

Email	l.a.kelley@imperial.ac.uk
Description	P60472
Date	Thu Jan 5 12:06:50 GMT 2012
Unique Job ID	9d691c5baa51da38

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

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

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

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

	250												
Sequence	R	R	F	G	G	T	E	P	G	D	E	T	A
Secondary structure	[Alpha Helix]												
SS confidence	[High]												
Disorder	[Disordered]												
Disorder confidence	[Low]												

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand