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Description	P46119
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Protein structure and disorder analysis for two protein segments. The top segment (residues 1-60) shows a high-confidence secondary structure (SS) with a continuous alpha-helical region. The bottom segment (residues 70-90) shows a lower-confidence SS with a helical region followed by a disordered region. Disorder analysis is shown as a bar chart below the SS, with red indicating high confidence in disorder and green indicating high confidence in structure.


Segment 1 (Residues 1-60):

- Sequence:** MRAIGKLPKGVLLIEFIGMMLLAVALLSVSDSLPEPF SRPEVQILMI FLGVLLMLPAA
- Secondary structure (SS):** High confidence alpha-helical region (green) from residue 1 to 60.
- Disorder:** High confidence in disorder (red) from residue 1 to 60.


Segment 2 (Residues 70-90):


- Sequence:** VVVILQVAKRLAPQLMNRPPQYSR SEREKDNDANH
- Secondary structure (SS):** High confidence alpha-helical region (green) from residue 70 to 80, followed by a disordered region (red) from residue 81 to 90.
- Disorder:** High confidence in disorder (red) from residue 70 to 80, followed by high confidence in structure (green) from residue 81 to 90.

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand