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The figure displays the protein structure and analysis for the full-length protein (1-140) and two fragments (1-60 and 70-120). The protein is composed of three alpha-helices and a C-terminal region. The full-length protein structure is shown in green, with the C-terminal region in blue. The fragments are shown in blue. The analysis tracks include Sequence, Secondary structure, SS confidence, Disorder, and Disorder confidence.

Full-length protein (1-140):

Sequence: M Q Y A K P V T L N V E E C D R L S F L P Y L F G N D F L Y A E A Y V Y A L A Q K M M P E Y Q G G F W H F I R L P D G G

Secondary structure: [Alpha-helices 1-3, C-terminal region]

SS confidence: [Confidence scores for secondary structure]

Disorder: [Disorder prediction]

Disorder confidence: [Disorder confidence scores]

Fragment 1 (1-60):

Sequence: M Q Y A K P V T L N V E E C D R L S F L P Y L F G N D F L Y A E A Y V Y A L A Q K M M P E Y Q G G F W H F I R L P D G G

Secondary structure: [Alpha-helices 1-3]

SS confidence: [Confidence scores for secondary structure]

Disorder: [Disorder prediction]

Disorder confidence: [Disorder confidence scores]

Fragment 2 (70-120):

Sequence: G Y M M P D G D R F H L V N G E N W F D R T V S A D A A G I I L T S L V I N R Q L W L Y H D S G D V G L T Q L Y R M R D


Secondary structure: [Alpha-helices 4-5]

SS confidence: [Confidence scores for secondary structure]


Disorder: [Disorder prediction]


Disorder confidence: [Disorder confidence scores]

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand