IMGVR_UVIG_290930379 4_000010|290930379 4|290930379 4|392331 2-396737 8:28,464-35,035 (+)
Tem PhD_cluster_32267:26,388-34,086 (+)
uvig_530249:20,930-27,014 (+)
IMGVR_UVIG_256151124 1_000001|256151124 1|256154223 1|293501-334144:27,218-33,281 (+)
IMGVR_UVIG_272175549 6_000006|272175549 6|272180092 2|367946 8-372033 5:26,153-34,164 (+)
uvig_2626:29,262-36,890 (+)
TemPhD_cluster_17569:26,918-32,622 (+)
IMGVR_UVIG_254104782 8_000002|254104782 8|254117428 1|20564-59701:24,940-31,684 (+)
MGV-GENOME-022858 5:14,312-26,638 (+)
TemPhD_cluster_42250:24,959-35,007 (+)
uvig_135 8:18,655-28,113 (+)
uvig_586835:27,767-37,310 (+)
TemPhD_cluster_27758:26,360-34,254 (+)
uvig_187611:25,377-34,264 (+)
IMGVR_UVIG_273448277 9_000005|273448277 9|273468532 3|49721-91746:25,150-35,740 (+)
5000 nt
DNA, RNA and nucleotide metabolism
unknown function
defence (DefenseFinder)
defence (PADLOC)
virulence (VFDB)
integration and excision
head and packaging
moron, auxiliary metabolic gene and host takeover
other
tail
transcription regulation
tRNA
deoxyribonucleoside 5' monophosphate phosphatase (CDS_0037)
transcriptional repressor (CDS_0046)
transcriptional regulator (CDS_0048)
primosomal protein (CDS_0044)
HNH endonuclease (CDS_0049)
IMGVR_UVIG_26903162 96_000007|26903162 96|26903712 63|85844-127189:28,090-38,204 (+)
DEDDh_I_II_III_IV_V_VI_1 (CDS_0034)
excisionase (CDS_0033)
replication initiation O-like (CDS_0038)
primosomal protein (CDS_0037)
MGV-GENOME-025992:24,413-29,273 (+)
baseplate wedge subunit (CDS_0021)
short tail fiber protein (CDS_0022)
transcriptional regulator (CDS_0034)
MTase_II_00118 Type (CDS_0037)
primosomal protein (CDS_0037)
integrase (CDS_0027)
uvig_354635:15,773-29,281 (+)
tail fiber assembly (CDS_0023)
tail fiber protein (CDS_0022)
Menshen_Nsn A (CDS_0025)
transcriptional regulator (CDS_0030)
baseplate protein (CDS_0020)
tail fiber protein (CDS_0022)
transcriptional regulator (CDS_0030)
MGV-GENOME-021990:14,724-24,844 (+)
transcriptional regulator (CDS_0040)
Tem PhD_cluster_41899:25,857-33,496 (+)
transcriptional regulator (CDS_0037)
primosomal protein (CDS_0040)
Tem PhD_cluster_49865:24,441-32,288 (+)
Tem PhD_cluster_36715_:25,337-32,988 (+)
transcriptional regulator (CDS_0041)
uvig_116805:25,619-32,425 (+)
transcriptional regulator (CDS_0039)
Tem PhD_cluster_36704_:24,725-31,726 (+)
transcriptional regulator (CDS_0037)
IMGVR_UVIG_277826108 9_000001|277826108 9|277827796 6:27,263-33,610 (+)
transcriptional regulator (CDS_0033)
MGV-GENOME-024771 4:19,438-25,785 (+)
primosomal protein (CDS_0045)
IMGVR_UVIG_277826164 1_000011|277826164 1|277833216 4|80811-121448:27,381-34,766 (+)
Tem PhD_cluster_61378:25,408-31,755 (+)
tail fiber assembly (CDS_0024)
LPS (VF0124) (CDS_0026)
transcriptional regulator (CDS_0031)
short tail fiber protein (CDS_0023)
uvig_110542:15,482-29,803 (+)
transcriptional regulator (CDS_0030)
IMGVR_UVIG_272175570 9_000003|272175570 9|272180130 8|327509 1-331850:26,994-35,163 (+)