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| Description | P0CF72 |
| Date | Thu Jan 5 11:31:35 GMT 2012 |
| Unique Job ID | f8ec8c8971ab64ed |


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 (SS) represented by green alpha-helices, confidence (red bar), Disorder (green bar), and Disorder confidence (red bar). The bottom panel shows a zoomed-in view of residues 70-90 with sequence ERELEMSTEIARLKRQLAERDEELAI LQKAATYFAKRLK. It includes the same four tracks for this region.

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand