

Red Pentagon) Domain ID: D41

Description: helix-turn-helix transcriptional regulator

on: No Hit

otation: CxxCG_CxxCG_HTH superfamily

Annotation: MqsA_antitoxin ; Antitoxin component of bacterial toxin-antitoxin system, MqsA (98.42%)

Annotation: Uncharacterized HTH-type transcriptional regulator ygiT (98.96%)

Predicted Antitoxin (Blue Pentagon) Domain ID: M89

-- RefSeq Protein Description: DUF4352 domain-containing protein

-- TADB Annotation: No Hit

-- NCBI-CDD Annotation: DUF4352 superfamily

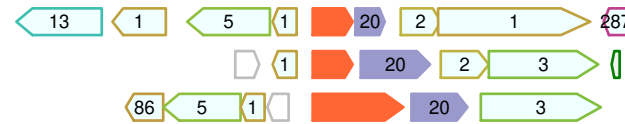
Annotation: MqsA_antitoxin ; Antitoxin component of bacterial toxin-antitoxin system, MqsA (98.42%)

Annotation: Protein - conserved hypothetical (98.70%)

entibacillus amyloliquefaciens (TA-Like pair: WP_068446673.1 & WP_156418718.1)

Salinicoccus albus DSM 19776 (TA-Like pair: WP_020006329.1 & WP_020006330.1)

Lactococcus phage phiLC3 (TA-Like pair: NP_996677.1 & NP_996676.1)



Description of flanking genes, please check this link: http://130.235.240.53/NetFlax/Mini_pdfs/M89.txt