

Email	l.a.kelley@imperial.ac.uk
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Figure 1 displays the schematic representation of the protein structure of the human protein, showing the sequence, secondary structure, and confidence scores across three segments.

Segment 1 (Residues 1-60):

- Sequence:** MKRKRLFLLASLLPMFALAGNKVNTTLPGGNMQFQGVIIAETCRIEAGDKQMTVNMGQIS
- Secondary structure:** Alpha-helices (green) and beta-strands (blue).
- SS confidence:** High confidence (red) across the segment.
- Disorder:** High disorder (green) across the segment.
- Disorder confidence:** High confidence (red) across the segment.


Segment 2 (Residues 70-120):

- Sequence:** SNRFHAVGEDSAPVPFVILHRECVSVSERVGVAFHGVADGKNPDVLSVGE GPGIATNIG
- Secondary structure:** Alpha-helices (green) and beta-strands (blue).
- SS confidence:** High confidence (red) across the segment.
- Disorder:** High disorder (green) across the segment.
- Disorder confidence:** High confidence (red) across the segment.


Segment 3 (Residues 130-180):


- Sequence:** VALFDDEGNLVPINRPPANWKRLYSGSTSLHFI AKYRATGRRVTGGI ANAQAWFSLTYQ
- Secondary structure:** Alpha-helices (green) and beta-strands (blue).
- SS confidence:** High confidence (red) across the segment.
- Disorder:** High disorder (green) across the segment.
- Disorder confidence:** High confidence (red) across the segment.

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand