

CP83031.1:1177611-118064 <i>Aerobacterium tumefaciens</i>	rcnB [TPS F1LH F1LH F1L_haemagg_2 F1L_haemagg_2 DUF682 Ntox] ? POTRA_2 Sh1b
CEV8202.1 G057738.1:1323997-3338025 <i>Micromonospora</i> sp. ATCC 39149	Syme_toxin [Imp PT-COG541]
CNC40383.1 CPW01000082.1:221729-229922 <i>Yersinia frederiksenii</i>	Cdi [Imp TPS DUF637 DUF637 PT-VEIN Ntox] ? Ema UFF823
EEPT2273.1 G057738.1:1323997-3338025 <i>Micromonospora</i> sp. ATCC 39149	ODE_Tnp_ISL3 [Imp SPVb R5 R5 Ntox] ? BetaPropeller Laminin Laminin Phage_integrase Phage_int_SAM_3
EEV090234.1 AABP001000076.1:1-11986 <i>Escherichia coli</i>	R5 Ntox [Imp SGRA-S4]
HE77460.1 DPL41000611.1:1-2382 <i>Lysinibacillus</i> sp.	R5b Ntox [Imp R5b]
HH24079.1 D1W0100082.1:1182-3718 <i>Xanthomonas vesicola</i> pv. zee	[Imp Ntox] ? DUF682
KW78877.1 LW001000811.1:77777-82207 <i>Aerobacterium tumefaciens</i> str. 86	? [Imp F1L_haemagg_2 Ntox] ? ? ?
KWV63172.1 L1W01000816.1:594-3171 <i>Neotrophomonas maltophilia</i>	? ? ? Ntox ?
NB0645074.1 JAC0401000047.1:172-2067 <i>Thalassotalea piscium</i>	[Imp R5 Ntox] ? Toxin_YnfY PrIF_antitoxin
NB1762482.1 J0UTP01000003.1:482983-408994 <i>Burkholderiales</i> bacterium	151b_1521 fvn Sigma70_r4_2 R5 DUF637 DUF637 Ntox [Imp ?
NB1903885.1 JALC001000045.1:1-1760 <i>Enterobacter asburiae</i>	Ntox [Imp ?
NB17714333.1 JADP001000064.1:828-3211 <i>Proteus hauseri</i>	Ntox [Imp ?
NB1770961.1 JAN7001000088.1:2915-6056 <i>Candidatus Nephthysia benettiae</i>	Bac_Luciferase [Imp Ntox] ? R5b ?
NB1645004.1 JADJ001000003.1:1032211-1038791 <i>Suprapiraceae</i> bacterium	? TrxA_AAD_bipr Ntox [Imp R5N
NB1640073.1 JADJ001000001.1:511233-523837 <i>Paraproteobacteria</i> bacterium	Asp_Glu_race DIS1N BetaPropeller unv0v_ASPIc SPVb BetaPropeller R5_repeat R5 R5 R5_repeat R5 Ntox ? G5bH PMI_type1_hel PMI_type1_cat ? ?
NB1640034.1 JADJ001000002.1:2188760-2193483 <i>Paraproteobacteria</i> bacterium	OnpK_HTH ? Ntox ? TPAA-synt_2b_HGTP_anticondop Q3-1_PPpI
NB1770961.1 JADJ001000043.1:5784-12049 <i>Acidobacteria</i> bacterium	R5b [Imp R5 Ntox] ? ?
NB1124183.1 HUS0200023.1:139-5517 OC5126 cluster Bacterium	q5p_X2 PG_binding_1 q511 Ntox ? ?
NB10080215.1 JAFN05010000239.1:2466-4456 <i>Mycobacterium</i> sp.	? [Imp R5b Ntox] ? ? R5b
NB1070372.1 JATW0010000192.1:68-1996 <i>Bacteroidales</i> bacterium	R5 LysH Ntox [Imp ?
NB11374895.1 JAU0P01000024.1:1126-1758 <i>Cardiobacteriaceae</i> bacterium	Ntox [Imp Imp Sell
NB1780322.1 JAFB001000011.1:1-3279 <i>Chromobacterium violaceum</i>	Ntox [Imp Imp Imp
NB1045096.1 JAH001000003.1:137749-348974 <i>Serratia liquefaciens</i>	U1_nuc_hydro [Imp Ntox] ? ? ?
NB1050777.1 JAGF001000070.1:238-4418 <i>Temerizicola</i> larvae	Ntox [Imp Syme_toxin
NB1076208.1 JAFAT010000060.1:8235-10579 <i>Verruconiscrobota</i> bacterium	THRM7b [Imp R5b Ntox]
NA1A2963.1 R0001000057.1:1-11289 <i>Salmonella enterica</i> subsp. <i>enterica</i>	R5b Ntox [Imp Imp Tox_A0b
NC250488.1 V0001000010.1:1-948 <i>Dickeya dianthicola</i>	Ntox [Imp Imp Syme_toxin
NML32388.1 JABF0201000009.1:4872-9321 <i>Paraburkholderia antisibatica</i>	? [Imp Ntox] ? ? ?
NH71254.1 V0001000016.1:174991-78264 <i>Vibrio parvulus</i>	Fera_12 R5b R5 DUF637 PT-TC Ntox [Imp Imp
NQ07107.1 JABP001000024.1:489-2819 <i>Flavobacteriaceae</i> bacterium	v2_Tnp_2n_Tnp_1591 HTH
NQ07107.1 JABP001000024.1:489-2819 <i>Flavobacteriaceae</i> bacterium	? [Imp R5 R5 Ntox] ? [Imp Tox-Arc
NRA0101.1 JABP001000024.1:489-2819 <i>Flavobacteriaceae</i> bacterium	HTN_Tnp_1 PT-TC Ntox [Imp ?
CA000001.1 LY001000023.1:1135894-149452 <i>Rhodobacteraceae</i> bacterium	Thioredoxin_a CoA_transf_3 TPS F1LH F1LH F1LH F1LH Ntox [Imp ?
CEK0141.1 J1W01000013.1:1589-2485 <i>Galibacterium</i> genomsp. 3	[Imp ? Ntox]
CC050006.1 M1W01000016.1:17532-9914 <i>[Bacteroides]</i> felis	DEAM-Dot [Imp ? ? ? Ntox]
PI00007.1 P0001000004.1:36196-38929 <i>Conservatibacter flavescens</i>	? [Imp DUF682 Ntox] ? R5b Ntox2
QV70006.1 CP07533.1:128796-232121 <i>Serratia marcescens</i>	Adhesin_E [Imp Ntox] ? ? PT-VEH
R170007.1 Q0001000028.1:9546-14269 <i>Nakamurella silvestris</i>	Peripla_BP_4 B1R-HTH R5b Metalloproteinase [Imp PadH-HTH Vir_act_alpha_c
R170007.1 Q0001000028.1:9546-14269 <i>Nakamurella silvestris</i>	Sigma70_r2 Sigma70_r4_2 [Imp HPase-PVC Ntox] ? 3BP_dG1ase Ema Ema
SAL0073.1 FC001000008.1:113-2861 <i>Caballeron pterocanthos</i>	Ntox [Imp ?
SE07001.1 P001000014.1:152084-59027 <i>Bosae latyrus</i>	? TFR Ntox ? Ab_Amino_oxidase
SEP0003.1 P001000018.1:133665-41529 <i>Mycolatopsis saxifolensis</i>	ABC_tran [Imp ALF ALF ALF ALF ALF ALF Ntox] ? ? TRNA-synt_2b_HGTP_anticondop
SH00002.1 P001000044.1:21147-36721 <i>Actinobacillus paucisporus</i>	? BetaPropeller BetaPropeller SPVb R5 R5 Ntox [Imp ? ? ODE_Tnp_1_asec
SI000001.1 FS001000001.1:865845-870494 <i>Burkholderia</i> sp. G0332	adh_short_C2 ? DUF631 R5 R5 R5 Ntox [Imp Imp
SLA0001.1 UG01000001.1:1382693-1388425 <i>Neisseria gonorrhoeae</i>	? R5b R5b Ntox [Imp DMP2 ?
TAK2207.1 SC001000027.1:61185-64198 <i>Chloroflexus</i> bacterium	q5p_X2 Ntox [Imp Ntox] ? Transposase_muc
WP_009000001.1 N2_001301000006.1:1-11008 <i>Neisseria meningitidis</i>	[Imp ? Ntox]
WP_009000001.1 N2_001301000006.1:1-11008 <i>Neisseria meningitidis</i>	? PAC DUF631 R5 R5 R5_repeat R5 R5_repeat R5 R5_repeat R5 Ntox [Imp R5b Secre_Ca2b TPase Phage_CATRA Caspase
WP_009000001.1 N2_	