


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|---------------|-----------------------------|
| Email         | l.a.kelley@imperial.ac.uk   |
| Description   | Q46891                      |
| Date          | Thu Jan 5 12:35:33 GMT 2012 |
| Unique Job ID | ef7c62b1c0f971e0            |


This section displays the full-length protein structure prediction results. The protein is 250 residues long. The sequence is shown in a color-coded format. The secondary structure is represented by green cylinders (alpha-helices) and blue arrows (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, 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, green for low). The disorder scores are marked with '???' at positions 1, 2, and 3.


| Sequence  | Secondary structure                                | SS confidence                            | Disorder                            | Disorder confidence                            |
|---|--|--|-------------------------------------|--|
| 1 M P R F A A N L S M M F T E V P F I E R F A A A R K A G F D A V E F L F P Y N Y S T L Q I Q K Q L E Q N H L T L A L F N   | [Secondary structure diagram for residues 1-60]    | [SS confidence bar for residues 1-60]    | [Disorder bar for residues 1-60]    | [Disorder confidence bar for residues 1-60]    |
| 70 T A P G D I N A G E W G L S A L P G R E H E A H A D I D L A L E Y A L A L N C E Q V H V M A G V V P A G E D A E R Y R A  | [Secondary structure diagram for residues 70-120]  | [SS confidence bar for residues 70-120]  | [Disorder bar for residues 70-120]  | [Disorder confidence bar for residues 70-120]  |
| 130 V F I D N I R Y A A D R F A P H G K R I L V E A L S P G V K P H Y L F S S Q Y Q A L A I V E E V A R D N V F I Q L D T F | [Secondary structure diagram for residues 130-180] | [SS confidence bar for residues 130-180] | [Disorder bar for residues 130-180] | [Disorder confidence bar for residues 130-180] |
| 190 H A Q K V D G N L T H L I R D Y A G K Y A H V Q I A G L P D R H E P D D G E I N Y P W L F R L F D E V G Y Q G W I G C E | [Secondary structure diagram for residues 190-240] | [SS confidence bar for residues 190-240] | [Disorder bar for residues 190-240] | [Disorder confidence bar for residues 190-240] |
| 250 Y K P R G L T E E G L G W F D A W R   | [Secondary structure diagram for residues 250]     | [SS confidence bar for residues 250]     | [Disorder bar for residues 250]     | [Disorder confidence bar for residues 250]     |

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand