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Description	P0AEX3
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Protein structure analysis of the full-length protein (1-420) showing sequence, secondary structure, SS confidence, disorder, and disorder confidence across 10 segments.

Segment	Sequence	Secondary structure	SS confidence	Disorder	Disorder confidence
1	MAESTVTADS KLTSSDTRRR I WAI VGASSGNLVEWFDYVYSFCSLYFAHIFFP SGNTTT	Alpha helix	High	Disordered	High
2	QLLQTAGVF AAGFLMRPI GGWLFGR I ADKHGRKKSM LLSV CMMCFGSLVI ACLPGYETI G	Alpha helix	High	Disordered	High
3	TWAPALLLL LARLFQGLSVGG EYGT SATYMS EVAVEGRKGFYASFQYVTLI GGQLLALLVV	Alpha helix	High	Disordered	High
4	VVLQHTMEDAALRE WGWRI P FALGAVLAVV ALWLRRLQ LDETSQQETRALKEAGSLKGLWR	Alpha helix	High	Disordered	High
5	NRRAFI MVLGFTAAGSLCFYTF TTYMQKYL VNTAGMHANVASGI MTAALFVFM LIQPLIG	Alpha helix	High	Disordered	High
6	ALSDKI GRRTSMLCFGSLAAI FTVPI LSA LQNVSSPYAAFGLVMCALLI VSFYTSI SGIL	Alpha helix	High	Disordered	High
7	KAEMFPAQVRALGVGLSYAVANAI FGGSAEYVALSL KSI GMETAFFWYV TLMMAVVAFLVS	Alpha helix	High	Disordered	High

