

ed Toxin (Red Pentagon) Domain ID: D36
eq Protein Description: Txe/YoeB family addiction module toxin
B Annotation: toxin yoeB SCO2236; PubMed: 22431991
-CDD Annotation: ParE_toxin superfamily
ed(Pfam) Annotation: YafQ_toxin ; Bacterial toxin of type II toxin-antitoxin system, YafQ (97.12%)
ed(PDB) Annotation: YoeB/ParE (97.63%)

Predicted Antitoxin (Blue Pentagon) Domain ID: M84
-- RefSeq Protein Description: hypothetical protein
-- TADB Annotation: No Hit
-- NCBI-CDD Annotation: No Hit
-- HHpred(Pfam) Annotation: No Hit
-- HHpred(PDB) Annotation: No Hit

Brumimicrobium aurantiacum (TA-Like pair: WP_116879848.1 & WP_116879849.1)
aliscomenobacter hydrossis DSM 1100 (TA-Like pair: WP_013768292.1 & WP_013768293.1)



description of flanking genes, please check this link: http://130.235.240.53/NetFlax/Mini_pdfs/M84.txt