

Email	l.a.kelley@imperial.ac.uk
Description	P52598
Date	Thu Jan 5 12:05:48 GMT 2012
Unique Job ID	84eaf723c5f09b38

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


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

	130	140	150	160	170	180																																																						
Sequence	T	N	D	F	V	I	A	D	Y	L	L	D	N	S	N	C	T	I	I	H	T	G	G	A	V	C	R	E	N	R	S	C	V	G	E	A	A	A	T	M	L	R	S	L	M	I	D	Q	A	F	I	S	A	S	S	W	S	V	R	G
Secondary structure																																																												
SS confidence																																																												
Disorder																																																												
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
	190	200	210	220	230	240																																																						
Sequence	I	S	T	P	A	E	D	K	V	T	V	K	R	A	I	A	S	A	S	R	Q	R	V	L	V	C	D	A	T	K	Y	G	Q	V	A	T	W	L	A	L	P	L	S	E	F	D	Q	I	I	T	D	D	G	L	P	E	S	A	S	R
Secondary structure																																																												
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
	250														
Sequence	A	L	V	K	Q	D	L	S	L	L	V	A	K	N	E
Secondary structure															
SS confidence															
Disorder															
Disorder confidence															

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand