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Figure 1 displays the secondary structure prediction of the protein sequence. The sequence is shown in a color-coded font (M: blue, E: green, R: red, N: blue, K: green, L: red, A: blue, R: red, Q: blue, I: green, I: green, D: red, T: blue, C: green, L: red, E: green, M: blue, T: blue, R: red, L: green, G: blue, L: green, N: blue, Q: blue, G: blue, T: blue, A: blue, G: blue, N: blue, V: blue, S: blue, V: blue, R: blue, Y: blue, Q: blue, D: blue, G: blue, M: blue, L: blue, I: blue, T: blue, P: blue, T: blue, G: blue, I: blue, P: blue, Y: blue, E: blue, K: blue, L: blue, T: blue, E: blue, S: blue, H: blue, I: blue, V: blue, F: blue, I: blue, D: blue, G: blue). The secondary structure is represented by green alpha-helices and blue beta-strands. The SS confidence is shown as a red bar, and the Disorder confidence is shown as a blue bar. The Disorder prediction is marked with question marks.

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand