

Email	l.a.kelley@imperial.ac.uk
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The figure displays the protein structure and disorder analysis for the human protein sequence (1-600 residues). The analysis is presented in a series of tracks for each segment of the protein.

Segment 1 (Residues 1-60):

- Sequence:** MNYRYRAMTQDQGKLGQGIIDANDERQARLRRLREEGLFLLDIRPQKSSGVKTRRPRI SHSE
- Secondary structure:** Shows a blue arrow indicating a long loop/turn, followed by a series of green arrows indicating alpha-helices.
- SS confidence:** A red bar indicating high confidence in the secondary structure prediction.
- Disorder:** A line with question marks indicating regions of potential disorder.
- Disorder confidence:** A bar with various colors indicating the confidence in the disorder prediction.

Segment 2 (Residues 70-120):

- Sequence:** LTLFTRQLATLSAAALPLEESLAVIGQQSSNKRLGDLVNQVRSAILLEGHP LSDALQHFPT
- Secondary structure:** Shows a series of green arrows indicating alpha-helices.
- SS confidence:** A red bar indicating high confidence in the secondary structure prediction.
- Disorder:** A line with question marks indicating regions of potential disorder.
- Disorder confidence:** A bar with various colors indicating the confidence in the disorder prediction.

Segment 3 (Residues 130-180):

- Sequence:** LFDLSLYRTL VKAGEKSGLLAPVLEKLADYNENRQKIRSKLIQSLIYPCMLTTVAIGVVI
- Secondary structure:** Shows a series of green arrows indicating alpha-helices.
- SS confidence:** A red bar indicating high confidence in the secondary structure prediction.
- Disorder:** A line with question marks indicating regions of potential disorder.
- Disorder confidence:** A bar with various colors indicating the confidence in the disorder prediction.

Segment 4 (Residues 190-240):

- Sequence:** LLTAVVPKITEQFVHMKQQLPLSTRILLGLSDTLQRTGPTLLATVFI VAVGF WLWLKRGN
- Secondary structure:** Shows a series of green arrows indicating alpha-helices.
- SS confidence:** A red bar indicating high confidence in the secondary structure prediction.
- Disorder:** A line with question marks indicating regions of potential disorder.
- Disorder confidence:** A bar with various colors indicating the confidence in the disorder prediction.

Segment 5 (Residues 250-300):

- Sequence:** NRHRFHAMLLRVALIGPLICAINSARYLRTLSSLQSSGVPLLDGMNLSLSTENNLEIRQR
- Secondary structure:** Shows a series of green arrows indicating alpha-helices.
- SS confidence:** A red bar indicating high confidence in the secondary structure prediction.
- Disorder:** A line with question marks indicating regions of potential disorder.
- Disorder confidence:** A bar with various colors indicating the confidence in the disorder prediction.

Segment 6 (Residues 310-360):

- Sequence:** LANAAENV RQGN SIHLSLEQTAFPPMMLYMVASGEKSGQLGTL MVRAADNQETLQQNRI
- Secondary structure:** Shows a series of green arrows indicating alpha-helices.
- SS confidence:** A red bar indicating high confidence in the secondary structure prediction.
- Disorder:** A line with question marks indicating regions of potential disorder.
- Disorder confidence:** A bar with various colors indicating the confidence in the disorder prediction.

Segment 7 (Residues 370-390):

- Sequence:** ALTLSIFEPALII TMALI VLFIVVSVLQPLLQLNSMI N
- Secondary structure:** Shows a series of green arrows indicating alpha-helices.
- SS confidence:** A red bar indicating high confidence in the secondary structure prediction.
- Disorder:** A line with question marks indicating regions of potential disorder.
- Disorder confidence:** A bar with various colors indicating the confidence in the disorder prediction.

Confidence Key
High(9)  Low (0)
? Disordered
 Alpha helix
 Beta strand