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Protein structure and analysis of the N-terminal region of the protein. The image shows three segments of the protein sequence, each with its corresponding secondary structure, sequence confidence, disorder, and disorder confidence. The first segment (residues 1-60) shows a series of alpha-helices and a beta-strand. The second segment (residues 70-120) shows a series of alpha-helices and a beta-strand. The third segment (residues 130-140) shows a series of alpha-helices. The analysis includes sequence, secondary structure, sequence confidence, disorder, and disorder confidence for each segment.

Residue Range	Sequence	Secondary Structure	Sequence Confidence	Disorder	Disorder Confidence
1 - 60	MYCTVKELIRDLDTDPDSECVFAVVLTRGDVRIHQDWNLSDDLEETVMQRLDDAF EY	Alpha-helices and beta-strand	High	Disordered	High
70 - 120	GADVSI VHDVVRELMEEKRASRQVTVPVAVMLEKVMALAGSEMKRLYAVGSENGGDGDAFV	Alpha-helices and beta-strand	High	Disordered	High
130 - 140	REEREAMDVVLLQALDGEHMS	Alpha-helices	High	Disordered	High

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand