

Email	l.a.kelley@imperial.ac.uk
Description	P24186
Date	Thu Jan 5 11:41:04 GMT 2012
Unique Job ID	c4ecd5957a54fbe1

Sequence: MAAKI I DGKTI AQQVRSEVAQKVQARI AAGLRAPGLAVVLVGSNPASQIYVASKRKACEE

Secondary structure: [Blue arrow] [Green helix] [Blue arrow] [Green helix]

SS confidence: [Red bar]

Disorder confidence: [Blue bar]

Sequence: VGFVRSYDL PETTSEAELELIDTLNADNTIDGILVQLPLPAGIDNVKVLERIHDPDKDV

Secondary structure: [Blue arrow] [Green helix] [Blue arrow] [Green helix]

SS confidence: [Red bar]

Disorder confidence: [Blue bar]

Sequence: DGFHPYNVGRLCQRAPRLRPCTPRGIVTLLERYNIDTFGLNAVVI GASNI VGRPMSMELL

Secondary structure: [Green helix] [Green helix] [Blue arrow] [Green helix]

SS confidence: [Red bar]

Disorder confidence: [Blue bar]

Sequence: LAGCTTTVTHRF TKNLRHHVENADLLIVAVGKPGFIPGDWIKEGAIVIDVGINRLENGKV

Secondary structure: [Green helix] [Blue arrow] [Green helix] [Blue arrow] [Green helix] [Blue arrow]

SS confidence: [Red bar]

Disorder confidence: [Blue bar]


Sequence: VGDDVVFEDAAKRASYITPVPGGVGPM TVATLI ENT LQACV EYHDPQDE

Secondary structure: [Blue arrow] [Green helix] [Blue arrow] [Green helix]


SS confidence: [Red bar]


Disorder confidence: [Blue bar]

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand