

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: AcrIIC3, CRISPR-associated endonuclease Cas9 (94.84%)

Predicted Antitoxin (Blue Pentagon) Domain ID: M133  
-- RefSeq Protein Description: hypothetical protein  
-- TADB Annotation: No Hit  
-- NCBI-CDD Annotation: No Hit  
-- HHpred(Pfam) Annotation: No Hit  
-- HHpred(PDB) Annotation: No Hit

Rhodopirellula maiorica SM1 (TA-Like pair: WP\_008706615.1 & WP\_008706613.1)  
Rosistilla carotiformis (TA-Like pair: WP\_145091528.1 & WP\_197452342.1)



\*\* For the description of flanking genes, please check this link: [http://130.235.240.53/NetFlax/Mini\\_pdfs/M133.txt](http://130.235.240.53/NetFlax/Mini_pdfs/M133.txt)