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Description	P323231
Date	Thu Jan 5 11:51:31 GMT 2012
Unique Job ID	0271cbe961755b89

Protein structure visualization of the full-length protein (1-420). The structure is shown as a ribbon diagram with green alpha-helices and loops. The sequence is displayed in a color-coded font above the structure. Below the sequence, four tracks provide additional information: Secondary structure (SS), confidence, Disorder, and Disorder confidence.

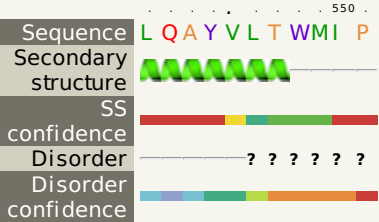
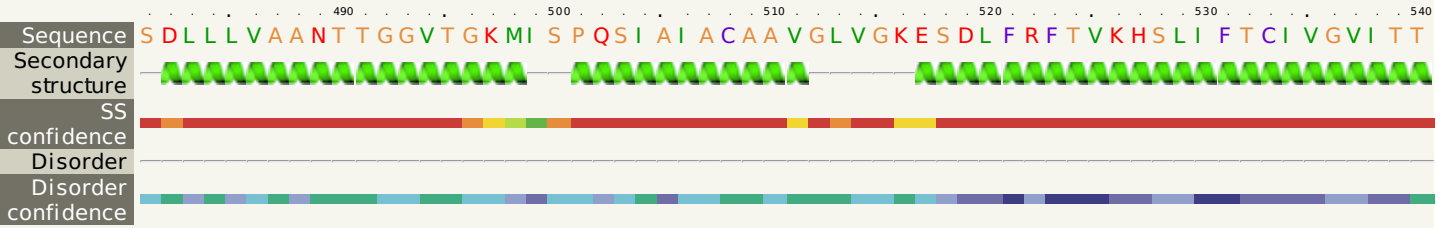
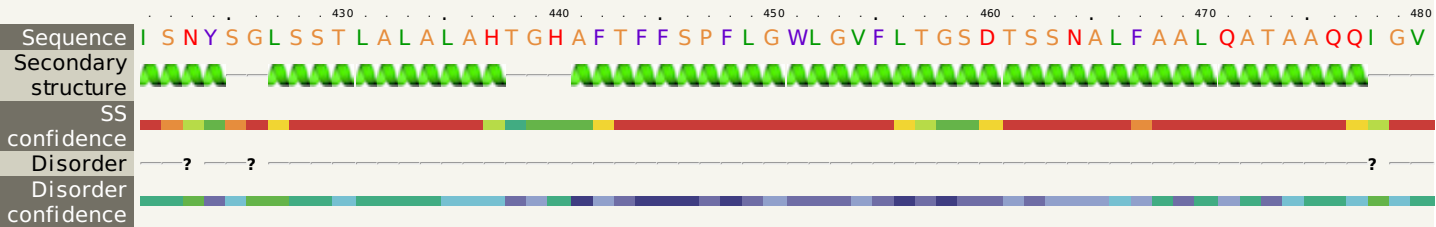
**Sequence:** MNLWQQNYDPAGNIWLSALI LFFFAALIKLKLKGYVAASWTVAILALAVALLFYKM  
 PVANALASVVYGFYGLWPIAWIIAAVFVYKISVKTGQFDIIRSSILSITPDQRLQMLI  
 VGFCFGAFLEGAAGFGAPVAITALLVGLGFKPLYAAGLCLIVNTAPVAFGAMGIPILVA  
 GQVTGIDSFIEGQMVGRQLPFMTIIVLFWIMAIMDGWRGKETWPAVVVAGGSFAIAQYL  
 SSNFI GP ELPDIISLVSLCLTLFLKRWQPVRVFRFGDLGASQVDMTLAHTGYTAGQVL  
 RAWTPFLFLTATVTLSIPPFKALFASGALYEWVINIPVPYLDKLVARMPPVVSEATAY  
 AAVFKFDWF SATGTAILFAALLSIVWLKMKPSDAISTFGSTLKEALALPIYSIGMVLAFAF

**Secondary structure (SS):** The structure is primarily composed of alpha-helices, with some loops and turns. The helices are represented by green cylinders, and the loops are represented by green lines.

**confidence:** The confidence track shows the reliability of the secondary structure prediction, with colors ranging from red (high confidence) to blue (low confidence).

**Disorder:** The disorder track indicates regions of the protein that are predicted to be disordered, with colors ranging from red (high disorder) to blue (low disorder).

**Disorder confidence:** The disorder confidence track shows the reliability of the disorder prediction, with colors ranging from red (high confidence) to blue (low confidence).



Confidence Key  
High(9) Low (0)  
? Disordered  
Alpha helix  
Beta strand