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Protein structure and disorder analysis of the full-length protein (1-410 aa). The figure displays the full sequence in 10 blocks of 10 amino acids each. For each block, the secondary structure (SS) is shown as a green ribbon (alpha-helix) or blue arrow (beta-strand). The confidence of the SS prediction is shown as a red bar. The disorder of the protein is shown as a green bar, and the confidence of the disorder prediction is shown as a blue bar. The protein is highly disordered, with only a few short alpha-helices and beta-strands. The disorder confidence is high (blue) for most of the protein, indicating a high degree of disorder.

Block	Sequence (1-10 aa)	SS	SS confidence	Disorder	Disorder confidence
1	MKQWI AALLL MLI P	α-helix	High	Disordered	High
2	GVQAAP QKVTLMVDD	β-strand	High	Disordered	High
3	VPVAQVLQALAEQEKL	α-helix	High	Disordered	High
4	NLVVSPDVS	β-strand	High	Disordered	High
5	GTVSL	β-strand	High	Disordered	High
6	HLTDVPWKQALQTVVK	α-helix	High	Disordered	High
7	SAGLI TRQEGNLSVHSI	β-strand	High	Disordered	High
8	AWQNNNI ARQEA	α-helix	High	Disordered	High
9	EQARQAANLPLENR	β-strand	High	Disordered	High
10		β-strand	High	Disordered	High
11	SITLQYADAGELAKAGEKLL	α-helix	High	Disordered	High
12	SAKGSMTVDKRTNRL	β-strand	High	Disordered	High
13	LLRDNKTALSALEQVVAQMDLP	α-helix	High	Disordered	High
14	VGQ	β-strand	High	Disordered	High
15	VELSAHI VTI NEKSLRELGVK	β-strand	High	Disordered	High
16	WTLADAQHAGGVGQVTT	β-strand	High	Disordered	High
17	LGSDL SVATATTHVGFNI	β-strand	High	Disordered	High
18	GRIN	β-strand	High	Disordered	High
19	GRLLDLELSALEEQKQQLDII	β-strand	High	Disordered	High
20	ASPRLLASHLQPASIKQGEI	β-strand	High	Disordered	High
21	PYQVSSGESGATSVEFKEA	β-strand	High	Disordered	High
22	VLGMEVPTPTVLQKGRI	β-strand	High	Disordered	High
23	RLKLHISQNVPGQVLQ	β-strand	High	Disordered	High
24	QADGEVLAI DKQEIETQVEVKS	β-strand	High	Disordered	High
25	GETLAL	β-strand	High	Disordered	High
26	GGI FTRKNKS GQDSVPL	β-strand	High	Disordered	High
27	LGDI PWF	β-strand	High	Disordered	High
28	GQLFRHDKGEDERREL	β-strand	High	Disordered	High
29	VVFITPRLVSS	β-strand	High	Disordered	High
30	SE	β-strand	High	Disordered	High

Confidence Key
High(9)  Low (0)
? Disordered
 Alpha helix
 Beta strand