

Email	l.a.kelley@imperial.ac.uk
Description	P07026
Date	Thu Jan 5 11:00:07 GMT 2012
Unique Job ID	ce222fa25bfdf73e

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

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

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


  

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


  


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

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand