





Email	l.a.kelley@imperial.ac.uk
Description	Q57049
Date	Thu Jan 5 12:37:12 GMT 2012
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Protein structure and disorder analysis for the protein sequence: ME G W R V L N K L A N L R I D Q N K S G T T A F H A G K F F R F P A F F S D P T V N S L F S H T E A F S Q F F G G H F.

The analysis includes the following tracks:


- Sequence:** ME G W R V L N K L A N L R I D Q N K S G T T A F H A G K F F R F P A F F S D P T V N S L F S H T E A F S Q F F G G H F.
- Secondary structure:** Visualization of alpha-helices (red) and beta-strands (blue).
- SS confidence:** Confidence scores for secondary structure prediction, ranging from 0.00 (blue) to 0.99 (red).
- Disorder:** Prediction of intrinsically disordered regions, marked with question marks.
- Disorder confidence:** Confidence scores for disorder prediction, ranging from 0.00 (blue) to 0.99 (red).

Sequence	I	A
Secondary structure		
SS confidence		
Disorder		
Disorder confidence		

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand