

ted Toxin (Red Pentagon) Domain ID: D119
Seq Protein Description: type II toxin-antitoxin system YafQ family toxin
B Annotation: No Hit
BI-CDD Annotation: ParE_toxin superfamily
red(Pfam) Annotation: ParE-like_toxin ; ParE-like toxin of type II bacterial toxin-antitoxin system (67.65%)
red(PDB) Annotation: conserved hypothtcal protein HP0894 (77.01%)

Predicted Antitoxin (Blue Pentagon) Domain ID: M218
-- RefSeq Protein Description: hypothetical protein
-- TADB Annotation: No Hit
-- NCBI-CDD Annotation: No Hit
-- HHpred(Pfam) Annotation: DUF6447 ; Family of unknown function (DUF6447) (58.61%)
-- HHpred(PDB) Annotation: Kinesin heavy chain, Microtubule-associated protein RP/EB (74.46%)

Helicobacter labetoulli (TA-Like pair: WP_121726328.1 & WP_121726329.1)
Campylobacter troglodytis (TA-Like pair: WP_142693986.1 & WP_142693987.1)



the description of flanking genes, please check this link: http://130.235.240.53/NetFlax/Mini_pdfs/M218.txt