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The figure displays the protein structure and analysis for the full-length protein (1-120) and a fragment (130-140). The protein is shown in a ribbon diagram, with the full-length protein (1-120) in blue and the fragment (130-140) in red. The structure is composed of several alpha-helices and beta-strands. The analysis tracks include:


- Sequence:** MVGYHQTNQKTDTGKTLTRRPVLVDHNRLPEGSRGRLAVAVAGDHPAAVQVTMTLVNDTG (1-60) and FDPVFSGSI AESWRQQPCTPSYCCDWEAATMLRAFP LAKKGEGRARLP SLYASF GKLG ET (61-120).
- Secondary structure:** Represented by green cylinders for alpha-helices and blue arrows for beta-strands.
- SS confidence:** A bar chart showing the confidence of the secondary structure prediction, with colors indicating different confidence levels.
- Disorder:** A bar chart showing the predicted disorder of the protein, with colors indicating different confidence levels.
- Disorder confidence:** A bar chart showing the confidence of the disorder prediction, with colors indicating different confidence levels.

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand