

ViennaRNA

Web Service

Alignment

Secondary Structure Prediction

Protein Disorder

Analysis

Conservation

Fetch DB References

JNet Secondary Structure Prediction

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

RNAAliFold Prediction

Change RNAAliFold settings...

Modify settings for the RNAAliFold prediction. Use this to hide or show different results of the RNA calculation, and change RNA folding parameters

- CCUUUGU--UAAGGGU----- UUG
-- CUAUGA-CCUAG-G----- GGU
-- GUUCA---CAC----- AGC
- UGUCUCU-UGGAC-GC----- CUGGU
- CCAGCCU--GCUG-GU----- GGG
- CCUUUGU--UAAG-GA----- CUUG
- AGGGUCA---UCC-A----- GAAA