Predicted Toxin (Red Pentagon) Domain ID: D119
-- RefSeq Protein Description: type II toxin-antitoxin system YafQ family toxin
-- TADB Annotation: No Hit

-- NCBI-CDD Annotation: ParE_toxin superfamily
-- HHpred(Pfam) Annotation: ParE-like_toxin; ParE-like toxin of type II bacterial toxin-antitoxin system (67.65%)

-- HHpred(PDB) Annotation: conserved hypothtical protein HP0894 (77.01%)

Bifidobacterium vespertilionis (TA-Like pair: WP_150352967.1 & WP_150352966.1)
Bifidobacterium moukalabense (TA-Like pair: WP_034875233.1 & WP_051428909.1)

Bifidobacterium eulemuris (TA-Like pair: WP_158217217.1 & WP_158217216.1)

Bifidobacterium catulorum (TA-Like pair: WP_158275798.1 & WP_158275797.1)

36 (98 4 16 27

98 8

97 1 98 1

499 217 98 2

21 14

Predicted Antitoxin (Blue Pentagon) Domain ID: M202
-- 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/Mini_pdfs/M202.txt