

Email	l.a.kelley@imperial.ac.uk
Description	P76423
Date	Thu Jan 5 12:22:57 GMT 2012
Unique Job ID	9168401db90bf9e2

Sequence: MQLVLLGSAQSAHALHLFHQHSPLVHCMTNDVVQTF TANTLLALGAS PAMVIETEEASQF

Secondary structure: [Helices and Strands]

SS confidence: [Color-coded bar]

Disorder: [Disordered regions]

Disorder confidence: [Color-coded bar]

Sequence: AAIASALLINVGTLTQPRAQAMRAAVEQAKSSQTPWTLDPVAVGALDYRRHFHELLSF K

Secondary structure: [Helices and Strands]

SS confidence: [Color-coded bar]

Disorder: [Disordered regions]

Disorder confidence: [Color-coded bar]

Sequence: PAAIRGNASEIMALAGIANGRGVDTTDAAANAI PAAQTLARETGAI VVV TGEMDYVT DG

Secondary structure: [Helices and Strands]

SS confidence: [Color-coded bar]

Disorder: [Disordered regions]

Disorder confidence: [Color-coded bar]

Sequence: HRIIGIHGGDPLMTKVVG TG CALSAVVAA CCALPGDTLENVASACHWMKQAGERAVARSE

Secondary structure: [Helices and Strands]

SS confidence: [Color-coded bar]

Disorder: [Disordered regions]

Disorder confidence: [Color-coded bar]

Sequence: GPGSFVPHFLDALWQLTQEVQA


Secondary structure: [Helices and Strands]

SS confidence: [Color-coded bar]


Disorder: [Disordered regions]


Disorder confidence: [Color-coded bar]

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand