replisome organizer (CDS_0043)
Druantia_II__DruM (CDS_0046)
RusA-like Holliday junction resolvase (CDS_0058)
uvig_147530 35896:44871(+)
Druantia_II__DruM (CDS_0053)
IMGVR_UViG_3300045988_091814|3300045988|Ga0495776_078082 38526:46310(+)
Druantia_II__DruM (CDS_0041)
uvig_479879 32176:39819(+)
Druantia_II__DruM (CDS_0043)
MGV-GENOME-0301188 33562:41538(+)
Druantia_II__DruM (CDS_0043)
uvig_31530 33096:44457(+)
RM_Type_II__Type_II_MTases__Type_II_MTases_FAM_0 (CDS_0039)
MGV-GENOME-0290101 30370:39280(+)
Druantia_II__DruM (CDS_0046)
CDS_0048
CDS_0050
CDS_0052
CDS_0054
CDS_0047
CDS_0049
CDS_0051
CDS_0053
SAMN09781697_a1_ct4791 33689:42066(+)
RM_Type_II__Type_II_MTases__Type_II_MTases_FAM_0 (CDS_0045)
CDS_0050
CDS_0047
CDS_0051
CDS_0046
CDS_0049
CDS_0052
MGV-GENOME-0299884 33660:41327(+)
RM_Type_II__Type_II_MTases__Type_II_MTases_FAM_0 (CDS_0044)
MGV-GENOME-0301150 33834:45136(+)
RM_Type_II__Type_II_MTases__Type_II_MTases_FAM_0 (CDS_0043)
uvig_171125 33739:45096(+)
MGV-GENOME-0295633 33970:41499(+)
IMGVR_UViG_3300045988_040465|3300045988|Ga0495776_097928 33061:41107(+)
Druantia_II__DruM (CDS_0044)
MGV-GENOME-0300084 34078:43100(+)
Druantia_II__DruM (CDS_0044)
uvig_259339 35257:46261(+)
Druantia_II__DruM (CDS_0042)
uvig_548315 31341:42332(+)
Druantia_II__DruM (CDS_0043)
uvig_92675 33508:44564(+)
IMGVR_UViG_3300045988_041342|3300045988|Ga0495776_132551 33685:40757(+)
IMGVR_UViG_3300045988_068457|3300045988|Ga0495776_016020 34093:40712(+)
SAMN09781789_a1_ct7961 35530:45303(+)
RM_Type_II__Type_II_MTases__Type_II_MTases_FAM_0 (CDS_0045)
DNA methyltransferase (CDS_0044)
IMGVR_UViG_3300029427_000265|3300029427|Ga0243889_1000030|131412-178253 34033:45009(+)
CDS_0047
CDS_0048
CDS_0049
IMGVR_UViG_3300045988_077170|3300045988|Ga0495776_053895 34615:42969(+)
CDS_0049
CDS_0050
CDS_0055
MGV-GENOME-0297897 35269:44165(+)
uvig_592138 34747:44657(+)
SspBCDE__SspD (CDS_0043)
MGV-GENOME-0302053 34372:45752(+)
MGV-GENOME-0290566 32850:42566(+)
uvig_575746 34100:41584(+)
uvig_580743 34776:41081(+)
MGV-GENOME-0301938 35753:42135(+)
IMGVR_UViG_3300030319_000335|3300030319|Ga0244979_100190|26001-70400 34034:42428(+)
uvig_589200 35129:45091(+)
CDS_0036
CDS_0035
CDS_0038
Druantia_II__DruM (CDS_0033)
CDS_0037
MGV-GENOME-0267878 27682:34417(+)
Druantia_II__DruM (CDS_0046)
MGV-GENOME-0300912 34678:41919(+)
CDS_0028
MGV-GENOME-0238055 24372:28964(+)
Druantia_II__DruM (CDS_0044)
IMGVR_UViG_3300045988_057111|3300045988|Ga0495776_097832 32567:40693(+)
IMGVR_UViG_3300045988_047432|3300045988|Ga0495776_098546 34220:44412(+)
uvig_3616 34181:45069(+)
IMGVR_UViG_3300045988_044387|3300045988|Ga0495776_185777 34181:44152(+)
MGV-GENOME-0250603 24541:30887(+)
MGV-GENOME-0301741 34842:41685(+)
MGV-GENOME-0246190 22932:31922(+)
SspBCDE__SspD (CDS_0037)
CDS_0039
MGV-GENOME-0276025 28500:35851(+)
RecE-like recombination exonuclease (CDS_0038)
uvig_48569 32718:39323(+)
SspBCDE__SspD (CDS_0043)
uvig_282935 32722:40450(+)
Druantia_II__DruM (CDS_0046)
uvig_326083 34095:42224(+)
Druantia_II__DruM (CDS_0041)
uvig_277695 31205:39333(+)
MGV-GENOME-0295461 34159:43975(+)
SspBCDE__SspD (CDS_0047)
Rampelli_SM_29_NODE_135_length_48302_cov_11.561403 35548:43015(+)
4000 nt
DNA, RNA and nucleotide metabolism
other
unknown function
Defence