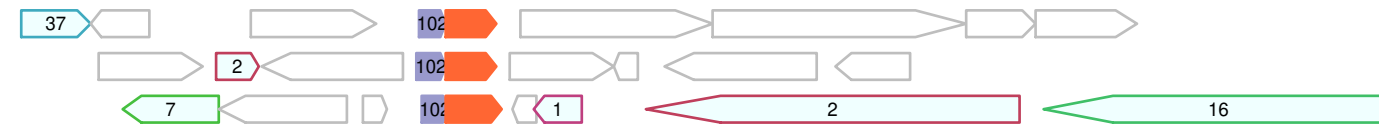


Predicted Toxin (Red Pentagon) Domain ID: D47
 -- RefSeq Protein Description: type II toxin-antitoxin system VapC family toxin
 -- TADB Annotation: No Hit
 -- NCBI-CDD Annotation: No Hit
 -- HHpred(Pfam) Annotation: No Hit
 -- HHpred(PDB) Annotation: No Hit

Predicted Antitoxin (Blue Pentagon) Domain ID: M150
 -- RefSeq Protein Description: hypothetical protein
 -- TADB Annotation: No Hit
 -- NCBI-CDD Annotation: No Hit
 -- HHpred(Pfam) Annotation: No Hit
 -- HHpred(PDB) Annotation: CdbA (54.27%)

Halomicrobium mukohataei DSM 12286 (TA-Like pair: WP_015762338.1 & WP_015762337.1)
 Halonotius roseus (TA-Like pair: WP_142442177.1 & WP_142442176.1)
 Salisaeta longa DSM 21114 (TA-Like pair: WP_028567283.1 & WP_022836913.1)



** For the description of flanking genes, please check this link: http://130.235.240.53/NetFlax/Mini_pdfs/M150.txt