amidase (CDS_0017)
NinG/ Rap DNA junction specific endonuclease (CDS_0035)
OTU_2487 16213:30748(+)
IMGVR_UViG_2609460318_000002|2609460318|2609562437|31601-70446 16213:30644(+)
CDS_0019
CDS_0020
CDS_0021
MGV-GENOME-0250822 16213:24124(+)
TemPhD_cluster_41892 17065:33105(+)
IMGVR_UViG_2558309115_000001|2558309115|2558321523|416128-457047 17065:32724(+)
ivig_2550 17065:30734(+)
IMGVR_UViG_2568526105_000005|2568526105|2568547507|1-37573 17065:29377(+)
MGV-GENOME-0242690 17065:25235(+)
multidrug efflux MATE transporter NorM (NF000263.1) (CDS_0027)
HAD-superfamily hydrolase (CDS_0022)
uvig_572430 17065:39601(+)
IMGVR_UViG_3300029458_000042|3300029458|Ga0244161_100443 16213:28430(+)
MGV-GENOME-0250847 16171:26137(+)
IMGVR_UViG_3300029724_000112|3300029724|Ga0245182_100004|91210-128561 16254:28743(+)
tail fiber protein (CDS_0013)
DNA methyltransferase (CDS_0029)
MGV-GENOME-0245346 12551:25843(+)
CDS_0019
CDS_0020
CDS_0021
MGV-GENOME-0296528 12551:23998(+)
IMGVR_UViG_3300029757_000057|3300029757|Ga0243876_100404 12551:30393(+)
IMGVR_UViG_3300029684_000310|3300029684|Ga0245194_100019|165894-205212 12551:31309(+)
IMGVR_UViG_3300029545_000111|3300029545|Ga0245039_100048|1058-40373 12551:31307(+)
RosmerTA__RmrA_2677172838 (CDS_0020)
endolysin (CDS_0017)
IMGVR_UViG_7000000317_000221|7000000317|SRS014459_WUGC_scaffold_92721 12650:27691(+)
RosmerTA__RmrA_2677172838 (CDS_0020)
IMGVR_UViG_3300007742_000435|3300007742|Ga0105758_1000216|36161-75495 12650:26857(+)
IMGVR_UViG_2687453203_000003|2687453203|2687453642|5130749-5171730 17067:30194(+)
IMGVR_UViG_3300029198_000123|3300029198|Ga0168723_100020|12990-50181 17067:26404(+)
IMGVR_UViG_3300039340_000057|3300039340|Ga0169801_00011|164503-204120 16213:28205(+)
TemPhD_cluster_24454 16269:28509(+)
IMGVR_UViG_3300039345_000017|3300039345|Ga0169235_0007|193279-231480 16269:24508(+)
IMGVR_UViG_3300029076_000066|3300029076|Ga0169233_100205|18494-55785 12650:23598(+)
IMGVR_UViG_3300045988_136877|3300045988|Ga0495776_167723 16269:22672(+)
CDS_0019
CDS_0020
MGV-GENOME-0212026 16213:20985(+)
uvig_313006 12650:29553(+)
IMGVR_UViG_2576861300_000005|2576861300|2576881918|1773-39297 16170:29328(+)
ivig_356 16170:25354(+)
RosmerTA__RmrA_2600853143 (CDS_0021)
uvig_440246 12551:28037(+)
CDS_0020
CDS_0019
CDS_0021
MGV-GENOME-0287850 17065:24749(+)
uvig_212608 17107:27617(+)
9000 nt
DNA, RNA and nucleotide metabolism
lysis
other
tail
unknown function
Defence
AMR