

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: M175 -- RefSeq Protein Description: hypothetical protein -- TADB Annotation: No Hit -- NCBI-CDD Annotation: No Hit -- HHpred(Pfam) Annotation: DUF6076 ; Family of unknown function (DUF6076) (92.91%) -- HHpred(PDB) Annotation: No Hit
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Virgibacillus proomii (TA-Like pair: WP_077319073.1 & WP_077319072.1)
 Sporolactobacillus putidus (TA-Like pair: WP_188803388.1 & WP_188803386.1)

