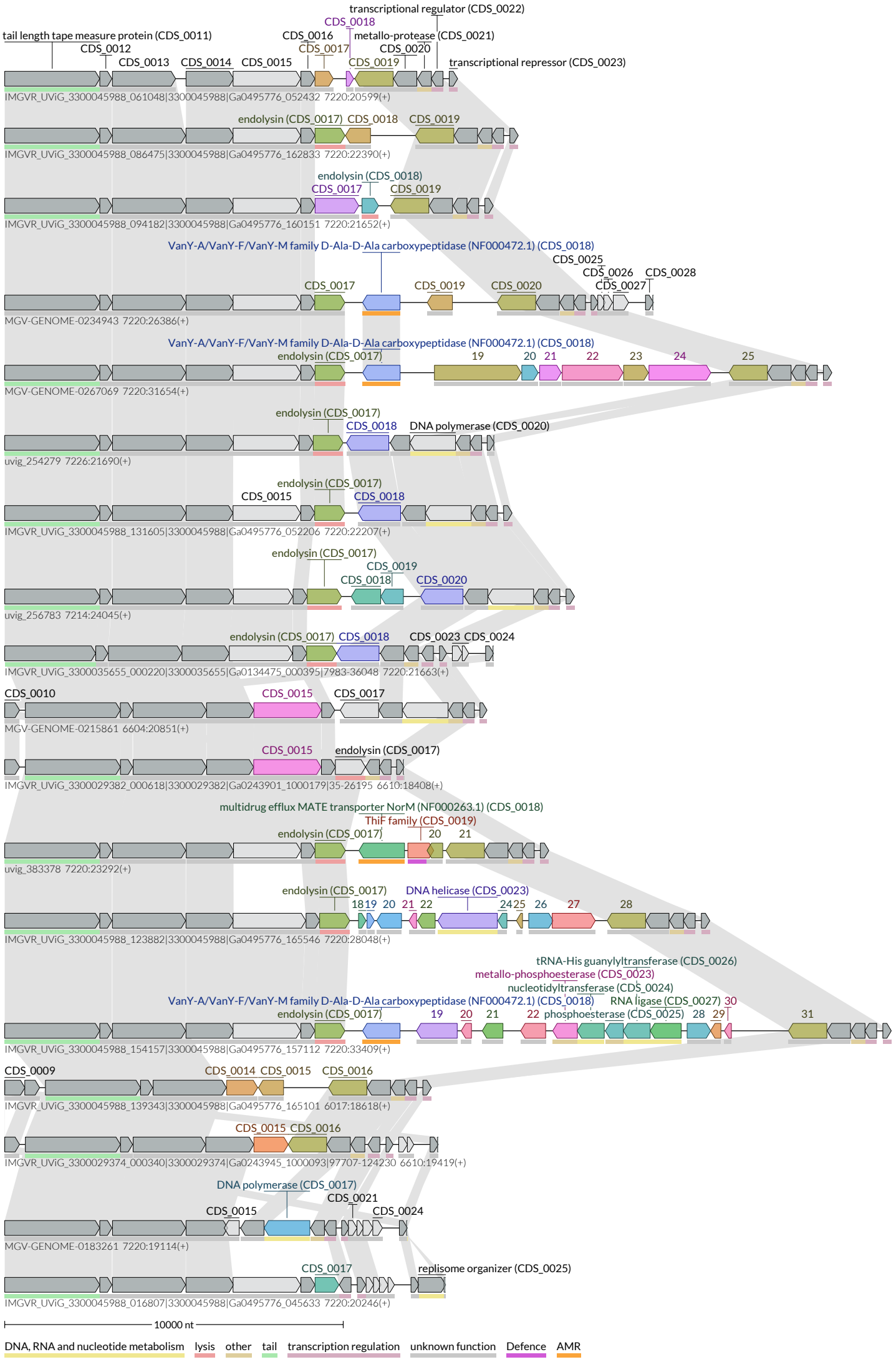
transcriptional regulator (CDS_0022)
CDS_0018
tail length tape measure protein (CDS_0011)
CDS_0016
metallo-protease (CDS_0021)
CDS_0012
CDS_0017
CDS_0020
CDS_0013
CDS_0014
CDS_0015
CDS_0019
transcriptional repressor (CDS_0023)
IMGVR_UViG_3300045988_061048|3300045988|Ga0495776_052432 7220:20599(+)
endolysin (CDS_0017)
CDS_0018
CDS_0019
IMGVR_UViG_3300045988_086475|3300045988|Ga0495776_162833 7220:22390(+)
endolysin (CDS_0018)
CDS_0017
CDS_0019
IMGVR_UViG_3300045988_094182|3300045988|Ga0495776_160151 7220:21652(+)
VanY-A/VanY-F/VanY-M family D-Ala-D-Ala carboxypeptidase (NF000472.1) (CDS_0018)
CDS_0025
CDS_0026
CDS_0028
CDS_0017
CDS_0019
CDS_0020
CDS_0027
MGV-GENOME-0234943 7220:26386(+)
VanY-A/VanY-F/VanY-M family D-Ala-D-Ala carboxypeptidase (NF000472.1) (CDS_0018)
endolysin (CDS_0017)
19
20
21
22
23
24
25
MGV-GENOME-0267069 7220:31654(+)
endolysin (CDS_0017)
CDS_0018
DNA polymerase (CDS_0020)
uvig_254279 7226:21690(+)
endolysin (CDS_0017)
CDS_0015
CDS_0018
IMGVR_UViG_3300045988_131605|3300045988|Ga0495776_052206 7220:22207(+)
endolysin (CDS_0017)
CDS_0019
CDS_0018
CDS_0020
uvig_256783 7214:24045(+)
endolysin (CDS_0017)
CDS_0018
CDS_0023
CDS_0024
IMGVR_UViG_3300035655_000220|3300035655|Ga0134475_000395|7983-36048 7220:21663(+)
CDS_0010
CDS_0015
CDS_0017
MGV-GENOME-0215861 6604:20851(+)
CDS_0015
endolysin (CDS_0017)
IMGVR_UViG_3300029382_000618|3300029382|Ga0243901_1000179|35-26195 6610:18408(+)
multidrug efflux MATE transporter NorM (NF000263.1) (CDS_0018)
ThiF family (CDS_0019)
endolysin (CDS_0017)
20
21
uvig_383378 7220:23292(+)
endolysin (CDS_0017)
DNA helicase (CDS_0023)
18
19
20
21
22
24
25
26
27
28
IMGVR_UViG_3300045988_123882|3300045988|Ga0495776_165546 7220:28048(+)
tRNA-His guanylyltransferase (CDS_0026)
metallo-phosphoesterase (CDS_0023)
nucleotidyltransferase (CDS_0024)
VanY-A/VanY-F/VanY-M family D-Ala-D-Ala carboxypeptidase (NF000472.1) (CDS_0018)
RNA ligase (CDS_0027)
30
endolysin (CDS_0017)
19
20
21
22
phosphoesterase (CDS_0025)
28
29
31
IMGVR_UViG_3300045988_154157|3300045988|Ga0495776_157112 7220:33409(+)
CDS_0009
CDS_0014
CDS_0015
CDS_0016
IMGVR_UViG_3300045988_139343|3300045988|Ga0495776_165101 6017:18618(+)
CDS_0015
CDS_0016
IMGVR_UViG_3300029374_000340|3300029374|Ga0243945_1000093|97707-124230 6610:19419(+)
DNA polymerase (CDS_0017)
CDS_0021
CDS_0015
CDS_0024
MGV-GENOME-0183261 7220:19114(+)
CDS_0017
replisome organizer (CDS_0025)
IMGVR_UViG_3300045988_016807|3300045988|Ga0495776_045633 7220:20246(+)
10000 nt
DNA, RNA and nucleotide metabolism
lysis
other
tail
transcription regulation
unknown function
Defence
AMR