



Email	l.a.kelley@imperial.ac.uk
Description	P0AB40
Date	Thu Jan 5 11:14:34 GMT 2012
Unique Job ID	7d9604fe5b842cc0


Figure 1 displays the protein structure and disorder analysis of the N-terminal region of the protein. The figure is divided into two panels. The top panel shows the full sequence (1-60) with secondary structure (SS) and disorder confidence. The bottom panel shows a zoomed-in view of the region from residue 70 to 80, highlighting the predicted alpha-helices and the disorder confidence. The sequence is MKNVKTLLAAAILSSMSFASF AAVEVQSTPE GQKVG TIS ANAGTNL G S L E E Q L A Q K A D E. The secondary structure shows two alpha-helices (green) and a loop (blue). The disorder confidence is high (red) for the first 10 residues and the last 10 residues, and low (green) for the middle 40 residues.

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand