EvolvR Web-app & R package

Query protein *AccNum, Fasta*

Molecular characterization of protein(s)

Identify homologs

Molecular characterization of homologs
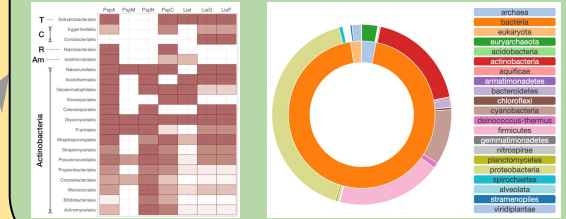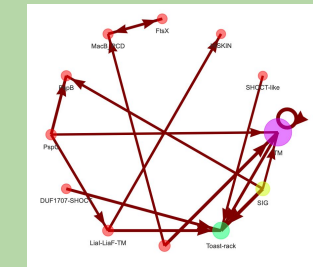
Phyletic spread

Proximity network

Structure-based sequence alignment Phylogenetic tree

Domain Architectures

Domain1 Domain2

Genomic Contexts

SIG TM D3 Domain1 Domain2 TM D4