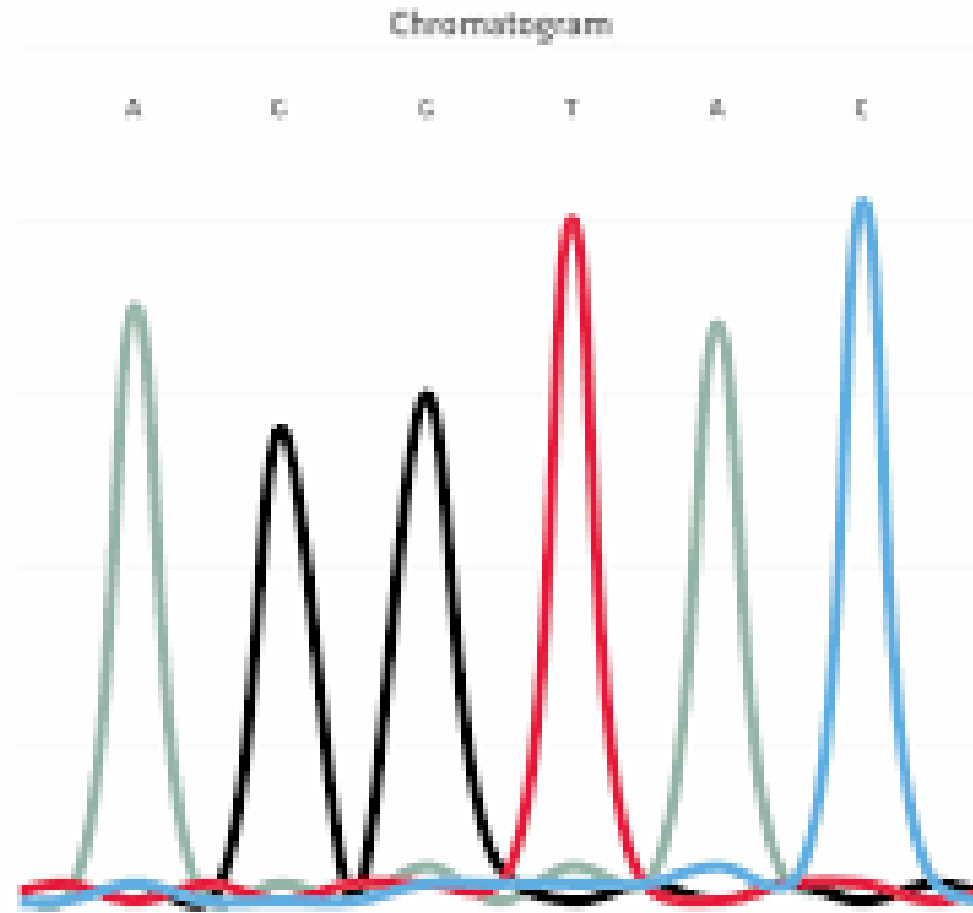


Sanger Sequencing

The Gold Standard in DNA Sequencing...

When have a doubt just sequence it out...



By Mohsen Al-Saleh

17/02/2025

Outlines:

- Introduction/Background
- Principles/Key Components
- Procedure/Steps
- Applications, Advantages, Limitations, Conclusion

DNA Sequencing:

-Determining the exact sequence of nucleotides, or bases, in a DNA molecule.

- **A sequencing can be done by different methods including:**
 1. Maxam – Gilbert sequencing (chemical degradation method). (1977)
 2. Sanger sequencing (dideoxy chain-termination method). (1977)
 3. High-throughput sequencing technologies (NGS). (2000)

-What is Sanger Sequencing?

- 1-Method to determine nucleotide sequence of DNA.
- 2-Relies on chain-terminating dideoxynucleotides (ddNTPs).

-Historical Context:

- 1-Developed by Frederick Sanger (1977).
- 2-Key role in the Human Genome Project (1990).

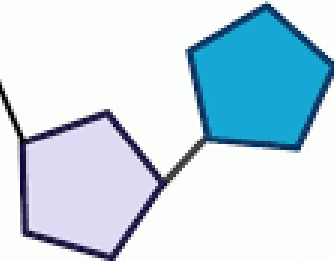
-Significance:

High accuracy; remains a benchmark for validating genetic data.

Phosphate

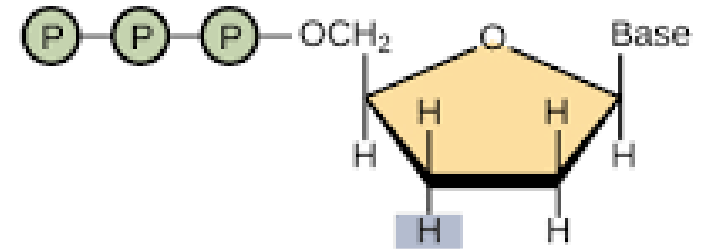


Sugar



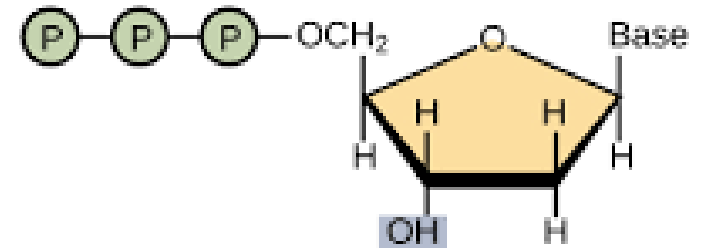
Nitrogenous
Base

Nucleotide



Dideoxynucleotide (ddNTP)

Sanger



Deoxynucleotide (dNTP)

DNA

Key Principles

-Chain Termination:

ddNTPs lack 3'-OH group, halting DNA synthesis.

-Components:

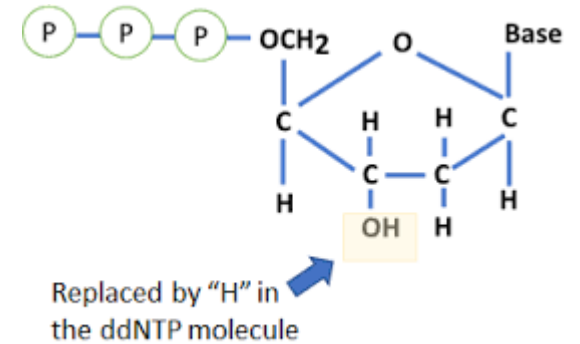
DNA template, primer, DNA polymerase, dNTPs, fluorescently labeled ddNTPs.

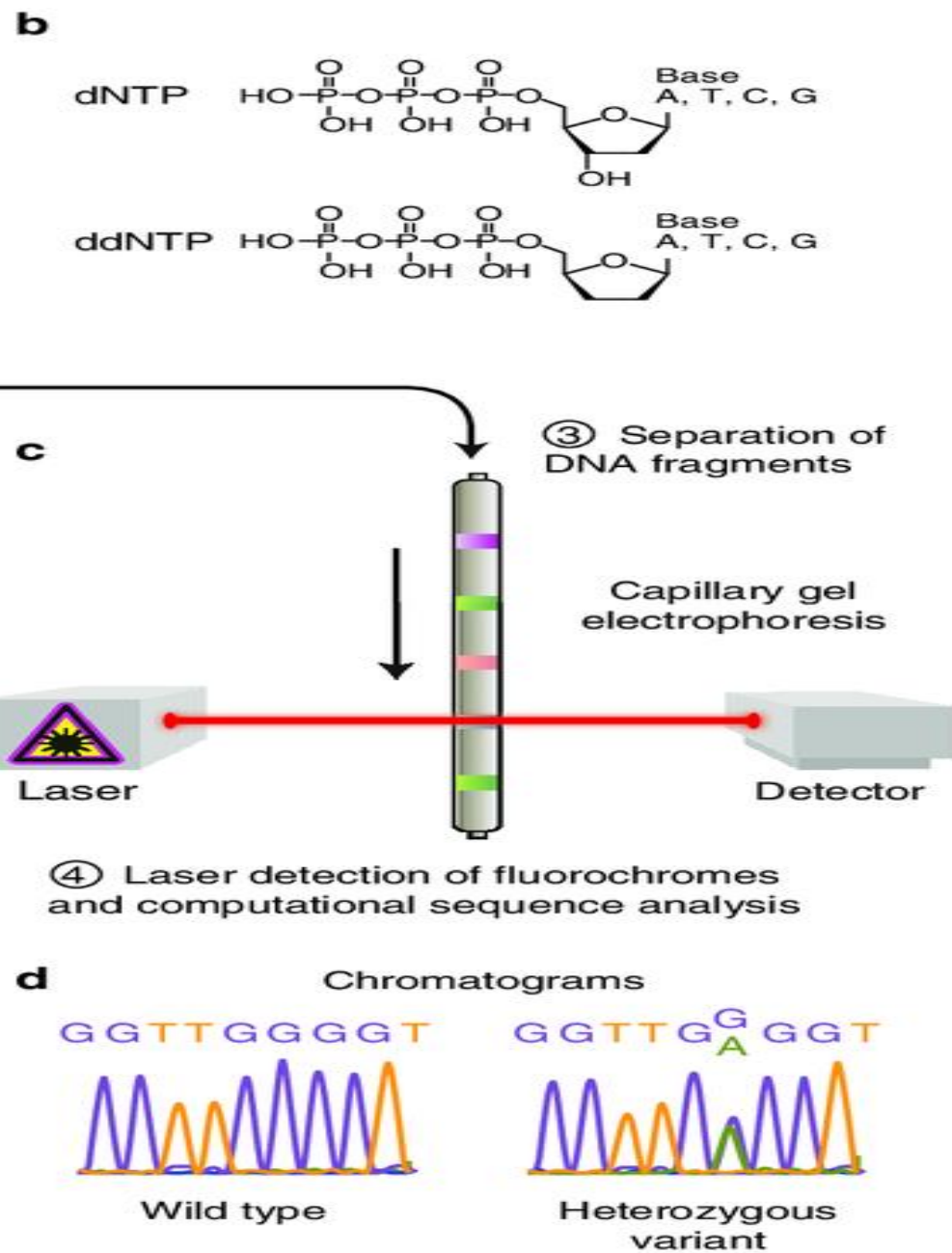
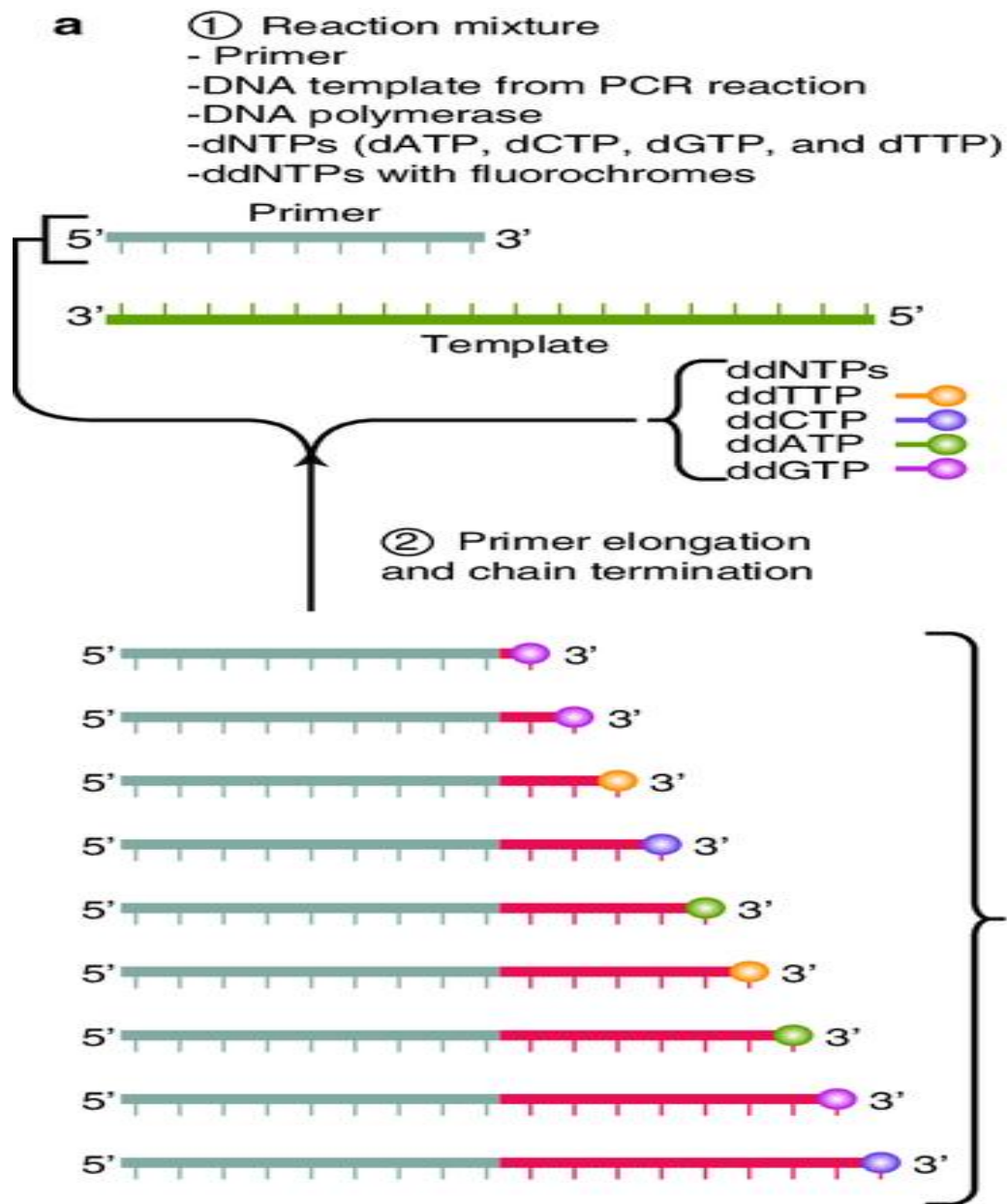
-Detection:

Fluorescent tags on ddNTPs (each base with a unique color). Laser beam

-Separation:

Capillary electrophoresis sorts DNA fragments by size.





Procedure Overview

-Template Preparation:

Isolate target DNA.

-Cycle Sequencing Reaction:

Linear amplification with ddNTPs and dNTPs.

-Denaturation:

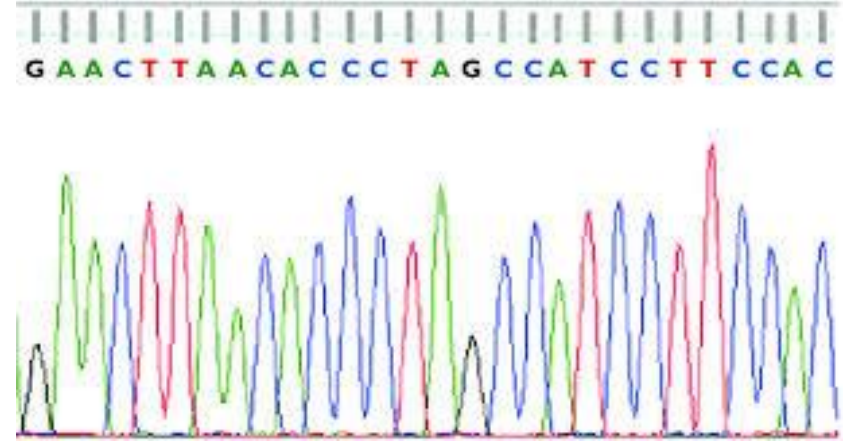
Separate DNA strands.

-Electrophoresis:

Fragments separated by size in a capillary tube.

-Detection & Analysis:

Laser excites fluorescent tags; software generates chromatogram.



Applications, Pros/Cons, and Conclusion

-Applications:

Genetic disease diagnosis, forensic analysis, microbial ID, mutation detection.

-Advantages:

High accuracy (~99.99%), reliable for short reads (up to 1,000 bp).

-Limitations:

Low throughput, costly/time-consuming for large genomes.

-Conclusion:

Still vital for small-scale projects and validating the (NGS) results.



	Sanger Sequencing	NGS
Accuracy	99.9%	99-99.9%
Cost Effectiveness	< 20 samples	> 20 samples
Speed < 20 Samples	Fast	Slow
Speed >20 Samples	Slow	Fast
Sensitivity	15-20%	1%
Sample Coverage	1 Read/Sample (300-850 bp)	Billions of Reads/Sample (Up to 16 Tera bite)