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Description	P68398
Date	Thu Jan 5 12:11:02 GMT 2012
Unique Job ID	b0836fb5cf88052e

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

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

SS confidence: [Confidence scores]

Disorder: [Disorder predictions]

Disorder confidence: [Disorder confidence scores]

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

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

SS confidence: [Confidence scores]

Disorder: [Disorder predictions]

Disorder confidence: [Disorder confidence scores]

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


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

SS confidence: [Confidence scores]


Disorder: [Disorder predictions]


Disorder confidence: [Disorder confidence scores]

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand