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The figure displays the N-terminal region of the protein, with the sequence: MT K K K A H K P G S A T I A L N K R A R H E Y F I E E E F E A G L A L Q G W E V K S L R A G K A N I S D S Y V L L R D. The secondary structure is shown as a blue arrow (alpha-helix) and a green zigzag (beta-strand). The disorder prediction is shown as a red bar (disordered) and a green bar (ordered). The disorder confidence is shown as a bar with red (high) and green (low) segments. The disorder prediction is shown as a bar with red (high) and green (low) segments.

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand