

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.

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

Download Prediction as PNG