


Email	l.a.kelley@imperial.ac.uk
Description	P10442
Date	Thu Jan 5 11:32:18 GMT 2012
Unique Job ID	9e8edf6b7ccedacd


This panel provides a comprehensive view of the protein structure prediction results for the full-length protein. It displays the sequence, secondary structure, and confidence scores across the entire length of the protein, from residue 1 to 180. The sequence is shown in a color-coded format, and the secondary structure is represented by green arrows indicating alpha-helices and blue arrows indicating beta-strands. The confidence scores are shown as a bar chart below the sequence, with colors indicating the level of confidence (red for high, yellow for medium, and green for low). The disorder scores are shown as a bar chart below the confidence scores, with colors indicating the level of disorder (red for high, yellow for medium, and green for low).


Residue	Sequence	Secondary structure	SS confidence	Disorder	Disorder confidence
1	M		High	Low	High
10	I		High	Low	High
20	V		High	Low	High
30	A		High	Low	High
40	P		High	Low	High
50	S		High	Low	High
60	E		High	Low	High
70	K		High	Low	High
80	L		High	Low	High
90	M		High	Low	High
100	A		High	Low	High
110	D		High	Low	High
120	P		High	Low	High
130	K		High	Low	High
140	V		High	Low	High
150	A		High	Low	High
160	Q		High	Low	High
170	H		High	Low	High
180	T		High	Low	High

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand