AAR03010.1
CP009458:1:1220408-1224793
Pluralibacter gergoviae
HTM_21:rve:rve
Rdh
?
?
Tox-HNHGIY
AAX70024.1
CP003776.1:4011949-4010042
Chryseobacterium gallinarum
Mur1N-PTD Acetylase
Tox-HNHGIY
AUP70284.1
CP025776.1:383124-388328
Enterobacter sp. EA-2
Ecp
EAS15605.1
AAON:1010000114.1:46-2606
Salmonella enterica
Tfb
Ant
HNHGIY
?
Rdh
EBA20750A1
AAEBEN01000002.1:79145-80540
Salmonella enterica
ApaK_toxin
Tfb
Ant
HNHGIY
EBA70866G1
DAEFAA01000158.1:305-3502
Escherichia coli
?
Tfb
Ant
HNHGIY
Rdh
HEA17151.1
UNUMG:1001236.1:1-1191
Pseudoalteromonas prydzensis
Tox-HNHGIY
TPR_Mal
Tfb
KQ002065.1
LRQHE:1000011.1:1005960-1006409
Chryseobacterium sp. Leaf394
Tfb
Sup
Tox-HNHGIY
?
?
KVE5K301.1
LDWJ:0100039.1:269159-271192
Burkholderia vietnamensis
Tox-PAAR-like
Tox-ABHYDROLASE3
Abh
Tox-HNHGIY
?
Tfb
WBU14G000.1
JACJCB:1000007.1:158-3701
Pseudoalteromonas sp. SG43-8
SUKA
Tfb
Tox-HNHGIY
Tfb
Tfb
WBU5G4G0.1
JAAHAB:1000011.1:82235-90977
Treponema sp.
CBM_4B_Alpha-amylase Isoamylase
4MDC_MDC_RNA-Helicase
Tox-HNHGIY
?
AMP-binding
WBU9Q04A1.
JACQAM:1000006.1:388-11345
Burkholderia bacterium
Tfb
?
Tox-HNHGIY
Cadherin
?
WBU5T12E0.1
JACDMA:1000050.1:11627-16164
Burkholderia bacterium
?
TPR_Mal
Tox-HNHGIY
WBU9X6G0A1.1
JACDMA:1000050.1:13526-56021
Acidobacteria bacterium
?
IG-Glucodextran_B-Glucodextran_B-Glucodextran_B-Glucodextran_B-Glucodextran_B-Glucodextran_B-Glucodextran_B-Glucodextran_B
Big_3
Big_3
Carboxypep_reg
Big_3
DUF5353_MMS_MDS
Rdh
Tox-HNHGIY
Tfb
IG_Tfb
WBU9C9T0A1
JAEMEE:1000003.1:406416-412098
Flavobacteria bacterium
?
Sup
Tox-HNHGIY-2
Tfb
?
?
WBU9A04A1.
JAEUDA:1000009.1:26245-41167
Leptospirales bacterium
?
?
Tox-HNHGIY
Ribonuclease
?
?
WBU8J4R0E1
JAEUDJ:0100039.1:2899-8725
Blastocatella bacterium
ABC_trim
CT-kinase STY-kinase
PT-TS
Tox-HNHGIY
?
CpA_binding_2
WBU8K3G0.1
JAEUDL:0100126.1:1-4520
Gemmatacer sp.
Sup
DUF2169
Tox-PAAR-like
HNHGIY
?
?
WBUPD5Q4A1
JACPPH:1000033.1:2369-8854
Latrys sp. LI14
DUF6351
DUF3836
RMS_repeat
RMS_repeat
RMS_repeat
RMS_repeat
RMS_repeat
Tox-Lin
RMS
RMS
HNHGIY
?
?
WBF1H0R0A.1
JAGAKZ:1000009.1:1248-5082
Oscillospiraceae bacterium
Tox-HNH-DHHW
Tfb
CBM
HNHGIY
RMS
DNRT
Tox-Tnh
WBS1K0E0E0.1
JAFPYU:0100006.1:225848-238850
Bacteroidetes bacterium
MFTase-PG
Sup
HNHGIY
CIS_Tfb
?
?
WBS0R0D4A1.1
JAGZMB:1000002.1:3583-9284
Ruminococcus sp.
Glycl-like
Tfb
RMS_repeat
RMS_repeat
RMS_repeat
RMS_repeat
RMS_repeat
RMS
RMS
RMS
HNHGIY
?
WBS0T7R0A.1
JAGMAB:1000050.1:479-4704
Tenacibaculum sp.
Phage_GPD
Phage_base_V
DUF2169
DUF4280
DNRT
Tox-HNHGIY
Tfb
Tfb
WBV0R0R0E.1
JAFAUY:1000032.1:1-3551
Verrucomicrobiota bacterium
Tox-HNHGIY
Sup
NPCBM-Methyltransf_2a
WBX2R0R0E.1
JAWMVU:0100001.1:6472-16268
Cytophaginaceae bacterium
?
Tfb
MFTase-PVC
Rhs
HNHGIY
PVC-AAA
?
WBV8R7R2E1.1
JAIIPT:0100001.1:1186-6637
Gemmatacae bacterium
?
STY-kinase LMK_LBh
SEC-C
DNRT
Tox-HNHGIY
Tfb
?
WCC7R0D0E.1
JAQZJE:0100030.1:183-3633
Crocinitomix sp.
?
RMS
RMS
HNHGIY
WAP4R0G1.1
WTE4G:0100007.1:88520-82739
Gillisella sp. Lep-135
?
?
DNRT
Tox-HNHGIY
?
Tfb
NJL1R7A6.1
JAAXRA:0100004.1:508-3004
Mitrospira sp.
HNHGIY
?
Rhs
HNHGIY
NLE3T0D6.1
JAATZP:0100002.1:332-2351
Propionibacterium sp.
AHBE_1
Tfb
HNHGIY
?
NVZ5Z7A4.1
JACAWW:0100002.1:40-346
Pseudomonas gingeri
HNHGIY
?
NWWR0C7L.1
JACAWW:0100004.1:401232-406213
Pseudomonas gingeri
?
?
Tox-HNHGIY
?
Tox-CELAC
4NH-DHHW
OE2R9W8.1
LRDME:0000050.1:180-10649
Bugaella sp. MH02
GlcD_B GlcD_B GlcD_B