



Email	l.a.kelley@imperial.ac.uk
Description	P75777
Date	Wed Jan 25 15:21:02 GMT 2012
Unique Job ID	7b6a3872973be2ff


	1	10	20	30	40	50	60
Sequence	MMKKP VVI GL AVVVLAAVVG GY WWY QSRQD NGLT LYG NV DI RTV NLS F RV GGRV E SLAV						
Secondary structure							
SS confidence							
Disorder	? ? ? ? ? ? ? ?						
Disorder confidence							
	70	80	90	100	110	120	
Sequence	DE GDAI KAG QV L GEL DHK P Y E I ALMQAKAG V SVA QAQ Y DL ML AGY R NEE I AQAAA AV KQA						
Secondary structure							
SS confidence							
Disorder	? ? ? ?						
Disorder confidence							
	130	140	150	160	170	180	
Sequence	QAAYDY AQNF YNR QQGL WKS RTI SANDLEN ARSS RDQAQAT LKSAQDKL RQY RSGNREQD						
Secondary structure							
SS confidence							
Disorder	? ? ? ? ? ? ? ?						
Disorder confidence							
	190	200	210	220	230	240	
Sequence	IAQAKASLEQAQAQLAQAE LNLQDSTLI APSDGTLL TRAVEP GTVLNEGGTVFTVSLTRP						
Secondary structure							
SS confidence							
Disorder	?						
Disorder confidence							
	250	260	270	280	290	300	
Sequence	VWVRAYVDERNL DQAQPG RKVLL YTDGRPD KPYHGQI GFVSPTAE FTPKT VETPDLRTDL						
Secondary structure							
SS confidence							
Disorder	? ? ? ? ? ? ? ? ? ?						
Disorder confidence							
	310	320	330				
Sequence	VYRLRI VVT DADDAL RQGMP VTVQFGDEAGHE						
Secondary structure							
SS confidence							
Disorder	? ? ? ? ? ? ? ?						
Disorder confidence							

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand