

Email	l.a.kelley@imperial.ac.uk
Description	P42912
Date	Thu Jan 5 12:02:11 GMT 2012
Unique Job ID	e2592cfc9f0fcfb6

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


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

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


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


	250										
Sequence	L	H	S	N	F	I	C	L	I	N	T
Secondary structure	[Loop]										
SS confidence	[Confidence Scale]										
Disorder	[Disorder Prediction]										
Disorder confidence	[Disorder Confidence Scale]										

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand