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Protein structure analysis of the full-length protein (1-300) and its fragments. The analysis includes sequence, secondary structure, SS confidence, disorder, and disorder confidence for each segment.

Full-length protein (1-300):

- Sequence: MDALSRLLMLNAPQGTIDKNCVLGSDWQLPHGAGELSVIRWHALTQGAAKLEMP TGEI FT
- Secondary structure: [Alpha-helices and beta-strands]
- SS confidence: [Confidence scores]
- Disorder: [Disorder scores]
- Disorder confidence: [Disorder confidence scores]

Fragment 1 (1-60):

- Sequence: LRP GNVVLLP QNSAHRLSHVDNESTCIVCGTLRLQHSARYFLTSLPETLFLAPVNHSEY
- Secondary structure: [Alpha-helices and beta-strands]
- SS confidence: [Confidence scores]
- Disorder: [Disorder scores]
- Disorder confidence: [Disorder confidence scores]

Fragment 2 (61-120):

- Sequence: NWLREAI PFL QQESRSAMP GVDALCSQICATFFTLAVREWI AQVNTEKNI LSLLLHPRLG
- Secondary structure: [Alpha-helices and beta-strands]
- SS confidence: [Confidence scores]
- Disorder: [Disorder scores]
- Disorder confidence: [Disorder confidence scores]

Fragment 3 (121-180):

- Sequence: AVIQQMLEMP GHAWTVESLASIAHMSRASFAQLFRDVS GTTPLAVLTKLRLQIAAQMF SR
- Secondary structure: [Alpha-helices and beta-strands]
- SS confidence: [Confidence scores]
- Disorder: [Disorder scores]
- Disorder confidence: [Disorder confidence scores]

Fragment 4 (181-240):

- Sequence: ETL P VVVI AESVGYASESSFHKAFVREFGCTPGEYRERV RQLAP
- Secondary structure: [Alpha-helices and beta-strands]
- SS confidence: [Confidence scores]
- Disorder: [Disorder scores]
- Disorder confidence: [Disorder confidence scores]

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand