

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: M50

-- RefSeq Protein Description: helix-turn-helix transcriptional regulator

-- TADB Annotation: No Hit

-- NCBI-CDD Annotation: No Hit

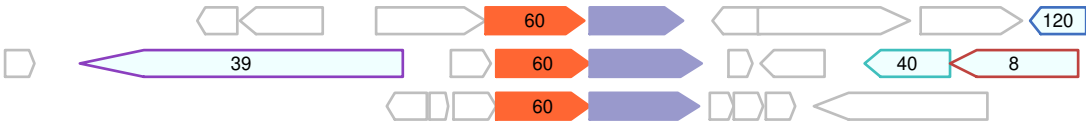
-- HHpred(Pfam) Annotation: No Hit

-- HHpred(PDB) Annotation: No Hit

*Bifidobacterium anseris* (TA-Like pair: WP\_133120871.1 & WP\_026645311.1)

*Microbacterium immunditiarum* (TA-Like pair: WP\_179486474.1 & WP\_179486475.1)

*Tessaracoccus flavescens* (TA-Like pair: WP\_077351602.1 & WP\_152024736.1)



\*\* For the description of flanking genes, please check this link: [http://130.235.240.53/NetFlax/Mini\\_pdfs/M50.txt](http://130.235.240.53/NetFlax/Mini_pdfs/M50.txt)