



Email	l.a.kelley@imperial.ac.uk
Description	P77326
Date	Thu Jan 5 12:27:45 GMT 2012
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
Protein structure and disorder analysis for the full-length protein (1-60). The top panel shows the full sequence (MI LDRHI T S R G Y V C M R P A T I A P L T P Y D K W D G E K V W V T D T E A Q H S V A V D A A E A Q R Q S L I D T A) with a secondary structure prediction (SS) showing alpha-helices (green) and a beta-strand (blue). The bottom panel shows the disorder analysis (Disorder) with a confidence scale (red to green) and a disorder prediction (Disorder confidence) showing regions of high and low confidence.

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand