

Predicted Toxin (Red Pentagon) Domain ID: D128

-- RefSeq Protein Description: type II toxin-antitoxin system VapC family toxin

-- TADB Annotation: Possible toxin VapC10; PubMed: 20011113

-- NCBI-CDD Annotation: PIN_VapC4-5_FitB-like

-- HHpred(Pfam) Annotation: DUF4411 ; Domain of unknown function (DUF4411) (97.07%)

-- HHpred(PDB) Annotation: tRNA(fMet)-specific endonuclease VapC (97.99%)

Predicted Antitoxin (Blue Pentagon) Domain ID: M197

-- RefSeq Protein Description: hypothetical protein

-- TADB Annotation: No Hit

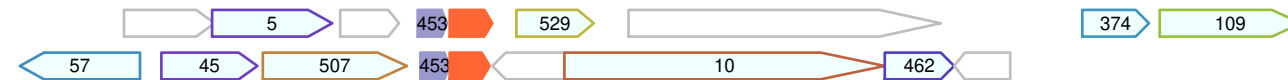
-- NCBI-CDD Annotation: No Hit

-- HHpred(Pfam) Annotation: No Hit

-- HHpred(PDB) Annotation: No Hit

Curvibacter gracilis ATCC BAA 807 (TA-Like pair: WP_027478545.1 & WP_043330983.1)

Leptothrix mobilis (TA-Like pair: WP_130480851.1 & WP_130481000.1)



** For the description of flanking genes, please check this link: http://130.235.240.53/NetFlax/Mini_pdfs/M197.txt