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Description	C6Y4B8
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The figure displays protein structure and disorder analysis for the full-length protein (1-120) and a fragment (1-4). The top section shows the full-length protein with residue numbers 1, 10, 20, 30, 40, 50, and 60. The bottom section shows the fragment with residue numbers 1, 2, 3, and 4. Each section includes the following tracks:

- Sequence:** Amino acid sequence (e.g., MYNTYAIIVHVS... for the full-length protein).
- Secondary structure:** Alpha-helices (green cylinders) and beta-strands (green arrows).
- SS confidence:** Confidence score for secondary structure (red = high, yellow = medium, green = low).
- Disorder:** Disordered regions (indicated by question marks).
- Disorder confidence:** Confidence score for disorder (red = high, yellow = medium, green = low).

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand