



Email	l.a.kelley@imperial.ac.uk
Description	P33227
Date	Thu Jan 5 11:51:26 GMT 2012
Unique Job ID	80bc17b1bbe0f284


Protein structure and disorder analysis of the full-length protein (1-120). The image shows three segments of the protein: residues 1-60, 70-120, and 130-160. Each segment displays the amino acid sequence, secondary structure (alpha-helices and beta-strands), sequence confidence, and disorder prediction. The full-length protein is predicted to be disordered, with no significant structural elements identified.

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand