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Protein structure and disorder analysis for two proteins. The top protein (60 residues) has a sequence MKII SKRRAMTI YRQHPE SRIFRYCTGKYQWHGSVCHYTGRDVPDIAGVLA VYAEERRQDR. It shows three alpha-helices (blue arrows) and a C-terminal beta-strand region (green zig-zag). The bottom protein (10 residues) has a sequence NGPYTCLMSITLN. It shows one alpha-helix (blue arrow). Both proteins have disorder confidence scores indicated by colored bars and question marks.

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand