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Description	Q9JMT1
Date	Thu Jan 5 12:38:06 GMT 2012
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The figure displays two panels of protein analysis. The top panel shows the full-length protein (1-60) with its sequence, secondary structure (SS), and disorder. The bottom panel shows a fragment (70-110) with its sequence, secondary structure (SS), and disorder. The sequences are color-coded by amino acid type. The secondary structure is represented by green cylinders. The disorder is indicated by a red line with question marks.


Top Panel (Full-length protein, 1-60):

- Sequence:** MFQADAYGGYRALYESGRI TEAACMAHARRKI HDVHARAPT DVTTEALQRI SELYAI EVA
- Secondary structure:** A blue arrow indicates a signal peptide. The protein has several alpha-helices (green cylinders) and loops.
- SS confidence:** A bar chart showing confidence levels for the secondary structure.
- Disorder:** A red line with question marks indicating regions of disorder.
- Disorder confidence:** A bar chart showing confidence levels for the disorder prediction.


Bottom Panel (Fragment, 70-110):


- Sequence:** VRGCSAEQRLAARKARPAPLMQSLYDWI QTQMKT LARHSDTAKAFAYLLQ
- Secondary structure:** The fragment contains several alpha-helices (green cylinders) and loops.
- SS confidence:** A bar chart showing confidence levels for the secondary structure.
- Disorder:** A red line with question marks indicating regions of disorder.
- Disorder confidence:** A bar chart showing confidence levels for the disorder prediction.

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand