

Email	l.a.kelley@imperial.ac.uk
Description	P39384
Date	Thu Jan 5 12:00:17 GMT 2012
Unique Job ID	f4c268a9b0ab3567

	1	10	20	30	40	50	60
Sequence	MSLVTDLPALFDQFSEARQTGFLTVMDLKERGIPLVGTCTFMPQEIPMAAGAVVVS LCS						
Secondary structure							
SS confidence							
Disorder	? ? ?						
Disorder confidence							

  

	70	80	90	100	110	120
Sequence	TSDETIEEAEKDLPRNLCPLIKSSYGFGKTDKCPYFYFSDLVVGETTCDGKKKMYEYMAE					
Secondary structure						
SS confidence						
Disorder	? ? ?					
Disorder confidence						

  

	130	140	150	160	170	180
Sequence	FKPVHVMQLPNSVKDDASRALWKAEMLRRLQKTVEERFGHEISEDALRDAIALKNRERRAL					
Secondary structure						
SS confidence						
Disorder						
Disorder confidence						

  

	190	200	210	220	230	240
Sequence	ANFYHLGQLNPPALSGSDILKVVYGATFRFDKEALINELDAMTARVRQQWEEGQRLDPRP					
Secondary structure						
SS confidence						
Disorder	? ? ? ? ? ?					
Disorder confidence						

  

	250	260	270	280	290	300
Sequence	RILITGCPIMGAAEKVVRAIEENGWVVGYNCTGAKATEQCVAETGDVYDALADKYLAI					
Secondary structure						
SS confidence						
Disorder						
Disorder confidence						

  

	310	320	330	340	350	360
Sequence	GCSCVSPNDQRLKMLSQMVVEEYQVDGVVDVILQACHTYAVESLAIKRHVRQQHNIPIYIAI					
Secondary structure						
SS confidence						
Disorder	? ?					
Disorder confidence						

  

	370	380
Sequence	ETDYSTSDVGQLSTRVAAFI EML	
Secondary structure		
SS confidence		
Disorder	? ? ? ? ?	
Disorder confidence		

Confidence Key  
High(9)  Low (0)  
? Disordered  
 Alpha helix  
 Beta strand