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Description	P0CF71
Date	Thu Jan 5 11:31:34 GMT 2012
Unique Job ID	ae0d3ddb3ea1543ad

Protein structure and disorder analysis for the full-length protein (1-60). The sequence is MT KTVSTSK KPRKQHSPEFRSEALKLAERI GVTAAARELSLYESQLYNWRSKQQNQQTSS. The secondary structure is shown as a green ribbon model. The confidence score is indicated by a red bar. The disorder score is indicated by a bar with question marks. The disorder confidence is indicated by a bar with question marks.

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand