Review of NGS in clinic

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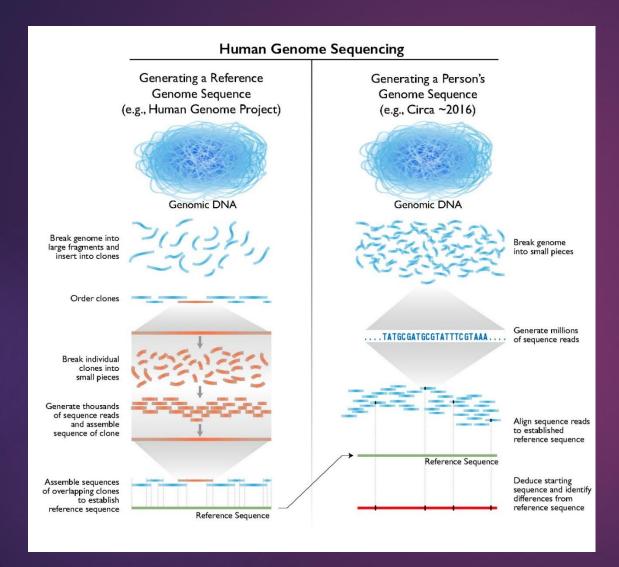
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Content of Lecture

- 1. Review NGS platforms
- 2. Review Library Prep
- 3. Review Genomic Testing

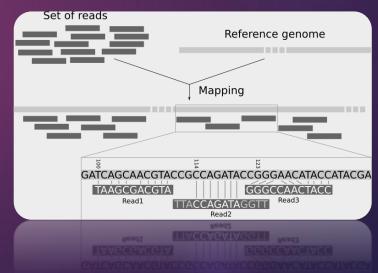
Giải trình tự gen thế hệ mới (NGS)



De novo assembly



Mapping to reference



Giải trình tự gen thế hệ mới (NGS): Có hệ gen tham chiếu

De novo assembly

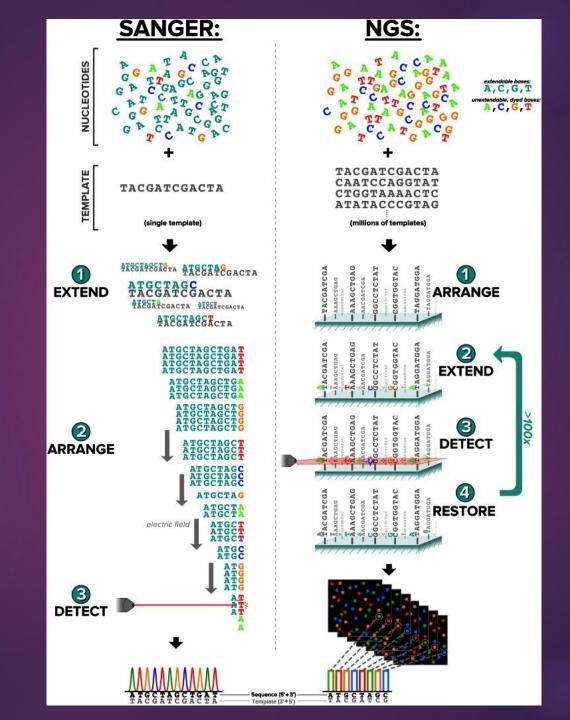


Mapping to reference



Giải trình tự gen thế hệ mới (NGS): giải trình tự song song

Read1:CTCGAATACG

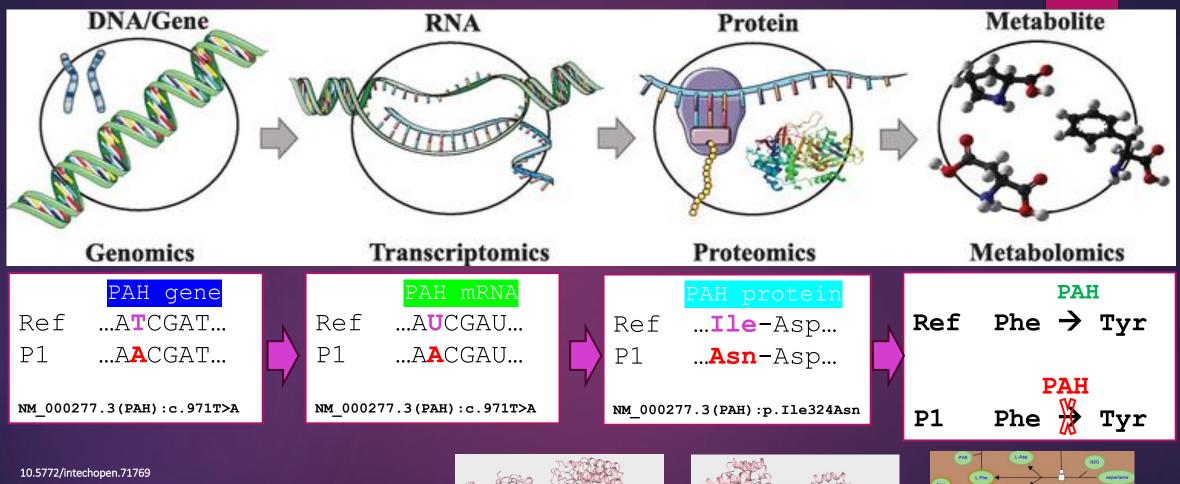


Read1:CTCGAATACG
Read2:CTCGAATACG
Read3:CTCGAATACG
Read4:CTCGAATACG
Read5:CGCGAATACG
Read5:CGCGAATACG
Read6:CGCGAATACG
Read7:CGCGACTACG
Read8:CGCGAATACG

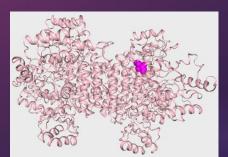
NGS vs Sanger

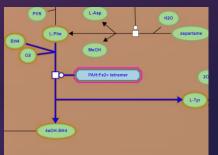


Quan hệ nhân quả: Biến thể gen và bệnh di truyền









Introduction to NGS platforms

- ► Sanger sequencing:
 - □ https://youtu.be/XtblMQ-Al0g?si=Q0CEaWzNvXDOIQWO



- Onso (Pacbio): https://youtu.be/tFmNO5rVoeM?si=ft4S8N9tbJ1We-GA
- ☐ GeneMind: https://youtu.be/2SLemogG8v4?si=G qIGS8WjtcAvwV7
- □ Illumina: https://youtu.be/-Kw5ov-orzU?si=1izVr8sZk6wPn4ii
- ☐ MGI: https://youtu.be/4 h4wNlyfk0?si=xrMMF9EEbj-5H19
- □ Ion Torrent (Thermal Fisher): https://youtu.be/qso3tdp0IPo?si=ACOAbgHL2dJl0cM8
- □ Aviti (Element Biosciences): https://youtu.be/pSwWofrRu3Y?si=A9Hid5fvLdcyN8rq
- Long read
 - Revio (Pacbio): https://youtu.be/mXcDYKnIDE8?si=QUvHRCWHlba7uF5N
 - □ ONT: https://youtu.be/MFVcj4VZHuU?si=hyRf8jU3WZRXcQF













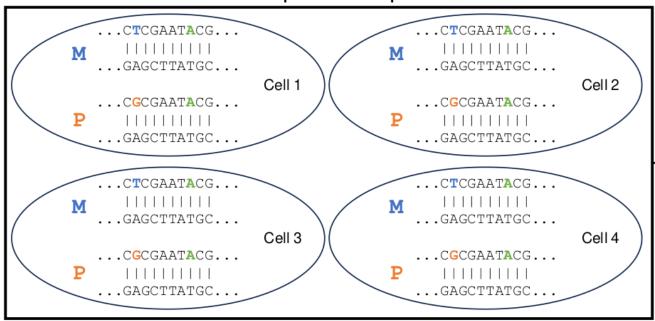




Quy trình XÉT NGHIỆM gen bằng phương pháp giải trình tự thế hệ mới (NGS)

Các bước trong XÉT NGHIỆM gen bằng phương pháp giải trình tự thế hệ mới (1)

A patient Sample



Fastq file

DNA extraction

2. Library Prep

3. Sequencing

Read1:CTCGAATACG

Read2:CTCGAATACG

Read3:CTCGAATACG

Read4:CTCGAATACG

Read5:CGCGAATACG

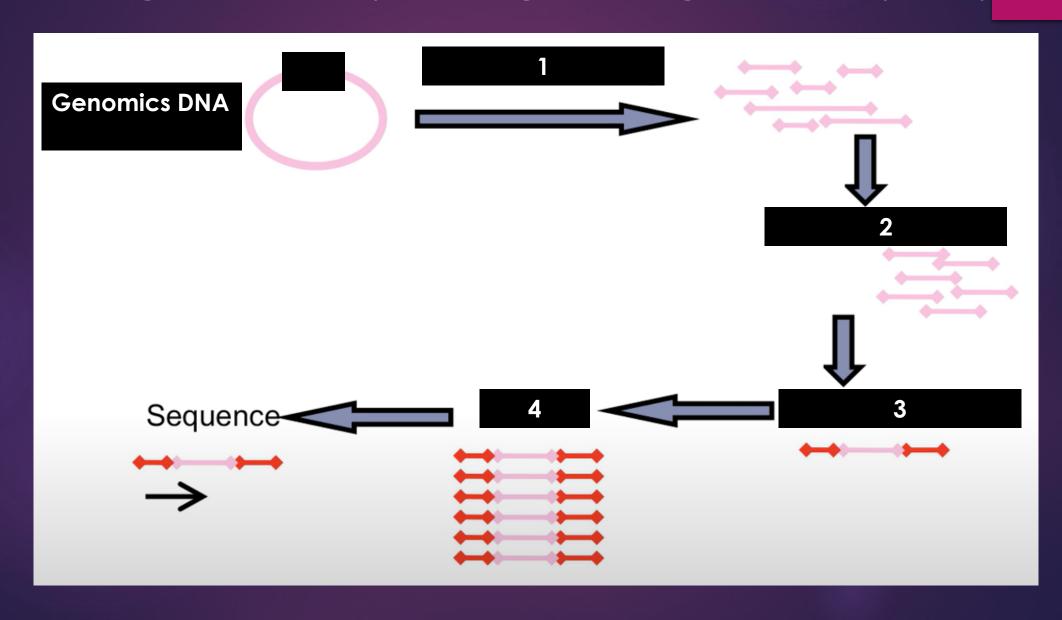
Read6:CGCGAATACG

Read7:CGCGACTACG

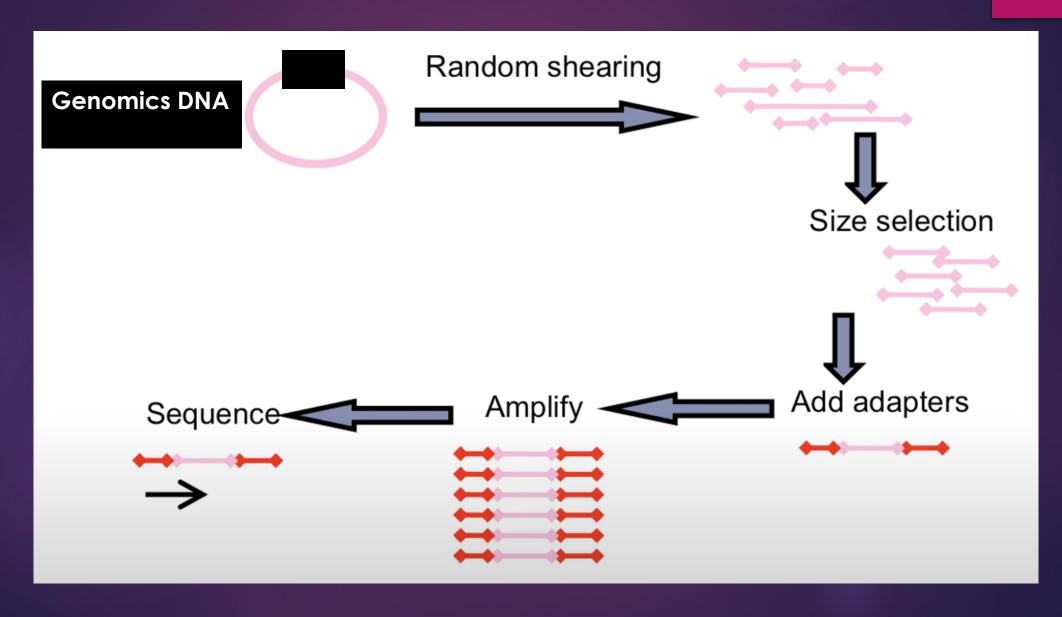
Read8: CGCGAATACG

- 4. Bioinformatics Analysis
 - 4.1 QCing Fastq
 - 4.2 Mapping to reference
 - 4.3 Calling variants
 - 4.4 Annotating variants

Whole genome sequencing – Shotgun Library Prep

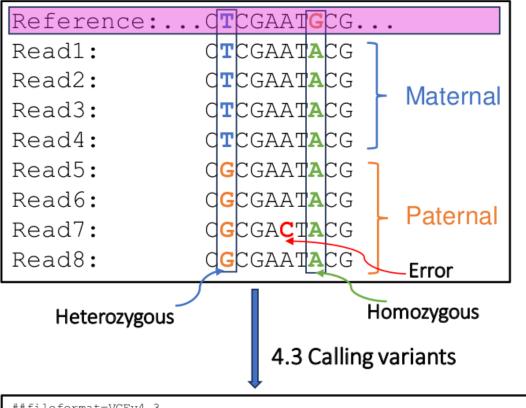


Whole genome sequencing – Shotgun Library Prep



Các bước trong XÉT NGHIỆM gen bằng phương pháp giải trình tự thế hệ mới (2)

4.2 Mapping reads to reference



 $\frac{\text{ANN=G}|\text{stop_gained}|\text{HIGH}|\text{OR4F5}|\text{ENSG00000186092}|\text{transcript}|\text{ENST0}|\\ 0000641515.2|\text{protein_coding}|3/3|\text{c.}822T>G|\text{p.Trp274*}|882/2618|822/981|274/326||\text{Pathogenic}|$

ANN=A | frameshift_variant | HIGH | ZSWIM2 | ENSG00000163012 | transcrip t | ENST00000295131.3 | protein_coding | 9/9 | c.1238G>A | p.lle413 | 1293/2451 | 1238/1902 | 413/633 | | ; LOF=(ZSWIM2 | ENSG00000163012 | 1 | 1.00)

4.4 Annotating variants

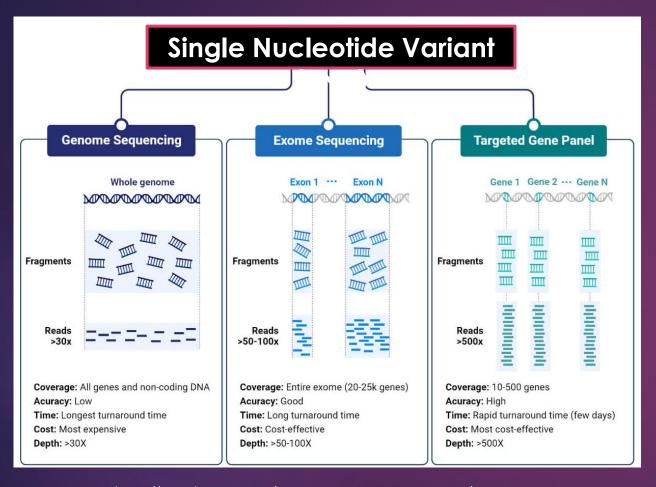
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##FORMAT=<ID=GT, Number=1, Type=String, Description="Genotype">
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##FORMAT=<ID=DP, Number=1, Type=Integer, Description="Read Depth">
##FORMAT=<ID=AD, Number=2, Type=Integer, Description="Read depth for each allele">
#CHROM POS
                                            REF
                                                                                                             Sample1
                                                     ALT
                                                                 OUAL
                                                                           FILTER
                                                                                        FORMAT
           14370
                          rs6054257
                                                                 129
                                                                                                              0/1:48:8:4,4
20
                                                      G
                                                                            PASS
                                                                                         GT:GO:DP:AD
20
           17330
                                                                 150
                                                                                                              1/1:49:8:8,8
                                                                             PASS
                                                                                         GT:GQ:DP:AD
```

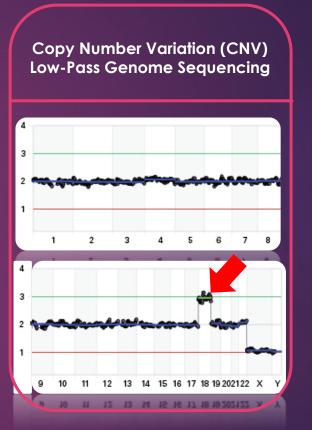
Hội đồng Hệ Gen: phiên giải và hội chẩn nh<mark>ững</mark> biến thể trong báo cáo kết quả NGS

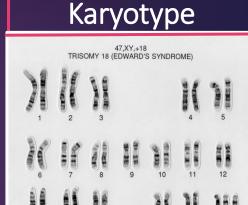


https://ggba.swiss/en/the-first-medical-genomics-center-opens-in-geneva/

Ứng dụng giải trình tự gen thế hệ mới trong lâm s<mark>àng</mark>







Xin chân thành cảm ơn!

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