

Email	l.a.kelley@imperial.ac.uk
Description	P0AA86
Date	Thu Jan 5 11:12:11 GMT 2012
Unique Job ID	0c73a6363d0a388c

Sequence: MKRKVLLIPLIIFLAIAAALLWQLARNAEGDDPTNLESA LIGKPVPKFRL ESLDNP GQFY

Secondary structure: [Alpha-helices and beta-strands represented by green and blue cylinders/arrows]

SS: [Solvent accessibility bar]

confidence: [Confidence scores for secondary structure]

Disorder: [Disorder prediction with question marks]

Disorder confidence: [Disorder confidence bar]

Sequence: QADVLTQGKPVLLNVWATWCP TCRAEHQYLNQLSAQGI R VVG MNYKDDRQKAI SWL KELG

Secondary structure: [Alpha-helices and beta-strands represented by green and blue cylinders/arrows]

SS: [Solvent accessibility bar]

confidence: [Confidence scores for secondary structure]

Disorder: [Disorder prediction with question marks]

Disorder confidence: [Disorder confidence bar]

Sequence: NPYALSLFDG D GMLGLDLGVYGAPETFLIDNGIIRYRHAGDLNPRVWEE EIKPLWEKYS

Secondary structure: [Alpha-helices and beta-strands represented by green and blue cylinders/arrows]

SS: [Solvent accessibility bar]

confidence: [Confidence scores for secondary structure]

Disorder: [Disorder prediction with question marks]

Disorder confidence: [Disorder confidence bar]

Sequence: KEAAQ

Secondary structure: [Alpha-helices and beta-strands represented by green and blue cylinders/arrows]


SS: [Solvent accessibility bar]

confidence: [Confidence scores for secondary structure]


Disorder: [Disorder prediction with question marks]


Disorder confidence: [Disorder confidence bar]

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand