Input

Clinical data

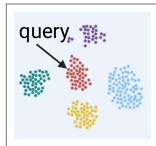Genome

Transcriptome

clinTALL
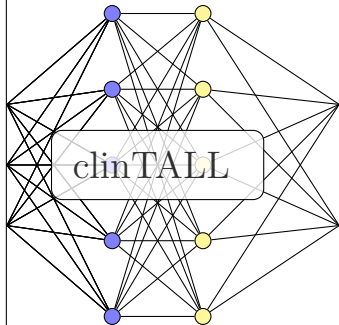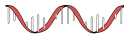
Output

Subtype prediction

query

Risk stratification

Low

Intermediate

query   High

Survival

Time