

Predicted Toxin (Red Pentagon) Domain ID: D57

-- RefSeq Protein Description: type II toxin-antitoxin system RelE/ParE family toxin

-- TADB Annotation: toxin relE like VCA0478; PubMed: 25897030

-- NCBI-CDD Annotation: RelE

-- HHpred(Pfam) Annotation: Toxin\_YhaV ; Toxin with endonuclease activity, of toxin-antitoxin system (97.35%)

-- HHpred(PDB) Annotation: mRNA interferase RelE (97.84%)

Predicted Antitoxin (Blue Pentagon) Domain ID: M95

-- RefSeq Protein Description: hypothetical protein

-- TADB Annotation: No Hit

-- NCBI-CDD Annotation: No Hit

-- HHpred(Pfam) Annotation: TPR\_15 ; Tetratricopeptide repeat (88.38%)

-- HHpred(PDB) Annotation: Type 4 fimbrial biogenesis protein (93.11%)

Desulfovibrio putialis DSM 16056 (TA-Like pair: WP\_027189548.1 & WP\_027189547.1)

Desulfovibrio psychrotolerans (TA-Like pair: WP\_174411108.1 & WP\_174411109.1)



\*\* For the description of flanking genes, please check this link: [http://130.235.240.53/NetFlax/Mini\\_pdfs/M95.txt](http://130.235.240.53/NetFlax/Mini_pdfs/M95.txt)