



Email	l.a.kelley@imperial.ac.uk
Description	P23898
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
Figure 1 displays the protein structure and disorder analysis for the 160-residue protein. The protein is shown in three segments: residues 1-60, 70-120, and 130-160. The amino acid sequence is shown at the top, followed by the secondary structure prediction (SS) and disorder prediction (Disorder) for each segment. The SS is represented by green helices and blue arrows, while the Disorder is represented by red and blue bars. The confidence score for each prediction is shown at the bottom of each segment.

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand