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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 in green, beta-strands in blue), solvent accessibility (SS) scores, disorder predictions, and disorder confidence scores. The sequence is color-coded by amino acid type: M, N, Q, T, L, S, S, F, G, T, P, F, E, R, V, E, N, A, L, A, A, L, R, E, G, R, G, V, M, V, L, D, D, E, D, R, E, N, E, G, D, M, I, F, P, A, E, T, M, T, V, E, Q, M, A, L, T, I, R.

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand