

Email	l.a.kelley@imperial.ac.uk
Description	P0ABL1
Date	Thu Jan 5 11:15:45 GMT 2012
Unique Job ID	e43e923068ffec71

Sequence: MSVLRSLLTAGVLA SGLLWSLNGITATPAAQASDDRYEVTQQRNPDAACLDCHKPDT EGM

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

SS confidence: [Color scale from red to green]

Disorder: [Color scale from red to green]

Disorder confidence: [Color scale from red to green]

Sequence: HGKHASVINPNNKL PVTCTNCHGQSPQHR EGVKDV MRFN EPMYKVGEQNSVCM SCHLPE

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

SS confidence: [Color scale from red to green]

Disorder: [Color scale from red to green]

Disorder confidence: [Color scale from red to green]

Sequence: QLQKAFWP HDVHVTKVACASCHSLHPQQDTMQT L S DKGRIKI CVDC HS DQRTNP NFPAS

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

SS confidence: [Color scale from red to green]

Disorder: [Color scale from red to green]

Disorder confidence: [Color scale from red to green]

Sequence: VPLLKEQP


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

SS confidence: [Color scale from red to green]


Disorder: [Color scale from red to green]


Disorder confidence: [Color scale from red to green]

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand