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%)

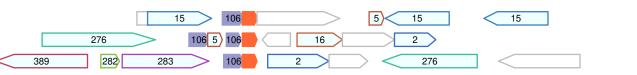
Helicobacter cetorum MIT 00 7128 (TA-Like pair: WP_014661883.1 & WP_014661884.1)

Helicobacter pylori 26695 (TA-Like pair: WP_001134157.1 & WP_000955518.1)

Helicobacter cetorum MIT 00 7128 (TA-Like pair: WP_014660237.1 & WP_014660238.1)

Predicted Antitoxin (Blue Pentagon) Domain ID: M192

- -- RefSeq Protein Description: type II toxin-antitoxin system antitoxin
- -- TADB Annotation: antitoxin DinJ HP0895; PubMed: 24060809
- -- NCBI-CDD Annotation: No Hit
- -- HHpred(Pfam) Annotation: No Hit
- -- HHpred(PDB) Annotation: Recifin modulatory peptide (52.06%)



^{**} For the description of flanking genes, please check this link: http://130.235.240.53/NetFlax/Mini pdfs/M192.txt