

Input

sequence

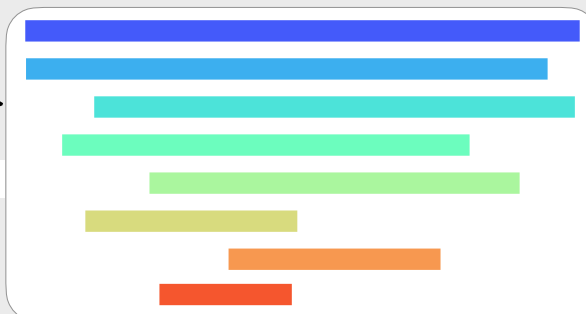


ColabFold

MMseqs2-based

UniRef

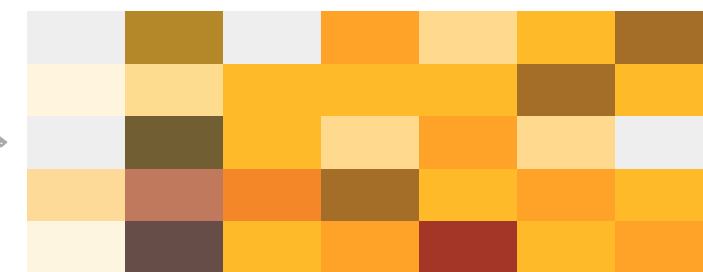
Env



Multiple Sequence
Alignment

Prediction

GEMME

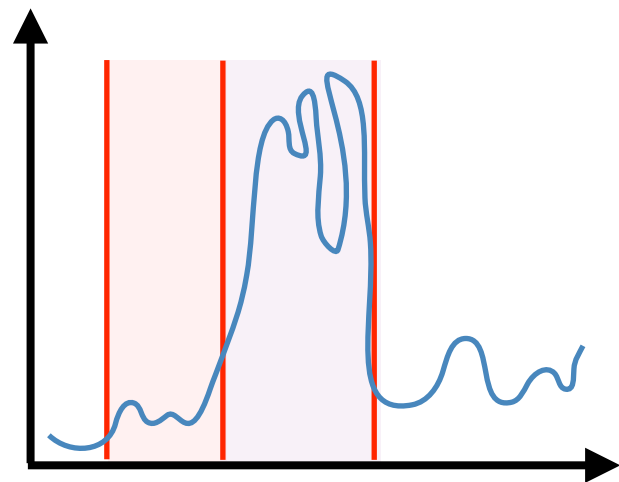


Mutational Landscape

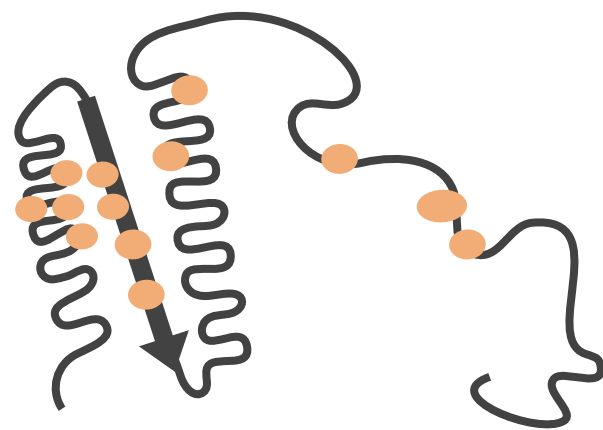
Post-processing



AlphaFoldDB
structure

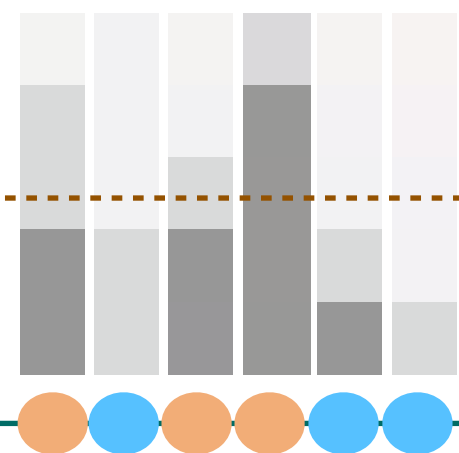


Sensitivity Profile



3D Structural
Mapping

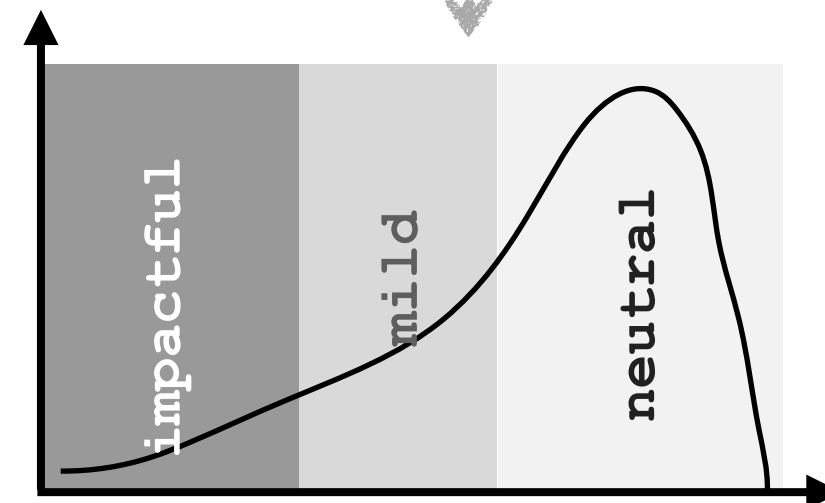
50%



sensitive tolerant
Residue Classification



Residue Confidence



Variant Classification