CDS_0029
cysH_IV-B_1 (CDS_0038)
CDS_0028
CDS_0034
CDS_0030
CDS_0033
CDS_0037
CDS_0026
CDS_0031
CDS_0036
CDS_0040
CDS_0027
CDS_0032
CDS_0035
CDS_0039
MGV-GENOME-0223806 21779:32194(+)
CDS_0033
CDS_0031
CDS_0034
CDS_0032
MGV-GENOME-0219962 21779:31544(+)
uvig_336017 21476:31469(+)
CDS_0035
CDS_0033
CDS_0039
CDS_0034
CDS_0036
CDS_0040
MGV-GENOME-0229268 24955:33174(+)
CDS_0039
CDS_0035
CDS_0036
CDS_0040
MGV-GENOME-0224750 24955:32356(+)
pyrophosphatase (CDS_0043)
amidase (CDS_0046)
uvig_226773 21519:44519(+)
pyrophosphatase (CDS_0041)
amidase (CDS_0044)
MGV-GENOME-0291051 21519:40788(+)
Paris_II__AriA_AAA21__All_AAA21_ali (CDS_0035)
pyrophosphatase (CDS_0038)
amidase (CDS_0041)
MGV-GENOME-0285609 21813:40755(+)
pyrophosphatase (CDS_0040)
amidase (CDS_0043)
MGV-GENOME-0269338 21922:40143(+)
pyrophosphatase (CDS_0039)
amidase (CDS_0042)
MGV-GENOME-0318963 21922:37925(+)
AAA domain, putative AbiEii toxin, Type IV TA system (CDS_0043)
pyrophosphatase (CDS_0039)
MGV-GENOME-0266232 21585:39551(+)
AAA domain, putative AbiEii toxin, Type IV TA system (CDS_0042)
pyrophosphatase (CDS_0038)
MGV-GENOME-0270907 21211:40457(+)
AAA domain, putative AbiEii toxin, Type IV TA system (CDS_0033)
MGV-GENOME-0228823 21213:33079(+)
MGV-GENOME-0222046 21211:31905(+)
AAA domain, putative AbiEii toxin, Type IV TA system (CDS_0041)
pyrophosphatase (CDS_0037)
MGV-GENOME-0260383 21813:38513(+)
cas2_I_II_III_IV_V_VI_6 (CDS_0039)
cas1_I_II_III_IV_V_VI_5 (CDS_0038)
AAA domain, putative AbiEii toxin, Type IV TA system (CDS_0047)
uvig_572100 21813:60241(+)
AAA domain, putative AbiEii toxin, Type IV TA system (CDS_0038)
MGV-GENOME-0315806 21834:38813(+)
uvig_334207 22712:33541(+)
CDS_0035
CDS_0034
CDS_0037
MGV-GENOME-0298243 23210:30774(+)
pyrophosphatase (CDS_0038)
amidase (CDS_0041)
MGV-GENOME-0263470 21162:39060(+)
RosmerTA__RmrA_2677172838 (CDS_0041)
MGV-GENOME-0320545 22482:39727(+)
RosmerTA__RmrA_2677172838 (CDS_0039)
DNA methyltransferase (CDS_0031)
uvig_135752 21414:38174(+)
10000 nt
lysis
other
unknown function
Defence