

Predicted Toxin (Red Pentagon) Domain ID: D132

-- RefSeq Protein Description: DUF3173 family protein

-- TADB Annotation: No Hit

-- NCBI-CDD Annotation: DUF3173 superfamily

-- HHpred(Pfam) Annotation: DUF3173 ; Domain of unknown function (DUF3173) (97.67%)

-- HHpred(PDB) Annotation: No Hit

Predicted Antitoxin (Blue Pentagon) Domain ID: M207

-- RefSeq Protein Description: Rep protein

-- TADB Annotation: No Hit

-- NCBI-CDD Annotation: PHA00330 superfamily

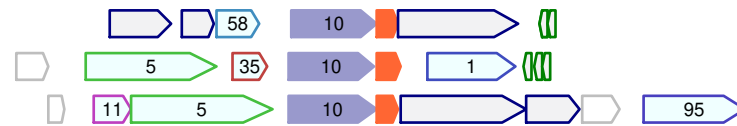
-- HHpred(Pfam) Annotation: DUF1424 ; Putative rep protein (DUF1424) (97.34%)

-- HHpred(PDB) Annotation: No Hit

Streptococcus castoreus DSM 17536 (TA-Like pair: WP_027969876.1 & WP_027969877.1)

Enterococcus rivorum (TA-Like pair: WP_069699890.1 & WP_069699891.1)

Lactococcus hircilactis (TA-Like pair: WP_153495902.1 & WP_010783659.1)



description of flanking genes, please check this link: http://130.235.240.53/NetFlax/Mini_pdfs/M207.txt