

CUJ05009.1 CYSS0100001.1:2603727-2608317 <i>Janthinobacterium</i> sp. CG23_2	DUF3349 ? RHS_repeat RHS_repeat RHS_repeat RHS_repeat RHS_repeat RHS Tox-ParBL1 ? ?
EBJ7918586.1 AAFTVL010000172.1:191-1409 <i>Salmonella enterica</i>	RHS Tox-ParBL1
EDS3300093.1 ANXB010000038.1:1-9825 <i>Salmonella enterica</i> subsp. <i>enterica</i> serovar <i>Umbadah</i>	DUF2169 ? RHS Tox-ParBL1 ? ?
ELP24735.1 ANKX01000052.1:99-1643 <i>Pantoea agglomerans</i> 299R	RHS Tox-ParBL1 ?
HCA06239.1 DOTS01000005.1:347274-356654 <i>Chryseobacterium</i> sp.	Peptidase_S41 ? DUF6531 RHS RHS RHS ParBC ? Phage_GPD Phage_base_V
MBC1768434.1 JAARUX010000003.1:271580-274539 <i>Listeria welshimeri</i>	DUF3130 ? WXG Tox-ParBL1 ? PTS_2-RNA
MBE7709502.1 SUTY01000018.1:21461-25949 <i>Cyanobacteria bacterium</i> SIG32	PTS_2-RNA ? Tox-ParBL1 ? PBP_dimer Transpeptidase
MBF0298341.1 JADGAM010000036.1:1472-11730 <i>Oligoflexia bacterium</i>	SWACOS STY-kinase ? FiLH Tox-ParBL1 ? SSP160
MBN2176102.1 JAFGBM010000001.1:75904-77378 <i>Demequinaceae bacterium</i>	PTS_2-RNA ? Tox-ParBL1
MBP5433123.1 JAGCWY010000106.1:55810-71054 <i>Ruminococcus</i> sp.	MscS_TM ? IG DUF6531 RHS_repeat RHS_repeat RHS_repeat RHS RHS_repeat RHS RHS_repeat RHS Tox-ParBL1 ? Lectin_C Dockerin_1
MBQ3302155.1 JAFBUB010000470.1:1-1447 <i>Eggerthellaceae bacterium</i>	DUF4417 Tox-ParBL1 ? RdRP
MBR1751139.1 JAFVAH010000112.1:25874-27961 <i>Ruminococcus</i> sp.	? ? HINT Tox-ParBL1 Deaminase GST_N_3
MQY12446.1 WEGJ01000006.1:203172-211505 <i>Streptomyces smaragdinus</i>	CoaE ? T7SS_signal DUF6531 RHS RHS RHS Tox-ParBL1 PAC2 Amidase_2
MUZ283934.1 WPH001000010.1:32835-35070 <i>Agrobacterium vitis</i>	RHS ? Tox-ParBL1 RHS ?
NJM80722.1 JAURU010000340.1:2229-4141 <i>Flavobacterium</i> sp.	PTS_2-RNA ? RHS Tox-ParBL1
NQZ38483.1 JABSPD010000039.1:643-2271 <i>Crocinitomicaceae bacterium</i>	RHS Tox-ParBL1 ? Tox-ART-HYD2
NUU69037.1 JABWC010000018.1:60094-64163 <i>Enterobacteriaceae bacterium</i> BIT-123	RHS ? RHS Tox-ParBL1 ? Imm50
OPX41871.1 MZGX01000048.1:28-2946 <i>Ruminiclostridium hungatei</i>	RHS Tox-ParBL1 ? RHS
PIT20138.1 MDVE01000059.1:1-5322 <i>Snodgrassella communis</i>	PTS_2-RNA Imm-SUKH-SMI1 PT-VENN Tox-ParBL1 ? PT-VENN
PZT91336.1 QFNA01000013.1:820-7509 <i>Citricrobium</i> sp.	Peptidase_M1_N Peptidase_M1 Leuk-A4-hydro_C MarR-HTH PTS_2-RNA FiL_haemagg_2 Tox-ParBL1
RHP98161.1 QUEU01000077.1:730-3247 <i>Firmicutes bacterium</i> AMS9-13	PTS_2-RNA ? RHS_repeat RHS_repeat RHS Tox-ParBL1
RML87211.1 RBNL01001659.1:1-542 <i>Pseudomonas syringae</i> pv. <i>maculicola</i>	Tox-ParBL1
SEB04148.1 FNQ01000021.1:1-7334 <i>Lonsdalea quercina</i>	FiLH FiL_haemagg_2 FiL_haemagg_2 FiL_haemagg_2 FiL_haemagg_2 PT-VENN Tox-ParBL1 SMI1_KNR4 PTS_2-RNA
SHI24935.1 FOXV01000030.1:1-1578 <i>Sporobacter termitidis</i> DSM 10068	Tox-ParBL1 ? PTS_2-RNA
TAL06778.1 SCTL01000065.1:61-4058 <i>Verrucomicrobia bacterium</i>	? ? RHS RHS PT-COG5412 Tox-ParBL1
WP_010582396.1 NZ_JH636434.1:1036801-1041040 <i>Schlesneria patudicola</i> DSM 18645	HINT ? HINT Tox-ParBL1 ? ?
WP_010769430.1 NZ_ASWE01000001.1:859856-862974 <i>Enterococcus phoenicicola</i> ATCC BAA-412	PTS_2-RNA ? Tox-ParBL1 WXG PT-TG OsmC
WP_206437638.1 NZ_CAAAF010000159.1:4344-6982 <i>Candidatus Protofrankia datisciae</i>	? ? Tox-ParBL1
WP_016469385.1 NZ_KE150472.1:82777-89591 <i>Streptomyces</i> sp. <i>HPH0547</i>	? WXG DUF6531 RHS RHS RHS_repeat RHS Tox-ParBL1 ? ? ?
WP_026150765.1 NZ_BARF010000050.1:44265-58608 <i>Streptomyces prunicolor</i> NBRC 13075	? Subtilisin ? Tetr_C_1 Tetr-HTH DUF6531 LevanB LamG RHS RHS RHS_repeat LamininG LamininG RHS RHS RHS_repeat RHS Tox-ParBL1
WP_100139032.1 NZ_MEIP01000013.1:203636-205887 <i>Snodgrassella alvi</i>	SMI1_KNR4 PT-VENN Tox-ParBL1 CdiI PTS_2-RNA
WP_206743637.1 NZ_JAIY01000006.1:132967-143556 <i>Kitasatospora</i> sp. <i>MBT66</i>	GntR-HTH UTRA ROK CitB-HTH ? ? DUF6531 RHS RHS RHS_repeat RHS_repeat RHS_repeat RHS RHS_repeat RHS_repeat RHS Tox-ParBL1
WP_139120346.1 NZ_CP025698.1:4781334-4785910 <i>Serratia marcescens</i>	Imm-Barstar ? ? ? Tox-ParBL1
WP_049664114.1 NZ_IFXJ01000005.1:289657-294122 <i>Lysinibacillus xylanilyticus</i>	RHS RHS Tox-ParBL1 Imm-SUKH-SMI1 ? ?
WP_078968535.1 NZ_LGEE01000263.1:7823-14174 <i>Streptomyces</i> sp. <i>NRRL F-6491</i>	? ? DEATH SseB T7SS_signal DUF6531 RHS RHS RHS_repeat RHS_repeat RHS_repeat RHS Tox-ParBL1
WP_157121691.1 NZ_BDCN01000009.1:120748-130526 <i>Nocardia violaceofusca</i> NBRC 14427	YbaB_DNA_bd WXG ? WXG Tox-ART-RSE SMAD TOXIN-PL Tox-ParBL1 ReLA-SpoT ?
WP_176704513.1 NZ_LRUN01000109.1:1-2387 <i>Pseudomonas bananamidigenes</i>	? ? RHS Tox-ParBL1
WP_067581410.1 NZ_JABMC2010000002.1:136034-144255 <i>Nocardia terpenica</i>	? ? WXG DUF5585 Tox-ParBL1 ReLA-SpoT T7SS_ESX_EspC ?
WP_089832762.1 NZ_FNB01000003.1:411677-419082 <i>Chitinophaga filiformis</i>	? FN3 WXG HINT Tox-ParBL1 ? Imm-SUKH-SMI1
WP_090002510.1 NZ_FMUG010000008.1:67460-70641 <i>Lysinibacillus fusiformis</i>	ARG Tox-ParBL1 ? ? RHS
WP_093858781.1 NZ_LT629775.1:1159123-1170533 <i>Streptomyces</i> sp. <i>TLI_053</i>	RibD_C DUF6531 RHS RHS RHS RHS_repeat RHS_repeat RHS_repeat RHS_repeat RHS_repeat RHS Tox-ParBL1 GSDH ? CBM_6
WP_141243305.1 NZ_NEVL01000003.1:202020-208046 <i>Bordetella genomsp. 1</i>	T6SS_TssG ClpABN-AAA ClpABC-AAA ? PTS_2-RNA Tox-ParBL1
WP_154212913.1 NZ_CP046027.1:2522921-2531331 <i>Neisseria brasiliensis</i>	Imm-SUKH-SMI1 ? PTS_2-RNA FiLH FiL_haemagg_2 FiL_haemagg_2 PT-VENN Tox-ParBL1 FiL_haemagg_2 SHOCT
WP_212768901.1 NZ_CP065143.1:63874-72465 <i>Pectobacterium versatile</i>	DcrB ? PrePAARTM PAAR_motif PAAR DUF6531 RHS RHS_repeat RHS_repeat RHS_repeat RHS Tox-ParBL1 Phage_GPD VgrG Toast-rack RHS
WP_106098274.1 NZ_CP027229.1:1539099-1545694 <i>Capnocytophaga</i> sp. <i>oral</i> taxon 878	? DUF6531 RHS RHS RHS_repeat RHS Tox-ParBL1 ? ? ?
WP_205692747.1 NZ_QOH001000025.1:13960-19239 <i>Lacrimispora amygdalina</i>	PTS_2-RNA ? DUF4280 Imm1 Phage_GPD Tox-ParBL1
WP_211338973.1 NZ_SNZE01000066.1:352-1833 <i>Hydromonas duriensis</i>	? RHS Tox-ParBL1
WP_169051682.1 NZ_SJZD01000005.1:49066-52352 <i>Flavobacterium</i> sp. <i>H122</i>	NADAR ? ? RHS RHS Tox-ParBL1
WP_196303695.1 NZ_LDUT01000075.1:145-1416 <i>Chromobacterium violaceum</i>	? RHS Tox-ParBL1
WP_139798835.1 NZ_FWXD01000016.1:496-4221 <i>Andreporetia lacus</i> DSM 23236	? MPTase Tox-ParBL1 ? Transposase_20 MPTase
WP_144994393.1 NZ_CP036281.1:1724341-1726548 <i>Polystyrenella longa</i>	? ? Tox-ParBL1 ? ?
WP_149694807.1 NZ_FRCN01000003.1:467435-474555 <i>Chitinophaga</i> sp. <i>CF418</i>	Sigma70_r4_2 Sigma70_r2 DUF5009 DUF6531 RHS RHS_repeat RHS Tox-ParBL1 ? Endonuclease_5
WP_167230982.1 NZ_VUYU010000030.1:75428-82200 <i>Massilia rubra</i>	Bac_export_3 Bac_export_1 DUF6861 DUF6531 RHS_repeat RHS_repeat RHS_repeat RHS RHS Tox-ParBL1 ? ?
WP_210429397.1 NZ_JAAMW2010000041.1:34892-45074 <i>Paenibacillus dendritiformis</i>	SUKH_3 Imm24 Phage_GPD Phage_base_V DUF6531 RHS_repeat RHS_repeat RHS RHS RHS RHS RHS AHH RHS Tox-ParBL1 PTS_2-RNA
WP_184478450.1 NZ_JACHIV010000001.1:2027587-2033469 <i>Saccharopolyspora gloriosae</i>	T7SS_ESX_EspC WXG DUF6531 RHS RHS RHS Tox-ParBL1 ? ?
WP_201256142.1 NZ_JAEDK2010000064.1:47727-57837 <i>Pseudomonas</i> sp. <i>TH03</i>	? DUF6531 RHS_repeat RHS RHS_repeat RHS Tox-ParBL1 DUF4123 Phage_GPD VgrG DUF2345 ?
WP_20609767.1 NZ_CP065053.1:166678-177591 <i>Massilia antarctica</i>	Bac_export_3 Bac_export_1 DUF6861 DUF6531 RHS_repeat RHS RHS RHS Tox-ParBL1 ? DUF6861 DUF6531 RHS_repeat RHS
WP_208100708.1 NZ_JAGEKB010000002.1:681999-684952 <i>Lachnoanaerobaculum</i> sp. <i>Marseille-04761</i>	Transposase_20 DEDD Tnp_IS110 PTS_2-RNA ? Tox-ParBL1 ?