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The figure displays two protein structure models and their corresponding analysis tracks. The top model represents the full-length protein (1-60), showing a green alpha-helical domain (residues 10-20), a blue beta-strand domain (residues 30-40), and a green alpha-helical domain (residues 50-60). The bottom model represents a protein fragment (70-100), showing a green alpha-helical domain (residues 70-80), a blue beta-strand domain (residues 90-100), and a green alpha-helical domain (residues 110-120). The analysis tracks below each model include: Sequence (residue numbers 1-60 and 70-100), Secondary structure (SS) (green for alpha-helix, blue for beta-strand), confidence (red for high, yellow for medium, green for low), Disorder (red for disordered, green for ordered), and Disorder confidence (red for high, green for low).

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand