

Predicted Antitoxin (Blue Pentagon) Domain ID: D95
-- RefSeq Protein Description: site-specific integrase
-- TADB Annotation: No Hit
-- NCBI-CDD Annotation: Phage_int_SAM_3_DNA_BRE_C superfamily
-- HHpred(Pfam) Annotation: Phage_integr_3 ; Archaeal phage integrase (97.47%)
-- HHpred(PDB) Annotation: Int protein/DNA Complex (99.04%)

Predicted Toxin (Red Pentagon) Domain ID: M170
-- RefSeq Protein Description: Ltp family lipoprotein
-- TADB Annotation: No Hit
-- NCBI-CDD Annotation: rplD superfamily_SAP
-- HHpred(Pfam) Annotation: SAP_new25 ; SAP domain-containing new25 (88.16%)
-- HHpred(PDB) Annotation: N-ADA 10 (DNA METHYLPHOSPHOTRIESTER REPAIR (95.12%)

Megamonas funiformis (TA-Like pair: WP_008538044.1 & WP_008538045.1)
Staphylococcus schweitzeri (TA-Like pair: WP_047549462.1 & WP_047549465.1)
Tepidiforma bonchosmolovskayae (TA-Like pair: WP_192498061.1 & WP_158066073.1)
Jeotgalibaca porci (TA-Like pair: WP_166063663.1 & WP_166063662.1)

