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Figure 1 displays the protein structure and disorder analysis of the human protein p115. The figure is divided into three panels, each showing a different region of the protein. Each panel includes the amino acid sequence, secondary structure (alpha-helices and beta-strands), solvent accessibility (SS), and disorder confidence scores.

Panel 1 (Residues 1-60):

- Sequence:** MSQT VHFQGNPVT VANSI PQAGSKAQTFTLVAKDLS DVTLGQFAGKRKVLNI FPSIDTGV
- Secondary structure:** Shows several alpha-helices (green cylinders) and beta-strands (blue arrows).
- SS (Solvent Accessibility):** A bar chart showing the relative solvent accessibility of each residue.
- Disorder:** A bar chart showing the disorder confidence scores for each residue.


Panel 2 (Residues 70-120):

- Sequence:** CAASVRKF NQLATEI DNTVVLCSADLPFAQSRFCGAEGLNNVITLSTFRNAEFLQAYGV
- Secondary structure:** Shows several alpha-helices (green cylinders) and beta-strands (blue arrows).
- SS (Solvent Accessibility):** A bar chart showing the relative solvent accessibility of each residue.
- Disorder:** A bar chart showing the disorder confidence scores for each residue.


Panel 3 (Residues 130-160):


- Sequence:** A IADGPLKGLAARAVVVIDENDNVI FSQLVDEI TTEPDYEAALAVLKA
- Secondary structure:** Shows several alpha-helices (green cylinders) and beta-strands (blue arrows).
- SS (Solvent Accessibility):** A bar chart showing the relative solvent accessibility of each residue.
- Disorder:** A bar chart showing the disorder confidence scores for each residue.

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand