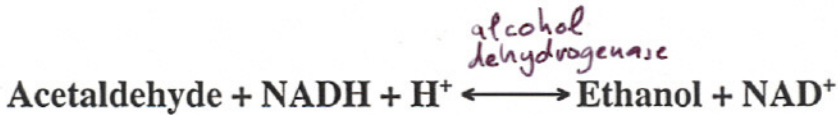
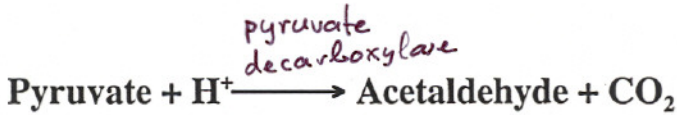


figure 15-3

Three possible catabolic fates of the pyruvate formed in glycolysis. Pyruvate also serves as a precursor in many anabolic reactions, not shown here.

FERMENTATION (S)

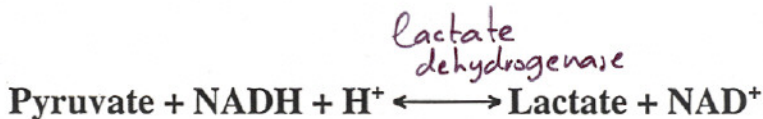
1. Fermentation to alcohol (alcoholic fermentation), in yeast and some microorganisms



Overall Reaction:



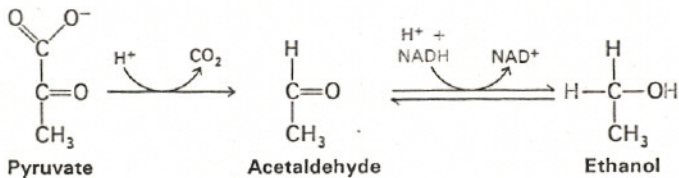
2. Fermentation to lactate, in muscle cells and some microorganisms



Overall Reaction:



Fermentation to alcohol:



Fermentation to lactate:

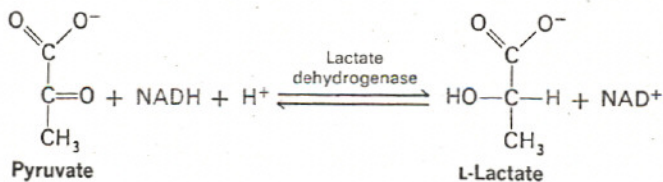
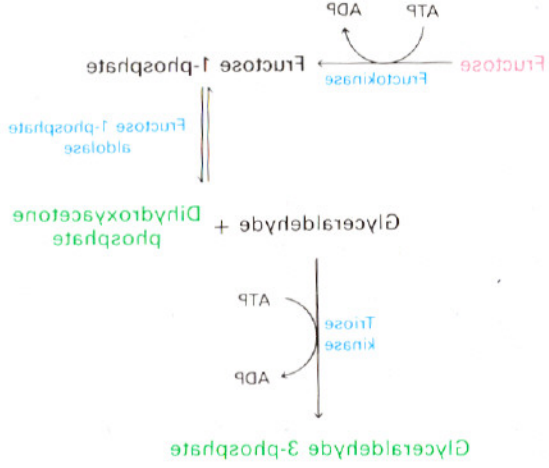
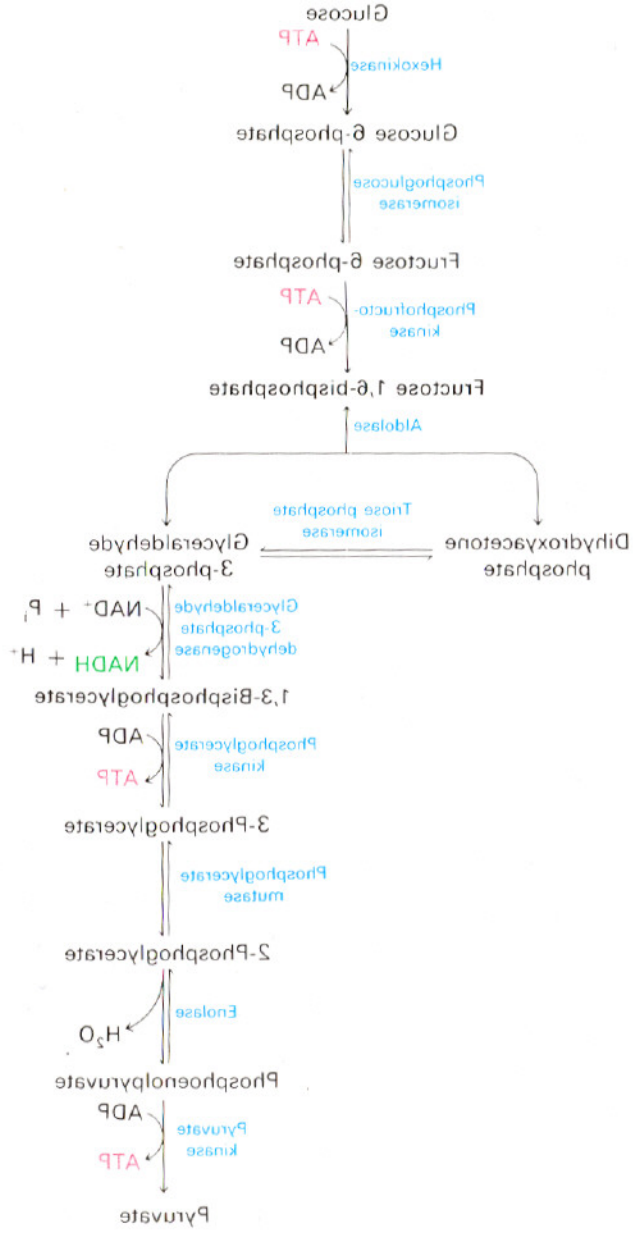


Figure 19-4, page 490; Figure 19-5, page 491



PHOSPHOFRUCTOKINASE

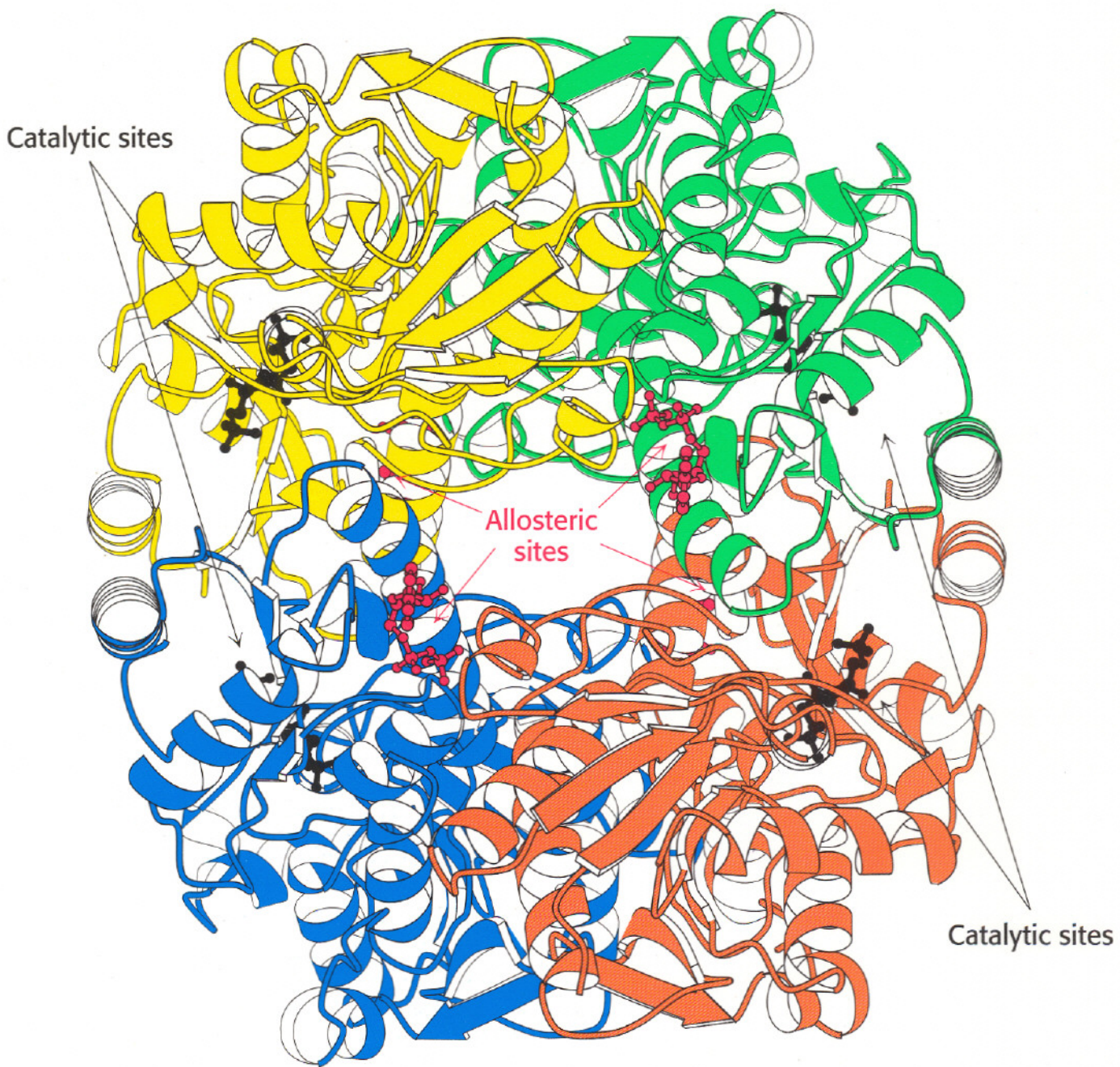
Phosphofructokinase is the most important control element in the mammalian glycolytic pathway

Solved X- Ray Structure

- Homotetramer (340 kDa)
- Has two conformation states, R and T, that are in equilibrium

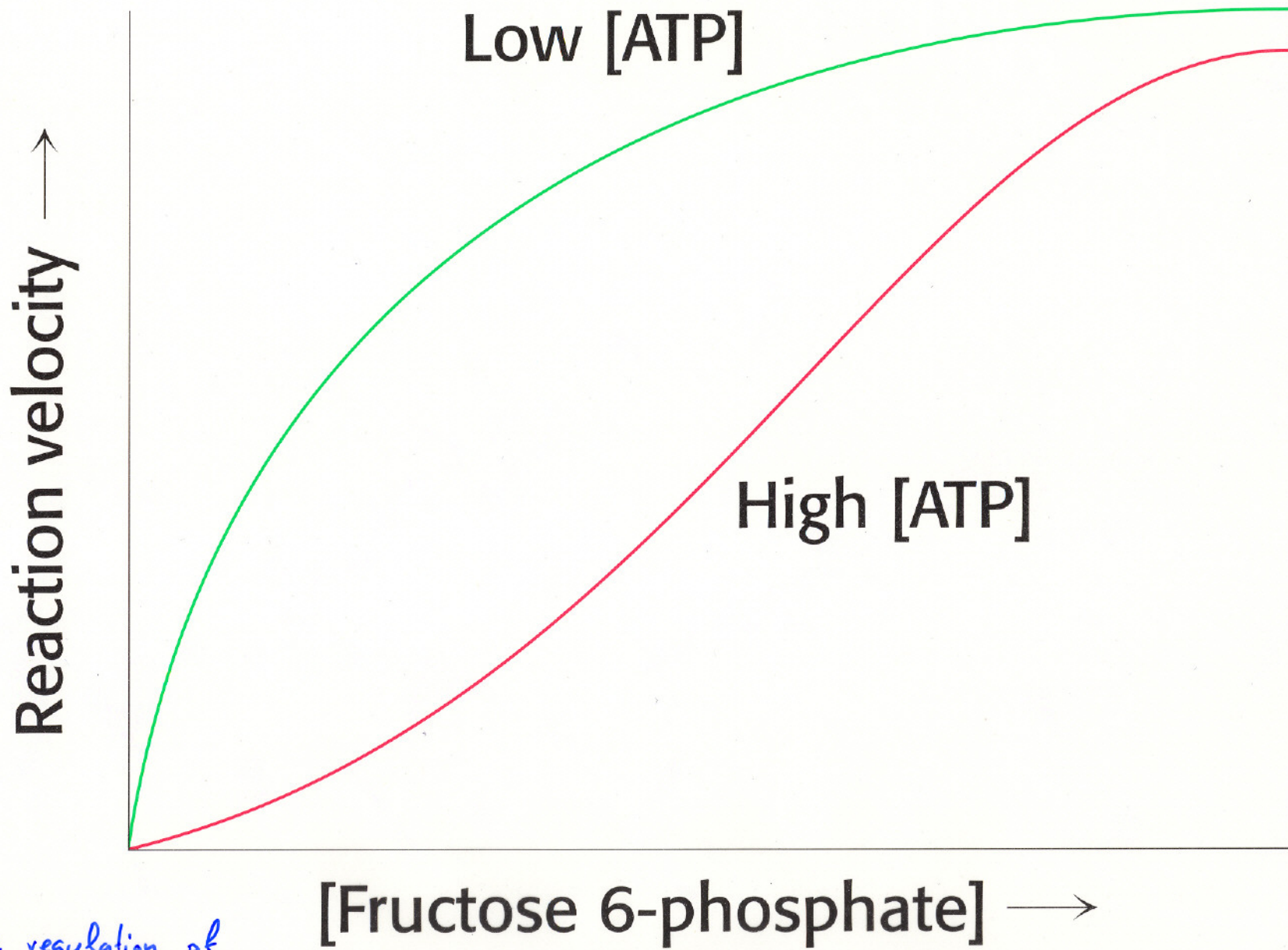
INHIBITORS

1. ATP
2. H^+
3. Citrate (intermediate of TCA Cycle)



Structure of phosphofructokinase

Figure 16-16
Stryer, Tymoczko, & Berg, BIOCHEMISTRY, Fifth Edition.
Copyright © 2002 by W. H. Freeman and Company.



*Allosteric regulation of
phosphofructokinase*

Figure 16-17
Stryer, Tymoczko, & Berg, BIOCHEMISTRY, Fifth Edition.
Copyright © 2002 by W. H. Freeman and Company.

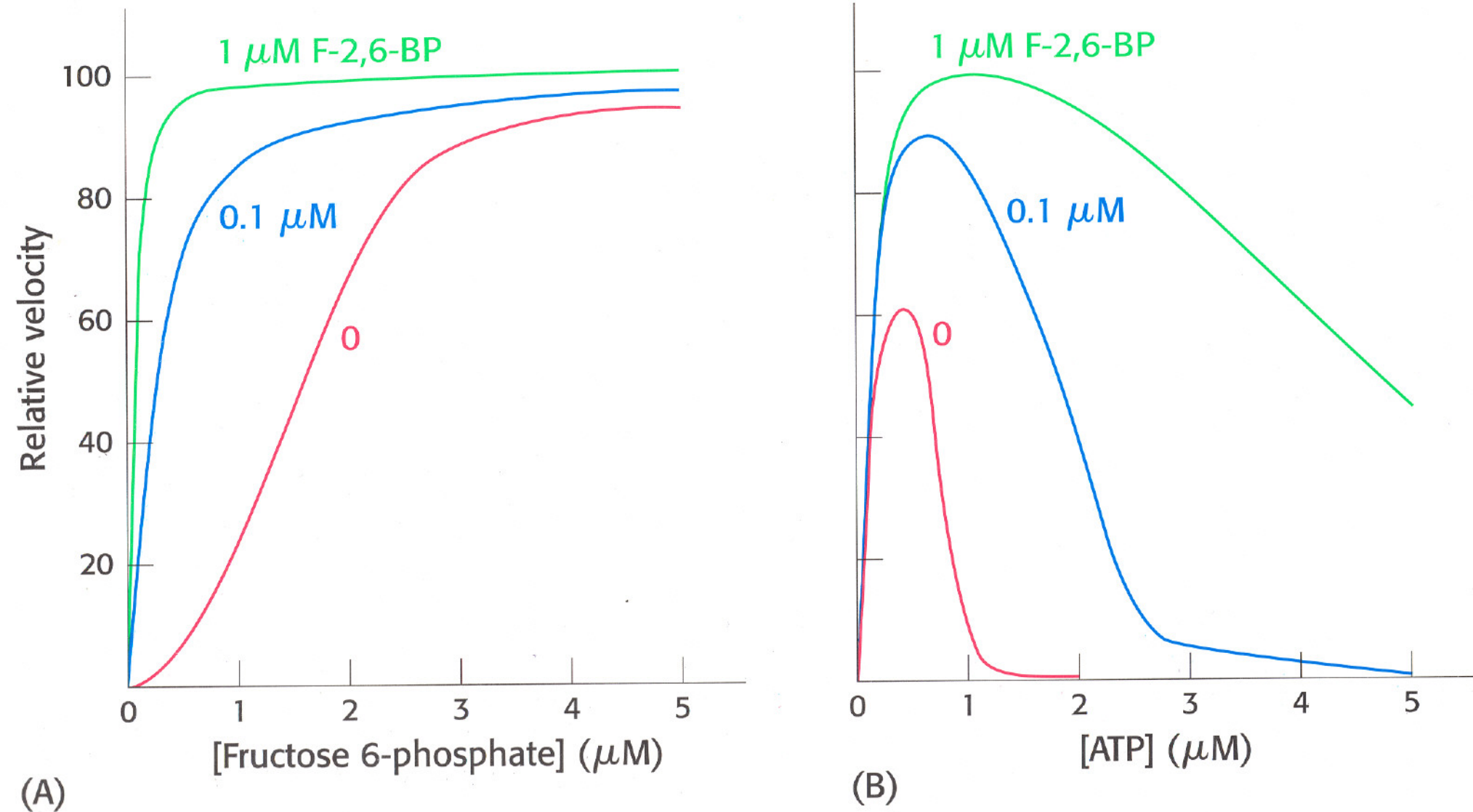
ACTIVATORS

1. AMP, ADP

2. Fructose 2, 6- bisphosphate

(example of Feedforward stimulation)

- **ATP is both a substrate and an allosteric inhibitor**
- **Each enzyme subunit has two binding sites for ATP, a substrate site, and an inhibitor site**
- **The substrate site binds ATP equally well in either conformation(R or T); the inhibitor site binds ATP almost exclusively in the T state**
- **The other substrates preferentially bind to the R state**



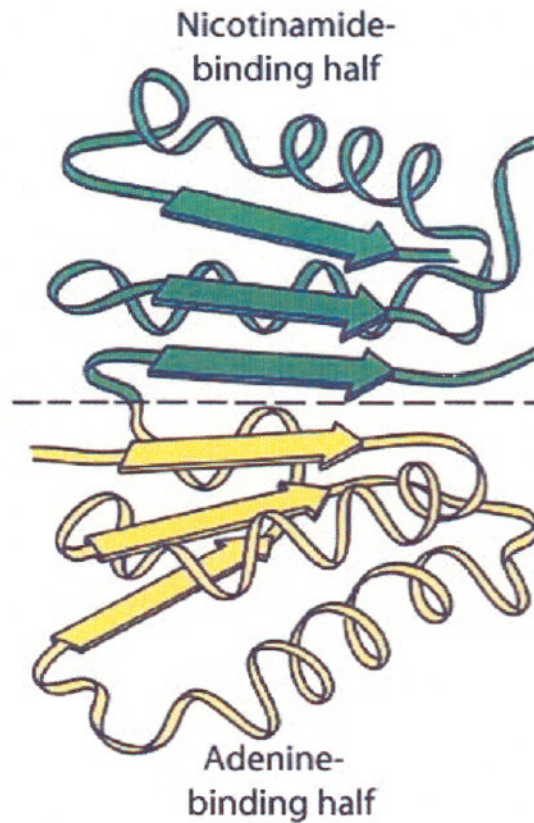
Activation of phosphofructokinase by fructose 2,6-bisphosphate
 The inhibitory effect of ATP is reversed

Conformation of hexokinase: Induced fit



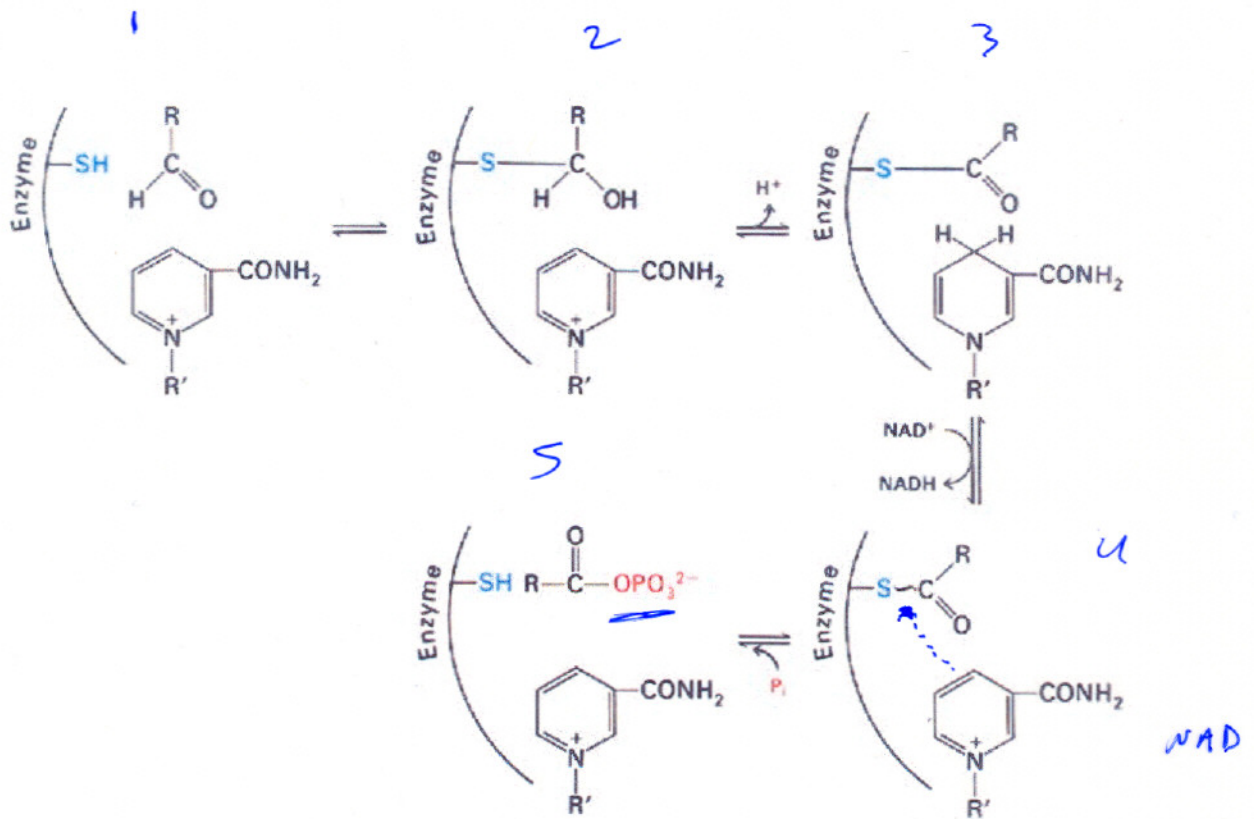
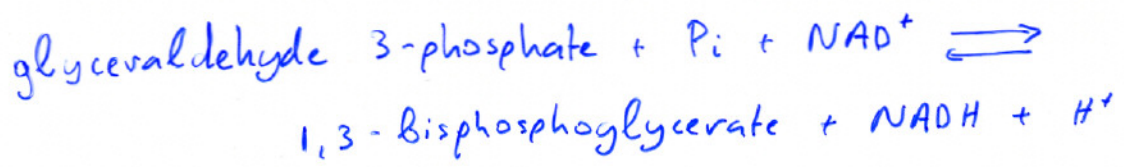
Figure 19-14, page 499

Schematic diagram of the NAD^+ -binding region in dehydrogenases



- alcohol dehydrogenase
- lactate dehydrogenase
- malate dehydrogenase
- glyceraldehyde 3-phosphate dehydrogenase

Fig 19-12



Catalytic mechanism of
glyceraldehyde 3-phosphate dehydrogenase

Fig 14-18

GLUCOSE TRANSPORTERS

- Integral membrane proteins of mammalian cells
- Named GLUT 1 to 5
- Single polypeptide chain ~ 500 a. a.
- 12 transmembrane α -helices

I. GLUT 1 & GLUT 3

- Present in nearly all mammalian cells
- Responsible for basal glucose uptake
- Continually transport glucose at an essential constant rate

II. GLUT 5

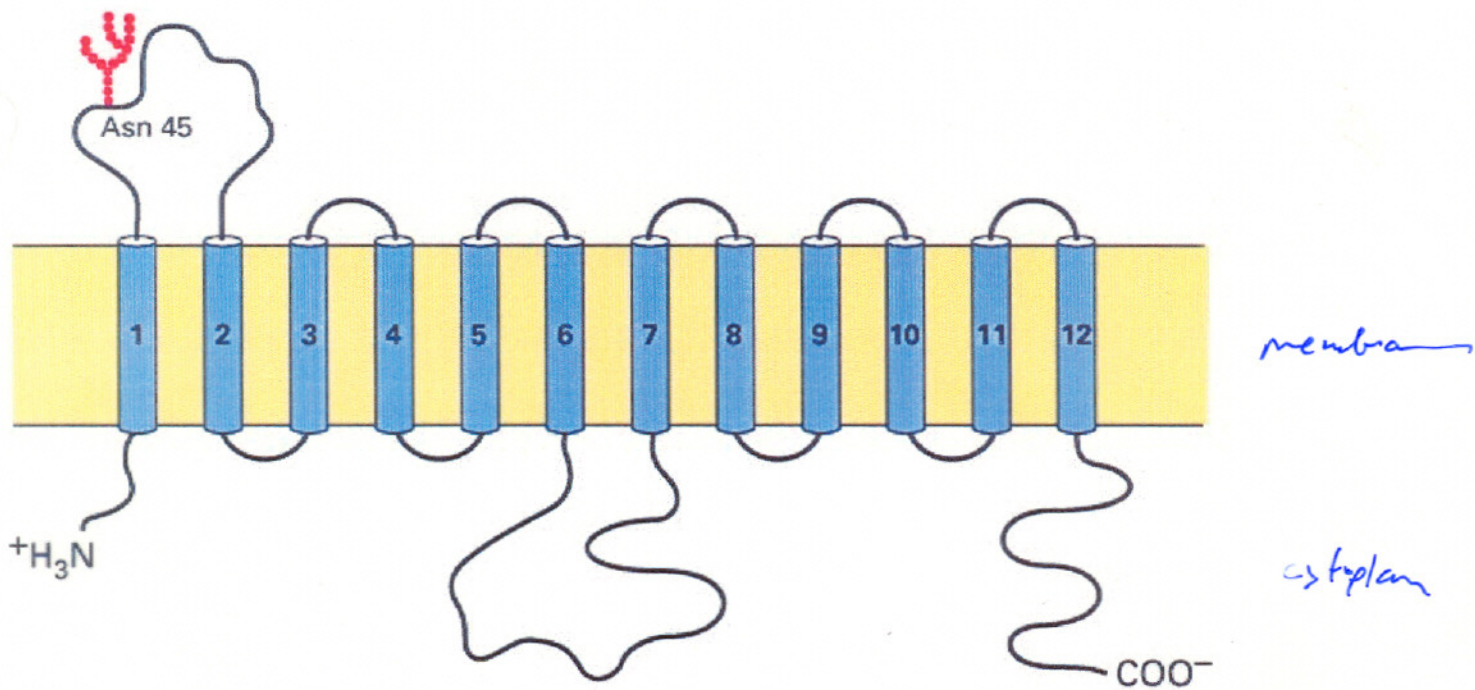
- **Present in the small intestine**
- **Works in tandem with the Na⁺– glucose symporter in the absorption of glucose from the gut**
- **Releases glucose into bloodstream**

III. GLUT 2

- **Present in liver and pancreatic β cells**

IV. GLUT 4

- **Mediates entry of glucose into muscle and fat cells**



BREWING BEER

Beer is made by alcohol fermentation of the carbohydrates in cereal grains (seeds) by yeast glycolytic enzymes

PROCEDURE:

- 1. Malting:** The seeds are allowed to germinate until they form the hydrolytic enzymes required to break down their polysaccharides. The product is called “malt”
- 2. Preparing the wort:** The malt is mixed with water and then mashed or crushed. Digestion of the polysaccharides. Separation of the cell matter from the liquid wort.
- 3. Adding the yeast cells:** First aerobic glycolysis and citric acid cycle. When oxygen is consumed, anaerobic fermentation
- 4. Adjustment of the amount of foam of head of the beer.**

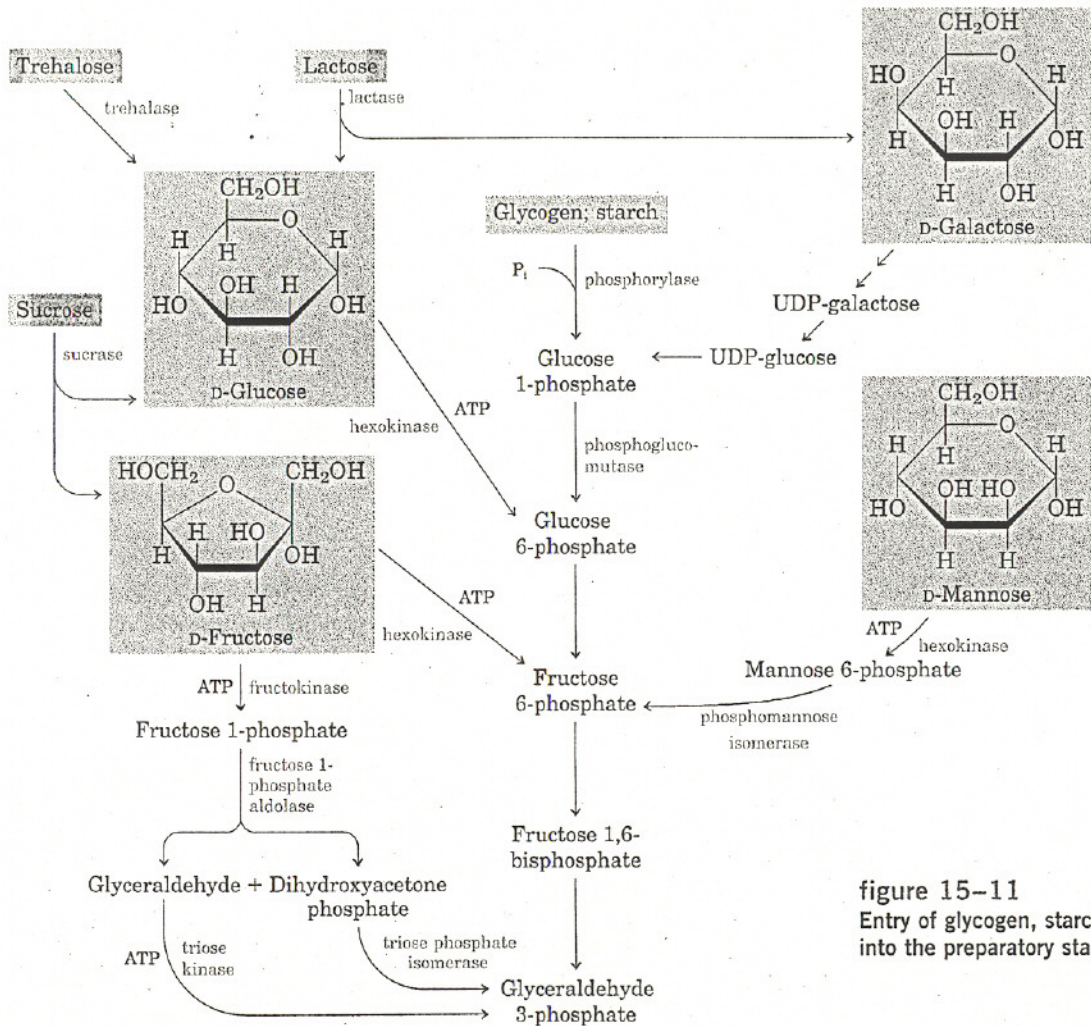
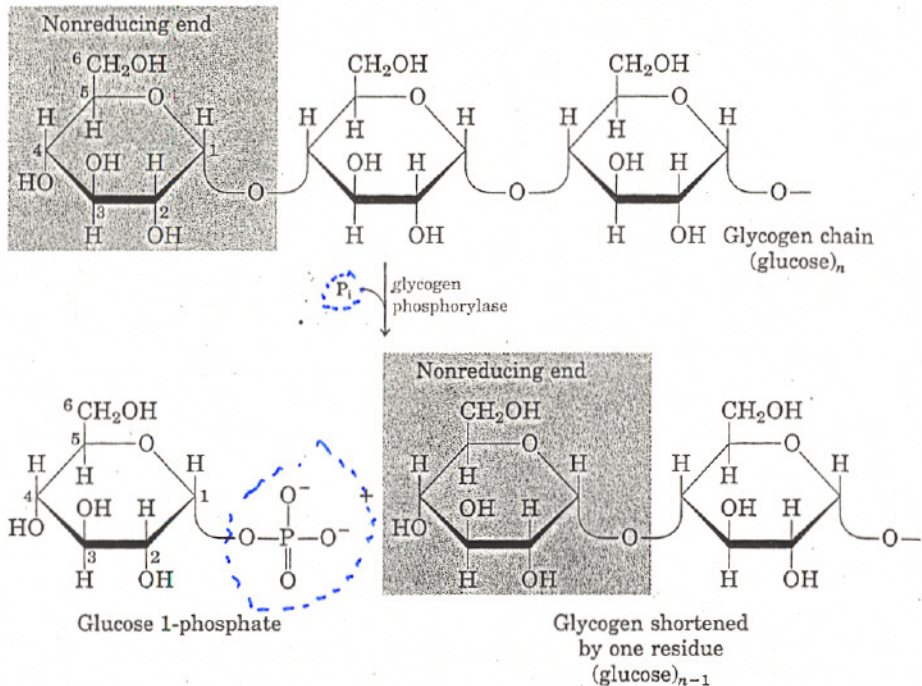


figure 15-11

Entry of glycogen, starch, disaccharides, and hexoses into the preparatory stage of glycolysis.

Figure 15-12

Removal of a terminal glucose residue from the nonreducing end of a glycogen chain by glycogen phosphorylase. This process is repetitive; the enzyme removes successive glucose residues until it reaches a fourth glucose unit from a branch point (see Figure 15-13). Amylopectin is degraded in a similar fashion by starch phosphorylase.



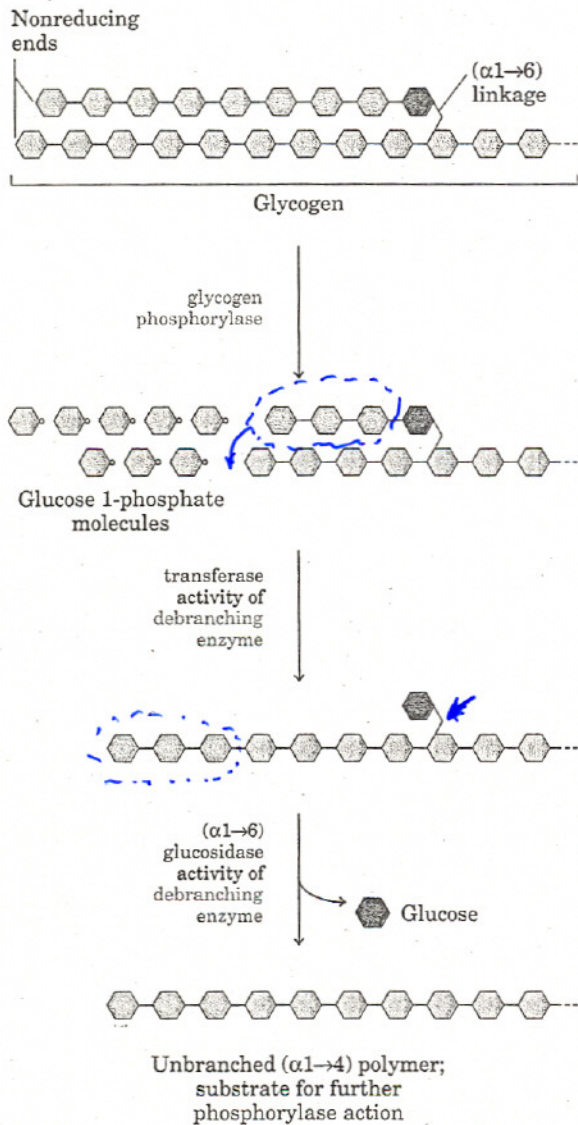


figure 15-13

Glycogen breakdown near an ($\alpha 1 \rightarrow 6$) branch point.

Following the sequential removal of terminal glucose residues by glycogen phosphorylase (see Fig. 15-12), glucose residues near a branch are removed in a two-step process that requires a bifunctional "debranching enzyme." First, the transferase activity of the enzyme shifts a block of three glucose residues from the branch to a nearby nonreducing end, to which they are re-attached in ($\alpha 1 \rightarrow 4$) linkage. The single glucose residue remaining at the branch point, in ($\alpha 1 \rightarrow 6$) linkage, is then released as free glucose by the debranching enzyme's ($\alpha 1 \rightarrow 6$) glucosidase activity. The glucose residues are shown here in shorthand form.