

Email	l.a.kelley@imperial.ac.uk
Description	P0AGF6
Date	Thu Jan 5 11:28:58 GMT 2012
Unique Job ID	b3d53038500c20f7

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

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

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

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

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

	310	320																											
Sequence	I	Q	N	R	K	T	V	S	I	I	S	G	G	N	I	D	L	S	R	V	S	Q	I	T	G	F	V	D	A
Secondary structure	[Loop]										[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