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Protein structure analysis of the N-terminal region of the protein. The figure is divided into three horizontal panels, each showing a different segment of the protein sequence (residues 1-60, 70-120, and 130-140). Each panel displays the amino acid sequence, secondary structure (alpha-helices and beta-strands), solvent accessibility (SS), confidence scores, and disorder predictions. The first panel (residues 1-60) shows a long alpha-helix. The second panel (residues 70-120) shows a mix of alpha-helices and beta-strands. The third panel (residues 130-140) shows a short alpha-helix. The confidence scores and disorder predictions are shown as colored bars below the sequence.

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand