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|---------------|-----------------------------|
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| Description   | O14468                      |
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**Protein 1 (Top):**

- Sequence:** M S T E K L E A S E E P Q A P L A N T S E T N S I K G D T E N I V T V F D L A N E I E K S L K D V Q R Q M K E N D D E F
- Secondary structure:** Alpha-helical region (residues 1-28) and a disordered region (residues 29-60).
- SS confidence:** High confidence in the alpha-helical region, lower confidence in the disordered region.
- Disorder:** Disordered region (residues 29-60).
- Disorder confidence:** High confidence in the disordered region.

**Protein 2 (Bottom):**

- Sequence:** S R S I Q A I E D K L N K M S R
- Secondary structure:** Alpha-helical region (residues 1-11) and a disordered region (residues 12-14).
- SS confidence:** High confidence in the alpha-helical region, lower confidence in the disordered region.
- Disorder:** Disordered region (residues 12-14).
- Disorder confidence:** High confidence in the disordered region.

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand