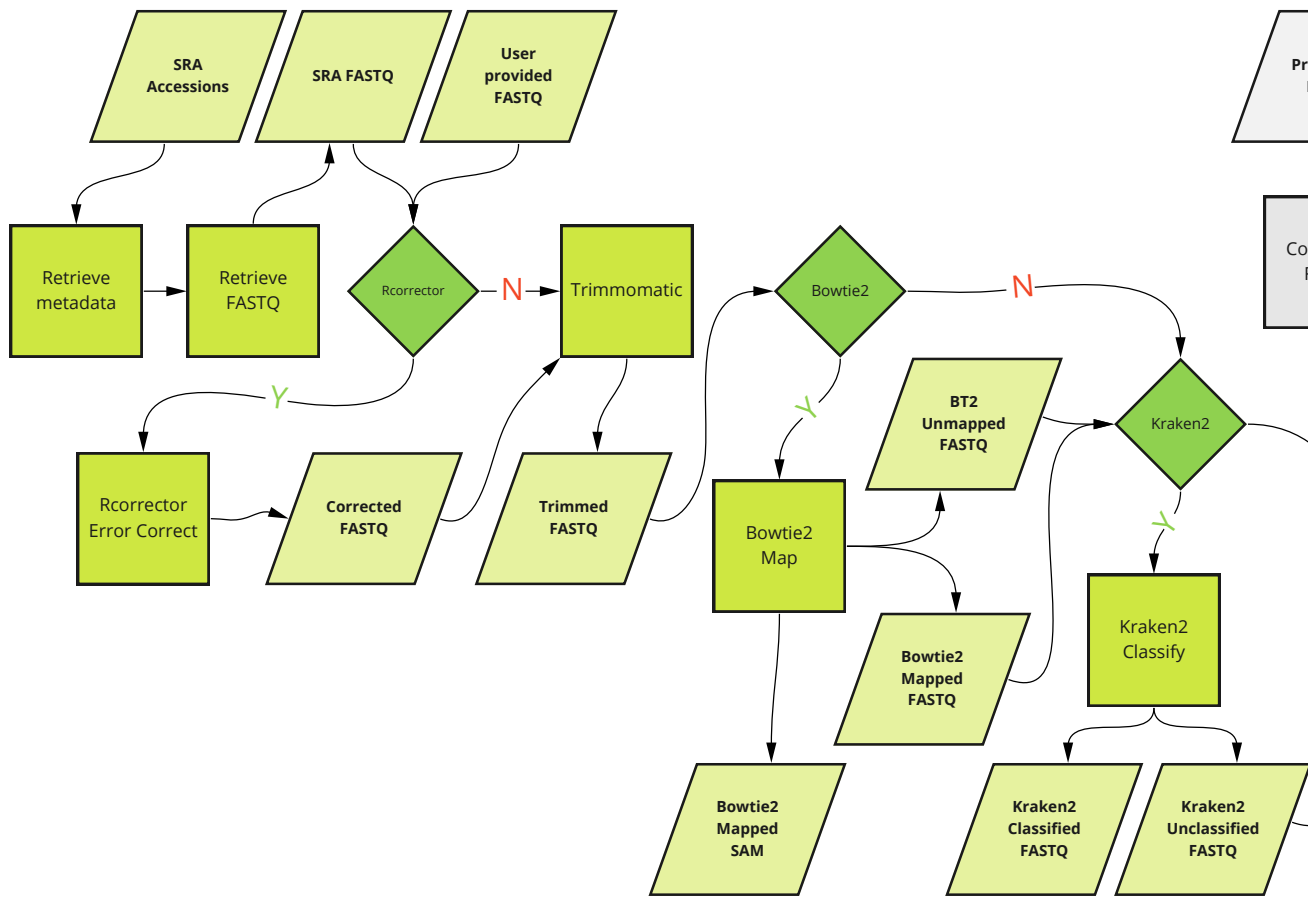
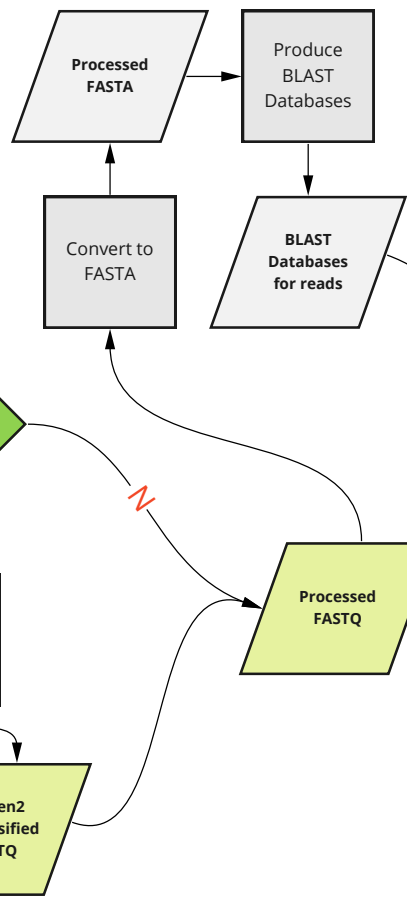


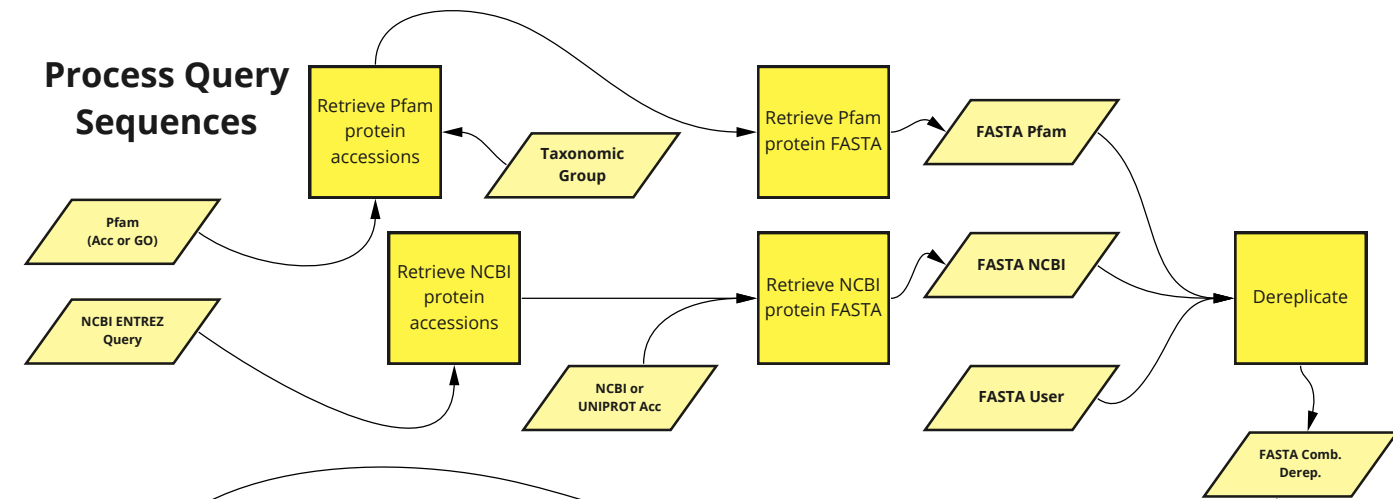
Process RNA-seq Reads



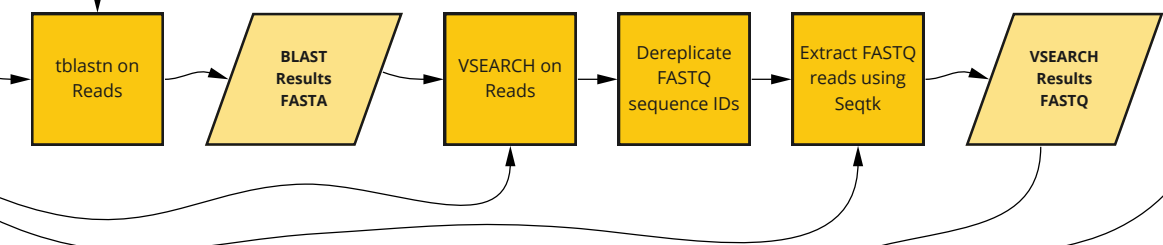
Produce BLAST Databases for Reads



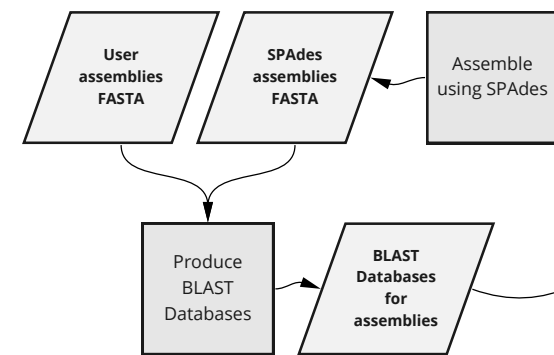
Process Query Sequences



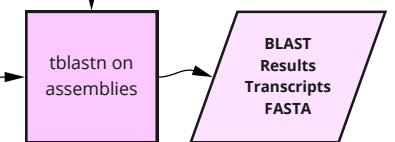
Search RNA-seq Reads



Assemble Reads



Search Assemblies



Annotate

