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| Email | l.a.kelley@imperial.ac.uk |
| Description | P23841 |
| Date | Thu Jan 5 11:39:47 GMT 2012 |
| Unique Job ID | 1c005f08b152a89f |

Sequence: MERVYRTDLKLLRYFLAVAEELHFGRAAARLNMSQPPLSIHIKELLENQLGTQLFIRHSRS

Secondary structure: [Alpha-helices and beta-strands]

SS confidence: [High confidence (red) for most of the protein]

Disorder: [Disordered regions (purple) and ordered regions (green)]

Disorder confidence: [High confidence (red) for most of the protein]

Sequence: VVLTHAGKILMEESRRLLVNANNVLARI EQIGRGEAGRI ELGVVGTAMVGRMRPVMRRFL

Secondary structure: [Alpha-helices and beta-strands]

SS confidence: [High confidence (red) for most of the protein]

Disorder: [Disordered regions (purple) and ordered regions (green)]

Disorder confidence: [High confidence (red) for most of the protein]

Sequence: RENPNVDVLFREKMPAMQMALLERRELDAGIWRMATEPPTGFTSLRLHESAFVLVAMPEEH

Secondary structure: [Alpha-helices and beta-strands]

SS confidence: [High confidence (red) for most of the protein]

Disorder: [Disordered regions (purple) and ordered regions (green)]

Disorder confidence: [High confidence (red) for most of the protein]

Sequence: HLLSSFSTVPLEALRDEYFVTMPPVYTDWDFLQRCVQQVGFSPVVI REVNEPQTVLAMVSM

Secondary structure: [Alpha-helices and beta-strands]

SS confidence: [High confidence (red) for most of the protein]

Disorder: [Disordered regions (purple) and ordered regions (green)]

Disorder confidence: [High confidence (red) for most of the protein]

Sequence: GIGITLIADSYAQMNWPGVIFRPLKQRI PADLYIVYETQQVTPAMVKLLAALTQ

Secondary structure: [Alpha-helices and beta-strands]

SS confidence: [High confidence (red) for most of the protein]

Disorder: [Disordered regions (purple) and ordered regions (green)]

Disorder confidence: [High confidence (red) for most of the protein]

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand