Predicted Antitoxin (Blue Pentagon) Domain ID: D148

- -- RefSeq Protein Description: Arc family DNA-binding protein
- -- TADB Annotation: No Hit
- -- NCBI-CDD Annotation: PRK13982 superfamily
- -- HHpred(Pfam) Annotation: RHH 7; Transcriptional regulator, RHH-like, CopG (87.89%)
- -- HHpred(PDB) Annotation: TRAFFICKING PROTEIN B, TRAFFICKING PROTEIN (96.39%) -- HHpred(PDB) Annotation: VapC toxin, VapB antitoxin (96.48%)

Predicted Toxin (Red Pentagon) Domain ID: M225

- -- RefSeq Protein Description: type II toxin-antitoxin system VapC family toxin
- -- TADB Annotation: Possible toxin VapC28 Contains PIN domain; PubMed: 20011113
- -- NCBI-CDD Annotation: PIN MtVapC28-VapC30-like
- -- HHpred(Pfam) Annotation: DUF4411; Domain of unknown function (DUF4411) (93.79%)

Methylobacter marinus A45 (TA-Like pair: WP 020160769.1 & WP 020160768.1) Crenothrix polyspora (TA-Like pair: WP 179210127.1 & WP 087146051.1) Oceanibaculum pacificum (TA-Like pair: WP 157089183.1 & WP 067551640.1) Caulobacter mirabilis (TA-Like pair: WP 099621603.1 & WP 099624164.1) Pseudopropionibacterium massiliense (TA-Like pair: WP 130875551.1 & WP 130875552.1) Thiothrix caldifontis (TA-Like pair: WP 175517889.1 & WP 093067428.1)



^{**} For the description of flanking genes, please check this link: http://130.235.240.53/NetFlax/Mini pdfs/M225.txt