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Figure 1 displays the protein structure and disorder analysis of the human protein. The top panel shows the full-length protein (1-60) with its sequence, secondary structure (SS), confidence, and disorder. The bottom panel shows a zoomed-in view of the C-terminal region (70-110) with similar annotations. The protein is composed of several alpha-helices (green) and beta-strands (blue). The disorder analysis indicates regions of high and low confidence, with some regions marked as disordered (red/yellow) and others as ordered (green/blue).

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand