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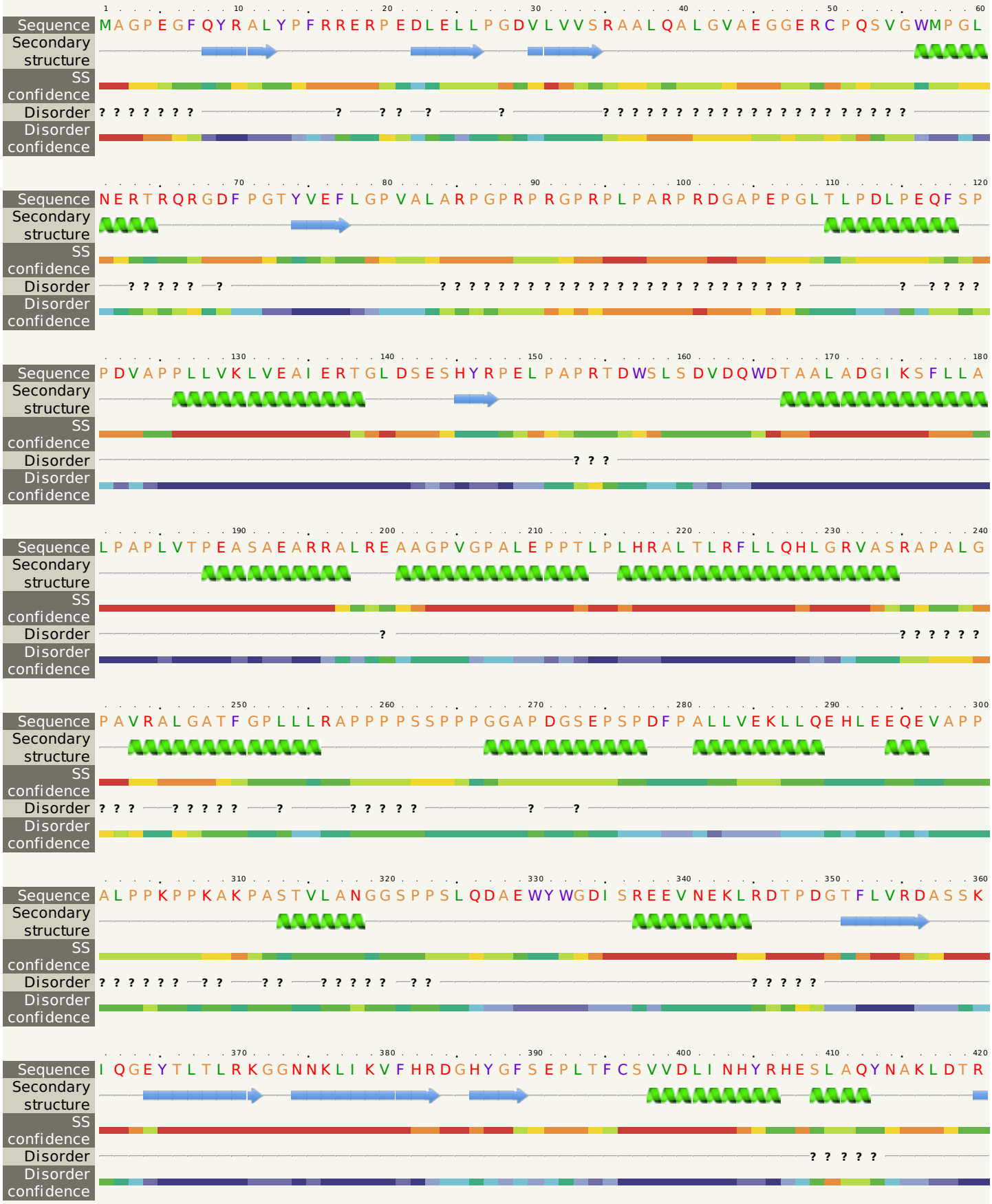
Description	O00459
Date	Fri May 25 09:44:30 BST 2012
Unique Job ID	0ae9a79684a9af6d

Secondary structure and disorder prediction



Sequence
Secondary structure
SS confidence
Disorder
Disorder confidence
430 440 450 460 470 480
L L Y P V S K Y Q Q D Q I V K E D S V E A V G A Q L K V Y H Q Q Y Q D K S R E Y D Q L Y E E Y T R T S Q E L Q M K R T A
490 500 510 520 530 540
I E A F N E T I K I F E E Q G Q T Q E K C S K E Y L E R F R R E G N E K E M Q R I L L N S E R L K S R I A E I H E S R T
550 560 570 580 590 600
K L E Q Q L R A Q A S D N R E I D K R M N S L K P D L M Q L R K I R D Q Y L V W L T Q K G A R Q K K I N E W L G I K N E
610 620 630 640 650 660
T E D Q Y A L M E D E D D L P H H E E R T W Y V G K I N R T Q A E E M L S G K R D G T F L I R E S S Q R G C Y A C S V V
670 680 690 700 710 720
V D G D T K H C V I Y R T A T G F G F A E P Y N L Y G S L K E L V L H Y Q H A S L V Q H N D A L T V T L A H P V R A P G
Sequence
Secondary structure
SS confidence
Disorder
Disorder confidence
P G P P P A A R
Confidence Key
High(9) Low (0)
? Disordered
Alpha helix
Beta strand