

Email	l.a.kelley@imperial.ac.uk
Description	P37329
Date	Thu Jan 5 11:55:15 GMT 2012
Unique Job ID	3bedeef5e41dc2fb

Sequence: MARKWLNLFAGAALSFAVAGNALADEGKITTVFAAASLTNAMAQDIATQFKKEKGV DVVSS F

Secondary structure: [Alpha-helix] [Beta-strand] [Alpha-helix] [Beta-strand]

SS confidence: [Bar chart]

Disorder: [Bar chart]

Disorder confidence: [Bar chart]

Sequence: ASSSTLARQIEAGAPADLFISADQKWM DYAVDKKAI DTATRQTLLGNSLVVVPKASVQK

Secondary structure: [Alpha-helix] [Beta-strand] [Alpha-helix] [Beta-strand]

SS confidence: [Bar chart]

Disorder: [Bar chart]

Disorder confidence: [Bar chart]

Sequence: DFTIDSKTNWTSLLNGGRLAVGDPEHVPAGIYAKEALQKLGAWDTLSPKLAPAEDVVRGAL

Secondary structure: [Alpha-helix] [Beta-strand] [Alpha-helix] [Beta-strand]

SS confidence: [Bar chart]

Disorder: [Bar chart]

Disorder confidence: [Bar chart]

Sequence: ALVERNEAPLGLIVYGSDAVASKGVKVVATFPEDSHKKVEYPVAVVEGHNNATVKAFYDYL

Secondary structure: [Alpha-helix] [Beta-strand] [Alpha-helix] [Beta-strand]

SS confidence: [Bar chart]

Disorder: [Bar chart]

Disorder confidence: [Bar chart]

Sequence: KGPQA AEI FKR YGFTIK


Secondary structure: [Alpha-helix] [Beta-strand]

SS confidence: [Bar chart]


Disorder: [Bar chart]


Disorder confidence: [Bar chart]

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand