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Protein structure prediction results for the protein sequence MT PGGQAQI G NVDLV KQLNS AAVYRLI DQY GPI SRI QI AE QSQLAPASVT KI TRQLI ERG. The figure displays the sequence, secondary structure, SS confidence, disorder, and disorder confidence across the entire length of the protein (1-60).




Sequence: MT PGGQAQI G NVDLV KQLNS AAVYRLI DQY GPI SRI QI AE QSQLAPASVT KI TRQLI ERG

Secondary structure: [Alpha-helices and loops represented by arrows]

SS confidence: [Confidence scores for secondary structure prediction]

Disorder: [Disordered regions indicated by red bars]

Disorder confidence: [Confidence scores for disorder prediction]

Confidence Key
High(9)  Low (0)
? Disordered
 Alpha helix
 Beta strand