

Lecture # 9

Exam next week!

**Chapters 10, 11 & 12 - Cell division,
Genetics & DNA Replication**

This week:

**Why we believe in DNA,
DNA Replication, PCR & DNA
sequencing**



3232

DNA
TESTING LAB INC
PATERNITY TESTING

Who's the Daddy?
1 (800) 8-1-1-11

Who's the Daddy?
1 (800) 8-1-1-11

Who's the Daddy?
1 (800) 8-1-1-11

3232
Join Us
Monday - Friday
11:00 am - 4:00 pm
Saturday - Sunday
10:00 pm - 1:00 pm
Phone Number
501-2766

DNA

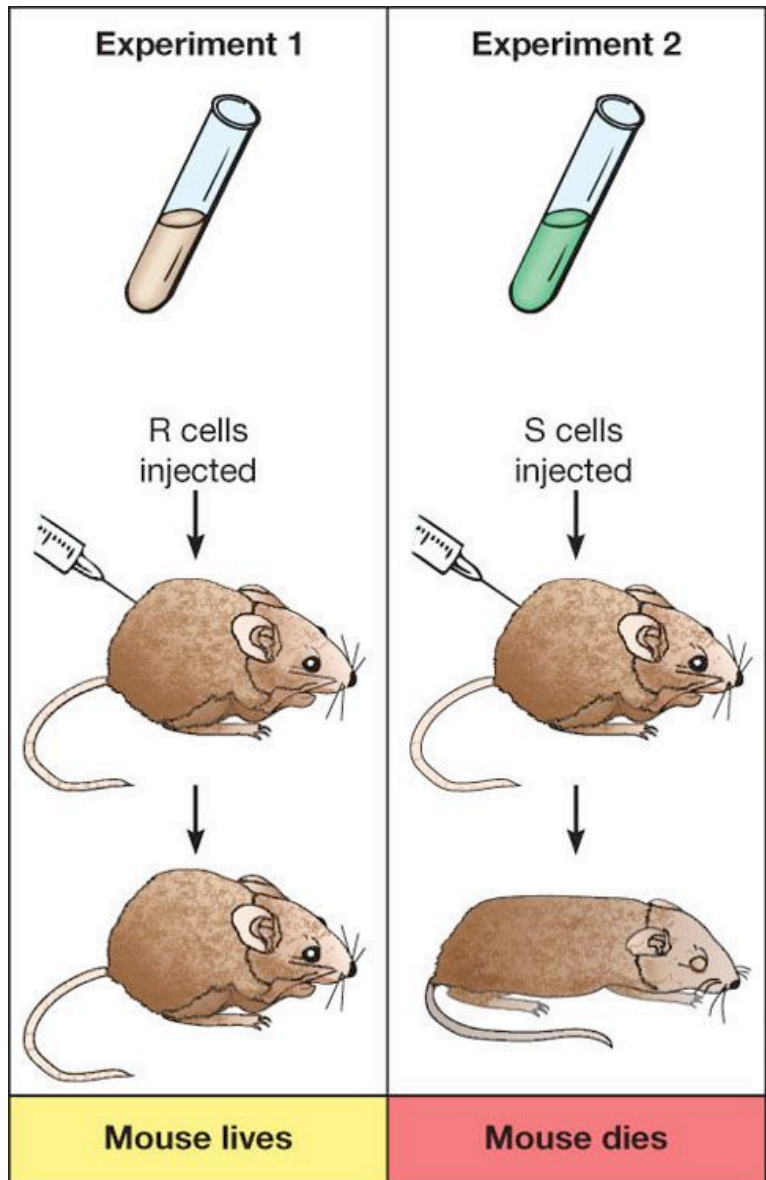
TESTING LAB INC

PATERNITY TESTING

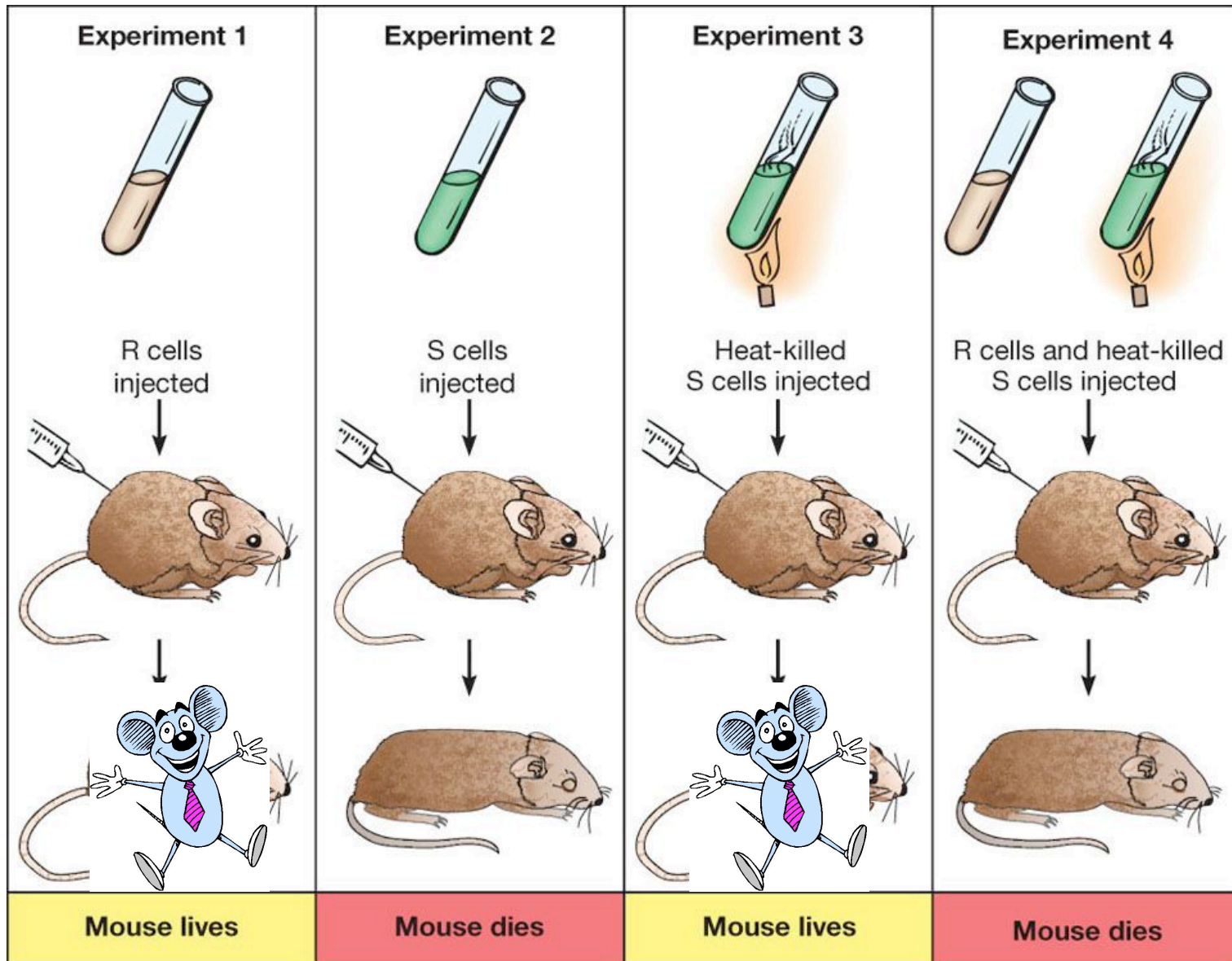
Who's the Daddy?
1 (800) **R-U-MY KID**

Who's the Daddy?
1 (800) **R-U-MY KID**

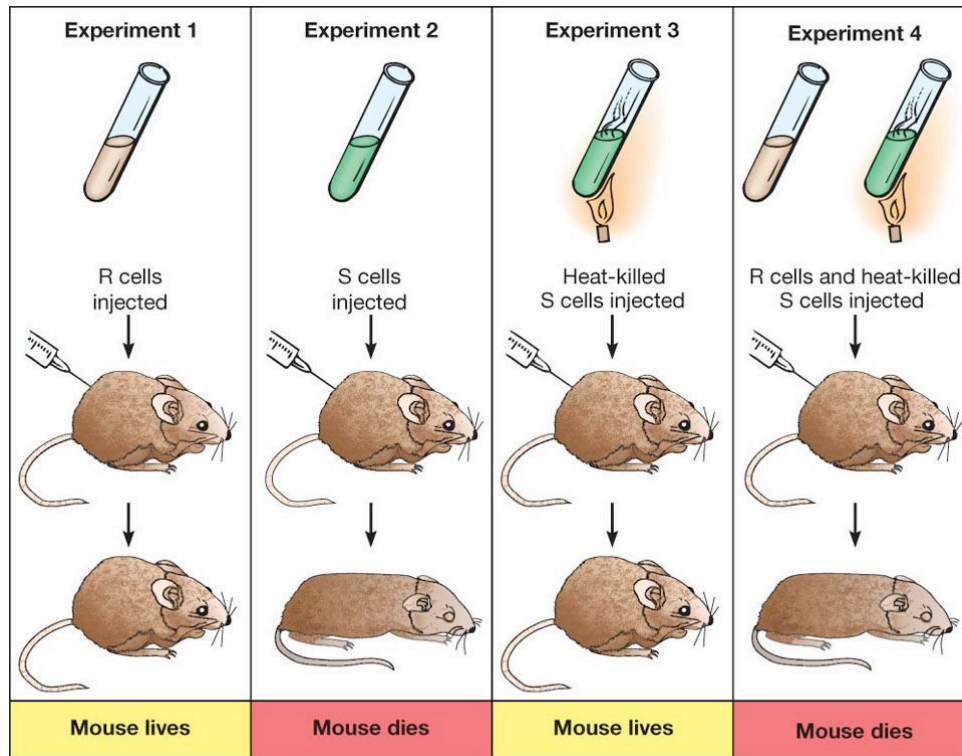
Frederick Griffith (1928) - bacterial transformation



Frederick Griffith - bacterial transformation



Frederick Griffith - bacterial transformation

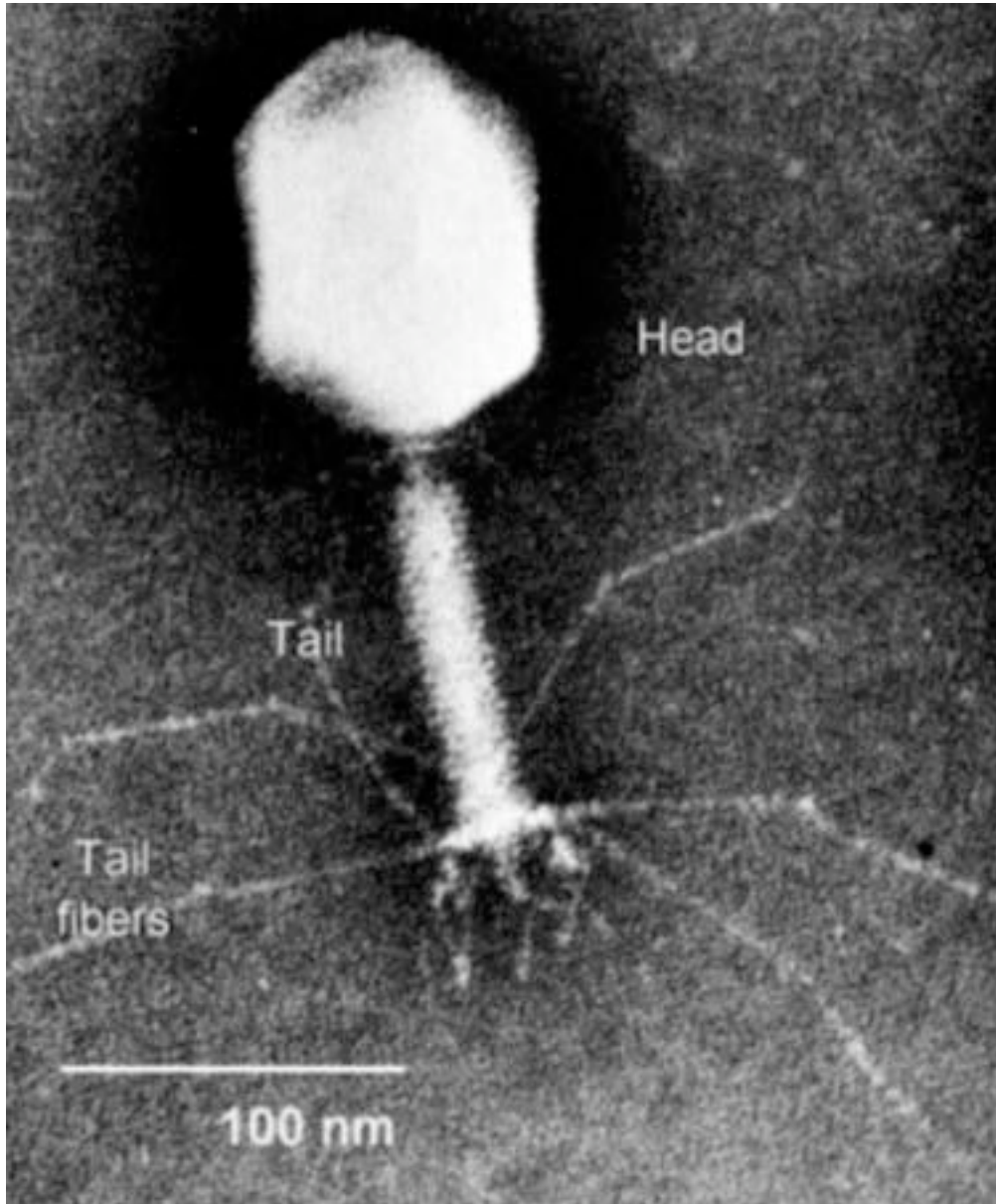


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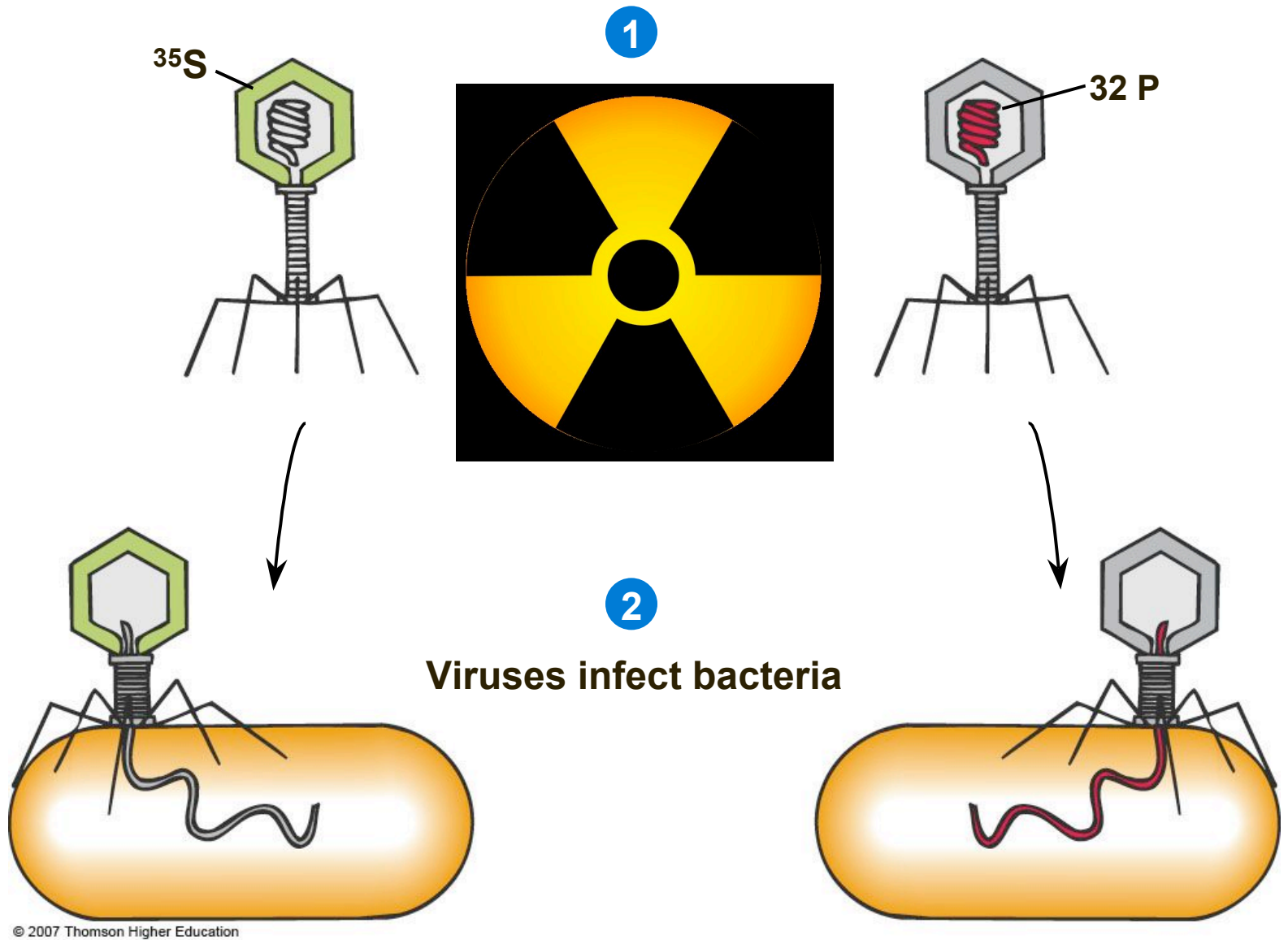
**Conclusion:
Something
travels from
the dead S
cells to the R
cells
(NB, proteins
tend to get
destroyed by
heat)**

Further support by Oswald Avery (1940s)

T4 Bacteriophage

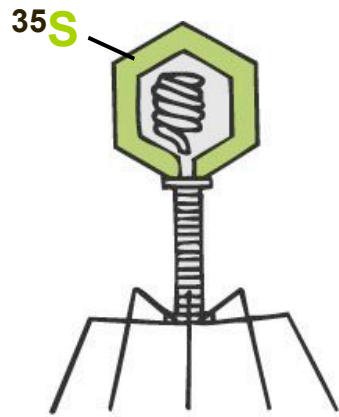


<http://www.bio.davidson.edu/Courses/Molbio/MolStudents/spring2003/Keogh/plasmids.html>

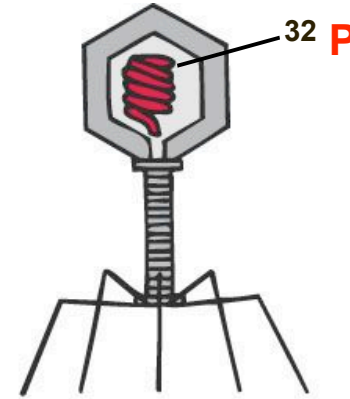


Hershey/Chase - Virus Labeling

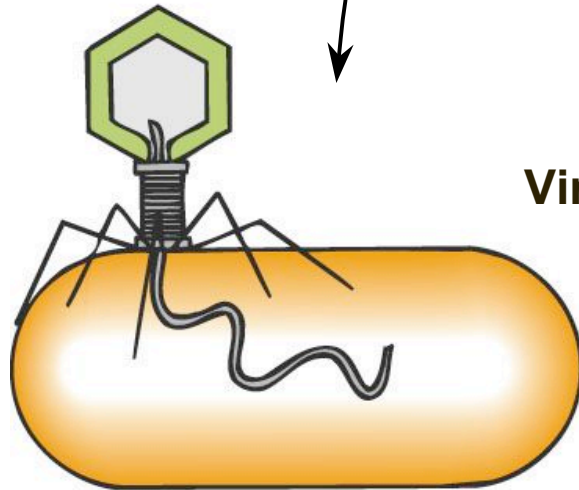
Sulphur is found only in some amino acids (not in DNA)



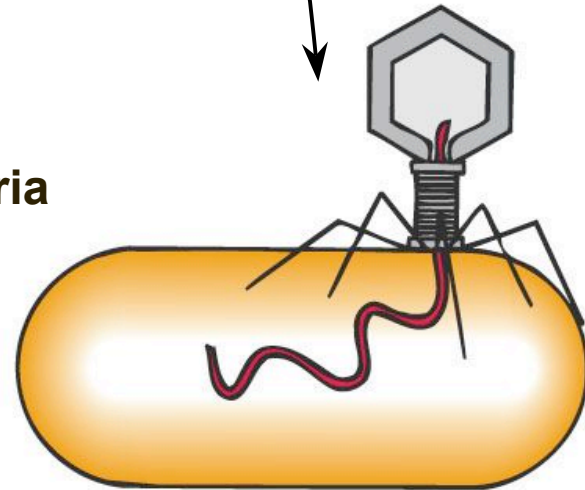
1
Bacterial viruses grown in ^{35}S to label protein coat or ^{32}P to label DNA



Phosphorus is found in all DNA (and not in amino acids)



2
Viruses infect bacteria



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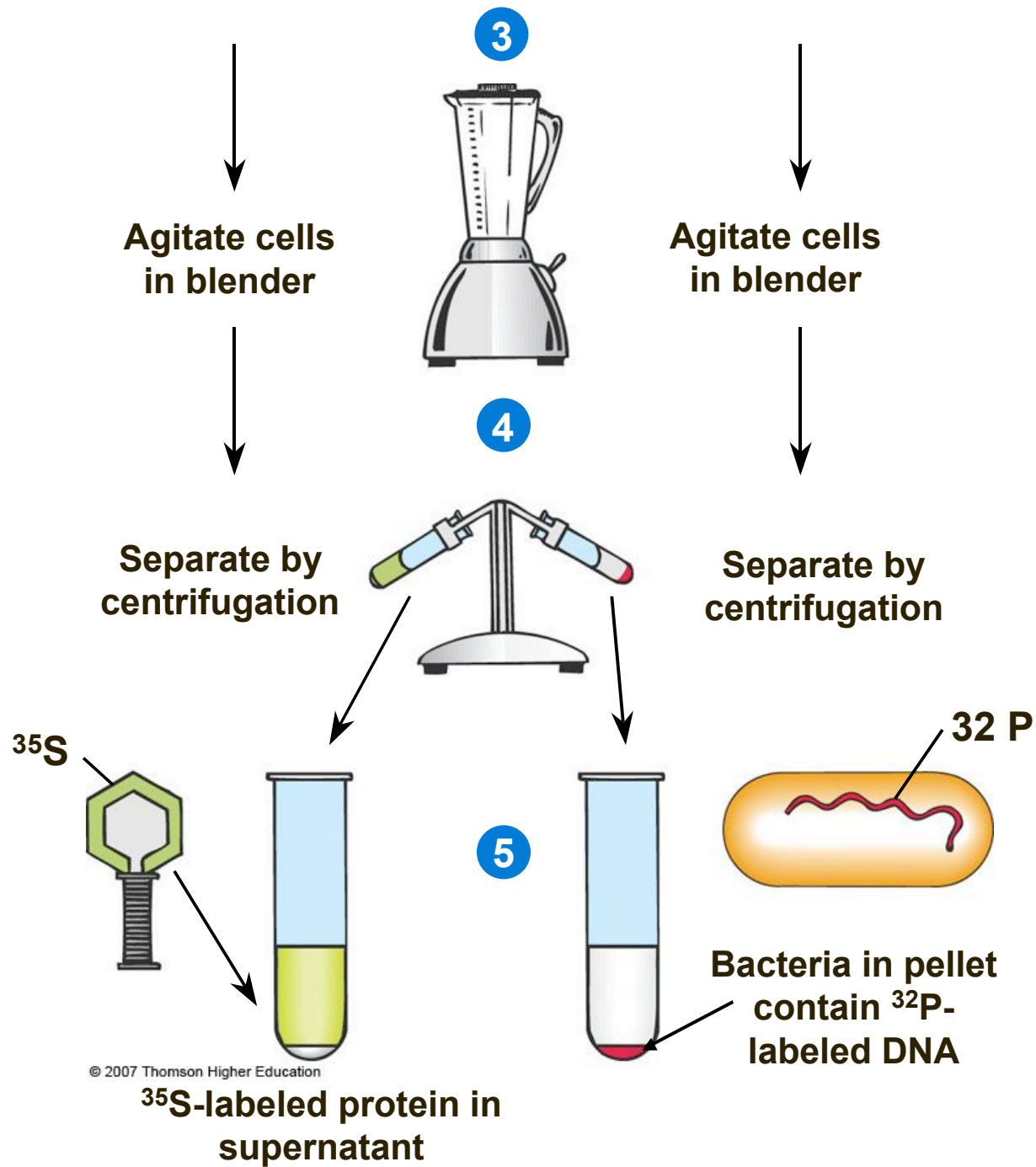
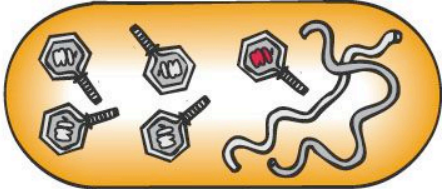
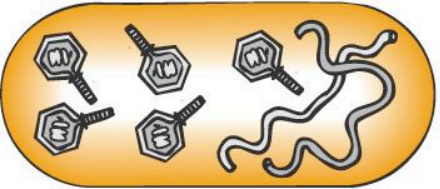


Fig. 12-2, p. 262

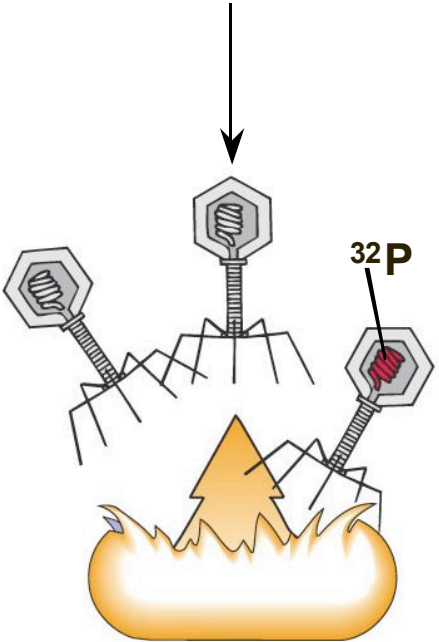
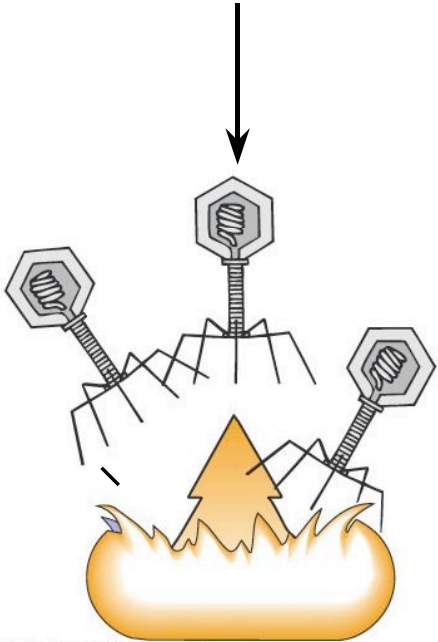
6

Viral reproduction
inside bacterial cells
from pellet



7

Cell lysis



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Fig. 12-2, p. 262

Other clues to DNAs role and function

Chargaff (1949) - Quantitative relationships among DNA bases

The amount of each DNA base (A,T,C or G) varies from organism to organism, BUT

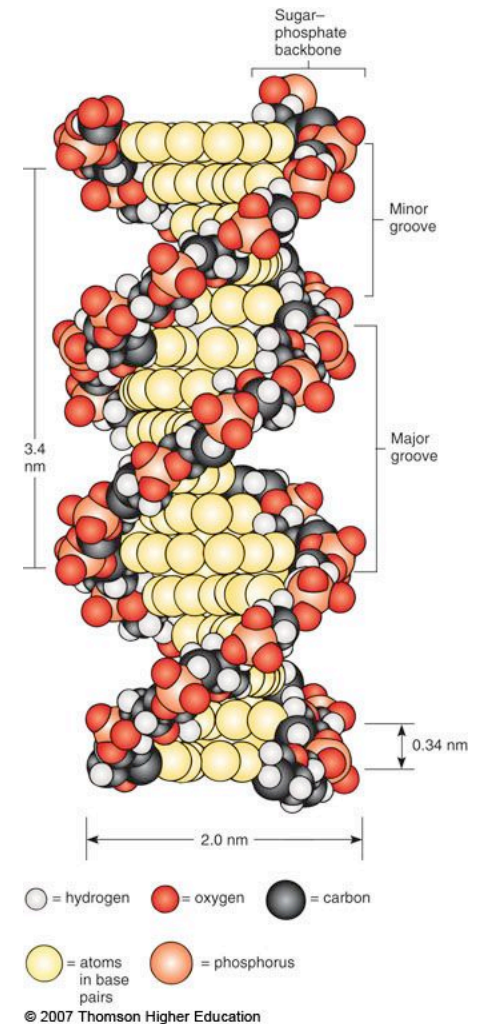
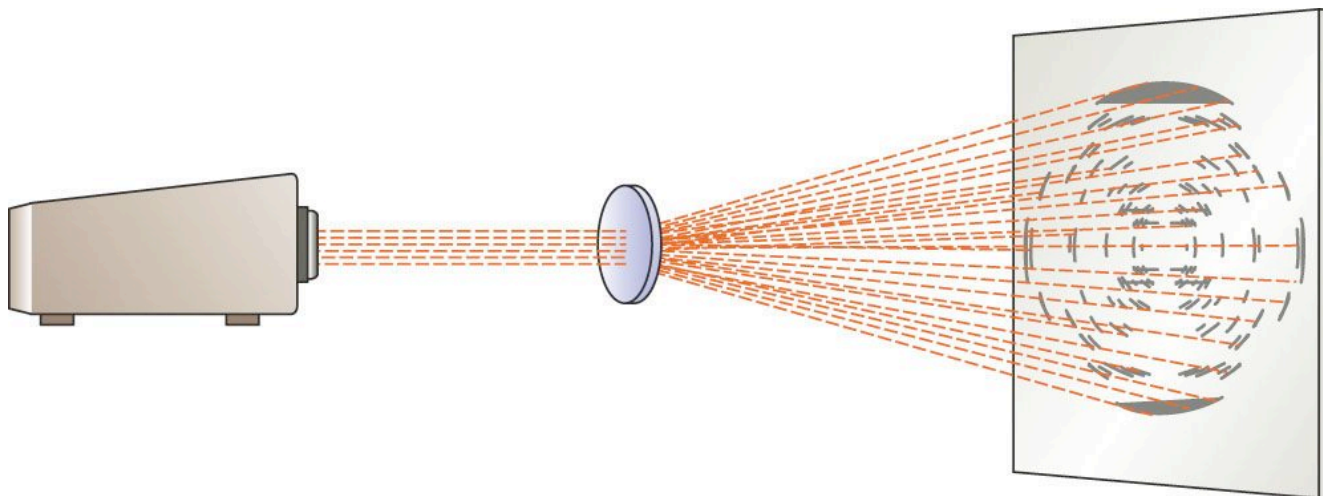
**The amount of A = the amount of T;
The amount of G = the amount of C**

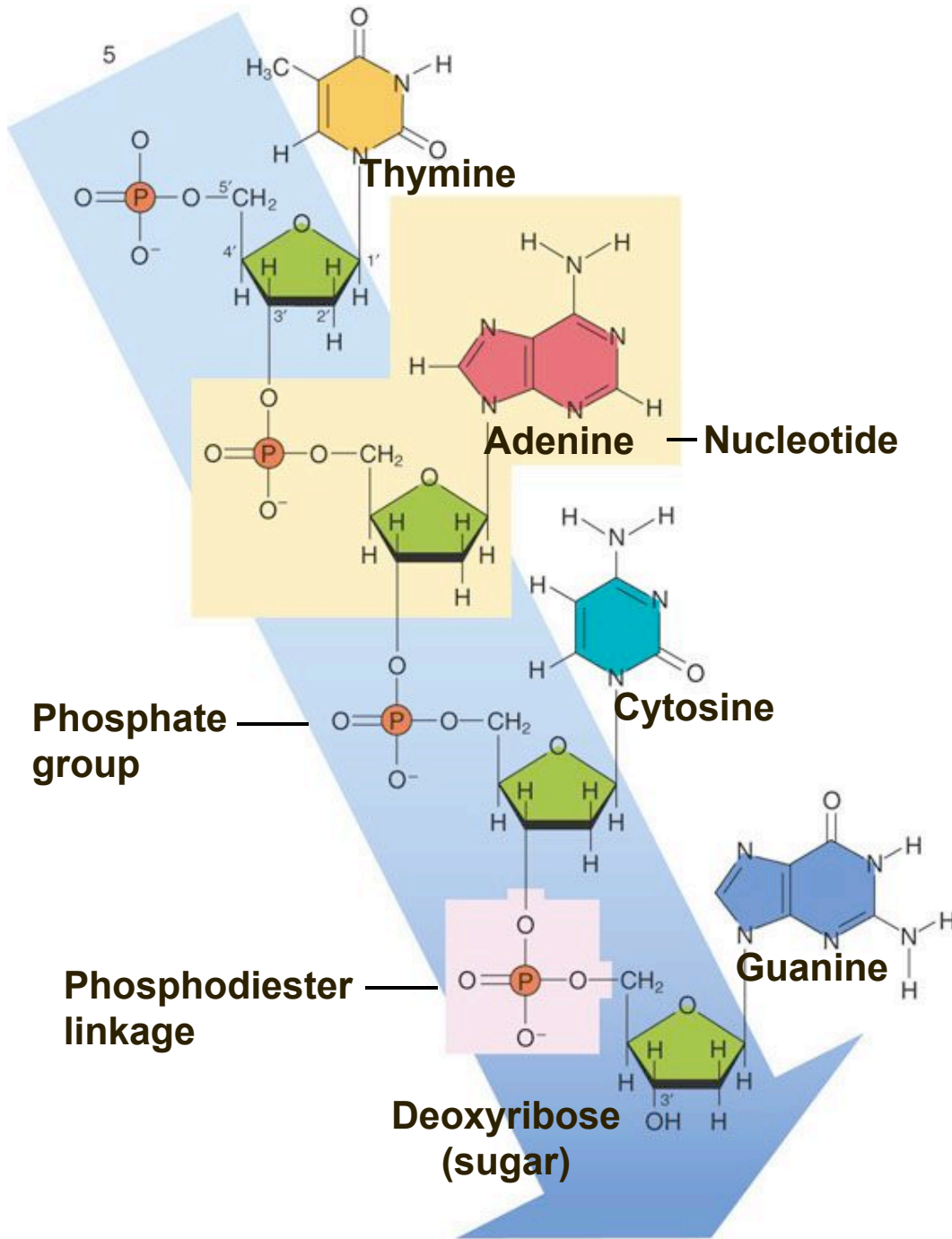
Early 1950s, Franklin, Watson, Crick, Wilkins

X-ray diffraction studies led to the

Double Helix

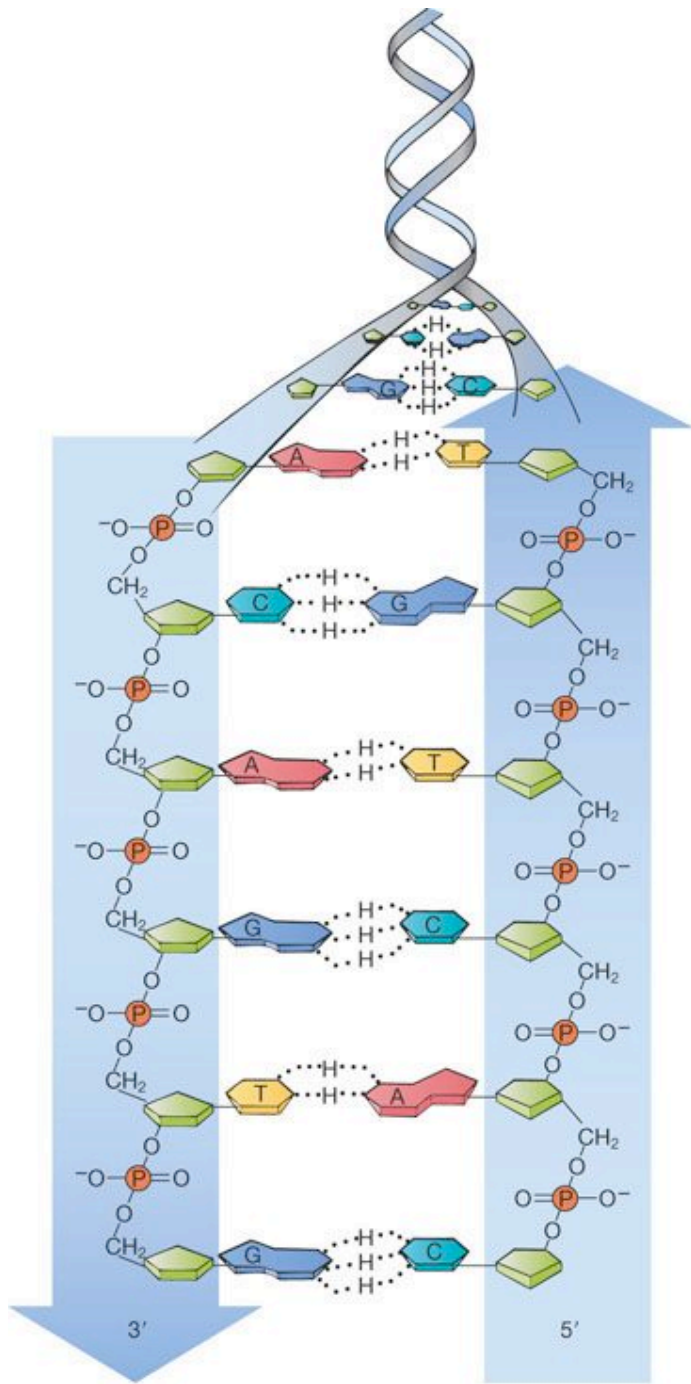
model





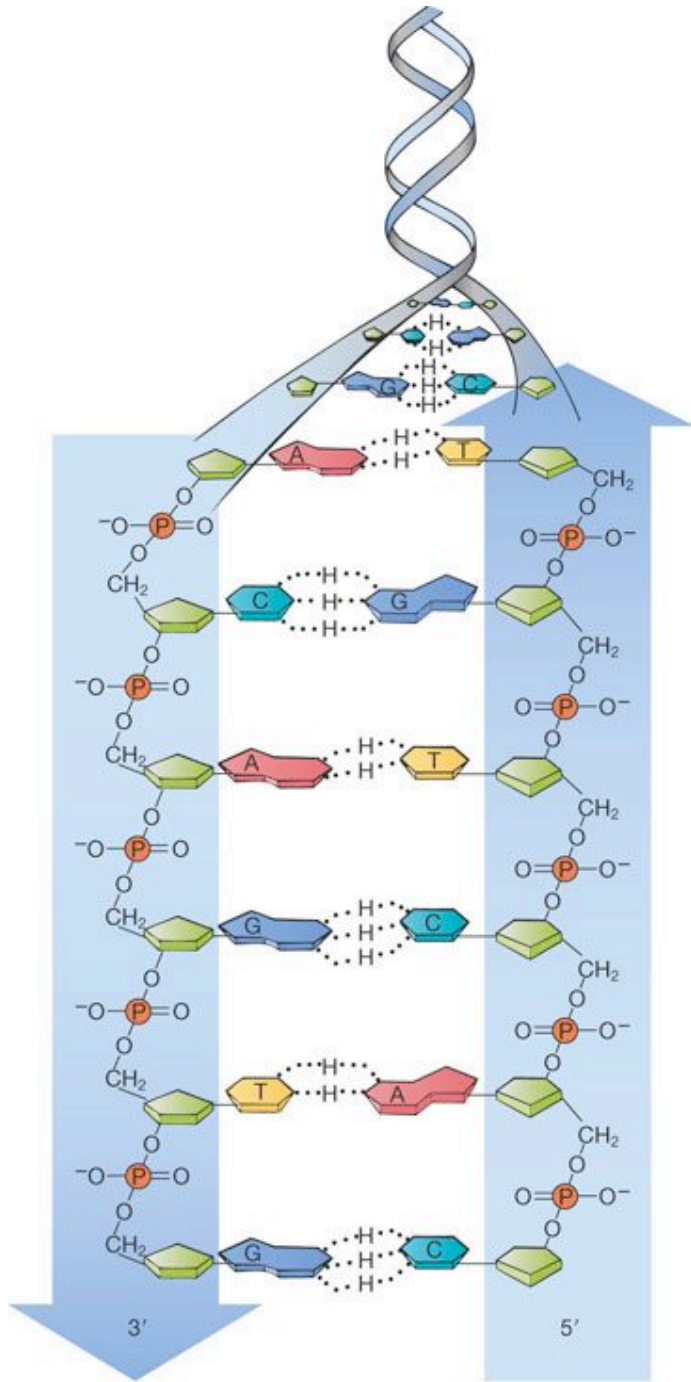
Each strand of the double helix is a polymer, with an orientation

Strands are **antiparallel**



Each strand of the double helix is a polymer, with an orientation

Strands are **antiparallel**



The bases are held to each other with 2 or 3 hydrogen bonds

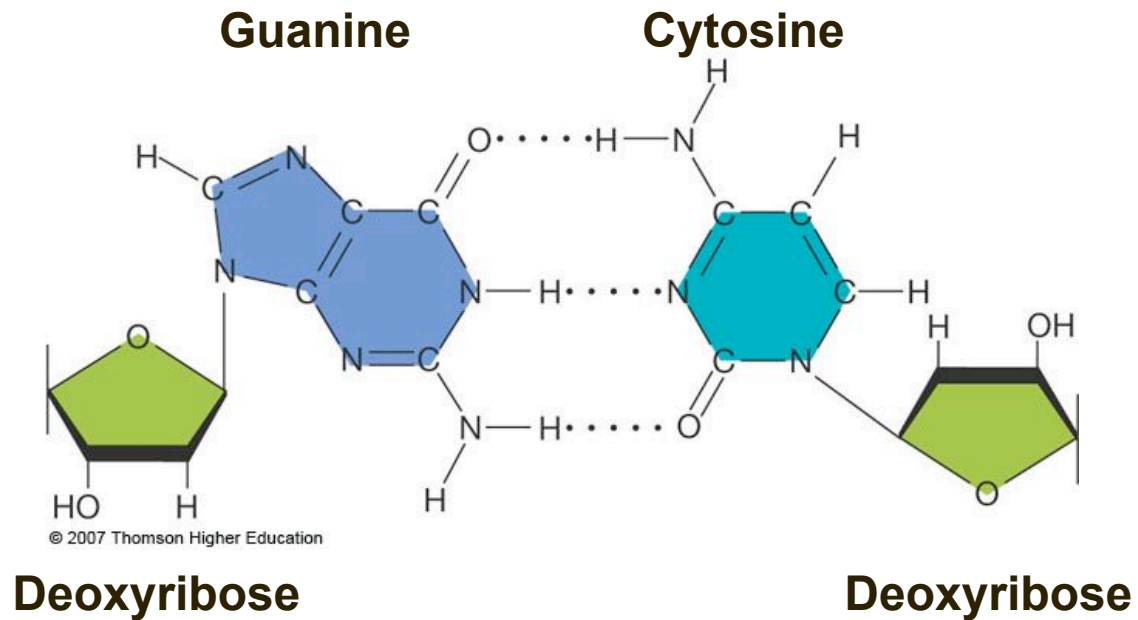
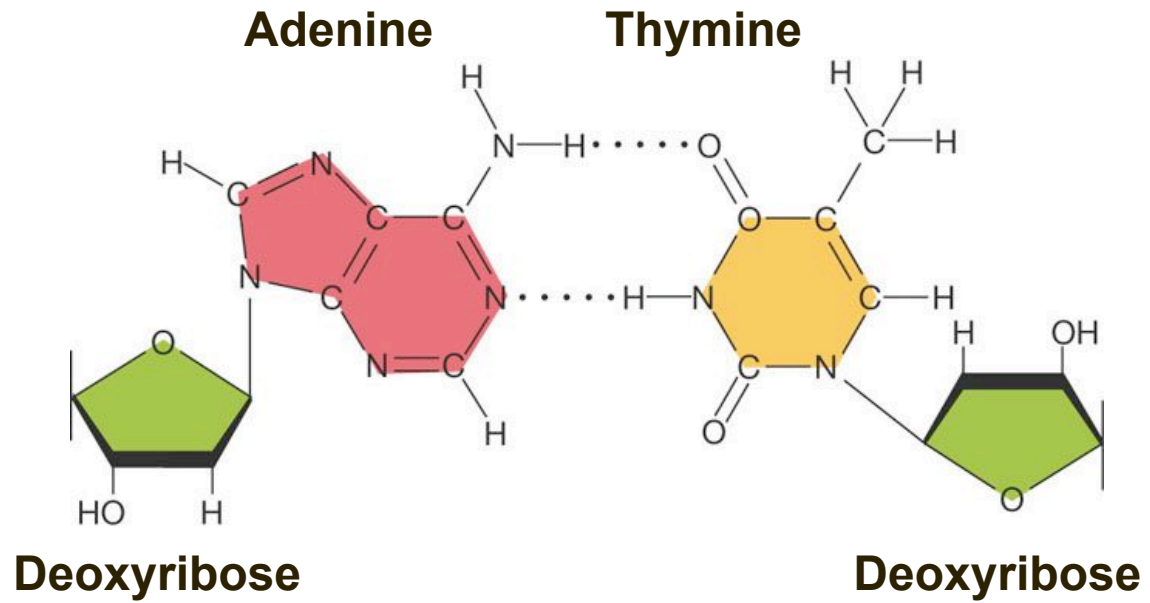
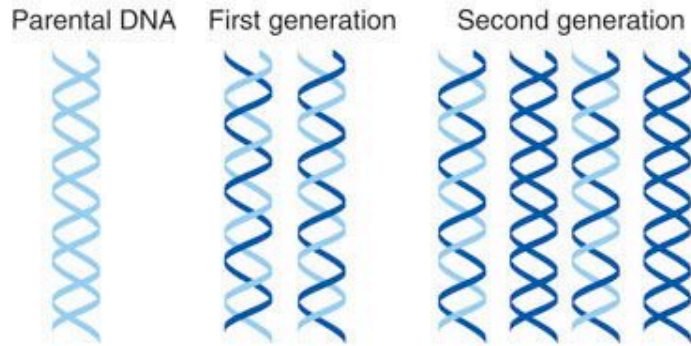


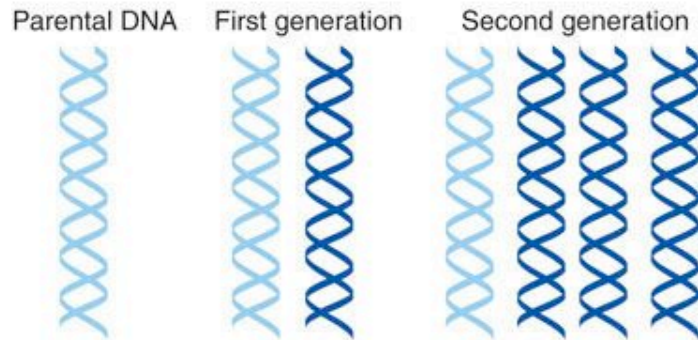
Fig. 12-6b, p. 267

(a) Hypothesis 1: Semiconservative replication

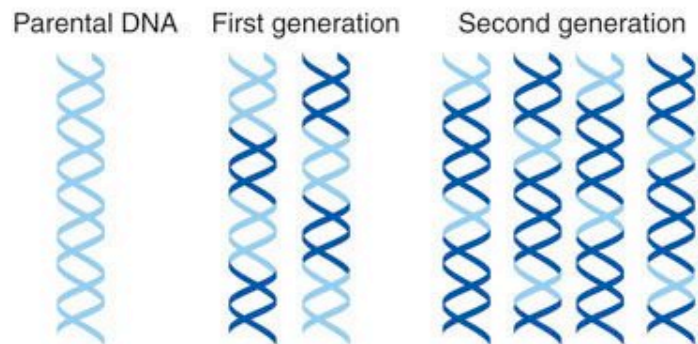


DNA Replication is
semi-conservative

(b) Hypothesis 2: Conservative replication



(c) Hypothesis 3: Dispersive replication



(a) Hypothesis 1: Semiconservative replication

Parental DNA



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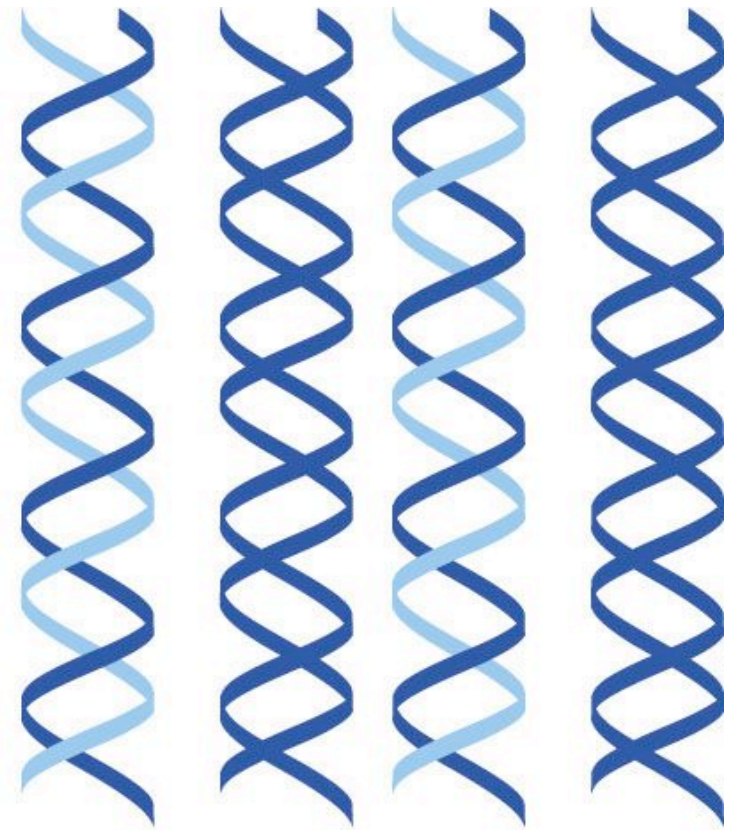
A B

First generation



C A B D

Second generation



A B

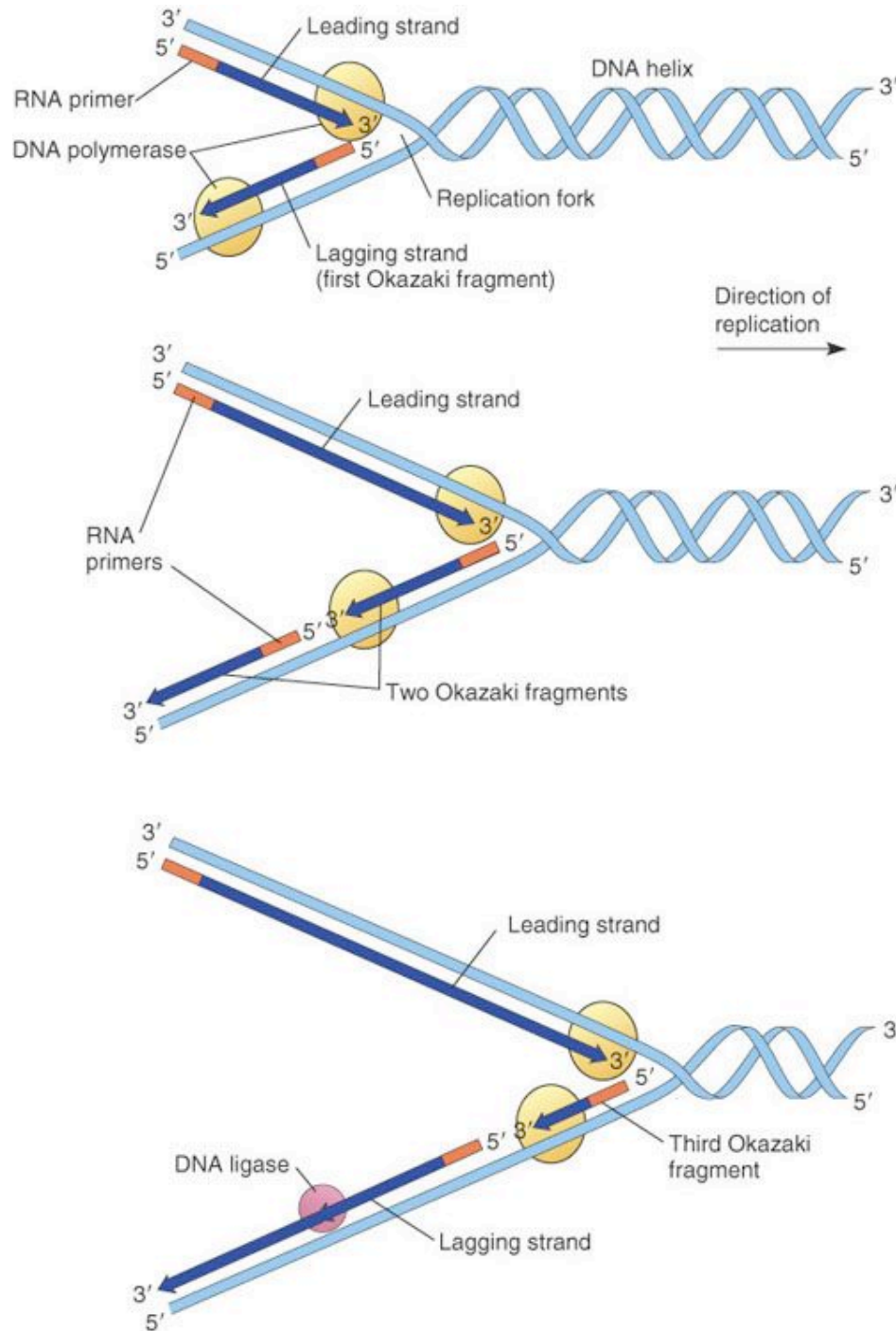
Fig. 12-7a, p. 268

TABLE 12-3**Proteins Involved in DNA Replication**

Enzyme	Function
Helicase	Opens the double helix at replication forks by disrupting the hydrogen bonds that hold the two strands together.
Single-strand binding (SSB) protein	Binds to single strands of DNA and prevents the helix from re-forming before it can be used as a template for replication.
Topoisomerase	Breaks one or both DNA strands, preventing excessive coiling during replication, and then rejoins them in a more relaxed configuration.
DNA polymerase	Links nucleotide subunits to form a new DNA strand from a DNA template.
DNA primase	Synthesizes short RNA primers on the lagging strand. Begins replication of the leading strand.
DNA ligase	Links Okazaki fragments by joining the 3' end of the new DNA fragment to the 5' end of the adjoining DNA.
Telomerase	Lengthens telomeric DNA by adding repetitive nucleotide sequences to the ends of eukaryotic chromosomes.

TABLE 12-3**Proteins Involved in DNA Replication**

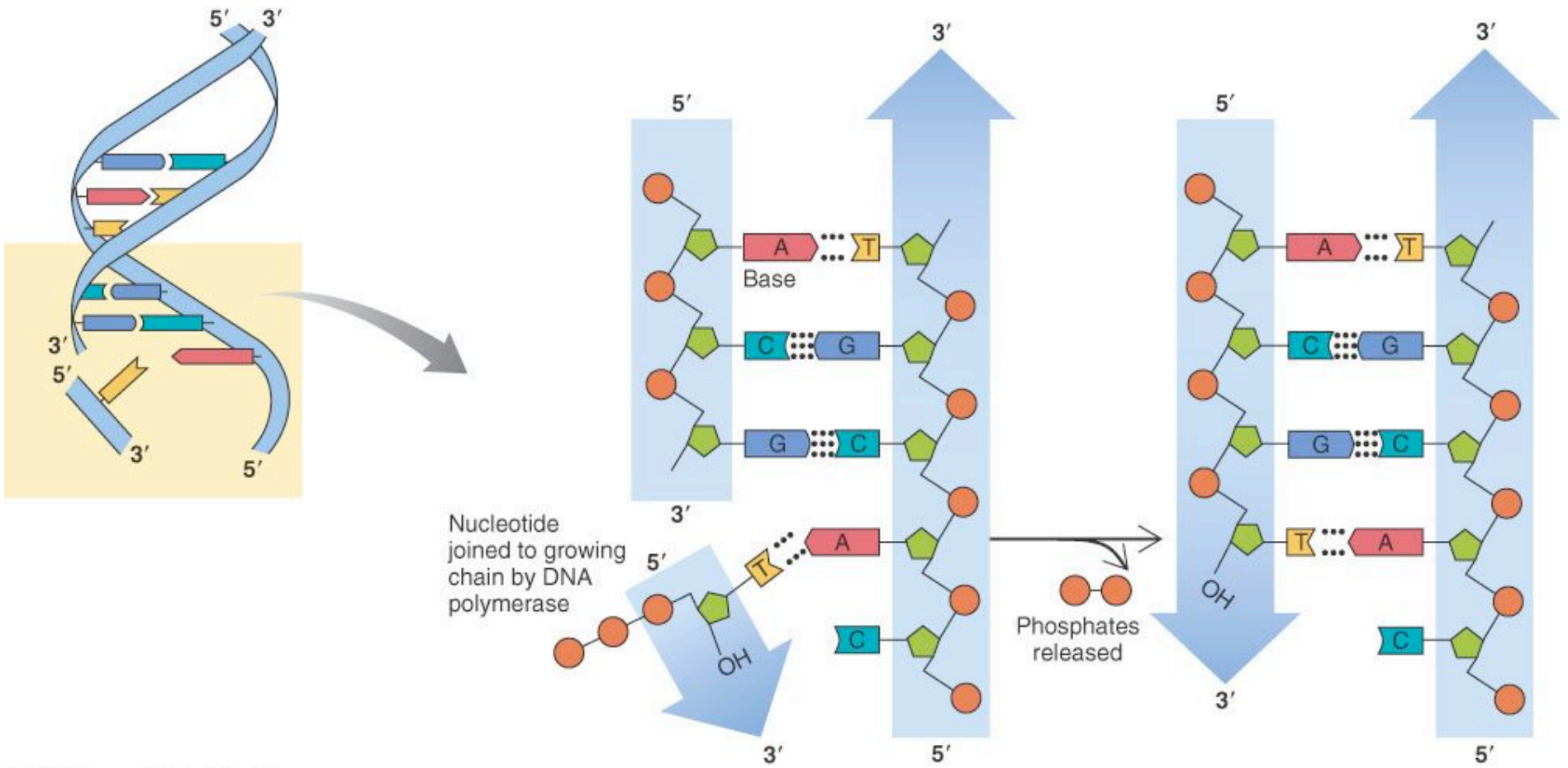
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1 Leading strand is synthesized continuously in direction toward replication fork; lagging strand is synthesized in direction away from replication fork. Both strands require RNA primer for initiation of synthesis because DNA can be elongated only by addition to 3' end of existing polynucleotide strand.

2 Lagging strand is synthesized as short Okazaki fragments. Okazaki fragment synthesis begins with the synthesis of RNA primer. Note that first Okazaki fragment synthesized is now at far left.

3 After each Okazaki fragment has been elongated by DNA polymerase, RNA primer is degraded, gaps are filled in with DNA, and adjoining fragments are linked by DNA ligase.



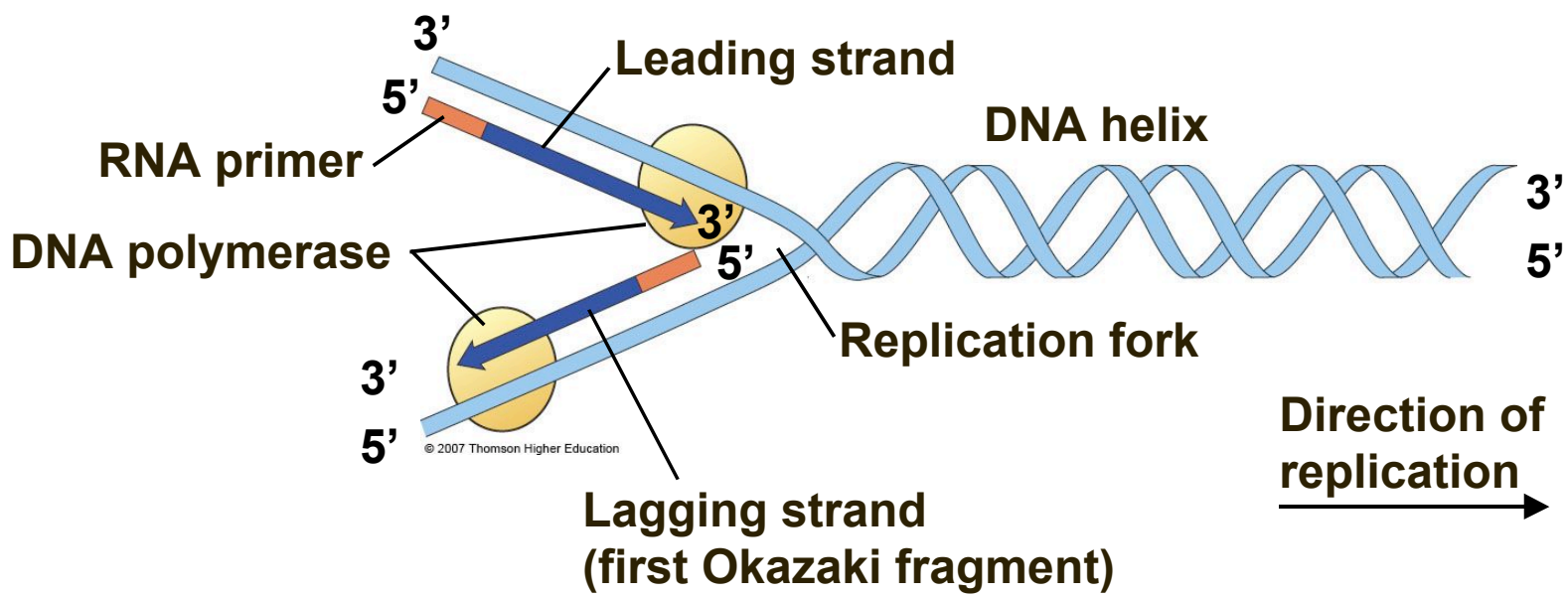


Fig. 12-12a, p. 273

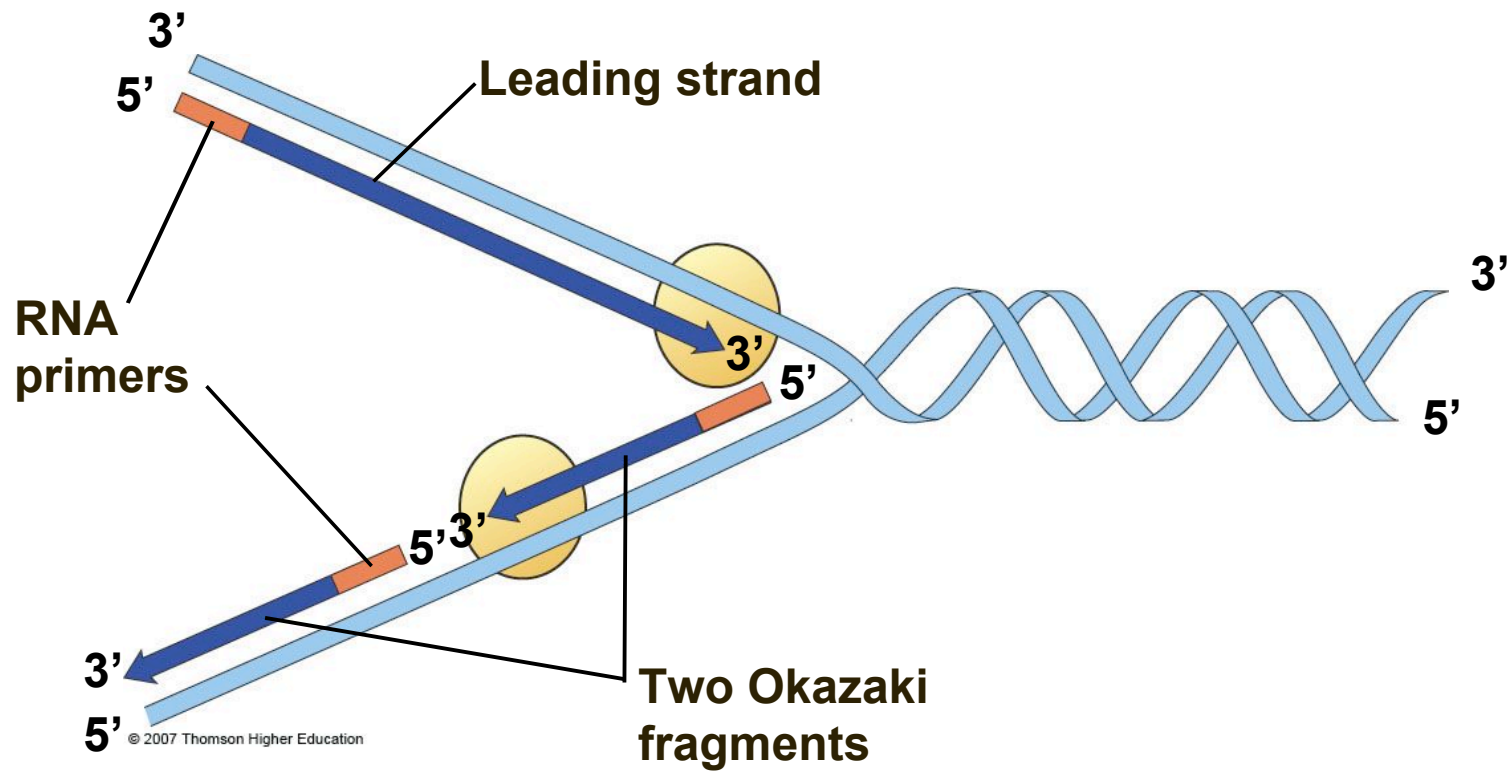


Fig. 12-12b, p. 273

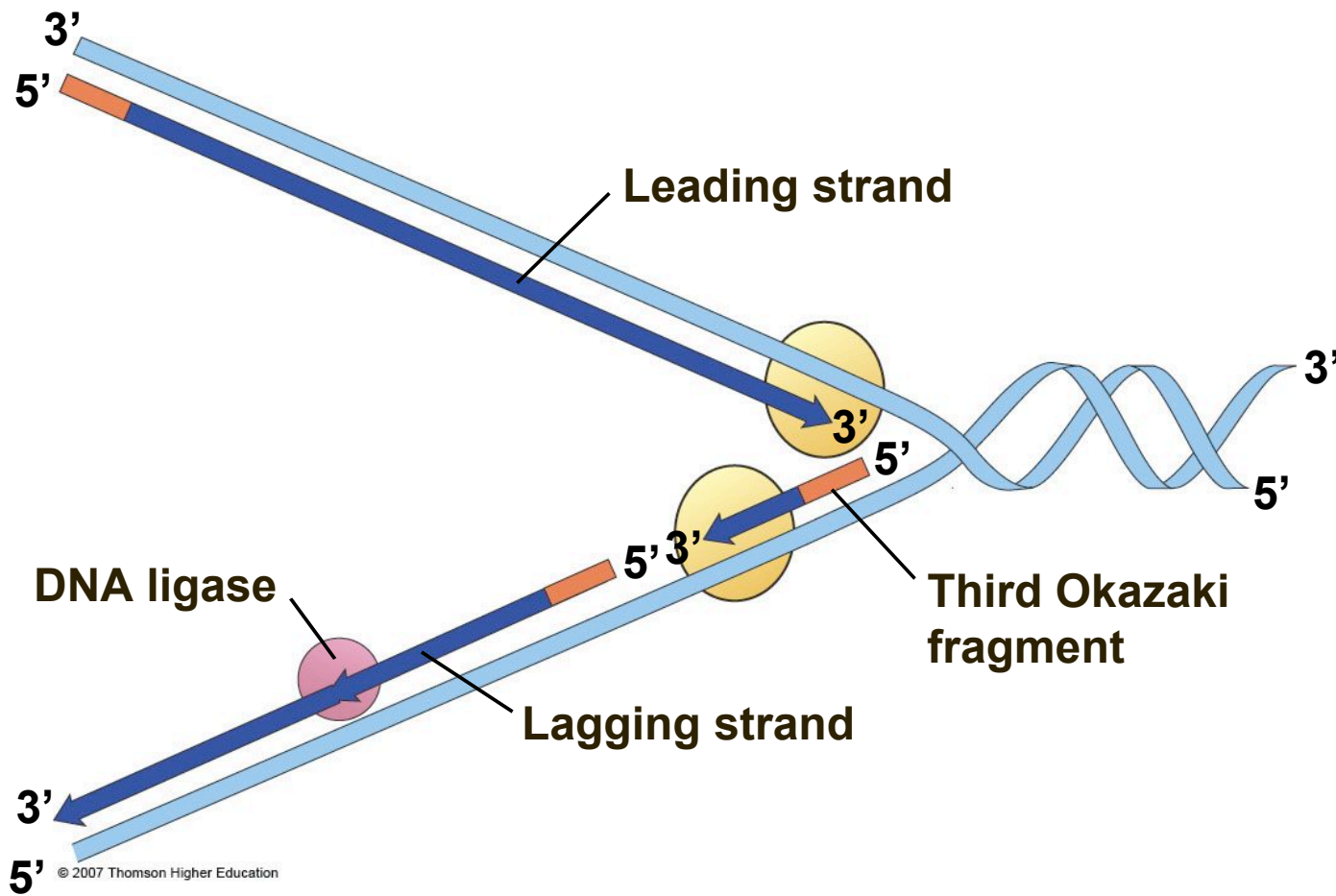


Fig. 12-12c, p. 273

Replication Rules

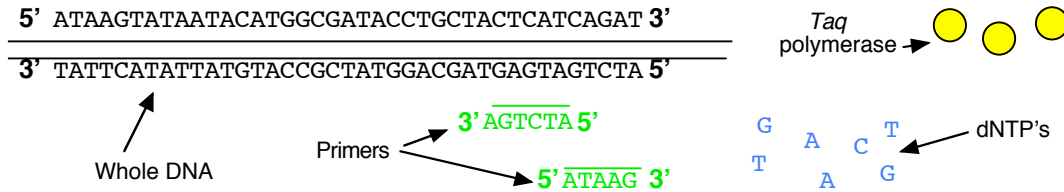
New bases can only be added on to the **3'** end of the growing strand

A DNA or RNA primer is necessary

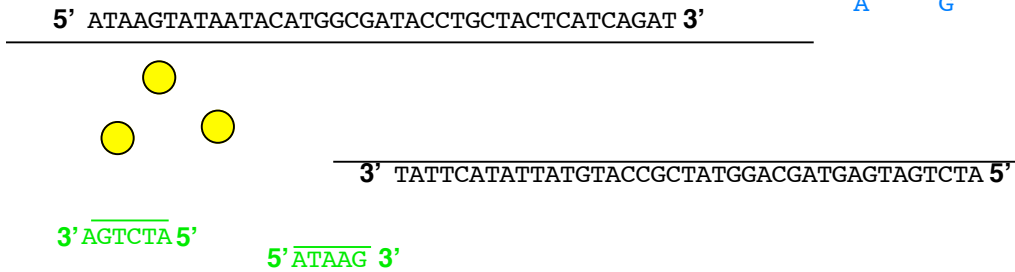
G always pairs with C, T with A

Strands are always antiparallel

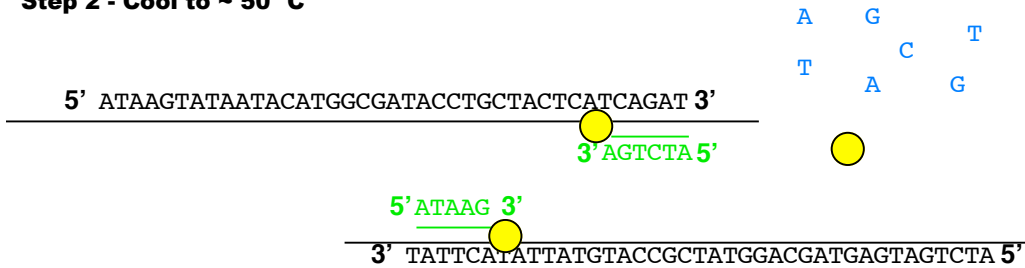
Replication is Semi-conservative



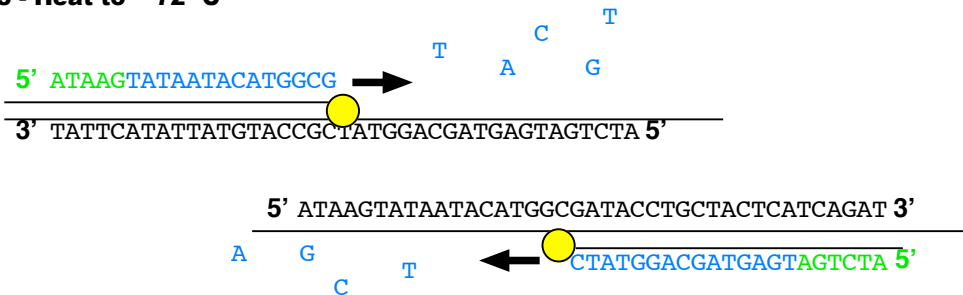
Step 1 - Heat to ~ 94° C



Step 2 - Cool to ~ 50° C



Step 3 - Heat to ~ 72° C



PCR - uses a polymerase from a high temp. bacteria to make copies of DNA, *in vitro*

DNA Sequencing...

AATAAGACCG
TTATTCTGGCTAGTAGTTAGTTAGGCCCTGACCTTCAA

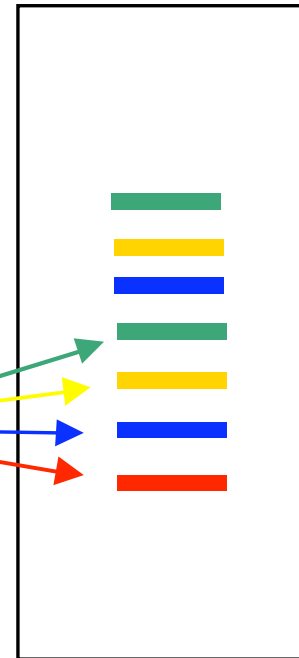
G = a **dideoxy** nucleotide, the chain cannot be extended

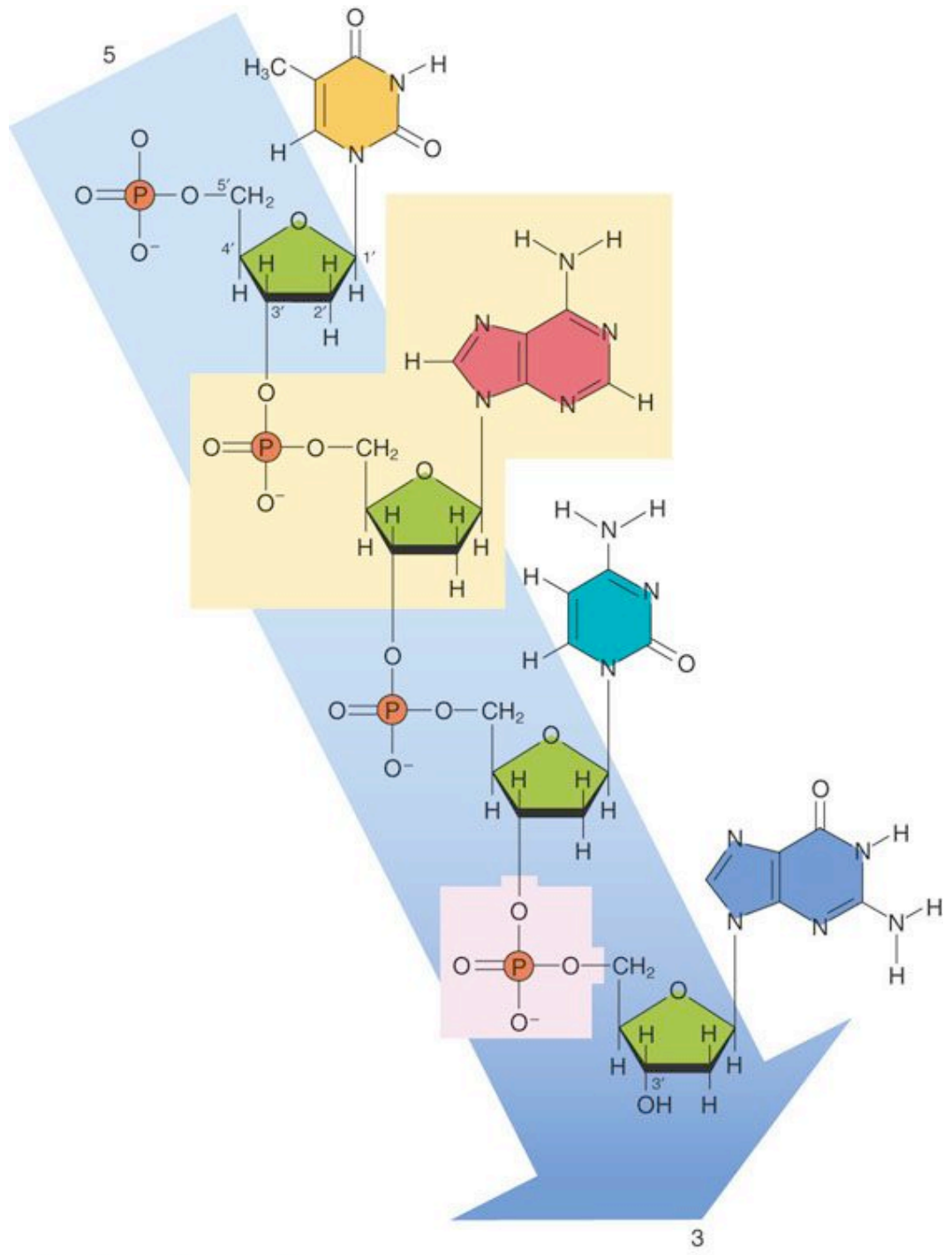
AATAAGACCG

AATAAGACCGA

AATAAGACCGAT

AATAAGACCGATC





DNA Sequencing...

AATAAGACCG
TTATTCTGGCTAGTAGTTAGTTAGGCCCTGACCTTCAA

G = a **dideoxy** nucleotide, the chain cannot be extended

AATAAGACCG

AATAAGACCGA

AATAAGACCGAT

AATAAGACCGATC

