

# Microbial Metabolism

- Sum of all biochemical reactions
- Catabolic and anabolic
- Enzymes:
  - Catalyze reactions
  - Lower energy of activation
  - Do not change equilibrium
  - Specific
  - Cofactors or coenzymes
- Oxidation reduction reactions

# Patterns of Energy Production

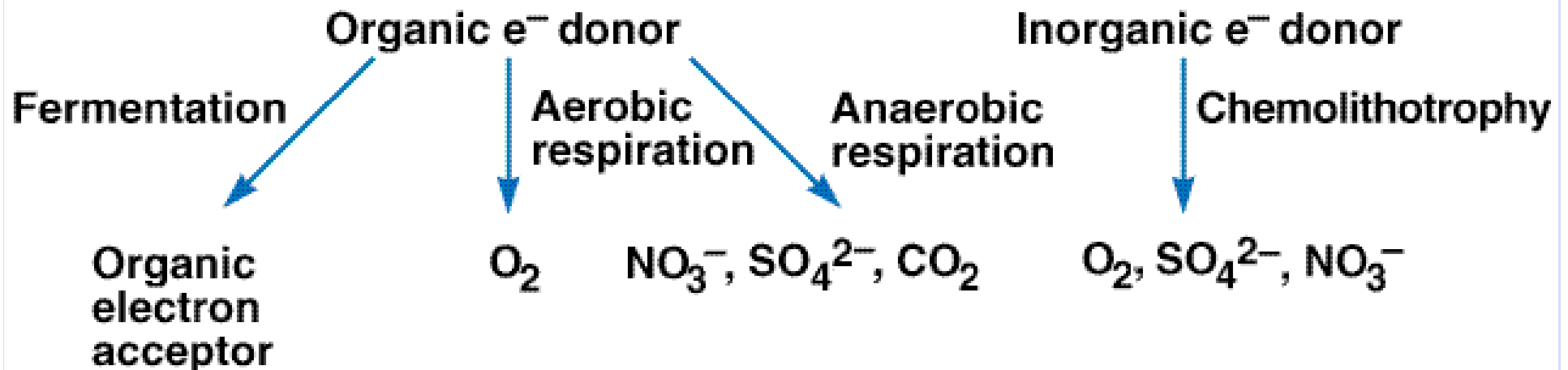


Fig. 9.2

# Three Stages of Catabolism

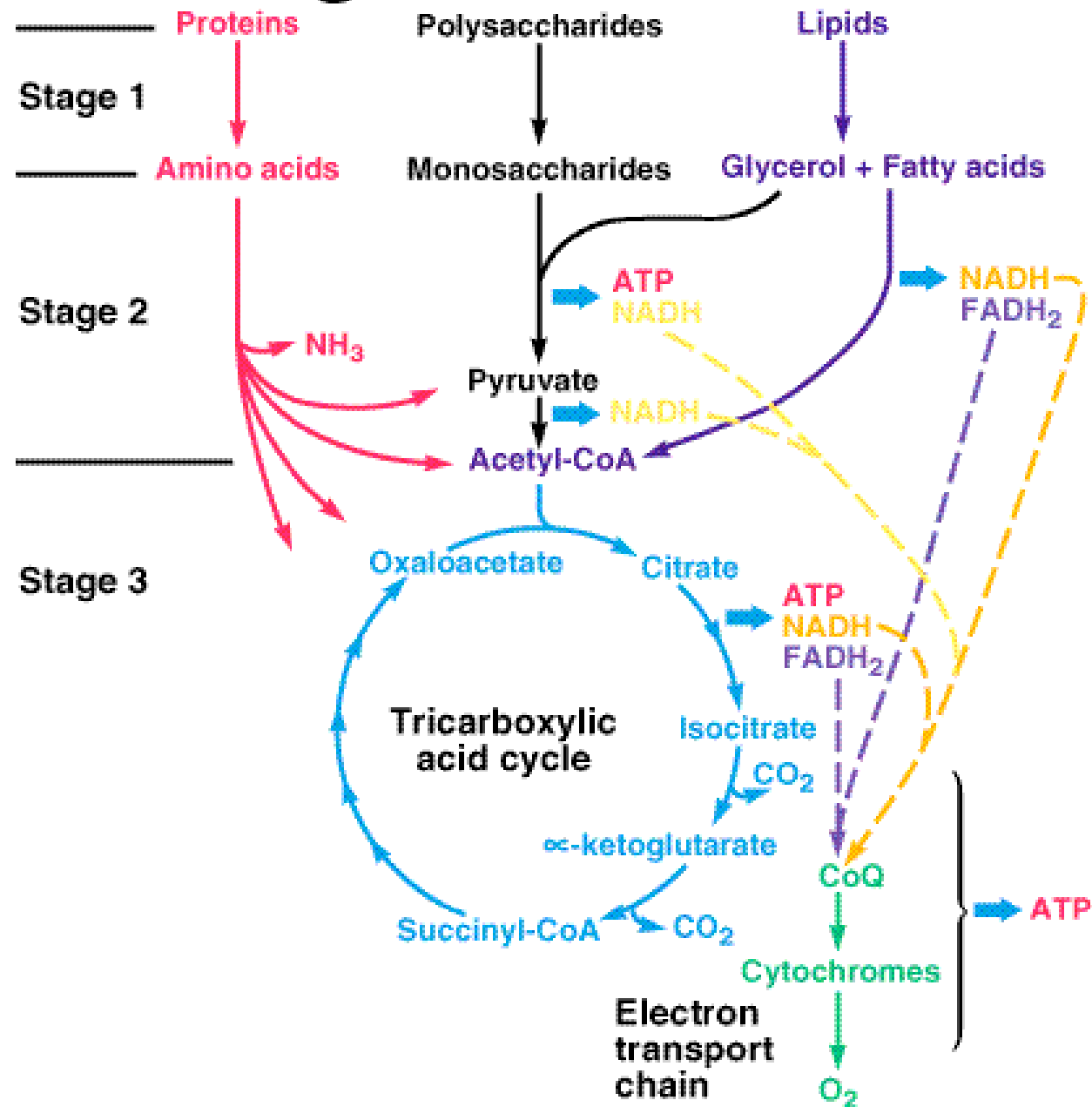


Fig. 9.3

# Glycolysis

4 ATPs  
-2 ATPs  

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2 ATPs

2 NADH  
2 Pyruvates

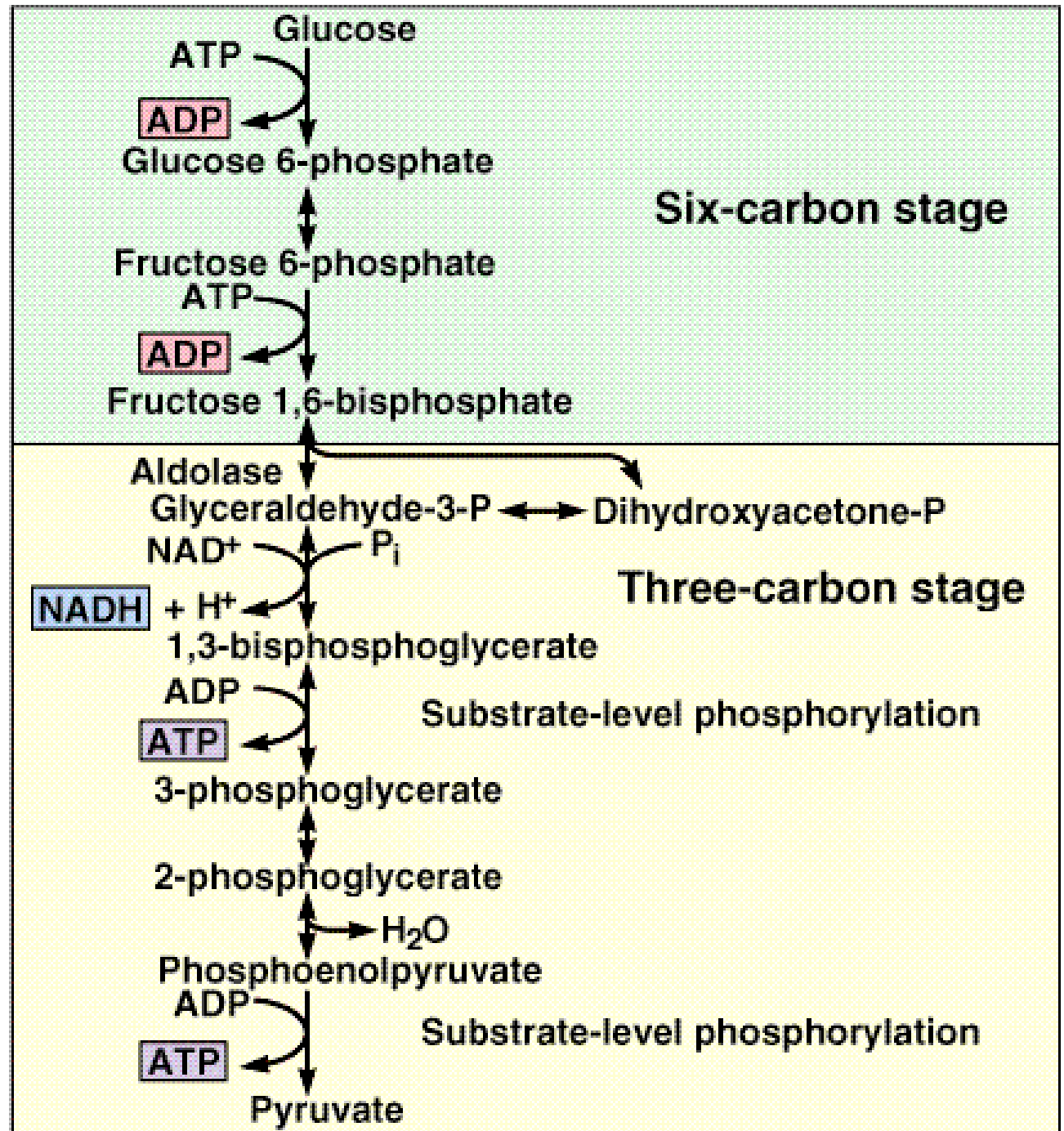
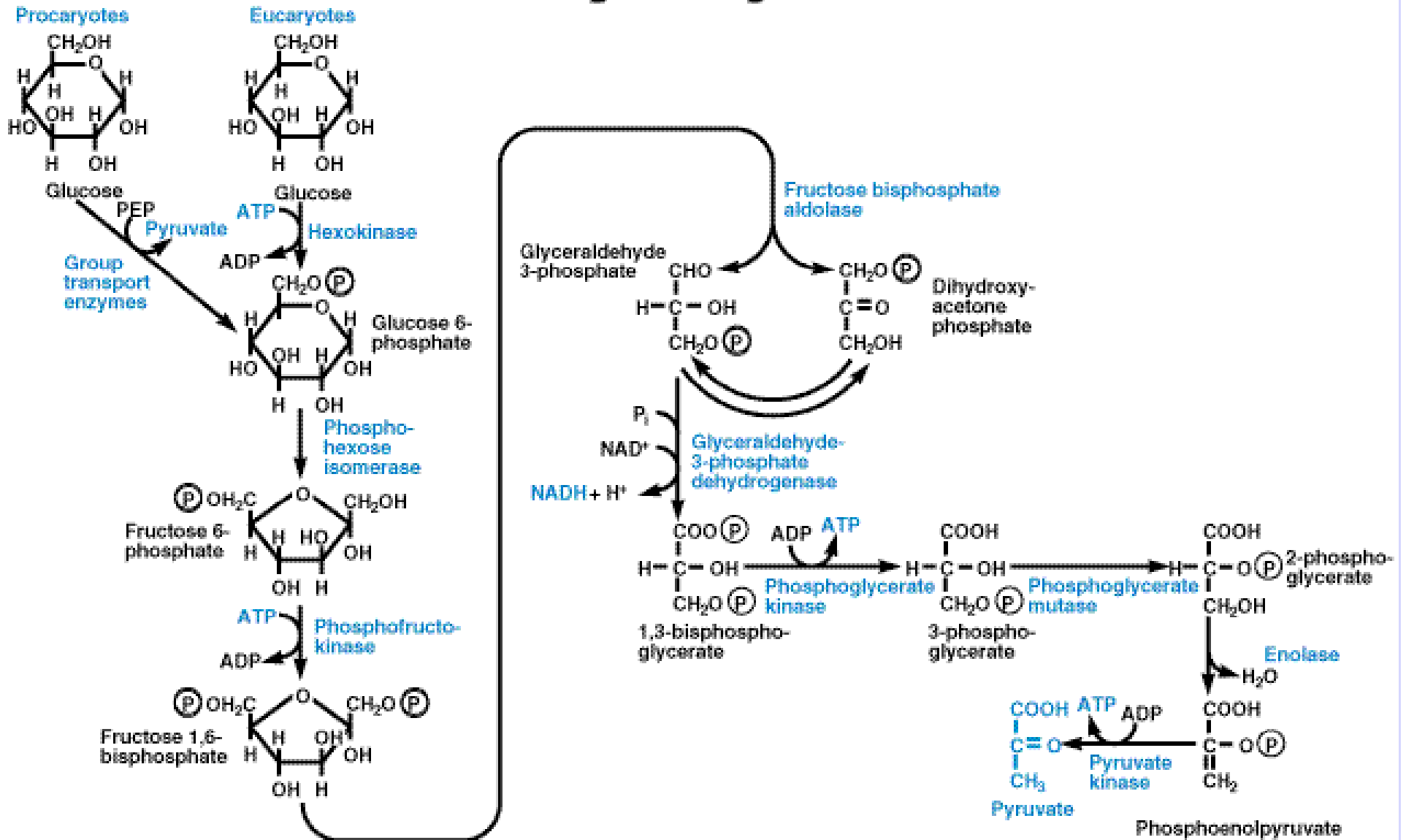


Fig. 9.5

# Glycolysis



# Bacterial PTS Transport

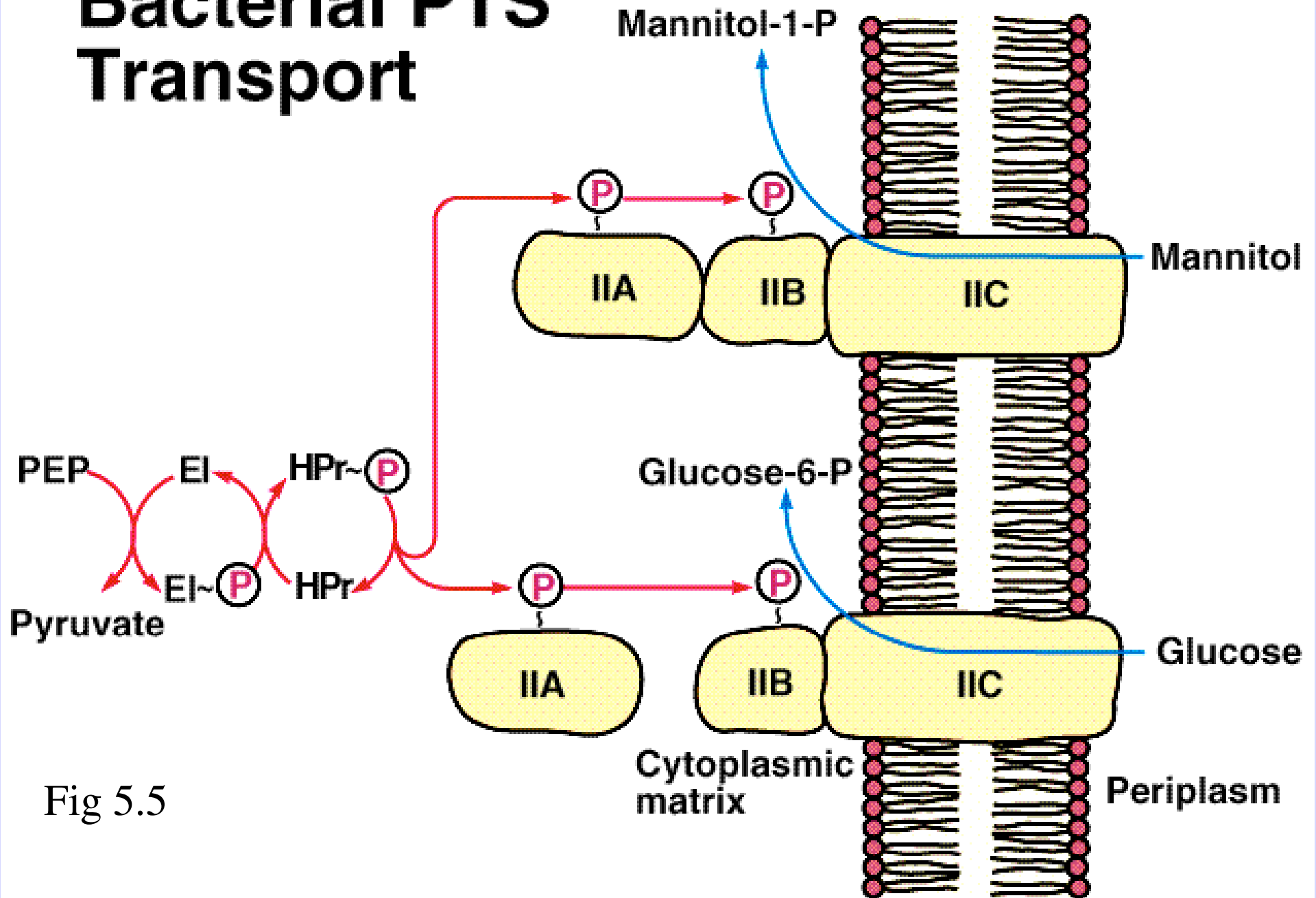


Fig 5.5

# Three Stages of Catabolism

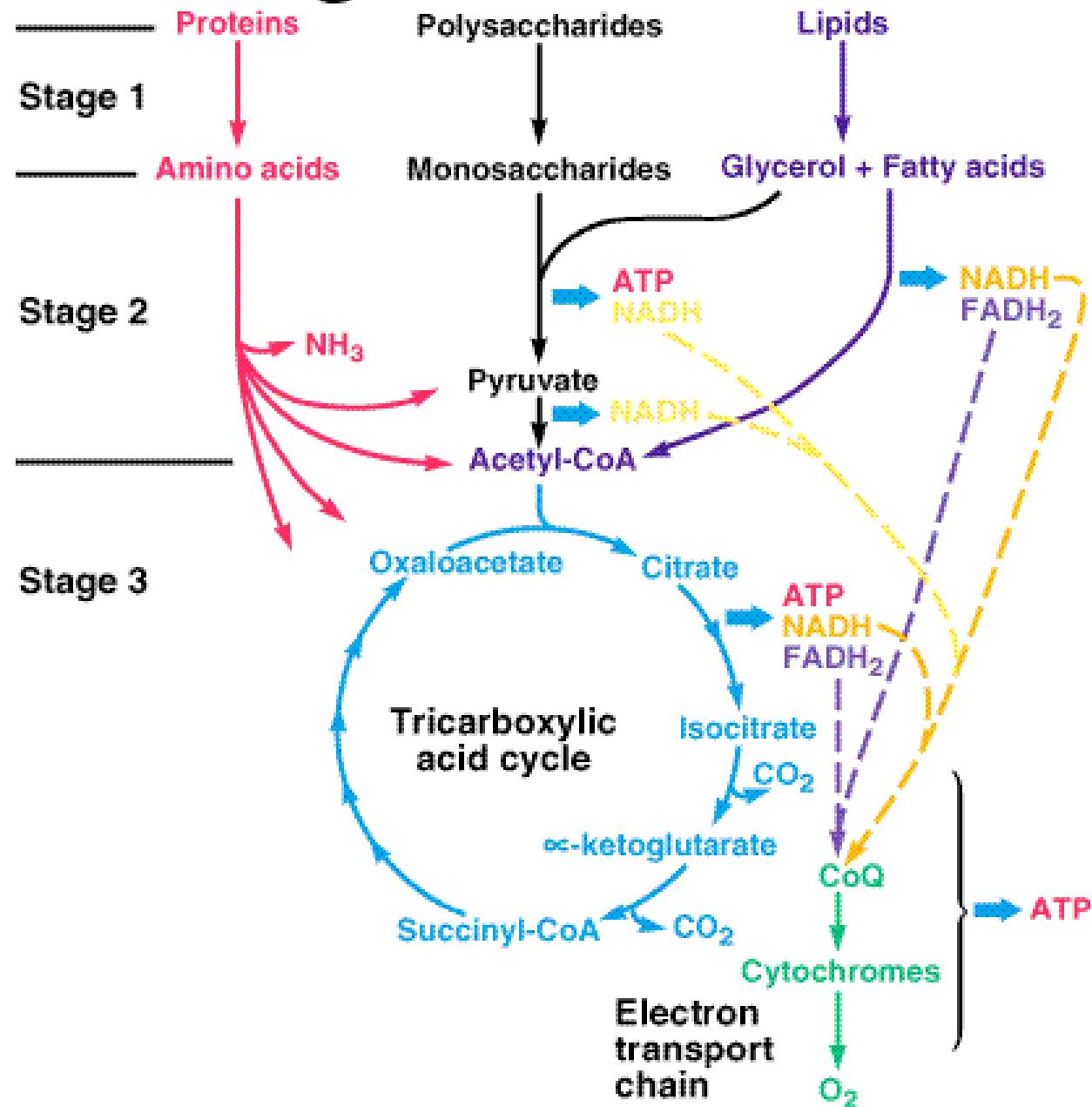


Fig. 9.3

# Entner-Doudoroff Pathway

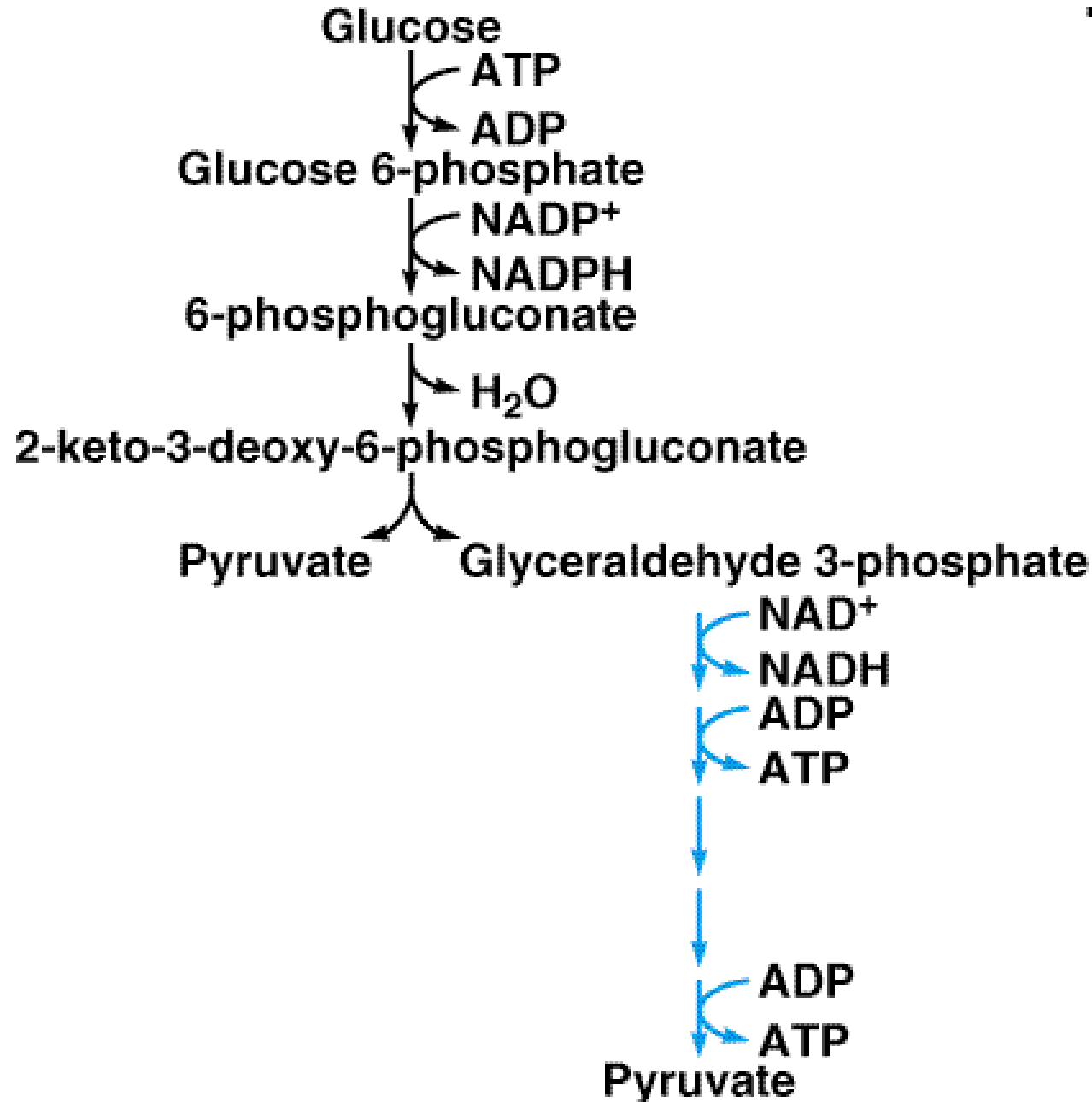


Fig 9.8



# Pentose Phosphate Pathway

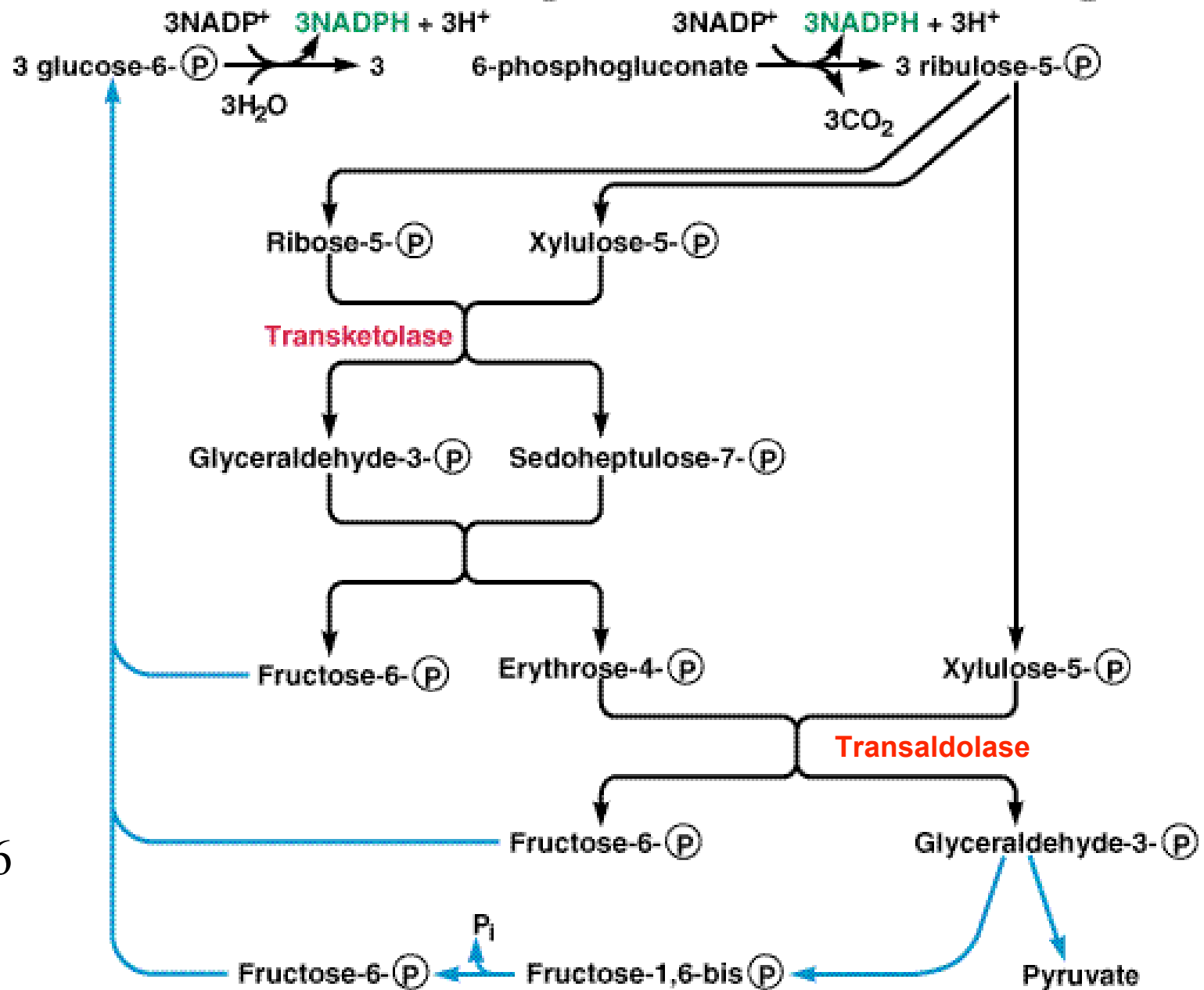
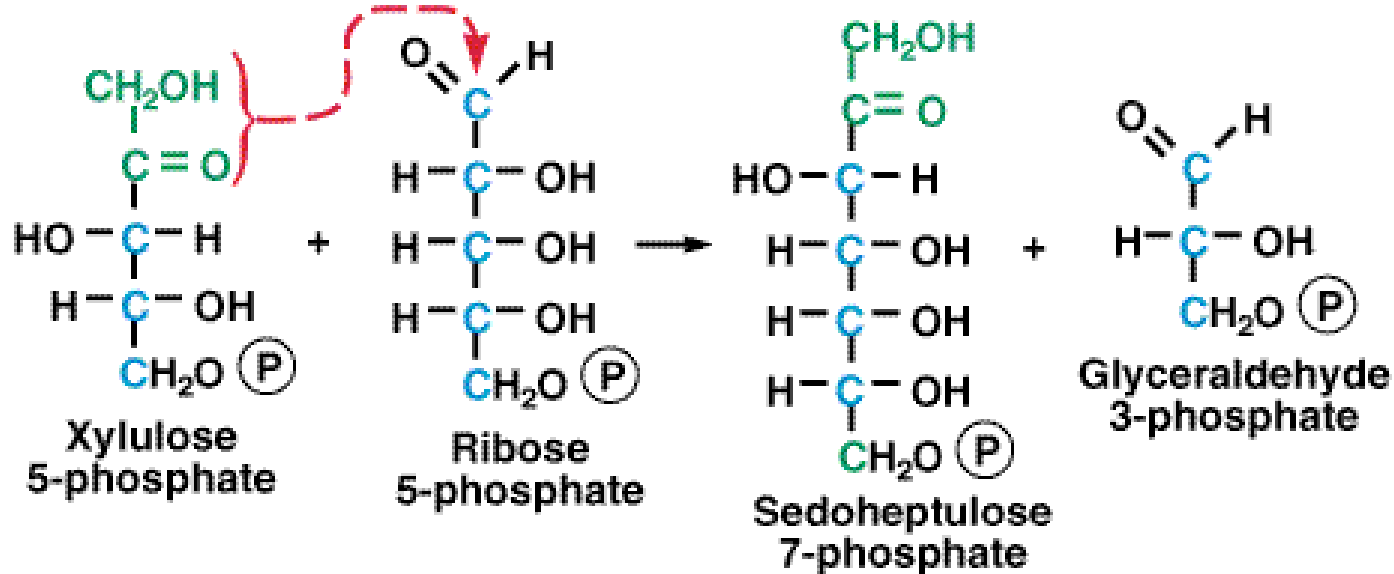


Fig 9.6

# Transketolase & Transaldolase

## The transketolase reaction



## The transaldolase reaction

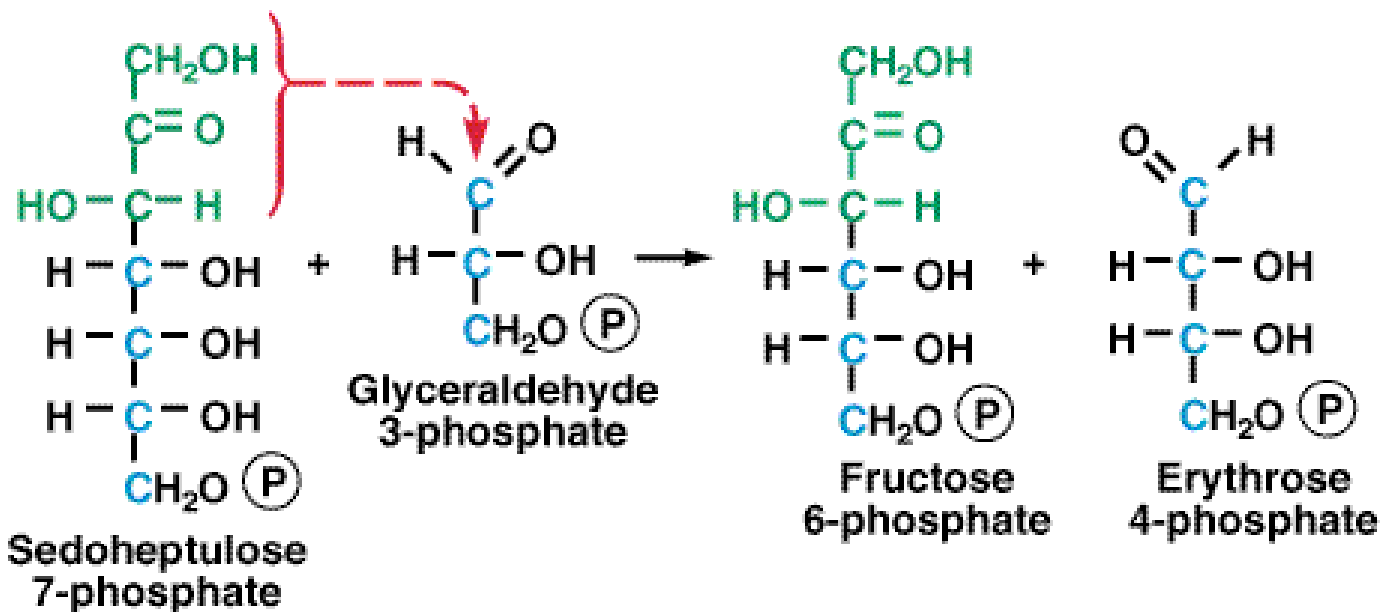


Fig. 9.7

# Three Stages of Catabolism

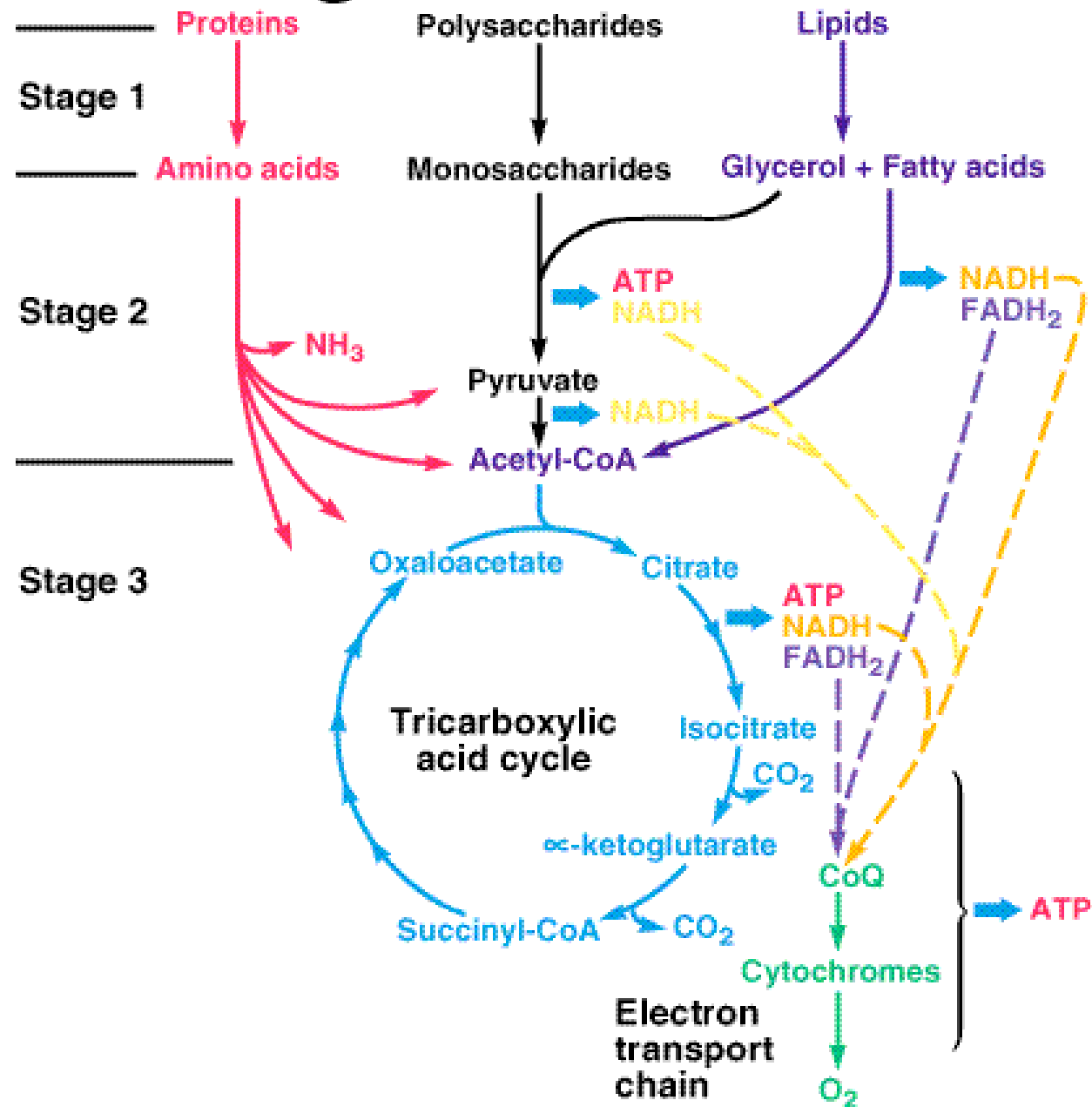


Fig 9.3

# Fatty Acid $\beta$ -oxidation

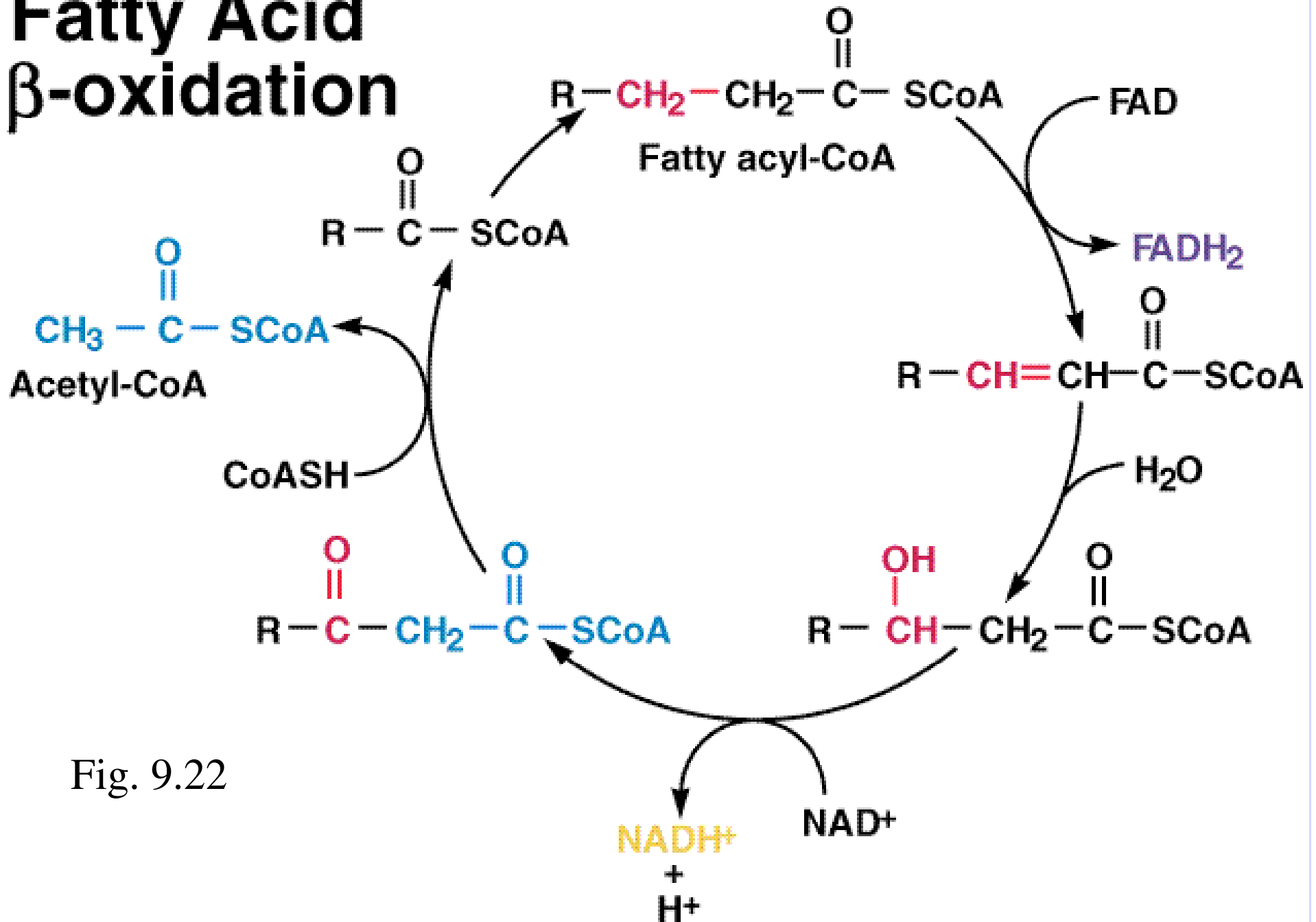


Fig. 9.22

# Tricarboxylic Acid Cycle

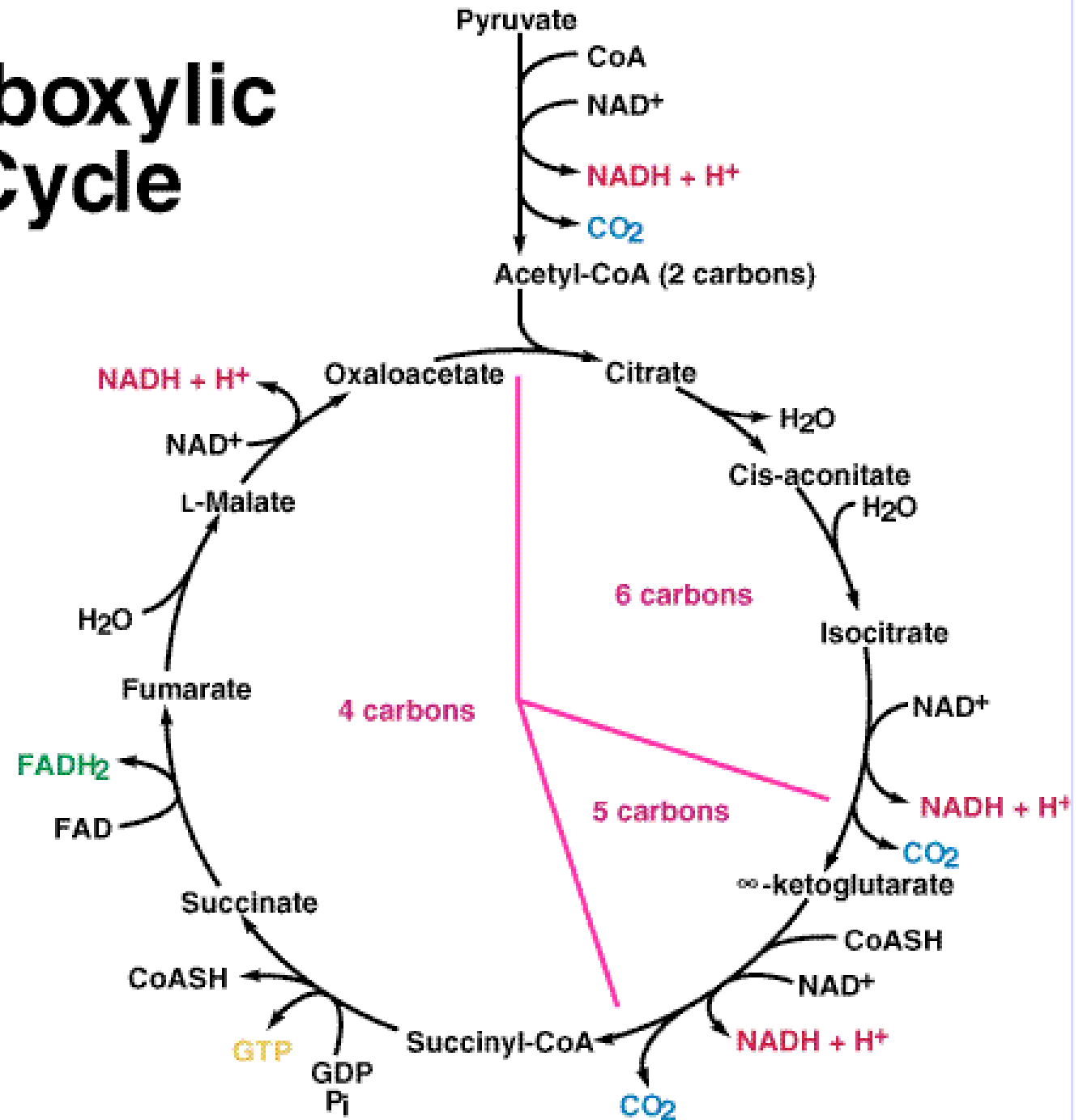
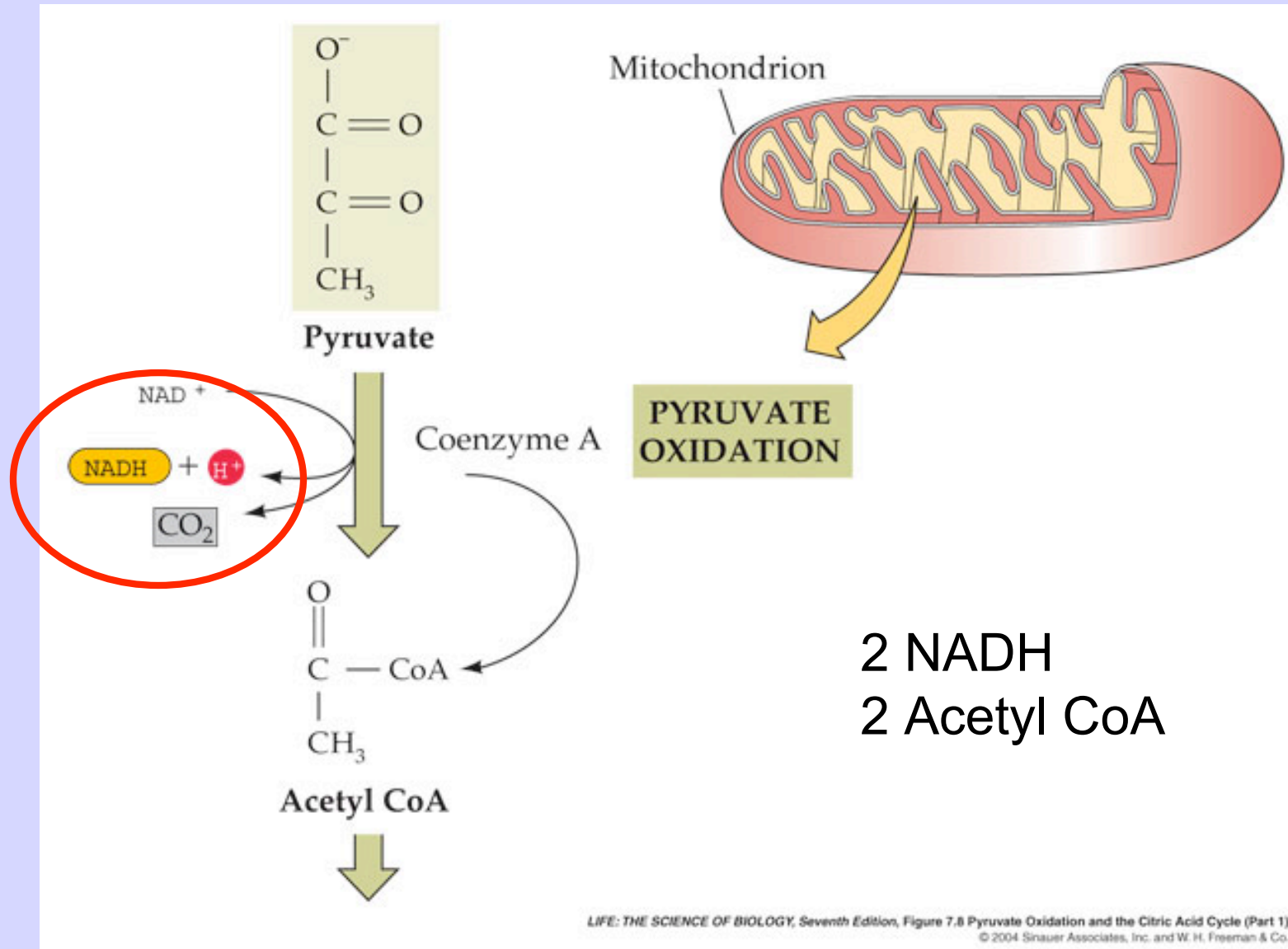


Fig. 9.12



# Tricarboxylic Acid Cycle

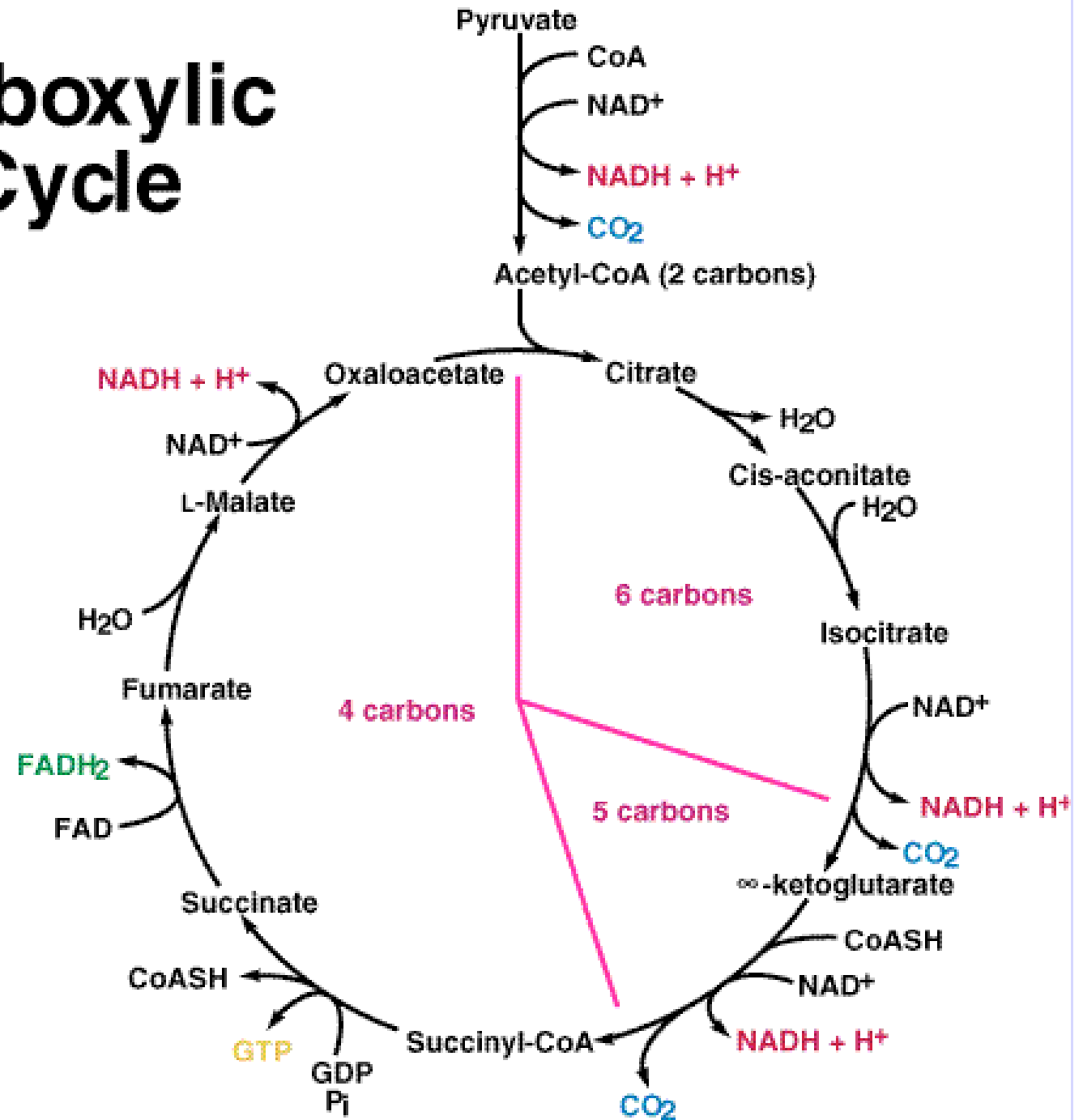
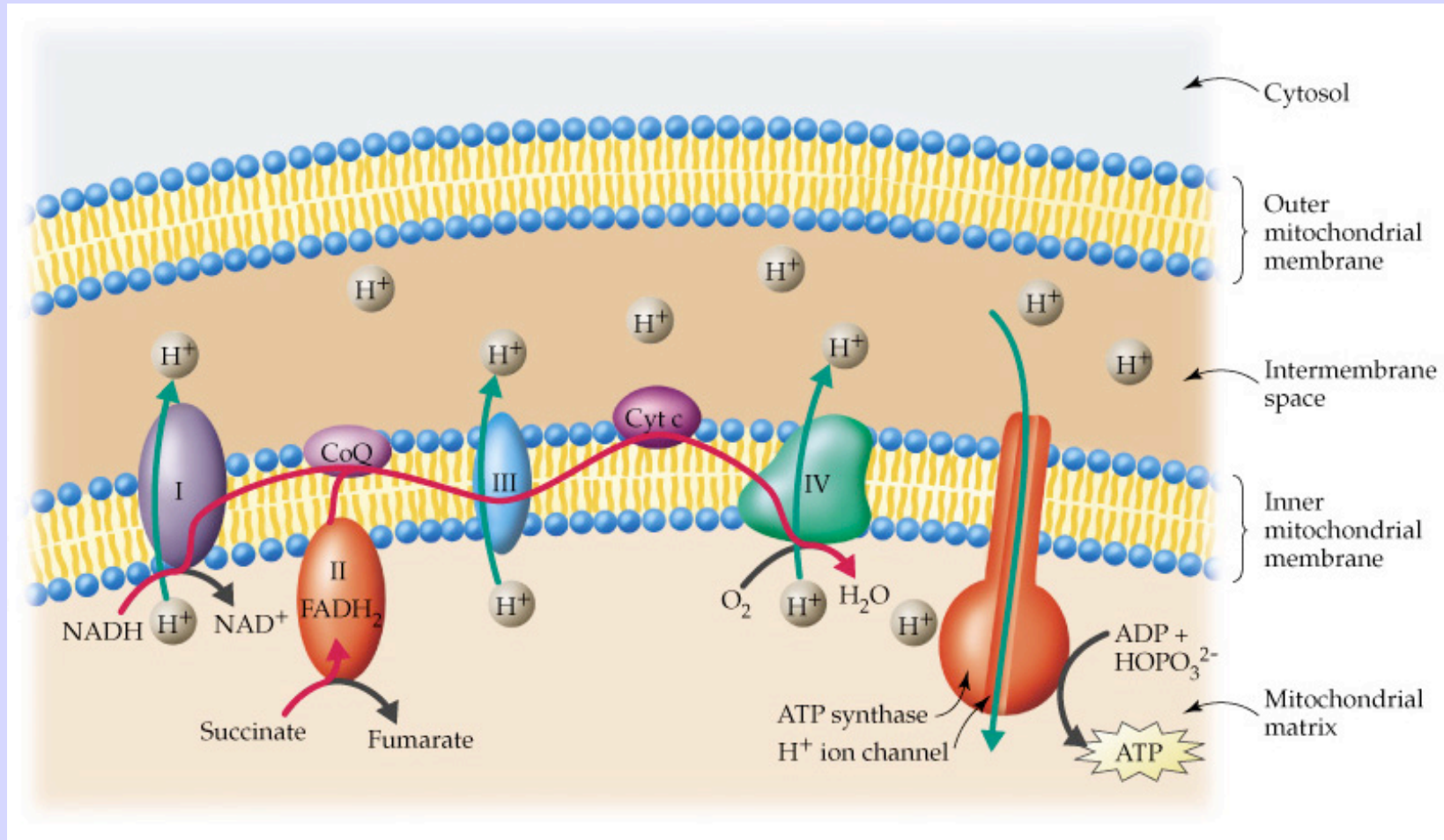
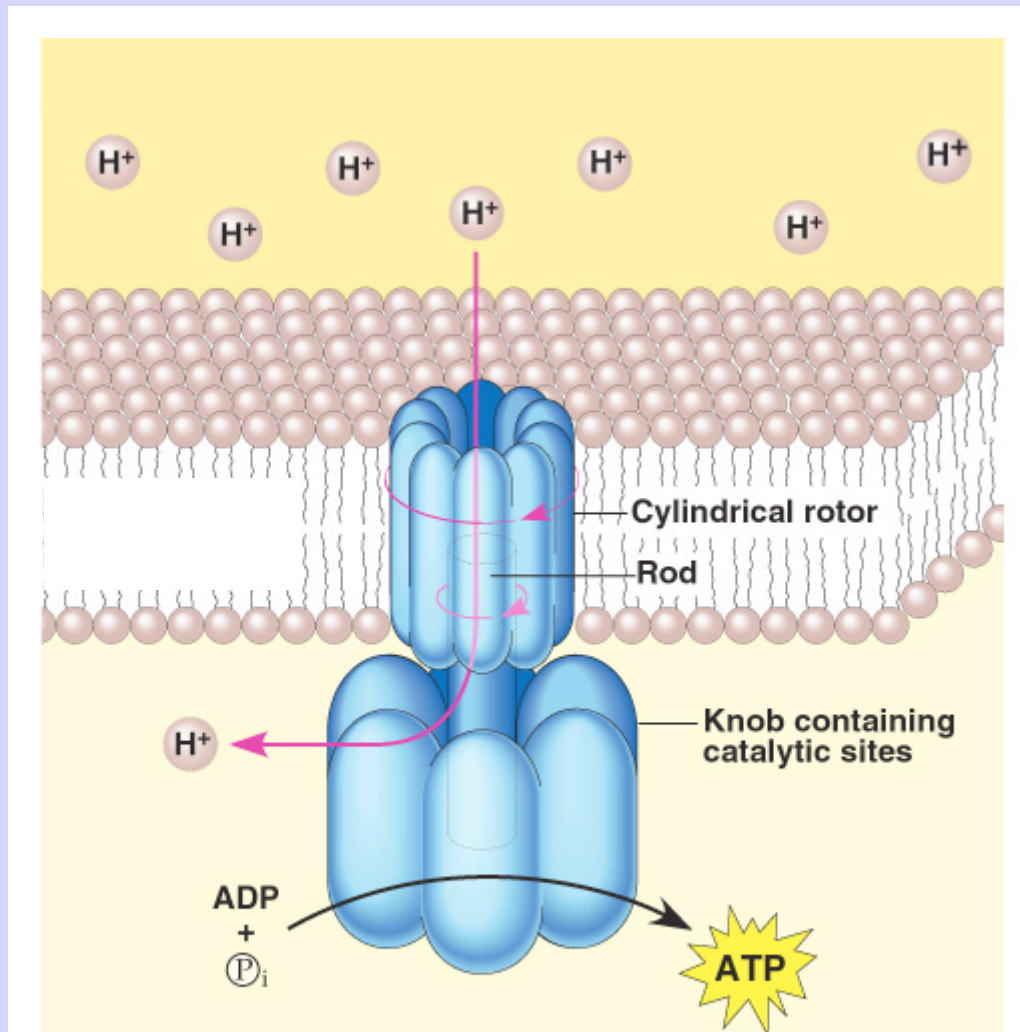


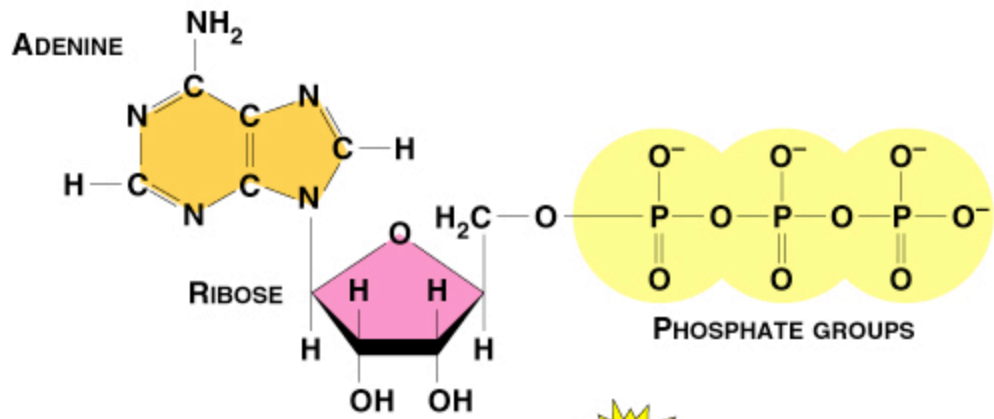
Fig. 9.12

# Electron Transport Chain









(a) Adenosine triphosphate

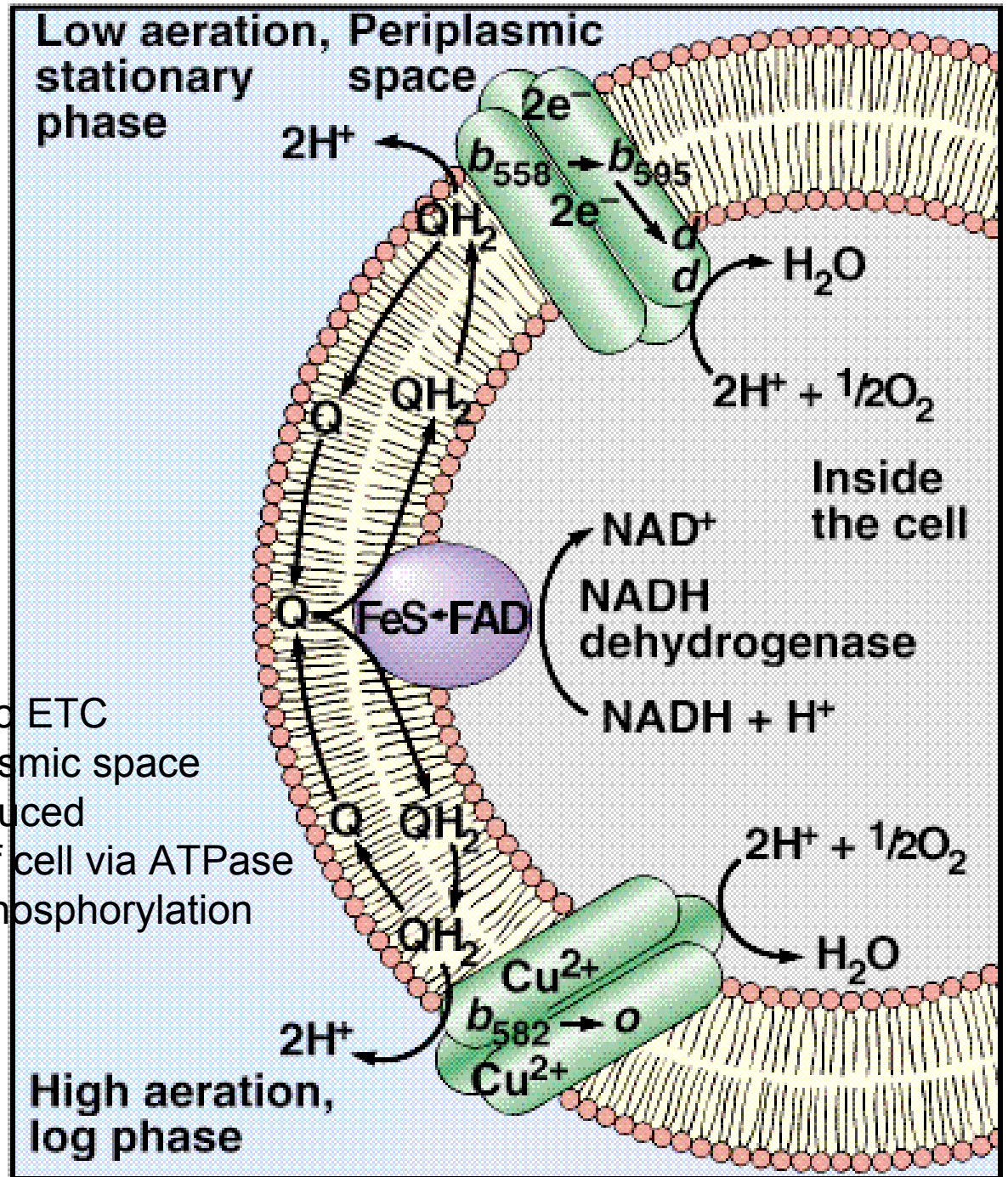


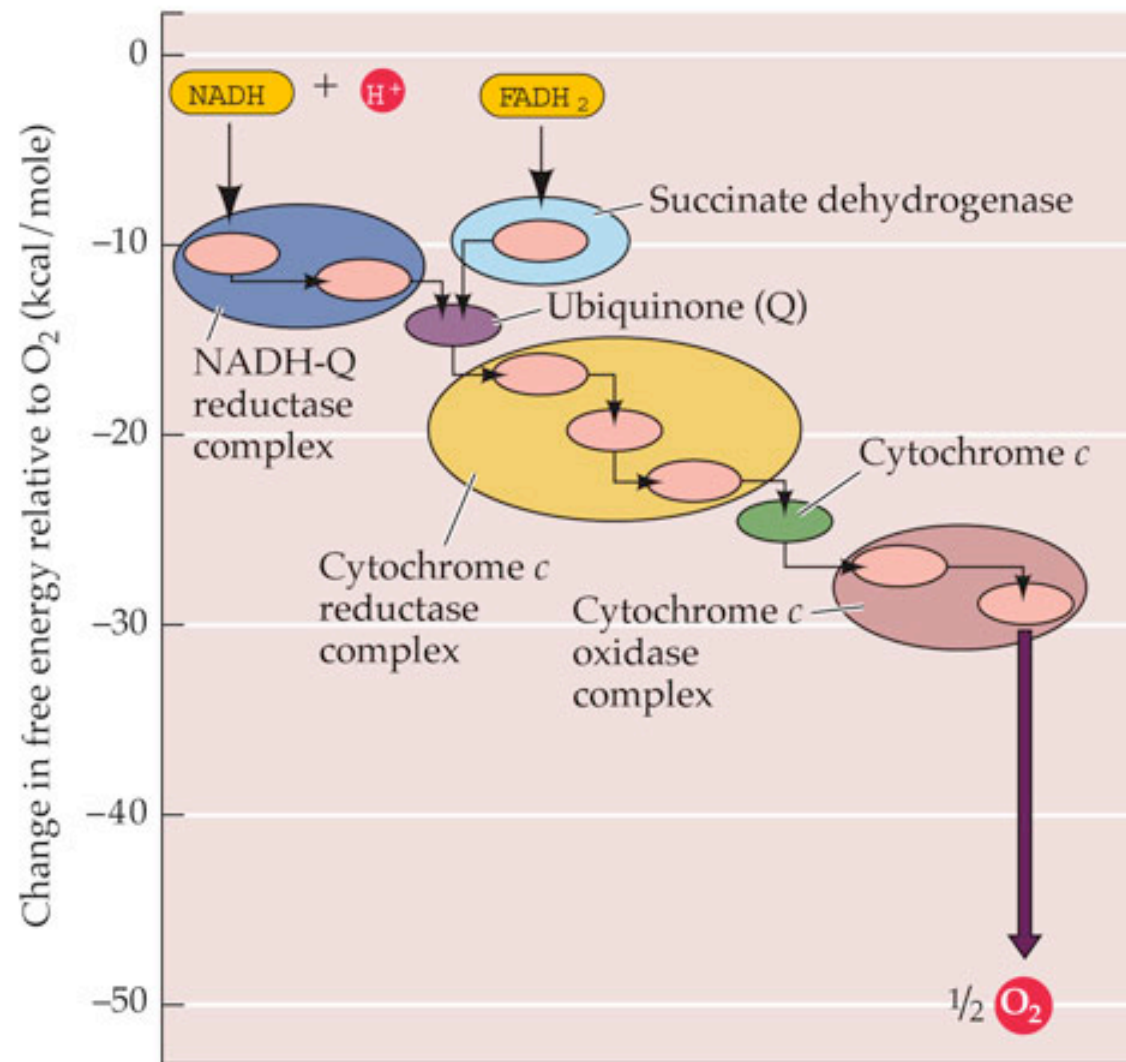
(b) Hydrolysis of ATP

# Respiratory Chain of *E. coli*

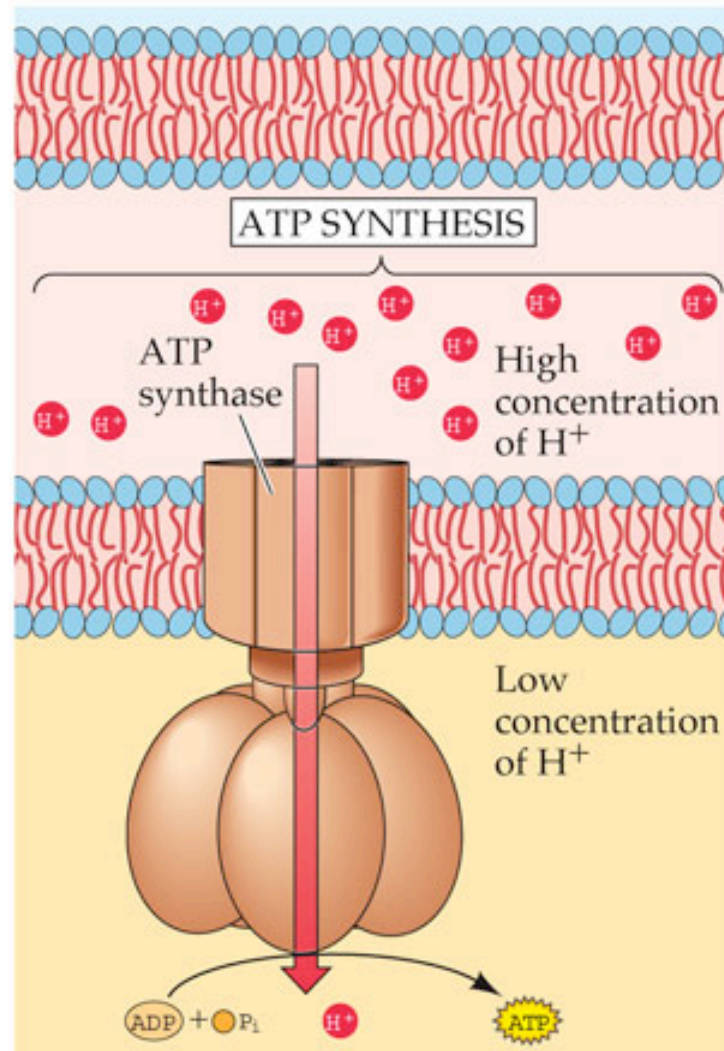
- Electron carrier transfer  $e^-$  ( $H^+$ ) to ETC
- Protons are pumped into periplasmic space
- Electrochemical gradient is produced
- Protons “travel back” to inside of cell via ATPase
- ATP is produced via oxidative phosphorylation
- $O_2$  is reduced to  $H_2O$

Fig. 9.15





LIFE: THE SCIENCE OF BIOLOGY, Seventh Edition, Figure 7.11 The Complete Respiratory Chain  
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## Animations

**TABLE 9.3** Some Electron Acceptors  
Used in Respiration

	Electron Acceptor	Reduced Products	Examples of Microorganisms
Aerobic	$O_2$	$H_2O$	All aerobic bacteria, fungi, protozoa, and algae
Anaerobic	$NO_3^-$	$NO_2^-$	Enteric bacteria
	$NO_3^-$	$NO_2^-$ , $N_2O$ , $N_2$	<i>Pseudomonas</i> and <i>Bacillus</i>
	$SO_4^{2-}$	$H_2S$	<i>Desulfovibrio</i> and <i>Desulfotomaculum</i>
	$CO_2$	$CH_4$	All methanogens
	$S^0$	$H_2S$	<i>Desulfuromonas</i> and <i>Thermoproteus</i>
	$Fe^{3+}$	$Fe^{2+}$	<i>Pseudomonas</i> and <i>Bacillus</i>

# Three Stages of Catabolism

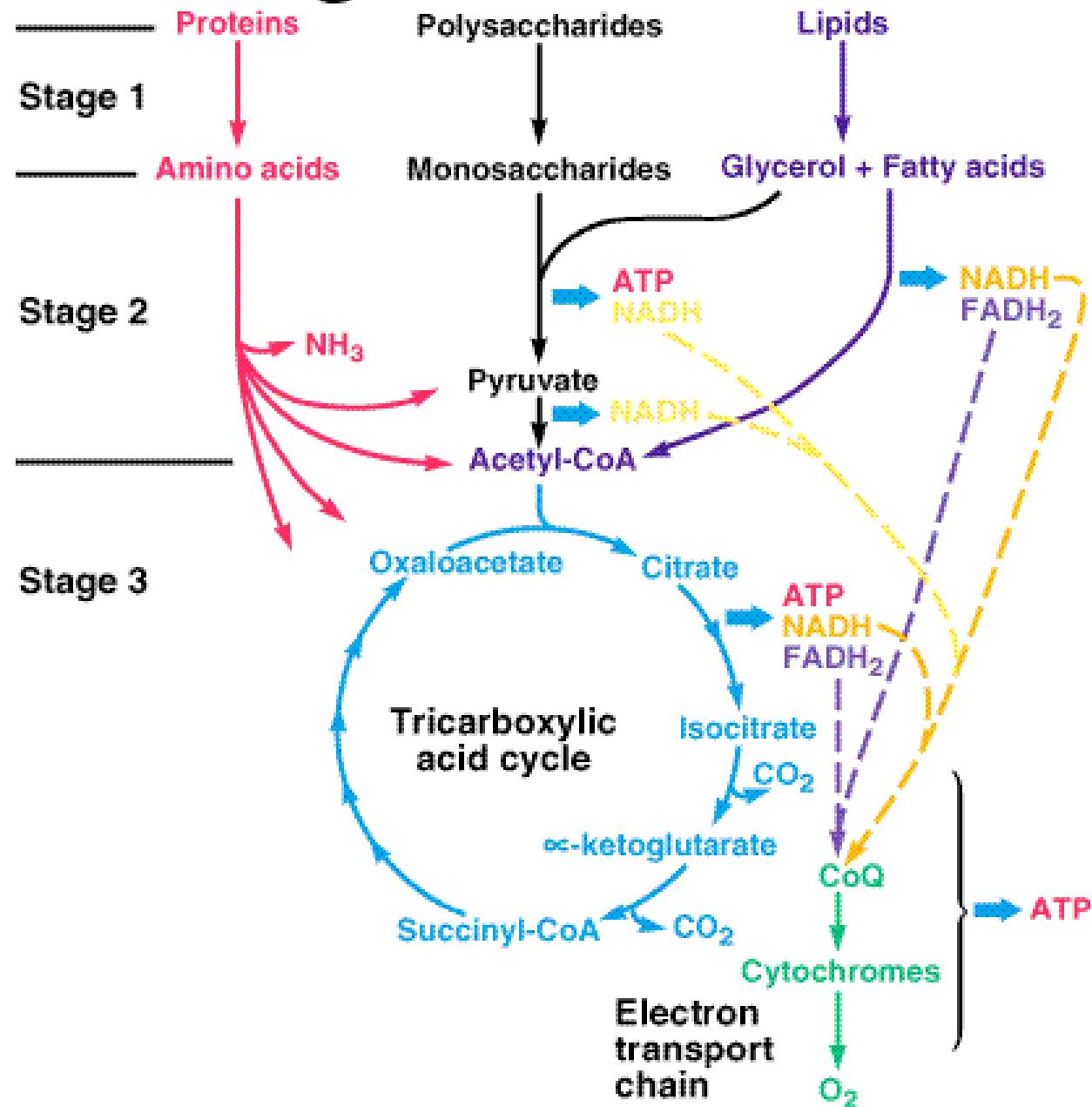
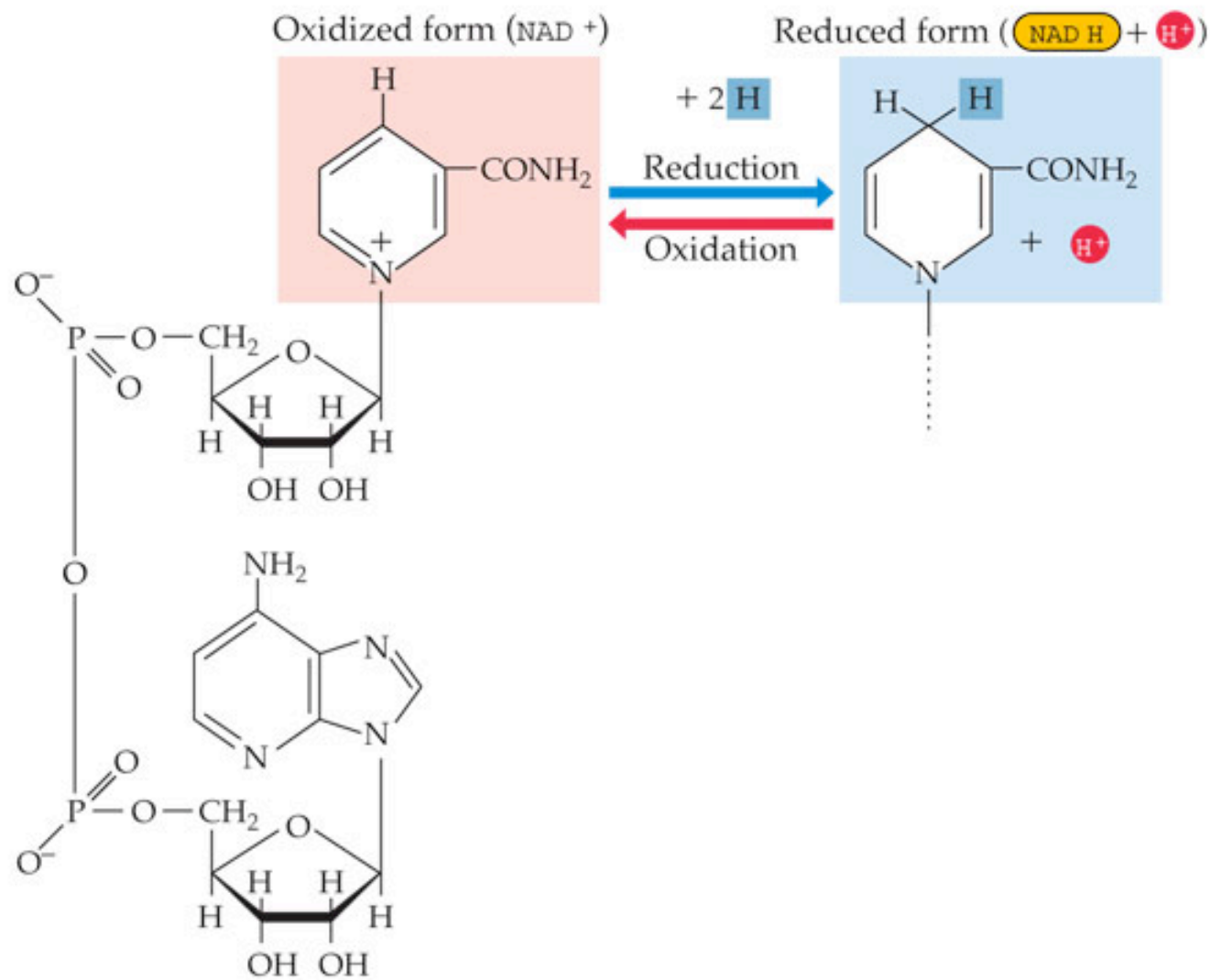


Fig. 9.3





# NADH Oxidation in Fermentation

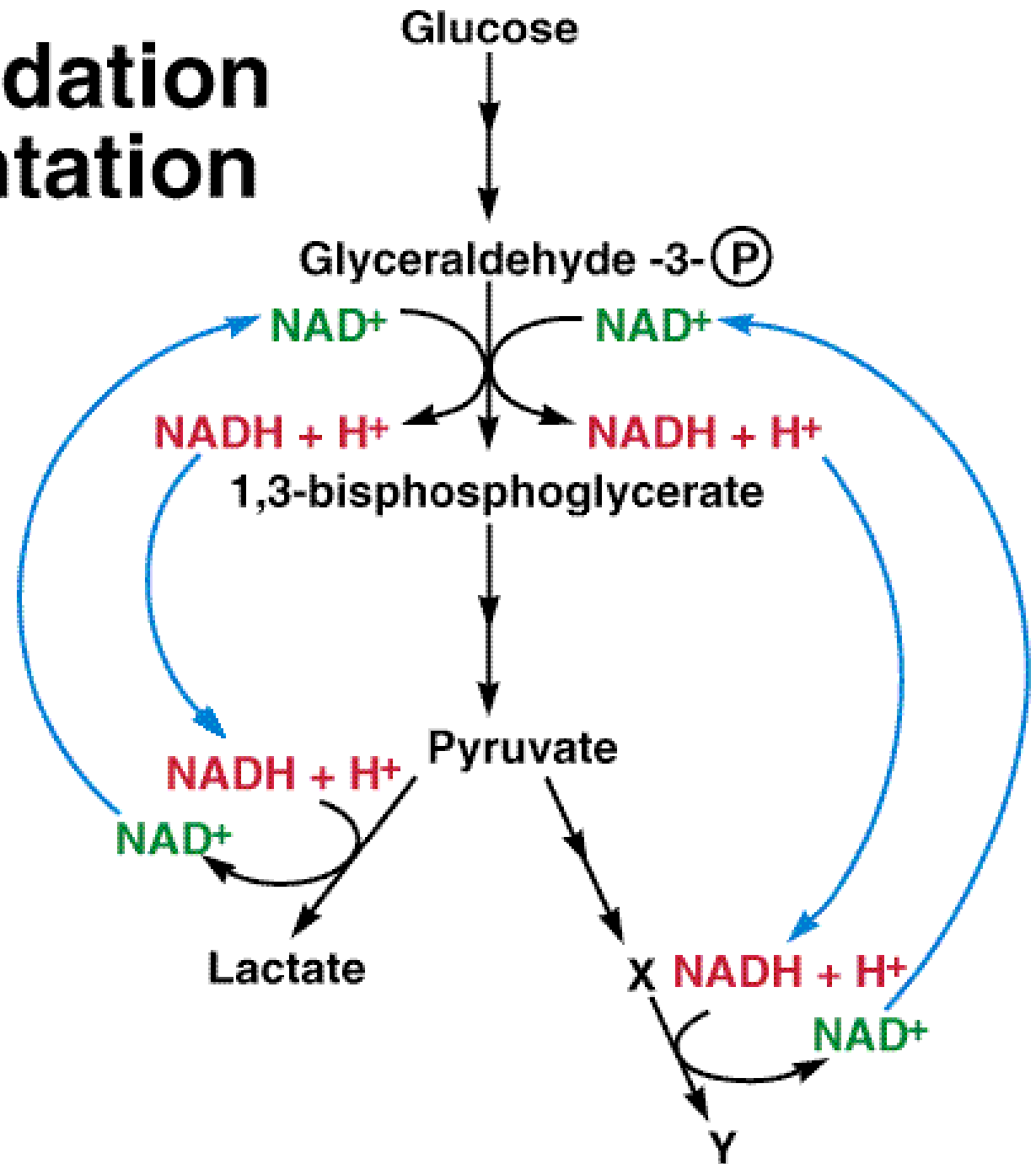


Fig. 9.9

# Microbial Fermentations

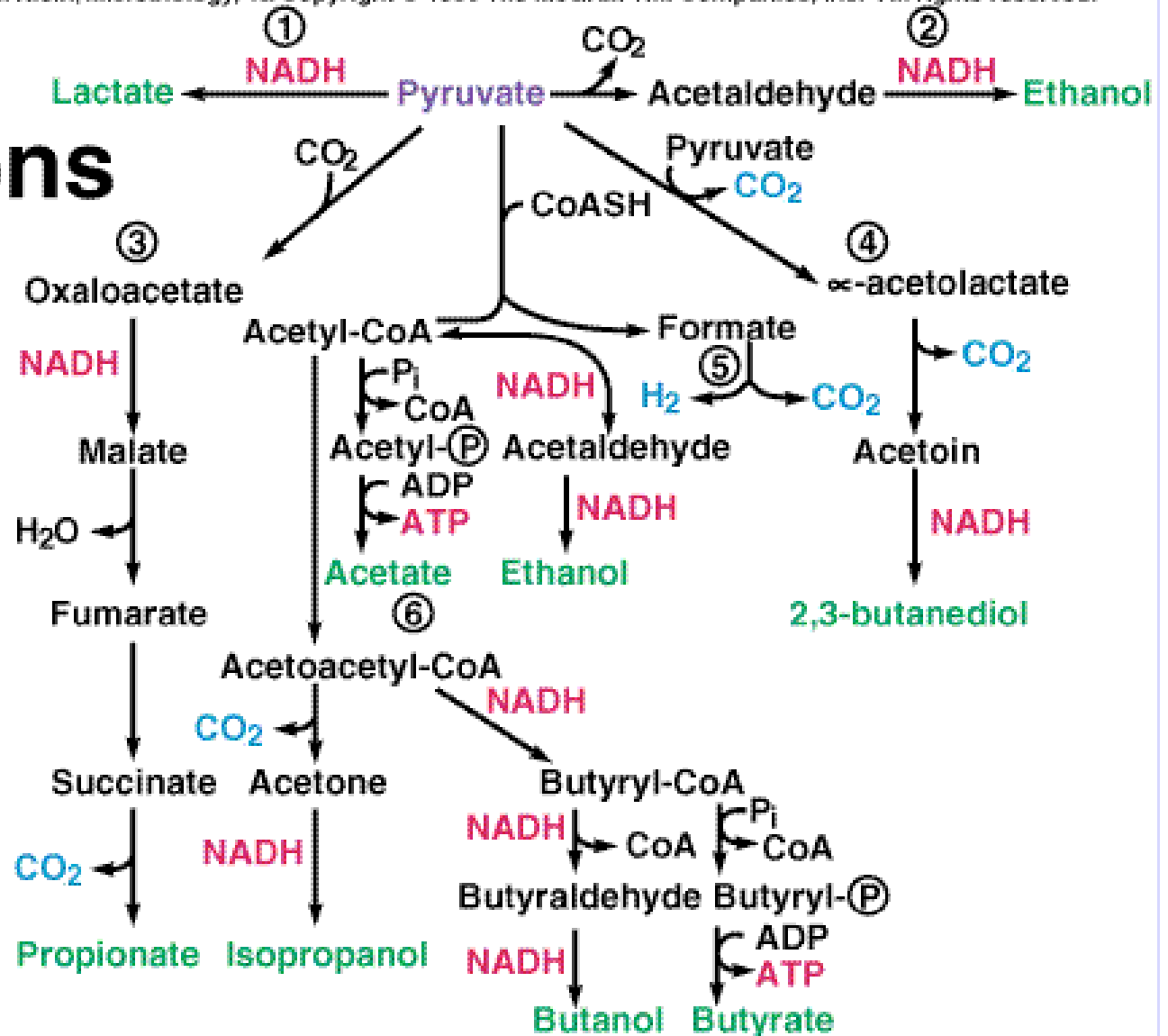
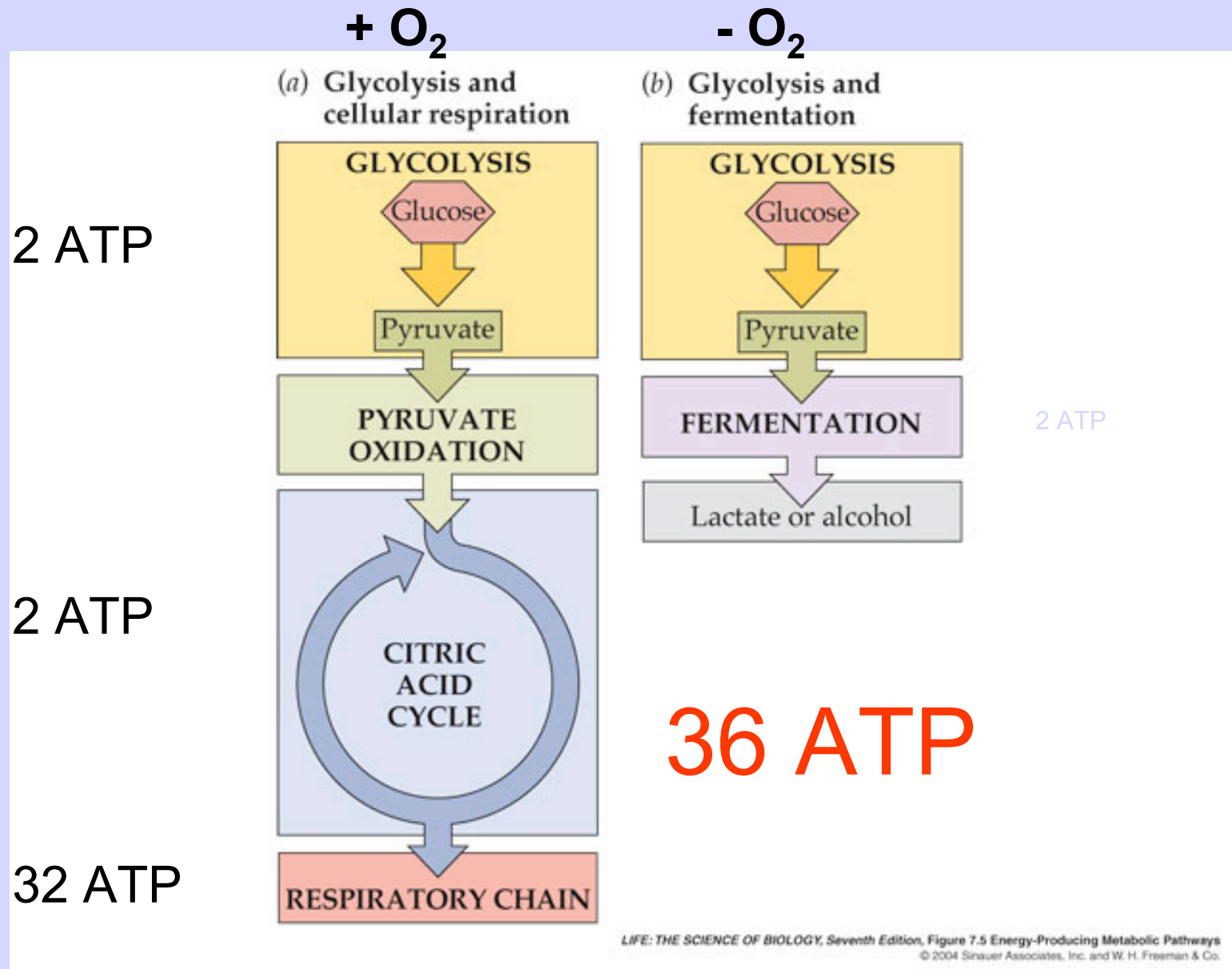

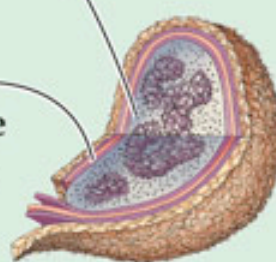


Fig. 9.10

1. Lactic acid bacteria (*Streptococcus*, *Lactobacillus*), *Bacillus*
2. Yeast, *Zymomonas*
3. Propionic acid bacteria (*Propionibacterium*)
4. *Enterobacter*, *Serratia*, *Bacillus*
5. Enteric bacteria (*Escherichia*, *Enterobacter*, *Salmonella*, *Proteus*)
6. *Clostridium*



## 7.1 Cellular Locations for Energy Pathways in Eukaryotes and Prokaryotes

EUKARYOTES	PROKARYOTES
 <p><b>External to mitochondrion</b> Glycolysis Fermentation</p> <p><b>Inside mitochondrion</b> Inner membrane Pyruvate oxidation Respiratory chain Matrix Citric acid cycle</p>	 <p><b>In cytoplasm</b> Glycolysis Fermentation Citric acid cycle</p> <p><b>On inner face of plasma membrane</b> Pyruvate oxidation Respiratory chain</p>

# Patterns of Energy Production

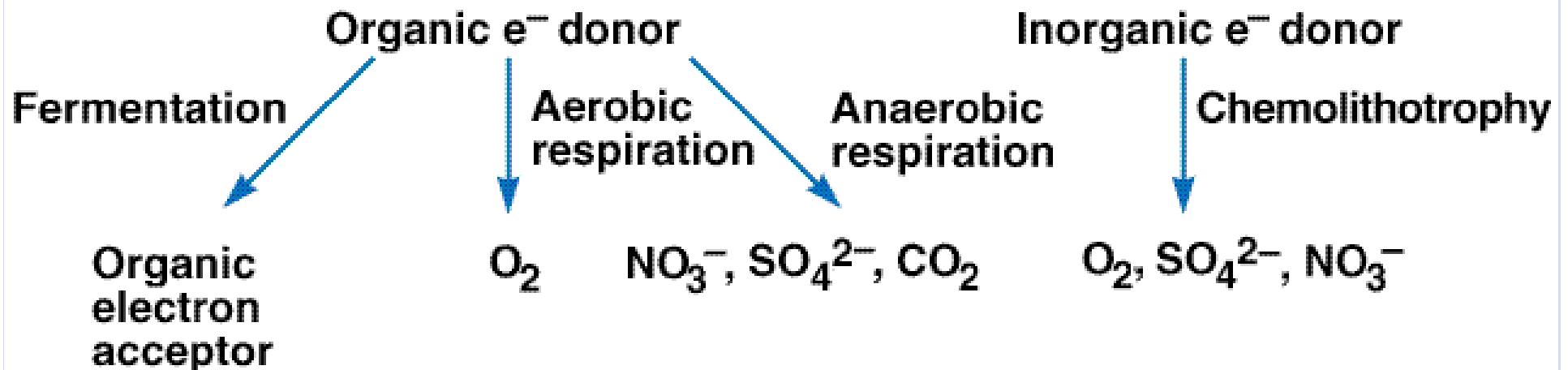


Fig. 9.2