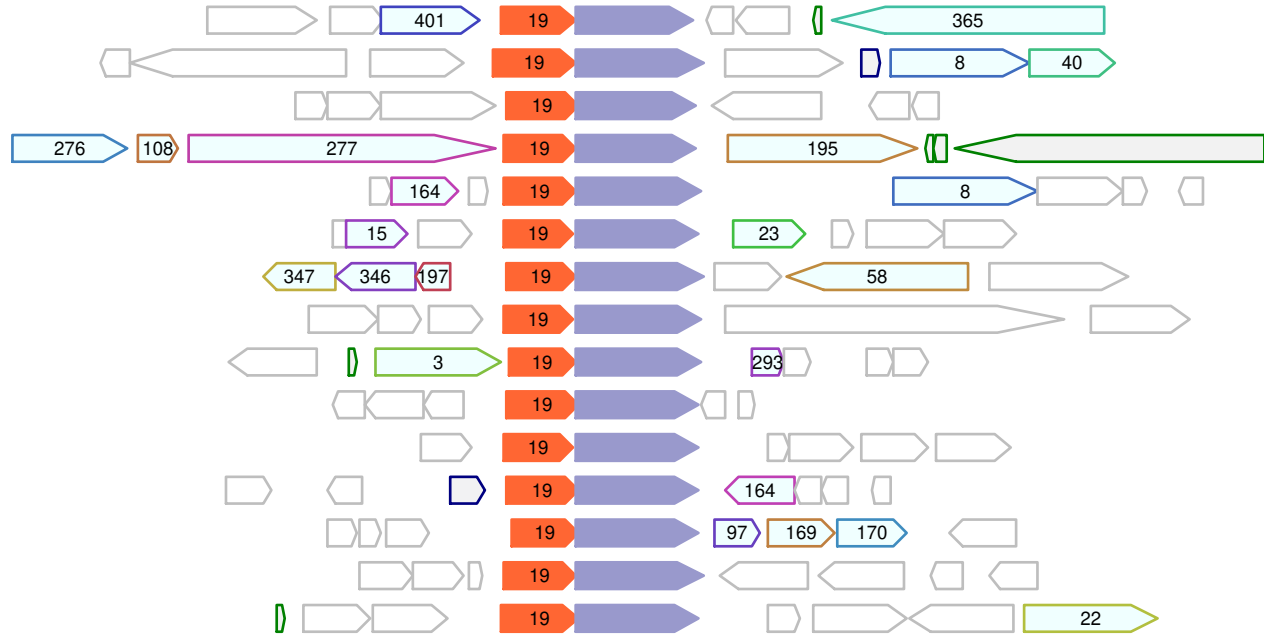


Predicted Antitoxin (Blue Pentagon) Domain ID: D25
-- RefSeq Protein Description: ImmA/IrrE family metallo-endopeptidase
-- TADB Annotation: No Hit
-- NCBI-CDD Annotation: Peptidase_M78
-- HHpred(Pfam) Annotation: Peptidase_M78 ; IrrE N-terminal-like domain (99.11%)
-- HHpred(PDB) Annotation: IRRE PROTEIN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

Predicted Toxin (Red Pentagon) Domain ID: D58
-- RefSeq Protein Description: hypothetical protein
-- TADB Annotation: No Hit
-- NCBI-CDD Annotation: No Hit
-- HHpred(Pfam) Annotation: No Hit
-- HHpred(PDB) Annotation: No Hit

Lactobacillus zhaoyuanensis (TA-Like pair: WP_125549515.1 & WP_125549516.1)
Myxococcus macrosporus DSM 14697 (TA-Like pair: WP_157757469.1 & WP_095957613.1)
Lactobacillus mulengensis (TA-Like pair: WP_125766101.1 & WP_125692940.1)
Deinococcus geothermalis DSM 11300 (TA-Like pair: WP_011531030.1 & WP_011531031.1)
Thermoanaerobacterium fracticalcicis (TA-Like pair: WP_034424705.1 & WP_034424706.1)
Kroppenstedtia guangzhouensis (TA-Like pair: WP_009710633.1 & WP_009710634.1)
Hyphomonas oceanitis SCH89 (TA-Like pair: WP_035537784.1 & WP_035537781.1)
Sphingomonas carotinifaciens (TA-Like pair: WP_149681075.1 & WP_149681074.1)
Salinicola lusitanus (TA-Like pair: WP_157958441.1 & WP_110601531.1)
Paenibacillus popilliae ATCC 14706 (TA-Like pair: WP_148277814.1 & WP_006284618.1)
Clostridioides difficile VL 0167 (TA-Like pair: WP_074138752.1 & WP_074138751.1)
Brevibacillus halotolerans (TA-Like pair: WP_181747381.1 & WP_181747382.1)
Sporomusa malonica (TA-Like pair: WP_084578464.1 & WP_084578463.1)
Bacillus cereus group sp N17 (TA-Like pair: WP_016109209.1 & WP_016109208.1)
Stenotrophomonas chelatiphaga (TA-Like pair: WP_152983335.1 & WP_083491848.1)



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