endolysin (CDS_0023)
CDS_0027
CDS_0026
CDS_0020
CDS_0022
CDS_0025
CDS_0029
CDS_0031
CDS_0033
CDS_0035
CDS_0021
CDS_0024
CDS_0028
CDS_0030
CDS_0032
CDS_0034
IMGVR_UViG_3300045988_163384|3300045988|Ga0495776_144050 16361:24158(+)
CDS_0027
CDS_0026
CDS_0028
MGV-GENOME-0236215 16361:23413(+)
FS_Sma__Sma (CDS_0028)
26
27
29
30
IMGVR_UViG_3300045988_187515|3300045988|Ga0495776_112404|221-37956 16361:26585(+)
CDS_0027
CDS_0026
CDS_0028
IMGVR_UViG_3300045988_167544|3300045988|Ga0495776_062324 16352:24524(+)
CDS_0027
CDS_0026
CDS_0028
IMGVR_UViG_3300045988_152224|3300045988|Ga0495776_177383 16361:23768(+)
CDS_0027
CDS_0026
CDS_0028
MGV-GENOME-0240204 16361:23497(+)
DNA invertase (CDS_0029)
26
27
28
30
31
IMGVR_UViG_3300045988_161753|3300045988|Ga0495776_085553 16361:27287(+)
26
27
28
29
30
IMGVR_UViG_3300007356_000013|3300007356|Ga0104763_100522 16361:26982(+)
CDS_0026
IMGVR_UViG_3300029438_000364|3300029438|Ga0243733_1000210|10501-46675 16361:24417(+)
26
27
28
29
30
31
uvig_6133 16361:27560(+)
RT_III-AM_III-B_2 (CDS_0029)
26
27
28
30
31
IMGVR_UViG_3300029747_000375|3300029747|Ga0245228_100154|43919-81853 16361:26177(+)
MutT/NUDIX hydrolase (CDS_0031)
FS_Sma__Sma (CDS_0028)
26
27
29
30
32
33
34
35
36
37
38
MGV-GENOME-0280721 16298:31163(+)
CDS_0019
CDS_0025
IMGVR_UViG_3300045988_169679|3300045988|Ga0495776_160692 16037:22621(+)
RT_III-AM_III-B_2 (CDS_0018)
CDS_0017
endolysin (CDS_0023)
CDS_0024
IMGVR_UViG_3300045988_167591|3300045988|Ga0495776_159771 14195:21941(+)
CDS_0027
CDS_0026
CDS_0024
CDS_0028
IMGVR_UViG_3300045988_155996|3300045988|Ga0495776_100961 16037:23413(+)
26
27
28
29GAPS1__GAPS1 (CDS_0030)
31
32
33
34
SRS011084_a1_ct9091_vs1 16361:31701(+)
co-chaperonin GroES (CDS_0033)
GroEL (VF0594) (CDS_0034)
endolysin (CDS_0025)
28
29
30
31
32
35
uvig_205392 16586:28310(+)
CDS_0029
CDS_0037
CDS_0028
CDS_0032
CDS_0034
CDS_0038
IMGVR_UViG_3300045988_168124|3300045988|Ga0495776_100580 16987:23669(+)
CDS_0030
MGV-GENOME-0285244 16978:24083(+)
CDS_0027
CDS_0026
CDS_0028
ABC transporter (VF0818) (CDS_0029)
CDS_0032
uvig_167235 16361:27356(+)
CDS_0025
CDS_0026
CDS_0029
MGV-GENOME-0229451 16484:22066(+)
CDS_0028
MGV-GENOME-4407469 16484:21860(+)
CDS_0025
IMGVR_UViG_3300045988_153961|3300045988|Ga0495776_145150 16361:21907(+)
CDS_0022
CDS_0021
CDS_0026
CDS_0027
IMGVR_UViG_3300014963_000946|3300014963|Ga0134523_1000377|73589-109821 16037:24474(+)
6000 nt
DNA, RNA and nucleotide metabolism
lysis
other
transcription regulation
unknown function
Defence
Virulence