

Email	l.a.kelley@imperial.ac.uk
Description	P0A8I3
Date	Thu Jan 5 11:07:54 GMT 2012
Unique Job ID	9c3777e0b9be5303

Sequence: MLILISP AKTLDYQSPLTTTTRYTLPELLDNSQQLIHEARKLTTPPQISTLMRISDKLAGIN

Secondary structure: [Alpha-helices and loops]

SS confidence: [Bar chart]

Disorder: [Bar chart]

Disorder confidence: [Bar chart]

Sequence: AARFHDWQPDFTPANARQAILAFKGDVYTG LQAETFSEDDFDFAQQHLRMLSGLYGVL RP

Secondary structure: [Alpha-helices and loops]

SS confidence: [Bar chart]

Disorder: [Bar chart]

Disorder confidence: [Bar chart]

Sequence: LDLMQPPYRLEMGIRLENARGKLDLYQFWGDIITNKLNEALAAQGDNVVINLASDEYFKSVK

Secondary structure: [Alpha-helices and loops]

SS confidence: [Bar chart]

Disorder: [Bar chart]

Disorder confidence: [Bar chart]

Sequence: PKKLNAEIIKPVFLDEKNGKFKIISFYAKKARGLMSRFIENRLTKPEQLTGFNSEGYFF

Secondary structure: [Alpha-helices and loops]

SS confidence: [Bar chart]

Disorder: [Bar chart]

Disorder confidence: [Bar chart]

Sequence: DEDSSNGELVKRYEQR


Secondary structure: [Alpha-helices and loops]

SS confidence: [Bar chart]


Disorder: [Bar chart]


Disorder confidence: [Bar chart]

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand