



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Description	P0AD53
Date	Thu Jan 5 11:20:02 GMT 2012
Unique Job ID	c9d6c874125ba49c


Protein structure and disorder analysis for the first two domains of the protein. The top section shows the first 60 residues (MYLRPDEVARVLEKVGFTVDVVTQKAYGYRRGENYVYVNRREARMGRTALVIHPTLKERS). The bottom section shows residues 70-110 (TLAEPASDIKTC DHYQQFPLYLAGERHEHYGI PHGFSSRVALER YLNGLFGEAS). Both sections include sequence, secondary structure (SS), SS confidence, disorder, and disorder confidence tracks.

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand