Phyre2

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Description	P0A9Q5
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Secondary structure and disorder prediction

Sequence (1–60): M S W I E R I K S N I T P T R K A S I P E G V W T K C D S C G Q V L Y R A E L E R N L E V C P K C D H H M R M T A R N R

Sequence (61–120): L H S L L D E G S L V E L G S E L E P K D V L K F R D S K K Y K D R L A S A Q K E T G E K D A L V V M K G T L Y G M P V

Sequence (121–180): V A A A F E F A F M G G S M G S V V G A R F V R A V E Q A L E D N C P L I C F S A S G G A R M Q E A L M S L M Q M A K T

Sequence (181–240): S A A L A K M Q E R G L P Y I S V L T D P T M G G V S A S F A M L G D L N I A E P K A L I G F A G P R V I E Q T V R E K

Sequence (241–300): L P P G F Q R S E F L I E K G A I D M I V R R P E M R L K L A S I L A K L M N L P A P N P E A P R E G V V V P P V P D Q

Sequence (301–304): E P E A

Rows for each block: Sequence, Secondary structure, SS confidence, Disorder, Disorder confidence

Confidence Key

High(9) … Low (0)

? Disordered

Alpha helix

Beta strand