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Description	P0A986
Date	Thu Jan 5 11:09:36 GMT 2012
Unique Job ID	86d55ab126e295af


Protein 1 (Top):

- Sequence:** MSN K MT G L V K V F N P E K G F G F I T P K D G S K D V F V H F S A I Q S N D F K T L E N Q E V E F G I E N G P K
- Secondary structure:** Four alpha-helices (blue arrows) spanning residues 1-15, 18-25, 28-38, and 50-60.
- SS confidence:** High confidence (red) for most regions, with some lower confidence (green, yellow, blue) segments.
- Disorder:** Mostly ordered (dashed line), with some disorder (question marks) at the N-terminus and C-terminus.
- Disorder confidence:** High confidence (dark blue) for the ordered regions, with some lower confidence (green, yellow, blue) segments.


Protein 2 (Bottom):


- Sequence:** G P A A V H V V A L
- Secondary structure:** One alpha-helix (blue arrow) spanning residues 1-6.
- SS confidence:** High confidence (red) for the helix, with some lower confidence (green, yellow, blue) segments.
- Disorder:** Mostly ordered (dashed line), with some disorder (question marks) at the N-terminus and C-terminus.
- Disorder confidence:** High confidence (dark blue) for the ordered regions, with some lower confidence (green, yellow, blue) segments.

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand