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Protein structure and disorder analysis for the first three domains of the protein. The figure shows sequence alignments, secondary structure predictions (green bars), solvent accessibility (SS) confidence (red/orange/green bars), and disorder predictions (colored bars) for domains 1, 2, and 3. Domain 1 (residues 1-60) is a beta-barrel. Domain 2 (residues 70-120) is an alpha-helix. Domain 3 (residues 130-140) is a short segment.

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand