

# 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 blue.

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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