

Predicted Toxin (Red Pentagon) Domain ID: D59
-- RefSeq Protein Description: putative toxin-antitoxin system toxin component, PIN_Tox
-- 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: M153
-- RefSeq Protein Description: hypothetical protein
-- TADB Annotation: No Hit
-- NCBI-CDD Annotation: No Hit
-- HHpred(Pfam) Annotation: Polysacc_synt_4 ; Polysaccharide biosynthesis (81.62%)
-- HHpred(PDB) Annotation: Uncharacterized protein (61.18%)

Conexibacter woesei DSM 14684 (TA-Like pair: WP_012936638.1 & WP_012936639.1)
Conexibacter arvalis (TA-Like pair: WP_183345150.1 & WP_183344747.1)



For a detailed description of flanking genes, please check this link: http://130.235.240.53/NetFlax/Mini_pdfs/M153.txt