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Description	P03825
Date	Thu Jan 5 10:58:02 GMT 2012
Unique Job ID	05ac18d10b5a5eab

The figure displays three protein structure models and their corresponding analysis tracks. The top model represents the full-length protein (1-120), showing a continuous alpha-helical structure. The middle model shows a fragment (1-20) with a blue arrow indicating a transition to a beta-strand structure. The bottom model shows a fragment (130-140) with a beta-strand structure. Each model is accompanied by tracks for Sequence, Secondary structure, SS confidence, Disorder, and Disorder confidence.

**Protein 1 (1-120):**

- Sequence: M F E F Y I A A R E Q K E T G H P G I F S R Q K H S T I I Y V I C L L L I C L W F A G M V L V G G Y A R Q L W V L W I V
- Secondary structure: Alpha-helical structure (green cylinders).
- SS confidence: High confidence (red/orange).
- Disorder: Low disorder (mostly dashes).
- Disorder confidence: High confidence (mostly dashes).


**Protein 2 (1-20):**

- Sequence: K A E V T V E A E T P A F K Q S T Q H Y F F K K Q P L P V V E S V E E E D D P G V A V E N A P S S S E D E E N T V E E S
- Secondary structure: Alpha-helical structure (green cylinders) transitioning to a beta-strand (blue arrow).
- SS confidence: High confidence (red/orange).
- Disorder: Low disorder (mostly dashes).
- Disorder confidence: High confidence (mostly dashes).


**Protein 3 (130-140):**


- Sequence: E E K A G L R E R V K N A L N E L E R
- Secondary structure: Beta-strand structure (green cylinders).
- SS confidence: High confidence (red/orange).
- Disorder: Low disorder (mostly dashes).
- Disorder confidence: High confidence (mostly dashes).

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand