



Email	l.a.kelley@imperial.ac.uk
Description	P00894
Date	Thu Jan 5 10:57:05 GMT 2012
Unique Job ID	0a18f202f7ad3100


Figure 1 displays the secondary structure prediction of the protein sequence. The sequence is shown in a color-coded font (M, R, R, I, L, S, V, L, L, E, N, E, S, G, A, L, S, R, V, I, G, L, F, S, Q, R, G, Y, N, I, E, S, L, T, V, A, P, T, D, D, P, T, L, S, R, M, T, I, Q, T, V, G, D, E, K, V, L, E, Q, I). The secondary structure is represented by blue arrows (alpha-helices) and green cylinders (beta-sheets). The SS confidence is shown as a red bar, and the Disorder confidence is shown as a blue bar. The Disorder prediction shows regions of high confidence (blue) and low confidence (red/yellow).

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand