

Email	l.a.kelley@imperial.ac.uk
Description	P0A759
Date	Thu Jan 5 11:04:54 GMT 2012
Unique Job ID	07b07a623071775b

	1	10	20	30	40	50	60
Sequence	MRLIPLTTAEQVGKWAARHIVNRI	NAFKPTADRP	FVLGLPTGGTP	MTTYKALVEMHKAGQ			
Secondary structure	[Diagram showing alpha helices and beta strands]						
SS confidence	[Color scale bar]						
Disorder	[Disorder prediction bar]						
Disorder confidence	[Disorder confidence bar]						

	70	80	90	100	110	120
Sequence	VSFKHVVTFN	MDEYVGLPKEHPESYYS	FMRNFFDHVDI	PAENINLLNGNAPDI	DAECRQ	
Secondary structure	[Diagram showing alpha helices and beta strands]					
SS confidence	[Color scale bar]					
Disorder	[Disorder prediction bar]					
Disorder confidence	[Disorder confidence bar]					

	130	140	150	160	170	180
Sequence	YEEKIRSYGKIHLFMGGV	GN DGHIAFNEP	ASSLASRTRI	KT LTHDTRVANSRFF	DNDVNQ	
Secondary structure	[Diagram showing alpha helices and beta strands]					
SS confidence	[Color scale bar]					
Disorder	[Disorder prediction bar]					
Disorder confidence	[Disorder confidence bar]					

	190	200	210	220	230	240
Sequence	VPKYALT	VGV GTLLDAEEVMI	LVLGSQKALALQA	AVEGCVNHMWTISCLQLHPKAIMV	CD	
Secondary structure	[Diagram showing alpha helices and beta strands]					
SS confidence	[Color scale bar]					
Disorder	[Disorder prediction bar]					
Disorder confidence	[Disorder confidence bar]					

	250	260
Sequence	EPSTMELKVKTLRYFNELEAENIKGL	
Secondary structure	[Diagram showing alpha helices and beta strands]	
SS confidence	[Color scale bar]	
Disorder	[Disorder prediction bar]	
Disorder confidence	[Disorder confidence bar]	

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand