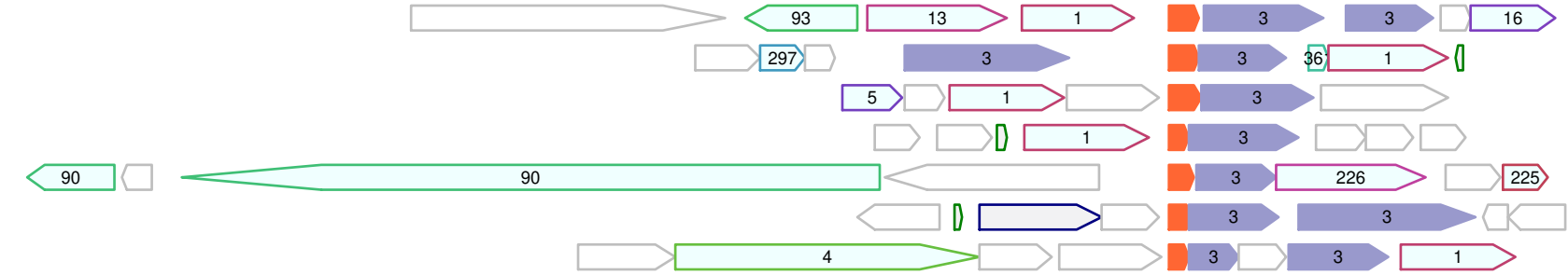


Predicted Toxin (Red Pentagon) Domain ID: D127
-- RefSeq Protein Description: helix-turn-helix domain-containing protein
-- TADB Annotation: No Hit
-- NCBI-CDD Annotation: HTH_17
-- HHpred(Pfam) Annotation: DUF3853 ; Protein of unknown function (DUF3853) (95.77%)
-- HHpred(PDB) Annotation: Excisionase from transposon Tn916 (96.33%)

Predicted Antitoxin (Blue Pentagon) Domain ID: M213
-- RefSeq Protein Description: AAA family ATPase
-- TADB Annotation: No Hit
-- NCBI-CDD Annotation: No Hit
-- HHpred(Pfam) Annotation: DUF5906 ; Family of unknown function (DUF5906) (96.71%)
-- HHpred(PDB) Annotation: Primase (98.69%)

Formosa agariphila KMM 3901 (TA-Like pair: WP_038529998.1 & WP_038529995.1)
Frankia casuarinae (TA-Like pair: WP_011435644.1 & WP_011435645.1)
Flavobacterium cyanobacteriorum (TA-Like pair: WP_094414397.1 & WP_094414395.1)
Paracoccus aminophilus JCM 7686 (TA-Like pair: WP_084621074.1 & WP_020951112.1)
Thermogutta terrifontis (TA-Like pair: WP_095415920.1 & WP_095415919.1)
Micavibrio aeruginosavorus ARL 13 (TA-Like pair: WP_041794067.1 & WP_081463081.1)
Deinococcus irradiatisoni (TA-Like pair: WP_109827990.1 & WP_109827989.1)



** For the description of flanking genes, please check this link: http://130.235.240.53/NetFlax/Mini_pdfs/M213.txt