VanR-ABDEGLN family response regulator transcription factor (NF033117.2) (NBMDJDKA_00002)
Putative septation protein SpoVG (NBMDJDKA_00004)
Adaptive-response sensory-kinase SasA (NBMDJDKA_00001)
ssDNA thymidine ADP-ribosyltransferase, DarTG__DarT (NBMDJDKA_00011)
DarTG__DarG (NBMDJDKA_00010)
IMGPR_plasmid_3300029505_000119 291:17375(+)
ssDNA thymidine ADP-ribosyltransferase, DarTG__DarT (CDGPEKBO_00029)
Phosphoadenosine phosphosulfate reductase (CDGPEKBO_00032)
CDGPEKBO_00028
CDGPEKBO_00026
CDGPEKBO_00024
DarTG__DarG (CDGPEKBO_00030)
CDGPEKBO_00025
CDGPEKBO_00023
IMGPR_plasmid_3300029096_000171 23315:34967(-)
Phosphoadenosine phosphosulfate reductase (MCOLBOAI_00012)
RnlAB__RnlB (MCOLBOAI_00015)
MCOLBOAI_00013
MCOLBOAI_00016
IMGPR_plasmid_3300029028_000111 11166:22184(+)
NCLLCOGK_00004
IMGPR_plasmid_3300029007_000071 6060:16368(+)
VanR-ABDEGLN family response regulator transcription factor (NF033117.2) (EBMCFGGG_00042)
Putative septation protein SpoVG (EBMCFGGG_00040)
Adaptive-response sensory-kinase SasA (EBMCFGGG_00043)
IMGPR_plasmid_3300029741_000080 32312:47707(-)
VanR-ABDEGLN family response regulator transcription factor (NF033117.2) (JKLEAHMN_00006)
Paris_I__AriA_AAA15__AriA_AAA15_ali (JKLEAHMN_00004)
Putative septation protein SpoVG (JKLEAHMN_00008)
Adaptive-response sensory-kinase SasA (JKLEAHMN_00005)
PqqA peptide cyclase (JKLEAHMN_00002)
IMGPR_plasmid_3300029558_000160 70:18607(+)
VanR-ABDEGLN family response regulator transcription factor (NF033117.2) (MDIACNHP_00042)
Paris_I__AriA_AAA15__AriA_AAA15_ali (MDIACNHP_00043)
Putative septation protein SpoVG (MDIACNHP_00037)
Adaptive-response sensory-kinase SasA (MDIACNHP_00041)
IMGPR_plasmid_3300006746_000075 30437:48426(-)
Siderophore-dependent iron uptake system (VF0738) (JNOKKEHG_00007)
putative peptide export permease protein YydJ (JNOKKEHG_00006)
Accessory gene regulator protein B (JNOKKEHG_00002)
Putative septation protein SpoVG (JNOKKEHG_00011)
S-adenosylmethionine synthase (JNOKKEHG_00003)
GTP 3',8-cyclase (JNOKKEHG_00005)
IMGPR_plasmid_3300035651_000089 275:18984(+)
Siderophore-dependent iron uptake system (VF0738) (BHNFDJDM_00001)
Putative septation protein SpoVG (BHNFDJDM_00005)
IMGPR_plasmid_3300029453_000092 9:14345(+)
Putative septation protein SpoVG (HBAGPBFH_00007)
Hopanoid C-3 methylase (HBAGPBFH_00001)
Ferredoxin (HBAGPBFH_00003)
IMGPR_plasmid_3300029057_000117 987:17736(+)
IS630 family transposase ISSac1 (AIOBHJOI_00005)
GTP 3',8-cyclase (AIOBHJOI_00002)
Putative septation protein SpoVG (AIOBHJOI_00009)
GTP 3',8-cyclase (AIOBHJOI_00006)
GTP 3',8-cyclase (AIOBHJOI_00003)
IMGPR_plasmid_3300029721_000149 1245:21410(+)
AINIMIPF_00039
IMGPR_plasmid_3300007210_000163 33044:39295(-)
ADLPIPCA_00001
IMGPR_plasmid_3300029404_000219 116:5972(+)
RT_III-AM_III-B_2 (GFCLMOLL_00041)
GFCLMOLL_00040
IMGPR_plasmid_3300023716_000099 32804:44359(-)
MDAMBKCP_00001
IMGPR_plasmid_3300029425_000270 63:10230(+)
8000 nt
Defence
Virulence
AMR