

MS alignment of Uniref50 Sequences - FASTA

Calculate Web Service

Sort 120 130 ▶

Calculate Tree ▶

Pairwise Alignments...

Principal Component Analysis

Autocalculate Consensus

140 150

Average Distance Using % Identity

Neighbour Joining Using % Identity

Average Distance Using BLOSUM62

Neighbour Joining using BLOSUM62

KVTSGSVNDDGSELDLDDQIKEGWVLTCAAYPTSDVTIETHKEEELTA-
 KVTSGSVNDDGSELDLDDQIKEGWVLTCAAYPVSDVTIETHKEEELTA-
 KVVSGSVDQSDQSFLDDDDQIAEGFVLTCAAYPTSDVTIETHREEDMV--
 KVVSGSVDQSDQSFLDDDDQIAEGFVLTCAAYPTSDVTIETHKEEELV--