

# **Chapter 16:** **DNA Structure & Replication**

**1. DNA Structure**

**2. DNA Replication**

# **1. DNA Structure**

**Chapter Reading – pp. 313-318**

# **Genetic Material: Protein or DNA?**

**Until the early 1950's no one knew for sure, but it was generally thought that protein was the genetic material. Why?**

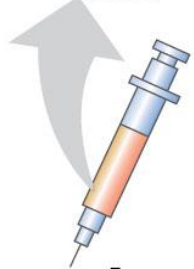
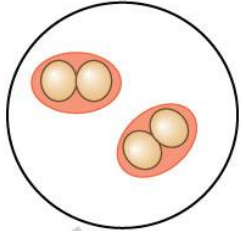
- protein is made of 20 different amino acids**
- DNA is made of only 4 different nucleotides**
- protein could theoretically store more info**
  - a “20 letter alphabet” vs a “4 letter alphabet”**
  - it was assumed that life was so complex, therefore a “bigger alphabet” was necessary to somehow encode it!**

**Some classic experiments would prove otherwise...**

# Transformation of Bacteria

## EXPERIMENT

Living S cells  
(control)

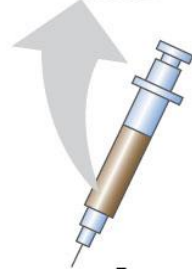
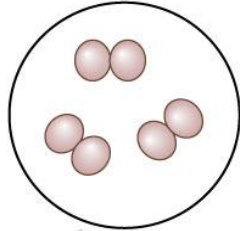


RESULTS

Mouse dies



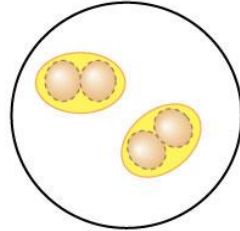
Living R cells  
(control)



Mouse healthy



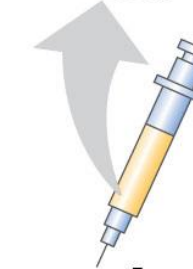
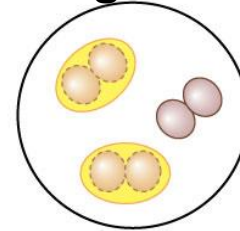
Heat-killed  
S cells  
(control)



Mouse healthy



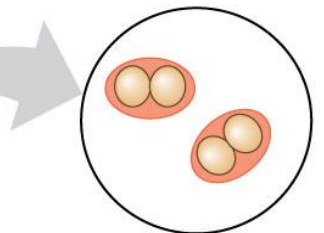
Mixture of  
heat-killed  
S cells and  
living R cells



Mouse dies



Demonstrated  
the transfer of  
a genetic trait  
between  
different  
bacteria.  
The nature of  
that genetic  
material was  
still unknown.



Living S cells

(Frederick Griffith, 1928)

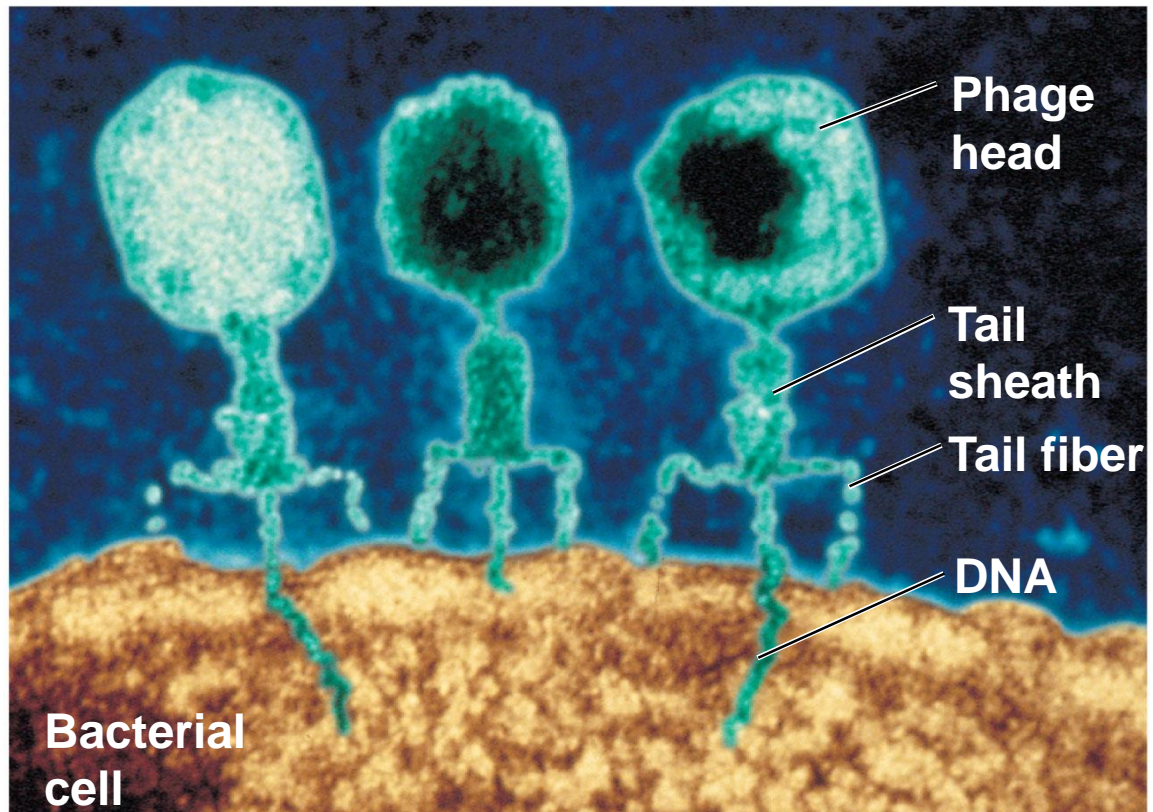
# What is the Genetic Material of Bacteriophages?

**Bacteriophages are viruses that infect bacteria.**

- consist of a protein capsid which contains DNA

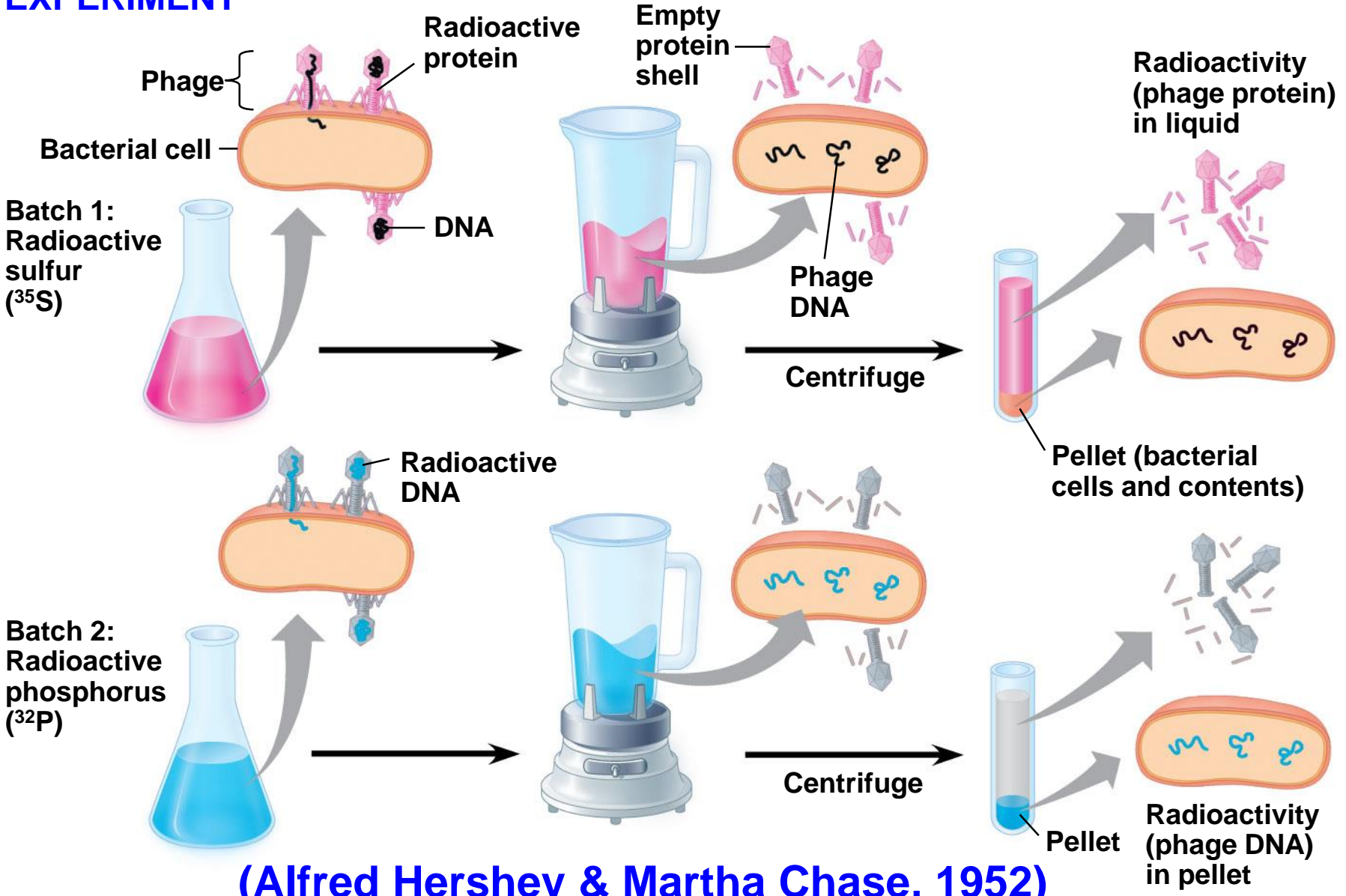
**What enters the bacterial host cell, viral protein, DNA, or both?**

- whatever enters the host cell should be the genetic material



# Bacteriophage Genetic Material is DNA

## EXPERIMENT

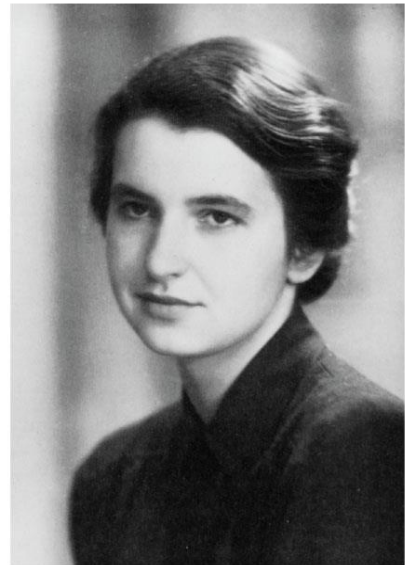


(Alfred Hershey & Martha Chase, 1952)

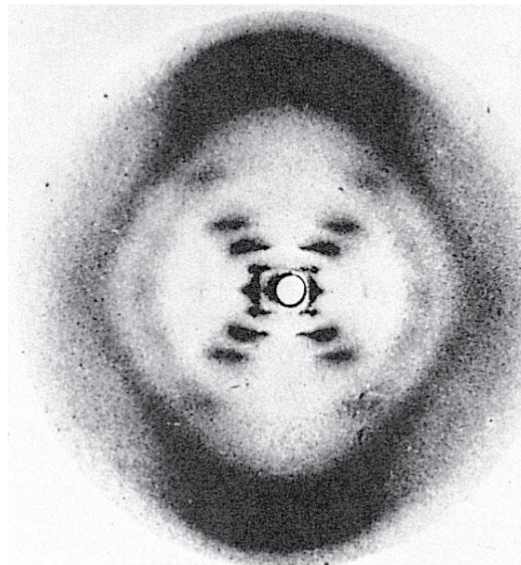
# The Discovery of DNA Structure

Using the technique of x-ray crystallography, Rosalind Franklin, James Watson & Francis Crick figured out the structure of DNA

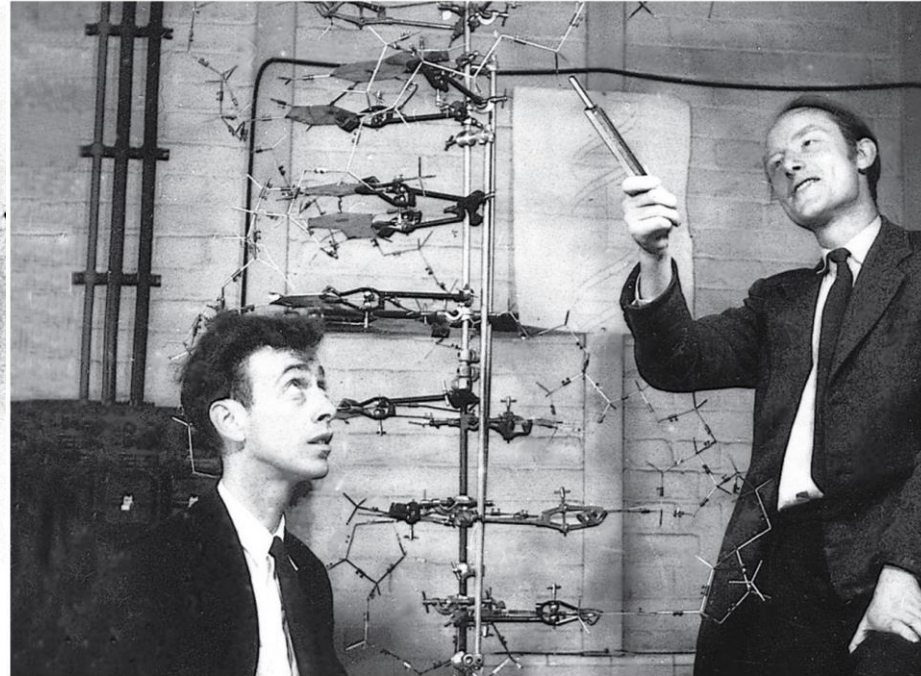
- Watson & Crick used the X-ray diffraction data of Rosalind Franklin to deduce the structure of DNA



(a) Rosalind Franklin



(b) Franklin's X-ray diffraction photograph of DNA

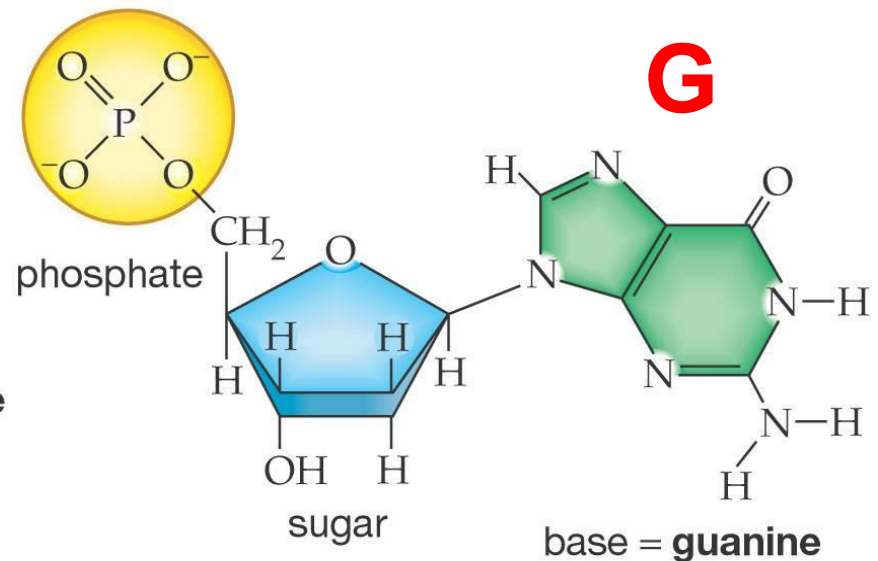
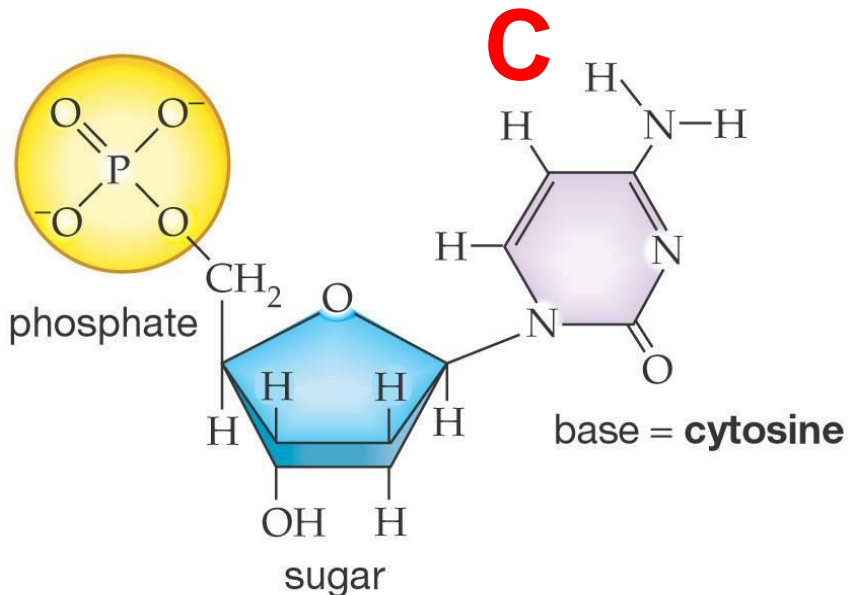
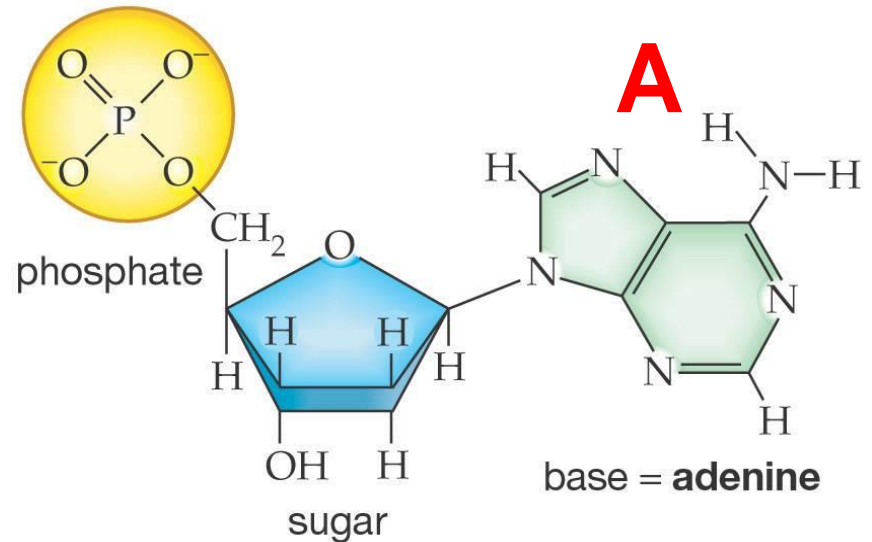
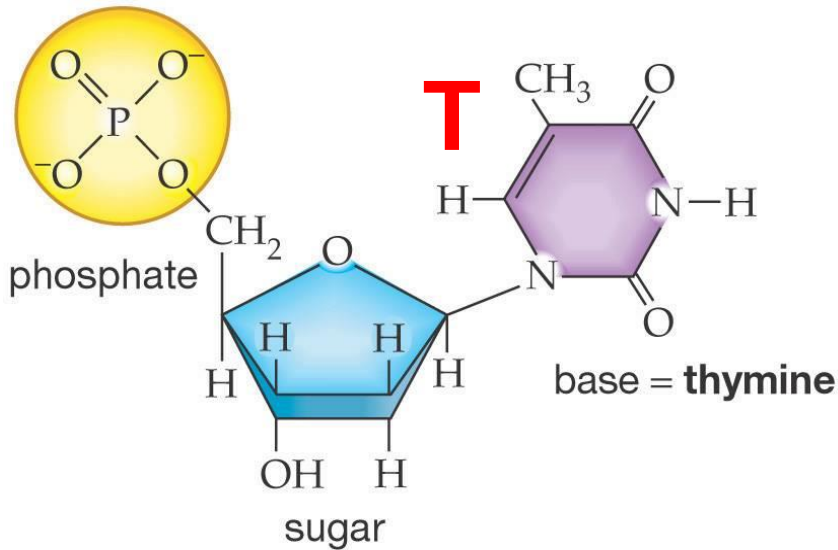


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# DNA: a Polymer of 4 Nucleotides

pyrimidines

purines





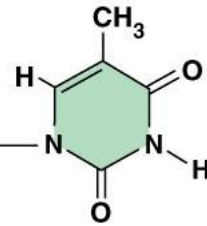
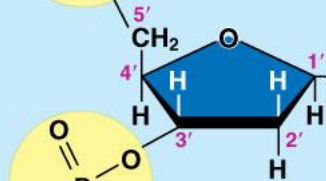
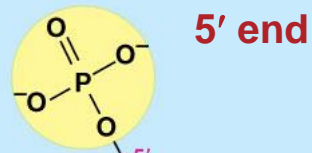
## Sugar-phosphate backbone

## Nitrogenous bases

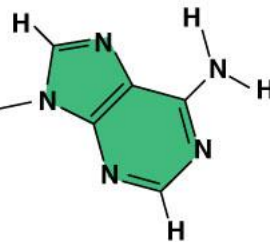
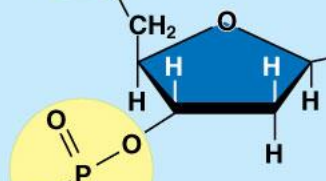
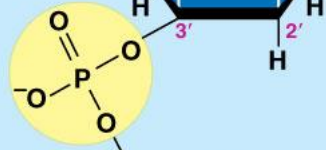
# Structure of a DNA Strand

A single DNA polymer or “strand” consists of a sugar-phosphate backbone with the bases project out.

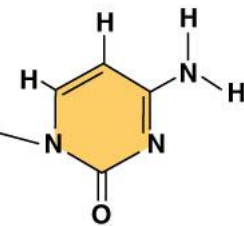
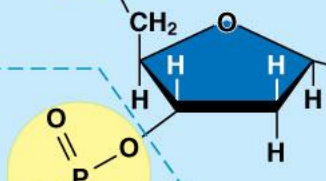
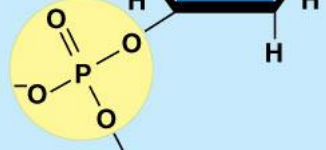
The ends of a DNA strand are different, with one end having a free 5' phosphate, and the other having a free 3' hydroxyl group



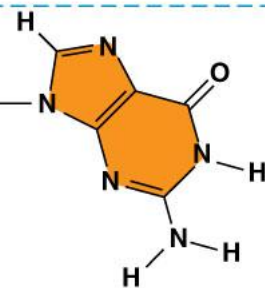
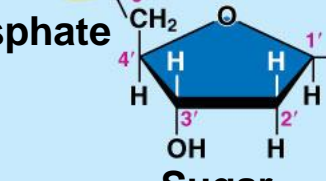
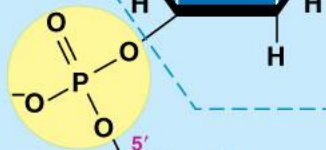
Thymine (T)



Adenine (A)



Cytosine (C)



Guanine (G)

Phosphate

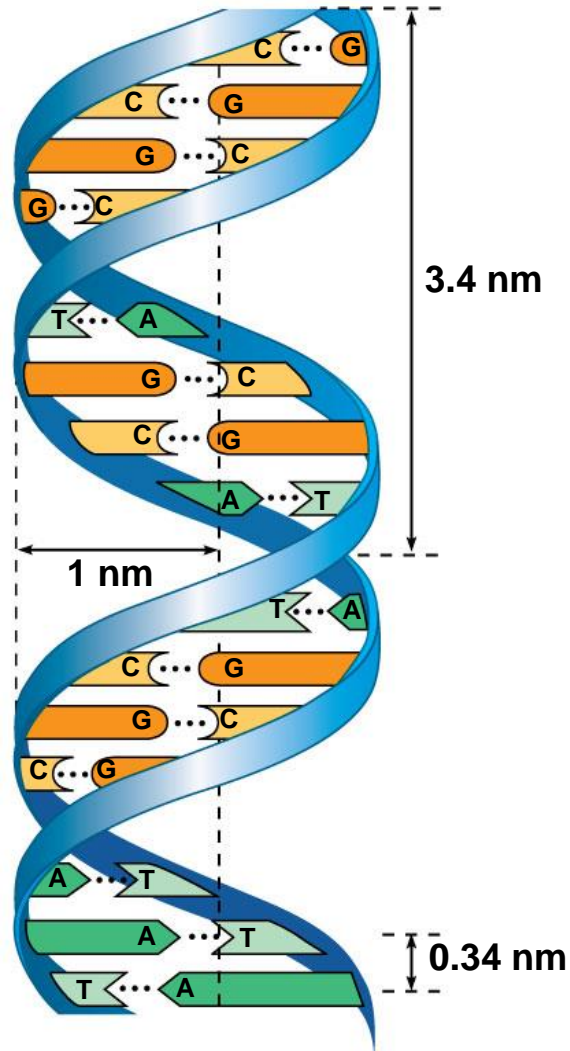
Sugar  
(deoxyribose)

Nitrogenous base

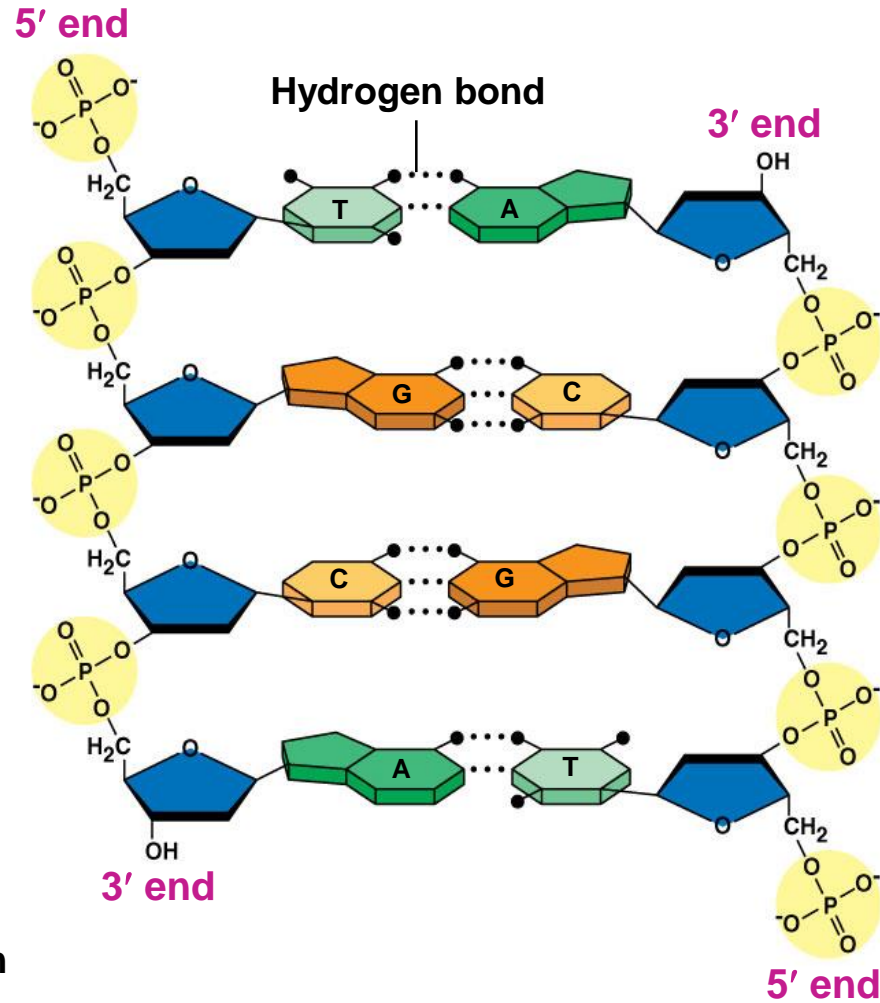
DNA nucleotide 3' end

# Structure of Double-stranded DNA

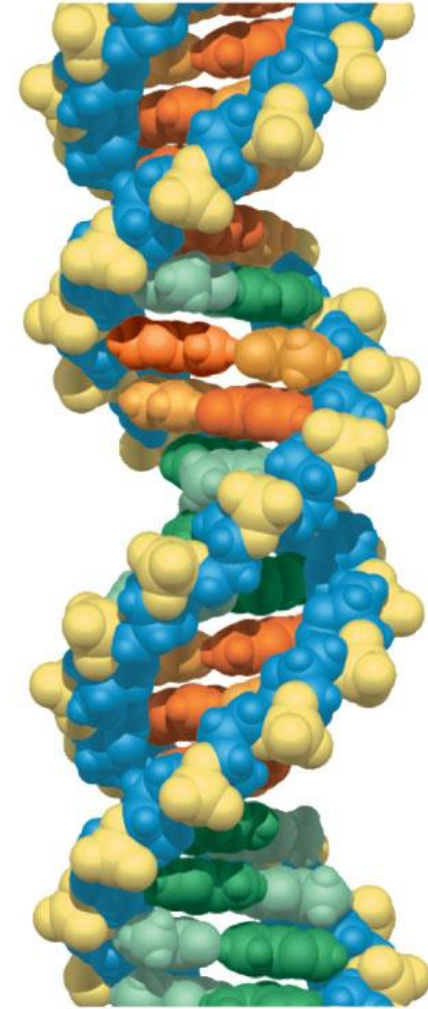
- the 2 strands are anti-parallel and interact via base pairs



(a) Key features of DNA structure



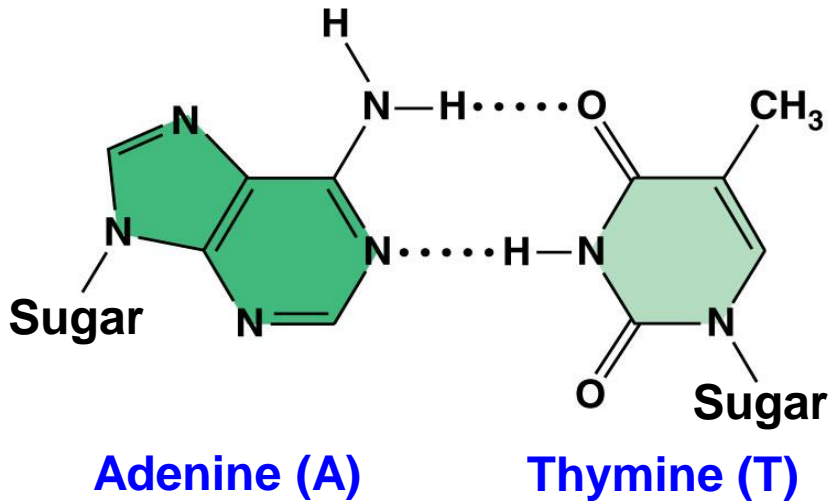
(b) Partial chemical structure



(c) Space-filling model

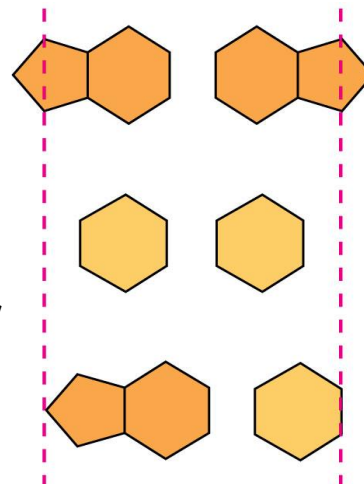
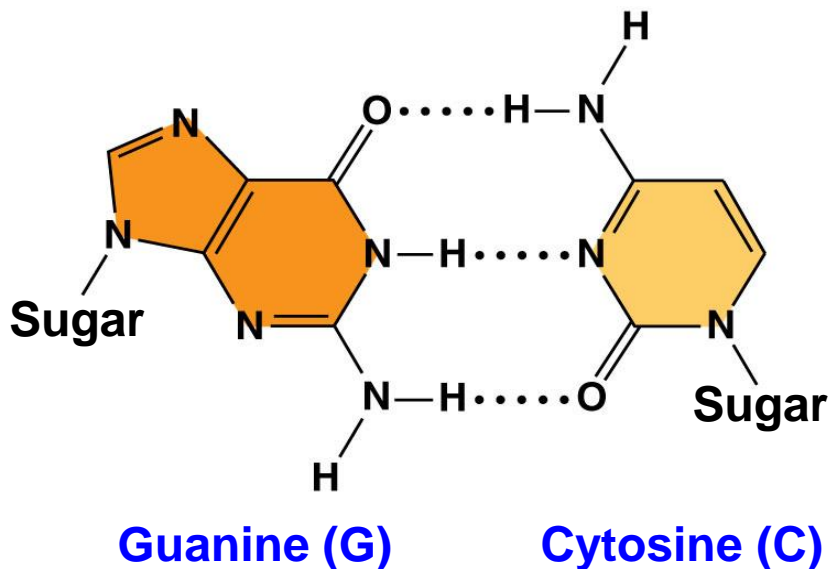
# DNA “Base-Pairing”

Base pairs are held together by hydrogen bonds.



## Why only A:T and C:G?

- the position of chemical groups involved in H-Bonds
- the size of the bases (purine & pyrimidine)



Purine + purine: too wide

Pyrimidine + pyrimidine: too narrow

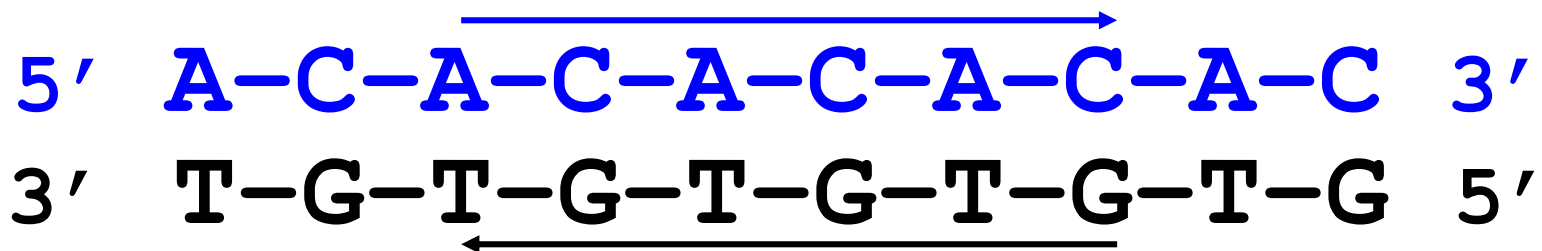
Purine + pyrimidine: width consistent with X-ray data

# The DNA “Sequence”

The DNA sequence is the linear order of nucleotides in a DNA strand:

- each DNA strand in the double helix has its own sequence
- the sequences in each strand are considered as complementary to each other
  - they differ, but “fit just right” with each other
  - ea strand will “fit” with only 1 complementary strand

e.g.



# **2. DNA Replication**

**Chapter Reading – pp. 318-330**

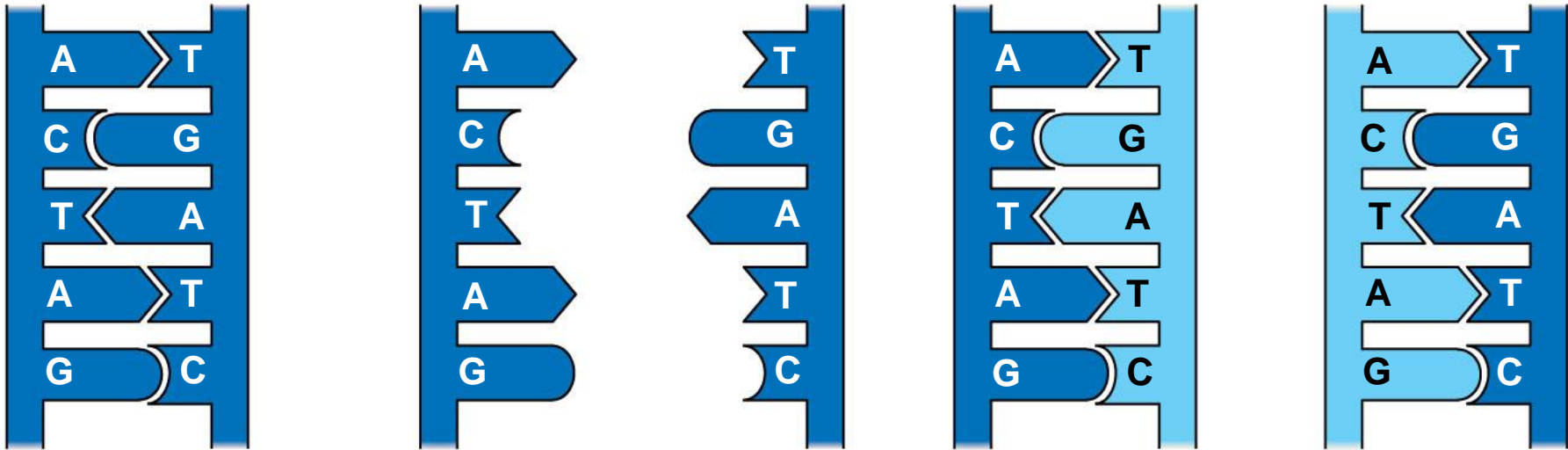
# How is DNA Replicated?

Every time a cell reproduces (i.e., divides) it must replicate its chromosomes (DNA) during S phase.

The process of DNA replication was originally proposed to depend on the rules of base pairing:

- A:T & T:A , C:G & G:C
- the sequence of one strand dictates the sequence of the other
- each strand of the double helix could serve as a template to make a complementary strand

# Model for DNA Replication



(a) Parent molecule

(b) Separation of strands

(c) "Daughter" DNA molecules, each consisting of one parental strand and one new strand

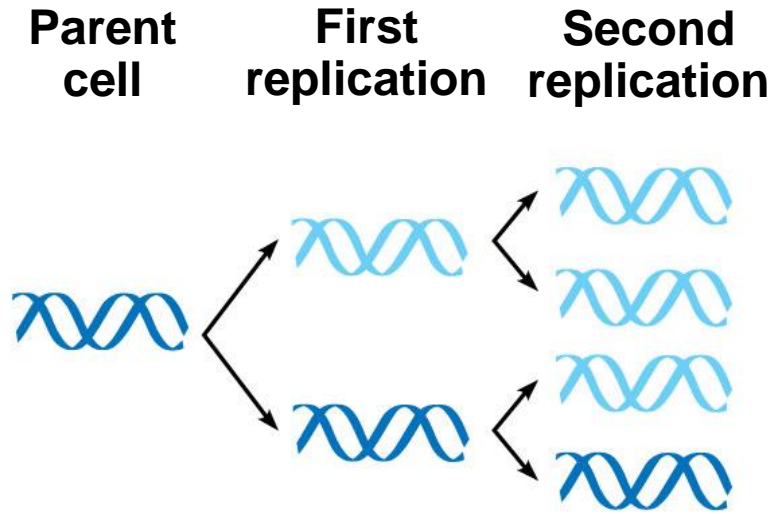
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The semiconservative model of DNA replication proposed that each original strand serves as a template to produce a new complementary strand.

- note that ea original strand ends up in a different molecule

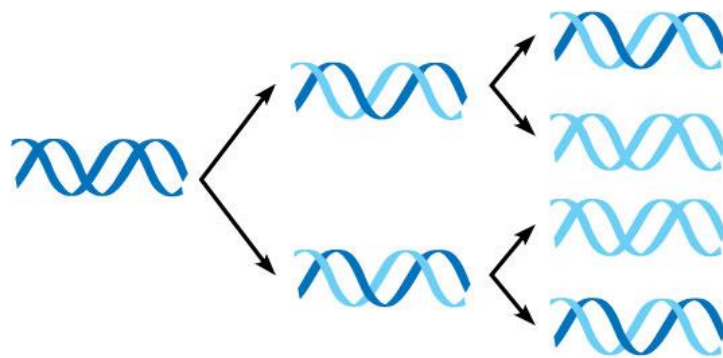
# Other Models

(a) Conservative model



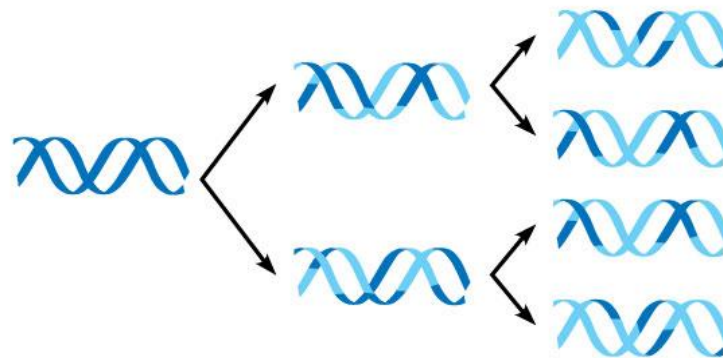
**CONSERVATIVE**  
The original DNA strands stay together

(b) Semiconservative model



**SEMICONSERVATIVE**  
The original DNA strands remain intact in separate molecules

(c) Dispersive model



**DISPERSIVE**  
The original DNA strands are dispersed among the all daughter strands



# Testing the Models

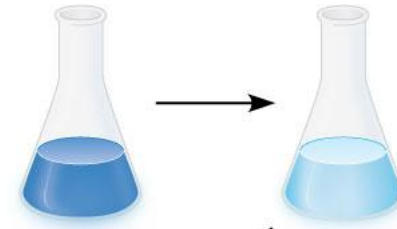
In this experiment, bacteria with DNA containing the “heavy” isotope  $^{15}\text{N}$  were allowed to reproduce in medium containing lighter  $^{14}\text{N}$ .

Density-gradient centrifugation revealed that DNA replication is semiconservative.

**Matthew Meselson & Franklin Stahl, 1958**

## EXPERIMENT

1 Bacteria cultured in medium with  $^{15}\text{N}$  (heavy isotope)



2 Bacteria transferred to medium with  $^{14}\text{N}$  (lighter isotope)

## RESULTS

3 DNA sample centrifuged after first replication



4 DNA sample centrifuged after second replication



Less dense  
More dense

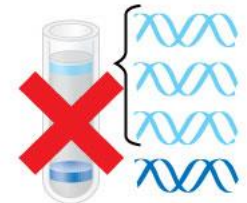
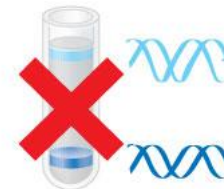
## CONCLUSION

Predictions:

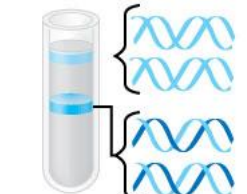
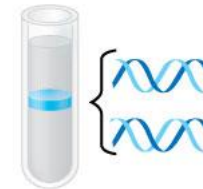
First replication

Second replication

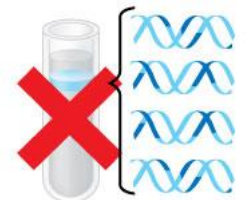
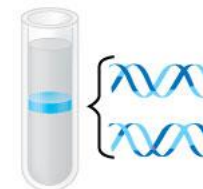
Conservative model



Semiconservative model

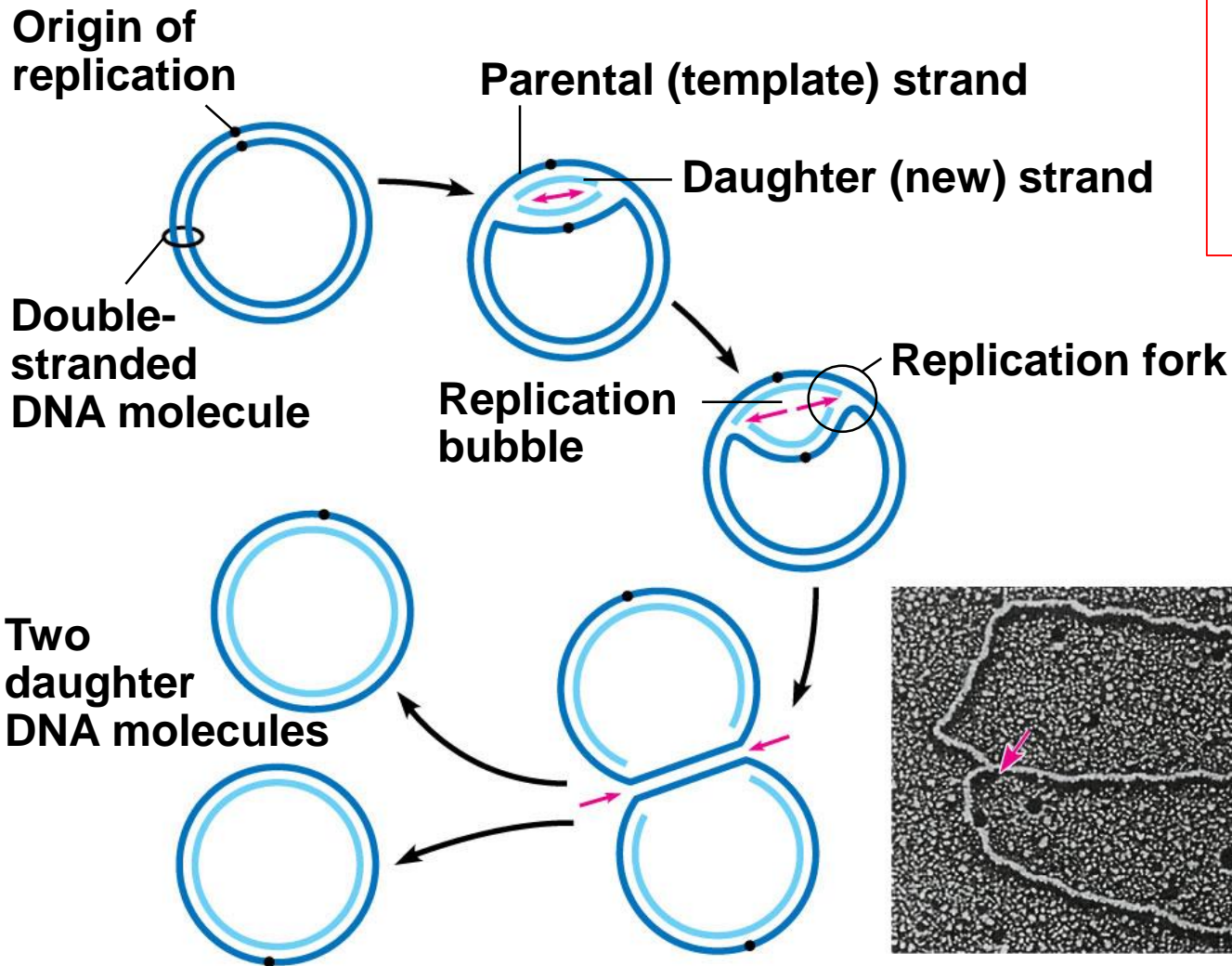


Dispersive model

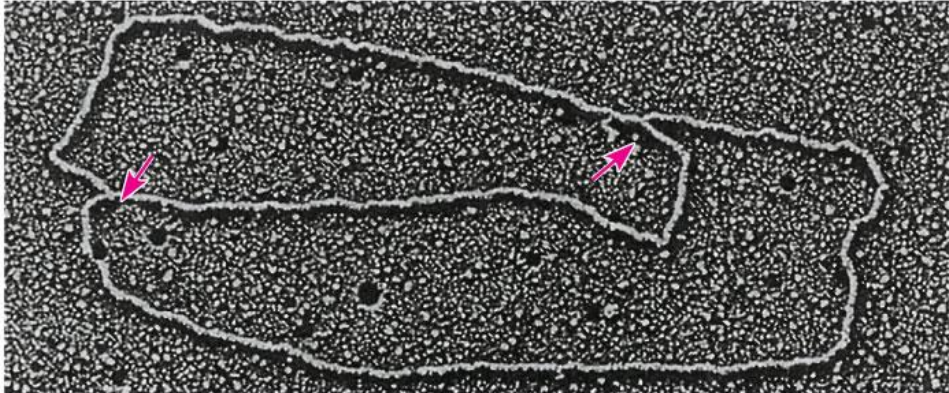


# DNA Replication in Bacteria

## (a) Origin of replication in an *E. coli* cell



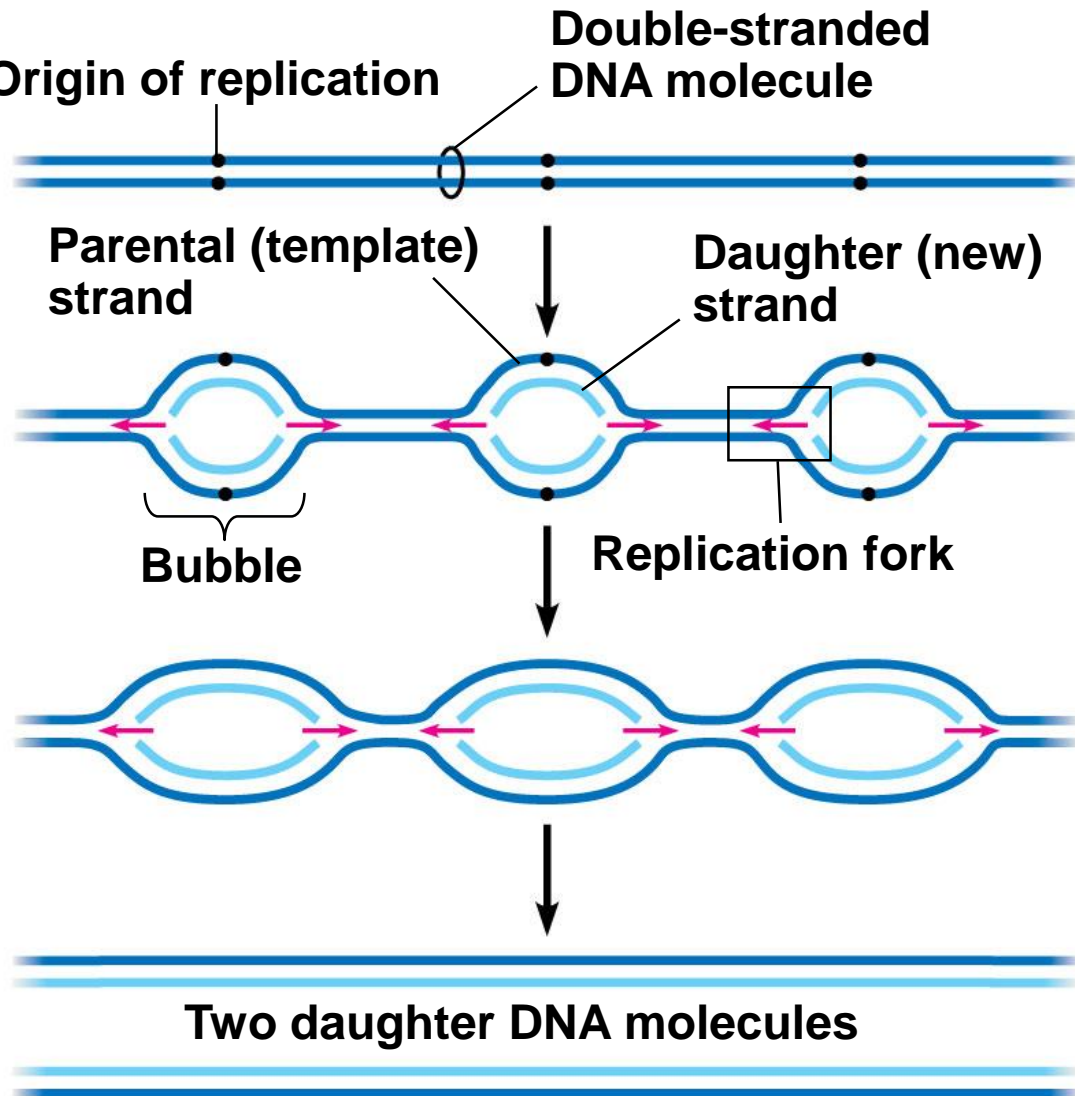
**Initiation of DNA replication requires an origin of replication**



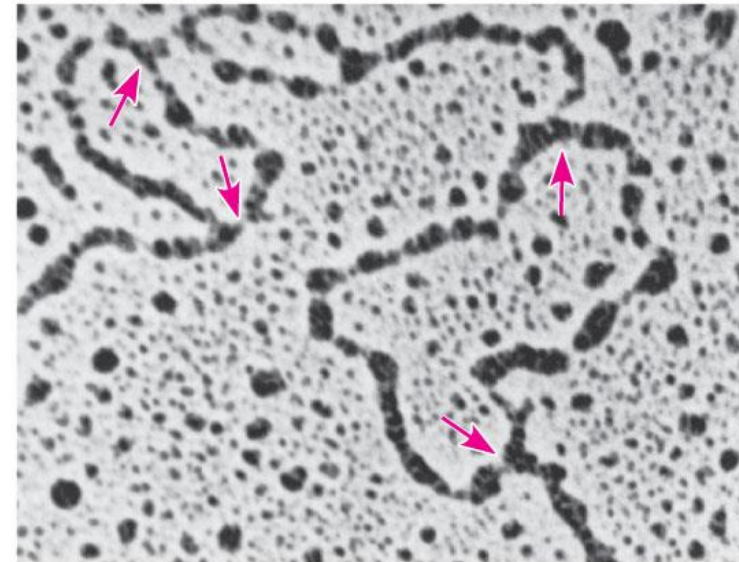
0.5 μm

# DNA Replication in Eukaryotes

## (b) Origins of replication in a eukaryotic cell



**Eukaryotic DNA replication requires multiple origins of replication**



0.25 μm

# Overview of DNA Replication

## Overview

Leading strand

Origin of replication

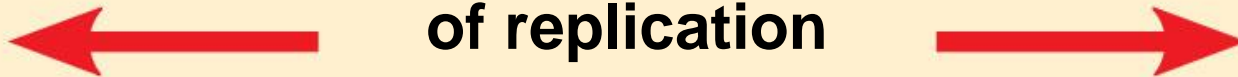
Lagging strand

Primer

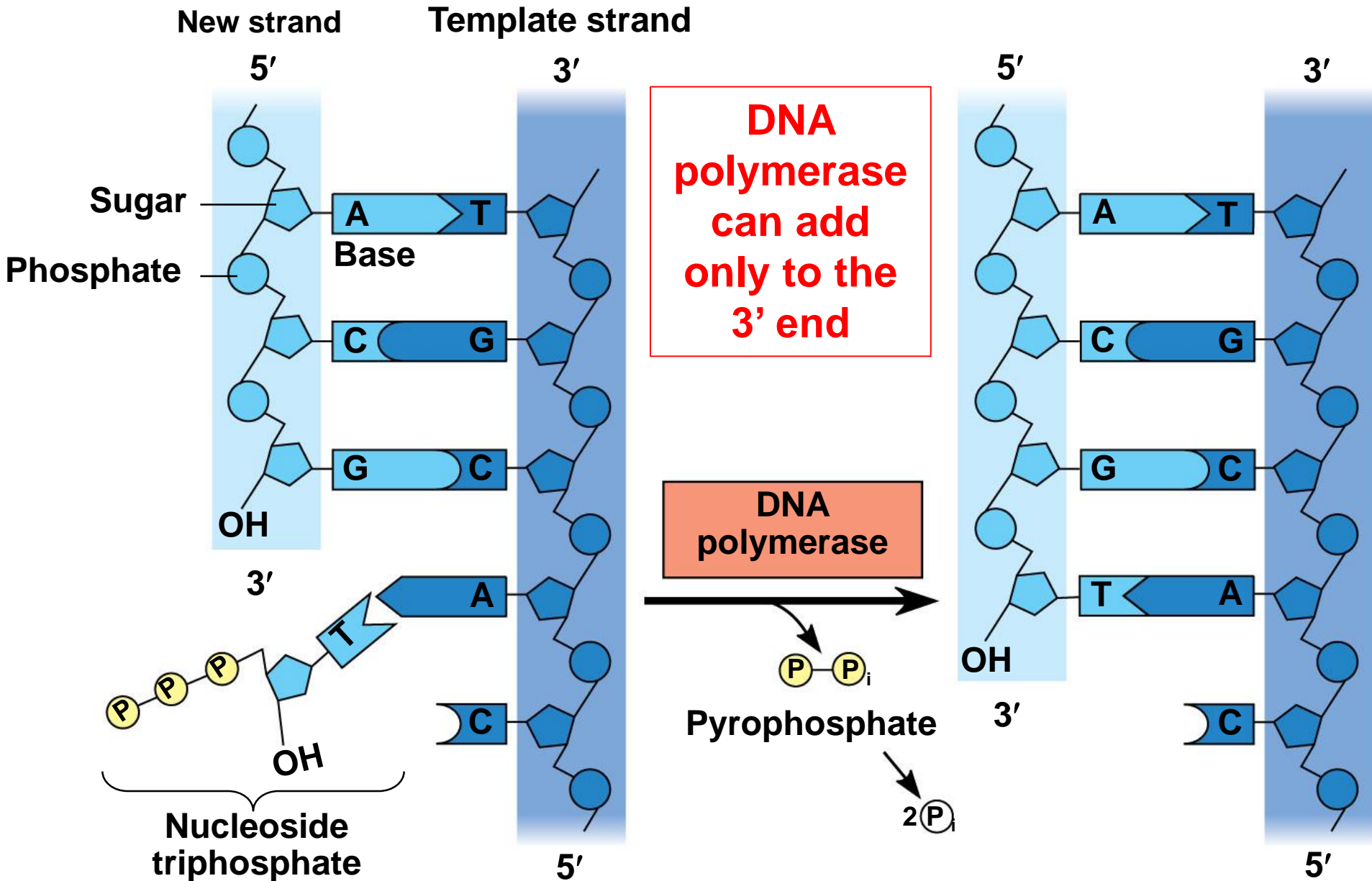
Lagging strand

Leading strand

Overall directions of replication



# DNA Replication proceeds 5' to 3'

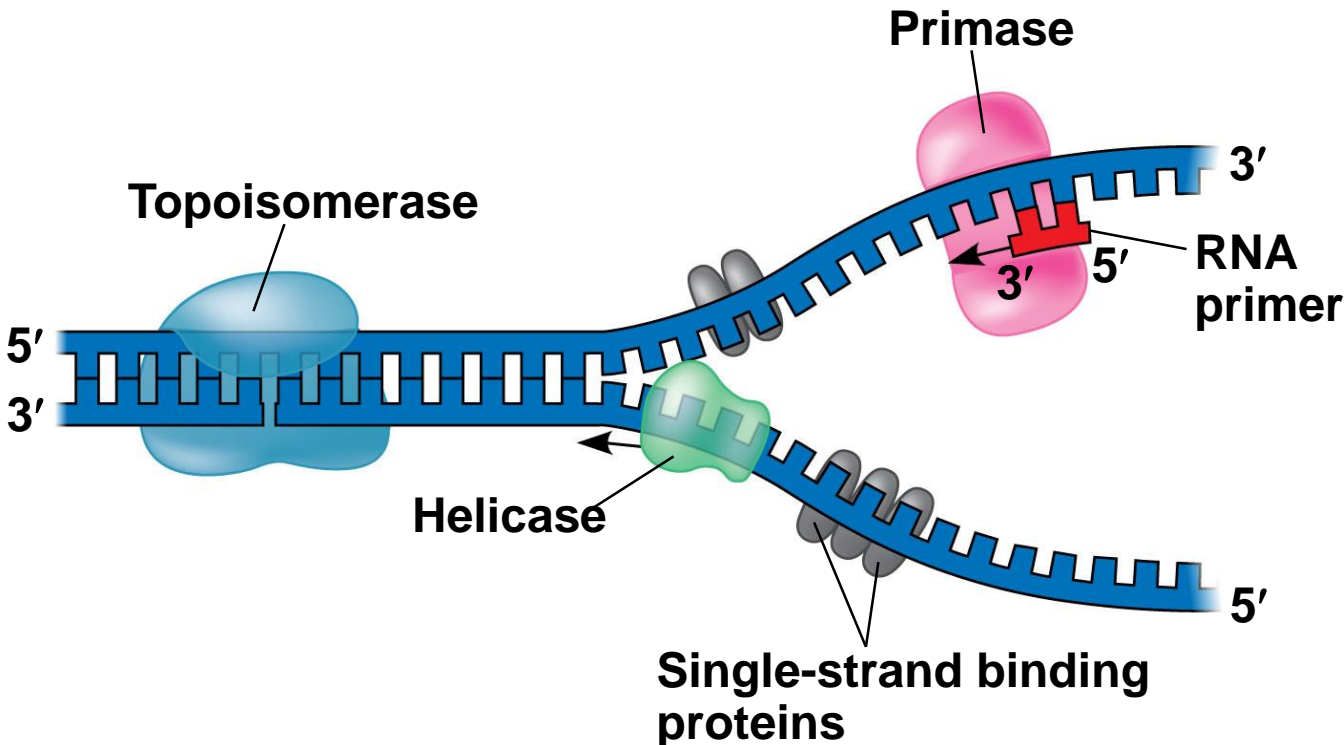


# Enzymes involved in DNA Replication

DNA Polymerase – synthesizes new DNA

Helicase – unwinds DNA double helix

Topoisomerase – relieves tension due to DNA unwinding

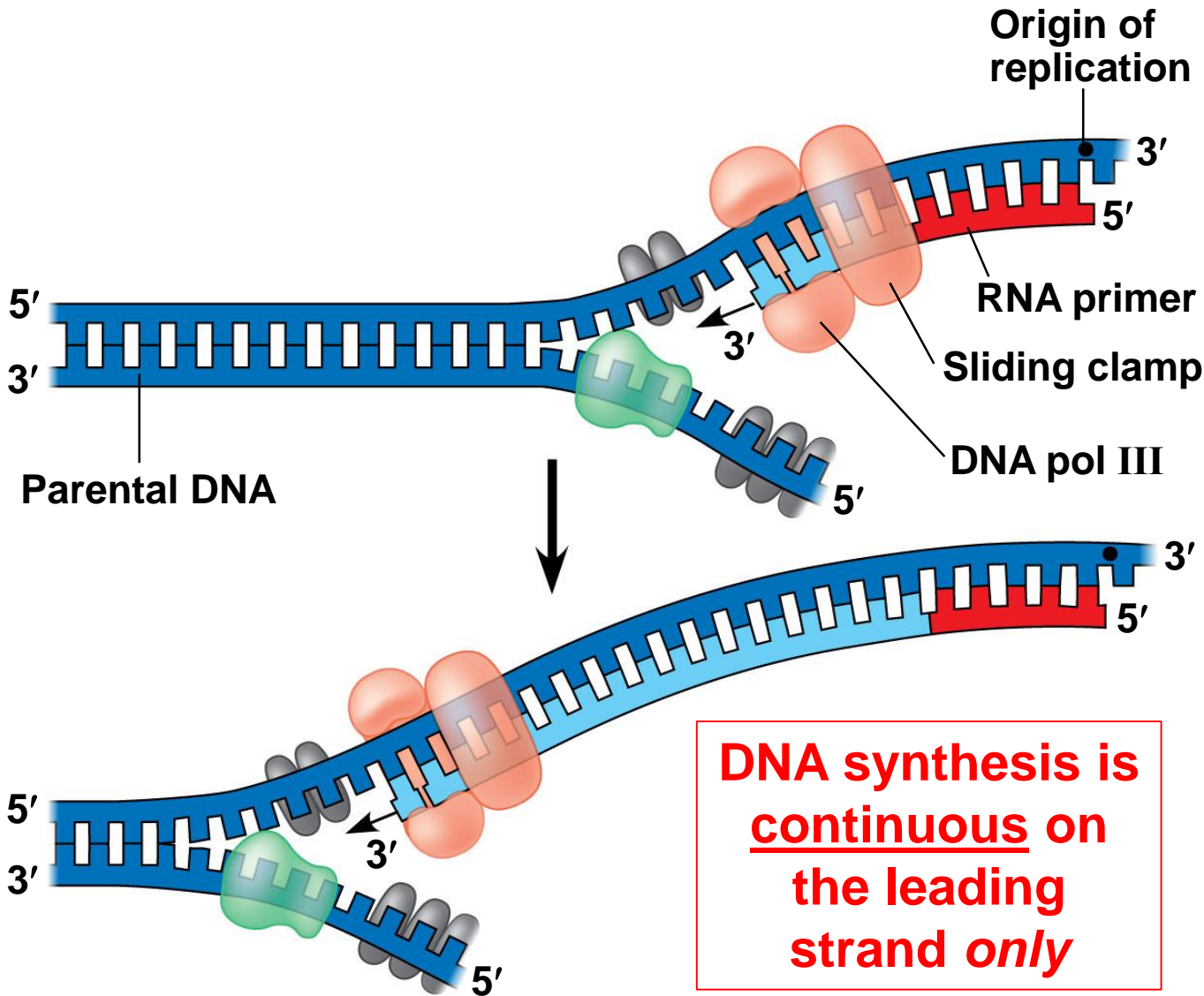


Primase –  
makes short  
RNA primers

DNA Ligase –  
connects DNA  
fragments

# Leading Strand DNA Synthesis

Proceeds toward unwinding replication fork:



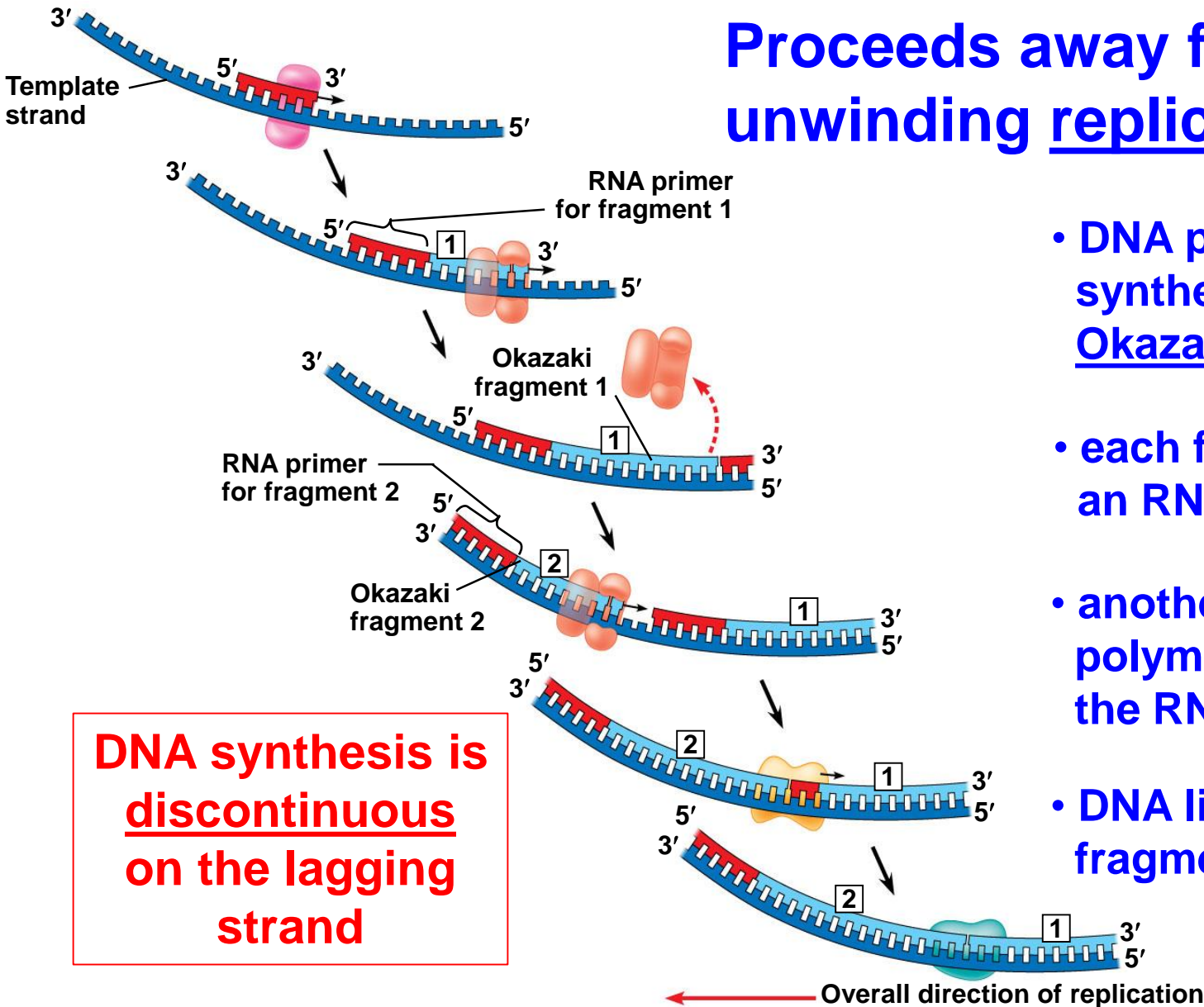
DNA polymerase can only synthesize DNA in a 5' to 3' direction.

DNA polymerase requires an RNA primer which it can extend in a continuous manner toward the unwinding replication fork

**DNA synthesis is continuous on the leading strand *only***

# Lagging Strand DNA Synthesis

Proceeds away from the unwinding replication fork:

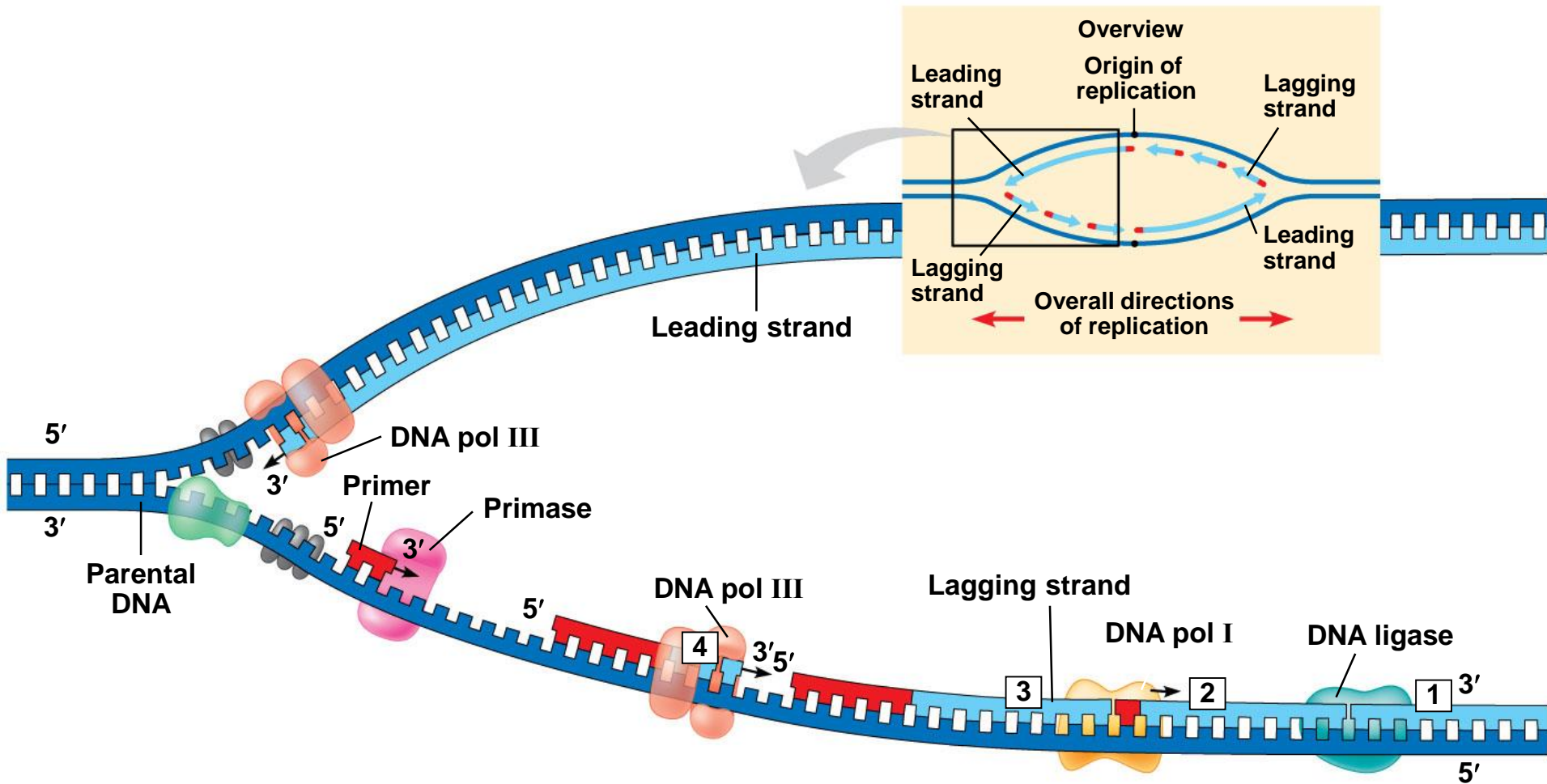


**DNA synthesis is discontinuous on the lagging strand**

- DNA polymerase synthesizes DNA in Okazaki fragments
- each fragment requires an RNA primer
- another DNA polymerase will replace the RNA with DNA
- DNA ligase will link the fragments together



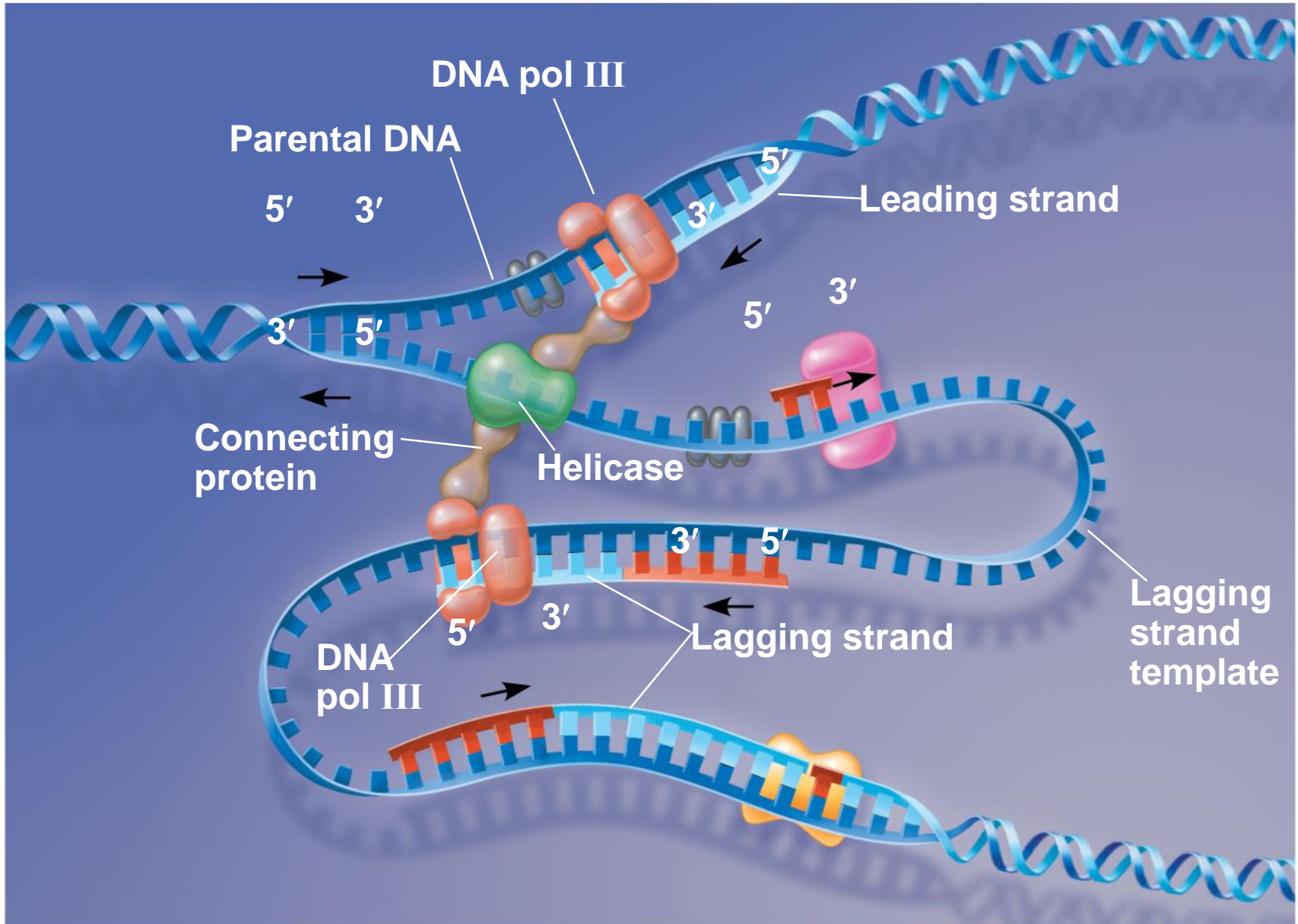
# Summary of DNA Replication



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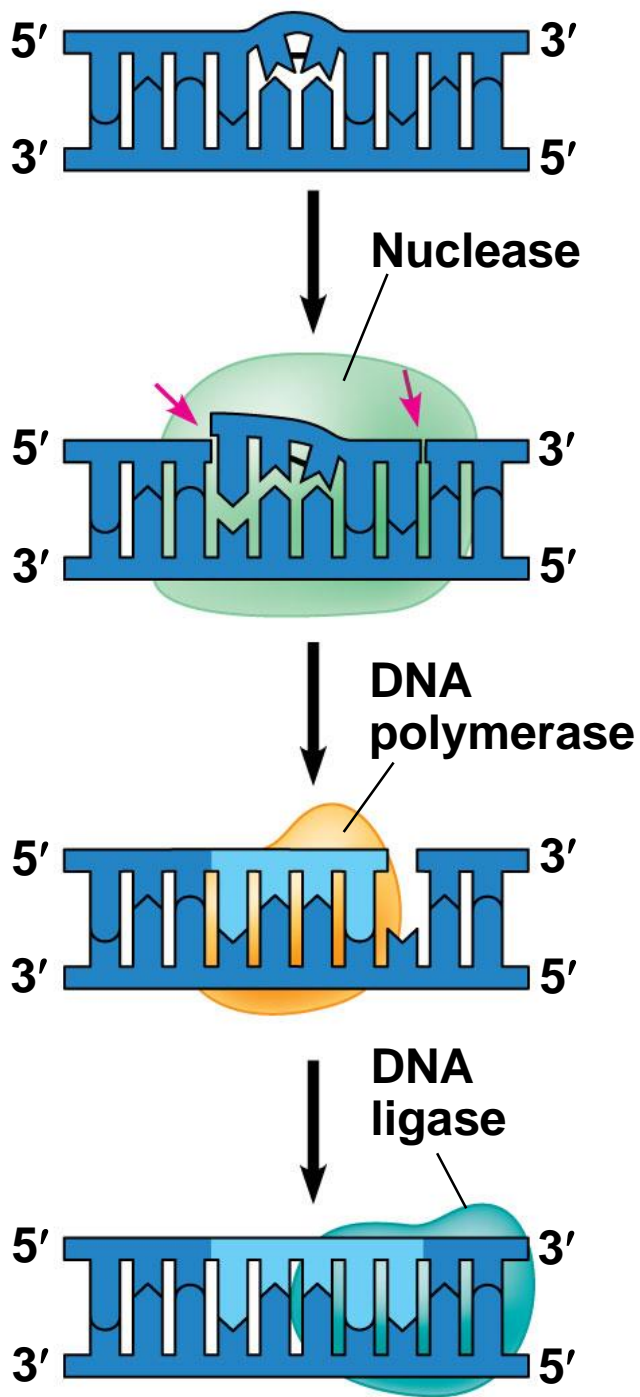
**DNA replication proceeds in this manner in ALL living organisms.**

# Current Model of DNA Replication



# DNA Repair

When DNA is damaged it is essential that the DNA is repaired so it can be replicated and expressed properly.

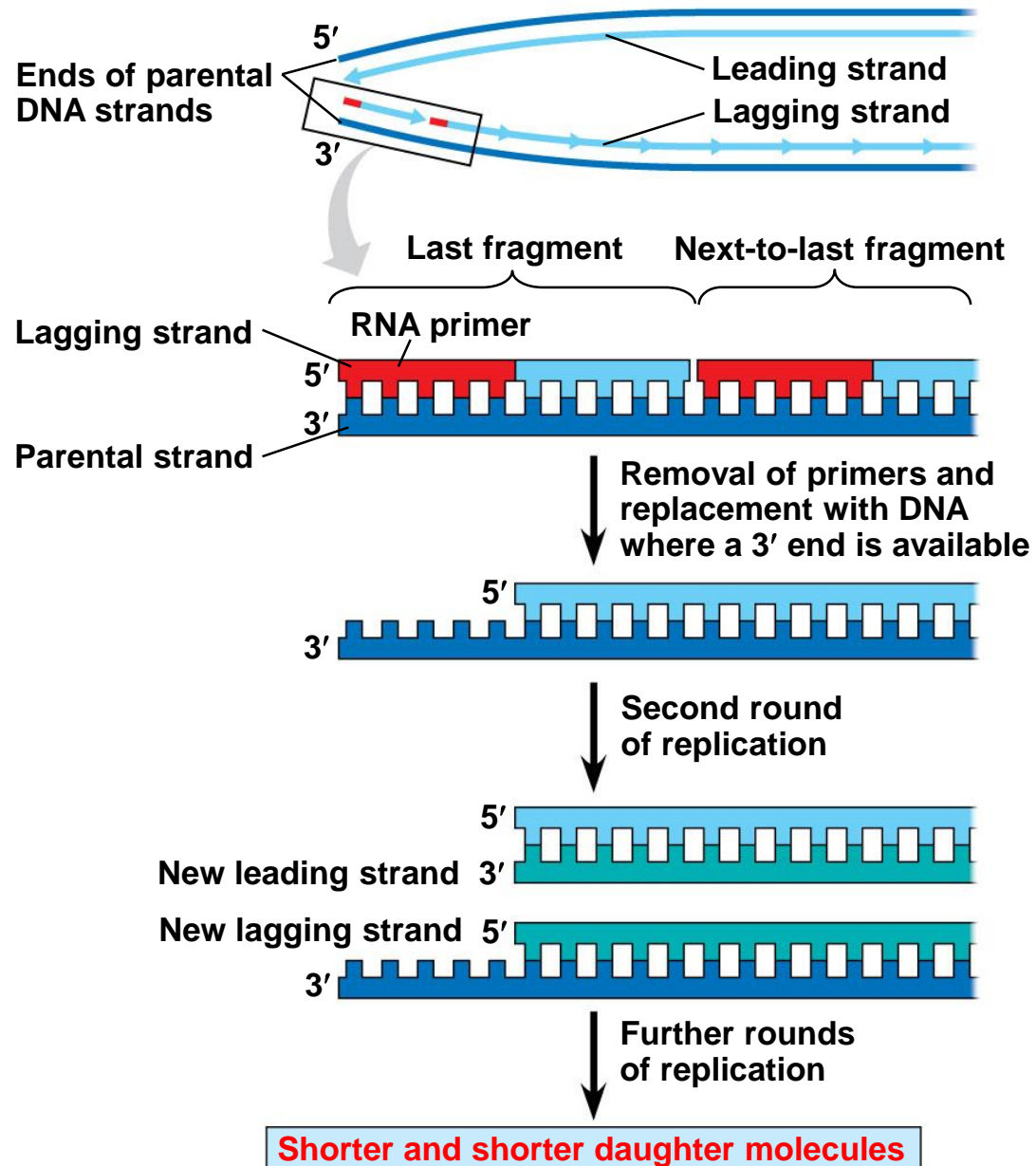


- special enzymes recognize and remove the damaged portion of DNA
- a DNA polymerase will fill in the gap
- DNA ligase will then connect the newly made DNA to the adjacent strand

# The Problem with Telomeres

The ends of linear chromosomes, the telomeres, cannot be completely copied on the lagging strand.

This results in progressive shortening of the chromosome every time it is replicated.



**Telomerase will solve this problem in certain cell types**



# ...more on Chromatin

**Chromatin refers to the complex of DNA and histone proteins in eukaryotic nuclei:**

- **chromosomal DNA wraps around histone proteins to form structures called nucleosomes that look like “beads on a string”**
- **different parts of a chromosome can be in various states of “packing”**

**EUCHROMATIN – loosely packed DNA**

**HETEROCHROMATIN – tightly packed DNA**

# Key Terms for Chapter 16

- bacterial transformation, bacteriophage
- x-ray crystallography
- pyrimidine, purine, base-pair, complementary
- double helix, anti-parallel, sugar-phosphate backbone
- DNA replication, template
- conservative, semiconservative, dispersive
- DNA polymerase, helicase, topoisomerase, primase, DNA ligase
- leading, lagging strand; continuous, discontinuous
- Okazaki fragment, telomere, telomerase
- chromatin, nucleosome, euchromatin, heterochromatin

**Relevant  
Chapter  
Questions  
1-7, 9**