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Description	P16528
Date	Thu Jan 5 11:35:13 GMT 2012
Unique Job ID	1814da404c78f9b0

Protein structure prediction results for the protein sequence MVAPIPAKRGKRKPAVATAPATGQVQSLTRGLKLLLEWI AESNGSVALT ELAQAGLPNSTT. The figure displays the sequence, secondary structure, confidence, disorder, and disorder confidence across the entire length of the protein (1-60).

Sequence: MVAPIPAKRGKRKPAVATAPATGQVQSLTRGLKLLLEWI AESNGSVALT ELAQAGLPNSTT

Secondary structure:

SS confidence:

Disorder:

Disorder confidence:

Sequence: HRL LTTMQQQGFVRQVGELGHWAIGA HAFMVGSSFLQSRNLLAIVHPILRNLMEESETV

Secondary structure:

SS confidence:

Disorder:

Disorder confidence:

Sequence: NMAVL DQSDHEAIIIDQVQCTHLMRMSAPIGGKLP MHASGAGKAF LAQLSEEQVT KLLHR

Secondary structure:

SS confidence:

Disorder:

Disorder confidence:

Sequence: KGLHAYTHATLVSPVHLKEDLAQTRKRGYSFDDEEHALGLRCLAACIFDEHREPF AAI SI

Secondary structure:

SS confidence:

Disorder:

Disorder confidence:

Sequence: SGPI SRI TDDRVT EFGAMVI KAAKEVT LAYGGMR

Secondary structure:

SS confidence:

Disorder:

Disorder confidence:

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand