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Protein structure and disorder analysis for two protein segments. The top segment (residues 1-60) shows a highly disordered protein with a complex secondary structure. The bottom segment (residues 70-120) also shows a disordered protein with a complex secondary structure. The analysis includes sequence, secondary structure, SS confidence, Disorder, and Disorder confidence for each segment.

Segment 1 (Residues 1-60):

- Sequence:** MNAVLSVQGASAPVKKKSFFSKFTRLNMLRLARAVIPAAVLMFMFPQLAMAAGSSGQDLM
- Secondary structure:** [Complex structure with many loops and turns]
- SS confidence:** [High confidence across most residues]
- Disorder:** [Highly disordered, indicated by many question marks]
- Disorder confidence:** [High confidence in disorder prediction]

Segment 2 (Residues 70-120):

- Sequence:** ASGNTTVKATFGKDSVVVKVVVLAEVLVGAVMYYMMTKNVKFLAGFAIISVFI AVGMVVVG
- Secondary structure:** [Complex structure with many loops and turns]
- SS confidence:** [High confidence across most residues]
- Disorder:** [Highly disordered, indicated by many question marks]
- Disorder confidence:** [High confidence in disorder prediction]

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand