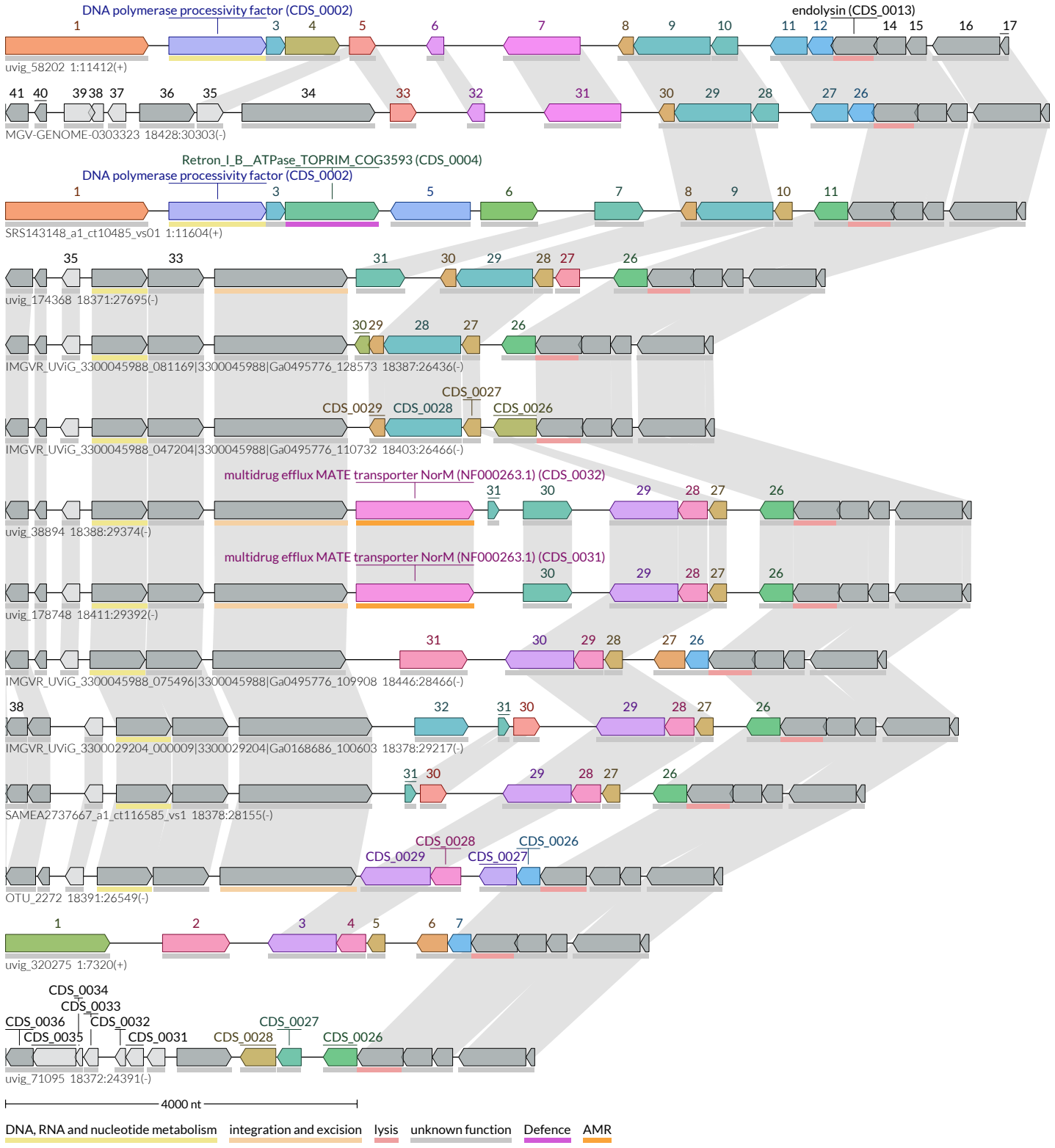
DNA polymerase processivity factor (CDS_0002)
endolysin (CDS_0013)
uvig_58202 1:11412(+)
MGV-GENOME-0303323 18428:30303(-)
Retron_I_B__ATPase_TOPRIM_COG3593 (CDS_0004)
DNA polymerase processivity factor (CDS_0002)
SRS143148_a1_ct10485_vs01 1:11604(+)
uvig_174368 18371:27695(-)
IMGVR_UViG_3300045988_081169|3300045988|Ga0495776_128573 18387:26436(-)
CDS_0027
CDS_0029
CDS_0028
CDS_0026
IMGVR_UViG_3300045988_047204|3300045988|Ga0495776_110732 18403:26466(-)
multidrug efflux MATE transporter NorM (NF000263.1) (CDS_0032)
uvig_38894 18388:29374(-)
multidrug efflux MATE transporter NorM (NF000263.1) (CDS_0031)
uvig_178748 18411:29392(-)
IMGVR_UViG_3300045988_075496|3300045988|Ga0495776_109908 18446:28466(-)
IMGVR_UViG_3300029204_000009|3300029204|Ga0168686_100603 18378:29217(-)
SAMEA2737667_a1_ct116585_vs1 18378:28155(-)
CDS_0028
CDS_0026
CDS_0029
CDS_0027
OTU_2272 18391:26549(-)
uvig_320275 1:7320(+)
CDS_0034
CDS_0033
CDS_0036
CDS_0032
CDS_0035
CDS_0031
CDS_0027
CDS_0028
CDS_0026
uvig_71095 18372:24391(-)
4000 nt
DNA, RNA and nucleotide metabolism
integration and excision
lysis
unknown function
Defence
AMR