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Description	P15373
Date	Thu Jan 5 11:34:46 GMT 2012
Unique Job ID	192fa70646c8c03e

Figure 1 displays the schematic representation of the protein structure of the human protein. The protein is composed of two main domains, each with a beta-barrel structure. The first domain (residues 1-60) is followed by a flexible linker (residues 61-70) and the second domain (residues 71-110). The figure shows the sequence, secondary structure (SS), confidence, disorder, and disorder confidence for both segments.

Segment 1 (Residues 1-60):

- Sequence:** MPANARSHAVLT TTESKVTIRGQTTPAPVREALKLKPGQDSIHYEILPGGQVFMCR LGDE
- Secondary structure (SS):** The structure shows two beta-barrel domains connected by a flexible linker. The first domain is residues 1-30, and the second is residues 31-60.
- confidence:** A bar chart showing confidence levels across the sequence.
- Disorder:** A bar chart showing disorder levels across the sequence.
- Disorder confidence:** A bar chart showing disorder confidence levels across the sequence.

Segment 2 (Residues 70-110):

- Sequence:** QEDHTMNAFLRF L DADI QN NPQKT RPFNI QQGKKLVAGMDVNI DDEIGDDE
- Secondary structure (SS):** The structure shows a single beta-barrel domain (residues 71-110).
- confidence:** A bar chart showing confidence levels across the sequence.
- Disorder:** A bar chart showing disorder levels across the sequence.
- Disorder confidence:** A bar chart showing disorder confidence levels across the sequence.

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand