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Figure 1 displays the protein structure and disorder analysis for the C-terminal region of the protein. The figure is organized into three main sections, each corresponding to a different range of residues.

Section 1 (Residues 1-60):

- Sequence:** M E S P L G S D L A R L V R I W R A L I D H R L K P L E L T Q T H W V T L H N I H Q L P P D Q S Q I Q L A K A I G I E Q
- Secondary structure:** The structure is predominantly disordered, with some local order indicated by green bars.
- Disorder:** The disorder analysis shows a high degree of disorder across the entire segment, indicated by a red bar.

Section 2 (Residues 70-120):

- Sequence:** P S L V R T L D Q L E E K G L I S R Q T C A S D R R A K R I K L T E K A E P L I S E M E A V I N K T R A E I L H G I S A
- Secondary structure:** The structure shows a mix of ordered regions (green bars) and disordered regions (blue arrows).
- Disorder:** The disorder analysis shows a mix of high (red) and low (blue) disorder across the segment.

Section 3 (Residues 130-140):


- Sequence:** E E L E Q L I T L I A K L E H N I I E L Q A K G
- Secondary structure:** The structure is highly ordered, with a continuous green bar.
- Disorder:** The disorder analysis shows a low degree of disorder across the entire segment, indicated by a blue bar.

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand