

Email	I.a.kelley@imperial.ac.uk
Description	P75982
Date	Thu Jan 5 12:16:55 GMT 2012
Unique Job ID	824b61ccc411dea3

Sequence: MDVTNDDYIRLLSALLPPGPVWSASDPAIAGAAPSLTRVHQRADALMRELDPRRTTTELIN

Secondary structure: [Alpha-helices and loops]

SS: [Secondary structure representation]

confidence: [Confidence bar]

Disorder: [Disorder prediction bar]

Disorder confidence: [Disorder confidence bar]

Sequence: RWERLCGLPDECIPAGTQTLRQRQQLDAKVNLAGGINEDFYLAQLAALGRPDATITRYD

Secondary structure: [Alpha-helices and loops]

SS: [Secondary structure representation]

confidence: [Confidence bar]

Disorder: [Disorder prediction bar]

Disorder confidence: [Disorder confidence bar]

Sequence: KSTFTCS SACTDAVN APEWRY YWQVNMPATTNSTWMT CGDPCDSALRI WGD TVVECVLNK

Secondary structure: [Alpha-helices and loops]

SS: [Secondary structure representation]

confidence: [Confidence bar]

Disorder: [Disorder prediction bar]

Disorder confidence: [Disorder confidence bar]

Sequence: LCPSHTYVIFKYPE

Secondary structure: [Alpha-helices and loops]

SS: [Secondary structure representation]

confidence: [Confidence bar]

Disorder: [Disorder prediction bar]

Disorder confidence: [Disorder confidence bar]

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand