

Chapter 5 Metabolism: *Aerobic Respiration*

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 Bio 139 Fall 2006
 Office Hours: Mondays & Wednesdays,
 8:30-10:00 AM

Some figures taken from Krogh *Biology: A Guide to the Natural World*

Bacterial Metabolism: 3 pathways to extract energy from glucose

- **Glycolysis**, followed by either
- **Fermentation**
 - oxidize NADH, usually without producing ATP
 - does NOT use oxygen
 - or
- **Aerobic Respiration**
(Krebs cycle, electron transport, oxidative phosphorylation)

To respire aerobically, a bacterium needs:

- Oxygen
- A cytochrome system
 - Indirectly tested for with *catalase test*
- (For facultative anaerobes that can ferment), a low concentration (<0.1%) of a fermentable sugar
 - If the concentration is higher, the organism will ferment it even in the presence of O₂ (TSI test!!!) because it takes fewer enzymes/fewer reactions
 - If the bacteria CANNOT ferment the sugar, they will use it aerobically at ANY concentration, high or low

And of course, all relevant enzymes for the various biochemical pathways.

Aerobic Respiration: steps

- **Production of acetyl-CoA** from pyruvate
- **Krebs Cycle**: also called
 - Tricarboxylic acid cycle
 - TCA cycle
 - Citric acid cycle
- **Electron Transport**
 - Redox reactions with energy transfer
- **Oxidative phosphorylation**
 - Formation of ATP from ADP + Pi

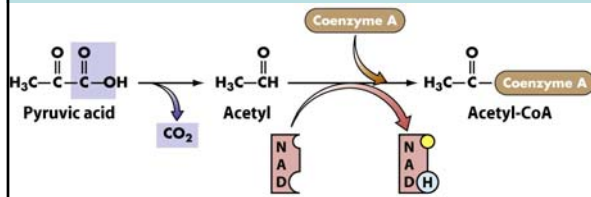
Key points:

- Done **INSTEAD** of fermentation
- Pyruvic acid/pyruvate is the **starting material**
- NADH is **oxidized** to NAD⁺ (in the end)
- Loads of **ATP** is produced
- **Oxygen** is the terminal electron acceptor, reduced to water

Acetyl-CoA

- Pyruvate is *decarboxylated* into a 2-carbon unit (with release of CO₂)
- Product is oxidized (NADH produced) and bound to **Coenzyme A** to make the starting product for TCA cycle,

★ Acetyl-CoA



★ Coenzyme A

Coenzyme A is *not* a protein

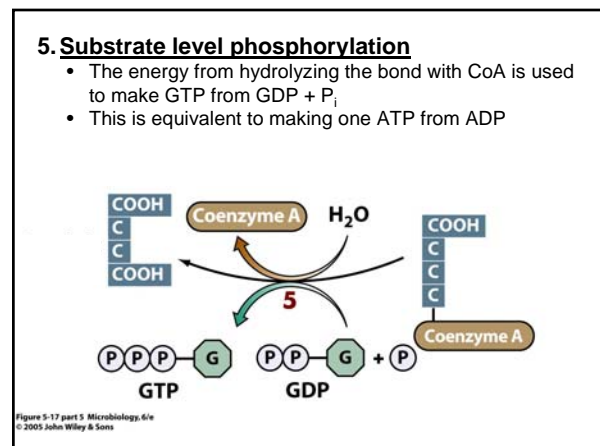
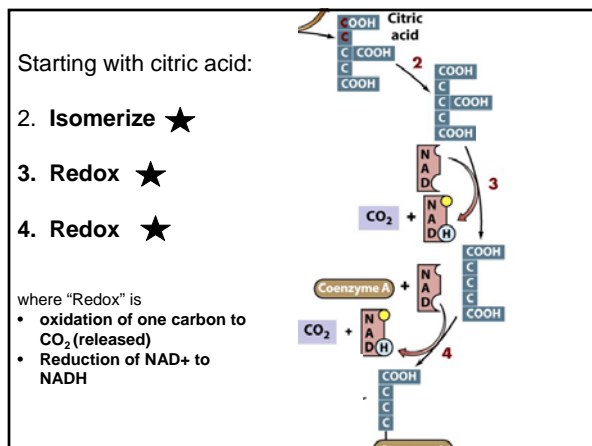
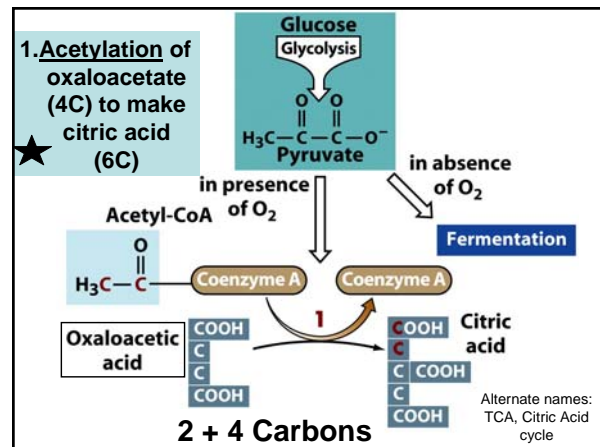
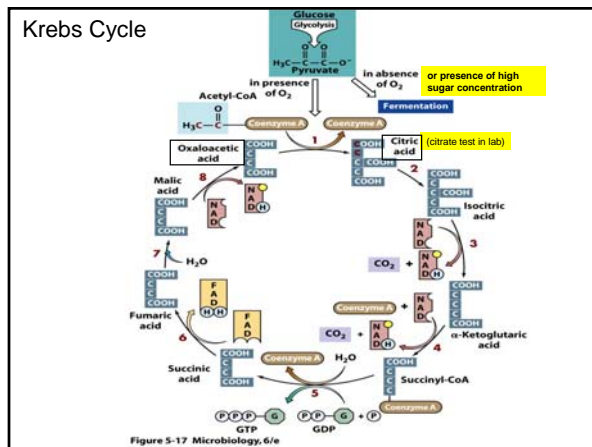
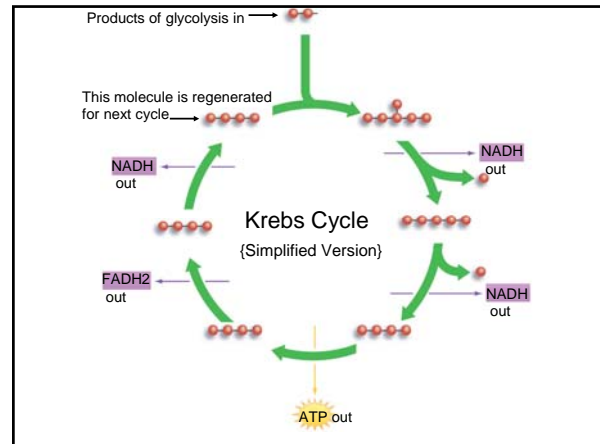
It is basically ADP with the vitamin **pantothenic acid** bound to it

It functions in multiple biochemical pathways as an "activated" (high energy) *carrier* of acetyl (2 carbon) groups (**acetyl CoA**)

Pyruvate is converted into an "activated" Acetyl-CoA molecule for oxidation through the Krebs cycle

Krebs Cycle: What happens

1. Those **2 Carbons** in the acetyl group (where did they come from originally?) will be **oxidized** to CO_2
2. **Electron carriers** (NAD^+ and FAD) will be **reduced** to NADH and FADH_2
3. **Substrate level phosphorylation**
4. **Regeneration of critical cycle component oxaloacetate**



The carbons derived from glucose are GONE.

These reactions **regenerate** the crucial starting material **oxaloacetate**, allowing another **cycle** to begin.

Additional energy (reducing potential) is harvested by redox reactions:

6. FAD to FADH₂

8. NAD⁺ to NADH

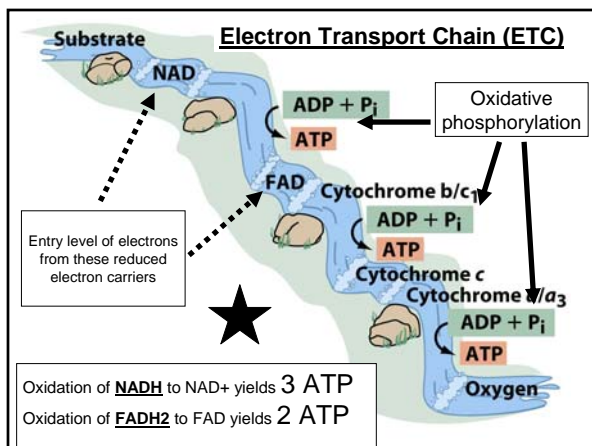
Energy extracted *after* glycolysis

Substrate-level phosphorylation:	Per glucose molecule:
Krebs 1 GTP	= 2 GTP
Reducing Power:	
Acetyl-CoA Production 1 NADH	= 8 NADH
Krebs 3 NADH	
1 FADH ₂	= 2 FADH ₂

Where does it happen? Aerobic Respiration & the endosymbiotic theory

All cells:	
Glycolysis	Cytoplasm
Bacteria:	
Krebs	Cytoplasm
ETC	Cell membrane
Eukaryotes:	
Krebs	Mitochondrial matrix
ETC	Inner mitochondrial membrane

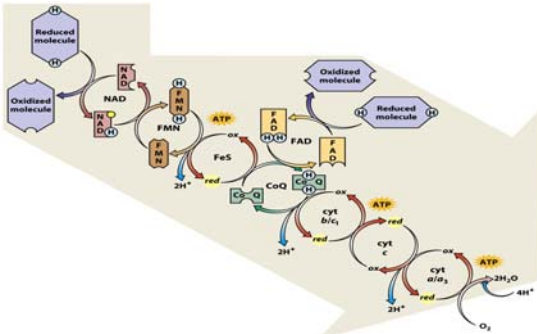
- ### Electron transport & Oxidative Phosphorylation
- ★ Goal: Extract *useful* energy (ATP) by oxidizing reduced electron carriers
 - Energy-carrying electrons of NADH & FADH₂ are passed through a series of electron carriers
 - At each passage, some energy is released
 - At **three** points in the chain of electrons, enough energy is released to power production of ATP from ADP + P_i
 - Last electron acceptor in the chain: Oxygen



- ### Carriers of the Electron Transport Chain
- Each species has a different set of specific carrier molecules but always including the following groups:
1. **Flavin mononucleotides** (derivative of vitamin riboflavin)
 2. **Iron sulfide** (FeS) centers
 3. **Coenzyme Q** (actually a lipid)
 - ★ 4. **Cytochromes** (proteins with iron-porphyrin groups like heme)
Catalase test correlates with presence of cytochromes
Oxidase test detects a specific one, cytochrome c

This figure is NOT helpful except to illustrate this point:

The ETC is a series of linked redox reactions, with multiple energy drops (some captured to make ATP), ending in oxygen being reduced to water



About those energy drops...

Where does the energy go? ★

It is *not* used to make ATP directly
(that would be substrate-level phosphorylation)

- Molecules of the ETC are in the cell membrane of bacteria
- The energy released by electron transfers is used to pump protons across the membrane (out of the cell) by active transport

This generates a proton (hydrogen ion, H⁺) gradient

Proton Motive Force

- Energy is stored in the proton gradient
 - Higher [H⁺] outside than inside: concentration gradient
 - Outside of membrane is + charged relative to inside: electrical gradient

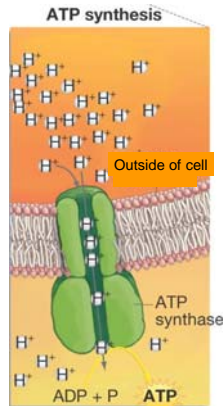
Collectively, this generates a proton motive force ★

Chemiosmosis

- The energy stored in the proton gradient is used to make ATP in a process called chemiosmosis
- Protons are allowed *back* into the cell through ATP synthase complex (a transmembrane protein channel)
- Energy released from neutralizing the proton gradient is used to synthesize ATP from ADP + P_i

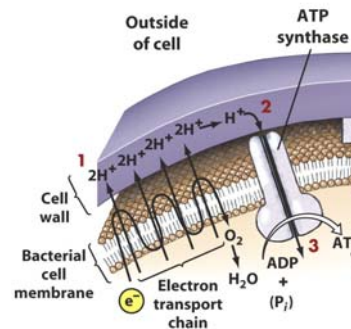
Chemiosmosis

- Hydrogen ions “fall” DOWN the concentration /electrical gradient through ATP synthase protein
- Energy of “falling” is coupled to ATP production



Chemiosmosis:

1. Proton gradient is generated by ETC
2. Gradient is released by ATP synthase & coupled to ATP production



Toxins that affect aerobic respiration

★ Cyanide & Azide

- Affect function of cytochromes
- Block electron transport
- Inhibits growth of catalase + bacteria
- Catalase – bacteria unaffected

★ Uncoupling agents

- "uncouple" dissipation of the proton gradient from ATP synthesis
- The energy stored in the gradient is wasted
- Example: dinitrophenol

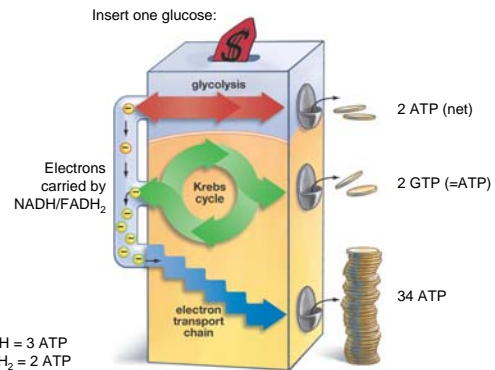


TABLE 5.2

Energy Captured in ATP Molecules from a Glucose Molecule by Anaerobic and Aerobic Metabolism in Prokaryotes

Prokaryotic Metabolic Process	Number of ATP Molecules	
	Anaerobic Conditions	Aerobic Conditions
<i>Glycolysis</i>		
Substrate level	4	4
Hydrogen to NAD	0	6
<i>Pyruvate to Acetyl-CoA</i>		
Hydrogen to NAD	0	6
<i>Krebs Cycle</i>		
Substrate level	0	2
Hydrogen to NAD	0	18
Hydrogen to FAD	0	4
<i>Less Energy for Phosphorylation</i>	-2	-2
Total	2	38

Table 5.2 Microbiology 6/e
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Aerobic metabolism & Growth

Bacteria using aerobic respiration for ATP production grow **FASTER**

(strict aerobes, facultative anaerobes in the *presence* of oxygen)

than

Bacteria relying on glycolysis & fermentation

(indifferent/aerotolerant, facultative anaerobes in the *absence* of oxygen)

- ★ You can see this in colony size: catalase – bacteria will generally grow more slowly / have smaller colonies than catalase + bacteria under the same aerobic conditions

TABLE 5.3

A Comparison of Metabolic Processes

	Glycolysis	Fermentation	Krebs Cycle*	Electron Transport Chain
Location	In cytoplasm	In cytoplasm	Prokaryotes: in cytoplasm Eukaryotes: in the mitochondrial matrix	Prokaryotes: in cell membrane Eukaryotes: in inner mitochondrial membranes
Oxygen Conditions	Anaerobic; oxygen is not required; does not stop, however, if oxygen is present	Without O ₂ ; presence of oxygen will cause it to stop	Acrobic	Acrobic
Starting Molecule(s)	1 glucose (6C)	Various substrate molecules go through glycolysis, yielding 2 pyruvic acid	2 pyruvic acid	6 O ₂
Ending Molecules	2 pyruvic acid (3C) 2 NADH	Various, depending on which form of fermentation occurs, e.g., ethanol, lactic acid, CO ₂ , acetic acid	6 CO ₂ 8 NADH 2 FADH	6 H ₂ O
Amount of ATP Produced	4 ATP (net 2 ATP)	Various, depending on which form of fermentation occurs, usually 2 or 3 ATP; always far less than is produced in aerobic respiration	2 GTP (= 2 ATP)	34 ATP

*Includes the pyruvic acid → acetyl-CoA step.

Table 5.3 Microbiology 6/e
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★ (unless a high concentration of fermentable sugar is present)

Anaerobic Respiration

★ An electron transport chain is used

- Final electron acceptor is NOT O₂

★ NO₂⁻, NO₃⁻, SO₄²⁻ are reduced instead

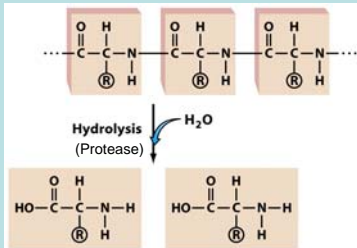
- Produce fewer ATP than aerobic respiration (but better than fermentation)

Protein catabolism

Proteins can be broken down for energy

First step:

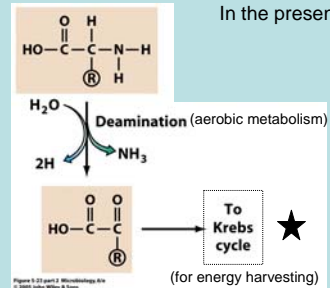
★ **Hydrolysis into amino acids**
by **proteases**



Next step for **aerobic** catabolism of proteins:

★ **Deamination**

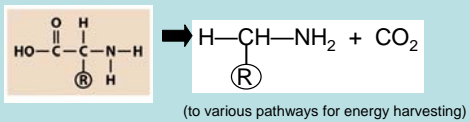
In the presence of oxygen!



For **anaerobic** catabolism of proteins:

★ **Decarboxylation**

In the absence of oxygen!



Deamination and decarboxylation reactions are catalyzed by enzymes that are **amino acid-specific**

Enzyme expression is **species specific**