



Email	l.a.kelley@imperial.ac.uk
Description	P19636
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Figure 1: Structural analysis of the protein sequence. The figure displays the protein sequence (MDQKQIEEIVRSVMASMGQAAPAPSEAKCATTNCAAPVTS ESCALDLGSAEAKAWIGVEN) and its predicted secondary structure (SS) and disorder. The sequence is shown in a color-coded format, with residues grouped into segments. The secondary structure is represented by green cylinders (alpha-helices) and blue arrows (beta-strands). The disorder is indicated by a red line with green segments. The disorder confidence is shown as a bar chart below the disorder line. The disorder is predicted to be high in the N-terminal region (residues 1-10) and low in the C-terminal region (residues 50-60).

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand