

Toxin (Red Pentagon) Domain ID: D127

Protein Description: helix-turn-helix domain-containing protein

Annotation: No Hit

DD Annotation: HTH_17

(Pfam) Annotation: DUF3853 ; Protein of unknown function (DUF3853) (95.77%)

(PDB) Annotation: Excisionase from transposon Tn916 (96.33%)

Predicted Antitoxin (Blue Pentagon) Domain ID: M200

Protein Description: Rha family transcriptional regulator

-- TADB Annotation: No Hit

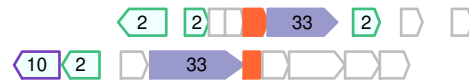
-- NCBI-CDD Annotation: Phage_pRha

(Pfam) Annotation: ORF6N ; ORF6N domain (89.35%)

(PDB) Annotation: Mitochondrial ribonuclease P protein 3 (52.29%)

rosarcina globispora (TA-Like pair: WP_053433947.1 & WP_053433948.1)

lavonifractor plautii (TA-Like pair: WP_065535737.1 & WP_065534228.1)



h of flanking genes, please check this link: http://130.235.240.53/NetFlax/Mini_pdfs/M200.txt