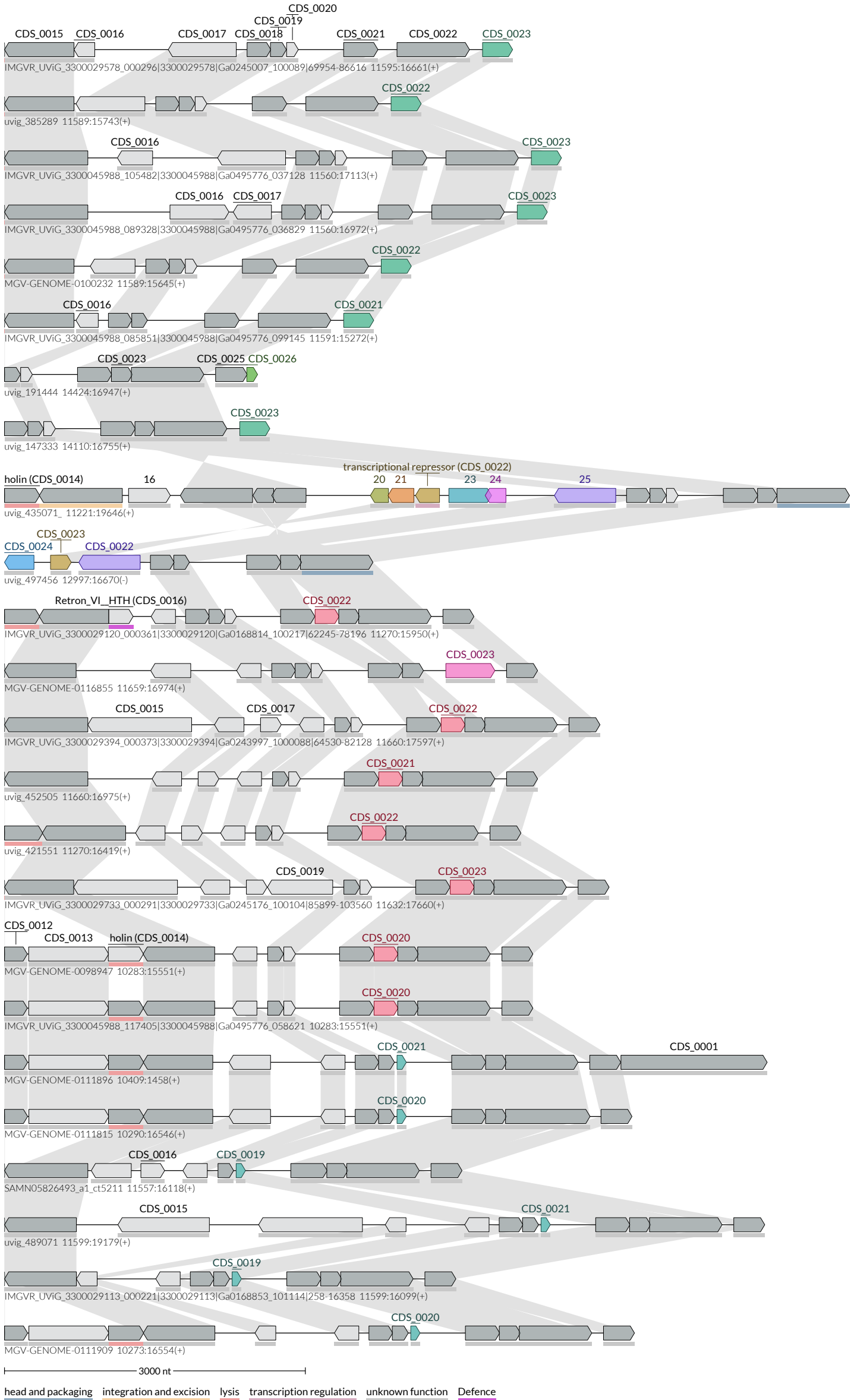
CDS_0020
CDS_0019
CDS_0015
CDS_0016
CDS_0017
CDS_0018
CDS_0021
CDS_0022
CDS_0023
IMGVR_UViG_3300029578_000296|3300029578|Ga0245007_100089|69954-86616 11595:16661(+)
CDS_0022
uvig_385289 11589:15743(+)
CDS_0016
CDS_0023
IMGVR_UViG_3300045988_105482|3300045988|Ga0495776_037128 11560:17113(+)
CDS_0016
CDS_0017
CDS_0023
IMGVR_UViG_3300045988_089328|3300045988|Ga0495776_036829 11560:16972(+)
CDS_0022
MGV-GENOME-0100232 11589:15645(+)
CDS_0016
CDS_0021
IMGVR_UViG_3300045988_085851|3300045988|Ga0495776_099145 11591:15272(+)
CDS_0023
CDS_0025
CDS_0026
uvig_191444 14424:16947(+)
CDS_0023
uvig_147333 14110:16755(+)
transcriptional repressor (CDS_0022)
holin (CDS_0014)
16
20
21
23
24
25
uvig_435071_ 11221:19646(+)
CDS_0023
CDS_0024
CDS_0022
uvig_497456 12997:16670(-)
Retron_VI__HTH (CDS_0016)
CDS_0022
IMGVR_UViG_3300029120_000361|3300029120|Ga0168814_100217|62245-78196 11270:15950(+)
CDS_0023
MGV-GENOME-0116855 11659:16974(+)
CDS_0015
CDS_0017
CDS_0022
IMGVR_UViG_3300029394_000373|3300029394|Ga0243997_1000088|64530-82128 11660:17597(+)
CDS_0021
uvig_452505 11660:16975(+)
CDS_0022
uvig_421551 11270:16419(+)
CDS_0019
CDS_0023
IMGVR_UViG_3300029733_000291|3300029733|Ga0245176_100104|85899-103560 11632:17660(+)
CDS_0012
CDS_0013
holin (CDS_0014)
CDS_0020
MGV-GENOME-0098947 10283:15551(+)
CDS_0020
IMGVR_UViG_3300045988_117405|3300045988|Ga0495776_058621 10283:15551(+)
CDS_0021
CDS_0001
MGV-GENOME-0111896 10409:1458(+)
CDS_0020
MGV-GENOME-0111815 10290:16546(+)
CDS_0016
CDS_0019
SAMN05826493_a1_ct5211 11557:16118(+)
CDS_0015
CDS_0021
uvig_489071 11599:19179(+)
CDS_0019
IMGVR_UViG_3300029113_000221|3300029113|Ga0168853_101114|258-16358 11599:16099(+)
CDS_0020
MGV-GENOME-0111909 10273:16554(+)
3000 nt
head and packaging
integration and excision
lysis
transcription regulation
unknown function
Defence