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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: MS RPT I I I NDLDAERI D I L L E Q P A Y A G L P I A D A L N A E L D R A Q M C S P E E M P H D V V T M N S R V
- Secondary structure: Multiple alpha helices and beta strands.
- SS confidence: High confidence across the segment.
- Disorder: Low disorder (mostly red).
- Disorder confidence: High confidence across the segment.


Segment 2 (Residues 70-120):

- Sequence: K F R N L S D G E V R V R T L V Y P A K M T D S N T Q L S V M A P V G A A L L G L R V G D S I H W E L P G G V A T H L E
- Secondary structure: Several alpha helices and beta strands.
- SS confidence: High confidence across the segment.
- Disorder: Low disorder (mostly red).
- Disorder confidence: High confidence across the segment.


Segment 3 (Residues 130-140):


- Sequence: V L E L E Y Q P E A A G D Y L L
- Secondary structure: A single beta strand.
- SS confidence: High confidence across the segment.
- Disorder: Low disorder (mostly red).
- Disorder confidence: High confidence across the segment.

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand