


Email	l.a.kelley@imperial.ac.uk
Description	P37681
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Protein structure and disorder analysis for the protein sequence MLI NRNI VAL FAL PF MAS AT A SEL SI GAGA AY NES PY RGY NENT KAI PLI SYEGDT FYVR. The figure shows four panels: Sequence, Secondary structure, SS confidence, Disorder, and Disorder confidence. The protein is 60 residues long. The secondary structure is shown as a blue ribbon diagram. The SS confidence is shown as a red bar. The Disorder is shown as a green bar. The Disorder confidence is shown as a green bar. The protein is predicted to be disordered.


Residue	Sequence	Secondary structure	SS confidence	Disorder	Disorder confidence
1	M				
2	L				
3	I				
4	N				
5	R				
6	N				
7	I				
8	V				
9	A				
10	L				
11	F				
12	A				
13	L				
14	P				
15	F				
16	M				
17	A				
18	S				
19	A				
20	T				
21	A				
22	S				
23	E				
24	L				
25	S				
26	I				
27	G				
28	A				
29	G				
30	A				
31	A				
32	Y				
33	N				
34	E				
35	S				
36	P				
37	Y				
38	R				
39	G				
40	N				
41	E				
42	N				
43	T				
44	K				
45	A				
46	I				
47	P				
48	L				
49	I				
50	S				
51	Y				
52	E				
53	G				
54	D				
55	T				
56	F				
57	Y				
58	V				
59	R				
60					

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand