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Figure 1 displays the protein structure and disorder analysis for the protein. The figure is organized into three main sections, each showing a different range of residues (1-60, 70-120, and 130-180). Each section includes the following information:


- Sequence:** The amino acid sequence for the specified range.
- Secondary structure:** A representation of the protein's secondary structure, showing alpha-helices (green cylinders) and beta-strands (blue arrows).
- SS confidence:** A bar chart indicating the confidence in the secondary structure prediction, with colors ranging from blue (low confidence) to red (high confidence).
- Disorder:** A bar chart indicating the predicted disorder of the protein, with colors ranging from blue (low disorder) to red (high disorder).
- Disorder confidence:** A bar chart indicating the confidence in the disorder prediction, with colors ranging from blue (low confidence) to red (high confidence).


The protein structure analysis shows that the protein is primarily composed of alpha-helices and beta-strands, with a high degree of disorder. The disorder confidence is generally high, particularly in the regions shown in red.

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand