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
Protein structure and disorder analysis for the protein sequence: MKILLAAALTSSFMLVGCTPRI EVAAPKEPI T I NMNVKIEHEI I KADKDV EELL ETRSDL.

The analysis includes the following tracks:


- Sequence:** MKILLAAALTSSFMLVGCTPRI EVAAPKEPI T I NMNVKIEHEI I KADKDV EELL ETRSDL
- Secondary structure (SS):** Represented by green arrows (alpha-helix) and blue arrows (beta-strand). The structure shows a series of alpha-helices from residue 1 to 15, followed by a beta-strand at residue 16, and then a series of alpha-helices from residue 30 to 55.
- SS confidence:** A bar chart showing the confidence of the secondary structure prediction. The confidence is generally high, with some lower confidence regions (yellow and red) at residues 1-15 and 30-55.
- Disorder:** A bar chart showing the disorder of the protein. The protein is mostly ordered (green), with some disordered regions (yellow and red) at residues 1-15 and 30-55.
- Disorder confidence:** A bar chart showing the confidence of the disorder prediction. The confidence is generally high, with some lower confidence regions (yellow and red) at residues 1-15 and 30-55.


Sequence	F
Secondary structure	—
SS confidence	■
Disorder	?
Disorder confidence	■

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand