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Description	P76227
Date	Thu Jan 5 12:20:53 GMT 2012
Unique Job ID	3633b1ccb8893e31

The figure displays protein structure and disorder analysis for the N-terminal region of the protein, showing residues 1 to 60 (top) and 70 to 90 (bottom).


Top Panel (Residues 1-60):

- Sequence:** MSRALFAVVLAFPLIALANPHYRPPDVEVNVPP E V F S SGGQSAQPC T Q C C V Y Q DQNYSEGA
- Secondary structure:** Shows alpha-helices (green) and beta-strands (blue).
- SS confidence:** A bar chart showing confidence levels for the sequence-sequence comparison.
- Disorder:** A bar chart showing disorder levels, with question marks indicating regions of uncertainty.
- Disorder confidence:** A bar chart showing confidence levels for the disorder prediction.


Bottom Panel (Residues 70-90):


- Sequence:** V I K A E G I L L Q C Q R D D K T L S T N P L V W R R V K P
- Secondary structure:** Shows alpha-helices (green) and beta-strands (blue).
- SS confidence:** A bar chart showing confidence levels for the sequence-sequence comparison.
- Disorder:** A bar chart showing disorder levels, with question marks indicating regions of uncertainty.
- Disorder confidence:** A bar chart showing confidence levels for the disorder prediction.

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand