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The figure displays three segments of protein analysis. Each segment includes the amino acid sequence, secondary structure (SS), SS confidence, disorder, and disorder confidence.

- Segment 1 (Residues 1-60):** Sequence: MLHTANPVIKHKAGLLNLAEELSNVSKACKIMGVSRDFTFYRYRELVAEGGVDAQINRSRR. SS shows multiple alpha-helices. Disorder is predicted in several regions.
- Segment 2 (Residues 70-120):** Sequence: APNLKNRTDEATEQAVVDYAVAFPTHGQHRASNELRKQGVFISDSGVRSVLWLLHNLNLK. SS shows alpha-helices and a beta-strand (indicated by a blue arrow). Disorder is predicted in several regions.
- Segment 3 (Residues 121-130):** Sequence: RRY. SS shows a short alpha-helix. Disorder is predicted.

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand