

ART38309.1 KX576139.1:14993-21372 <i>uncultured bacterium</i>	← Rieseke Rhs Rhs Tlde1 ? Rhs IPTL-CTERM Rhs TseH
CBG80105.1 FN543502.1:2480973-2484627 <i>Citrobacter rodentium ICC168</i>	← Ferredoxin-beta-grasp ? Ntox24 ? LZ_Tnp_IS66 zf-IS66 DDE_Tnp_IS66 DDE_Tnp_IS66_C
EBV0231387.1 AAHDY1010000050.1:1-1247 <i>Salmonella enterica subsp. salamae serovar Sofia</i>	← ? Rhs Ntox24
ECJ586653.1 AA1YV010000002.1:238129-247314 <i>Salmonella enterica subsp. salamae</i>	← MatE MatE ? DUF6531 Rhs Rhs_repeat Rhs Rhs Rhs_repeat Rhs Ntox24 ? Rhs Tox-JAB2
EF124099.1 GL349411.1:274086-282891 <i>Neisseria sp. oral taxon 014 str. F0314</i>	← YbaB_DNA_bd Alpha-amylase Malt_amylase_C Tox-PAAR-Like DUF6531 Rhs Rhs Rhs Ntox24 Pentapeptide_4 ?
EXB27792.1 JEW01000001.1:568794-571453 <i>Acinetobacter baumannii 1437282</i>	← ? Ntox24 ? ? ?
HBB7028235.1 DADVZ010000001.1:467770-475389 <i>Escherichia coli</i>	← ? ? Phage_GPD COOHNH2Ligase DUF5445 GST_N_2 GST_C_2
HCK4345624.1 DAIWS010000006.1:1-3233 <i>Pseudomonas aeruginosa</i>	← Rhs Ntox24 ? Ntox24 ? ?
HCT30801.1 DPJ801000007.1:101247-110182 <i>Moraxellaceae bacterium</i>	← S1_2 S1_2 Cvfb_WH ? DUF6531 Rhs_repeat Rhs_repeat Rhs Rhs Rhs Rhs Ntox24 GDDEF Mbetalac Beta-Casp RMMBL
KAF1009578.1 WDD001000001.1:194215-206011 <i>Luteibacter sp.</i>	← CoxA Imm DUF6531 Rhs_repeat Rhs_repeat Rhs Rhs Ntox24 Dak1 Dak2 Peptidase_59
KNE06404.1 AENW82000003.1:2-6867 <i>Aggregatibacter aphrophilus ATCC 33389</i>	← DUF2345 DUF6531 Rhs_repeat Rhs Rhs_repeat Rhs Rhs Ntox24 ? Adenylsucc_synt
KPA17598.1 JPD01000570.1:13621-17102 <i>Candidatus Magnetomorum sp. HK-1</i>	← PemK_toxin ? Ntox24 ? ?
KRW62034.1 LKKX01000001.1:213151-218934 <i>Pseudomonas sp. TTU014-080ASC</i>	← Rhs Rhs Rhs Rhs Ntox37 ? Rhs Ntox24 ? TetR_C_24 TetR-HTH
MBA3114137.1 AAGWY020000055.1:244881-247892 <i>Salmonella enterica</i>	← ? ? Rhs Ntox24
MBD9658560.1 JACYU0010000016.1:1-1251 <i>Oxalobacteraceae sp. CFBP 13730</i>	← Rhs_repeat ? Rhs Ntox24
MBE0535677.1 JACZK0010000083.1:49481-52877 <i>Phycisphaerae bacterium</i>	← SM-MscS ? Rhs Ntox24 ? ?
MBI3408315.1 JACZP010000048.1:404-1479 <i>Planctomycetota bacterium</i>	← Rhs Ntox24
MBI389506.1 JACF010000039.1:11079-15164 <i>Gammaproteobacteria bacterium</i>	← HTH ? Rhs Ntox24 Rhs_repeat Rhs DUF6531 Rhs
MBI4802861.1 JACYT010000092.1:60128-63985 <i>Elusimicrobia bacterium</i>	← ? ? Ntox24 SIS GKR- Like_C ab-Hydrolase
MBI9019390.1 JAE106010000170.1:3837-7772 <i>Phycisphaerae bacterium</i>	← PDZ PDZ Trypsin ? Rhs Ntox24
MBK8915058.1 JADJXS010000001.1:2641560-2651675 <i>Phycisphaerales bacterium</i>	← ? HTH_33 DDE_3 Rhs Rhs Rhs Ntox24 ? Rhs
MBK9260594.1 JADJXL010000010.1:942-7084 <i>Polyangiaceae bacterium</i>	← SF2-DUF3427A EcoEI_R_C ? Ntox24 ? ?
MBK9262824.1 JADJXL010000018.1:646214-648402 <i>Polyangiaceae bacterium</i>	← ? ? Ntox24 ? ?
MBQ9094965.1 JAFTE010000043.1:7383-9049 <i>Akkermansia sp.</i>	← ? TIM-GlycoH Rhs Ntox24
MBR0968506.1 JAFCKV010000039.1:18479-16122 <i>Bradyrhizobium diazoefficiens</i>	← ? ? DUF6531 Rhs Rhs Rhs Rhs Rhs Ntox24 ? ?
MBX3251153.1 JAHBW010000188.1:153-5862 <i>Myxococcales bacterium</i>	← GidB TrmB SLH Ntox24 DUF2914 baeRF1
MBZ0150428.1 JAI0JB010000156.1:1-2034 <i>Planctomycetota bacterium</i>	← Rhs Ntox24 ?
MCC2103186.1 JAGRKJ010000127.1:1-1922 <i>Hyphomicrobiales bacterium</i>	← Rhs Ntox24 HINI Tox-REase-9
MCC7330937.1 JADY20010000157.1:1-1204 <i>Gammaproteobacteria bacterium</i>	← Rhs Ntox24 ? ?
NGO37790.1 JAAKYA010000001.1:95-724 <i>Limisphaera ngatamarikiensis</i>	← Rhs Ntox24
NOT8217.1 JABFR010000001.1:1900644-1909610 <i>Lysobacter sp.</i>	← AAA_19 UvrD_C_2 GTP_cyclohydrol DUF6531 Rhs Rhs_repeat Rhs_repeat Rhs Rhs Ntox24 ? Rick-17kDa-Anti
NY269815.1 JACCKB010000154.1:825-3022 <i>Spartinivicius marinus</i>	← HTH ? Rhs
PIT08410.1 MDVA010000033.1:714-2578 <i>Snodgrassella communis</i>	← Rhs Stox-39 ? Ntox24
PNC69062.1 PJKR010000008.1:1795-3583 <i>Akkermansia muciniphila</i>	← TOX-HTH Imm Rhs Ntox24
PWB18185.1 QETI010000007.1:159099-176441 <i>Comamonas sp. JNW</i>	← ? Big_7 Big_6 Big_7 CarboxypepD-reg-IG ASH-Ig CarboxypepD-reg-IG Rhs_repeat Rhs_repeat Rhs Rhs Rhs Ntox24 ? Lipoprotein_9
QBY31975.1 CP038008.1:1250793-1259792 <i>Citrobacter rodentium</i>	← HIPEP Bac_export_2 ? CheZ Rhs Ntox24
QGM98373.1 CP044331.1:2830072-2839534 <i>Methylocystis parvus</i>	← DUF2336 HTH_18 DUF6531 Rhs Rhs Rhs Rhs Rhs_repeat Rhs Ntox24 ? Rhs
QIA75974.1 CP040863.1:131498-140654 <i>Rodentibacter caecimuris</i>	← Phage_GPD VgrG DUF2345 DUF6531 Rhs Rhs Rhs Stox-59 ? Rhs Ntox24 ? Rhs
QIA75982.1 CP040863.1:143663-147935 <i>Rodentibacter caecimuris</i>	← Ntox24 ? Rhs Ntox24 Rhs_repeat Rhs prok-SAA ?
RPO23810.1 NSOK01000012.1:204272-209690 <i>Pseudomonas aeruginosa</i>	← CN_hydrolase ? Rhs Ntox24
RYY80261.1 SECL01000005.1:10341-13209 <i>Moraxellaceae bacterium</i>	← Rhs ? Rhs Nmad4 Htase_MTS_N
SDD76281.1 FMY001000025.1:2255-9405 <i>Paraburkholderia lycopersici</i>	← ? DUF6714 Rhs Ntox24 Toprim AAA_29 Phage-portal SbcC_Walker_B
SEG15533.1 FNVK010000037.1:15318-18966 <i>Nitrosospora multiformis ATCC 25196</i>	← Phage_integrase Phage_int_SAM_1 ? Rhs_repeat Rhs_repeat Rhs Ntox24
TAN40871.1 SCQE010000093.1:11134-13031 <i>Nitrospirae bacterium</i>	← Rhs_repeat Rhs ? Rhs Ntox24
TWT45278.1 SMLU010000001.1:2110025-2115989 <i>Phycisphaerae bacterium RAS1</i>	← Rhs ? Rhs Ntox24 Dockerin_1 HTH_38 TraK-HTH rve
VTX59384.1 CABFLM010000039.1:45-521 <i>uncultured Aggregatibacter sp.</i>	← Ntox24
VTX84710.1 CABFLM010000083.1:1101-7415 <i>uncultured Aggregatibacter sp.</i>	← SbcC_ABC_tran SbcC UvrA_DNA-bind UvrA_inter ? Rhs Ntox24 Pentapeptide Pentapeptide_4 Rhs
WP_212749466.1 NZ_KB849164.1:1220164-1223481 <i>Acinetobacter bohemicus ANC 3994</i>	← MerR-HTH ? ? Stox-2 Ntox24
WP_050693340.1 NZ_CP012067.1:1207888-1216856 <i>Aggregatibacter aphrophilus</i>	← DUF484 ? ? Phage_GPD VgrG DUF2345 DUF6531 Rhs_repeat Rhs Rhs Rhs Ntox24 Rhs Ntox24
WP_224229830.1 NZ_CP043403.1:1185470-1195341 <i>Xanthomonas oryzae pv. oryzicola</i>	← DUF6869 CoxE DUF6531 Rhs Rhs_repeat Rhs_repeat Rhs Rhs Ntox24 DUF772 DDE_Tnp_1 HTH_Tnp_1 HTH_21 rve_3
WP_049876215.1 NC_010162.1:2484882-2491817 <i>Sorangium cellulosum So ce56</i>	← ? ? FA_hydroxylase Metallopeptidase Ntox24 ?
WP_021761170.1 NC_022444.1:2542886-2547658 <i>Megalodesulfobivrio gigas DSM 1382 = ATCC 19364</i>	← ? Rhs Ntox24 DDE_Tnp_1 DUF772 Rhs ?
WP_10009876.1 NZ_MDUY01000021.1:2704-5865 <i>Snodgrassella communis</i>	← ? ? ? ? Rhs Ntox24
WP_196247953.1 NZ_JP3001000036.1:1-4161 <i>Gallibacterium anatis</i>	← ? Rhs Rhs_repeat Rhs Ntox24 Rhs Tox-DUF4105
WP_228064786.1 NZ_JADGL010000018.1:45863-55275 <i>Muribacter muris</i>	← T6SS_HCP Phage_GPD VgrG DUF6531 Rhs_repeat Rhs_repeat Rhs Rhs Tox-PLA2 ? Rhs Ntox24 RxD-HxH-H
WP_06758200.1 NZ_CP016895.1:3454923-3468774 <i>Acinetobacter larvae</i>	← ? Rhs_repeat Rhs_repeat Rhs_repeat Rhs_repeat Rhs Rhs Ntox24 Phage_GPD VgrG T6SS_Vgr DUF2345 HSP90 HIN-HTH HTH_Tnp_1 HTH_32 rve rve_2
WP_139271159.1 NZ_MPI001000021.1:1-1646 <i>Atopomonas hussainii</i>	← ? T2SSC Rhs Ntox24
WP_081306407.1 NZ_JTEI01000328.1:989-3642 <i>Xanthomonas euvesicatoria</i>	← Rhs Ntox24 ? Rhs
WP_089462763.1 NZ_NKFR01000004.1:1409032-1418557 <i>Burkholderia sp. AU27893</i>	← Rhs Rhs Ntox24 T6SS_TssF ? Phage_GPD VgrG DUF6531 Rhs Rhs_repeat Rhs Rhs
WP_108984754.1 NZ_BFBRO1000004.1:124021-128247 <i>Candidatus Phycosocius bacilliformis</i>	← Rhs ? Rhs Ntox24 Redoxin Thiolase_N Thiolase_C
WP_151980300.1 NZ_BKYM01000003.1:212534-216506 <i>Acinetobacter guerae</i>	← ? Phage_holin_3_3 Stox-56 LPAM_1 Ntox24 ?
WP_131688269.1 NZ_QRUB01000002.1:195113-201083 <i>Psychrobacter pygocelis</i>	← Rhs Ntox37 ? Rhs Rhs ? Rhs
WP_142854889.1 NZ_FXW01000010.1:30915-34117 <i>Aliroseovarius halocynthiae</i>	← Rhs ? ANK Rhs Ntox24 Rhs
WP_158860101.1 NZ_VOLC01000015.1:103607-110539 <i>Lunatibacter saliacus</i>	← Metallopeptidase Metallopeptidase PG_binding_1 ? Metallopeptidase Ntox24 Metallopeptidase Ntox42-Colicin_5 PVC-AA
WP_164027455.1 NZ_CP040863.1:143663-147935 <i>Rodentibacter caecimuris</i>	← Ntox24 ? Rhs Ntox24 Rhs_repeat Rhs prok-SAA ?
WP_172240493.1 NZ_JABME010000001.1:1412366-1422548 <i>Permethianbacter fluminis</i>	← AMP-binding_C AMP-binding ACAS_N TerC DUF6531 Rhs_repeat Rhs_repeat Rhs_repeat Rhs_repeat Rhs Rhs Ntox24 ? BetaPropeller
WP_187082436.1 NZ_JACOR010000005.1:362447-368805 <i>Ramlibacter albus</i>	← AMP-binding_C AMP-binding ? Rhs Ntox24 DUF6531 Rhs_repeat Rhs Catalase
WP_193033916.1 NZ_JADBL010000028.1:83361-84849 <i>Enterobacter cloacae complex sp. P15RS</i>	← ? ? Rhs Ntox24
WP_22727247.1 NZ_JACNM0010000086.1:167-598 <i>Roseobacter weihaiensis</i>	← Ntox24
WP_423440941.1 NZ_JBNVQY010000001.1:460162-466664 <i>Limisphaera sp. VF-2</i>	← AhpC-TSA Thioredoxin_8 Rhs Rhs Ntox24 Rhs ?