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Description	P75894
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Figure 1 displays the structural analysis of the protein sequence, showing the sequence, secondary structure, and confidence scores across three segments.

Segment 1 (Residues 1-60):

- Sequence: MNEAVSPGALSTLFTDARTHNGWRETPVSDETLREIYALMKWGPTSANCSPARIVFTRTA
- Secondary structure: Alpha helix (residues 1-15), Alpha helix (residues 25-35), Alpha helix (residues 45-55), Alpha helix (residues 58-60)
- SS confidence: High (red bar)
- Disorder: Low (blue bar)
- Disorder confidence: High (red bar)

Segment 2 (Residues 61-120):

- Sequence: EGKERLRPALSSGNLQKTLTAPVTAIVAWDSEFYERLPLLFPHGDARSWFTSSPQLAEET
- Secondary structure: Alpha helix (residues 61-75), Alpha helix (residues 80-90), Alpha helix (residues 95-105), Alpha helix (residues 110-120)
- SS confidence: Medium (yellow bar)
- Disorder: Low (blue bar)
- Disorder confidence: High (red bar)


Segment 3 (Residues 121-180):

- Sequence: AFRNSSMQAAYLIVACRALGLDTGPMMSGFDRQHVDDAFFTGSTLKSNNLLINIGYGDSSKL
- Secondary structure: Alpha helix (residues 121-135), Alpha helix (residues 140-150), Alpha helix (residues 155-165), Alpha helix (residues 170-180)
- SS confidence: Low (green bar)
- Disorder: Low (blue bar)
- Disorder confidence: High (red bar)


Segment 4 (Residues 181-240):


- Sequence: YARLPRLSFEEACGLL
- Secondary structure: Alpha helix (residues 181-195), Alpha helix (residues 200-210)
- SS confidence: Low (green bar)
- Disorder: Low (blue bar)
- Disorder confidence: High (red bar)

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand