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|---------------|-----------------------------|
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| Description   | P26281                      |
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| Unique Job ID | 65164d545bfb16c3            |

Protein structure prediction results for a protein of length 160. The image shows three segments of the protein, each with its sequence, secondary structure, and confidence scores for SS, Disorder, and Disorder confidence.

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

- Sequence: M T V A Y I A I G S N L A S P L E Q V N A A L K A L G D I P E S H I L T V S S F Y R T P P L G P Q D Q P D Y L N A A V A
- Secondary structure: Blue arrows indicate alpha-helices at residues 10-15 and 35-45. Green cylinders indicate beta-strands from residue 18 to 32.
- SS confidence: High confidence (red) for most residues, with some lower confidence (yellow/green) at the boundaries of the helices and between helices.
- Disorder: Low disorder (dark blue) across the entire segment.
- Disorder confidence: High confidence (dark blue) across the entire segment.


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

- Sequence: L E T S L A P E E L L N H T Q R I E L Q Q G R V R K A E R W G P R T L D L D I M L F G N E V I N T E R L T V P H Y D M K
- Secondary structure: Blue arrows indicate alpha-helices at residues 75-80, 105-110, 115-120, and 125-130. Green cylinders indicate beta-strands from residue 78 to 92.
- SS confidence: High confidence (red) for most residues, with some lower confidence (yellow/green) at the boundaries of the helices and between helices.
- Disorder: Low disorder (dark blue) across the entire segment.
- Disorder confidence: High confidence (dark blue) across the entire segment.


**Segment 3 (Residues 130-160):**


- Sequence: N R G F M L W P L F E I A P E L V F P D G E M L R Q I L H T R A F D K L N K W
- Secondary structure: Green cylinders indicate beta-strands from residue 132 to 142 and 152 to 160. Blue arrows indicate alpha-helices at residues 145-150 and 155-160.
- SS confidence: High confidence (red) for most residues, with some lower confidence (yellow/green) at the boundaries of the helices and between helices.
- Disorder: Low disorder (dark blue) across the entire segment.
- Disorder confidence: High confidence (dark blue) across the entire segment.

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand