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Protein structure and disorder analysis of the human protein sequence: MKSKVLALLIPALLAAGAAHAAEVYNKDG NKL DLYGKVDGLHYFSDNSAKDGDQSYARLG.

The figure displays the protein structure and disorder analysis for the human protein sequence. The sequence is shown in the top row, with residues numbered 1 to 60. The secondary structure is shown in the second row, with blue arrows indicating alpha-helices and green cylinders indicating beta-strands. The solvent accessibility (SS) is shown in the third row, with a color scale from red (high) to blue (low). The confidence scores are shown in the fourth row, with a color scale from red (high) to blue (low). The disorder analysis is shown in the fifth row, with question marks indicating disordered regions.

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Confidence Key
High(9)  Low (0)
? Disordered
 Alpha helix
 Beta strand