

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

Panel 1 (Residues 1-60):

- Sequence:** M T E M A K G S V T H Q R L I A L L S Q E G A D F R V V T H E A V G K C E A V S E I R G T A L G Q G A K A L V C K V K G
- Secondary structure:** Shows alpha helices (green cylinders) and beta strands (blue arrows).
- SS:** Solvent accessibility scores.
- confidence:** Confidence scores for the structure.
- Disorder:** Disorder predictions (red and yellow bars).
- Disorder confidence:** Confidence scores for the disorder predictions.

Panel 2 (Residues 70-120):

- Sequence:** N G V N Q H V L A I L A A D Q Q A D L S Q L A S H I G G L R A S L A S P A E V D E L T G C V F G A I P P F S F H P K L K
- Secondary structure:** Shows alpha helices (green cylinders) and beta strands (blue arrows).
- SS:** Solvent accessibility scores.
- confidence:** Confidence scores for the structure.
- Disorder:** Disorder predictions (red and yellow bars).
- Disorder confidence:** Confidence scores for the disorder predictions.

Panel 3 (Residues 130-160):

- Sequence:** L V A D P L L F E R F D E I A F N A G M L D K S V I L K T A D Y L R I A Q P E L V N F R R T A
- Secondary structure:** Shows alpha helices (green cylinders) and beta strands (blue arrows).
- SS:** Solvent accessibility scores.
- confidence:** Confidence scores for the structure.
- Disorder:** Disorder predictions (red and yellow bars).
- Disorder confidence:** Confidence scores for the disorder predictions.

➡ Beta strand