

# 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 green.



DOWNLOAD PREDICTION