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Figure 1: Structural analysis of the protein sequence. The figure displays four segments of the protein sequence (1-60, 70-120, 130-180, and 190-210) with corresponding secondary structure predictions (alpha-helices and beta-strands) and disorder analysis. The disorder analysis includes a 'Disorder' track with question marks indicating predicted disorder and a 'Disorder confidence' track showing the confidence level. The protein is shown to be highly disordered, with only a few short alpha-helices and beta-strands predicted.

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand