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Figure 1 displays the protein structure and disorder analysis of the 60-residue protein. The top panel shows the full sequence (MESNFI DWHPADII AGLRKKGTS MAAESRRNGLSSSTLANALSRPWPKGE MII AKALGTD) with secondary structure elements (alpha-helices and beta-strands) and confidence scores. The bottom panel shows a zoomed-in view of residues 70-90 (P WV I WPS RY HDP QT H E F I DRT Q L MRS Y T K P K K) with similar annotations. Disorder is indicated by red bars and question marks.

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand