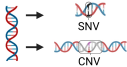


Input

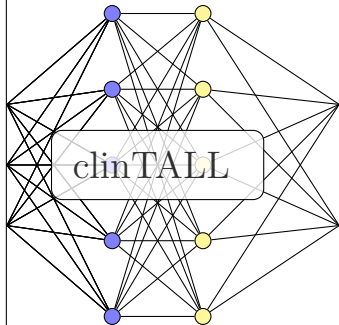
Clinical data



Genome

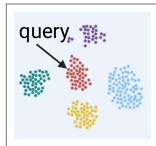


Transcriptome



Output

Subtype prediction



Risk stratification

