

พันธุวิศวกรรมและการโคลนนิ่ง

DNA fingerprinting

Gel electrophoresis

DNA sequencing

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หลักสูตรชีววิทยา

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Polymerase Chain Reaction (PCR)

- Template DNA
- DNA polymerase
- Primers (forward and reverse)
- Deoxynucleoside triphosphates (dNTPs: dATP, dCTP, dGTP, and dTTP)
- Required cofactor for activity of DNA polymerases: Mg^{2+}
- Buffer: suitable chemical environment for activity of DNA polymerase



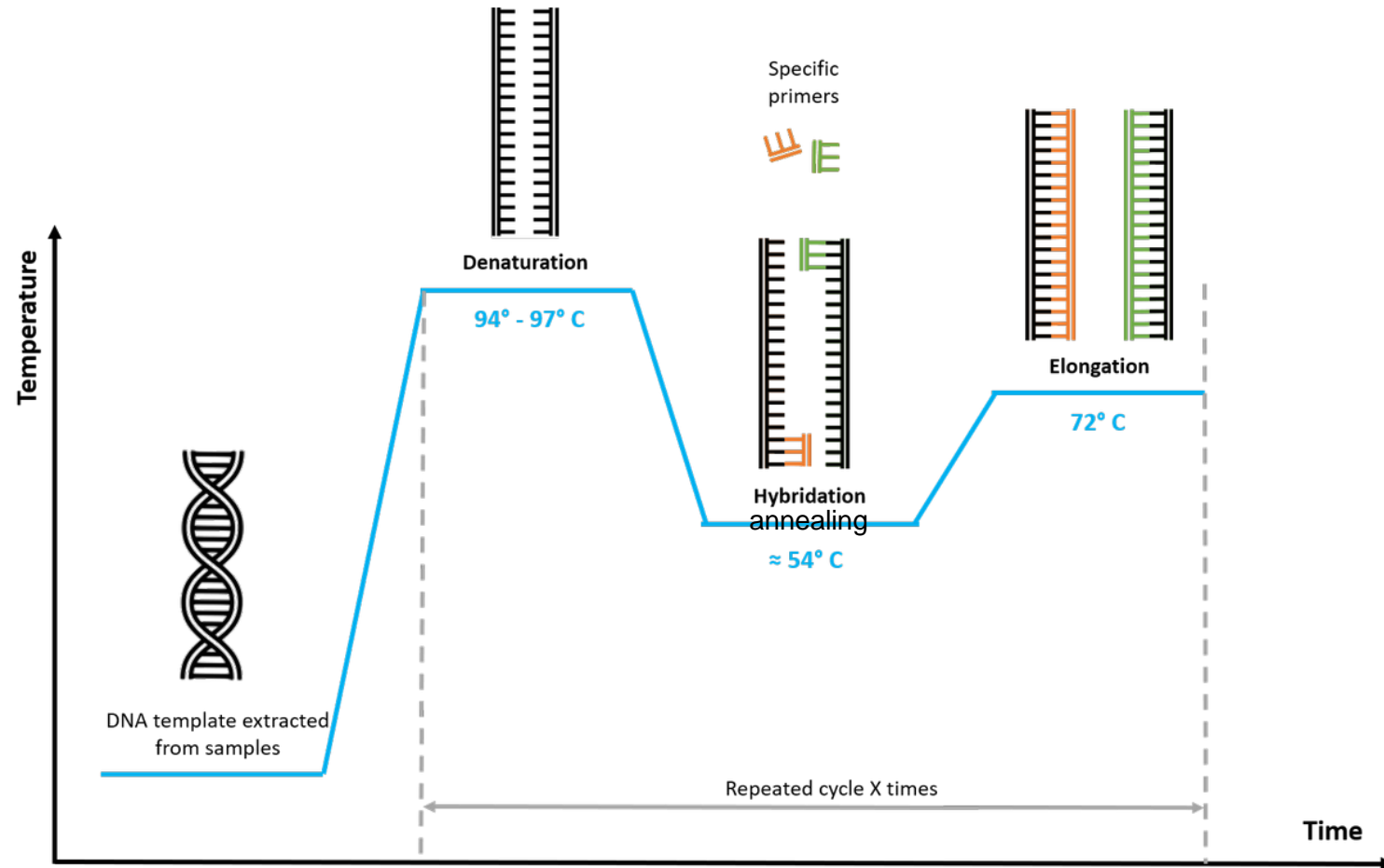
Polymerase Chain Reaction (PCR)

PCR thermocycler



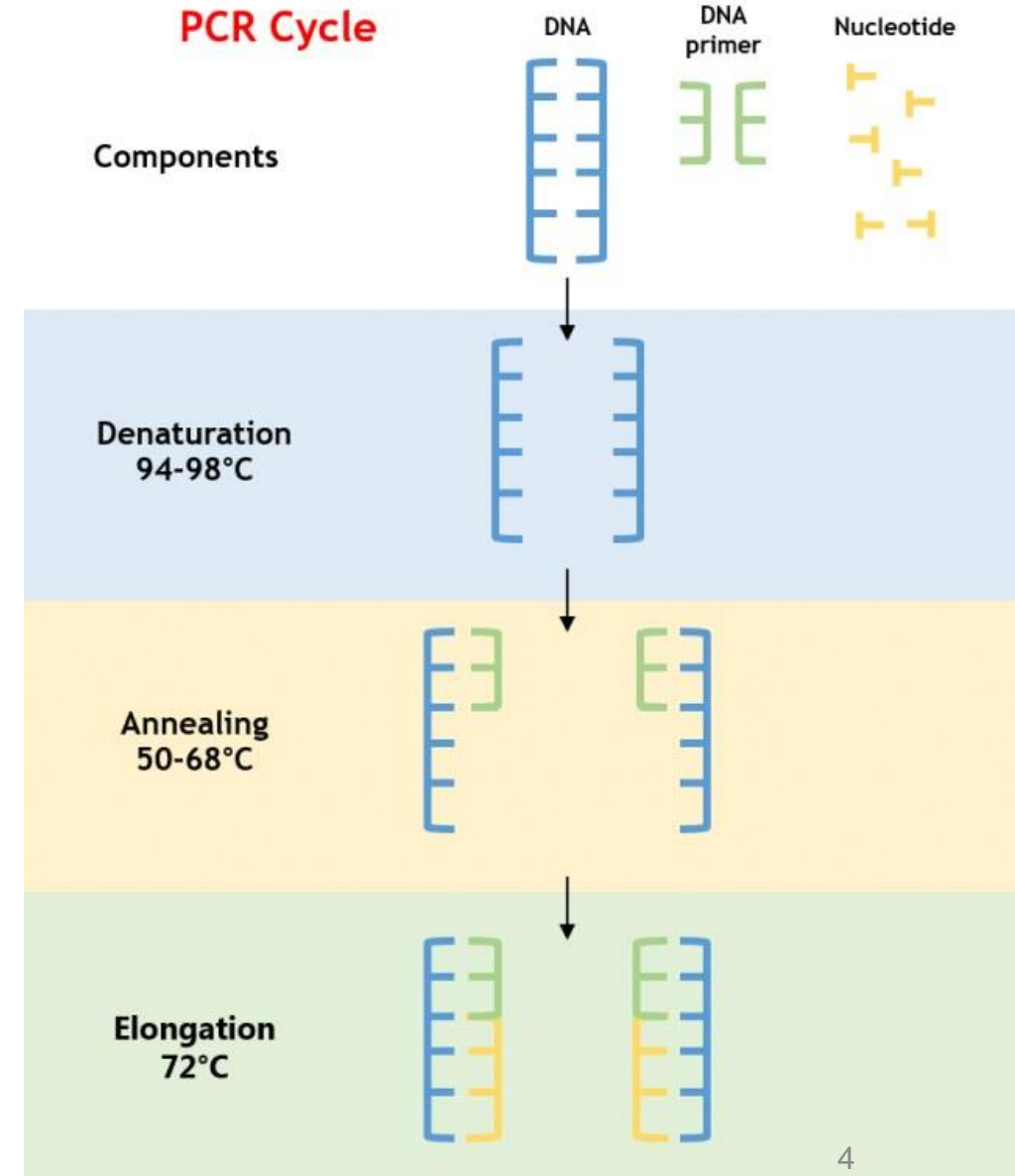
Polymerase Chain Reaction (PCR)

Steps

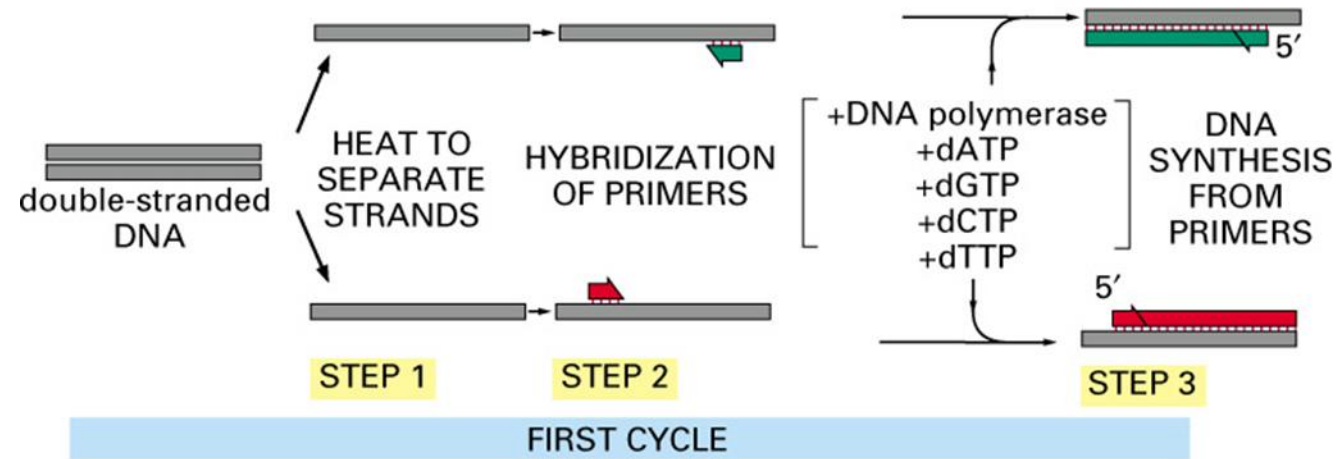


PCR Cycle

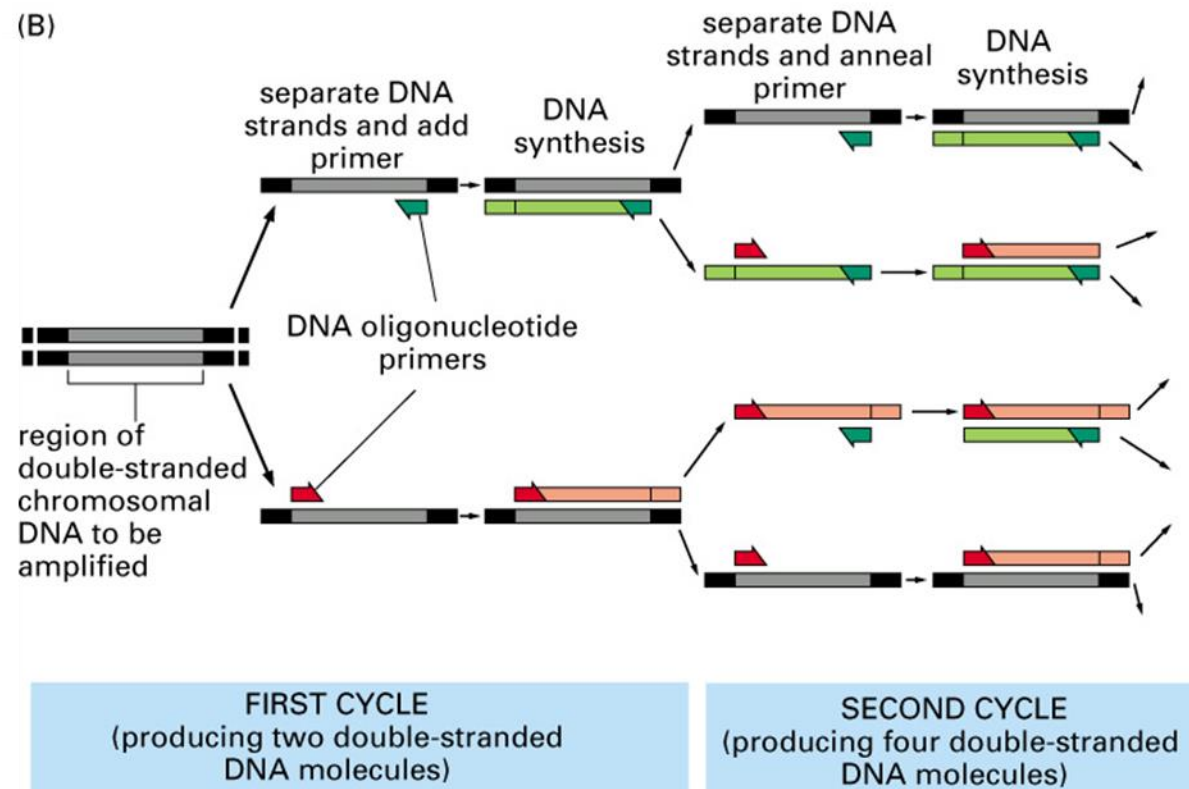
Components



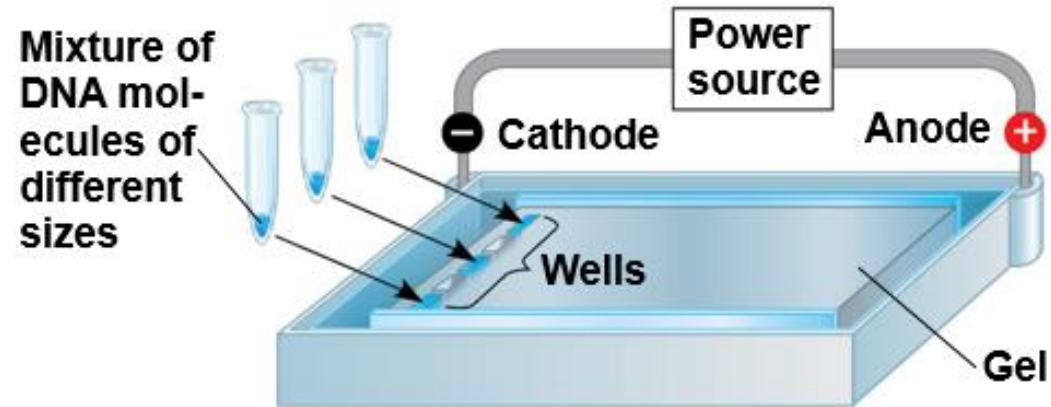
(A)



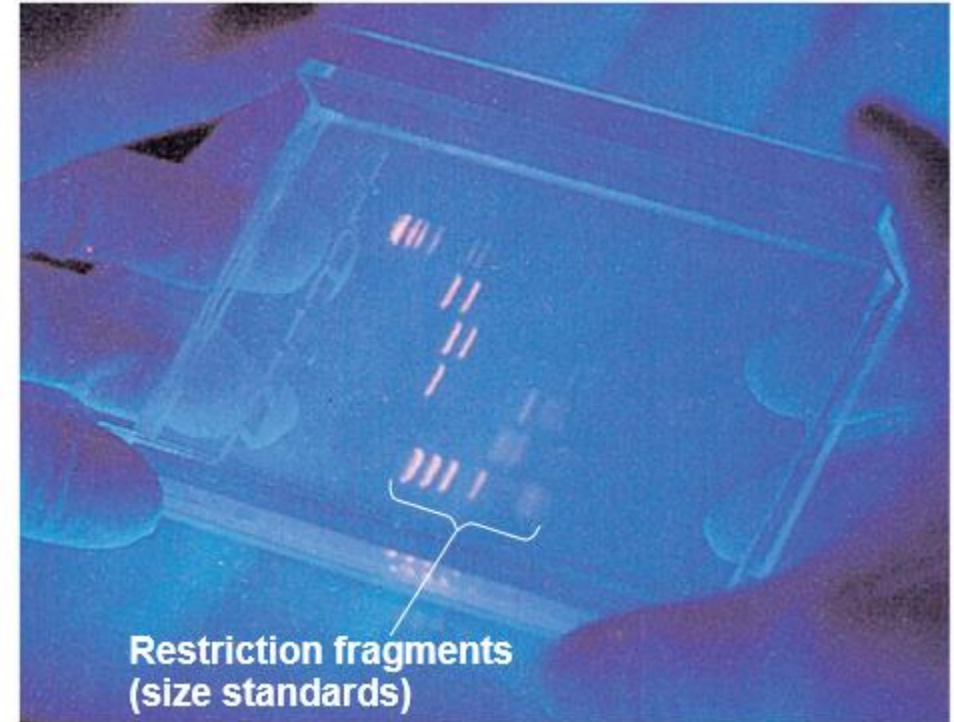
(B)



Gel Electrophoresis



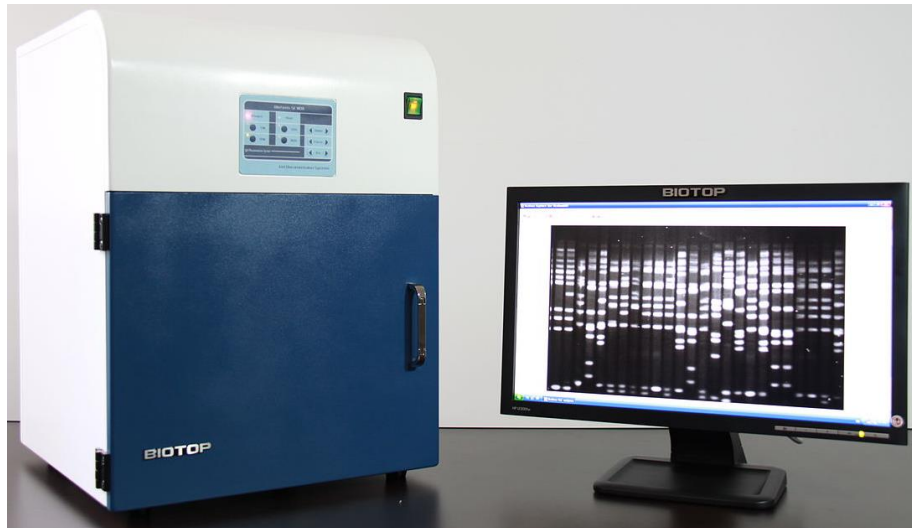
a) Negatively charged DNA molecules will move toward the positive electrode.

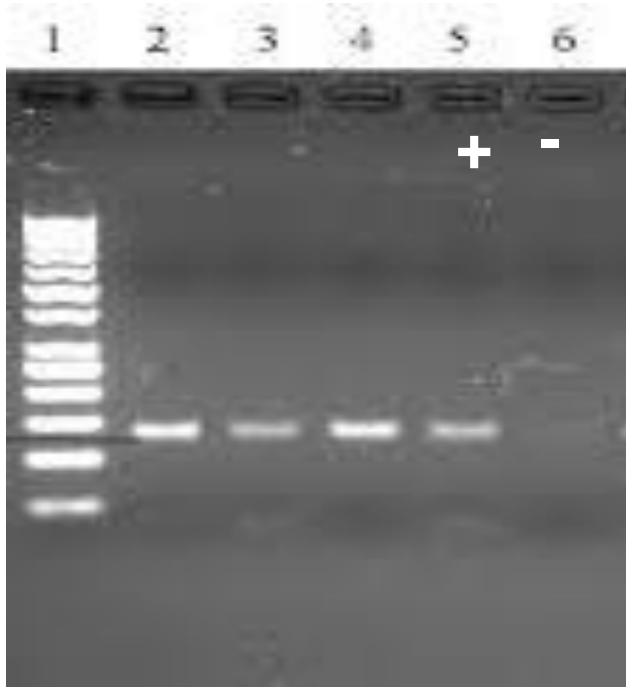


b) Shorter molecules are slowed down less than longer ones, so they move faster through the gel.

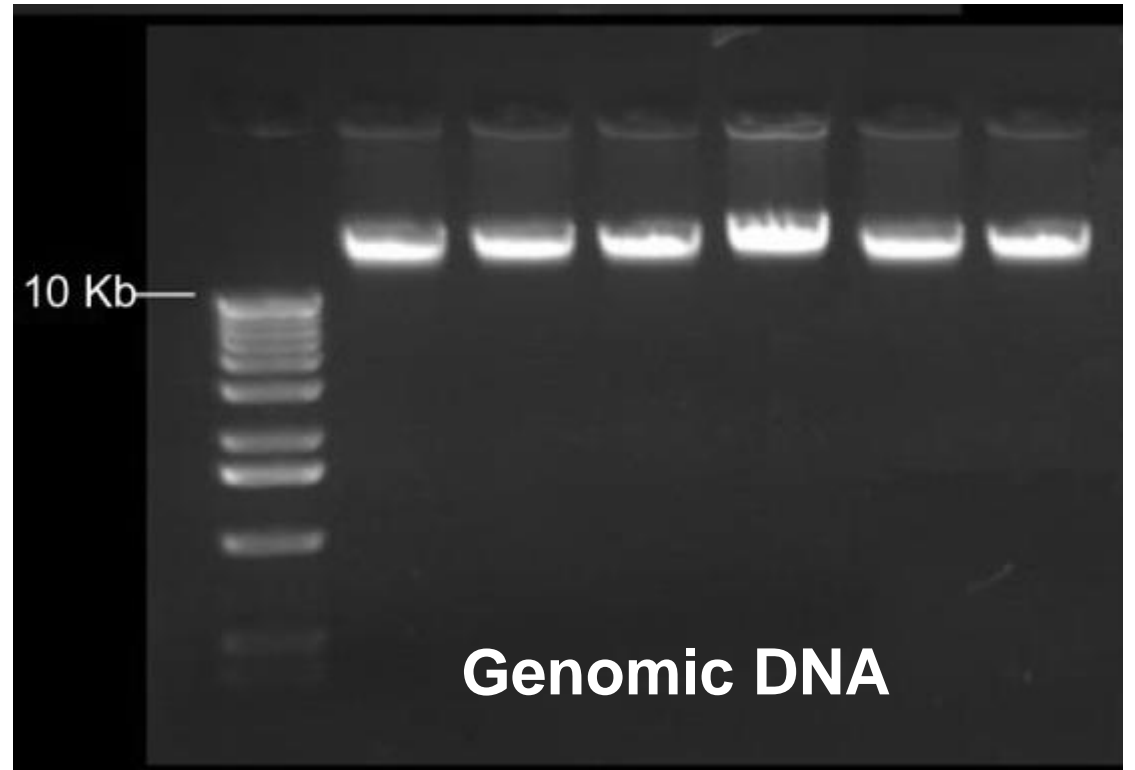


Gel documentation system



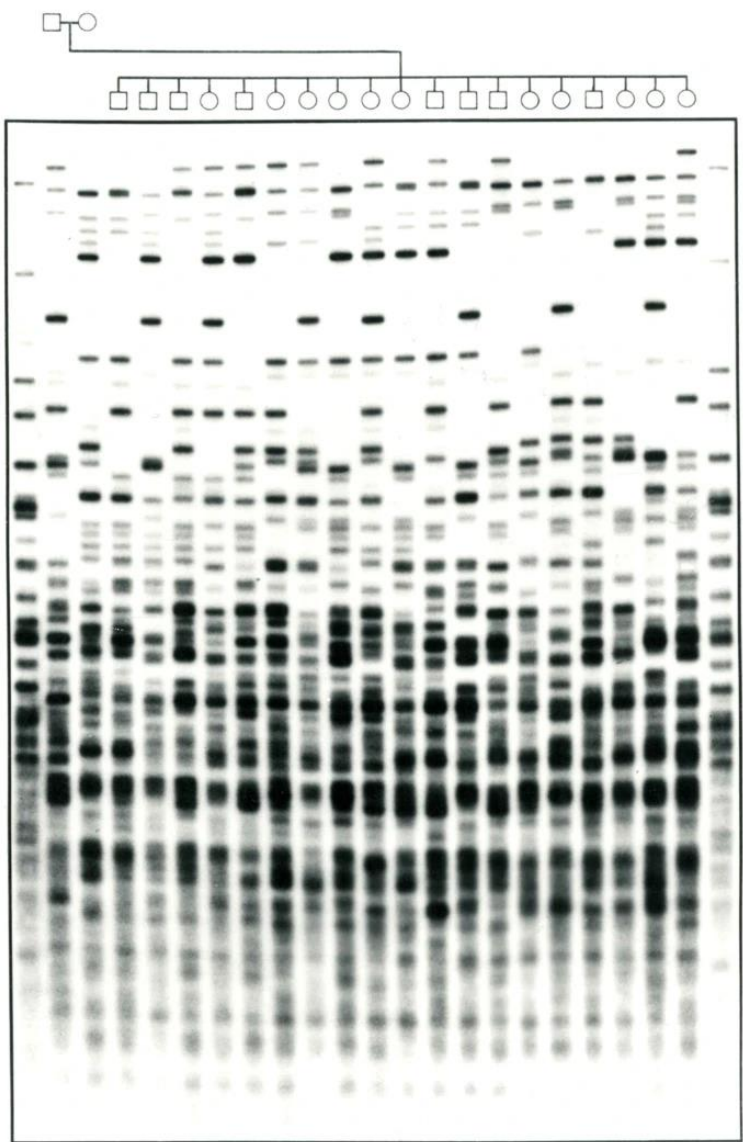


**PCR product:
Amplicon**

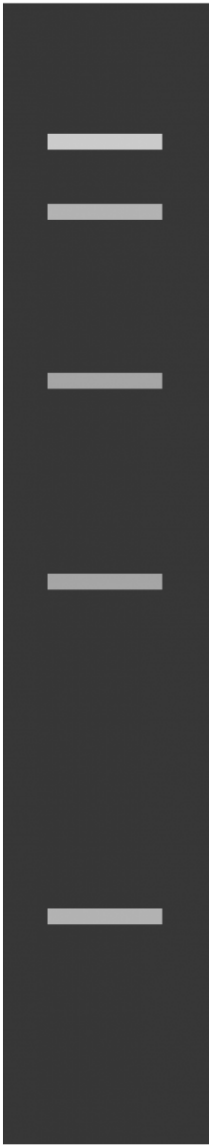


Genomic DNA

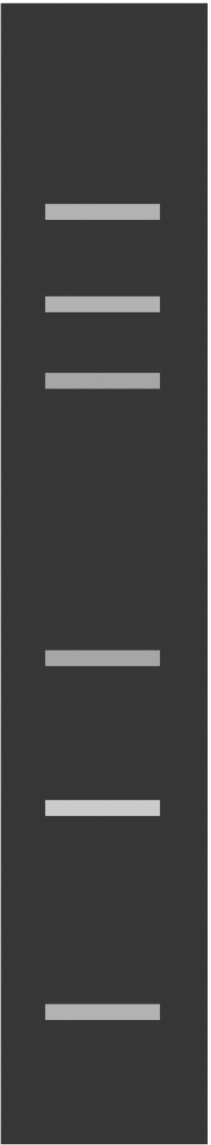
DNA fingerprinting



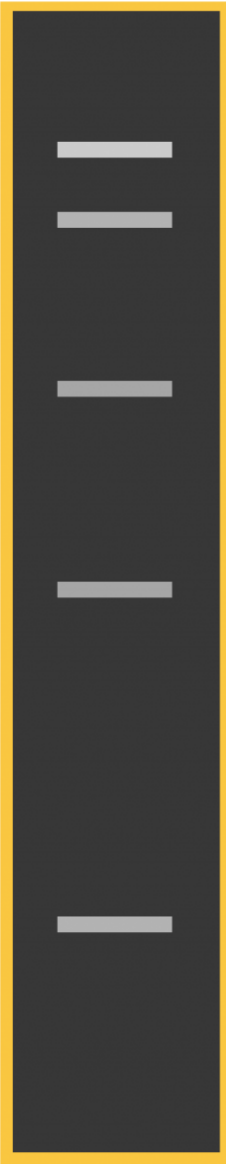
crime
scene



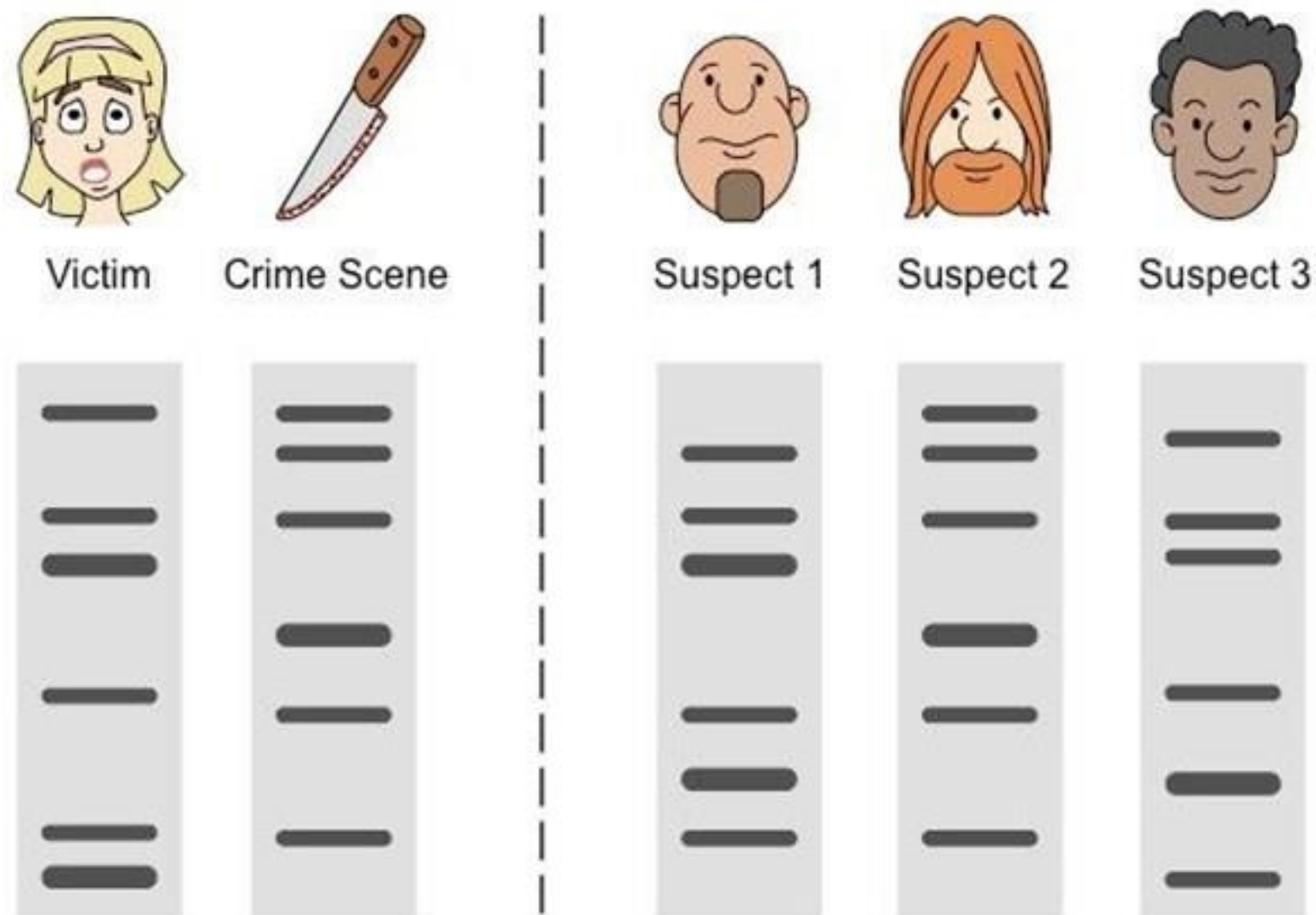
suspect 1



suspect 2



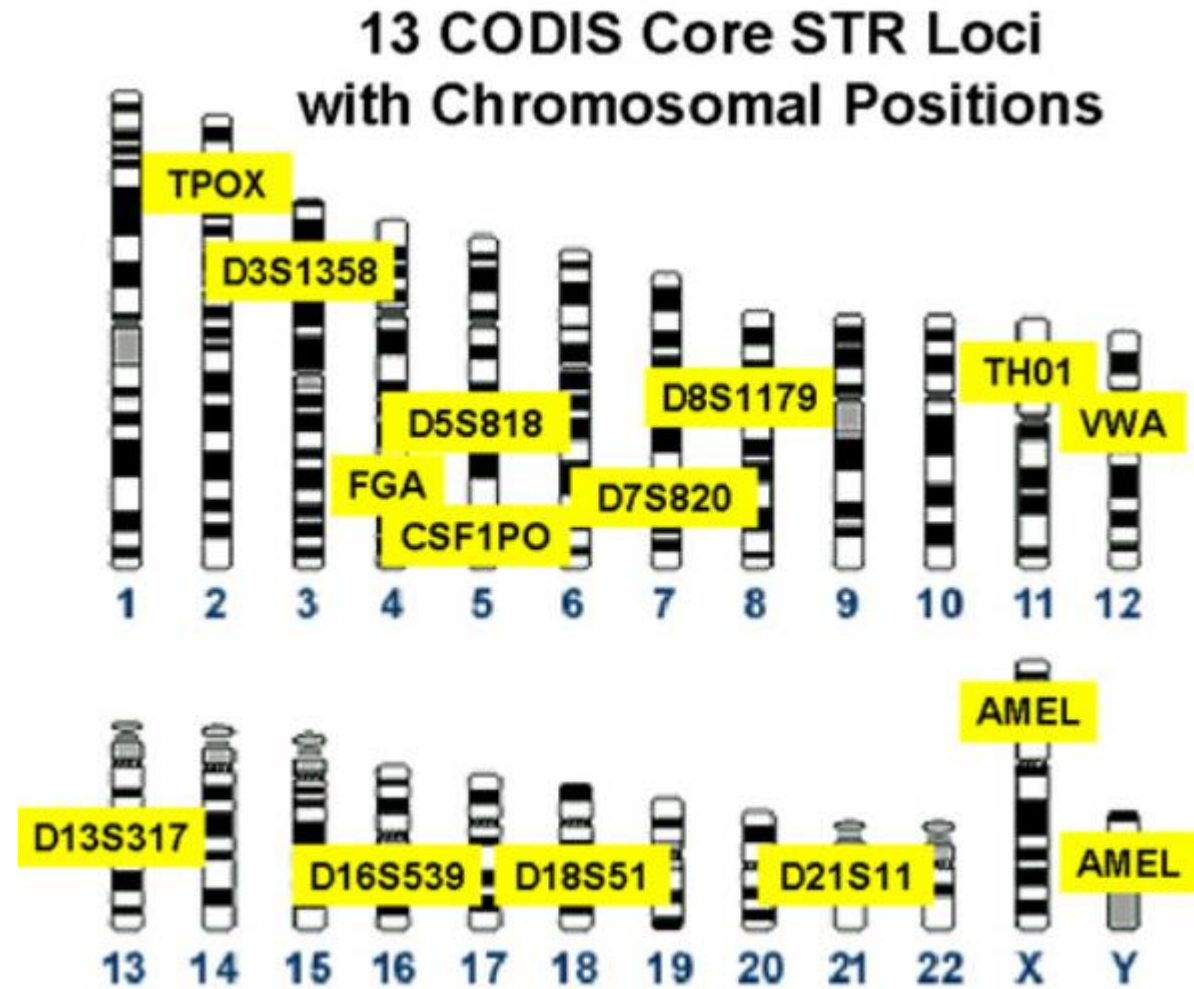
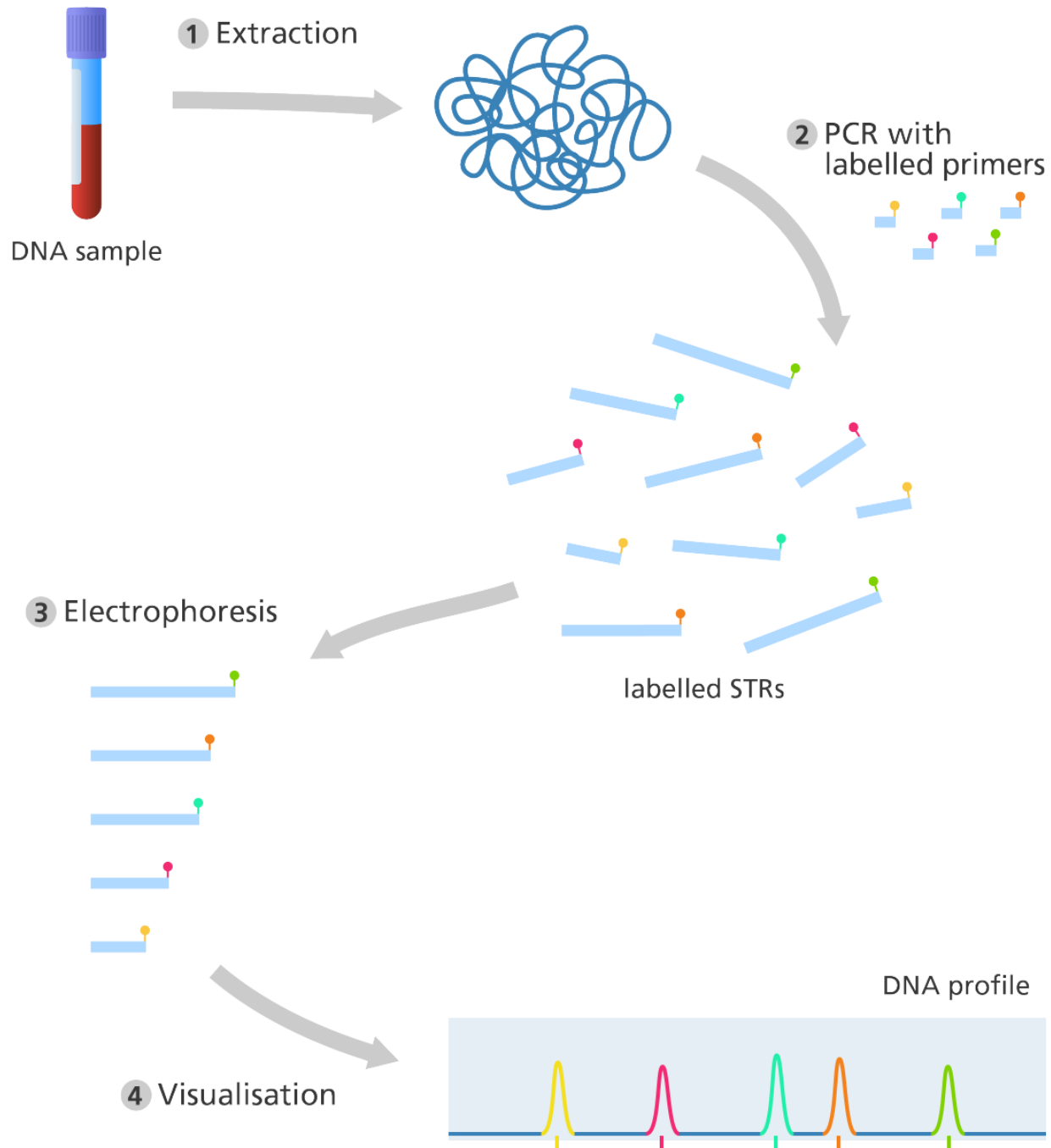
DNA fingerprinting



DNA fingerprinting

DNA repeats

- [illegible]



TPOX

Other Names	Chromosomal Location
hTPO, TPO UniSTS: 240638	2p25.3 ; intron 10 of human thyroid peroxidase gene Chr 2; 1.472 Mb (May 2004, NCBI build 35)

98 bp

102 bp

106 bp

110 bp

113 bp

114 bp

118 bp

122 bp

123 bp

125 bp

126 bp

130 bp

134 bp

135 bp

138 bp

142 bp

146 bp



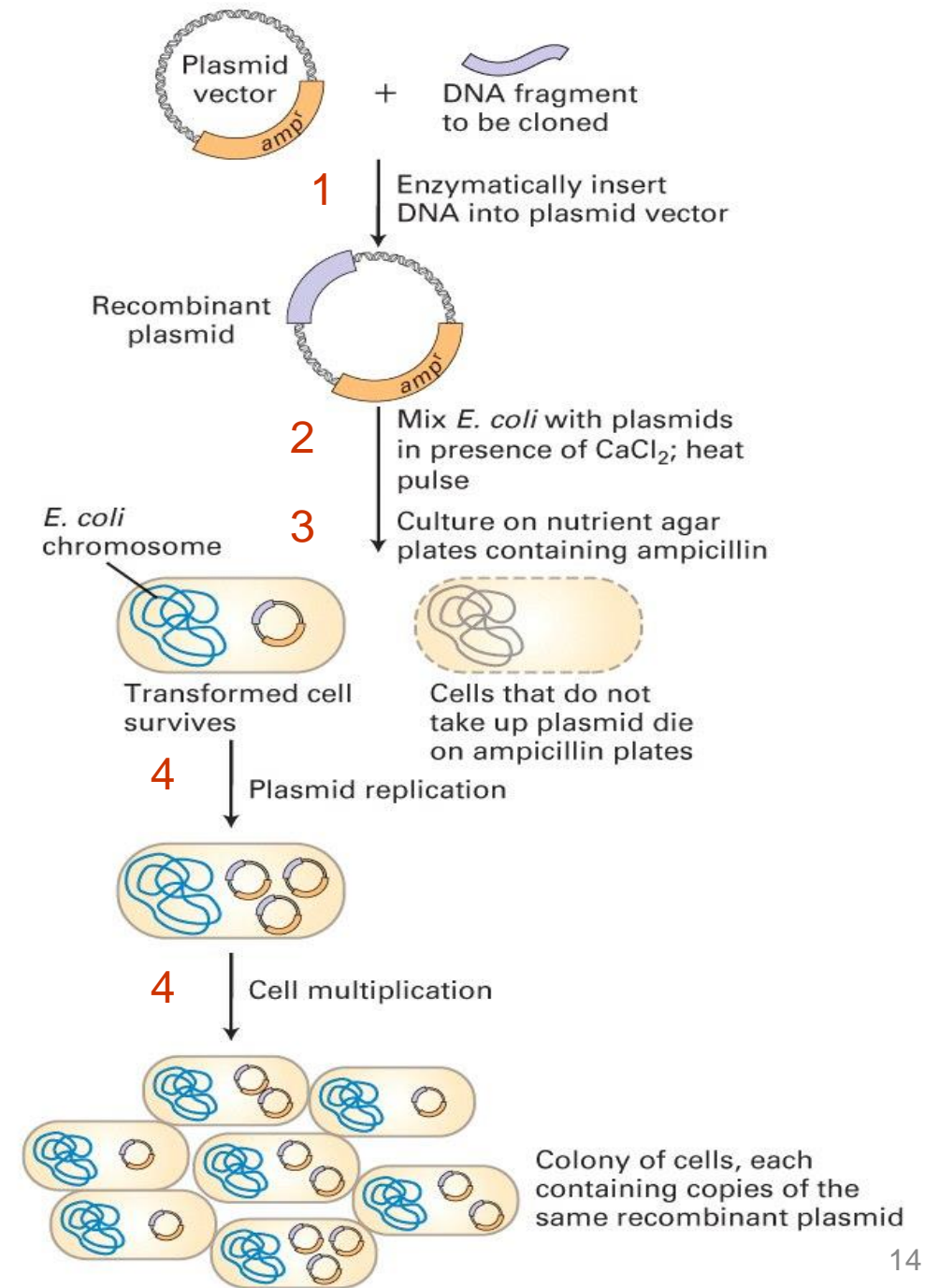
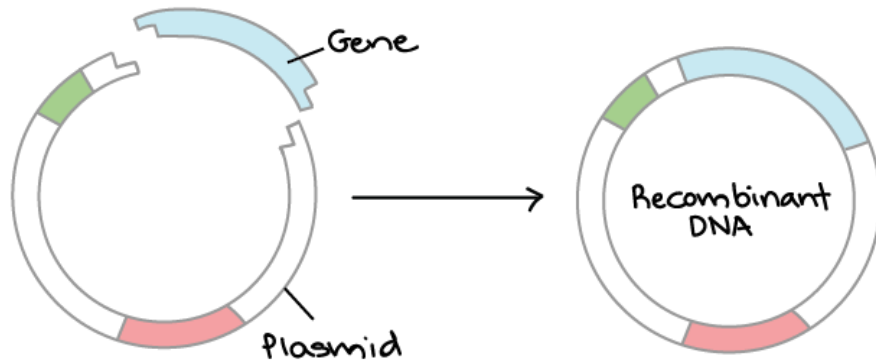
HOME UPDATES INFO + Human STRs +

FBI CODIS Core STR Loci:

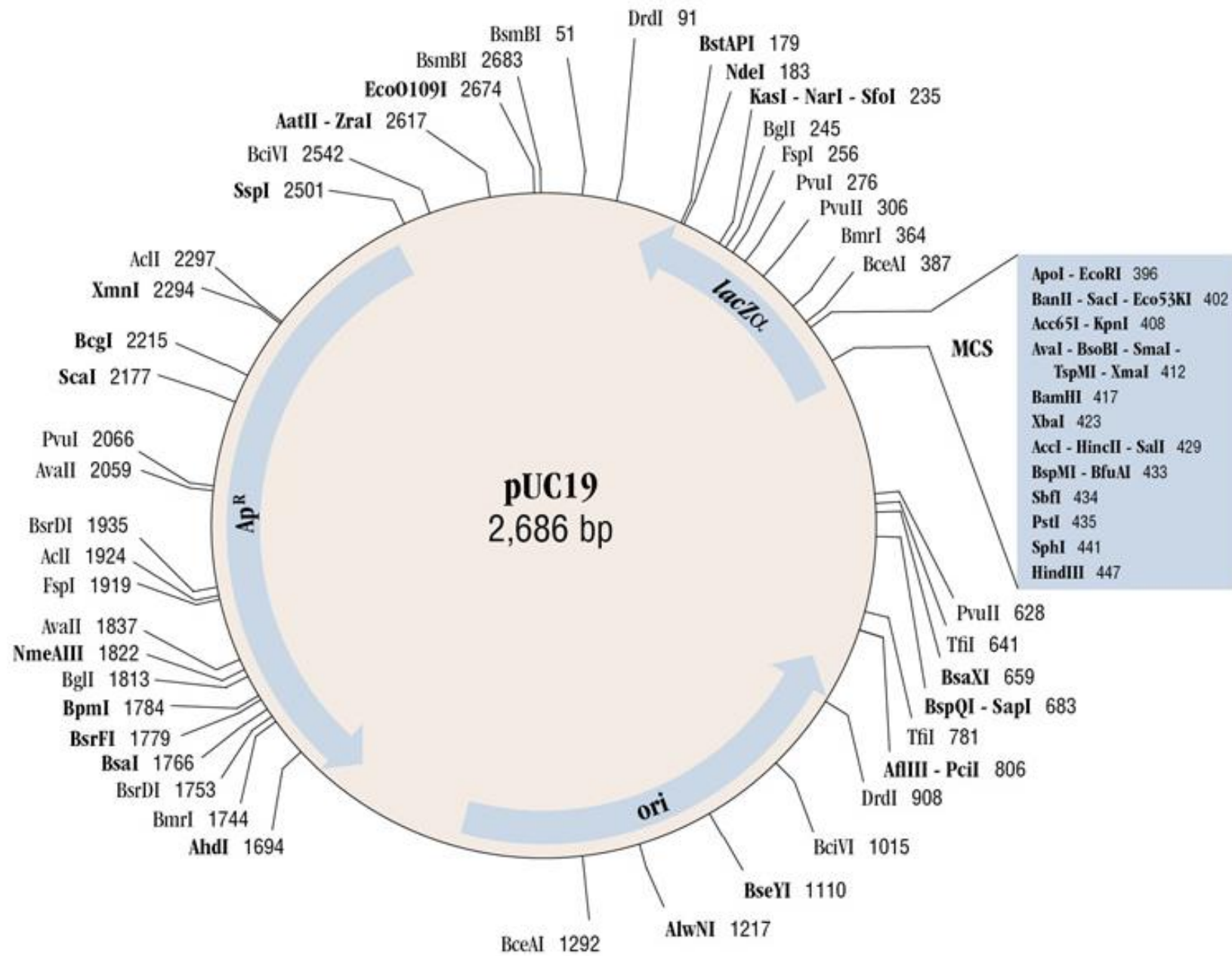


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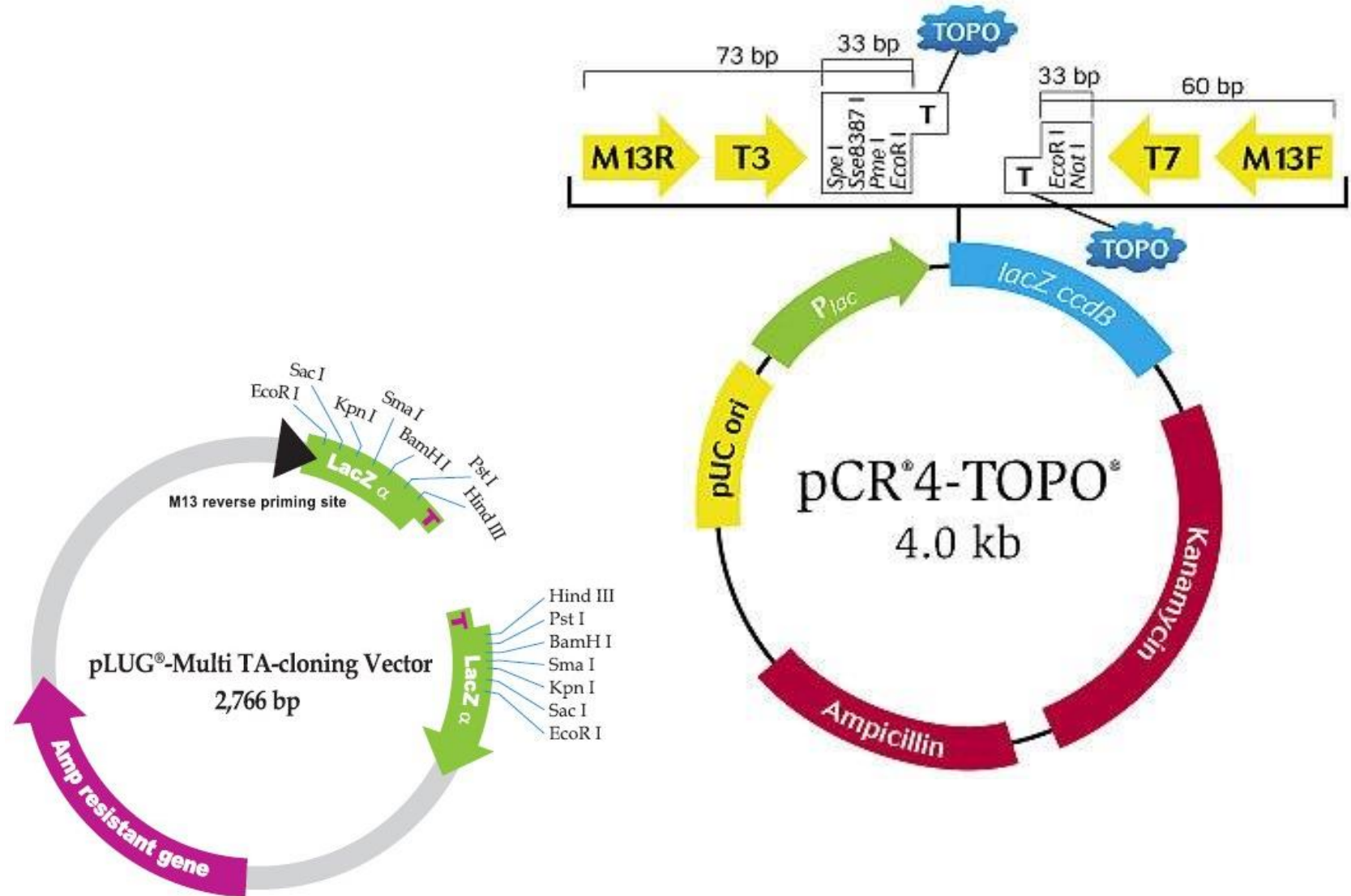
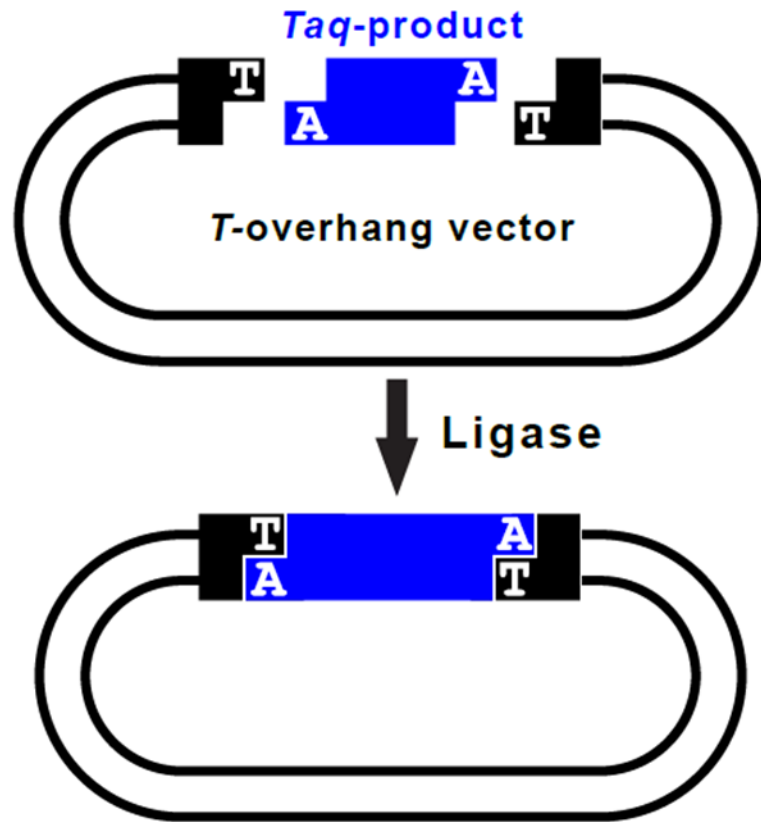
DNA cloning







DNA cloning

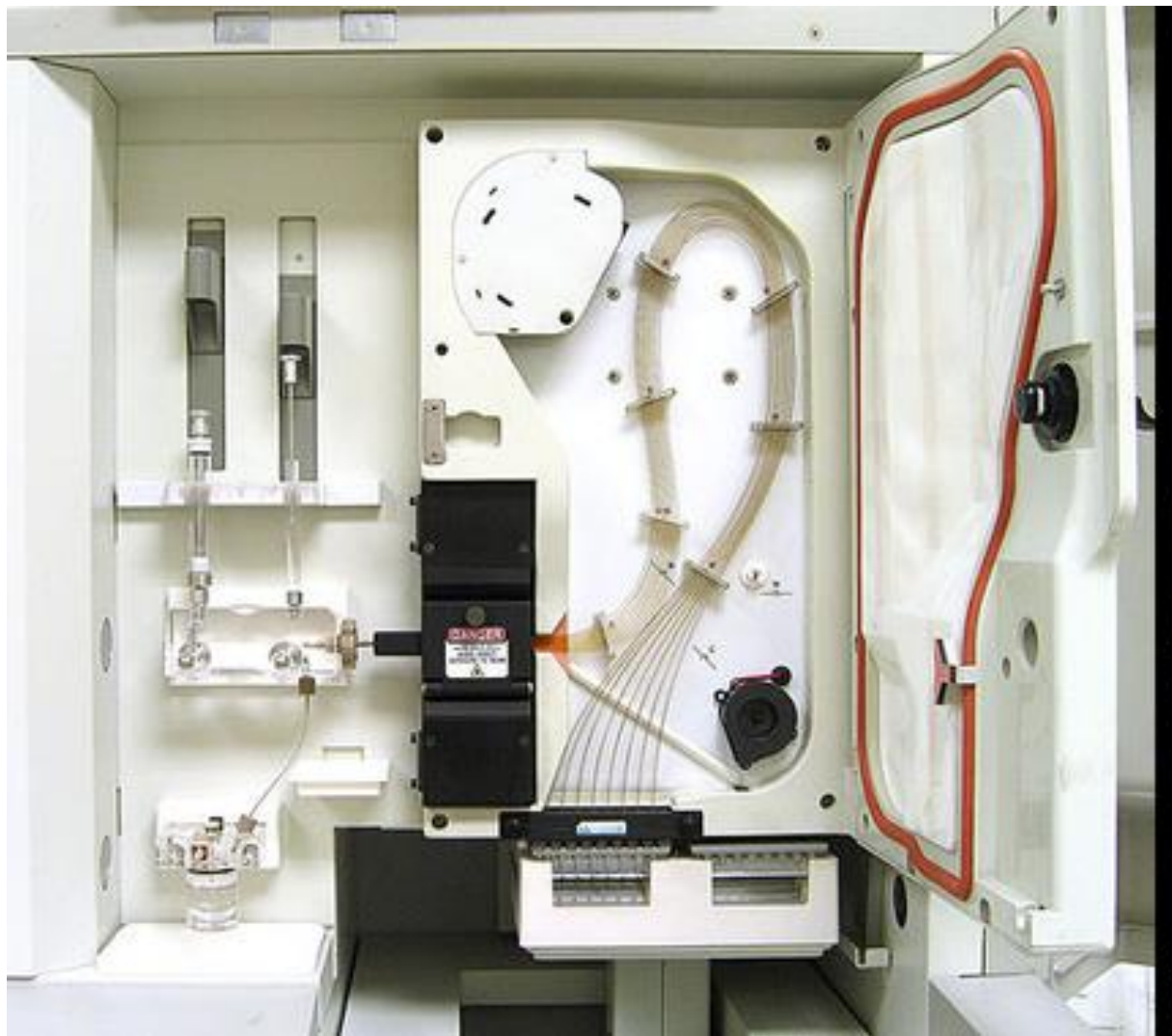


How do you sequence DNA?

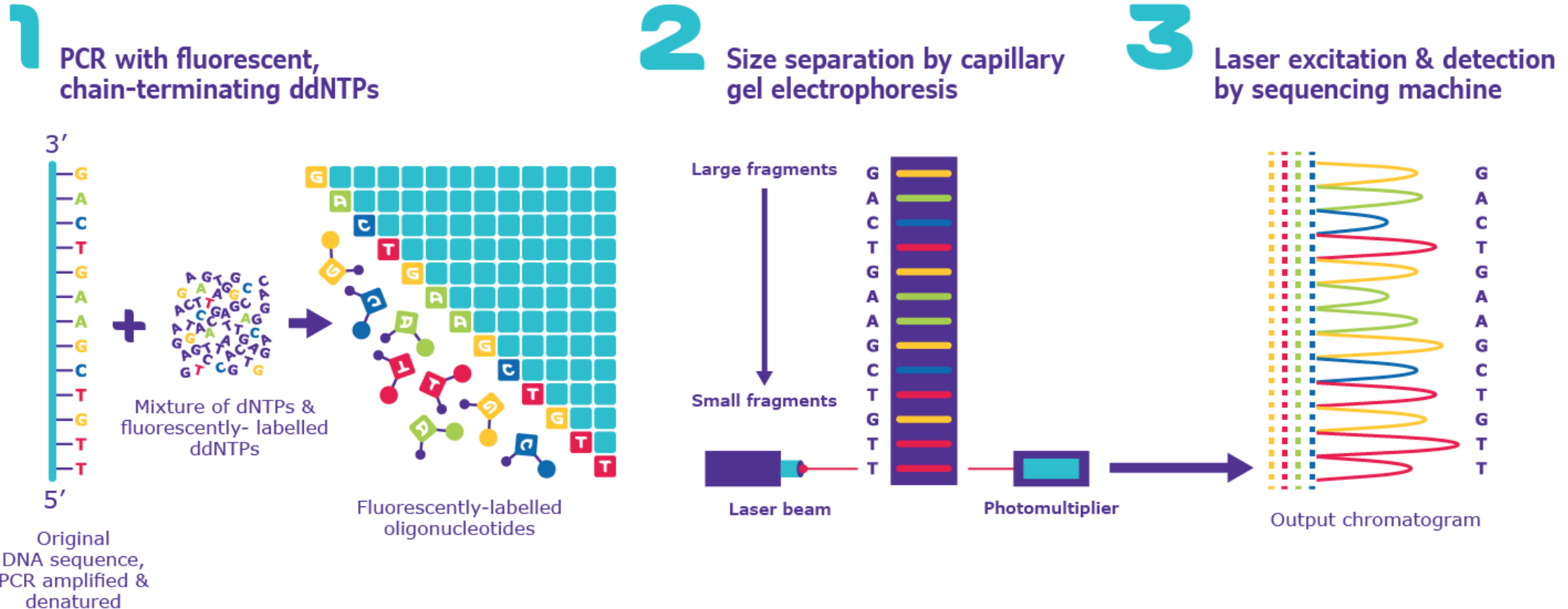
- Sanger (**dideoxy**, enzymatic) – developed by Frederick Sanger and is still used today with little change to the basic method.

High-throughput DNA sequencing

- **Next-generation sequencing**



1. All 4 fluorescently-labeled ddNTPs are used in 1 reaction, each a different “color”
2. Fragments are separated in matrix-filled capillary tubes, 1 capillary per reaction
3. Laser detects fluorescence automatically as each fragment exits capillary
4. Computer software “calls bases” and processes sequence files



Sequencing chromatogram

