

adapt_MexA-NCBIFAM multidrug efflux RND transporter periplasmic adaptor subunit MexA (NF033834.1) (AKLOBHHC_00101)
 putative HTH-type transcriptional regulator YtcD (AKLOBHHC_00105)
 HTH-type transcriptional regulator SarZ (AKLOBHHC_00098)
 Toluene efflux pump membrane transporter TtgE (AKLOBHHC_00100)
 4,5-DOPA dioxygenase extradiol (AKLOBHHC_00106)
 Transcriptional regulator SlyA (AKLOBHHC_00110)
 FarAB (VF0450) (AKLOBHHC_00111)
 IMGPR_plasmid_3300025059_000096: 104,359-116,383 (+)
 putative HTH-type transcriptional regulator YtcD (JFPLPDGI_00104)
 IMGPR_plasmid_3300006790_000233: 104,355-111,873 (+)
 High-affinity branched-chain amino acid transport system permease protein LivH (HONGJMAG_00002)
 putative HTH-type transcriptional regulator YtcD (HONGJMAG_00011)
 ABC-type heme transporter (VF0739) (HONGJMAG_00005)
 4,5-DOPA dioxygenase extradiol (HONGJMAG_00010)
 IS6 family transposase ISPpa9 (HONGJMAG_00007)
 mads_8 (HONGJMAG_00004)
 IMGPR_plasmid_3300045789_000256: 1,314-18,113 (-)
 putative HTH-type transcriptional regulator YtcD (KCHALAGA_00104)
 ABC-type heme transporter (VF0739) (KCHALAGA_00110)
 4,5-DOPA dioxygenase extradiol (KCHALAGA_00105)
 IS6 family transposase ISPpa9 (KCHALAGA_00108)
 mads_8 (KCHALAGA_00111)
 IMGPR_plasmid_3300046449_000140: 104,377-119,253 (+)
 putative HTH-type transcriptional regulator YtcD (OBANJBOG_00104)
 Thioldisulfide interchange protein CycY (OBANJBOG_00121)
 Thioldisulfide interchange protein TlpA (OBANJBOG_00120)
 Thioldisulfide interchange protein DsbD (OBANJBOG_00116)
 4,5-DOPA dioxygenase extradiol (OBANJBOG_00105)
 IS6 family transposase ISRssp3 (OBANJBOG_00112)
 Transcriptional regulator SlyA (OBANJBOG_00109)
 Putative permease (OBANJBOG_00115)
 FarAB (VF0450) (OBANJBOG_00110)
 Multicopper oxidase MmcO (OBANJBOG_00125)
 Sigma L (VF0855) (OBANJBOG_00130)
 SigE (VF0295) (OBANJBOG_00136)
 Cytochrome c6 (OBANJBOG_00134)
 Paa (VF0194) (OBANJBOG_00132)
 TnsC_O_V-K (OBANJBOG_00140)
 IMGPR_plasmid_3300037195_000035: 104,298-141,521 (+)
 putative HTH-type transcriptional regulator YtcD (BDPPFABB_00036)
 Thioldisulfide interchange protein CycY (BDPPFABB_00019)
 Thioldisulfide interchange protein TlpA (BDPPFABB_00020)
 Thioldisulfide interchange protein DsbD (BDPPFABB_00024)
 4,5-DOPA dioxygenase extradiol (BDPPFABB_00035)
 Transcriptional regulator SlyA (BDPPFABB_00031)
 Multicopper oxidase MmcO (BDPPFABB_00015)
 IS6 family transposase ISRssp3 (BDPPFABB_00028)
 Putative permease (BDPPFABB_00025)
 FarAB (VF0450) (BDPPFABB_00030)
 Sigma L (VF0855) (BDPPFABB_00009)
 SigE (VF0295) (BDPPFABB_00003)
 Paa (VF0194) (BDPPFABB_00007)
 Cytochrome c6 (BDPPFABB_00005)
 IMGPR_plasmid_3300030334_000085: 784-34,334 (-)
 putative HTH-type transcriptional regulator YtcD (LPBPBBLD_00046)
 Thioldisulfide interchange protein CycY (LPBPBBLD_00029)
 Thioldisulfide interchange protein TlpA (LPBPBBLD_00030)
 Thioldisulfide interchange protein DsbD (LPBPBBLD_00034)
 4,5-DOPA dioxygenase extradiol (LPBPBBLD_00045)
 IS6 family transposase ISRssp3 (LPBPBBLD_00038)
 Transcriptional regulator SlyA (LPBPBBLD_00041)
 Multicopper oxidase MmcO (LPBPBBLD_00025)
 Putative permease (LPBPBBLD_00035)
 FarAB (VF0450) (LPBPBBLD_00040)
 Sigma L (VF0855) (LPBPBBLD_00019)
 SigE (VF0295) (LPBPBBLD_00013)
 Cytochrome c6 (LPBPBBLD_00015)
 Segregation and condensation protein B (LPBPBBLD_00007)
 Nucleoid occlusion protein (LPBPBBLD_00003)
 TnsC_O_V-K (LPBPBBLD_00009)
 IMGPR_plasmid_3300030100_000069: 411-44,301 (-)
 10000 nt
 defence (PADLOC) virulence (VFDB) AMR (AMRFinderPlus)