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Description	P0A8J4
Date	Thu Jan 5 11:07:59 GMT 2012
Unique Job ID	aaf469dec18f14a3

Protein structure and disorder analysis of the first two domains of the protein.


Top Section (Residues 1-60):

- Sequence:** MKTKLNEELLEFPPTFTYKVMGQALPELV DQVVEVVQRHAPGDYTPTVKPSSKGN YHSVSI
- Secondary structure (SS):** Shows two alpha-helices (green) and several loops (blue). The first helix is between residues 5-10, and the second is between residues 15-25.
- SS confidence:** A bar chart showing confidence levels for the SS predictions.
- Disorder:** A bar chart showing regions of predicted disorder (red) and ordered regions (blue).
- Disorder confidence:** A bar chart showing confidence levels for the disorder predictions.


Bottom Section (Residues 1-54):


- Sequence:** T I N A T H I E Q V E T L Y E E L G K I D I V R M V L
- Secondary structure (SS):** Shows two alpha-helices (green) and a loop (blue). The first helix is between residues 5-10, and the second is between residues 15-25.
- SS confidence:** A bar chart showing confidence levels for the SS predictions.
- Disorder:** A bar chart showing regions of predicted disorder (red) and ordered regions (blue).
- Disorder confidence:** A bar chart showing confidence levels for the disorder predictions.

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand