

Predicted Toxin (Red Pentagon) Domain ID: D126
-- RefSeq Protein Description: type II toxin-antitoxin system PemK/MazF family toxin
-- TADB Annotation: No Hit
-- NCBI-CDD Annotation: YlqF_related_GTPase superfamily_ABC_ABC_ChvD superfamily
-- HHpred(Pfam) Annotation: PRK ; Phosphoribulokinase / Uridine kinase family (97.23%)
-- HHpred(PDB) Annotation: BH3686 (97.31%)

Predicted Antitoxin (Blue Pentagon) Domain ID: M214
-- RefSeq Protein Description: MarR family transcriptional regulator
-- TADB Annotation: No Hit
-- NCBI-CDD Annotation: HTH_ARSR
-- HHpred(Pfam) Annotation: DsrD ; Dissimilatory sulfite reductase D (DsrD) (96.52%)
-- HHpred(PDB) Annotation: Ethanolamine utilization protein eutK (96.99%)

Natronomonas pharaonis DSM 2160 (TA-Like pair: WP_011324426.1 & WP_011324427.1)
Halorubrum halodurans (TA-Like pair: WP_094534489.1 & WP_094534487.1)
Halalkaliarchaeum desulfuricum (TA-Like pair: WP_119817940.1 & WP_119817937.1)
Natronobacterium texcoconense (TA-Like pair: WP_090385125.1 & WP_090385127.1)

