



Email	l.a.kelley@imperial.ac.uk
Description	P0ACW6
Date	Thu Jan 5 11:19:17 GMT 2012
Unique Job ID	1dc402a5b0f4f22e


Protein structure and disorder analysis for the first protein (1-60). The sequence is MFPEYRDLSRLKNENPRFMSLFDKHNKL DHEI ARKEGSDGRGYNAE VVRMKKQKLQLKD. The secondary structure is shown as a green ribbon model. The disorder analysis shows a high confidence disorder region from residue 1 to 40, followed by a low confidence region from residue 41 to 60.

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand