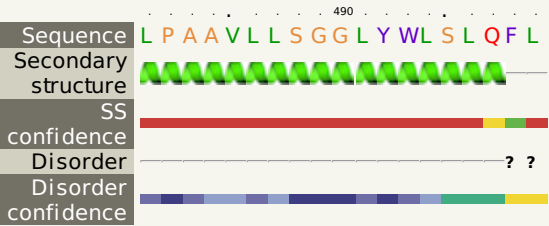
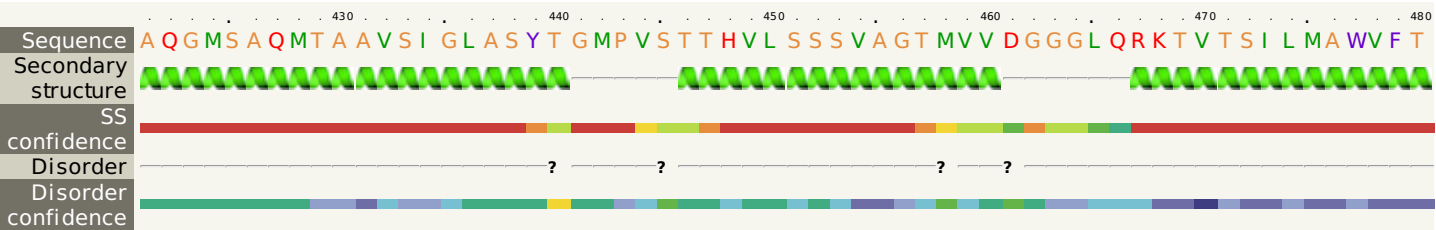


Email	l.a.kelley@imperial.ac.uk
Description	P0AFJ7
Date	Thu Jan 5 11:26:31 GMT 2012
Unique Job ID	a89691be470dab9e

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



Confidence Key

High(9) Low (0)

? Disordered

Alpha helix

Beta strand