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Description	P69910
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Unique Job ID	8c42f0d3e272b728

Protein structure prediction results for the protein sequence MDKKQVTDLRS... showing sequence, secondary structure, SS confidence, disorder, and disorder confidence across 420 residues.

Sequence: MDKKQVTDLRS ELLDSRF GA KSI STI AESKR FPLHE MRDDVAFQI INDELYLDGNARQNL

Secondary structure: [Alpha-helices and loops represented by green cylinders and blue arrows]

SS confidence: [Confidence scores for secondary structure prediction]

Disorder: [Disorder prediction scores]

Disorder confidence: [Confidence scores for disorder prediction]

Sequence: ATFCQTWDDEN VHKLMDLSI NKNWI DKEEYPQSAAI DLRCVNMVADL WHAPAPKNGQAVG

Secondary structure: [Alpha-helices and loops represented by green cylinders and blue arrows]

SS confidence: [Confidence scores for secondary structure prediction]

Disorder: [Disorder prediction scores]

Disorder confidence: [Confidence scores for disorder prediction]

Sequence: TNTIGSSEACMLGGMAMKWR WRKRME AAGKPTDKPNLVC GPVQI CWHKFARYWDVELREI

Secondary structure: [Alpha-helices and loops represented by green cylinders and blue arrows]

SS confidence: [Confidence scores for secondary structure prediction]

Disorder: [Disorder prediction scores]

Disorder confidence: [Confidence scores for disorder prediction]

Sequence: PMRPGQLFMDPKRMI EACDENTI GVVPTFGVTYTGNYEFP QPLHDALDKFQADTGI DIDI DM

Secondary structure: [Alpha-helices and loops represented by green cylinders and blue arrows]

SS confidence: [Confidence scores for secondary structure prediction]

Disorder: [Disorder prediction scores]

Disorder confidence: [Confidence scores for disorder prediction]

Sequence: HIDAASGGFLAPFVAPDI VWDFRLPRVKSI SASGHKFG L APLGCGWVI WRDEEALPQELV

Secondary structure: [Alpha-helices and loops represented by green cylinders and blue arrows]

SS confidence: [Confidence scores for secondary structure prediction]

Disorder: [Disorder prediction scores]

Disorder confidence: [Confidence scores for disorder prediction]

Sequence: FNV DYLGGQI GTFAI NFSRPAGQVI AQYYEFLRLGREGYT KVQNASYQVAAYLADEIAKL

Secondary structure: [Alpha-helices and loops represented by green cylinders and blue arrows]

SS confidence: [Confidence scores for secondary structure prediction]

Disorder: [Disorder prediction scores]

Disorder confidence: [Confidence scores for disorder prediction]

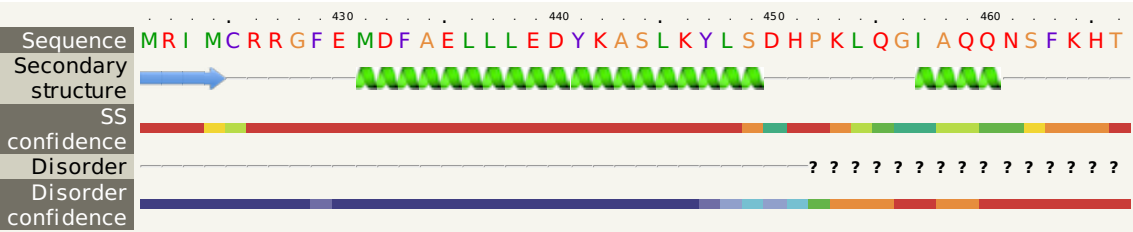
Sequence: GPYEFICTGRPDEGI PAVCFKLKDGEDPGYTLYDLSE R LRLRGWQVPAFTLGGEATDII VV

Secondary structure: [Alpha-helices and loops represented by green cylinders and blue arrows]

SS confidence: [Confidence scores for secondary structure prediction]

Disorder: [Disorder prediction scores]

Disorder confidence: [Confidence scores for disorder prediction]



Confidence Key

High(9) [red bar] [orange bar] [yellow bar] [green bar] [light blue bar] [dark blue bar] Low (0)

? Disordered

Alpha helix

Beta strand