

Email	l.a.kelley@imperial.ac.uk
Description	P77704
Date	Thu Jan 5 12:31:54 GMT 2012
Unique Job ID	628b6a540ad34177

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

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

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

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

	250	260	270																																			
Sequence	V	K	E	N	Q	D	Q	P	F	L	H	L	E	R	E	V	R	K	A	V	K	E	R	A	L	E	K	I	K	L	F	G	S	D	G	K	A	E
Secondary structure	[Alpha Helix]										[Alpha Helix]										[Alpha Helix]																	
SS confidence	[Confidence Scale]																																					
Disorder	[Disorder Prediction]																																					
Disorder confidence	[Disorder Confidence]																																					

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand