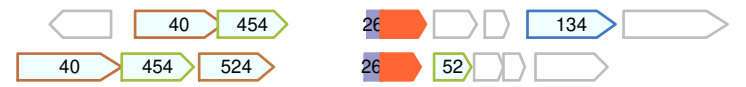


Predicted Toxin (Red Pentagon) Domain ID: D126
-- RefSeq Protein Description: type II toxin-antitoxin system PemK/MazF family toxin
-- TADB Annotation: No Hit
-- NCBI-CDD Annotation: YlqF_related_GTPase superfamily_ABC_ABC_ChvD superfamily
-- HHpred(Pfam) Annotation: PRK ; Phosphoribulokinase / Uridine kinase family (97.23%)
-- HHpred(PDB) Annotation: BH3686 (97.31%)

Predicted Antitoxin (Blue Pentagon) Domain ID: M217
-- RefSeq Protein Description: hypothetical protein
-- TADB Annotation: No Hit
-- NCBI-CDD Annotation: No Hit
-- HHpred(Pfam) Annotation: No Hit
-- HHpred(PDB) Annotation: No Hit

Moorella thermoacetica ATCC 39073 (TA-Like pair: WP_053104316.1 & WP_162490271.1)
Desulfallas alcoholivorax DSM 16058 (TA-Like pair: WP_027365107.1 & WP_169726466.1)



description of flanking genes, please check this link: http://130.235.240.53/NetFlax/Mini_pdfs/M217.txt