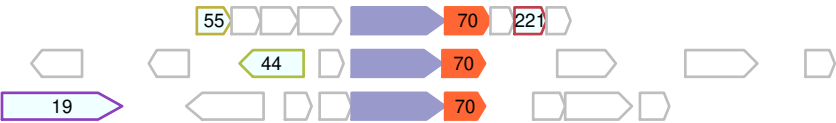


Predicted Antitoxin (Blue Pentagon) Domain ID: D34
-- RefSeq Protein Description: DNA-binding protein
-- TADB Annotation: No Hit
-- NCBI-CDD Annotation: No Hit
-- HHpred(Pfam) Annotation: DUF4248 ; Domain of unknown function (DUF4248) (61.41%)
-- HHpred(PDB) Annotation: HTH-type transcriptional regulator GlnR (71.69%)

Predicted Toxin (Red Pentagon) Domain ID: M71
-- RefSeq Protein Description: hypothetical protein SEA_YEEZY_45
-- TADB Annotation: No Hit
-- NCBI-CDD Annotation: No Hit
-- HHpred(Pfam) Annotation: No Hit
-- HHpred(PDB) Annotation: No Hit

Gordonia phage Yeezy (TA-Like pair: YP_009304373.1 & YP_009304374.1)
Mycobacterium dioxanotrophicus (TA-Like pair: WP_087075880.1 & WP_087075882.1)
Mycolicibacterium litorale (TA-Like pair: WP_185292775.1 & WP_185292774.1)



For the description of flanking genes, please check this link: http://130.235.240.53/NetFlax/Mini_pdfs/M71.txt