

Protein Disorder Prediction - Individual Project

This server uses deep learning methods to examine the intrinsic disorder in protein sequences.

Protein sequences can be found from the following websites: Protein Data Bank and UniProt.

To enter another sequence: [return to home page](#)

Name of entered sequence: >sp|P03265|DNB2_ADE05

Positions in the sequence that have been identified as being disordered are coloured red, ordered residues are blue.



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