

Web Service

Alignment

Secondary Structure Prediction

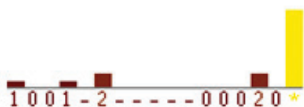
Protein Disorder

Analysis

Conservation

Fetch DB References

-----A  
GLKS - GTARGG RVTAMA  
-----A  
GLKS - STARGG RVTAMA  
GLKS - STARGG RVTAMA  
GRSA - SSRR - -RLRAQA  
KPLA - APMRRQLLRAQA



<http://www.compbio.dundee.ac.uk/jabaws>

Tcoffee with Defaults

Edit settings and run...

Run Tcoffee with preset

Probcons with Defaults

Edit settings and run...

Muscle with Defaults

Edit settings and run...

Run Muscle with preset

Mafft with Defaults

Edit settings and run...

Run Mafft with preset

MSAprobs with Defaults

Edit settings and run...

GLprobs with Defaults

Edit settings and run...

Clustal

Realign with Clustal

ClustalO

Realign with ClustalO