## **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: >splP03265|DNB2\_ADE05

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

							_																																													D		
-																																																				M		
F	S	N	P	P	V	L	I	K	10	j K	(	i G	K	R	T	V	R I	R I	N	ΙE	D	D	P	V	A	R	G I	M	R	T (	Q E	3 E	E	E	E	P 3	S	Ξ /	E	S	E	Ι	T	V ]	<b>M</b> :	N I	P J	LS	s V	Ρ	Ι	V	S	A
W	Е	$\mathbf{K}$	G	M	E	Α.	Αl	R A	\ L	L N	1 L	K	Y	Н	$\mathbf{v}$	D	N I	) I	. K	A	N	F	K	L	L	<b>P</b> !	D	Q	V	E	ΑL	ے A	A	$\mathbf{v}$	<b>C</b> 1	K 7	ΓV	V I	L N	Ε	E	Н	R	G	L	Q I	L 7	Г	FT	S	K	K	T	F
V	T	M	M	G	R	F :	L (	Q A	Y	L	, ر	) S	F	A	Е	V	Τ ?	Y	H	Н	E	P	T	G	C	<b>A</b> :	L	W	L	Н	R C	CA	E	Ι	E	G I	ΞΙ	Ŀ	C	L	Н	G	S	I	M	I	N I	K	3 H	V	Ι	Ε.	M	D
																																																				F		
E	$\mathbf{G}$	A	$\mathbf{K}$	Α	Q	V.	A 1	F	ζ	) I	K	A	F	M	Q	A	L	Y	N	I A	Q	T	G	Н	G	H	L	L	M	PΙ	LR	C	E	C	N	S	K I	9 (	3 H	Α	P	$\mathbf{F}$	L	G	R	Q I	L J	PΙ	ζL	. T	P	F	A	L
S	$\mathbf{N}$	A	$\mathbf{E}$	D	L	D.	ΑI	) I	l	S	S	) K	S	V	L	A	S	V	H	I P	A	L	I	V	F	Q	C	C	N	P	VΥ	R	N	S	R.	Α (	Q (	3 (	G G	P	N	C	D	F.	K	I	S I	A I	PD	L	L	N	A	L
$\mathbf{V}$	M	V	$\mathbf{R}$	S	L	W	S I	ΕΝ	N F	7	E	EL	P	R	M	V	V I	P E	F	K	W	S	T	K	H	Q `	Y	R	N	V S	SI	. P	V	Α	Н	S I	) [	A F	Q S	N	P	$\mathbf{F}$	D	F										

Download Prediction as PNG