

Email	l.a.kelley@imperial.ac.uk
Description	P31062
Date	Thu Jan 5 11:46:59 GMT 2012
Unique Job ID	fe19f50d71ff2b31

Sequence: MEV NKKQLADI F GAS I RTI QNWQE QGMP VL RGGGKGNEVL Y DSA AVI K WY A ERDAEI ENE

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

SS confidence: [Bar chart showing confidence scores for each residue]

Disorder: [Bar chart showing disorder scores for each residue]

Disorder confidence: [Bar chart showing confidence scores for each residue]

Sequence: KLRREVEELLQASETDLP GTI EYERHRLTRAQADAQELKNARDSA EVVETAFCTFVLSR

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

SS confidence: [Bar chart showing confidence scores for each residue]

Disorder: [Bar chart showing disorder scores for each residue]

Disorder confidence: [Bar chart showing confidence scores for each residue]

Sequence: IAGEIASILDGIP LSVQRRFP ELENRHVDF LKRDI I KAMNKAAALDELI PGL LSEYI EQS

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

SS confidence: [Bar chart showing confidence scores for each residue]

Disorder: [Bar chart showing disorder scores for each residue]

Disorder confidence: [Bar chart showing confidence scores for each residue]

Sequence: G


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

SS confidence: [Bar chart showing confidence scores for each residue]


Disorder: [Bar chart showing disorder scores for each residue]


Disorder confidence: [Bar chart showing confidence scores for each residue]

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand