

Email	l.a.kelley@imperial.ac.uk
Description	P09152
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Protein structure prediction results for a protein of length 480. The figure displays the amino acid sequence, predicted secondary structure (alpha helices and beta strands), and disorder confidence scores across the entire length of the protein.

Sequence: MSKFLDRFRYFKQKGETFADGHGQLLNTNRDWEDGYRQRWQHDKIVRSTHGVNCTGSCSW
 Secondary structure: [Alpha helices and beta strands indicated by arrows and cylinders]
 SS confidence: [Color-coded scale from 0 to 100]
 Disorder: [Disorder prediction line with question marks]
 Disorder confidence: [Color-coded scale from 0 to 100]

Sequence: KIYVKNGLVTWETQQTDPRTTRPDLPNHEPRGCPRGASYSWYLYSANRLKYPPMRKRLMK
 Secondary structure: [Alpha helices and beta strands indicated by arrows and cylinders]
 SS confidence: [Color-coded scale from 0 to 100]
 Disorder: [Disorder prediction line with question marks]
 Disorder confidence: [Color-coded scale from 0 to 100]

Sequence: MWREAKALHSDPVEAWASIIDADKAKSFQARGRGGFVRSSWQEVNELIAASNVTYTIKN
 Secondary structure: [Alpha helices and beta strands indicated by arrows and cylinders]
 SS confidence: [Color-coded scale from 0 to 100]
 Disorder: [Disorder prediction line with question marks]
 Disorder confidence: [Color-coded scale from 0 to 100]

Sequence: YGPDRAVAGFSPIPAMSMVSYASGARYLSLIGGTCLSFYDWYCDLPPASPQTWGEQTDVPE
 Secondary structure: [Alpha helices and beta strands indicated by arrows and cylinders]
 SS confidence: [Color-coded scale from 0 to 100]
 Disorder: [Disorder prediction line with question marks]
 Disorder confidence: [Color-coded scale from 0 to 100]

Sequence: SADWYNSSYIIAWGSNVPQTRTPDAHFFTEVRYKGTKTVAVTPDYAEIAKLCDLWLAPKQ
 Secondary structure: [Alpha helices and beta strands indicated by arrows and cylinders]
 SS confidence: [Color-coded scale from 0 to 100]
 Disorder: [Disorder prediction line with question marks]
 Disorder confidence: [Color-coded scale from 0 to 100]

Sequence: GTDAAMALAMGHVMLREFHLDNPSQYFTDYVRRYTDMPMLVMLEERDGYAAGRMLRAAD
 Secondary structure: [Alpha helices and beta strands indicated by arrows and cylinders]
 SS confidence: [Color-coded scale from 0 to 100]
 Disorder: [Disorder prediction line with question marks]
 Disorder confidence: [Color-coded scale from 0 to 100]

Sequence: LVDALGQENNPEWKTVAFNTNGEMVAPNGSIGFRWGEKKGKWNLEQRDGKTGEETELQLSL
 Secondary structure: [Alpha helices and beta strands indicated by arrows and cylinders]
 SS confidence: [Color-coded scale from 0 to 100]
 Disorder: [Disorder prediction line with question marks]
 Disorder confidence: [Color-coded scale from 0 to 100]

Sequence: LGSQDEIAEVGFPHYGGDGT EHFNKVELENVLLHKLVPVKRLQLADGSTALVTTVYDLTLA
 Secondary structure: [Alpha helices and beta strands indicated by arrows and cylinders]
 SS confidence: [Color-coded scale from 0 to 100]
 Disorder: [Disorder prediction line with question marks]
 Disorder confidence: [Color-coded scale from 0 to 100]





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