

Predicted Toxin (Red Pentagon) Domain ID: D127  
-- RefSeq Protein Description: helix-turn-helix domain-containing protein  
-- TADB Annotation: No Hit  
-- NCBI-CDD Annotation: HTH\_17  
-- HHpred(Pfam) Annotation: DUF3853 ; Protein of unknown function (DUF3853) (95.77%)  
-- HHpred(PDB) Annotation: Excisionase from transposon Tn916 (96.33%)

Predicted Antitoxin (Blue Pentagon) Domain ID: M184  
-- RefSeq Protein Description: hypothetical protein  
-- TADB Annotation: No Hit  
-- NCBI-CDD Annotation: No Hit  
-- HHpred(Pfam) Annotation: No Hit  
-- HHpred(PDB) Annotation: UNCHARACTERIZED PROTEIN 114 (53.85%)

Pseudodesulfovibrio profundus (TA-Like pair: WP\_097013702.1 & WP\_097011833.1)  
Desulfovibrio senegalensis (TA-Like pair: WP\_151151597.1 & WP\_151151596.1)

