

Email	l.a.kelley@imperial.ac.uk
Description	P42601
Date	Thu Jan 5 12:01:46 GMT 2012
Unique Job ID	3e317dddf4f0c908

	1	10	20	30	40	50	60																																																					
Sequence	M	N	T	V	G	T	P	L	L	W	G	G	F	A	V	V	V	A	I	M	L	A	I	D	L	L	L	Q	G	R	R	G	A	H	A	M	T	M	K	Q	A	A	A	V	S	L	V	W	V	T	L	S	L	L	F	N	A	A	F	W
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	W	Y	L	V	Q	T	E	G	R	A	V	A	D	P	Q	A	L	A	F	L	T	G	Y	L	I	E	K	S	L	A	V	D	N	V	F	V	W	L	M	L	F	S	Y	F	S	V	P	A	A	L	Q	R	R	V	L	V	Y	G	V	L
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
	130	140	150	160	170	180																																																					
Sequence	G	A	I	V	L	R	T	I	M	I	F	T	G	S	W	L	I	S	Q	F	D	W	I	L	Y	I	F	G	A	F	L	F	T	G	V	K	M	A	L	A	H	E	D	E	S	G	I	G	D	K	P	L	V	R	W	L	R	G	H
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	190	200	210	220	230	240																																																						
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	250	260	270	280	290	300																																																						
Sequence	N	L	F	A	I	L	G	L	R	A	M	Y	F	L	L	A	G	V	A	E	R	F	S	M	L	K	Y	G	L	A	V	I	L	V	F	I	G	I	K	M	L	I	V	D	F	Y	H	I	P	I	A	V	S	L	G	V	V	F	G	I
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	310	320																			
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Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand