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Description	P0A763
Date	Wed Jan 25 15:20:17 GMT 2012
Unique Job ID	767f0c28fa1fe21e

Protein structure prediction results for three protein segments. Each segment shows the amino acid sequence, secondary structure (alpha helices and beta strands), sequence confidence, disorder, and disorder confidence.

Segment 1 (1-60):

- Sequence: MAIERTFSIKPNAVAKNVI GNIFARFEAAGFKI VGTKMLHLTVEQARGFYAEHDGKPPF
- Secondary structure: Alpha helices (green) and beta strands (blue) are shown.
- SS confidence: High confidence (red) across the entire segment.
- Disorder: No disorder predicted (all red).
- Disorder confidence: High confidence (red) across the entire segment.


Segment 2 (70-120):

- Sequence: DGLVEFM TSGPI VVS VLE GENAVQRHRDLL GATNPANALAGTLRADYADSLTENGTHGSD
- Secondary structure: Alpha helices (green) and beta strands (blue) are shown.
- SS confidence: High confidence (red) across the entire segment.
- Disorder: No disorder predicted (all red).
- Disorder confidence: High confidence (red) across the entire segment.


Segment 3 (130-140):


- Sequence: SVESAAREIAYFFGEGEVCPRT R
- Secondary structure: Alpha helices (green) and beta strands (blue) are shown.
- SS confidence: High confidence (red) across the entire segment.
- Disorder: No disorder predicted (all red).
- Disorder confidence: High confidence (red) across the entire segment.

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand