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Description	A6X993
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Panel 1 (Residues 1-60):
 Sequence: MVISSVFSAFADIPSVYLISVNCGARELWLTITTSIHFVSLREQRYEFLANSLGERTSSLK
 Secondary structure: Helices at residues 1-10, 12-15, 17-25, 27-38, 40-60.
 Disorder confidence: High disorder (red) at residues 1-10, 12-15, 17-25, 27-38, 40-60.

Panel 2 (Residues 70-120):
 Sequence: SQEEMVSVSRNGKQLLEASFFRLGKHVHLNFLPFENTFEMALRNDQIFCTSILFTCYI
 Secondary structure: Helices at residues 70-85, 87-95, 97-120.
 Disorder confidence: High disorder (red) at residues 70-85, 87-95, 97-120.

Panel 3 (Residues 130-140):
 Sequence: QSFSL LISNFFIAIEVSR SFF
 Secondary structure: Helices at residues 130-135, 137-140.
 Disorder confidence: High disorder (red) at residues 130-135, 137-140.

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand