

Email	l.a.kelley@imperial.ac.uk
Description	Q9JMR6
Date	Thu Jan 5 12:37:50 GMT 2012
Unique Job ID	6e7a4450c3df15c1

	1	10	20	30	40	50	60																																																					
Sequence	M	A	I	D	I	F	Q	S	L	K	N	C	V	F	D	L	Q	R	A	D	V	Q	N	Y	Q	Q	P	L	K	Q	L	A	R	L	L	N	S	E	N	L	Q	S	V	N	A	H	L	T	R	N	V	E	L	D	T	F	L	A	R	S
Secondary structure	[Alpha Helix]										[Alpha Helix]										[Alpha Helix]										[Alpha Helix]																													
SS confidence	[Confidence Scale]																																																											
Disorder	[Disorder Prediction]																																																											
Disorder confidence	[Disorder Confidence Scale]																																																											
	70	80	90	100	110	120																																																						
Sequence	E	D	T	E	S	S	M	A	G	S	A	V	L	Q	W	P	D	E	P	A	D	I	L	G	L	K	L	L	I	E	K	M	A	D	D	N	N	F	S	F	N	F	C	H	T	F	F	Y	D	R	N	I	I	E	S	I	R	K	F	
Secondary structure	[Alpha Helix]										[Alpha Helix]										[Alpha Helix]																																							
SS confidence	[Confidence Scale]																																																											
Disorder	[Disorder Prediction]																																																											
Disorder confidence	[Disorder Confidence Scale]																																																											
	130	140	150	160	170	180																																																						
Sequence	T	S	S	L	V	A	P	F	V	R	D	Y	Q	L	Y	V	E	N	Q	H	D	P	E	P	A	V	F	R	P	V	S	R	K	I	F	I	V	H	G	H	D	N	D	A	L	Q	S	V	A	R	F	I	S	R	I	G	L	E	E	I
Secondary structure	[Alpha Helix]										[Alpha Helix]										[Alpha Helix]																																							
SS confidence	[Confidence Scale]																																																											
Disorder	[Disorder Prediction]																																																											
Disorder confidence	[Disorder Confidence Scale]																																																											
	190	200	210	220	230	240																																																						
Sequence	I	L	S	E	R	P	D	G	S	R	T	I	I	E	K	F	E	A	E	S	G	D	V	S	F	A	I	V	L	M	T	P	D	D	S	G	S	A	L	A	S	E	S	T	R	L	R	A	R	Q	N	V	L	Y	E	L	G	Y	F	A
Secondary structure	[Alpha Helix]	[Alpha Helix]										[Alpha Helix]										[Alpha Helix]																																						
SS confidence	[Confidence Scale]																																																											
Disorder	[Disorder Prediction]																																																											
Disorder confidence	[Disorder Confidence Scale]																																																											
	250	260	270	280	290																																																							
Sequence	G	K	L	G	R	G	K	V	L	V	L	R	K	G	D	I	E	I	P	S	D	L	A	G	V	L	Y	T	E	L	D	E	H	G	G	W	K	R	K	L	L	S	E	L	A	Y	A	G	V	P	F	D	K	D	K	A	L	S	A	
Secondary structure	[Alpha Helix]	[Alpha Helix]										[Alpha Helix]										[Alpha Helix]																																						
SS confidence	[Confidence Scale]																																																											
Disorder	[Disorder Prediction]																																																											
Disorder confidence	[Disorder Confidence Scale]																																																											

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand