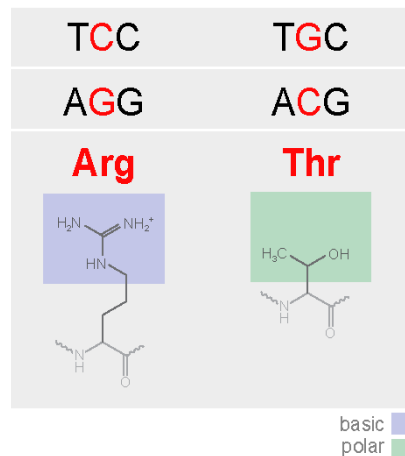


## CONCEPT: GENOME EVOLUTION

### Mutations in Evolution

- Accumulations in **mutations** (DNA changes) promote genome \_\_\_\_\_
  - **Point mutations** are changes in a single nucleotide pair
    - Can be beneficial or detrimental to the organisms
    - Arise from errors in DNA replication
  - Mutations can occur in the gene, or in regulatory DNA
    - Gene: effects activity, interaction, stability and is easy to spot
    - Regulatory DNA: effects how genes are expressed and are not so easy to find

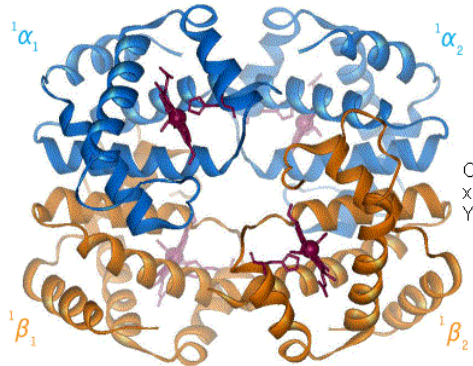
**EXAMPLE:** Point mutations can affect amino acid sequence



### Gene Duplication in Evolution

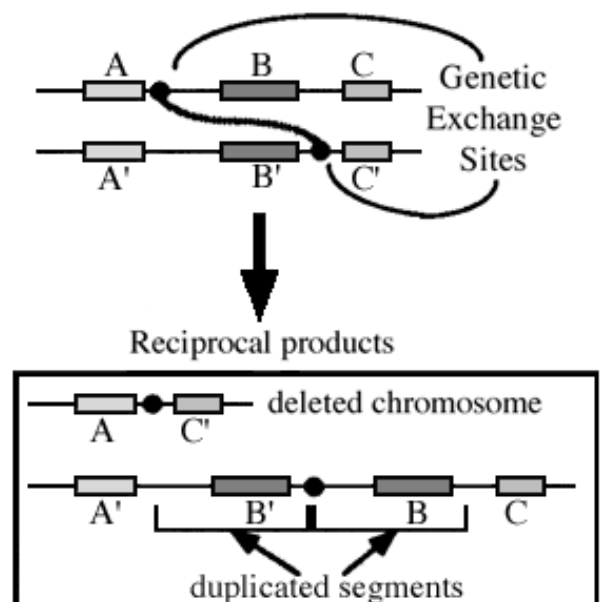
- **Gene duplication** is a main driver of genomic evolution
  - **Gene families** are groups of genes with similar sequences but \_\_\_\_\_ functions
    - Once a gene is duplicated, each copy is free to accumulate mutations that result in different functions
    - Gene families are the result of gene duplication

**EXAMPLE:** Structure of hemoglobin demonstrates gene families



- Gene duplication arises from \_\_\_\_\_ *crossing over* during mitosis
  - Misalignment of chromosomes during *homologous recombination* can lead to lopsided genetic exchange
  - Result: one chromosome with extra gene copy and one with no copy
- **Pseudogenes** are duplicated genes that have lost their functional ability but remain in the genome
  - *Processed pseudogenes* occur by changing a mRNA to a DNA and integrating it into a chromosome
- **Whole genome duplication** is when the entire genome of an organisms is copied and retained in a single cell
  - **Polyploidization** (whole genome duplications) are common in fungi and plants

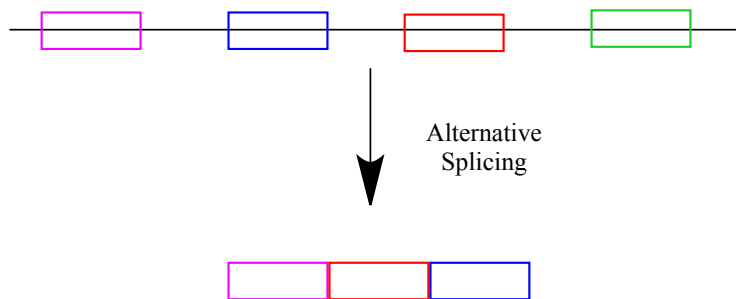
**EXAMPLE:** Unequal crossing over results in duplicated genes



## Introns and Splicing in Evolution

- The presence of *introns* allows for gene shuffling and gene evolution
  - Genes of most organisms contain *Introns* and *Exons* (not completely universal ex: Histone proteins)
    - **Introns** are noncoding regions of a gene that are cut out during gene processing
    - **Exons** are the coding regions of a gene
  - **Alternative splicing** is the combining of exons from one gene in new orders (occurs in 50-90% of human genes)
    - Produce *isoforms* which are different forms of the same protein produced through alternative splicing
  - **Exon shuffling** is the combining of exons from two \_\_\_\_\_ genes
    - Can also occur if exons are duplicated or moved to different genomic location

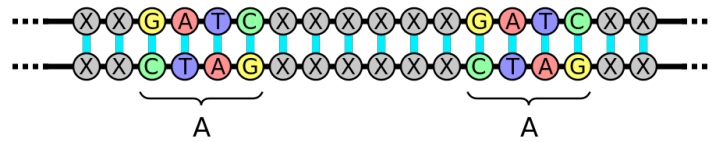
### **EXAMPLE:** Model of alternative splicing



## Repetitive DNA sequences in Evolution

- **Repetitive DNA sequences** have evolved in the genomes and are extremely common
  - **Simple-sequence repeats** are arrays of thousands of copies of a short sequence (1-500 nucleotide)
    - *Drosophila*: ACAAACT
    - Not transcribed, and contain no genetic information
  - **Mobile genetic elements** are DNA sequences that can \_\_\_\_\_ through the genome
    - Contain repetitive DNA flanking protein coding regions
    - **Transposons** are mobile genetic elements and can move through RNA or DNA intermediates
    - Insert anywhere in a gene, and can effect gene structure or regulation

**EXAMPLE:** Example of sequence repeats in the genetic code



**PRACTICE:**

1. Which of the following is not a driver of genome evolution?
  - a. Mutations
  - b. Gene duplication
  - c. Alternative Splicing
  - d. Histone modifications

2. Which of the following genomic changes are most likely to cause pseudogenes?

- a. Point Mutations
- b. Gene Duplications
- c. Exon Shuffling
- d. Transposons

3. Protein isoforms are created through which process?

- a. Point Mutations
- b. Alternative Splicing
- c. Exon Shuffling
- d. Simple Sequence Repeats