Predicted Toxin (Red Pentagon) Domain ID: D169

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

-- TADB Annotation: VapC PAE0151; PubMed: 22539524

-- NCBI-CDD Annotation: PIN Pae0151-like

-- HHpred(Pfam) Annotation: PIN; PIN domain (96.66%)

-- HHpred(PDB) Annotation: VapC toxin, VapB antitoxin (96.60%)

Predicted Antitoxin (Blue Pentagon) Domain ID: M228

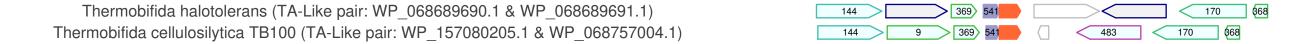
-- RefSeq Protein Description: hypothetical protein

-- TADB Annotation: No Hit

-- NCBI-CDD Annotation: No Hit

-- HHpred(Pfam) Annotation: RHH 7; Transcriptional regulator, RHH-like, CopG (60.78%)

-- HHpred(PDB) Annotation: Putative uncharacterized protein (78.42%)



For the description of flanking genes, please check this link: http://130.235.240.53/NetFlax/Mini_pdfs/M228.txt