Predicted Antitoxin (Blue Pentagon) Domain ID: D109

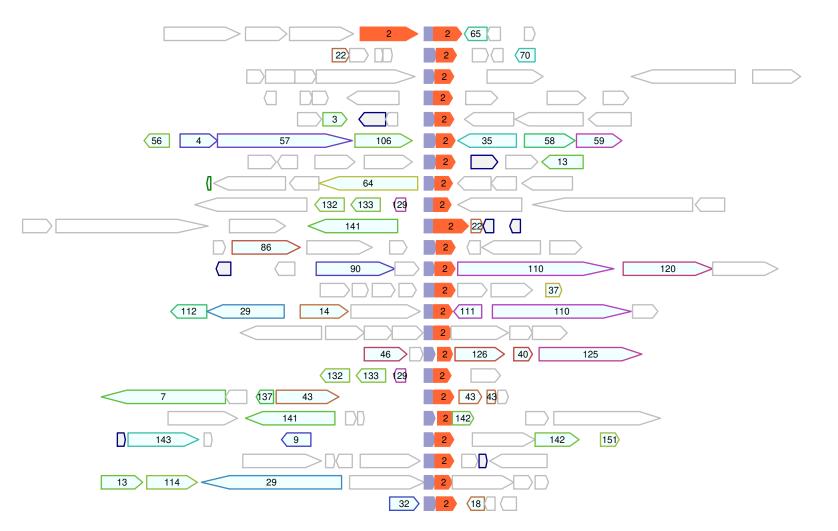
- -- RefSeq Protein Description: DUF2281 domain-containing protein
- -- TADB Annotation: No Hit
- -- NCBI-CDD Annotation: DUF2281
- -- HHpred(Pfam) Annotation: DUF2281; Protein of unknown function (DUF2281) (86.73%)
- -- HHpred(PDB) Annotation: No Hit

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Nostoc punctiforme PCC 73102 (TA-Like pair: WP_041566668.1 & WP_012413205.1)
          Salinibacter ruber M8 (TA-Like pair: WP 043551965.1 & WP 013061203.1)
 Thermodesulfatator indicus DSM 15286 (TA-Like pair: WP 013907266.1 & WP 013907265.1)
      Turneriella parva DSM 21527 (TA-Like pair: WP 041949526.1 & WP 014804586.1)
      Thioflavicoccus mobilis 8321 (TA-Like pair: WP 015279851.1 & WP 157633648.1)
Desulfotignum phosphitoxidans DSM 13687 (TA-Like pair: WP 006966977.1 & WP 006966976.1)
       Salisaeta longa DSM 21114 (TA-Like pair: WP 022834907.1 & WP 022834908.1)
        Thioalkalivibrio versutus (TA-Like pair: WP 018144158.1 & WP 018938063.1)
      Nostoc calcicola FACHB 389 (TA-Like pair: WP 073641033.1 & WP 073641024.1)
  Candidatus Thiodictyon syntrophicum (TA-Like pair: WP_100918642.1 & WP_100918643.1)
           Salinivenus iranica (TA-Like pair: WP 103038611.1 & WP 103038610.1)
      Methylobacter tundripaludum (TA-Like pair: WP 104421948.1 & WP 104421949.1)
       Coraliomargarita sinensis (TA-Like pair: WP 110132118.1 & WP 110132117.1)
        Halomonas alimentaria (TA-Like pair: WP 161432797.1 & WP 161432796.1)
        Sulfurimicrobium lacus (TA-Like pair: WP 173060420.1 & WP 173060423.1)
          Halomonas cerina (TA-Like pair: WP 183327648.1 & WP 183327646.1)
  Desmonostoc muscorum LEGE 12446 (TA-Like pair: WP 073641033.1 & WP 193923958.1)
        Thiohalocapsa halophila (TA-Like pair: WP 200234967.1 & WP 200234965.1)
        Thiohalocapsa halophila (TA-Like pair: WP 200239822.1 & WP 200239820.1)
        Thiohalocapsa halophila (TA-Like pair: WP 200239865.1 & WP 200239863.1)
   Thiohalospira halophila DSM 15071 (TA-Like pair: WP 093426914.1 & WP 093426915.1)
         Sulfurivirga caldicuralii (TA-Like pair: WP 074201307.1 & WP 074201308.1)
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Aliifodinibius sediminis (TA-Like pair: WP 142716086.1 & WP 142716087.1)

Predicted Toxin (Red Pentagon) Domain ID: D126

- -- RefSeg Protein Description: type II toxin-antitoxin system PemK/MazF family toxin
- -- TADB Annotation: No Hit
- -- NCBI-CDD Annotation: YlqF related GTPase superfamily ABC ABC ChvD superfamily
- -- HHpred(Pfam) Annotation: PRK; Phosphoribulokinase / Uridine kinase family (97.23%)
- -- HHpred(PDB) Annotation: BH3686 (97.31%)



<sup>\*\*</sup> For the description of flanking genes, please check this link: http://130.235.240.53/NetFlax/T8 pdfs/D126.txt