Central Dogma of Biology Procedures

- 1. Replication
- 2. Transcription
- 3. Translation

Replication

- 1. **Purpose**: Replication produces two identical copies of a DNA double-stranded molecule, so that they can be inherited by the two daughter cells of a parent cell.
- 2. **Procedure**: Each strand serves as a template for the synthesis of its complementary strand in an antiparallel orientation, following Chargaff's rule.

Transcription

- 1. **Purpose**: Transcription produces a messenger RNA (mRNA) copy of the gene's DNA coding strand. mRNAs are expendable molecules that will be translated into polypetide chains in the cytoplasm.
- 2. **Procedure**: The non-coding DNA strand of the gene serves as a template for the synthesis of mRNA, according to Chargaff's rule.

Translation

- 1. **Purpose**: Translation connects amino acids, forming a polypeptide chain, using the information that is stored in DNA coding regions.
- 2. Procedure: Ribosomes facilitate the formation of the peptide bonds between amino acids. Transfer RNAs (tRNAs) read the mRNA's sequence, 3 nitrogenous bases (codons) at a time. They pair their anti-codon with the next available codon, ensuring the proper positioning of the corresponding amino acid. The Codon - Anti-codon pairing, in anti-parallel orientation, follows the Chargaff's rule.

DNA Nitrogenous Bases Table

| Table 1A: DNA Nitrogenous bases | | | | | |
|---------------------------------|--------------|------------|--|--|--|
| | Name | Category | | | |
| 1. | Adenine (A) | Purine | | | |
| 2. | Thymine (T) | Pyrimidine | | | |
| 3. | Guanine (G) | Purine | | | |
| 4. | Cytosine (C) | Pyrimidine | | | |

RNA Nitrogenous Bases Table

| Table 1B: RNA Nitrogenous bases | | | | | | |
|---------------------------------|--------------|------------|--|--|--|--|
| | Name | Category | | | | |
| 1. | Adenine (A) | Purine | | | | |
| 2. | Uracil (U) | Pyrimidine | | | | |
| 3. | Guanine (G) | Purine | | | | |
| 4. | Cytosine (C) | Pyrimidine | | | | |

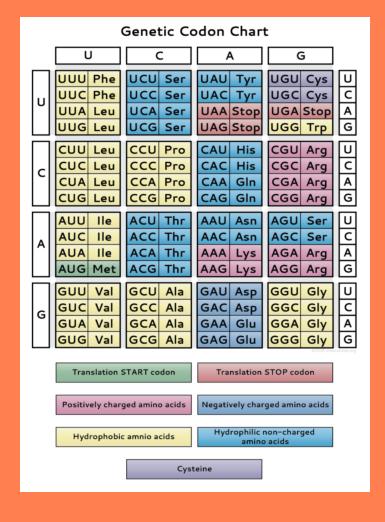
Chargaff's Rule of Complementary Nitrogenous Bases Pairing

According to Chargaff's rule the complementary pairings are:

- 1. A pairs with U via 2 Hydrogen bonds.
- 2. A pairs with T via 2 Hydrogen bonds.
- 3. G pairs with C via 3 Hydrogen bonds.

Standard Genetic Code Table

Image taken from: sciencenotes.org



Central Dogma of Biology Summary Table

| Table 2: DNA Coding Sequence Replication, | Transcription and Trar | nslation (Central Dogma) |
|--|------------------------|--------------------------|
| <i>y</i> , , , , , , , , , , , , , , , , , , , | | ` , |

| Procedure | Macromolecule | Starting Edge | Starting Codon | Codon 1 | Codon 2 | Termination Codon | Ending Edge |
|---------------|----------------------|------------------|-------------------|------------|------------|----------------------|----------------|
| | DNA coding | 5' | ATG | GAG | CTC | TAA | 3' |
| Replication | DNA complementary | 3' | TAC | СТС | GAG | ATT | 5' |
| Transcription | mRNA | 5' | AUG | GAG | CUC | UAA | 3' |
| Translation | Polypeptide | H ₂ N | Met | Glu | Leu | - | COOH |

The End

For more information check your eclass!