MTase_II_00058 Type (DIFFAGNO_00010)
IMGPR_plasmid_3300008082_000012: 7,719-15,804 (+)
IMGPR_plasmid_3300007648_000029: 355-5,544 (+)
IMGPR_plasmid_3300006302_000006: 38,634-44,871 (-)
MTase_II_00058 Type (GEGKGNPM_00044)
IMGPR_plasmid_3300006743_000010: 38,383-46,030 (-)
MTase_II_00058 Type (ANGCPAMJ_00002)
IMGPR_plasmid_3300008144_000014: 437-8,335 (+)
IMGPR_plasmid_7000000449_000023: 114-4,877 (+)
IMGPR_plasmid_3300008089_000037: 37-6,185 (+)
IMGPR_plasmid_7000000008_000005: 20,473-24,085 (-)
IMGPR_plasmid_3300008611_000025: 37,960-43,083 (-)
IMGPR_plasmid_3300008273_000019: 3,981-7,593 (+)
IMGPR_plasmid_3300008056_000007: 347-7,054 (+)
IMGPR_plasmid_3300007284_000012: 33,943-39,505 (-)
Phosphoenolpyruvate carboxylase (CAJBDEEC_00046)
IMGPR_plasmid_3300008534_000023: 40,704-44,954 (-)
IMGPR_plasmid_3300008518_000026: 38,424-43,450 (-)
IMGPR_plasmid_3300007995_000013: 39,104-44,556 (-)
IMGPR_plasmid_7000000205_000009: 21,209-24,232 (-)
IMGPR_plasmid_2684622627_000003: 269-3,783 (+)
IMGPR_plasmid_7000000623_000011: 216-4,521 (+)
IMGPR_plasmid_3300006259_000024: 3,701-9,482 (+)
IMGPR_plasmid_2826184818_000002: 268-5,804 (+)
IMGPR_plasmid_3300006328_000014: 40,296-45,409 (-)
IMGPR_plasmid_3300008270_000008: 24,850-29,607 (-)
IMGPR_plasmid_3300008700_000019: 323-6,062 (+)
IMGPR_plasmid_3300007663_000008: 31,701-36,998 (-)
IMGPR_plasmid_7000000650_000007: 813-6,250 (+)
IMGPR_plasmid_3300008474_000004: 39,904-47,279 (-)
IMGPR_plasmid_3300007967_000006: 40,672-47,905 (-)
IMGPR_plasmid_3300007810_000006: 366-6,895 (+)
IMGPR_plasmid_3300008626_000031: 324-6,163 (+)
IMGPR_plasmid_7000000005_000039: 1,063-6,760 (+)
IMGPR_plasmid_3300008142_000025: 35,500-45,010 (-)
Lipid II:glycine glycyltransferase (BPCNJLOK_00002)
IMGPR_plasmid_3300007753_000012: 72-6,524 (+)
IMGPR_plasmid_3300007782_000014: 41,016-46,373 (-)
IMGPR_plasmid_3300008483_000020: 29,573-34,181 (-)
IMGPR_plasmid_3300008637_000013: 3,088-7,696 (+)
MTase_II_00058 Type (FCHKAAJJ_00004)
IMGPR_plasmid_3300008682_000015: 1,370-7,662 (+)
MTase_II_00058 Type (BIPPJEJI_00034)
IMGPR_plasmid_3300006295_000017: 28,577-34,869 (-)
MTase_II_00058 Type (MPGCJLKP_00038)
IMGPR_plasmid_3300013749_000009: 32,164-38,380 (-)
IMGPR_plasmid_7000000261_000005: 33,568-39,188 (-)
IMGPR_plasmid_3300006249_000011: 32,407-37,509 (-)
High-affinity branched-chain amino acid transport ATP-binding protein LivF (IOAMGOLC_00001)
IMGPR_plasmid_3300008630_000015: 24-5,181 (+)
IMGPR_plasmid_3300008140_000020: 394-4,695 (+)
IMGPR_plasmid_3300008078_000003: 28,595-34,358 (-)
IMGPR_plasmid_3300014133_000005: 403-6,166 (+)
IMGPR_plasmid_7000000713_000004: 32,765-38,840 (-)
Glycogen phosphorylase (PECLPNFM_00001)
IMGPR_plasmid_3300008454_000026: 36-4,350 (+)
IMGPR_plasmid_2558860933_000001: 40,751-45,393 (-)
IMGPR_plasmid_3300007208_000025: 38,076-43,781 (+); 1-2,600 (+)
IMGPR_plasmid_2660238679_000003: 20,096-31,368 (-)
IMGPR_plasmid_3300008734_000027: 14,809-27,239 (+)
IMGPR_plasmid_3300006254_000024: 11,166-23,231 (+)
IMGPR_plasmid_2826272898_000001: 22,619-35,863 (-)
IMGPR_plasmid_3300008281_000017: 584-6,150 (+)
IMGPR_plasmid_2534682068_000001: 42,132-46,427 (+); 1-5,289 (+)
5000 nt
defence (PADLOC)