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Description	P0A7A9
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Protein structure prediction results for three protein segments. Each segment shows the amino acid sequence, secondary structure (alpha helices and beta strands), sequence confidence, disorder, and disorder confidence.

**Segment 1 (Residues 1-60):**

- Sequence: MSLLNVPAGKDLPEDIYVVI EIPANADPI KYEIDKESGALFVDRFMSTAMFYPCNYGYI N
- Secondary structure: Three alpha helices (blue arrows) at residues 10-25, 30-45, and 45-60.
- SS confidence: High confidence (red) across the segment.
- Disorder: Low confidence (blue) across the segment.
- Disorder confidence: High confidence (red) across the segment.


**Segment 2 (Residues 70-120):**

- Sequence: HTLSLDGDPVDVLVPTPYPLQPGSVIRCRPVGVLLKMTDEAGEDAKLVAVPHSKLSKEYDH
- Secondary structure: Three alpha helices (blue arrows) at residues 75-90, 95-110, and 115-120.
- SS confidence: High confidence (red) across the segment.
- Disorder: Low confidence (blue) across the segment.
- Disorder confidence: High confidence (red) across the segment.


**Segment 3 (Residues 130-170):**


- Sequence: IKDVNDLPELLKAQIAHF FEHYKDLEKGGKVVKEGWENAEAAKAEIVASF ERAKNK
- Secondary structure: A beta sheet (green zigzag) at residues 130-145, followed by an alpha helix (blue arrow) at residues 150-165, and another beta sheet (green zigzag) at residues 165-170.
- SS confidence: High confidence (red) across the segment.
- Disorder: Low confidence (blue) across the segment.
- Disorder confidence: High confidence (red) across the segment.

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand