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Protein structure prediction results for the protein sequence: MNNEQRGVALLI VLMLLLALMAALAADMTLS FHSQLQRTRQVNHHLQRQYDIELAEKLLALA S LTQDVKDNDRQTTLQQYWAQPQQLQLEDGNTVKWQLRDAQHCFNLNALAKISDDPLASP DFPAQVFSALLINAGIDRGNTDEIVQSIADYIDVDDSPRFHGAEDSFYQSQTPPRHSAHQ MLFLTGELRQIKGITENIYQRLIPYVCVLP TTELSI NLNMLTENDI PLFRALFLNNITDA DARVLLQKRPREGWLTTDAFLYWAQQDFSGVKPLVAQVKRHLFPYSRYFTLSTESISDEQ SQQWQSHIFFNRKQQSAQIYRRTLQLY

The figure displays six blocks of analysis, each corresponding to a segment of the protein. Each block includes the sequence, secondary structure (SS) prediction, confidence scores, and disorder predictions. The secondary structure is shown as a ribbon diagram with alpha-helices in green and beta-strands in blue. The confidence scores are represented by a color scale from red (low) to green (high). The disorder predictions are shown as a bar chart with red indicating high disorder and green indicating low disorder.

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand