TIGRFAMs: mercury resistance system periplasmic binding protein MerP (TIGR02052.1) (CDS_0040)
PRK|PRK13747.1: broad-spectrum mercury transporter MerE (NF010310.2) (CDS_0036)
mercury resistance co-regulator MerD (NF033783.1) (CDS_0037)
organomercurial lyase MerB (NF033555.1) (CDS_0038)
TIGRFAMs: mercury(II) reductase (TIGR02053.1) (CDS_0039)
PlcR-PapR quorum sensing (VF0660) (CDS_0035)
transposase (CDS_0034)
transposase (CDS_0043)
TemPhD_cluster_57480 15887:36057(+)
TemPhD_cluster_57748 15887:34731(+)
TIGRFAMs: mercury resistance system periplasmic binding protein MerP (TIGR02052.1) (CDS_0039)
TemPhD_cluster_26821 15887:34618(+)
TemPhD_cluster_57965 15887:33864(+)
TIGRFAMs: mercury resistance system periplasmic binding protein MerP (TIGR02052.1) (CDS_0041)
TemPhD_cluster_57475 15523:36123(+)
TemPhD_cluster_58079 15523:33273(+)
TemPhD_cluster_58033 15887:33520(+)
TemPhD_cluster_16782 15887:33705(+)
TemPhD_cluster_58025 15887:33486(+)
TIGRFAMs: mercury resistance system periplasmic binding protein MerP (TIGR02052.1) (CDS_0035)
TemPhD_cluster_18301 15887:30215(+)
TemPhD_cluster_46473 16372:34495(+)
TemPhD_cluster_39379_ 16372:36725(+)
TemPhD_cluster_26801 15887:29637(+)
TemPhD_cluster_46818 17201:31905(+)
CDS_0033
TemPhD_cluster_48222 15523:25946(+)
CDS_0034
TemPhD_cluster_32029 15887:23731(+)
TIGRFAMs: mercury(II) reductase (TIGR02053.1) (CDS_0039)
transposase (CDS_0047)
TemPhD_cluster_46118 17201:40729(+)
ribonucleotide reductase large subunit (CDS_0039)
transposase (CDS_0046)
TemPhD_cluster_57136 16419:38814(+)
9000 nt
DNA, RNA and nucleotide metabolism
integration and excision
unknown function
Virulence
AMR