

Predicted Toxin (Red Pentagon) Domain ID: D8
-- RefSeq Protein Description: hypothetical protein
-- TADB Annotation: ToxPhRel2
-- NCBI-CDD Annotation: NT_Pol-beta-like superfamily
-- HHpred(Pfam) Annotation: RelA_SpoT ; Region found in RelA / SpoT proteins (96.52%)
-- HHpred(PDB) Annotation: GTP pyrophosphokinase YjbM (96.86%)

Predicted Antitoxin (Blue Pentagon) Domain ID: M28
-- RefSeq Protein Description: cold shock domain-containing protein
-- TADB Annotation: No Hit
-- NCBI-CDD Annotation: S1_like superfamily
-- HHpred(Pfam) Annotation: S1-like ; S1-like (96.99%)
-- HHpred(PDB) Annotation: RNA chaperone, negative regulator of (98.62%)

Acinetobacter guerrae (TA-Like pair: WP_152876609.1 & WP_152876608.1)
Acinetobacter marinus (TA-Like pair: WP_092619006.1 & WP_092619003.1)



** For the description of flanking genes, please check this link: http://130.235.240.53/NetFlax/Mini_pdfs/M28.txt