

PCR, DNA Fingerprinting, and DNA Sequencing

- **Polymerase Chain Reaction** is DNA Replication in a Test Tube
 - PCR copies a specific DNA sequence
 - Can take a very small amount of DNA and make millions of copies for testing.
 - PCR requires **primers** that can target certain sections of DNA.
 - The **three steps** of PCR
 - Denaturation (split into single strands using heat)
 - primers attach
 - Extension (DNA polymerase makes complimentary strand)
 - Steps are repeated over and over to amplify the DNA.
- **Electrophoresis**
 - An **agrose gel** (jello-like) allows pieces of DNA to pass through it in response to an electrical charge.
 - DNA is negatively charged and is attracted to the positive electrode.
 - Shorter pieces move further/faster.
 - Stain or some other method is used to see the bands.
- DNA Fingerprints Can Identify Individuals
 - Focusing on highly variable sections of DNA
 - Sections can vary in sequence or in length.
 - Used in forensics to rule out suspects or as evidence against a suspect.
 - **RFLP (restriction fragment length polymorphism)**
 - Differences in sequences alter restriction enzyme cutting sites
 - Different cutting sites make different lengths of DNA pieces.
 - **VNTRs (Variable-number tandem repeats)**
 - Differences in length is caused by repeated sections of DNA bounded by restriction enzyme sites
 - More repeats equals a longer DNA segment
 - **STRs (Short tandem repeats)**
 - Really short VNTRs (2,3,4,5)
 - Have primer sites on either side for PCR
 - Used for PCR analysis in forensic work to generate DNA profiles (**Combined DNA Index System, CODIS**)
 - Paternity Test
 - Match markers between mother and child. Rest of the markers must come from father.
- **DNA Sequencing**
 - Figuring out the bases on a strand of DNA.
 - Used in the Human Genome Project to sequence all of human DNA.
 - Have sequenced other organisms
 - Large-scale genome sequencing is automated and uses fluorescent dye-labeled dideoxynucleotides.

Population Genetics

- A **population** is a group of individuals with a common set of genes that lives in the same geographic area and can or does interbreed.
- A population's **gene pool** is all of the alleles present in that population. Due to population dynamics, the gene pool can change over time.
- The Hardy-Weinberg Law Describes the Relationship between Allele Frequencies and Genotype Frequencies in an Ideal Population
 - The **Hardy-Weinberg law** makes two predictions:
 - (1) the frequency of the alleles in the gene pool does not change over time; and
 - (2) after one generation of random mating, the genotype frequencies for two alleles can be calculated as $p^2 + 2pq + q^2 = 1$ where p equals the frequency of allele A and q is the frequency of allele a

- The Hardy-Weinberg model assumes
 - 1.that there is no selection,
 - 2.that no new alleles arise from mutation,
 - 3.that there is no migration into or out of the population,
 - 4.that the population is large, and
 - 5.that random mating occurs.
- An example of how the Hardy-Weinberg law can be applied to humans is analysis of susceptibility to HIV-1 infection based on the genotype for the *CCR5* HIV-1 receptor gene.
- Ex. Percent of the population that carries the CF gene.