



Email	l.a.kelley@imperial.ac.uk
Description	P0AA04
Date	Thu Jan 5 11:11:42 GMT 2012
Unique Job ID	8b5b30cafa3af001


Figure 1 displays the protein structure and disorder analysis of the C-terminal region of the protein. The top panel shows the full sequence (MFQQEV TITAPNGLHTRPAAQFVKEAKGFTSEITVTSNGKSASAKSLFKLQTLGLTQGTV) with secondary structure elements (SS) and disorder confidence. The bottom panel shows a zoomed-in view of the C-terminal region (VTISAEGEDEQKAVEHLVKLMAELE) with secondary structure elements and disorder confidence. The protein is shown as a blue ribbon structure with green helices and red loops. The disorder confidence is indicated by a color scale from red (high) to blue (low).

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand