

Email	l.a.kelley@imperial.ac.uk
Description	Q46943
Date	Thu Jan 5 12:36:04 GMT 2012
Unique Job ID	2cba8e09e0fc3e17

Protein structure analysis of the N-terminal region of the protein. The analysis is divided into three segments: residues 1-60, 70-120, and 130-160. Each segment displays the amino acid sequence, secondary structure (alpha-helices and beta-strands), sequence confidence (SS), disorder, and disorder confidence. The N-terminal region (1-60) shows a high degree of disorder and low confidence. The middle region (70-120) shows a more structured domain with several alpha-helices and beta-strands. The C-terminal region (130-160) shows a highly structured domain with a large alpha-helix and a beta-strand.

➡ Beta strand