

Email	l.a.kelley@imperial.ac.uk
Description	P52126
Date	Wed Jan 25 15:20:57 GMT 2012
Unique Job ID	4d25e83e8027c027

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

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

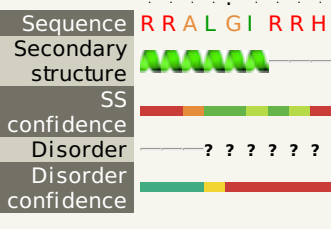
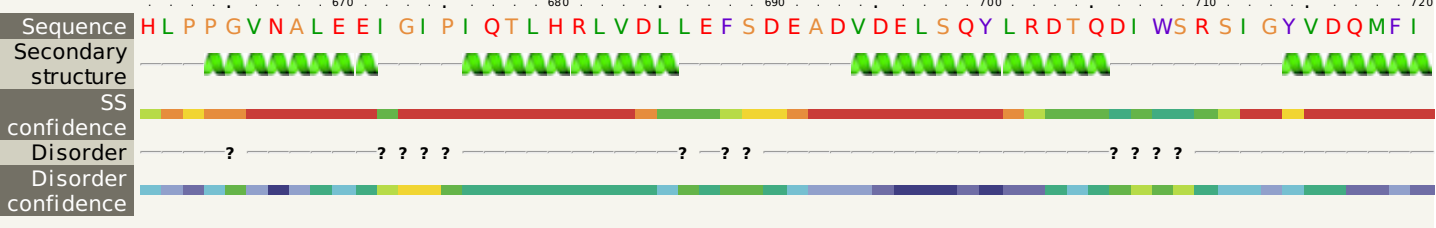
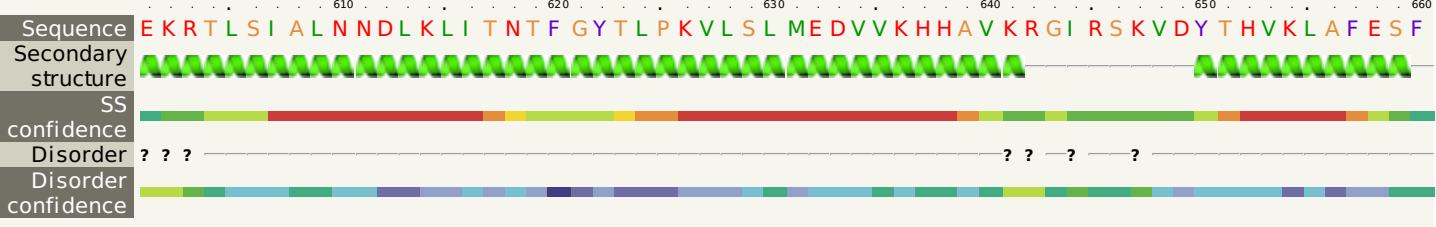
	130	140	150	160	170	180																																																						
Sequence	S	A	P	T	S	M	G	K	S	A	I	V	D	S	L	L	G	M	G	T	L	K	R	L	V	L	V	V	P	T	V	A	L	A	D	E	T	R	R	R	L	Q	E	R	F	G	D	R	Y	Q	I	I	H	H	S	S	Q	V	C	H
Secondary structure	[Alpha Helix]						[Alpha Helix]						[Alpha Helix]						[Alpha Helix]																																									
SS confidence	[Confidence Scale]																																																											
Disorder	[Disorder Prediction]																																																											
Disorder confidence	[Disorder Confidence]																																																											

	190	200	210	220	230	240																																																						
Sequence	S	D	Q	A	V	Y	V	L	T	Q	E	R	V	N	E	R	D	D	I	V	D	I	D	L	F	V	I	D	E	F	Y	K	L	A	F	R	Q	L	K	S	G	D	I	D	H	Q	D	E	R	V	I	E	L	N	I	A	L	S	K	L
Secondary structure	[Alpha Helix]						[Alpha Helix]						[Alpha Helix]						[Alpha Helix]																																									
SS confidence	[Confidence Scale]																																																											
Disorder	[Disorder Prediction]																																																											
Disorder confidence	[Disorder Confidence]																																																											

	250	260	270	280	290	300																																																						
Sequence	L	K	V	S	R	Q	F	Y	L	T	G	P	F	V	N	S	I	R	G	L	E	K	L	G	Y	P	H	T	F	V	S	T	D	F	N	T	V	A	L	D	V	K	T	F	G	I	K	A	N	D	D	K	A	K	L	K	A	L	G	E
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SS confidence	[Confidence Scale]																																																											
Disorder	[Disorder Prediction]																																																											
Disorder confidence	[Disorder Confidence]																																																											

	310	320	330	340	350	360																																																						
Sequence	I	A	H	A	C	V	D	A	T	I	I	Y	C	K	S	P	T	V	A	G	L	V	A	R	E	L	I	R	L	G	H	G	T	P	T	E	N	P	H	V	D	V	V	S	E	E	F	D	A	D	W	D	Y	T	V	A	L	R	N	G
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Disorder	[Disorder Prediction]																																																											
Disorder confidence	[Disorder Confidence]																																																											

	370	380	390	400	410	420																																																						
Sequence	I	G	L	H	F	G	A	L	P	R	A	L	Q	Q	Y	T	A	D	Q	F	N	A	G	K	L	R	F	L	L	C	T	S	T	I	I	E	G	V	N	T	I	A	K	N	V	V	I	Y	D	N	R	D	G	T	R	S	I	D	K	F
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Confidence Key
High(9) Low (0)
? Disordered
Alpha helix
Beta strand