

Email	l.a.kelley@imperial.ac.uk
Description	P0AC26
Date	Thu Jan 5 11:16:59 GMT 2012
Unique Job ID	de90825a16b7392b

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

	70	80	90	100	110	120																																																						
Sequence	M	G	A	T	F	G	I	A	L	T	L	V	I	I	A	G	S	E	L	F	T	G	H	T	M	F	L	T	F	G	V	K	A	G	S	I	S	H	G	Q	M	W	A	I	L	P	Q	T	W	L	G	N	L	V	G	S	V	F	V	A
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Disorder	[Disorder prediction]																																																											
Disorder confidence	[Disorder confidence scale]																																																											

	130	140	150	160	170	180																																																					
Sequence	M	L	Y	S	W	G	G	S	L	L	P	V	D	T	S	I	V	H	S	V	A	L	A	K	T	T	A	P	A	M	V	L	F	F	K	G	A	L	C	N	W	L	V	C	L	A	I	W	M	A	L	R	T	E	G	A	A	K	F
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	190	200	210	220	230	240																																																						
Sequence	I	A	I	W	W	C	L	L	A	F	I	A	S	G	Y	E	H	S	I	A	N	M	T	L	F	A	L	S	W	F	G	N	H	S	E	A	Y	T	L	A	G	I	G	H	N	L	L	W	V	T	L	G	N	T	L	S	G	A	V	F
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	250	260																										
Sequence	M	G	L	G	Y	W	Y	A	T	P	K	A	N	R	P	V	A	D	K	F	N	Q	T	E	T	A	A	G
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Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand