



Email	l.a.kelley@imperial.ac.uk
Description	P0CF69
Date	Thu Jan 5 11:31:29 GMT 2012
Unique Job ID	2eed4a21b974df58


Protein structure and disorder analysis for the full-length protein (1-60). The top panel shows the full sequence MTKTVSTSKKPKRKQHSPEFRSEALKLAERIGVTAARELSLYESQLYNWRSKQQNQQTSS. Below the sequence are four tracks: Secondary structure (green alpha-helices), SS (red line), confidence (yellow/green bars), and Disorder (red line with question marks). The bottom panel shows a zoomed-in view of residues 70-90: ERELEMSTEIARLKRQLAERDEELAI LQKAATYFAKRLK. It includes the same four tracks for this region.

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand