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Figure 1 displays the protein structure and disorder analysis for the protein. The figure is organized into three panels, each showing a different segment of the protein sequence (residues 1-60, 70-120, and 130-180).

Each panel includes the following information:


- Sequence:** The amino acid sequence for the segment.
- Secondary structure:** The predicted secondary structure elements (alpha-helices and beta-strands) shown as green cylinders and blue arrows, respectively.
- SS confidence:** The confidence score for the secondary structure prediction, represented by a color scale from red (high confidence) to blue (low confidence).
- Disorder:** The predicted disorder regions, indicated by red and blue bars.
- Disorder confidence:** The confidence score for the disorder prediction, represented by a color scale from red (high confidence) to blue (low confidence).


The protein structure is shown as a blue ribbon model, with green cylinders representing alpha-helices and red cylinders representing beta-strands. The disorder analysis shows regions of high confidence (red) and low confidence (blue) for the protein structure.

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand