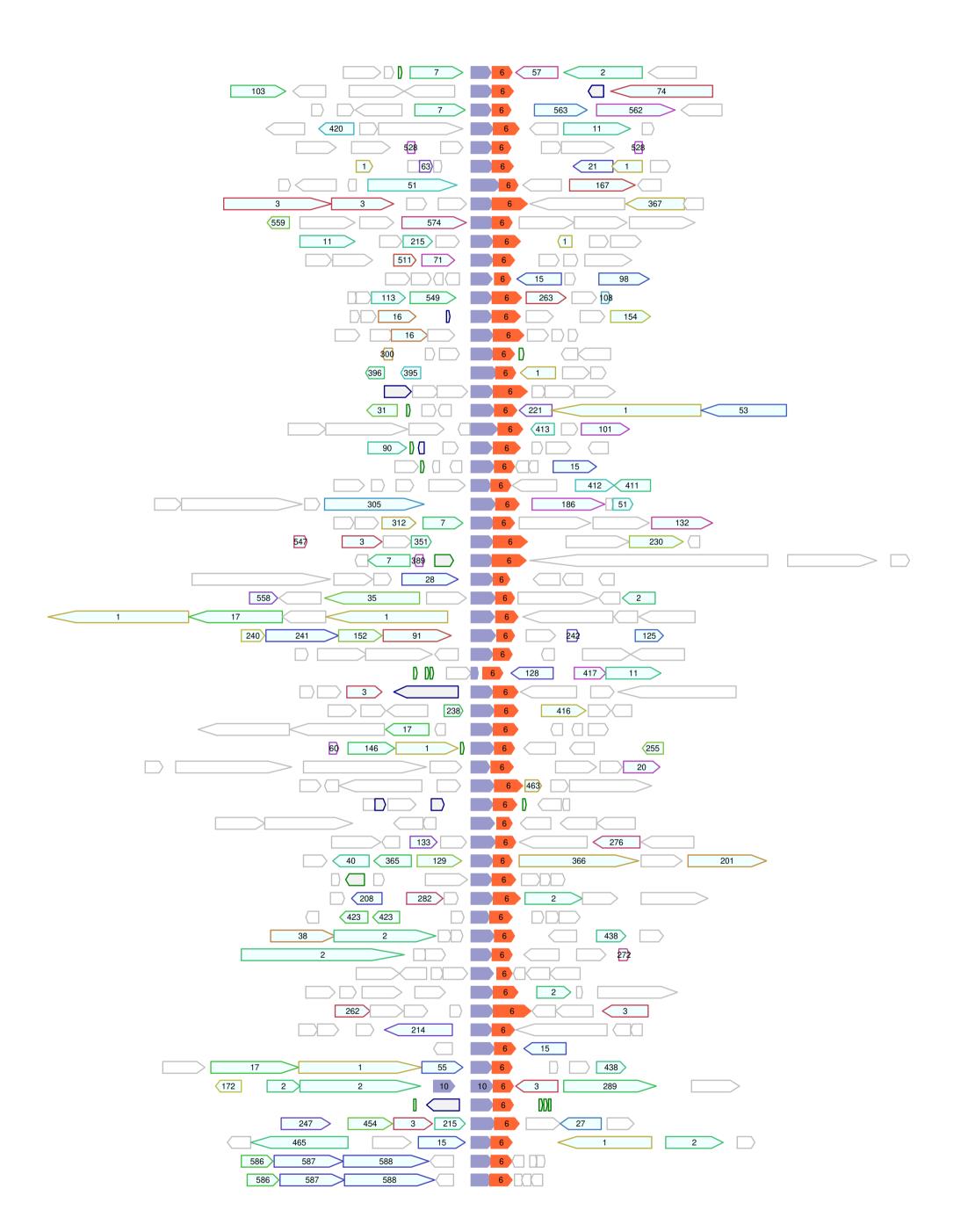
## Antitoxin (Blue Pentagon) Domain Name: PanA

Aminobacterium colombiense DSM 12261 (TA-Like pair: WP 013048536.1 & WP 013048535.1) Actinocorallia herbida (TA-Like pair: WP\_123667508.1 & WP\_148086129.1) Saccharospirillum salsuginis (TA-Like pair: WP\_189608847.1 & WP\_189609096.1) Companilactobacillus heilongjiangensis (TA-Like pair: WP\_048698801.1 & WP\_048698805.1) Spirosoma fluviale (TA-Like pair: WP 097125246.1 & WP 097125245.1) Lactobacillus paragasseri (TA-Like pair: WP 144842118.1 & WP 144842119.1) Kamptonema formosum PCC 6407 (TA-Like pair: WP 007353925.1 & WP 007353924.1) Phocaeicola salanitronis DSM 18170 (TA-Like pair: WP\_004293873.1 & WP\_004293874.1) Salinarimonas ramus (TA-Like pair: WP 188914613.1 & WP 188914614.1) Companilactobacillus huachuanensis (TA-Like pair: WP\_137610633.1 & WP\_137610634.1) Clostridium cadaveris NLAE zl G419 (TA-Like pair: WP\_074844973.1 & WP\_074844970.1) Flavobacterium johnsoniae UW101 (TA-Like pair: WP 044047495.1 & WP 085950303.1) Cohnella xylanilytica (TA-Like pair: WP 185133921.1 & WP 185134006.1) Clostridium intestinale (TA-Like pair: WP\_181603774.1 & WP\_181601387.1) Paenibacillus brasilensis (TA-Like pair: WP 152382191.1 & WP 152380867.1) Solimonas variicoloris DSM 15731 (TA-Like pair: WP 051101888.1 & WP 154654880.1) Candidatus Liberibacter solanacearum (TA-Like pair: WP\_103847170.1 & WP\_103847169.1) Shewanella inventionis (TA-Like pair: WP 188740533.1 & WP 188740532.1) Nafulsella turpanensis ZLM 10 (TA-Like pair: WP 017732543.1 & WP 017732542.1) Flavobacterium tibetense (TA-Like pair: WP 113988438.1 & WP 113988437.1) Trichococcus patagoniensis (TA-Like pair: WP 108031431.1 & WP 108031432.1) Chryseobacterium oleae (TA-Like pair: WP 090023811.1 & WP 090023812.1) Pedobacter namyangjuensis (TA-Like pair: WP 113651985.1 & WP 113651984.1) Prevotella melaninogenica (TA-Like pair: WP\_004359498.1 & WP\_004359499.1) Lysinibacillus massiliensis 4400831 CIP 108448 CCUG 49529 (TA-Like pair: WP\_036177549.1 & WP\_052126195.1) Vibrio palustris (TA-Like pair: WP 077311932.1 & WP 077311930.1) Chryseobacterium geocarposphaerae (TA-Like pair: WP\_100377045.1 & WP\_100376784.1) Candidatus Phytoplasma solani (TA-Like pair: WP 122225589.1 & WP 023161225.1) Salinarimonas ramus (TA-Like pair: WP 188911154.1 & WP 188911152.1) Hymenobacter psychrotolerans DSM 18569 (TA-Like pair: WP\_073282534.1 & WP\_139252166.1) Spiroplasma citri (TA-Like pair: WP 171283747.1 & WP 071937179.1) Alteripontixanthobacter maritimus (TA-Like pair: WP\_115365767.1 & WP\_115365768.1) Legionella busanensis (TA-Like pair: WP 115332045.1 & WP 115332044.1) Chryseobacterium frigidisoli (TA-Like pair: WP\_090080310.1 & WP\_090080309.1) Muricauda pacifica (TA-Like pair: WP\_106143626.1 & WP\_106143627.1) Massilia aurea (TA-Like pair: WP\_123070264.1 & WP\_148044035.1) Lactobacillus japonicus (TA-Like pair: WP 054396215.1 & WP 054396213.1) Lactobacillus nodensis DSM 19682 JCM 14932 NBRC 107160 (TA-Like pair: WP\_082609511.1 & WP\_025023194.1) Dyadobacter fermentans DSM 18053 (TA-Like pair: WP\_015810302.1 & WP\_015810301.1) Enterococcus plantarum (TA-Like pair: WP 111248349.1 & WP 111248348.1) Novosphingobium rosa NBRC 15208 (TA-Like pair: WP\_084455213.1 & WP\_068083796.1) Nitrolancea hollandica Lb (TA-Like pair: WP\_008476557.1 & WP\_008476559.1) Roseateles aquatilis (TA-Like pair: WP 088385116.1 & WP 088385117.1) endosymbiont GvMRE of Glomus versiforme (TA-Like pair: WP\_147410129.1 & WP\_147410128.1) Megasphaera elsdenii DSM 20460 (TA-Like pair: WP\_014015785.1 & WP\_014015786.1) Candidatus Phytoplasma ziziphi (TA-Like pair: WP 121463974.1 & WP 121463973.1) Nonlabens ponticola (TA-Like pair: WP\_126447207.1 & WP\_126447205.1) Sphaerospermopsis aphanizomenoides BCCUSP55 (TA-Like pair: WP\_200327304.1 & WP\_200327305.1) Rhizobium acidisoli (TA-Like pair: WP 054182632.1 & WP 054182752.1) Candidatus Phytoplasma pruni (TA-Like pair: WP\_178733983.1 & WP\_178733982.1) Roseateles aquatilis (TA-Like pair: WP 088388490.1 & WP 141101007.1) Acidobacterium capsulatum ATCC 51196 (TA-Like pair: WP 041839411.1 & WP 148215087.1) Bacteroides uniformis (TA-Like pair: WP 196031200.1 & WP 196031201.1) Algibacter wandonensis (TA-Like pair: WP\_133965460.1 & WP\_133965462.1) Lactobacillus shenzhenensis LY 73 (TA-Like pair: WP\_022529529.1 & WP\_155828593.1) Lactobacillus melliventris (TA-Like pair: WP\_187156394.1 & WP\_187156395.1) Chitinophaga skermanii (TA-Like pair: WP\_111597956.1 & WP\_111597957.1) Occallatibacter savannae (TA-Like pair: WP 162601179.1 & WP 109485033.1) Escherichia coli MOD1 EC5558 (TA-Like pair: WP 089622957.1 & WP 077635758.1) Salmonella phage vB SemP Emek (TA-Like pair: YP\_006560565.1 & YP\_006560566.1)

- Predicted Toxin (Red Pentagon) Domain ID: D3
  -- RefSeg Protein Description: hypothetical protein
- -- TADB Annotation: No Hit
- -- NCBI-CDD Annotation: No Hit
- -- HHpred(Pfam) Annotation: No Hit
- -- HHpred(PDB) Annotation: FACT complex subunit SSRP1 (65.51%)



<sup>\*\*</sup> For the description of flanking genes, please check this link: http://130.235.240.53/NetFlax/T8 pdfs/D3.txt