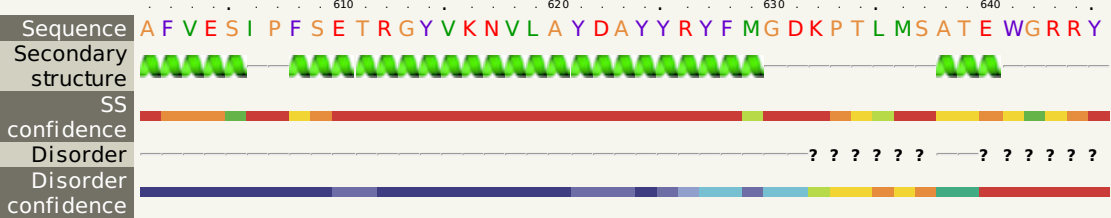
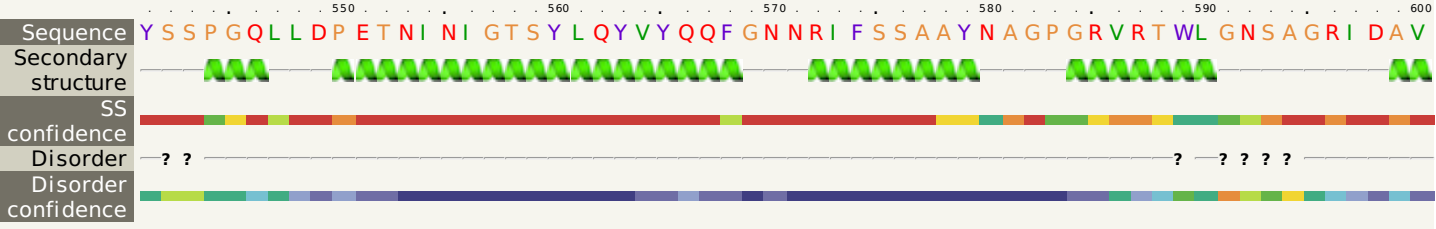
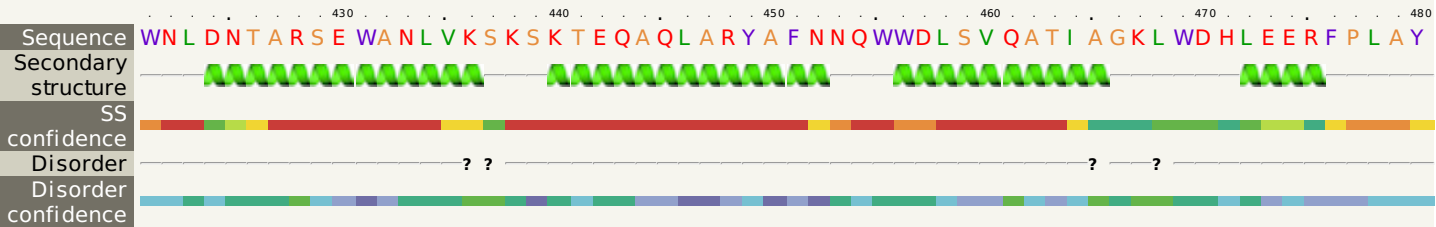


Email	l.a.kelley@imperial.ac.uk
Description	P0AGC3
Date	Thu Jan 5 11:28:44 GMT 2012
Unique Job ID	e4cf65c76a35f8f3

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Sequence	M	E	K	A	K	Q	V	T	W	R	L	L	A	G	V	C	L	L	T	V	S	S	V	A	R	A	D	S	L	D	E	Q	R	S	R	Y	A	Q	I	K	Q	A	W	D	N	R	Q	M	D	V	V	E	Q	M	M	P	G	L	K	
Secondary structure																																																												
SS confidence																																																												
Disorder																																																												
Disorder confidence																																																												
	70	80	90	100	110	120																																																						
Sequence	D	Y	P	L	Y	P	P	L	E	Y	R	Q	I	T	D	D	L	M	N	Q	P	A	V	T	V	T	N	F	V	R	A	N	P	T	L	P	P	A	R	T	L	Q	S	R	F	V	N	E	L	A	R	R	E	D	W	R	G	L	L	A
Secondary structure																																																												
SS confidence																																																												
Disorder																																																												
Disorder confidence																																																												
	130	140	150	160	170	180																																																						
Sequence	F	S	P	E	K	P	G	T	T	E	A	Q	C	N	Y	Y	A	K	W	N	T	G	Q	S	E	E	A	W	Q	G	A	K	E	L	W	L	T	G	K	S	Q	P	N	A	C	D	K	L	F	S	V	W	R	A	S	G	K	Q	D	
Secondary structure																																																												
SS confidence																																																												
Disorder																																																												
Disorder confidence																																																												
	190	200	210	220	230	240																																																						
Sequence	P	L	A	Y	L	E	R	I	R	L	A	M	K	A	G	N	T	G	L	V	T	V	L	A	G	Q	M	P	A	D	Y	Q	T	I	A	S	A	I	I	S	L	A	N	N	P	N	T	V	L	T	F	A	R	T	T	G	A	T	D	F
Secondary structure																																																												
SS confidence																																																												
Disorder																																																												
Disorder confidence																																																												
	250	260	270	280	290	300																																																						
Sequence	T	R	Q	M	A	V	A	F	A	S	V	A	R	Q	D	A	E	N	A	R	L	M	I	P	S	L	A	Q	A	Q	Q	L	N	E	D	Q	I	Q	E	L	R	D	I	V	A	W	R	L	M	G	N	D	V	T	D	E	Q	A	K	
Secondary structure																																																												
SS confidence																																																												
Disorder																																																												
Disorder confidence																																																												
	310	320	330	340	350	360																																																						
Sequence	W	R	D	D	A	I	M	R	S	Q	S	T	S	L	I	E	R	R	V	R	M	A	L	G	T	G	D	R	R	G	L	N	T	W	L	A	R	L	P	M	E	A	K	E	K	D	E	W	R	Y	W	Q	A	D	L	L	E	R	G	
Secondary structure																																																												
SS confidence																																																												
Disorder																																																												
Disorder confidence																																																												
	370	380	390	400	410	420																																																						
Sequence	R	E	A	E	A	K	E	I	L	H	Q	L	M	Q	R	G	F	Y	P	M	V	A	A	Q	R	I	G	E	E	Y	E	L	K	I	D	K	A	P	Q	N	V	D	S	A	L	T	Q	G	P	E	M	A	R	V	R	E	L	M	Y	
Secondary structure																																																												
SS confidence																																																												
Disorder																																																												
Disorder confidence																																																												



Confidence Key
High(9) [High] [High] [High] [High] [High] [High] [High] [High] [High] Low (0)
? Disordered
[Alpha helix] Alpha helix
[Beta strand] Beta strand