

Predicted Toxin (Red Pentagon) Domain ID: D59

-- RefSeq Protein Description: putative toxin-antitoxin system toxin component, PIN family

-- TADB Annotation: No Hit

-- NCBI-CDD Annotation: PIN_SF superfamily

-- HHpred(Pfam) Annotation: PIN_9 ; PIN like domain (96.80%)

-- HHpred(PDB) Annotation: Uncharacterized protein TTHA0540 (97.56%)

Predicted Antitoxin (Blue Pentagon) Domain ID: M148

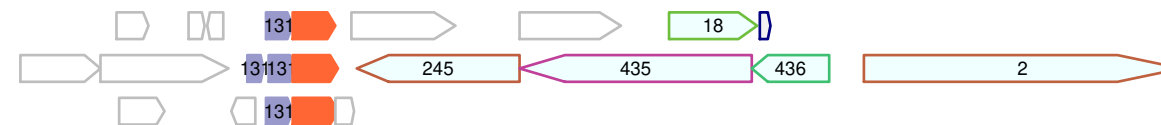
-- RefSeq Protein Description: hypothetical protein

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-- TADB Annotation: No Hit
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-- NCBI-CDD Annotation: No Hit

-- HHpred(Pfam) Annotation: No Hit

-- HHpred(PDB) Annotation: Cohesin subunit SA-1 (50.15%)



Belnapia rosea (TA-Like pair: WP_090664455.1 & WP_176849710.1)

Roseomonas stagni DSM 19981 (TA-Like pair: WP_092956603.1 & WP_175533757.1)

Paracraurococcus ruber (TA-Like pair: WP_133222111.1 & WP_133222110.1)

** For the description of flanking genes, please check this link: http://130.235.240.53/NetFlax/Mini_pdfs/M148.txt