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Description	P0A8I1
Date	Thu Jan 5 11:07:53 GMT 2012
Unique Job ID	57e552270fdb4cf9

Protein structure analysis of the full-length protein (1-120) and two fragments (1-60 and 70-120). The analysis includes sequence, secondary structure, SS confidence, disorder, and disorder confidence.

Full-length protein (1-120):

- Sequence:** MS GTLLAFDFGT KSI GVA VGQRITGTARPLPAIKAQDGP DWNI IERLLKEWQPEDII VG
- Secondary structure:** [Diagram showing alpha-helices and beta-strands]
- SS confidence:** [Bar chart showing confidence levels]
- Disorder:** [Diagram showing disordered regions]
- Disorder confidence:** [Bar chart showing confidence levels]


Fragment 1 (1-60):

- Sequence:** MS GTLLAFDFGT KSI GVA VGQRITGTARPLPAIKAQDGP DWNI IERLLKEWQPEDII VG
- Secondary structure:** [Diagram showing alpha-helices and beta-strands]
- SS confidence:** [Bar chart showing confidence levels]
- Disorder:** [Diagram showing disordered regions]
- Disorder confidence:** [Bar chart showing confidence levels]


Fragment 2 (70-120):


- Sequence:** LPLNMDGTEQPLTARARKFANRIHGRFGVEVKLHDERLSTVEARSGLFEQGGYRALNKGK
- Secondary structure:** [Diagram showing alpha-helices and beta-strands]
- SS confidence:** [Bar chart showing confidence levels]
- Disorder:** [Diagram showing disordered regions]
- Disorder confidence:** [Bar chart showing confidence levels]

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand