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Protein structure prediction results for three different protein segments. Each segment is analyzed for its sequence, secondary structure, SS confidence, disorder, and disorder confidence.

Segment 1 (Residues 1-60):

- Sequence: M S A E F M V I C K K I L F R N C V I V S L F V F T Y N T W A Q C N N N I K I M R K Y E S E G K Y T V R N L V K N K A I
- Secondary structure: Alpha helix (green) and Beta strand (blue) regions.
- SS confidence: High confidence (red) for the alpha helix and beta strand regions.
- Disorder: Low confidence (green) for the alpha helix and beta strand regions.
- Disorder confidence: High confidence (red) for the alpha helix and beta strand regions.


Segment 2 (Residues 70-120):

- Sequence: A L E L A E I Y V K N R Y G Q D A A E E E K P Y E I T E L T T S V W V E G T I H S D Q I A G G V F I I E I G K N D G R I
- Secondary structure: Alpha helix (green) and Beta strand (blue) regions.
- SS confidence: High confidence (red) for the alpha helix and beta strand regions.
- Disorder: Low confidence (green) for the alpha helix and beta strand regions.
- Disorder confidence: High confidence (red) for the alpha helix and beta strand regions.


Segment 3 (Residues 1-5):


- Sequence: L N F G H G K
- Secondary structure: Alpha helix (green) and Beta strand (blue) regions.
- SS confidence: High confidence (red) for the alpha helix and beta strand regions.
- Disorder: Low confidence (green) for the alpha helix and beta strand regions.
- Disorder confidence: High confidence (red) for the alpha helix and beta strand regions.

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand