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Protein structure prediction results for three protein segments. Each segment shows the amino acid sequence, secondary structure (alpha helices and beta strands), solvent accessibility (SS), confidence scores, and disorder predictions.

Segment 1 (Residues 1-60):

- Sequence: MELTTRTLPAARKHIALVAHDHCKQMLMSVVERHQPLLEQHVLYATGTTGNLISRATGMNV
- Secondary structure: Alpha helices (green) and a beta strand (blue).
- SS: Solvent accessibility scores.
- confidence: Confidence scores for each residue.
- Disorder: Disorder predictions (mostly negative).
- Disorder confidence: Confidence scores for disorder predictions.

Segment 2 (Residues 70-120):

- Sequence: NAMLSGPMGGDQQVGALISEGKIDVLIFFWDPLNAVPHDPDKALLRLATVWNI PVATNV
- Secondary structure: Alpha helices (green) and a beta strand (blue).
- SS: Solvent accessibility scores.
- confidence: Confidence scores for each residue.
- Disorder: Disorder predictions (mostly negative).
- Disorder confidence: Confidence scores for disorder predictions.

Segment 3 (Residues 130-150):

- Sequence: ATADFI IQSPHFNDAVDILIPDYQRYLADRLK
- Secondary structure: Alpha helices (green).
- SS: Solvent accessibility scores.
- confidence: Confidence scores for each residue.
- Disorder: Disorder predictions (mostly negative).
- Disorder confidence: Confidence scores for disorder predictions.

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand