

Email	l.a.kelley@imperial.ac.uk
Description	P0A9C9
Date	Thu Jan 5 11:09:54 GMT 2012
Unique Job ID	b9a816cb73286308

	1	10	20	30	40	50	60																																																					
Sequence	M	R	R	E	L	A	I	E	F	S	R	V	T	E	S	A	A	L	A	G	Y	K	W	L	G	R	G	D	K	N	T	A	D	G	A	A	V	N	A	M	R	I	M	L	N	Q	V	N	I	D	G	T	I	V	I	G	E	G	E	I
Secondary structure	[Alpha Helix]										[Alpha Helix]										[Alpha Helix]																																							
SS confidence	[Confidence Scale]																																																											
Disorder	?										?																																																	
Disorder confidence	[Disorder Confidence Scale]																																																											

	70	80	90	100	110	120																																																						
Sequence	D	E	A	P	M	L	Y	I	G	E	K	V	G	T	G	R	G	D	A	V	D	I	A	V	D	P	I	E	G	T	R	M	T	A	M	G	Q	A	N	A	L	A	V	L	A	V	G	D	K	G	C	F	L	N	A	P	D	M	Y	M
Secondary structure	[Alpha Helix]										[Alpha Helix]										[Alpha Helix]																																							
SS confidence	[Confidence Scale]																																																											
Disorder																																																												
Disorder confidence	[Disorder Confidence Scale]																																																											

	130	140	150	160	170	180																																																						
Sequence	E	K	L	I	V	G	P	G	A	K	G	T	I	D	L	N	L	P	L	A	D	N	L	R	N	V	A	A	A	L	G	K	P	L	S	E	L	T	V	T	I	L	A	K	P	R	H	D	A	V	I	A	E	M	Q	Q	L	G	V	R
Secondary structure	[Alpha Helix]										[Alpha Helix]										[Alpha Helix]																																							
SS confidence	[Confidence Scale]																																																											
Disorder																																																												
Disorder confidence	[Disorder Confidence Scale]																																																											

	190	200	210	220	230	240																																																						
Sequence	V	F	A	I	P	D	G	D	V	A	A	S	I	L	T	C	M	P	D	S	E	V	D	V	L	Y	G	I	G	G	A	P	E	G	V	V	S	A	A	V	I	R	A	L	D	G	D	M	N	G	R	L	L	A	R	H	D	V	K	G
Secondary structure	[Alpha Helix]										[Alpha Helix]										[Alpha Helix]																																							
SS confidence	[Confidence Scale]																																																											
Disorder																																																												
Disorder confidence	[Disorder Confidence Scale]																																																											


  

	250	260	270	280	290	300																																																						
Sequence	D	N	E	E	N	R	R	I	G	E	Q	E	L	A	R	C	K	A	M	G	I	E	A	G	K	V	L	R	L	G	D	M	A	R	S	D	N	V	I	F	S	A	T	G	I	T	K	G	D	L	L	E	G	I	S	R	K	G	N	I
Secondary structure	[Alpha Helix]										[Alpha Helix]										[Alpha Helix]																																							
SS confidence	[Confidence Scale]																																																											
Disorder	?																																																											
Disorder confidence	[Disorder Confidence Scale]																																																											


  


	310	320	330																																	
Sequence	A	T	T	E	T	L	L	I	R	G	K	S	R	T	I	R	R	I	Q	S	I	H	Y	L	D	R	K	D	P	E	M	Q	V	H	I	L
Secondary structure	[Alpha Helix]										[Alpha Helix]																									
SS confidence	[Confidence Scale]																																			
Disorder	?																																			
Disorder confidence	[Disorder Confidence Scale]																																			

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand