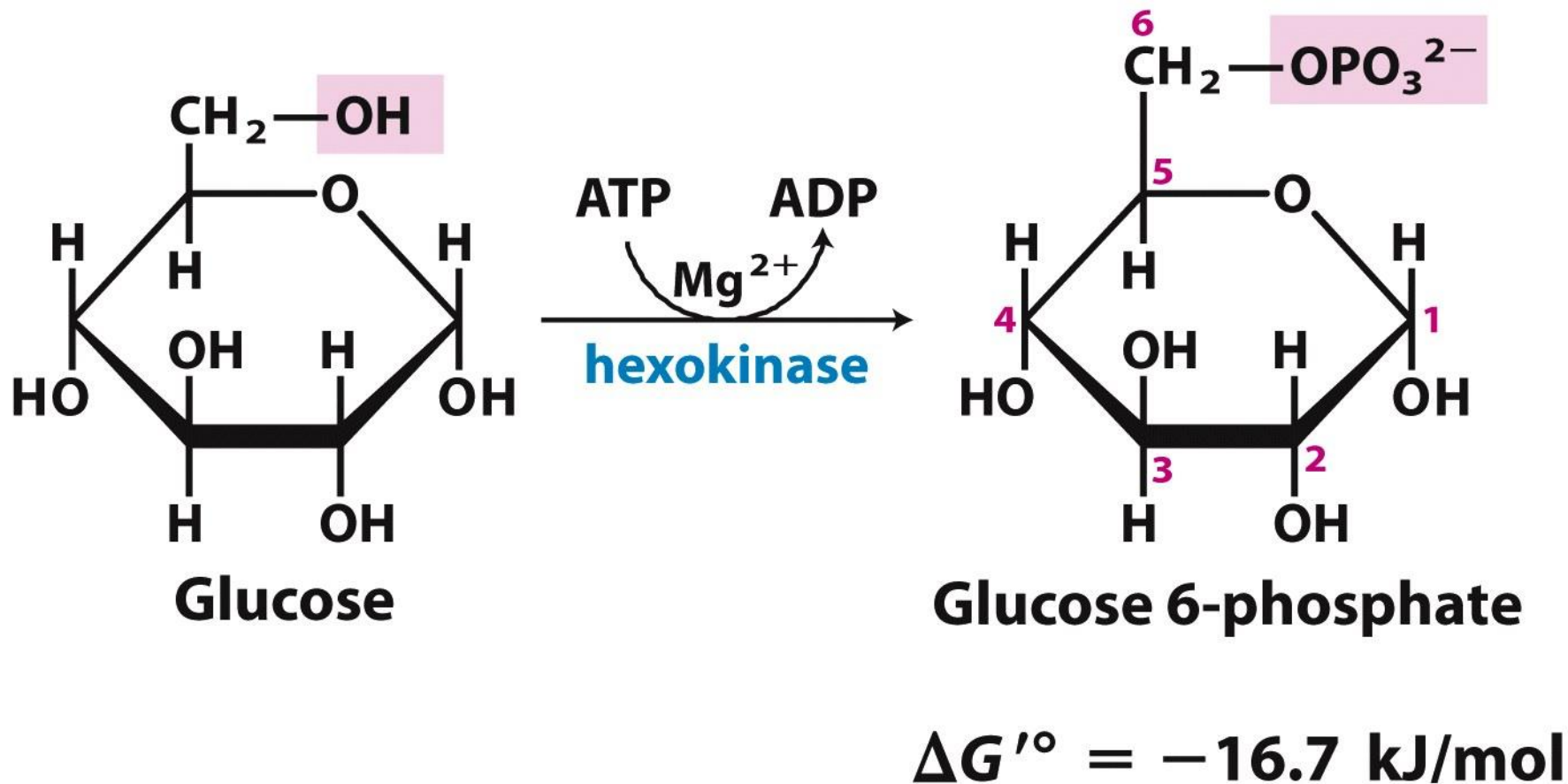
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Arthur Harden
1865–1940



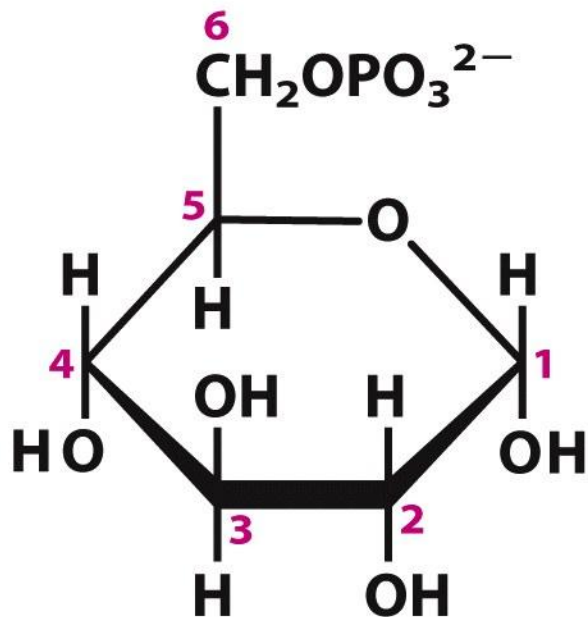
William Young
1878–1942



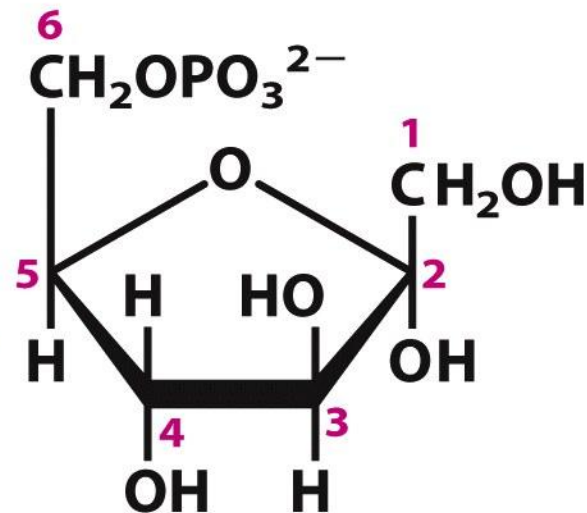
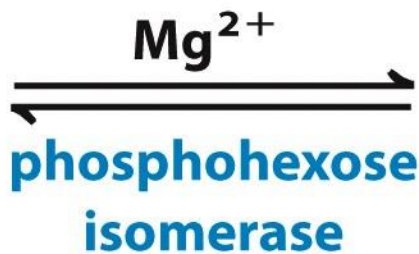
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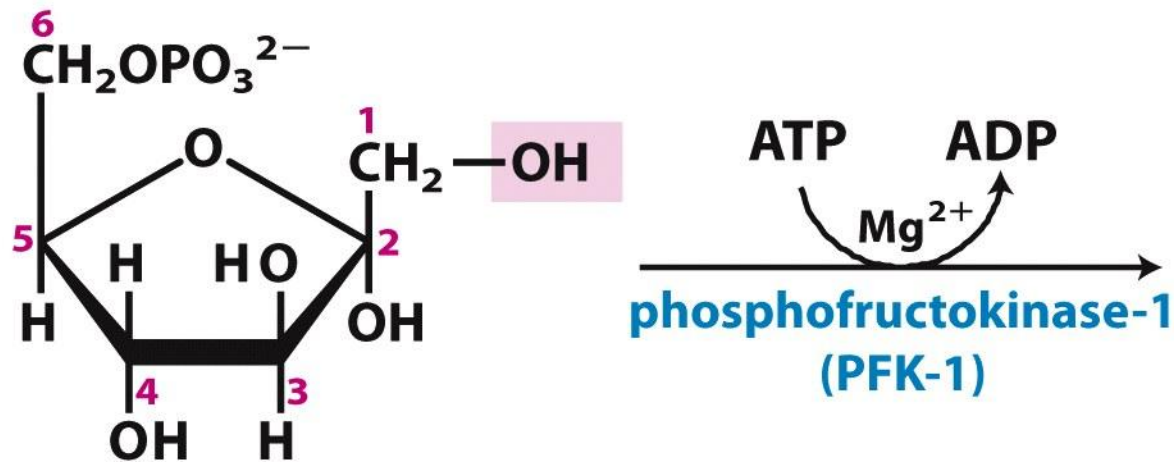


Glucose 6-phosphate

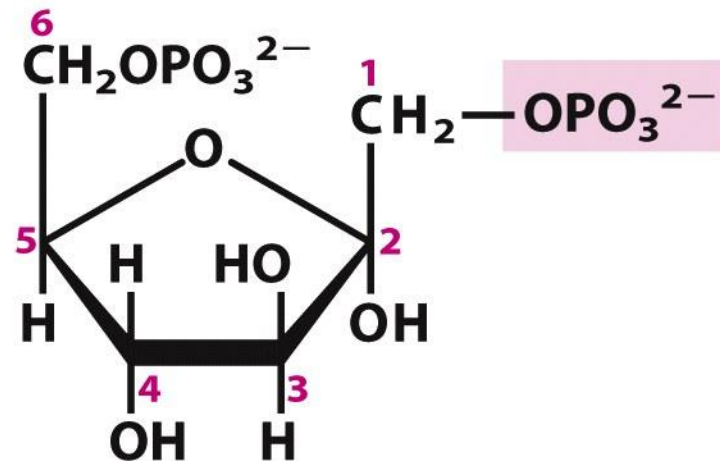


Fructose 6-phosphate

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



Fructose 6-phosphate



Fructose 1,6-bisphosphate

$$\Delta G'^{\circ} = -14.2 \text{ kJ/mol}$$

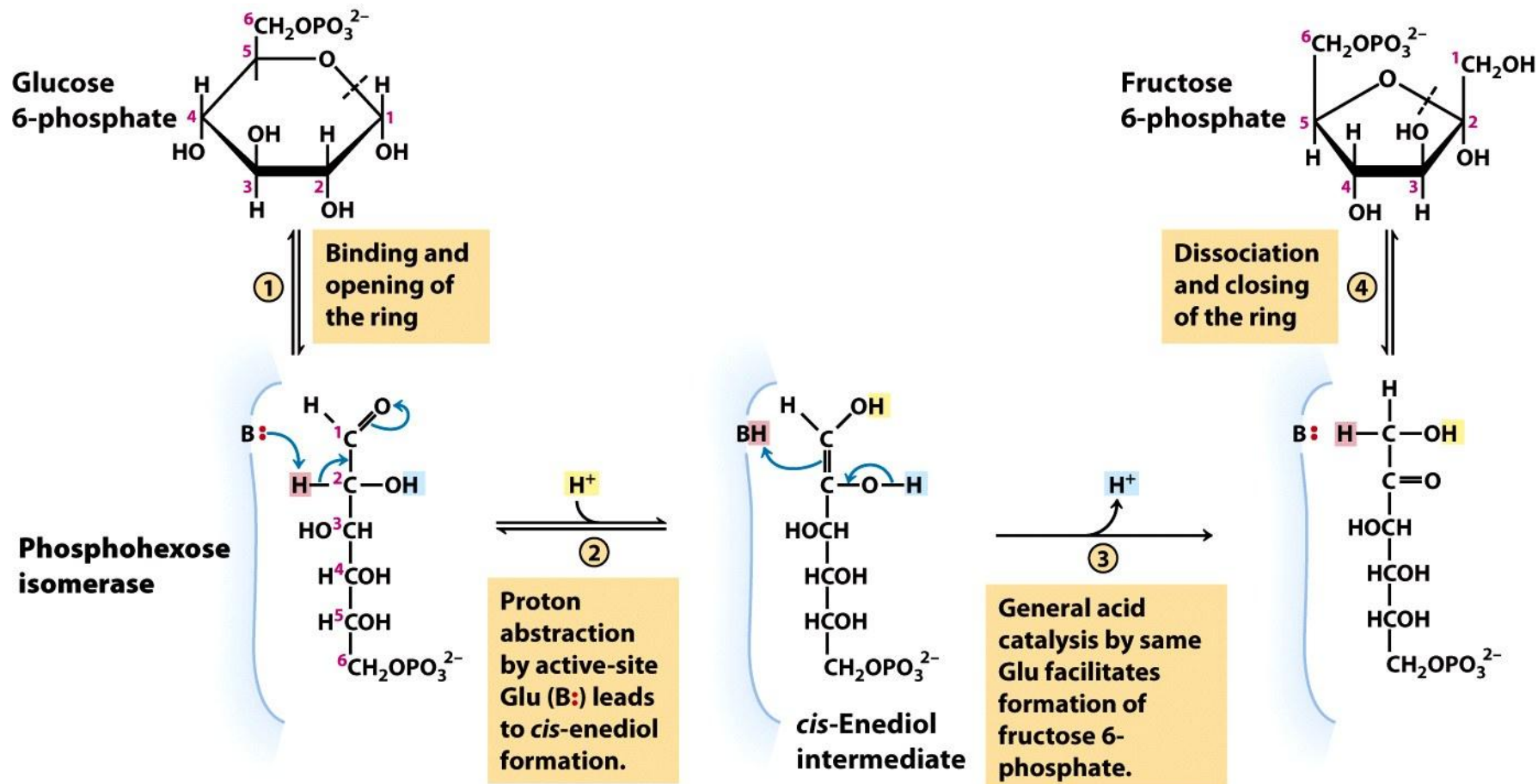
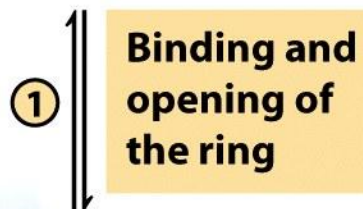
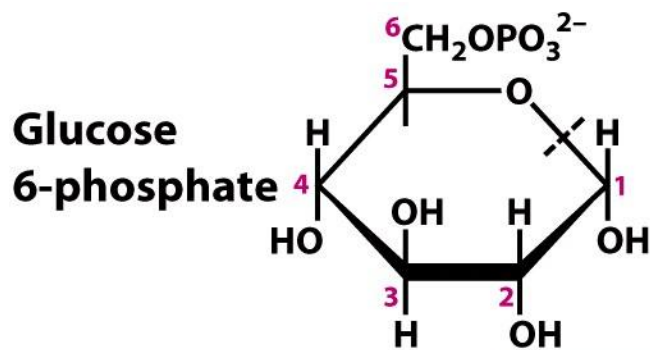


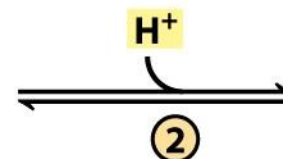
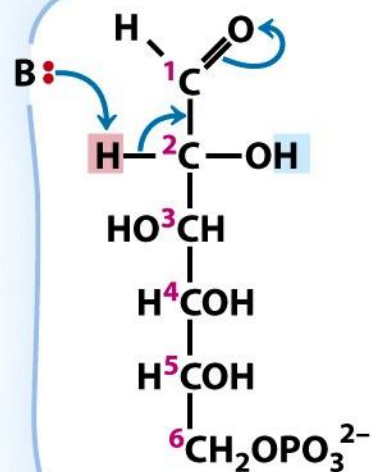
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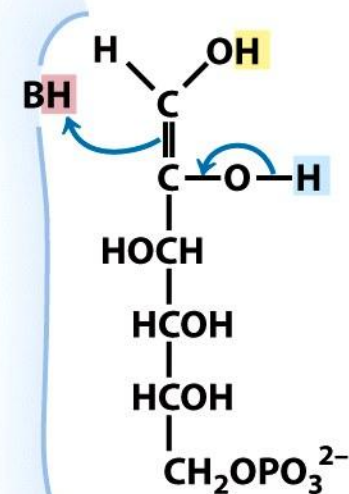
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Phosphohexose isomerase



Proton abstraction by active-site Glu (B:) leads to *cis*-enediol formation.



cis-Enediol intermediate

Figure 14-4 part 1

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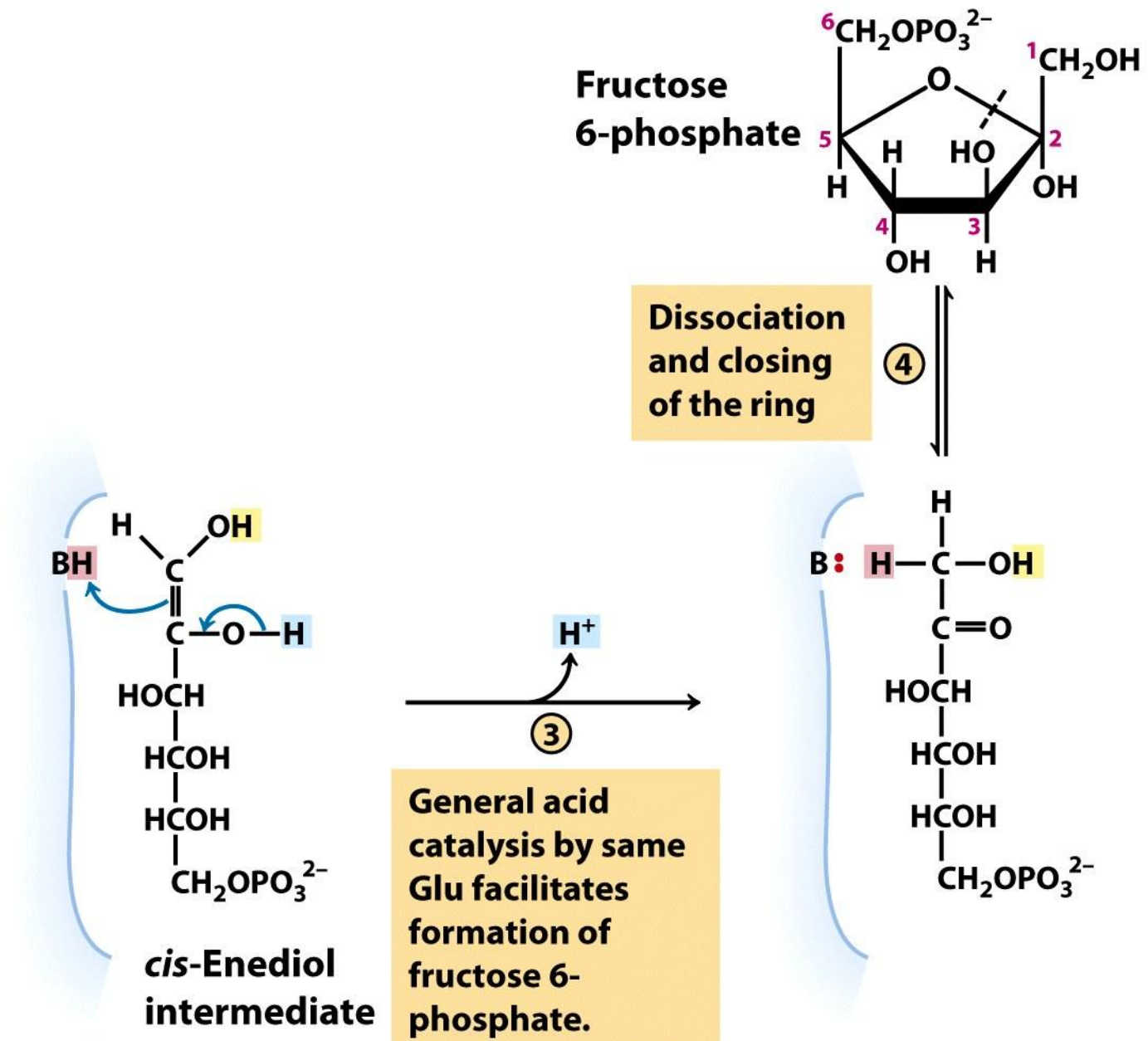
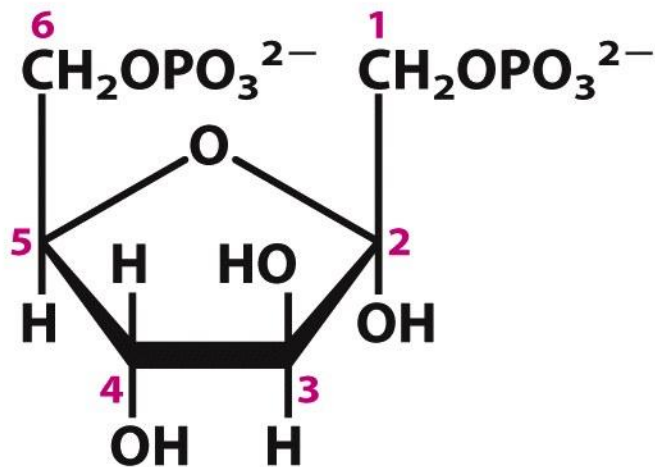


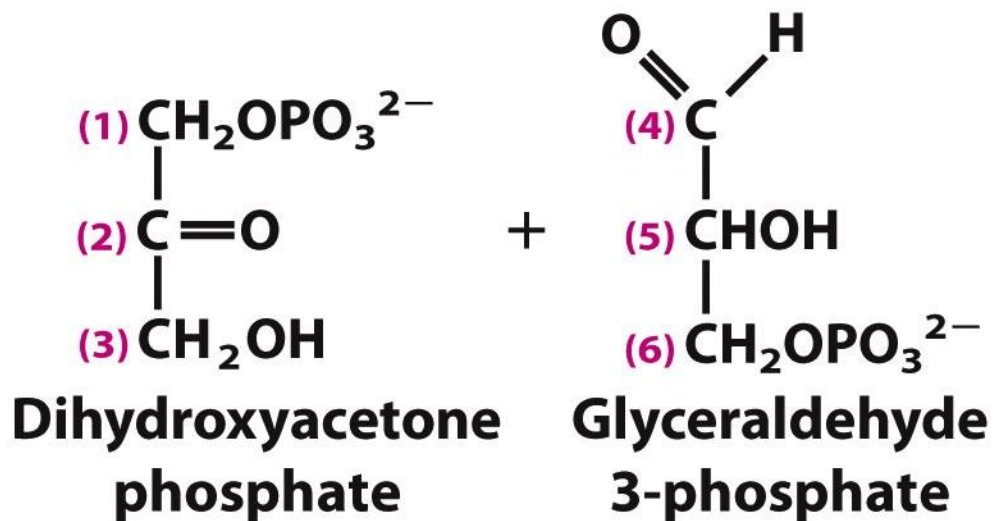
Figure 14-4 part 2

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Fructose 1,6-bisphosphate



$$\Delta G'^{\circ} = 23.8 \text{ kJ/mol}$$

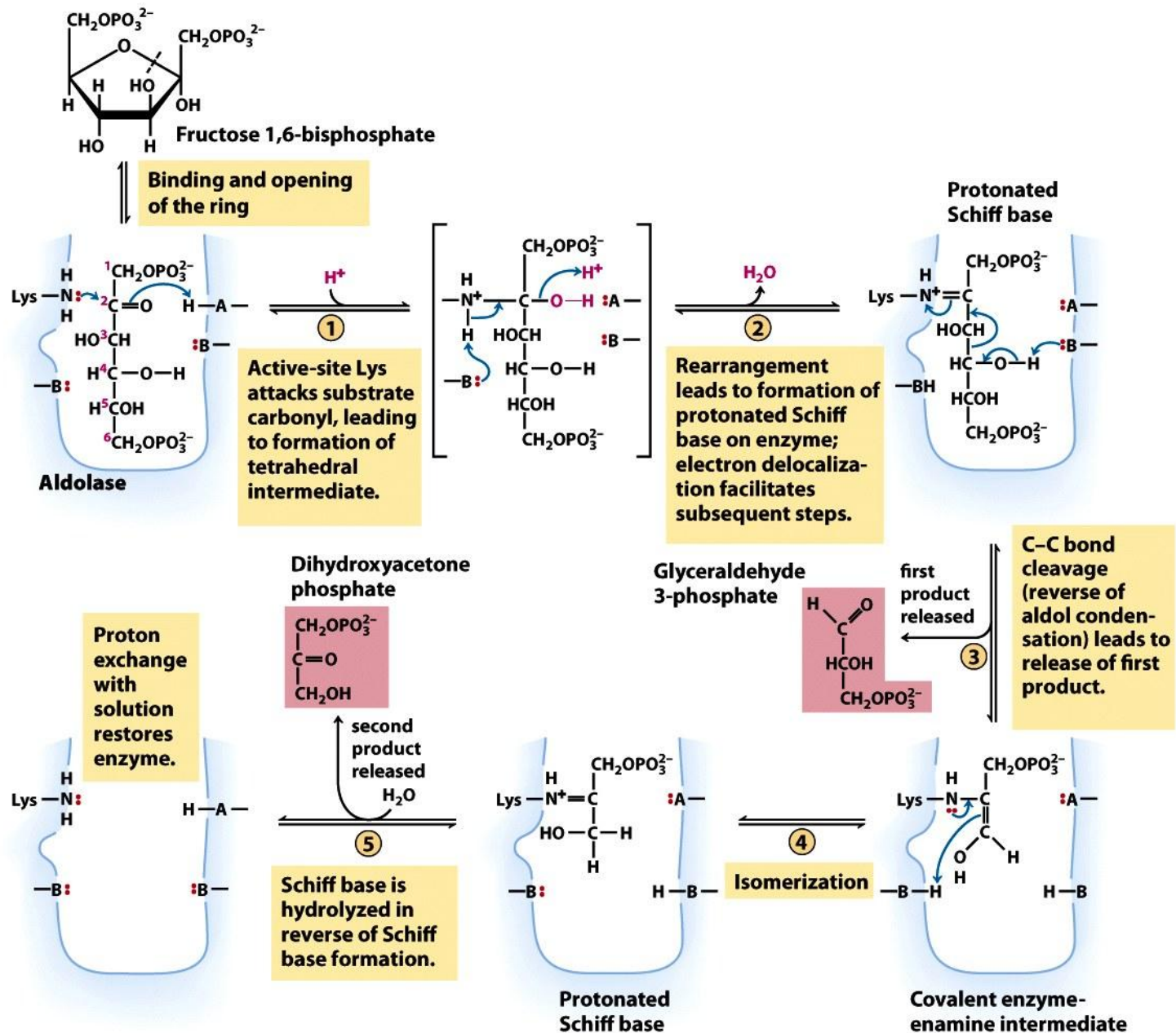


Figure 14-5

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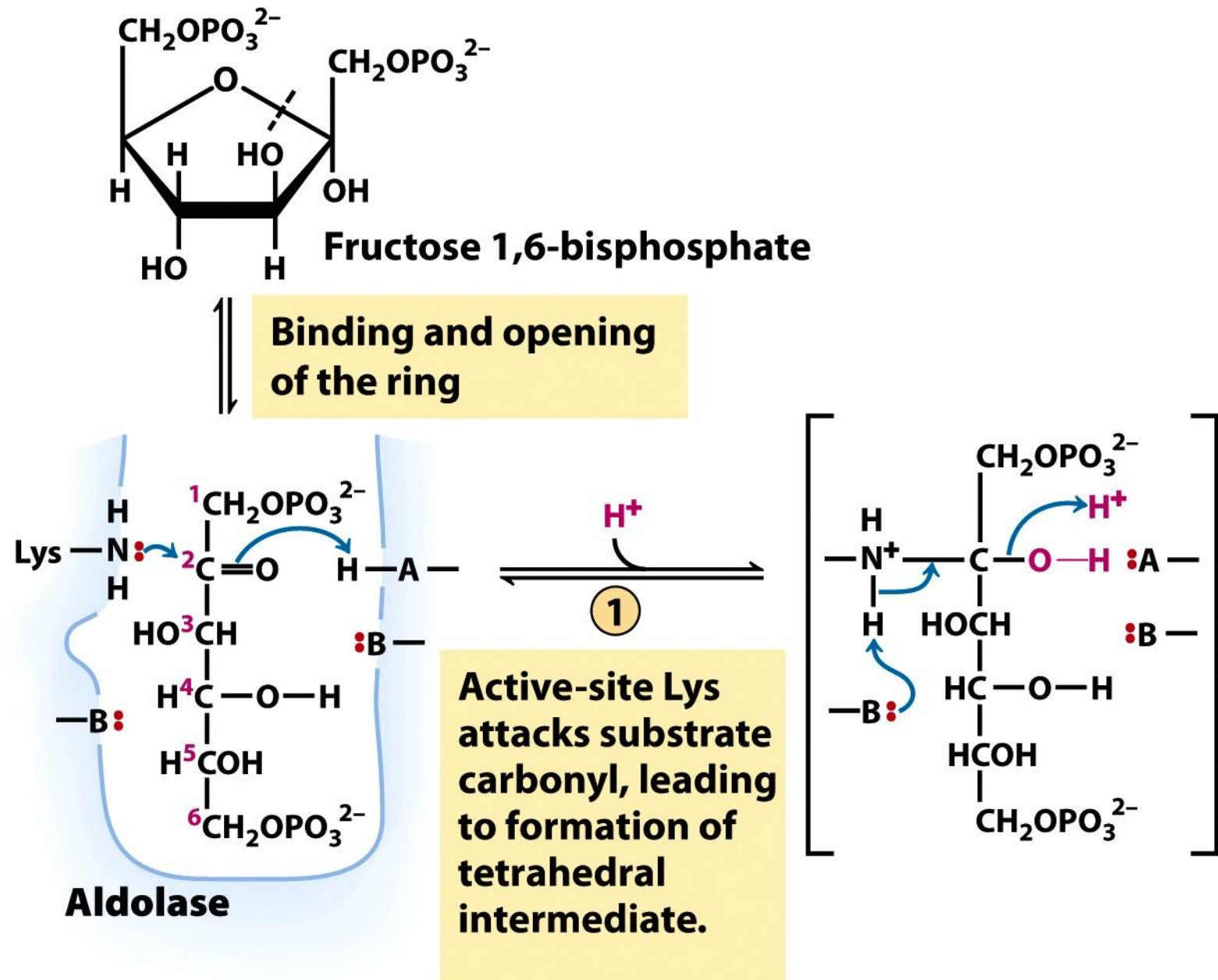
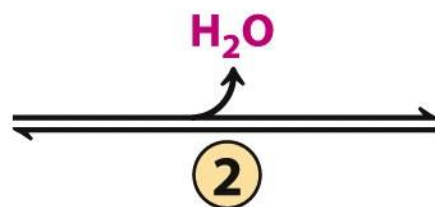
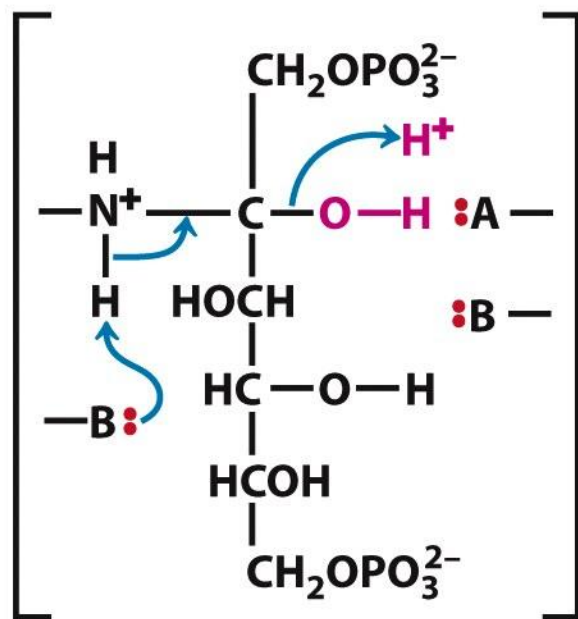


Figure 14-5 part 1

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Rearrangement leads to formation of protonated Schiff base on enzyme; electron delocalization facilitates subsequent steps.

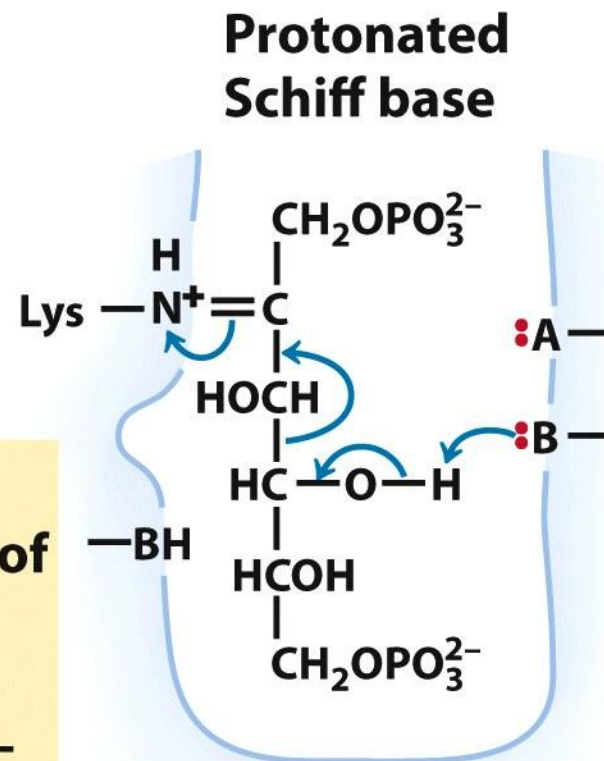
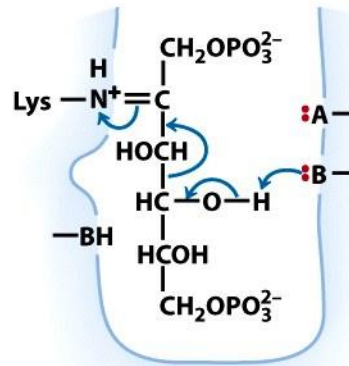


Figure 14-5 part 2

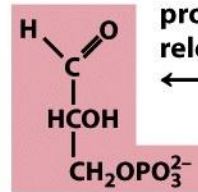
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Protonated Schiff base



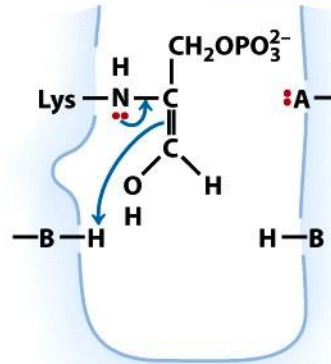
Glyceraldehyde 3-phosphate



first
product
released

3

C-C bond
cleavage
(reverse of
aldol conden-
sation) leads to
release of first
product.



Covalent enzyme- enamine intermediate

Figure 14-5 part 3

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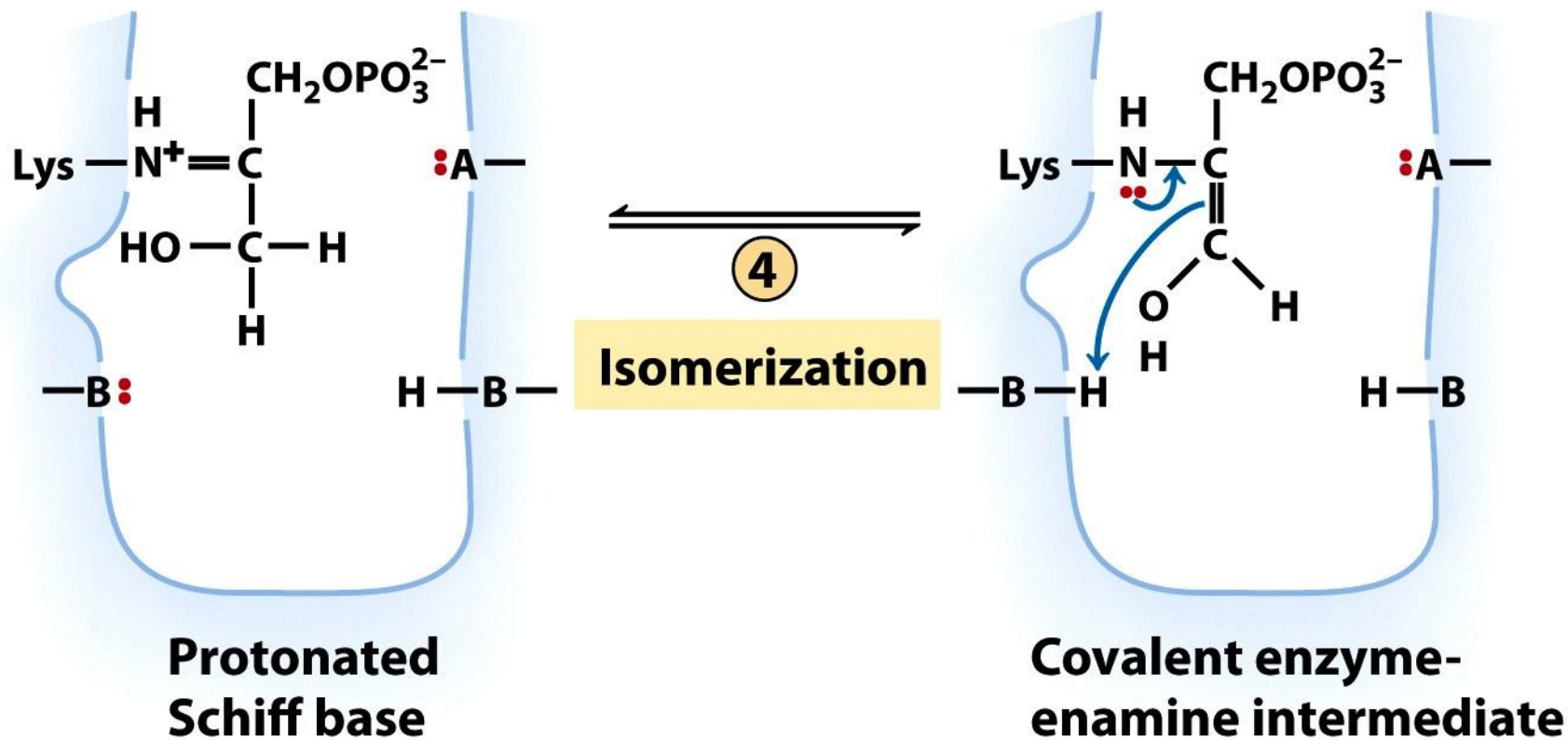


Figure 14-5 part 4
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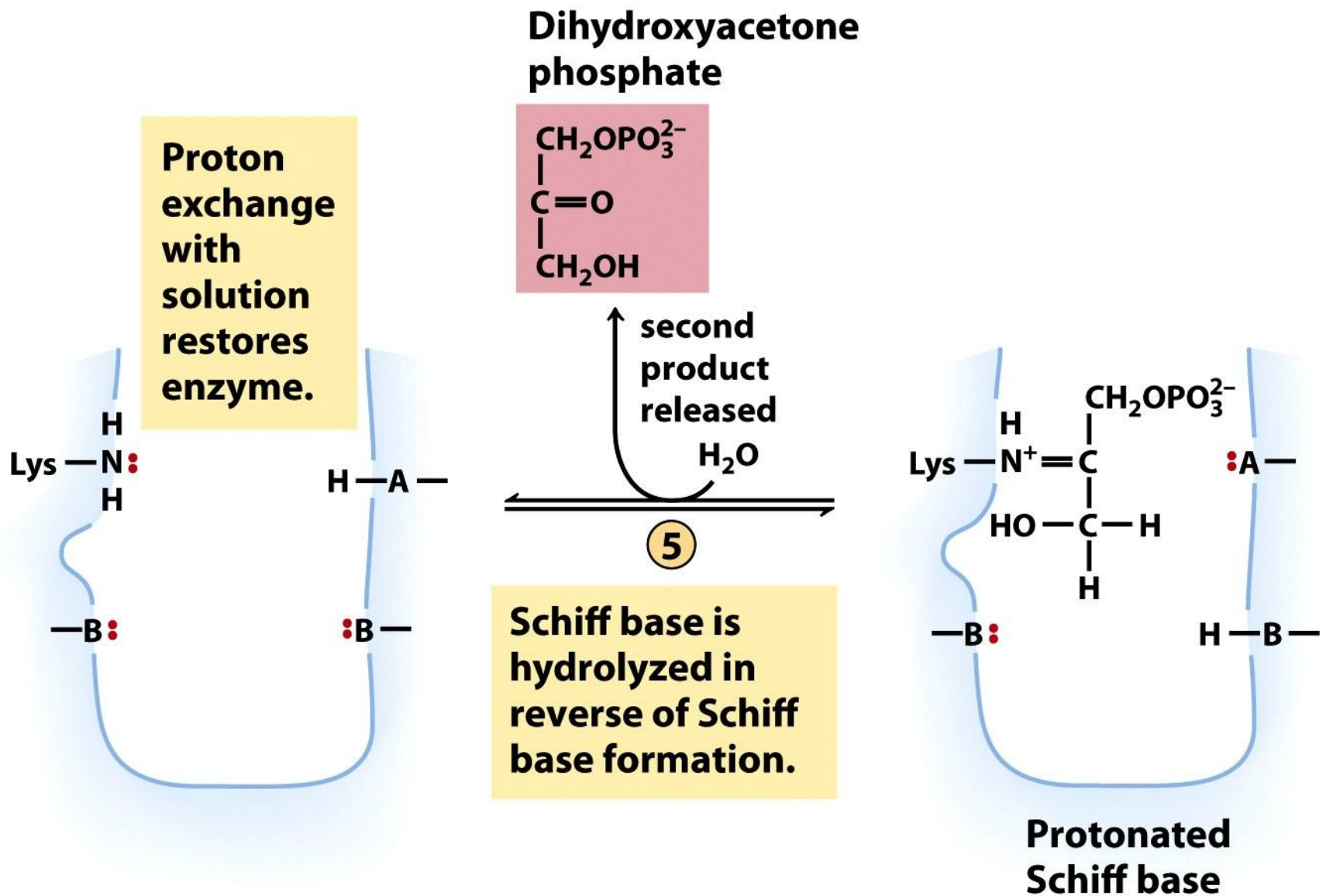
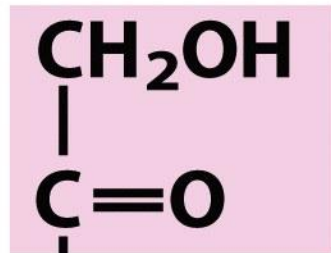
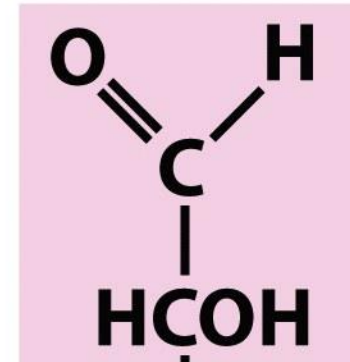


Figure 14-5 part 5
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**Dihydroxyacetone
phosphate**



**Glyceraldehyde
3-phosphate**

$$\Delta G'^{\circ} = 7.5 \text{ kJ/mol}$$

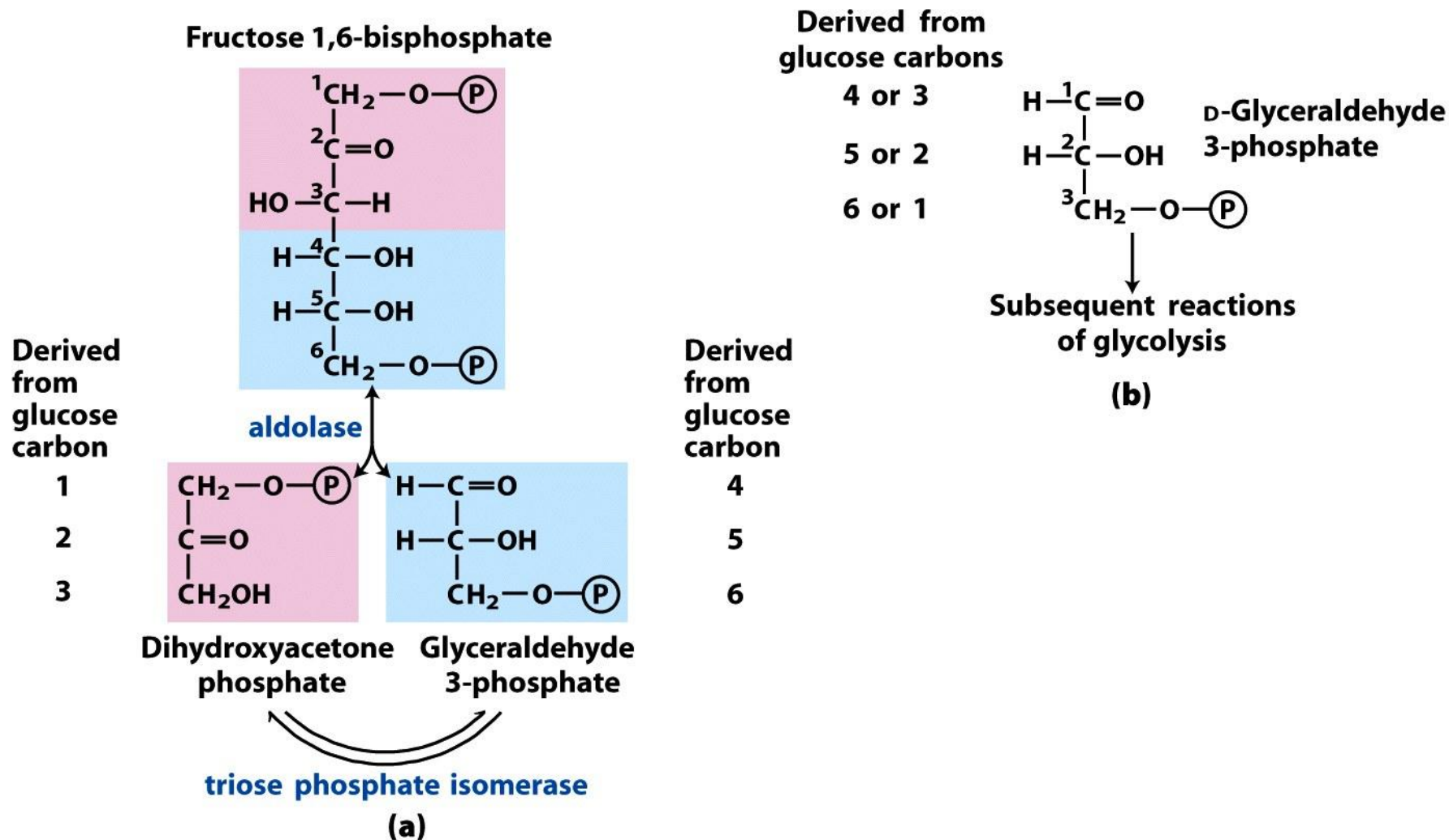


Figure 14-6

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Fructose 1,6-bisphosphate

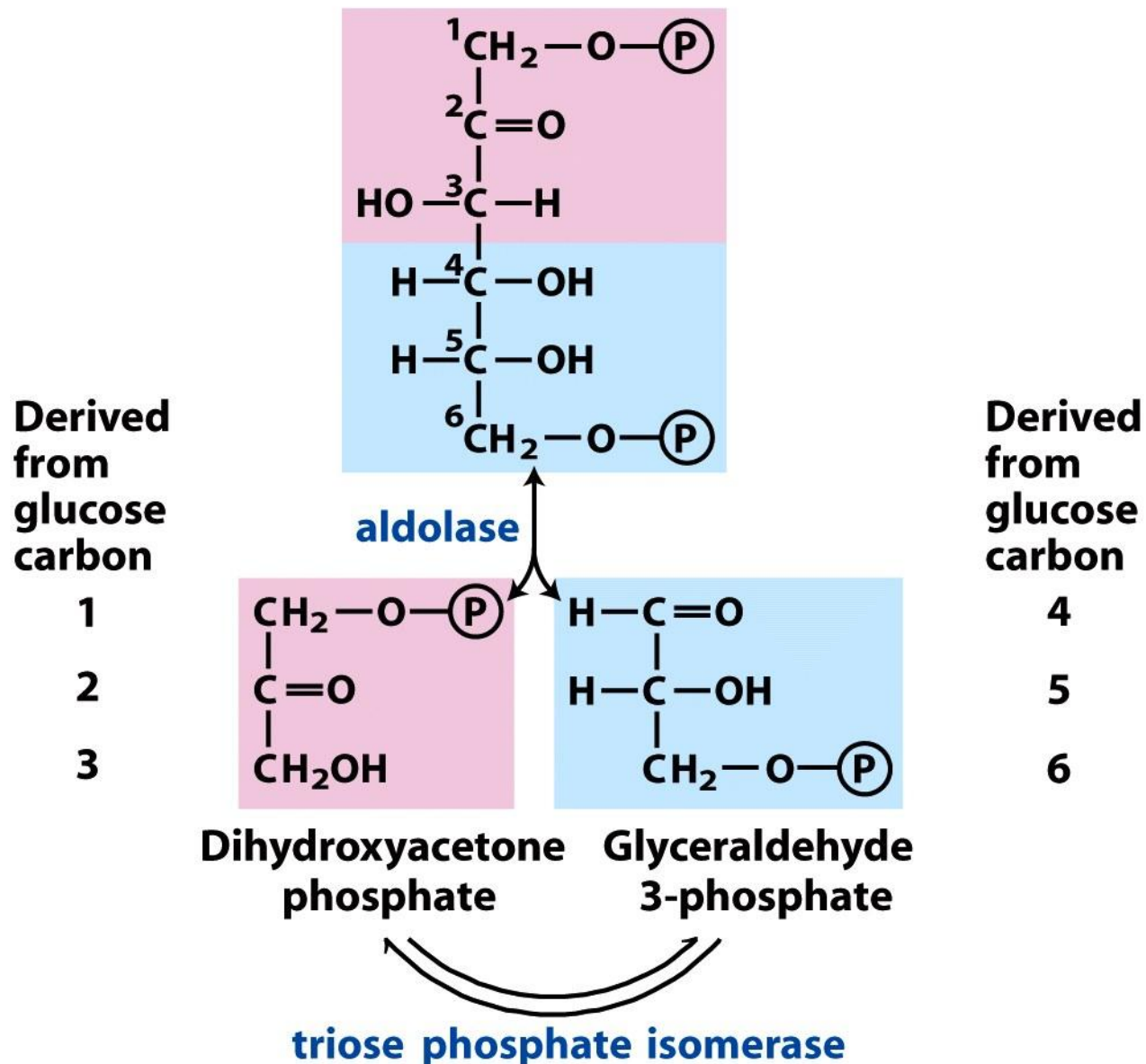


Figure 14-6a

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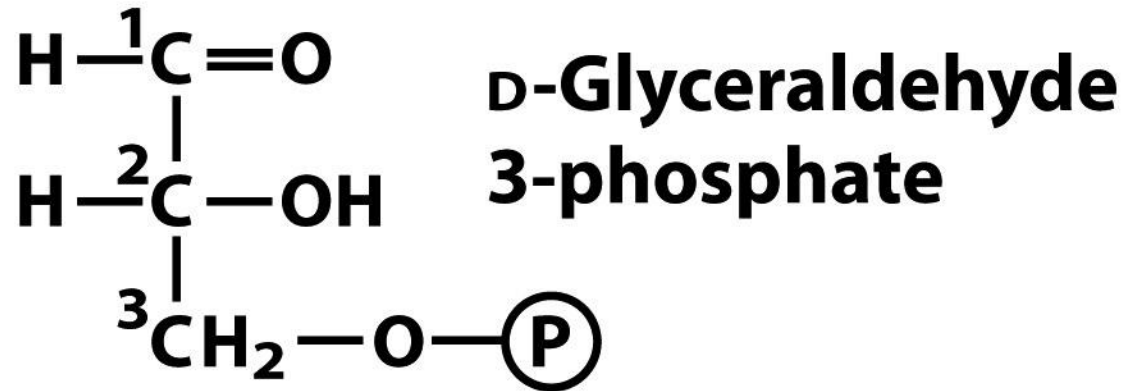
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**Derived from
glucose carbons**

4 or 3

5 or 2

6 or 1

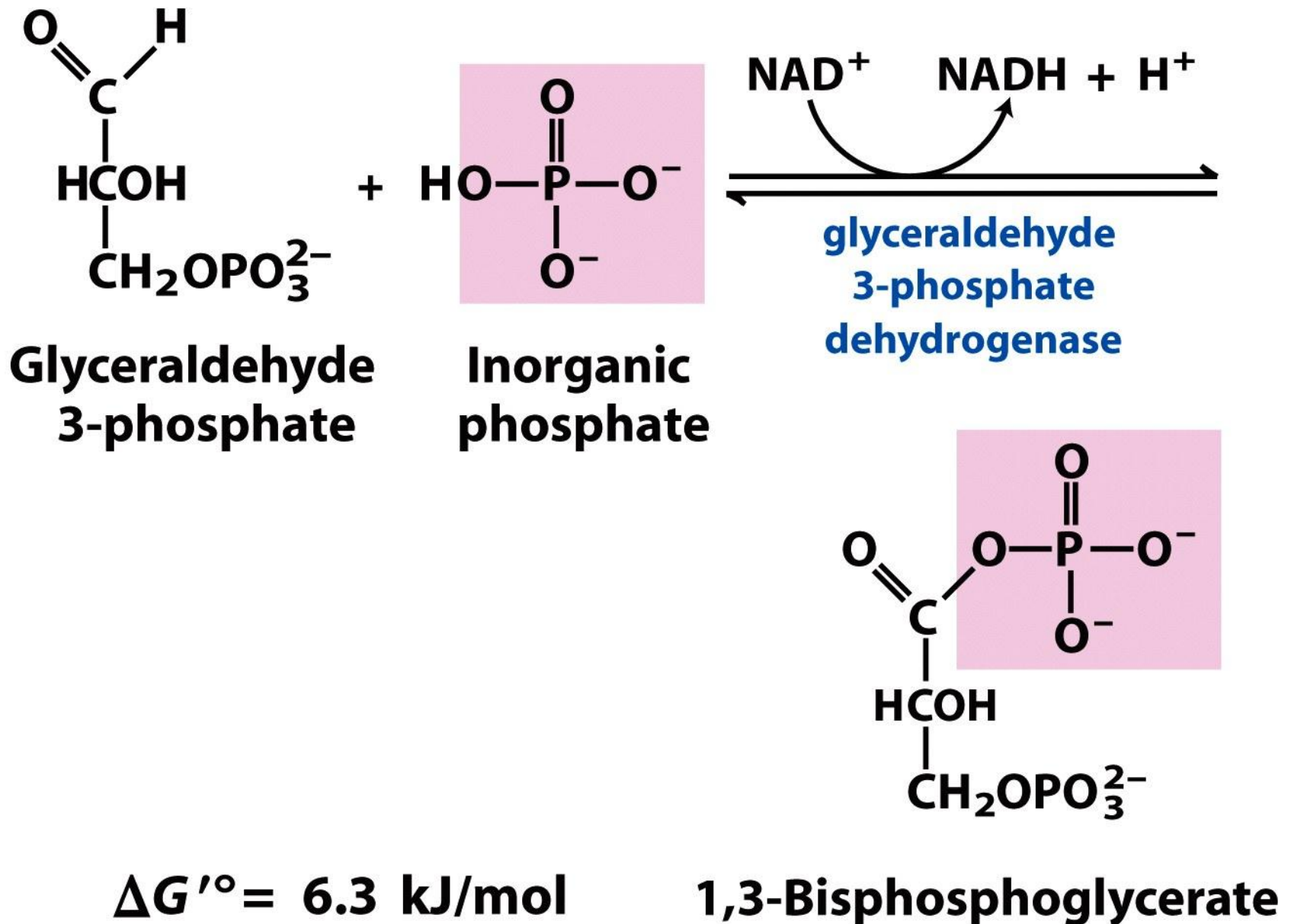


↓
**Subsequent reactions
of glycolysis**

Figure 14-6b

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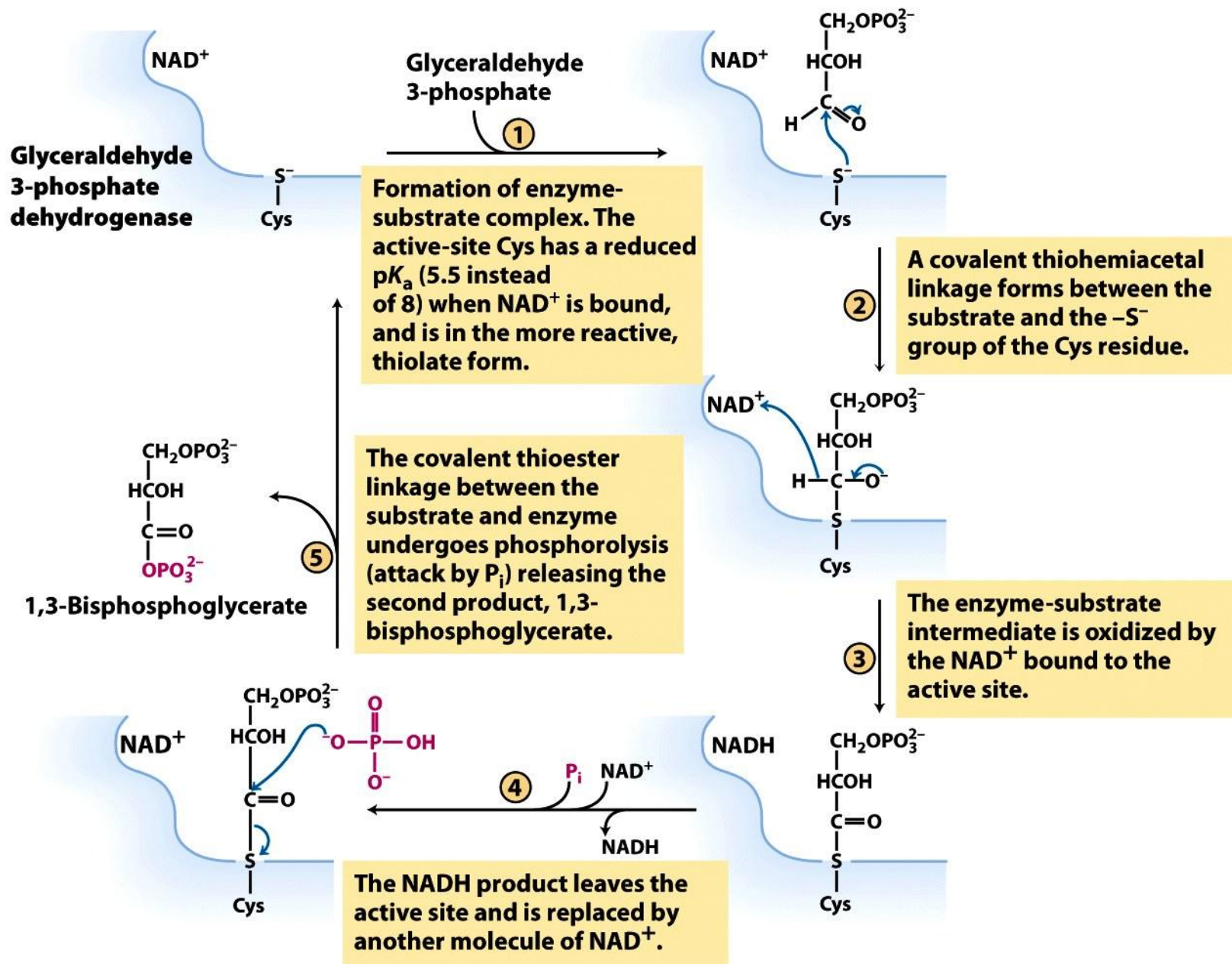


Figure 14-7

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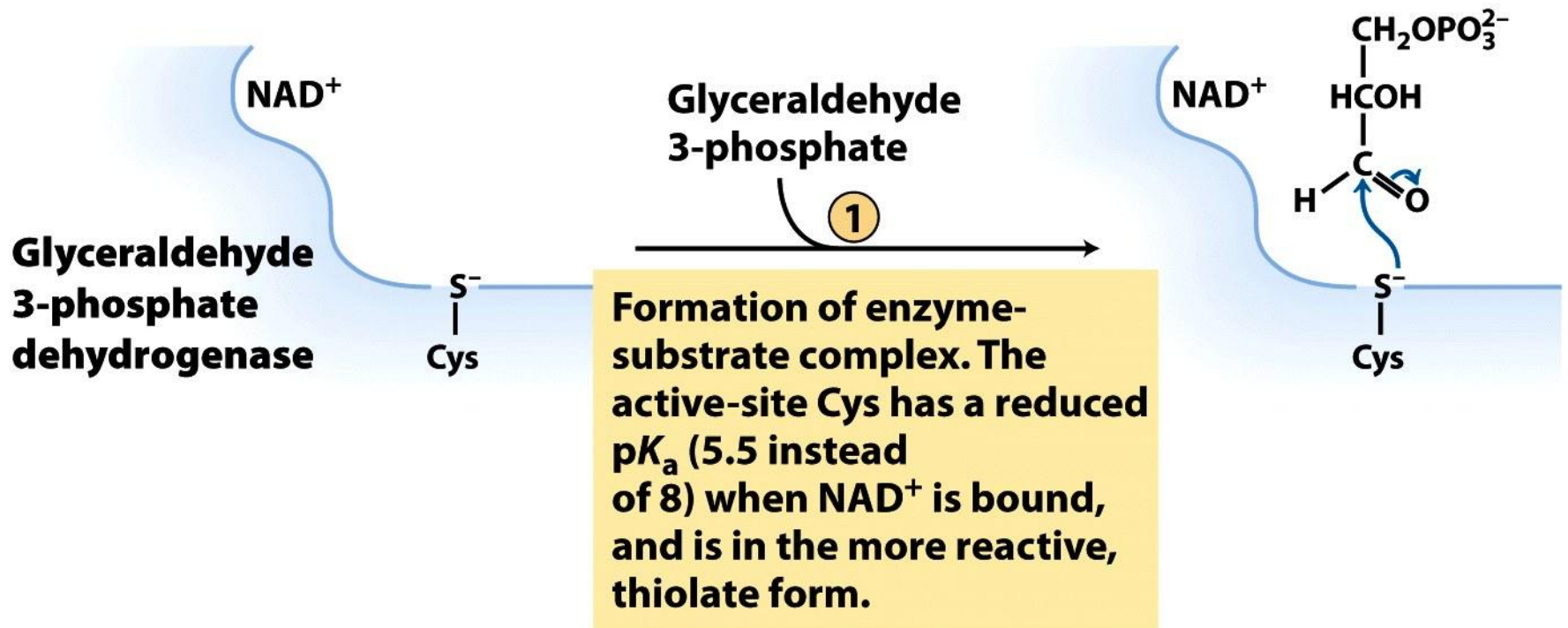


Figure 14-7 part 1

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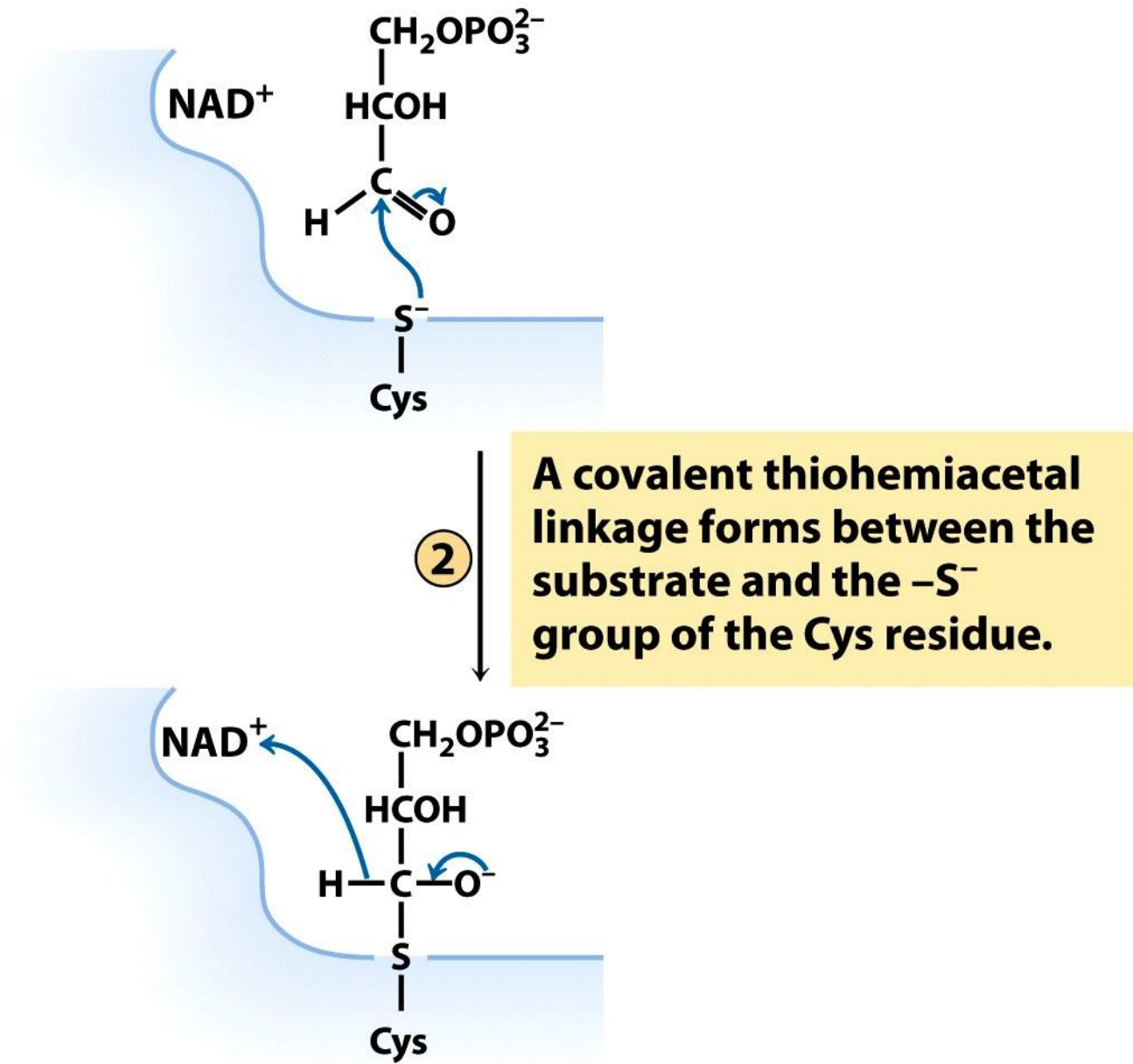


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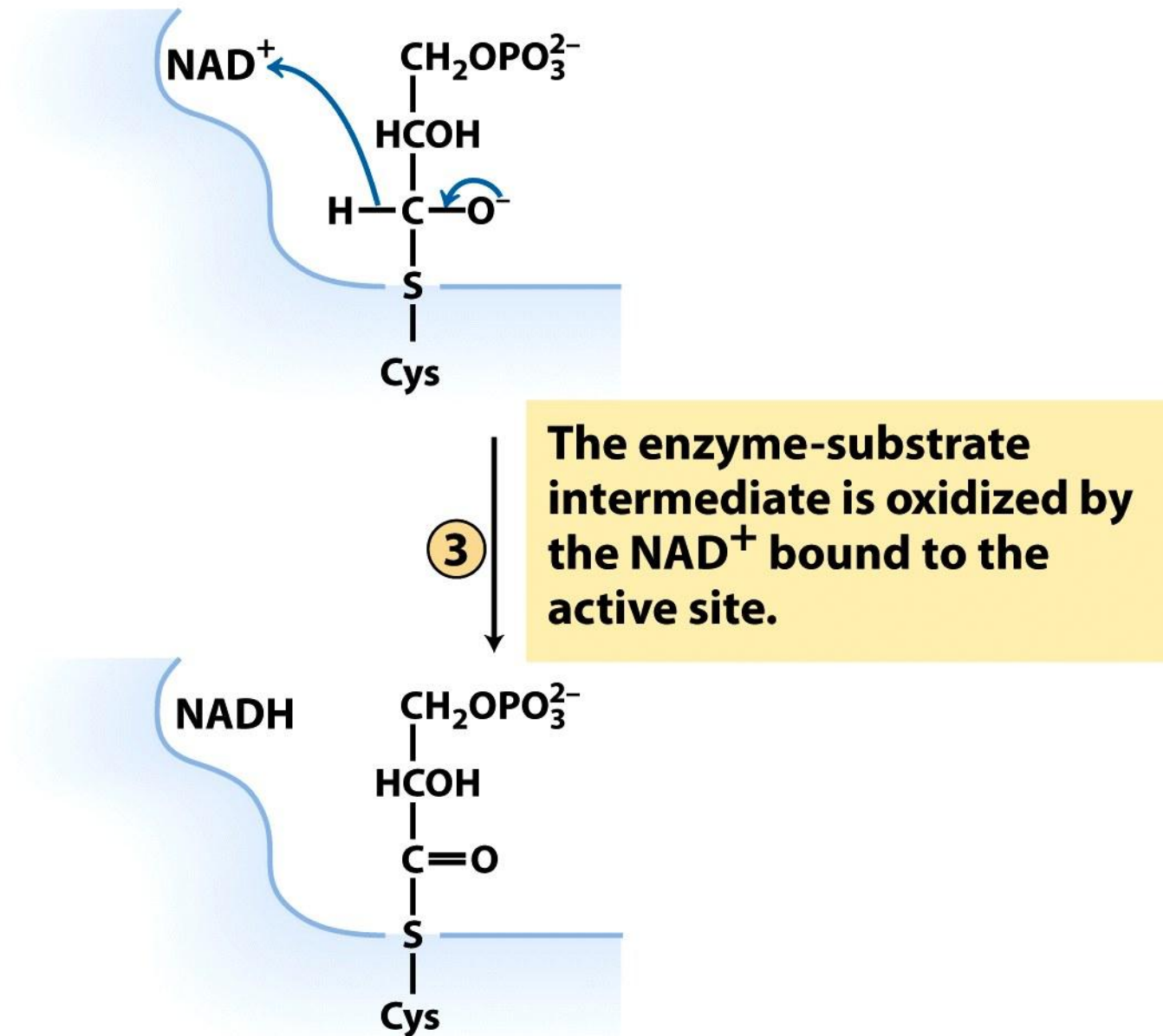


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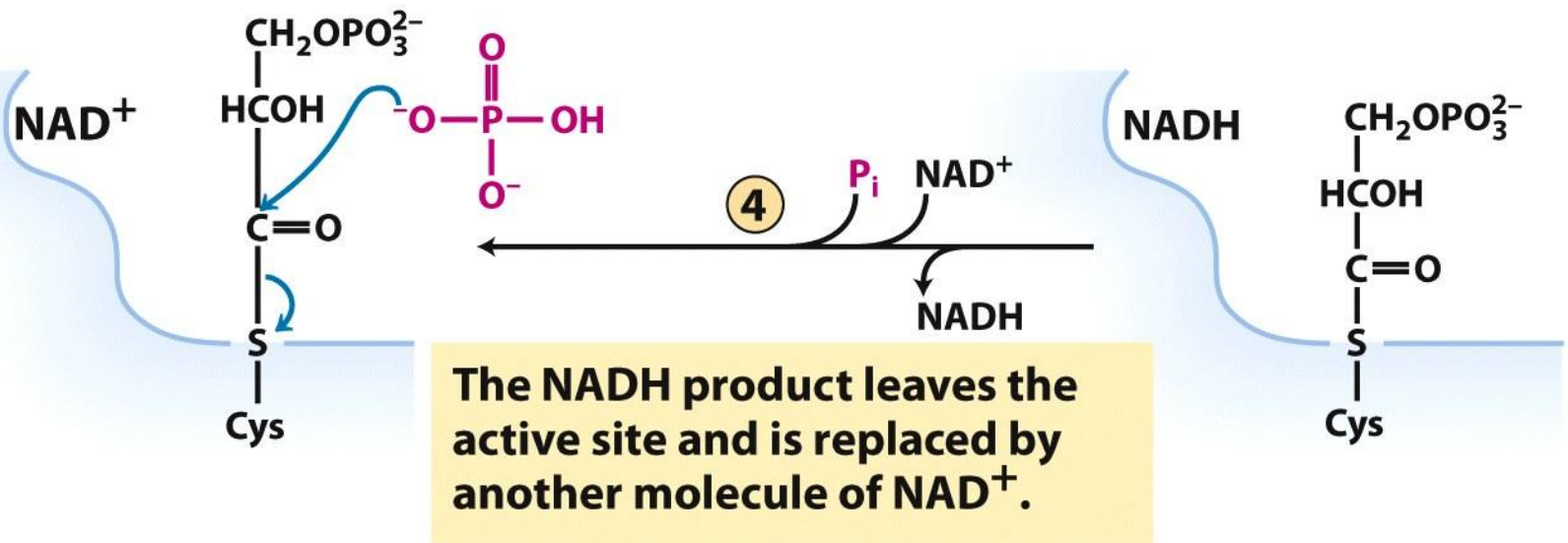


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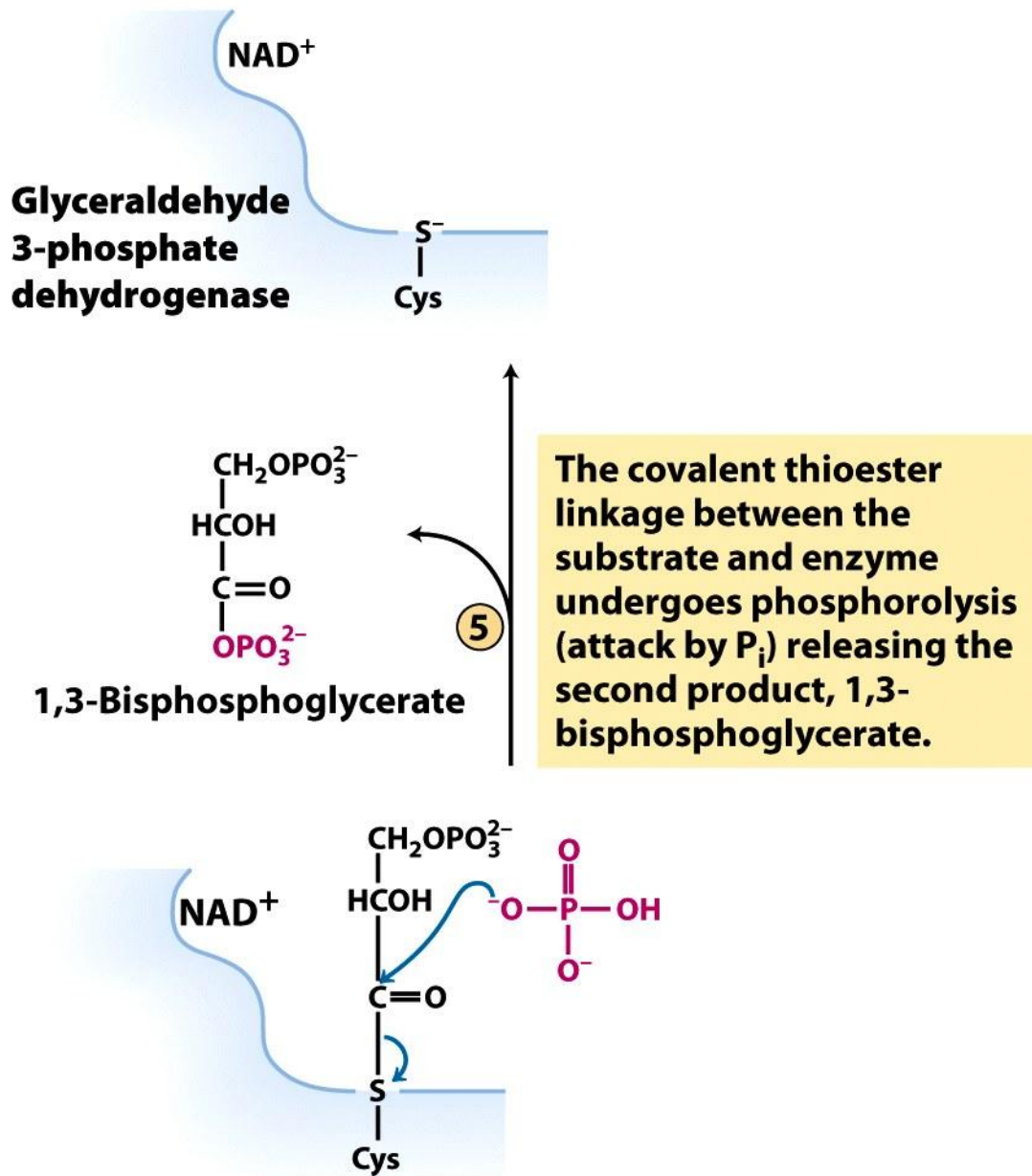
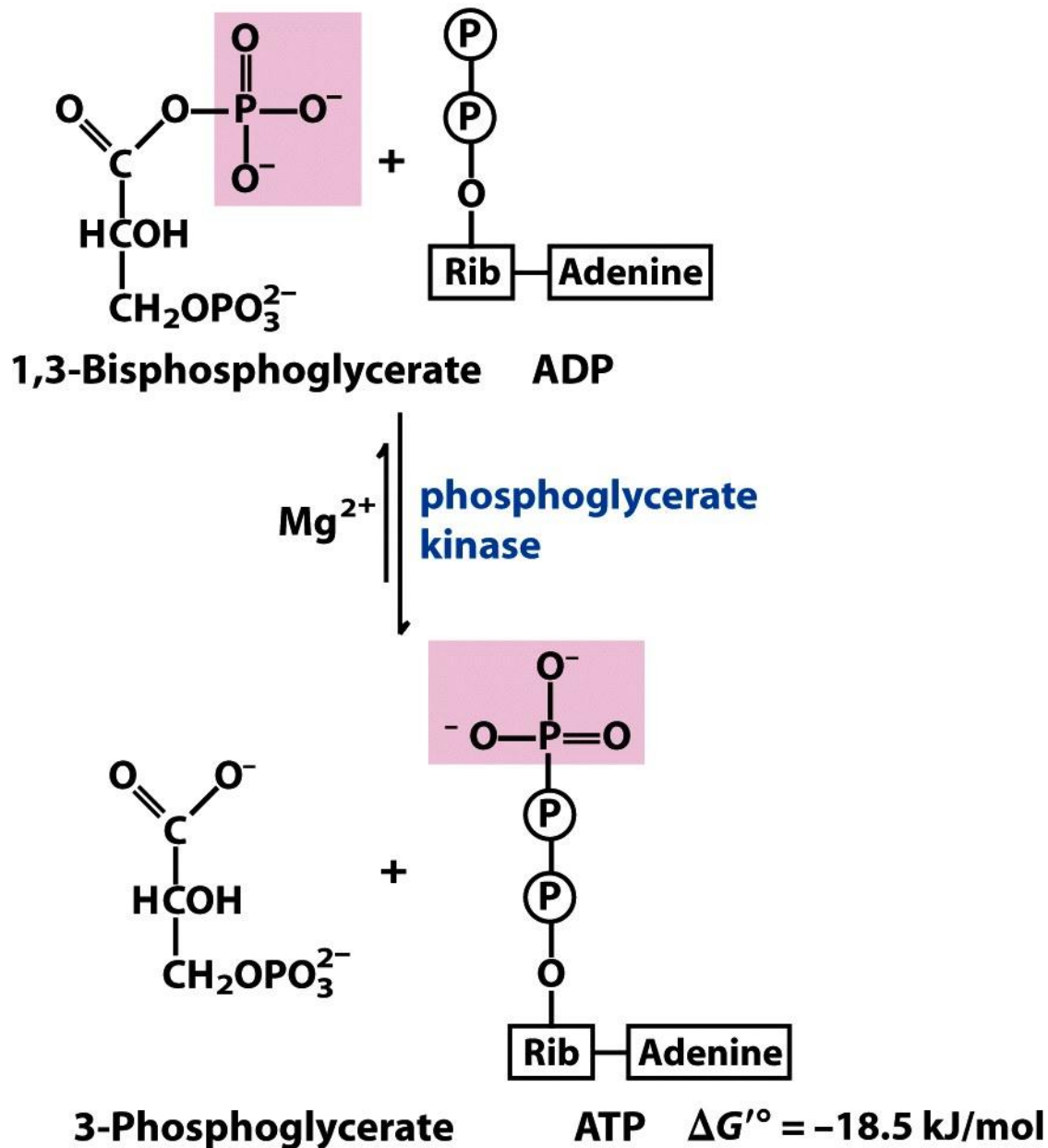
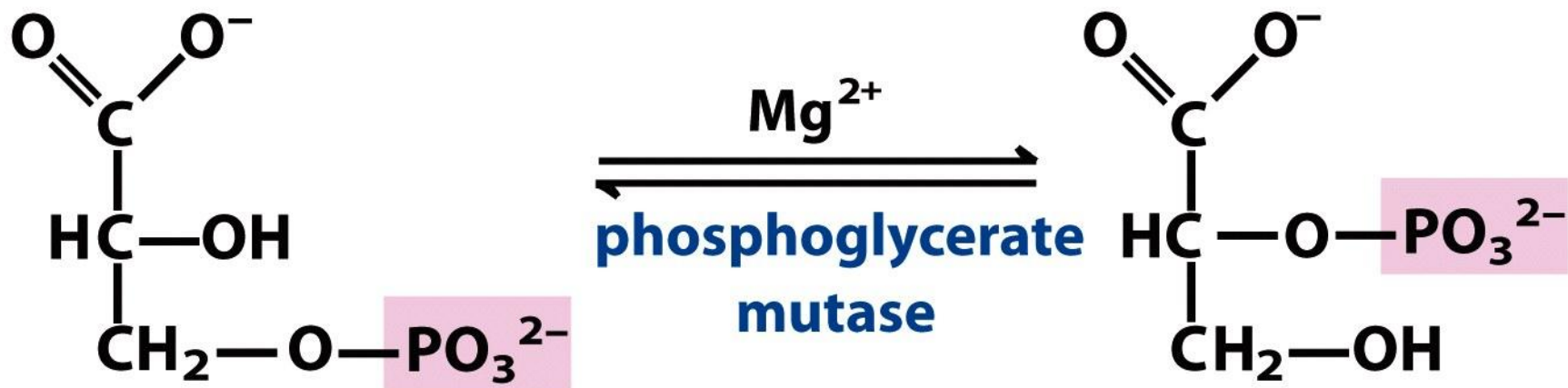


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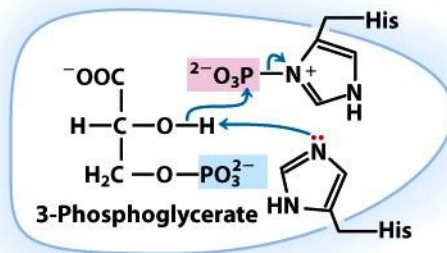


3-Phosphoglycerate

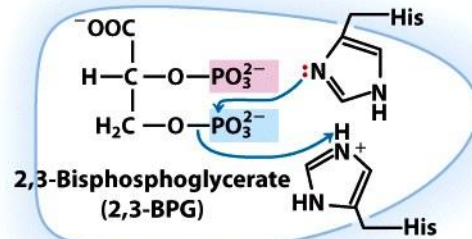
2-Phosphoglycerate

$$\Delta G'^{\circ} = 4.4 \text{ kJ/mol}$$

Phosphoglycerate mutase



1
Phosphoryl transfer occurs between an active-site His and C-2 (OH) of the substrate. A second active-site His acts as general base catalyst.



2
Phosphoryl transfer from C-3 of the substrate to the first active-site His. The second active-site His acts as general acid catalyst.

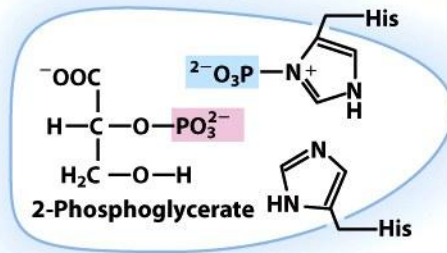
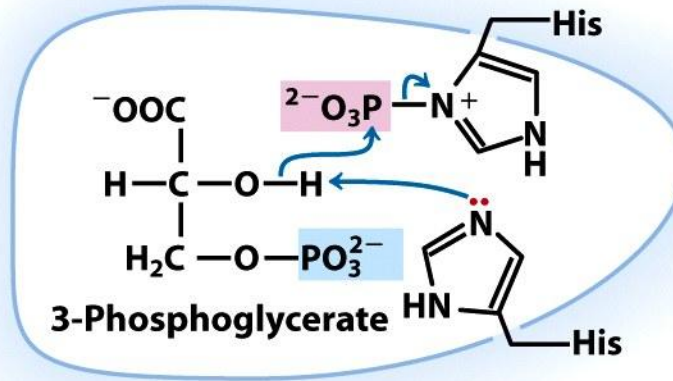


Figure 14-8

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Phosphoglycerate mutase



1

Phosphoryl transfer occurs between an active-site His and C-2 (OH) of the substrate. A second active-site His acts as general base catalyst.

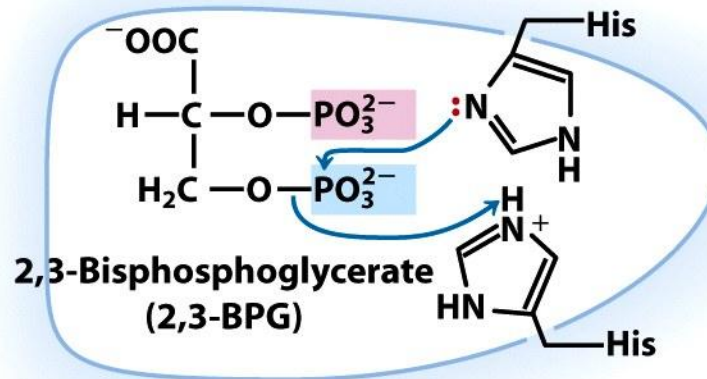
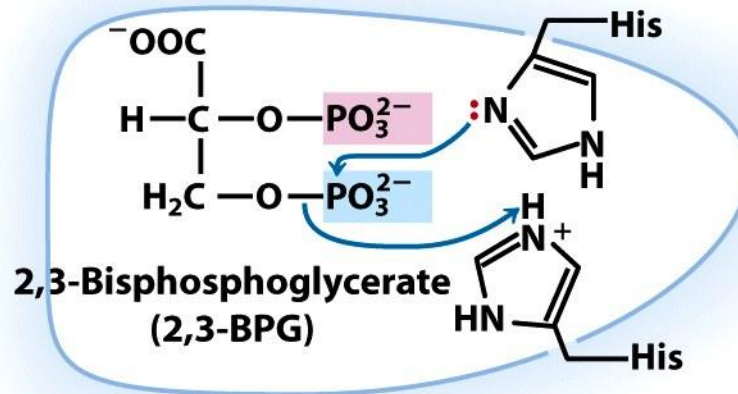


Figure 14-8 part 1

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2

Phosphoryl transfer from C-3 of the substrate to the first active-site His. The second active-site His acts as general acid catalyst.

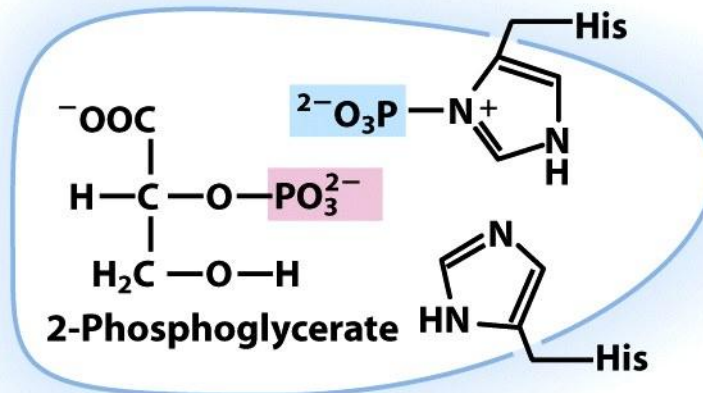
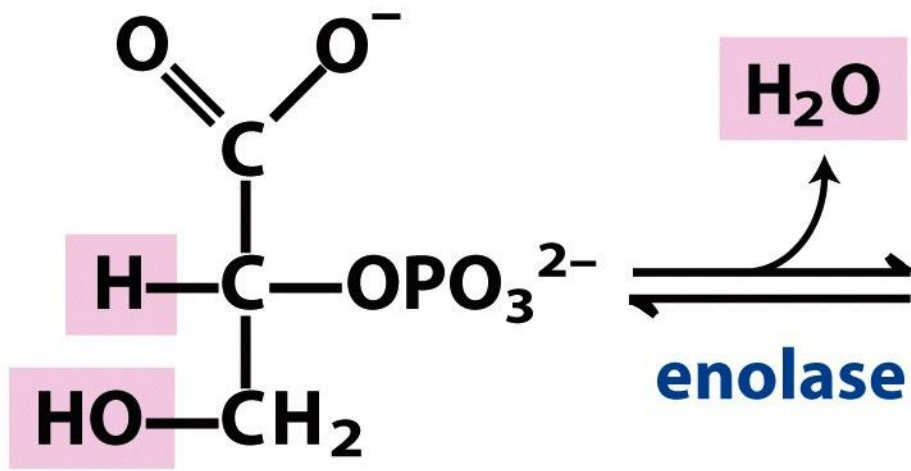


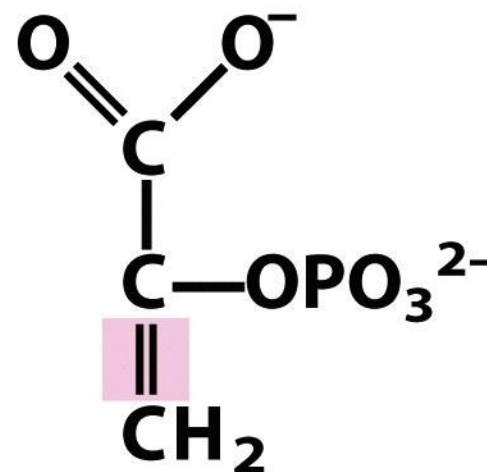
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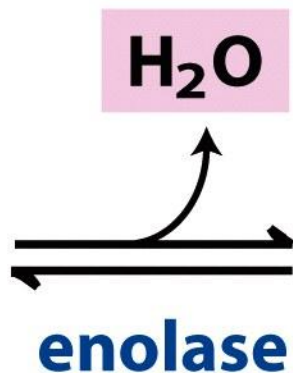
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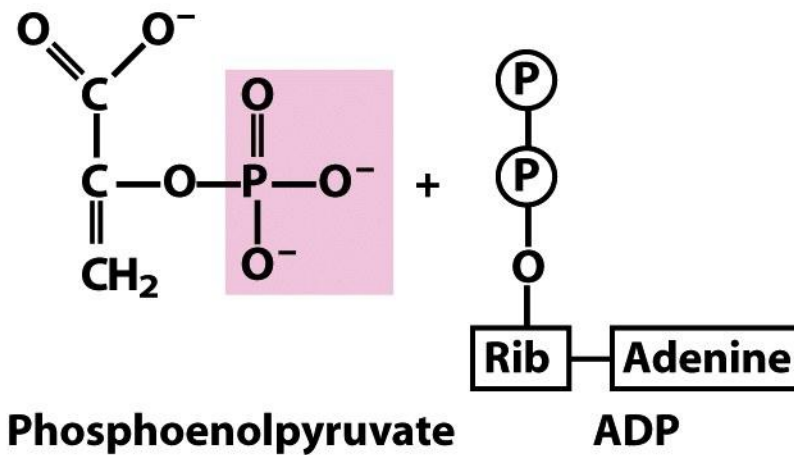
2-Phosphoglycerate



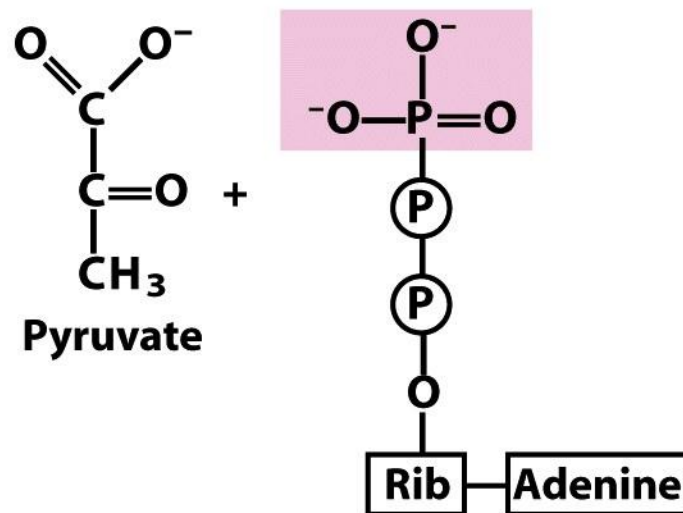
Phosphoenolpyruvate



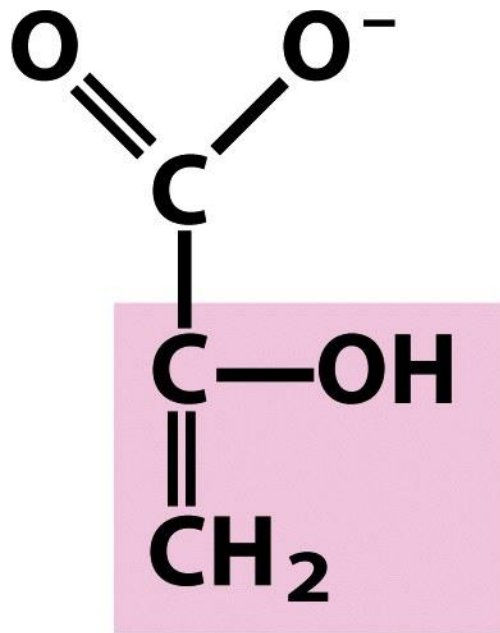
$$\Delta G'^{\circ} = 7.5 \text{ kJ/mol}$$



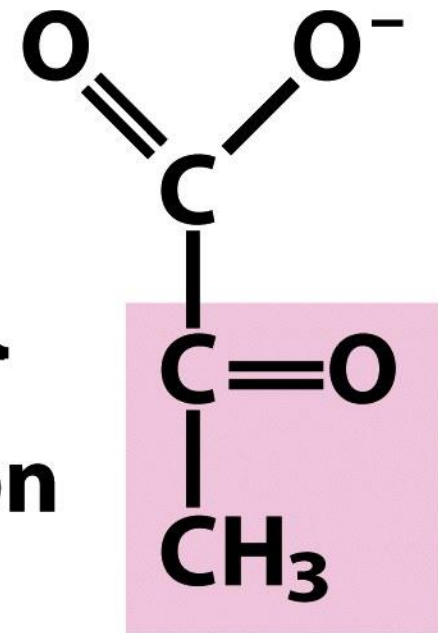
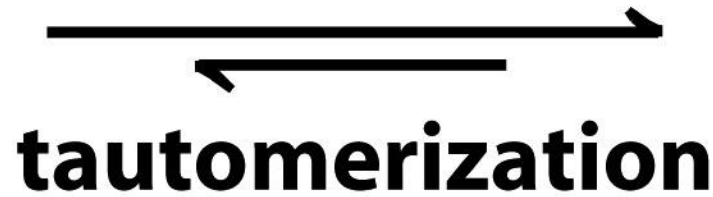
$\text{Mg}^{2+}, \text{K}^{+}$ **pyruvate kinase**



$$\Delta G'^{\circ} = -31.4 \text{ kJ/mol}$$



**Pyruvate
(enol form)**



**Pyruvate
(keto form)**

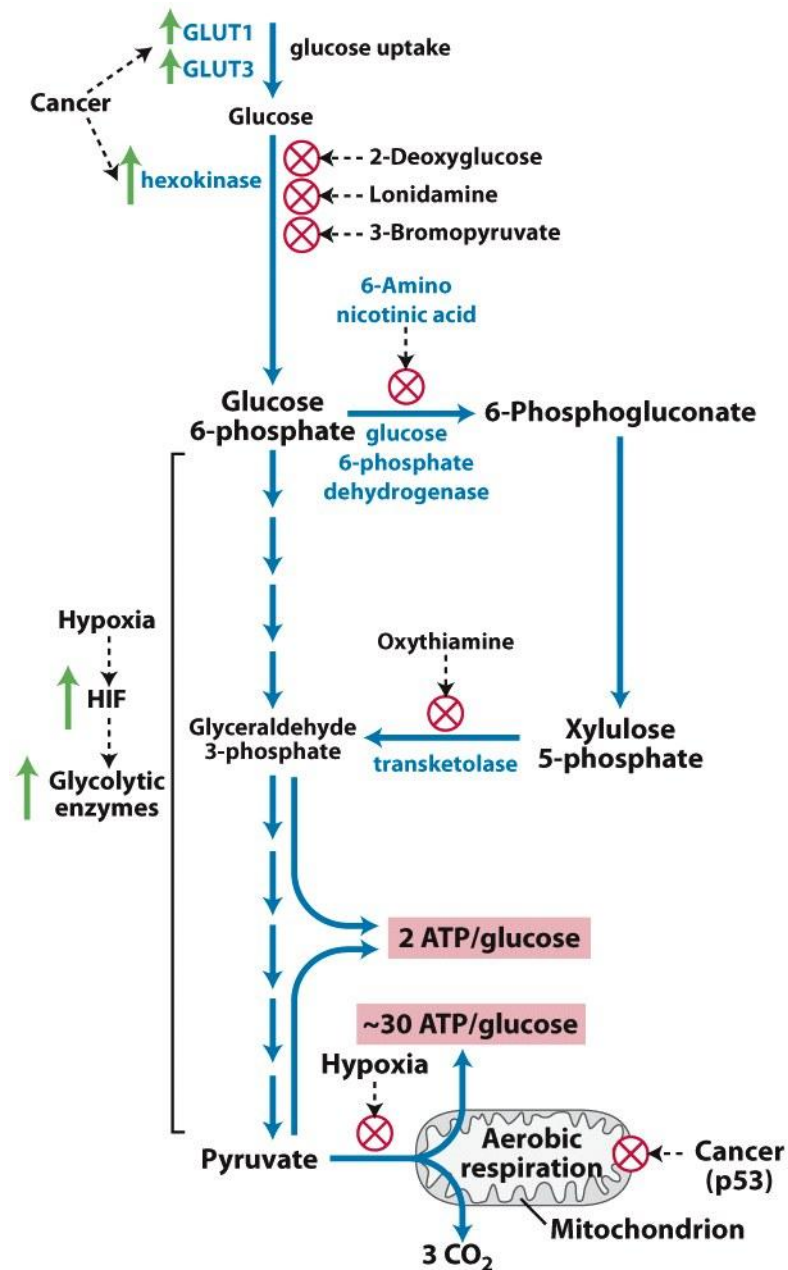


Otto Warburg, 1883–1970

Unnumbered 14 p539

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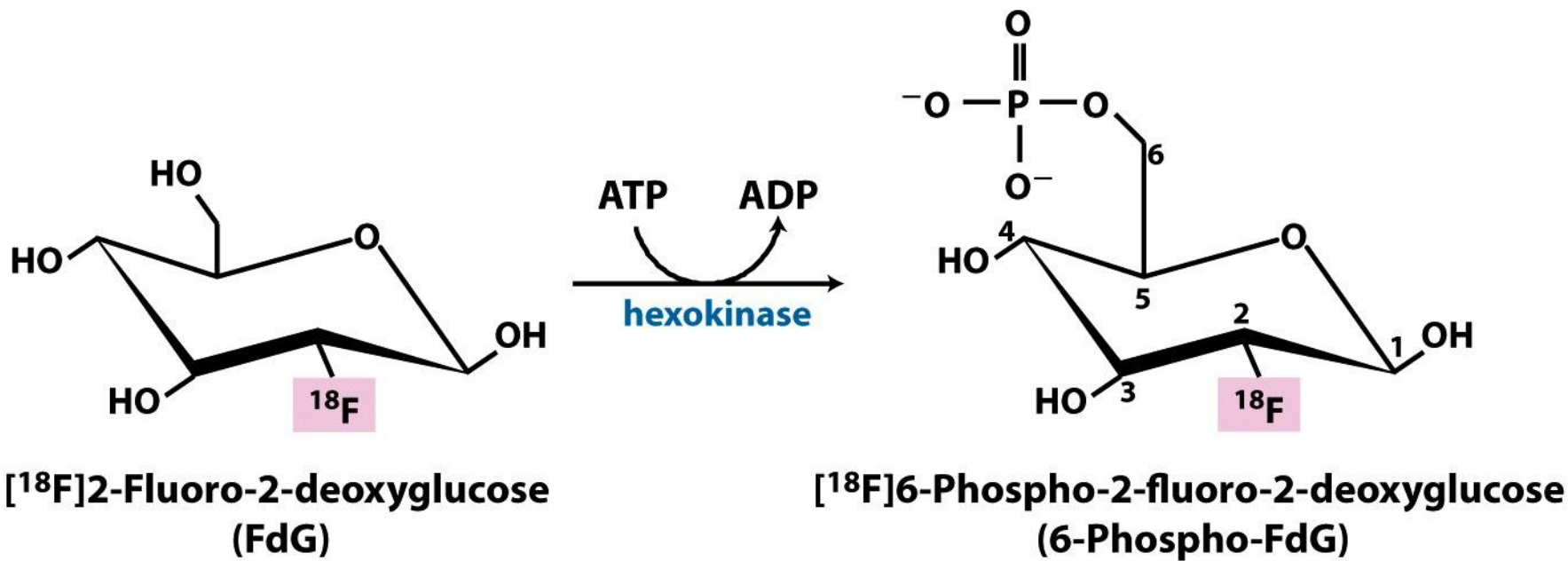
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Box 14-1 figure 1

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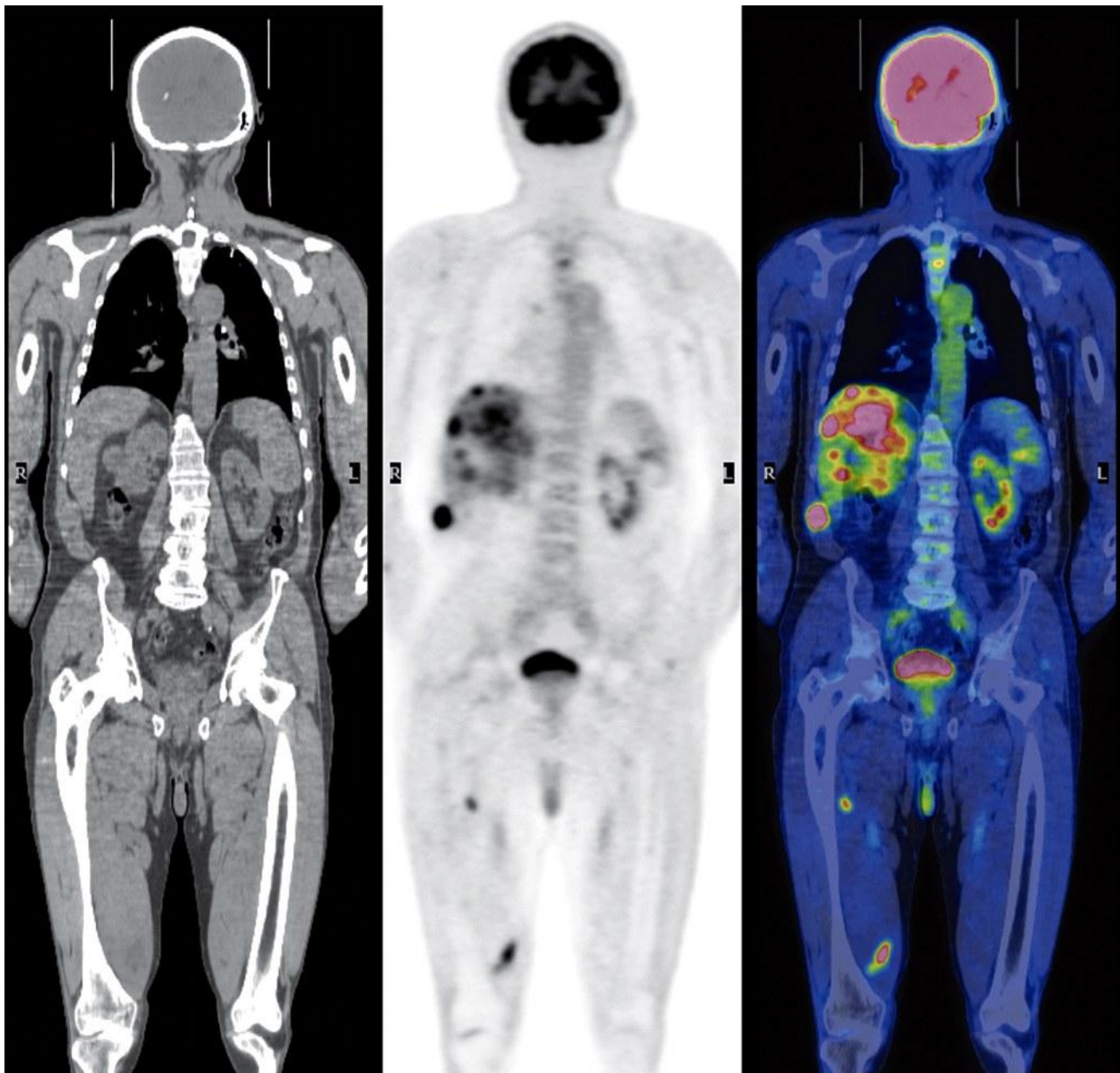
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Box 14-1 figure 2

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Box 14-1 figure 3

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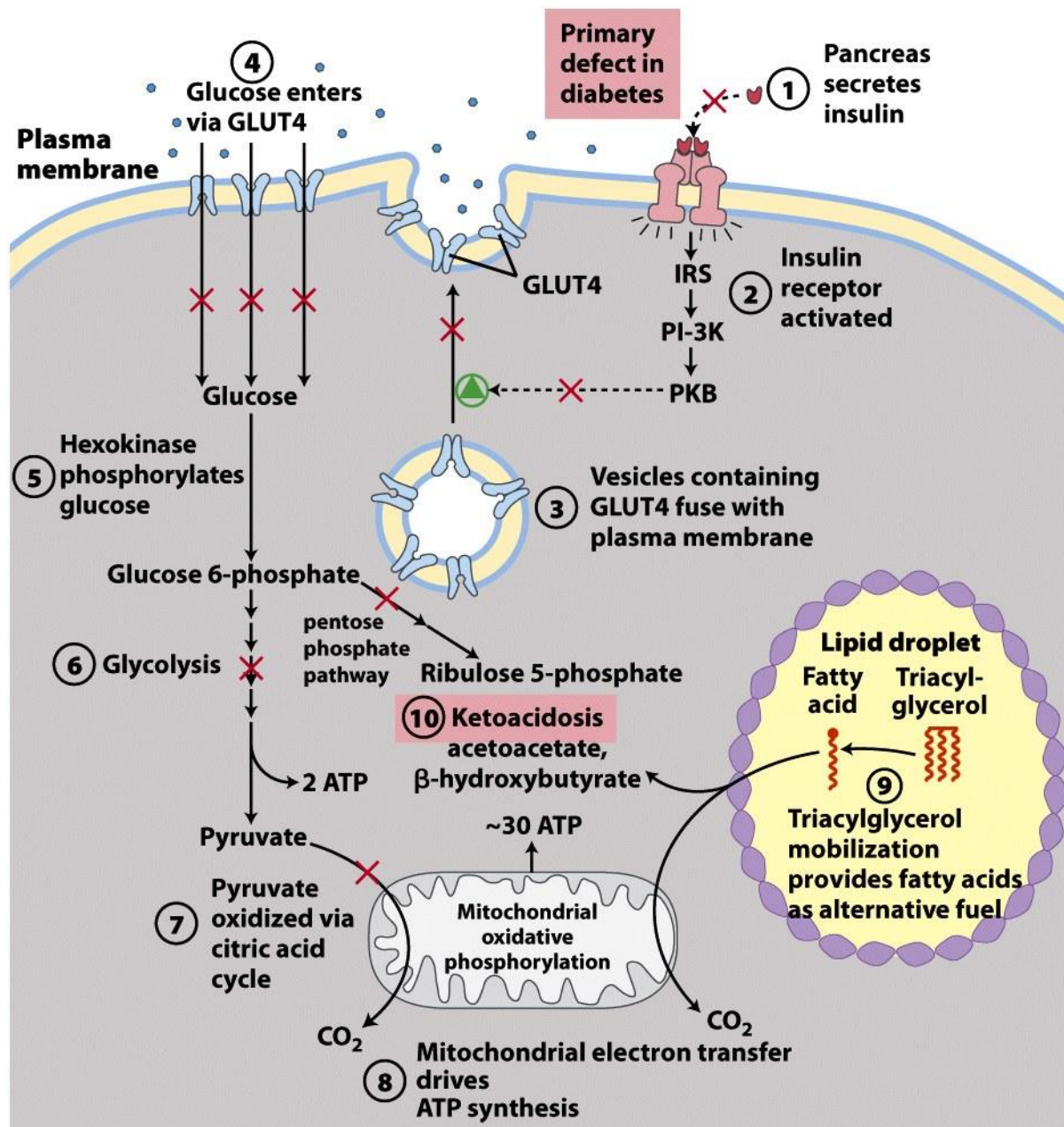


Figure 14-9

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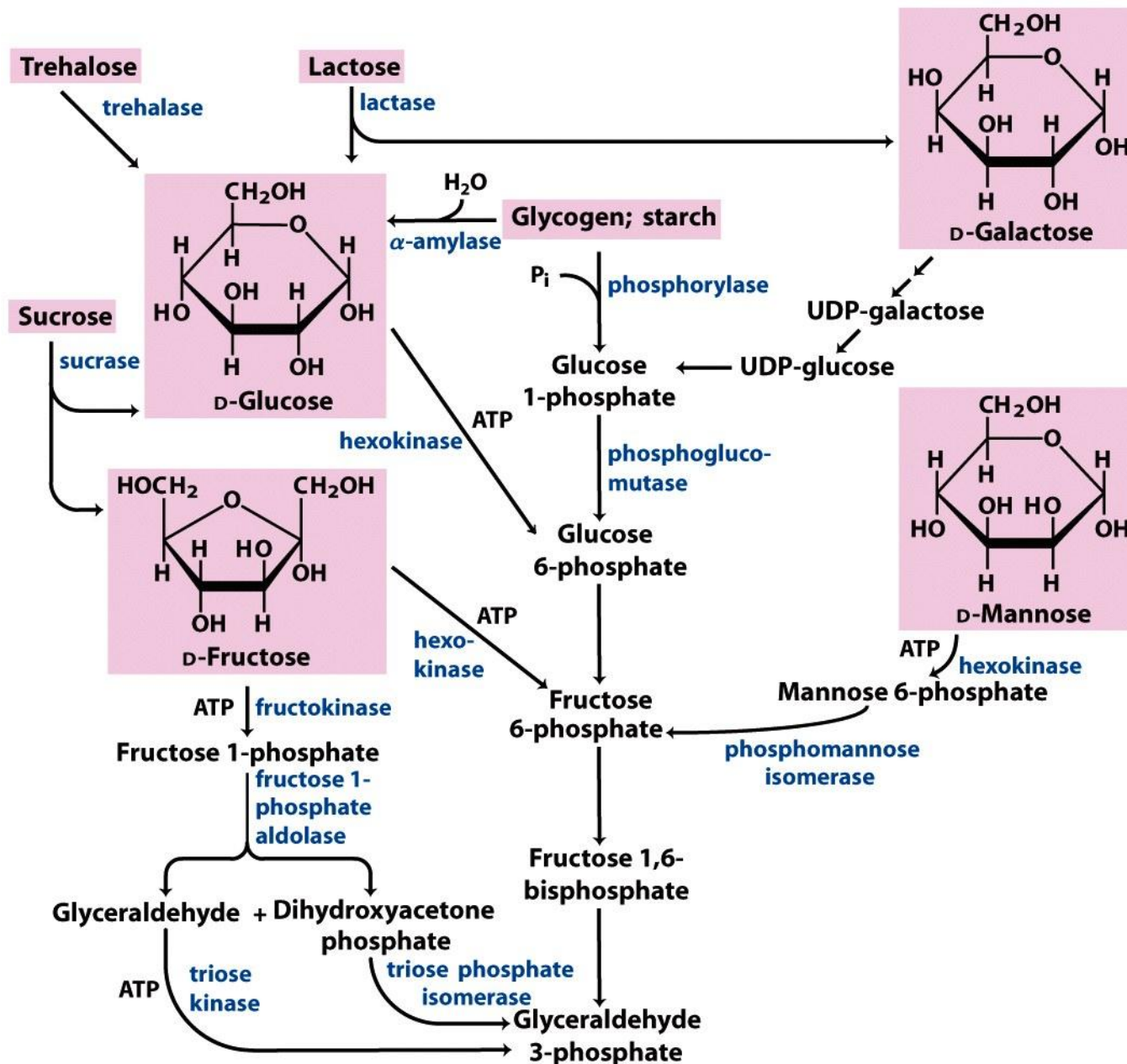
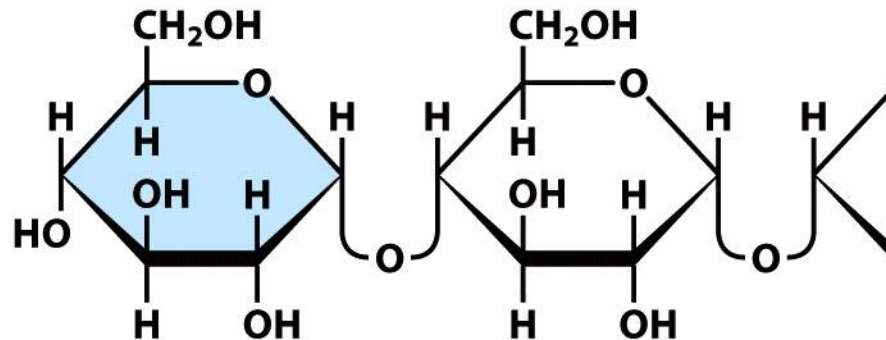


Figure 14-10

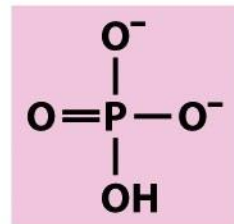
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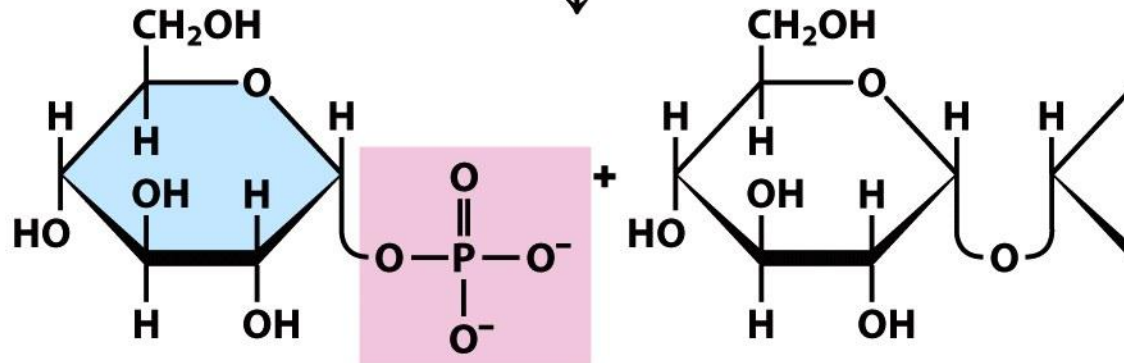
Nonreducing end



Glycogen (starch)
***n* glucose units**



glycogen (starch)
phosphorylase



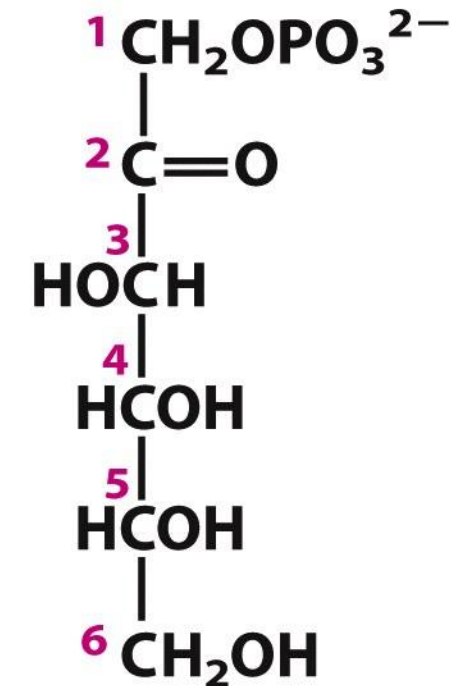
Glucose
1-phosphate

Glycogen (starch)
(*n*-1) glucose units

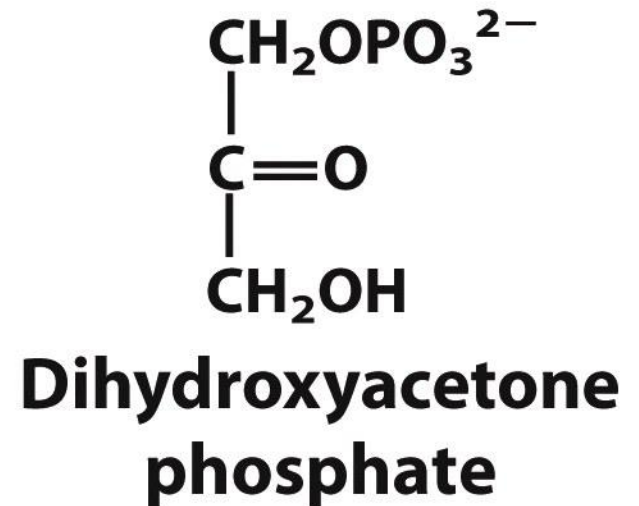
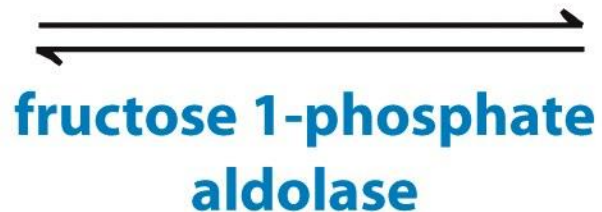
Figure 14-11

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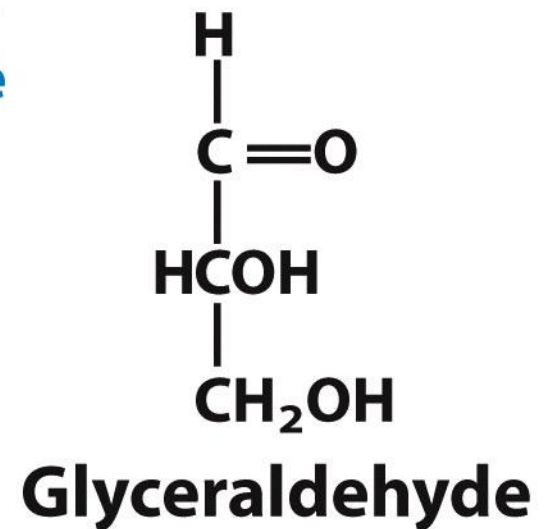
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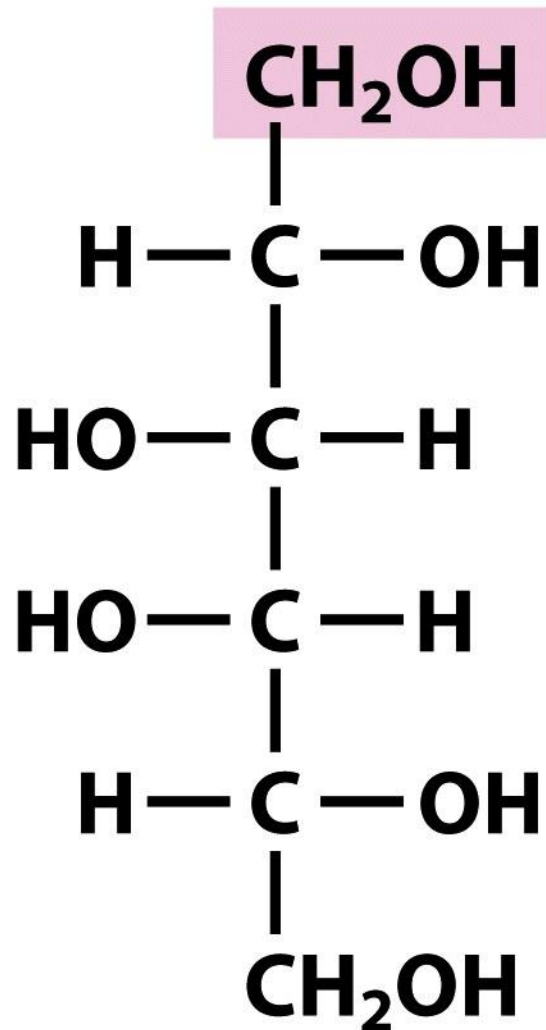


Fructose 1-phosphate



+





D-Galactitol

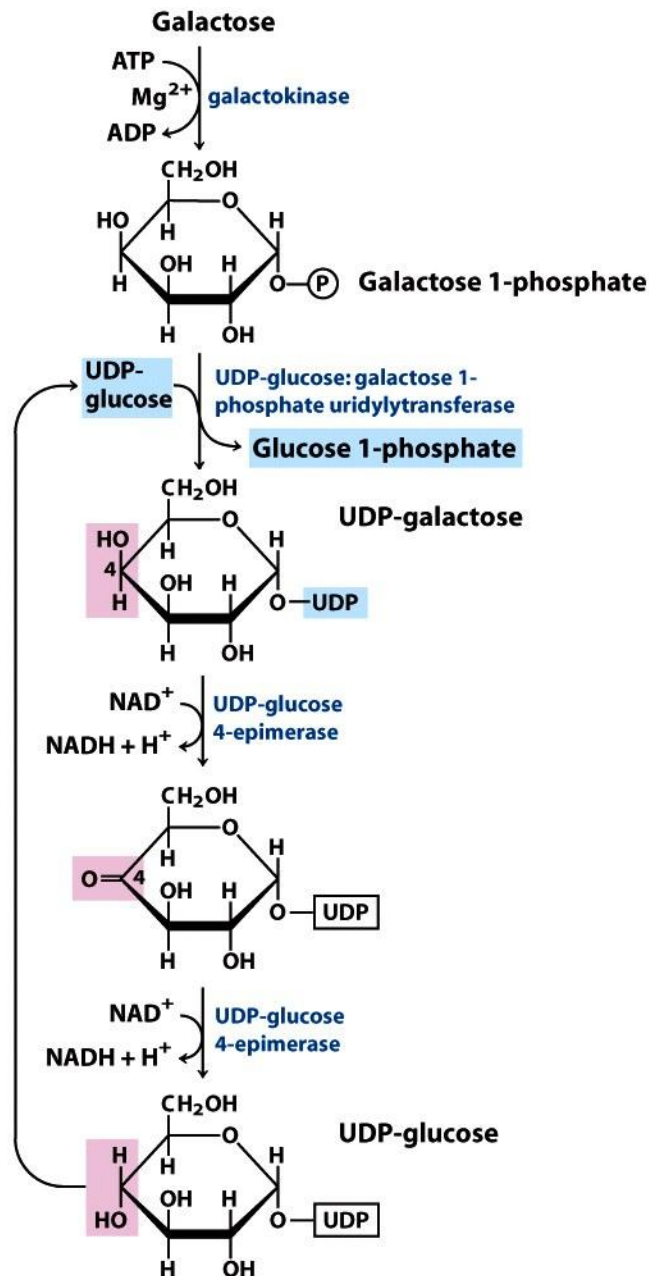


Figure 14-12

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Galactose

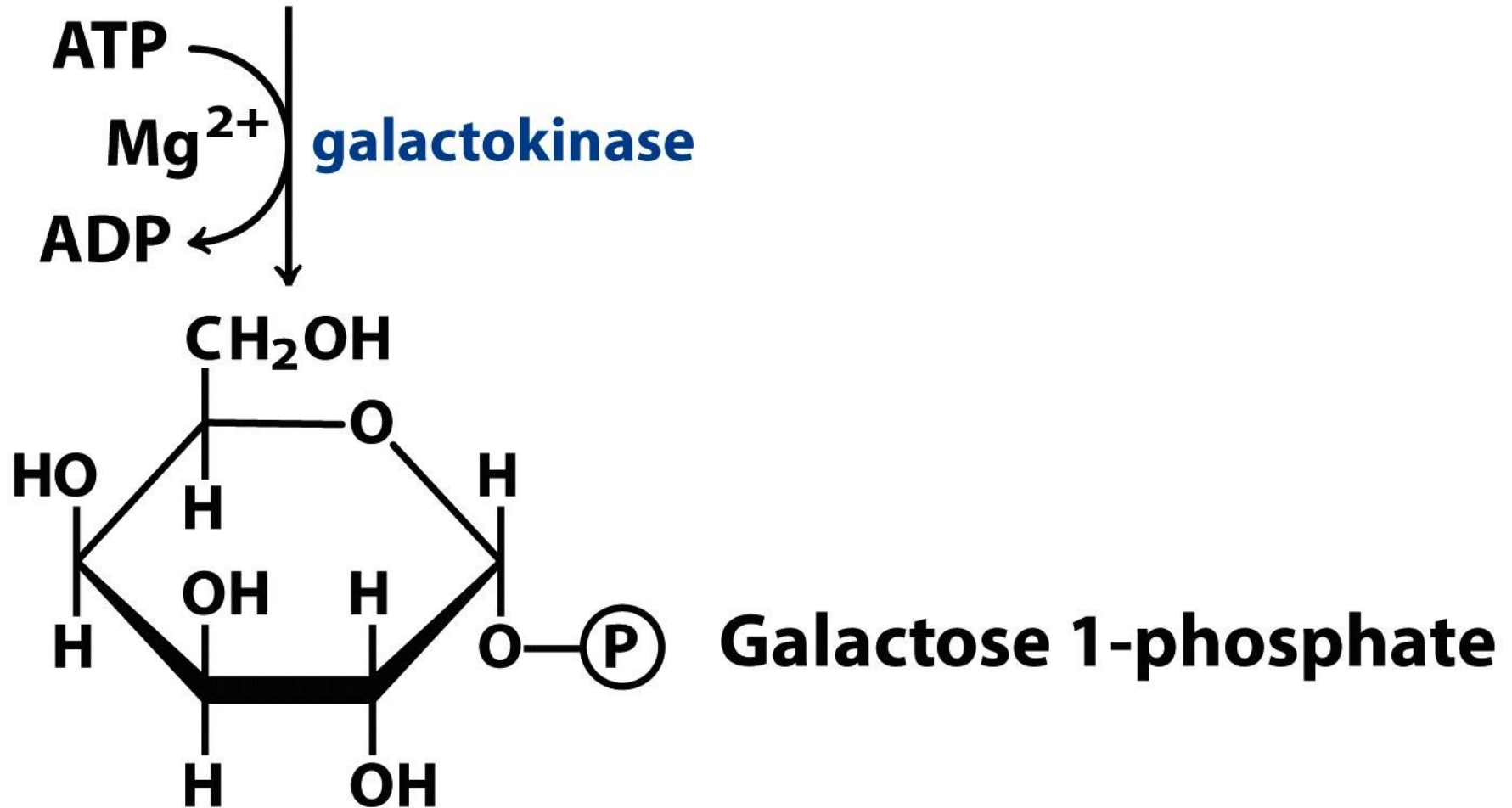


Figure 14-12 part 1

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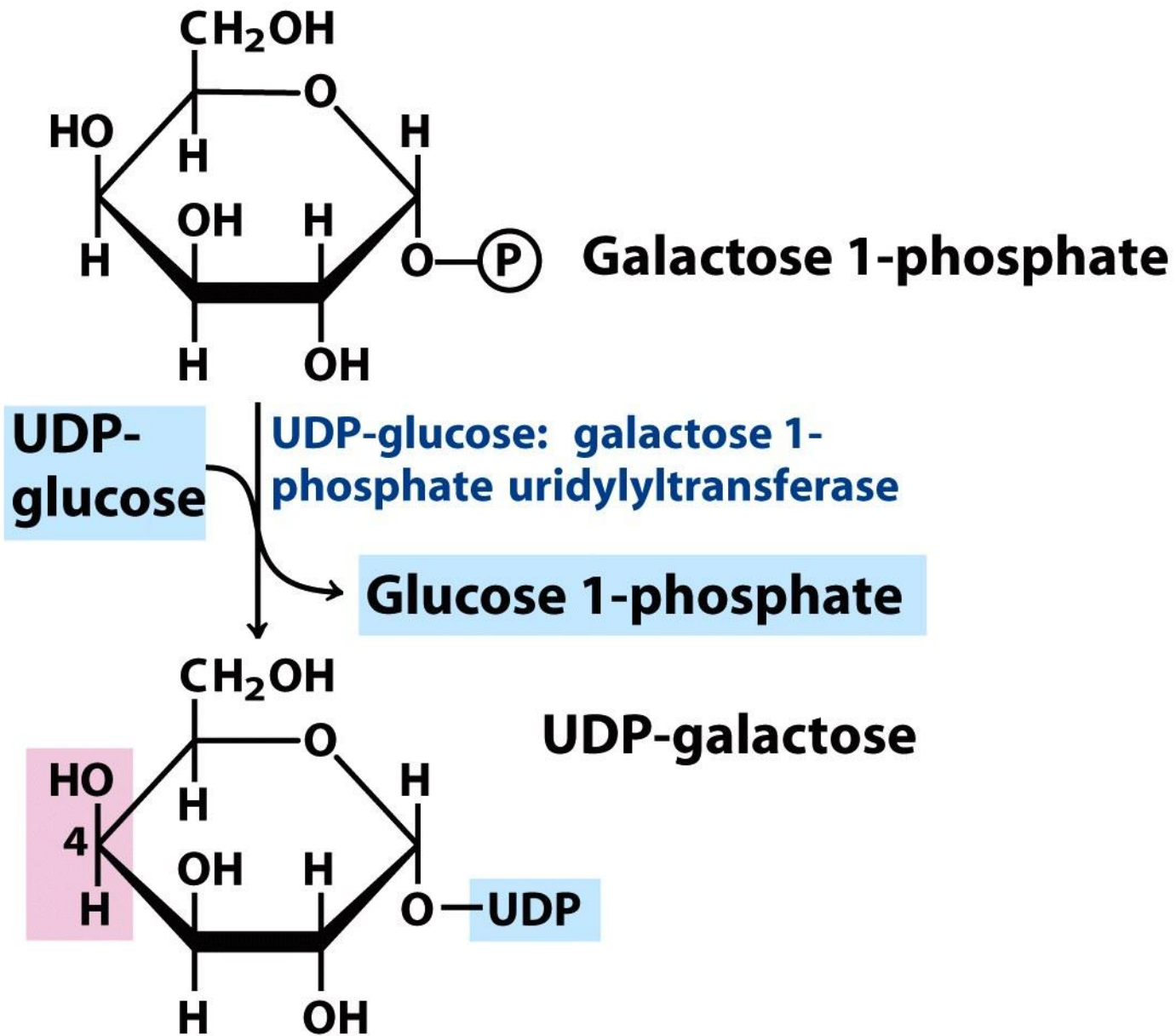


Figure 14-12 part 2

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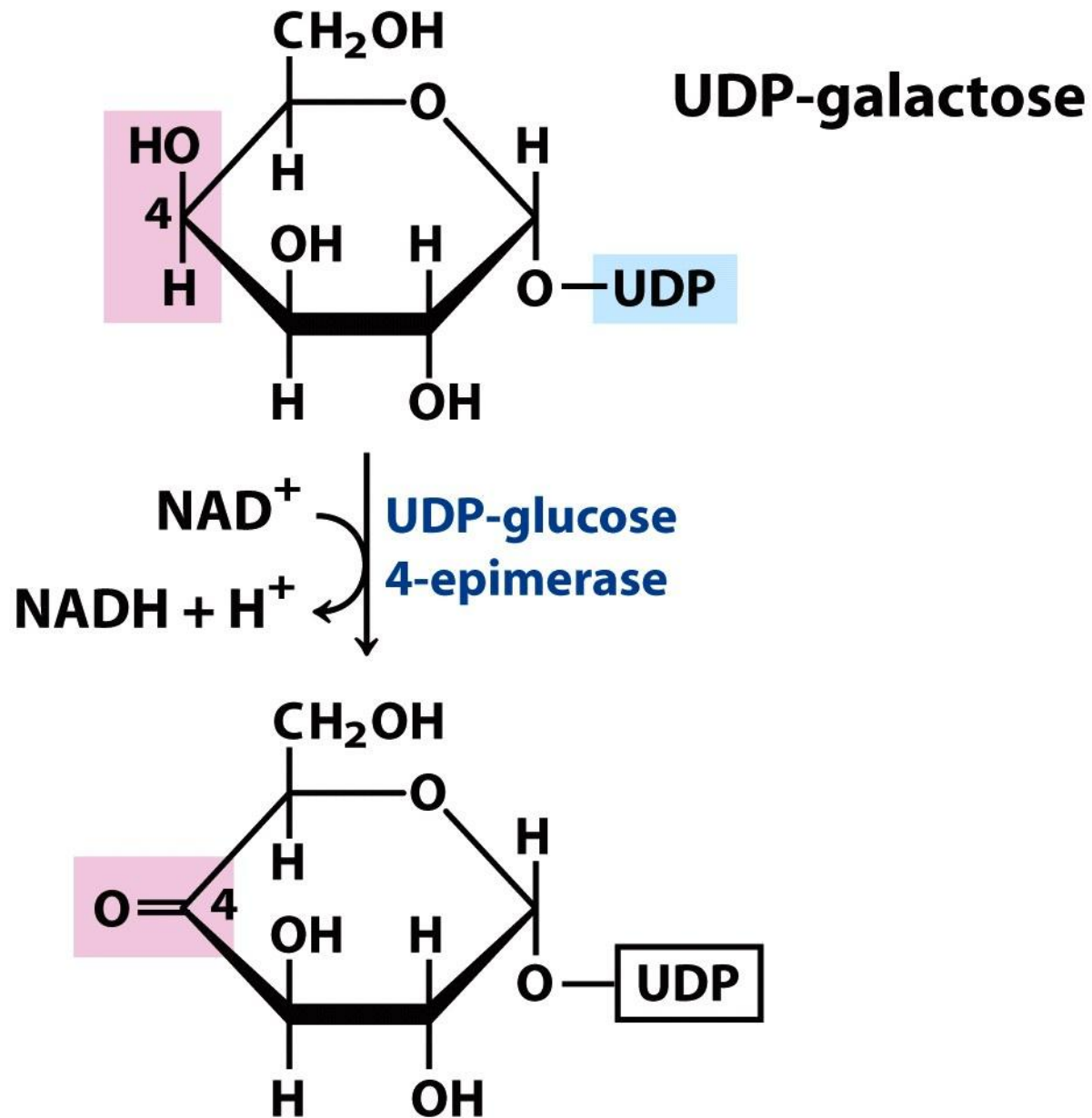


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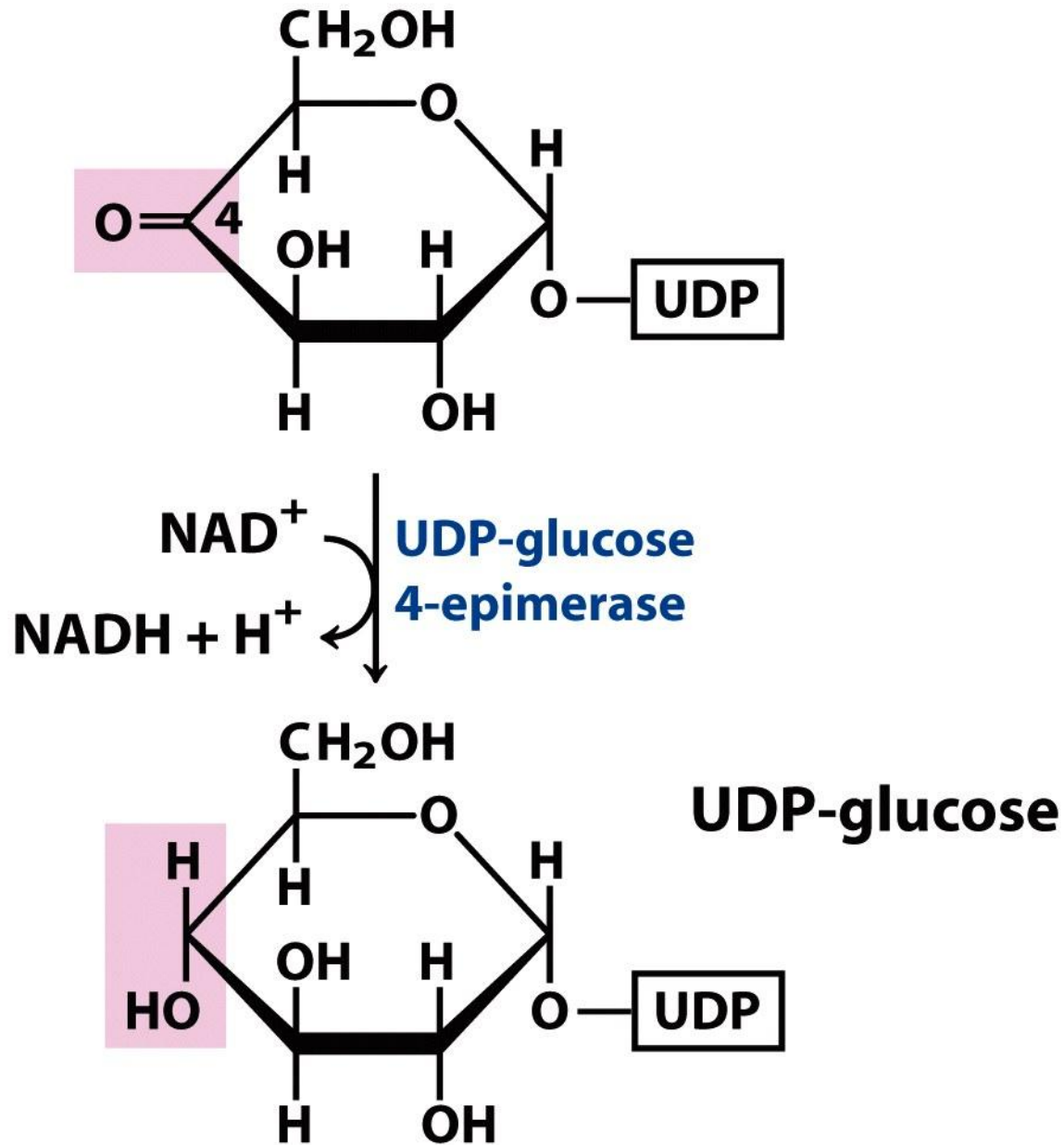
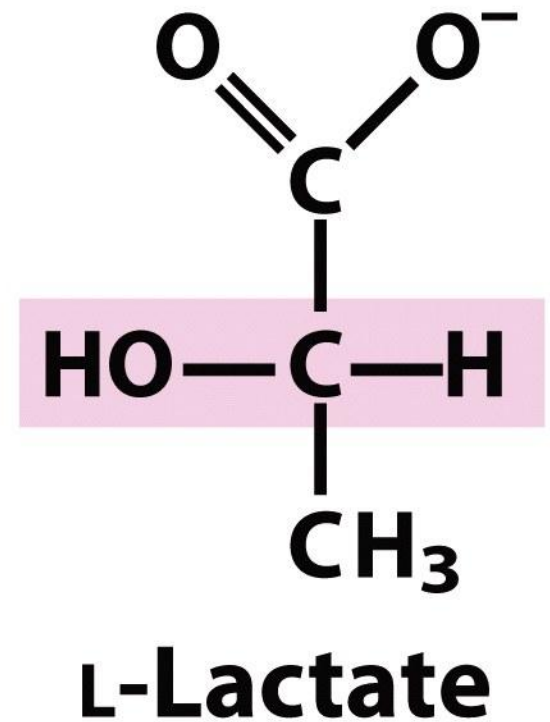
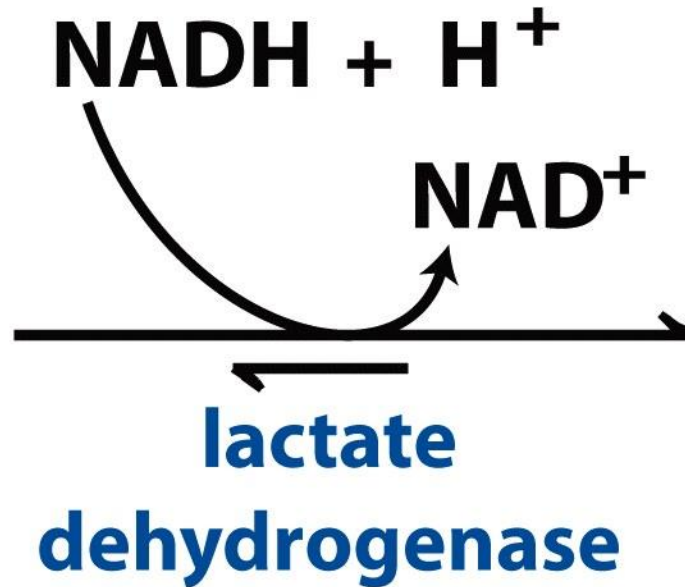
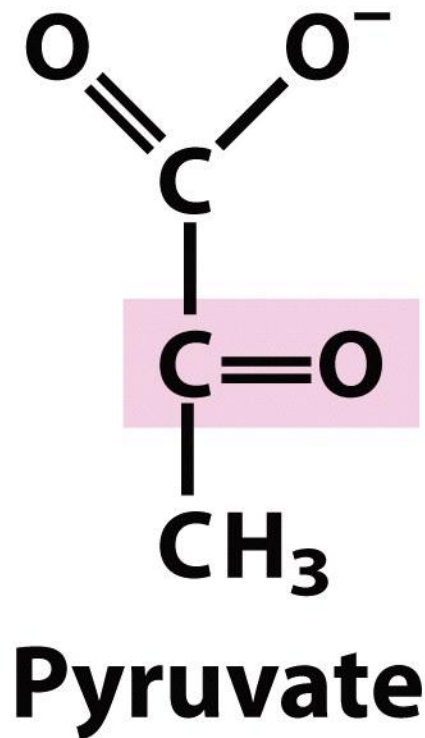
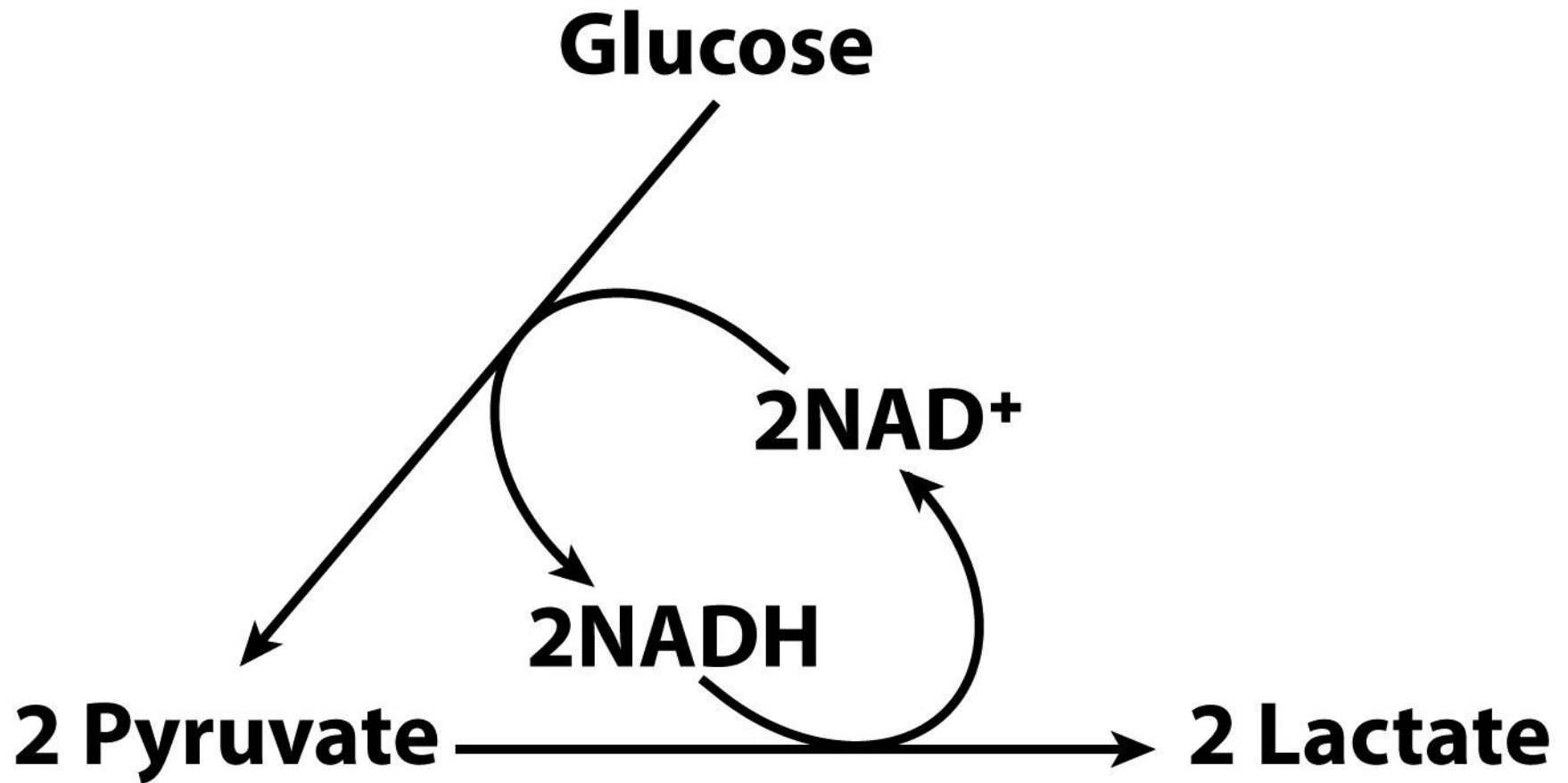


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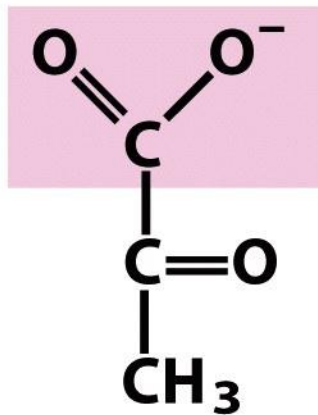
$$\Delta G'^{\circ} = - 25.1 \text{ kJ/mol}$$



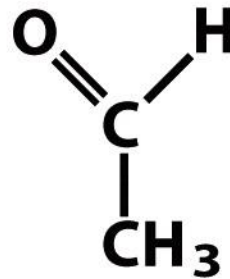
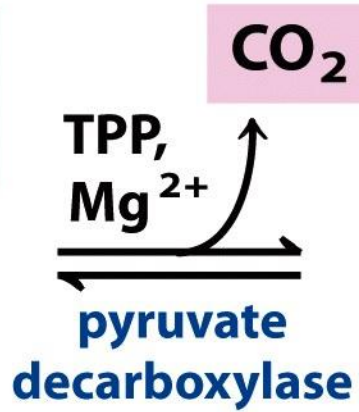
Unnumbered 14 p547b

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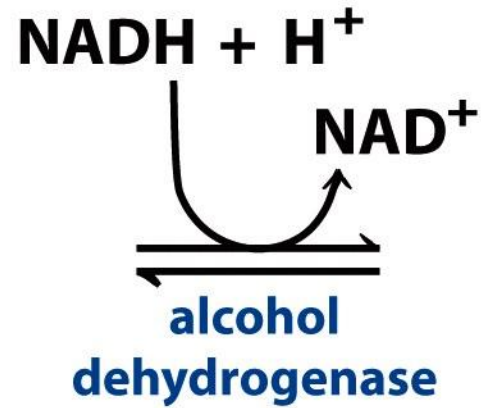
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Pyruvate



Acetaldehyde



Ethanol

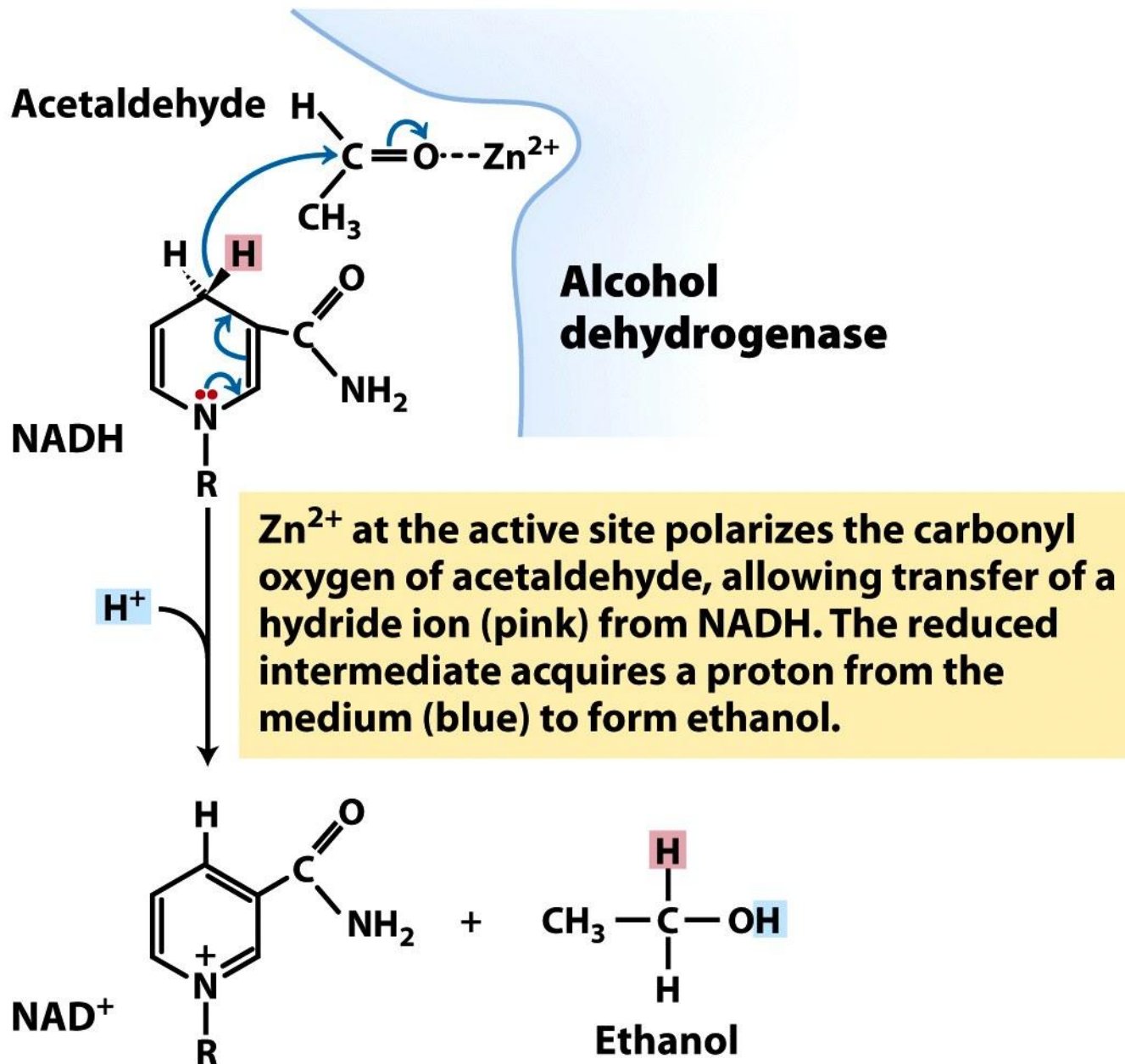
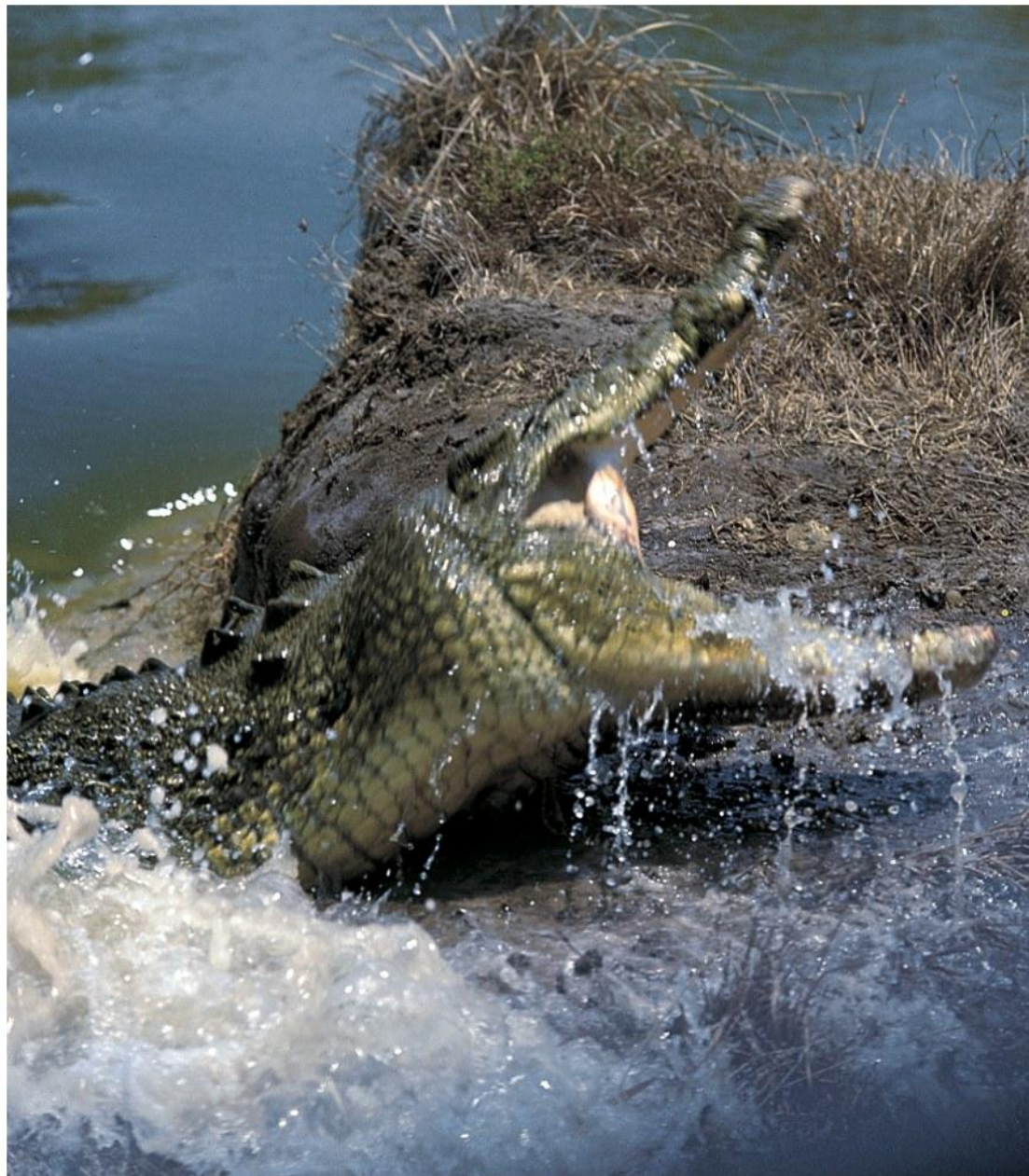


Figure 14-13

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Box 14-2 figure 1

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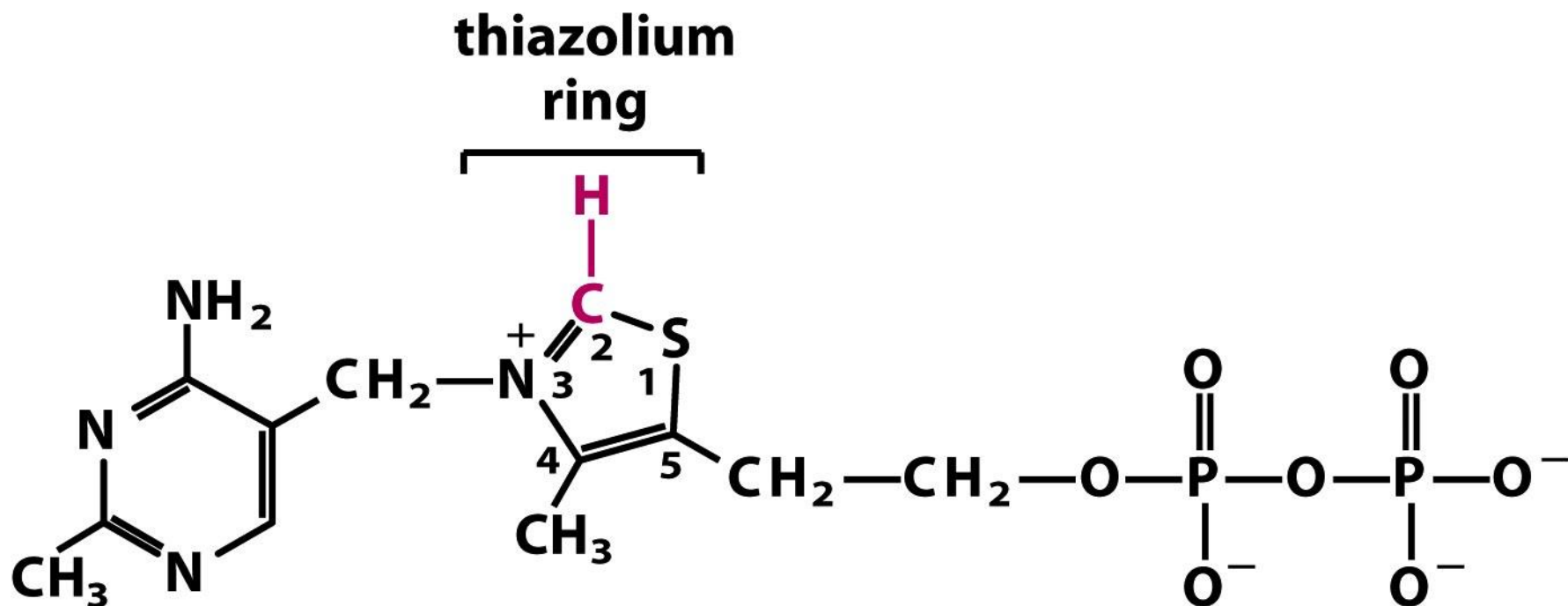
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Box 14-3 figure 1

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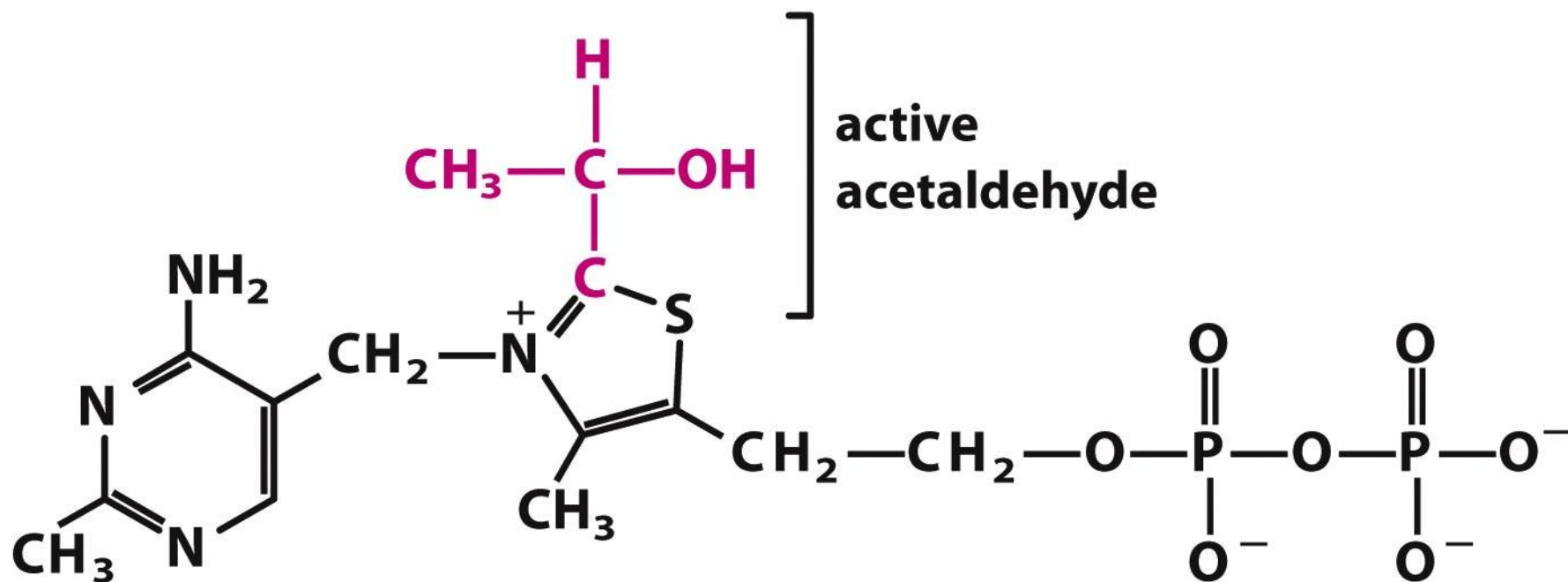


Thiamine pyrophosphate (TPP)

Figure 14-14a

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Hydroxyethyl thiamine pyrophosphate

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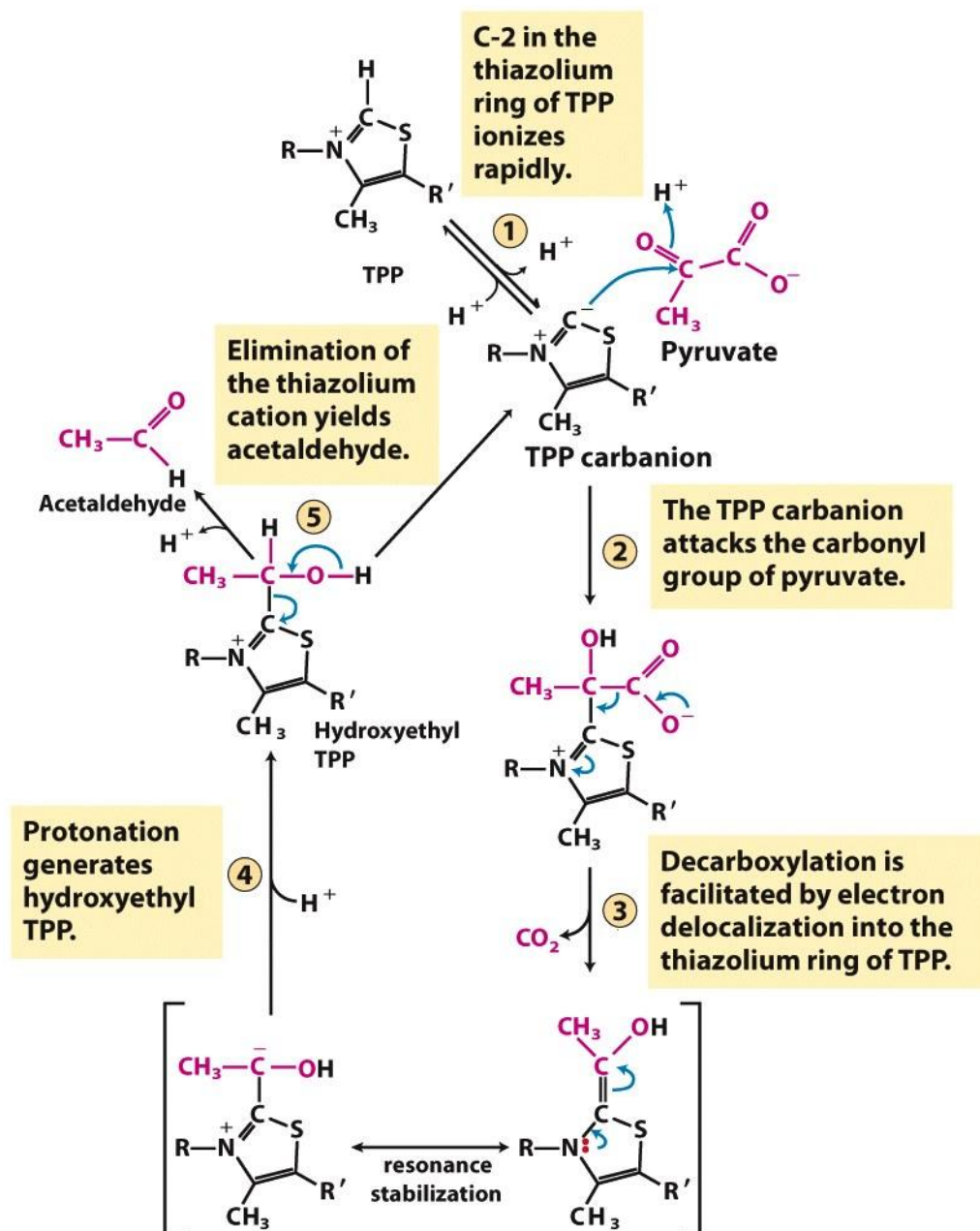


Figure 14-14c

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TABLE 14–1 Some TPP-Dependent Reactions			
Enzyme	Pathway(s)	Bond cleaved	Bond formed
Pyruvate decarboxylase	Ethanol fermentation	$\begin{array}{c} \text{O} \qquad \text{O} \\ \parallel \quad \diagup \\ \text{R}^1\text{—C—C} \\ \qquad \diagdown \\ \qquad \text{O}^- \end{array}$	$\begin{array}{c} \text{O} \\ \parallel \\ \text{R}^1\text{—C} \\ \qquad \diagdown \\ \qquad \text{H} \end{array}$
Pyruvate dehydrogenase α -Ketoglutarate dehydrogenase	Synthesis of acetyl-CoA Citric acid cycle	$\begin{array}{c} \text{O} \qquad \text{O} \\ \parallel \quad \diagup \\ \text{R}^2\text{—C—C} \\ \qquad \diagdown \\ \qquad \text{O}^- \end{array}$	$\begin{array}{c} \text{O} \\ \parallel \\ \text{R}^2\text{—C} \\ \qquad \diagdown \\ \qquad \text{S—CoA} \end{array}$
Transketolase	Carbon-assimilation reactions Pentose phosphate pathway	$\begin{array}{c} \text{O} \quad \text{OH} \\ \parallel \quad \\ \text{R}^3\text{—C—C—R}^4 \\ \qquad \\ \qquad \text{H} \end{array}$	$\begin{array}{c} \text{O} \quad \text{OH} \\ \parallel \quad \\ \text{R}^3\text{—C—C—R}^5 \\ \qquad \\ \qquad \text{H} \end{array}$

Table 14-1

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TABLE 14–2 **Free-Energy Changes of Glycolytic Reactions in Erythrocytes**

Glycolytic reaction step	$\Delta G'^{\circ}$ (kJ/mol)	ΔG (kJ/mol)
1 Glucose + ATP \longrightarrow glucose 6-phosphate + ADP	−16.7	−33.4
2 Glucose 6-phosphate \rightleftharpoons fructose 6-phosphate	1.7	0 to 25
3 Fructose 6-phosphate + ATP \longrightarrow fructose 1,6-bisphosphate + ADP	−14.2	−22.2
4 Fructose 1,6-bisphosphate \rightleftharpoons dihydroxyacetone phosphate + glyceraldehyde 3-phosphate	23.8	−6 to 0
5 Dihydroxyacetone phosphate \rightleftharpoons glyceraldehyde 3-phosphate	7.5	0 to 4
6 Glyceraldehyde 3-phosphate + P_i + NAD^+ \rightleftharpoons 1,3-bisphosphoglycerate + $NADH$ + H^+	6.3	−2 to 2
7 1,3-Bisphosphoglycerate + ADP \rightleftharpoons 3-phosphoglycerate + ATP	−18.8	0 to 2
8 3-Phosphoglycerate \rightleftharpoons 2-phosphoglycerate	4.4	0 to 0.8
9 2-Phosphoglycerate \rightleftharpoons phosphoenolpyruvate + H_2O	7.5	0 to 3.3
10 Phosphoenolpyruvate + ADP \longrightarrow pyruvate + ATP	−31.4	−16.7

Note: $\Delta G'^{\circ}$ is the standard free-energy change, as defined in Chapter 13 (pp. 491–492). ΔG is the free-energy change calculated from the actual concentrations of glycolytic intermediates present under physiological conditions in erythrocytes, at pH 7. The glycolytic reactions bypassed in gluconeogenesis are shown in red. Biochemical equations are not necessarily balanced for H or charge (p. 501).

Table 14-2

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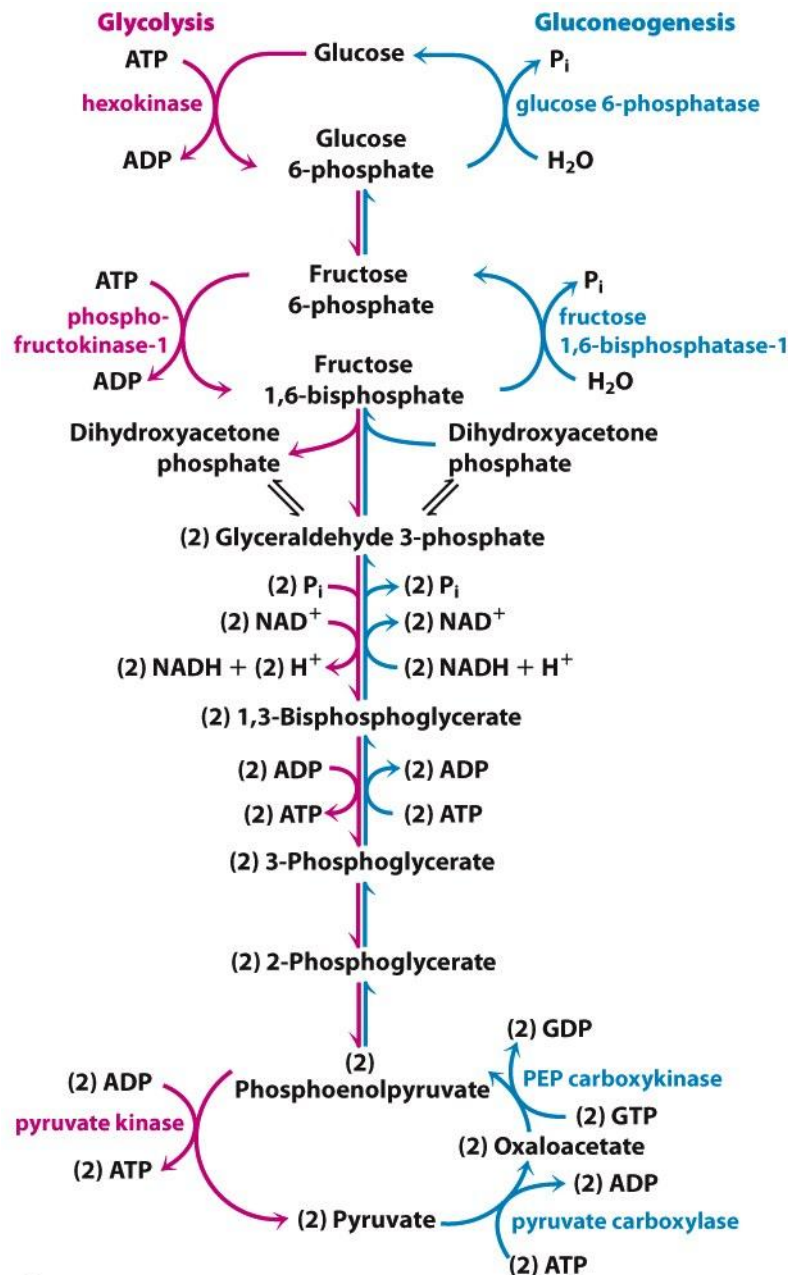


Figure 14-16

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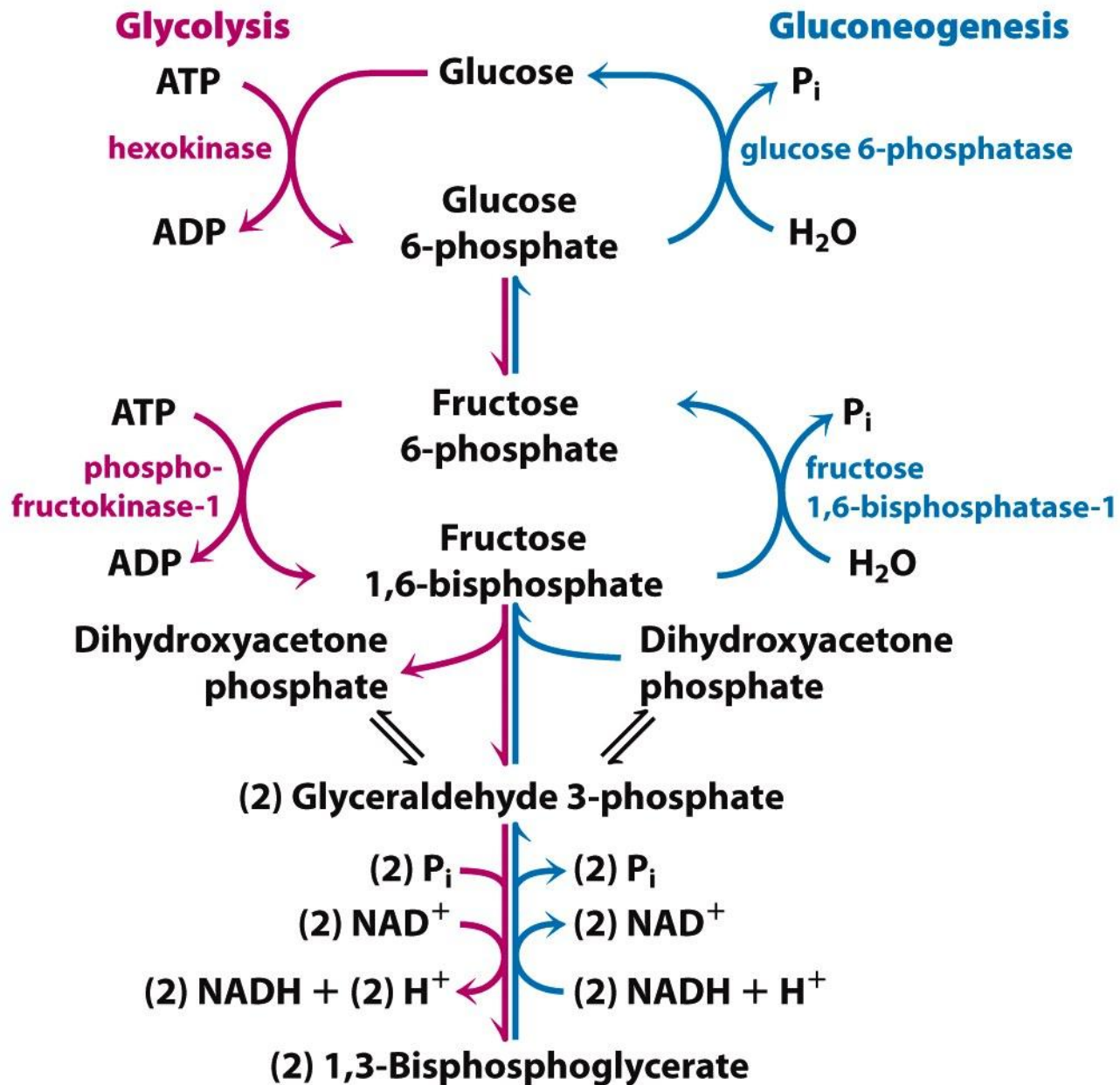


Figure 14-16 part 1

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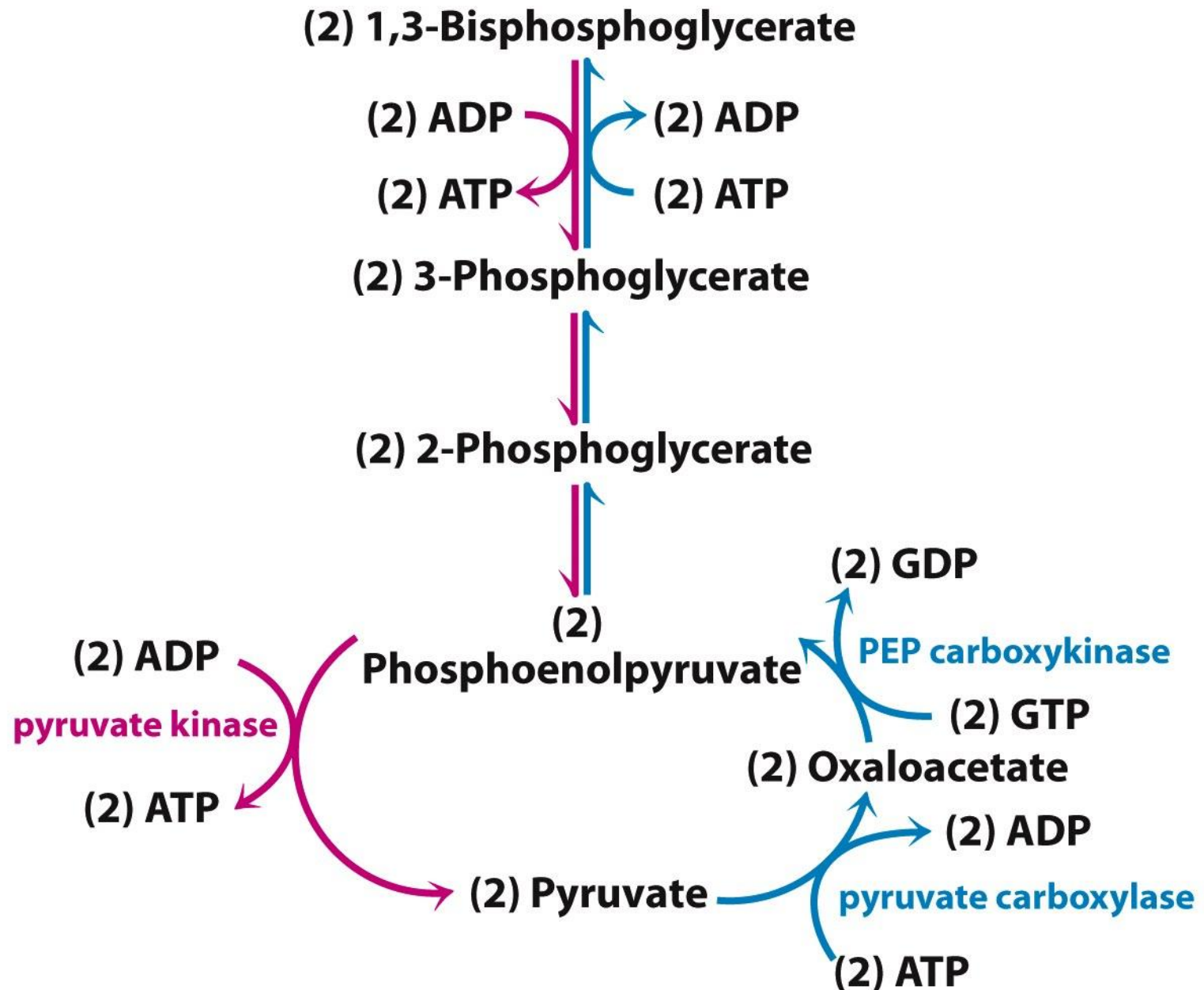


Figure 14-16 part 2
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Bicarbonate

Pyruvate

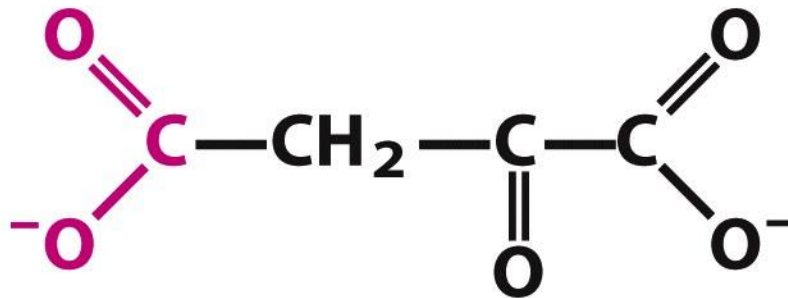


**pyruvate
carboxylase**

ATP

biotin

ADP + P_i



Oxaloacetate

Figure 14-17a

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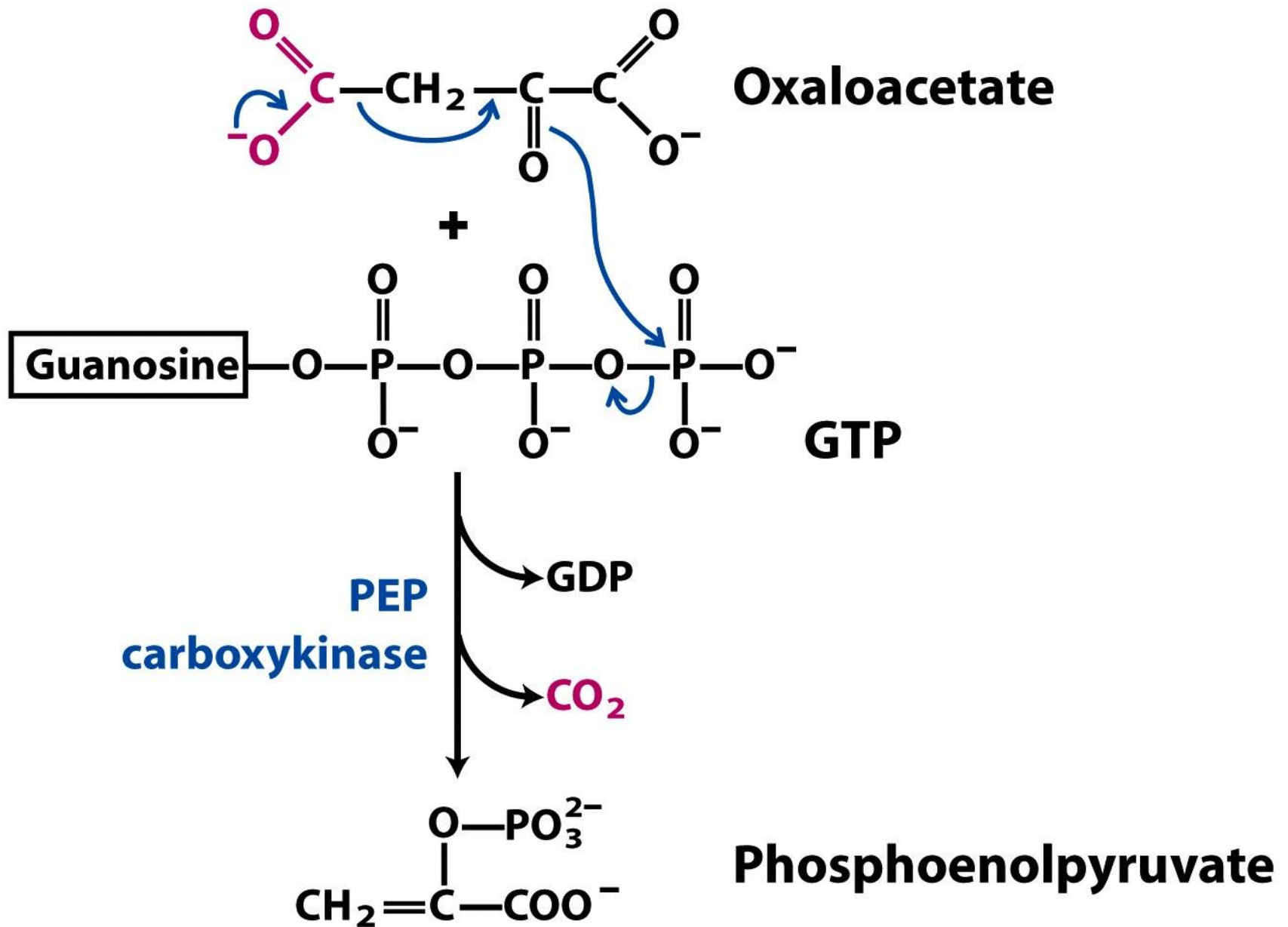


Figure 14-17b

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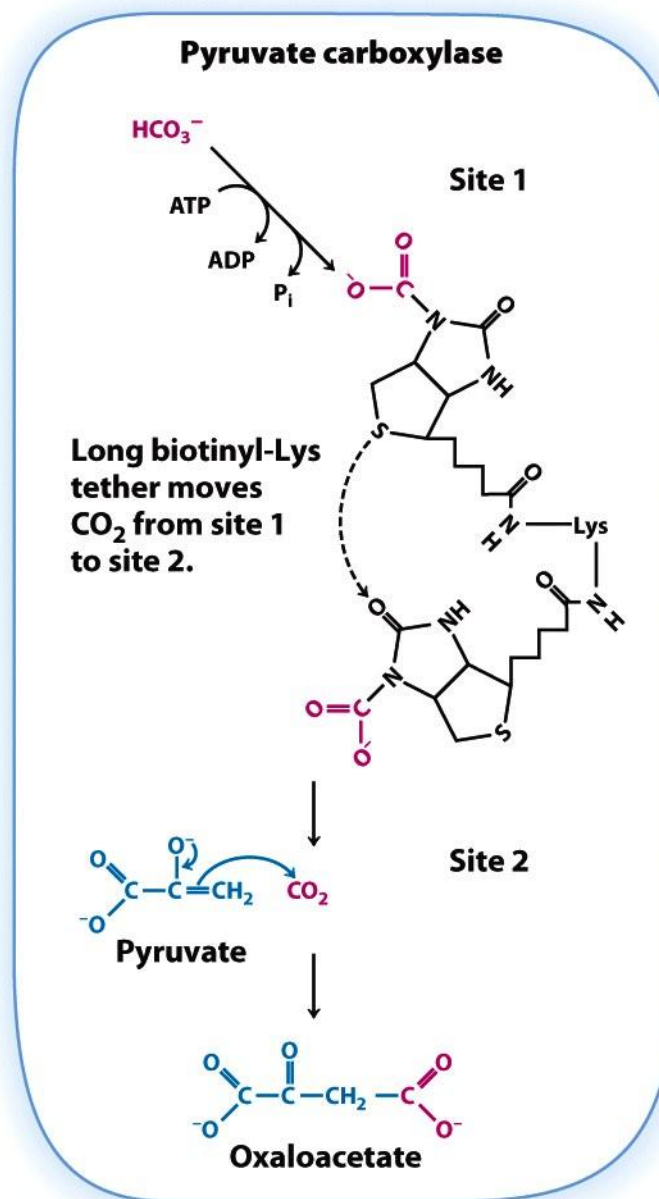


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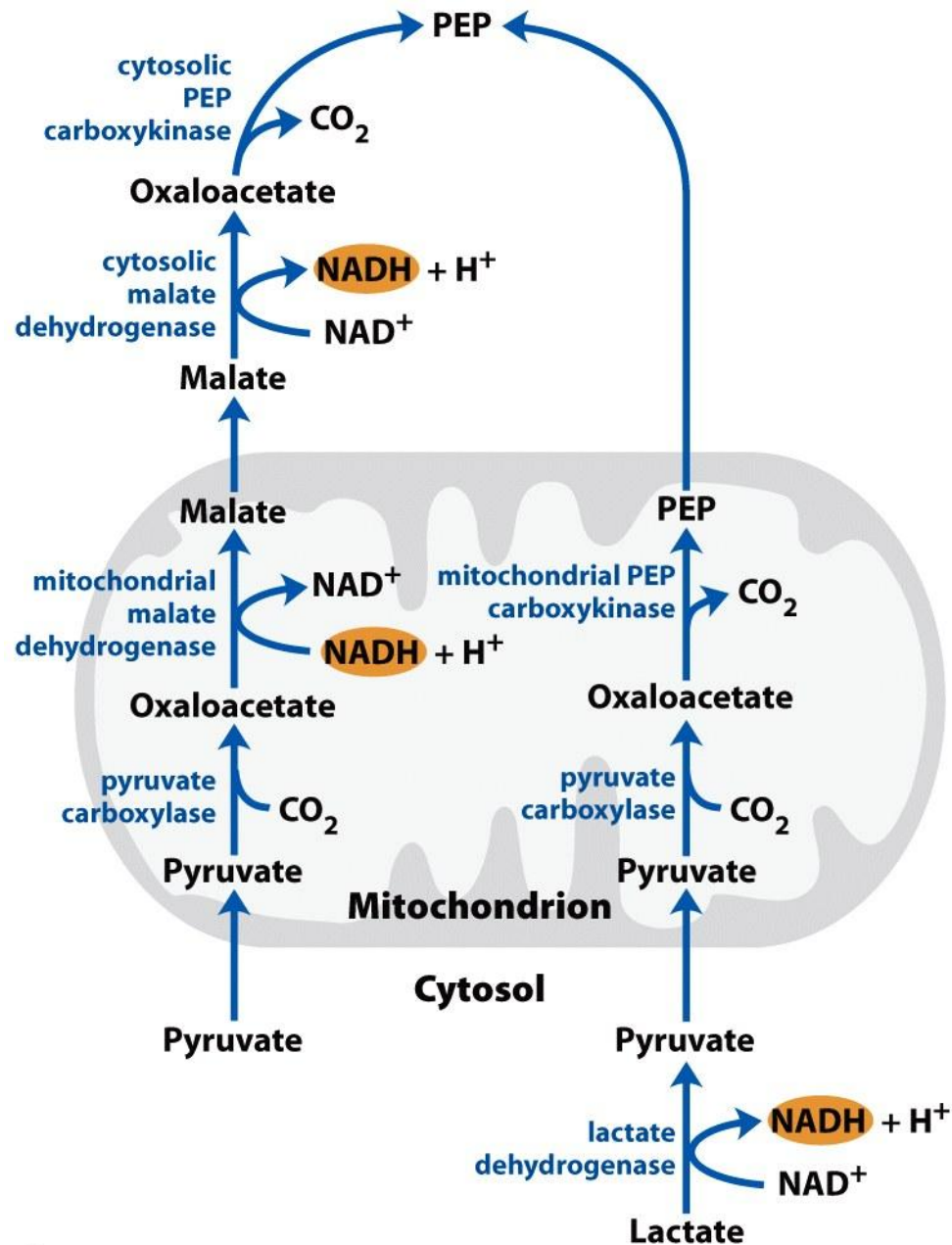


Figure 14-19

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TABLE 14–3 Sequential Reactions in Gluconeogenesis Starting from Pyruvate

Pyruvate + HCO₃[−] + ATP → oxaloacetate + ADP + P_i	×2
Oxaloacetate + GTP ⇌ phosphoenolpyruvate + CO₂ + GDP	×2
Phosphoenolpyruvate + H ₂ O ⇌ 2-phosphoglycerate	×2
2-Phosphoglycerate ⇌ 3-phosphoglycerate	×2
3-Phosphoglycerate + ATP ⇌ 1,3-bisphosphoglycerate + ADP	×2
1,3-Bisphosphoglycerate + NADH + H ⁺ ⇌ glyceraldehyde 3-phosphate + NAD ⁺ + P _i	×2
Glyceraldehyde 3-phosphate ⇌ dihydroxyacetone phosphate	
Glyceraldehyde 3-phosphate + dihydroxyacetone phosphate ⇌ fructose 1,6-bisphosphate	
Fructose 1,6-bisphosphate → fructose 6-phosphate + P_i	
Fructose 6-phosphate ⇌ glucose 6-phosphate	
Glucose 6-phosphate + H₂O → glucose + P_i	
Sum: 2 Pyruvate + 4ATP + 2GTP + 2NADH + 2H⁺ + 4H₂O → glucose + 4ADP + 2GDP + 6P_i + 2NAD⁺	

Note: The bypass reactions are in red; all other reactions are reversible steps of glycolysis. The figures at the right indicate that the reaction is to be counted twice, because two three-carbon precursors are required to make a molecule of glucose. The reactions required to replace the cytosolic NADH consumed in the glyceraldehyde 3-phosphate dehydrogenase reaction (the conversion of lactate to pyruvate in the cytosol or the transport of reducing equivalents from mitochondria to the cytosol in the form of malate) are not considered in this summary. Biochemical equations are not necessarily balanced for H and charge (p. 501).

Table 14-3

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TABLE 14-4**Glucogenic Amino Acids, Grouped by Site of Entry****Pyruvate****Alanine****Cysteine****Glycine****Serine****Threonine****Tryptophan*** **α -Ketoglutarate****Arginine****Glutamate****Glutamine****Histidine****Proline****Succinyl-CoA****Isoleucine*****Methionine****Threonine****Valine****Fumarate****Phenylalanine*****Tyrosine*****Oxaloacetate****Asparagine****Aspartate**

Note: All these amino acids are precursors of blood glucose or liver glycogen, because they can be converted to pyruvate or citric acid cycle intermediates. Of the 20 common amino acids, only leucine and lysine are unable to furnish carbon for net glucose synthesis.

***These amino acids are also ketogenic (see Fig. 18-21).**

Table 14-4

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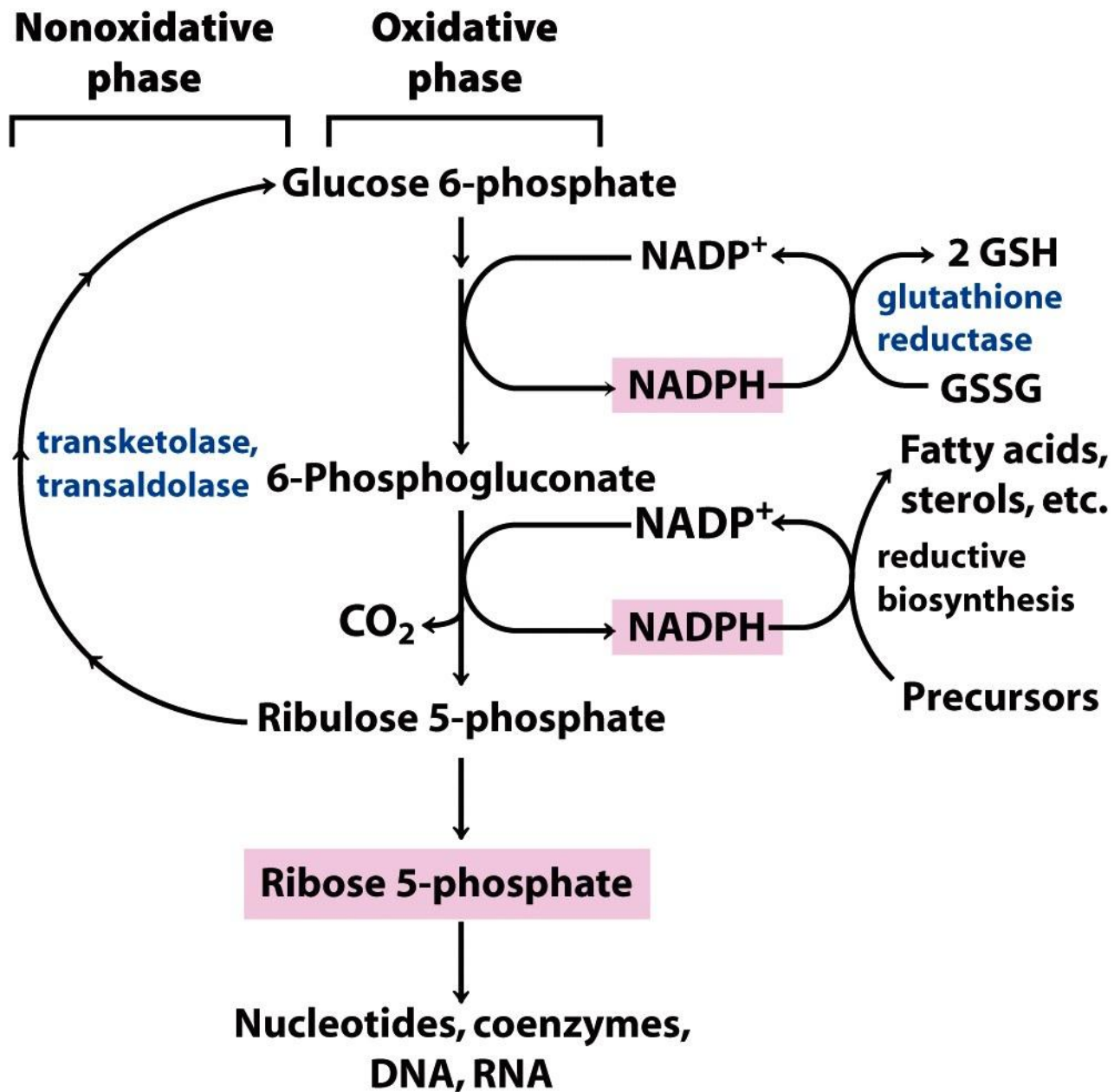
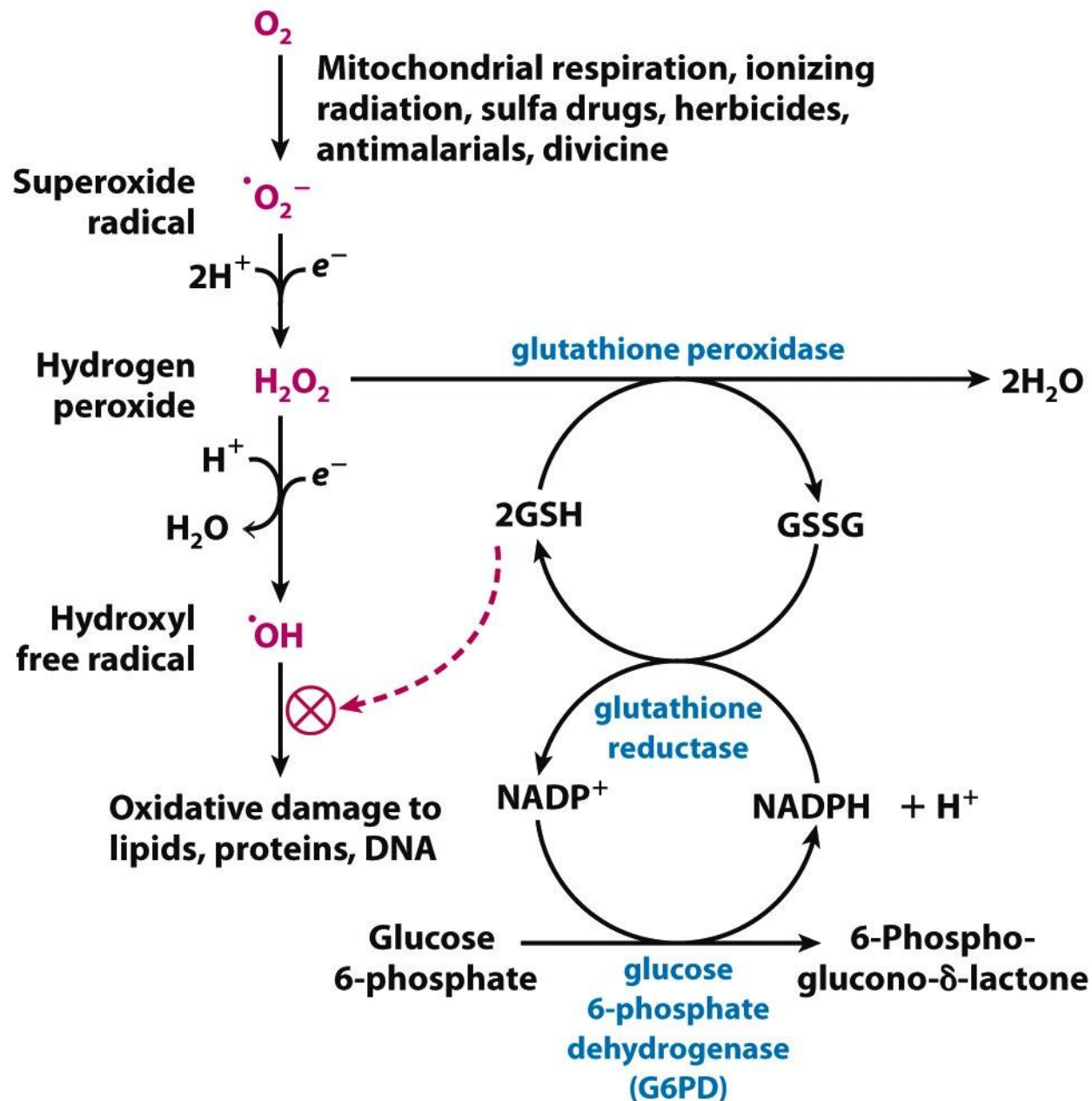


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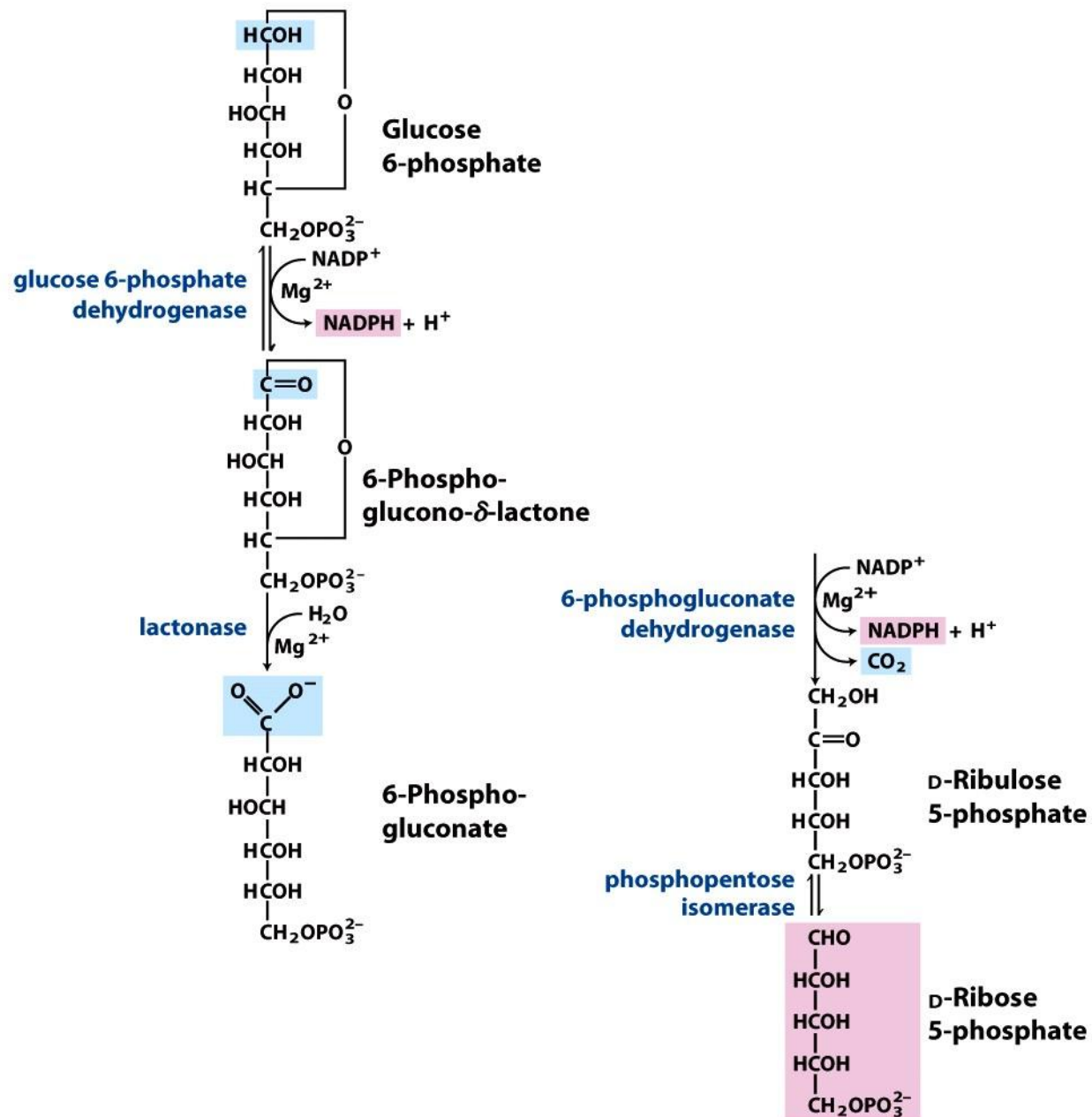


Figure 14-21

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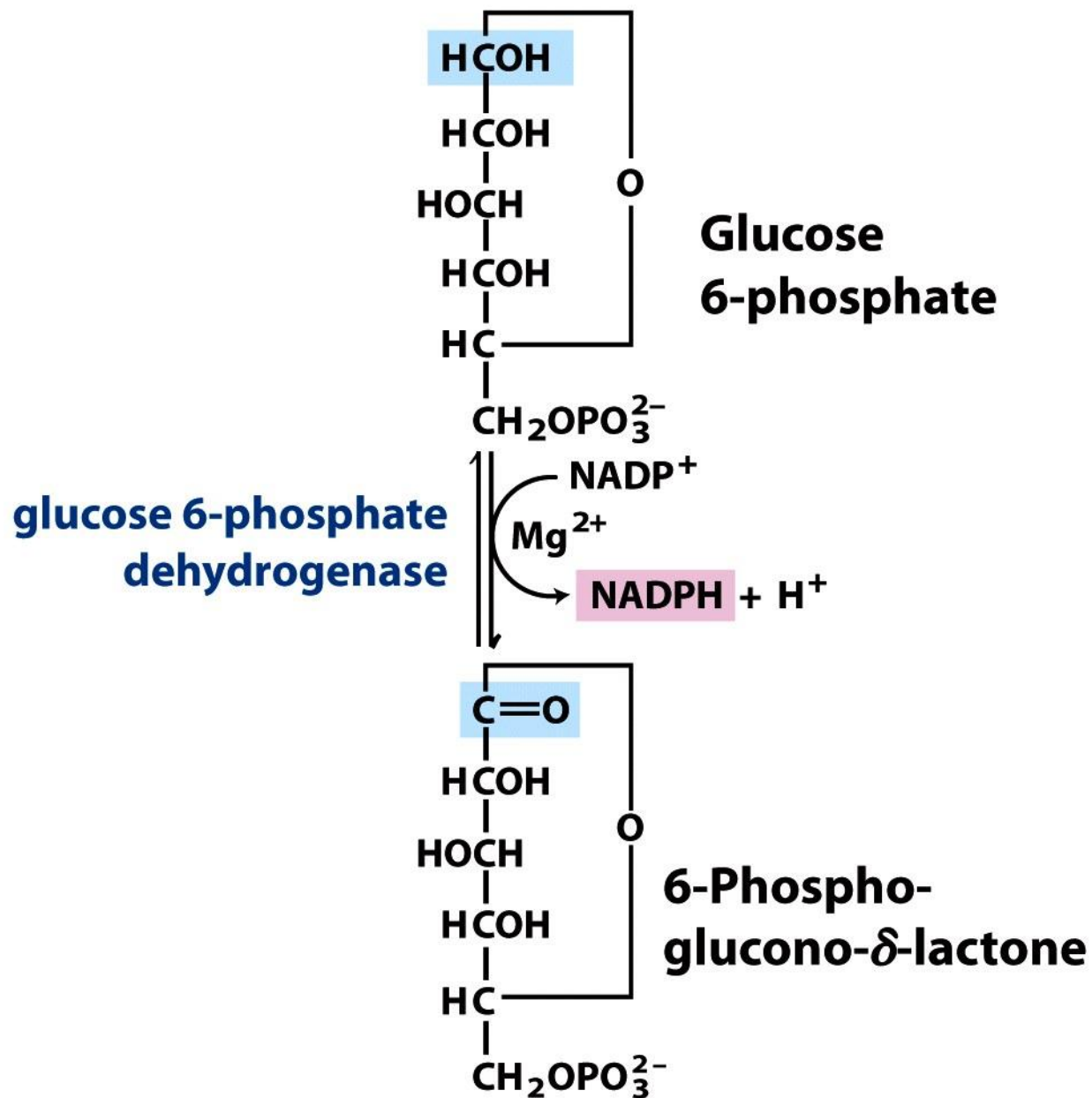


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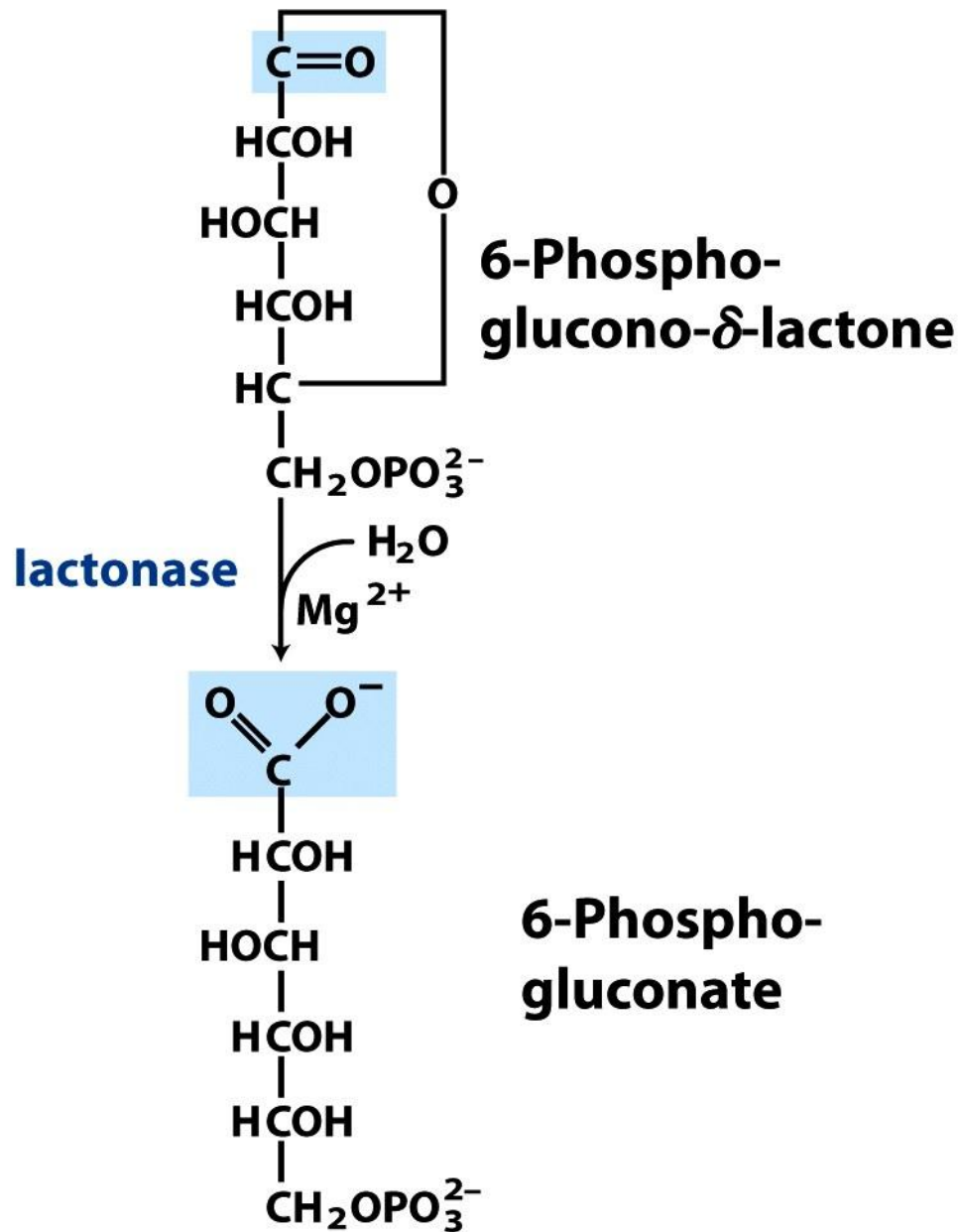


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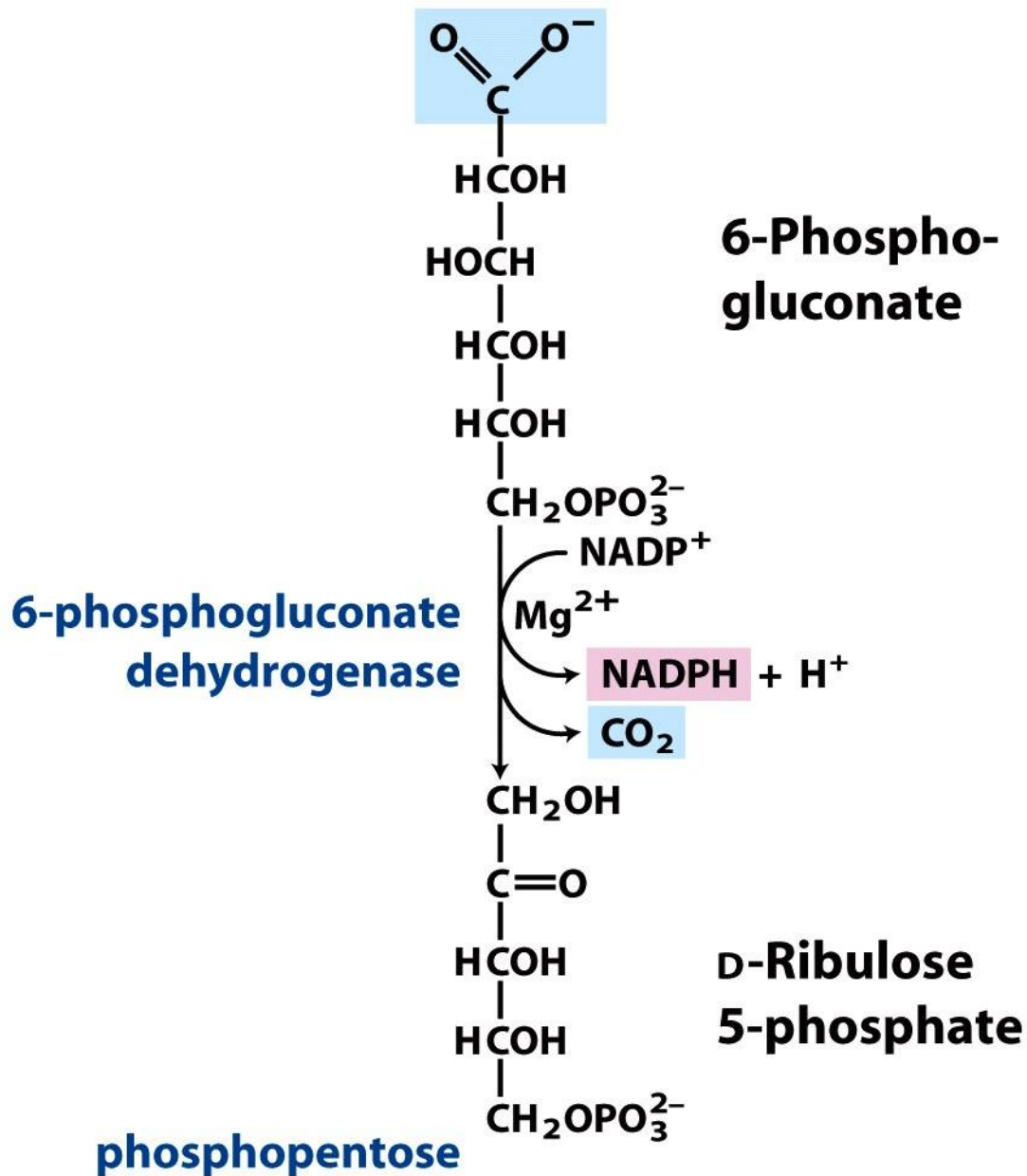


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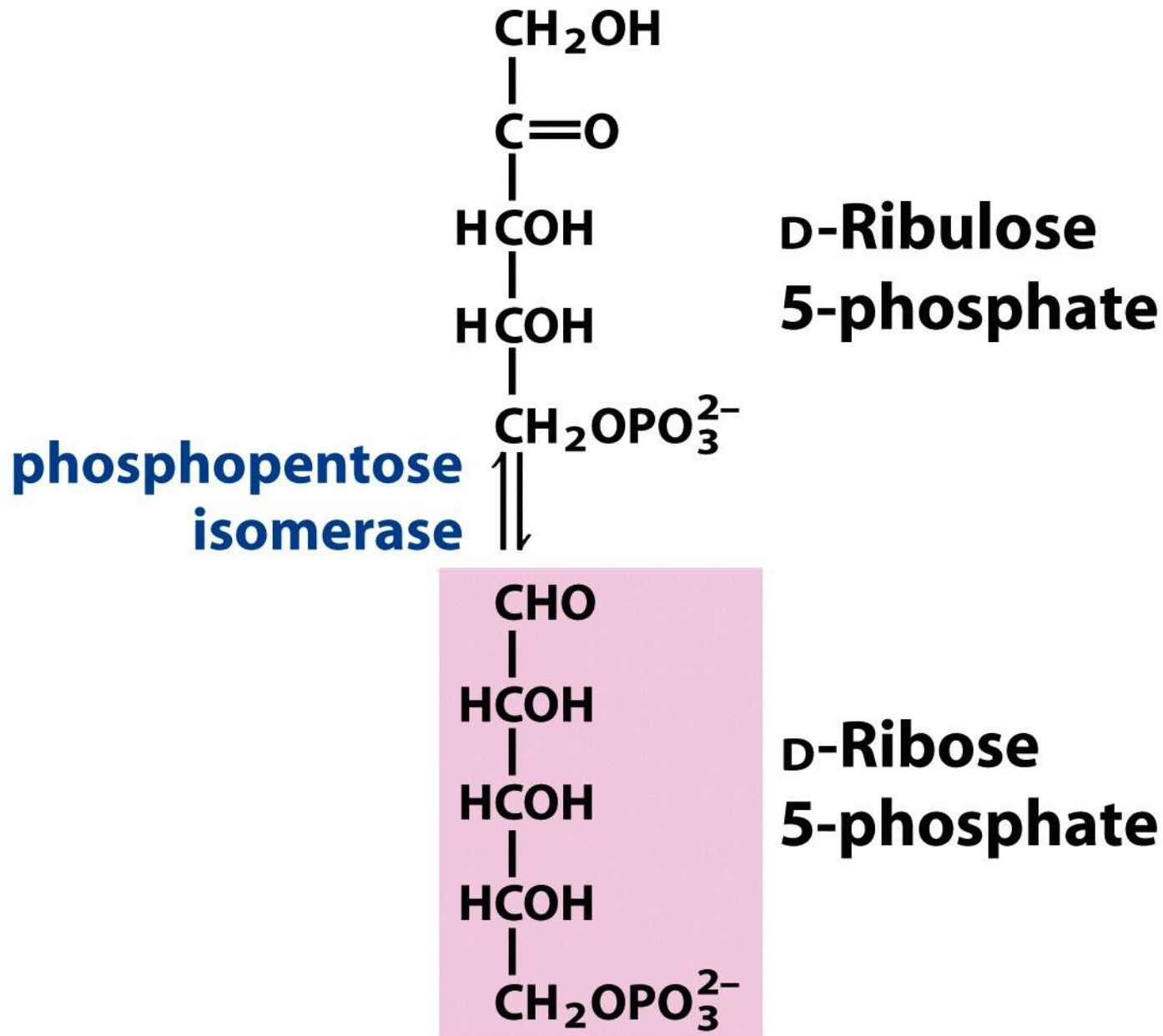
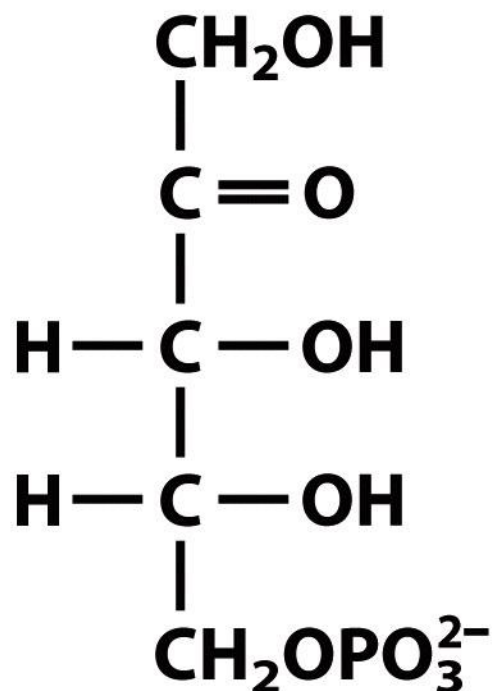


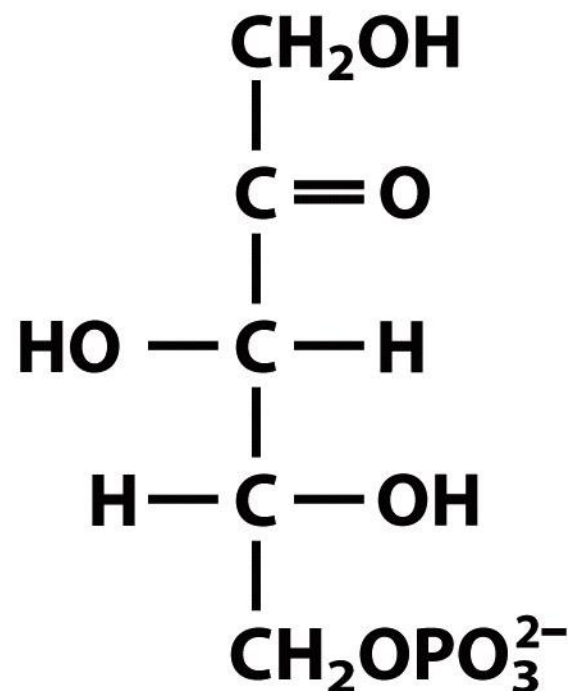
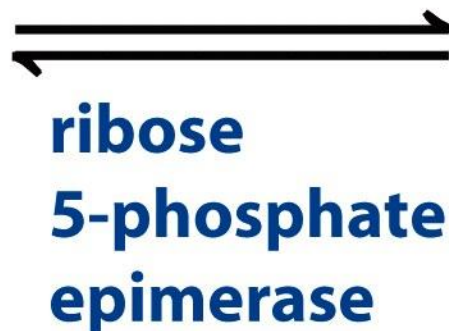
Figure 14-21 part 4

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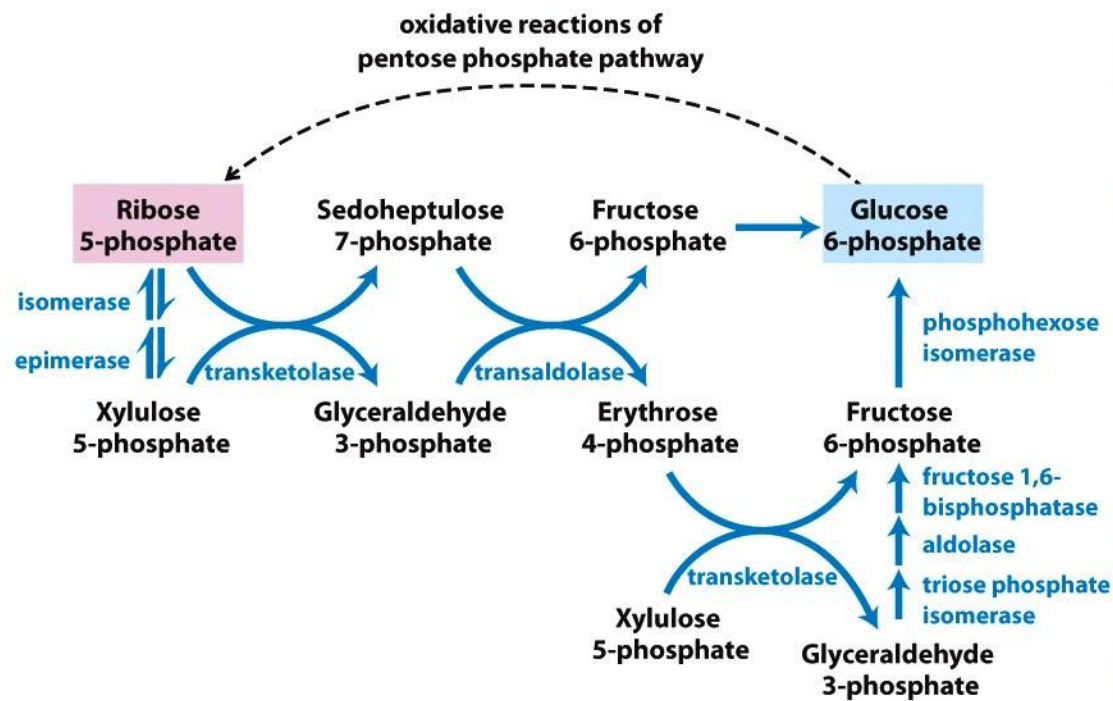
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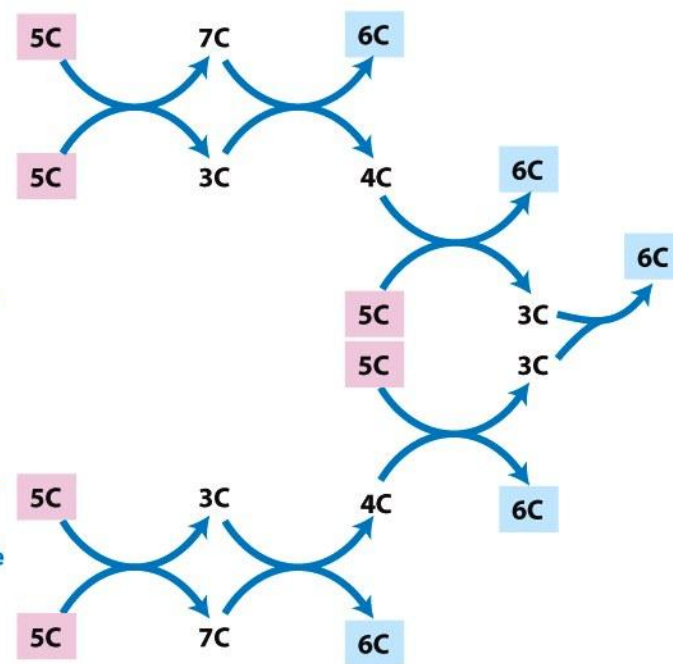
**Ribulose
5-phosphate**



Xylulose 5-phosphate



(a)



(b)

Figure 14-22

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**oxidative reactions of
pentose phosphate pathway**

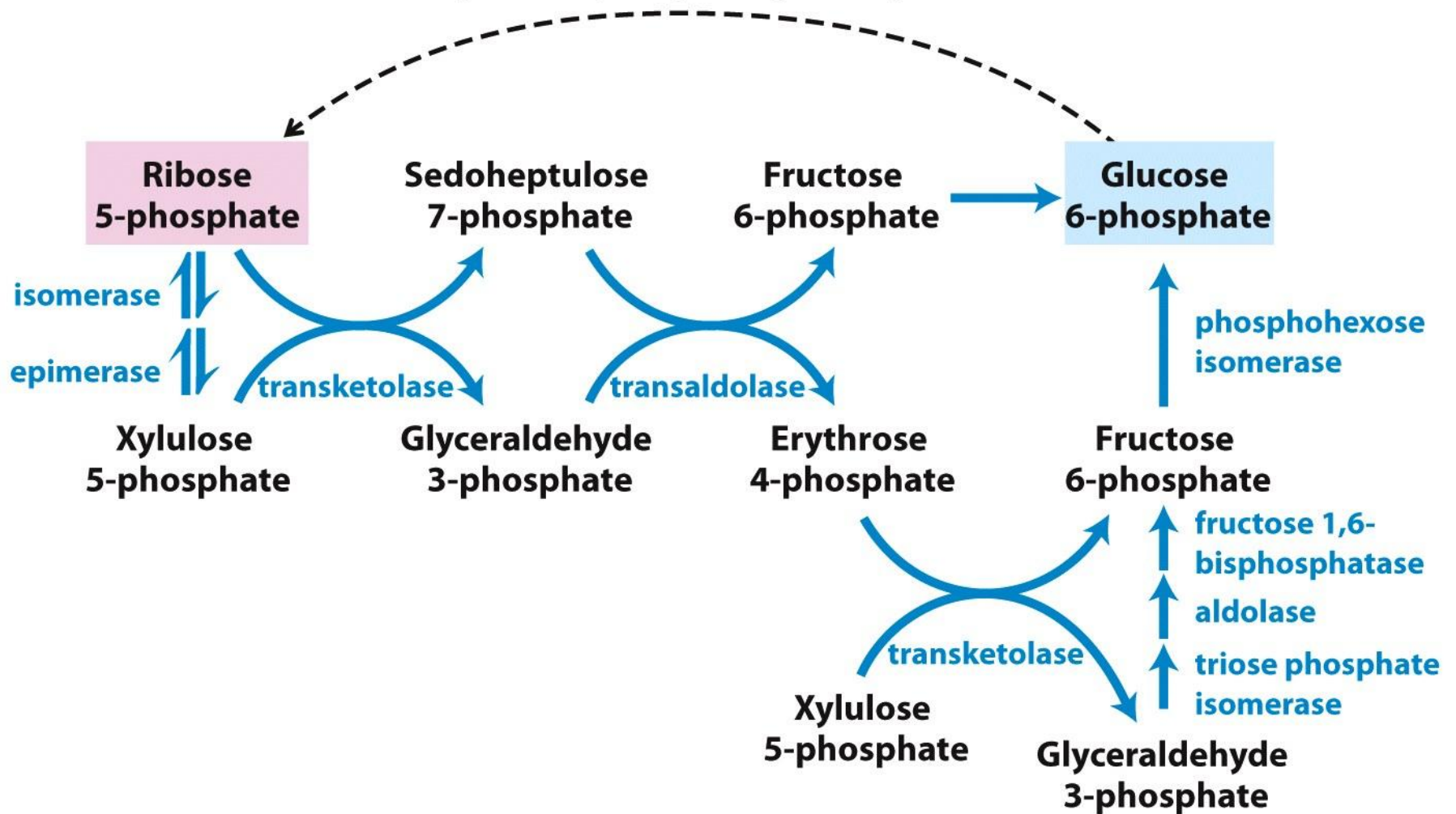


Figure 14-22a

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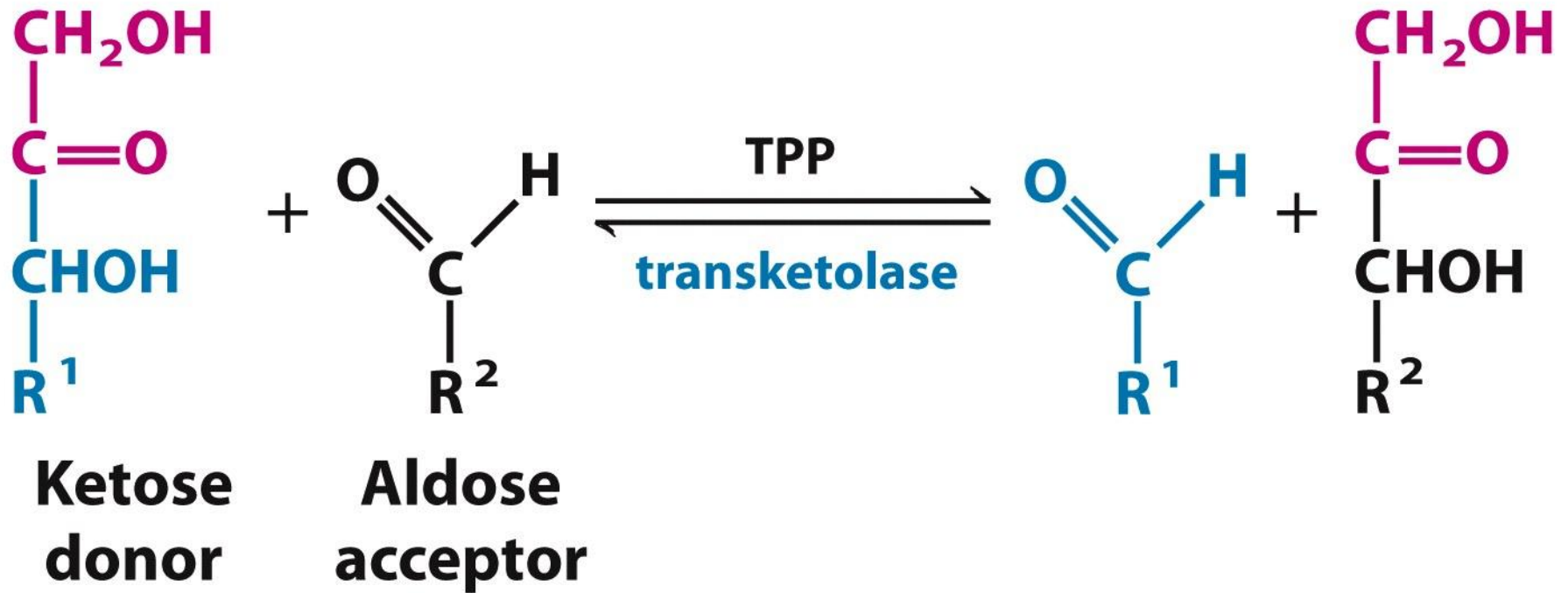


Figure 14-23a

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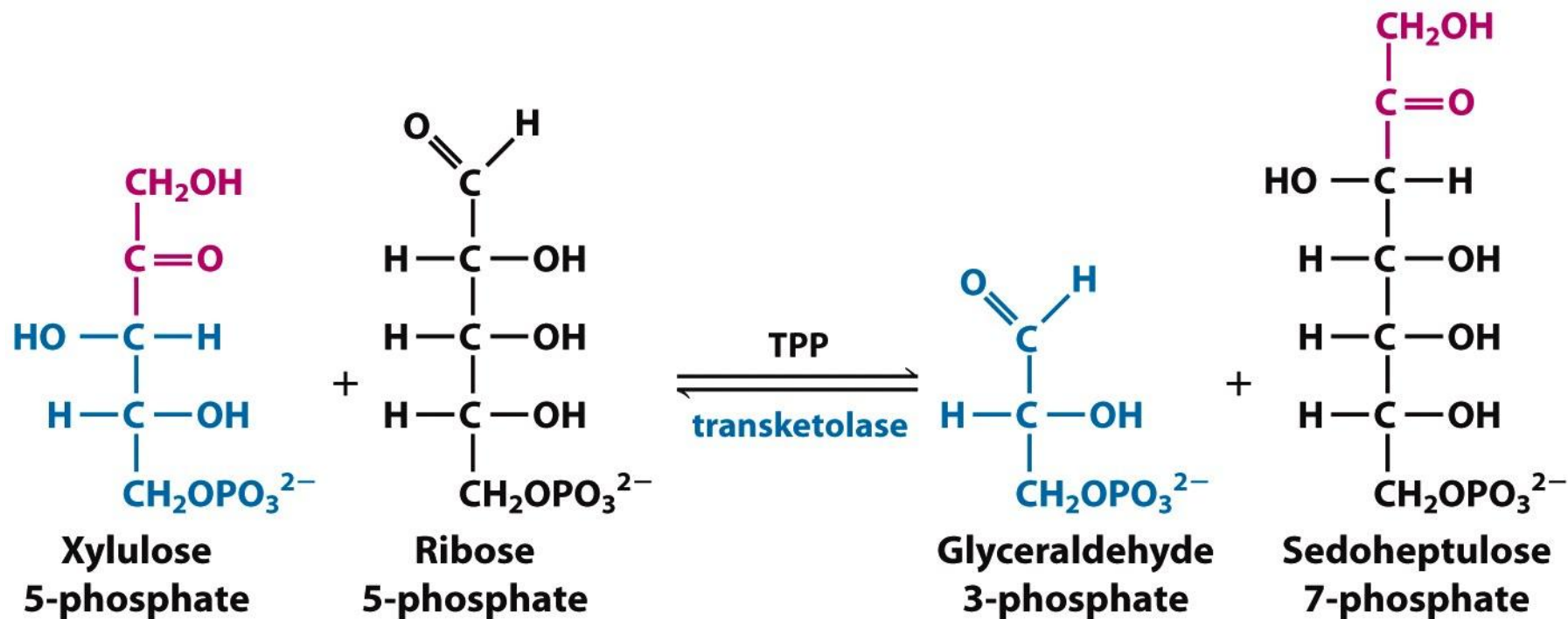


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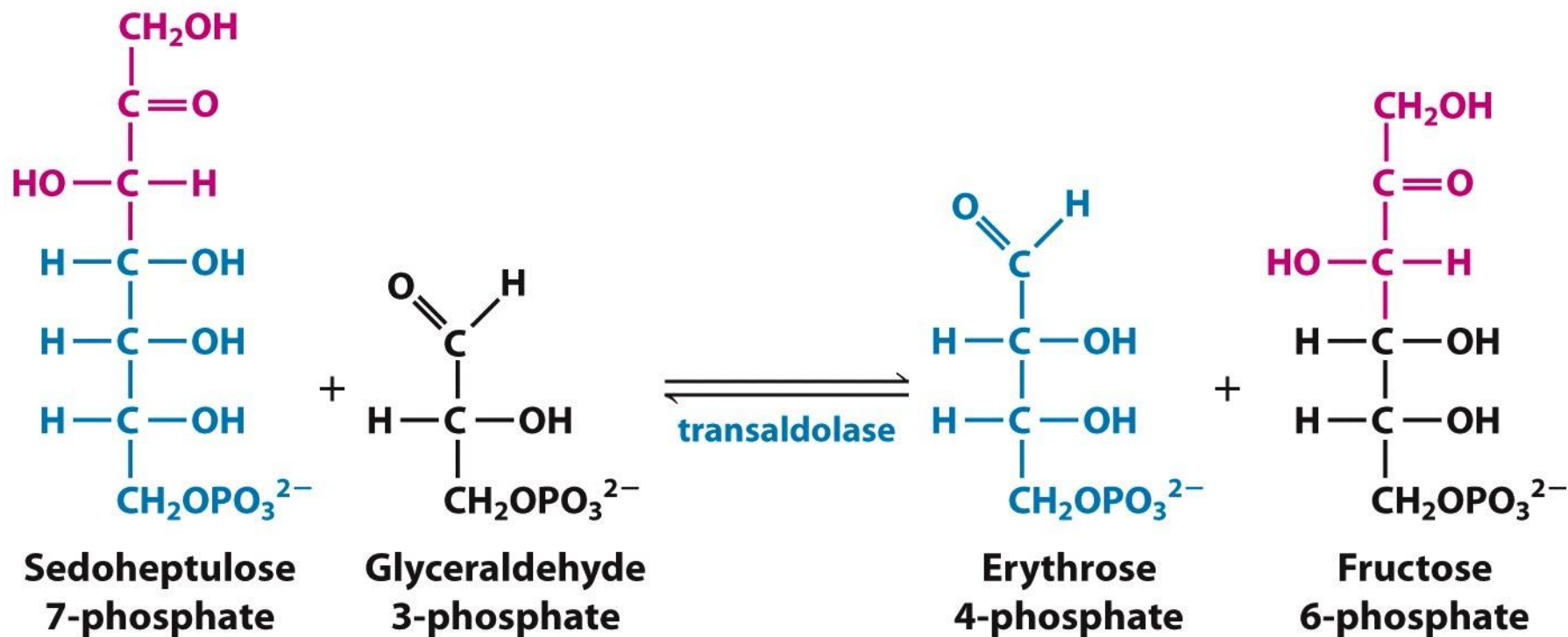


Figure 14-24

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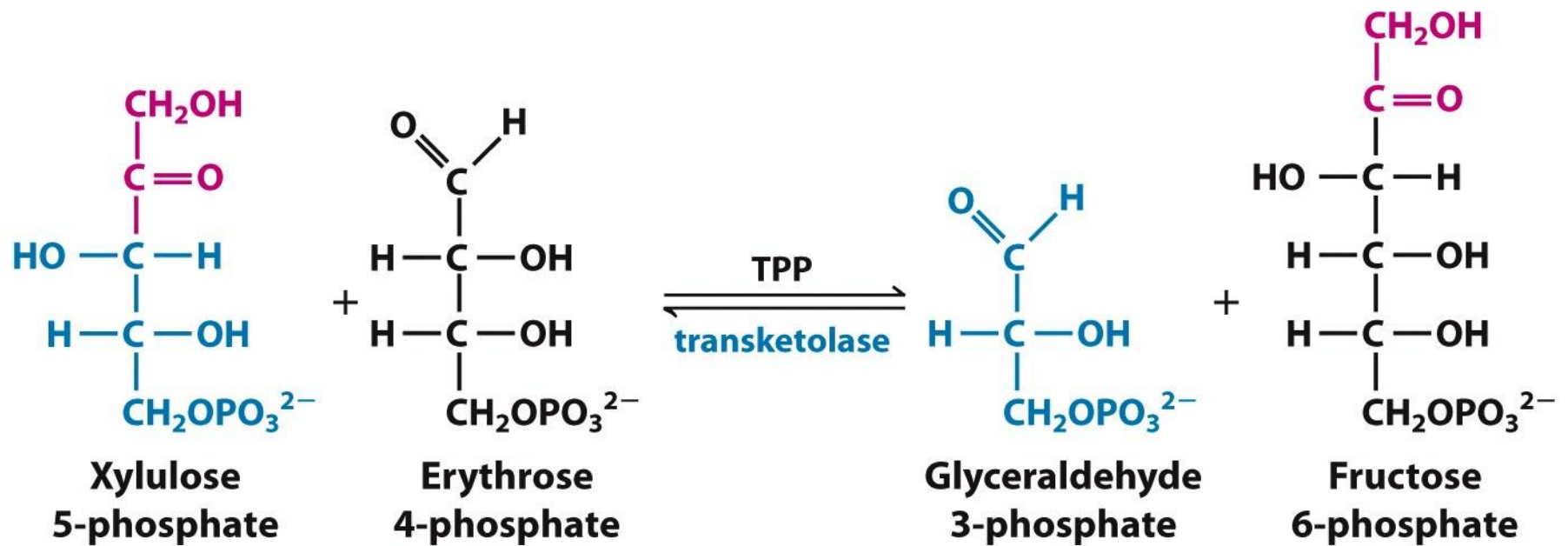
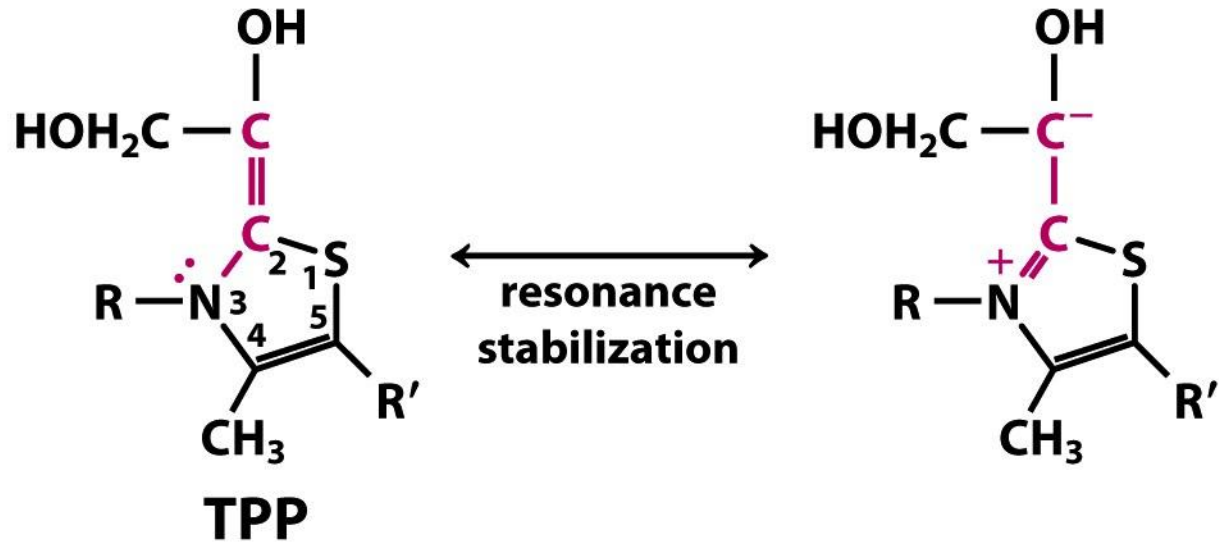


Figure 14-25

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(a) Transketolase



(b) Transaldolase

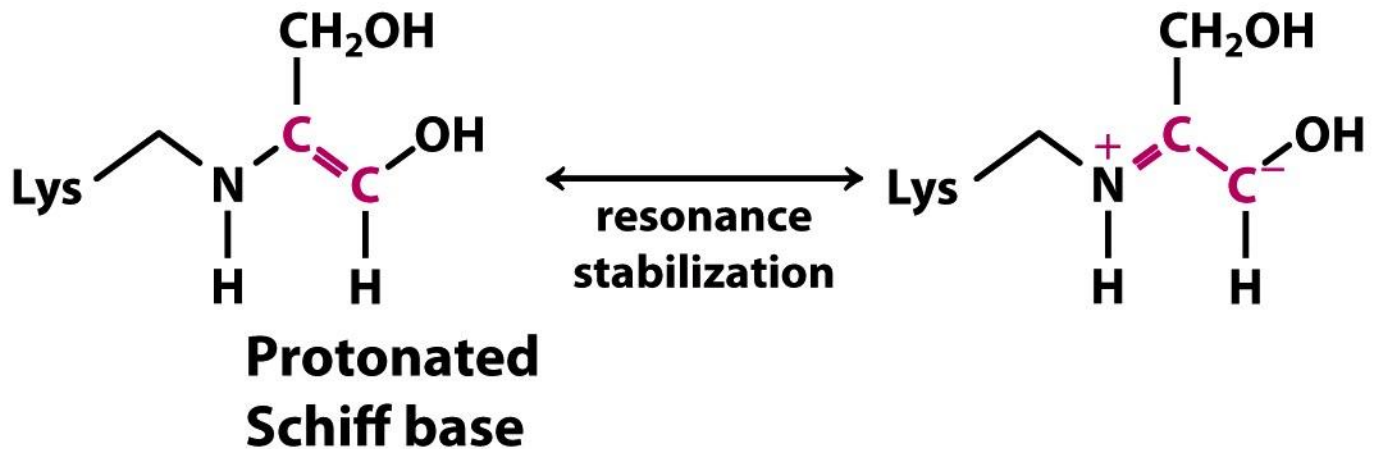


Figure 14-26

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Transketolase

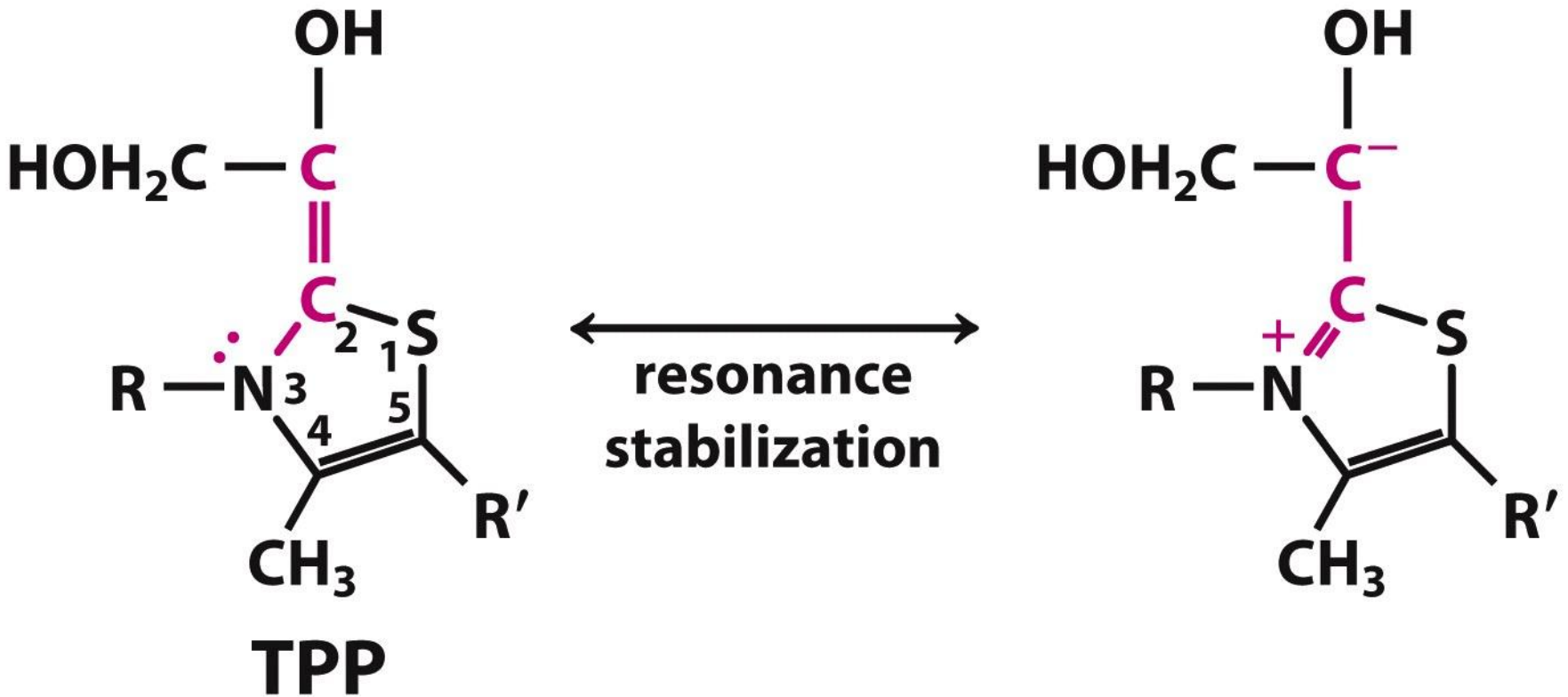


Figure 14-26a

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Transaldolase

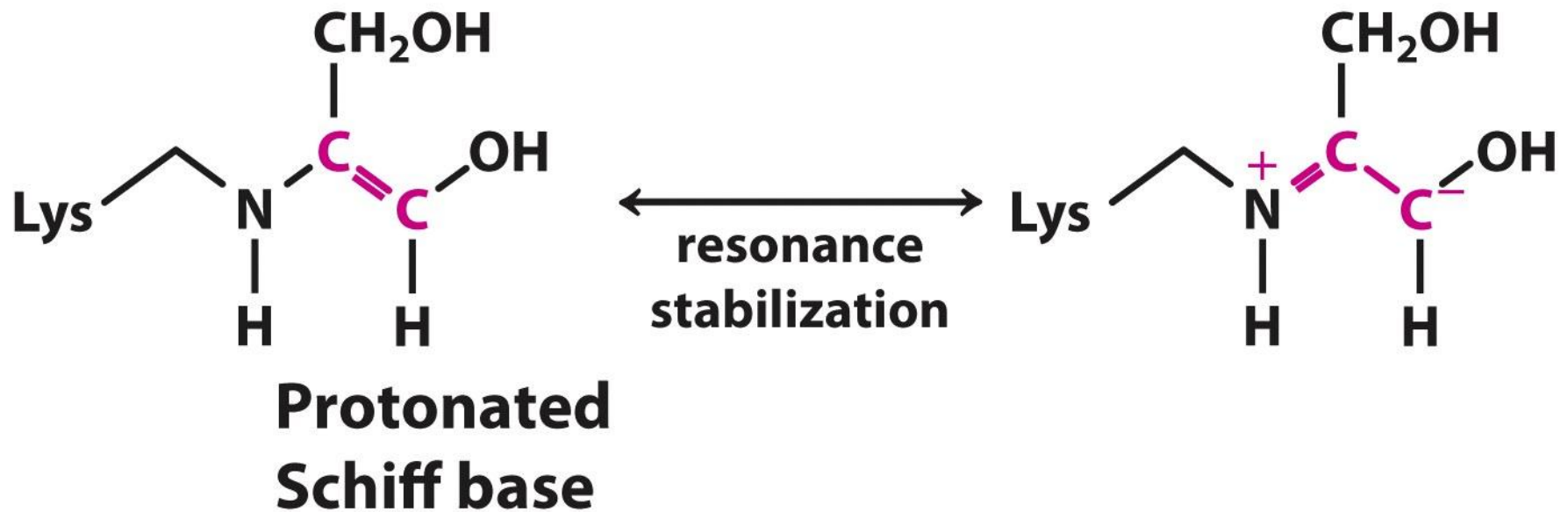


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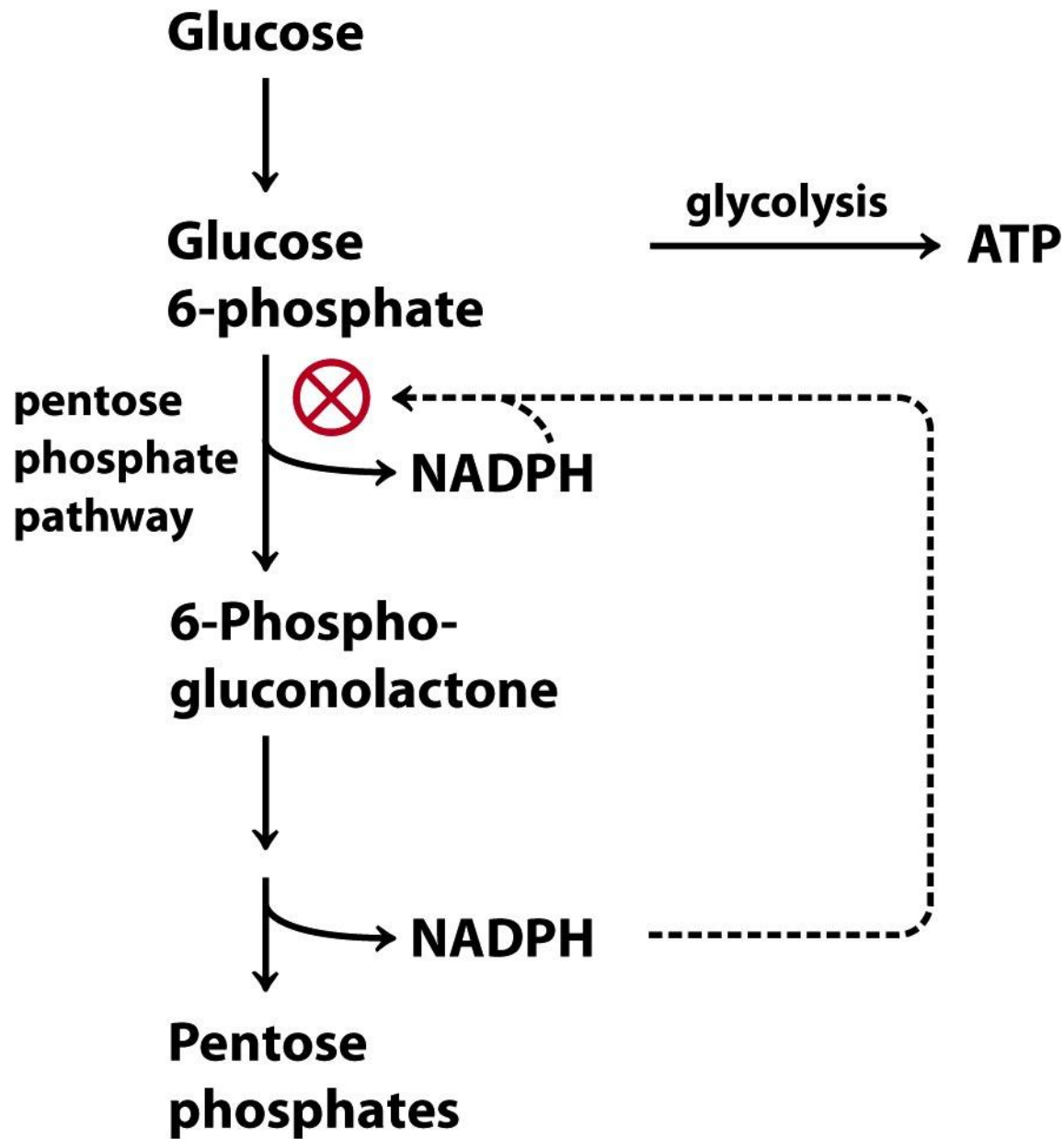


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