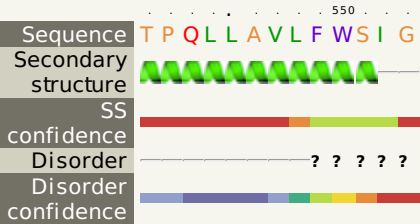
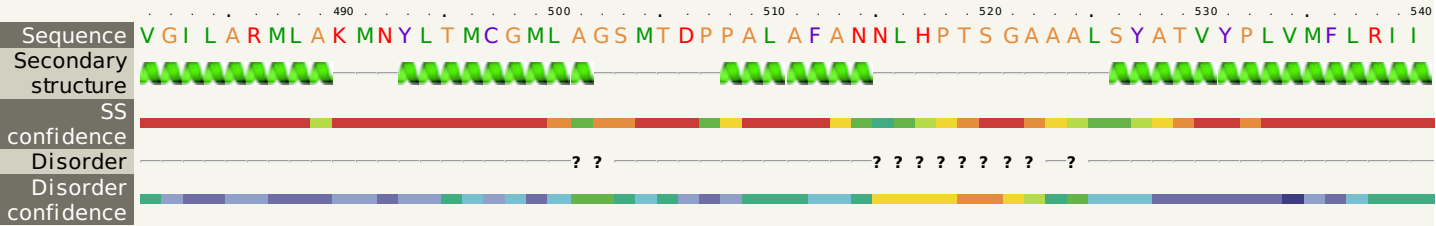
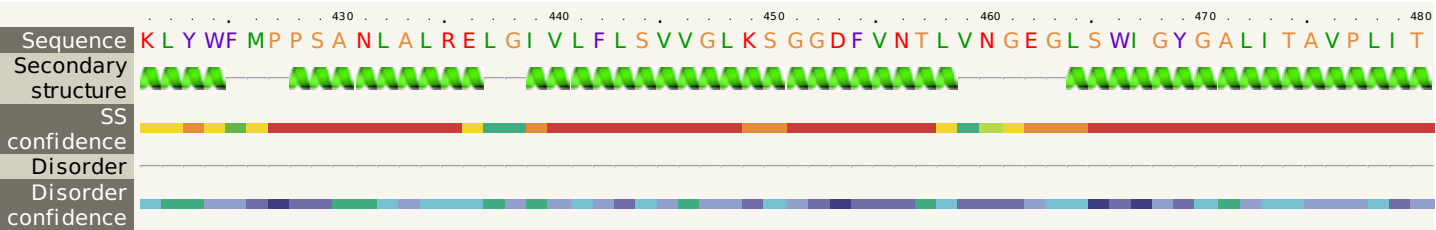


Email	l.a.kelley@imperial.ac.uk
Description	P60872
Date	Thu Jan 5 12:07:13 GMT 2012
Unique Job ID	5843cebe03b081a4

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



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