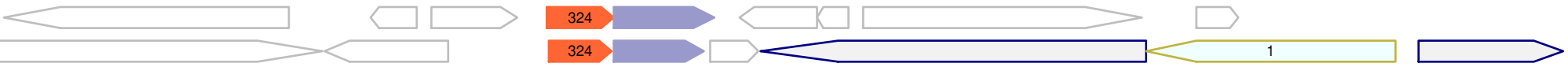Predicted Antitoxin (Blue Pentagon) Domain ID: D25
-- RefSeq Protein Description: ImmA/IrrE family metallo-endopeptidase
-- TADB Annotation: No Hit
-- NCBI-CDD Annotation: Peptidase_M78
-- HHpred(Pfam) Annotation: Peptidase_M78 ; IrrE N-terminal-like domain (99.11%)
-- HHpred(PDB) Annotation: IRRE PROTEIN (99.28%)

Predicted Toxin (Red Pentagon) Domain ID: M51
-- RefSeq Protein Description: hypothetical protein
-- TADB Annotation: No Hit
-- NCBI-CDD Annotation: No Hit
-- HHpred(Pfam) Annotation: No Hit
-- HHpred(PDB) Annotation: No Hit

Actinophytocola xanthii (TA-Like pair: WP_075125644.1 & WP_143229456.1)
Corynebacterium gottingense (TA-Like pair: WP_147453267.1 & WP_122085995.1)

** For the description of flanking genes, please check this link: http://130.235.240.53/NetFlax/Mini_pdfs/M51.txt