EvolvR Web-app & R package

Query protein *AccNum, Fasta*

Molecular characterization of protein(s)

Identify homologs

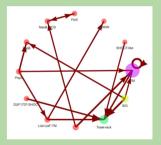
Molecular characterization of homologs

Domain Architectures Domain D

Phyletic spread



Proximity network



Structure-based sequence alignment Phylogenetic tree



