

Email	l.a.kelley@imperial.ac.uk
Description	P36566
Date	Thu Jan 5 11:53:21 GMT 2012
Unique Job ID	59b625f519ac4f7c

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


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

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


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


	250	260																			
Sequence	Y	I	T	L	G	R	Y	I	H	V	T	A	R	K	P	Q	S	K	D	K	V
Secondary structure																					
SS confidence																					
Disorder																					
Disorder confidence																					

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand