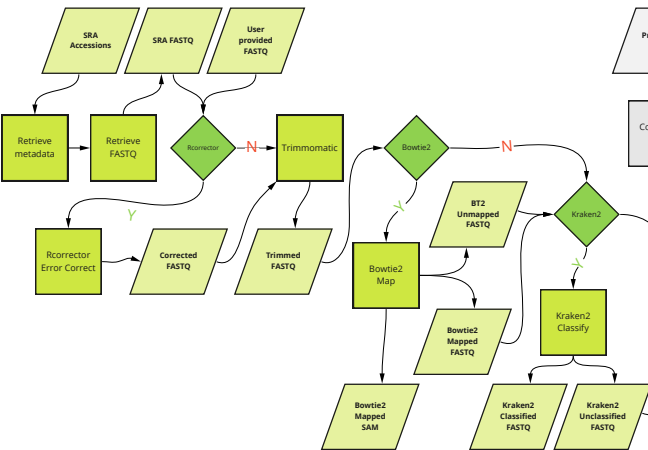
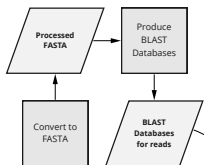


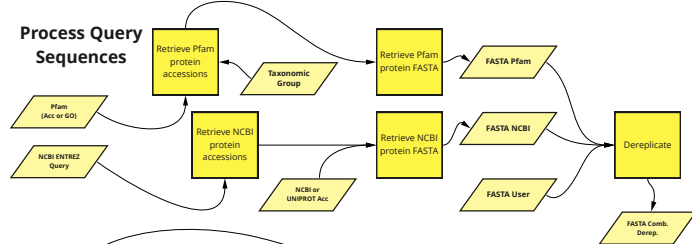
## Process RNA-seq Reads



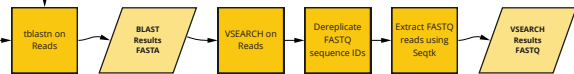
## Produce BLAST Databases for Reads



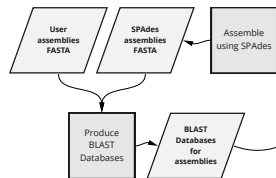
## Process Query Sequences



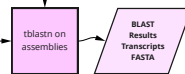
## Search RNA-seq Reads



## Assemble Reads



## Search Assemblies



## Annotate

