transcriptional repressor (CDS_0045)
CDS_0046
DNA polymerase exonuclease subunit (CDS_0044)
CDS_0047 CDS_0049
CDS_0041
CDS_0042
CDS_0043
CDS_0048
CDS_0050
CDS_0051
uvig_456003 31666:38850(+)
CDS_0030
CDS_0036
IMGVR_UViG_3300029373_000337|3300029373|Ga0243914_100688|8416-44357 22867:30708(+)
Retron_VI__HTH (CDS_0035)
CDS_0028
CDS_0034
uvig_489054 20772:30633(+)
CDS_0036
IMGVR_UViG_3300007105_000269|3300007105|Ga0102630_100234|664-36302 22843:30733(+)
CDS_0028
CDS_0032 CDS_0033
CDS_0035
IMGVR_UViG_3300023489_000055|3300023489|Ga0257030_100587 22081:29478(+)
CDS_0035
IMGVR_UViG_3300029419_000355|3300029419|Ga0243913_100781|2623-37454 22079:29598(+)
DNA polymerase exonuclease subunit (CDS_0048)
Retron_VI__HTH (CDS_0050)
transcriptional repressor (CDS_0049)
IMGVR_UViG_3300029794_000469|3300029794|Ga0243854_1000639|6236-48981 30339:37536(+)
DNA polymerase exonuclease subunit (CDS_0034)
Retron_VI__HTH (CDS_0036)
transcriptional repressor (CDS_0035)
Kang_0013_ST001_A_NODE_47_length_36276_cov_40.723613 22048:31044(+)
CDS_0050
uvig_486634 35093:41762(+)
CDS_0002
uvig_495425 50:3191(+)
CDS_0030
CDS_0035 CDS_0036
CDS_0038
IMGVR_UViG_3300029438_000326|3300029438|Ga0243733_1000281|20571-56783 21757:30774(+)
CDS_0035 CDS_0036
uvig_378864 21897:31003(+)
CDS_0034 CDS_0035
IMGVR_UViG_3300014553_000179|3300014553|Ga0134451_100012|57065-92268 21897:29432(+)
transcriptional repressor (CDS_0036)
CDS_0034
CDS_0027
CDS_0035
CDS_0037
IMGVR_UViG_3300029789_000298|3300029789|Ga0243734_1000383|2589-37690 20881:30201(+)
toxin-antitoxin system HicB-like (CDS_0034)
toxin (CDS_0033)
DNA polymerase (CDS_0035)
CDS_0028
OTU_3006 20953:30856(+)
3000 nt
DNA, RNA and nucleotide metabolism
integration and excision
lysis
moron, auxiliary metabolic gene and host takeover
transcription regulation
unknown function
Defence