

Email	l.a.kelley@imperial.ac.uk
Description	P76268
Date	Wed Jan 25 15:21:07 GMT 2012
Unique Job ID	084091e64e979788

	1	10	20	30	40	50	60																																																					
Sequence	M	A	N	A	D	L	D	K	Q	P	D	S	V	S	S	V	L	K	V	F	G	I	L	Q	A	L	G	E	E	R	E	I	G	I	T	E	L	S	Q	R	V	M	M	S	K	S	T	V	Y	R	F	L	Q	T	M	K	T	L	G	Y
Secondary structure											[Alpha Helix]										[Alpha Helix]										[Alpha Helix]																													
SS confidence	[Confidence Scale]																																																											
Disorder	[Disorder Prediction]																																																											
Disorder confidence	[Disorder Confidence Scale]																																																											


	70	80	90	100	110	120																																																						
Sequence	V	A	Q	E	G	E	S	E	K	Y	S	L	T	L	K	F	E	L	G	A	R	A	L	Q	N	V	D	L	I	R	S	A	D	I	Q	M	R	E	L	S	R	L	T	K	E	T	I	H	L	G	A	L	D	E	D	S	I	V	Y	
Secondary structure	[Alpha Helix]										[Alpha Helix]										[Alpha Helix]										[Alpha Helix]																													
SS confidence	[Confidence Scale]																																																											
Disorder	[Disorder Prediction]																																																											
Disorder confidence	[Disorder Confidence Scale]																																																											

	130	140	150	160	170	180																																																						
Sequence	I	H	K	I	D	S	M	Y	N	L	R	M	Y	S	R	I	G	R	R	N	P	L	Y	S	T	A	I	G	K	V	L	L	A	W	R	D	R	D	E	V	K	Q	I	L	E	G	V	E	Y	K	R	S	T	E	R	T	I	T	S	T
Secondary structure	[Alpha Helix]										[Alpha Helix]										[Alpha Helix]										[Alpha Helix]																													
SS confidence	[Confidence Scale]																																																											
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
	190	200	210	220	230	240																																																						
Sequence	E	A	L	L	P	V	L	D	Q	V	R	E	Q	G	Y	G	E	D	N	E	E	Q	E	E	G	L	R	C	I	A	V	P	V	F	D	R	F	G	V	V	I	A	G	L	S	I	S	F	P	T	L	R	F	S	E	E	R	L	Q	E
Secondary structure	[Alpha Helix]										[Alpha Helix]										[Alpha Helix]										[Alpha Helix]																													
SS confidence	[Confidence Scale]																																																											
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
	250	260																					
Sequence	Y	V	A	M	L	H	T	A	A	R	K	I	S	A	Q	M	G	Y	H	D	Y	P	F
Secondary structure	[Alpha Helix]																						
SS confidence	[Confidence Scale]																						
Disorder	[Disorder Prediction]																						
Disorder confidence	[Disorder Confidence Scale]																						

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand