Predicted Antitoxin (Blue Pentagon) Domain ID: D153

-- RefSeq Protein Description: hypothetical protein

-- TADB Annotation: No Hit

-- NCBI-CDD Annotation: No Hit

-- HHpred(Pfam) Annotation: DUF2281; Protein of unknown function (DUF2281)H@ed(5%)m) Annotation: DNA_pol3_beta_3; DNA polymerase III beta subunit, C-terminal domain (98.83%)

-- NCBI-CDD Annotation: beta clamp

-- TADB Annotation: No Hit

Predicted Toxin (Red Pentagon) Domain ID: M224

-- RefSeg Protein Description: DNA polymerase III subunit beta

-- HHpred(PDB) Annotation: uncharacterized protein yiiF (67.92%) -- HHpred(PDB) Annotation: No Hit

Brachyspira murdochii DSM 12563 (TA-Like pair: WP_013113460.1 & WP_013113461.1)

Brachyspira suanatina (TA-Like pair: WP_048595210.1 & WP_014487070.1)

Brachyspira hyodysenteriae (TA-Like pair: WP_012670783.1 & WP_047113408.1)

or the description of flanking genes, please check this link: http://130.235.240.53/NetFlax/Mini_pdfs/M224.txt