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Protein structure and disorder analysis of the C-terminal region (residues 1-420). The figure displays the amino acid sequence, secondary structure (SS), confidence, disorder, and disorder confidence for each residue.

Sequence: MNATAAQTKSLEWLNRLRANPKIPLIVAGSAAVAVMVALILWAKAPDYRTLFSNLSDDQG
 Secondary structure: [Alpha-helices and beta-strands]
 SS confidence: [Bar chart]
 Disorder: [Red line with question marks]
 Disorder confidence: [Bar chart]

Sequence: GAI VSQLTQMNI PYRFSEASGAIEVPADKVVHELRLRLAQQGLPKGGAVGF ELLDQEKFGI
 Secondary structure: [Alpha-helices and beta-strands]
 SS confidence: [Bar chart]
 Disorder: [Red line with question marks]
 Disorder confidence: [Bar chart]

Sequence: SQFSEQVNYQRAL EGE LSR TIETI GPVKGARVHLAMPKPSLVREQKSPSASVTVNLLPG
 Secondary structure: [Alpha-helices and beta-strands]
 SS confidence: [Bar chart]
 Disorder: [Red line with question marks]
 Disorder confidence: [Bar chart]

Sequence: RALDEGQISAI VHLVSSAVAGLPPGNVTLV DQGGHLLTQSNTSGRDLNDAQLKYASDVEG
 Secondary structure: [Alpha-helices and beta-strands]
 SS confidence: [Bar chart]
 Disorder: [Red line with question marks]
 Disorder confidence: [Bar chart]

Sequence: RIQRRIEAILSPI VGN GNIHAQVTAQLDFASKEQTEEQYRPNGDESHAALRSRQLNESEQ
 Secondary structure: [Alpha-helices and beta-strands]
 SS confidence: [Bar chart]
 Disorder: [Red line with question marks]
 Disorder confidence: [Bar chart]

Sequence: SSGGYPGGVPGALSNQPAPANNAPISTPPANQNNRQQQASTTSNSGPRSTQRNETSNYEV
 Secondary structure: [Alpha-helices and beta-strands]
 SS confidence: [Bar chart]
 Disorder: [Red line with question marks]
 Disorder confidence: [Bar chart]

Sequence: DRTIRHTKMNVGDVQRLSVAVVVNYKTLPLDGKPLPLSNEQMKQIEDLTREAMGFSSEKRGD
 Secondary structure: [Alpha-helices and beta-strands]
 SS confidence: [Bar chart]
 Disorder: [Red line with question marks]
 Disorder confidence: [Bar chart]

