

Email	l.a.kelley@imperial.ac.uk
Description	P0AB85
Date	Thu Jan 5 11:14:52 GMT 2012
Unique Job ID	e9b76d24707119c9

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

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

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

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

	250	260	270	280	290	300																																																						
Sequence	V	Q	A	V	V	D	I	N	G	H	G	I	S	T	S	G	S	Y	R	N	Y	Y	E	L	D	G	K	R	L	S	H	V	I	D	P	Q	T	G	R	P	I	E	H	N	L	V	S	V	T	V	I	A	P	T	A	L	E	A	D	A
Secondary structure	[Alpha Helix]										[Alpha Helix]										[Alpha Helix]																																							
SS confidence	[Confidence Scale]																																																											
Disorder	[Disorder Prediction]																																																											
Disorder confidence	[Disorder Confidence]																																																											

	310	320	330	340	350																																																							
Sequence	W	D	T	G	L	M	V	L	G	P	E	K	A	K	E	V	V	R	R	E	G	L	A	V	Y	M	I	T	K	E	G	D	S	F	K	T	W	M	S	P	Q	F	K	S	F	L	V	S	E	K	N									
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