

Email	l.a.kelley@imperial.ac.uk
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Figure 1 displays the schematic representation of the protein structure of the human protein. The protein is composed of several alpha-helices and beta-strands. The figure shows three segments of the protein sequence (1-60, 70-120, and 130-170) with corresponding secondary structure predictions (SS), confidence scores, and disorder predictions. The protein is composed of several alpha-helices and beta-strands. The first segment (1-60) shows a long alpha-helix (residues 10-25) and a long beta-strand (residues 30-45). The second segment (70-120) shows a long alpha-helix (residues 75-95) and a long beta-strand (residues 100-115). The third segment (130-170) shows a long alpha-helix (residues 135-155) and a long beta-strand (residues 160-170).

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand