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Protein structure and disorder analysis for two proteins. The top protein (60 residues) has a complex secondary structure with multiple alpha-helices and loops, and a high disorder confidence score. The bottom protein (11 residues) has a simple secondary structure with one alpha-helix and one loop, and a low disorder confidence score.

Protein 1 (Top):

- Sequence: MKP L S S A I A A A L I L F S A Q G V A E Q T T Q P V V T S C A N V V V V P P S Q E H P P F D L N H M G T G S D K S D
- Secondary structure: Multiple alpha-helices (green) and loops (blue).
- SS confidence: High confidence (red/yellow/green).
- Disorder: High disorder confidence (red/yellow/green).
- Disorder confidence: High disorder confidence (red/yellow/green).


Protein 2 (Bottom):


- Sequence: A L G V P Y Y N Q H A M
- Secondary structure: One alpha-helix (green) and one loop (blue).
- SS confidence: High confidence (red/yellow/green).
- Disorder: Low disorder confidence (blue/purple).
- Disorder confidence: Low disorder confidence (blue/purple).

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand