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: >T0644

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

M K F L K F S L L T A V L L S V V F A F S S C G D D D D T G Y L P P S Q A I Q D A L K K L Y P N A T A I K W E Q K G V Y Y V A D C Q A D G R E K E V W F D A N A N W L M T E T E L N S I N N L P P A V L T A F M E S S Y N N W V V D D V V I L E Y P N E P S T E F V V T V E Q G K K V D L Y F S E G G G L L H E K D V T N G D D T H W P R V

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