

Email	l.a.kelley@imperial.ac.uk
Description	P37194
Date	Thu Jan 5 11:55:00 GMT 2012
Unique Job ID	33ca6ece8a835ac5

Sequence: MNMTK GALI LSL SFL LAACSSI PQNI KGNN QPDI QKS FVAV VHN QPGL YVG QQAR FGGKVI

Secondary structure: [Alpha-helices and beta-strands]

SS: [Secondary structure elements]

confidence: [Confidence bar]

Disorder: [Disorder prediction]

Disorder confidence: [Disorder confidence bar]

Sequence: NVI NGKTDTLLEI SVLP LDSY AKPDIEANY QGRLL ARQS GFLDPV NYRNHFVTILGTI QG

Secondary structure: [Alpha-helices and beta-strands]

SS: [Secondary structure elements]

confidence: [Confidence bar]

Disorder: [Disorder prediction]

Disorder confidence: [Disorder confidence bar]

Sequence: EQPGFI NKVPYNFLEV NMQGI QVWHLREV VNTTYNLWDYGYGAF WPEPGWGAPYYTNAV S

Secondary structure: [Alpha-helices and beta-strands]

SS: [Secondary structure elements]

confidence: [Confidence bar]

Disorder: [Disorder prediction]

Disorder confidence: [Disorder confidence bar]

Sequence: QVTPELVK

Secondary structure: [Alpha-helices and beta-strands]


SS: [Secondary structure elements]

confidence: [Confidence bar]


Disorder: [Disorder prediction]


Disorder confidence: [Disorder confidence bar]

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand