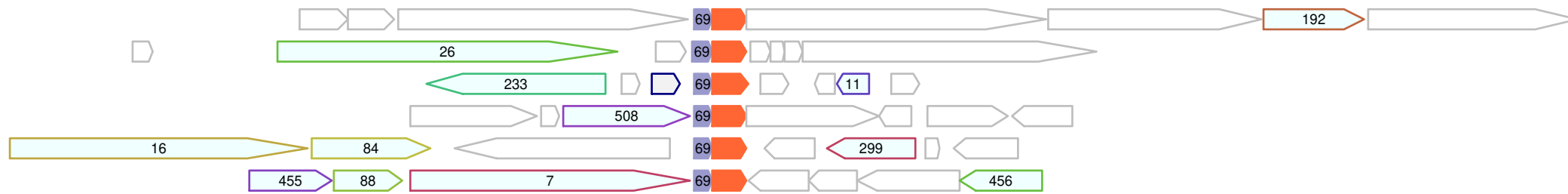
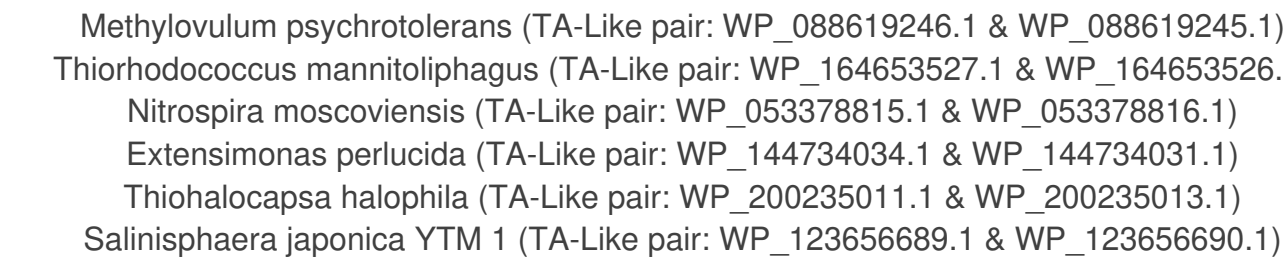


Predicted Toxin (Red Pentagon) Domain ID: D169  
 -- RefSeq Protein Description: type II toxin-antitoxin system VapC family toxin  
 -- TADB Annotation: VapC PAE0151; PubMed: 22539524  
 -- NCBI-CDD Annotation: PIN\_Pae0151-like  
 -- HHpred(Pfam) Annotation: PIN ; PIN domain (96.66%)  
 -- HHpred(PDB) Annotation: VapC toxin, VapB antitoxin (96.60%)

Predicted Antitoxin (Blue Pentagon) Domain ID: M231  
 -- RefSeq Protein Description: hypothetical protein  
 -- TADB Annotation: No Hit  
 -- NCBI-CDD Annotation: No Hit  
 -- HHpred(Pfam) Annotation: RHH\_6 ; Ribbon-helix-helix domain (93.67%)  
 -- HHpred(PDB) Annotation: Putative uncharacterized protein (93.79%)



\*\* For the description of flanking genes, please check this link: [http://130.235.240.53/NetFlax/Mini\\_pdfs/M231.tx](http://130.235.240.53/NetFlax/Mini_pdfs/M231.tx)