

Email	l.a.kelley@imperial.ac.uk
Description	P64606
Date	Thu Jan 5 12:09:53 GMT 2012
Unique Job ID	5c09f02f15470891

	1	10	20	30	40	50	60																																																					
Sequence	M	L	L	N	A	L	A	S	L	G	H	K	G	I	K	T	L	R	T	F	G	R	A	G	L	M	L	F	N	A	L	V	G	K	P	E	F	R	K	H	A	P	L	L	V	R	Q	L	Y	N	V	G	V	L	S	M	L	I	I	V
Secondary structure	[Alpha-helices represented by green cylinders]																																																											
SS confidence	[Confidence scale from 0 to 1]																																																											
Disorder	[Disorder prediction]																																																											
Disorder confidence	[Confidence scale from 0 to 1]																																																											


	70	80	90	100	110	120																																																					
Sequence	V	S	G	V	F	I	G	M	V	L	G	L	Q	G	Y	L	V	L	T	T	Y	S	A	E	T	S	L	G	M	L	V	A	L	S	L	R	E	L	G	P	V	V	A	A	L	L	F	A	G	R	A	G	S	A	L	T	A	E	I
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	130	140	150	160	170	180																																																					
Sequence	G	L	M	R	A	T	E	Q	L	S	S	M	E	M	M	A	V	D	P	L	R	R	V	I	S	P	R	F	W	A	G	V	I	S	L	P	L	L	T	V	I	F	V	A	V	G	I	W	G	S	L	V	G	V	S	W	K	G	I
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	190	200	210	220	230	240																																																						
Sequence	D	S	G	F	F	W	S	A	M	Q	N	A	V	D	W	R	M	D	L	V	N	C	L	I	K	S	V	V	F	A	I	T	V	T	W	I	S	L	F	N	G	Y	D	A	I	P	T	S	A	G	I	S	R	A	T	T	R	T	V	V
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	250	260																		
Sequence	H	S	S	L	A	V	L	G	L	D	F	V	L	T	A	L	M	F	G	N
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Disorder	[Disorder prediction]																			
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Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand