

Predicted Toxin (Red Pentagon) Domain ID: D126

-- RefSeq Protein Description: type II toxin-antitoxin system PemK/MazF family toxin

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-- TADB Annotation: No Hit
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-- NCBI-CDD Annotation: YlgF 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: M206

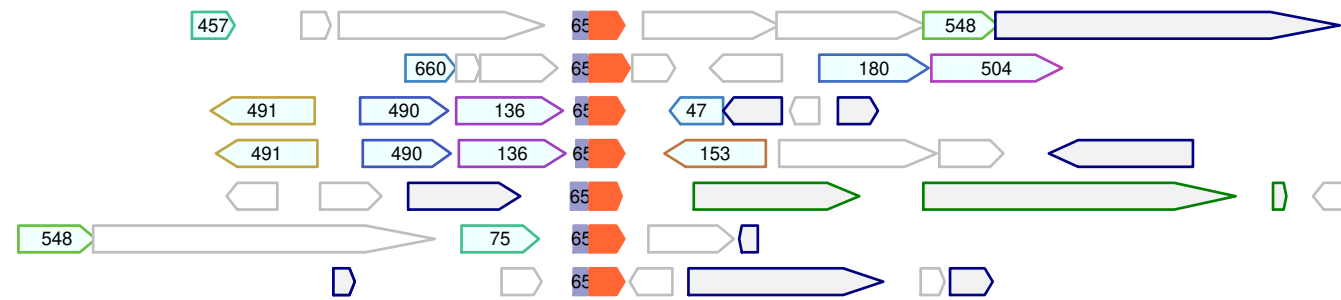
-- RefSeq Protein Description: hypothetical protein

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-- TADB Annotation: No Hit
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-- NCBI-CDD Annotation: No Hit

-- HHpred(Pfam) Annotation: DUF4258 ; Domain of unknown function (DUF4258) (61.10%)

-- HHpred(PDB) Annotation: negative strand RNA virus, polymerase (72.04%)



\*\* For the description of flanking genes, please check this link: [http://130.235.240.53/NetFlax/Mini\\_pdfs/M206.txt](http://130.235.240.53/NetFlax/Mini_pdfs/M206.txt)