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Figure 1 displays the protein structure and disorder analysis of the human protein. The figure is organized into four horizontal panels, each representing a different segment of the protein sequence (residues 1-60, 70-120, 130-180, and 190-200). Each panel includes the following information:

- Sequence:** The amino acid sequence for the segment, with residues numbered above the letters.
- Secondary structure:** A diagram showing the predicted secondary structure elements (alpha-helices in green, beta-strands in blue) and loops (grey).
- SS confidence:** A bar chart representing the confidence in the secondary structure prediction, with colors corresponding to different confidence levels.
- Disorder:** A bar chart representing the predicted disorder of the protein, with colors corresponding to different disorder levels.
- Disorder confidence:** A bar chart representing the confidence in the disorder prediction, with colors corresponding to different confidence levels.

The protein is shown to be highly disordered, with many regions of low confidence in the disorder analysis. The secondary structure is primarily composed of alpha-helices, with some beta-strands and loops interspersed.

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand