



Email	l.a.kelley@imperial.ac.uk
Description	P75991
Date	Thu Jan 5 12:17:03 GMT 2012
Unique Job ID	9960ace92267e39e


Protein structure and disorder analysis for the first two proteins. The top protein (60 residues) has a sequence MHQNSVTLDSAGAITRYFAKANLHTQQETLGEIVTEILKDGRLSRKSLCAKLLCRLEHA. It shows a secondary structure with two alpha-helices and a large disordered region. The bottom protein (20 residues) has a sequence TGE EEQKHYNALIGLLFE. It shows a secondary structure with one alpha-helix and a large disordered region.

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand