

Email	l.a.kelley@imperial.ac.uk
Description	P0A7V0
Date	Thu Jan 5 11:06:16 GMT 2012
Unique Job ID	2f820f40d1a95dca

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

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

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


  

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


  


	250
Sequence	E
Secondary structure	
SS confidence	
Disorder	?
Disorder confidence	

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand