

De l'hélice à l'électron.



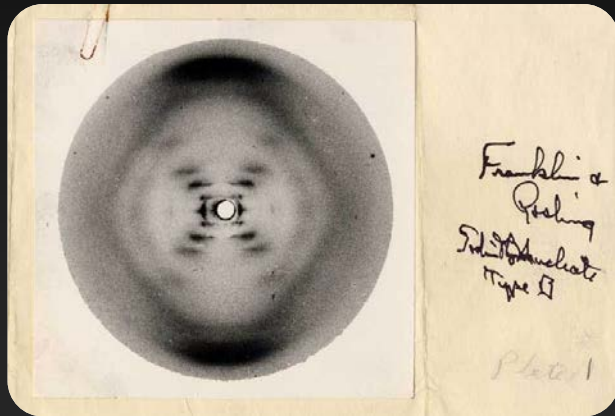
Dr. P. Beckers
Service de Génétique
Laboratoire de Biologie Moléculaire

Une situation surréaliste...

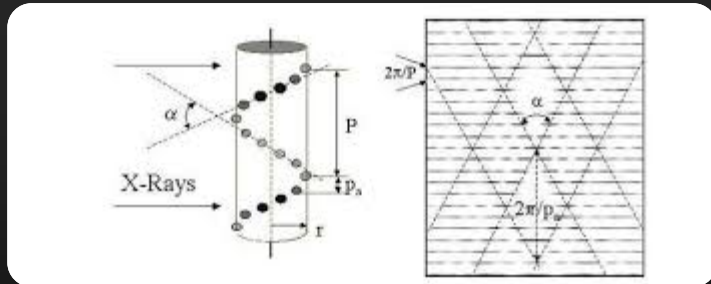
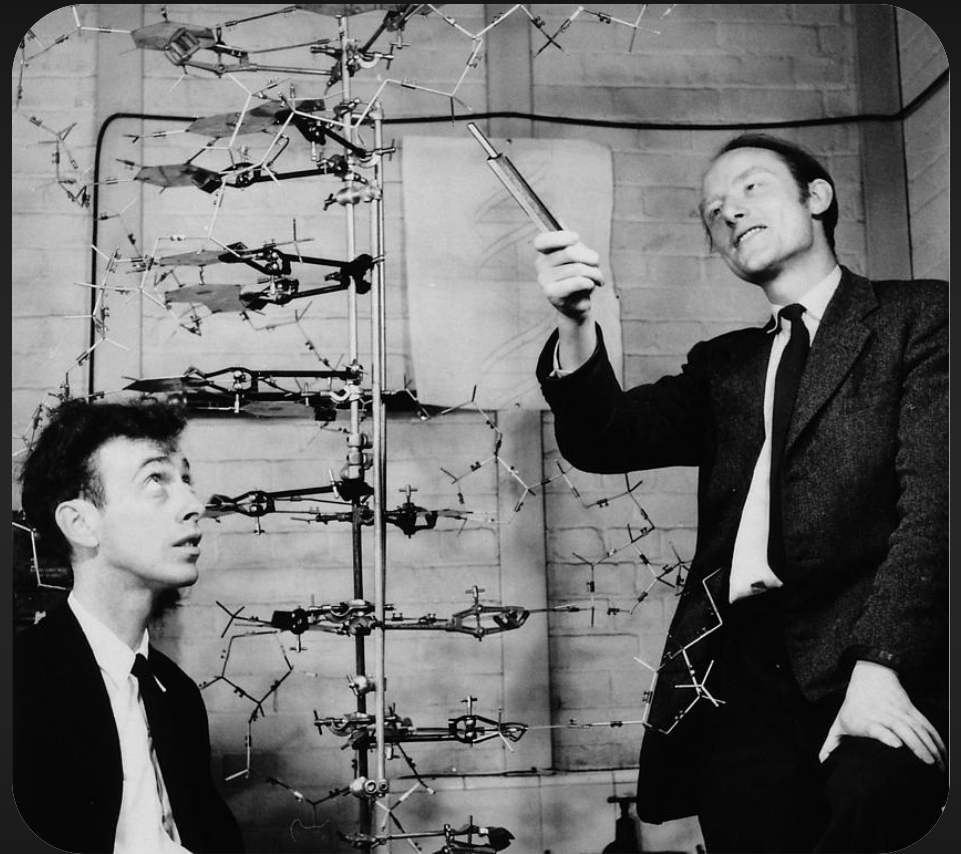
- Caryotypes
- FISH
- CGH
- MLPA
- Sanger
- NGS
- Nanopore



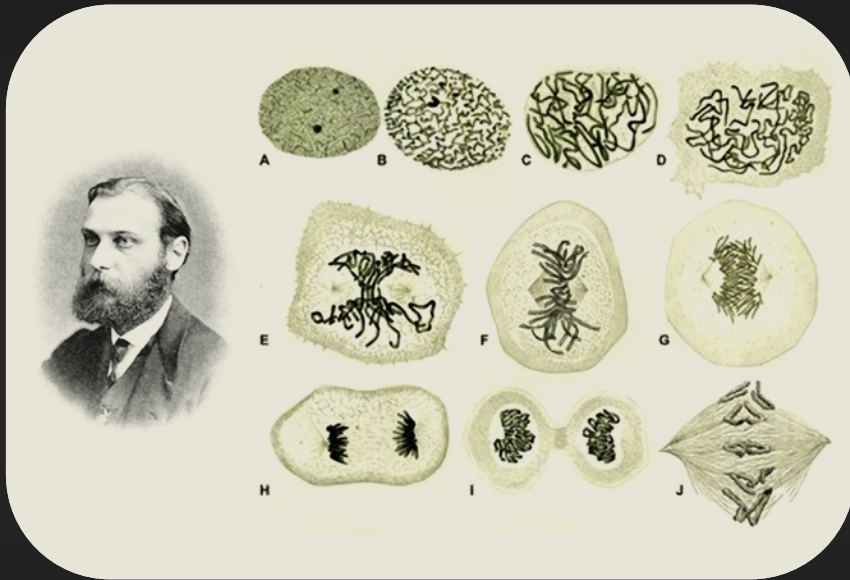
Watson et Crick - 1953



Maurice Wilkins et Rosalind Franklin



Au commencement...



Au sein de l'œuf, la matrice...

- **L'ADN** peut être extrait :
 - Des cellules nucléées (leucocytes)
 - Des cellules tumorales.
 - **Attention** : un patient ayant reçu une **greffe** de moëlle, nécessité le prélèvement d'autres cellules somatiques que les leucocytes...

1. Cytogénétique

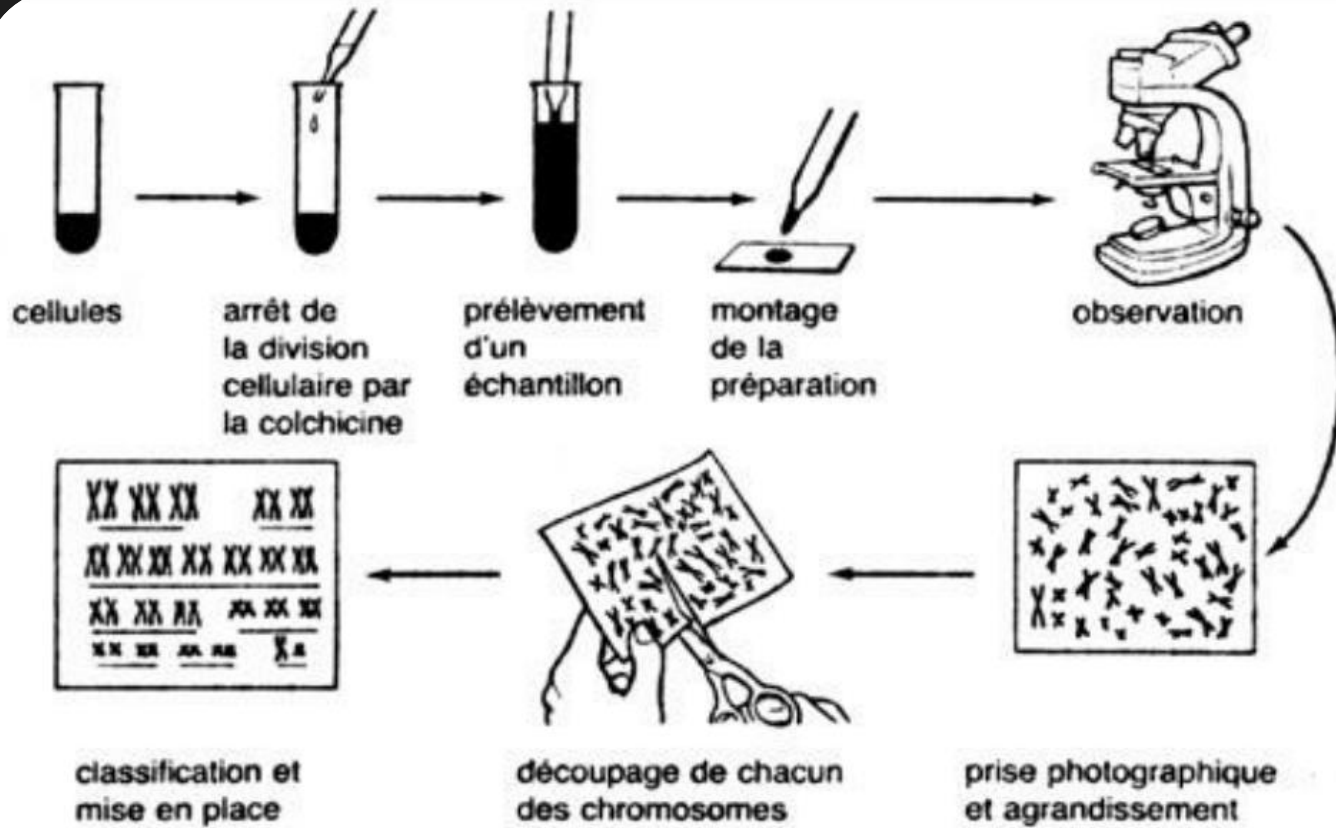
2. Génétique Moléculaire



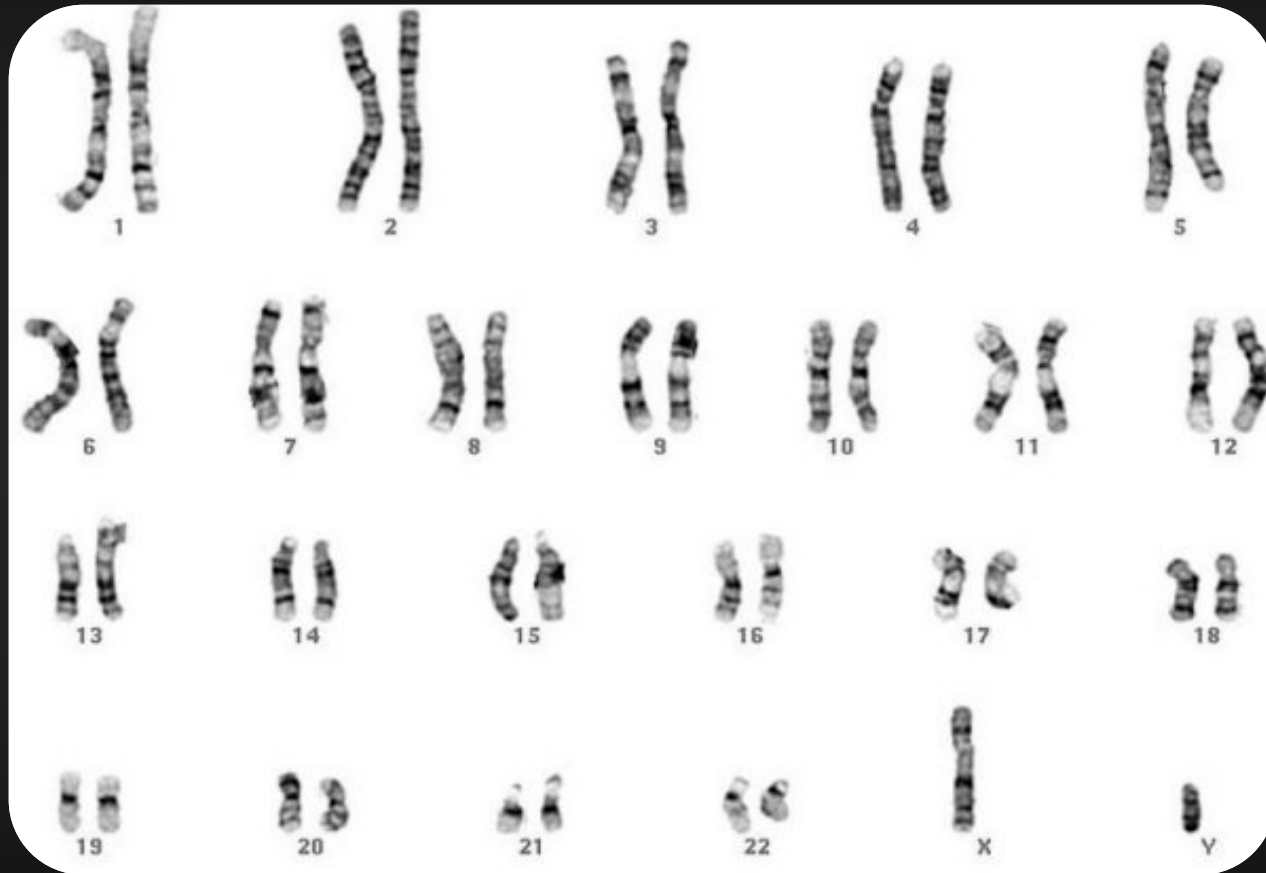
1. Cytogénétique

- Cytogénétique **conventionnelle** = Etude des chromosomes de la cellule en métaphase, par le :
 - **Caryotype**
- Cytogénétique **moléculaire** = Etude de fragments chromosomiques avec des sondes moléculaires fluorescentes.
 - **FISH** (*Fluorescence In Situ Hybridization*)
- Cytogénétique **moléculaire** = Etude de fragments chromosomiques par comparaison avec une séquence de référence.
 - **CGH** (*Comparative Genomic Hybridization*)

Caryotype



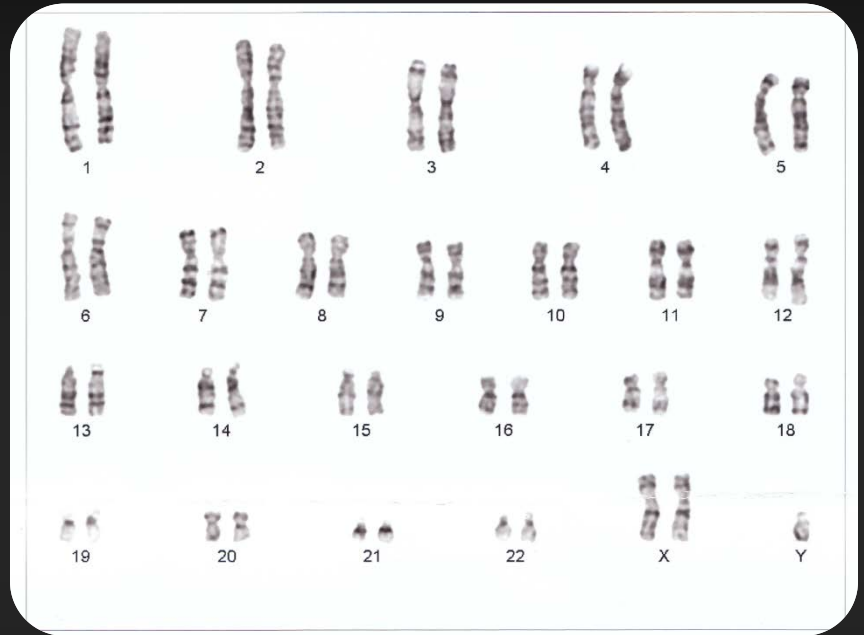
Caryotype



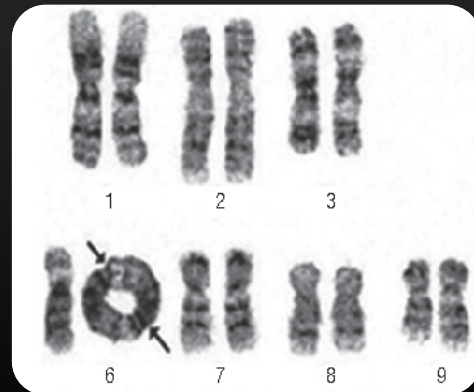
Caryotype



Turner

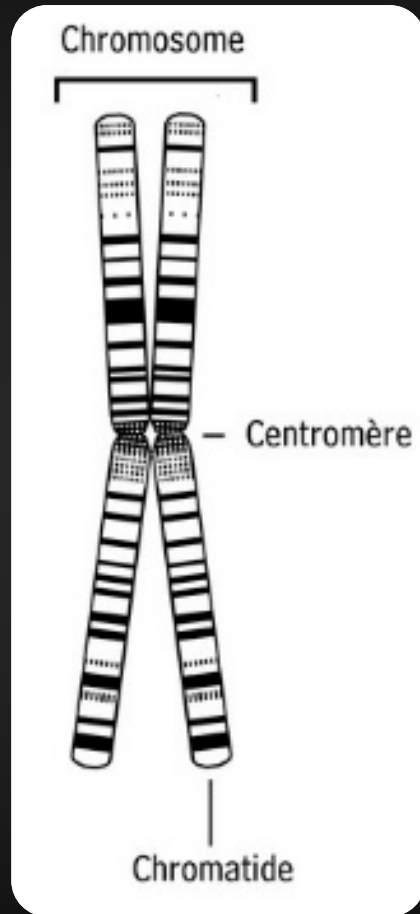


Klinefelter



O Ring

Caryotype



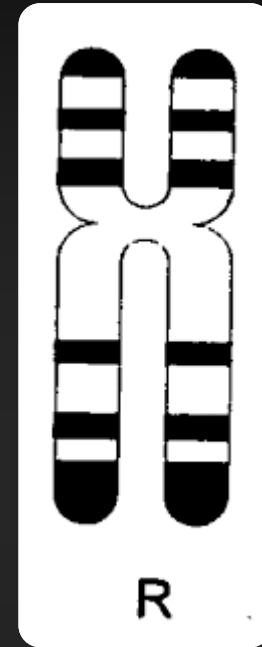
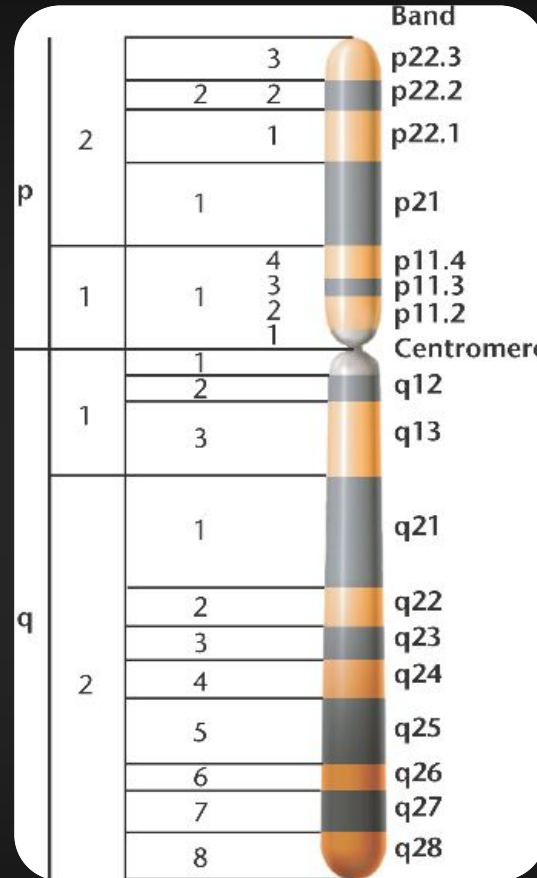
Caryotype



G

Bandes G
Extrémités claires

ADN riche en paires A-T



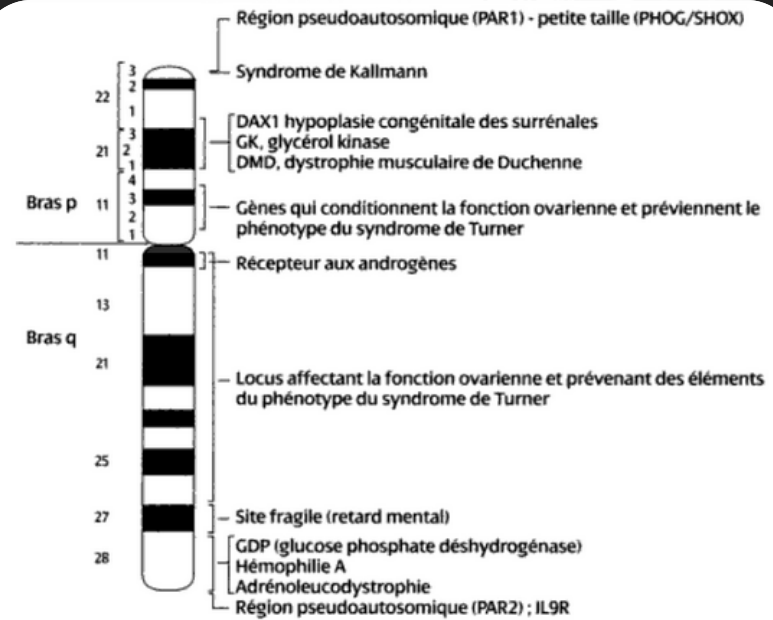
R

Bandes R (Reverse)
Extrémités foncées
ADN riche en paires G-C

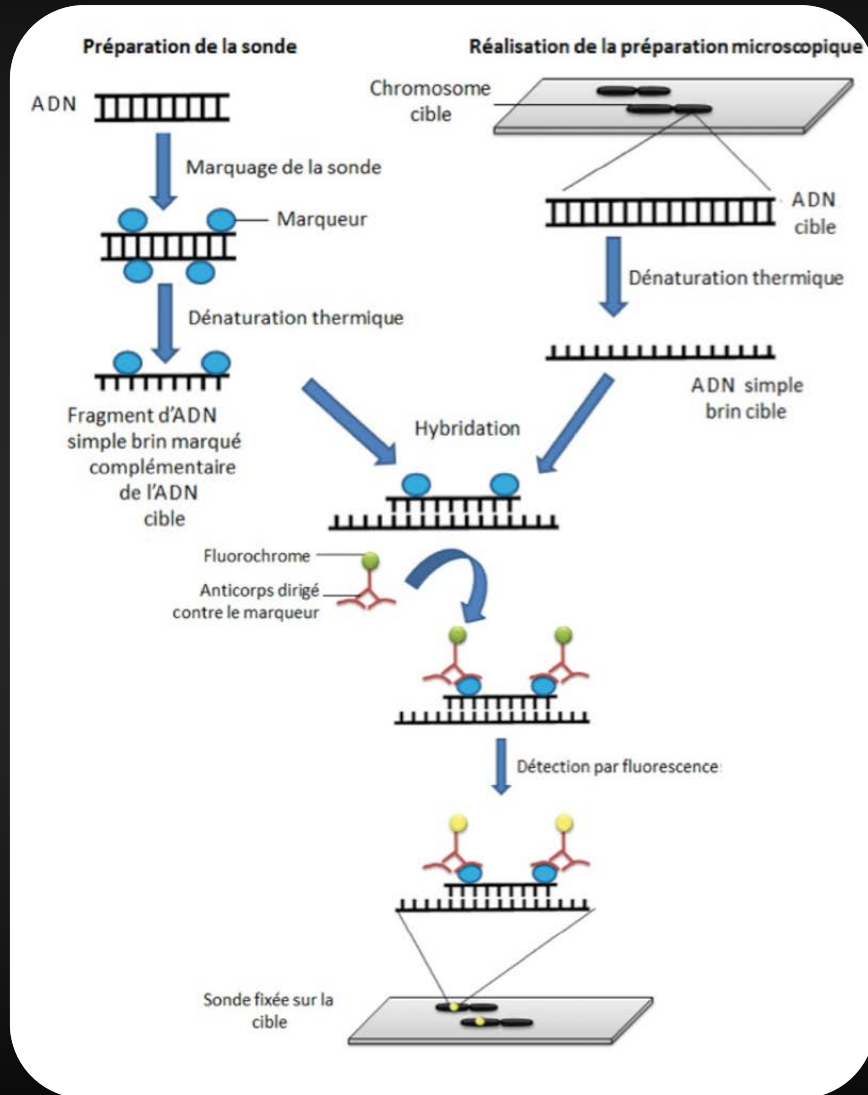
Caryotype



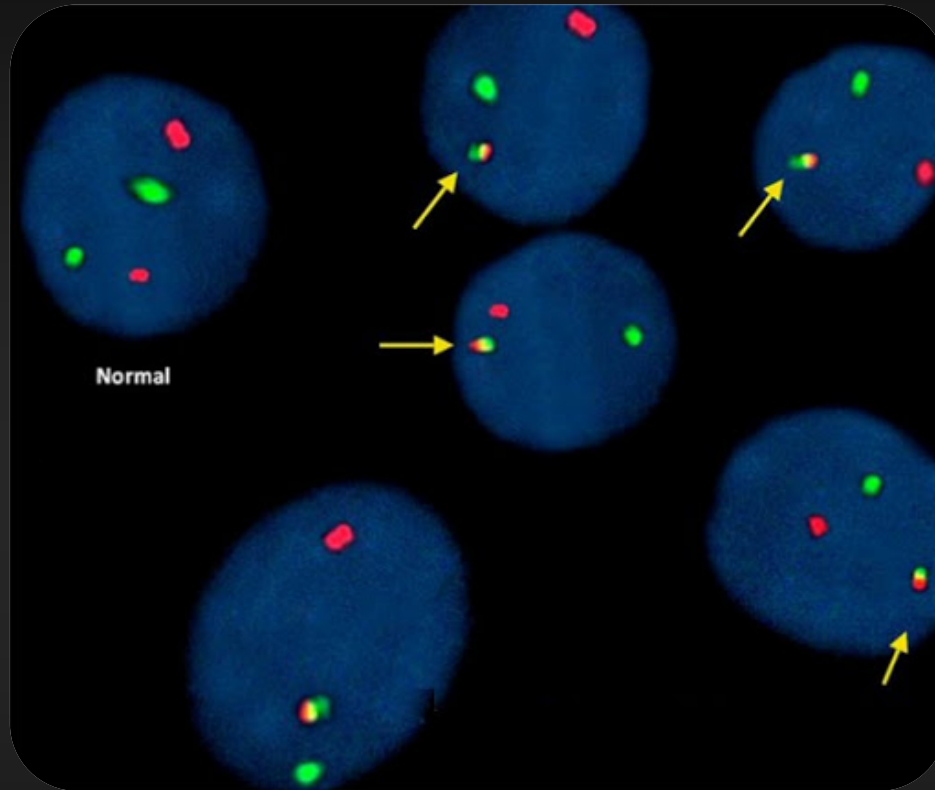
Exemple Chromosome X



Fluorescence In Situ Hybridization (FISH)

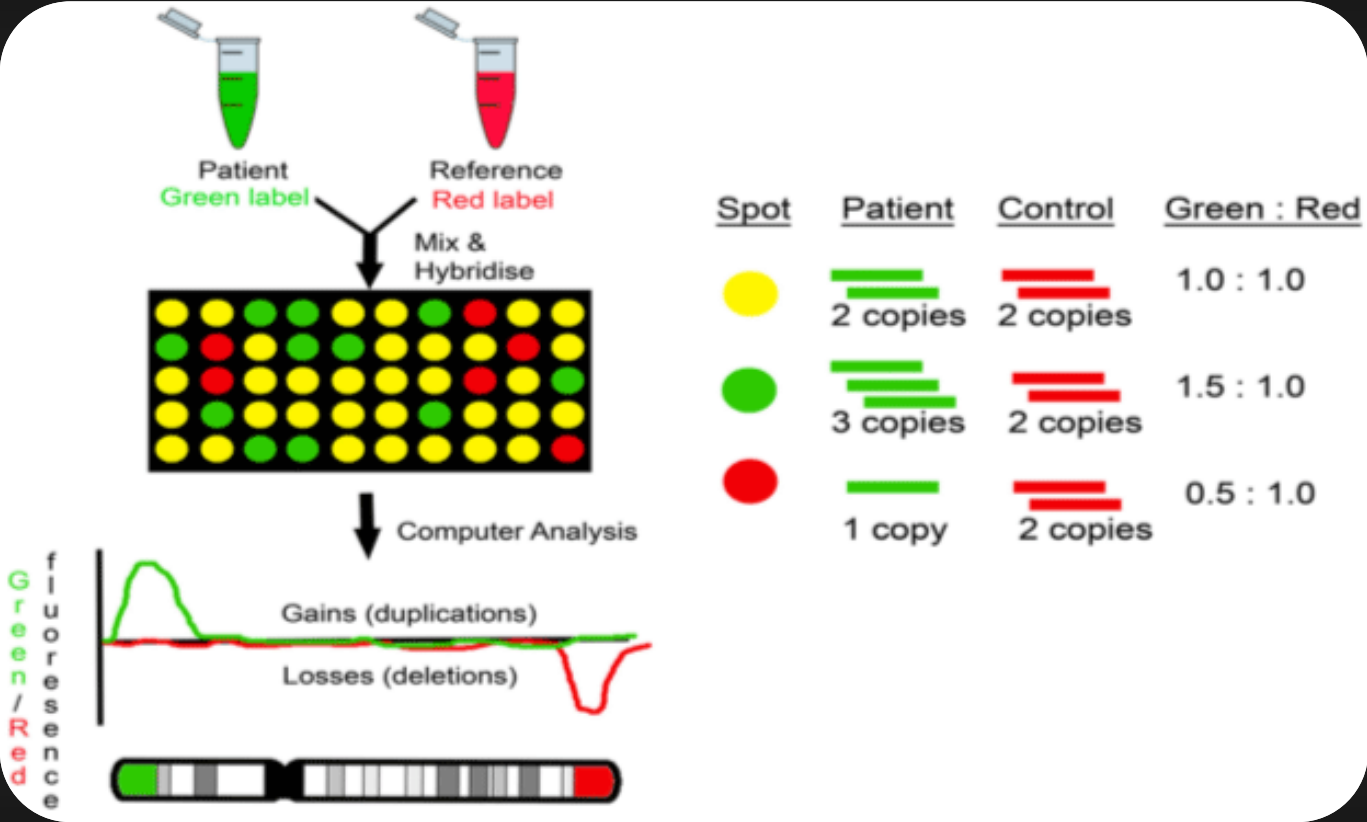


Fluorescence In Situ Hybridization (FISH)

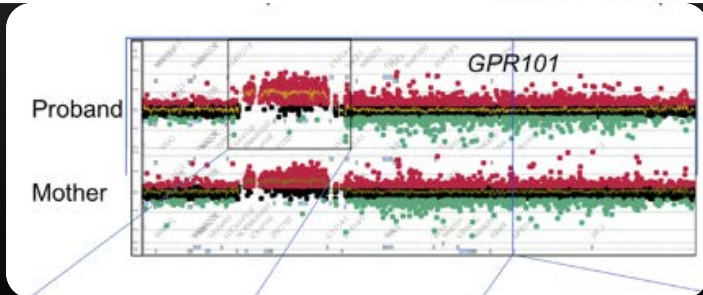
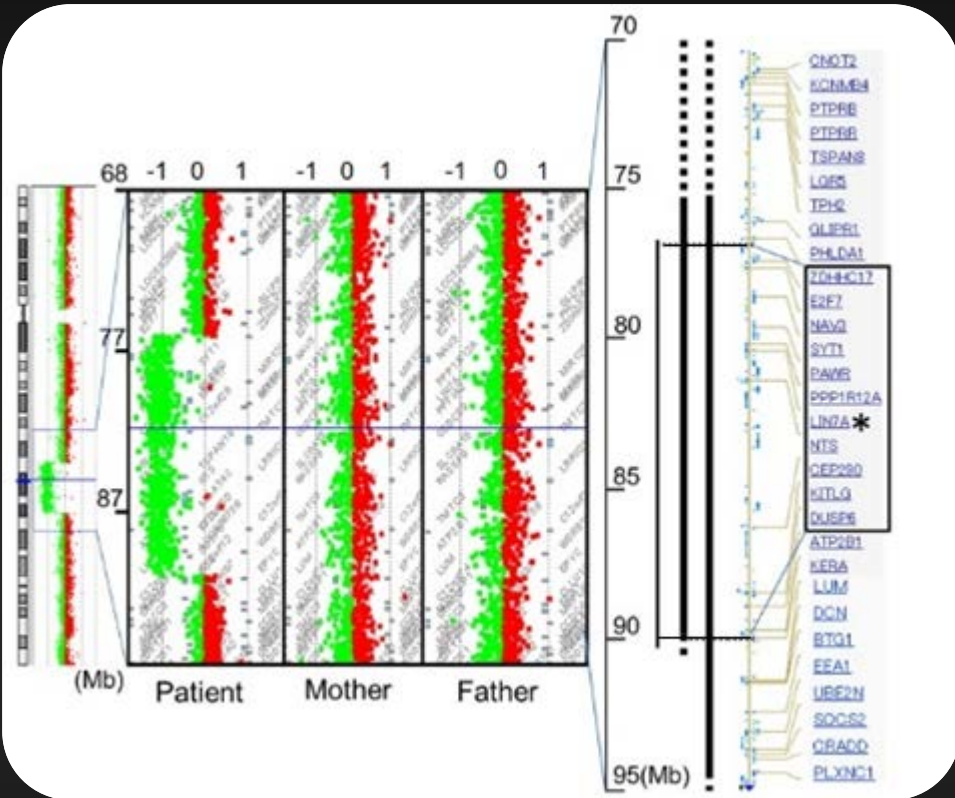


Translocation (7;12)

array-Comparative Genomic Hybridization (aCGH)



array-Comparative Genomic Hybridization (aCGH)



En résumé...



Caryotype
Full Génome
Basse résolution
5 à 10 Mb



FISH
Site Spécifique
Meilleure résolution
100 à 300 kb



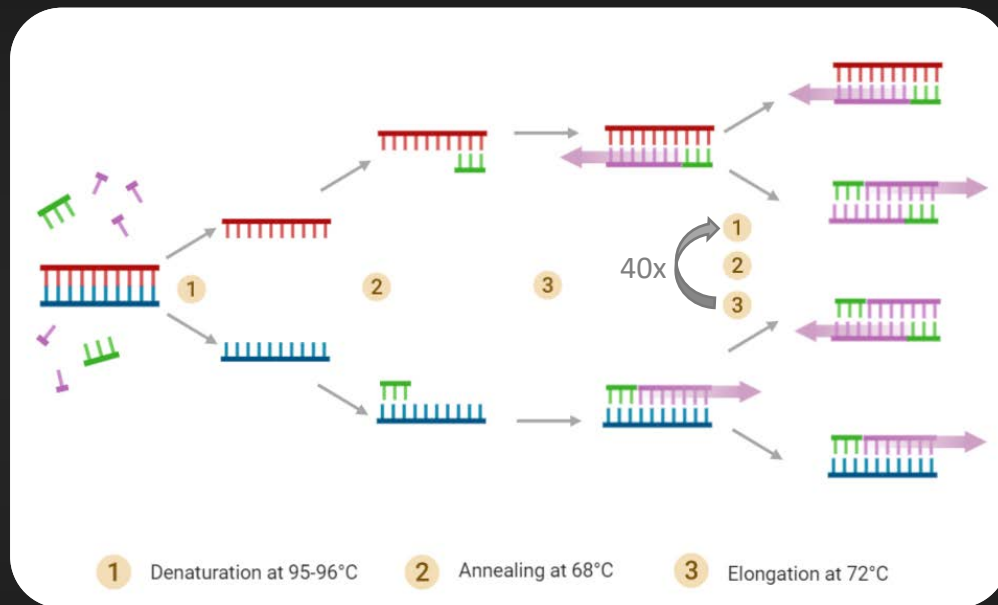
aCGH
Full Génome
Haute résolution
5 à 500 kb

2. Génétique Moléculaire

- **Génétique moléculaire** = Etude de l'ADN, recherche de variants dans des gènes, après extraction de l'ADN des cellules nucléées
- MLPA
- Séquençage Sanger
- NGS
 - Nanopore

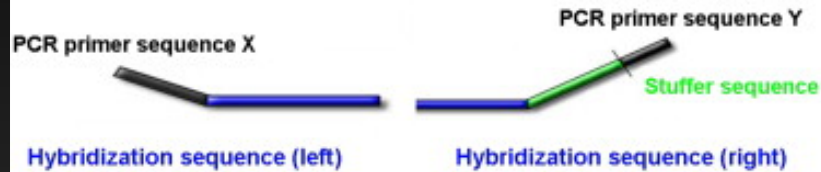


La Polymérase Chain Reaction (PCR)



La multiplex ligation-dependent probe amplification (MLPA)

1. Denaturation and Hybridization



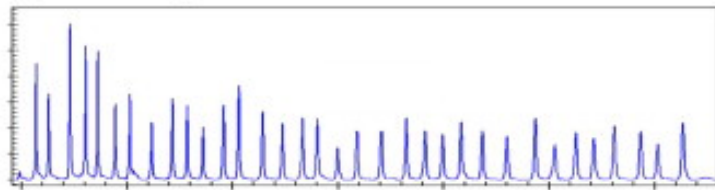
2. Ligation



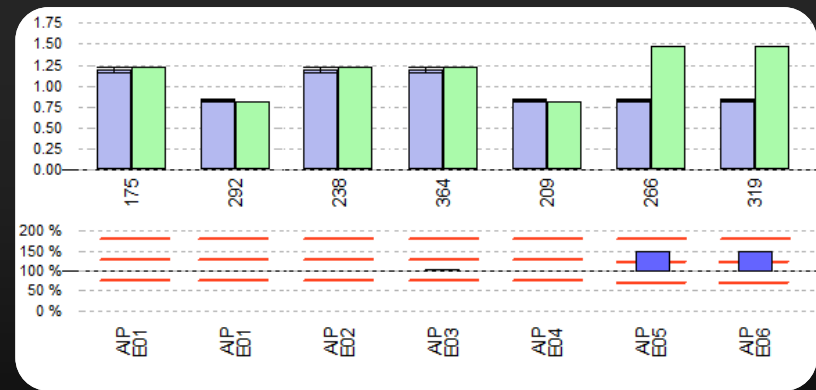
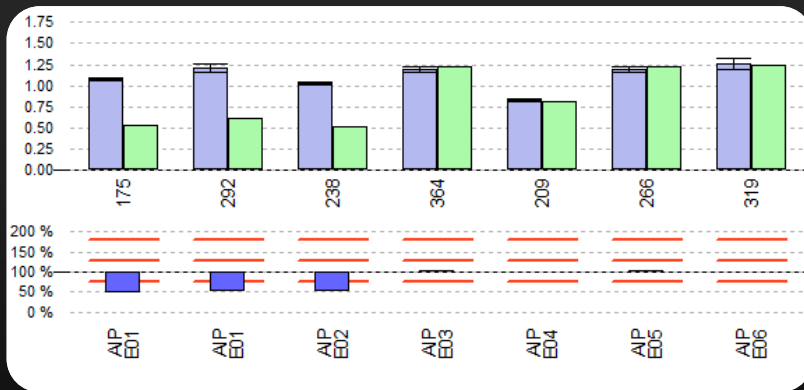
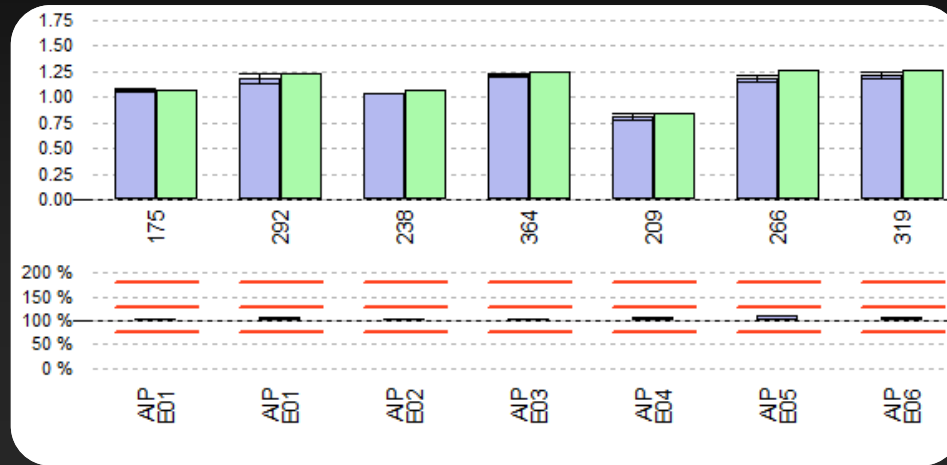
3. PCR with universal primers X and Y exponential amplification of ligated probes only



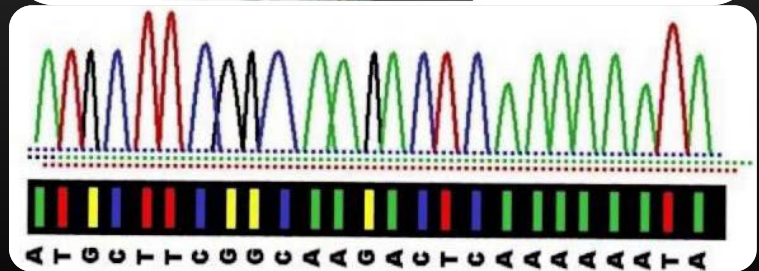
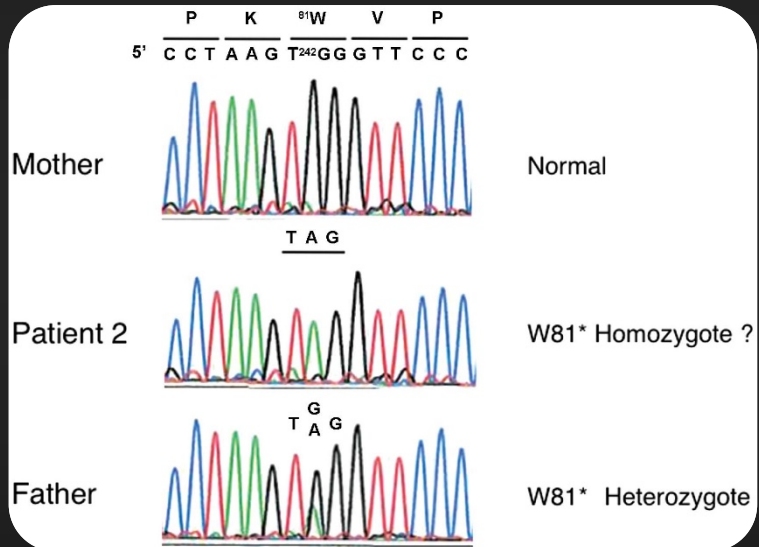
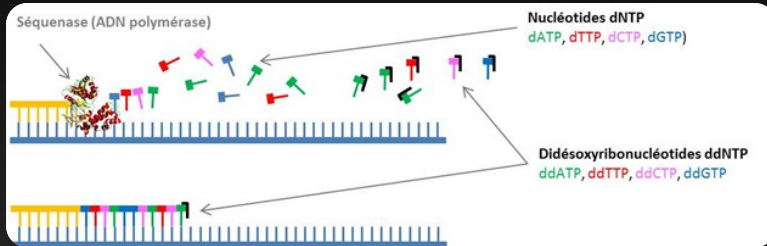
4. Fragment analysis



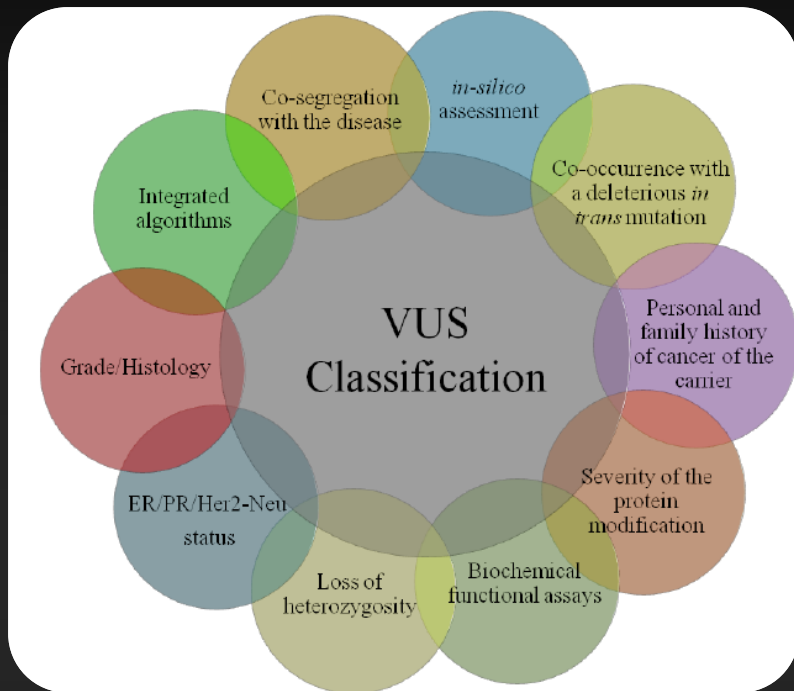
La multiplex ligation-dependent probe amplification (MLPA)



Séquençage Sanger

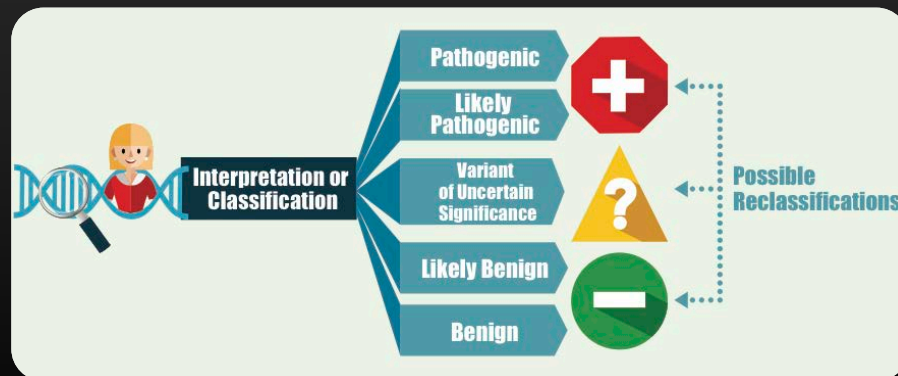


Interprétation des variants



Recommandations de l'American College of Medical Genetics (2015)

- Faisceau d'arguments
 - **En faveur de la pathogénicité**
 - PVS (Pathogenic Very Strong) argument très fort
 - PS (Pathogenic Strong) argument fort
 - PM (Pathogenic Moderate) argument moyen
 - PP (Pathogenic Poor or supporting) argument faible
 - **En faveur du caractère bénin**
 - BA (Benign stand Alone) argument suffisant
 - BS (Benign Strong) argument fort
 - BP (Benign Poor or supporting) argument faible

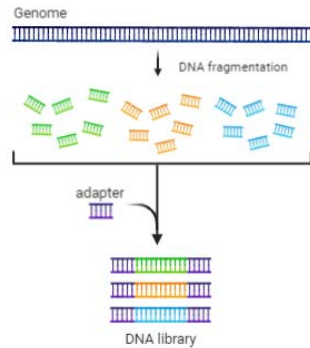


Seconde évaluation des variants

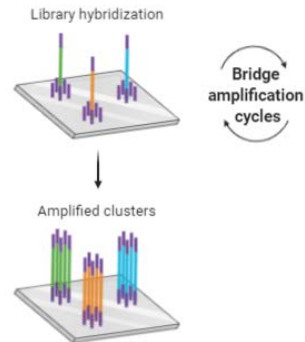
For variants with a prior score of:	when re-evaluated, variant score:				Odds (up)
	n	Moved Down	Stayed Same	Moved Up	
(likely benign)	737	← 287 (38.9%)	444 (60.2%)	→ 6 (0.8%)	.02
(VUS, but suggesting benign)	957	← 318 (33.2%)	619 (64.7%)	→ 20 (2.1%)	.06
(VUS)	776	← 153 (19.7%)	518 (66.8%)	→ 105 (13.5%)	.70
(VUS, but suggesting pathogenic)	427	← 22 (5.2%)	341 (79.8%)	→ 64 (15.0%)	2.88
(likely pathogenic)	220	← 4 (1.8%)	186 (84.5%)	→ 30 (13.6%)	7.56

Next Generation Sequencing (NGS)

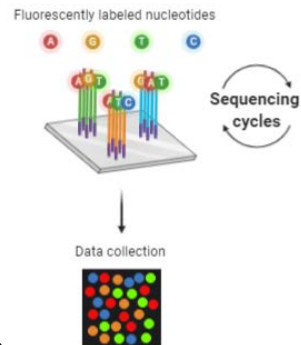
1 Library preparation



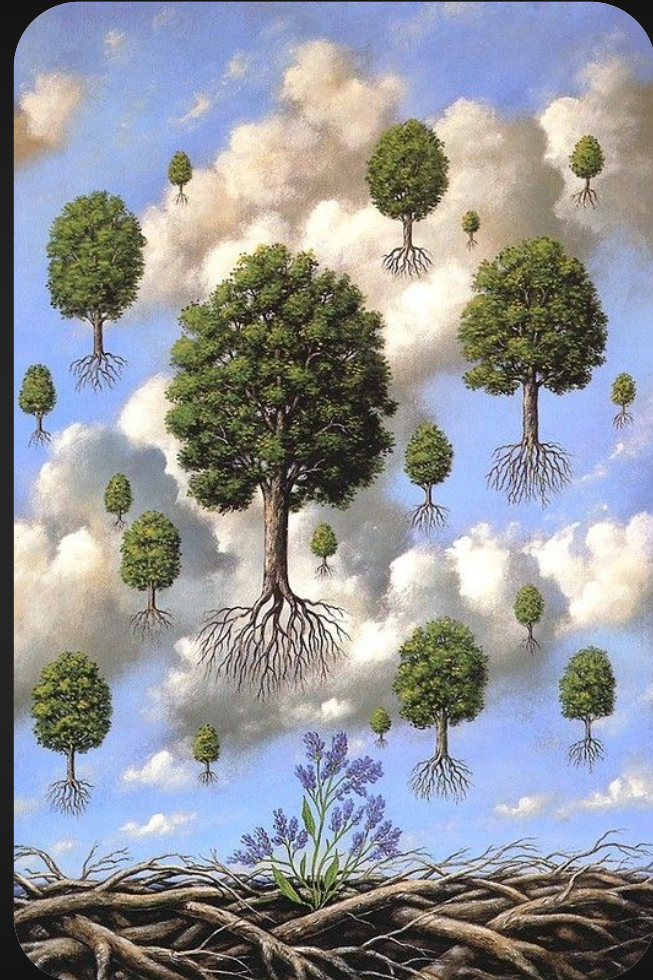
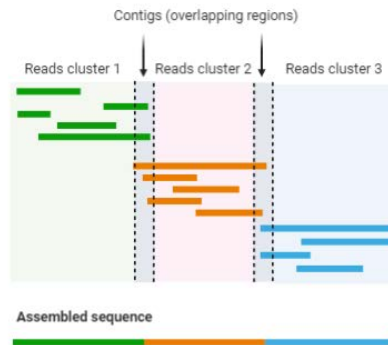
2 DNA library bridge amplification



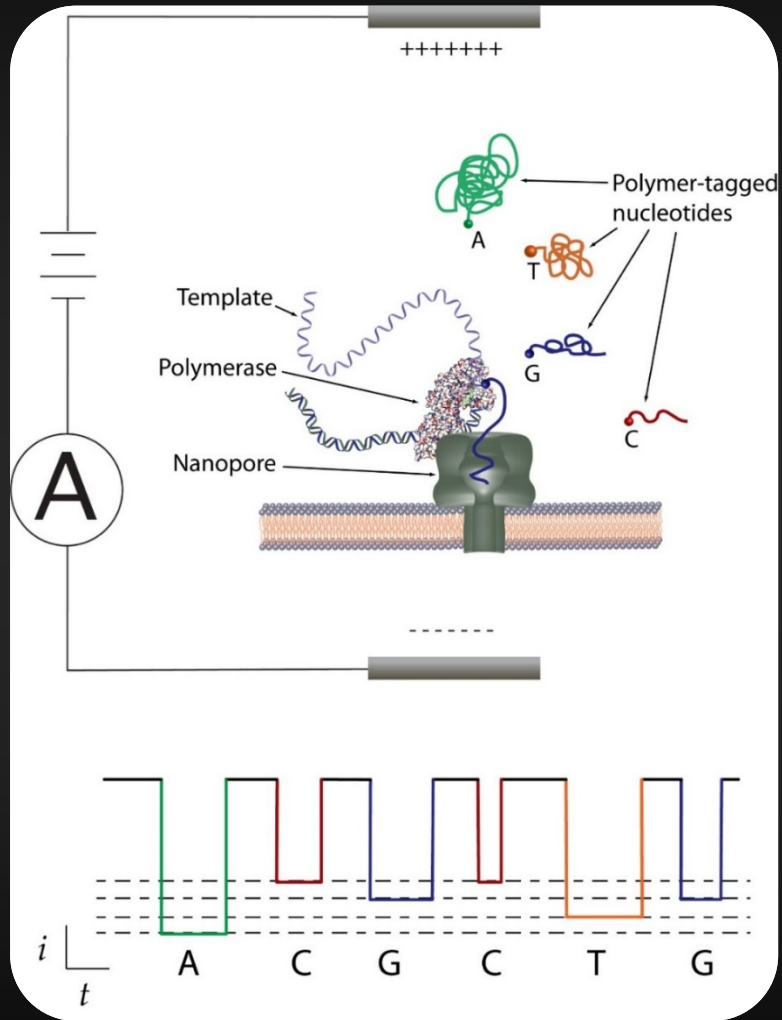
3 DNA library sequencing



4 Alignment and data analysis



Nanopore



En résumé...



Sanger



MLPA



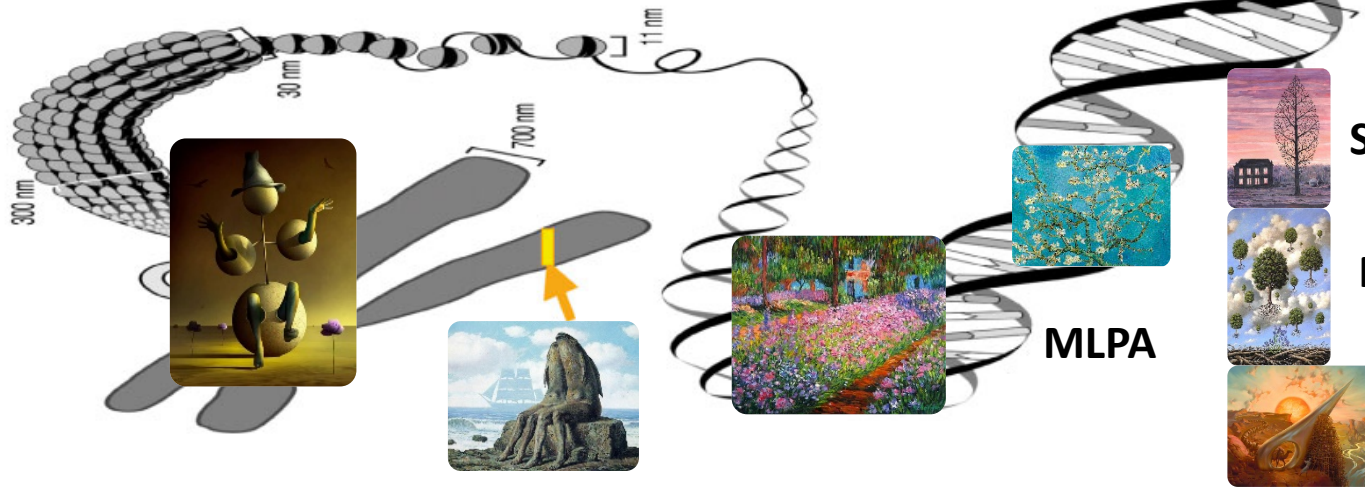
NGS



Nanopore

Chromosomes

ADN



Caryotype conventionnel

FISH

Caryotype moléculaire (puce à ADN)

Génétique moléculaire

Sanger

NGS

Nano

Seuil de détection

5 Mb

250 kb

5 kb

1 b

3×10^9 bases

Remerciements

Merci à tous pour votre attention



« Les passions sont les vents qui enflent les voiles du navire ;
elles le submergent quelquefois, mais sans elles il ne pourrait voguer. »

Voltaire