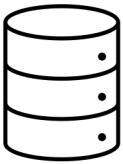


Custom protein
reference
(fasta file)

indexing

Database



Generate n-mers

Peptide	ID	Length
AVGHILYT	sp Q32FT	8
VGHILYTR	sp Q32FT	8

Alignments

Filtered peptides