

Email	l.a.kelley@imperial.ac.uk
Description	P75895
Date	Thu Jan 5 12:15:46 GMT 2012
Unique Job ID	c61568e18a7a3907

	1	10	20	30	40	50	60																																																				
Sequence	M	K	L	S	L	S	P	P	Y	A	D	A	P	V	V	V	L	I	S	G	L	G	G	S	G	S	Y	W	L	P	Q	L	A	V	L	E	Q	E	Y	Q	V	V	C	Y	D	Q	R	G	T	G	N	N	P	D	T	L	A	E	D
Secondary structure	→			→			α			→																																																	
SS confidence																																																											
Disorder	?	?	?																												?	?																											
Disorder confidence																																																											


	70	80	90	100	110	120																																																					
Sequence	Y	S	I	A	Q	M	A	E	L	H	Q	A	L	V	A	A	G	I	E	H	Y	A	V	V	G	H	A	L	G	A	L	V	G	M	Q	L	A	L	D	Y	P	A	S	V	T	V	L	I	S	V	N	G	W	L	R	I	N	A	H
Secondary structure	α										→										α										→										α																		
SS confidence																																																											
Disorder																															?	?	?																										
Disorder confidence																																																											

	130	140	150	160	170	180																																																				
Sequence	T	R	R	C	F	Q	V	R	E	R	L	L	Y	S	G	G	A	Q	A	W	V	E	A	Q	P	L	F	L	Y	P	A	D	W	M	A	A	R	A	P	R	L	E	A	E	D	A	L	A	H	F	Q	G	K	N	N	L	L	R
Secondary structure	α										α										α										α										α																	
SS confidence																																																										
Disorder																															?	?										?	?	?	?	?												
Disorder confidence																																																										


	190	200	210	220	230	240																																																					
Sequence	R	L	N	A	L	K	R	A	D	F	S	H	H	A	D	R	I	R	C	P	V	Q	I	C	A	S	D	D	L	L	V	P	T	A	C	S	S	E	L	H	A	A	L	P	D	S	Q	K	M	V	M	P	Y	G	G	H	A	C	N
Secondary structure	α					α					→										α										→										α																		
SS confidence																																																											
Disorder																																																											
Disorder confidence																																																											


	250	260																																
Sequence	V	T	D	P	E	T	F	N	A	L	L	N	G	L	A	S	L	L	H	H	R	E	A	A	L									
Secondary structure	α					α																												
SS confidence																																		
Disorder																										?	?	?	?	?	?	?	?	?
Disorder confidence																																		

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand