

EAA6174270.1
AAACAW010000083.1:1-1886
Salmonella enterica subsp. enterica serovar Hadar

MBA4156983.1
JACDHR010000047.1:361-6678
Gemmatimonadetes bacterium

MBK9260817.1
JADJXL010000015.1:7805-12210
Polyangiaceae bacterium

MCE4057325.1
JAJTTH010000010.1:257206-259193
Pseudomonas sp. Au-Pse12

MCG8700439.1
JAKSAV010001307.1:1-2153
Bacteroidales bacterium

MCR6678449.1
JANPXH010000073.1:1-1081
Escherichia marmotae

NCQ34600.1
JAACRO010000228.1:486-5468
bacterium Bacteria.

NIA00676.1
WHJI010000069.1:13717-21895
Massilia sp. CCM 8734

WP_010760834.1
NZ_CP172890.1:1202304-1205776
Enterococcus sp. DIV0724b

WP_024361236.1
NZ_AUOZ010000006.1:2497-5955
Lysinibacillus sphaericus KCTC 3346

WP_074905024.1
NZ_FONN010000034.1:16914-23344
Paenibacillus algorifonticola

WP_065565647.1
NZ_LZGL010000018.1:21956-25220
Gilliamella sp. wkb112

WP_073290058.1
NZ_FRCP010000020.1:51518-63372
Anaerosporbacter mobilis DSM 15930

WP_129538978.1
NZ_CP033068.1:2220585-2223822
Flavobacterium sp. 140616W15

WP_162309229.1
NZ_AP019400.1:1143574-1147666
Cohnella abietis

WP_228458479.1
NZ_BJYJ010000069.1:1-2535
Chryseobacterium hagamense

WP_151144873.1
NZ_WAGX010000005.1:736921-741648
Candidatus Galacturonibacter soehngeni

WP_181345119.1
NZ_JABXNL010000007.1:352345-354999
Listeria monocytogenes

WP_185636473.1
NZ_JAARZD010000012.1:75208-79334
Listeria ivanovii

WP_228055720.1
NZ_JADEWZ010000022.1:29797-41067
Lusitaniella coriacea LEGE 07157

WP_255561391.1
NZ_CP048263.1:1039107-1041402
Gilliamella sp. ESL0443

WP_236857756.1
NZ_CP084580.1:1157226-1160713
Chryseobacterium sp. MEB0G06

RHS STox-14* ? DUF6756

AAA-ToxA ? MPTase-PVC STox-14* ? Metallopeptidase Aspzincin_M35

? ? RCC1_2 STox-14* ? Y2_Tnp Zn_Tnp_IS91

? ? RHS STox-14*

? RHS RHS STox-14*

RHS STox-14* ?

PVC-AAA DUF4157 STox-14* ?

Phage_GPD VgrG Gp5_C DcrB PrePAARTM PAAR_motif DUF6531 RHS RHS RHS RHS_repeat RHS STox-14* ? ?

Bactofilin ? STox-14* ? ?

Tox-AHH STox-14* ? ? ?

? ? RHS RHS HINT STox-14* RHS_repeat RHS_repeat ?

RHS Ntox12 STox-14* SUKH_5 SUKH-SMI1 ?

4HB_MCP_1 HAMP MA ? FlgD_ig DUF6531 RHS_repeat CBM_3 CBM_3 CBM_3 RHS_repeat RHS_repeat RHS_repeat RHS_repeat RHS RHS_repeat RHS RHS STox-14* ? RHS

Imm11 ? STox-14* ? Ntox41

Melibiose Glyco_hydro_36C ? Imm73 STox-14* HINT

? LINKER Tox-NucA STox-14*

? ? STox-14* Phage_integrase DEDD_Tnp_IS110 Transposase_20

? STox-14* ? ? Imm38

? STox-14* ? ? Ntox29

? ? ? ? HfIG IG IG He_PIG Big_7 IG DUF6531 RHS_repeat RHS RHS RHS RHS_repeat RHS HINT STox-14*

? ? ? ? STox-14*

? ? HTH_3 RHS STox-14* ?