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Description	P0AG27
Date	Thu Jan 5 11:27:57 GMT 2012
Unique Job ID	5f665930ab34ae8b

1 10 20 30 40 50 60

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

Secondary structure

SS confidence

Disorder ? ? ? ? ?

Disorder confidence

70 80 90 100 110 120

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

Secondary structure

SS confidence

Disorder ? ? ? ? ?

Disorder confidence

130 140

Sequence Q V F V L K E G V A G W A G E N L P L V R G K

Secondary structure

SS confidence

Disorder ? ? ? ? ? ? ?

Disorder confidence

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand