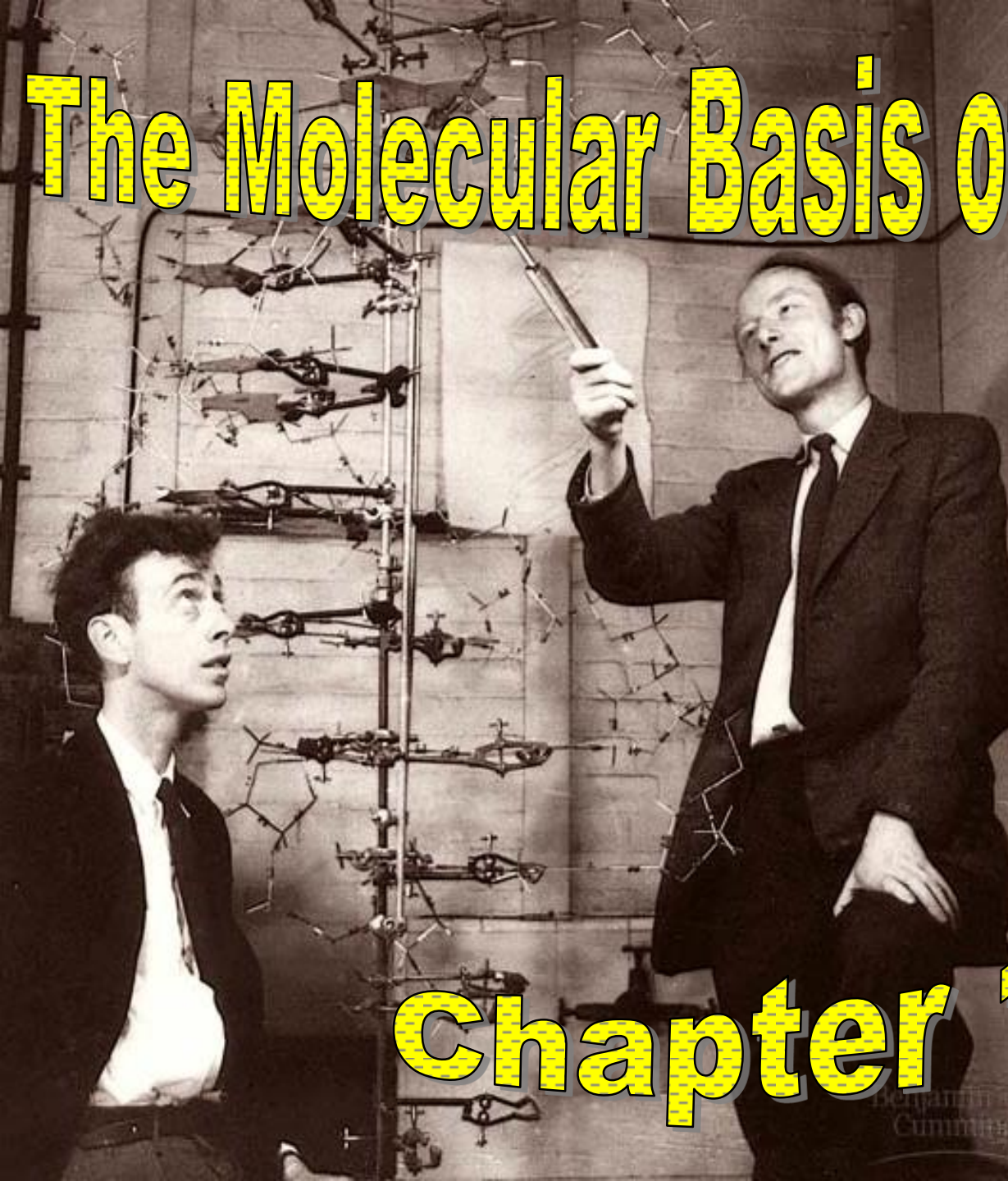
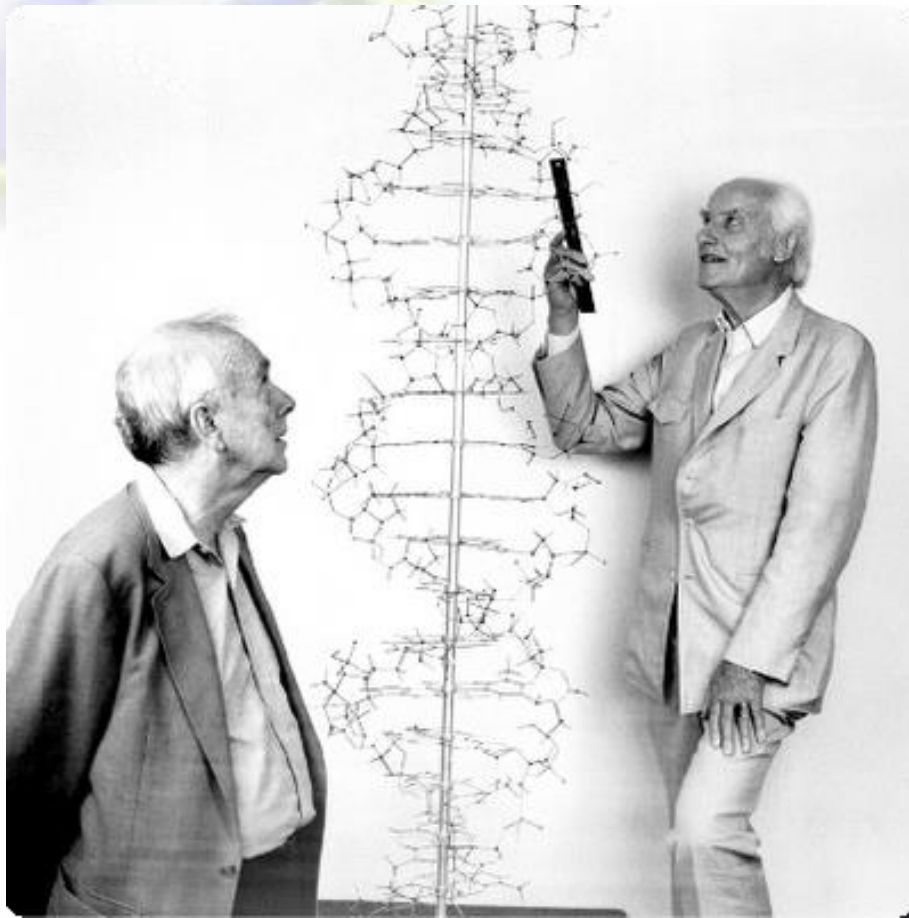


The Molecular Basis of Inheritance



Chapter 16



James D. Watson

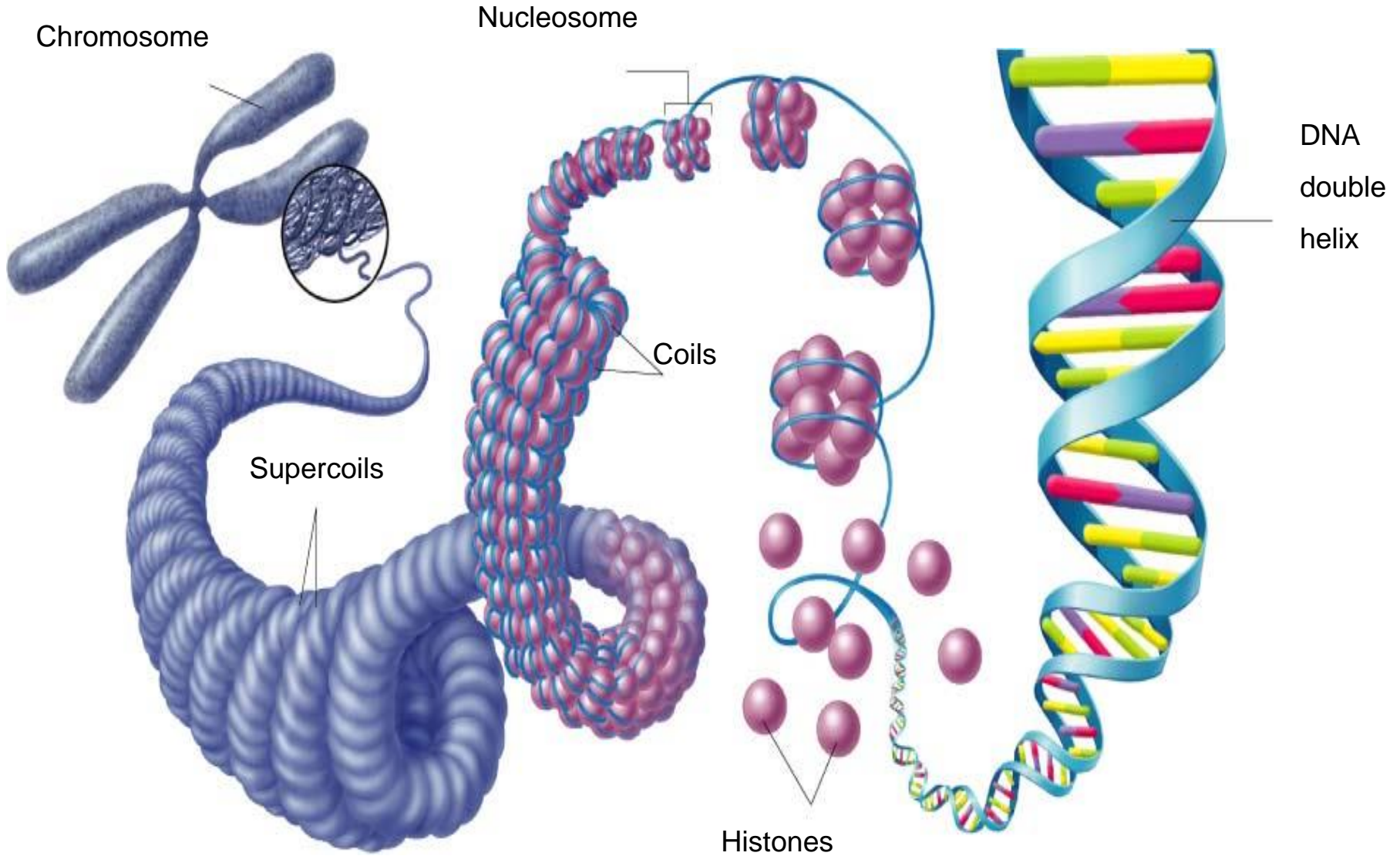
Unnumbered 8 p177
Laboratory Principles of Biochemistry, 6th Edition
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**Francis Crick,
1916–2004**

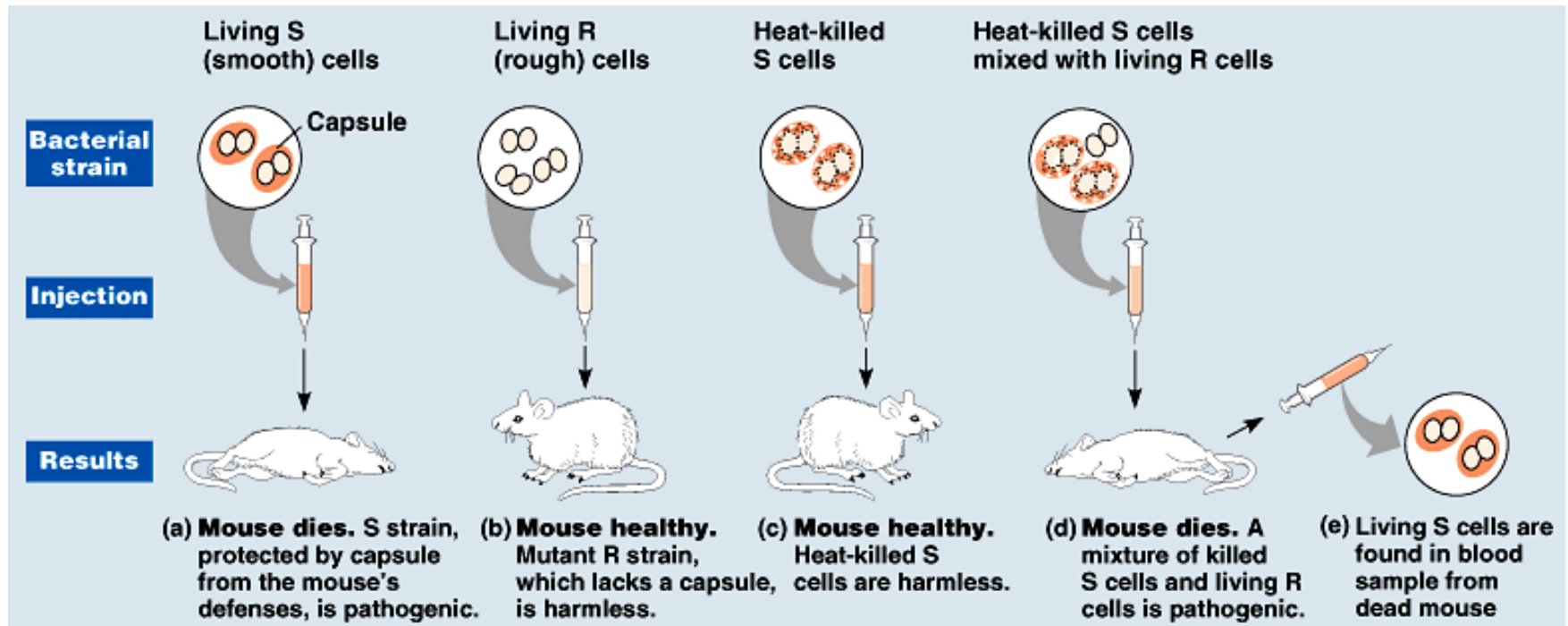


Chromosomes



Evidence That DNA Can Transform Bacteria

- Frederick Griffith's experiment 1928
- Griffith called the phenomenon *transformation* – a change in genotype and phenotype due to the assimilation of external DNA by a cell.



Conclusion:

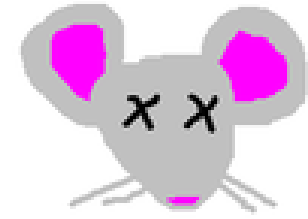
Of all the molecules tested, only DNA changed the nonvirulent bacteria to become virulent; DNA can change a bacteria's genotype, thus changing its phenotype
transformation

Evidence That DNA Can Transform Bacteria

- Avery, MacLeod, and McCarty
- Used a similar experiment to Griffith's
- One extra step was to take isolated DNA from the heat killed virulent bacteria and mixed it with non-encapsulated bacteria, which killed the mouse
- This confirmed that DNA was the material that transformed the avirulent bacteria

Procedures:

1. The encapsulated, virulent bacteria is injected into the mouse. The mouse dies.



2. The non-encapsulated, avirulent bacteria is injected into the mouse. The mouse lives.



3. The heat-killed, virulent bacteria is injected into the mouse. The mouse lives.








4. The heat-killed, virulent bacteria is mixed with the non-encapsulated avirulent bacteria and injected into the mouse. The mouse dies.



5. The DNA extracted from the heat-killed, virulent bacteria is mixed with the non-encapsulated avirulent bacteria and injected into the mouse. The mouse dies.

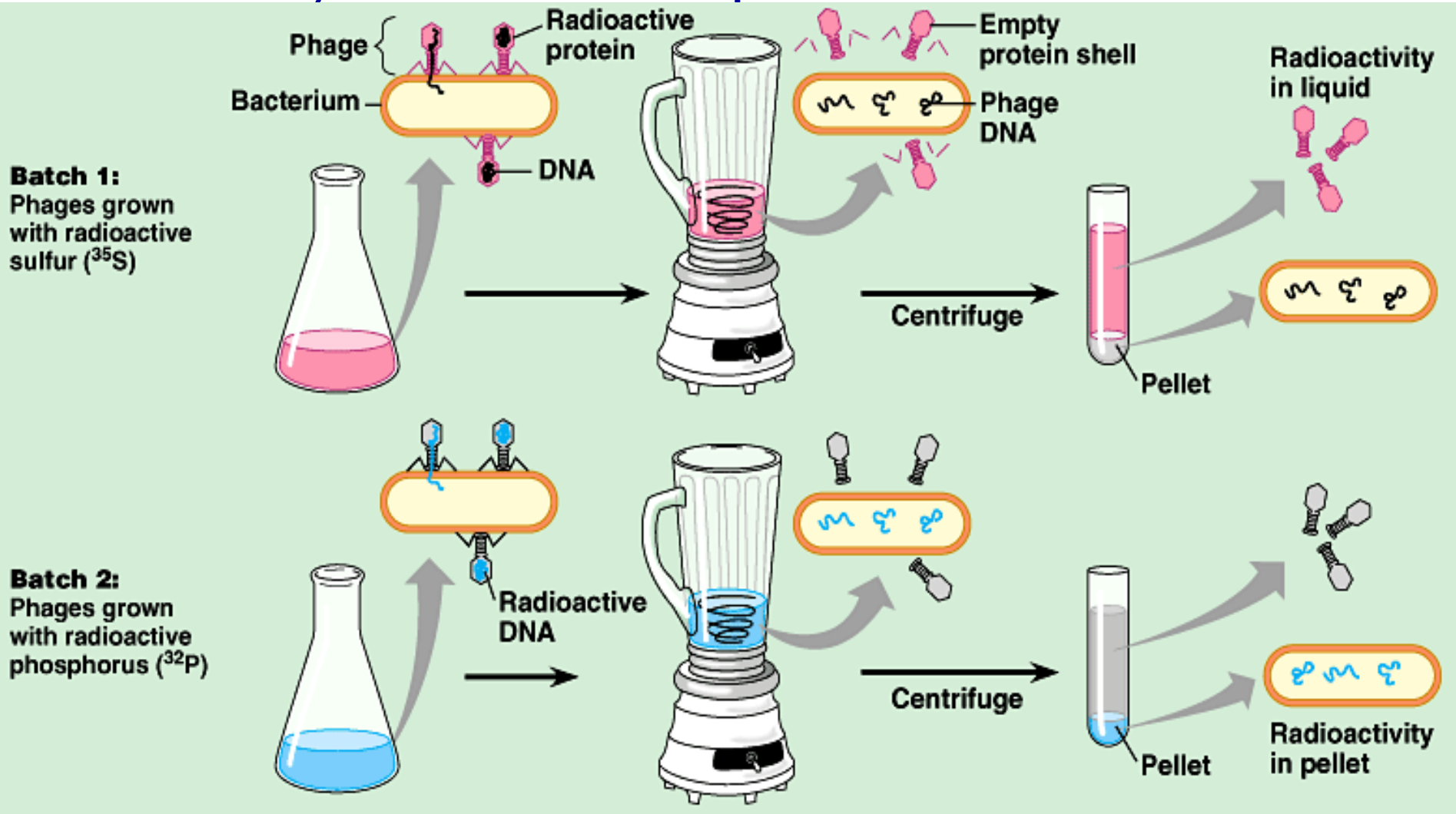


Legend:

-  encapsulated, virulent bacteria
-  heat-killed, virulent bacteria
-  DNA extracted from 
-  non-encapsulated, avirulent bacteria

Evidence That DNA Can Transform Bacteria

- Hershey and Chase experiment 1952



Conclusion:

Injected viral DNA provides genetic information that makes the cells produce more viruses

Evidence that nucleic acids, not proteins were the genetic material

Chargaff's Experiment



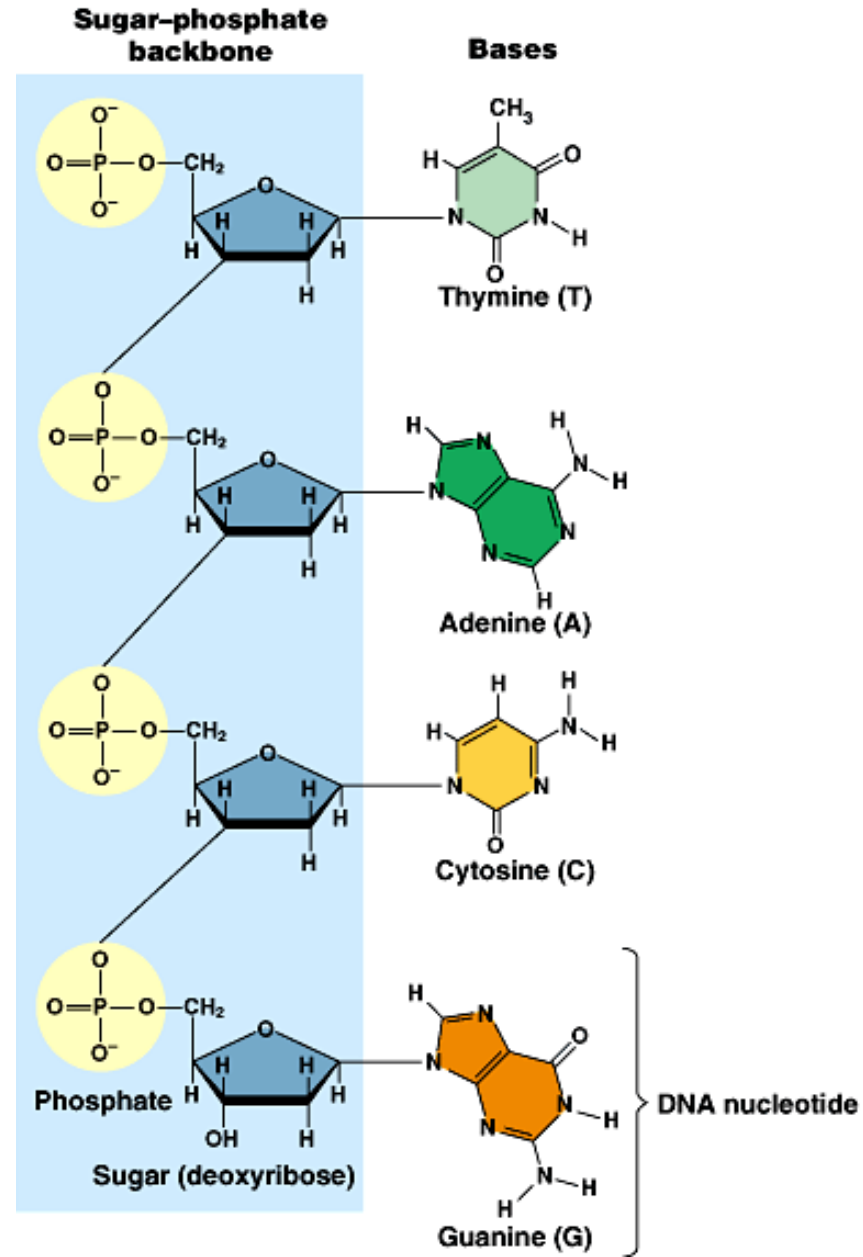
- The composition of DNA varied from one species to another, in particular in the relative amounts of the bases A, C, T, G
- In any DNA, the number of A's was the same as the number of T's; number of C's was equal to the number of G's
- What did this mean?
Chargaff couldn't say

Additional Evidence

- Diploid sets of chromosomes have twice as much DNA as the haploid sets found in the gametes of the same organism
- Ratio of DNA bases varies from species to species but are all present in the same amount

$$A = T$$

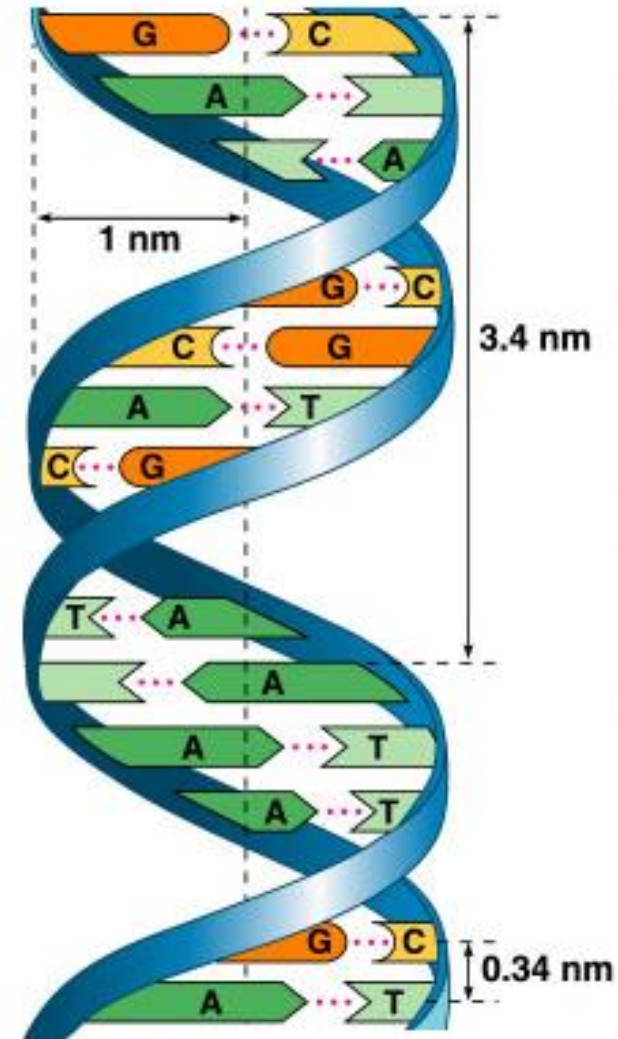
$$C = G$$



James Watson and Francis Crick

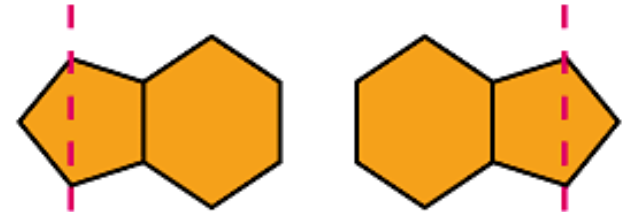
Using Rosalind Franklin's X ray diffraction image of DNA, they deduced:

- Helical shape – double stranded
- 10 layers of base pairs per turn
- One full turn every 3.4 nm
- Width of the helix (suggest 2 strands)
- Spacing of the nitrogenous bases

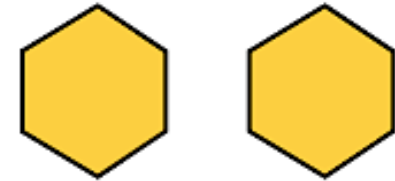


James Watson and Francis Crick

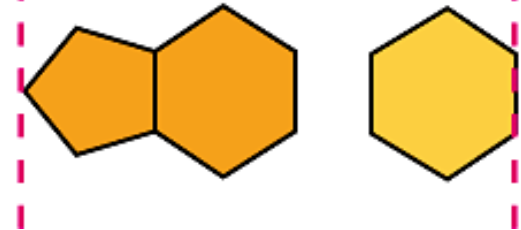
Purine + purine: too wide



Pyrimidine + pyrimidine: too narrow



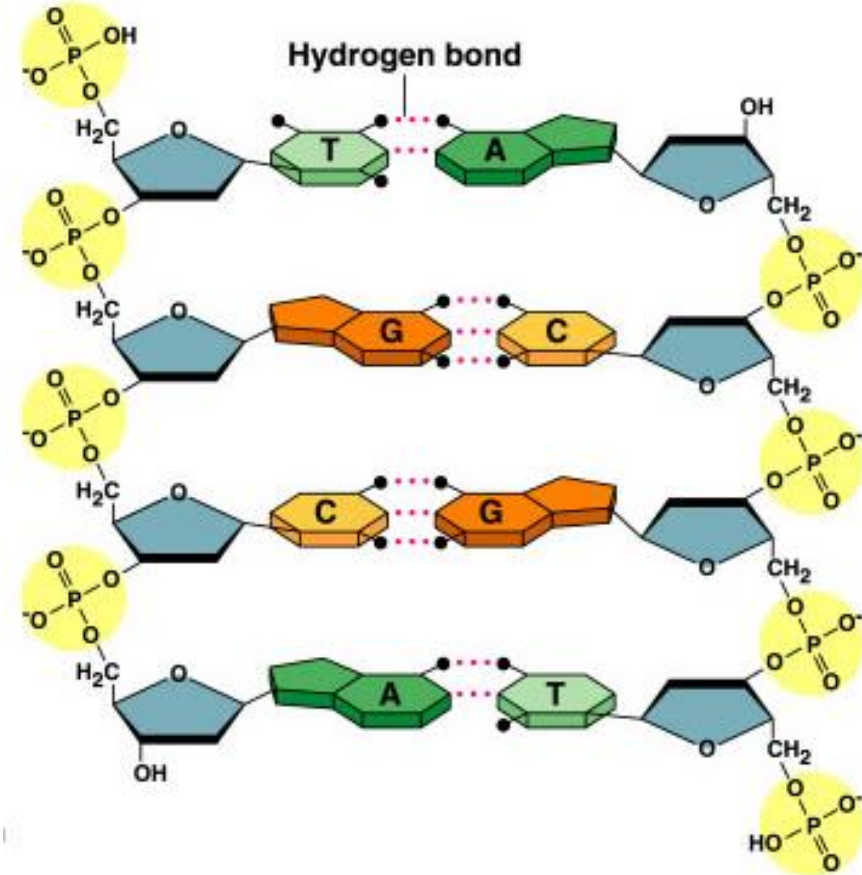
Purine + pyrimidine: width consistent with X-ray data



James Watson and Francis Crick

Structure of DNA

- Purines (Adenine & Guanine) always bond with pyrimidines (Thymine & Cytosine)
- Adenine forms 2 hydrogen bonds with thymine
- G forms 3 hydrogen bonds with cytosine

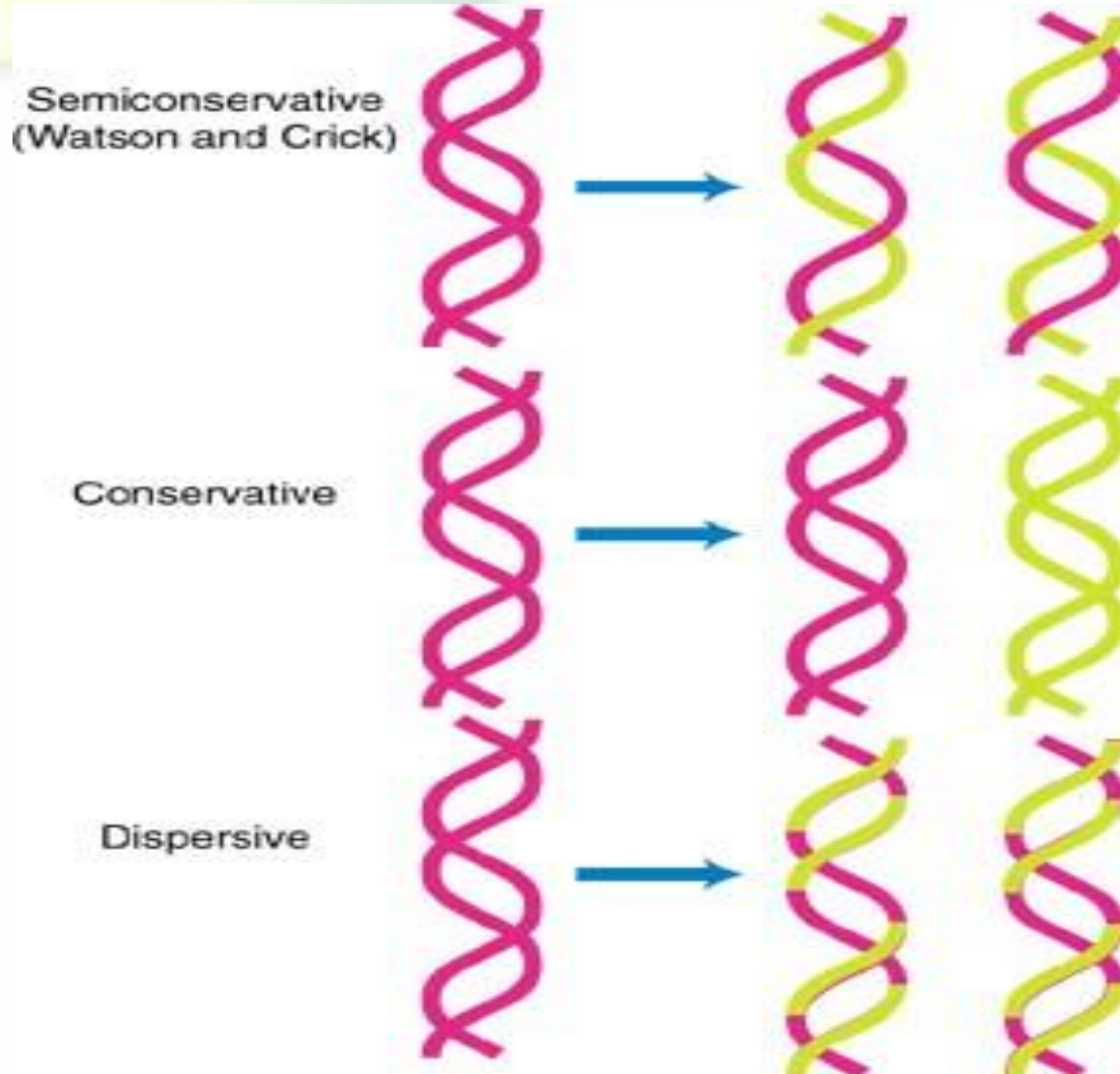




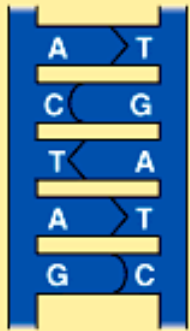
How is DNA Replicated?

Conservative,
Semiconservative or Dispersive?

3 Hypotheses for Replication



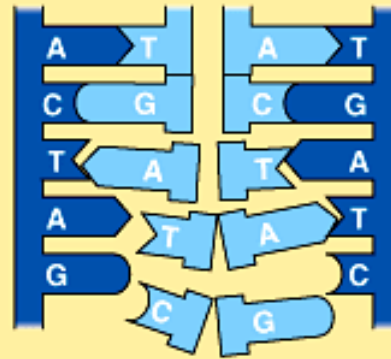
Semi-conservative DNA Replication



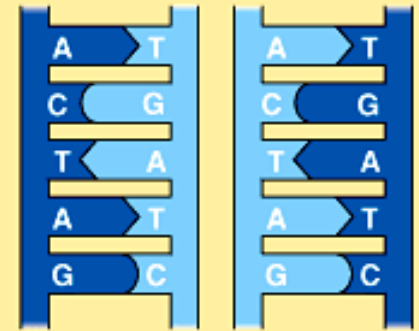
(a) The parent molecule has two complementary strands of DNA. Each base is paired by hydrogen bonding with its specific partner, A with T and G with C.



(b) The first step in replication is separation of the two DNA strands.

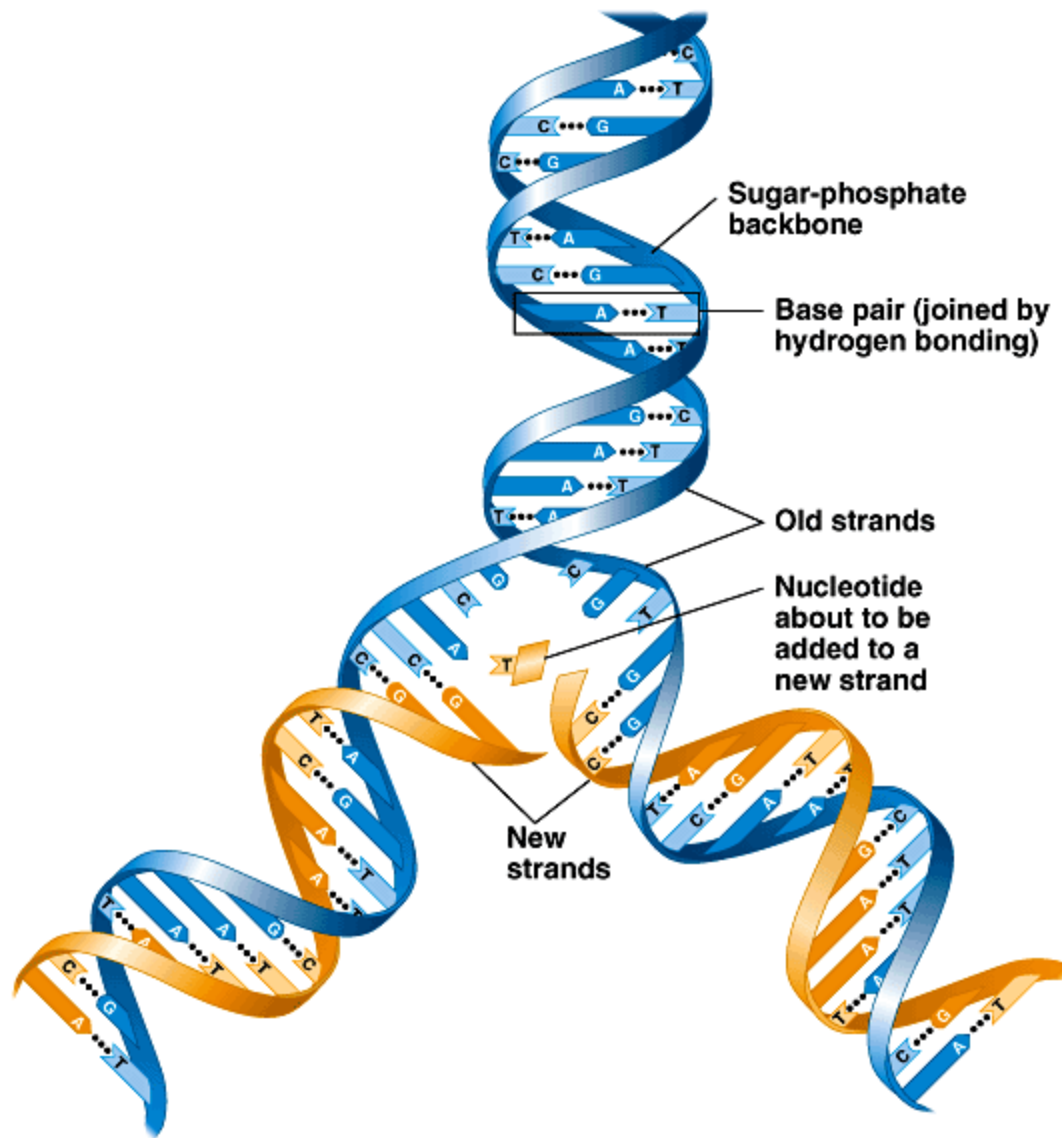


(c) Each parental strand now serves as a template that determines the order of nucleotides along a new complementary strand.



(d) The nucleotides are connected to form the sugar-phosphate backbones of the new strands. Each "daughter" DNA molecule consists of one parental strand and one new strand.

DNA Replication



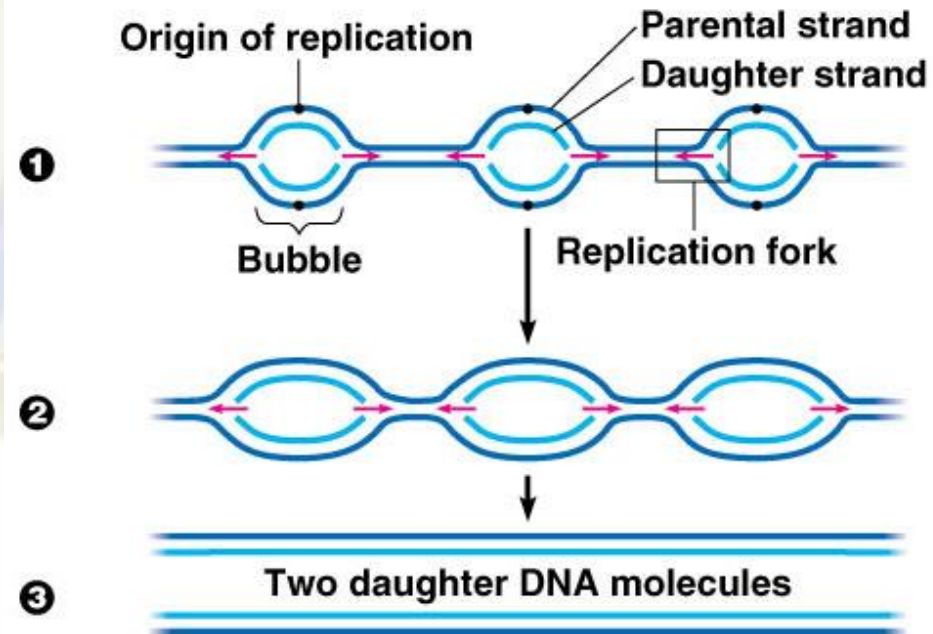
DNA Replication



- A human cell has 6 billion base pairs on 46 chromosomes
- There is one DNA molecule per chromosome
- DNA replication is so exact that there is 1 error per 1 billion nucleotides
- DNA replication utilizes more than a dozen enzymes and other proteins

Origin of Replication

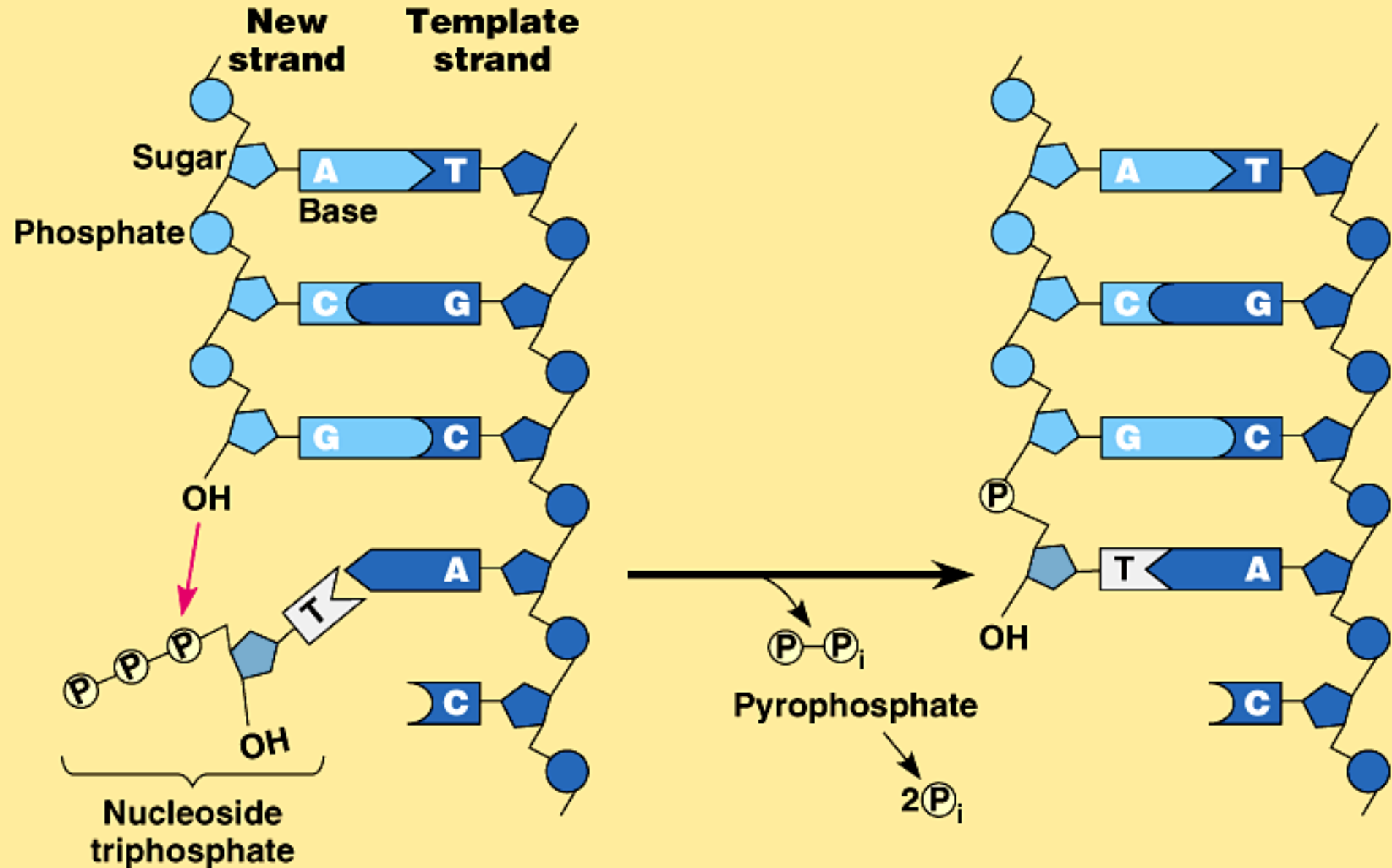
- Proteins recognize the sequence and attach to the DNA
- Replication occurs in both directions and at each end produces a replication fork where the new strand of DNA is elongated



Elongating a New DNA Strand

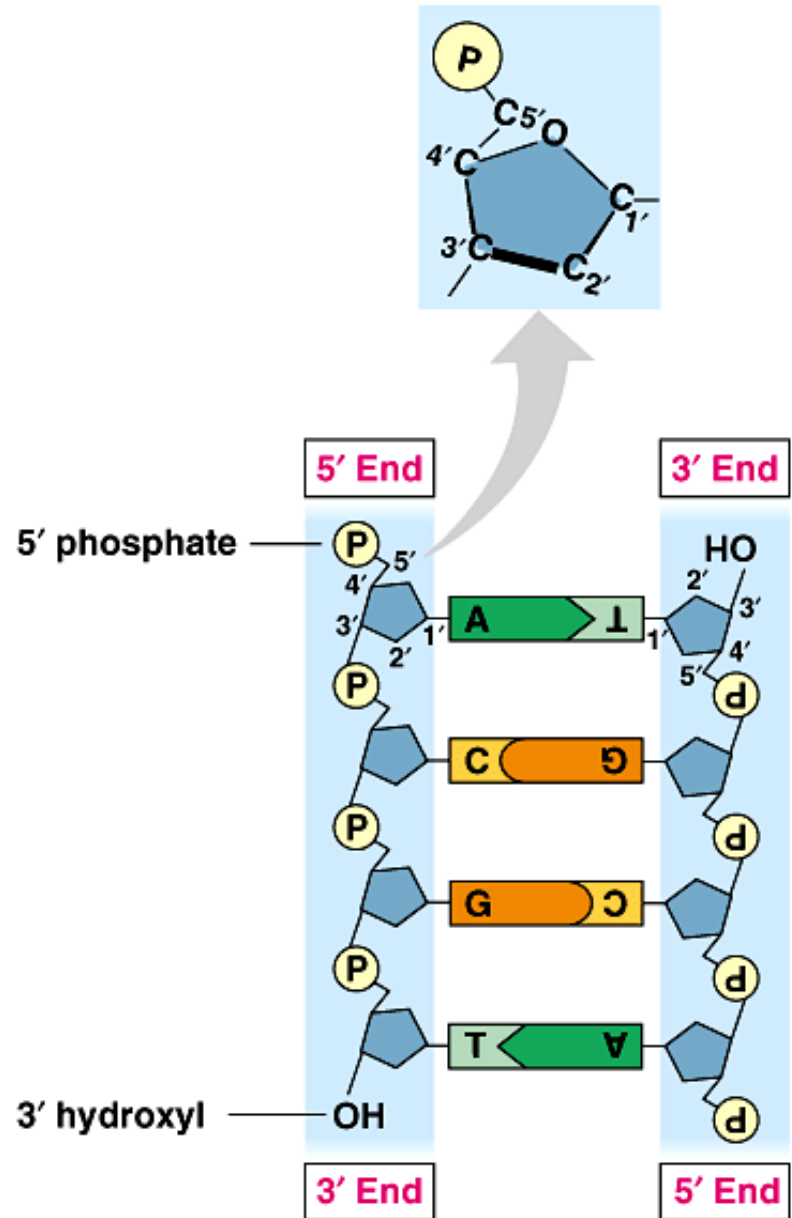
- The enzyme called DNA polymerase catalyses nucleotide attachment
- The rate of elongation is about 50 nucleotides per second in human cells
- Nucleotides that serve as substrates for DNA polymerase are triphosphates (tails have an unstable cluster of negative charges) as monomers join the DNA strand – 2 phosphate groups (called pyrophosphates) hydrolysis to 2 P_i and provides energy

Elongating a New DNA Strand



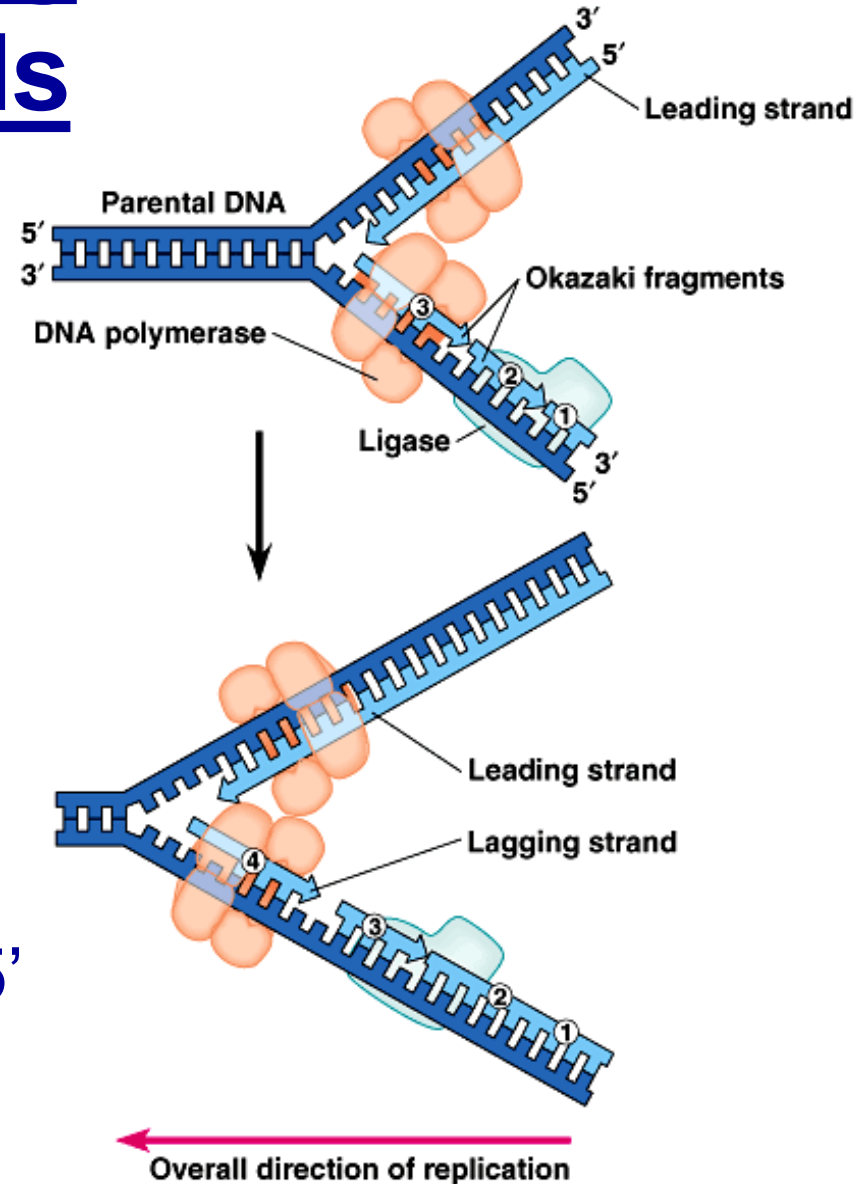
Antiparallel Arrangement

- DNA polymerase adds nucleotides only to the free 3' end
- Elongation occurs 5' → 3'



Synthesis of Leading and Lagging Strands

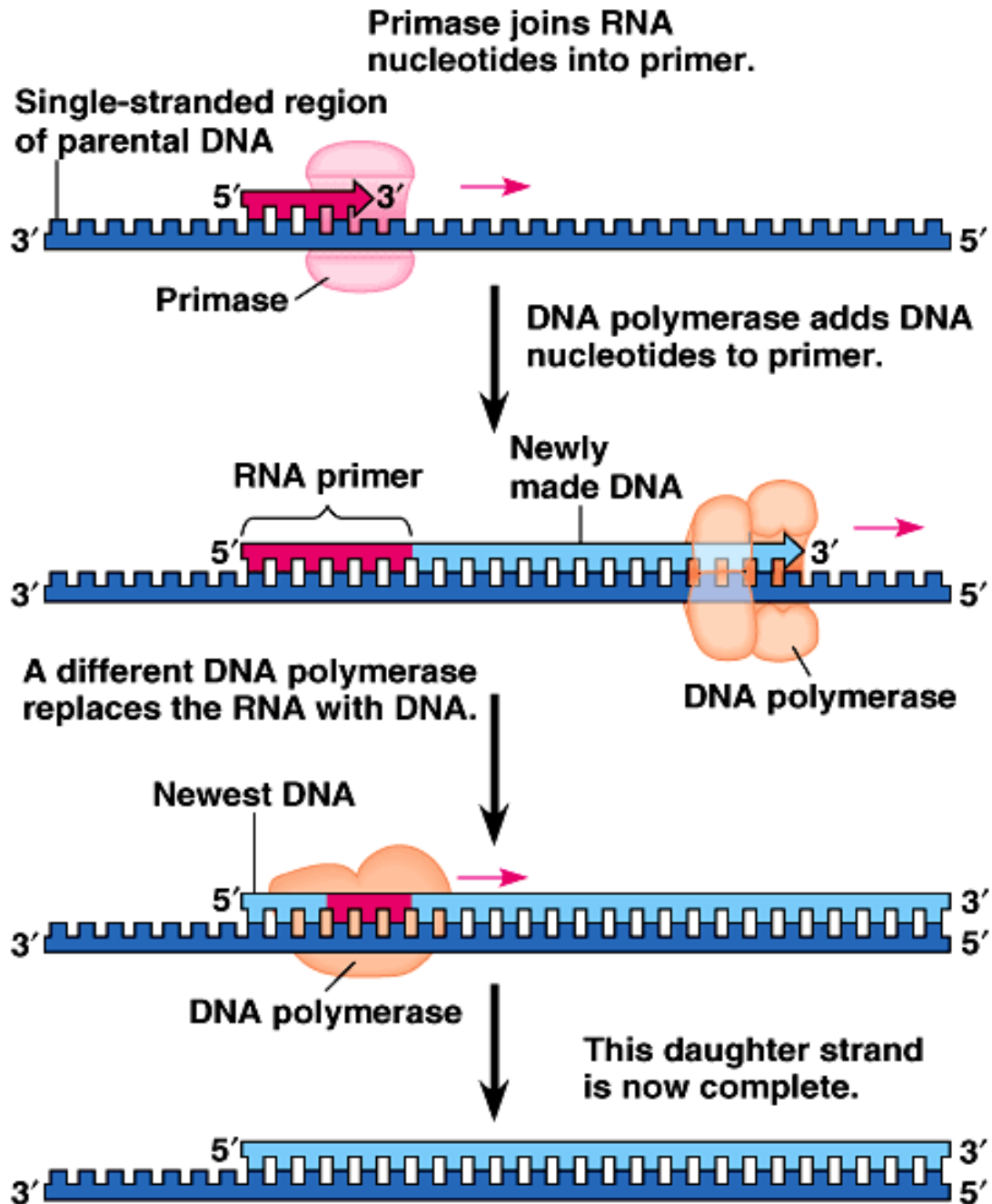
- *Leading Strand* – DNA copied in the 5' → 3' direction
- *Lagging Strand* – DNA copied in the opposite directions away from the replication fork
 - *Okazaki fragments* – short fragments that are synthesized in the 3' → 5' direction
 - Joined together by DNA ligase



Priming DNA Synthesis

- The start of a new chain of DNA is a short stretch of RNA called a *primer*
- The enzyme *primase* joins RNA nucleotides to make the primer
- A different DNA polymerase replaces the RNA primer with DNA

Priming DNA Synthesis

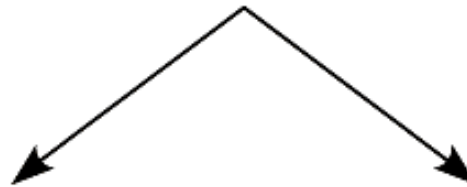


Proteins in DNA Replication

Initiation of replication

Double helix unwinds, providing single-stranded DNA templates

Helicases and single-strand binding proteins



Synthesis of leading strand

Synthesis of lagging strand

Priming

Primase

Priming for Okazaki fragment

Primase

Elongation

DNA polymerase

Elongation of fragment

DNA polymerase

Replacement of RNA primer by DNA

DNA polymerase

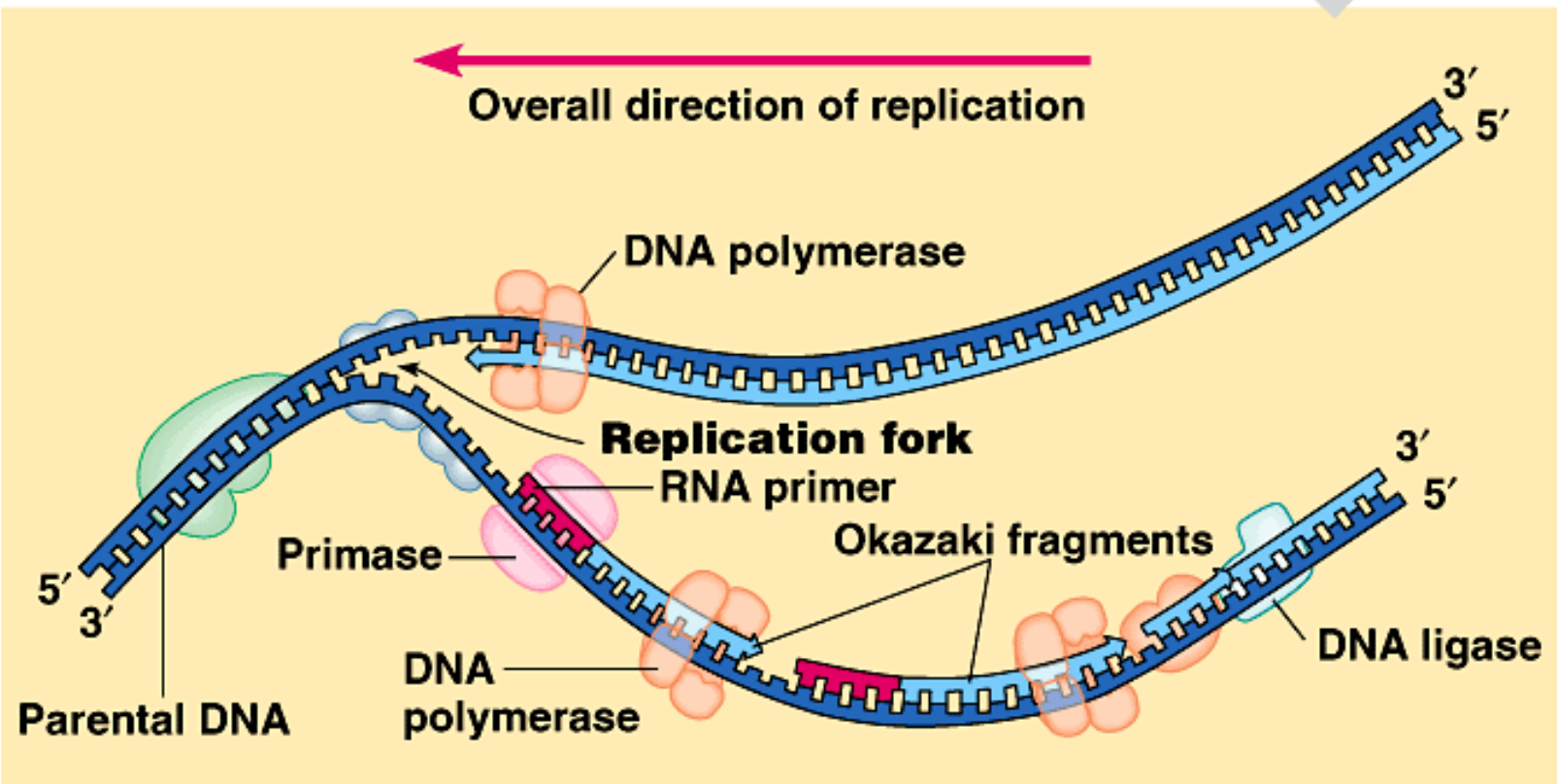
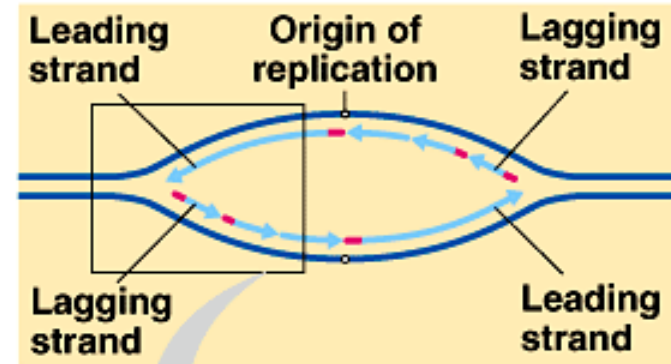
Replacement of RNA primer by DNA

DNA polymerase

Joining of fragments

Ligase

A Summary of DNA Replication

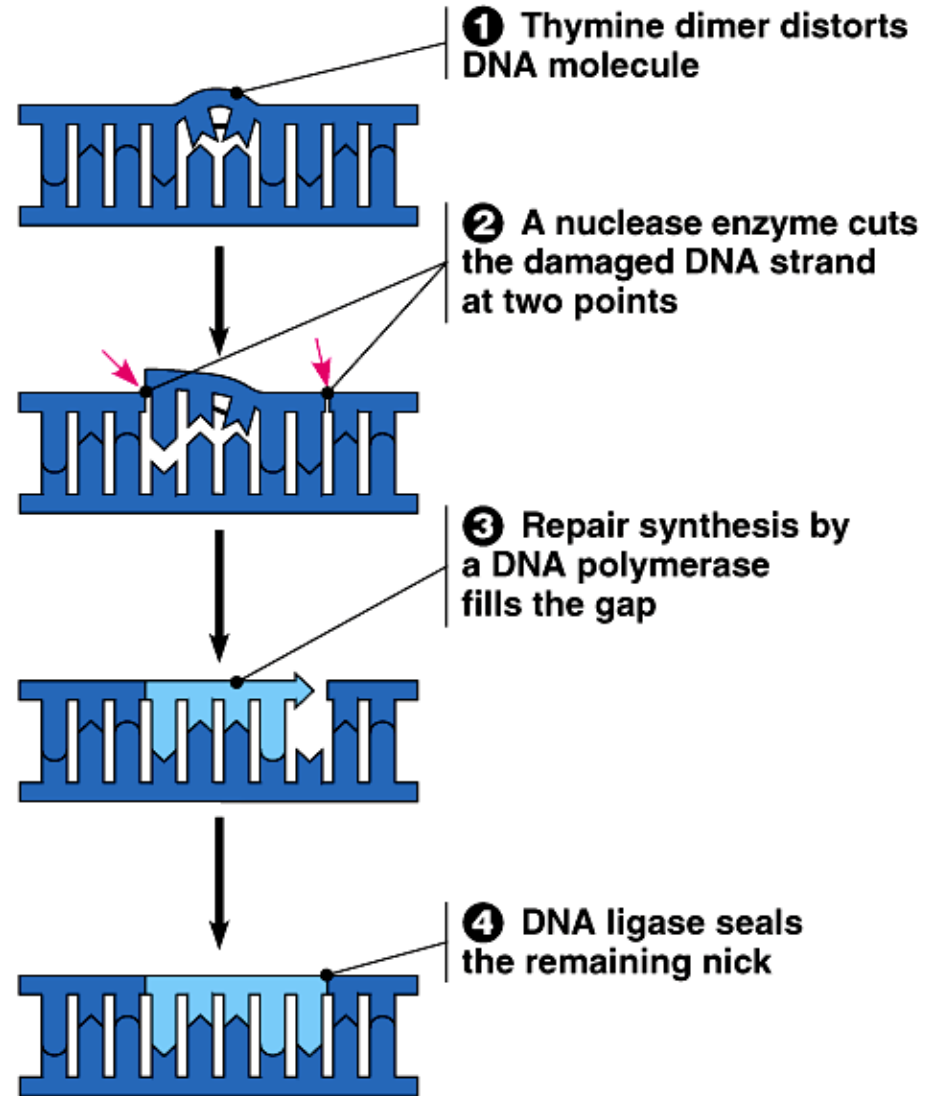


Enzymes Proofread DNA

- DNA polymerase proofreads each nucleotide against its template – if incorrect, it removes and replaces it.
- *Mismatch repairs* – cells use special enzymes to fix incorrectly paired nucleotides.
- DNA requires frequent repair due to environmental factors:
 - Radioactive emissions
 - X-rays
 - UV light
 - Spontaneous chemical changes
- Cells continually monitor and repair its genetic material using 130 enzymes

Enzymes Proofread DNA

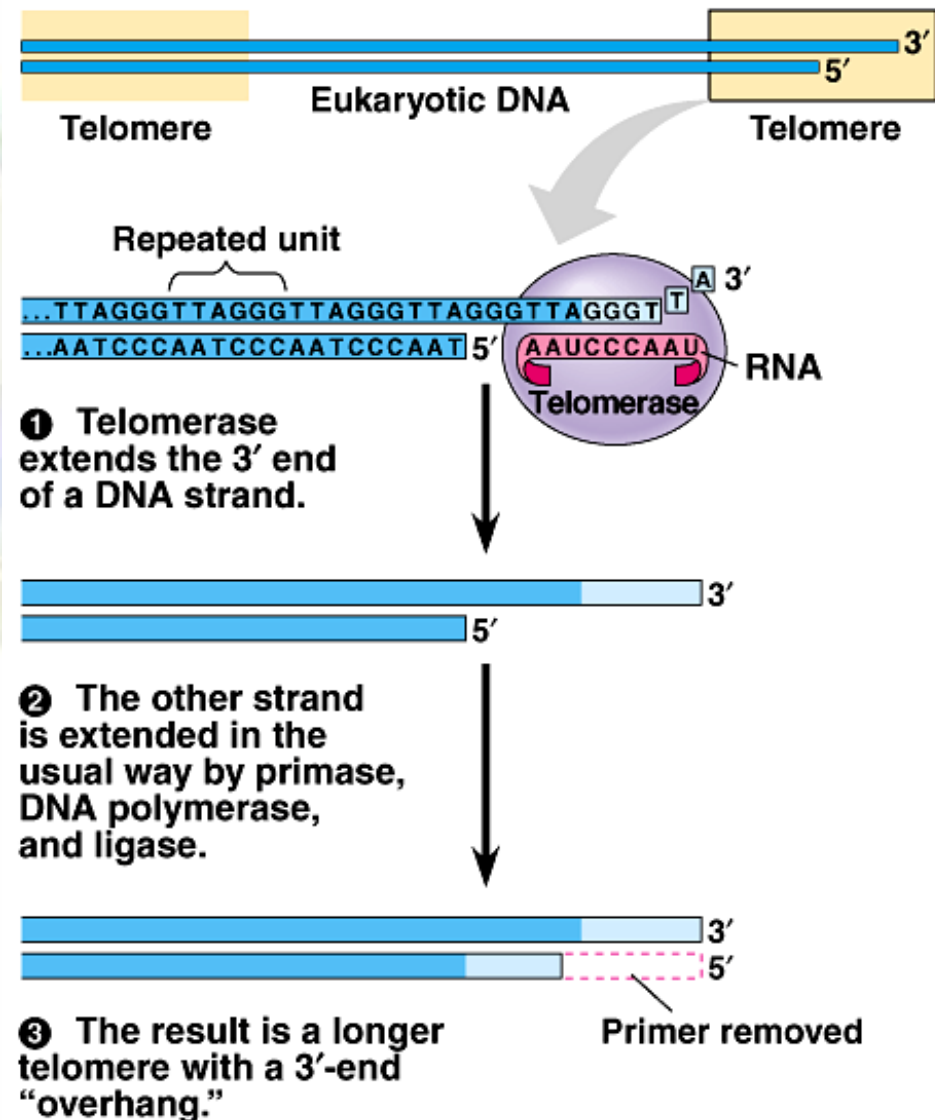
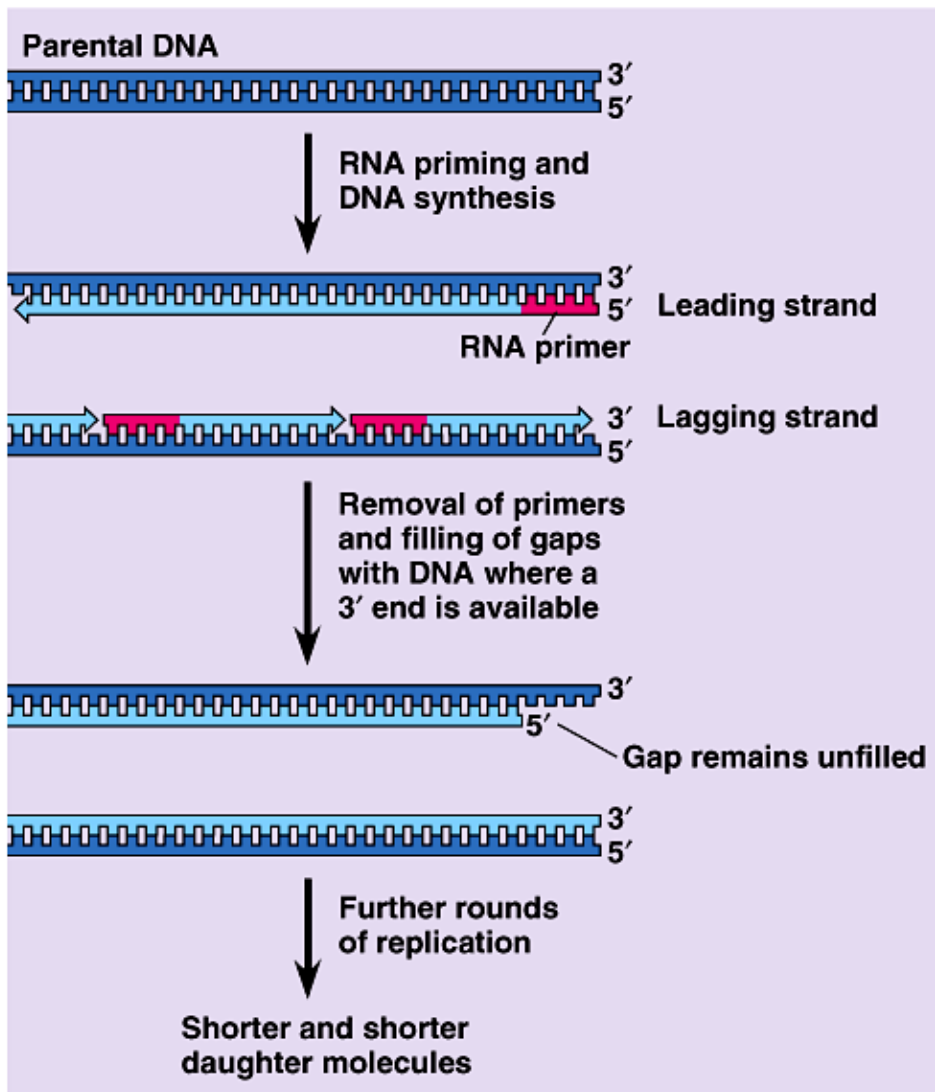
- A *Nuclease* (DNA polymerase and/or ligase) – may cut out a strand of damaged DNA called *nucleotide excision repair*



Ends Are Replicated By a Special Mechanism

- Since there is no way to complete the 5' ends of daughter DNA, replication could produce shorter and shorter strands
- Special sequences called *telomeres* at ends contain multiple repetitions of one short sequence
- Telomeres are not present in most multicellular organisms.
- *Telomerase* – catalyzes lengthening of telomeres: may be a limiting factor in the life span of certain tissues

Ends Are Replicated By Special Mechanism



Telomeres

