

Email	l.a.kelley@imperial.ac.uk
Description	P08337
Date	Thu Jan 5 11:01:08 GMT 2012
Unique Job ID	c59d8386228ef464

Protein structure prediction results for three protein segments. Each segment is shown with its amino acid sequence, secondary structure (arrows for helices, coils for loops), solvent accessibility (SS) (colored bars), disorder (colored bars), and disorder confidence (colored bars). The first two segments are longer, with the second segment having residue numbers 1-60 and 70-120 marked. The third segment is shorter, with residue numbers 1-6 marked.

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand