

Email	l.a.kelley@imperial.ac.uk
Description	Q46920
Date	Thu Jan 5 12:35:53 GMT 2012
Unique Job ID	da6b85928219f7a4

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

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

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

	250	260	270	280																																																								
Sequence	L	R	F	C	Q	P	E	K	L	S	V	Y	A	R	Y	T	R	R	G	L	D	I	N	P	W	R	S	N	S	D	F	V	P	S	T	T	R	L	V	R	Q																			
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