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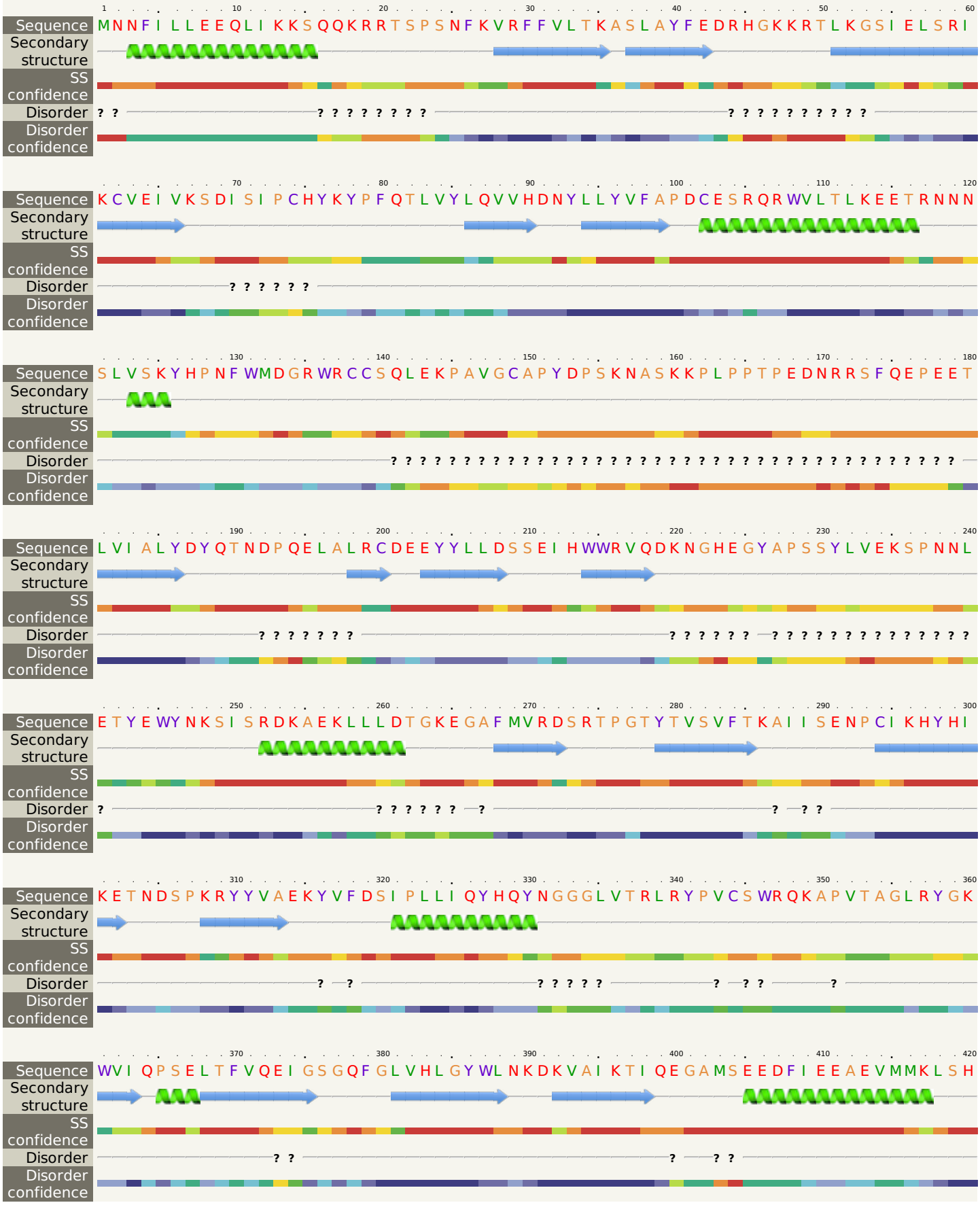
Description	Q03526
Date	Wed Jul 10 14:09:50 BST 2013
Unique Job ID	1958dd69f9d7474b

Secondary structure and disorder prediction



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