NinB/ Orf homologous recombination mediator (CDS_0026)
AAA domain, putative AbiEii toxin, Type IV TA system (CDS_0025)
DNA polymerase III theta subunit (CDS_0021)
transcriptional repressor (CDS_0029)
replication initiation protein (CDS_0018)
DNA methyltransferase (CDS_0023)
CDS_0028
CDS_0019
CDS_0020
CDS_0022
CDS_0024
CDS_0027
CDS_0030
SAMN11031284_a1_ct4160 12778:20922(+)
CDS_0021
CDS_0023
AAA domain, putative AbiEii toxin, Type IV TA system (CDS_0024)
uvig_394461 13547:21694(+)
AAA domain, putative AbiEii toxin, Type IV TA system (CDS_0025)
CDS_0022
CDS_0024
CDS_0026
TemPhD_cluster_51282 12788:20961(+)
AAA domain, putative AbiEii toxin, Type IV TA system (CDS_0024)
CDS_0016
CDS_0021
CDS_0023
CDS_0025
uvig_395143 11624:19990(+)
CDS_0025
CDS_0026
TemPhD_cluster_62130 13575:21732(+)
CDS_0022
CDS_0024
CDS_0025
CDS_0026
uvig_369681 12805:21213(+)
replication initiation O-like (CDS_0017)
CDS_0022
CDS_0023
CDS_0028
TemPhD_cluster_24912 12042:20680(+)
transcriptional regulator (CDS_0024)
HNH endonuclease (CDS_0022)
RNA polymerase beta subunit (CDS_0021)
23
25
uvig_316948 11607:19411(+)
transcriptional regulator (CDS_0021)
HNH endonuclease (CDS_0019)
RNA polymerase beta subunit (CDS_0018)
20
22
MGV-GENOME-0220470 9448:17317(+)
transcriptional regulator (CDS_0028)
HNH endonuclease (CDS_0026)
RNA polymerase beta subunit (CDS_0025)
27
29
TemPhD_cluster_62158 13391:21288(+)
transcriptional regulator (CDS_0023)
DNA methyltransferase (CDS_0022)
CDS_0024
TemPhD_cluster_62069 14306:21911(+)
RNA polymerase beta subunit (CDS_0025)
transcriptional regulator (CDS_0027)
CDS_0023
HNH endonuclease (CDS_0026)
CDS_0028
TemPhD_cluster_62006 14388:21944(+)
RNA polymerase beta subunit (CDS_0024)
transcriptional regulator (CDS_0026)
HNH endonuclease (CDS_0025)
CDS_0027
MGV-GENOME-0244572 13579:21357(+)
RNA polymerase beta subunit (CDS_0020)
transcriptional regulator (CDS_0022)
HNH endonuclease (CDS_0021)
CDS_0023
uvig_317110 10981:18470(+)
RNA polymerase beta subunit (CDS_0025)
transcriptional regulator (CDS_0027)
HNH endonuclease (CDS_0026)
CDS_0028
TemPhD_cluster_37291_ 15241:22779(+)
RNA polymerase beta subunit (CDS_0026)
transcriptional regulator (CDS_0028)
HNH endonuclease (CDS_0027)
CDS_0029
TemPhD_cluster_33482 13078:20594(+)
transcriptional regulator (CDS_0027)
PD-Lambda-5__PD-Lambda-5_B (CDS_0026)
CDS_0025
CDS_0028
TemPhD_cluster_28125 13883:21616(+)
transcriptional regulator (CDS_0019)
PD-Lambda-5__PD-Lambda-5_B (CDS_0018)
CDS_0017
CDS_0020
HNH endonuclease (CDS_0022)
uvig_370140 6997:15169(+)
transcriptional regulator (CDS_0025)
PD-Lambda-5__PD-Lambda-5_B (CDS_0023)
CDS_0022
CDS_0024
MGV-GENOME-0239617 12843:20925(+)
RNA polymerase beta subunit (CDS_0025)
transcriptional regulator (CDS_0027)
CDS_0022
CDS_0023
HNH endonuclease (CDS_0026)
uvig_394936 12170:20024(+)
HNH endonuclease (CDS_0030)
TemPhD_cluster_37458_ 16111:21252(+)
transcriptional regulator (CDS_0028)
HNH endonuclease (CDS_0026)
RNA polymerase beta subunit (CDS_0025)
CDS_0027
TemPhD_cluster_43490 13880:21820(+)
transcriptional regulator (CDS_0027)
HNH endonuclease (CDS_0025)
RNA polymerase beta subunit (CDS_0024)
CDS_0026
uvig_215665 13432:21533(+)
CII-like transcriptional activator (CDS_0016)
CDS_0022
TemPhD_cluster_37593_ 12298:20050(+)
21
22
23
24
transcriptional regulator (CDS_0025)
TemPhD_cluster_51254 13028:21238(+)
DNA methyltransferase (CDS_0024)
DNA methyltransferase (CDS_0025)
transcriptional regulator (CDS_0026)
TemPhD_cluster_43517 12543:21671(+)
3000 nt
DNA, RNA and nucleotide metabolism
other
transcription regulation
unknown function
Defence