

Email	l.a.kelley@imperial.ac.uk
Description	P28721
Date	Thu Jan 5 11:45:12 GMT 2012
Unique Job ID	d2da4da8fea7da08

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


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

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


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


	250													
Sequence	I	K	L	D	G	N	V	T	I	S	V	Q	Y	L
Secondary structure														
SS confidence														
Disorder														
Disorder confidence														

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand