

Email	l.a.kelley@imperial.ac.uk
Description	P39361
Date	Thu Jan 5 11:59:52 GMT 2012
Unique Job ID	1a4433c0867070c8

	1	10	20	30	40	50	60																																																					
Sequence	M	S	Q	R	P	D	R	I	K	Q	M	L	H	Y	L	W	Q	H	R	H	L	S	T	Q	Q	A	M	E	L	F	G	Y	A	E	A	T	V	R	R	D	F	Q	Y	I	V	N	Q	Y	P	G	M	I	R	G	H	G	C	L	D	
Secondary structure																																																												
SS confidence																																																												
Disorder																																																												
Disorder confidence																																																												


	70	80	90	100	110	120																																																						
Sequence	F	D	D	S	T	D	D	K	E	Y	V	F	D	V	K	R	T	L	Q	S	V	A	K	R	E	I	A	A	L	A	R	T	M	I	K	D	G	D	C	F	F	L	D	S	G	S	T	C	L	E	L	A	K	C	L	A	D	A	R	V
Secondary structure																																																												
SS confidence																																																												
Disorder																																																												
Disorder confidence																																																												

	130	140	150	160	170	180																																																						
Sequence	K	V	I	C	N	D	I	K	I	A	N	E	L	G	C	F	P	H	V	E	S	Y	I	I	G	G	L	I	R	P	G	Y	F	S	V	G	E	S	L	A	E	M	I	N	A	F	S	V	E	R	A	F	I	S	C	D	A	L	S	
Secondary structure																																																												
SS confidence																																																												
Disorder																																																												
Disorder confidence																																																												


	190	200	210	220	230	240																																																						
Sequence	L	E	T	G	I	T	N	A	T	M	F	E	V	G	V	K	T	R	I	I	Q	R	S	R	E	V	I	L	M	A	D	H	S	K	F	D	A	V	E	P	H	A	V	A	T	L	S	C	I	K	T	I	I	S	D	S	G	L	P	E
Secondary structure																																																												
SS confidence																																																												
Disorder																																																												
Disorder confidence																																																												


	250	260																		
Sequence	T	I	A	Q	R	Y	Q	R	A	G	C	Q	L	F	L	P	H	S	I	K
Secondary structure																				
SS confidence																				
Disorder																				
Disorder confidence																				

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand