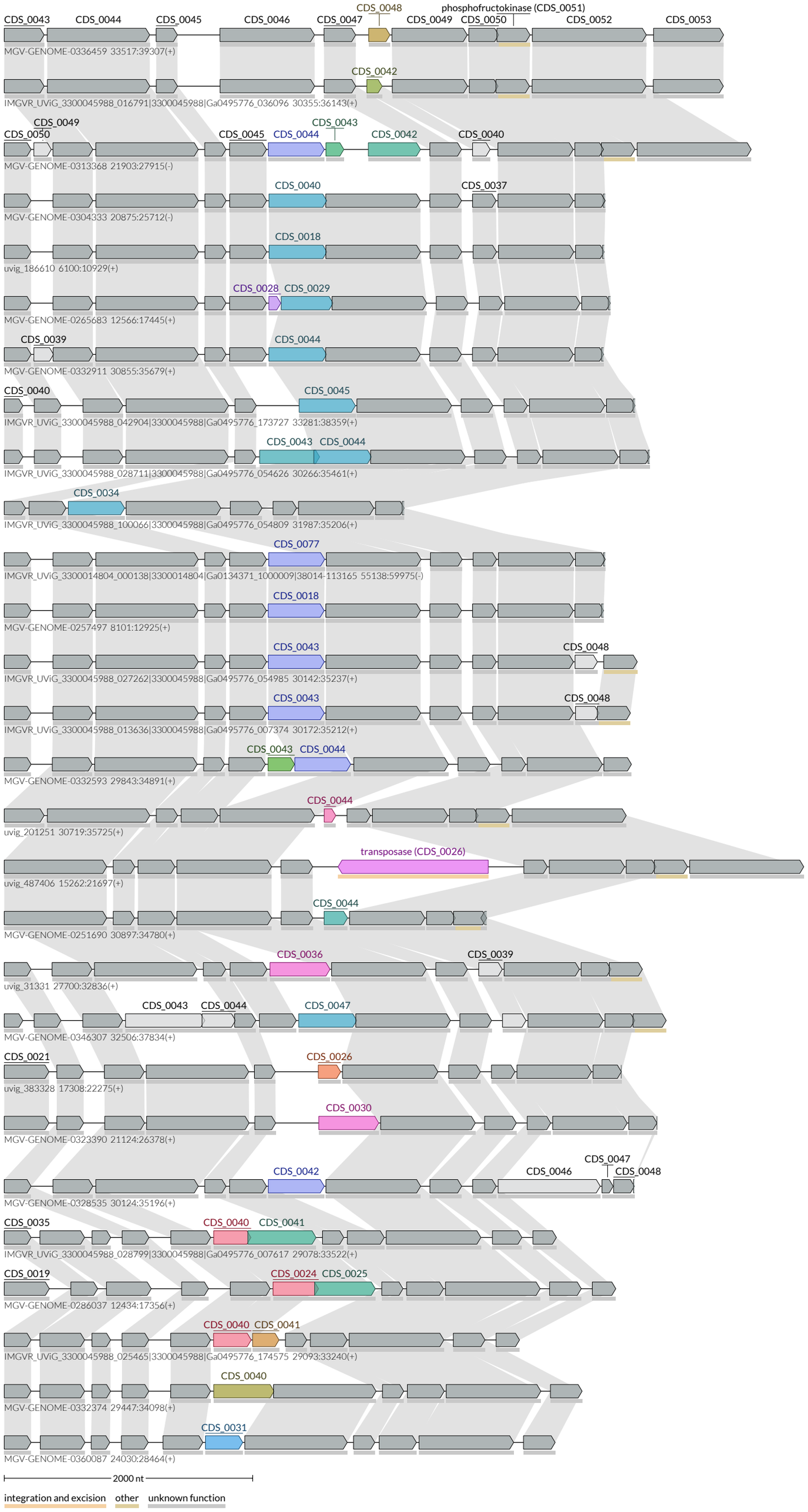
CDS_0043
CDS_0044
CDS_0045
CDS_0046
CDS_0047
CDS_0048
CDS_0049
phosphofructokinase (CDS_0051)
CDS_0050
CDS_0052
CDS_0053
MGV-GENOME-0336459 33517:39307(+)
CDS_0042
IMGVR_UViG_3300045988_016791|3300045988|Ga0495776_036096 30355:36143(+)
CDS_0049
CDS_0050
CDS_0043
CDS_0045
CDS_0044
CDS_0042
CDS_0040
MGV-GENOME-0313368 21903:27915(-)
CDS_0040
CDS_0037
MGV-GENOME-0304333 20875:25712(-)
CDS_0018
uvig_186610 6100:10929(+)
CDS_0028
CDS_0029
MGV-GENOME-0265683 12566:17445(+)
CDS_0039
CDS_0044
MGV-GENOME-0332911 30855:35679(+)
CDS_0040
CDS_0045
IMGVR_UViG_3300045988_042904|3300045988|Ga0495776_173727 33281:38359(+)
CDS_0043
CDS_0044
IMGVR_UViG_3300045988_028711|3300045988|Ga0495776_054626 30266:35461(+)
CDS_0034
IMGVR_UViG_3300045988_100066|3300045988|Ga0495776_054809 31987:35206(+)
CDS_0077
IMGVR_UViG_3300014804_000138|3300014804|Ga0134371_1000009|38014-113165 55138:59975(-)
CDS_0018
MGV-GENOME-0257497 8101:12925(+)
CDS_0043
CDS_0048
IMGVR_UViG_3300045988_027262|3300045988|Ga0495776_054985 30142:35237(+)
CDS_0043
CDS_0048
IMGVR_UViG_3300045988_013636|3300045988|Ga0495776_007374 30172:35212(+)
CDS_0043
CDS_0044
MGV-GENOME-0332593 29843:34891(+)
CDS_0044
uvig_201251 30719:35725(+)
transposase (CDS_0026)
uvig_487406 15262:21697(+)
CDS_0044
MGV-GENOME-0251690 30897:34780(+)
CDS_0036
CDS_0039
uvig_31331 27700:32836(+)
CDS_0043
CDS_0044
CDS_0047
MGV-GENOME-0346307 32506:37834(+)
CDS_0021
CDS_0026
uvig_383328 17308:22275(+)
CDS_0030
MGV-GENOME-0323390 21124:26378(+)
CDS_0047
CDS_0042
CDS_0046
CDS_0048
MGV-GENOME-0328535 30124:35196(+)
CDS_0035
CDS_0040
CDS_0041
IMGVR_UViG_3300045988_028799|3300045988|Ga0495776_007617 29078:33522(+)
CDS_0019
CDS_0024
CDS_0025
MGV-GENOME-0286037 12434:17356(+)
CDS_0040
CDS_0041
IMGVR_UViG_3300045988_025465|3300045988|Ga0495776_174575 29093:33240(+)
CDS_0040
MGV-GENOME-0332374 29447:34098(+)
CDS_0031
MGV-GENOME-0360087 24030:28464(+)
2000 nt
integration and excision
other
unknown function