



Email	l.a.kelley@imperial.ac.uk
Description	P75967
Date	Thu Jan 5 12:16:34 GMT 2012
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
Figure 1: Protein structure and disorder analysis of the human protein. The figure displays four segments of the protein sequence (1-60, 70-120, 130-180, and 190-220) with corresponding secondary structure predictions (SS), confidence scores, and disorder analysis. The protein is shown as a ribbon diagram with alpha-helices in green and beta-strands in blue. The disorder analysis includes a disorder confidence scale from 0 to 100, with regions of high confidence (red) and low confidence (blue/green).

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand