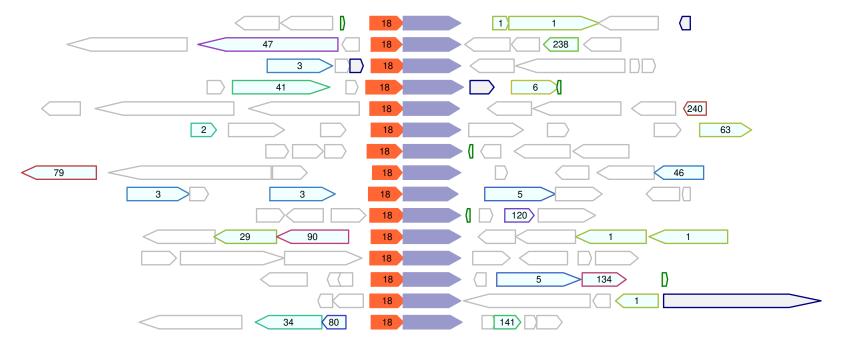
Predicted Antitoxin (Blue Pentagon) Domain ID: D25

- -- 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%)

Prevotella fusca JCM 17724 (TA-Like pair: WP_025078931.1 & WP_025078930.1)
Chryseolinea serpens (TA-Like pair: WP_073139059.1 & WP_073139057.1)
Serratia proteamaculans (TA-Like pair: WP_153857737.1 & WP_153857738.1)
Sulfitobacter indolifex (TA-Like pair: WP_132447099.1 & WP_132447101.1)
Terracidiphilus gabretensis (TA-Like pair: WP_058188294.1 & WP_157439895.1)
Lewinella xylanilytica (TA-Like pair: WP_104421149.1 & WP_170067763.1)
Gimesia alba (TA-Like pair: WP_145218147.1 & WP_145218145.1)
Pyxidicoccus trucidator (TA-Like pair: WP_164008485.1 & WP_164008483.1)
Streptacidiphilus carbonis (TA-Like pair: WP_042407513.1 & WP_052433198.1)
Hymenobacter actinosclerus (TA-Like pair: WP_143069666.1 & WP_177189648.1)
Sphingomonas ginsenosidivorax (TA-Like pair: WP_147081784.1 & WP_147081782.1)
Acinetobacter nosocomialis (TA-Like pair: WP_000951724.1 & WP_000067934.1)
Paenibacillus lentus (TA-Like pair: WP_125081097.1 & WP_125081096.1)
Dyadobacter psychrotolerans (TA-Like pair: WP_131962612.1 & WP_131962614.1)
Cytophaga hutchinsonii ATCC 33406 (TA-Like pair: WP 041932258.1 & WP 011584875.1)

Predicted Toxin (Red Pentagon) Domain ID: D56

- -- RefSeq Protein Description: hypothetical protein
- -- TADB Annotation: No Hit
- -- NCBI-CDD Annotation: No Hit
- -- HHpred(Pfam) Annotation: No Hit
- -- HHpred(PDB) Annotation: No Hit



^{**} For the description of flanking genes, please check this link: http://130.235.240.53/NetFlax/T8_pdfs/D56.txt