

Predicted Antitoxin (Blue Pentagon) Domain ID: D25

-- RefSeq Protein Description: ImmA/IrrE family metallo-endopeptidase

-- TADB Annotation: No Hit

-- NCBI-CDD Annotation: Peptidase_M78

-- HHpred(Pfam) Annotation: Peptidase_M78 ; IrrE N-terminal-like domain (99.11%)

-- HHpred(PDB) Annotation: IRRE PROTEIN (99.28%)

Predicted Toxin (Red Pentagon) Domain ID: M67

-- RefSeq Protein Description: hypothetical protein

-- TADB Annotation: No Hit

-- NCBI-CDD Annotation: No Hit

-- HHpred(Pfam) Annotation: DUF1918 ; Domain of unknown function (DUF1918) (67.85%)

-- HHpred(PDB) Annotation: No Hit

Dehalobacterium formicoaceticum (TA-Like pair: WP_089609003.1 & WP_089609002.1)

Agathobaculum butyriciproducens (TA-Like pair: WP_008396548.1 & WP_008396551.1)



For the description of flanking genes, please check this link: http://130.235.240.53/NetFlax/Mini_pdfs/M67.txt