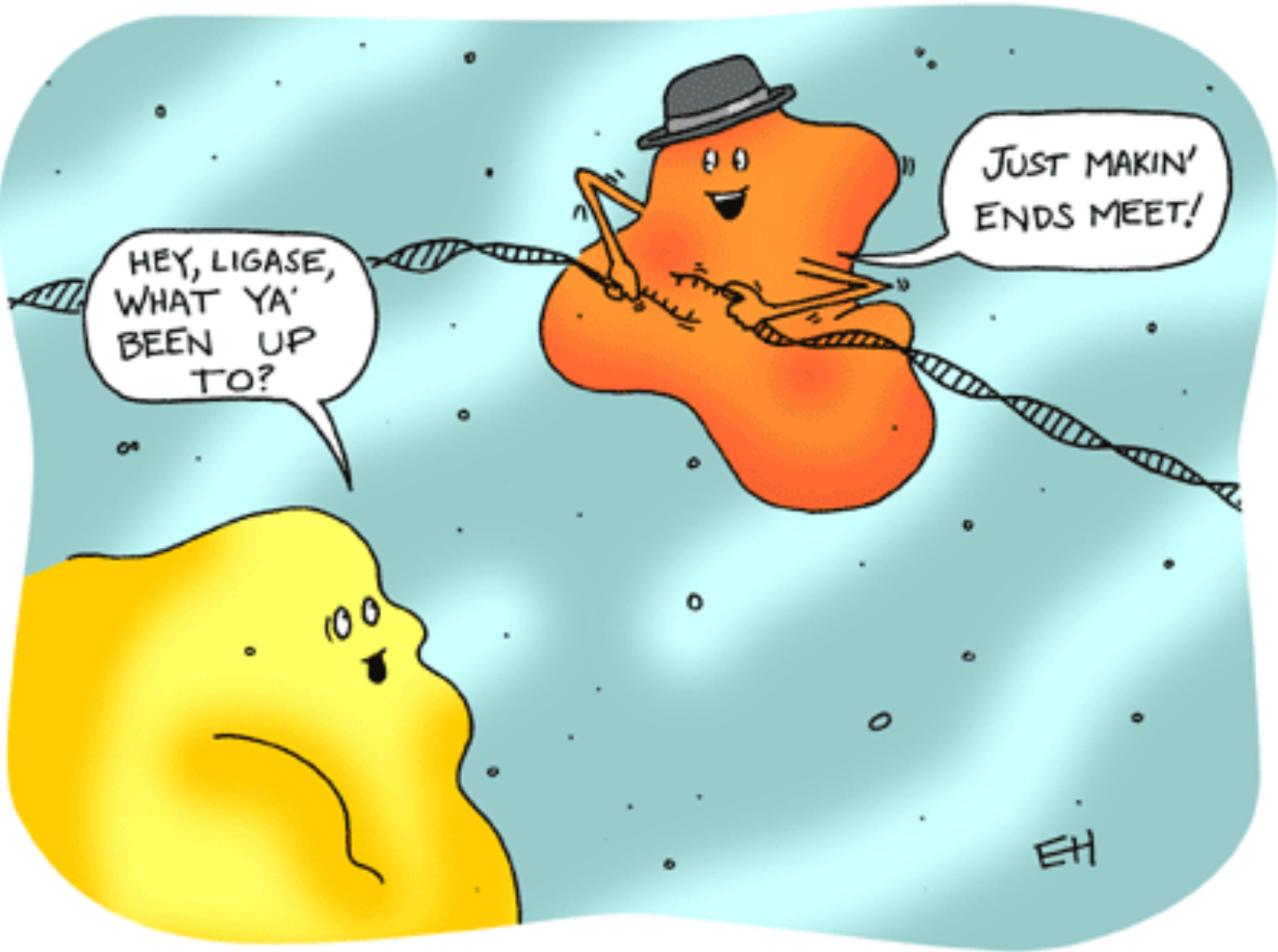

HEY, LIGASE, WHAT YA' BEEN UP TO?
JUST MAKIN' ENDS MEET!
EH

# THE MOLECULAR BASIS OF INHERITANCE

Chapter 13

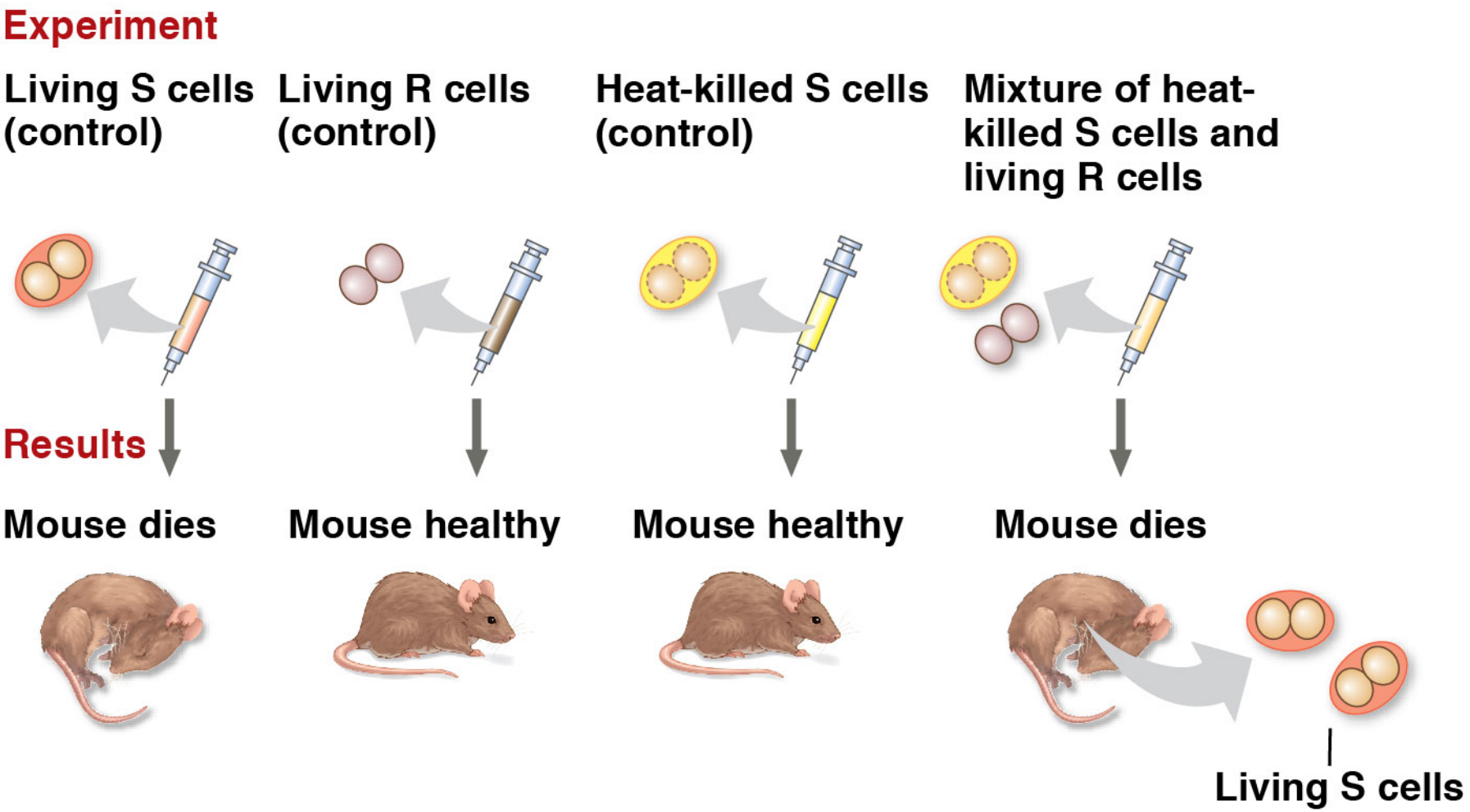
# What you must know

- The structure of DNA.
- The knowledge about DNA gained from the work of Griffith; Avery, MacLeod, and McCarty; Hershey and Chase; Wilkins and Franklin; and Watson and Crick.
- That replication is semiconservative and occurs 5' to 3'.
- The roles of DNA polymerase, ligase, helicase, and topoisomerase in replication.
- The general differences between bacterial chromosomes and eukaryotic chromosomes.
- How DNA is packaged can affect gene expression.

# Problem:

Is the genetic material of organisms made of DNA or proteins?

# Frederick Griffith (1928)



# Frederick Griffith (1928)

Conclusion: living R bacteria transformed into deadly S bacteria by unknown, heritable substance

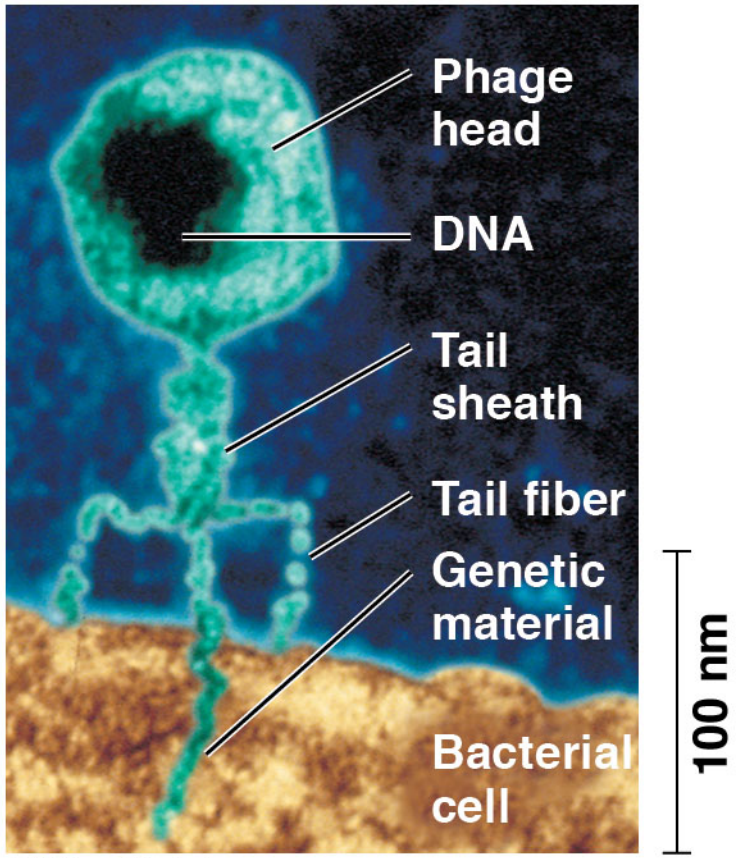
**Avery, McCarty, MacLeod (1944)**

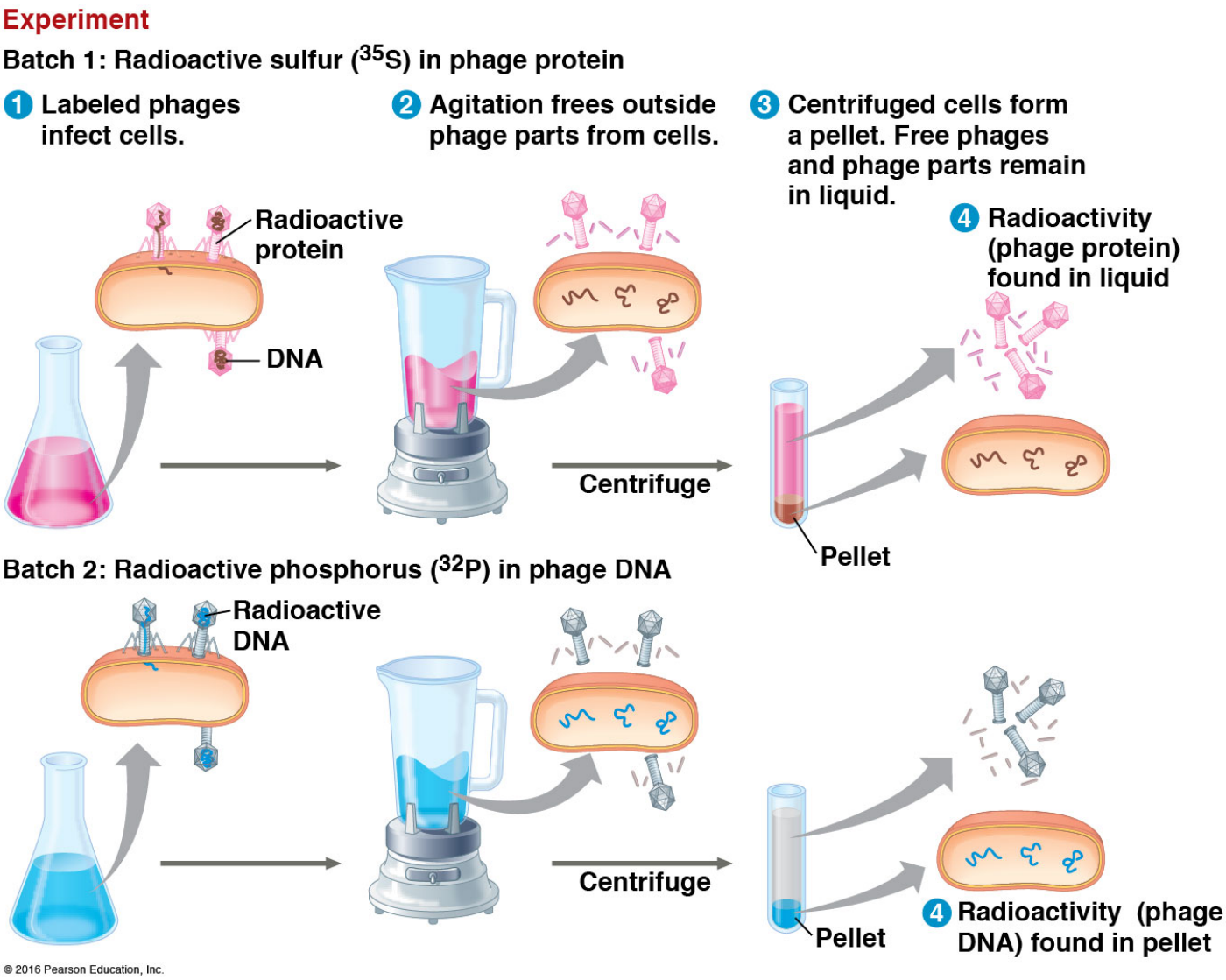
- Tested DNA, RNA, & proteins in heat-killed pathogenic bacteria
- Discovered that the transforming agent was DNA

# Hershey and Chase (1952)

- Bacteriophages: virus that infects bacteria; composed of DNA and protein

**Protein = radiolabel S**
**DNA = radiolabel P**






**Conclusion**: DNA entered infected bacteria → DNA must be the genetic material!

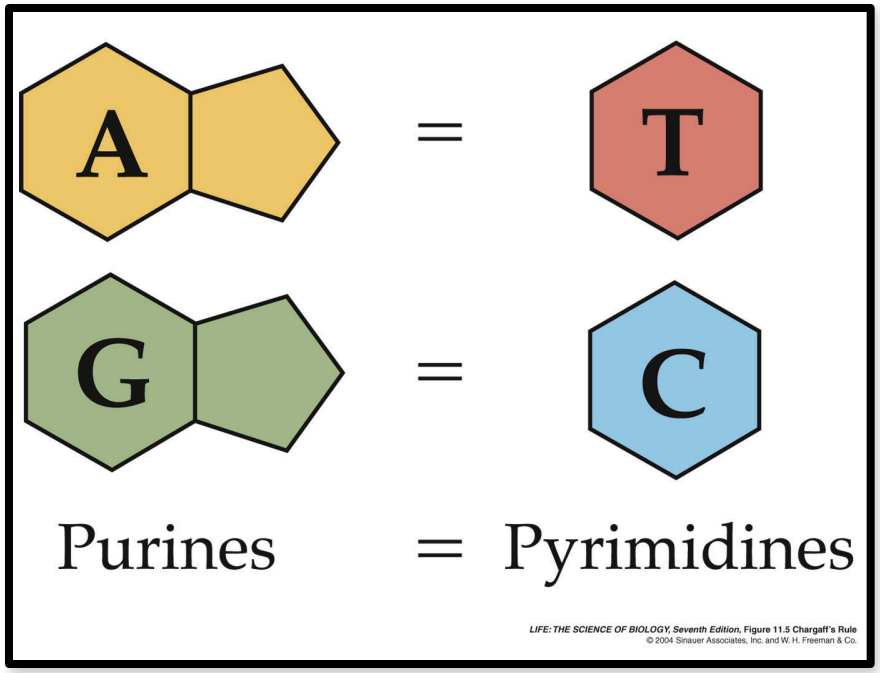
# Problem:

What is the structure of DNA?

# Edwin Chargaff (1947)

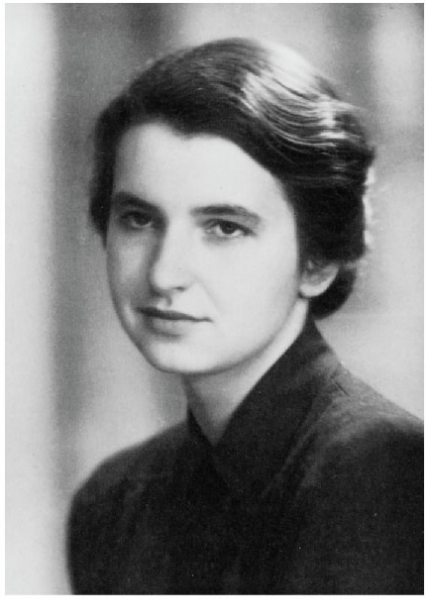
Chargaff's Rules:

- DNA composition varies between species
- Ratios: %A = %T and %G = %C

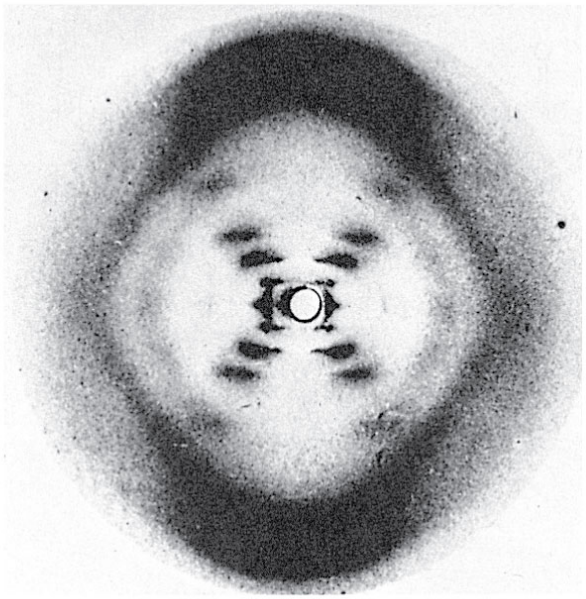


# Rosalind Franklin (1950's)

- Worked with Maurice Wilkins
- X-ray crystallography = images of DNA
- Provided measurements on chemistry of DNA



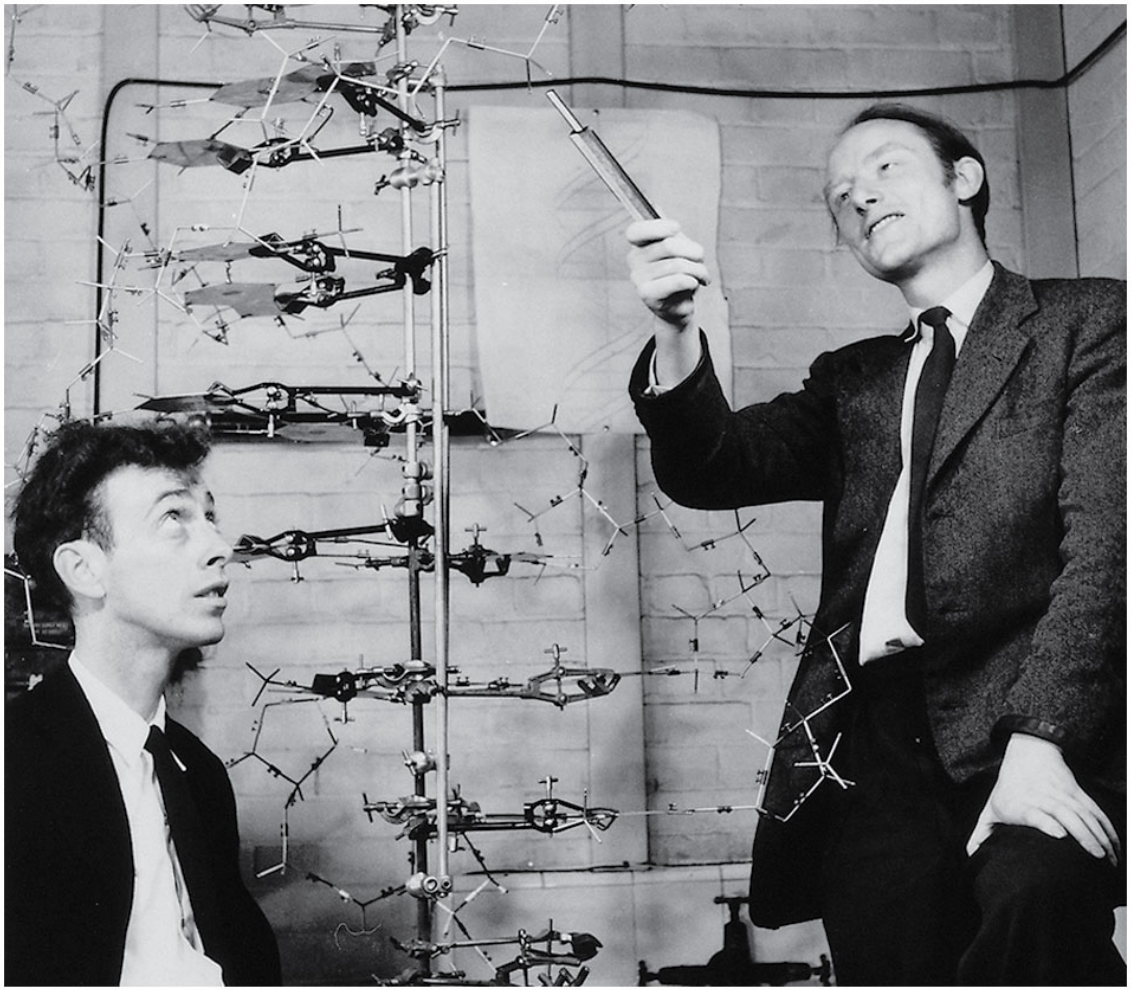
(a) Rosalind Franklin



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

# James Watson & Francis Crick (1953)

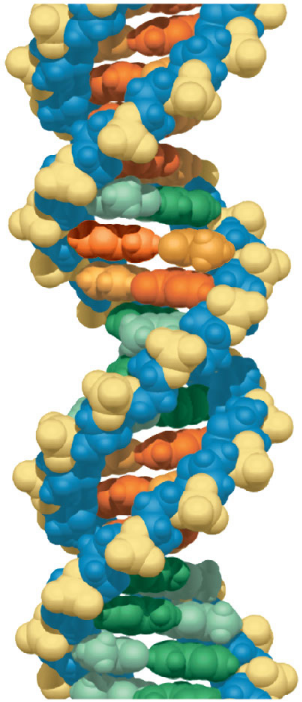
- Discovered the double helix by building models to conform to Franklin's X-ray data and Chargaff's Rules.



# DNA = **Double Helix**

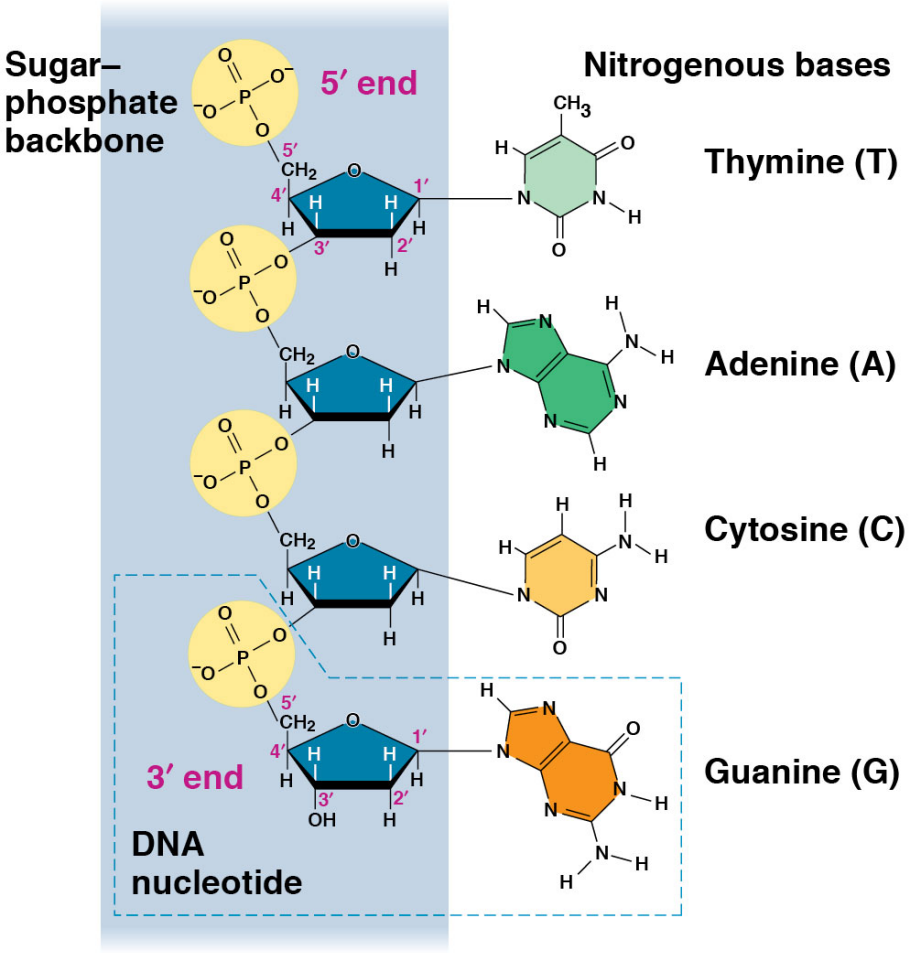
"Backbone" = sugar + phosphate

"Rungs" = nitrogenous bases



(c) Space-filling model





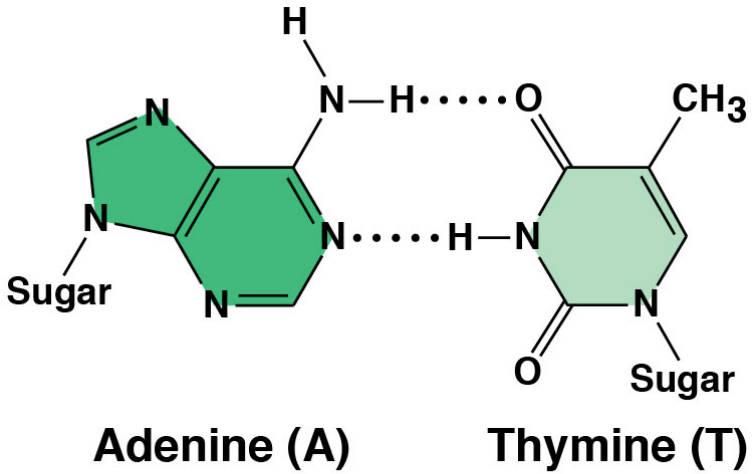
# Nitrogenous Bases

- Adenine (A)
- Guanine (G)

purine

- Thymine (T)
- Cytosine (C)

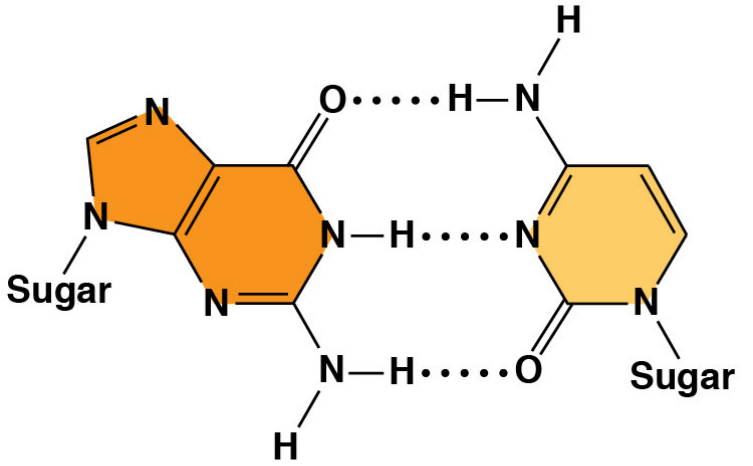
pyrimidine




Adenine (A) Thymine (T)

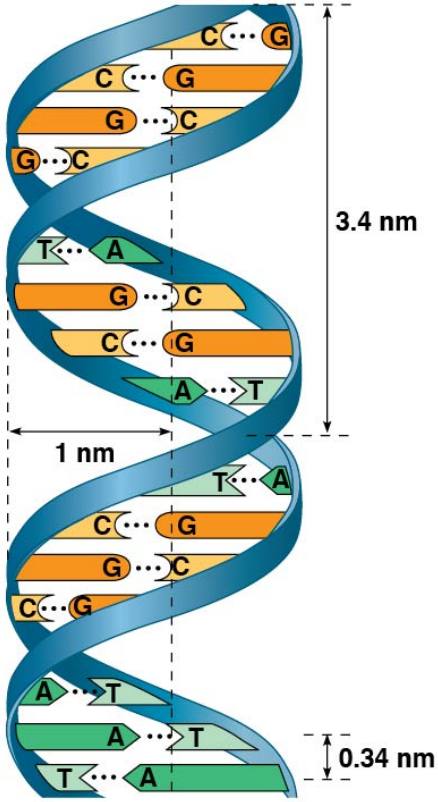
Pairing:

- Purine + Pyrimidine
  - A = T
  - G ≡ C



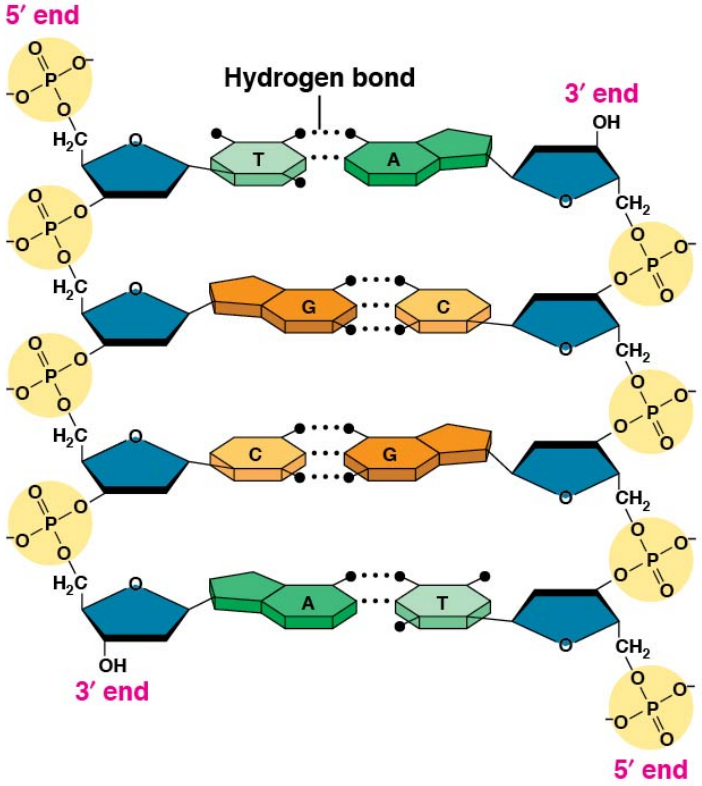

Guanine (G) Cytosine (C)

# Hydrogen Bonds

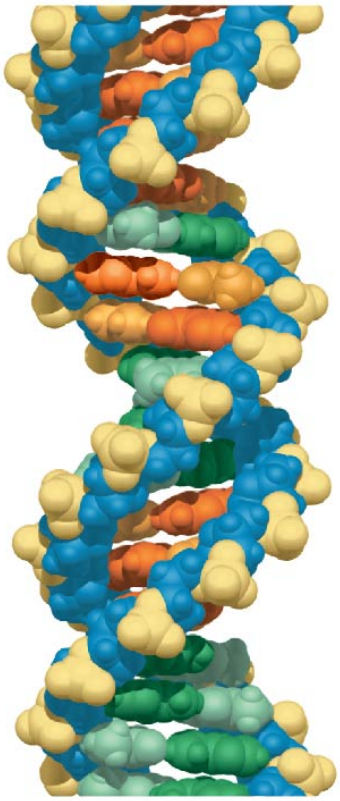



(a) Key features of DNA structure






(b) Partial chemical structure

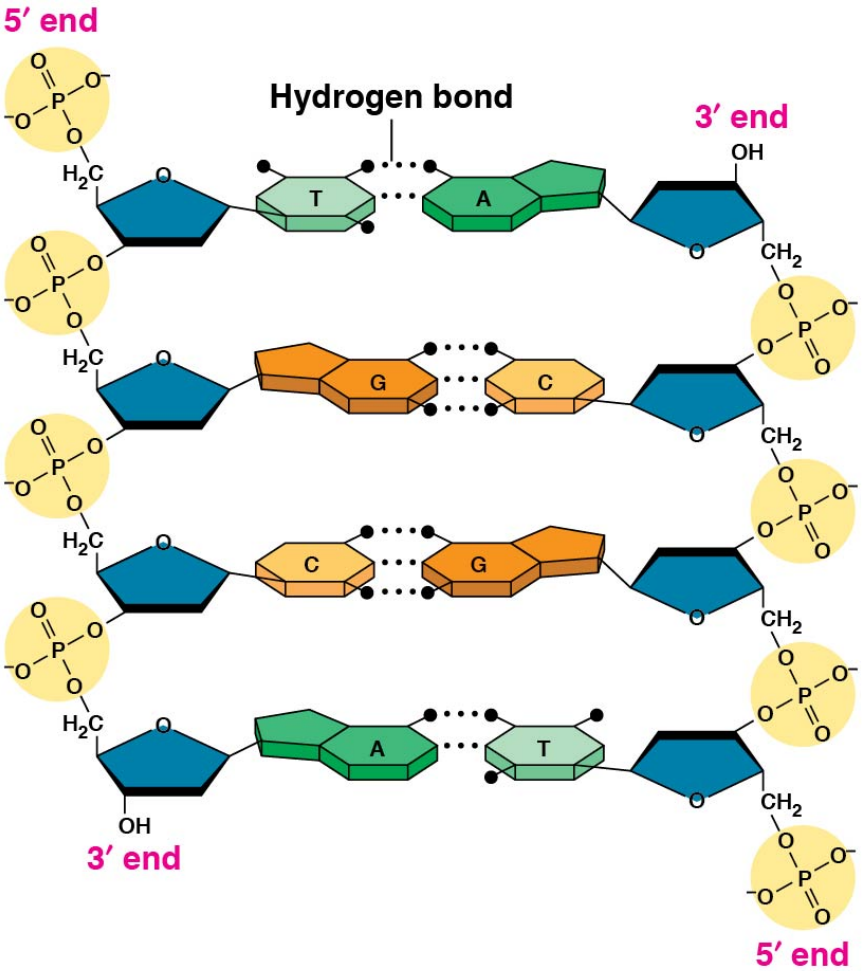


(c) Space-filling model

**Hydrogen bonds** between base pairs of the two strands hold the molecule together like a zipper.

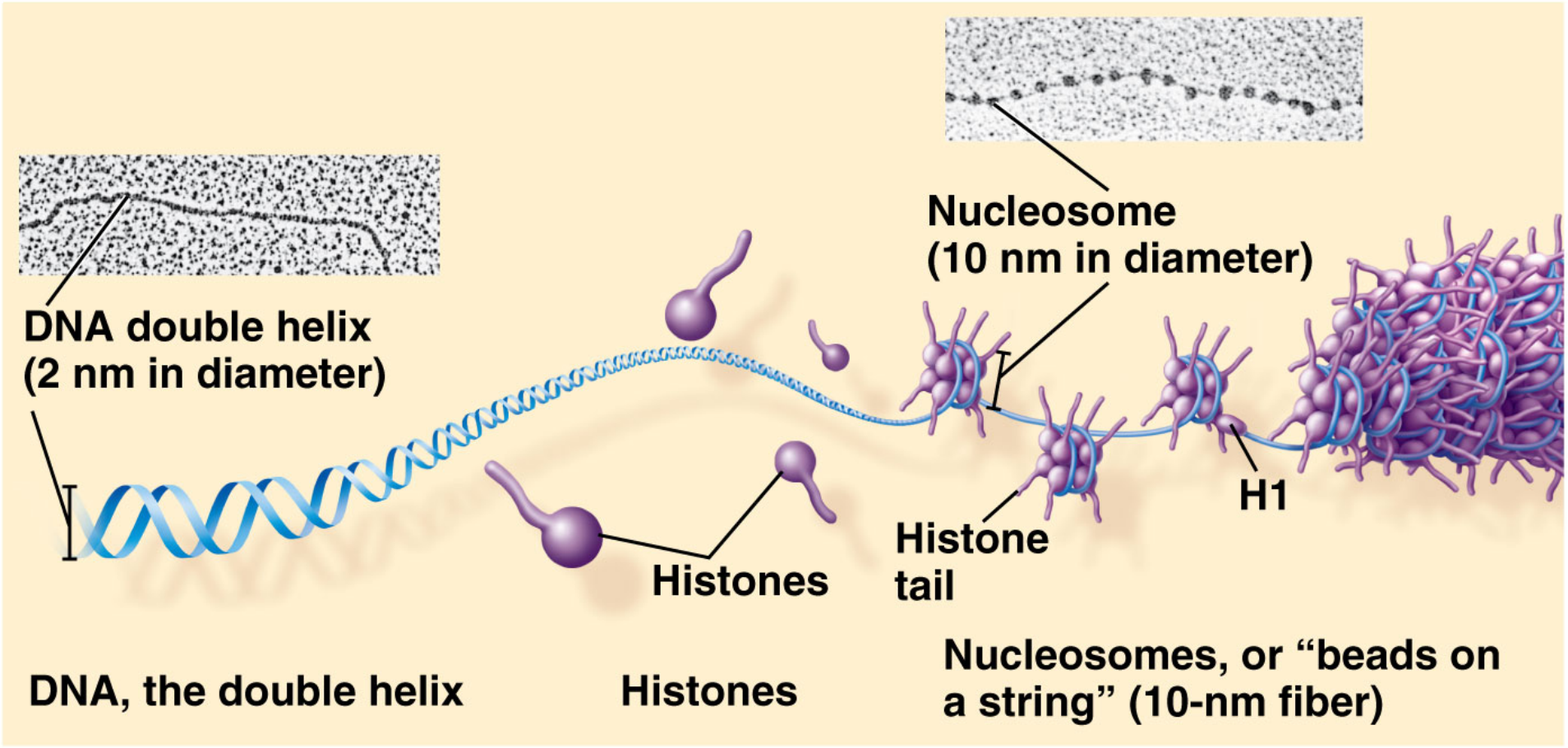
# DNA strands are **Antiparallel**

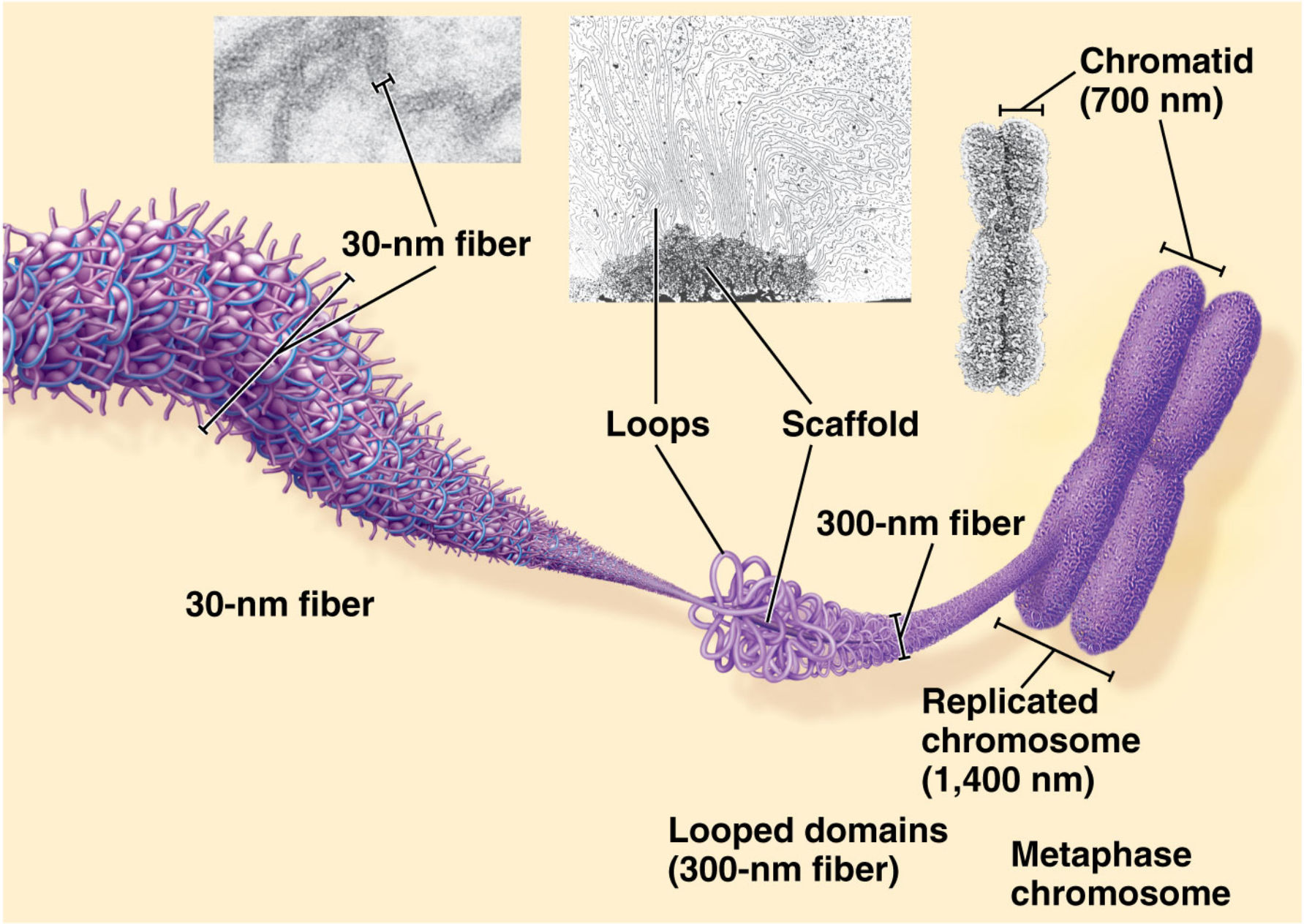
One strand (5'→ 3'), other strand runs in opposite, upside-down direction (3' → 5')




(b) Partial chemical structure

# How DNA is packaged





# DNA Comparison

## Prokaryotic DNA

- Double-stranded
- Circular
- One chromosome
- In cytoplasm
- Supercoiled DNA (nucleoid)
- No histones

## Eukaryotic DNA

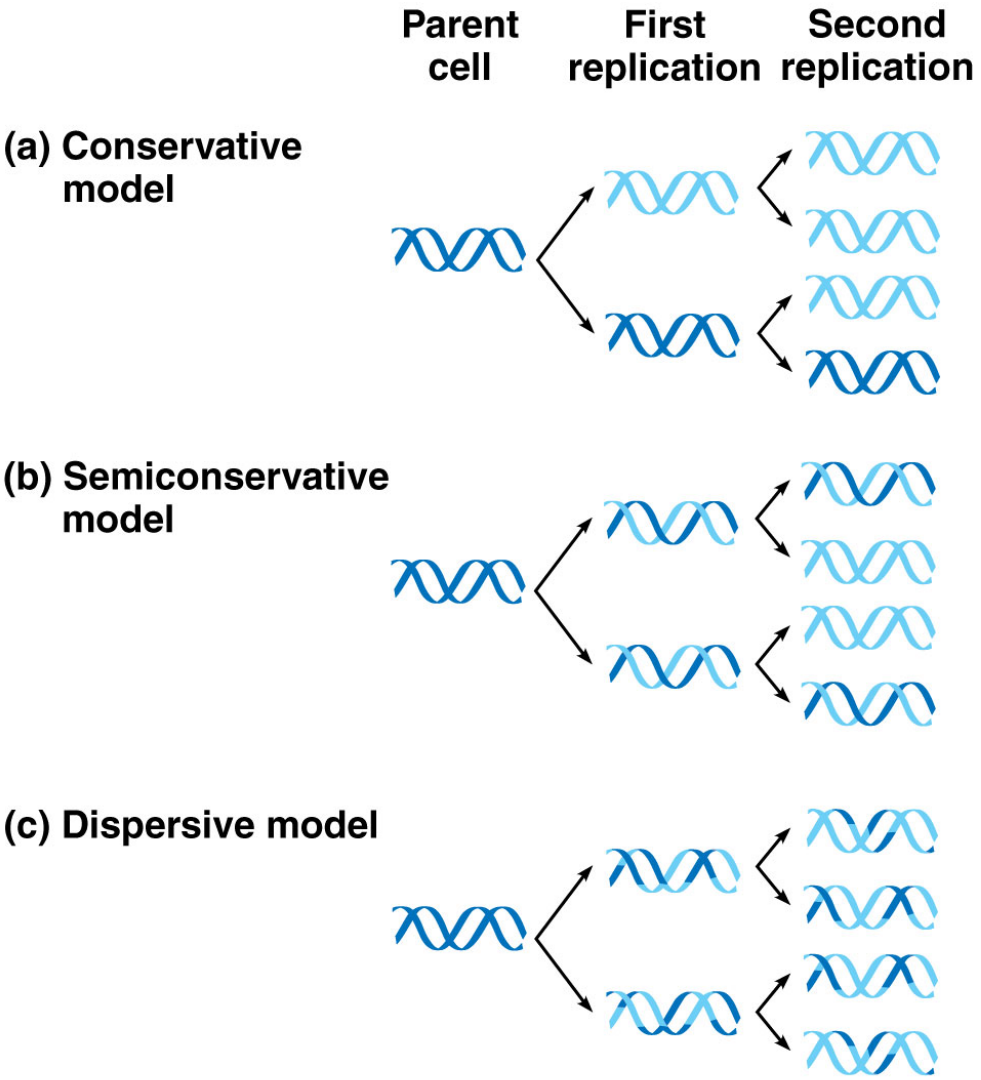
- Double-stranded
- Linear
- Usually 1+ chromosomes
- In nucleus
- Chromatin = DNA wrapped around histones (proteins)

# Problem:

How does DNA replicate?

# **Replication**: Making DNA from existing DNA

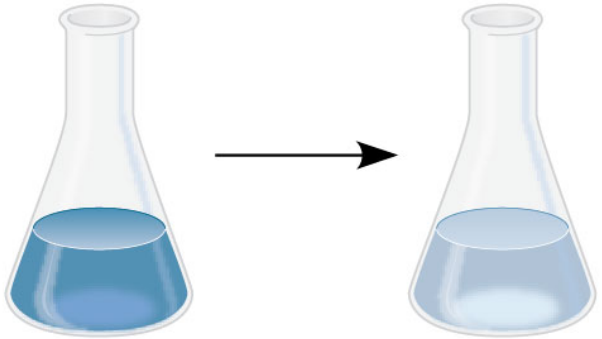
3 alternative models of DNA replication



# Meselson & Stahl

## Experiment

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



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

## Results

3. DNA sample centrifuged after first replication

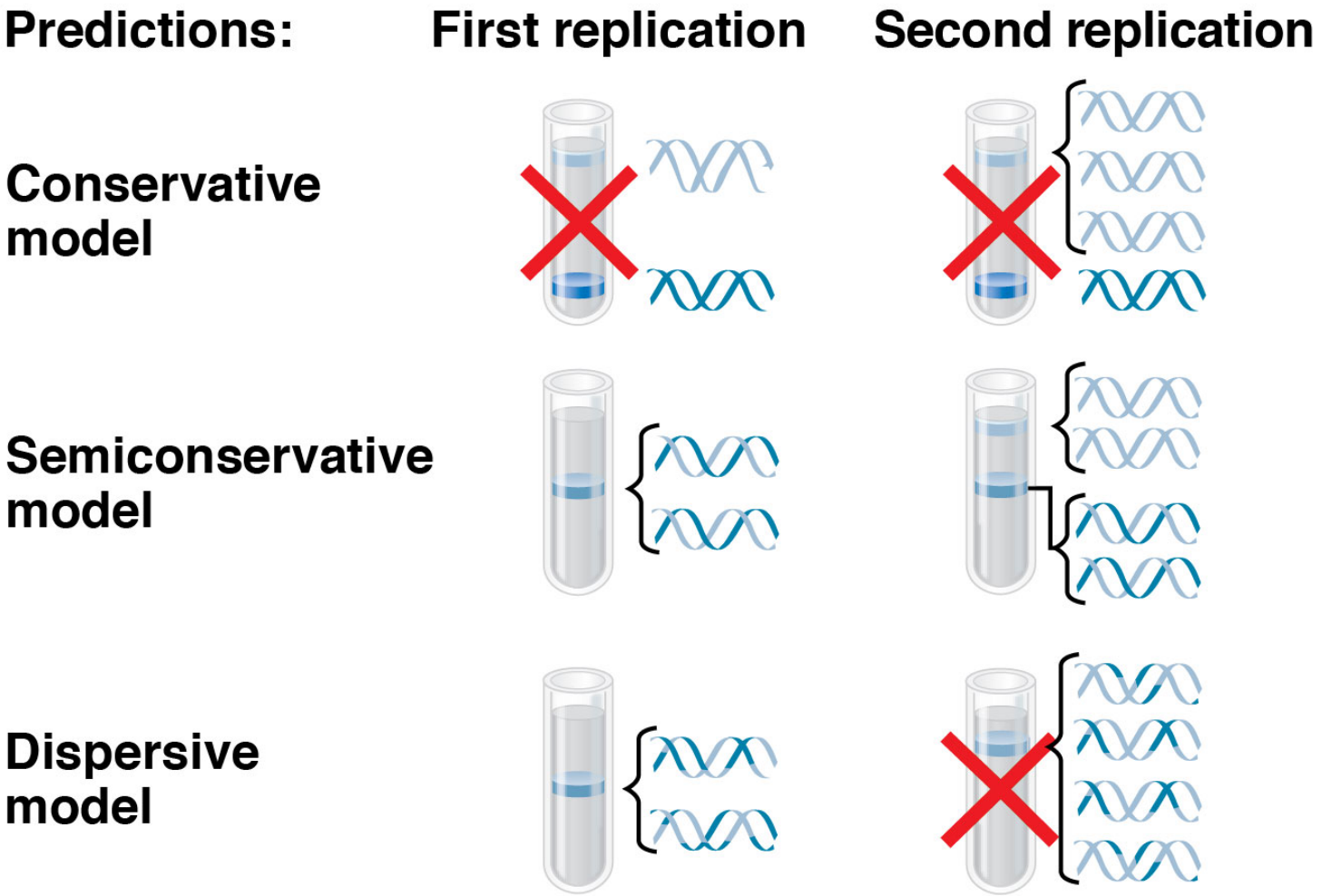
4. DNA sample centrifuged after second replication

Less dense

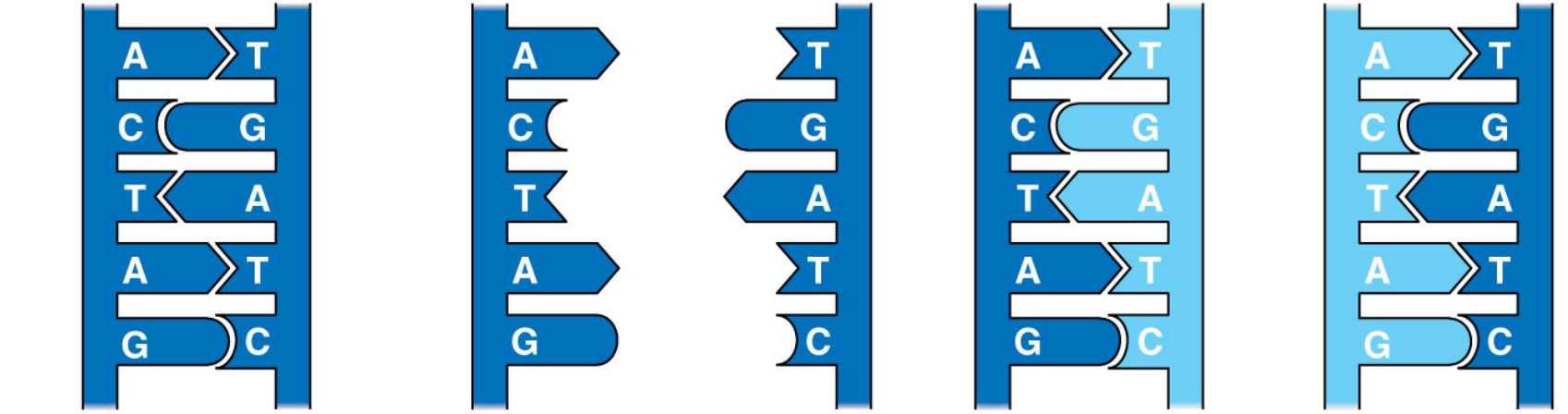
More dense

# Meselson & Stahl

## Conclusion



# Replication is semiconservative



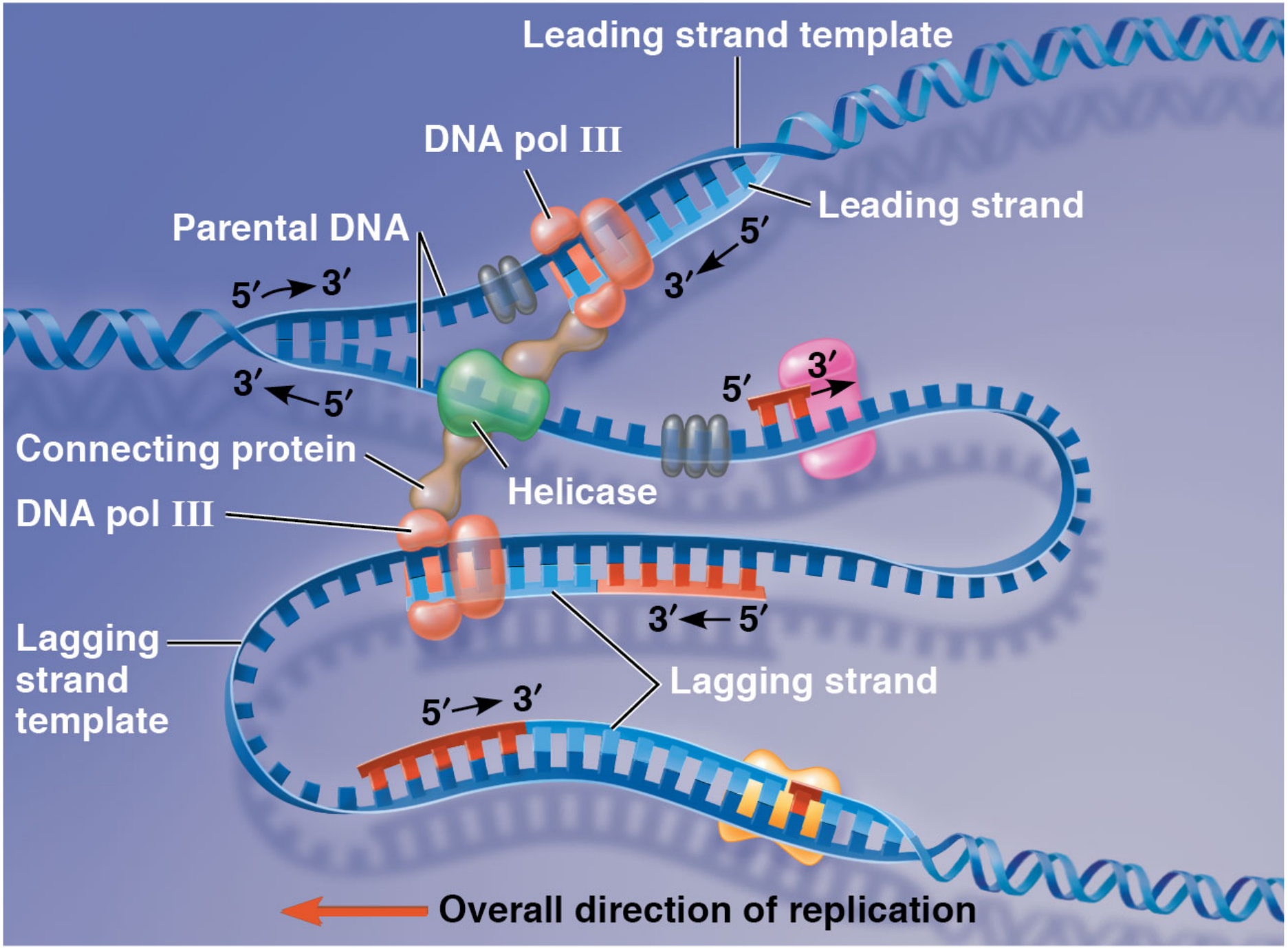

(a) Parent molecule

(b) Separation of strands

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

# DNA Replication Video

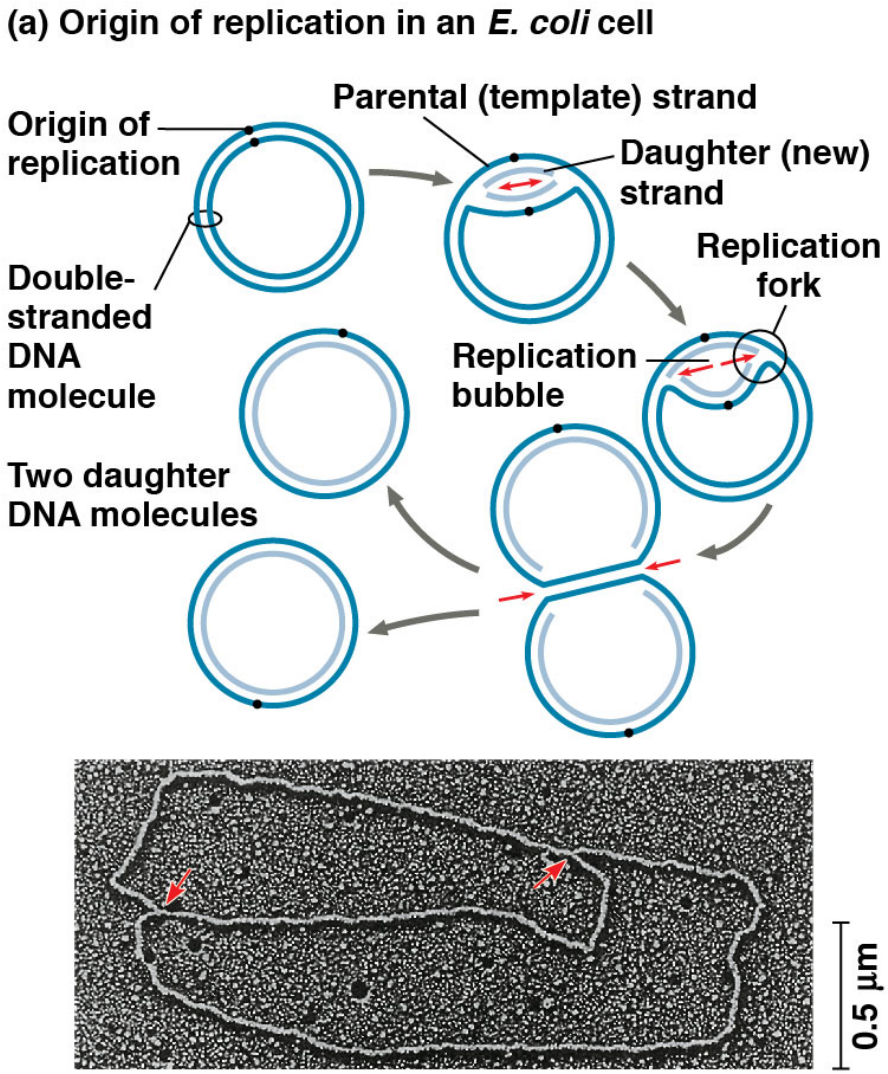
http://www.youtube.com/watch?v=4jtmOZalvS0&feature=related



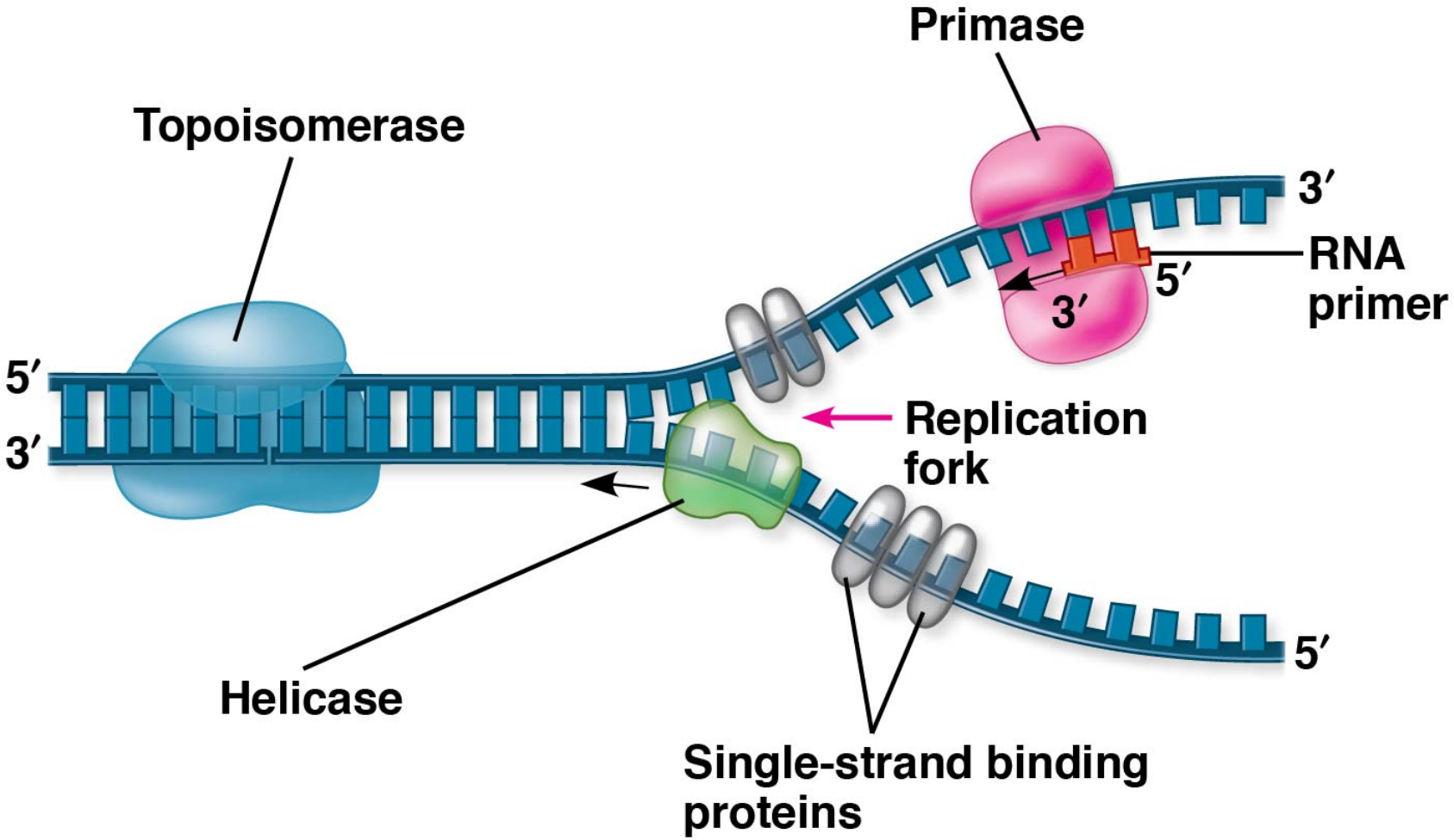
# Major Steps of Replication:

1. **Helicase:** unwinds DNA at *origins of replication*
2. Initiation proteins separate 2 strands → forms *replication bubble*
3. **Topoisomerase:** relieves overwinding strain ahead of replication forks by breaking, swiveling, rejoining DNA strands
4. **Primase:** puts down RNA primer to start replication
5. **DNA polymerase III:** adds complimentary bases to *leading strand* (new DNA is made 5' → 3')
6. *Lagging strand* grows in 3'→5' direction by the addition of *Okazaki fragments*
7. **DNA polymerase I:** replaces RNA primers with DNA
8. **DNA ligase:** seals fragments together

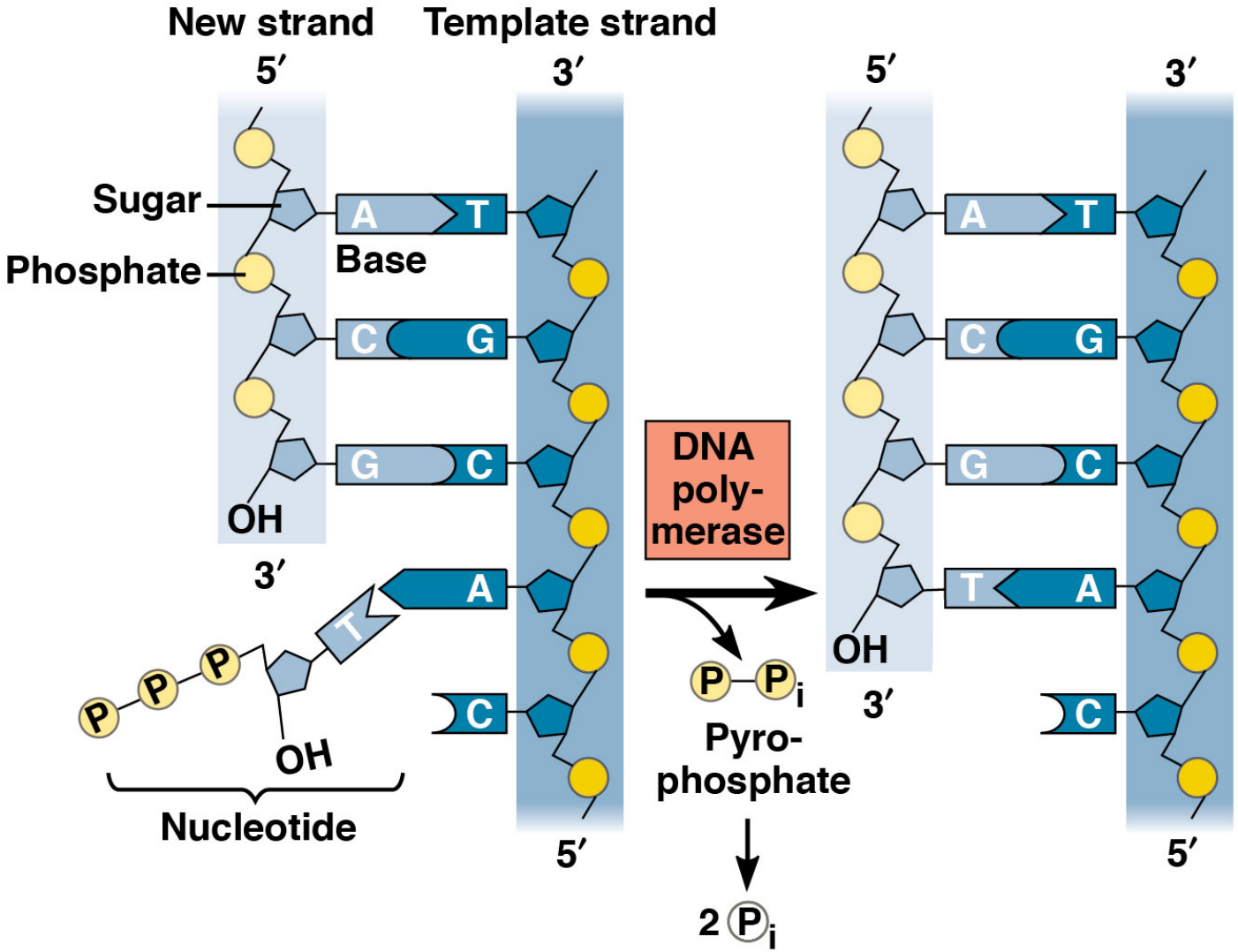
# 1. Helicase unwinds DNA at *origins of replication* and creates *replication forks*



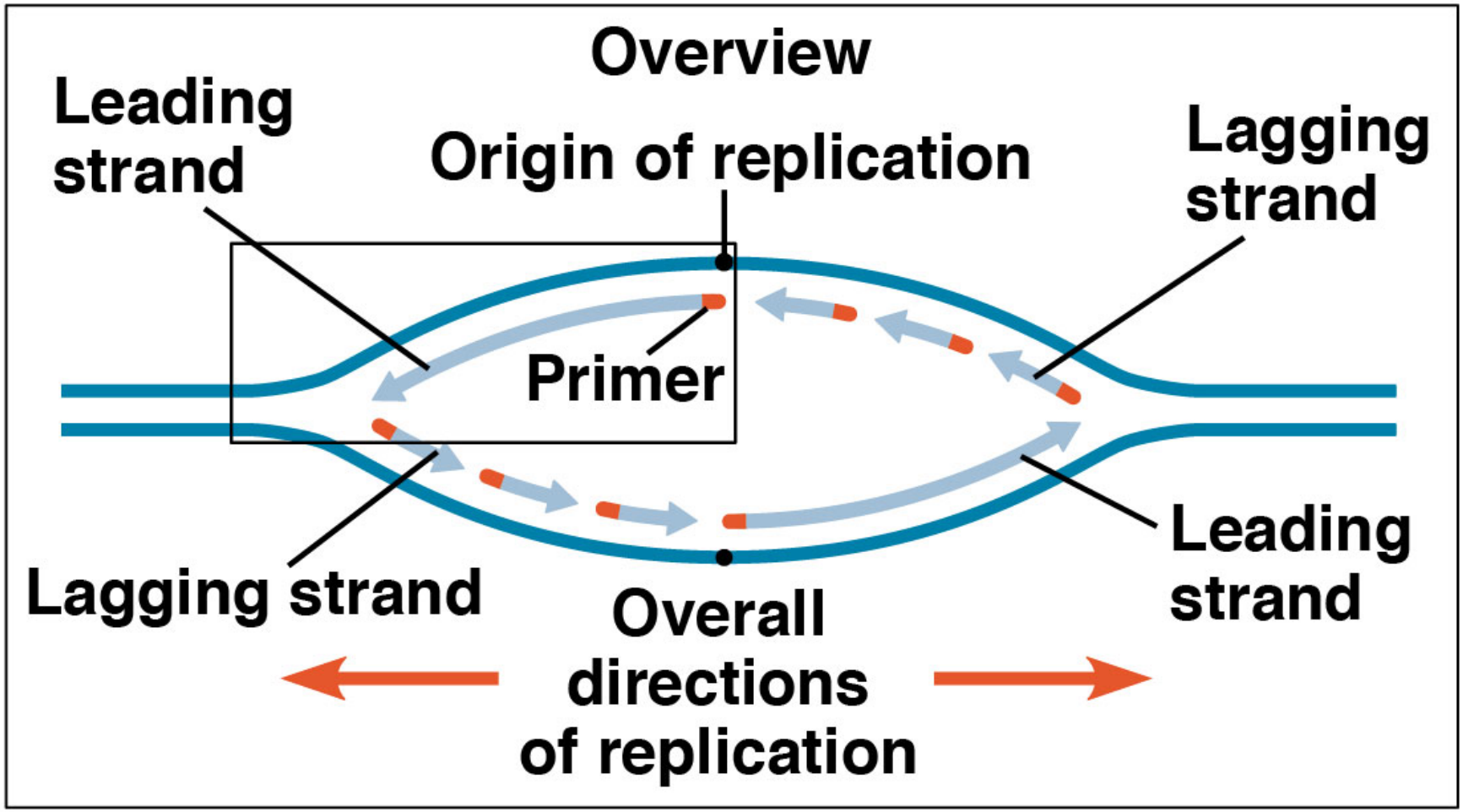
# 4. Primase adds *RNA primer*



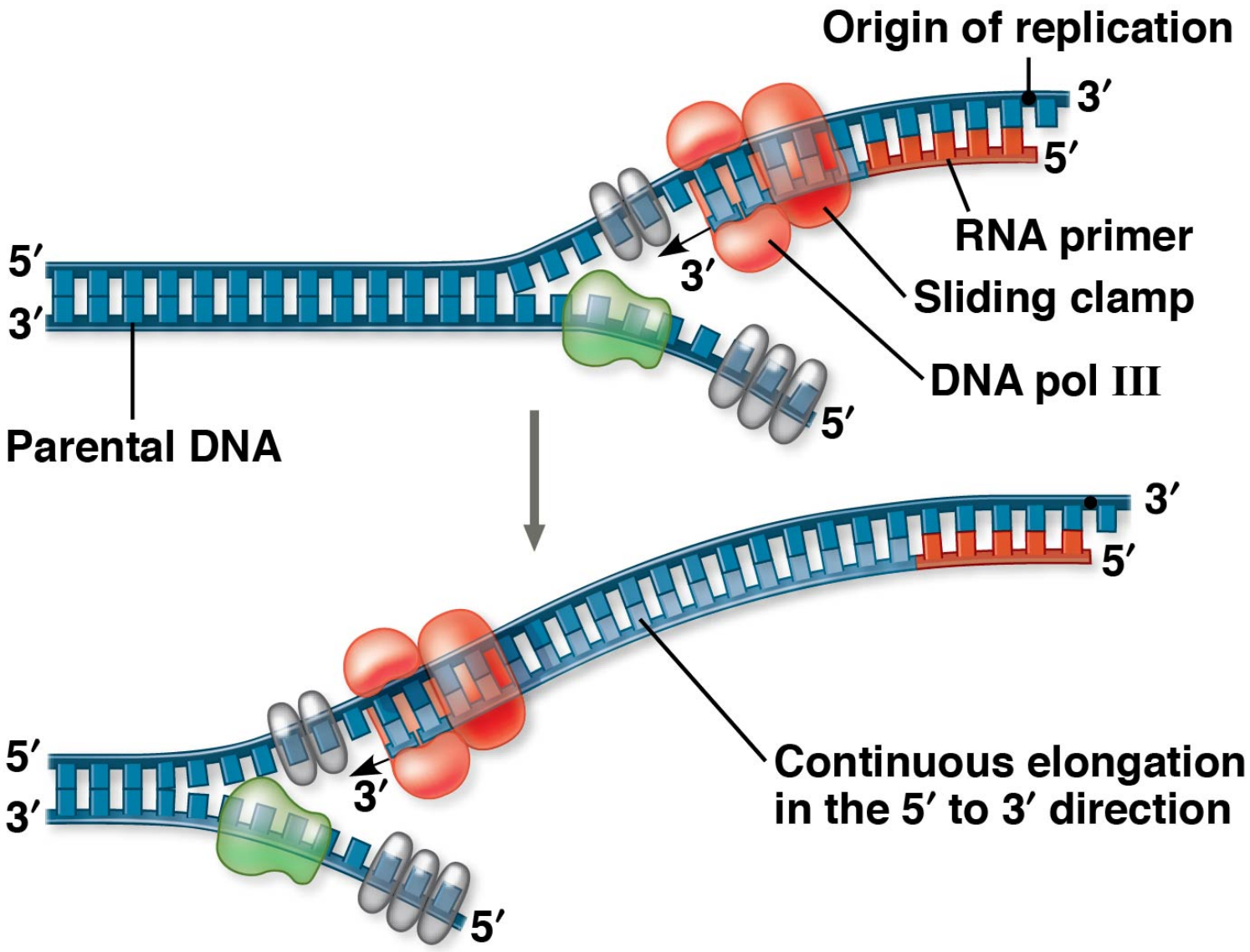
# 5. DNA polymerase III adds nucleotides in 5'→3' direction on *leading strand*



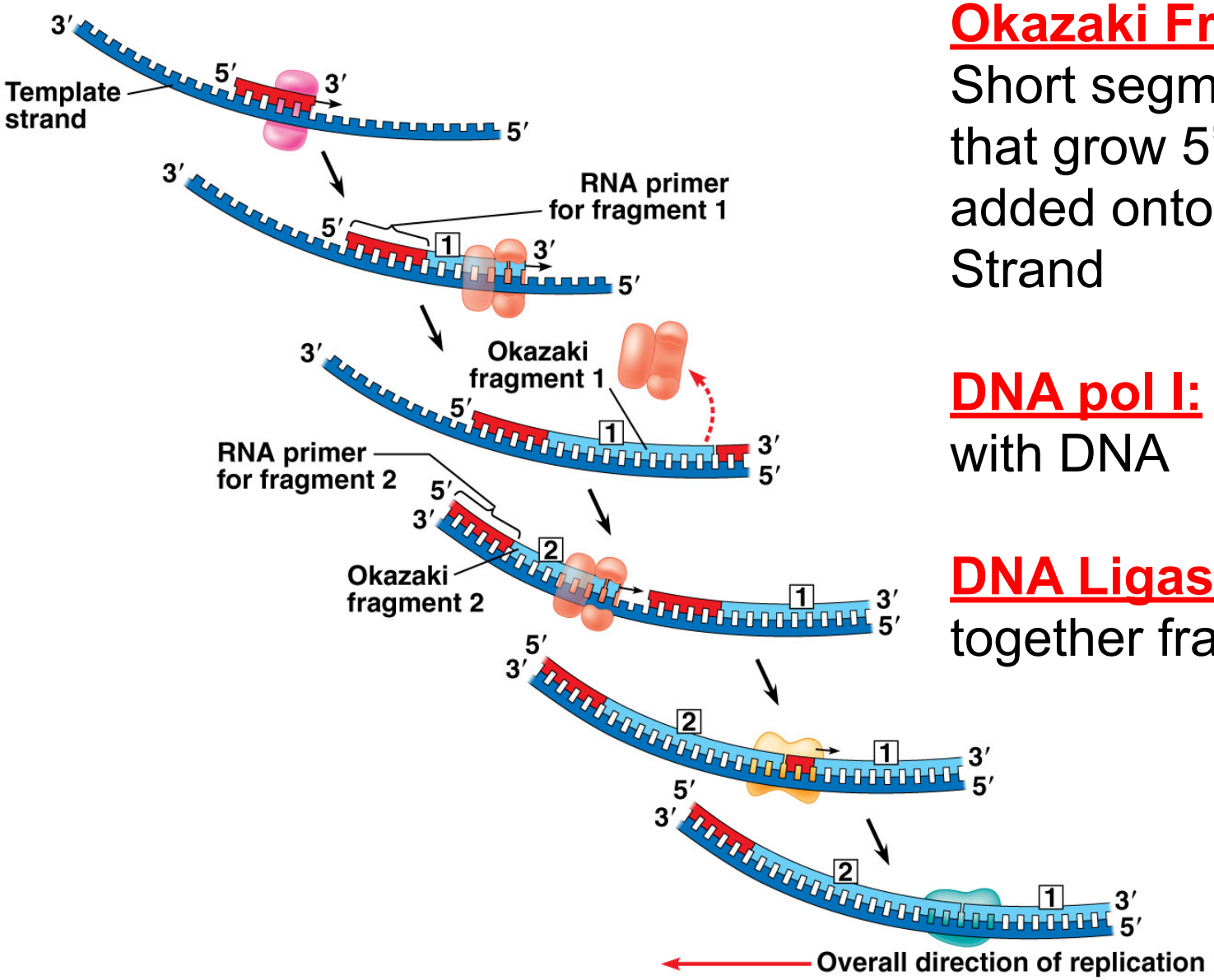
# *Leading strand* vs. *Lagging strand*



# Replication on leading strand



# Replication on lagging strand

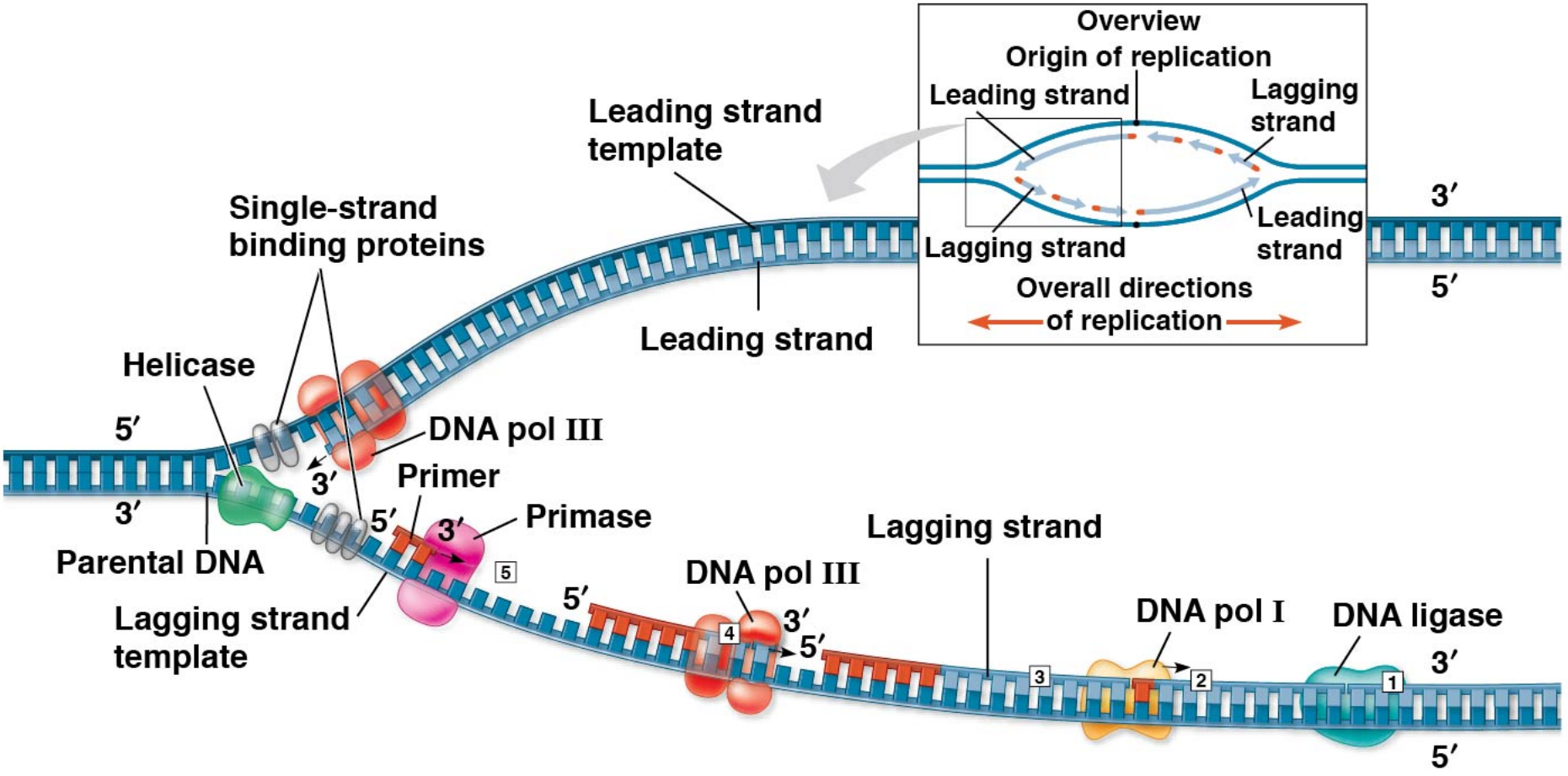



**Okazaki Fragments**: Short segments of DNA that grow 5'→3' that are added onto the Lagging Strand

**DNA pol I:** replace RNA with DNA

**DNA Ligase**: seals together fragments

# Summary of DNA Replication



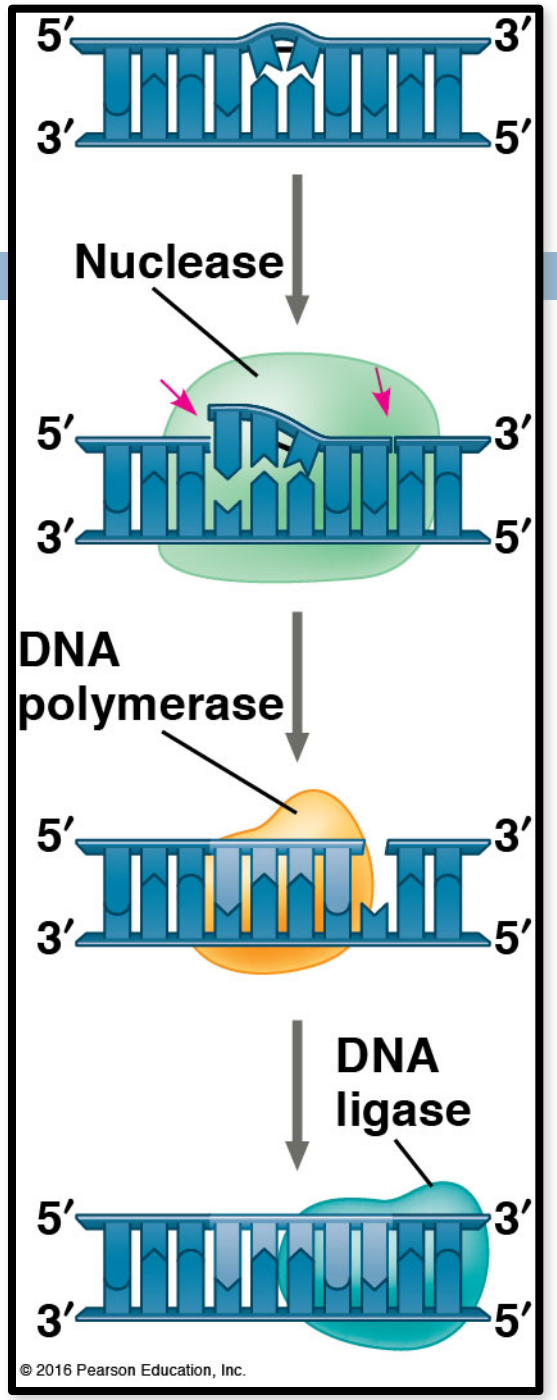
# Proofreading and Repair

- DNA polymerases proofread as bases added
- Errors:
  - Pairing errors: 1 in 100,000 nucleotides
  - Complete DNA: 1 in 10 billion nucleotides

- Mismatch repair: special enzymes fix incorrect pairings

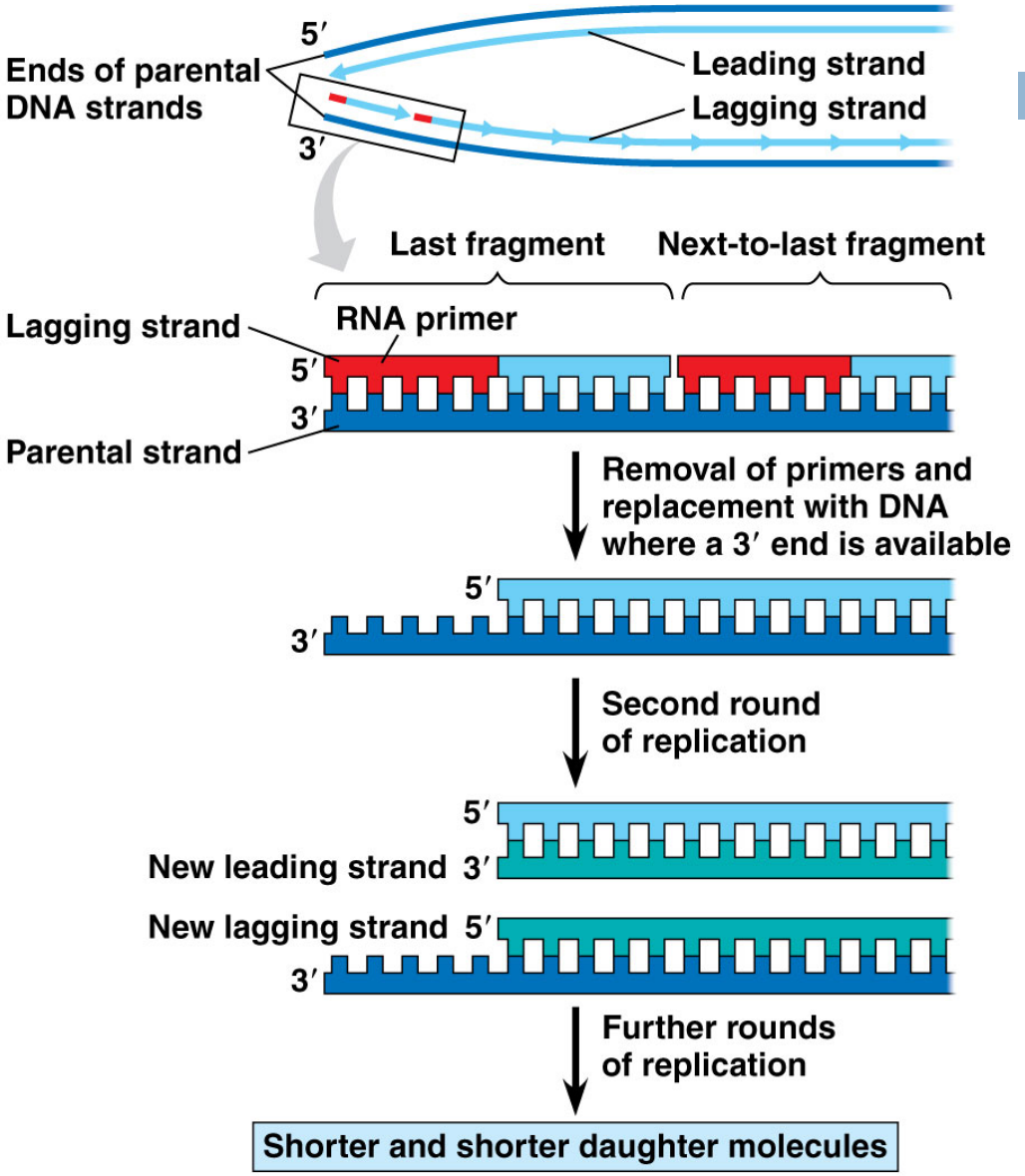
# Nucleotide Excision Repair

- **Nucleases** cut damaged DNA
- DNA poly and ligase fill in gaps



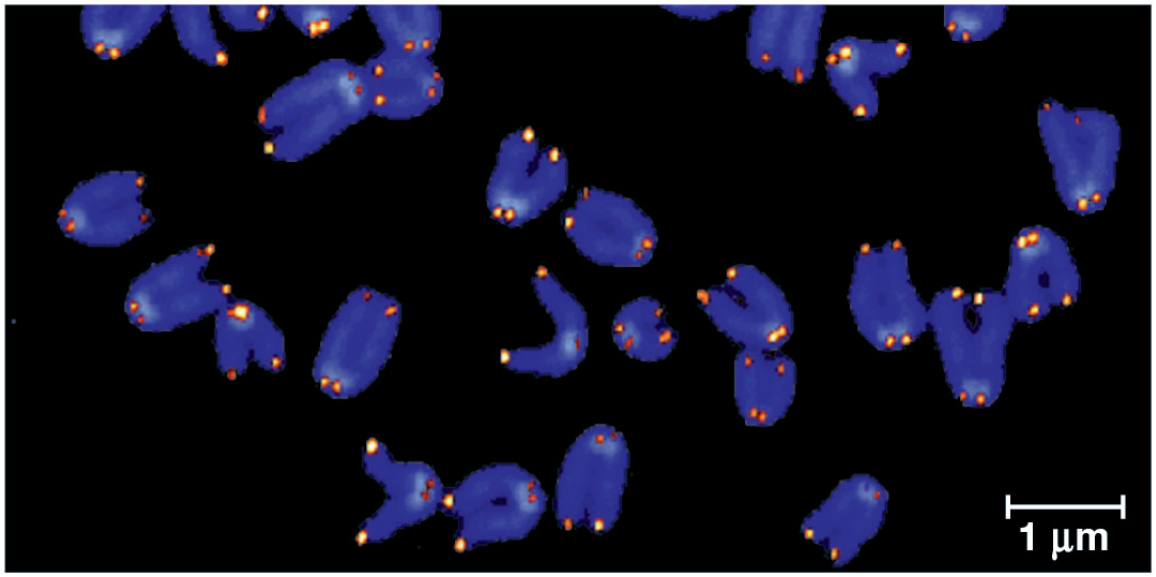
# Problem at the 5’ End

- DNA poly only adds nucleotides to 3’ end
- No way to complete 5’ ends of daughter strands
- Over many replications, DNA strands will grow shorter and shorter



# **Telomeres**: repeated units of short nucleotide sequences (TTAGGG) at ends of DNA

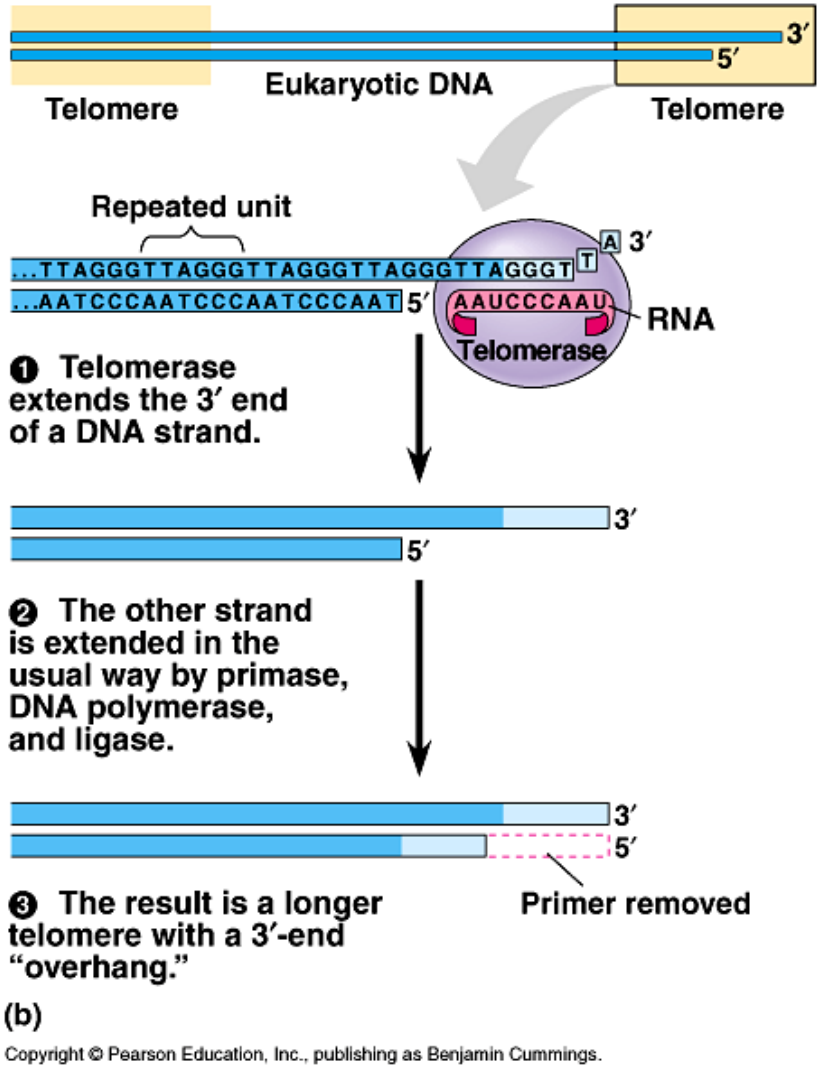
- Telomeres "cap" ends of DNA to postpone erosion of genes at ends (TTAGGG)
- **Telomerase**: enzyme that adds to telomeres
  - Eukaryotic germ cells, cancer cells






Telomeres stained orange at the ends of mouse chromosomes

# Telomeres & Telomerase



# BioFlix: DNA Replication

http://media.pearsoncmg.com/bc/bc_0media_bio/bioflix/bioflix.htm?8apdnarep