


Email	dan.maclean@tsl.ac.uk
Description	Undefined
Date	Fri Jul 2 15:08:57 BST 2021
Unique Job ID	b42df8c823c60664

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Sequence	M	A	K	S	S	F	K	I	S	N	P	L	E	A	R	M	S	E	S	S	R	I	R	E	K	Y	P	D	R	I	P	V	I	V	E	K	A	G	Q	S	D	V	P	D	I	D	K	K	Y	L	V	P	A	D	L	T	V	G	Q	
Secondary structure											[Alpha Helix]										[Alpha Helix]										[Alpha Helix]																													
SS confidence	[Color scale bar]																																																											
Disorder	[Disorder prediction bar]																																																											
Disorder confidence	[Disorder confidence bar]																																																											


	70	80	90	100	110	120																																																						
Sequence	F	V	Y	V	V	R	K	R	I	K	L	G	A	E	K	A	I	F	V	F	V	K	N	T	L	P	P	T	A	A	L	M	S	A	I	Y	E	E	H	K	D	E	D	G	F	L	Y	M	T	Y	S	G	E	N	T	F	G	S	L	T
Secondary structure	[Alpha Helix]										[Alpha Helix]										[Alpha Helix]										[Alpha Helix]																													
SS confidence	[Color scale bar]																																																											
Disorder	[Disorder prediction bar]																																																											
Disorder confidence	[Disorder confidence bar]																																																											


	1	2
Sequence	V	A
Secondary structure	[Alpha Helix]	
SS confidence	[Color scale bar]	
Disorder	[Disorder prediction bar]	
Disorder confidence	[Disorder confidence bar]	

Confidence Key

High(9)  Low (0)

? Disordered (25%)

 Alpha helix (33%)

 Beta strand (23%)