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| Email | l.a.kelley@imperial.ac.uk |
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Figure 1: Sequence and structural analysis of the protein. The figure displays four segments of the protein sequence (1-60, 70-120, 130-180, and 190-230) with corresponding secondary structure predictions (alpha-helices in green, beta-strands in blue), solvent accessibility (SS) plots, confidence scores, and disorder predictions. The protein is shown to have a complex structure with multiple alpha-helices and beta-strands, and is predicted to be disordered in several regions.

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand