PRK|PRK13752.1: Hg(II)-responsive transcriptional regulator (NF010315.0) (GNKMIDAK_00004)
mercury resistance system transport protein MerF (NF033565.0) (GNKMIDAK_00007)
TIGRFAMs: mercury(II) reductase (TIGR02053.1) (GNKMIDAK_00008)
Mercuric transport protein MerT (GNKMIDAK_00005)
Conjugal transfer protein TraG (GNKMIDAK_00003)
IMGPR_plasmid_3300037195_000052 50:12844(+)
PRK|PRK13752.1: Hg(II)-responsive transcriptional regulator (NF010315.0) (CBMEDGIE_00046)
mercury resistance system transport protein MerF (NF033565.0) (CBMEDGIE_00043)
TIGRFAMs: mercury(II) reductase (TIGR02053.1) (CBMEDGIE_00042)
Mercuric transport protein MerT (CBMEDGIE_00045)
Conjugal transfer protein TraG (CBMEDGIE_00047)
IMGPR_plasmid_3300042269_000086 32851:44976(-)
PRK|PRK13752.1: Hg(II)-responsive transcriptional regulator (NF010315.0) (EAOHELKA_00004)
mercury resistance system transport protein MerF (NF033565.0) (EAOHELKA_00007)
TIGRFAMs: mercury(II) reductase (TIGR02053.1) (EAOHELKA_00008)
Mercuric transport protein MerT (EAOHELKA_00005)
Conjugal transfer protein TraG (EAOHELKA_00003)
IMGPR_plasmid_3300030108_000012 449:13097(+)
PRK|PRK13752.1: Hg(II)-responsive transcriptional regulator (NF010315.0) (FPJDDGGC_00046)
mercury resistance system transport protein MerF (NF033565.0) (FPJDDGGC_00043)
TIGRFAMs: mercury(II) reductase (TIGR02053.1) (FPJDDGGC_00042)
Mercuric transport protein MerT (FPJDDGGC_00045)
Conjugal transfer protein TraG (FPJDDGGC_00047)
IMGPR_plasmid_3300037991_000009 32758:46731(-)
mercury resistance system transport protein MerF (NF033565.0) (NGGJJKAH_00002)
TIGRFAMs: mercury(II) reductase (TIGR02053.1) (NGGJJKAH_00003)
DNA primase TraC (NGGJJKAH_00011)
IMGPR_plasmid_3300042256_000063 368:9859(+)
PRK|PRK13752.1: Hg(II)-responsive transcriptional regulator (NF010315.0) (IILDHNGH_00045)
mercury resistance system transport protein MerF (NF033565.0) (IILDHNGH_00042)
Nickel and cobalt resistance protein CnrB (IILDHNGH_00051)
Outer membrane protein TolC (IILDHNGH_00056)
TIGRFAMs: mercury(II) reductase (TIGR02053.1) (IILDHNGH_00041)
csa3_I_II_III_IV_V_VI_2 (IILDHNGH_00049)
Sulfide-quinone reductase (IILDHNGH_00053)
Mercuric transport protein MerT (IILDHNGH_00044)
Thioredoxin 2 (IILDHNGH_00054)
Conjugal transfer protein TraG (IILDHNGH_00046)
Swarming motility protein SwrC (IILDHNGH_00050)
IMGPR_plasmid_3300042275_000202 32545:53741(-)
NEPGBOCK_00026
NEPGBOCK_00024
NEPGBOCK_00025
IMGPR_plasmid_2582580772_000006 16221:23764(-)
OHECIICD_00001
OHECIICD_00002
IMGPR_plasmid_3300037784_000221 252:5528(+)
ONHAGCNK_00026
ONHAGCNK_00025
ONHAGCNK_00027
IMGPR_plasmid_3300003979_000047 19095:25463(-)
AGBENIBK_00040
AGBENIBK_00041
IMGPR_plasmid_3300009779_000196 32846:38206(-)
BOBLGAFF_00002
BOBLGAFF_00001
IMGPR_plasmid_3300001732_000026 123:5187(+)
DJAFNOIJ_00027
DJAFNOIJ_00026
DJAFNOIJ_00025
IMGPR_plasmid_3300036764_000380 19552:24977(-)
Quinohemoprotein alcohol dehydrogenase ADH IIB (IJIHKAAC_00052)
putative enoyl-CoA hydratase echA8 (IJIHKAAC_00048)
Acetyl-CoA:oxalate CoA-transferase (IJIHKAAC_00051)
Coniferyl aldehyde dehydrogenase (IJIHKAAC_00045)
3-ketoacyl-CoA thiolase (IJIHKAAC_00049)
Vitamin B12 transporter BtuB (IJIHKAAC_00043)
Mycobactin (VF0299) (IJIHKAAC_00046)
Conjugal transfer protein TraG (IJIHKAAC_00041)
FadE29 (VF0845) (IJIHKAAC_00050)
putative oxidoreductase (IJIHKAAC_00044)
IMGPR_plasmid_3300017702_000073 31906:57831(-)
putative enoyl-CoA hydratase echA8 (BAAIAJAG_00002)
Coniferyl aldehyde dehydrogenase (BAAIAJAG_00005)
Vitamin B12 transporter BtuB (BAAIAJAG_00007)
Mycobactin (VF0299) (BAAIAJAG_00004)
Conjugal transfer protein TraG (BAAIAJAG_00009)
putative oxidoreductase (BAAIAJAG_00006)
IMGPR_plasmid_3300017706_000048 551:18614(+)
10000 nt
Defence
Virulence
AMR