



Email	l.a.kelley@imperial.ac.uk
Description	P37005
Date	Thu Jan 5 11:54:16 GMT 2012
Unique Job ID	14a50a95ece1fa63


This figure displays the full protein structure prediction results for the protein sequence. The sequence is shown in three segments: 1-60, 70-120, and 130-180. The predicted secondary structure is shown as green cylinders (alpha-helices) and blue arrows (beta-strands). The confidence is shown as a red bar, and the disorder is shown as a blue bar. The sequence is: M R I T I I L V A P A R A E N I G A A A R A M K T M G F S D L R I V D S Q A H L E P A T R V V A H G S G D I I D N I K V F P T L A E S L H D V D F T V A T T A R S R A K Y H Y A T P V E L V P L L E E K S S V M S H A A L V F G R E D S G L T N E E L A L A D V L T G V P M V A D Y P S L N L G Q A V M V Y C Y Q L A T L I Q Q P A K S D A T A D Q H Q L Q A L R E R A M T L L T T L A V A D D I K L V D W L Q Q R L G L L E Q R D T A M L H R L L H D I E K N I T K.

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand