

Email	l.a.kelley@imperial.ac.uk
Description	P0AAD2
Date	Thu Jan 5 11:12:35 GMT 2012
Unique Job ID	50277a1f282ad8a2

Protein structure analysis of the full-length protein (1-410 aa). The figure displays the amino acid sequence, secondary structure, and disorder predictions for the entire protein. The sequence is color-coded by physicochemical properties. The secondary structure is shown as a green ribbon model. The disorder prediction is shown as a red line with green segments indicating regions of low disorder confidence. The disorder confidence is shown as a green line with red segments indicating regions of high disorder confidence. The disorder prediction is shown as a red line with green segments indicating regions of low disorder confidence. The disorder confidence is shown as a green line with red segments indicating regions of high disorder confidence.

Confidence Key
High(9)  Low (0)
? Disordered
 Alpha helix
 Beta strand