



Email	l.a.kelley@imperial.ac.uk
Description	P60723
Date	Thu Jan 5 12:07:06 GMT 2012
Unique Job ID	7d2d6f710a02719a


Figure 1 displays the secondary structure prediction of the protein sequence. The sequence is shown in a color-coded format (M, E, L, V, L, K, D, A, Q, S, A, L, T, V, S, E, T, T, F, G, R, D, F, N, E, A, L, V, H, Q, V, V, V, A, Y, A, A, G, A, R, Q, G, T, R, A, Q, K, T, R, A, E, V, T, G, S, G, K, K, P, W). The secondary structure is represented by a blue arrow indicating a beta-strand and green cylinders representing alpha-helices. The SS confidence is shown as a bar chart with red (high confidence) and yellow (low confidence) segments. The Disorder and Disorder confidence are also shown as bar charts.

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand