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The figure displays two protein analysis panels. The top panel shows a protein of 60 residues. The sequence is M A A N A F V R A R I D E D L K N Q A A D V L A G M G L T I S D L V R I T L T K V A R E K A L P F D L R E P N Q L T I Q. The secondary structure is represented by a blue arrow for the first 10 residues and green helices for the rest. The SS confidence is shown as a red bar with some yellow and green segments. The Disorder is indicated by question marks for the first 10 residues. The Disorder confidence is shown as a blue bar with some green and yellow segments. The bottom panel shows a protein of 15 residues. The sequence is S I K N S E A G I D V H K A K D A D D L F D K L G I. The secondary structure is represented by green helices. The SS confidence is shown as a red bar with some green and yellow segments. The Disorder is indicated by question marks for the first 10 residues. The Disorder confidence is shown as a blue bar with some green and yellow segments.

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand