



Email	l.a.kelley@imperial.ac.uk
Description	P75884
Date	Thu Jan 5 12:15:36 GMT 2012
Unique Job ID	d1268bede0ac6d86


Figure 1 displays the protein structure and disorder analysis of the human protein encoded by LOC101929. The figure shows the full-length protein (608 amino acids) and three fragments (1-60, 70-120, and 130-180) in ribbon representation. Below each structure, a series of tracks show sequence, secondary structure, SS confidence, disorder, and disorder confidence. The full-length protein is shown in green, while the fragments are shown in blue. The disorder tracks indicate regions of predicted disorder with varying confidence levels.

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand