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Figure 1 displays the protein structure and disorder analysis of the human protein. The protein is shown as a blue ribbon structure with green helices and red loops. The disorder analysis shows regions of high confidence (red) and low confidence (yellow/green).


The protein structure is shown in three panels, each representing a different segment of the protein. The panels are labeled 1, 2, and 3, corresponding to the sequence ranges 1-60, 70-120, and 130-180, respectively.

Each panel displays the following information:


- Sequence:** The amino acid sequence for the segment.
- Secondary structure:** The predicted secondary structure elements (SS) for the segment.
- confidence:** The confidence score for the secondary structure prediction.
- Disorder:** The predicted disorder status for the segment.
- Disorder confidence:** The confidence score for the disorder prediction.


The protein structure is shown as a blue ribbon structure with green helices and red loops. The disorder analysis shows regions of high confidence (red) and low confidence (yellow/green).

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand