

Email	l.a.kelley@imperial.ac.uk
Description	P0CF88
Date	Thu Jan 5 11:31:46 GMT 2012
Unique Job ID	02075b3a88d9a7ed

	1	10	20	30	40	50	60																																																					
Sequence	M	R	R	T	F	T	A	E	E	K	A	S	V	F	E	L	W	K	N	G	T	G	F	S	E	I	A	N	I	L	G	S	K	P	G	T	I	F	T	M	L	R	D	T	G	G	I	K	P	H	E	R	K	R	A	V	A	H	L	T
Secondary structure											A					A					A																																							
SS confidence											0.80					0.80					0.80																																							
Disorder	?															?																																												
Disorder confidence	0.95										0.95					0.95					0.95																																							

	70	80	90	100	110	120																																																					
Sequence	L	S	E	R	E	E	I	R	A	G	L	S	A	K	M	S	I	R	A	I	A	T	A	L	N	R	S	P	S	T	I	S	R	E	V	Q	R	N	R	G	R	R	Y	K	A	V	D	A	N	N	R	A	N	R	M	A	K	R	P
Secondary structure	A					A					A					A																																											
SS confidence	0.80					0.80					0.80					0.80																																											
Disorder	?					?										?																																											
Disorder confidence	0.95					0.95					0.95					0.95																																											

	130	140	150	160	170	180																																																						
Sequence	K	P	C	L	L	D	Q	N	L	P	L	R	K	L	V	L	E	K	L	E	M	K	W	S	P	E	Q	I	S	G	W	L	R	R	T	K	P	R	Q	K	T	L	R	I	S	P	E	T	I	Y	K	T	L	Y	F	R	S	R	E	A
Secondary structure	A					A					A																																																	
SS confidence	0.80					0.80					0.80																																																	
Disorder											?																																																	
Disorder confidence	0.95					0.95					0.95																																																	

	190	200	210	220	230	240																																																						
Sequence	L	H	H	L	N	I	Q	H	L	R	R	S	H	S	L	R	H	G	R	R	H	T	R	K	G	E	R	G	T	I	N	I	V	N	G	T	P	I	H	E	R	S	R	N	I	D	N	R	R	S	L	G	H	W	E	G	D	L	V	S
Secondary structure	A					A					A																																																	
SS confidence	0.80					0.80					0.80																																																	
Disorder	?					?					?																																																	
Disorder confidence	0.95					0.95					0.95																																																	

	250	260	270	280	290	300																																																					
Sequence	G	T	K	N	S	H	I	A	T	L	V	D	R	K	S	R	Y	T	I	I	V	R	L	R	G	K	D	S	V	S	V	N	Q	A	L	T	D	K	F	L	S	P	S	E	L	R	K	S	L	T	W	D	R	G	M	E	L	A	R
Secondary structure						A					A					A																																											
SS confidence						0.80					0.80					0.80																																											
Disorder																																																											
Disorder confidence																																																											

	310	320	330	340	350	360																																																						
Sequence	H	L	E	F	T	V	S	T	G	V	K	V	Y	F	C	D	P	Q	S	P	W	Q	R	G	T	N	E	N	T	N	G	L	I	R	Q	Y	F	P	K	K	T	C	L	A	Q	Y	T	Q	H	E	L	D	L	V	A	A	Q	L	N	N
Secondary structure	A					A					A					A					A																																							
SS confidence	0.80					0.80					0.80					0.80					0.80																																							
Disorder						?					?																																																	
Disorder confidence						0.95					0.95					0.95																																												

	370	380																					
Sequence	R	P	R	K	T	L	K	F	K	T	P	K	E	I	I	E	R	G	V	A	L	T	D
Secondary structure	A					A																	
SS confidence	0.80					0.80																	
Disorder	?					?																	

Confidence Key
High(9)  Low (0)
? Disordered
 Alpha helix
 Beta strand