



Email	l.a.kelley@imperial.ac.uk
Description	P76555
Date	Thu Jan 5 12:24:33 GMT 2012
Unique Job ID	21e79aa7a1ba43c0


Figure 1: Protein structure and disorder analysis of the human protein. The figure displays three segments of the protein sequence (1-60, 70-120, and 130-180) with corresponding secondary structure predictions (SS), confidence scores, and disorder analysis. The protein is shown as a blue ribbon structure with green helices and red loops. The disorder analysis includes a disorder score (0-1) and a disorder confidence score (0-1). The protein is predicted to be disordered in several regions, including the N-terminal region (1-60) and the C-terminal region (130-180).

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand