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Protein structure and disorder analysis of the N-terminal region of the protein. The figure shows three segments of the protein sequence (residues 1-60, 70-120, and 130-180) with corresponding secondary structure predictions (alpha-helices and beta-strands) and disorder analysis results. The disorder analysis includes confidence scores for disorder and disorder confidence, as well as disorder confidence scores. The protein is shown to be highly disordered, with many regions of low confidence for disorder prediction.

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand