



Email	l.a.kelley@imperial.ac.uk
Description	P37767
Date	Thu Jan 5 11:57:35 GMT 2012
Unique Job ID	5e9a5d6e870b1ed9


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

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand