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Description	P0A9L3
Date	Thu Jan 5 11:10:37 GMT 2012
Unique Job ID	87b3f95f1b9a5906

1 10 20 30 40 50 60

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

Secondary structure

SS

confidence

Disorder

Disorder confidence

70 80 90 100 110 120

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

Secondary structure

SS

confidence

Disorder

Disorder confidence

130 140 150 160 170 180

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

Secondary structure

SS

confidence

Disorder

Disorder confidence

190 200

Sequence Y G E R G A G A S I P P F S T L V F E V E L L E I L

Secondary structure


SS

confidence


Disorder


Disorder confidence

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand