ParA-like partition protein (CDS_0039)
transcriptional regulator (CDS_0036)
RT_III-AM_III-B_2 (CDS_0028)
ParB-like partition protein (CDS_0038)
transposase (CDS_0033)
transposase (CDS_0037)
Retron_VI__HTH (CDS_0050)
DarB-like antirestriction (CDS_0034)
IMGVR_UViG_3300029700_000076|3300029700|Ga0245207_100383 23400:49615(+)
ParA-like partition protein (CDS_0038)
transcriptional regulator (CDS_0035)
ParB-like partition protein (CDS_0037)
transposase (CDS_0036)
transposase (CDS_0033)
MGV-GENOME-0341382 23400:44747(+)
ParA-like partition protein (CDS_0057)
transcriptional regulator (CDS_0054)
ParB-like partition protein (CDS_0056)
integrase (CDS_0061)
SoFic__SoFic (CDS_0042)
transposase (CDS_0033)
transposase (CDS_0055)
DarB-like antirestriction (CDS_0052)
MGV-GENOME-0363531 23400:65140(+)
ParA-like partition protein (CDS_0057)
transcriptional regulator (CDS_0054)
ParB-like partition protein (CDS_0056)
SoFic__SoFic (CDS_0042)
transposase (CDS_0033)
transposase (CDS_0055)
DNA methyltransferase (CDS_0052)
IMGVR_UViG_3300045988_152239|3300045988|Ga0495776_158574 23400:65140(+)
CDS_0033
CDS_0034
SAMEA2737747_b1_ct60_vs2 23400:36313(+)
CDS_0033
CDS_0035
CDS_0034
CDS_0036
CDS_0044
IMGVR_UViG_3300045988_126434|3300045988|Ga0495776_037242 23400:35489(+)
ParB-like partition protein (CDS_0036)
ParA-like partition protein (CDS_0035)
DarB-like antirestriction (CDS_0046)
glycopeptide resistance transcriptional regulator VanU (NF000090.1) (CDS_0056)
transcriptional regulator (CDS_0055)
IMGVR_UViG_3300045988_186729|3300045988|Ga0495776_122560|1-84165 23400:61862(+)
DNA primase (CDS_0040)
ParB-like partition protein (CDS_0036)
ParA-like partition protein (CDS_0035)
DarB-like antirestriction (CDS_0046)
glycopeptide resistance transcriptional regulator VanU (NF000090.1) (CDS_0055)
transcriptional regulator (CDS_0054)
integrase (CDS_0058)
uvig_467692 23400:58960(+)
ParB-like partition protein (CDS_0036)
ParA-like partition protein (CDS_0035)
DNA methyltransferase (CDS_0046)
metal-dependent phosphohydrolase (CDS_0057)
IMGVR_UViG_3300045988_128573|3300045988|Ga0495776_072274 23400:51715(+)
AAC(3)-I family aminoglycoside 3-N-acetyltransferase (NF033083.0) (CDS_0037)
Sigma M (VF0822) (CDS_0034)
OTU_615 23400:38417(+)
MGV-GENOME-0329210 23400:35940(+)
10000 nt
DNA, RNA and nucleotide metabolism
integration and excision
lysis
moron, auxiliary metabolic gene and host takeover
other
transcription regulation
unknown function
Defence
Virulence
AMR