



| | |
|---------------|-----------------------------|
| Email | l.a.kelley@imperial.ac.uk |
| Description | P18393 |
| Date | Thu Jan 5 11:36:56 GMT 2012 |
| Unique Job ID | 1accfe2fc9a91c8d |


Protein structure and disorder analysis for the first protein (residues 1-60). The sequence is: M A F S N P F D D P Q G A F Y I L R N A Q G Q F S L W P Q Q C V L P A G W D I V C Q P Q S Q A S C Q Q W L E A H W R T L. The secondary structure shows two alpha-helices (green) and two beta-strands (blue). The SS confidence is high (red/yellow). The Disorder confidence is low (dark blue). The disorder analysis shows two regions of high disorder (red/yellow) at the N-terminus and C-terminus.

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand