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.

Please enter your protein sequence in FASTA format:

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>sp|P03265|DNB2_ADE05 DNA-binding protein OS=Human adenovirus C serotype 5 OX=28285
GN=DBP PE=1 SV=1
MASREEEQRETTPERGRGAARRPPTMEDVSSPSPSPPPPRAPPKKRMRRRIESEDEEDSS
QDALVPRTPSPRPSTSAADLAIAPKKKKKRPSPKPERPPSPEVIVDSEEEREDVALQMVG
FSNPPVLIKHGKGGKRTVRRLNEDDPVARGMRTQEEEEEPSEAESEITVMNPLSVPIVSA
WEKGMEAARALMDKYHVDNDLKANFKLLPDQVEALAAVCKTWLNEEHRGLQLTFTSKKTF
VTMMGRFLQAYLQSFAEVTYKHHEPTGCALWLHRCAEIEGELKCLHGSIMINKEHVIEMD
VTSENGQRALKEQSSKAKIVKNRWGRNVVQISNTDARCCVHDAACPANQFSGKSCGMFFS
EGAKAQVAFKQIKAFMQALYPNAQTGHGHLLMPLRCECNSKPGHAPFLGRQLPKLTPFAL
SNAEDLDADLISDKSVLASVHHPALIVFQCCNPVYRNSRAQGGGPNCDFKISAPDLLNAL
VMVRSLWSENFTELPRMVVPEFKWSTKHQYRNVSLPVAHSDARQNPFDF
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Submit Sequence