

Email	l.a.kelley@imperial.ac.uk
Description	Q47537
Date	Thu Jan 5 12:36:50 GMT 2012
Unique Job ID	2599b706bb0d7216

	1	10	20	30	40	50	60
Sequence	MAI SSRNTLLAALAFI AFQAQAVNVTVAYQTS AEPAKV AQADNTFAKESGATVDWRKFDS						
Secondary structure							
SS confidence							
Disorder	? ? ? ? ? ? ? ? ? ?						
Disorder confidence							
	70	80	90	100	110	120	
Sequence	GASIVRALASGDVQIGNLGSSPLAVAASQQVPI EVFLLASKLGNSEALVV KKTISKPEDL						
Secondary structure							
SS confidence							
Disorder							
Disorder confidence							
	130	140	150	160	170	180	
Sequence	IGKR IAVPFI STTHYSLLAALKHWGI KPGQVEI VNLQPPAII AA WQRGDI DGAYV WAPAV						
Secondary structure							
SS confidence							
Disorder							
Disorder confidence							
	190	200	210	220	230	240	
Sequence	NALEKDGKVL TDSEQVGQWGAPTLDVWVVRKDF AEKHPEVVKAF AKSAI DAQQPYI ANPD						
Secondary structure							
SS confidence							
Disorder	?						
Disorder confidence							
	250	260	270	280	290	300	
Sequence	VWLKQPENI SKLARLSGVP EGDVPGLVKGNTYLT PQQQTAE LTGPVNKAI I DTAQFLKEQ						
Secondary structure							
SS confidence							
Disorder	? ? ? ? ? ? ? ?						
Disorder confidence							
	310	320					
Sequence	GKVP AVANDY SQYVTSR FVQ						
Secondary structure							
SS confidence							
Disorder	?						
Disorder confidence							

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand