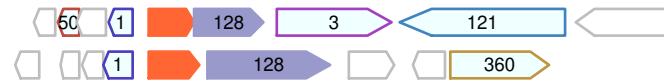


(Red Pentagon) Domain ID: D41
in Description: helix-turn-helix transcriptional regulator
ation: No Hit
nnotation: CxxCG_CxxCG_HTH superfamily
) Annotation: MqsA_antitoxin ; Antitoxin component of bacterial toxin-antitoxin system (Stam) MqsA (98.42%)
) Annotation: Uncharacterized HTH-type transcriptional regulator ygiT (98.36%)

Predicted Antitoxin (Blue Pentagon) Domain ID: M138
-- RefSeq Protein Description: DUF4428 domain-containing protein
-- TADB Annotation: No Hit
-- NCBI-CDD Annotation: SHOCT_DUF4428
Hlxines (Stam) MqsA (98.42%)
) Annotation: DUF4755 ; Domain of unknown function (DUF4755) (97.08%)
) Annotation: Anti-sigma-G factor Gin (78.55%)

arnobacterium pleistocenium FTR1 (TA-Like pair: WP_051910429.1 & WP_035051623.1)
Pyramidobacter pisolens W5455 (TA-Like pair: WP_083798230.1 & WP_009164221.1)



Description of flanking genes, please check this link: http://130.235.240.53/NetFlax/Mini_pdfs/M138.txt