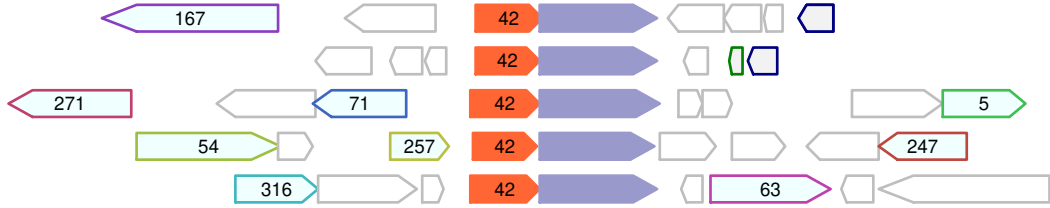


Predicted Antitoxin (Blue Pentagon) Domain ID: D25
-- RefSeq Protein Description: ImmA/IrrE family metallo-endopeptidase
-- TADB Annotation: No Hit
-- NCBI-CDD Annotation: Peptidase_M78
-- HHpred(Pfam) Annotation: Peptidase_M78 ; IrrE N-terminal-like domain (99.11%)
-- HHpred(PDB) Annotation: IRRE PROTEIN (99.28%)

Predicted Toxin (Red Pentagon) Domain ID: M62
-- RefSeq Protein Description: hypothetical protein
-- TADB Annotation: No Hit
-- NCBI-CDD Annotation: No Hit
-- HHpred(Pfam) Annotation: No Hit
-- HHpred(PDB) Annotation: No Hit

Shewanella litoralis (TA-Like pair: WP_160054222.1 & WP_160054221.1)
Ideonella dechloratans (TA-Like pair: WP_151123594.1 & WP_151123595.1)
Photobacterium phosphoreum (TA-Like pair: WP_107273565.1 & WP_107273564.1)
Acinetobacter baumannii (TA-Like pair: WP_000658761.1 & WP_001095788.1)
Alcanivorax gelatiniphagus (TA-Like pair: WP_138772642.1 & WP_138772641.1)



** For the description of flanking genes, please check this link: http://130.235.240.53/NetFlax/Mini_pdfs/M62.txt