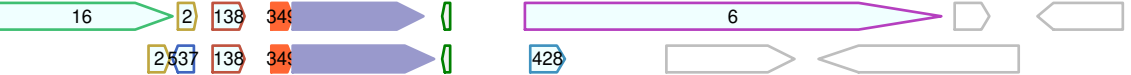


Predicted Antitoxin (Blue Pentagon) Domain ID: D95
-- RefSeq Protein Description: site-specific integrase
-- TADB Annotation: No Hit
-- NCBI-CDD Annotation: Phage_int_SAM_3_DNA_BRE_C superfamily
-- HHpred(Pfam) Annotation: Phage_integr_3 ; Archaeal phage integrase (97.47%)
-- HHpred(PDB) Annotation: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

Frankia torreyi (TA-Like pair: WP_044886766.1 & WP_044886736.1)
Frankia saprophytica (TA-Like pair: WP_007508760.1 & WP_007508759.1)

Predicted Toxin (Red Pentagon) Domain ID: M162
-- RefSeq Protein Description: hypothetical protein
-- TADB Annotation: No Hit
-- NCBI-CDD Annotation: No Hit
-- HHpred(Pfam) Annotation: Thioredoxin_4 ; Thioredoxin (80.76%)
-- HHpred(PDB) Annotation: Hypothetical protein (82.63%)



** For the description of flanking genes, please check this link: http://130.235.240.53/NetFlax/Mini_pdfs/M162.txt