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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:** MLRA FARLLRLI CFSRRTLKI ACLLLLVAGATIL I ADRV MVNASKQLTWS DVNAV PARNV
- Secondary structure:** [Alpha-helices and loops]
- SS confidence:** [Color-coded bar]
- Disorder:** [Color-coded bar]
- Disorder confidence:** [Color-coded bar]

Segment 2 (Residues 70-120):

- Sequence:** GLLLGARPGNRYFTRRI DTA AALYHAGKV K WLLVSGDNGRK NYDEASGMQ QALI AKGVPA
- Secondary structure:** [Alpha-helices and loops]
- SS confidence:** [Color-coded bar]
- Disorder:** [Color-coded bar]
- Disorder confidence:** [Color-coded bar]


Segment 3 (Residues 130-180):

- Sequence:** KVI FCDYAGFSTLDSVVRAKKVF GENHIT I SQEFHNQRAI WLA KQYGI DAI GFNAPDLN
- Secondary structure:** [Alpha-helices and loops]
- SS confidence:** [Color-coded bar]
- Disorder:** [Color-coded bar]
- Disorder confidence:** [Color-coded bar]


Segment 4 (Residues 190-230):


- Sequence:** MKHGFYTQLREKLARVSAVI DAKI LHRQPKYLGPSVMI GPFSEHGCPAQK
- Secondary structure:** [Alpha-helices and loops]
- SS confidence:** [Color-coded bar]
- Disorder:** [Color-coded bar]
- Disorder confidence:** [Color-coded bar]

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand