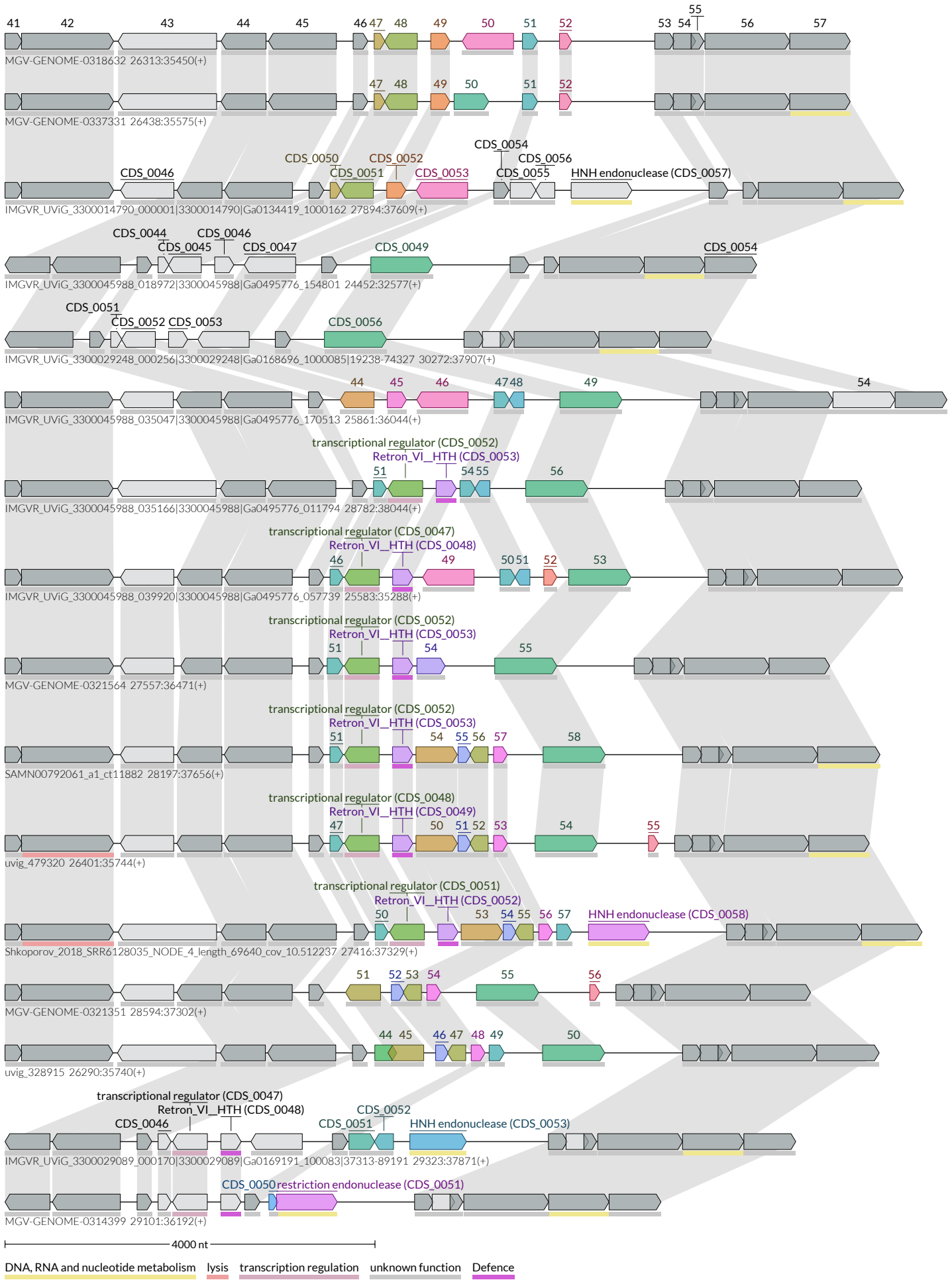
MGV-GENOME-0318632 26313:35450(+)
MGV-GENOME-0337331 26438:35575(+)
CDS_0046
CDS_0050
CDS_0051
CDS_0052
CDS_0053
CDS_0054
CDS_0055
CDS_0056
HNH endonuclease (CDS_0057)
IMGVR_UViG_3300014790_000001|3300014790|Ga0134419_1000162 27894:37609(+)
CDS_0044
CDS_0045
CDS_0046
CDS_0047
CDS_0049
CDS_0054
IMGVR_UViG_3300045988_018972|3300045988|Ga0495776_154801 24452:32577(+)
CDS_0051
CDS_0052 CDS_0053
CDS_0056
IMGVR_UViG_3300029248_000256|3300029248|Ga0168696_1000085|19238-74327 30272:37907(+)
IMGVR_UViG_3300045988_035047|3300045988|Ga0495776_170513 25861:36044(+)
transcriptional regulator (CDS_0052)
Retron_VI_HTH (CDS_0053)
IMGVR_UViG_3300045988_035166|3300045988|Ga0495776_011794 28782:38044(+)
transcriptional regulator (CDS_0047)
Retron_VI_HTH (CDS_0048)
IMGVR_UViG_3300045988_039920|3300045988|Ga0495776_057739 25583:35288(+)
transcriptional regulator (CDS_0052)
Retron_VI_HTH (CDS_0053)
MGV-GENOME-0321564 27557:36471(+)
transcriptional regulator (CDS_0052)
Retron_VI_HTH (CDS_0053)
SAMN00792061_a1_ct11882 28197:37656(+)
transcriptional regulator (CDS_0048)
Retron_VI_HTH (CDS_0049)
uvig_479320 26401:35744(+)
transcriptional regulator (CDS_0051)
Retron_VI_HTH (CDS_0052)
HNH endonuclease (CDS_0058)
Shkoporov_2018_SRR6128035_NODE_4_length_69640_cov_10.512237 27416:37329(+)
MGV-GENOME-0321351 28594:37302(+)
uvig_328915 26290:35740(+)
transcriptional regulator (CDS_0047)
Retron_VI_HTH (CDS_0048)
CDS_0046
CDS_0052
CDS_0051
HNH endonuclease (CDS_0053)
IMGVR_UViG_3300029089_000170|3300029089|Ga0169191_100083|37313-89191 29323:37871(+)
CDS_0050 restriction endonuclease (CDS_0051)
MGV-GENOME-0314399 29101:36192(+)
4000 nt
DNA, RNA and nucleotide metabolism
lysis
transcription regulation
unknown function
Defence