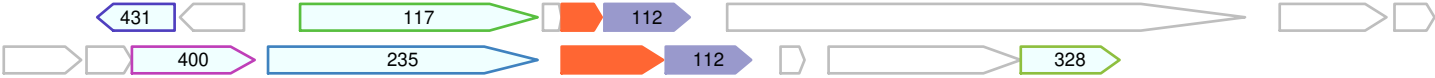


Predicted Toxin (Red Pentagon) Domain ID: D6
-- RefSeq Protein Description: type II toxin-antitoxin system death-on-curing family toxin
-- TADB Annotation: No Hit
-- NCBI-CDD Annotation: Fic superfamily_Fic
-- HHpred(Pfam) Annotation: Fic ; Fic/DOC family (87.67%)
-- HHpred(PDB) Annotation: Death on curing protein, Prevent (97.17%)

Predicted Antitoxin (Blue Pentagon) Domain ID: M29
-- RefSeq Protein Description: pyrroline-5-carboxylate reductase
-- TADB Annotation: No Hit
-- NCBI-CDD Annotation: PRK11880
-- HHpred(Pfam) Annotation: P5CR_dimer ; Pyrroline-5-carboxylate reductase dimerisation (99.52%)
-- HHpred(PDB) Annotation: Pyrroline-5-carboxylate reductase (99.78%)

Ilyomonas limi (TA-Like pair: WP_137262377.1 & WP_137262376.1)
Anaerorhabdus furcosa (TA-Like pair: WP_078711762.1 & WP_078711761.1)



** For the description of flanking genes, please check this link: http://130.235.240.53/NetFlax/Mini_pdfs/M29.txt