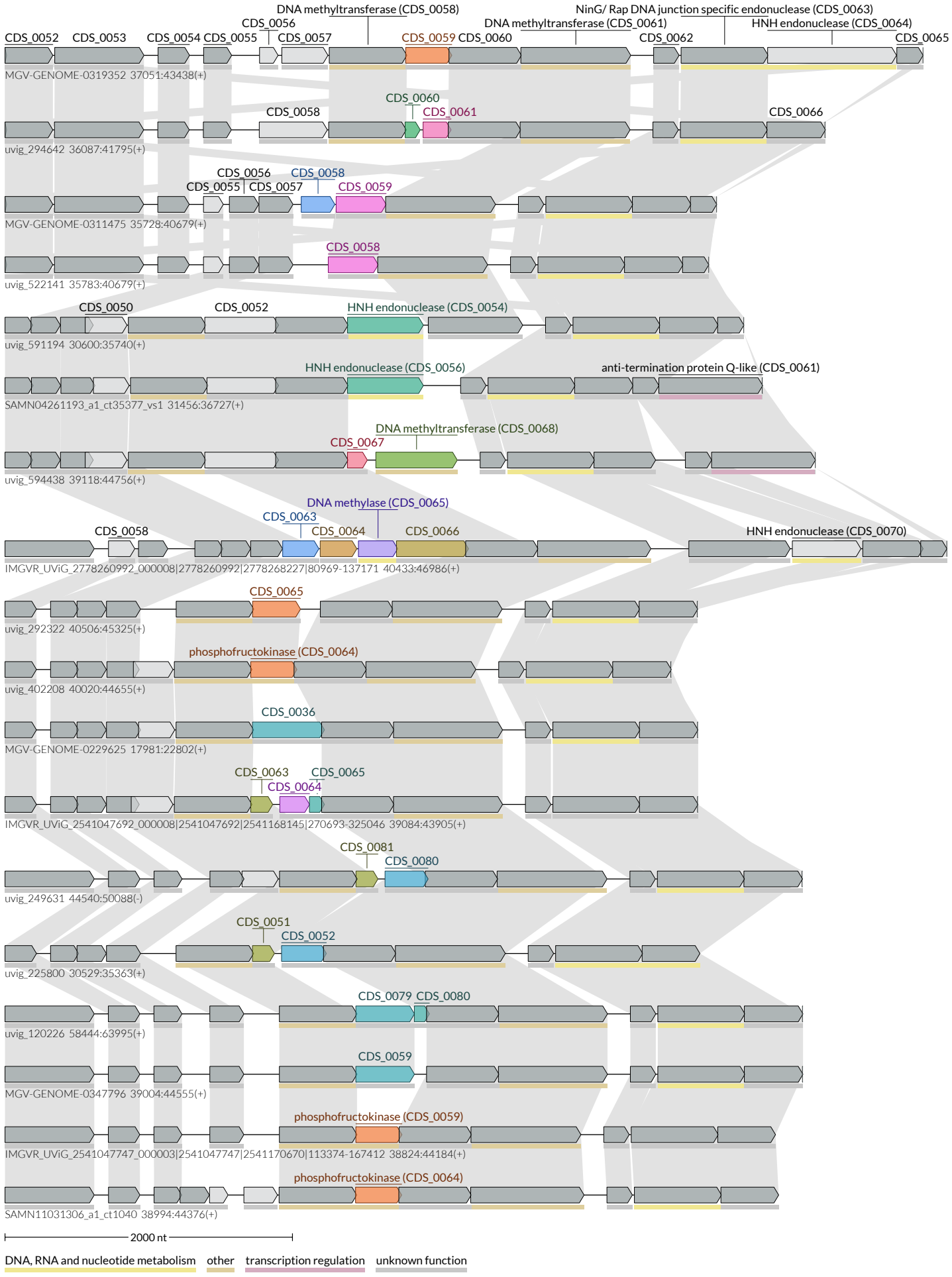
DNA methyltransferase (CDS_0058)
NinG/ Rap DNA junction specific endonuclease (CDS_0063)
CDS_0056
DNA methyltransferase (CDS_0061)
HNH endonuclease (CDS_0064)
CDS_0052
CDS_0053
CDS_0054
CDS_0055
CDS_0057
CDS_0059
CDS_0060
CDS_0062
CDS_0065
MGV-GENOME-0319352 37051:43438(+)
CDS_0060
CDS_0058
CDS_0061
CDS_0066
uvig_294642 36087:41795(+)
CDS_0056
CDS_0058
CDS_0055
CDS_0057
CDS_0059
MGV-GENOME-0311475 35728:40679(+)
CDS_0058
uvig_522141 35783:40679(+)
CDS_0050
CDS_0052
HNH endonuclease (CDS_0054)
uvig_591194 30600:35740(+)
HNH endonuclease (CDS_0056)
anti-termination protein Q-like (CDS_0061)
SAMN04261193_a1_ct35377_vs1 31456:36727(+)
DNA methyltransferase (CDS_0068)
CDS_0067
uvig_594438 39118:44756(+)
DNA methylase (CDS_0065)
CDS_0063
CDS_0058
CDS_0064
CDS_0066
HNH endonuclease (CDS_0070)
IMGVR_UViG_2778260992_000008|2778260992|2778268227|80969-137171 40433:46986(+)
CDS_0065
uvig_292322 40506:45325(+)
phosphofructokinase (CDS_0064)
uvig_402208 40020:44655(+)
CDS_0036
MGV-GENOME-0229625 17981:22802(+)
CDS_0063
CDS_0065
CDS_0064
IMGVR_UViG_2541047692_000008|2541047692|2541168145|270693-325046 39084:43905(+)
CDS_0081
CDS_0080
uvig_249631 44540:50088(-)
CDS_0051
CDS_0052
uvig_225800 30529:35363(+)
CDS_0079
CDS_0080
uvig_120226 58444:63995(+)
CDS_0059
MGV-GENOME-0347796 39004:44555(+)
phosphofructokinase (CDS_0059)
IMGVR_UViG_2541047747_000003|2541047747|2541170670|113374-167412 38824:44184(+)
phosphofructokinase (CDS_0064)
SAMN11031306_a1_ct1040 38994:44376(+)
2000 nt
DNA, RNA and nucleotide metabolism
other
transcription regulation
unknown function