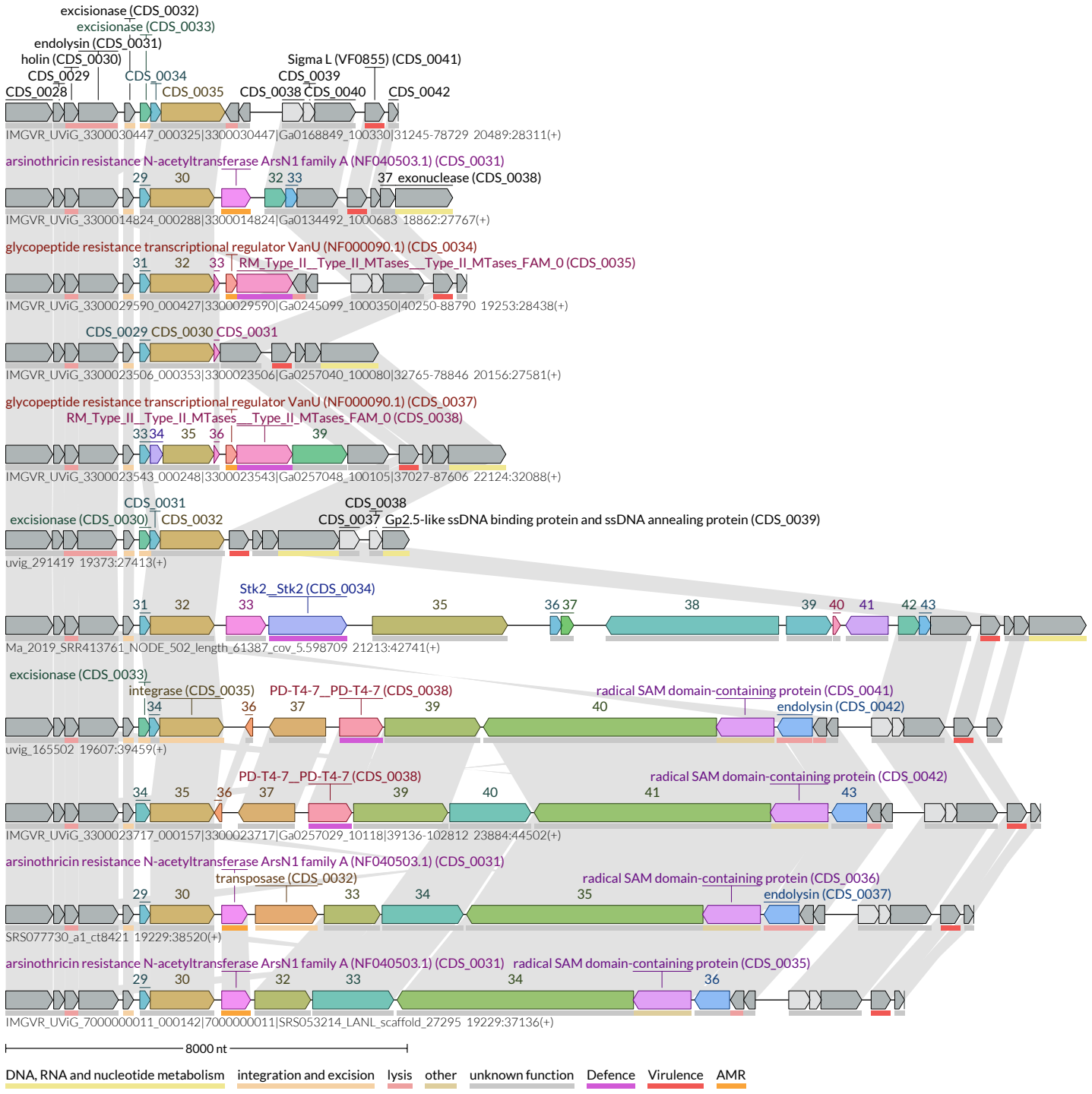
excisionase (CDS_0032)
excisionase (CDS_0033)
endolysin (CDS_0031)
holin (CDS_0030)
CDS_0029
CDS_0028
CDS_0034
CDS_0035
Sigma L (VF0855) (CDS_0041)
CDS_0039
CDS_0038
CDS_0040
CDS_0042
IMGVR_UViG_3300030447_000325|3300030447|Ga0168849_100330|31245-78729 20489:28311(+)
arsinothricin resistance N-acetyltransferase ArsN1 family A (NF040503.1) (CDS_0031)
29
30
32
33
37 exonuclease (CDS_0038)
IMGVR_UViG_3300014824_000288|3300014824|Ga0134492_1000683 18862:27767(+)
glycopeptide resistance transcriptional regulator VanU (NF000090.1) (CDS_0034)
31
32
33
RM_Type_II__Type_II_MTases__Type_II_MTases_FAM_0 (CDS_0035)
IMGVR_UViG_3300029590_000427|3300029590|Ga0245099_1000350|40250-88790 19253:28438(+)
CDS_0029
CDS_0030
CDS_0031
IMGVR_UViG_3300023506_000353|3300023506|Ga0257040_100080|32765-78846 20156:27581(+)
glycopeptide resistance transcriptional regulator VanU (NF000090.1) (CDS_0037)
RM_Type_II__Type_II_MTases__Type_II_MTases_FAM_0 (CDS_0038)
33
34
35
36
39
IMGVR_UViG_3300023543_000248|3300023543|Ga0257048_100105|37027-87606 22124:32088(+)
CDS_0031
excisionase (CDS_0030)
CDS_0032
CDS_0038
CDS_0037
Gp2.5-like ssDNA binding protein and ssDNA annealing protein (CDS_0039)
uvig_291419 19373:27413(+)
Stk2__Stk2 (CDS_0034)
31
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43
Ma_2019_SRR413761_NODE_502_length_61387_cov_5.598709 21213:42741(+)
excisionase (CDS_0033)
integrase (CDS_0035)
PD-T4-7__PD-T4-7 (CDS_0038)
radical SAM domain-containing protein (CDS_0041)
endolysin (CDS_0042)
34
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uvig_165502 19607:39459(+)
PD-T4-7__PD-T4-7 (CDS_0038)
radical SAM domain-containing protein (CDS_0042)
34
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43
IMGVR_UViG_3300023717_000157|3300023717|Ga0257029_10118|39136-102812 23884:44502(+)
arsinothricin resistance N-acetyltransferase ArsN1 family A (NF040503.1) (CDS_0031)
transposase (CDS_0032)
radical SAM domain-containing protein (CDS_0036)
endolysin (CDS_0037)
29
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SRS077730_a1_ct8421 19229:38520(+)
arsinothricin resistance N-acetyltransferase ArsN1 family A (NF040503.1) (CDS_0031)
radical SAM domain-containing protein (CDS_0035)
29
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IMGVR_UViG_7000000011_000142|7000000011|SRS053214_LANL_scaffold_27295 19229:37136(+)
8000 nt
DNA, RNA and nucleotide metabolism
integration and excision
lysis
other
unknown function
Defence
Virulence
AMR