MGV-GENOME-0233163 17287:26064(+)
endolysin (CDS_0024)
DNA primase (CDS_0026)
FtsK/SpoIIIE-like protein (CDS_0028)
MGV-GENOME-0183186 17287:24836(+)
FtsK/SpoIIIE-like protein (CDS_0027)
MGV-GENOME-0196804 17386:26269(+)
DNA primase (CDS_0027)
FtsK/SpoIIIE-like protein (CDS_0029)
MGV-GENOME-0191194 17287:26121(+)
DNA primase (CDS_0026)
FtsK/SpoIIIE-like protein (CDS_0028)
MGV-GENOME-0209268 17386:28487(+)
anaerobic ribonucleoside reductase large subunit (CDS_0025)
DNA primase/helicase (CDS_0028)
DNA primase (CDS_0031)
FtsK/SpoIIIE-like protein (CDS_0033)
MGV-GENOME-0202711 17286:28134(+)
DNA primase/helicase (CDS_0026)
DNA primase (CDS_0030)
FtsK/SpoIIIE-like protein (CDS_0032)
MGV-GENOME-0208486 17302:29156(+)
anaerobic ribonucleoside reductase large subunit (CDS_0025)
DNA primase/helicase (CDS_0028)
DNA primase (CDS_0032)
FtsK/SpoIIIE-like protein (CDS_0034)
MGV-GENOME-0231988 17389:28032(+)
DNA primase/helicase (CDS_0028)
DNA primase (CDS_0031)
FtsK/SpoIIIE-like protein (CDS_0033)
MGV-GENOME-0192223 17307:25672(+)
DNA primase (CDS_0025)
FtsK/SpoIIIE-like protein (CDS_0027)
MGV-GENOME-0233110 17287:30606(+)
NrdD-like anaerobic ribonucleotide reductase large subunit (CDS_0025)
DNA primase/helicase (CDS_0028)
DNA primase (CDS_0031)
FtsK/SpoIIIE-like protein (CDS_0033)
MGV-GENOME-0169929 18524:22893(+)
CDS_0025
CDS_0027
MGV-GENOME-0177398 17390:23148(+)
CDS_0026
CDS_0025
MGV-GENOME-0170763 17305:22981(+)
CDS_0025
CDS_0026
MGV-GENOME-0168832 17287:22703(+)
FtsK/SpoIIIE-like protein (CDS_0025)
CDS_0026
CDS_0027
MGV-GENOME-0174510 17305:23520(+)
CDS_0025
MGV-GENOME-0178649 17403:24141(+)
FtsK/SpoIIIE-like protein (CDS_0025)
CDS_0026
CDS_0027
MGV-GENOME-0176728 17390:23855(+)
FtsK/SpoIIIE-like protein (CDS_0025)
CDS_0026
CDS_0027
MGV-GENOME-0192259 17284:23099(+)
FtsK/SpoIIIE-like protein (CDS_0025)
CDS_0026
CDS_0027
MGV-GENOME-0222792 17305:23342(+)
FtsK/SpoIIIE-like protein (CDS_0029)
DarB-like antirestriction (CDS_0025)
DarB-like antirestriction (CDS_0027)
Reverse transcriptase (CDS_0026)
MGV-GENOME-0198497 17287:27390(+)
FtsK/SpoIIIE-like protein (CDS_0026)
CDS_0027
DarB-like antirestriction (CDS_0025)
CDS_0028
MGV-GENOME-0175371 17287:23647(+)
CDS_0025
SRS047433_a1_ct17824_vs1 16924:21205(+)
MGV-GENOME-0181371 17299:24554(+)
FtsK/SpoIIIE-like protein (CDS_0025)
CDS_0026
CDS_0027
MGV-GENOME-0175824 17389:22905(+)
FtsK/SpoIIIE-like protein (CDS_0025)
CDS_0026
CDS_0027
MGV-GENOME-0230507 17370:22673(+)
CDS_0026
CDS_0025
CDS_0027
MGV-GENOME-0173795 17370:22605(+)
MGV-GENOME-0200636 17287:24605(+)
FtsK/SpoIIIE-like protein (CDS_0029)
MGV-GENOME-0199949 17287:27634(+)
MGV-GENOME-0195223 17287:26798(+)
FtsK/SpoIIIE-like protein (CDS_0029)
MGV-GENOME-0196547 17284:27036(+)
RM_Type_III__Type_III_MTases___Type_III_MTases_FAM_0 (CDS_0028)
uvig_263731 17287:30819(+)
FtsK/SpoIIIE-like protein (CDS_0031)
MGV-GENOME-0185508 16605:25197(+)
CDS_0025
CDS_0024
CDS_0026
CDS_0027
Reverse transcriptase (CDS_0029)
FtsK/SpoIIIE-like protein (CDS_0030)
CDS_0031
MGV-GENOME-0194902 16605:26747(+)
tetracycline resistance ribosomal protection protein (NF012153.1) (CDS_0029)
CDS_0025
CDS_0027
CDS_0024
CDS_0026
CDS_0032
CDS_0033
MGV-GENOME-0197054 16628:27127(+)
CDS_0025
CDS_0027
CDS_0024
CDS_0026
CDS_0029
MGV-GENOME-0191315 16628:26138(+)
IlpA (VF0513) (CDS_0026)
FtsK/SpoIIIE-like protein (CDS_0033)
uvig_517937 17305:31710(+)
DarB-like antirestriction (CDS_0025)
DarB-like antirestriction (CDS_0027)
Reverse transcriptase (CDS_0026)
MGV-GENOME-0202250 17287:28051(+)
CDS_0032
CDS_0033
CDS_0034
CDS_0035
uvig_179975 23468:33235(+)
5000 nt
DNA, RNA and nucleotide metabolism
lysis
moron, auxiliary metabolic gene and host takeover
unknown function
Defence
Virulence
AMR