



MAMMOTH
SCIENCE



Unit 9 – DNA

- *Identify the components of DNA*
- *Know that traits are determined by proteins that are built according to instructions – coded in DNA*
- *Summarize the process of DNA replication*
- *Understand that enzymes proofread the newly synthesized DNA correcting mistakes*



Menu

History of DNA

Structure

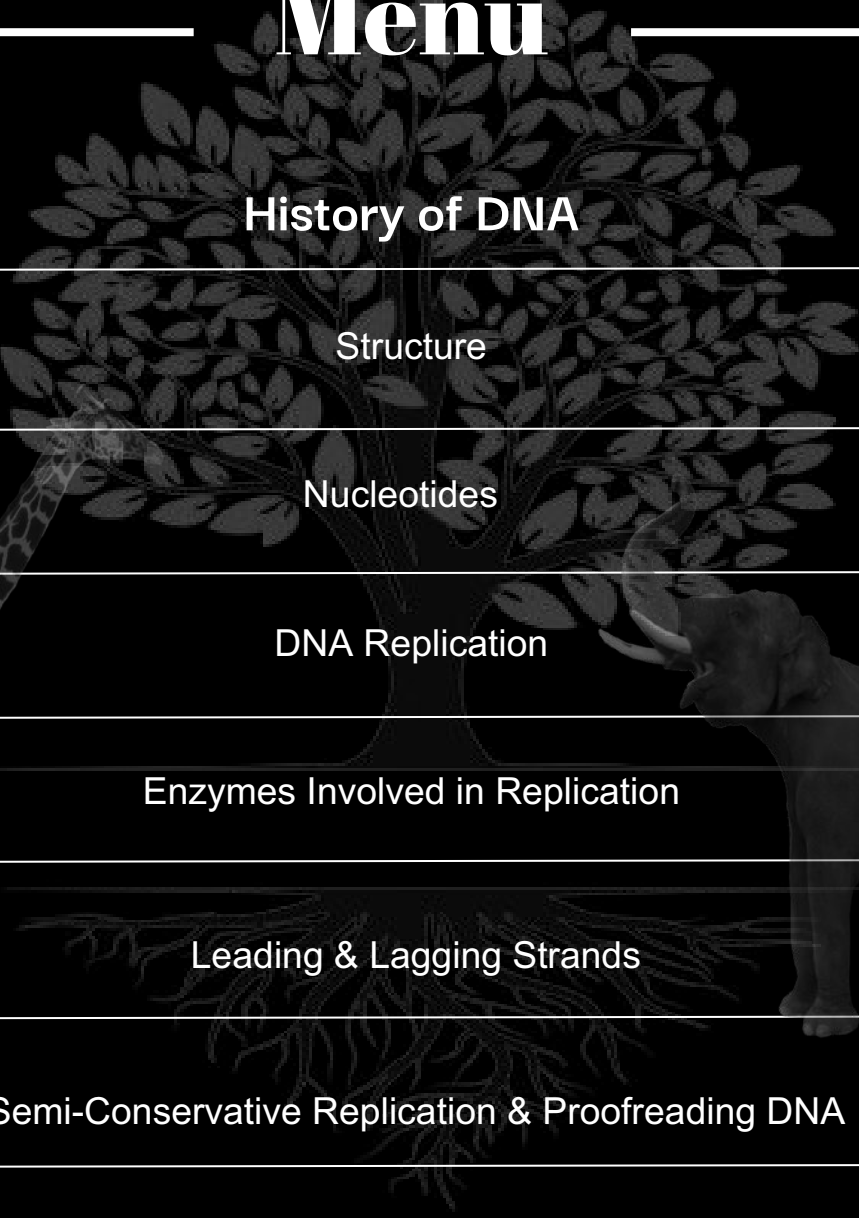
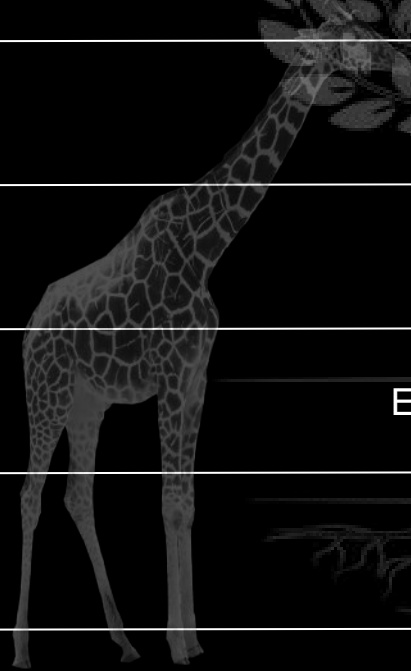
Nucleotides

DNA Replication

Enzymes Involved in Replication

Leading & Lagging Strands

Semi-Conservative Replication & Proofreading DNA



History of DNA



History of DNA

- Early scientists thought protein was the cell's hereditary material because it was more complex than DNA
- **Proteins** were composed of **20 different amino acids** in long **polypeptide** chains
- **Griffith Transformation:**
 - Fred Griffith worked with virulent S and non-virulent R strain Pneumococcus bacteria
 - He found that R strain could become virulent when it took in DNA from heat-killed S strain
 - **Study suggested that DNA was probably the genetic material**
- **Hershey & Chase**
 - **Chromosomes** are made of both **DNA** and **protein**
 - Experiments on bacteriophage viruses by Hershey & Chase proved that **DNA** was the cell's **genetic material**

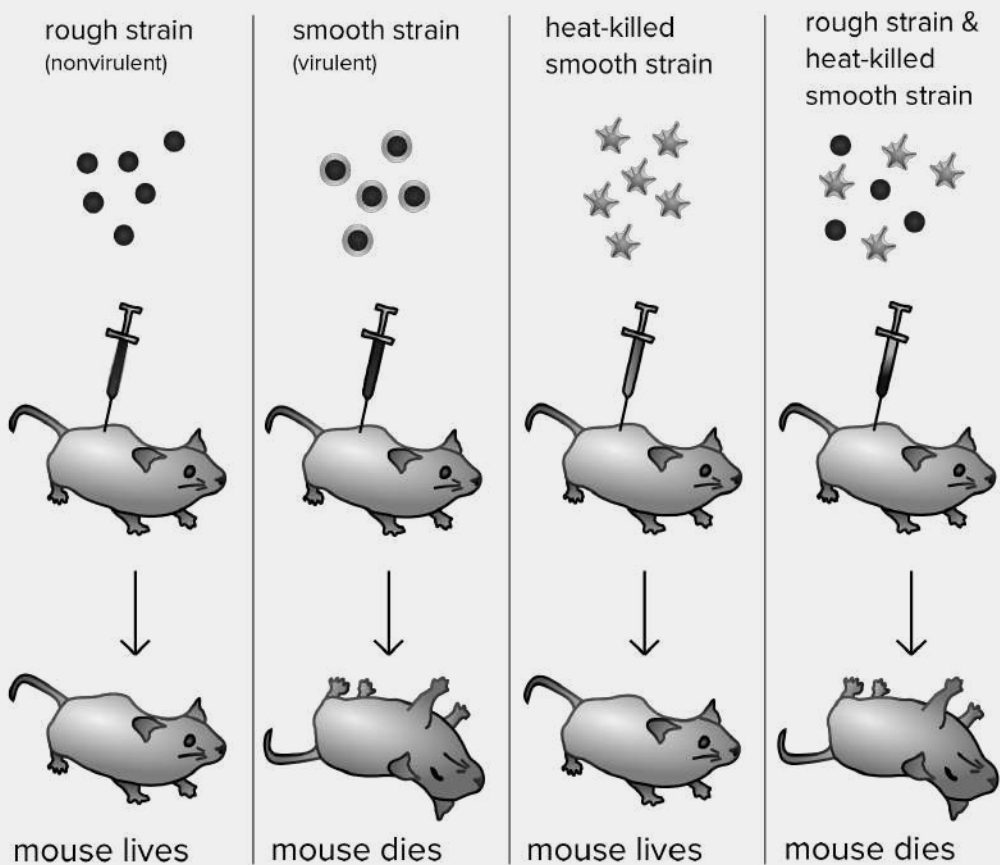
Frederick Griffith



Martha Chase /
Alfred Hershey

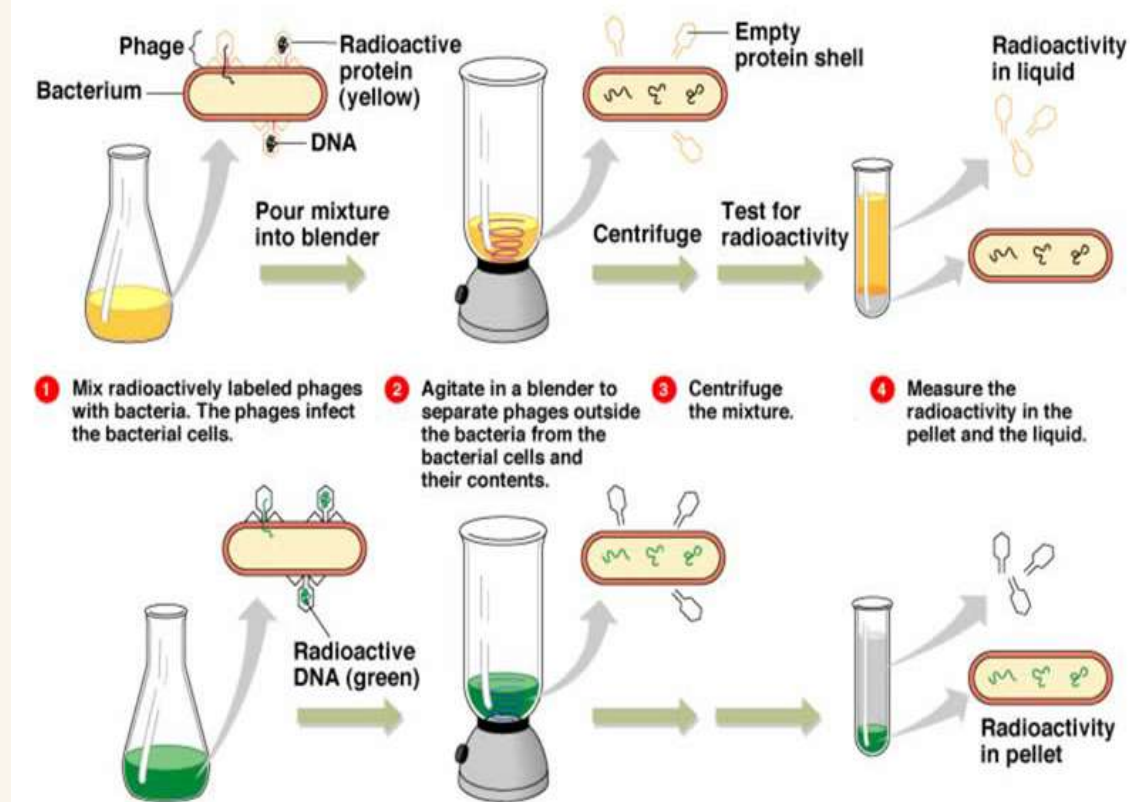


Experimentation

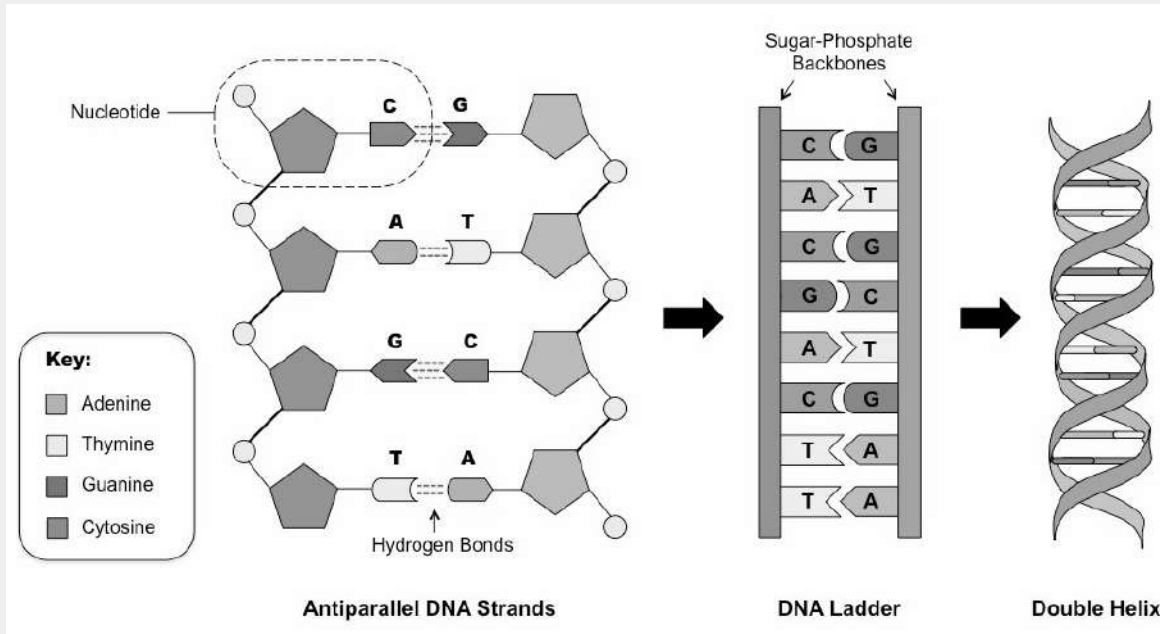


Griffith's Transformation

Hershey - Chase



DNA Structure



DNA Structure:

- Erwin Chargaff showed the amounts of the four bases on DNA (A,T,C,G)
- In a body or somatic cell:
 - A = 30.3%
 - T = 30.3%
 - G = 19.5%
 - C = 19.9%
- Chargaff's Rule:
 - Adenine must pair with Thymine
 - Guanine must pair with Cytosine



DNA Structure Continued



- The bases are held together by weak hydrogen bonds
- DNA's First Photograph
 - **Rosalind Franklin** took diffraction x-ray photographs of DNA crystals
 - In the 1950's, **Watson & Crick** built the first model of DNA using Franklin's X-rays



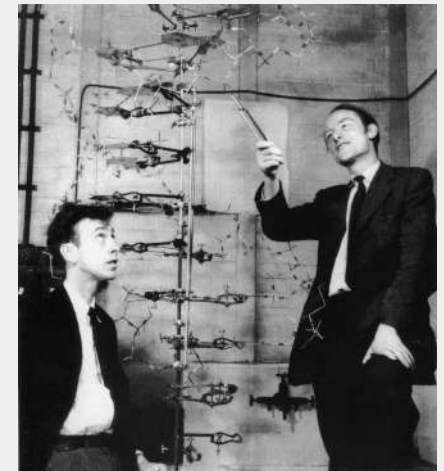
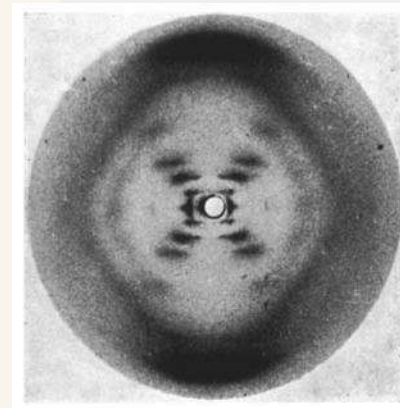
Francis Crick



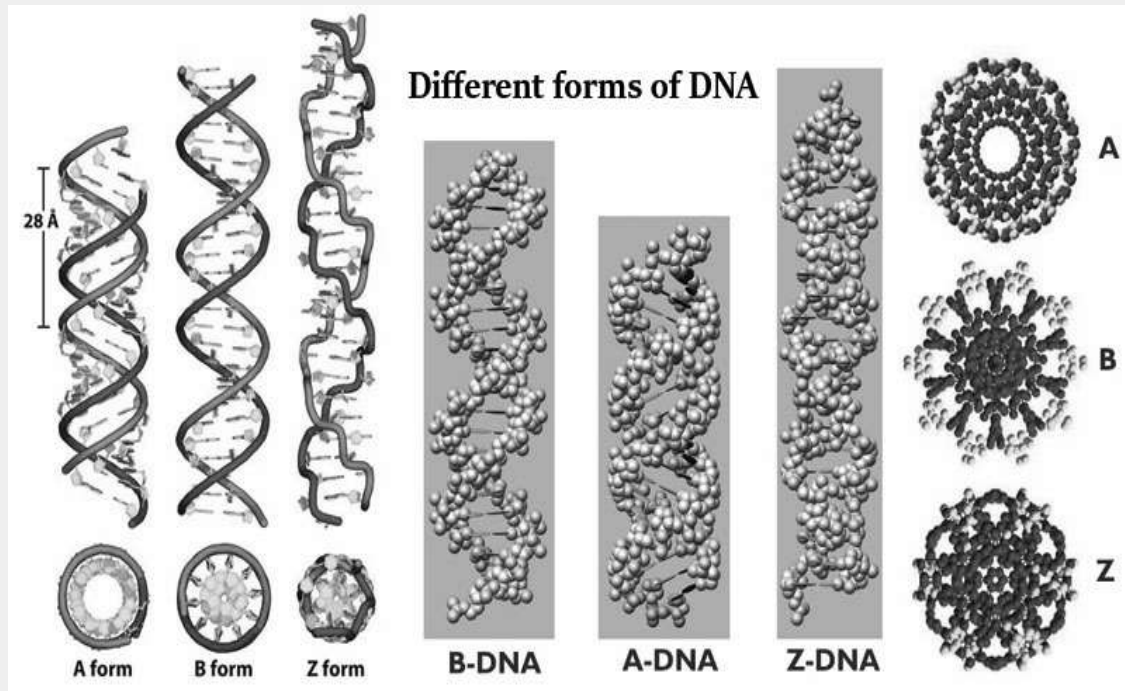
James Watson



Rosalind
Franklin



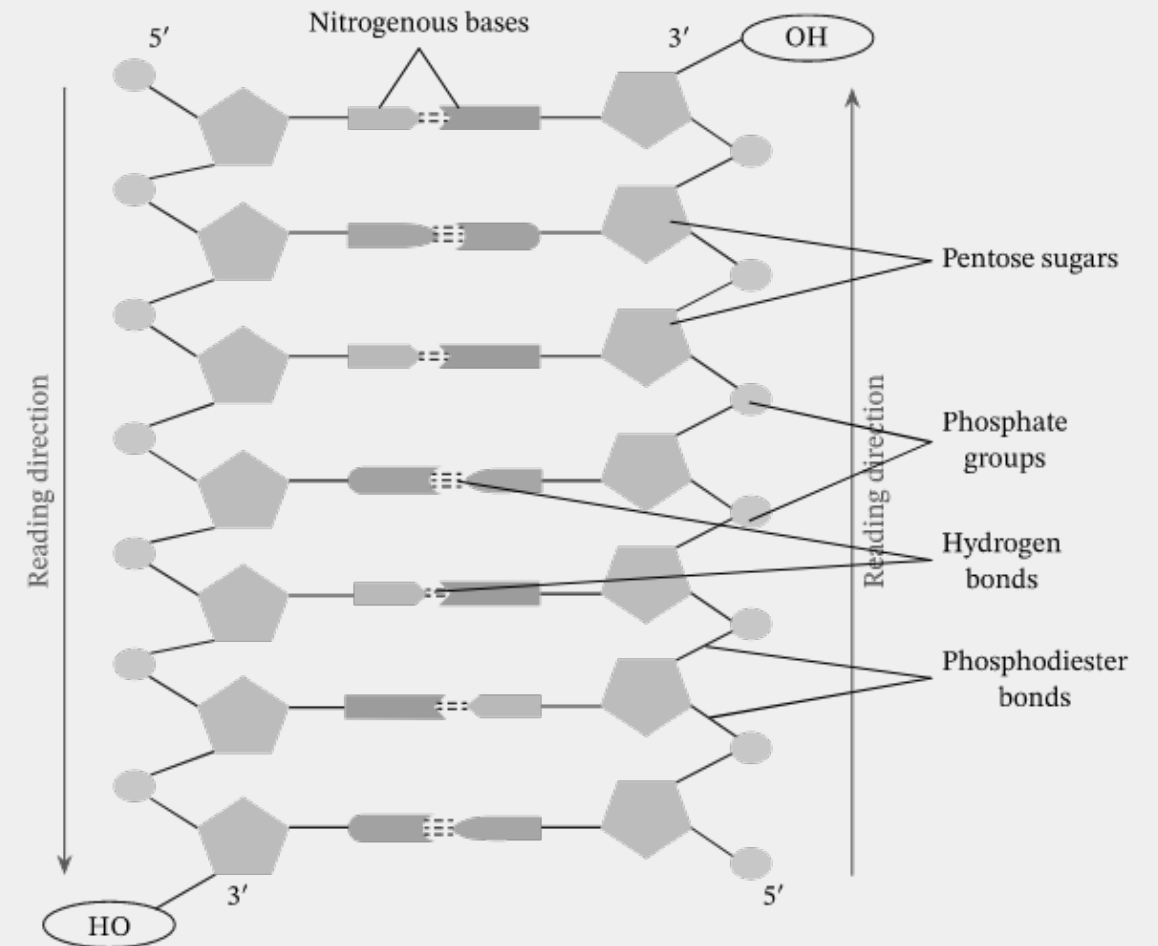
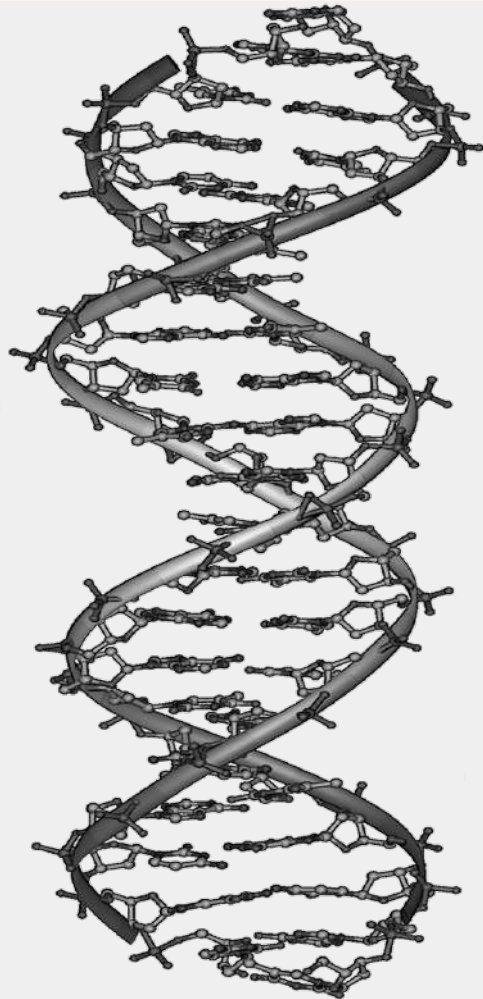
DNA Structure Continued



Structure

- Two strands coiled called a **double helix**
- Sides made of a **pentose sugar Deoxyribose** bonded to **phosphate (PO₄)** groups by **phosphodiester bonds**
- Center made of **nitrogen bases** bonded together by weak **hydrogen bonds**
- Helix
 - **Most DNA (B-DNA)** has a right-hand twist with 10 base pairs in a complete turn
 - Left twisted DNA is called Z-DNA or southpaw DNA
 - Hot spots occur where right and left twisted DNA meet producing mutations

DNA Structure Continued



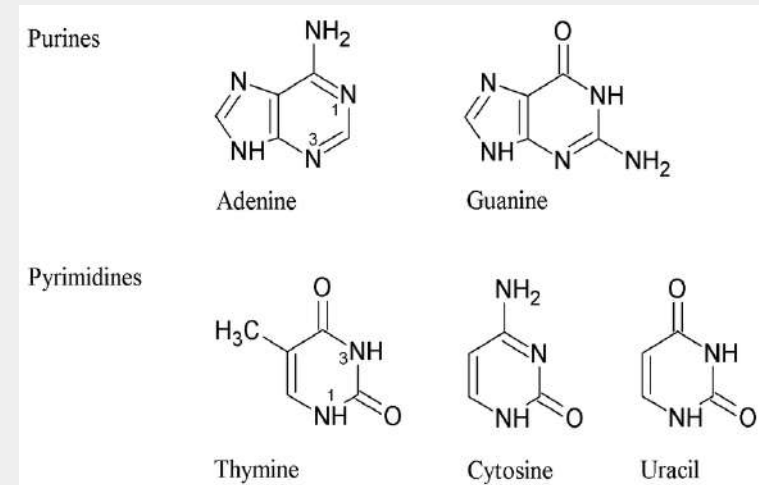
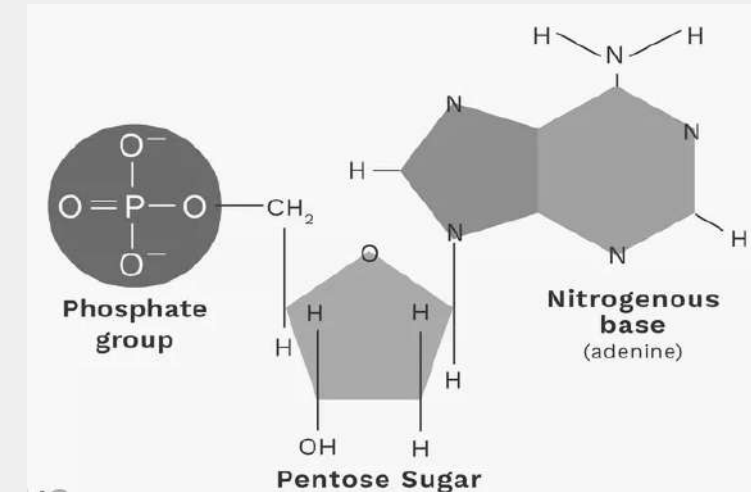
Nucleotides



Nucleotides

- DNA Stands for **Deoxyribonucleic acid**
- Made up of subunits called **nucleotides**
- **Nucleotide** made of:
 - **Phosphate group**
 - **5-carbon sugar**
 - **Nitrogenous base** (genetic code)
- **Double ring PURINES**
 - **Adenine (A)**
 - **Guanine (G)**

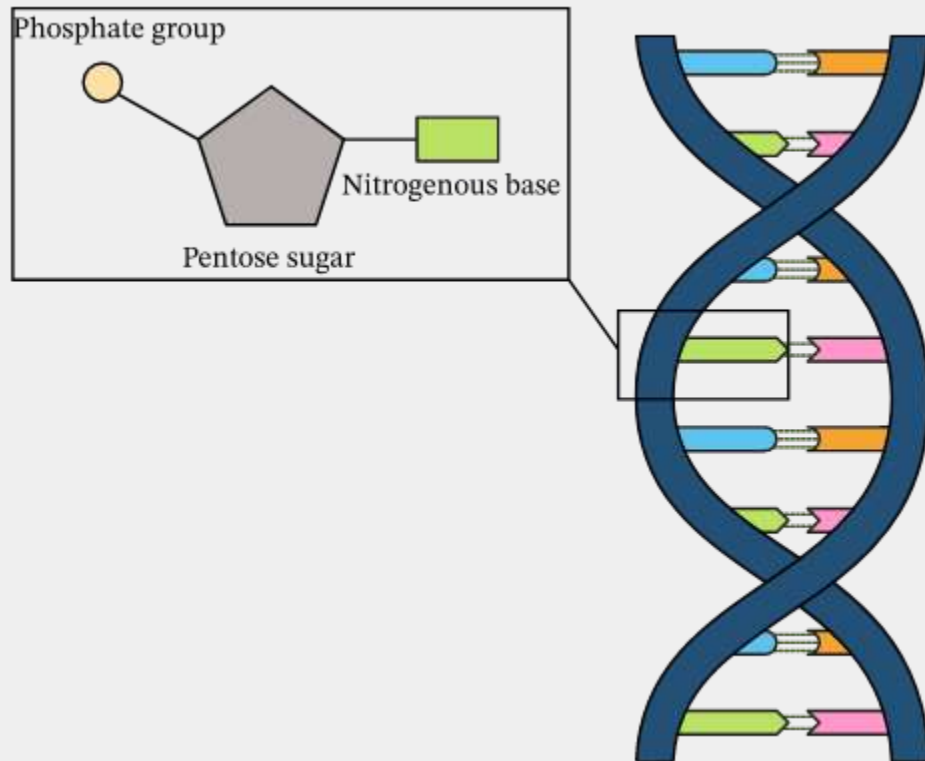
- **Single ring PYRIMIDINES**
 - **Thymine (T)**
 - **Cytosine (C)**
- **Base Pairings**
 - **Purines** only pair with **Pyrimidines**
 - **Three hydrogen bonds** required to bond **Guanine to Cytosine**
 - **Two hydrogen bonds** are required to bond **Adenine to Thymine**
 - These are what allows for DNA to be copied exactly



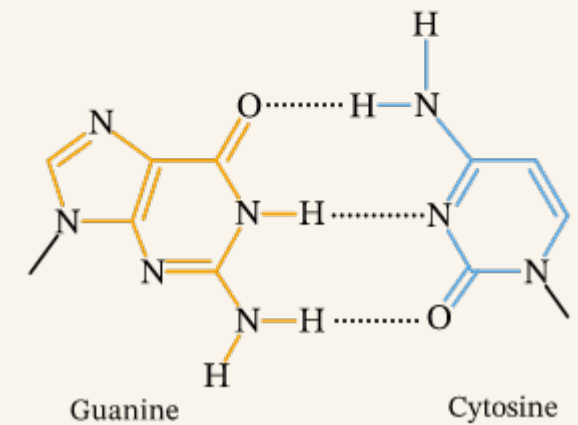
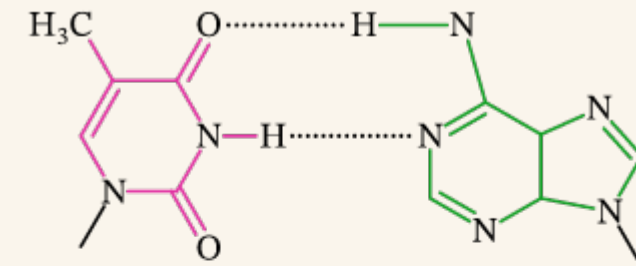
Nucleotides Continued



Nucleotide Structure



Base Pairing



DNA Replication



DNA Replication:

- DNA has to be **copied** before a cell **divides**
- DNA is copied during the **S or synthesis phase** of **interphase**
- New cells will need **identical DNA** strands

→ Occurs in the Nucleus of eukaryotes

→ Replication Fork -

- ❑ Begins at **Origins of Replication** - Two strands open forming **Replication Forks: (Y-shaped region)**

- ❑ New strands grow at the forks

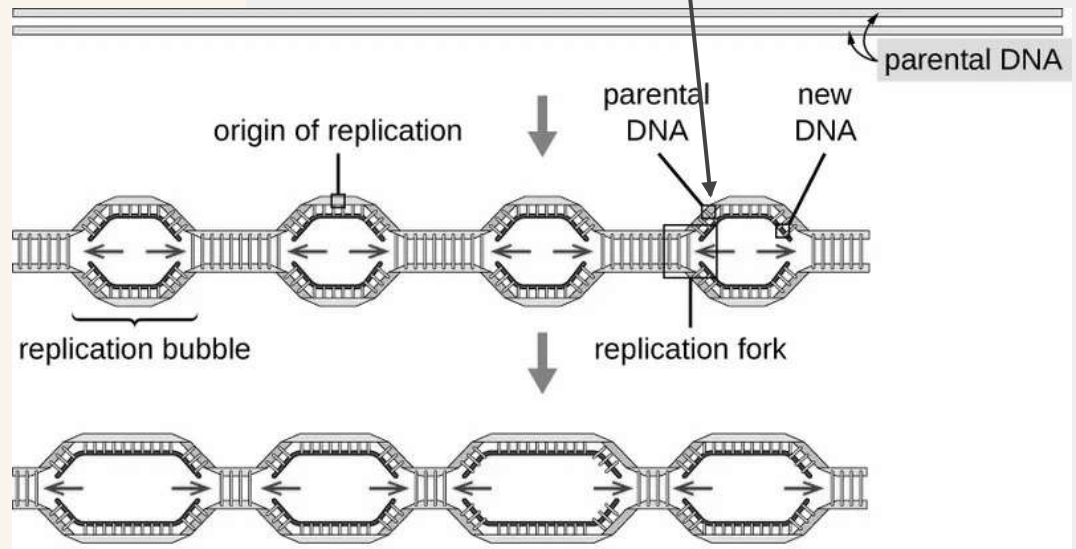
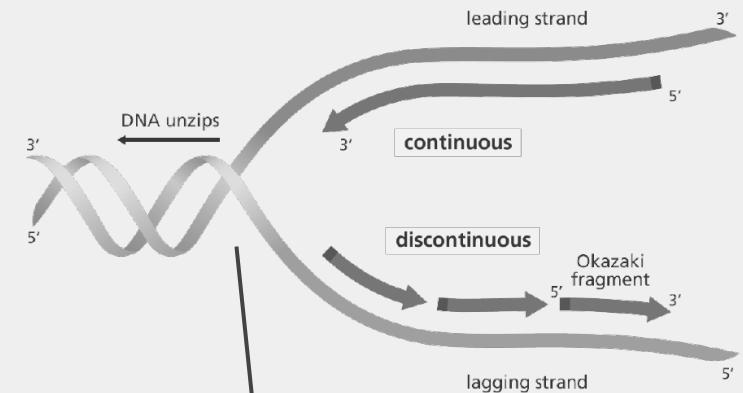
→ Replication Bubbles

- ❑ As the 2 DNA strands open at the origin, Replication Bubbles form

- ❑ **Eukaryotic** chromosomes have **MANY** bubbles

- ❑ **Prokaryotes** (bacteria) have a **single** bubble

DNA replication fork



Enzymes for Replication



01

Helicase

- Enzyme: **Helicase** **unwinds** and **separates** the 2 DNA strands by **breaking** the weak **hydrogen bonds**.
- **Single-Strand Binding Proteins** attach and **keep** the 2 DNA strands **separated** and **untwisted**

02

Topoisomerase

- **Topoisomerase**
→ Enzyme: **Topoisomerase** **attaches** to the 2 forks of the bubble to **relieve stress on the DNA molecule** as it **separates**

03

RNA Primase

- **RNA Primers and Primase**
→ Before new DNA strands can form, there must be **RNA primers present to start** the addition of new **nucleotides**
→ Primase is the enzyme that **synthesizes** the **RNA Primer**
→ **DNA polymerase** can then **add** the new **nucleotides**

04

DNA Polymerase

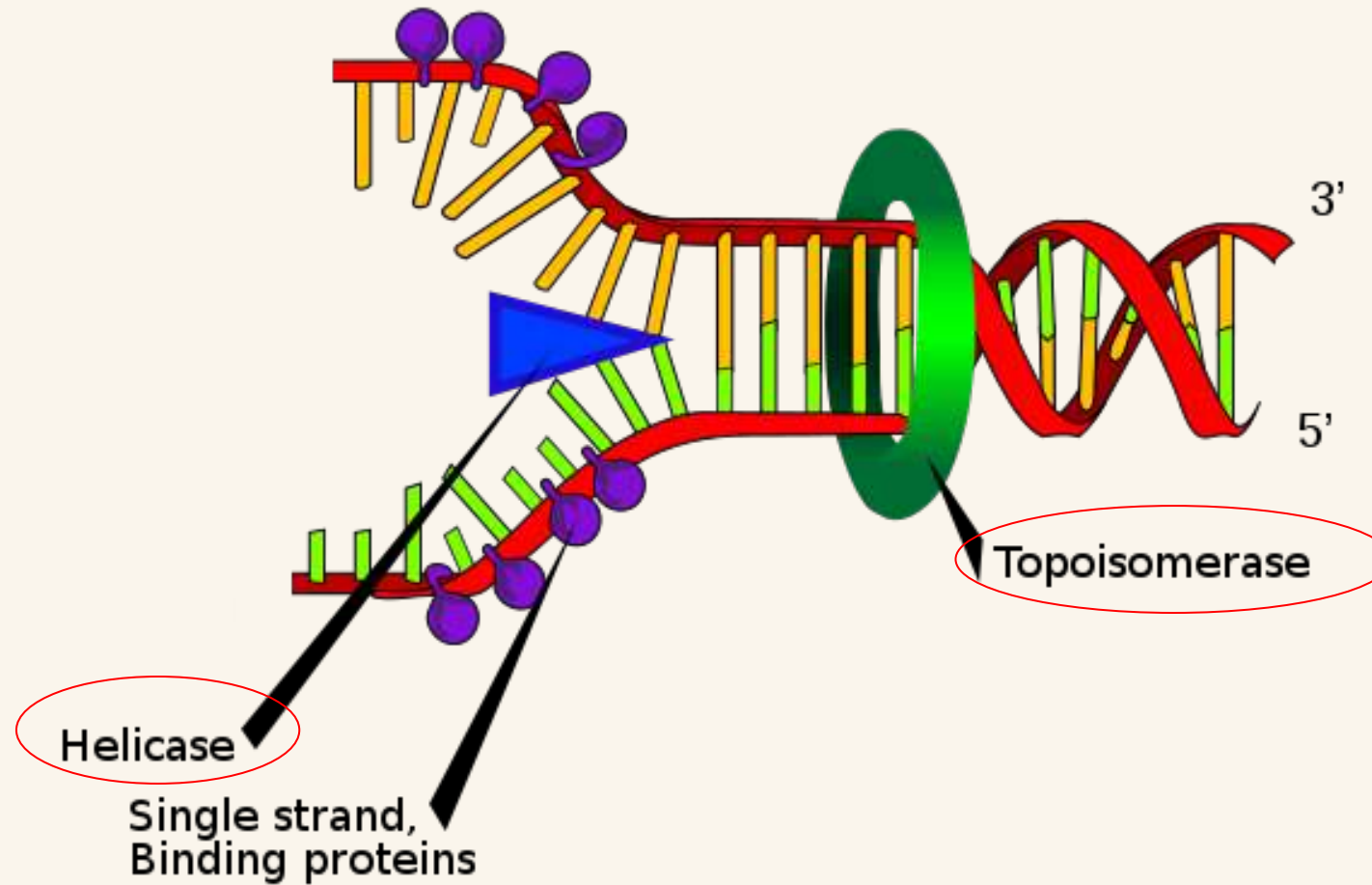
- **DNA Polymerase**
→ DNA polymerase can only add nucleotides to the 3' end of the DNA
→ This causes the **NEW strand** to be built in a 5' to 3' direction

05

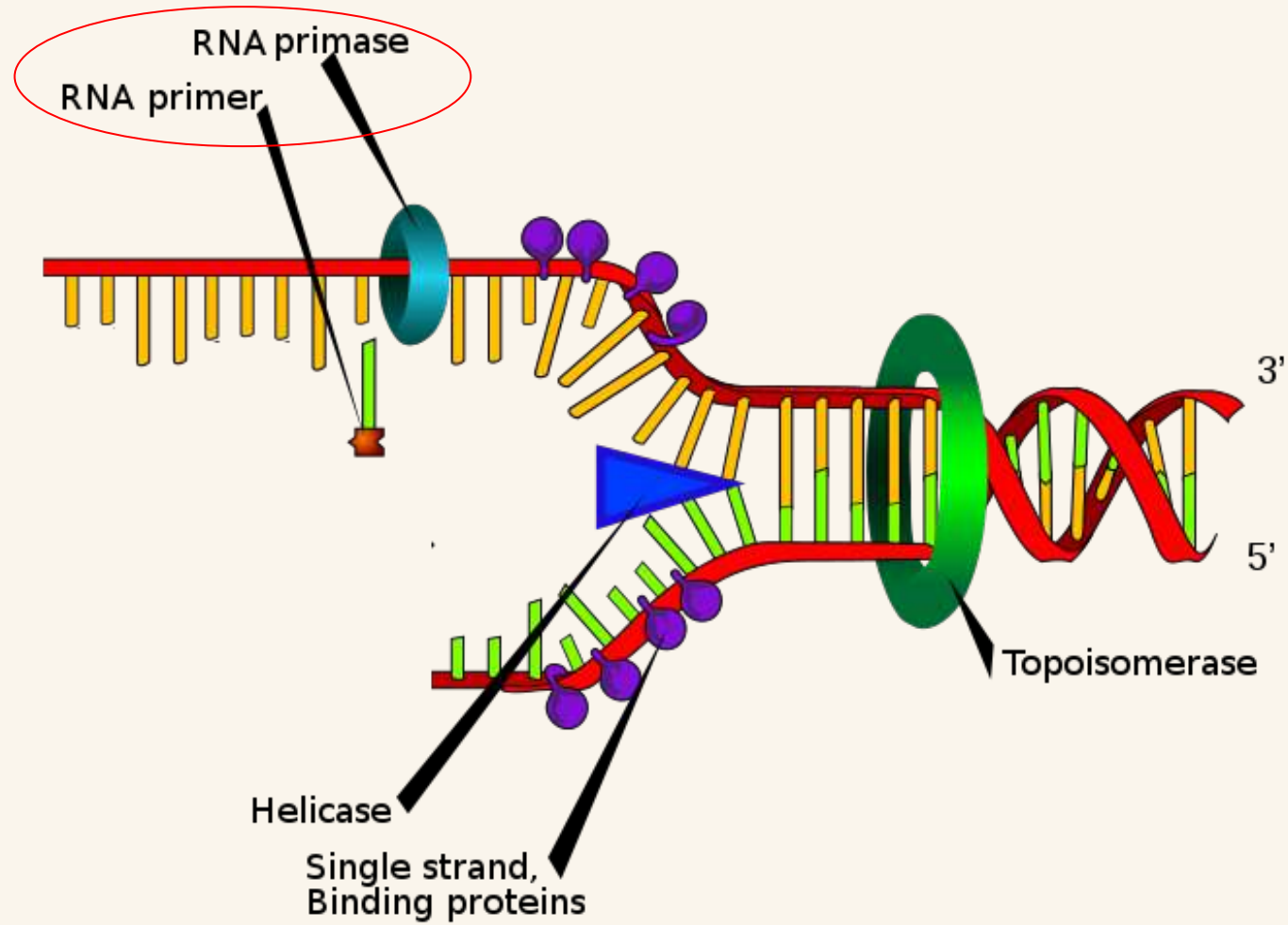
Ligase

- **DNA Ligase**
→ **Okazaki Fragments** - *series of short segments on the lagging strand*
→ Okazaki Fragments must be *joined together* by an *enzyme* – **DNA Ligase**

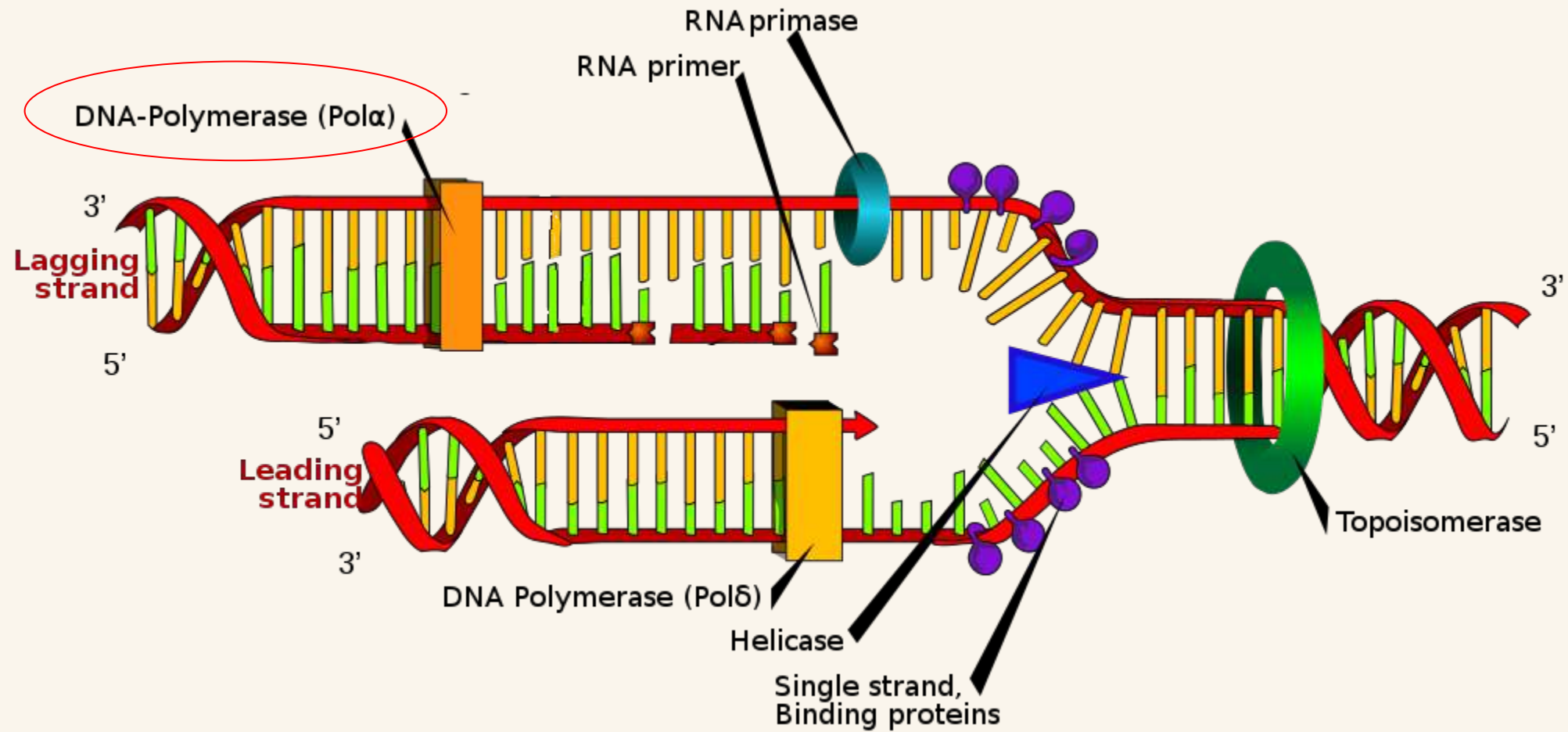
Helicase & Topoisomerase



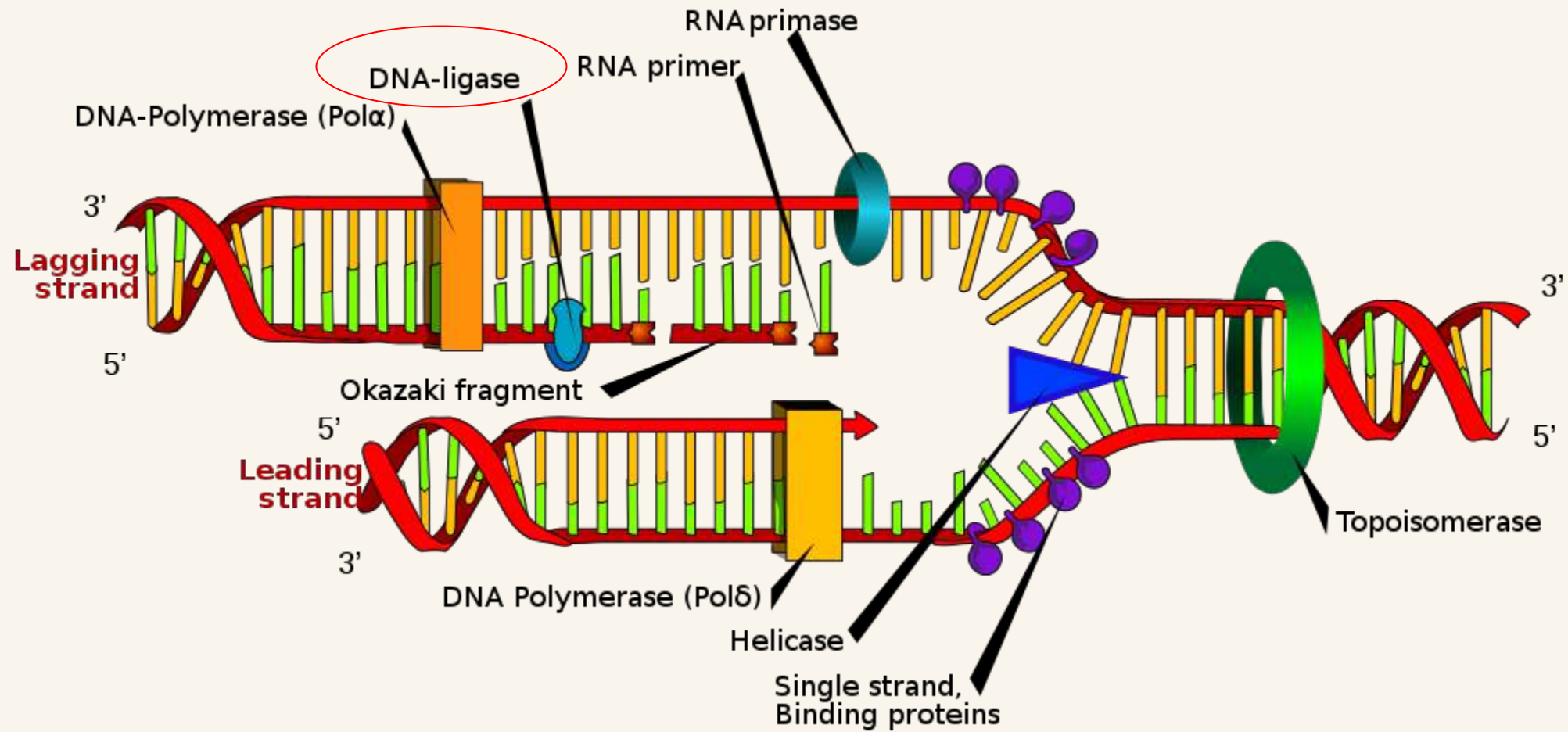
RNA Primase



DNA Polymerase



DNA Ligase



Leading & Lagging Strands

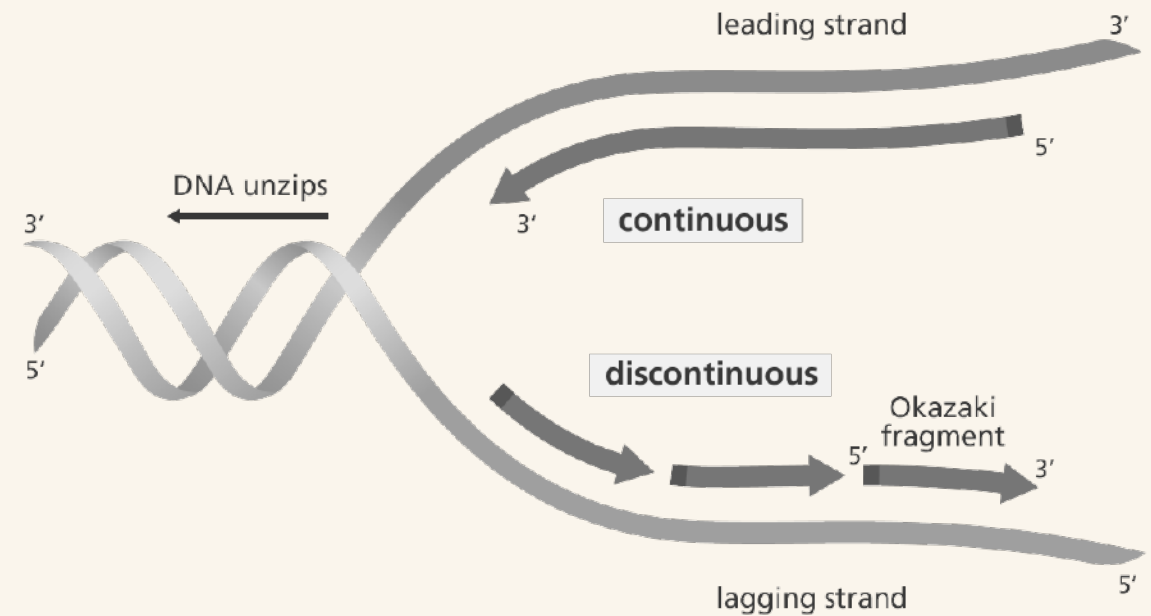


The Leading Strand - (continuous)

- is synthesized as a single strand from the point of origin toward the opening replication fork

Lagging Strand: (discontinuous)

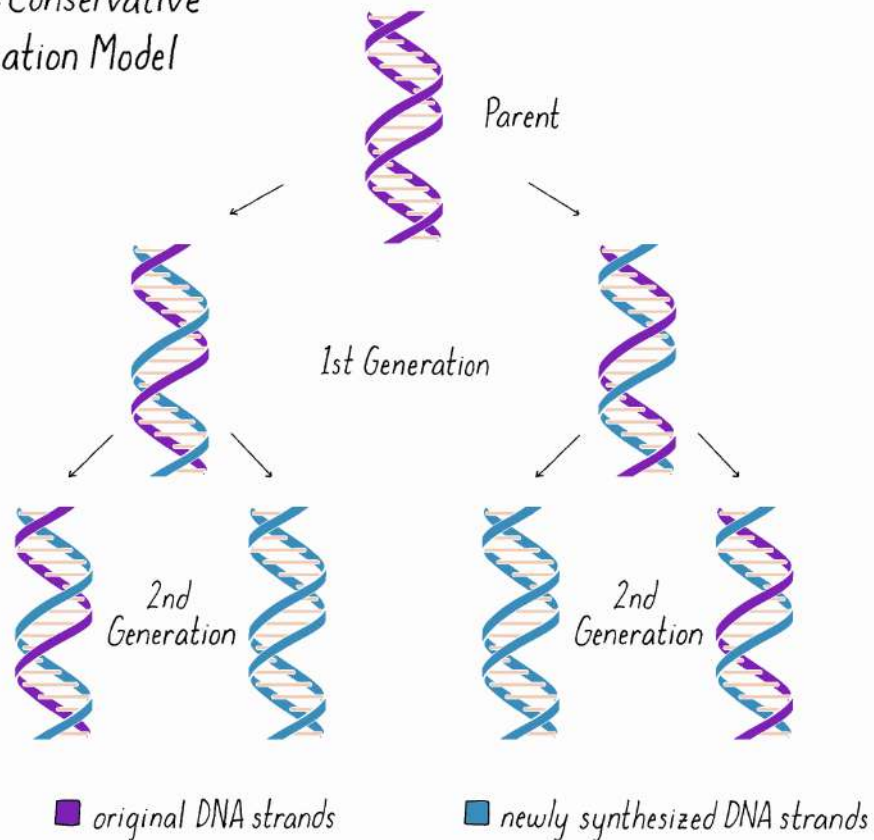
- The Lagging Strand is synthesized discontinuously against overall direction of replication
- This strand is made in MANY short segments It is replicated from the **replication fork toward the origin**
- **Okazaki Fragments** - series of short segments on the lagging strand



Semi-Conservative Replication & Proofreading



*Semi-Conservative
Replication Model*



Semiconservative Model for Replication

- Idea presented by Watson & Crick
- The two strands of the parental molecule separate, and each acts as a template for a new complementary strand
- New DNA consists of 1 PARENTAL (original) and 1 NEW strand of DNA

Proofreading DNA

- DNA polymerase initially makes about 1:10,000 base pairing errors
- Enzymes proofread and correct these mistakes
- The new error rate for DNA that has been proofread is 1 in 1 billion base pairing errors

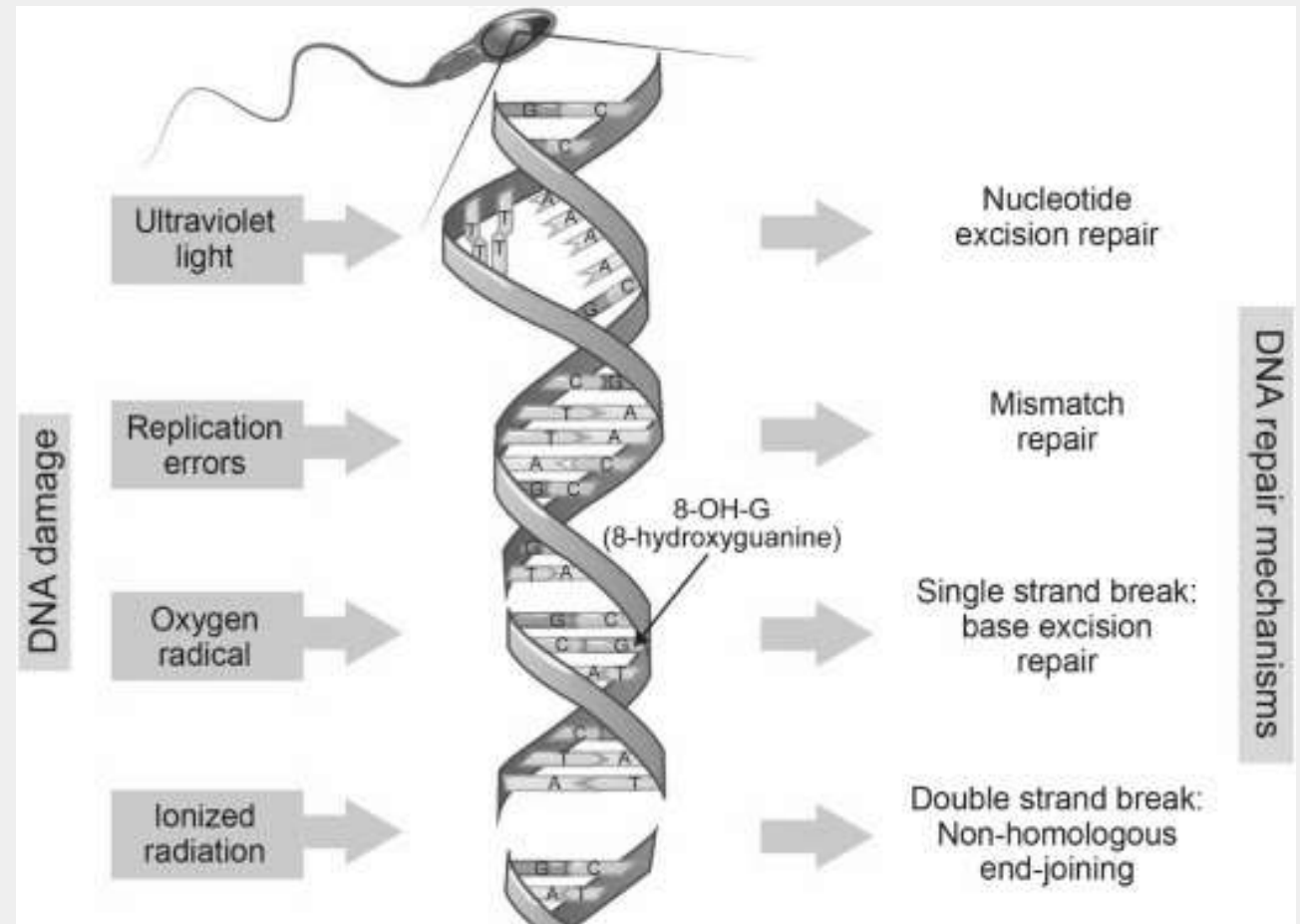


DNA Damage & Repair

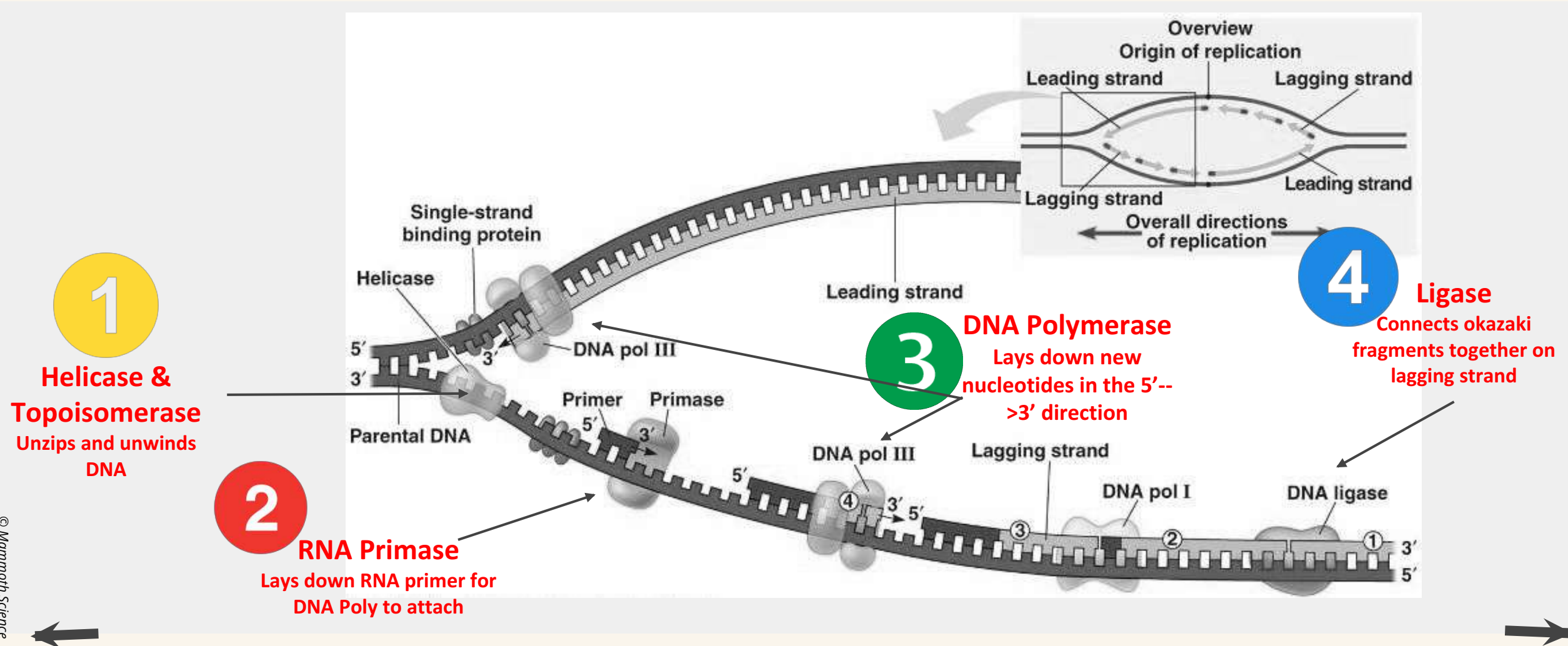


DNA Damage & Repair:

- Chemicals & ultraviolet radiation damage the DNA in our body cells
- Cells must continuously repair DAMAGED DNA
- Excision repair occurs when any of over 50 repair enzymes remove damaged parts of DNA
- DNA polymerase and DNA ligase replace and bond the new nucleotides together



DNA Replication Summary





Thank you!

Do you have any questions?

instructor@email.com

XXX-XXX-XXXX

