

BIOL 240: General Microbiology

Spring 2020 Rm. 23-203 MW, Mar. 9-11

<http://accounts.smccd.edu/staplesn/biol240/>

1. **Pre-Lab Writeup #9 due!**: ALL of Expt. 10. Prepare before each Monday's labs (for BOTH Mon. & Wed.)!! (*What? Why? How? are we doing in the lab? Question? HYPOTHESIS?*)
2. **0.5-1 hour of OPEN LAB *expected* each week. Goal: 8-16h by May!** ☺
3. **LAB: Bacteriophages!!**
4. **QUIZ #4 DUE Wed. night!!!**
5. **Study Guides & Objectives (see slides) due THIS WEEK!!** (Chs. 5b, 6, 7)!
6. **Read Chs. 6 & 7!! See Chs. 6 & 7 lecture ONLINE!!**
Obj's. due THIS week! (MT2 Review Sheet is UPDATED!!)
7. **Midterm #1 REVIEW SESSIONS THIS WEEK: Mon/Wed 2-3pm.... in LAB. Wed. 8:15am-9:35ish.** ☺ Come PREPARED with questions!
8. **Extra Credit Opportunity: Wed. evenings, 5-6 pm, (2/4-3/25/2020) in Bldg. 6, Room 102 – STEM SPEAKER Series. 1 page summary & reflection due (on CANVAS) the following week. (see also: YOUTUBE)**
 - <https://www.canadacollege.edu/stemcenter/speaker-series.php>
 - NOTE: YOU may upload MULTIPLE Speaker Summaries, but only I can see all of them. You can only see the last upload! It's OK!! ☺

1

REVIEW:

1. Diagram the investment and release of Energy and Carbon atoms from Glycolysis (through the Citric Acid Cycle).
2. Diagram or outline the process by which high energy electron carriers are produced by glycolysis, pyruvate oxidation, and the Krebs/TCA cycle.

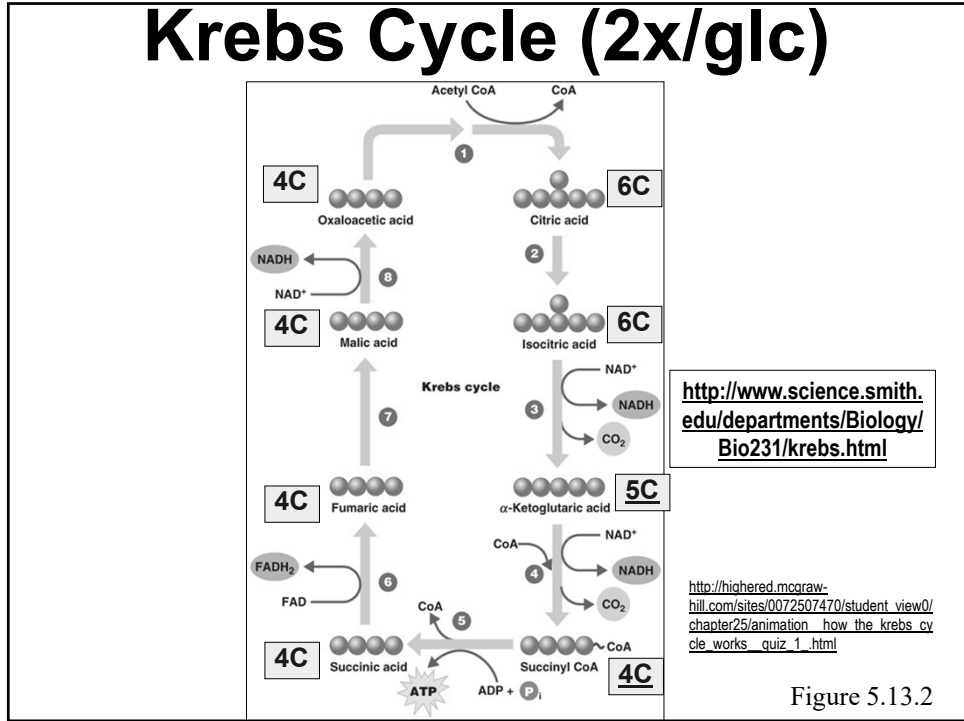
OBJECTIVES: Students should be able to:

1. Diagram how high energy electrons are used to produce ATP in the mitochondrial inner membrane (or bact. plasma mem).
2. Compare and contrast the energy inputs and outputs of Fermentation & Aerobic Respiration/Oxidative phosphorylation.
3. Explain how lipids and proteins are catabolized and energy harvested thru pathways shared with glucose metabolism.
4. Distinguish the carbon and energy sources for all of the trophisms: *chemo, photo, hetero, auto*, & comb'ns of each.
5. Diagram how catabolic and anabolic pathways can share intermediates to efficiently regulate energy storage, energy usage, and biosynthesis.
6. **Ch. 8:** Describe the **Central Dogma** of molecular genetics and the three processes that drive the flow of genetic information in an organism.
7. Describe several properties of DNA and the process of DNA replication that contribute to DNA's central role as the hereditary material.
8. Draw a replication fork and label 5 enzymes involved in DNA replication. Describe the function of each enzyme.

❖ **These questions are your HOMEWORK between classes!!!**
➢ **DUE (and/or Study Guide questions) WED. at the start of Lab!!**

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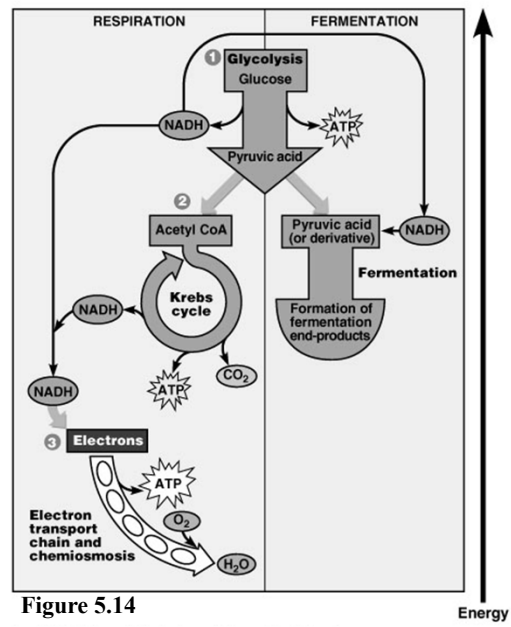
Krebs Cycle (2x/glc)



3

D. The Electron Transport Chain

- A series of carrier molecules that are, in turn, oxidized and reduced as electrons are passed down the chain.
- Energy released can be used to produce ATP by **ChemiOsmosis**.



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E. Chemi-osmosis

- Each NADH → ~3 ATP
- Each FADH₂ → ~2 ATP

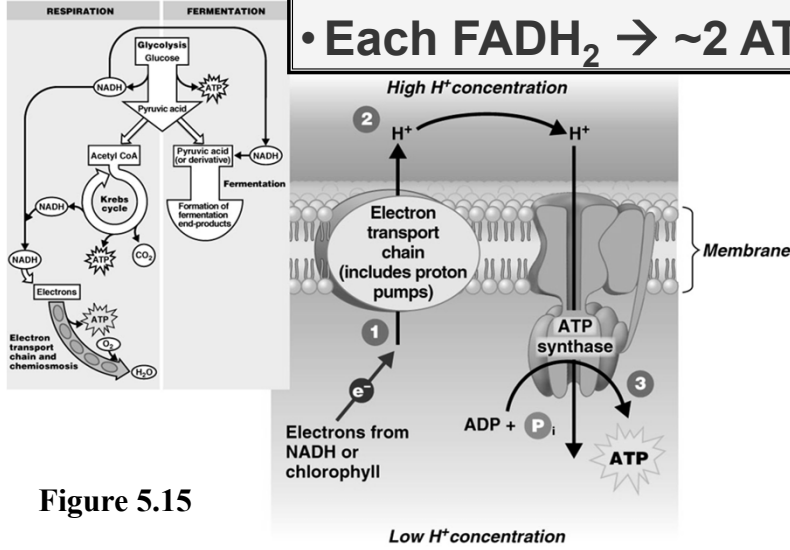


Figure 5.15

➤ Create gradient of [H⁺] and electric charge — the Proton-Motive Force.

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Chemi-osmosis

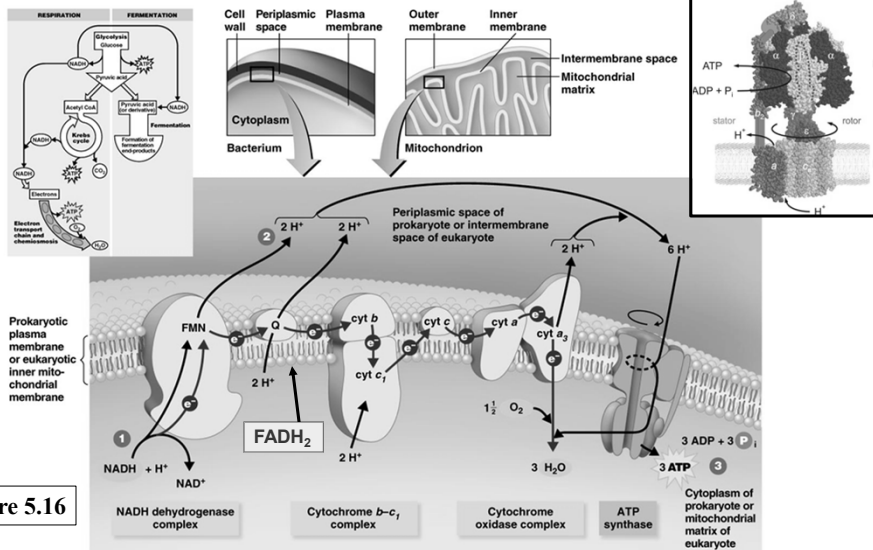


Figure 5.16

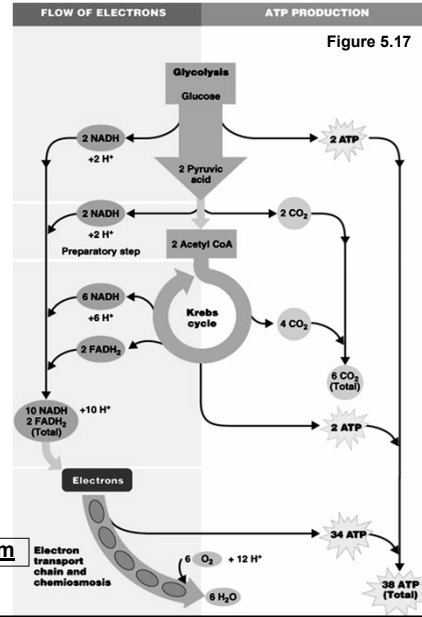
• Stronger Electron-Acceptors/Oxidizers →→

<http://vcell.ndsu.nodak.edu/animations/etc/movie.htm>

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5.6) Respiration

- **Aerobic respiration:** The final electron acceptor in the electron transport chain is molecular oxygen (O_2).
- *****ETC + Chemiosmosis = Oxidative Phosphorylation!!*****
- **Anaerobic respiration:** The final electron acceptor in the electron transport chain is not O_2 .
 - Yields less energy than aerobic respiration
 - only part of the Krebs cycle operates w/out O_2 .



<http://vcell.ndsu.nodak.edu/animations/etc/movie.htm>
<http://www.science.smith.edu/departments/Biology/Bio231/krebs.html>
<http://www.science.smith.edu/departments/Biology/Bio231/etc.html>

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A. Aerobic Respiration


<u>Pathway</u>	<u>Eukaryote</u>	<u>Prokaryote</u>
Glycolysis	Cytoplasm	Cytoplasm
Intermediate step (Pyruvate Ox'n)	Mito. Inner Memb	Cytoplasm
Krebs cycle	Mitochondrial matrix	Cytoplasm
ETC	Mitochondrial inner membrane	Plasma membrane

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- **Energy produced (types)** – from complete oxidation of 1 glucose using aerobic respiration

Pathway	ATP produced	NADH produced	FADH ₂ produced
Glycolysis	2	2	0
Pyruvate Oxidation	0	2	0
Krebs cycle	2	6	2
Total	4	10	2

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- ❖ **ATP produced (chemical processes)** from complete oxidation of 1 glucose using aerobic respiration. 

Pathway	By Substrate-Level Phosphorylation	By Oxidative Phosphorylation	
		From <u>NADH</u>	From <u>FADH₂</u>
Glycolysis	2	6	0
Pyruvate Oxidation	0	6	0
Krebs cycle	2	18	4
<u>Total</u>	<u>4</u>	<u>30</u>	<u>4</u>

- **38** ATPs are produced in *prokaryotes*, **36** in *Euk.*

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Aerobic Metabolic Yield, Summary (Eukaryotes):

NET:

1. Glyc = 2 ATP (4 made, 2 invested)
 $\underline{2 \text{ NADHH} \rightarrow 4 \text{ ATP}}$ (use 2 ATP on transport into mito IM!!)
 2. Pyr.Ox. = $2 \text{ NADHH} \rightarrow 6 \text{ ATP}$
 3. TCA = $6 \text{ NADHH} \rightarrow 18 \text{ ATP}$
 $2 \text{ GTP} \rightarrow 2 \text{ ATP}$
 $\underline{2 \text{ FADH}_2 \rightarrow 4 \text{ ATP}}$
- 36 ATP total NET** from Glycolysis
& Oxidative Respiration

<http://vcell.ndsu.nodak.edu/animations/atpgradient/movie.htm>

http://www.wiley.com/legacy/college/boyer/0470003790/animations/electron_transport/electron_transport.swf

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B. Anaerobic Respiration

= truly *RESPIRATION*, and NOT glycolysis/fermentation!

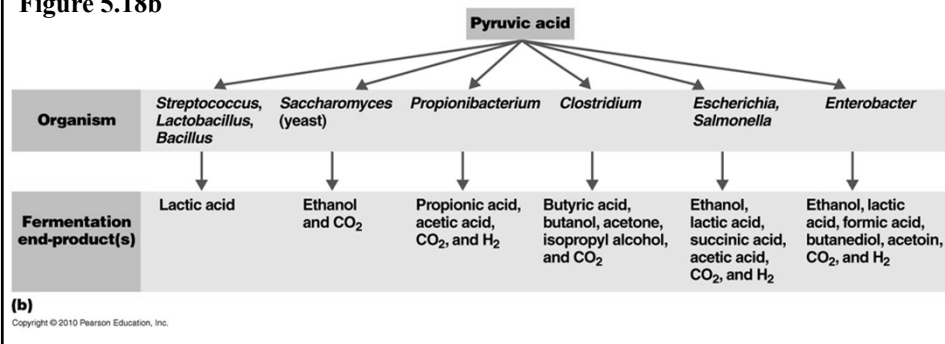
<u>Electron acceptor</u>	<u>Products</u>
NO_3^-	$\text{NO}_2^-; \text{N}_2 + \text{H}_2\text{O}$
SO_4^{2-}	$\text{H}_2\text{S} + \text{H}_2\text{O}$
CO_3^{2-}	$\text{CH}_4 + \text{H}_2\text{O}$

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5.7) Fermentation

1. Releases energy from oxidation of organic molecules
2. Does not require oxygen
3. Does not use the Krebs cycle or ETC
4. Uses an organic molecule as the final electron acceptor

Figure 5.18b

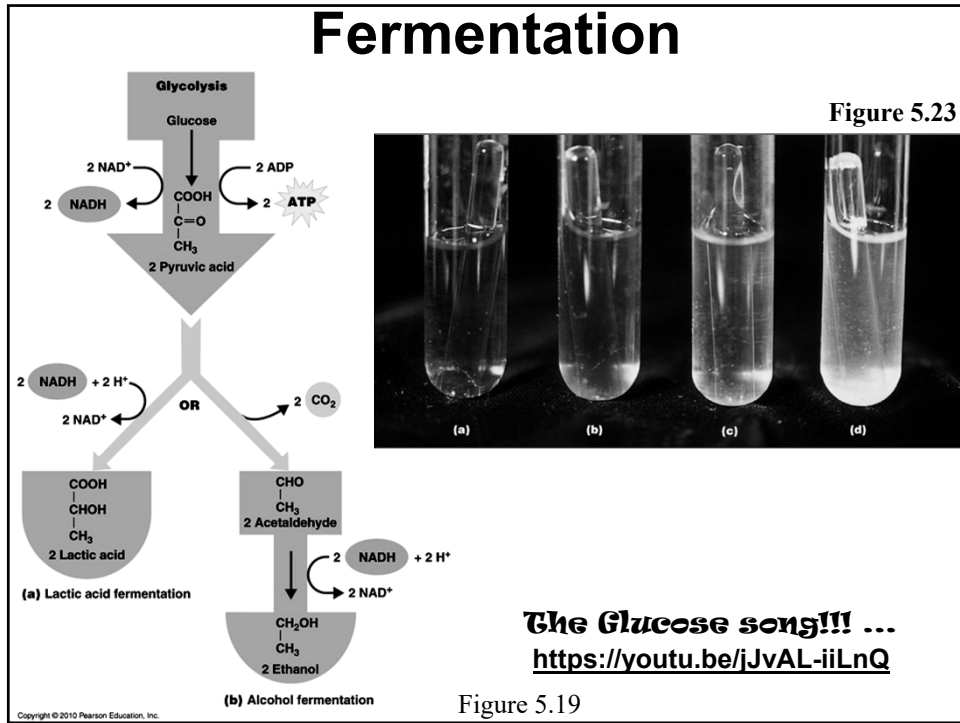


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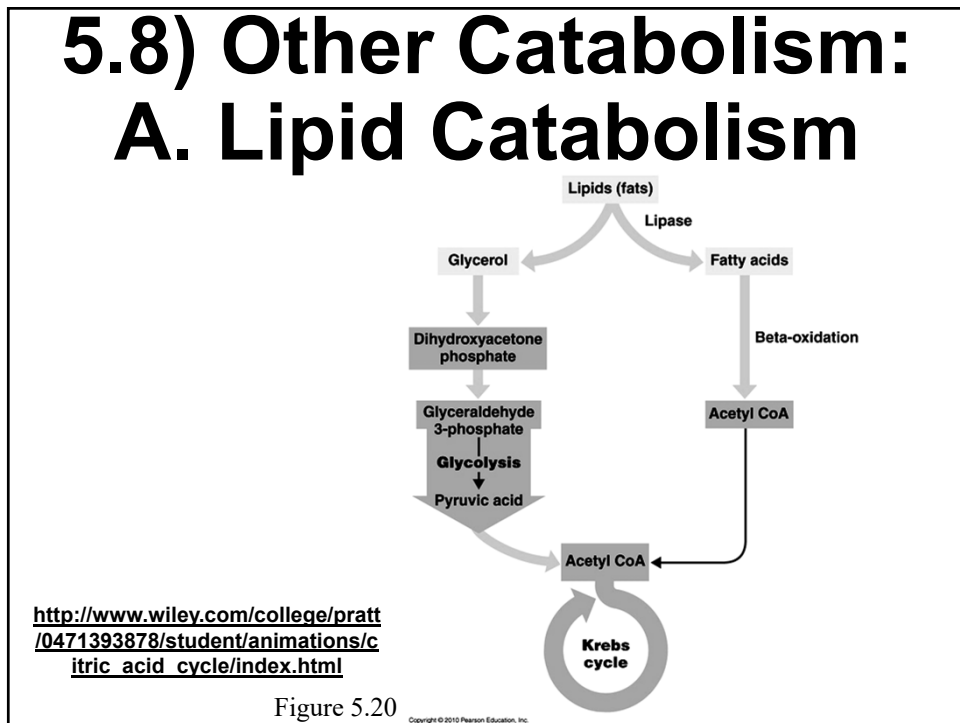
Types of Fermentation

- A. Alcohol fermentation** - Produces ethyl alcohol + CO₂
- B. Lactic acid fermentation** - Produces lactic acid.
- **Homolactic fermentation** - Produces lactic acid only.
 - **Heterolactic fermentation** - Produces lactic acid and other compounds (eg: acetoin).

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B. Protein Catabolism

Protein $\xrightarrow{\text{Extracellular proteases}}$ Amino acids

Deamination (\rightarrow acid), decarboxylation (\rightarrow amine), dehydrogenation \rightarrow Organic acid \rightarrow Krebs cycle

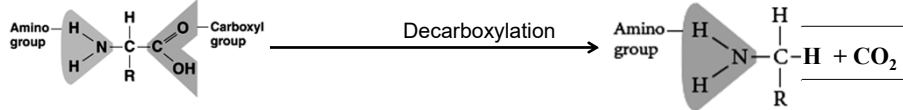
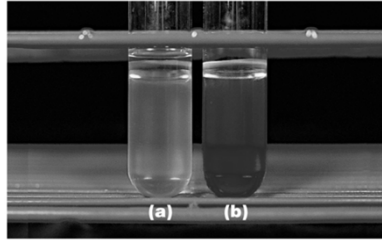


Figure 5.22

<http://www.wiley.com/legacy/college/boyer/0470003790/animations/tca/tca.htm>

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** Biochemical tests **

- Used to identify bacteria.

Dichotomous

Key:

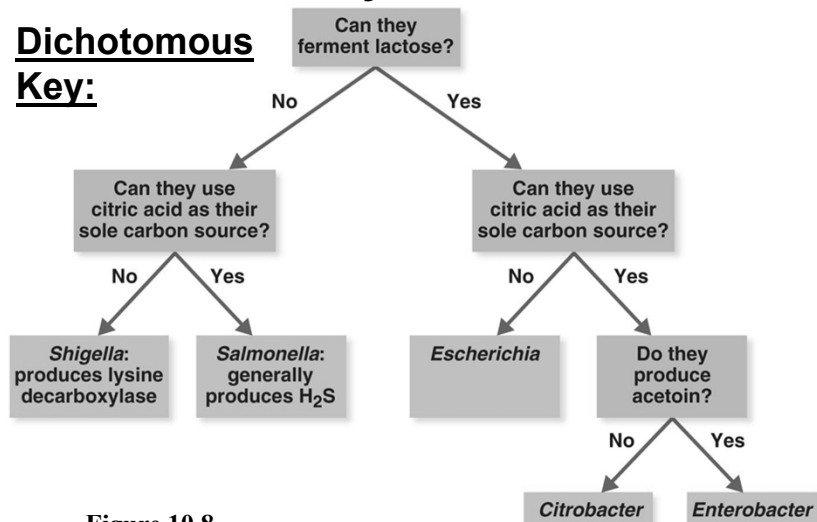


Figure 10.8

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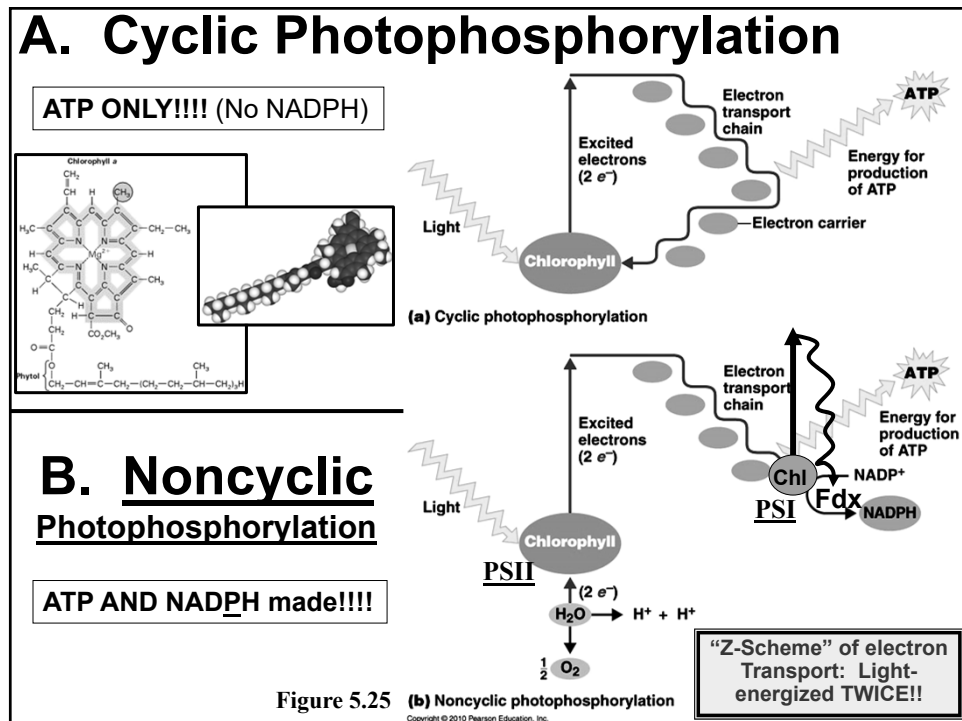
5.9) Photosynthesis

- **Photo:** Conversion of light energy into chemical energy (ATP).
 - Light-dependent (light) reactions
- **Synthesis:** Fixing carbon into organic molecules.
 - Light-independent (dark) reaction, Calvin-Benson cycle
 - **Oxygenic:**

$$6 \text{ CO}_2 + 12 \text{ H}_2\text{O} + \text{Light energy} \rightarrow \text{C}_6\text{H}_{12}\text{O}_6 + 6 \text{ O}_2 + 6 \text{ H}_2\text{O}$$
 - **Anoxygenic:**

$$\text{CO}_2 + 2 \text{ H}_2\text{S} + \text{Light energy} \rightarrow [\text{CH}_2\text{O}] + 2 \text{ S} + \text{H}_2\text{O}$$

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Noncyclic Pathway

“Z-Scheme” of electron Transport: Light-energized TWICE!!

<http://vcell.ndsu.nodak.edu/animations/photosynthesis/movie.htm>
<http://www.science.smith.edu/departments/Biology/Bio231/ltrxn.html>
<http://www.fw.vt.edu/dendro/forestbiology/photosynthesis.swf>
<http://www.stolaf.edu/people/giannini/flashanimat/metabolism/photosynthesis.swf>

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C. Calvin-Benson Cycle: Carbon-Fixation!

Figure 5.26

- **SYNTHESIS!!! = “Dark Reactions”**
- Using ATP & NADPH made in the light reactions!!

Step-By-Step Narration:
http://www.cells.de/cellseng/1medienarchiv/Zellfunktionen/Memb_Vorg/Photosynthese/Dunkel_u_Saerke/Calvin-Benson-Zyklus/index.jsp

**<http://www.science.smith.edu/departments/Biology/Bio111/calvin.html>

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Characteristic	Eukaryotes		Prokaryotes	
	Algae, Plants	Cyanobacteria	Green Bacteria	Purple Bacteria
Substance That Reduces CO ₂	H atoms of H ₂ O	H atoms of H ₂ O	Sulfur, sulfur compounds, H ₂ gas	Sulfur, sulfur compounds, H ₂ gas
Oxygen Production	Oxygenic	Oxygenic (and anoxygenic)	Anoxygenic	Anoxygenic
Type of Chlorophyll	Chlorophyll <i>a</i>	Chlorophyll <i>a</i>	Bacteriochlorophyll <i>a</i>	Bacteriochlorophyll <i>a</i> or <i>b</i>
Site of Photosynthesis	Chloroplasts with thylakoids	Thylakoids	Chlorosomes	Chromatophores
Environment	Aerobic	Aerobic (and anaerobic)	Anaerobic	Anaerobic

****Study figure 5. 28!! ****
-- nutritional flow chart of organisms**

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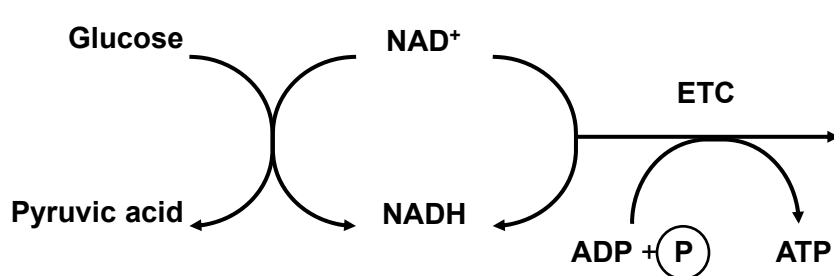
- ***Halobacterium*** uses **bacteriorhodopsin**,
 - not chlorophyll,
 - to generate electrons for a *chemiosmotic proton pump*.
 - Different pools drying, with different salinities.
 - Different pigmented species of *Halobacteria* grow in diff't [salt]s.



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5.10) Chemotrophs

- Use energy from chemicals.
 - ***Chemoheterotroph***.



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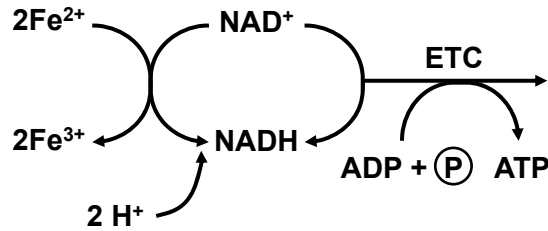
    graph LR
      Glucose --> Pyruvic_acid[Pyruvic acid]
      Pyruvic_acid --> NADH
      NADH --> ETC[ETC]
      ETC --> ATP
      ADP_P[ADP + P] --> ATP
      NADH --> NADp[NAD+]
      NADp --> NADH
    
```

- Energy is used in anabolism.

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Chemotrophs

- Use energy from chemicals.
 - **Chemoautotroph**, *Thiobacillus ferrooxidans*

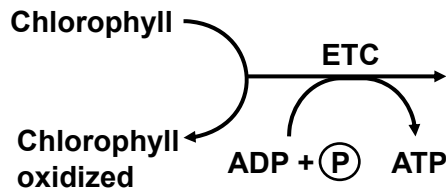


- Energy used in the **Calvin-Benson** cycle to fix CO_2 .

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5.11) Phototrophs

- Use light energy.



- **Photoautotrophs** use energy in the Calvin-Benson cycle to fix CO_2 .
- **Photoheterotrophs** use energy.

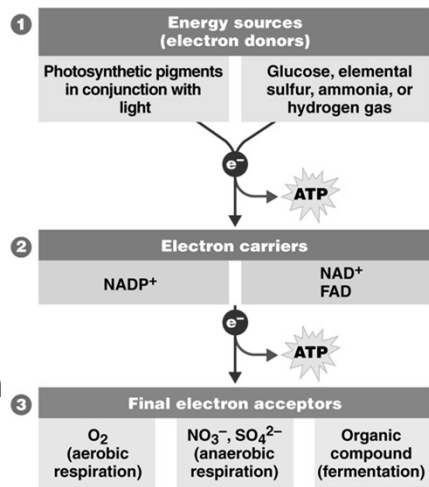


Figure 5.27

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Metabolic Diversity Among Organisms

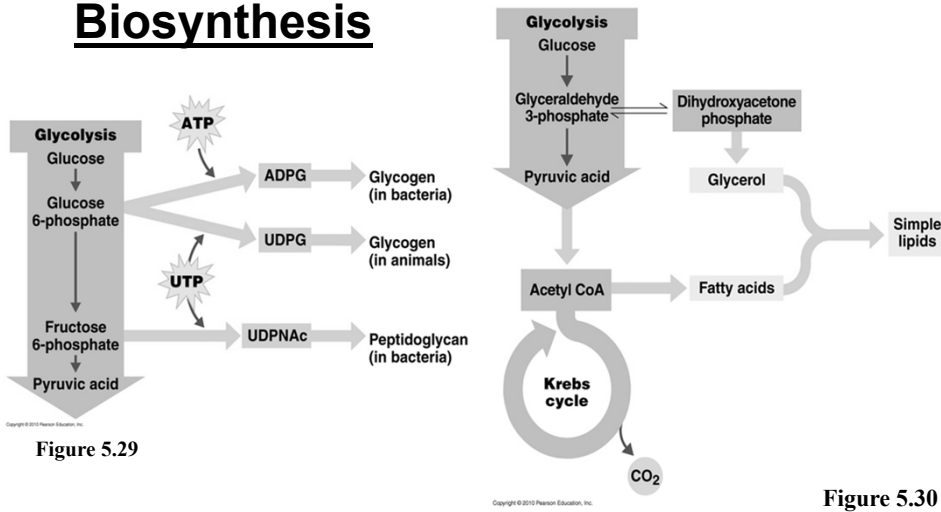
Nutritional type	Energy source	Carbon source	Example
Photoautotroph	Light	CO ₂	Oxygenic: Cyanobacteria plants. Anoxygenic: Green, purple bacteria.
Photoheterotroph	Light	Organic compounds	Green, purple nonsulfur bacteria.
Chemoautotroph	Chemical	CO ₂	Iron-oxidizing bacteria.
Chemoheterotroph	Chemical	Organic compounds	Fermentative bacteria. Animals, protozoa, fungi, bacteria.

**** Study figure 5.28!! ****

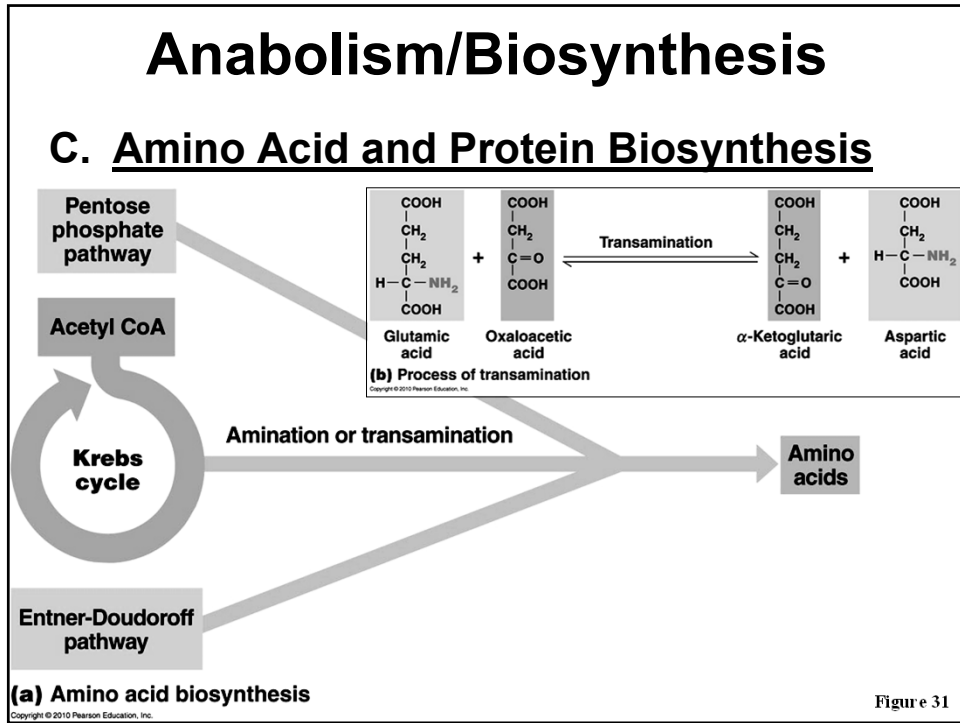
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5.12) Anabolism: Metabolic Pathways of Energy Use

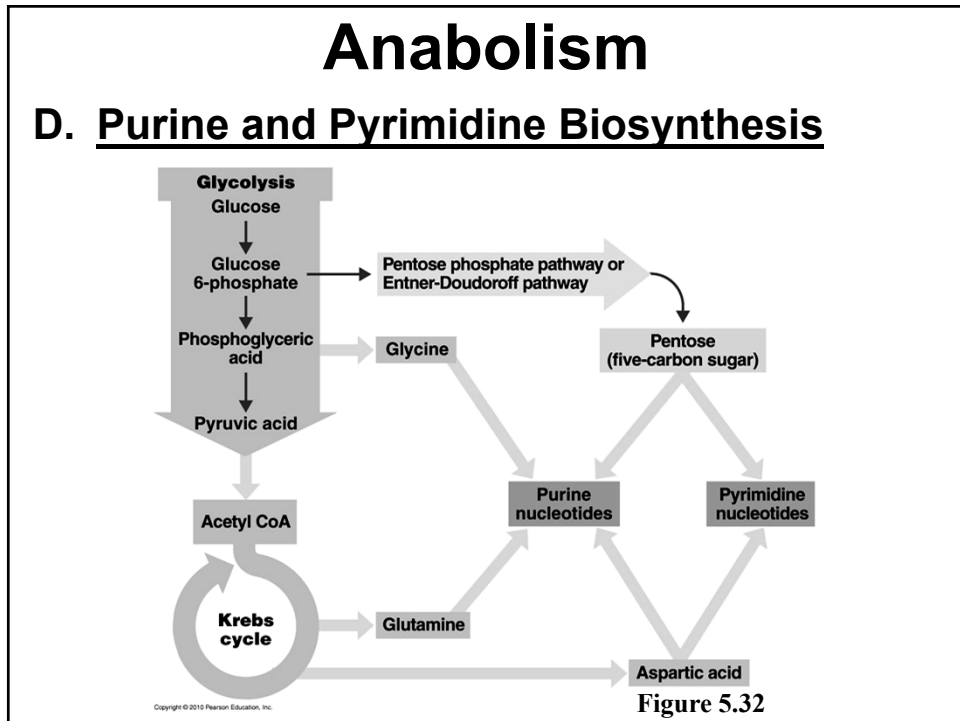
A. Polysaccharide Biosynthesis B. Lipid Biosynthesis



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5.13) Amphibolic pathways

- Are metabolic pathways that have both catabolic and anabolic functions.

Glucose, Glucose!!
<https://youtu.be/jJvAL-iiLnQ>

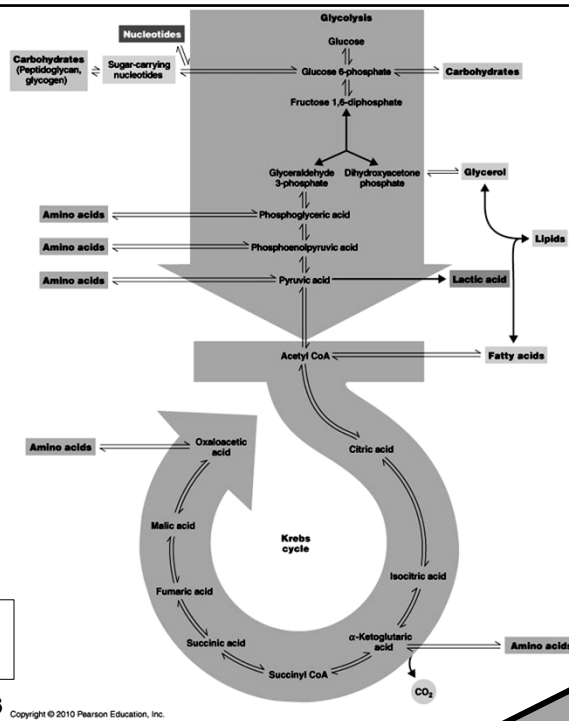
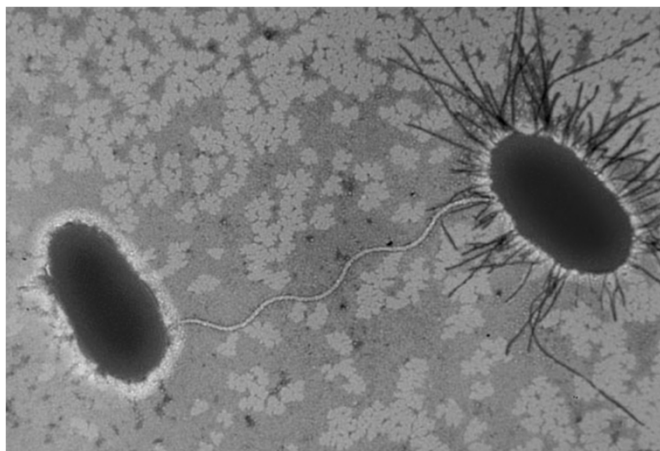


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Chapter 8 Microbial Genetics



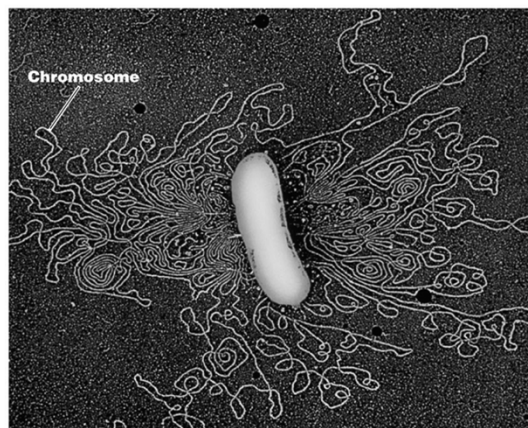
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Terminology

1. **Genetics:** Study of what genes are, how they carry information, how information is expressed, and how genes are replicated; “the science of heredity”
2. **Gene:** Segment of DNA that encodes a functional product, usually a protein
3. **Genome** = All of the genetic material in a cell
4. **Genomics** = Molecular study of genomes
5. **Genotype** = Specific forms of genes in an organism
 - Types of alleles present.
6. **Phenotype** = *physical characteristics resulting from expression of the genes*

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E. coli

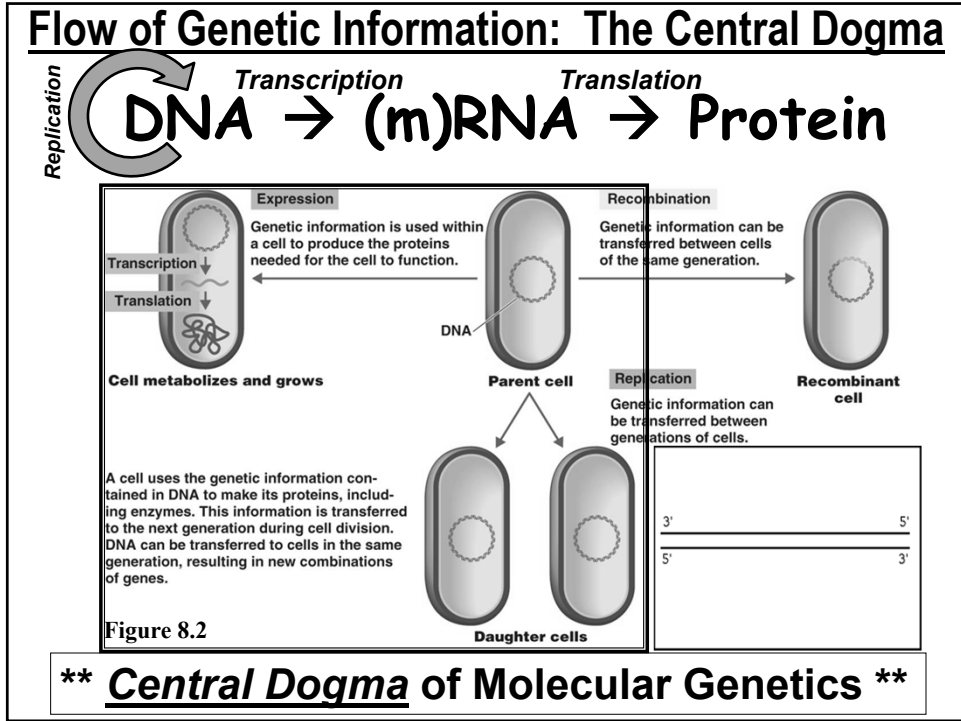


(a) The tangled mass and looping strands of DNA emerging from this disrupted *E. coli* cell are part of its single chromosome.

The Model Organism for molecular biology.

Figure 8.1a

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8.1) DNA

- Polymer of nucleotides: adenine, thymine, cytosine, guanine (**ATGC**)
- Double helix associated with proteins
- "Backbone" is deoxyribose-phosphate
- Strands held together by hydrogen bonds between **A=T** and **G=C**
- Strands are **Antiparallel**
 - **5' (PO₄) → 3' (OH) polarity**

Figure 8.3

(b) The two strands of DNA are antiparallel. The sugar-phosphate backbone of one strand is upside down relative to the backbone of the other strand. Turn the book upside down to demonstrate this.

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A. DNA Structure

- **Semi-Conservative Replication:**
 - Each “parental strand” serves as *template*...
 - for synthesis of a new “daughter strand”
 - Following **complementary base-pairing** rules
 - **A=T**
 - **G≡C**
 - Rules allow one to predict 2nd strand sequence from 1st!!!

KEY

- T Thymine
- A Adenine
- C Cytosine
- G Guanine
- Deoxyribose sugar
- Phosphate

1 The double helix of the parental DNA separates as weak hydrogen bonds between the nucleotides on opposite strands break in response to the action of replication enzymes.

2 Hydrogen bonds form between new complementary nucleotides and each strand of the parental template to form new base pairs.

3 Enzymes catalyze the formation of sugar-phosphate bonds between sequential nucleotides on each resulting daughter strand.

(a) The replication fork.

Figure 8.3
DNA strand forming

http://www.fed.cuhk.edu.hk/~johnson/teaching/genetics/animations/dna_replication.htm
http://www.uic.edu/classes/cmeng/cmeng521/DNA_replication.html

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B. DNA Replication

New Strand

Template Strand

= DNA-directed DNA synthesis.

Hydrolysis of the phosphate bonds provides the energy for the reaction.

http://www.fed.cuhk.edu.hk/~johnson/teaching/genetics/animations/dna_replication.htm

http://www.uic.edu/classes/cmeng/cmeng521/DNA_replication.html

When a nucleoside triphosphate bonds to the sugar, it loses two phosphates.

- **5' → 3' synthesis.**
- **Template read 3' → 5'.**

<http://www.courses.fas.harvard.edu/~biotext/animations/replication1.swf>
Figure 8.5

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DNA Replication Proteins

- Open helix at **Origin (Ori)**, lay-down primers:
 - DNA Helicase** “melts” strands apart, breaking H-bonds
 - Single-Strand Binding Proteins (**SSB**) keep template strands apart.
 - RNA Primase** lays down first several nucleotides (RNA!!) – gives “starting block” (free 3'-OH) to begin actual DNA synthesis. [RNA Primers are removed later!]
- DNA = copied by **DNA polymerase III** (Dpol3)
- In the **5' → 3'** direction – new nucleotides added to the 3' hydroxyl (-OH) group on deoxyribose in the growing strand – Initiated by an **RNA primer** (RNA Primase enz.)
- Leading strand** synthesized **continuously**
 - follows fork (1/fork; 2/ “bubble”)
- Lagging strand** synthesized **discontinuously**
 - Opposite to fork movement → **Okazaki fragments** (unsealed lagging pieces)
- RNA primers are removed and Okazaki fragments joined by **DNA polymerase I** & **DNA ligase** → “fill and seal” to finish job!!

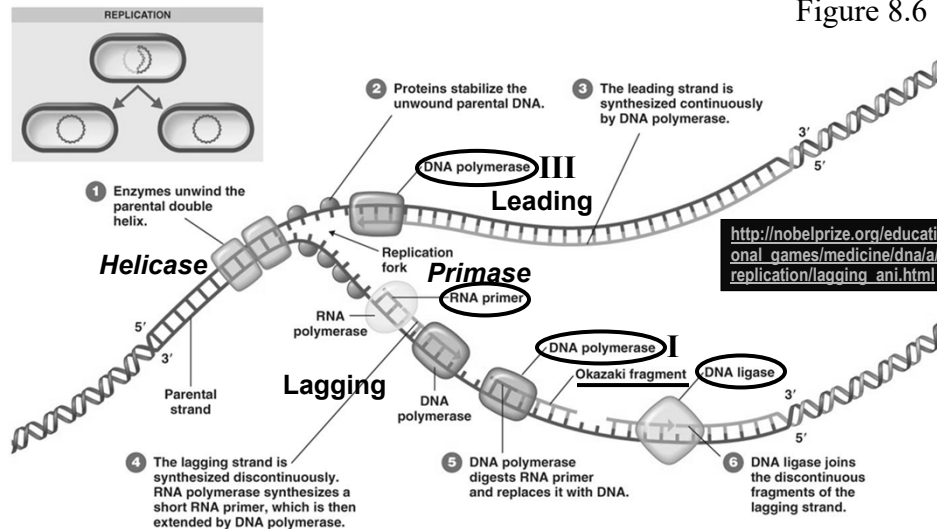
AMOEBASISTERS!!!!
<https://youtu.be/5qSrmeiWsuc>

- <http://www.stolaf.edu/people/giannini/flashanimat/molgenetics/dna-rna2.swf>
 Detailed: http://www.wiley.com/college/pratt/0471393878/student/animations/dna_replication/index.html

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C. DNA Replication Fork

Figure 8.6



http://nobelprize.org/educational_games/medicine/dna/replication/lagging_ani.html

<http://www.stolaf.edu/people/giannini/flashanimat/molgenetics/dna-rna2.swf>

http://highered.mcgraw-hill.com/sites/0072943696/student_view0/chapter3/animation_dna_replication_quiz_1.html
<http://www.ihonkyrk.com/DNAreplication.html> http://www.dnalc.org/resources/3d/DNAReplicationBasic_w_FX.html

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DNA replication is Semiconservative & Bidirectional

Figure 8.6
Bidirectional replication of a circular bacterial DNA molecule

http://www.dnalc.org/resources/3d/DNAReplicationBasic_w_FX.html

<http://www.hhmi.org/biointeractive/dna/animations.html>

http://www.wnorton.com/college/biology/mbio/animations/dna_replication.asp

<http://www.sciencemedia.com/website/demos/biochem/ecoli/Replication.html>

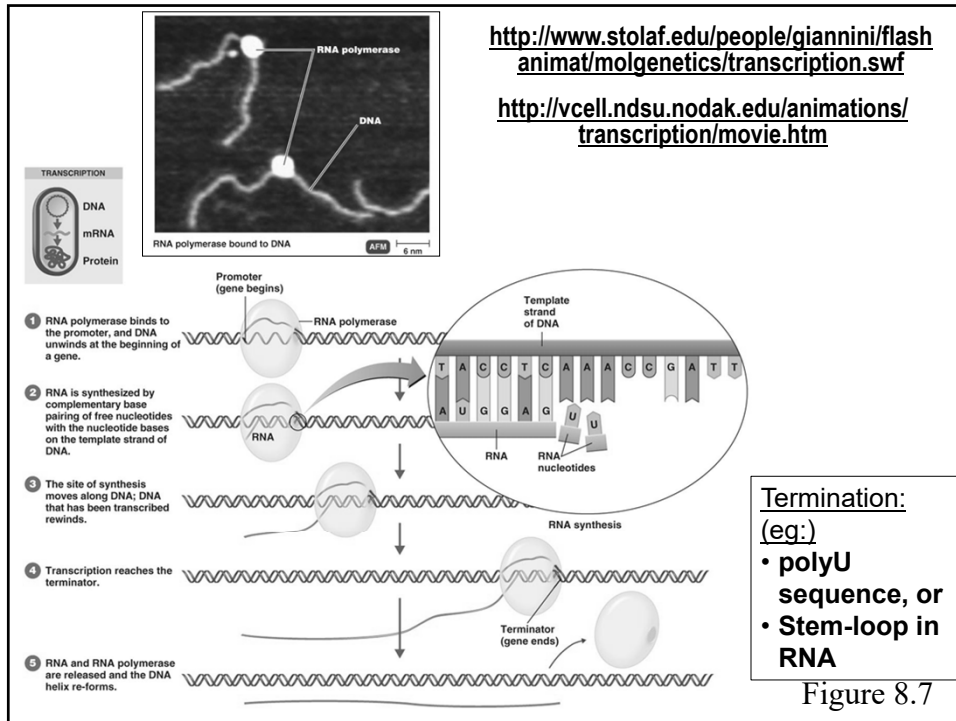
- = **DNA-directed DNA synthesis!!!**
- **2 forks: opposite directions!**
- Replication results in two daughter DNA duplexes.
 - each with one completely new strand, &
 - one old strand (parental strand)
 - = **“SEMI-CONSERVATIVE”**
- **ONLY synthesize 5' → 3'!!!**
- **Two replication forks** move in opposite directions =
- ❖ **“Replication Bubble”**

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8.2) Transcription: RNA Synthesis

1. DNA is transcribed to make RNA (**AUGC**)
 - a) **mRNA** = messenger RNA → translated to protein
 - b) **tRNA** = transfer RNA → brings amino acid to ribos.
 - c) **rRNA** = ribosomal RNA → makes up ribo; catalytic
2. Transcription begins when **RNA polymerase** binds to the **PROMOTER** sequence
 - Tells Rpol which strand, and where to start transcribing!!
3. Transcription proceeds in the **5' → 3' direction** (same in ALL nucleic acid synthesis!)
 - new nucleotides added to the 3' hydroxyl group on **ribose** in the growing strand
4. Transcription stops when it reaches the **Terminator Sequence** (often many U's or “hairpin loop”)
 - New RNA and Rpol fall off of DNA template.

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8.3) Translation

1. = **RNA-directed Protein synthesis!!!**
2. mRNA is translated in **Codons.**
 - 3 nucleotide "words"
3. Translation of mRNA begins at the **start codon: AUG.**
 - Tells ribosome where to start translating,
 - and sets the reading Frame!!!
4. Translation ends at a **STOP ("nonsense") codon: UAA, UAG, UGA.**
 - Do NOT encode an amino acid!!!

Transcription ↓
 Translation ↓

Cell metabolizes and grows

Figure 8.2

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