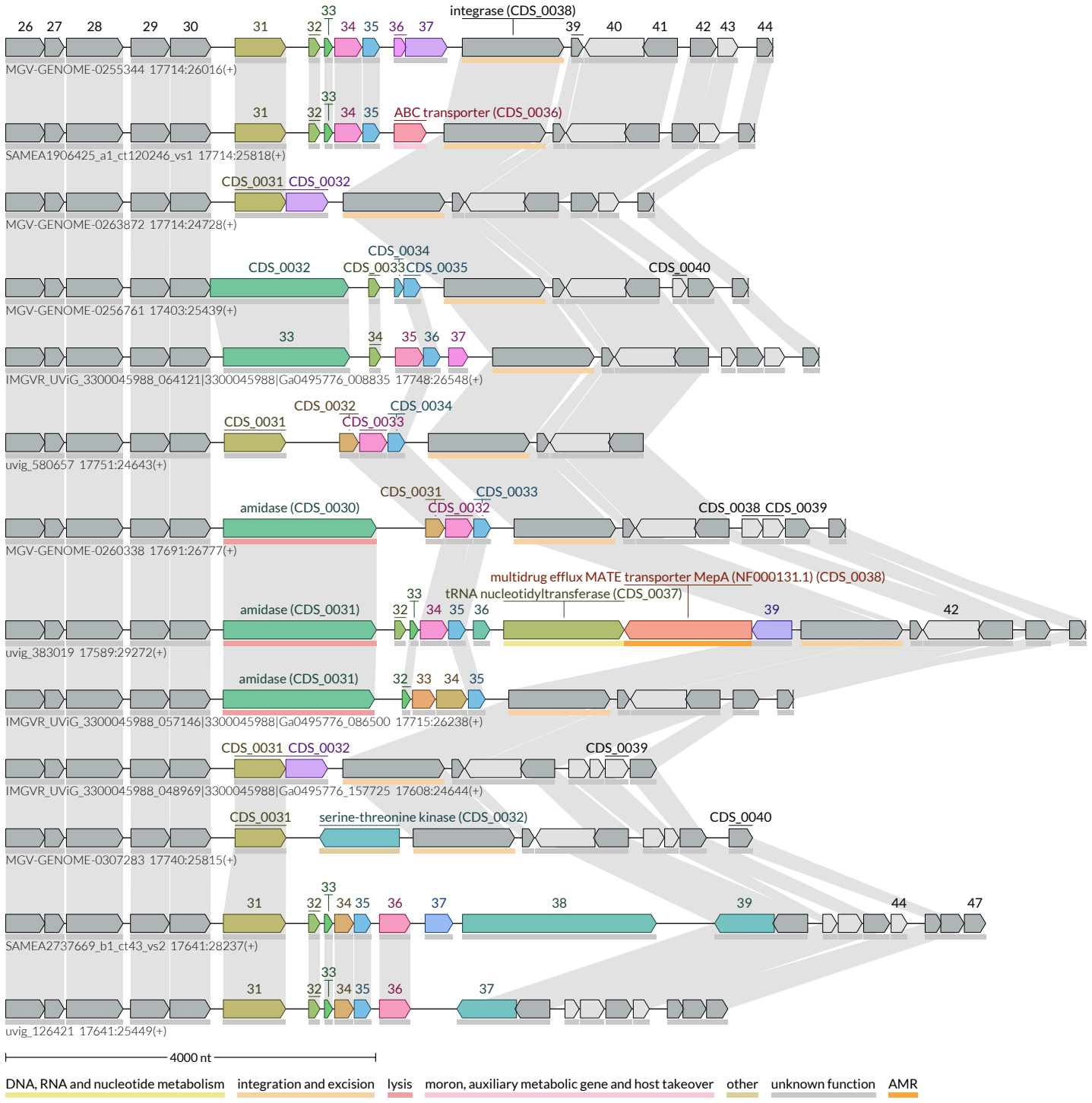
integrase (CDS_0038)
MGV-GENOME-0255344 17714:26016(+)
ABC transporter (CDS_0036)
SAMEA1906425_a1_ct120246_vs1 17714:25818(+)
CDS_0031 CDS_0032
MGV-GENOME-0263872 17714:24728(+)
CDS_0032 CDS_0033 CDS_0034 CDS_0035 CDS_0040
MGV-GENOME-0256761 17403:25439(+)
IMGVR_UViG_3300045988_064121|3300045988|Ga0495776_008835 17748:26548(+)
CDS_0031 CDS_0032 CDS_0033 CDS_0034
uvig_580657 17751:24643(+)
amidase (CDS_0030) CDS_0031 CDS_0032 CDS_0033 CDS_0038 CDS_0039
MGV-GENOME-0260338 17691:26777(+)
multidrug efflux MATE transporter MepA (NF000131.1) (CDS_0038)
tRNA nucleotidyltransferase (CDS_0037)
amidase (CDS_0031)
uvig_383019 17589:29272(+)
amidase (CDS_0031)
IMGVR_UViG_3300045988_057146|3300045988|Ga0495776_086500 17715:26238(+)
CDS_0031 CDS_0032 CDS_0039
IMGVR_UViG_3300045988_048969|3300045988|Ga0495776_157725 17608:24644(+)
CDS_0031 serine-threonine kinase (CDS_0032) CDS_0040
MGV-GENOME-0307283 17740:25815(+)
SAMEA2737669_b1_ct43_vs2 17641:28237(+)
uvig_126421 17641:25449(+)
4000 nt
DNA, RNA and nucleotide metabolism
integration and excision
lysis
moron, auxiliary metabolic gene and host takeover
other
unknown function
AMR