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|---|---|
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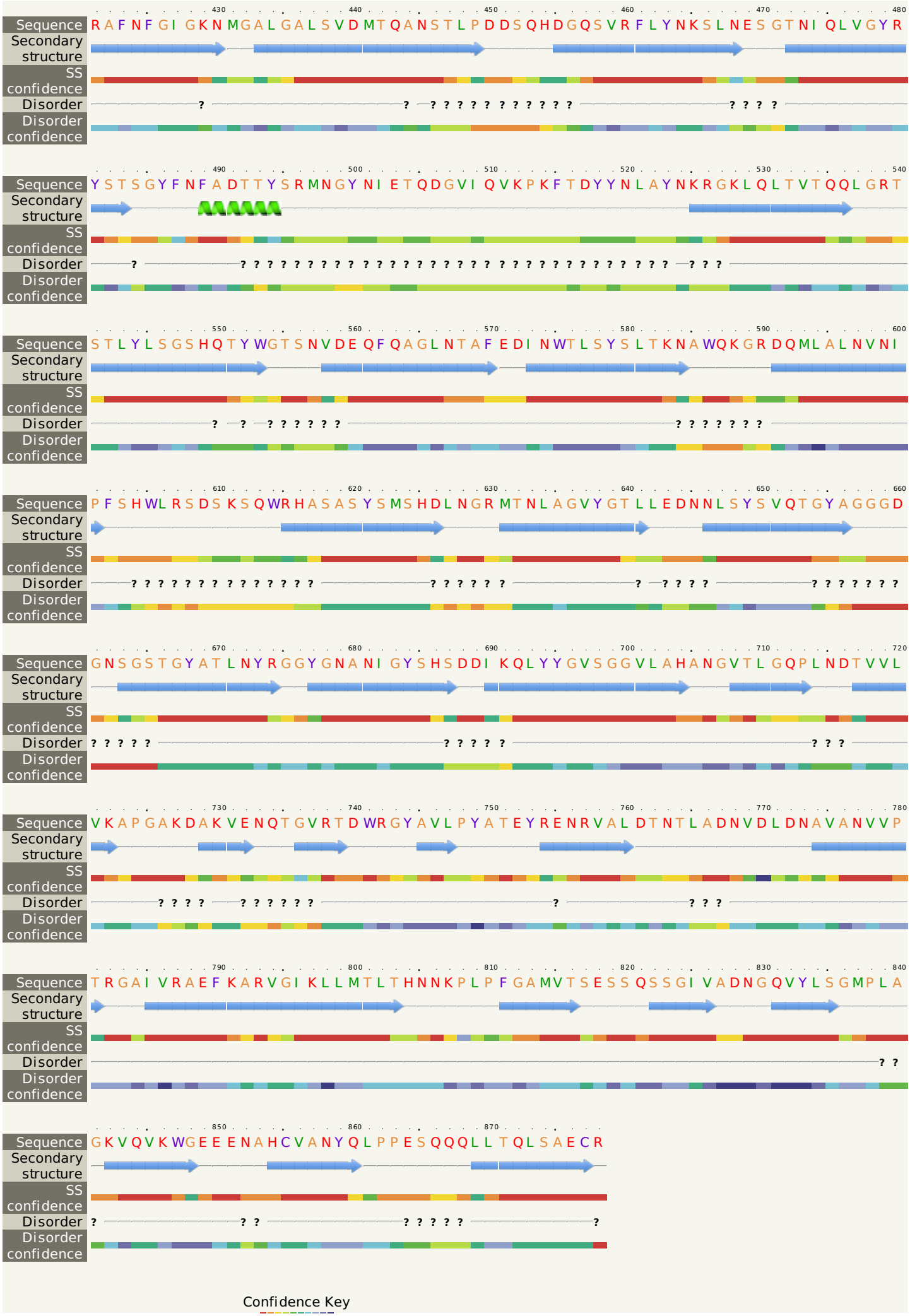
| Description | P30130 |
| Date | Thu Jan 5 11:45:50 GMT 2012 |
| Unique Job ID | ba69b6e503c1c479 |

## Secondary structure and disorder prediction


Sequence
Secondary structure
SS confidence
Disorder
Disorder confidence
430 440 450 460 470 480
R A F N F G I G K N M G A L G A L S V D M T Q A N S T L P D D S Q H D G Q S V R F L Y N K S L N E S G T N I Q L V G Y R
490 500 510 520 530 540
Y S T S G Y F N F A D T T Y S R M N G Y N I E T Q D G V I Q V K P K F T D Y Y N L A Y N K R G K L Q L T V T Q Q L G R T
550 560 570 580 590 600
S T L Y L S G S H Q T Y W G T S N V D E Q F Q A G L N T A F E D I N W T L S Y S L T K N A W Q K G R D Q M L A L N V N I
610 620 630 640 650 660
P F S H W L R S D S K S Q W R H A S A S Y S M S H D L N G R M T N L A G V Y G T L L E D N N L S Y S V Q T G Y A G G G D
670 680 690 700 710 720
G N S G S T G Y A T L N Y R G G Y G N A N I G Y S H S D D I K Q L Y Y G V S G G V L A H A N G V T L G Q P L N D T V V L
730 740 750 760 770 780
V K A P G A K D A K V E N Q T G V R T D W R G Y A V L P Y A T E Y R E N R V A L D T N T L A D N V D L D N A V A N V V P
790 800 810 820 830 840
T R G A I V R A E F K A R V G I K L L M T L T H N N K P L P F G A M V T S E S S Q S S G I V A D N G Q V Y L S G M P L A
850 860 870
G K V Q V K W G E E E N A H C V A N Y Q L P P E S Q Q Q L L T Q L S A E C R
Confidence Key

High(9) Low (0)

? Disordered

Alpha helix

Beta strand