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Description	P76575
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Protein 1 (Residues 1-60):

- Sequence:** M E L H C P Q C Q H V L D Q D N G H A R C R S C G E F I E M K A L C P D C H Q P L Q V L K A C G A V D Y F C Q H G H G L
- Secondary structure:** Alpha-helices (blue arrows) and beta-strands (green zig-zags).
- SS confidence:** High confidence (green/blue) across most of the segment.
- Disorder:** Low confidence (red/orange) at the N-terminus and C-terminus, high confidence (green/blue) in the middle.

Protein 2 (Residues 70-100):

- Sequence:** I S K K R V E F V L A
- Secondary structure:** Alpha-helix (blue arrow).
- SS confidence:** Low confidence (red/orange) at the N-terminus, high confidence (green/blue) in the middle.
- Disorder:** Low confidence (red/orange) at the N-terminus, high confidence (green/blue) in the middle.

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand