



Email	I.a.kelley@imperial.ac.uk
Description	P76502
Date	Thu Jan 5 12:23:39 GMT 2012
Unique Job ID	f55ae73e4baa3d17


Figure 1 displays the protein structure and disorder analysis for the protein. The protein is shown as a blue ribbon structure. The secondary structure (SS) is predicted using green cylinders (alpha-helices) and blue arrows (beta-strands). The confidence scores are shown as a red bar. The disorder analysis is shown as a grey bar with question marks indicating regions of uncertainty. The protein is composed of several alpha-helices and beta-strands, with a high degree of disorder in the N-terminal region.

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand