



Email	l.a.kelley@imperial.ac.uk
Description	P75826
Date	Thu Jan 5 12:14:42 GMT 2012
Unique Job ID	c3d70911400a2792


	1	10	20	30	40	50	60																																																					
Sequence	M	F	S	G	L	L	I	I	L	V	P	L	I	V	G	Y	L	I	P	L	R	Q	Q	A	A	L	K	V	I	N	Q	L	L	S	W	M	V	L	I	L	F	F	M	G	I	S	L	A	F	L	D	N	L	A	S	N	L	L	A	
Secondary structure																																																												
SS confidence																																																												
Disorder																																																												
Disorder confidence																																																												
	70	80	90	100	110	120																																																						
Sequence	I	L	H	Y	S	A	V	S	I	T	V	I	L	L	C	N	I	A	A	L	M	W	L	E	R	G	L	P	W	R	N	H	H	Q	Q	E	K	L	P	S	R	I	A	M	A	L	E	S	L	K	L	C	G	V	V	V	I	G	F	A
Secondary structure																																																												
SS confidence																																																												
Disorder																																																												
Disorder confidence																																																												
	130	140	150	160	170	180																																																						
Sequence	I	G	L	S	G	L	A	F	L	Q	H	A	T	E	A	S	E	Y	T	L	I	L	L	F	L	V	G	I	Q	L	R	N	N	G	M	T	L	K	Q	I	V	L	N	R	R	G	M	I	V	A	V	V	V	V	V	S	L	I		
Secondary structure																																																												
SS confidence																																																												
Disorder																																																												
Disorder confidence																																																												
	190	200	210	220	230	240																																																						
Sequence	G	G	L	I	N	A	F	I	L	D	L	P	I	N	T	A	L	A	M	A	S	G	F	G	W	V	S	L	S	G	I	L	L	T	E	S	F	G	P	V	I	G	S	A	A	F	F	N	D	L	A	R	E	L	I	A	I	M	L	I
Secondary structure																																																												
SS confidence																																																												
Disorder																																																												
Disorder confidence																																																												
	250	260	270	280	290																																																							
Sequence	P	G	L	I	R	R	S	R	S	T	A	L	G	L	C	G	A	T	S	M	D	F	T	L	P	V	L	Q	R	T	G	G	L	D	M	V	P	A	A	I	V	H	G	F	I	L	S	L	L	V	P	I	L	I	A	F	F	S	A	
Secondary structure																																																												
SS confidence																																																												
Disorder																																																												
Disorder confidence																																																												

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand