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Protein structure analysis of the full-length protein (1-410 aa). The figure displays the amino acid sequence, secondary structure, and confidence scores for the full-length protein. The sequence is color-coded by amino acid type. The secondary structure is shown as a green ribbon. The confidence scores are shown as a red bar. The disorder analysis is shown as a green bar with question marks indicating regions of low confidence.

Position	Sequence	Secondary structure	SS confidence	Disorder	Disorder confidence
1-60	MSLAKASLVTAASTLVKIGAGLLVGKLLAVSFGPAGLGLAANFRQLITVLGVLGAGIFN	Alpha-helix	High	Low	High
61-120	GVTKYVAQYHDNPQQLRRVVGTSSAMVLGFSTLMALVFLAAAPISQGLFGNTDYQGLVR	Alpha-helix	High	Low	High
121-180	LVALVQMGIAWGNLLALMKGFRDAAGNALSLIVGSLIGVLAYYVSYRLGGYEGALLGLA	Alpha-helix	High	Low	High
181-240	LIPALVVI PAAMLIKRGVILSYLKPSWDNGLAGQLSKFTLMALITSVTLVPVAYIMMRK	Alpha-helix	High	Low	High
241-300	LLAAQYSWDEVGIWQGVSSI SDAYLQFITASFVYLLPTLSRLTEKRDITREVVKSLKVF	Alpha-helix	High	Low	High
301-360	LPAAVAAASFTVWLLRDFAIWLLLSNKFTAMRDLFAWQLVGDVLKVGAYVFGYLVIAKASL	Alpha-helix	High	Low	High
361-410	RFYILAEV SQFTLLMVFAHWLIPAHGALGAAQAYMATYIVVFSLCCGVFLLWRRRA	Alpha-helix	High	Low	High

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