

Email	l.a.kelley@imperial.ac.uk
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The figure displays three panels of protein analysis for different regions of the protein. Each panel shows the amino acid sequence, secondary structure (alpha helices and beta strands), solvent accessibility (SS), disorder, and disorder confidence.

Panel 1 (Residues 1-60):

- Sequence:** MAQVAIFKEIFDQVRKDLNCELFYSELKRHNVS HYIYYLATDNIHIVLENDNTVLIKGLK
- Secondary structure:** Shows alpha helices (green) and beta strands (blue).
- SS:** Solvent accessibility scale from 0 to 100.
- Disorder:** Disorder scale from 0 to 100.
- Disorder confidence:** Disorder confidence scale from 0 to 100.


Panel 2 (Residues 70-120):

- Sequence:** KVVNVKFSRNTHLIENTSYDRLKSREITFQQYRENLA KAGVFRWITNIHEHKRYYYTFDNS
- Secondary structure:** Shows alpha helices (green) and beta strands (blue).
- SS:** Solvent accessibility scale from 0 to 100.
- Disorder:** Disorder scale from 0 to 100.
- Disorder confidence:** Disorder confidence scale from 0 to 100.


Panel 3 (Residues 130-140):


- Sequence:** LLFTESI QNTTQIFPR
- Secondary structure:** Shows alpha helices (green) and beta strands (blue).
- SS:** Solvent accessibility scale from 0 to 100.
- Disorder:** Disorder scale from 0 to 100.
- Disorder confidence:** Disorder confidence scale from 0 to 100.

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand