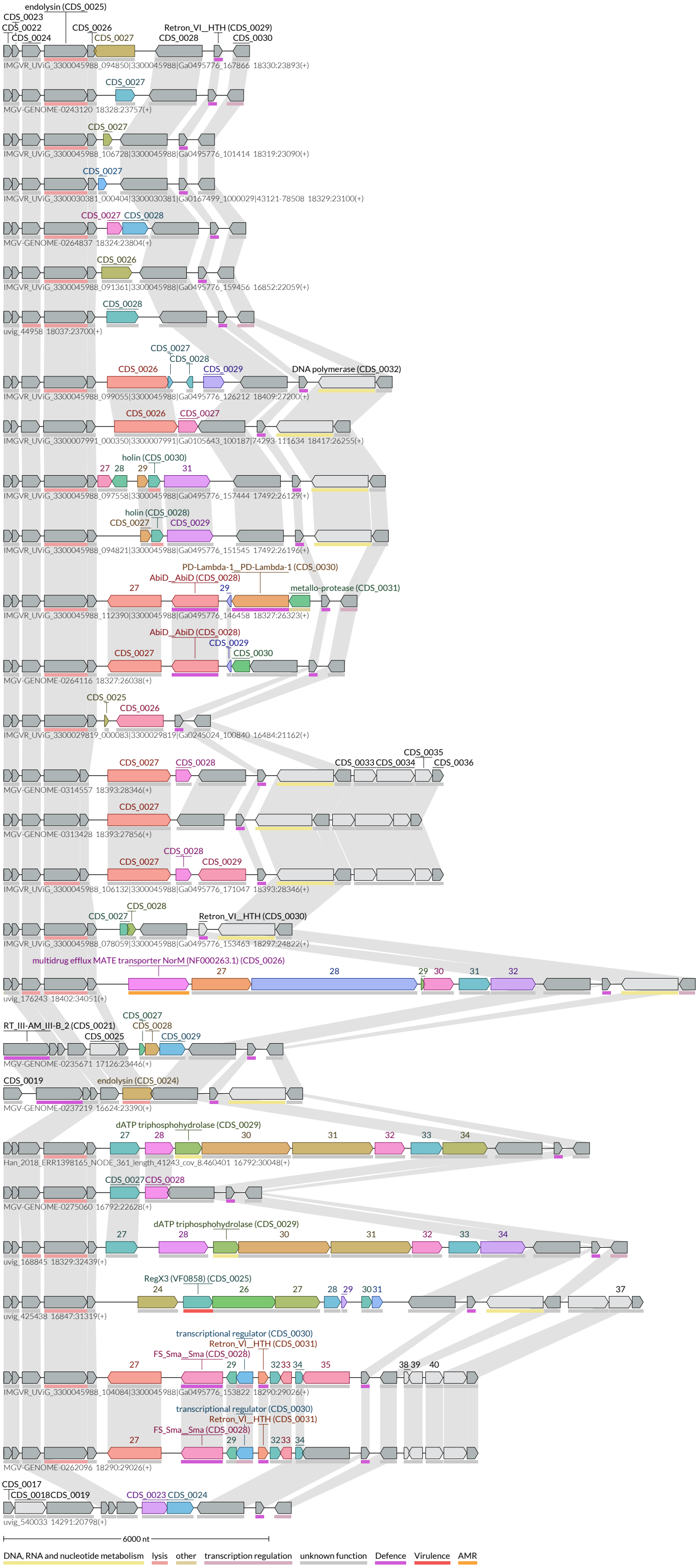
endolysin (CDS_0025)
CDS_0023
CDS_0022
CDS_0024
CDS_0026
CDS_0027
Retron_VI__HTH (CDS_0029)
CDS_0028
CDS_0030
IMGVR_UViG_3300045988_094850|3300045988|Ga0495776_167866 18330:23893(+)
CDS_0027
MGV-GENOME-0243120 18328:23757(+)
CDS_0027
IMGVR_UViG_3300045988_106728|3300045988|Ga0495776_101414 18319:23090(+)
CDS_0027
IMGVR_UViG_3300030381_000404|3300030381|Ga0167499_1000029|43121-78508 18329:23100(+)
CDS_0027 CDS_0028
MGV-GENOME-0264837 18324:23804(+)
CDS_0026
IMGVR_UViG_3300045988_091361|3300045988|Ga0495776_159456 16852:22059(+)
CDS_0028
uvig_44958 18037:23700(+)
CDS_0027
CDS_0028
CDS_0026
CDS_0029
DNA polymerase (CDS_0032)
IMGVR_UViG_3300045988_099055|3300045988|Ga0495776_126212 18409:27200(+)
CDS_0026
CDS_0027
IMGVR_UViG_3300007991_000350|3300007991|Ga0105643_100187|74293-111634 18417:26255(+)
holin (CDS_0030)
27
28
29
31
IMGVR_UViG_3300045988_097558|3300045988|Ga0495776_157444 17492:26129(+)
holin (CDS_0028)
CDS_0027
CDS_0029
IMGVR_UViG_3300045988_094821|3300045988|Ga0495776_151545 17492:26196(+)
PD-Lambda-1__PD-Lambda-1 (CDS_0030)
AbiD__AbiD (CDS_0028)
27
29
metallo-protease (CDS_0031)
IMGVR_UViG_3300045988_112390|3300045988|Ga0495776_146458 18327:26323(+)
AbiD__AbiD (CDS_0028)
CDS_0029
CDS_0027
CDS_0030
MGV-GENOME-0264116 18327:26038(+)
CDS_0025
CDS_0026
IMGVR_UViG_3300029819_000083|3300029819|Ga0245024_100840 16484:21162(+)
CDS_0035
CDS_0027
CDS_0028
CDS_0033
CDS_0034
CDS_0036
MGV-GENOME-0314557 18393:28346(+)
CDS_0027
MGV-GENOME-0313428 18393:27856(+)
CDS_0028
CDS_0027
CDS_0029
IMGVR_UViG_3300045988_106132|3300045988|Ga0495776_171047 18393:28346(+)
CDS_0028
CDS_0027
Retron_VI__HTH (CDS_0030)
IMGVR_UViG_3300045988_078059|3300045988|Ga0495776_153463 18297:24822(+)
multidrug efflux MATE transporter NorM (NF000263.1) (CDS_0026)
27
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32
uvig_176243 18402:34051(+)
CDS_0027
RT_III-AM_III-B_2 (CDS_0021)
CDS_0028
CDS_0025
CDS_0029
MGV-GENOME-0235671 17126:23446(+)
CDS_0019
endolysin (CDS_0024)
MGV-GENOME-0237219 16624:23390(+)
dATP triphosphohydrolase (CDS_0029)
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Han_2018_ERR1398165_NODE_361_length_41243_cov_8.460401 16792:30048(+)
CDS_0027
CDS_0028
MGV-GENOME-0275060 16792:22628(+)
dATP triphosphohydrolase (CDS_0029)
27
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uvig_168845 18329:32439(+)
RegX3 (VF0858) (CDS_0025)
24
26
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37
uvig_425438 16847:31319(+)
transcriptional regulator (CDS_0030)
Retron_VI__HTH (CDS_0031)
FS_Sma__Sma (CDS_0028)
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IMGVR_UViG_3300045988_104084|3300045988|Ga0495776_153822 18290:29026(+)
transcriptional regulator (CDS_0030)
Retron_VI__HTH (CDS_0031)
FS_Sma__Sma (CDS_0028)
27
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MGV-GENOME-0262096 18290:29026(+)
CDS_0017
CDS_0018
CDS_0019
CDS_0023
CDS_0024
uvig_540033 14291:20798(+)
6000 nt
DNA, RNA and nucleotide metabolism
lysis
other
transcription regulation
unknown function
Defence
Virulence
AMR