Mu Gam-like end protection (CDS_0011)
excisionase (CDS_0008)
CDS_0010
CDS_0009
exonuclease (CDS_0012)
CDS_0013
CDS_0014
CII-like transcriptional activator (CDS_0017)
transcriptional repressor (CDS_0016)
CDS_0015
replication initiation protein (CDS_0019)
CDS_0018
CDS_0020
CDS_0021
TemPhD_cluster_37400_ 6579:15975(+)
CDS_0030
CDS_0031
IMGVR_UViG_2609460296_000006|2609460296|2609560185|46103-85819 23220:32275(+)
CDS_0024
CDS_0029
uvig_579122 21112:30975(+)
CDS_0015
CDS_0016
transcriptional repressor (CDS_0017)
DNA polymerase III theta subunit (CDS_0023)
TemPhD_cluster_37565_ 8914:18159(+)
transcriptional regulator (CDS_0016)
CDS_0013
CDS_0015
replication initiation O-like (CDS_0019)
TemPhD_cluster_43236 6579:16494(+)
CII-like regulator (CDS_0038)
CDS_0035
CDS_0036
transcriptional regulator (CDS_0039)
IMGVR_UViG_2695420762_000002|2695420762|2695478602|2-41571 25944:34842(+)
CDS_0032
IMGVR_UViG_2681813156_000004|2681813156|2681859301|59-39593 23752:32444(+)
CII-like regulator (CDS_0034)
CDS_0031
CDS_0033
transcriptional regulator (CDS_0035)
IMGVR_UViG_2568526506_000006|2568526506|2568575144|93982-133452 23665:32719(+)
transcriptional regulator (CDS_0035)
CDS_0033
transcriptional regulator (CDS_0036)
uvig_199579 25284:34453(+)
transcriptional regulator (CDS_0036)
CDS_0035
IMGVR_UViG_2791355392_000003|2791355392|2791395105|193189-232994 25682:34516(+)
transcriptional regulator (CDS_0040)
transcriptional regulator (CDS_0039)
TemPhD_cluster_41098 26932:36364(+)
transcriptional regulator (CDS_0033)
transcriptional regulator (CDS_0032)
IMGVR_UViG_2571042453_000004|2571042453|2571114383|3-38982 23600:32666(+)
transcriptional regulator (CDS_0038)
transcriptional regulator (CDS_0037)
IMGVR_UViG_3300013917_000002|3300013917|Ga0120820_100006|118910-164303 28972:38000(+)
transcriptional regulator (CDS_0015)
transcriptional regulator (CDS_0014)
SAMN11031284_a1_ct4160 6815:15853(+)
transcriptional regulator (CDS_0014)
CDS_0013
MGV-GENOME-0241100 7105:15849(+)
transcriptional regulator (CDS_0034)
transcriptional repressor (CDS_0033)
uvig_316213 23181:32367(+)
transcriptional regulator (CDS_0015)
transcriptional repressor (CDS_0014)
CDS_0013
HNH endonuclease (CDS_0018)
TemPhD_cluster_43490 7235:17033(+)
transcriptional regulator (CDS_0035)
CDS_0034
IMGVR_UViG_2609460294_000006|2609460294|2609559921|185-41663 26520:35077(+)
transcriptional regulator (CDS_0012)
CII-like regulator (CDS_0011)
uvig_316948 5612:14700(+)
transcriptional regulator (CDS_0008)
CII-like regulator (CDS_0007)
uvig_370140 1253:10090(+)
CII-like regulator (CDS_0014)
CDS_0015
TemPhD_cluster_51282 6805:15890(+)
transcriptional regulator (CDS_0017)
Mu Gam-like end protection (CDS_0013)
CII-like regulator (CDS_0016)
TemPhD_cluster_43306 11061:19886(+)
transcriptional regulator (CDS_0009)
CDS_0001
CDS_0002
transcriptional regulator (CDS_0008)
uvig_314910 1:11273(+)
transcriptional regulator (CDS_0009)
CDS_0002
transcriptional regulator (CDS_0008)
MGV-GENOME-0220470 1:12523(+)
CDS_0036
IMGVR_UViG_2896813311_000004|2896813311|2896813311|1853217-1898311 26779:36489(+)
transcriptional regulator (CDS_0015)
CDS_0014
CDS_0016
TemPhD_cluster_62130 6997:16668(+)
transcriptional regulator (CDS_0010)
CDS_0009
CDS_0011
uvig_317110 4562:14047(+)
transcriptional regulator (CDS_0015)
CDS_0013
CDS_0014
transcriptional repressor (CDS_0016)
TemPhD_cluster_37291_ 8101:18334(+)
transcriptional regulator (CDS_0015)
transcriptional regulator (CDS_0014)
replication initiation O-like (CDS_0018)
TemPhD_cluster_43517 6803:16088(+)
transcriptional regulator (CDS_0014)
transcriptional regulator (CDS_0013)
TemPhD_cluster_24912 6618:15587(+)
transcriptional regulator (CDS_0015)
CII-like regulator (CDS_0014)
uvig_215665 7725:16977(+)
CII-like regulator (CDS_0014)
CDS_0015
TemPhD_cluster_62208 6615:16404(+)
transcriptional regulator (CDS_0014)
CDS_0013
TemPhD_cluster_51364 6646:16233(+)
transcriptional regulator (CDS_0015)
transcriptional repressor (CDS_0014)
TemPhD_cluster_43553 7701:17169(+)
transcriptional regulator (CDS_0015)
transcriptional regulator (CDS_0014)
TemPhD_cluster_62492 8177:17316(+)
transcriptional regulator (CDS_0015)
transcriptional regulator (CDS_0014)
TemPhD_cluster_62044 6579:16084(+)
transcriptional regulator (CDS_0016)
HNH endonuclease (CDS_0019)
CDS_0024 DNA methyltransferase (CDS_0025)
uvig_427543 7163:17124(+)
HNH endonuclease (CDS_0018)
CDS_0023
TemPhD_cluster_62158 7504:17517(+)
replication initiation O-like (CDS_0018)
CDS_0022
TemPhD_cluster_37593_ 7500:17043(+)
transcriptional regulator (CDS_0018)
transcriptional regulator (CDS_0017)
CDS_0016
CII-like transcriptional activator (CDS_0019)
TemPhD_cluster_24858 9060:18838(+)
Rz-like spanin (CDS_0005)
Mu Gam-like end protection (CDS_0010)
TemPhD_cluster_62012 4796:15326(+)
5000 nt
DNA, RNA and nucleotide metabolism
integration and excision
lysis
other
transcription regulation
unknown function