

Email	l.a.kelley@imperial.ac.uk
Description	P77580
Date	Thu Jan 5 12:30:45 GMT 2012
Unique Job ID	f248ccac7f3d09a4

	1	10	20	30	40	50	60
Sequence	MSKRKVAII GSGNI GTDLMI KILRHGQHLE MAVMVGIDPQSDGLARARRMGVATTHEGVII						
Secondary structure							
SS confidence							
Disorder	? ? ? ?						
Disorder confidence							
	70	80	90	100	110	120	
Sequence	GLMNMPEFADI DIVDATSAGAHVKNDAAAL REAKPDIRLI DLTAAI GPYCVPVVNLEAN						
Secondary structure							
SS confidence							
Disorder	? ? ?						
Disorder confidence							
	130	140	150	160	170	180	
Sequence	VDQLNVNMVTCGGQATIPMVAAVSRVARVHYAEII ASIASKSAGPGTRANIDEFTETTSR						
Secondary structure							
SS confidence							
Disorder	? ? ? ? ? ? ?						
Disorder confidence							
	190	200	210	220	230	240	
Sequence	AIEVVGGAAKGKAIIVLNP AEPPLMMRDTVYVLSDEASQDDIEASINEMAEAVQAYVPGY						
Secondary structure							
SS confidence							
Disorder	?						
Disorder confidence							
	250	260	270	280	290	300	
Sequence	RLKQRVQFEVIPQDKPVNLP GVGQFSGLKTAVVLEVEGA AHYLPAYAGNLDIMTSSALAT						
Secondary structure							
SS confidence							
Disorder	? ? ? ? ? ? ?						
Disorder confidence							
	310						
Sequence	AEKMAQSLARKAGEAA						
Secondary structure							
SS confidence							
Disorder	? ? ? ? ? ? ?						
Disorder confidence							

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand