

Email	l.a.kelley@imperial.ac.uk
Description	P69506
Date	Thu Jan 5 12:11:46 GMT 2012
Unique Job ID	a3c8380d748df140

Sequence: MAYRDQPLGELALSI PRASALFRKYDMDYCCGGKQTLARAAARKELDVEVIEAELAKLAE

Secondary structure: [Ribbon diagram showing alpha-helices and beta-strands]

SS confidence: [Color scale from 0.5 to 1.0]

Disorder: [Red line indicating predicted disorder]

Disorder confidence: [Color scale from 0.0 to 1.0]

Sequence: QPIEKDWRSAPLAEI IDHI VRYHDRHREQLPELILQATKVERVHADKPSVPKGLTKYLT

Secondary structure: [Ribbon diagram showing alpha-helices and beta-strands]

SS confidence: [Color scale from 0.5 to 1.0]

Disorder: [Red line indicating predicted disorder]

Disorder confidence: [Color scale from 0.0 to 1.0]

Sequence: MLHEELSSHMMKEEQILFPMIKQGMGSQAMGPISVMESEHDEAGELLEVIKHTTNNTTPP

Secondary structure: [Ribbon diagram showing alpha-helices and beta-strands]

SS confidence: [Color scale from 0.5 to 1.0]

Disorder: [Red line indicating predicted disorder]

Disorder confidence: [Color scale from 0.0 to 1.0]

Sequence: PEACTTWKAMYNGINELIDDLMDHISLENNVLFPRALAGE


Secondary structure: [Ribbon diagram showing alpha-helices and beta-strands]

SS confidence: [Color scale from 0.5 to 1.0]


Disorder: [Red line indicating predicted disorder]


Disorder confidence: [Color scale from 0.0 to 1.0]

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand