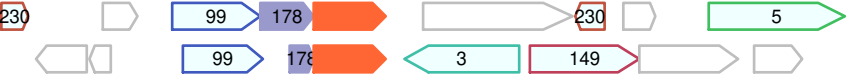


Predicted Toxin (Red Pentagon) Domain ID: D51  
-- RefSeq Protein Description: TIGR02646 family protein  
-- TADB Annotation: No Hit  
-- NCBI-CDD Annotation: HNHc superfamily  
-- HHpred(Pfam) Annotation: No Hit  
-- HHpred(PDB) Annotation: AcrIIIC3, CRISPR-associated endonuclease Cas9 (94.84%)

Predicted Antitoxin (Blue Pentagon) Domain ID: M78  
-- RefSeq Protein Description: hypothetical protein  
-- TADB Annotation: No Hit  
-- NCBI-CDD Annotation: No Hit  
-- HHpred(Pfam) Annotation: No Hit  
-- HHpred(PDB) Annotation: No Hit

*Stenotrophomonas panacihumi* (TA-Like pair: WP\_057643307.1 & WP\_146165801.1)  
*Pseudoxanthomonas indica* (TA-Like pair: WP\_188444687.1 & WP\_079723548.1)



For the description of flanking genes, please check this link: [http://130.235.240.53/NetFlax/Mini\\_pdfs/M78.txt](http://130.235.240.53/NetFlax/Mini_pdfs/M78.txt)