

| | |
|---------------|-----------------------------|
| Email | l.a.kelley@imperial.ac.uk |
| Description | P0A8H8 |
| Date | Thu Jan 5 11:07:52 GMT 2012 |
| Unique Job ID | 3060060ee479bb0b |

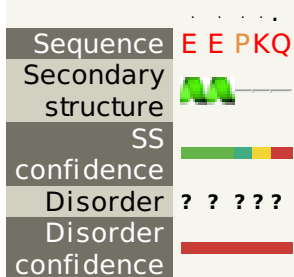
Sequence: MS ETI TVNCP TCGKT VVWGEI SPFRP FC S K R C Q L I D L G E W A A E E K R I P S S G D L S E S D D W S

Secondary structure: [Alpha-helices and a beta-strand are shown as green cylinders and arrows, respectively, starting around residue 25 and ending around residue 45.]


SS confidence: [Confidence scores for secondary structure are shown as a bar chart with colors ranging from red (low) to green (high).]

Disorder: [Disordered regions are indicated by question marks (?) above the sequence, starting from residue 1 to 10, 15 to 20, 25 to 30, 35 to 40, 45 to 50, and 55 to 60.]


Disorder confidence: [Confidence scores for disorder are shown as a bar chart with colors ranging from red (low) to green (high).]




Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand