

File Tools Help **Window****Close All**

Raise Associated Windows

Minimize Associated Windows

MuscleWS alignment of Uniref50 Sequences - FASTA

Average distance tree using BLOSUM62 on region from MuscleWS alignment of Uniref50 Sequences - FASTA

FER_CAPAA/1-97

FER_CAPAN/1-144

FER1_LYCES/1-144

SCAGK IAGGAVDQTDGNFLDDQLEEGWVLT CVAY
SCAGKVTACSVDSDCNEI LDEDOEAAACEVLT CVAY

Q93XJ9_SOLTU
FER1_PEA