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Protein structure and disorder analysis for the protein sequence MKKRIP...VNT. The figure displays the full sequence (1-420 residues) with secondary structure elements (alpha-helices and beta-strands) and disorder predictions. The sequence is color-coded by residue type, and the disorder analysis shows regions of high and low confidence.

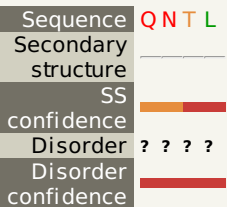
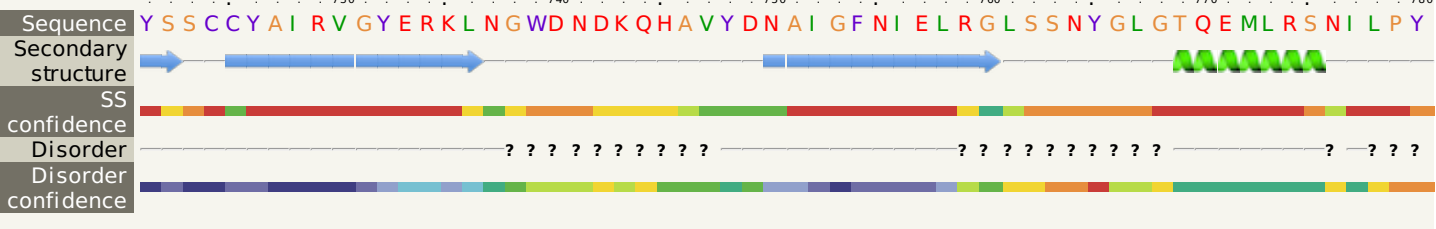
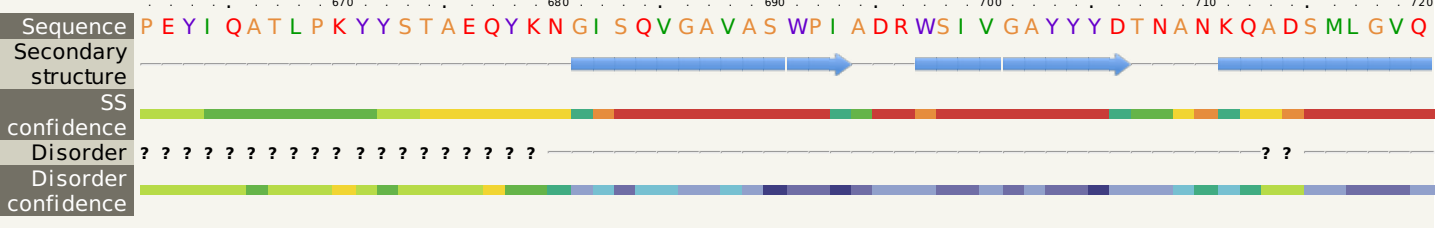
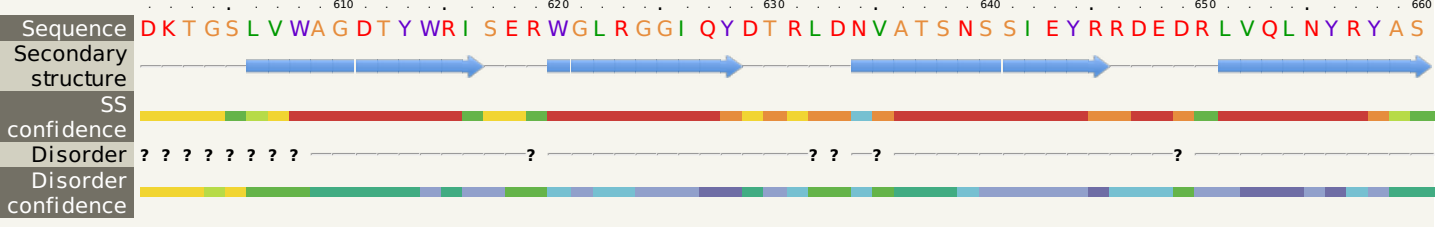
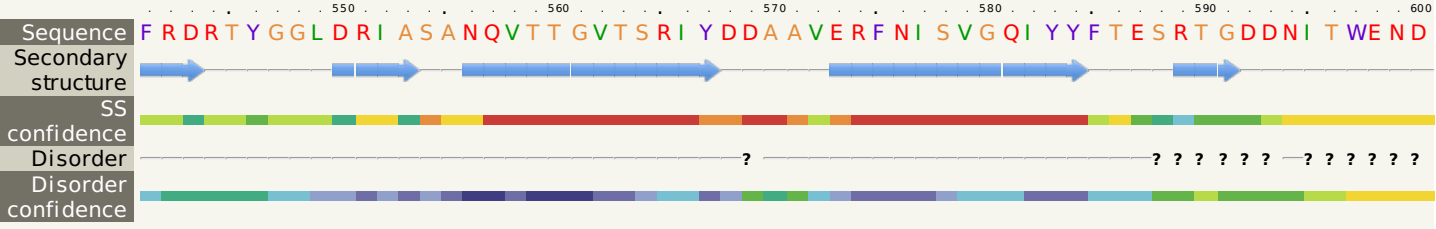
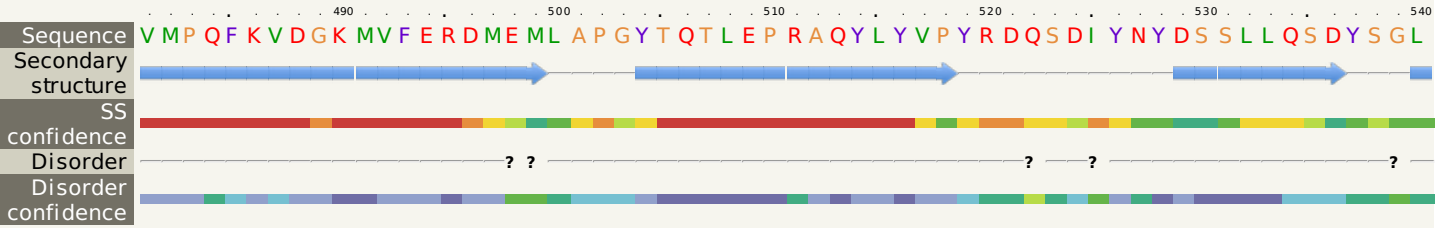
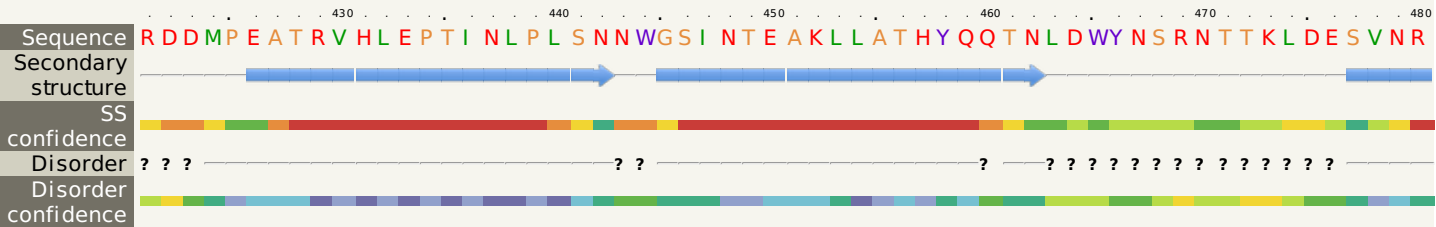
**Sequence:** MKKRIP...VNT

**Secondary structure:** [Alpha-helices and beta-strands are indicated by blue arrows and cylinders]

**SS confidence:** [Confidence scale for secondary structure]

**Disorder:** [Disorder prediction scale]

**Disorder confidence:** [Confidence scale for disorder prediction]



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