

Email	l.a.kelley@imperial.ac.uk
Description	Q46938
Date	Thu Jan 5 12:35:57 GMT 2012
Unique Job ID	4aad2dc4166e23c8

	1	10	20	30	40	50	60
Sequence	MDVRQSI HSAHAKTLDTQGLRNEFLVEKVFVADEYTMVYSHI DRIIVGGIMPTIKTVSVG						
Secondary structure							
SS confidence							
Disorder	? ? ? - ? ? ? - ? ? ? - ? ? ? - ? ? ? - ? ? ?						
Disorder confidence							

  

	70	80	90	100	110	120
Sequence	GEV GKQLGVSYFLERRELGVINIGGAGTITVDGQCYEIGHRDALYVGKGAKEVVFASIDT					
Secondary structure						
SS confidence						
Disorder	? ? ? ? - ? ? ? ? - ? ? ? ? - ? ? ? ? - ? ? ? ?					
Disorder confidence						

  

	130	140	150	160	170	180
Sequence	GTPAKFYFNCAPAHTTYP TKKVTPDEVSPVTLGDNLTSNRRTINKYFVPDVLETCLSMG					
Secondary structure						
SS confidence						
Disorder	? ? - ? ? ? ? ?					
Disorder confidence						

  

	190	200	210	220	230	240
Sequence	LTE LAPGNLWNTMPCHTHERRMEVYFYFNMDDDACV F HMMGQPQETRHI VMHNEQAVISP					
Secondary structure						
SS confidence						
Disorder						
Disorder confidence						

  

	250	260	270
Sequence	SWSI HSGVGT KAYTFI WGMVGENQV FDDMDHVAVKDLR		
Secondary structure			
SS confidence			
Disorder	? - ? ? ?		
Disorder confidence			

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand