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Description	P25741
Date	Thu Jan 5 11:42:28 GMT 2012
Unique Job ID	5a4a5ea86cfa6c97

1 10 20 30 40 50 60

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

Secondary structure

SS

confidence

Disorder

Disorder confidence

70 80 90 100 110 120

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

Secondary structure

SS

confidence

Disorder

Disorder confidence

130 140 150 160 170 180

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

Secondary structure

SS

confidence

Disorder

Disorder confidence

190 200 210 220 230 240

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

Secondary structure

SS

confidence

Disorder

Disorder confidence

250 260

Sequence L E Q N L L N M A S V K A E R I K E R T Q R K G L

Secondary structure


SS

confidence


Disorder


Disorder confidence

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand