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Figure 1 displays the protein structure and disorder analysis of the C-terminal region of the human protein. The figure is organized into four segments, each showing the sequence, secondary structure (SS), and disorder confidence.

Segment 1 (Residues 1-60):

- Sequence: M N R F V I A D S T L C I G C H T C E A A C S E T H R Q H G L Q S M P R L R V M L N E K E S A P Q L C H H C E D A P C A
- Secondary structure: SS (blue arrows and green helices)
- Disorder confidence: High (red/green bars)

Segment 2 (Residues 70-120):

- Sequence: V V C P V N A I T R V D G A V Q L N E S L C V S C K L C G I A C P F G A I E F S G S R P L D I P A N A N T P K A P P A P
- Secondary structure: SS (blue arrows and green helices)
- Disorder confidence: High (red/green bars)

Segment 3 (Residues 130-180):

- Sequence: P A P A R V S T L L D W V P G I R A I A V K C D L C S F D E Q G P A C V R M C P T K A L H L V D N T D I A R V S K R K R
- Secondary structure: SS (blue arrows and green helices)
- Disorder confidence: High (red/green bars)

Segment 4 (Residues 190-200):

- Sequence: E L T F N T D F G D L T L F Q Q A Q S G E A K
- Secondary structure: SS (blue arrow and green helices)
- Disorder confidence: Low (green bars)

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand