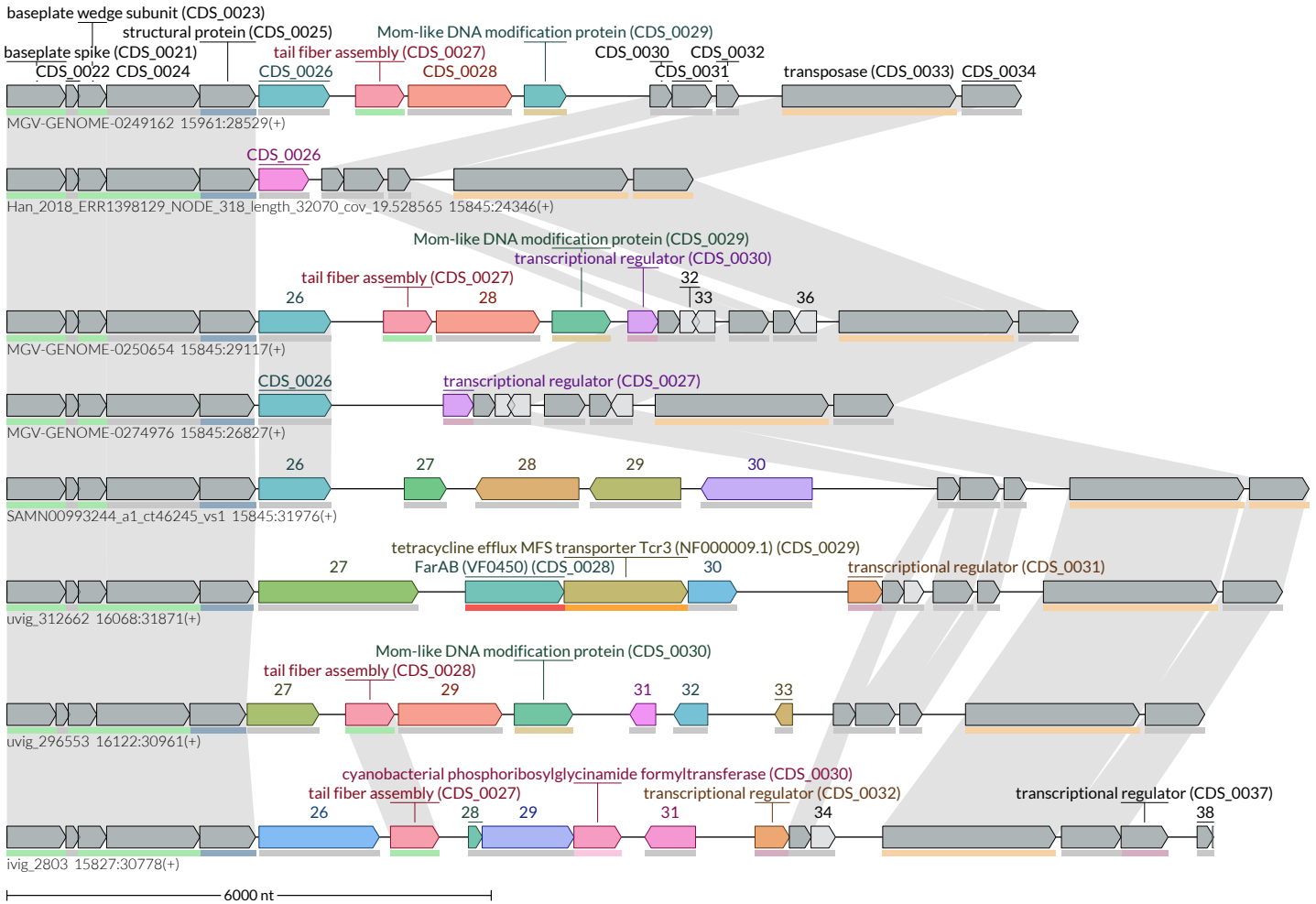
baseplate wedge subunit (CDS_0023)
structural protein (CDS_0025)
baseplate spike (CDS_0021)
CDS_0022
CDS_0024
Mom-like DNA modification protein (CDS_0029)
tail fiber assembly (CDS_0027)
CDS_0026
CDS_0028
CDS_0030
CDS_0032
CDS_0031
transposase (CDS_0033)
CDS_0034
MGV-GENOME-0249162 15961:28529(+)
CDS_0026
Han_2018_ERR1398129_NODE_318_length_32070_cov_19.528565 15845:24346(+)
Mom-like DNA modification protein (CDS_0029)
transcriptional regulator (CDS_0030)
tail fiber assembly (CDS_0027)
26
28
32
33
36
MGV-GENOME-0250654 15845:29117(+)
CDS_0026
transcriptional regulator (CDS_0027)
MGV-GENOME-0274976 15845:26827(+)
26
27
28
29
30
SAMN00993244_a1_ct46245_vs1 15845:31976(+)
tetracycline efflux MFS transporter Tcr3 (NF000009.1) (CDS_0029)
27
FarAB (VF0450) (CDS_0028)
30
transcriptional regulator (CDS_0031)
uvig_312662 16068:31871(+)
Mom-like DNA modification protein (CDS_0030)
tail fiber assembly (CDS_0028)
27
29
31
32
33
uvig_296553 16122:30961(+)
cyanobacterial phosphoribosylglycinamide formyltransferase (CDS_0030)
tail fiber assembly (CDS_0027)
transcriptional regulator (CDS_0032)
transcriptional regulator (CDS_0037)
26
28
29
31
34
38
ivig_2803 15827:30778(+)
6000 nt
DNA, RNA and nucleotide metabolism
head and packaging
integration and excision
moron, auxiliary metabolic gene and host takeover
other
tail
transcription regulation
unknown function
Virulence
AMR