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Protein structure and sequence analysis for two proteins. The top panel shows a full-length protein (60 residues) with a complex fold, including a large alpha-helical bundle and a C-terminal tail. The bottom panel shows a shorter protein (14 residues) with a simple alpha-helical structure. Both panels include sequence, secondary structure, SS confidence, disorder, and disorder confidence tracks.

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand