IMGVR_UViG_3300029340_000360|3300029340|Ga0243727_1000104|167-59026 30240:40807(+)
IMGVR_UViG_3300045988_083527|3300045988|Ga0495776_026620 31217:39663(+)
MGV-GENOME-0330964 31265:38676(+)
IMGVR_UViG_3300029445_000481|3300029445|Ga0243741_1000525|3-56610 31187:40257(+)
MGV-GENOME-0325658 31909:38896(+)
MGV-GENOME-0322435 31242:39081(+)
IMGVR_UViG_3300045988_069419|3300045988|Ga0495776_151609 31265:38676(+)
uvig_402667 31169:41723(+)
IMGVR_UViG_3300045988_067086|3300045988|Ga0495776_130683 30839:39462(+)
sporulation stage III protein D (CDS_0053)
integrase (CDS_0054)
uvig_98230 31425:41551(+)
IMGVR_UViG_3300045988_089587|3300045988|Ga0495776_005076 31477:39103(+)
MGV-GENOME-0324697 31409:38858(+)
IMGVR_UViG_3300045988_063469|3300045988|Ga0495776_183676 31359:38299(+)
uvig_580917 31182:39324(+)
MGV-GENOME-0335906 31829:39659(+)
IMGVR_UViG_3300045988_109741|3300045988|Ga0495776_151751 31422:41923(+)
IMGVR_UViG_3300045988_063255|3300045988|Ga0495776_130717 31896:39503(+)
MGV-GENOME-0292549 19304:28062(+)
radical SAM domain-containing protein (CDS_0061)
uvig_529275 31811:46290(+)
IMGVR_UViG_3300030040_000035|3300030040|Ga0134560_1000173 31371:39950(+)
uvig_374148 31371:39942(+)
IMGVR_UViG_3300045988_037170|3300045988|Ga0495776_026537 31371:40235(+)
Rampelli_10024_NODE_173_length_56009_cov_49.679254 31371:39906(+)
MGV-GENOME-0307711 31371:39303(+)
MazF-like growth inhibitor (CDS_0053)
transposase (CDS_0056)
IMGVR_UViG_3300029773_000317|3300029773|Ga0243856_1000101|46432-107600 31417:44732(+)
MazF-like growth inhibitor (CDS_0053)
transposase (CDS_0055)
IMGVR_UViG_3300029350_000245|3300029350|Ga0243859_100069|36963-98031 31417:44632(+)
transposase (CDS_0053)
IMGVR_UViG_3300029416_000334|3300029416|Ga0243860_100207|2-59478 31417:43040(+)
transposase (CDS_0052)
IMGVR_UViG_3300029411_000268|3300029411|Ga0243861_100177|140-58807 31417:42231(+)
transposase (CDS_0058)
IMGVR_UViG_3300029761_000088|3300029761|Ga0243857_100367 31417:48947(+)
MazF-like growth inhibitor (CDS_0075)
ParB-like partition protein (CDS_0076)
translation elongation factor (CDS_0053)
RtxA (VF0482) (CDS_0054)
DNA binding protein (CDS_0067)
transposase (CDS_0077)
integrase (CDS_0080)
uvig_452930 31417:69248(+)
sporulation stage III protein D (CDS_0054)
RNA polymerase sigma factor (CDS_0056)
RtxA (VF0482) (CDS_0058)
SigF (VF0294) (CDS_0055)
SAMEA1906523_a1_ct92878_vs1 31826:52298(+)
uvig_584377 30293:39387(+)
IMGVR_UViG_3300045988_065898|3300045988|Ga0495776_130278 30504:39110(+)
IMGVR_UViG_3300045988_052031|3300045988|Ga0495776_005060 30409:37707(+)
MGV-GENOME-0322298 30409:38122(+)
MGV-GENOME-0324031 30841:39951(+)
MazF-like growth inhibitor (CDS_0073)
ParB-like partition protein (CDS_0074)
Serine-threonine phosphatase stp (VF0787) (CDS_0056)
integrase (CDS_0068)
integrase (CDS_0077)
uvig_446203 30222:67433(+)
cyanobacterial phosphoribosylglycinamide formyltransferase (CDS_0053)
phospho-2-dehydro-3-deoxyheptonate aldolase (CDS_0058)
phosphoribosylaminoimidazole synthetase (CDS_0054)
DNA polymerase (CDS_0066)
uvig_243762 31665:55366(+)
10000 nt
DNA, RNA and nucleotide metabolism
integration and excision
moron, auxiliary metabolic gene and host takeover
other
transcription regulation
unknown function
Virulence