

Predicted Toxin (Red Pentagon) Domain ID: D53

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

-- TADB Annotation: Possible toxin ParE1; PubMed: 19016878

-- NCBI-CDD Annotation: ParE

-- HHpred(Pfam) Annotation: ParE_toxin ; ParE toxin of type II toxin-antitoxin system, parDE (97.21%)

-- HHpred(PDB) Annotation: Toxin protein parE-1, Antitoxin protein (98.97%)

Predicted Antitoxin (Blue Pentagon) Domain ID: M125

-- RefSeq Protein Description: hypothetical protein

-- TADB Annotation: No Hit

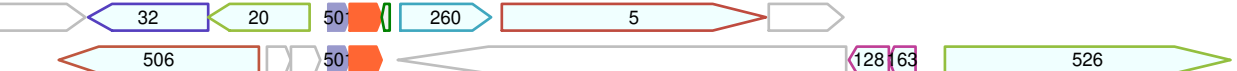
-- NCBI-CDD Annotation: No Hit

-- HHpred(Pfam) Annotation: No Hit

-- HHpred(PDB) Annotation: No Hit

Rheinheimera nanhaiensis E407 8 (TA-Like pair: WP_008224260.1 & WP_008224259.1)

Candidatus Methylobacter oryzae (TA-Like pair: WP_127029118.1 & WP_127029119.1)



** For the description of flanking genes, please check this link: http://130.235.240.53/NetFlax/Mini_pdfs/M125.txt