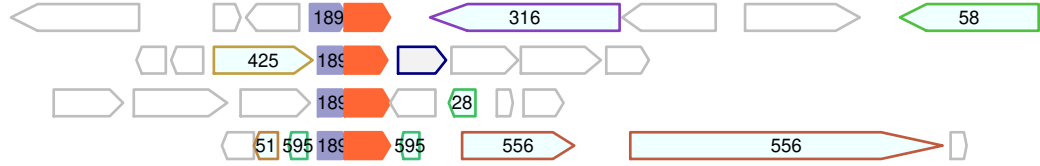


Predicted Toxin (Red Pentagon) Domain ID: D59
-- RefSeq Protein Description: putative toxin-antitoxin system toxin component, PIN family
-- TADB Annotation: No Hit
-- NCBI-CDD Annotation: PIN_SF superfamily
-- HHpred(Pfam) Annotation: PIN_9 ; PIN like domain (96.80%)
-- HHpred(PDB) Annotation: Uncharacterized protein TTHA0540 (97.56%)

Predicted Antitoxin (Blue Pentagon) Domain ID: M134
-- RefSeq Protein Description: hypothetical protein
-- TADB Annotation: No Hit
-- NCBI-CDD Annotation: No Hit
-- HHpred(Pfam) Annotation: Holin_BhlA ; BhlA holin family (51.48%)
-- HHpred(PDB) Annotation: No Hit

Haloferax gibbonsii (TA-Like pair: WP_193493623.1 & WP_193493625.1)
Haloplanus natans DSM 17983 (TA-Like pair: WP_049935889.1 & WP_049935890.1)
Halosimplex carlsbadense 2 9 1 (TA-Like pair: WP_049929868.1 & WP_006881968.1)
Natrarchaeobaculum aegyptiacum (TA-Like pair: WP_086888002.1 & WP_086888001.1)



** For the description of flanking genes, please check this link: http://130.235.240.53/NetFlax/Mini_pdfs/M134.txt