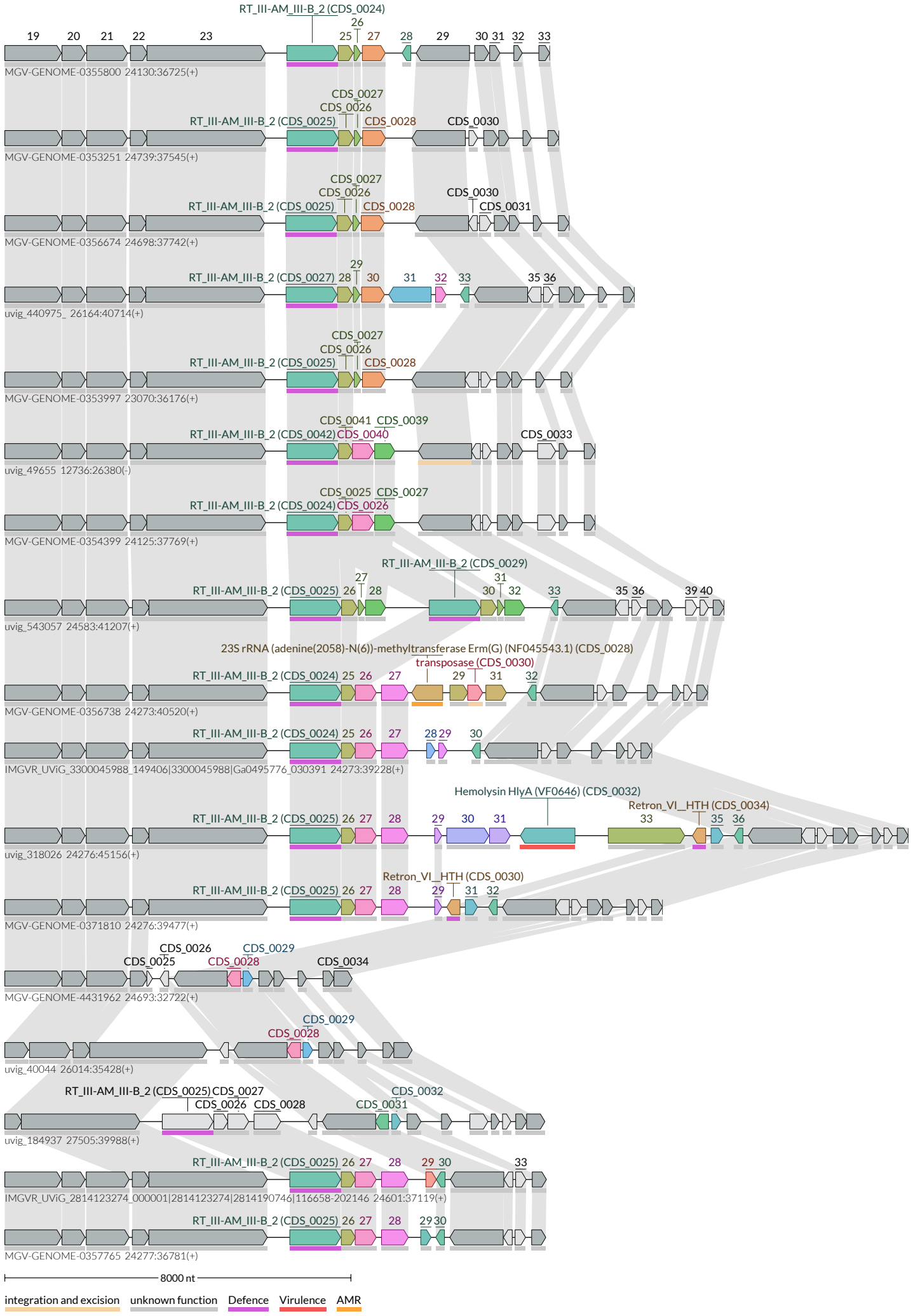
RT_III-AM_III-B_2 (CDS_0024)
19 20 21 22 23 25 26 27 28 29 30 31 32 33
MGV-GENOME-0355800 24130:36725(+)
CDS_0027
CDS_0026
RT_III-AM_III-B_2 (CDS_0025)
CDS_0028
CDS_0030
MGV-GENOME-0353251 24739:37545(+)
CDS_0027
CDS_0026
RT_III-AM_III-B_2 (CDS_0025)
CDS_0028
CDS_0030
CDS_0031
MGV-GENOME-0356674 24698:37742(+)
29
RT_III-AM_III-B_2 (CDS_0027) 28 30 31 32 33 35 36
uvig_440975_ 26164:40714(+)
CDS_0027
CDS_0026
RT_III-AM_III-B_2 (CDS_0025)
CDS_0028
MGV-GENOME-0353997 23070:36176(+)
CDS_0041 CDS_0039
RT_III-AM_III-B_2 (CDS_0042) CDS_0040
CDS_0033
uvig_49655 12736:26380(-)
CDS_0025 CDS_0027
RT_III-AM_III-B_2 (CDS_0024) CDS_0026
MGV-GENOME-0354399 24125:37769(+)
RT_III-AM_III-B_2 (CDS_0029)
27
31
RT_III-AM_III-B_2 (CDS_0025) 26 28 30 32 33 35 36 39 40
uvig_543057 24583:41207(+)
23S rRNA (adenine(2058)-N(6))-methyltransferase Erm(G) (NF045543.1) (CDS_0028)
transposase (CDS_0030)
RT_III-AM_III-B_2 (CDS_0024) 25 26 27 29 31 32
MGV-GENOME-0356738 24273:40520(+)
RT_III-AM_III-B_2 (CDS_0024) 25 26 27 28 29 30
IMGVR_UViG_3300045988_149406|3300045988|Ga0495776_030391 24273:39228(+)
Hemolysin HlyA (VF0646) (CDS_0032)
Retron_VI__HTH (CDS_0034)
RT_III-AM_III-B_2 (CDS_0025) 26 27 28 29 30 31 33 35 36
uvig_318026 24276:45156(+)
Retron_VI__HTH (CDS_0030)
RT_III-AM_III-B_2 (CDS_0025) 26 27 28 29 31 32
MGV-GENOME-0371810 24276:39477(+)
CDS_0026
CDS_0029
CDS_0025
CDS_0028
CDS_0034
MGV-GENOME-4431962 24693:32722(+)
CDS_0029
CDS_0028
uvig_40044 26014:35428(+)
RT_III-AM_III-B_2 (CDS_0025) CDS_0027
CDS_0032
CDS_0026 CDS_0028
CDS_0031
uvig_184937 27505:39988(+)
RT_III-AM_III-B_2 (CDS_0025) 26 27 28 29 30 33
IMGVR_UViG_2814123274_000001|2814123274|2814190746|116658-202146 24601:37119(+)
RT_III-AM_III-B_2 (CDS_0025) 26 27 28 29 30
MGV-GENOME-0357765 24277:36781(+)
8000 nt
integration and excision
unknown function
Defence
Virulence
AMR