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Protein structure and disorder analysis for the first protein (1-60). The sequence is MASVSI S C P S C S A T D G V V R N G K S T A G H Q R Y L C S H C R K T W Q L Q F T Y T A S Q P G T H Q K I I D M A. The secondary structure shows several alpha-helices (green) and beta-strands (blue). The SS confidence is high (red/yellow). The disorder confidence is low (blue/purple).

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand