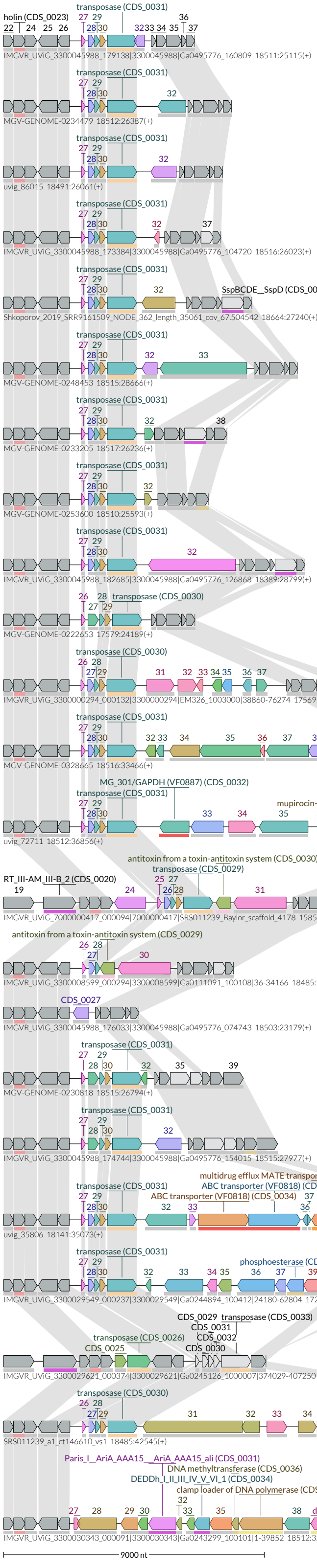
holin (CDS_0023)
transposase (CDS_0031)
IMGVR_UViG_3300045988_179138|3300045988|Ga0495776_160809 18511:25115(+)
MGV-GENOME-0234479 18512:26387(+)
uvig_86015 18491:26061(+)
IMGVR_UViG_3300045988_173384|3300045988|Ga0495776_104720 18516:26023(+)
SspBCDE__SspD (CDS_0037)
Shkoporov_2019_SRR9161509_NODE_362_length_35061_cov_67.504542 18664:27240(+)
MGV-GENOME-0248453 18515:28666(+)
MGV-GENOME-0233205 18517:26236(+)
MGV-GENOME-0253600 18510:25593(+)
IMGVR_UViG_3300045988_182685|3300045988|Ga0495776_126868 18389:28799(+)
transposase (CDS_0030)
MGV-GENOME-0222653 17579:24189(+)
IMGVR_UViG_3300000294_000132|3300000294|EM326_1003000|38860-76274 17569:29594(+)
MGV-GENOME-0328665 18516:33466(+)
MG_301/GAPDH (VF0887) (CDS_0032)
mupirocin-resistant isoleucine--tRNA ligase MupA (NF000257.1) (CDS_0038)
uvig_72711 18512:36856(+)
antitoxin from a toxin-antitoxin system (CDS_0030)
RT_III-AM_III-B_2 (CDS_0020)
IMGVR_UViG_7000000417_000094|7000000417|SRS011239_Baylor_scaffold_4178 15855:27797(+)
antitoxin from a toxin-antitoxin system (CDS_0029)
IMGVR_UViG_3300008599_000294|3300008599|Ga0111091_100108|36-34166 18485:26426(+)
CDS_0027
IMGVR_UViG_3300045988_176033|3300045988|Ga0495776_074743 18503:23179(+)
MGV-GENOME-0230818 18515:26794(+)
IMGVR_UViG_3300045988_174744|3300045988|Ga0495776_154015 18515:27977(+)
multidrug efflux MATE transporter MepA (NF000131.1) (CDS_0038)
ABC transporter (VF0818) (CDS_0035)
ABC transporter (VF0818) (CDS_0034)
uvig_35806 18141:35073(+)
phosphoesterase (CDS_0038)
IMGVR_UViG_3300029549_000237|3300029549|Ga0244894_100412|24180-62804 17292:31237(+)
CDS_0029
CDS_0031
CDS_0032
CDS_0030
transposase (CDS_0033)
transposase (CDS_0026)
CDS_0025
IMGVR_UViG_3300029621_000374|3300029621|Ga0245126_1000007|374029-407250 15881:24942(+)
antitoxin from a toxin-antitoxin system (CDS_0045)
thymidylate kinase (CDS_0043)
SRS011239_a1_ct146610_vs1 18485:42545(+)
Paris_I__AriA_AAA15__AriA_AAA15_ali (CDS_0031)
DNA methyltransferase (CDS_0036)
DEDDh_I_II_III_IV_V_VI_1 (CDS_0034)
clamp loader of DNA polymerase (CDS_0037)
dCMP deaminase (CDS_0039)
IMGVR_UViG_3300030343_000091|3300030343|Ga0243299_100101|1-39852 18512:32034(+)
9000 nt
DNA, RNA and nucleotide metabolism
integration and excision
lysis
moron, auxiliary metabolic gene and host takeover
other
unknown function
Defence
Virulence
AMR