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Description	C6Y4A7
Date	Sun Jul 8 11:45:05 BST 2012
Unique Job ID	f564deef9d77addc


Protein structure and disorder analysis for the full-length protein (1-100). The top panel shows the full-length protein (1-100) with sequence, secondary structure, SS confidence, disorder, and disorder confidence. The bottom panel shows a zoomed-in view of the C-terminal region (70-100) with the same metrics. The protein is primarily alpha-helical with some beta-strands. Disorder is predicted in several regions, particularly at the N-terminus and C-terminus.

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand