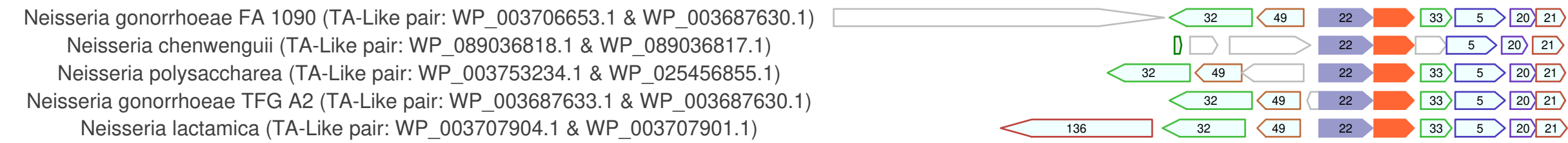


Predicted Toxin (Red Pentagon) Domain ID: D8
-- RefSeq Protein Description: hypothetical protein
-- TADB Annotation: ToxPhRel2
-- NCBI-CDD Annotation: NT_Pol-beta-like superfamily
-- HHpred(Pfam) Annotation: RelA_SpoT ; Region found in RelA / SpoT proteins (96.52%)
-- HHpred(PDB) Annotation: GTP pyrophosphokinase YjbM (96.86%)

Predicted Antitoxin (Blue Pentagon) Domain ID: M25
-- RefSeq Protein Description: 16S rRNA (adenine(1518)-N(6)/adenine(1519)-N(6))-dimethyltransferase RsmA
-- TADB Annotation: No Hit
-- NCBI-CDD Annotation: ksgA
-- HHpred(Pfam) Annotation: RrnaAD ; Ribosomal RNA adenine dimethylase (99.01%)
-- HHpred(PDB) Annotation: Ribosomal RNA small subunit methyltransferase (99.20%)



** For the description of flanking genes, please check this link: http://130.235.240.53/NetFlax/Mini_pdfs/M25.txt