CDS_0011
CDS_0010
tail length tape measure protein (CDS_0012)
CDS_0013
CDS_0014
CDS_0015
CDS_0016
CDS_0017
CDS_0018
CDS_0019
CDS_0020
CDS_0021
CDS_0022
amidase (CDS_0023)
MGV-GENOME-4429729 7245:23207(+)
MGV-GENOME-0269219 7244:23294(+)
IMGVR_UViG_3300045988_062540|3300045988|Ga0495776_070587 7262:23475(+)
IMGVR_UViG_3300026522_000059|3300026522|Ga0256406_1001298 7241:23288(+)
MGV-GENOME-0198754 7809:23373(+)
tail length tape measure protein (CDS_0021)
CDS_0027
IMGVR_UViG_3300045988_103340|3300045988|Ga0495776_098603 13269:28532(+)
CDS_0028
MGV-GENOME-0298383 13700:28687(+)
IMGVR_UViG_3300034415_000142|3300034415|Ga0168781_00065|138197-178949 7325:23382(+)
tail fiber protein (CDS_0018)
Han_2018_ERR1398246_NODE_252_length_39604_cov_100.587499 7314:23485(+)
CDS_0030
IMGVR_UViG_3300045988_075032|3300045988|Ga0495776_157310 17803:29962(+)
IMGVR_UViG_3300045988_075696|3300045988|Ga0495776_109812 11875:23623(+)
uvig_133027 6162:17879(-)
MGV-GENOME-0298295 18397:30752(+)
CDS_0023
uvig_171713 17095:29246(+)
MGV-GENOME-0188944 10311:21780(+)
MGV-GENOME-0207343 11872:23963(+)
CDS_0020
MGV-GENOME-0270755 11884:23971(+)
CDS_0029
CDS_0034
IMGVR_UViG_3300045988_092496|3300045988|Ga0495776_161572 17324:29607(+)
MGV-GENOME-0231315 13499:25172(+)
CDS_0019
uvig_574688 14687:23664(+)
CDS_0032
IMGVR_UViG_3300045988_089128|3300045988|Ga0495776_097059 18888:30619(+)
CDS_0026
uvig_484677 17219:28677(+)
CDS_0008
MGV-GENOME-0200891 1685:13416(-)
IMGVR_UViG_3300045988_063603|3300045988|Ga0495776_025023 11865:23605(+)
uvig_284122 17881:29612(+)
uvig_84860 11881:23199(+)
CDS_0014
SAMN06164579_a1_ct65036_vs1 5458:16759(-)
MGV-GENOME-0159160 10104:21432(+)
CDS_0007
MGV-GENOME-0192834 2541:14505(-)
MGV-GENOME-0177926 10112:22184(+)
uvig_410035 17175:29680(+)
CDS_0025
IMGVR_UViG_3300045988_067137|3300045988|Ga0495776_186778 16649:28611(+)
MGV-GENOME-0295783 11880:23540(+)
IMGVR_UViG_3300045988_083850|3300045988|Ga0495776_001095 11884:24295(+)
uvig_559992 11949:24575(+)
OTU_1951 11869:24154(+)
MGV-GENOME-0172874 12854:21377(+)
CDS_0036
MGV-GENOME-0300843 20205:31689(+)
uvig_302989 11878:23309(+)
MGV-GENOME-0353501 21871:31744(+)
MGV-GENOME-0280287 16178:24817(+)
MGV-GENOME-0197678 19294:27342(+)
CDS_0005
uvig_347612 27:8080(-)
IMGVR_UViG_3300045988_084050|3300045988|Ga0495776_163404 22281:31333(+)
RM_Type_II__Type_II_MTases___Type_II_MTases_FAM_0 (CDS_0038)
DNA helicase (CDS_0040)
MGV-GENOME-0264365 22704:35256(+)
IMGVR_UViG_3300045988_119649|3300045988|Ga0495776_098559 22704:31985(+)
6000 nt
DNA, RNA and nucleotide metabolism
lysis
tail
unknown function
Defence