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Protein structure and confidence analysis for two proteins. The top protein is 60 residues long, with a sequence: MSRKMTGIVKTFDGKSGKGLITPSDGRIDVQLHVSALNLRDAEEITGLRVEFCRI NGLR. It shows a secondary structure with four alpha-helices and a disordered region (red) from residue 1 to 10. The bottom protein is 70 residues long, with a sequence: GPSAANVYLS. It shows a secondary structure with one alpha-helix and a disordered region (red) from residue 1 to 10.

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand