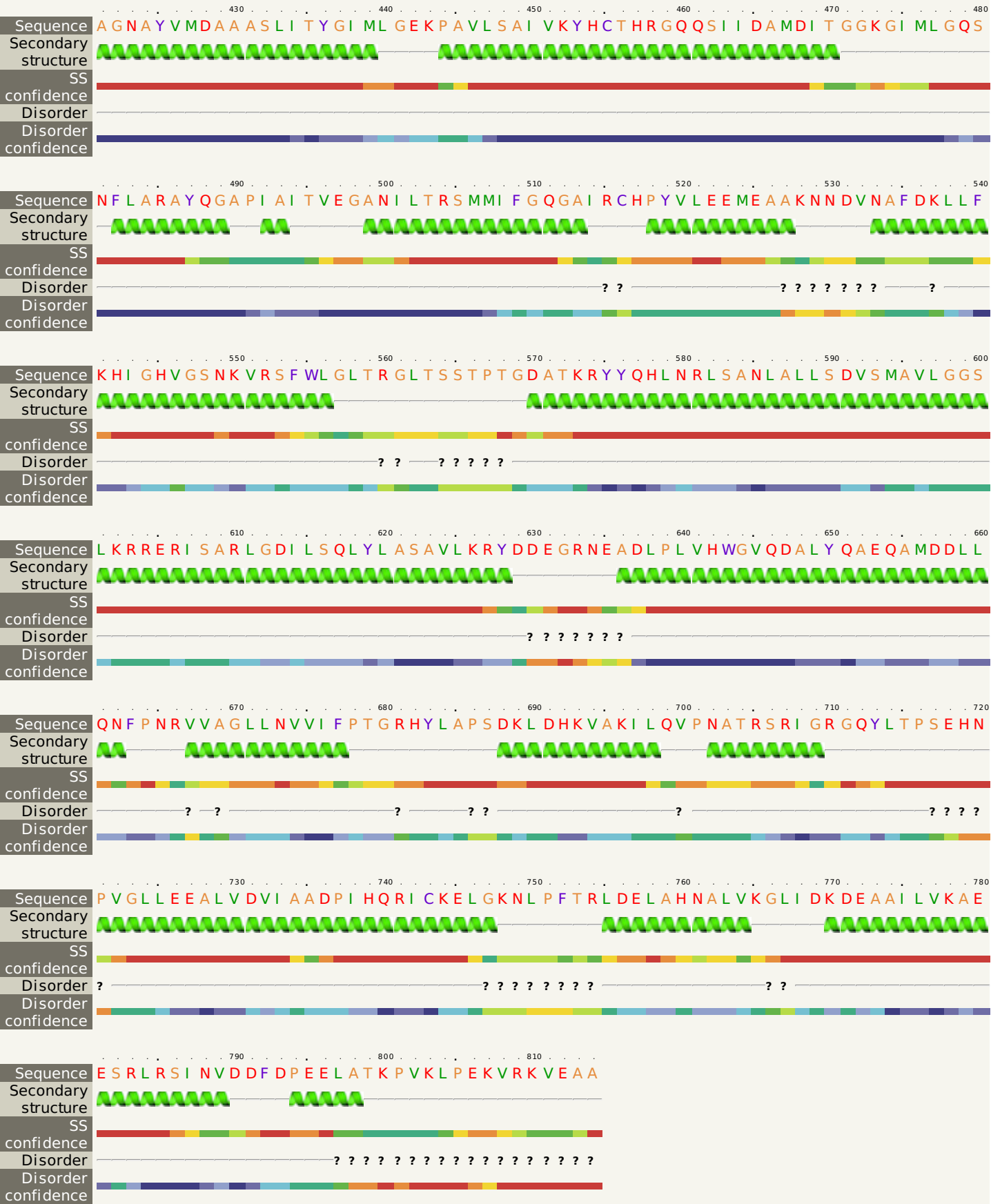


Email	l.a.kelley@imperial.ac.uk
Description	Q47146
Date	Thu Jan 5 12:36:21 GMT 2012
Unique Job ID	dacf1d00198aeb7d

	1	10	20	30	40	50	60																																																					
Sequence	M	M	I	L	S	I	L	A	T	V	V	L	L	G	A	L	F	Y	H	R	V	S	L	F	I	S	S	L	I	L	L	A	W	T	A	A	L	G	V	A	G	L	W	S	A	V	V	L	V	P	L	A	I	I	L	V	P	F	N	F
Secondary structure																																																												
SS confidence																																																												
Disorder																																																												
Disorder confidence																																																												
	70	80	90	100	110	120																																																						
Sequence	A	P	M	R	K	S	M	I	S	A	P	V	F	R	G	F	R	K	V	M	P	P	M	S	R	T	E	K	E	A	I	D	A	G	T	T	W	W	E	G	D	L	F	Q	G	K	P	D	W	K	K	L	H	N	Y	P	Q	P	R	L
Secondary structure																																																												
SS confidence																																																												
Disorder																																																												
Disorder confidence																																																												
	130	140	150	160	170	180																																																						
Sequence	T	A	E	E	Q	A	F	L	D	G	P	V	E	E	A	C	R	M	A	N	D	F	Q	I	T	H	E	L	A	D	L	P	P	E	L	W	A	Y	L	K	E	H	R	F	F	A	M	I	I	K	K	E	Y	G	G	L	E	F	S	A
Secondary structure																																																												
SS confidence																																																												
Disorder																																																												
Disorder confidence																																																												
	190	200	210	220	230	240																																																						
Sequence	Y	A	Q	S	R	V	L	Q	K	L	S	G	V	S	G	I	L	A	I	T	V	G	V	P	N	S	L	G	P	G	E	L	L	Q	H	Y	G	T	D	E	Q	K	D	H	Y	L	P	R	L	A	R	G	Q	E	I	P	C	F	A	L
Secondary structure																																																												
SS confidence																																																												
Disorder																																																												
Disorder confidence																																																												
	250	260	270	280	290	300																																																						
Sequence	T	S	P	E	A	G	S	D	A	G	A	I	P	D	T	G	I	V	C	M	G	E	W	Q	G	Q	Q	V	L	G	M	R	L	T	W	N	K	R	Y	I	T	L	A	P	I	A	T	V	L	G	L	A	F	K	L	S	D	P	E	K
Secondary structure																																																												
SS confidence																																																												
Disorder																																																												
Disorder confidence																																																												
	310	320	330	340	350	360																																																						
Sequence	L	L	G	G	A	E	D	L	G	I	T	C	A	L	I	P	T	T	T	P	G	V	E	I	G	R	R	H	F	P	L	N	V	P	F	Q	N	G	P	T	R	G	K	D	V	F	V	P	I	D	Y	I	I	G	G	P	K	M	A	G
Secondary structure																																																												
SS confidence																																																												
Disorder																																																												
Disorder confidence																																																												
	370	380	390	400	410	420																																																						
Sequence	Q	G	W	R	M	L	V	E	C	L	S	V	G	R	G	I	T	L	P	S	N	S	T	G	G	V	K	S	V	A	L	A	T	G	A	Y	A	H	I	R	R	Q	F	K	I	S	I	G	K	M	E	G	I	E	E	P	L	A	R	I
Secondary structure																																																												
SS confidence																																																												
Disorder																																																												
Disorder confidence																																																												



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