

Email	l.a.kelley@imperial.ac.uk
Description	P0ACR4
Date	Thu Jan 5 11:18:48 GMT 2012
Unique Job ID	81abf3aefe60a37d

	1	10	20	30	40	50	60																																																					
Sequence	M	H	I	T	L	R	Q	L	E	V	F	A	E	V	L	K	S	G	S	T	T	Q	A	S	V	M	L	A	L	S	Q	S	A	V	S	A	A	L	T	D	L	E	G	Q	L	G	V	Q	L	F	D	R	V	G	K	R	L	V	V	N
Secondary structure	[Alpha Helix]										[Alpha Helix]										[Alpha Helix]										[Coiled-coil]										[Coiled-coil]																			
SS confidence	[Confidence bars]																																																											
Disorder	[Disorder prediction]																																																											
Disorder confidence	[Disorder confidence bars]																																																											

  

	70	80	90	100	110	120																																																						
Sequence	E	H	G	R	L	L	P	R	A	L	A	L	L	E	Q	A	V	E	I	E	Q	L	F	R	E	D	N	G	A	I	R	I	Y	A	S	T	I	G	N	Y	I	L	P	A	V	I	A	R	Y	R	H	D	P	Q	L	P	I			
Secondary structure	[Alpha Helix]																				[Coiled-coil]										[Alpha Helix]										[Coiled-coil]																			
SS confidence	[Confidence bars]																																																											
Disorder	[Disorder prediction]																																																											
Disorder confidence	[Disorder confidence bars]																																																											

  

	130	140	150	160	170	180																																																						
Sequence	E	L	S	V	G	N	S	Q	D	V	M	Q	A	V	L	D	F	R	V	D	I	G	F	I	E	G	P	C	H	S	T	E	I	I	S	E	P	W	L	E	D	E	L	V	V	F	A	A	P	T	S	P	L	A	R	G	P	V	T	L
Secondary structure	[Coiled-coil]										[Alpha Helix]										[Coiled-coil]										[Alpha Helix]										[Alpha Helix]																			
SS confidence	[Confidence bars]																																																											
Disorder	[Disorder prediction]																																																											
Disorder confidence	[Disorder confidence bars]																																																											

  

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

  

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

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand