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Description	Q09746
Date	Sun Jul 8 11:56:56 BST 2012
Unique Job ID	b0e8ecbef6aa74c1

Secondary structure and disorder prediction

Sequence
Secondary structure
SS confidence
Disorder
Disorder confidence
430 440 450 460 470 480
L S K R G C V C K N C G F K C H A R C E L L V P A N C K N G E P E V A D D D A V D T S V T A T D D F D A S A S S S N A Y
490 500 510 520 530 540
E S Y R N T Y T D D M D S S S I Y Q T S L S N V K T E E T T P A E P A S K V D G V V L Y D F T G E H E G V I T A S E G Q
550 560 570 580 590 600
E F T L L E P D D G S G W V R V K I D G T D G L I P A S Y V K L N D E L N T S V T L D G D S S Y V K A L Y A Y T A Q S D
610 620 630 640
M E L S I Q E G D I I Q V T N R N A G N G W S E G I L N G V T G Q F P A N Y V T D V
Confidence Key
High(9)
Low (0)
? Disordered
Alpha helix
Beta strand