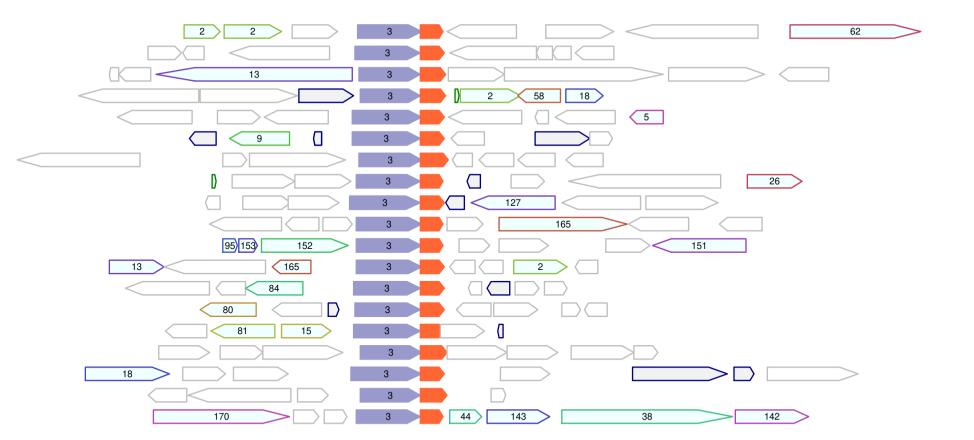
Predicted Toxin (Red Pentagon) Domain ID: D3

- -- RefSeq 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%)

Saprospira grandis str Lewin (TA-Like pair: WP 014373231.1 & WP 014373230.1) Galbibacter marinus (TA-Like pair: WP 008991521.1 & WP 008991522.1) Treponema vincentii F0403 (TA-Like pair: WP 016519637.1 & WP 016519638.1) Treponema medium ATCC 700293 (TA-Like pair: WP 016522799.1 & WP 016522800.1) Endozoicomonas numazuensis (TA-Like pair: WP 161787486.1 & WP 034842319.1) Gallibacterium genomosp 3 (TA-Like pair: WP 065237658.1 & WP 065237657.1) Pelagibaculum spongiae (TA-Like pair: WP 158527317.1 & WP 116687493.1) Pedobacter namyangiuensis (TA-Like pair: WP 113653598.1 & WP 113653597.1) Marinomonas pollencensis (TA-Like pair: WP 115899184.1 & WP 115899183.1) Chitinophaga barathri (TA-Like pair: WP 120514119.1 & WP 120514118.1) Chryseobacterium echinoideorum (TA-Like pair: WP 144281855.1 & WP 144281856.1) Arachidicoccus ginsenosidivorans (TA-Like pair: WP 158639534.1 & WP 146782772.1) Neptunomonas concharum (TA-Like pair: WP 170221765.1 & WP 138987103.1) Acinetobacter guerrae (TA-Like pair: WP 152878174.1 & WP 152878172.1) Advenella faeciporci (TA-Like pair: WP 189386082.1 & WP 189386081.1) Treponema zioleckii (TA-Like pair: WP 191017291.1 & WP 191017289.1) Williamwhitmania taraxaci (TA-Like pair: WP 092440549.1 & WP 092440547.1) Hydrogenimonas thermophila (TA-Like pair: WP 092913906.1 & WP 092913907.1) Chitinophaga jiangningensis (TA-Like pair: WP 073077276.1 & WP 073077275.1)

Predicted Antitoxin (Blue Pentagon) Domain ID: D29

- -- RefSea Protein Description: ATP-binding protein
- -- TADB Annotation: No Hit
- -- NCBI-CDD Annotation: AAA 21
- -- HHpred(Pfam) Annotation: AAA 21; AAA domain, putative AbiEii toxin, Type IV TA system (98.89%)
- -- HHpred(PDB) Annotation: DNA double-strand break repair rad50 (95.83%)



^{**} For the description of flanking genes, please check this link: http://130.235.240.53/NetFlax/T8 pdfs/D29.txt