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Figure 1 displays protein structure prediction results for three proteins, showing sequence, secondary structure, and confidence scores.

Protein 1 (Top): Sequence: MSKTIATENAPAAIGPYVQGVDLGNMIITSGQIPVNPKTGEVPADVAAQARQSLDNVKAII. Secondary structure: Alpha helix (residues 1-10), Alpha helix (residues 20-25), Alpha helix (residues 28-32), Alpha helix (residues 35-40), Alpha helix (residues 45-55). Confidence: SS (red), Disorder (green), Disorder confidence (blue).

Protein 2 (Middle): Sequence: VEAAGLKVGDIVKTTVFVKDLNDFATVNATYEAFFTEHNAFTPARSCVEVARLPKDVKIE. Secondary structure: Alpha helix (residues 1-10), Alpha helix (residues 20-25), Alpha helix (residues 30-40), Alpha helix (residues 50-60), Alpha helix (residues 70-80), Alpha helix (residues 90-100), Alpha helix (residues 110-120). Confidence: SS (red), Disorder (green), Disorder confidence (blue).


Protein 3 (Bottom): Sequence: IEAIAVRR. Secondary structure: Alpha helix (residues 1-5). Confidence: SS (red), Disorder (green), Disorder confidence (blue).

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand