



Email	l.a.kelley@imperial.ac.uk
Description	P77294
Date	Thu Jan 5 12:27:20 GMT 2012
Unique Job ID	e38bd1cdb42e00b1


Figure 1 displays the protein structure and disorder analysis of the human protein p115. The figure is divided into three horizontal panels, each showing a different segment of the protein. Each panel includes the amino acid sequence, secondary structure (represented by blue arrows for alpha-helices and green cylinders for beta-strands), solvent accessibility (SS, represented by a color scale from red to green), disorder (represented by a line with question marks), and disorder confidence (represented by a color scale from red to green). The top panel shows residues 1-60, the middle panel shows residues 70-120, and the bottom panel shows residues 130-160. The protein is composed of several alpha-helices and beta-strands, with varying levels of disorder and confidence throughout its length.

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand