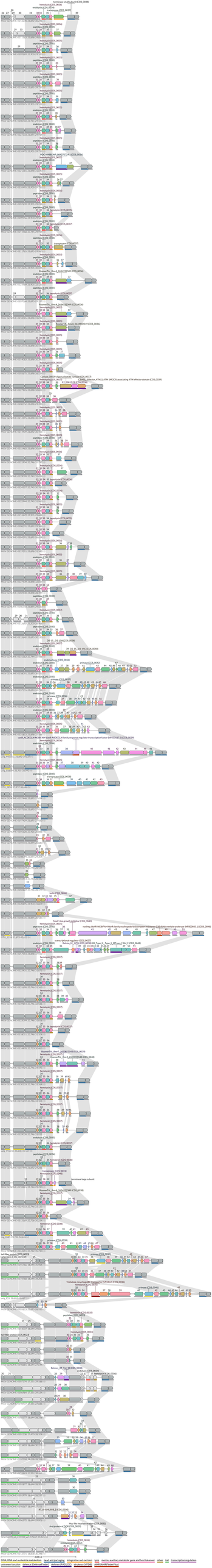

terminase small subunit (CDS_0038)
hemolysin (CDS_0036)
endolysin (CDS_0034)
transposase (CDS_0037)
MGV-GENOME-0213278: 21,892-30,246 (+)
hemolysin (CDS_0036)
peptidase (CDS_0034)
MGV-GENOME-0211162: 21,898-29,850 (+)
hemolysin (CDS_0035)
peptidase (CDS_0033)
MGV-GENOME-0209289: 21,900-29,523 (+)
hemolysin (CDS_0035)
peptidase (CDS_0033)
MGV-GENOME-0211245: 21,901-29,865 (+)
hemolysin (CDS_0035)
peptidase (CDS_0033)
MGV-GENOME-0210442: 21,895-29,727 (+)
hemolysin (CDS_0035)
peptidase (CDS_0033)
MGV-GENOME-0209225: 21,901-29,512 (+)
hemolysin (CDS_0035)
endolysin (CDS_0033)
MGV-GENOME-0208760: 21,911-29,431 (+)
endolysin (CDS_0033)
MGV-GENOME-0227303: 21,895-29,919 (+)
hemolysin (CDS_0035)
peptidase (CDS_0033)
MGV-GENOME-0210472: 21,901-29,732 (+)
PDC-M48B_WP_084275724.1 (CDS_0036)
hemolysin (CDS_0035)
endolysin (CDS_0033)
MGV-GENOME-0221381: 21,898-31,690 (+)
hemolysin (CDS_0035)
peptidase (CDS_0033)
MGV-GENOME-0210495: 21,900-29,735 (+)
hemolysin (CDS_0035)
peptidase (CDS_0033)
MGV-GENOME-0207639: 21,901-29,234 (+)
peptidase (CDS_0033)
hemolysin (CDS_0035)
MGV-GENOME-0209556: 21,900-29,567 (+)
endolysin (CDS_0035)
hemolysin (CDS_0037)
MGV-GENOME-0211568: 23,745-29,919 (+)
hemolysin (CDS_0036)
peptidase (CDS_0034)
transposase (CDS_0037)
MGV-GENOME-0211785: 21,895-29,960 (+)
hemolysin (CDS_0035)
peptidase (CDS_0033)
MGV-GENOME-0212235: 21,898-30,044 (+)
RosmerTA__RmrA_2634932349 (CDS_0036)
endolysin (CDS_0033)
MGV-GENOME-0223989: 21,901-32,125 (+)
peptidase (CDS_0035)
hemolysin (CDS_0037)
MGV-GENOME-0207064: 21,895-29,137 (+)
RosmerTA__RmrA_2634932349 (CDS_0036)
hemolysin (CDS_0033)
MGV-GENOME-0218585: 21,898-31,212 (+)
hemolysin (CDS_0035)
RosmerTA__RmrA_2634932349 (CDS_0036)
MGV-GENOME-0213673: 21,894-30,317 (+)
hemolysin (CDS_0035)
MGV-GENOME-0212773: 21,898-30,145 (+)
hemolysin (CDS_0035)
MGV-GENOME-0214082: 21,898-30,392 (+)
hemolysin (CDS_0035)
MGV-GENOME-0203675: 21,898-28,529 (+)
Cyclase_00020 oligonucleotide-cyclase (CDS_0037)
hemolysin (CDS_0035)
CBASS__Effector_4TM_S_4TM SMODS-associating 4TM effector domain (CDS_0039)
E2_00013 E2 (CDS_0038)
MGV-GENOME-0227965: 21,898-32,821 (+)
MGV-GENOME-0208552: 21,898-29,393 (+)
hemolysin (CDS_0035)
MGV-GENOME-0214195: 21,898-30,414 (+)
hemolysin (CDS_0035)
MGV-GENOME-0214409: 21,892-30,453 (+)
hemolysin (CDS_0035)
peptidase (CDS_0033)
MGV-GENOME-0215482: 21,898-30,647 (+)
hemolysin (CDS_0036)
MGV-GENOME-0221904: 23,746-31,797 (+)
hemolysin (CDS_0036)
MGV-GENOME-0229090: 23,742-32,418 (+)
hemolysin (CDS_0036)
MGV-GENOME-0234227: 23,742-30,761 (+)
hemolysin (CDS_0036)
MGV-GENOME-0222944: 23,742-31,963 (+)
hemolysin (CDS_0035)
MGV-GENOME-0212629: 21,901-29,688 (+)
hemolysin (CDS_0035)
MGV-GENOME-0208501: 21,892-29,386 (+)
MGV-GENOME-0204391: 21,892-28,654 (+)
hemolysin (CDS_0035)
MGV-GENOME-0219311: 21,898-31,343 (+)
hemolysin (CDS_0035)
endolysin (CDS_0033)
MGV-GENOME-0217035: 21,892-30,936 (+)
hemolysin (CDS_0035)
MGV-GENOME-0227939: 21,898-32,817 (+)
MGV-GENOME-0202930: 21,898-28,397 (+)
peptidase (CDS_0033)
MGV-GENOME-0206350: 21,895-29,020 (+)
hemolysin (CDS_0037)
peptidase (CDS_0035)
MGV-GENOME-0212692: 21,200-28,968 (+)
peptidase (CDS_0033)
MGV-GENOME-0212642: 21,895-30,120 (+)
DS-15__DS-15A (CDS_0038)
hemolysin (CDS_0037)
endolysin (CDS_0035)
DS-15__DS-15C (CDS_0040)
MGV-GENOME-0231362: 23,744-33,488 (+)
endonuclease (CDS_0036)
endolysin (CDS_0033)
primase (CDS_0042)
MGV-GENOME-0258091: 21,898-38,021 (+)
endolysin (CDS_0033)
primase (CDS_0037)
MGV-GENOME-0237555: 21,894-34,718 (+)
endolysin (CDS_0033)
primase (CDS_0036)
MGV-GENOME-0235279: 21,872-34,270 (+)
endolysin (CDS_0033)
MGV-GENOME-0223761: 21,900-32,090 (+)
hemolysin (CDS_0035)
MGV-GENOME-0231852: 21,898-33,588 (+)
vanR_ACDEGLN-NCBIFAM VanR-ABDEGLN family response regulator transcription factor (NF033117.2) (CDS_0039)
hemolysin (CDS_0036)
endolysin (CDS_0034)
uvig_445356_:21,892-39,637 (+)
hemolysin (CDS_0035)
uvig_118304: 21,955-33,993 (+)
peptidase (CDS_0035)
transposase (CDS_0038)
OTU_2896: 23,937-36,646 (+)
MGV-GENOME-0197448: 21,892-27,424 (+)
MGV-GENOME-0197518: 21,892-27,436 (+)
MGV-GENOME-0197429: 21,895-27,420 (+)
MGV-GENOME-0205628: 23,426-28,888 (+)
MGV-GENOME-0204514: 21,894-28,680 (+)
MGV-GENOME-0208094: 21,895-29,304 (+)
MGV-GENOME-0203179: 23,746-28,439 (+)
MGV-GENOME-0203413: 23,743-28,482 (+)
MGV-GENOME-0202822: 23,746-28,377 (+)
holin (CDS_0036)
MGV-GENOME-0220895: 21,900-31,609 (+)
MGV-GENOME-0204001: 21,895-28,585 (+)
MazF-like growth inhibitor (CDS_0040)
hemolysin (CDS_0035)
peptidase (CDS_0033)
nosihep_NshR-NCBIFAM NshR family nosiheptide/thiostrepton resistance 23S rRNA methyltransferase (NF000035.1) (CDS_0048)
OTU_1518: 21,893-43,612 (+)
transcriptional regulator (CDS_0037)
endolysin (CDS_0033)
Retron_VI__HTH (CDS_0038)
RM_Type_II__Type_II_MTases_FAM_2 (CDS_0048)
MGV-GENOME-0257514: 21,898-37,922 (+)
hemolysin (CDS_0037)
MGV-GENOME-0222431: 23,746-31,872 (+)
hemolysin (CDS_0037)
MGV-GENOME-0231698: 23,746-31,565 (+)
hemolysin (CDS_0037)
MGV-GENOME-0239814: 23,746-32,586 (+)
hemolysin (CDS_0037)
MGV-GENOME-0222099: 23,746-31,818 (+)
hemolysin (CDS_0037)
MGV-GENOME-0218011: 23,746-31,104 (+)
MGV-GENOME-0214478: 23,733-30,465 (+)
RosmerTA__RmrT_2600853143 (CDS_0039)
hemolysin (CDS_0037)
RosmerTA__RmrA_2600853143 (CDS_0040)
MGV-GENOME-0234924: 23,746-34,195 (+)
hemolysin (CDS_0037)
MGV-GENOME-0230233: 23,746-33,264 (+)
hemolysin (CDS_0037)
MGV-GENOME-0235259: 23,746-33,101 (+)
hemolysin (CDS_0037)
MGV-GENOME-0230736: 23,746-33,362 (+)
hemolysin (CDS_0037)
MGV-GENOME-0229030: 23,746-33,031 (+)
endolysin (CDS_0035)
hemolysin (CDS_0037)
uvig_155650: 23,604-31,167 (+)
peptidase (CDS_0034)
hemolysin (CDS_0036)
MGV-GENOME-0207944: 21,892-29,281 (+)
transposase (CDS_0041)
hemolysin (CDS_0040)
terminase large subunit
MGV-GENOME-0210865: 21,744-29,290 (+)
RosmerTA__RmrA_2634932349 (CDS_0038)
hemolysin (CDS_0037)
MGV-GENOME-0229553: 23,746-33,135 (+)
MGV-GENOME-0213169: 23,738-30,224 (+)
hemolysin (CDS_0038)
uvig_2681: 23,742-34,626 (+)
primase (CDS_0039)
MGV-GENOME-0244211: 23,742-35,730 (+)
tail fiber protein (CDS_0023)
tail protein (CDS_0022)
primase (CDS_0037)
MGV-GENOME-0245766: 18,644-35,964 (+)
MGV-GENOME-0274768: 18,644-34,208 (+)
Trehalose-recycling ABC transporter (VF0842) (CDS_0036)
hemolysin (CDS_0035)
primase (CDS_0041)
uvig_255: 18,644-41,087 (+)
MGV-GENOME-0205813: 21,895-28,925 (+)
hemolysin (CDS_0035)
peptidase (CDS_0033)
MGV-GENOME-0211007: 18,644-29,826 (+)
tail fiber protein (CDS_0023)
hemolysin (CDS_0034)
MGV-GENOME-4405332: 18,644-29,658 (+)
MGV-GENOME-0198674: 18,644-27,643 (+)
MGV-GENOME-0198681: 18,644-27,644 (+)
Retron__RT_Tot_10 (CDS_0030)
endolysin (CDS_0034)
hemolysin (CDS_0036)
MGV-GENOME-0207374: 17,151-29,188 (+)
MGV-GENOME-0206984: 17,151-29,124 (+)
MGV-GENOME-0198907: 17,151-27,681 (+)
MGV-GENOME-0202577: 18,644-28,338 (+)
MGV-GENOME-0201308: 17,470-28,104 (+)
MGV-GENOME-0213419: 18,644-30,272 (+)
MGV-GENOME-0202524: 18,628-28,328 (+)
MGV-GENOME-0259443: 23,912-38,261 (+)
MGV-GENOME-0201877: 18,644-28,208 (+)
MGV-GENOME-0201979: 18,644-28,227 (+)
MGV-GENOME-0198167: 18,644-27,551 (+)
RT_III-AM_III-B_2 (CDS_0030)
MGV-GENOME-0206272: 18,649-29,006 (+)
Hoc-like head decoration (CDS_0032)
Avd protein of DGR (CDS_0029)
SRS058070_a1_ct13313_vs1: 17,657-31,145 (+)
hemolysin (CDS_0034)
endolysin (CDS_0032)
MGV-GENOME-0213771: 18,838-30,348 (+)
8000 nt
DNA, RNA and nucleotide metabolism
head and packaging
integration and excision
lysis
moron, auxiliary metabolic gene and host takeover
other
tail
transcription regulation
unknown function
defence (DefenseFinder)
defence (PADLOC)
virulence (VFDB)
AMR (AMRFinderPlus)