RelE-like toxin (CDS_0022)
CDS_0016
CDS_0017
CDS_0018
CDS_0019
CDS_0020
CDS_0021
CDS_0023
CDS_0024
CDS_0025
CDS_0026
CDS_0027
CDS_0028
CDS_0029
CDS_0030
CDS_0031
CDS_0032
CDS_0033
IMGVR_UViG_3300029847_000261|3300029847|Ga0245281_100185|31331-73826 25005:34543(+)
IMGVR_UViG_3300029045_000208|3300029045|Ga0169661_100107|22087-64741 25018:34391(+)
IMGVR_UViG_3300029576_000203|3300029576|Ga0245100_100029|234266-276430 25013:35364(+)
transcriptional repressor (CDS_0022)
uvig_200269 25013:33390(+)
IMGVR_UViG_3300039323_000284|3300039323|Ga0169852_00071|16190-58271 24780:35463(+)
uvig_108717 24780:33687(+)
IMGVR_UViG_3300045988_153247|3300045988|Ga0495776_111045 26992:34012(+)
uvig_164029 25271:31883(+)
uvig_485529 24731:33384(+)
uvig_57354 32815:38825(+)
uvig_109315 29559:36145(+)
uvig_8620 17554:24886(+)
IMGVR_UViG_3300006747_000210|3300006747|Ga0101801_100067|57162-100797 30353:36046(+)
IMGVR_UViG_3300029671_000211|3300029671|Ga0245181_100001|143102-185596 28069:34453(+)
MGV-GENOME-0261228 25860:32427(+)
MGV-GENOME-0268855 25007:32467(+)
IMGVR_UViG_3300039356_000253|3300039356|Ga0169776_00069|31966-76397 32707:38593(+)
IMGVR_UViG_3300007975_000049|3300007975|Ga0114169_100162 25005:35391(+)
MGV-GENOME-0268828 25005:33168(+)
IMGVR_UViG_3300029610_000016|3300029610|Ga0245132_100566 29310:35707(+)
Kang_0009_ST003_A_NODE_18_length_41590_cov_138.296304 25007:34000(+)
IMGVR_UViG_3300045988_122925|3300045988|Ga0495776_179303 25031:31920(+)
IMGVR_UViG_3300029503_000204|3300029503|Ga0244781_100152|13400-53728 25042:32430(+)
IMGVR_UViG_2899900031_000001|2899900031|2899900042|89041-128315 25002:31376(+)
IMGVR_UViG_3300029614_000296|3300029614|Ga0245155_100039|76019-118175 25035:34563(+)
uvig_101249 25024:32239(+)
TemPhD_cluster_61717 24163:31403(+)
IMGVR_UViG_3300029580_000243|3300029580|Ga0245096_100012|194697-236227 25023:33633(+)
IMGVR_UViG_3300029661_000175|3300029661|Ga0245258_100023|273474-315375 25008:33816(+)
MGV-GENOME-0270443 24727:32143(+)
IMGVR_UViG_3300029715_000462|3300029715|Ga0245215_1000003|538033-580276 25005:34220(+)
MGV-GENOME-0252514 24971:31627(+)
IMGVR_UViG_3300029208_000195|3300029208|Ga0168729_100011|82527-124951 25042:34526(+)
IMGVR_UViG_3300029121_000170|3300029121|Ga0168811_100066|107026-146358 25041:31438(+)
uvig_284006 28797:34463(+)
MGV-GENOME-0293675 31668:36803(+)
CDS_0011
CDS_0012
MGV-GENOME-0211388 10010:21701(+)
uvig_56056 10031:21354(+)
holin (CDS_0014)
CDS_0015
CDS_0020
CDS_0021
uvig_194247 24794:31161(+)
CDS_0019
CDS_0022
IMGVR_UViG_3300029729_000226|3300029729|Ga0245191_100500|134-40277 24790:32735(+)
ivig_820 24787:31842(+)
transcriptional repressor (CDS_0021)
uvig_490745_ 23833:30016(+)
uvig_183442 35060:39807(+)
4000 nt
integration and excision
lysis
moron, auxiliary metabolic gene and host takeover
transcription regulation
unknown function