Phyre2

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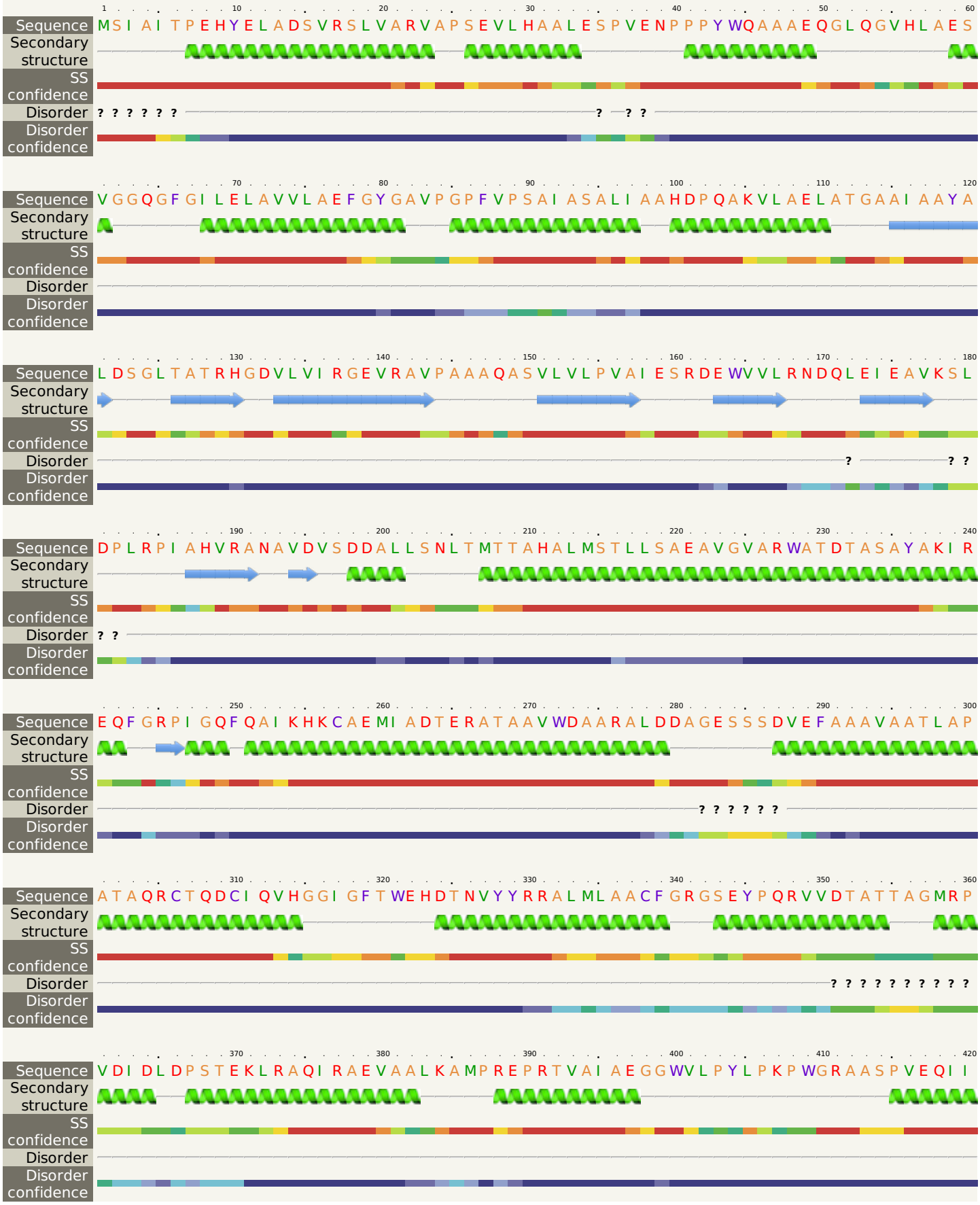
Description	RVBD0271c_(fadE6)_326266_328461
Date	Tue Jul 23 14:50:33 BST 2019
Unique Job ID	5f7bcf08513df6c3

Secondary structure and disorder prediction



Sequence
Secondary structure
SS confidence
Disorder
Disorder confidence
I A Q E F T A G R V K R P Q I A I A T W I V P S I V A F G T D N Q K Q R L L P P T F R G D I F W C Q L F S E P G A G S D
L A S L A T K A T R V D G G W R I T G Q K I W T T G A Q Y S Q W G A L L A R T D P S A P K H N G I T Y F L L D M K S E G
V Q V K P L R E L T G K E F F N T V Y L D D V F V P D E L V L G E V N R G W E V S R N T L T A E R V S I G G S D S T F L
P T L G E F V D F V R D Y R F E G Q F D Q V A R H R A G Q L I A E G H A T K L L N L R S T L L T L A G G D P M A P A A I
S K L L S M R T G Q G Y A E F A V S S F G T D A V I G D T E R L P G K W G E Y L L A S R A T T I Y G G T S E V Q L N I I
A E R L L G L P R D P
Confidence Key
High(9)
Low (0)
? Disordered (7%)
Alpha helix (55%)
Beta strand (14%)