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Protein structure analysis of the full-length protein (1-420 aa). The figure displays the amino acid sequence, secondary structure, and disorder analysis across the entire protein. The sequence is color-coded by physicochemical properties. The secondary structure is shown as green cylinders. The disorder analysis includes confidence scores for disorder and disorder confidence.

Position	Sequence	Secondary structure	Disorder	Disorder confidence
1-60	MSGDI LQT PDAPK PQQGALDN YFKI TARGST VRQEVLAGL TTF LAMVY SVI VVPGMLGKAG	Alpha-helix	Disordered	High
61-120	FPPAAVFVATCLVAGFGSLLMGLWANLPMAIGCAISLTAF TAFSLVLGQQI SVPVALGAV	Alpha-helix	Disordered	High
121-180	FLMGVIFTAISVTGVRTWILRNLPMGIAHGTGIGIGLFLLLIAANGVGMVINKNPIEGLPV	Alpha-helix	Disordered	High
181-240	ALGAF TSF PVMMSSLGLAVIFGLEKCRVPGGILLVIAISII GLIFDPAVKYHGLVAMPS	Alpha-helix	Disordered	High
241-300	LTGEDGKSLIFSLDIMGALQPTVLPSVLALVMTAVFDATGTIRAVAGQANLLDKDNQIIN	Alpha-helix	Disordered	High
301-360	GKGAL TS DSVSSI FSGLVGAAPAAVYIESAAGTAAGGKTGLTATVVGALFLLILFLSPLS	Alpha-helix	Disordered	High
361-420	FLIPGYATAPALMYVGLLMLSNVSKLDFNDFIDAMAGLVCAVFI VLTCNI VTGIMLG FVT	Alpha-helix	Disordered	High

