

Email	l.a.kelley@imperial.ac.uk
Description	P0A908
Date	Thu Jan 5 11:09:06 GMT 2012
Unique Job ID	ce480ab48932a7cd

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


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

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


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


	
Sequence	S	T	G	I	T	Y	K	F
Secondary structure								
SS confidence								
Disorder	? ?							
Disorder confidence								

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand