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Description	P31475
Date	Thu Jan 5 11:48:04 GMT 2012
Unique Job ID	582a8ff9e0e172e5

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

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

SS confidence: [Red bar]

Disorder confidence: [Green bar]

Disorder: [Blue bar]

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

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

SS confidence: [Red bar]

Disorder confidence: [Green bar]

Disorder: [Blue bar]

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

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

SS confidence: [Red bar]

Disorder confidence: [Green bar]

Disorder: [Blue bar]

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


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

SS confidence: [Red bar]


Disorder confidence: [Green bar]


Disorder: [Blue bar]

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand