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Description	Q47269
Date	Thu Jan 5 12:36:41 GMT 2012
Unique Job ID	0909771946e621ab

Sequence: M N L S Q D G I K L H R G N F T A I G R Q I Q P Y L E E G K C F R M V L K P W R E K R S L S Q N A L S H M W Y S E I S E

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

SS confidence: [Confidence bars]

Disorder: [Disorder prediction]

Disorder confidence: [Confidence bars]

Sequence: Y L I S R G K T F A T P A W V K D A L K H T Y L G Y E T K D L V D V V T G D I T T I Q S L R H T S D L D T G E M Y V F L

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

SS confidence: [Confidence bars]

Disorder: [Disorder prediction]

Disorder confidence: [Confidence bars]

Sequence: C K V E A W A M N I G C H L T I P Q S C E F Q L L R D K Q E A

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

SS confidence: [Confidence bars]

Disorder: [Disorder prediction]

Disorder confidence: [Confidence bars]

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand