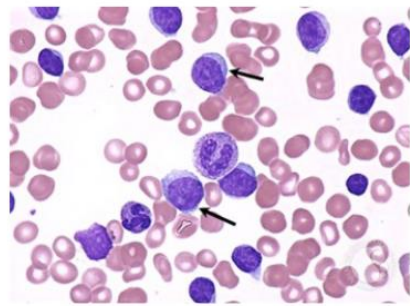
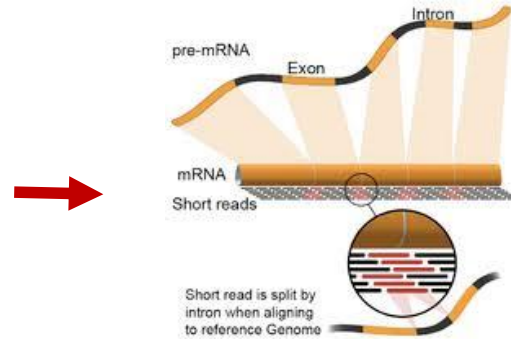


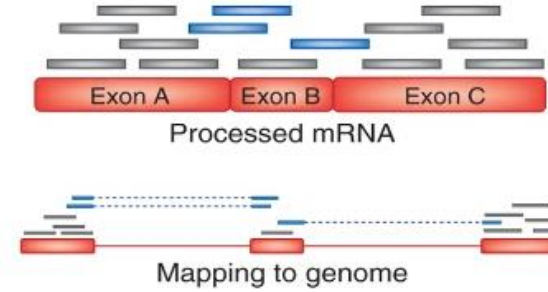
Project overview



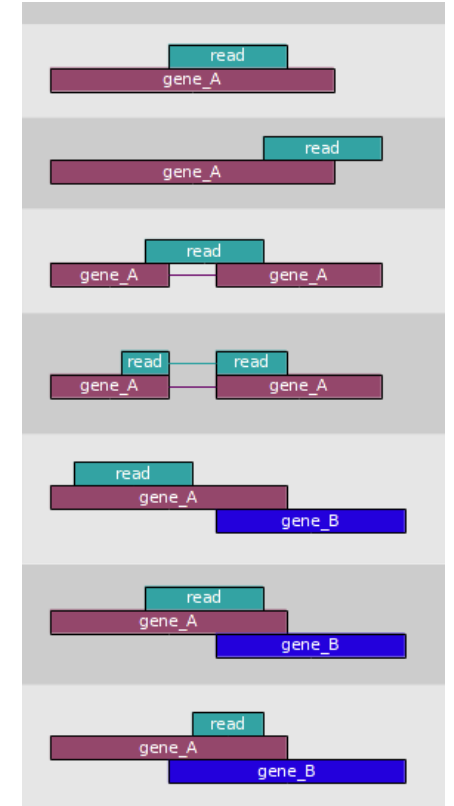
CLL blood sample



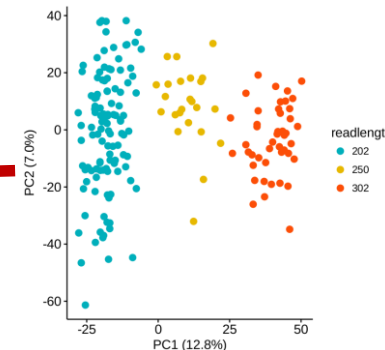
RNAseq



Mapping: STAR

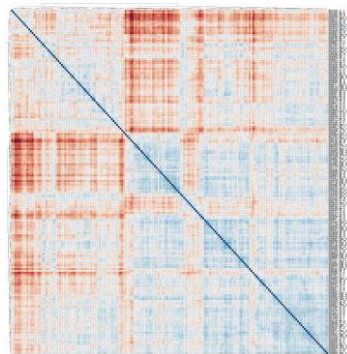


Counting: HtSeq



QC, EDA ..

Gene expression
Dataset (184 CLL)



Data Analysis (Differential
expression, Enrichment,
epistasis model..)

Biological
interpretation