

|               |                             |
|---------------|-----------------------------|
| Email         | l.a.kelley@imperial.ac.uk   |
| Description   | P64481                      |
| Date          | Thu Jan 5 12:08:48 GMT 2012 |
| Unique Job ID | 51f3c11526153b95            |

Sequence: MTAEGHLLFSIACAVFAKNAELTPVLAQGDWWHIVPSAILTCLLPDIDHPKSFLLGQRLKWW

Secondary structure: [Alpha-helices in green, Beta-strands in blue]

SS: [Solvent Accessibility Plot]

confidence: [Confidence Scores]

Disorder: [Disorder Prediction]

Disorder confidence: [Disorder Confidence]

Sequence: ISKPIARAFGHRGFTHSLLAVFALLATFYLVKVEGWFI PADALQGMVLGYLSHILADMLT

Secondary structure: [Alpha-helices in green, Beta-strands in blue]

SS: [Solvent Accessibility Plot]

confidence: [Confidence Scores]

Disorder: [Disorder Prediction]

Disorder confidence: [Disorder Confidence]

Sequence: PAGVPLLWPCRWRFRLPILVPQKGNQLERFICMALFVWSVWMPHSLPENSAVRWSSQMIN

Secondary structure: [Alpha-helices in green, Beta-strands in blue]

SS: [Solvent Accessibility Plot]

confidence: [Confidence Scores]

Disorder: [Disorder Prediction]

Disorder confidence: [Disorder Confidence]

Sequence: TLQIQFHRLLKHQVEY

Secondary structure: [Alpha-helices in green, Beta-strands in blue]


SS: [Solvent Accessibility Plot]

confidence: [Confidence Scores]


Disorder: [Disorder Prediction]


Disorder confidence: [Disorder Confidence]

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand