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The figure displays five rows of protein analysis data across four segments of a protein sequence:

- Segment 1 (Residues 1-60):** Sequence MQTEQQRAVTRLCIQCGFLFLQHGAESALVDELSSRLGRALGMDSVESSISSNAIVLTTI. Secondary structure features several alpha-helices and two long beta-strands.
- Segment 2 (Residues 70-120):** Sequence KDGGCLTSTRKNHDRGINMHVVTEVQHI VILAEHHLLDYKGV EKRFSQIQPLRYPRWLV A. Secondary structure includes one beta-strand followed by multiple alpha-helices.
- Segment 3 (Residues 130-180):** Sequence LMVG LSCACFC KLNNGGW DGA VI TFFASTTAMYIRQLLAQRHLHPQINFC LTAF AATTIS. Secondary structure consists of continuous alpha-helices.
- Segment 4 (Residues 190-240):** Sequence GLLL QLPTF SNTPTI AMAASV LL LVPGFP LI NAVADMF KGHI NTGLARWAI ASLL TLATC. Secondary structure continues with alpha-helices.
- Segment 5 (Residues 250-258):** Sequence VGVMALT IWGLRGVW. Secondary structure shows an alpha-helix.

Each segment has associated tracks for SS confidence (color-coded), Disorder (indicated by "?"), and Disorder confidence (color-coded).

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand