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Protein structure prediction results for the protein sequence MRVENNNVSGQNHDPEQIDLLI DLLVQLWRGKMTIIISVI VAI ALAI GYLAVAKEKWTSTA. The figure displays the sequence, secondary structure, SS confidence, disorder, and disorder confidence across the entire length of the protein (1-600).

Sequence: MRVENNNVSGQNHDPEQIDLLI DLLVQLWRGKMTIIISVI VAI ALAI GYLAVAKEKWTSTA

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

SS confidence: [Confidence scores for secondary structure prediction]

Disorder: [Disordered regions indicated by question marks]

Disorder confidence: [Confidence scores for disorder prediction]

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand