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The figure displays the protein structure of the full-length protein and three fragments. The full-length protein is shown at the top, with residues 1 to 220. The fragments are shown below, with residues 1 to 60, 70 to 120, and 130 to 180. Each fragment is represented by a sequence, secondary structure (SS), confidence, disorder, and disorder confidence.

Full-length protein (1-220):

Sequence: MLQVYLVRHGETQWNAERRIQGQSDSPLTAKGEQQAMQVATRAKELGITHTIISSDLGRTR

Secondary structure: [Diagram showing alpha-helices and beta-strands]

SS: [Diagram showing secondary structure elements]

confidence: [Diagram showing confidence scores]

Disorder: [Diagram showing disorder scores]

Disorder confidence: [Diagram showing disorder confidence scores]

Fragment 1 (1-60):

Sequence: RTAEIIAQACGCDIIFDSRLRELNMGVLEKRHIDSLTEEEENWRRQLVNGTVDGRIPEGE

Secondary structure: [Diagram showing alpha-helices and beta-strands]

SS: [Diagram showing secondary structure elements]

confidence: [Diagram showing confidence scores]

Disorder: [Diagram showing disorder scores]

Disorder confidence: [Diagram showing disorder confidence scores]

Fragment 2 (70-120):

Sequence: SMQELSDRVNAALES CRDLPQGS RPLLVS HGI ALGCLVSTILGLPAWAERRRLRLNC S I S

Secondary structure: [Diagram showing alpha-helices and beta-strands]

SS: [Diagram showing secondary structure elements]

confidence: [Diagram showing confidence scores]

Disorder: [Diagram showing disorder scores]

Disorder confidence: [Diagram showing disorder confidence scores]

Fragment 3 (130-180):

Sequence: RVDYQESLWLASGWV VETAGDI SHLDAPALDELQR

Secondary structure: [Diagram showing alpha-helices and beta-strands]


SS: [Diagram showing secondary structure elements]

confidence: [Diagram showing confidence scores]


Disorder: [Diagram showing disorder scores]


Disorder confidence: [Diagram showing disorder confidence scores]

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand