anti-repressor (CDS_0039)
CDS_0042
CDS_0041
CDS_0044
CDS_0037
CDS_0032
tail fiber protein (CDS_0030)
CDS_0031
endolysin (CDS_0033)
CDS_0034
CDS_0035
CDS_0036
CDS_0038
CDS_0040
CDS_0043
uvig_232012 23961:33907(+)
anti-repressor Ant (CDS_0037)
endolysin (CDS_0031)
uvig_214745 22338:32892(+)
anti-repressor Ant (CDS_0039)
MGV-GENOME-0290472 22921:33857(+)
anti-repressor Ant (CDS_0040)
IMGVR_UViG_3300045988_003668|3300045988|Ga0495776_012746 22952:33800(+)
anti-repressor Ant (CDS_0039)
uvig_146985 23534:32966(+)
transcriptional regulator (CDS_0038)
anti-repressor Ant (CDS_0041)
endolysin (CDS_0034)
uvig_438146 22952:33855(+)
IMGVR_UViG_3300045988_004145|3300045988|Ga0495776_014382 24272:34105(+)
TemPhD_cluster_29378 22472:32373(+)
MGV-GENOME-0291557 21998:31872(+)
MGV-GENOME-0296464 23891:33493(+)
superinfection exclusion Sie-like (CDS_0035)
CDS_0037
CDS_0039
endolysin (CDS_0033)
CDS_0036
CDS_0038
CDS_0040
uvig_37960 27415:34631(+)
IMGVR_UViG_3300045988_004211|3300045988|Ga0495776_020866 24501:33513(+)
MGV-GENOME-0290079 24446:33577(+)
IMGVR_UViG_3300045988_014613|3300045988|Ga0495776_016025 24230:34563(+)
MGV-GENOME-4421322 23549:32740(+)
anti-repressor (CDS_0042)
uvig_182926 23218:34179(+)
MADS_mad1 (CDS_0037)
anti-repressor Ant (CDS_0039)
IMGVR_UViG_3300045988_004208|3300045988|Ga0495776_012726 27466:35267(+)
MADS_mad1 (CDS_0039)
anti-repressor Ant (CDS_0043)
IMGVR_UViG_3300045988_005010|3300045988|Ga0495776_098273 24949:35638(+)
MGV-GENOME-0285986 25156:33565(+)
MGV-GENOME-0294452 27025:34517(+)
IMGVR_UViG_3300045988_015444|3300045988|Ga0495776_015643 27694:35889(+)
MGV-GENOME-0295761 27694:35889(+)
MGV-GENOME-0291061 27694:34843(+)
Retron_VI__HTH (CDS_0042)
uvig_251875 25430:35366(+)
uvig_333409 25517:35249(+)
Retron_VI__HTH (CDS_0039)
uvig_153482 23596:33652(+)
Retron_VI__HTH (CDS_0037)
IMGVR_UViG_3300045988_002919|3300045988|Ga0495776_139702 22339:31987(+)
Retron_VI__HTH (CDS_0043)
IMGVR_UViG_3300045988_008551|3300045988|Ga0495776_014469 25855:36142(+)
transcriptional regulator (CDS_0037)
Retron_VI__HTH (CDS_0038)
anti-repressor Ant (CDS_0041)
uvig_176887 24895:35339(+)
Retron_VI__HTH (CDS_0034)
IMGVR_UViG_3300045988_015154|3300045988|Ga0495776_014017 23704:33063(+)
IMGVR_UViG_3300045988_017351|3300045988|Ga0495776_137994 24796:33640(+)
MGV-GENOME-0293387 25830:36089(+)
transcriptional regulator (CDS_0041)
IMGVR_UViG_3300045988_004375|3300045988|Ga0495776_015325 25498:35109(+)
transcriptional regulator (CDS_0036)
uvig_384152 23353:33429(+)
MGV-GENOME-0297999 23310:33456(+)
MGV-GENOME-0295852 24016:34261(+)
transcriptional regulator (CDS_0035)
OTU_1240 23462:34063(+)
IMGVR_UViG_3300045988_008558|3300045988|Ga0495776_006662 23462:33820(+)
IMGVR_UViG_3300045988_008225|3300045988|Ga0495776_006659 23462:33714(+)
transcriptional regulator (CDS_0041)
anti-repressor Ant (CDS_0043)
SAMEA2737819_b1_ct123_vs2 26136:35534(+)
transcriptional regulator (CDS_0042)
SRS017521_a1_ct28933_vs01 31898:37418(+)
transcriptional regulator (CDS_0039)
IMGVR_UViG_3300045988_004251|3300045988|Ga0495776_137747 23701:33591(+)
MGV-GENOME-0294716 25943:36354(+)
4000 nt
DNA, RNA and nucleotide metabolism
integration and excision
lysis
moron, auxiliary metabolic gene and host takeover
tail
transcription regulation
unknown function
Defence