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Protein structure and analysis of the C-terminal region of the protein. The top panel shows the full-length protein (1-60) with sequence, secondary structure (SS), confidence, disorder, and disorder confidence. The bottom panel shows a zoomed-in view of the C-terminal region (70-110) with sequence, secondary structure (SS), confidence, disorder, and disorder confidence. The C-terminal region is predicted to be a beta-strand (SS) and is highly confident (confidence). It is also predicted to be disordered (disorder) with high confidence (disorder confidence).

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand