

AHN25716.1 CP007445.1:1261505-1263796 <i>Gilliamella apicola</i>	? RHS RHS STox-1 → Caspase → ?
ALB61113.1 CP012264.1:122569-126615 <i>Cronobacter condimenti</i> 1330	? ? RHS STox-1 → ? RHS
AMO46689.1 CP012487.1:298467-302661 <i>Enterobacter</i> sp. FY-07	Tox-ART-HYD3 ? RHS STox-55 → ? RHS Ntox41
EDH6256436.1 AAMIGI010000064.1:3002-4448 <i>Salmonella enterica</i> subsp. <i>enterica</i> serovar Amoutive	CARP-ART ? STox-1 →
EGT4347247.1 AAXVRU010000003.1:523641-525828 <i>Elizabethkingia anophelis</i>	? DUF2947 STox-55 →
HAZ3638659.1 DADNFX010000107.1:41-895 <i>Escherichia coli</i>	? RHS STox-1 →
MBQ2885716.1 JAFQRT010000364.1:320-3997 <i>Alphaproteobacteria bacterium</i>	GDResolvase ? RHS STox-1 →
MBQ6160948.1 JAFRPC010000055.1:66579-72591 <i>Oscillospiraceae bacterium</i>	BAR_3_WASP_bdg PrePAARTM HINT Tox-CompNuc DUF2947 BAR_3_WASP_bdg HINT STox-1 → Ntox44 AAA_13
MBS6632883.1 JAGZYA010000012.1:50295-53653 <i>Ruminococcus bicirculans</i> (ex Wegman et al. 2014)	? DUF2947 STox-1 → Imm-PA2201 DUF2185
MCM1006351.1 JAMP1L010000001.1:35674-38525 <i>Ruminococcus flavefaciens</i>	HINT Tox-REase-1 ? HINT STox-1 → SMI1_KNR4 Imm55
MCM1060151.1 JAMPHS010000028.1:132-1649 <i>Eubacterium</i> sp.	STox-1 → ? ?
MCR5599338.1 JAILNU010000009.1:20981-25519 <i>Ruminococcus</i> sp.	HINT DUF2947 Gly-zipper_YMG6 HINT STox-1 → ? ?
NJO16335.1 JAAUSR010000031.1:46319-49811 <i>Thioploca</i> sp.	DUF6484 DUF2345 DUF2169 STox-1 → DUF2947 ?
NLP45252.1 JAAYSI010000304.1:1-1773 <i>Peptococcaceae bacterium</i>	RHS STox-1 → ?
OYD06597.1 NOWF01000011.1:96439-100949 <i>Paludifilum halophilum</i>	Ntox35 ? PT-TG STox-1 → Cys_rich_CPCC ?
RZJ61347.1 SEBN01001116.1:146-2011 <i>Flavobacterium</i> sp.	DUF2947 RHS STox-1 →
SSV52102.1 UFOZ01000016.1:2714-5565 <i>Acinetobacter nosocomialis</i>	? Acetylase RHS STox-1 → ? SUKH-CF44
TDT51581.1 SOBB01000004.1:196127-204189 <i>Enterobacter</i> sp. AG5470	RHS STox-11 ? RHS STox-55 → ? PrePAARTM PAAR PAAR_motif DUF6531 RHS_repeat RHS_repeat RHS_repeat RHS_repeat RHS RHS_repeat RHS_repeat RHS STox-ART-HYD2
TDV44491.1 SOCRO1000014.1:146399-153681 <i>Pseudomonas graminis</i>	Abhydrolase_2 Hemerythrin DUF6531 RHS_repeat RHS RHS_repeat RHS STox-1 → DUF2947 ?
WP_050676853.1 NZ_CACSGV010000122.1:1-1660 <i>Acinetobacter baumannii</i>	? STox-1 → CdiI_2 ?
WP_006781831.1 NZ_CABL0010000106.1:159-4193 <i>Hungateella hathewayi</i>	Flg_new Flg_new Flg_new STox-1 → ? PAS_6 HTH_22 ? CR-REase-U1
WP_021348305.1 NZ_AVCQ01000016.1:321197-325328 <i>Elizabethkingia anophelis</i> 502	? ? DUF2947 STox-1 → ?
WP_083703168.1 NZ_FTMN01000012.1:45737-54122 <i>Marinobacterium stanieri</i>	PSBP DUF2947 Phage_GPD VgrG Gp5_C Amidase_2 PAAR_motif DUF6531 RHS_repeat RHS_repeat RHS RHS STox-55 →
WP_017662949.1 NZ_KB235960.1:107260-114768 <i>Baaleninema simplex</i> PCC 7105	PD Ntox1 STox-55 → DUF2947 ? Metallopeptidase Metallopeptidase
WP_056833818.1 NZ_LMSP01000016.1:30333-39303 <i>Paenibacillus</i> sp. Soil787	S4 HU-IHF DUF3238 DUF6756 DUF6531 RHS_repeat RHS RHS RHS RHS RHS_repeat RHS_repeat RHS_repeat RHS RHS_repeat RHS RHS STox-1 →
WP_228406599.1 NZ_QGTV01000037.1:9746-13465 <i>Chryseobacterium</i> sp. AG844	COXG DUF2947 ? RHS Peptidase_M10 STox-55 →
WP_211264673.1 NZ_CP007142.1:2775465-2778544 <i>Gyneuella sunshinyii</i> YC6258	? RHS DUF2947 ? RHS STox-1 →
WP_052658175.1 NZ_LN810019.1:3667617-3679658 <i>Shewanella algae</i>	? PAIREDC-HTH DDE_3 DUF2947 SPVB BetaPropeller TcdB_toxin_midN RHS RHS STox-55 → ?
WP_127891255.1 NZ_CP137208.1:743702-746366 <i>Citrobacter koseri</i>	? STox-1 STox-59 ? ?
WP_080614174.1 NZ_CP020452.2:2038931-2044088 <i>Neisseria mucosa</i>	? rve HTH_38 RHS RHS STox-1 → Imm24 Imm-PA2201
WP_092745643.1 NZ_FNZC01000018.1:1-7953 <i>Paracidovorax valerianellae</i>	DcrB PrePAARTM PAAR_motif DUF6531 RHS RHS_repeat RHS RHS STox-1 → VgrG T6SS_Vgr DUF2345 DUF2947 Tox-HNH-EHH
WP_141201743.1 NZ_NOWF01000011.1:96439-100949 <i>Paludifilum halophilum</i>	Ntox35 ? Cys_rich_CPCC ? PT-TG STox-1 →
WP_211233970.1 NZ_ATZJ01000005.1:181-1984 <i>Chitinilyticum litopenaei</i> DSM 21440	? Imm50 RHS STox-1 →
WP_149502777.1 NZ_CP035708.1:837293-845738 <i>Sphaerotilus sulfidivorans</i>	Phage_GPD VgrG DUF2345 DcrB PAAR DUF6531 RHS RHS RHS RHS STox-1 → DUF6444 ?
WP_211228151.1 NZ_AUCH01000055.1:1-977 <i>Azovibrio restrictus</i> DSM 23866	? RHS STox-1 →
WP_172298071.1 NZ_JABJU010000007.1:3252536-3258247 <i>Bacillus</i> sp. WMMC1349	? LXG PT-TG STox-1 → ? MarR-HTH Phage_integrase
WP_183724118.1 NZ_JACHW010000006.1:82149-84789 <i>Paraburkholderia bannensis</i>	STox-1 → DUF2947 PAAR_motif CHC2 DNAG_N HTH
WP_237719665.1 NZ_RPBS01000013.1:1-3226 <i>Cronobacter dublinensis</i>	RHS STox-1 DUF2947 ? Imm18