

Predicted Antitoxin (Blue Pentagon) Domain ID: D101

-- RefSeq Protein Description: ribbon-helix-helix protein, CopG family

-- TADB Annotation: No Hit

-- NCBI-CDD Annotation: No Hit

-- HHpred(Pfam) Annotation: RHH_3 ; Ribbon-helix-helix domain (78.28%)

-- HHpred(PDB) Annotation: Probable endoribonuclease MazF7 (92.78%)

Predicted Toxin (Red Pentagon) Domain ID: M166

-- RefSeq Protein Description: hypothetical protein

-- TADB Annotation: No Hit

-- NCBI-CDD Annotation: No Hit

-- HHpred(Pfam) Annotation: No Hit

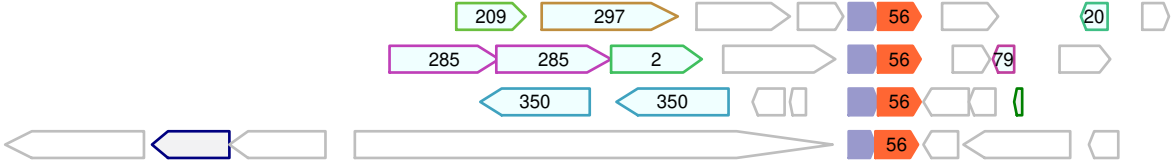
-- HHpred(PDB) Annotation: No Hit

Streptomyces klenkii (TA-Like pair: WP_120758691.1 & WP_120758692.1)

Streptomyces palmae (TA-Like pair: WP_135339453.1 & WP_135339454.1)

Streptomyces qinzhouensis (TA-Like pair: WP_146481064.1 & WP_146481063.1)

Actinomadura alba (TA-Like pair: WP_187241695.1 & WP_187241694.1)



** For the description of flanking genes, please check this link: http://130.235.240.53/NetFlax/Mini_pdfs/M166.txt