

Email	l.a.kelley@imperial.ac.uk
Description	P0ABC3
Date	Thu Jan 5 11:15:09 GMT 2012
Unique Job ID	7caf33bffc53f426

	1	10	20	30	40	50	60																																																					
Sequence	M	R	K	S	V	I	A	I	I	I	V	L	V	V	L	Y	M	S	V	F	V	V	K	E	G	E	R	G	I	T	L	R	F	G	K	V	L	R	D	D	D	N	K	P	L	V	Y	E	P	G	L	H	F	K	I	P	F	I	E	
Secondary structure																																																												
SS confidence																																																												
Disorder																																																												
Disorder confidence																																																												
	70	80	90	100	110	120																																																						
Sequence	T	V	K	M	L	D	A	R	I	Q	T	M	D	N	Q	A	D	R	F	V	T	K	E	K	K	D	L	I	V	D	S	Y	I	K	W	R	I	S	D	F	S	R	Y	L	A	T	G	G	G	D	I	S	Q	A	E	V	L	L	K	
Secondary structure																																																												
SS confidence																																																												
Disorder																																																												
Disorder confidence																																																												
	130	140	150	160	170	180																																																						
Sequence	R	K	F	S	D	R	L	R	S	E	I	G	R	L	D	V	K	D	I	V	T	D	S	R	G	R	L	T	L	E	V	R	D	A	L	N	S	G	S	A	G	T	E	D	E	V	T	T	P	A	A	D	N	A	I	A	E	A	E	
Secondary structure																																																												
SS confidence																																																												
Disorder																																																												
Disorder confidence																																																												
	190	200	210	220	230	240																																																						
Sequence	R	V	T	A	E	T	K	G	K	V	P	V	I	N	P	N	S	M	A	A	L	G	I	E	V	V	D	V	R	I	K	Q	I	N	L	P	T	E	V	S	E	A	I	Y	N	R	M	R	A	E	R	E	A	V	A	R	R	H	R	S
Secondary structure																																																												
SS confidence																																																												
Disorder																																																												
Disorder confidence																																																												
	250	260	270	280	290	300																																																						
Sequence	Q	G	Q	E	E	A	E	K	L	R	A	T	A	D	Y	E	V	T	R	T	L	A	E	A	E	R	Q	G	R	I	M	R	G	E	G	D	A	E	A	A	K	L	F	A	D	A	F	S	K	D	P	D	F	Y	A	F	I	R	S	L
Secondary structure																																																												
SS confidence																																																												
Disorder																																																												
Disorder confidence																																																												
	310	320	330																																																									
Sequence	R	A	Y	E	N	S	F	S	G	N	Q	D	V	M	V	M	S	P	D	S	D	F	F	R	Y	M	K	T	P	T	S	A	T	R																										
Secondary structure																																																												
SS confidence																																																												
Disorder																																																												
Disorder confidence																																																												

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand