uvig_348444:1-51,620 (+)
NC_049838.1:1-54,438 (+)
NC_049837.1:1-51,487 (+)
NC_049836.1:1-51,780 (+)
NC_047825.1:1-50,526 (+)
MGV-GENOME-031320 1:1-51,119 (+)
NC_047784.1:1-51,099 (+)
NC_049833.1:1-50,713 (+)
NC_049834.1:1-52,866 (+)
MGV-GENOME-030989 8:1-49,905 (+)
uvig_234015:1-48,858 (+)
uvig_329390:1-48,979 (+)
MGV-GENOME-031424 0:1-51,565 (+)
uvig_145376:1-49,469 (+)
MGV-GENOME-030890 7:1-49,513 (+)
uvig_255004:1-50,328 (+)
uvig_467799:1-49,631 (+)
NC_043469.1:1-51,543 (+)
MGV-GENOME-031360 5:1-51,290 (+)
MGV-GENOME-031281 9:1-50,972 (+)
uvig_215036:1-50,916 (+)
uvig_349911:1-34,594 (+)
2000 0 nt
defence (PADLOC)
anti-defence (dbAPIS)
uvig_354241:1-51,621 (+)
TemPhD_cluster_39656_:1-51,544 (+)
MGV-GENOME-030159 7:1-47,100 (+)
uvig_328591:1-47,402 (+)
MGV-GENOME-030270 2:1-47,436 (+)
MGV-GENOME-030670 4:1-48,774 (+)
MGV-GENOME-031121 4:1-50,348 (+)
MGV-GENOME-031135 6:1-50,403 (+)
MGV-GENOME-031042 6:1-50,071 (+)
uvig_283917:1-51,650 (+)
MGV-GENOME-031230 0:1-50,762 (+)
uvig_331247:1-51,311 (+)
MGV-GENOME-031364 8:1-51,311 (+)
uvig_284377:1-50,720 (+)
MGV-GENOME-044274 0:1-49,852 (+)
Ma_2019_SRR4137 10_NODE_378_length_5071 5_cov_48.086538:1-50,715 (+)
uvig_243694:1-50,577 (+)
MGV-GENOME-044251 27:1-50,621 (+)
MGV-GENOME-031192 2:1-50,626 (+)
MGV-GENOME-030950 8:1-49,738 (+)
MGV-GENOME-030847 6:1-49,366 (+)
MGV-GENOME-031540 :1-50,481 (+)
SAMEA2737751_a1_c15309:1-50,382 (+)
NC_049846.1:1-49,045 (+)
NC_048126.1:1-49,861 (+)
uvig_219619:1-47,173 (+)
MGV-GENOME-030183 0:1-47,173 (+)
MGV-GENOME-030166 3:1-47,118 (+)
uvig_574399:1-47,603 (+)
MGV-GENOME-030862 2:1-49,421 (+)
MGV-GENOME-030607 3:1-48,570 (+)
MGV-GENOME-030823 9:1-49,295 (+)
NC_031053.1:1-50,934 (+)
NC_028760.1:1-49,037 (+)
MGV-GENOME-025032 1:1-36,787 (+)
SAMN10080877_a1_c119236_v s1:1-30,632 (+)
NC_049844.1:1-43,094 (+)
uvig_394929:1-51,153 (+)
uvig_376089:1-49,147 (+)
uvig_132550:1-49,636 (+)
TemPhD_cluster_39856_:1-49,839 (+)
NC_049845.1:1-49,477 (+)
uvig_224277:1-49,371 (+)
MGV-GENOME-030849 5:1-49,371 (+)
TemPhD_cluster_31468:1-49,118 (+)
uvig_315584:1-49,173 (+)
TemPhD_cluster_31466:1-49,118 (+)
MGV-GENOME-030806 8:1-49,237 (+)
TemPhD_cluster_31467:1-49,118 (+)
MGV-GENOME-030739 2:1-48,992 (+)
NC_028786.1:1-49,462 (+)
NC_048162.1:1-49,427 (+)
NC_047907.1:1-50,249 (+)
NC_047850.1:1-49,807 (+)
NC_049841.1:1-49,935 (+)
NC_049847.1:1-49,916 (+)
MGV-GENOME-031104 9:1-50,279 (+)
MGV-GENOME-030899 8:1-49,548 (+)
NC_029099.1:1-49,797 (+)
MGV-GENOME-030873 5:1-49,458 (+)
MGV-GENOME-030849 5:1-49,373 (+)
uvig_279208:1-49,250 (+)
uvig_536741:1-49,871 (+)
uvig_535962:1-49,873 (+)
NC_049842.1:1-49,316 (+)
NC_028774.1:1-48,754 (+)
NC_048143.1:1-49,106 (+)
uvig_574762:1-50,405 (+)
uvig_278768:1-48,961 (+)
MGV-GENOME-030265 5:1-47,422 (+)
NC_048043.1:1-50,132 (+)
NC_049840.1:1-50,381 (+)
NC_048044.1:1-49,468 (+)
NC_048042.1:1-49,344 (+)
uvig_285149:1-48,646 (+)
MGV-GENOME-030586 9:1-48,507 (+)
NC_055956.1:1-49,276 (+)
MGV-GENOME-030544 9:1-48,360 (+)
uvig_130754:1-48,360 (+)
MGV-GENOME-030582 4:1-48,491 (+)
uvig_369684:1-49,299 (+)
NC_049848.1:1-49,835 (+)
NC_048024.1:1-49,292 (+)
NC_049839.1:1-50,241 (+)