

Email	l.a.kelley@imperial.ac.uk
Description	P0A884
Date	Thu Jan 5 11:07:10 GMT 2012
Unique Job ID	9d2bc356928fdd43

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


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

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

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


	250	260																						
Sequence	D	Y	R	F	E	D	F	E	I	E	G	Y	D	P	H	P	G	I	K	A	P	V	A	I
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