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Description	P0A6K3
Date	Thu Jan 5 11:03:20 GMT 2012
Unique Job ID	7083b07b6dc5bc13

Panel 1: Residues 1-60

Sequence: MSVLQVLHI PDERLRKVAKPVEEVNAEIQRIVDDMFETMYAEEGI GLAATQVDIHQRIIV

Secondary structure: Shows alpha helices (green cylinders) and beta strands (blue arrows). Helices are located around residues 10-15, 20-25, 30-40, and 50-55.

SS: Solvent accessibility scale from 0 (red) to 1 (green).

confidence: Confidence score scale from 0 (red) to 1 (green).

Disorder: Disorder prediction scale from 0 (red) to 1 (green).

Disorder confidence: Disorder confidence scale from 0 (red) to 1 (green).

Panel 2: Residues 70-120

Sequence: IDVSENRLDERLVLI NPELLEKSGETGIEEGCLSIPEQRALVPRAEKVKIRALDRDGKPFEE

Secondary structure: Shows alpha helices (green cylinders) and beta strands (blue arrows). Helices are located around residues 75-80, 85-90, 95-100, 105-110, and 115-120.

SS: Solvent accessibility scale from 0 (red) to 1 (green).

confidence: Confidence score scale from 0 (red) to 1 (green).

Disorder: Disorder prediction scale from 0 (red) to 1 (green).

Disorder confidence: Disorder confidence scale from 0 (red) to 1 (green).

Panel 3: Residues 130-160

Sequence: LEADGLLAICI QHEMDHLVGKLFMDYLSPLKQQRIRQKVEKLDRLKARA

Secondary structure: Shows alpha helices (green cylinders) and beta strands (blue arrows). Helices are located around residues 135-140 and 145-150.


SS: Solvent accessibility scale from 0 (red) to 1 (green).

confidence: Confidence score scale from 0 (red) to 1 (green).


Disorder: Disorder prediction scale from 0 (red) to 1 (green).


Disorder confidence: Disorder confidence scale from 0 (red) to 1 (green).

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand