

Predicted Toxin (Red Pentagon) Domain ID: D169
-- RefSeq Protein Description: type II toxin-antitoxin system VapC family toxin
-- TADB Annotation: VapC PAE0151; PubMed: 22539524
-- NCBI-CDD Annotation: PIN_Pae0151-like
-- HHpred(Pfam) Annotation: PIN ; PIN domain (96.66%)
-- HHpred(PDB) Annotation: VapC toxin, VapB antitoxin (96.60%)

Predicted Antitoxin (Blue Pentagon) Domain ID: M228
-- RefSeq Protein Description: hypothetical protein
-- TADB Annotation: No Hit
-- NCBI-CDD Annotation: No Hit
-- HHpred(Pfam) Annotation: RHH_7 ; Transcriptional regulator, RHH-like, CopG (60.78%)
-- HHpred(PDB) Annotation: Putative uncharacterized protein (78.42%)

Thermobifida halotolerans (TA-Like pair: WP_068689690.1 & WP_068689691.1)
Thermobifida cellulosilytica TB100 (TA-Like pair: WP_157080205.1 & WP_068757004.1)

