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Protein structure prediction results for the protein sequence MTRIKINARRIFSLLPFFFTSVHAEQTAAPAKPVTVEAKNETFAPQHPDQYLSWKATSEQSERVDALAE D PRLVILWAGYFPSRDYNKPRGHAFVTDVRET LRTGAPKNAEDGPLPMACWSCKSPDVARLIQKDGEDGYFHGKWARGGPEIVNNLGCADCHNTASPEFAKGKPELTL SRPYAARAMEAIGKPF EKAGRF DQQSMVCGQCHVEYYFDGKNKAVKFPWD DGMKVENMEQYYDKIAFSDWTNSLSKTPMLKAQHPEYETWTAGIHGKNNVTCIDCHMPKVQNAEGKLYTD HKI GNPFDNFAQTCANCHTQDKAALQKVVAERKQSI NDLKI KVEDQLVHAHF EAKAALDAGATEAEMKPIQDDIRHAQWRWDLAIA SHGI HMHAP E E GLRMLGTAMDKAADARTKLARLL.

The figure displays the sequence, secondary structure, SS confidence, Disorder, and Disorder confidence for each residue. The secondary structure is shown as a ribbon diagram with alpha-helices in green and beta-strands in blue. The SS confidence is shown as a bar chart with colors representing different confidence levels. Disorder is indicated by question marks, and Disorder confidence is shown as a bar chart.

