Predicted Antitoxin (Blue Pentagon) Domain ID: D31

- -- RefSeq Protein Description: type II toxin-antitoxin system Phd/YefM family antitoxin
- -- TADB Annotation: No Hit
- -- NCBI-CDD Annotation: PhdYeFM antitox
- -- HHpred(Pfam) Annotation: PhdYeFM antitox; Antitoxin Phd YefM, type II toxin-antitoxin system (93.89%)
- -- HHpred(PDB) Annotation: Antitoxin VapB46 (96.43%)

Thiocapsa marina 5811 (TA-Like pair: WP_007191265.1 & WP_199484193.1)

Mesorhizobium alhagi CCNWXJ12 2 (TA-Like pair: WP 008837863.1 & WP 008837864.1)

Pannonibacter phragmitetus (TA-Like pair: WP_058897849.1 & WP_058897850.1)

Labrenzia aggregata (TA-Like pair: WP_077294868.1 & WP_062492111.1)

Labrys okinawensis (TA-Like pair: WP_105865423.1 & WP_105865424.1)

Mesorhizobium ephedrae (TA-Like pair: WP_106773904.1 & WP_106773903.1)

Roseiarcus fermentans (TA-Like pair: WP 113888360.1 & WP 113888361.1)

Thiorhodococcus mannitoliphagus (TA-Like pair: WP_164655757.1 & WP_164655758.1)

Nitrospirillum iridis (TA-Like pair: WP 184798285.1 & WP 184797904.1)

Seohaeicola zhoushanensis (TA-Like pair: WP_189678944.1 & WP_189678943.1)

Thiocapsa roseopersicina (TA-Like pair: WP_007191265.1 & WP_093036301.1)

** For the description of flanking genes, please check this link: http://130.235.240.53/NetFlax/T8_pdfs/D64.txt

Predicted Toxin (Red Pentagon) Domain ID: D64

- -- RefSeq Protein Description: growth inhibitor PemK
- -- TADB Annotation: No Hit
- -- NCBI-CDD Annotation: No Hit
- -- HHpred(Pfam) Annotation: PemK_toxin; PemK-like, MazF-like toxin of type II toxin-antitoxin system (90.99%)
- -- HHpred(PDB) Annotation: Antitoxin MazE, Uncharacterized protein (95.03%)

