

Email	l.a.kelley@imperial.ac.uk
Description	P21363
Date	Thu Jan 5 11:38:10 GMT 2012
Unique Job ID	38be226bf0cea8cd

Sequence: MNRI EHYHDWLRDAHAMEKQAESML ESMASRI DNYPELRARI EQHLS ETKNQI VQLET I L

Secondary structure: [Alpha-helices and beta-strand representation]

SS confidence: [Confidence scores for secondary structure]

Disorder: [Disorder scores]

Disorder confidence: [Confidence scores for disorder]

Sequence: DRNDI SRSVI KDSMSKMAALGQSIGGI FPSDEI VKGSI SGYVFEQFEI ACYTSL LAAAKN

Secondary structure: [Alpha-helices and beta-strand representation]

SS confidence: [Confidence scores for secondary structure]

Disorder: [Disorder scores]

Disorder confidence: [Confidence scores for disorder]

Sequence: AGDTAS I PTI EAI LNEEKQMA D WLI QNI PQTTEK FLI RSETDGVEAKK


Secondary structure: [Alpha-helices and beta-strand representation]

SS confidence: [Confidence scores for secondary structure]


Disorder: [Disorder scores]


Disorder confidence: [Confidence scores for disorder]

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand