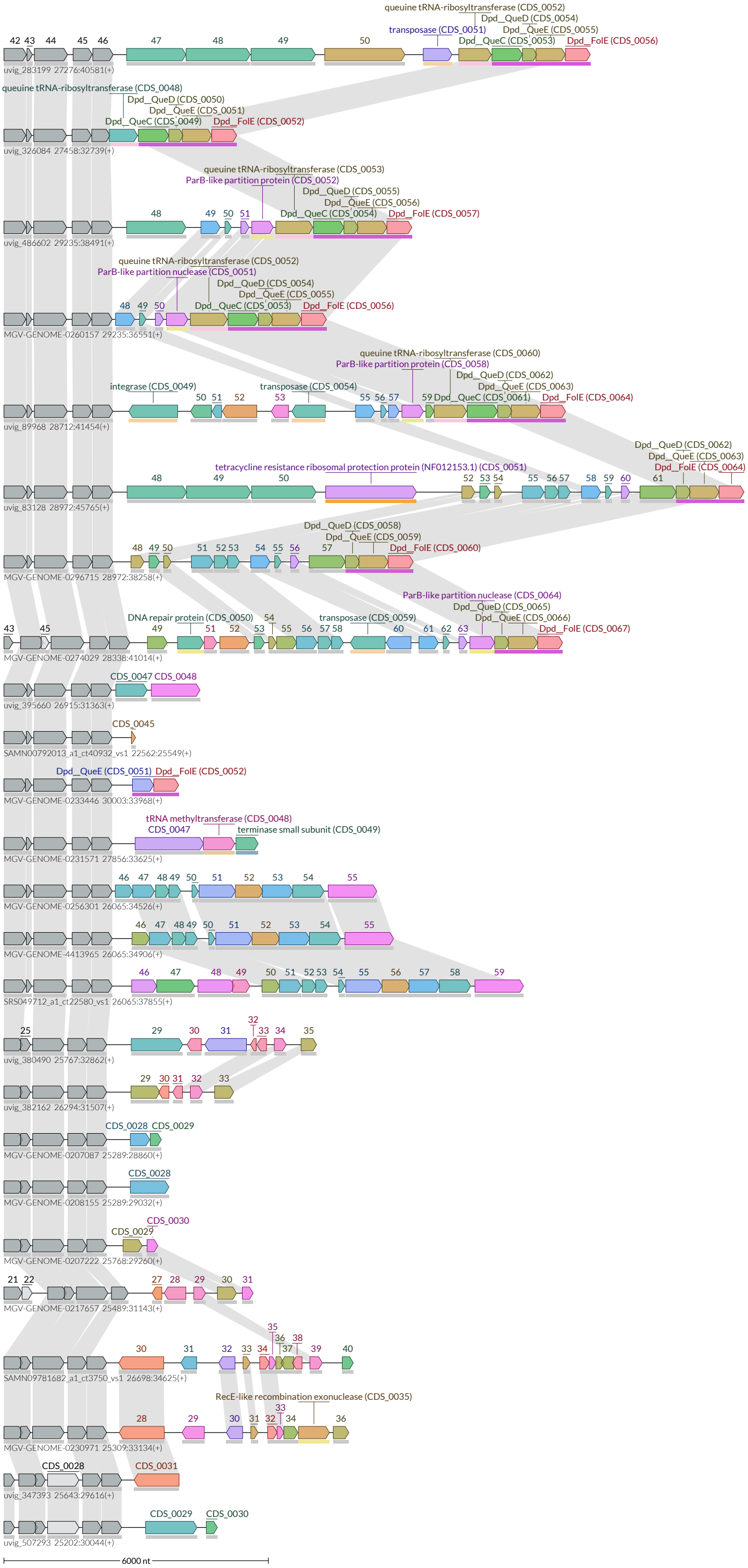
queuine tRNA-ribosyltransferase (CDS_0052)
Dpd__QueD (CDS_0054)
transposase (CDS_0051)
Dpd__QueE (CDS_0055)
Dpd__QueC (CDS_0053)
Dpd__FolE (CDS_0056)
uvig_283199 27276:40581(+)
queuine tRNA-ribosyltransferase (CDS_0048)
Dpd__QueD (CDS_0050)
Dpd__QueE (CDS_0051)
Dpd__QueC (CDS_0049)
Dpd__FolE (CDS_0052)
uvig_326084 27458:32739(+)
queuine tRNA-ribosyltransferase (CDS_0053)
ParB-like partition protein (CDS_0052)
Dpd__QueD (CDS_0055)
Dpd__QueE (CDS_0056)
Dpd__QueC (CDS_0054)
Dpd__FolE (CDS_0057)
uvig_486602 29235:38491(+)
queuine tRNA-ribosyltransferase (CDS_0052)
ParB-like partition nuclease (CDS_0051)
Dpd__QueD (CDS_0054)
Dpd__QueE (CDS_0055)
Dpd__QueC (CDS_0053)
Dpd__FolE (CDS_0056)
MGV-GENOME-0260157 29235:36551(+)
queuine tRNA-ribosyltransferase (CDS_0060)
ParB-like partition protein (CDS_0058)
Dpd__QueD (CDS_0062)
Dpd__QueE (CDS_0063)
integrase (CDS_0049)
transposase (CDS_0054)
Dpd__QueC (CDS_0061)
Dpd__FolE (CDS_0064)
uvig_89968 28712:41454(+)
Dpd__QueD (CDS_0062)
Dpd__QueE (CDS_0063)
Dpd__FolE (CDS_0064)
tetracycline resistance ribosomal protection protein (NF012153.1) (CDS_0051)
uvig_83128 28972:45765(+)
Dpd__QueD (CDS_0058)
Dpd__QueE (CDS_0059)
Dpd__FolE (CDS_0060)
MGV-GENOME-0296715 28972:38258(+)
ParB-like partition nuclease (CDS_0064)
Dpd__QueD (CDS_0065)
Dpd__QueE (CDS_0066)
DNA repair protein (CDS_0050)
transposase (CDS_0059)
Dpd__FolE (CDS_0067)
MGV-GENOME-0274029 28338:41014(+)
CDS_0047 CDS_0048
uvig_395660 26915:31363(+)
CDS_0045
SAMN00792013_a1_ct40932_vs1 22562:25549(+)
Dpd__QueE (CDS_0051) Dpd__FolE (CDS_0052)
MGV-GENOME-0233446 30003:33968(+)
tRNA methyltransferase (CDS_0048)
CDS_0047
terminase small subunit (CDS_0049)
MGV-GENOME-0231571 27856:33625(+)
MGV-GENOME-0256301 26065:34526(+)
MGV-GENOME-4413965 26065:34906(+)
SRS049712_a1_ct22580_vs1 26065:37855(+)
uvig_380490 25767:32862(+)
uvig_382162 26294:31507(+)
CDS_0028 CDS_0029
MGV-GENOME-0207087 25289:28860(+)
CDS_0028
MGV-GENOME-0208155 25289:29032(+)
CDS_0030
CDS_0029
MGV-GENOME-0207222 25768:29260(+)
MGV-GENOME-0217657 25489:31143(+)
SAMN09781682_a1_ct3750_vs1 26698:34625(+)
RecE-like recombination exonuclease (CDS_0035)
MGV-GENOME-0230971 25309:33134(+)
CDS_0028
CDS_0031
uvig_347393 25643:29616(+)
CDS_0029 CDS_0030
uvig_507293 25202:30044(+)
6000 nt
DNA, RNA and nucleotide metabolism
head and packaging
integration and excision
moron, auxiliary metabolic gene and host takeover
other
unknown function
Defence
AMR