Predicted Toxin (Red Pentagon) Domain ID: D45

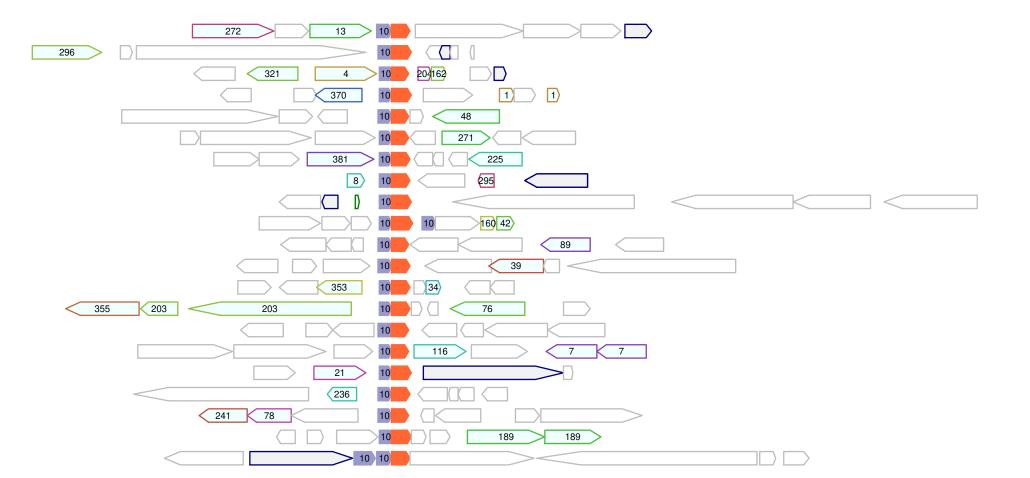
- -- RefSeg Protein Description: type II toxin-antitoxin system PemK/MazF family toxin
- -- TADB Annotation: ppGpp-regulated growth inhibitor ChpA/MazF; PubMed: 12718874
- -- NCBI-CDD Annotation: PemK toxin superfamily
- -- HHpred(Pfam) Annotation: PemK toxin; PemK-like, MazF-like toxin of type II toxin-antitoxin system (98.97%)
- -- HHpred(PDB) Annotation: PemK-like cell growth regulatory protein (98.65%)

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Caldanaerobacter subterraneus subsp tengcongensis MB4 (TA-Like pair: WP_011026253.1 & WP_031314082.1)
         Nostoc commune NIES 4072 (TA-Like pair: WP 109012346.1 & WP 109012347.1)
         Nitrosococcus halophilus Nc 4 (TA-Like pair: WP 013031413.1 & WP 013031412.1)
           Desulfosarcina widdelii (TA-Like pair: WP 155302160.1 & WP 197740481.1)
    Desulfobacterium vacuolatum DSM 3385 (TA-Like pair: WP 084068391.1 & WP 170923764.1)
              Dyadobacter soli (TA-Like pair: WP 143016787.1 & WP 090146479.1)
           Nitrospira moscoviensis (TA-Like pair: WP 053380659.1 & WP 053382105.1)
            Nitrosomonas aestuarii (TA-Like pair: WP 107796476.1 & WP 107796475.1)
     Candidatus Sulfopaludibacter sp SbA3 (TA-Like pair: WP 105497885.1 & WP 105497886.1)
   endosymbiont GvMRE of Glomus versiforme (TA-Like pair: WP 147410089.1 & WP 172575258.1)
   endosymbiont of Acanthamoeba sp UWC8 (TA-Like pair: WP 148299514.1 & WP 084173820.1)
      Candidatus Bipolaricaulis anaerobius (TA-Like pair: WP 157959537.1 & WP 122031668.1)
          Deferribacter autotrophicus (TA-Like pair: WP 149266883.1 & WP 149266955.1)
    Thermacetogenium phaeum DSM 12270 (TA-Like pair: WP 015049494.1 & WP 015049495.1)
              Dokdonia pacifica (TA-Like pair: WP 089372226.1 & WP 089372227.1)
              Rubritalea profundi (TA-Like pair: WP 105044183.1 & WP 105044873.1)
    Gloeocapsopsis crepidinum LEGE 06123 (TA-Like pair: WP 193935022.1 & WP 193935021.1)
        Alkalispirochaeta odontotermitis (TA-Like pair: WP 037561776.1 & WP 037561775.1)
       Candidatus Desulfofervidus auxilii (TA-Like pair: WP 066060368.1 & WP 066060367.1)
endosymbiont DhMRE of Dentiscutata heterogama (TA-Like pair: WP 195909698.1 & WP 195909697.1)
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endosymbiont GvMRE of Glomus versiforme (TA-Like pair: WP 147409633.1 & WP 147409634.1)

Predicted Antitoxin (Blue Pentagon) Domain ID: D102

- -- RefSeq Protein Description: ribbon-helix-helix protein, CopG family
- -- TADB Annotation: No Hit
- -- NCBI-CDD Annotation: NikR superfamily
- -- HHpred(Pfam) Annotation: RHH 8; Ribbon-Helix-Helix transcriptional regulator family (97.31%)
- -- HHpred(PDB) Annotation: mRNA interferase EndoA (99.10%)



^{**} For the description of flanking genes, please check this link: http://130.235.240.53/NetFlax/T8_pdfs/D102.txt