

Email	l.a.kelley@imperial.ac.uk
Description	P0ADV1
Date	Thu Jan 5 11:21:54 GMT 2012
Unique Job ID	7e2b64c05d90dac9

Sequence: MKFKTNKLSLNLVLASSLLAASI PAFAVTGD TDQPI HI ES DQQS LDMQGNVVTFTGNVI V

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

SS confidence: [Confidence scale]

Disorder: [Disorder scale]

Disorder confidence: [Disorder confidence scale]

Sequence: TQGTI KINADKVVVTRPGGEQGKEVI DGYGKPATFYQM QDNGKPV EGHAS QMHYELAKDF

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

SS confidence: [Confidence scale]

Disorder: [Disorder scale]

Disorder confidence: [Disorder confidence scale]

Sequence: VVLTGNAYLQQVDSNIKGDKITLVKEQKMQAFSDKGKRVTTVLVPSQLQDKNNKGGQT PA

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

SS confidence: [Confidence scale]

Disorder: [Disorder scale]

Disorder confidence: [Disorder confidence scale]

Sequence: QKKGN


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

SS confidence: [Confidence scale]


Disorder: [Disorder scale]


Disorder confidence: [Disorder confidence scale]

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand