

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 (97.44%)

-- HHpred(PDB) Annotation: Int protein/DNA Complex (99.04%)

Predicted Toxin (Red Pentagon) Domain ID: M177

-- RefSeq Protein Description: hypothetical protein

-- TADB Annotation: No Hit

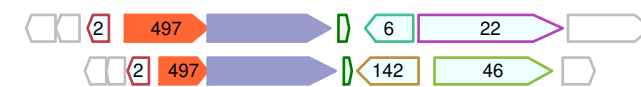
-- NCBI-CDD Annotation: No Hit

-- HHpred(Pfam) Annotation: MobC\_2 ; MobC-like protein (78.77%)

-- HHpred(PDB) Annotation: Bacterial mobilisation protein (MobC) (62.94%)

Jiella endophytica (TA-Like pair: WP\_134760235.1 & WP\_134760237.1)

Mesorhizobium prunaredense (TA-Like pair: WP\_077371409.1 & WP\_077371411.1)



cription of flanking genes, please check this link: [http://130.235.240.53/NetFlax/Mini\\_pdfs/M177.txt](http://130.235.240.53/NetFlax/Mini_pdfs/M177.txt)