

Email	l.a.kelley@imperial.ac.uk
Description	P77307
Date	Thu Jan 5 12:27:34 GMT 2012
Unique Job ID	806ffb3b4e4c7347

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


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

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


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


	250																		
Sequence	Y	R	K	F	Y	N	S	R	H	Q	L	V	V	T	Q	L	K	K	K
Secondary structure	[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