

Antitoxin (Blue Pentagon) Domain ID: D26

q Protein Description: ribbon-helix-helix protein, CopG family

Annotation: No Hit

CDD Annotation: No Hit

d(Pfam) Annotation: RHH\_7 ; Transcriptional regulator, RHH-like, CopG (83-27%)

d(PDB) Annotation: MNT REPRESSOR MUTANT WITH C-TERMINAL (PDB) Annotation: No Hit

Predicted Toxin (Red Pentagon) Domain ID: M48

-- RefSeq Protein Description: hypothetical protein SEA\_HEDWIG\_39

-- TADB Annotation: No Hit

-- NCBI-CDD Annotation: No Hit

copG (83-27%)

copG (83-27%)

Gordonia phage Hedwig (TA-Like pair: YP\_009289848.1 & YP\_009289847.1)

Gordonia oryzae (TA-Like pair: WP\_123925598.1 & WP\_123925596.1)



on of flanking genes, please check this link: [http://130.235.240.53/NetFlax/Mini\\_pdfs/M48.txt](http://130.235.240.53/NetFlax/Mini_pdfs/M48.txt)