

Email	l.a.kelley@imperial.ac.uk
Description	P76418
Date	Thu Jan 5 12:22:53 GMT 2012
Unique Job ID	17378e692b6701cd

	1	10	20	30	40	50	60																																																					
Sequence	M	K	T	E	R	I	L	G	A	L	Y	G	Q	A	L	G	D	A	M	G	M	P	S	E	L	W	P	R	S	R	V	K	A	H	F	G	W	I	D	R	F	L	P	G	P	K	E	N	N	A	A	C	Y	F	N	R	A	E	F	T
Secondary structure	[Alpha Helix]										[Alpha Helix]																																																	
SS confidence	[Confidence Scale]																																																											
Disorder	[Disorder Prediction]																																																											
Disorder confidence	[Disorder Confidence Scale]																																																											
	70	80	90	100	110	120																																																						
Sequence	D	D	T	S	M	A	L	C	L	A	D	A	L	L	E	R	E	G	K	I	D	P	D	L	I	G	R	N	I	L	D	W	A	L	R	F	D	A	F	N	K	N	V	L	G	P	T	S	K	I	A	L	N	A	I	R	D	G	K	P
Secondary structure	[Alpha Helix]										[Alpha Helix]										[Alpha Helix]																																							
SS confidence	[Confidence Scale]																																																											
Disorder	[Disorder Prediction]																																																											
Disorder confidence	[Disorder Confidence Scale]																																																											
	130	140	150	160	170	180																																																						
Sequence	V	A	E	L	E	N	N	G	V	T	N	G	A	A	M	R	V	S	P	L	G	C	L	L	P	A	R	D	V	D	S	F	I	D	D	V	A	L	A	S	S	P	T	H	K	S	D	L	A	V	A	G	A	V	V	I	A	W	A	I
Secondary structure	[Alpha Helix]					[Alpha Helix]										[Alpha Helix]										[Alpha Helix]																																		
SS confidence	[Confidence Scale]																																																											
Disorder	[Disorder Prediction]																																																											
Disorder confidence	[Disorder Confidence Scale]																																																											
	190	200	210	220	230	240																																																						
Sequence	S	R	A	I	D	G	E	S	W	S	A	I	V	D	S	L	P	S	I	A	R	H	A	Q	Q	K	R	I	T	T	F	S	A	S	L	A	A	R	L	E	I	A	L	K	I	V	R	N	A	D	G	T	E	S	A	S	E	Q	L	Y
Secondary structure	[Alpha Helix]					[Alpha Helix]										[Alpha Helix]										[Alpha Helix]																																		
SS confidence	[Confidence Scale]																																																											
Disorder	[Disorder Prediction]																																																											
Disorder confidence	[Disorder Confidence Scale]																																																											
	250	260	270	280	290	300																																																						
Sequence	Q	V	V	G	A	G	T	S	T	I	E	S	V	P	C	A	I	A	L	V	E	L	A	Q	T	D	P	N	R	C	A	V	L	C	A	N	L	G	G	D	T	D	T	I	G	A	M	A	T	A	I	C	G	A	L	H	G	V	N	A
Secondary structure	[Alpha Helix]					[Alpha Helix]										[Alpha Helix]										[Alpha Helix]																																		
SS confidence	[Confidence Scale]																																																											
Disorder	[Disorder Prediction]																																																											
Disorder confidence	[Disorder Confidence Scale]																																																											
	310	320	330																																																									
Sequence	I	D	P	A	L	K	A	E	L	D	A	V	N	Q	L	D	F	N	R	Y	A	T	A	L	A	K	Y	R	Q	Q	R	E	A	V																										
Secondary structure	[Alpha Helix]										[Alpha Helix]																																																	
SS confidence	[Confidence Scale]																																																											
Disorder	[Disorder Prediction]																														[Disorder Prediction]																													
Disorder confidence	[Disorder Confidence Scale]																														[Disorder Confidence Scale]																													

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand