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Protein structure and disorder analysis for two protein segments.

Top Segment (Residues 1-60):

- Sequence:** MLQS RNDHL RQTAL RNAHT PAL LTTLT E P QDRSLAI NNPQLAADVKTA WLKEDPS LLLF
- Secondary structure:** Alpha-helices (green) and a disordered region (blue arrow).
- SS confidence:** Color-coded bars representing confidence levels.
- Disorder:** Predicted disordered regions (dashed lines) and confidence (colored bars).

Bottom Segment (Residues 70-90):

- Sequence:** VEQPDL SLLRDLVKT GATRKI RSEARHRL EEKQ
- Secondary structure:** Alpha-helices (green) and a disordered region (blue arrow).
- SS confidence:** Color-coded bars representing confidence levels.
- Disorder:** Predicted disordered regions (dashed lines) and confidence (colored bars).

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand