

Email	l.a.kelley@imperial.ac.uk
Description	P0AF10
Date	Thu Jan 5 11:24:51 GMT 2012
Unique Job ID	c4523128c21f6b80

Sequence: MV DQAQDTL RPN NRRLSDMQATMEQTQAFENRVLERLNAGKTVRSFLITAVELLTEAVNLL

Secondary structure: [Ribbon diagram showing alpha-helices and loops]

SS: [Solvent accessibility plot]

confidence: [Red line plot]

Disorder: [Grey line with question marks]

Disorder confidence: [Colored bar plot]

Sequence: VLQVF RKDDY AVKYAVEPLL DGDGPL GDL SVRLKLI YGL GVI NRQEY EDA ELLMAL REEL

Secondary structure: [Ribbon diagram showing alpha-helices and loops]

SS: [Solvent accessibility plot]

confidence: [Red line plot]

Disorder: [Grey line with question marks]

Disorder confidence: [Colored bar plot]

Sequence: NHDGNEY AFT DDEI LGPFGELHCV AALPPP PQF EPADSSLY AMQI QRYQQAVRSTMVLSL

Secondary structure: [Ribbon diagram showing alpha-helices and loops]

SS: [Solvent accessibility plot]

confidence: [Red line plot]

Disorder: [Grey line with question marks]

Disorder confidence: [Colored bar plot]

Sequence: TELISKI SLKKAFQK

Secondary structure: [Ribbon diagram showing alpha-helices and loops]


SS: [Solvent accessibility plot]

confidence: [Red line plot]


Disorder: [Grey line with question marks]


Disorder confidence: [Colored bar plot]

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand