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The figure displays protein structure prediction results for the full-length protein (1-260) and three fragments. The results are organized into four panels, each corresponding to a different segment of the protein.

Panel 1 (Residues 1-60):

- Sequence:** M F D I G V N L T S S Q F A K D R D D V V A C A F D A G V N G L L I T G T N L R E S Q Q A Q K L A R Q Y S S C W S T A G
- Secondary structure:** Shows a blue arrow (coil) from residue 1 to 10, green cylinders (helix) from 10 to 20, green cylinders (helix) from 20 to 30, a blue arrow (coil) from 30 to 40, green cylinders (helix) from 40 to 50, and a blue arrow (coil) from 50 to 60.
- SS confidence:** A bar chart showing confidence levels across the segment.
- Disorder:** A bar chart showing disorder levels across the segment.
- Disorder confidence:** A bar chart showing confidence levels for disorder predictions.

Panel 2 (Residues 70-120):

- Sequence:** V H P H D S S Q W Q A A T E E A I I E L A A Q P E V V A I G E C G L D F N R N F S T P E E Q E R A F V A Q L R I A A D L
- Secondary structure:** Shows a blue arrow (coil) from residue 70 to 80, green cylinders (helix) from 80 to 90, a blue arrow (coil) from 90 to 100, and green cylinders (helix) from 100 to 120.
- SS confidence:** A bar chart showing confidence levels across the segment.
- Disorder:** A bar chart showing disorder levels across the segment.
- Disorder confidence:** A bar chart showing confidence levels for disorder predictions.

Panel 3 (Residues 130-180):

- Sequence:** N M P V F M H C R D A H E R F M T L L E P W L D K L P G A V L H C F T G T R E E M Q A C V A H G I Y I G I T G W V C D E
- Secondary structure:** Shows a blue arrow (coil) from residue 130 to 140, green cylinders (helix) from 140 to 150, a blue arrow (coil) from 150 to 160, green cylinders (helix) from 160 to 170, a blue arrow (coil) from 170 to 180, and green cylinders (helix) from 180 to 190.
- SS confidence:** A bar chart showing confidence levels across the segment.
- Disorder:** A bar chart showing disorder levels across the segment.
- Disorder confidence:** A bar chart showing confidence levels for disorder predictions.


Panel 4 (Residues 190-240):

- Sequence:** R R G L E L R E L L P L I P A E K L L I E T D A P Y L L P R D L T P K P S S R R N E P A H L P H I L Q R I A H W R G E D
- Secondary structure:** Shows green cylinders (helix) from residue 190 to 200, a blue arrow (coil) from 200 to 210, and green cylinders (helix) from 210 to 240.
- SS confidence:** A bar chart showing confidence levels across the segment.
- Disorder:** A bar chart showing disorder levels across the segment, with a question mark indicating uncertainty.
- Disorder confidence:** A bar chart showing confidence levels for disorder predictions.


Panel 5 (Residues 250-260):


- Sequence:** A A W L A A T T D A N V K T L F G I A F
- Secondary structure:** Shows green cylinders (helix) from residue 250 to 260.
- SS confidence:** A bar chart showing confidence levels across the segment.
- Disorder:** A bar chart showing disorder levels across the segment, with a question mark indicating uncertainty.
- Disorder confidence:** A bar chart showing confidence levels for disorder predictions.

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand