RM_Type_II__Type_II_MTases__Type_II_MTases_FAM_0 (CDS_0020)
DNA methyltransferase (CDS_0021)
PD-T4-9__PD-T4-9_C (CDS_0028)
MGV-GENOME-0338208 14456:39676(+)
SAMN05827106_a1_ct3868 14457:39476(+)
methyltransferase (CDS_0021)
IMGVR_UViG_3300030447_000343|3300030447|Ga0168849_100111|23225-87778 14457:39712(+)
MGV-GENOME-0341942 14457:40423(+)
uvig_229981 14457:42931(+)
MGV-GENOME-0339949 14457:38698(+)
MGV-GENOME-0338986 14457:39671(+)
IMGVR_UViG_3300045988_172345|3300045988|Ga0495776_163398 14457:39408(+)
IMGVR_UViG_3300045988_169465|3300045988|Ga0495776_090468 14457:37153(+)
MGV-GENOME-0351273 14456:54152(+)
PD-T4-9__PD-T4-9_C (CDS_0021)
DISARM_2__drmMII (CDS_0020)
MGV-GENOME-0329247 14457:34149(+)
CDS_0020
CDS_0021
CDS_0019
CDS_0022
uvig_110469 14214:33092(+)
CDS_0021
CDS_0023
CDS_0022
MGV-GENOME-0343964 15815:34033(+)
PD-T4-9__PD-T4-9_C (CDS_0023)
DISARM_2__drmMII (CDS_0022)
MGV-GENOME-0329711 28495:34513(+)
CDS_0019
CDS_0020
IMGVR_UViG_3300045988_172123|3300045988|Ga0495776_091102 14457:33269(+)
IMGVR_UViG_3300045988_172736|3300045988|Ga0495776_056561 14457:33400(+)
RM_Type_II__Type_II_MTases__Type_II_MTases_FAM_0 (CDS_0021)
DNA methyltransferase (CDS_0022)
CDS_0024
Ma_2019_SRR413634_NODE_297_length_62682_cov_13.622511 14646:37002(+)
IMGVR_UViG_3300045988_174355|3300045988|Ga0495776_090946 13723:34291(+)
CDS_0021
IMGVR_UViG_3300045988_173303|3300045988|Ga0495776_036073 14457:34969(+)
IMGVR_UViG_3300045988_175113|3300045988|Ga0495776_091078 13723:34388(+)
MGV-GENOME-0333878 14214:36049(+)
CDS_0025
MGV-GENOME-0333524 14143:34645(+)
IMGVR_UViG_3300007805_000215|3300007805|Ga0105637_100019|126040-190194 14457:38105(+)
IMGVR_UViG_3300045988_171981|3300045988|Ga0495776_150111 29491:36839(+)
MGV-GENOME-0338966 29249:37326(+)
CDS_0028
MGV-GENOME-0333267 13481:34333(+)
uvig_112427 13723:34874(+)
IMGVR_UViG_3300029052_000148|3300029052|Ga0169758_100053 13723:41838(+)
IMGVR_UViG_3300045988_172665|3300045988|Ga0495776_140196 13481:39439(+)
protease (CDS_0032)
IMGVR_UViG_3300045988_172100|3300045988|Ga0495776_140373 14457:39308(+)
MGV-GENOME-0335182 15271:38625(+)
MGV-GENOME-0333470 15271:35366(+)
IMGVR_UViG_3300007796_000315|3300007796|Ga0105889_100007|136173-200225 14143:38042(+)
OTU_609 14143:38093(+)
IMGVR_UViG_7000000334_000286|7000000334|SRS015217_WUGC_scaffold_7681|3-60787 26931:35034(+)
Heme receptor HutA (VF0629) (CDS_0022)
uvig_506190 13724:48466(+)
OTU_549 14066:37578(+)
nucleoside 2-deoxyribosyltransferase (CDS_0021)
MGV-GENOME-0329588 14658:36158(+)
uvig_423824 14683:36609(+)
IMGVR_UViG_3300045988_171790|3300045988|Ga0495776_111586 14066:36555(+)
IMGVR_UViG_3300045988_171926|3300045988|Ga0495776_090609 14215:35890(+)
MGV-GENOME-0333898 14215:35602(+)
uvig_238167 14215:36226(+)
IMGVR_UViG_3300045988_173504|3300045988|Ga0495776_102295 14698:37781(+)
uvig_380256 14457:51244(+)
10000 nt
DNA, RNA and nucleotide metabolism
moron, auxiliary metabolic gene and host takeover
other
unknown function
Defence
Virulence