

Email	l.a.kelley@imperial.ac.uk
Description	P75715
Date	Thu Jan 5 12:13:24 GMT 2012
Unique Job ID	c334039bb9a7cd3b

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

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

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

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

	250	260	270	280	290	300																																																						
Sequence	A	Y	L	T	M	R	L	E	A	S	A	V	S	G	Q	A	M	V	S	D	N	Q	D	L	G	F	I	V	A	D	Q	N	D	T	P	I	T	P	N	D	L	N	S	V	I	P	F	R	L	D	A	A	A	A	N	V	T	L	R	
Secondary structure																																																												
SS confidence																																																												
Disorder																																																												
Disorder confidence																																																												

	310	320																									
Sequence	A	W	P	I	S	I	T	G	Q	K	P	T	E	G	P	F	S	A	L	G	Y	L	R	V	D	Y	Q
Secondary structure																											
SS confidence																											
Disorder																											
Disorder confidence																											

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand