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Description	P75966
Date	Thu Jan 5 12:16:33 GMT 2012
Unique Job ID	27e34be6dea04f27

Figure 1 displays the schematic representation of the protein structure of the human protein, showing the sequence, secondary structure (SS), and disorder confidence scores across four segments of the protein (1-60, 70-120, 130-180, and 190-210).


The protein sequence is shown in the top row of each segment, with residues numbered. The secondary structure (SS) is represented by blue arrows (alpha-helices) and green cylinders (beta-strands). The disorder confidence scores are shown as a bar chart below the sequence, with colors indicating different levels of confidence (red for high, yellow for medium, green for low, and blue for very low).


The protein is highly disordered, with many regions of high confidence in disorder. The disorder confidence scores are shown as a bar chart below the sequence, with colors indicating different levels of confidence (red for high, yellow for medium, green for low, and blue for very low).

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand