



# Multiple Sequence Alignment

File Tools Alignment View Help

Phylogenetic Tree *Hide Display*

**Residue Selection** ▶

Number of residues per molecule

Print QR ordering

RMSD Tools ▶

Q Value

Sequence Identity

1asza2  
1b8aa2  
1efwa3  
1sera2

--EKE-ASEELR-  
-K--NGWWEPE-R