



Email	l.a.kelley@imperial.ac.uk
Description	P75747
Date	Thu Jan 5 12:13:44 GMT 2012
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
	1	10	20	30	40	50	60																																																					
Sequence	M	P	V	L	Q	W	G	M	L	C	V	L	S	L	L	S	I	G	F	L	A	V	H	L	P	A	A	L	L	G	P	M	I	A	G	I	I	F	S	M	R	G	I	T	L	Q	L	P	R	S	A	F	L	A	A	Q	A	I		
Secondary structure	[Alpha-helices 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	L	G	C	M	I	A	Q	N	L	T	G	S	I	L	T	T	L	A	V	N	W	P	I	V	L	A	I	L	L	V	T	L	S	S	A	I	V	G	V	L	L	V	R	Y	S	S	L	P	G	N	T	G	A	W	G	S	S	P	G	
Secondary structure	[Alpha-helices represented by green cylinders]																																																											
SS confidence	[Confidence scale from 0 to 1]																																																											
Disorder	[Disorder prediction]																																																											
Disorder confidence	[Disorder confidence scale]																																																											
	130	140	150	160	170	180																																																						
Sequence	G	A	A	M	V	A	M	A	Q	D	Y	G	A	D	I	R	L	V	A	F	M	Q	Y	L	R	V	L	F	V	A	G	A	V	L	V	T	R	M	M	L	G	D	N	A	E	A	V	N	Q	H	I	V	W	F	P	P	V	S		
Secondary structure	[Alpha-helices represented by green cylinders]																																																											
SS confidence	[Confidence scale from 0 to 1]																																																											
Disorder	[Disorder prediction]																																																											
Disorder confidence	[Disorder confidence scale]																																																											
	190	200	210	220	230	240																																																						
Sequence	I	N	L	L	T	I	L	L	A	V	V	A	G	T	V	G	C	L	L	R	L	P	S	G	T	M	L	I	P	M	L	A	G	A	V	L	Q	S	G	Q	L	I	T	I	E	L	P	E	V	L	L	A	M	A	Y	M	A	I	G	
Secondary structure	[Alpha-helices represented by green cylinders]																																																											
SS confidence	[Confidence scale from 0 to 1]																																																											
Disorder	[Disorder prediction]																																																											
Disorder confidence	[Disorder confidence scale]																																																											
	250	260	270	280	290	300																																																						
Sequence	W	R	I	G	L	G	F	D	K	Q	I	L	L	R	A	L	R	P	L	P	Q	I	L	L	S	I	F	A	L	L	A	I	C	A	G	M	A	W	G	L	T	R	F	M	H	I	D	F	M	T	A	Y	L	A	T	S	P	G	G	L
Secondary structure	[Alpha-helices represented by green cylinders]																																																											
SS confidence	[Confidence scale from 0 to 1]																																																											
Disorder	[Disorder prediction]																																																											
Disorder confidence	[Disorder confidence scale]																																																											
	310	320	330	340																																																								
Sequence	D	T	V	A	V	I	A	A	G	S	N	A	D	M	A	L	I	M	A	M	Q	T	L	R	L	F	S	I	L	L	T	G	P	A	I	A	R	F	I	S	T	Y	A	P	K	R	S	A												
Secondary structure	[Alpha-helices represented by green cylinders]																																																											
SS confidence	[Confidence scale from 0 to 1]																																																											
Disorder	[Disorder prediction]																																																											
Disorder confidence	[Disorder confidence scale]																																																											

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand