

Email	l.a.kelley@imperial.ac.uk
Description	P37683
Date	Thu Jan 5 11:56:58 GMT 2012
Unique Job ID	ca7572bffffb536e0

	1	10	20	30	40	50	60
Sequence	MDLLIILTYVAFAWMFKIFKIPVNKWTIPTAALGGIFIVSGLILLMNYNHPYTFKAQKA						
Secondary structure							
SS confidence							
Disorder	? ?						
Disorder confidence							
	70	80	90	100	110	120	
Sequence	VISIPVVPQVTGVVIEVTDKKNTLIIKKGEVLFRLDPTTRYQARVDRLMADIVTAEHKQRAL						
Secondary structure							
SS confidence							
Disorder	? ? ? ? ? ? ? ? ? ? ? ? ? ? ? ? ? ?						
Disorder confidence							
	130	140	150	160	170	180	
Sequence	GAELDEMAANTQQAKATRDKFAKEYQRYARGSQAKVNPFSERDIDVARQNYLAQEASVKS						
Secondary structure							
SS confidence							
Disorder	? ? ? ? ? ? ? ? ? ? ? ? ? ? ? ? ? ?						
Disorder confidence							
	190	200	210	220	230	240	
Sequence	SAAEQKQIQSQLDSLVLGEHSQIASLKAQLAEAKYNLEQTIVRAPSDGYVTQVLI R PGT Y						
Secondary structure							
SS confidence							
Disorder	? ? ? ? ? ? ? ? ? ? ? ? ? ? ? ? ? ?						
Disorder confidence							
	250	260	270	280	290	300	
Sequence	AASLPLRPVMVFI PDQKRQI VAQFRQNSLLRLAPGDDAEVVFNALPGKVFSGKLA AISPA						
Secondary structure							
SS confidence							
Disorder	? ? ? ? ? ? ? ? ? ? ? ? ? ? ? ? ? ?						
Disorder confidence							
	310	320	330	340	350	360	
Sequence	VPGGAYQSTGTLQTLNTAPGSDGVIATIELDEHTDLSALPDGIYAQVAVYSDHFSHVSVM						
Secondary structure							
SS confidence							
Disorder	? ? ? ? ? ? ? ? ? ? ? ? ? ? ? ? ? ?						
Disorder confidence							
	370						
Sequence	RKVLLRMTSVVHYLYLDH						
Secondary structure							
SS confidence							
Disorder	? ?						
Disorder confidence							

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