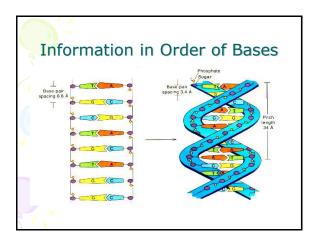


Function

- 1. DNA stores and transfers all of the GENETIC information of a cell.
- 2. DNA is the code for ALL proteins in the cells.
- 3. DNA→ mRNA→ amino acid order→ protein shape → protein function



Location

Prokaryotic Cells

- Bacteria → no nucleus
- DNA is stored in ONE CIRCULAR chromosome in the middle of the cytoplasm

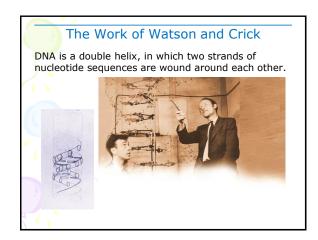
Eukaryotic Cells

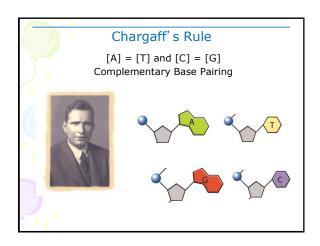
- Plant and animal cells
 → have nucleus
- DNA ALWAYS STORED INSIDE OF THE NUCLEAR MEMBRANE.
- Linear chromosome

Determining the Structure

- 1952 Hershey and Chase
 - determined DNA was genetic material
- 1953 James Watson and Francis Crick
 - Determined 3-d structure of DNA
 - -Based on work by Rosalind Franklin
 - Won Nobel Prize in 1962

Franklin's X-rays DNA is a helix. Likely two strands to the molecule Nitrogenous bases near the center of the molecule



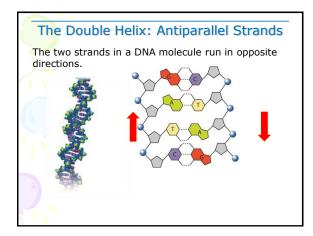


Structure

- 1. Polymer made of repeating nucleotides.
- 2. DNA has a DOUBLE HELIX shape
 - Spiral
 - Twisted ladder
- 3. One side of ladder is upside down compared to other

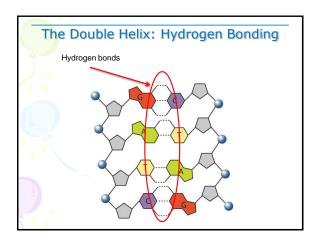
Structure

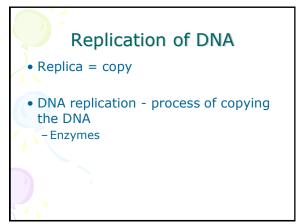
- 4. Two strands of nucleotides bonded together in the middle.
- 5. Bases follow the complementary base pairing rules
 - A ↔ T
 - -C ↔ G

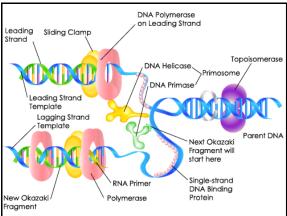


Bonding

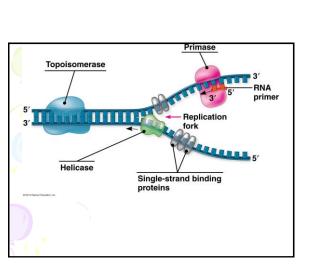
- 1. Strong **covalent** bonds hold together the sugar-phosphate backbone.
- 2. Weak **hydrogen** bonds hold together the complementary base pairs in the middle of the molecule.









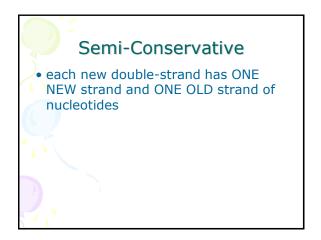


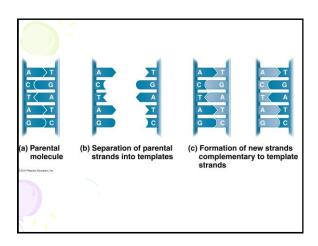
The Enzymes

- DNA polymerase (I and III): elongates new strands
- Helicase: Separates strands
- Primase: starts process of elongation (RNA)
- Ligase: Joins pieces together
- Single strand binding proteins: keeps strands from rejoining.
- Topoisomerase: prevents supercoiling

Process

- 1. Helicase separates the strands
- 2. SSBP keep strands from rejoining
- 3. Primase adds primer to start process
- DNA polymerase III reads each strand, making complementary base pairs.
- 5. DNA polymerase I replaces primer
- 6. Ligase joins ends of pieces





Replication • How do the two new pieces of DNA compare to each other? - identical • How do they compare to the original piece? - identical