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The figure displays protein structure and disorder analysis for the full-length protein (1-120) and two fragments (1-60 and 61-120). The analysis includes sequence, secondary structure, solvent accessibility (SS), confidence, disorder, and disorder confidence.

Full-length protein (1-120):

- Sequence:** MI L F V G F L L M E I V M P Q I S R T A L V P Y S A E Q M Y Q L V N D V Q S Y P Q Q F L P G C T G S R I L E S T P G Q M
- Secondary structure:** Blue arrows indicate alpha-helices and beta-strands. Green wavy lines indicate loops.
- SS confidence:** Red bar representing solvent accessibility confidence.
- Disorder:** Grey line with question marks indicating regions of predicted disorder.
- Disorder confidence:** Dark blue bar representing disorder confidence.


Fragment 1 (1-60):

- Sequence:** MI L F V G F L L M E I V M P Q I S R T A L V P Y S A E Q M Y Q L V N D V Q S Y P Q Q F L P G C T G S R I L E S T P G Q M
- Secondary structure:** Blue arrows indicate alpha-helices and beta-strands. Green wavy lines indicate loops.
- SS confidence:** Red bar representing solvent accessibility confidence.
- Disorder:** Grey line with question marks indicating regions of predicted disorder.
- Disorder confidence:** Dark blue bar representing disorder confidence.


Fragment 2 (61-120):


- Sequence:** T A A V D V S K A G I S K T F T T R N Q L T S N Q S I L M N L V D G P F K K L I G G W K F T P L S Q E A C R I E F H L D
- Secondary structure:** Blue arrows indicate alpha-helices and beta-strands. Green wavy lines indicate loops.
- SS confidence:** Red bar representing solvent accessibility confidence.
- Disorder:** Grey line with question marks indicating regions of predicted disorder.
- Disorder confidence:** Dark blue bar representing disorder confidence.

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand