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Description	P66899
Date	Thu Jan 5 12:10:21 GMT 2012
Unique Job ID	64e437e3879ae46e

Protein structure prediction results for the protein sequence MSVFSLKIDIANKFNGETSPFSSQSQAKLARQFHQKIAGYRPTPLCALDDLNLFGVK. The figure displays the sequence, secondary structure, SS confidence, disorder, and disorder confidence across the entire length of the protein (1-60).

**Sequence:** MSVFSLKIDIANKFNGETSPFSSQSQAKLARQFHQKIAGYRPTPLCALDDLNLFGVK

**Secondary structure:** The structure is shown as a series of green cylinders representing alpha-helices and blue arrows representing beta-strands. The structure is highly flexible, with many regions of disorder.

**SS confidence:** The confidence is shown as a bar chart with colors ranging from red (low confidence) to green (high confidence). The confidence is generally low, with some higher confidence regions.

**Disorder:** The disorder is indicated by a bar chart with colors ranging from red (disordered) to green (ordered). The protein is highly disordered, with many regions of disorder.

**Disorder confidence:** The confidence is shown as a bar chart with colors ranging from red (low confidence) to green (high confidence). The confidence is generally low, with some higher confidence regions.

Confidence Key  
High(9)  Low (0)  
? Disordered  
 Alpha helix  
 Beta strand