



Email	l.a.kelley@imperial.ac.uk
Description	P77607
Date	Thu Jan 5 12:30:58 GMT 2012
Unique Job ID	1d3a40cb7c11de8a


Figure 1 displays the protein structure and disorder analysis of the human protein. The protein sequence is shown in three segments: 1-60, 70-120, and 130-180. The secondary structure is represented by green cylinders (alpha-helices) and blue arrows (beta-strands). The confidence is shown as a red bar. The disorder analysis is shown as a green bar with question marks indicating regions of uncertainty. The protein is predicted to be disordered in several regions, including the N-terminal region (1-60), the C-terminal region (130-180), and a central region (70-120).

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand