



Email	l.a.kelley@imperial.ac.uk
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Protein structure and disorder analysis of the full-length protein (1-180 residues). The image shows three segments of the protein: residues 1-60, 70-120, and 130-180. Each segment displays the amino acid sequence, secondary structure (alpha-helices and beta-strands), solvent accessibility (SS), confidence scores, and disorder predictions. The protein is composed of several alpha-helices and beta-strands, with some regions predicted to be disordered.

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand