

CRG50531.1 CVMG01000014.1:330-1862 <i>Yersinia wautersii</i>	Transposase_mut Tle3 ? ?
ECA1950233.1 AAHT001000001.1:434629-441035 <i>Salmonella enterica subsp. enterica serovar Virchow</i>	DotU Phage_GPD VgrG T6SS_Vgr DUF2345 Tox-ABHYDROLASE2 DUF3274 Tle3 ? ?
ECF1703739.1 AAIKH501000061.1:6063-12580 <i>Salmonella enterica subsp. enterica</i>	Tox-PAAR-like Tox-ABHYDROLASE2 DUF3274 Tle3 ? Tox-ABHYDROLASE2 DUF3274 Tle3 DUF1525 TraU
EDR4080347.1 AANUB001000031.1:194-6884 <i>Salmonella enterica</i>	? Tox-PAAR-like ab-Hydrolase Tle3 Macro Tox-PAAR-like Tox-ABHYDROLASE2 DUF3274 Tle3
MBB1199422.1 RQJH01000002.1:52258-59897 <i>Enterobacteriaceae bacterium 89</i>	HTH_50 NtrC-AAA PAS ACT Phage_GPD VgrG T6SS_Vgr DUF2345 ab-Hydrolase Tle3 ? PAAR_motif
MBS1204473.1 JAGTSE010000009.1:64070-70478 <i>Proteobacteria bacterium</i>	Phage_GPD VgrG T6SS_Vgr DUF2345 Tox-ABHYDROLASE2 DUF3274 Tle3 ? ?
QLK62226.1 CP045665.1:3561829-3570906 <i>Enterobacteriaceae bacterium Kacie_13</i>	Glyco_transf_9 Phage_GPD VgrG T6SS_Vgr DUF2345 ab-Hydrolase Tle3 ? ?
QMB07944.1 CP056967.1:4450565-4458064 <i>Citrobacter freundii</i>	? ? Tox-ABHYDROLASE2 Tle3 Phage_GPD VgrG T6SS_Vgr DUF2345 CarboxypepD-reg-IG Lyx_isomer
WP_407199092.1 NZ_JBNAL010000001.1:4427354-4434847 <i>Citrobacter portucalensis</i>	Lyx_isomer ? ? Tox-ABHYDROLASE2 Tle3 Phage_GPD VgrG T6SS_Vgr DUF2345
WP_047058253.1 NZ_CP101017.1:3197477-3203247 <i>Klebsiella aerogenes</i>	AA_permease Gln-synt_N_2 Gln-synt_C Phage_GPD Tle3 ? ?
WP_126508535.1 NZ_RXQH01000007.1:39-11505 <i>Enterobacter hormaechei</i>	OEP OEP ACR_tran ? ? Phage_GPD VgrG T6SS_Vgr DUF2345 Tox-ABHYDROLASE2 DUF3274 Tle3
WP_023309373.1 NZ_CP034336.1:3257411-3262515 <i>Enterobacter asburiae</i>	HerA_C Tle3 ? DDE_Tnp_IS1 HTH_Tnp_IS1 Zn_Tnp_IS1 Tox-ABHYDROLASE2
WP_023492314.1 NZ_AYKS02000112.1:106722-113539 <i>Serratia sp. DD3</i>	DotU Phage_GPD VgrG T6SS_Vgr DUF2345 Tox-ABHYDROLASE2 Tle3 ? ?
WP_106876006.1 NZ_PYEP01000001.1:376960-384810 <i>Siccibacter turicensis</i>	Citrate_synt ? DUF707 Tox-ABHYDROLASE2 Tle3 Phage_GPD VgrG T6SS_Vgr DUF2345 CarboxypepD-reg-IG
WP_049294516.1 NZ_JVIC010000083.1:179866-186912 <i>Franconibacter helveticus</i>	Na_Ca_ex Tox-ABHYDROLASE2 Tle3 ? PAAR_motif Phage_GPD VgrG T6SS_Vgr DUF2345
WP_034493790.1 NZ_JMPI01000020.1:185097-190603 <i>Buttiauxella agrestis ATCC 33320</i>	? Tox-PAAR-like Tox-ABHYDROLASE2 DUF3274 Tle3 DSBH DSBH ? ?
WP_052130888.1 NZ_JRUQ01000102.1:20584-27463 <i>Erwinia typographi</i>	? 35exo Tox-ABHYDROLASE2 DUF3274 Tle3 Phage_GPD VgrG T6SS_Vgr DUF2345 PAAR_motif
WP_110025437.1 NZ_QGTS01000004.1:181117-187811 <i>Mangrovibacter plantisponsor</i>	PRA-CH PRA-PH Phage_GPD VgrG T6SS_Vgr DUF2345 ab-Hydrolase Tle3 PAAR_motif ?
WP_047066407.1 NZ_LDBY01000017.1:4174-9458 <i>Klebsiella aerogenes</i>	Gln-synt_N_2 Gln-synt_C AA_permease Tle3 ? ?
WP_046497042.1 NZ_CP011132.1:5419217-5429331 <i>Citrobacter amalonaticus Y19</i>	? Tle3 ? Autotransporter AC_1 PATR ESPR ?
WP_047608055.1 NZ_JUHL01000017.1:73749-80706 <i>Rahnella aquatilis</i>	PAAR_motif Tox-ABHYDROLASE2 Tle3 Phage_GPD VgrG T6SS_Vgr DUF2345 Fe-ADH ?
WP_048980141.1 NZ_JWAA01000053.1:6767-8625 <i>Enterobacter asburiae</i>	UPF0149 SEC-C Phage_GPD Tle3 ?
WP_058970807.1 NZ_BB0C01000021.1:56023-64226 <i>Type-D symbiont of Plautia stali</i>	LPAM_1 ab-Hydrolase Tle3 Phage_GPD VgrG T6SS_Vgr DUF2345 Abhydrolase_3 ?
WP_065559281.1 NZ_LZGN01000038.1:335536-343374 <i>Gilliamella sp. wkb308</i>	HTH ab-Hydrolase DUF3274 Tle3 ? ? Phage_GPD VgrG T6SS_Vgr DUF2345
WP_101077196.1 NZ_PITQ01000004.1:256730-263401 <i>Rahnella sp. AA</i>	? Tle3 Phage_GPD VgrG T6SS_Vgr DUF2345 PAAR_motif ?
WP_132964898.1 NZ_SLYT01000003.1:60374-66509 <i>Rahnella sp. Jub53</i>	Shikimate_dh_N MR_MLE_C ? Tle3 AAA_33
WP_150034503.1 NZ_VWUX01000002.1:214707-222564 <i>Pantoea sp. Bo_7</i>	PAAR_motif Tox-ABHYDROLASE2 Tle3 Phage_GPD VgrG T6SS_Vgr DUF2345 Sh1B POTRA_2 ?
WP_132964519.1 NZ_CAUX0010000004.1:311018-318300 <i>Rahnella variigena</i>	? Acetylase MarR-HTH Tox-ABHYDROLASE2 Tle3 PAAR_motif Phage_GPD VgrG T6SS_Vgr DUF2345
WP_132964519.1 NZ_SLYT01000002.1:408433-415715 <i>Rahnella sp. Jub53</i>	? Acetylase MarR-HTH Phage_GPD VgrG T6SS_Vgr DUF2345 Tox-ABHYDROLASE2 Tle3 PAAR_motif
WP_122096957.1 NZ_MJIB01000014.1:457353-464760 <i>Rahnella sp. Larv3_ips</i>	Phage_GPD VgrG T6SS_Vgr DUF2345 ? Sugar_tr Tox-ABHYDROLASE2 DUF3274 Tle3 ?
WP_130961908.1 NZ_SISY01000074.1:1-8050 <i>Hafnia paralvei</i>	ab-Hydrolase Tle3 Band-7 Flot Phage_GPD VgrG T6SS_Vgr DUF2345
WP_131390891.1 NZ_CP036514.1:1722349-1730664 <i>Hafnia alvei</i>	ClpABN-AAA ClpABC-AAA Phage_GPD VgrG T6SS_Vgr DUF2345 Tox-ABHYDROLASE2 DUF3274 Tle3 ? ?
WP_143440449.1 NZ_VJXS01000044.1:1-3765 <i>Klebsiella michiganensis</i>	T6SS_Vgr DUF2345 Tox-ABHYDROLASE2 DUF3274 Tle3 ?
WP_180246385.1 NZ_UNUE01000020.1:111-1706 <i>Enterobacter hormaechei</i>	? ? Tle3
WP_169402147.1 NZ_JAADJU010000003.1:20399-26248 <i>Rouxiiella aceris</i>	Peptidase_M55 LysR_substrate Tox-ABHYDROLASE2 DUF3274 Tle3 ? ?
WP_171936187.1 NZ_JABMCU010000068.1:1-2145 <i>Enterobacter sp. CM29</i>	Tle3 ? ?
WP_208725989.1 NZ_CP024637.1:654980-766 <i>Pantoea phytobeneficialis</i>	Transposase_20 PAAR_motif Tle3 HbpS-like ?