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Description	P46889
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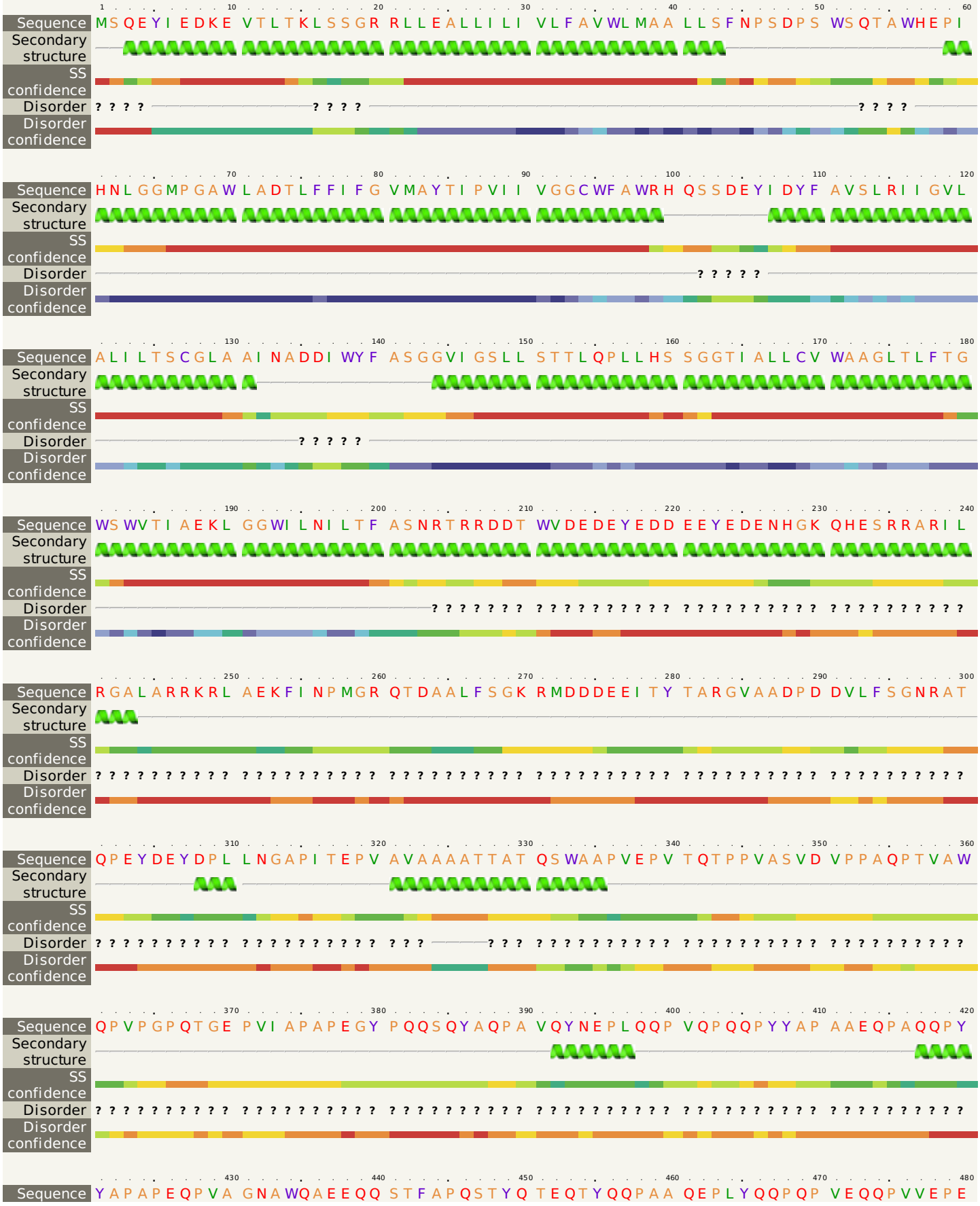
Unique Job ID	9ff5d5a8aaa0b07c

Secondary structure and disorder prediction



Secondary structure
SS confidence
Disorder
Disorder confidence
Sequence
PVVEETKPAR PPLYYFEEVE EKRAREREQL AAWYQPIPEP VKEPEPIKSS LKAPSVAAVP
PVEAAAAVSP LASGVKKATL ATGAAATVAA PVFSLANSGG PRPQVKEGIG PQLPRPKRIR
VPTRRELASY GIKLPSQRAA EEKAREAQRN QYDSGDQYND DEIDAMQQDE LARQFAQTQQ
QRYGEQYQHD VPVNAEDADA AAEAELARQF AQTQQQRYSG EQPAGANPFS LDDFEFSPMK
ALLDDGPHEP LFTPIVEPVQ QPQQPVAPQQ QYQQPQQPVP PQPQYQQPQQ PVAPQPQYQQ
PQQPVAPQQQ YQQPQQPVAP QQQYQQPQQP VAPQPQDTLL HPLLMRNGDS RPLHKPTTPL
PSLDLLTPPP SEVEPVDTFA LEQMARLVEA RLADFRIKAD VVNYSPGPVI TRFELNLAPG
VKAARISNLS RDLARSLSTV AVRVVEVIPG KPYVGLELPN KKRQTVYLRE VLDNAKFRDN

Disorder confidence
Sequence
Secondary structure
SS confidence
Disorder
Disorder confidence
970 980 990 1000 1010 1020
P S P L T V V L G K D I A G E P V V A D L A K M P H L L V A G T T G S G K S V G V N A M I L S M L Y K A Q P E D V R F I
1030 1040 1050 1060 1070 1080
M I D P K M L E L S V Y E G I P H L L T E V V T D M K D A A N A L R W C V N E M E R R Y K L M S A L G V R N L A G Y N E
1090 1100 1110 1120 1130 1140
K I A E A D R M M R P I P D P Y W K P G D S M D A Q H P V L K K E P Y I V V L V D E F A D L M M T V G K K V E E L I A R
1150 1160 1170 1180 1190 1200
L A Q K A R A A G I H L V L A T Q R P S V D V I T G L I K A N I P T R I A F T V S S K I D S R T I L D Q A G A E S L L G
1210 1220 1230 1240 1250 1260
M G D M L Y S G P N S T L P V R V H G A F V R D Q E V H A V V Q D W K A R G R P Q Y V D G I T S D S E S E G G A G G F D
1270 1280 1290 1300 1310 1320
G A E E L D P L F D Q A V Q F V T E K R K A S I S G V Q R Q F R I G Y N R A A R I I E Q M E A Q G I V S E Q G H N G N R
1330
E V L A P P P F D
Confidence Key
High(9) Low (0)
? Disordered
Alpha helix
Beta strand