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Description	P32162
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Protein structure and disorder analysis for two protein segments. The top segment (1-60) shows a sequence with several alpha-helices and beta-strands. The bottom segment (70-90) shows a sequence with a beta-strand and a loop. Both segments show high confidence in the secondary structure and low confidence in the disorder analysis.


Segment 1 (1-60):

- Sequence: MKDVVDKCS TKGCAI DI GTVI DNDNCTSKFSRFFATREEAESFMTKLKELAAATSSADEG
- Secondary structure: Alpha-helices (green) and beta-strands (blue) are shown. The structure is highly confident.
- SS confidence: High confidence (mostly red/yellow).
- Disorder: Low confidence (mostly green/yellow).
- Disorder confidence: High confidence (mostly red/yellow).


Segment 2 (70-90):


- Sequence: ASVAYKIKDL EGQVELDAAF TFS CQAEMIIFELSLRLA
- Secondary structure: Beta-strand (blue) and loop (green) are shown. The structure is highly confident.
- SS confidence: High confidence (mostly red/yellow).
- Disorder: Low confidence (mostly green/yellow).
- Disorder confidence: High confidence (mostly red/yellow).

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand