

Microbes as Energy Transducers

- The Metabolic Menu
- Metabolic Strategies
- Respiration & Fermentation
- Chemolithotrophy
- Photoautotrophy
- Biogeochemical Cycles
- Metabolism in Primitive Organisms

All major types of nutrition and metabolism evolved among prokaryotes: they are the ultimate biochemists

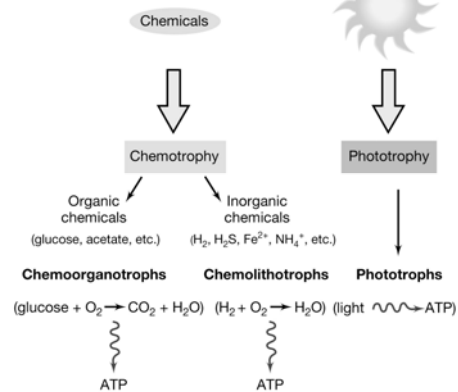
The prokaryotes exhibit some unique modes of nutrition as well as every type of nutrition found in eukaryotes.

Major Modes of Nutrition:

Prokaryotes exhibit a great diversity in how they obtain the necessary resources (**energy and carbon**) to synthesize organic compounds.

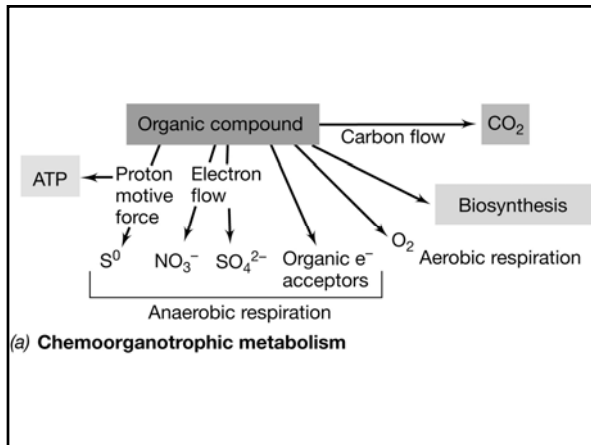
- Some obtain energy from light (**phototrophs**), while others use chemicals taken from the environment (**chemotrophs**).
- Many can utilize CO₂ as a carbon source (**autotrophs**) and others require at least one organic nutrient as a carbon source (**heterotrophs**).

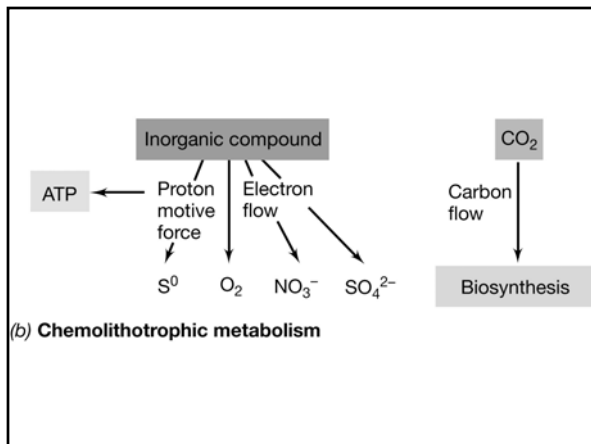
Energy Sources

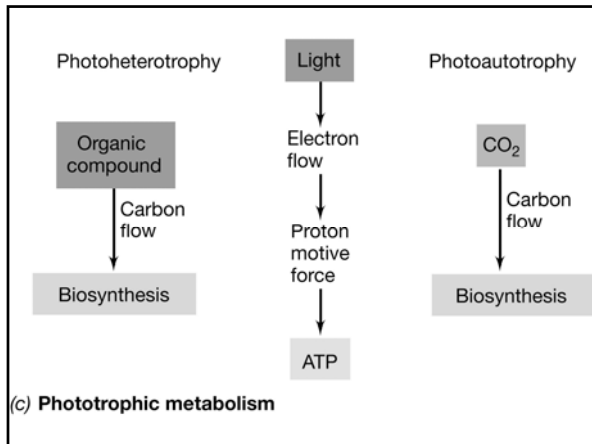


Depending upon the energy source **AND** the carbon source, prokaryotes have **four** possible nutritional modes:

1. **Photoautotrophs:** Use light energy to synthesize organic compounds from CO_2 – Includes the cyanobacteria. (Actually all photosynthetic eukaryotes fit in this category.)
2. **Chemoautotrophs:** Require only CO_2 as a carbon source and obtain energy by oxidizing inorganic compounds. This mode of nutrition is unique only to certain prokaryotes.
3. **Photoheterotrophs:** Use light to generate ATP from an organic carbon source. This mode of nutrition is unique only to certain prokaryotes.
4. **Chemoheterotrophs:** Must obtain organic molecules for energy and as a source of carbon. Found in many bacteria as well as most eukaryotes.

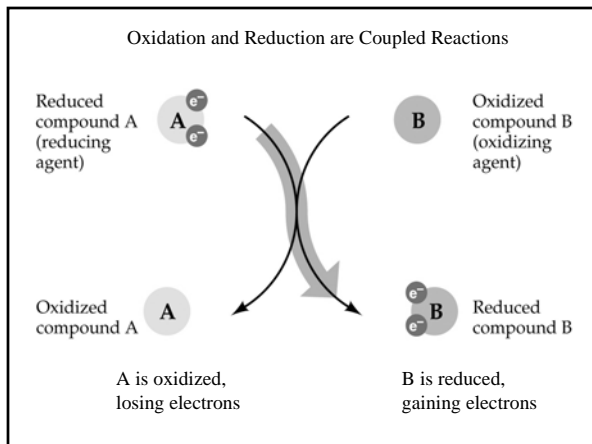


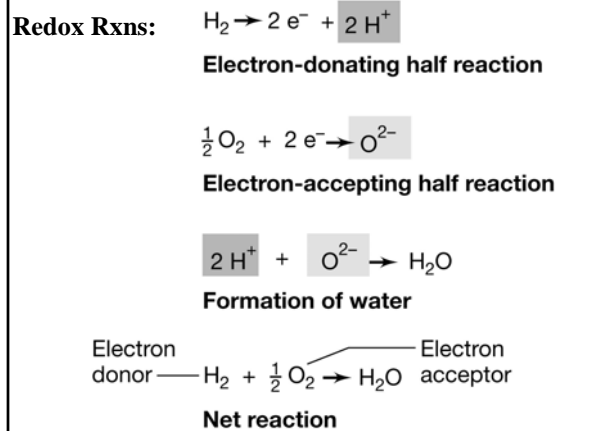


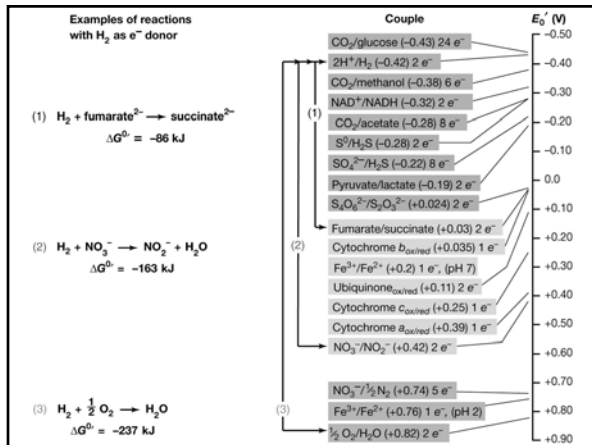


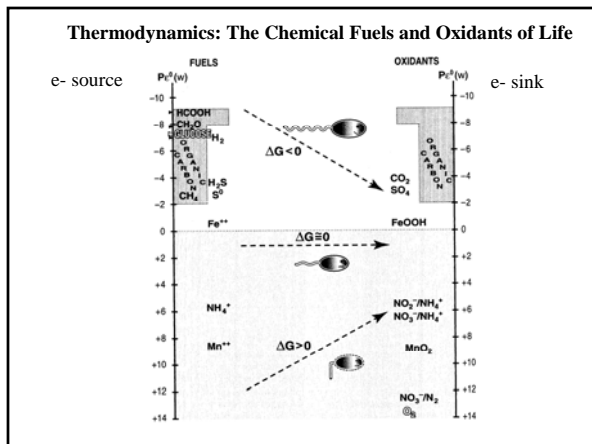
Metabolic Menu For Chemotrophs

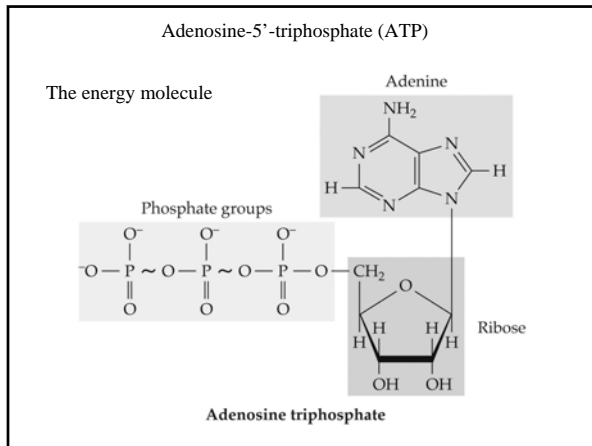
Potential Microbial Metabolic Processes			
e ⁻ donor	e ⁻ acceptor	C source	Organisms
Autolithotrophy			
H ₂	O ₂	CO ₂	Hydrogen oxidizers
HS ⁻ , S ₂ O ₃ ²⁻	O ₂	CO ₂	Sulfur oxidizers
Fe ²⁺	O ₂	CO ₂	Iron oxidizers
Mn ²⁺	O ₂	CO ₂	Manganese oxidizers
NH ₄ ⁺ , NO ₂ ⁻	O ₂	CO ₂	Nitrifiers
HS ⁻ , S ₂ O ₃ ²⁻	NO ₃ ⁻	CO ₂	Denitrifying S-oxidizers
H ₂	NO ₃ ⁻	CO ₂	Hydrogen oxidizers
H ₂	S ⁰ , SO ₄ ²⁻	CO ₂	Sulfate Reducers (SRBs)
H ₂	CO ₂	CO ₂	Methanogens & Acetogens
Heteroorganotrophy			
Org. C	O ₂	Org. C	Aerobic Heterotrophy
Org. C	NO ₃ ⁻	Org. C	Denitrifiers
Org. C	S ⁰ , SO ₄ ²⁻	Org. C	Sulfate Reducers (SRBs)
Org. C	Org. C	Org. C	Fermenters
Methylophony			
CH ₄ (C-1)	O ₂ , SO ₄ ²⁻	CH ₄ , CO ₂ , CO	Methane (C-1) oxidizers

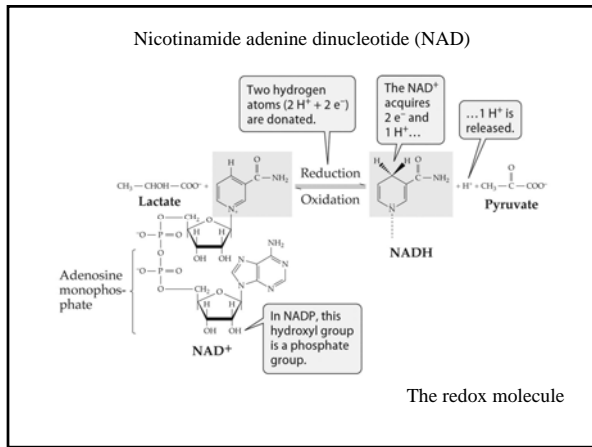


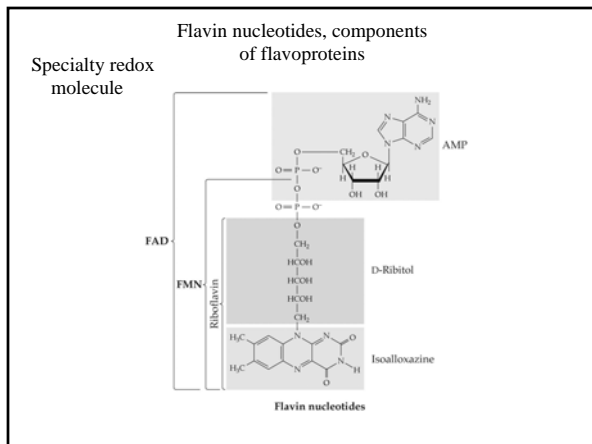


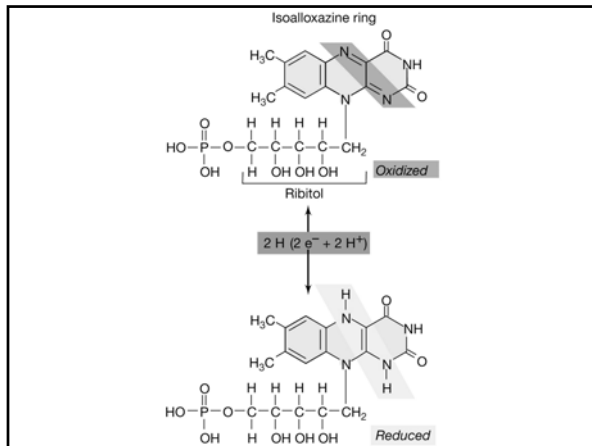


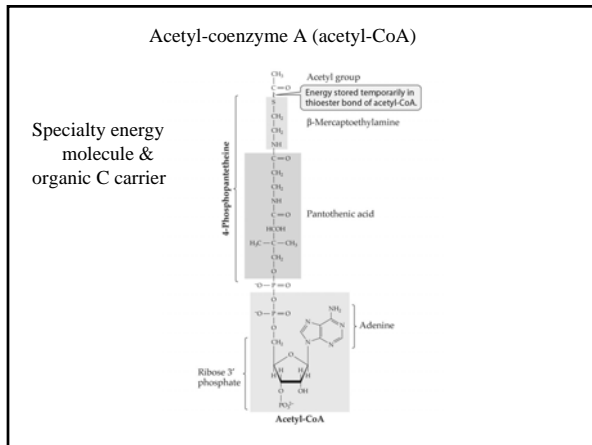


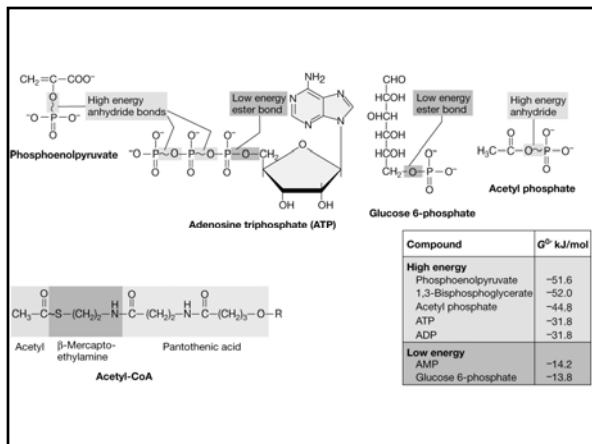


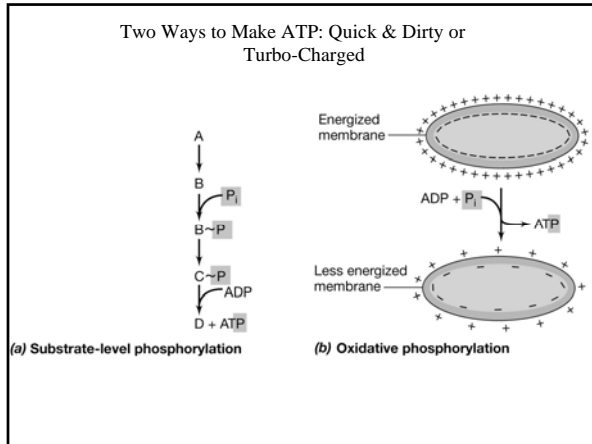






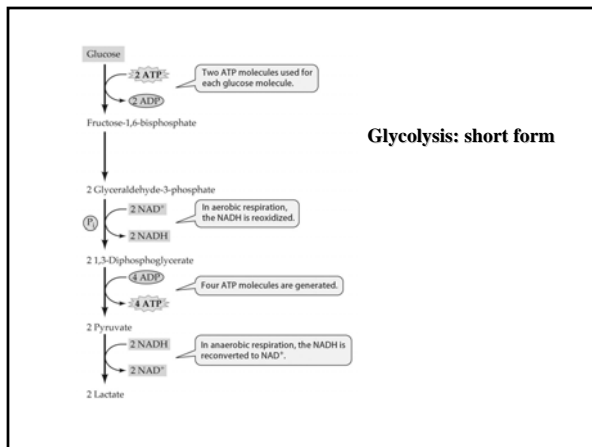


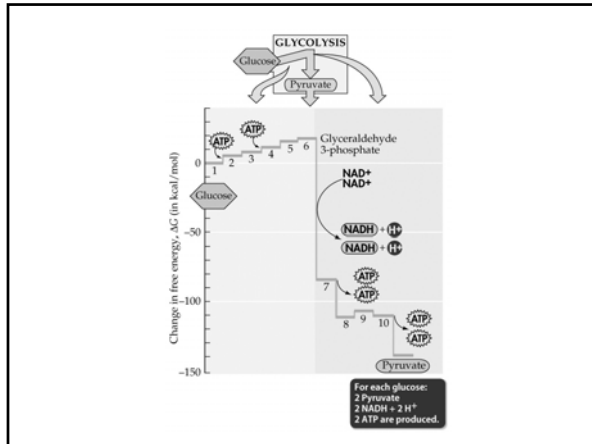


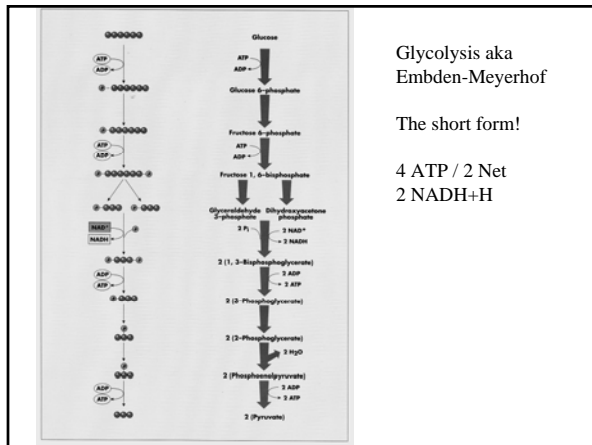


7.1 Cellular Locations for Energy Pathways in Eukaryotes and Prokaryotes

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



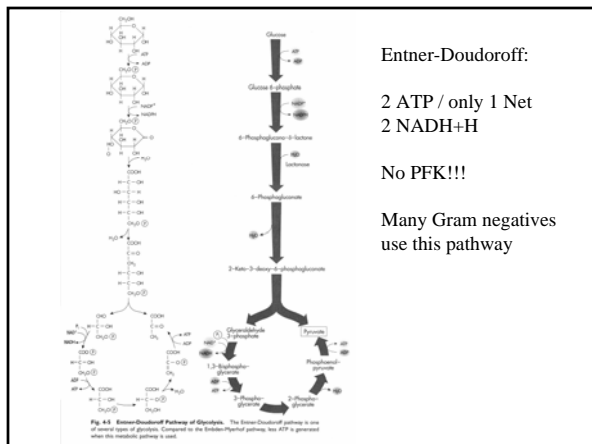




Glycolysis aka
Embden-Meyerhof

The short form!

4 ATP / 2 Net
2 NADH+H

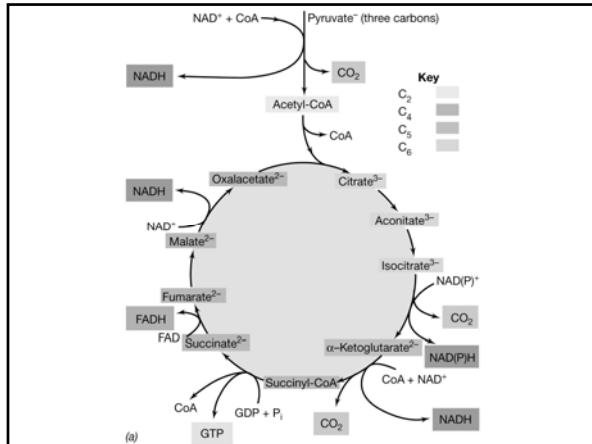


Entner-Doudoroff:

2 ATP / only 1 Net
2 NADH+H

No PFK!!!

Many Gram negatives
use this pathway



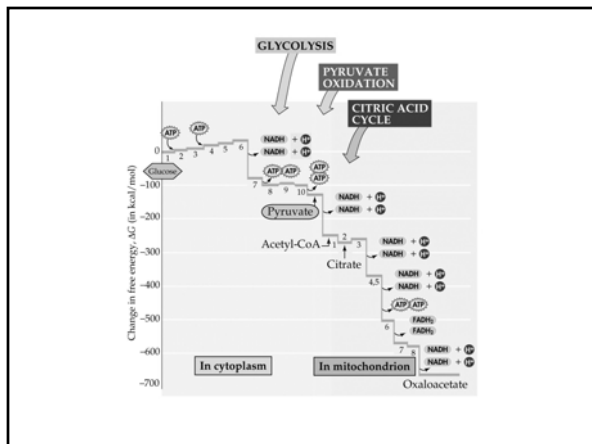
Overall reaction: $\text{Pyruvate}^- + 4 \text{NAD}^+ + \text{FAD} \rightarrow 3 \text{CO}_2 + 4 \text{NADH} + \text{FADH}$

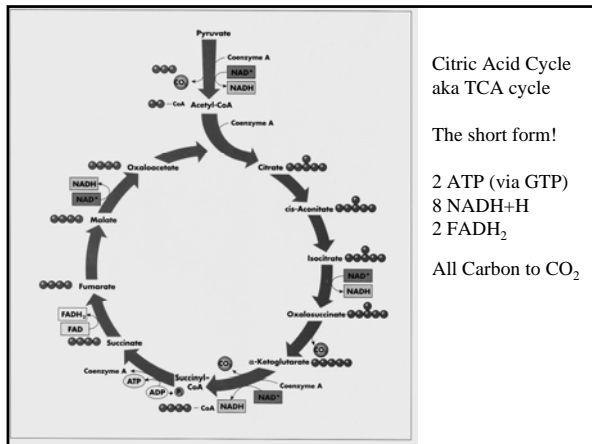
(1) Substrate-level phosphorylation $\text{GDP} + \text{P}_i \rightarrow \text{GTP}$
 $\text{GTP} + \text{ADP} \rightarrow \text{GDP} + \text{ATP}$

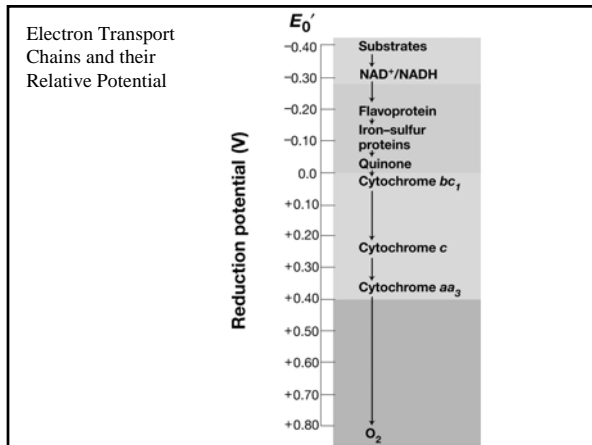
(2) Electron transport phosphorylation $4 \text{NADH} \equiv 12 \text{ATP}$
 $\text{FADH} \equiv 2 \text{ATP}$

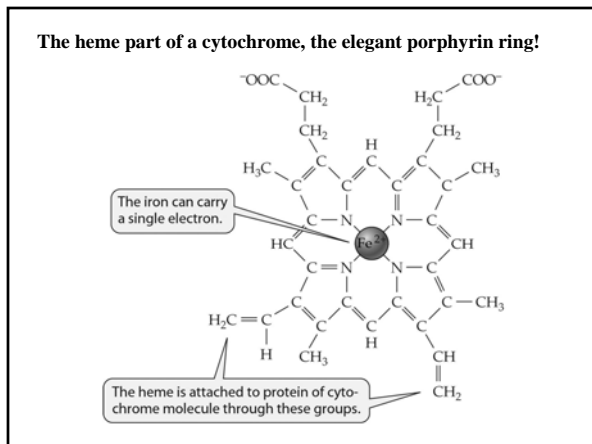
(3) Sum: CAC plus glycolysis $\rightarrow 38 \text{ATP per glucose}$

(b)

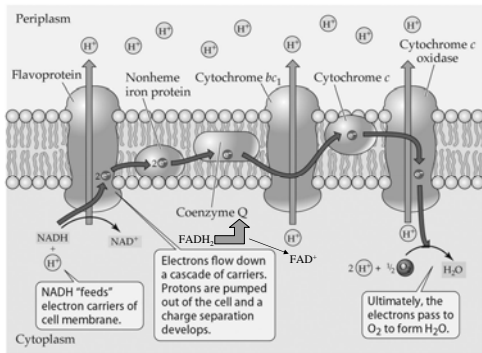




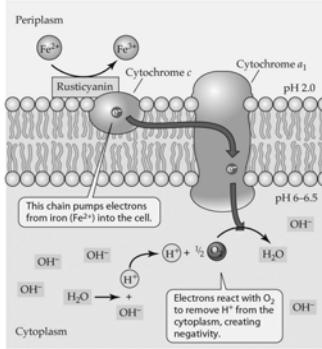




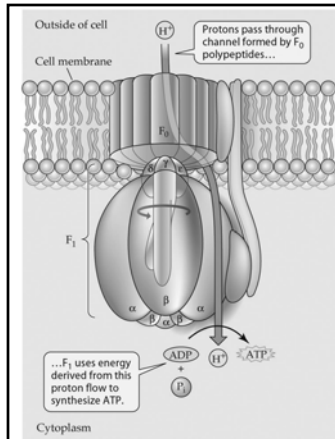
Electron transport chain in aerobic bacterium

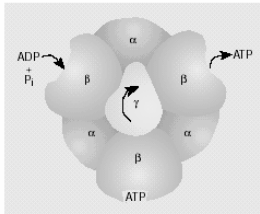


Abbreviated electron transport chain of an "iron-oxidizing" bacterium



ATP Synthase Structure & Function





F1 Subunit Topview

ATP Synthase acts as a rotary motor turning in 120 degree steps.

Figure 2 How rotation of the γ subunit drives catalysis. During ATP synthesis, rotation of the γ subunit causes sequential changes in the β subunits. A rotation of 120° changes the β subunit that binds ADP and P_i to a form with tightly bound ATP. The subunit with tightly bound ATP then changes to a form that releases ATP, and the third subunit prepares to bind another ADP and P_i .

Table 9.2 ATP Yield from the Aerobic Oxidation of Glucose by Eucaryotic Cells

Glycolytic Pathway	
Substrate-level phosphorylation (ATP)	2 ATP ^a
Oxidative phosphorylation with 2 NADH	6 ATP
2 Pyruvate to 2 Acetyl-CoA	
Oxidative phosphorylation with 2 NADH	6 ATP
Tricarboxylic Acid Cycle	
Substrate-level phosphorylation (GTP)	2 ATP
Oxidative phosphorylation with 6 NADH	18 ATP
Oxidative phosphorylation with 2 FADH ₂	4 ATP
Total Aerobic Yield	38 ATP

^aATP yields are calculated with an assumed P/O ratio of 3.0 for NADH and 2.0 for FADH₂.

Fermentation – Key Features

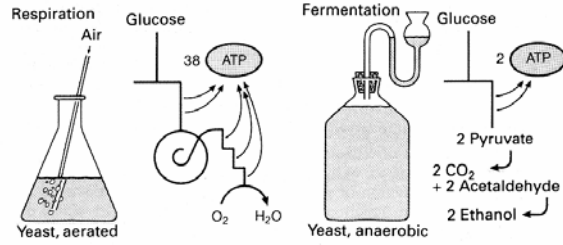
- (1) Substrate-level phosphorylation is the rule*.
- (2) Always anaerobic (even when some O₂ might be around).
- (3) No externally supplied terminal electron acceptor.

Many types.... 2 major themes

- (1) NADH+H⁺ gets oxidized to NAD⁺
- (2) Electron acceptor is usually **Pyruvate** or its derivative.

*Rules are always meant to be broken!

Pasteur Effect: ~20X more biomass when aerated



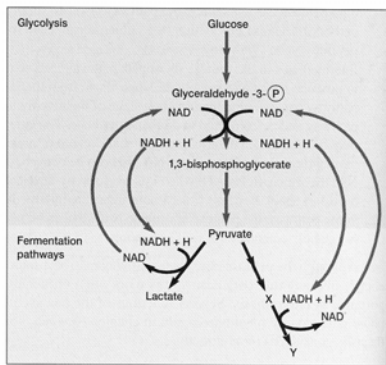
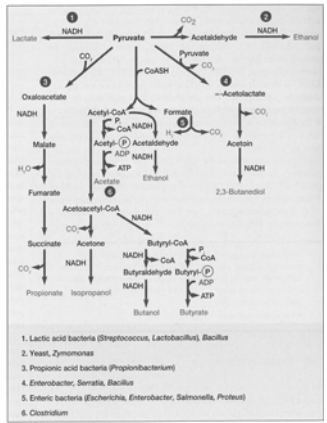


Figure 9.9 Reoxidation of NADH During Fermentation. NADH from glycolysis is reoxidized by being used to reduce pyruvate or a pyruvate derivative (X). Either lactate or reduced product Y result.

Figure 9.10 Some Common Microbial Fermentations. Only pyruvate fermentations are shown for the sake of simplicity; many other organic molecules can be fermented. Most of these pathways have been simplified by deletion of one or more steps and intermediates. Pyruvate and major end products are shown in color.



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

