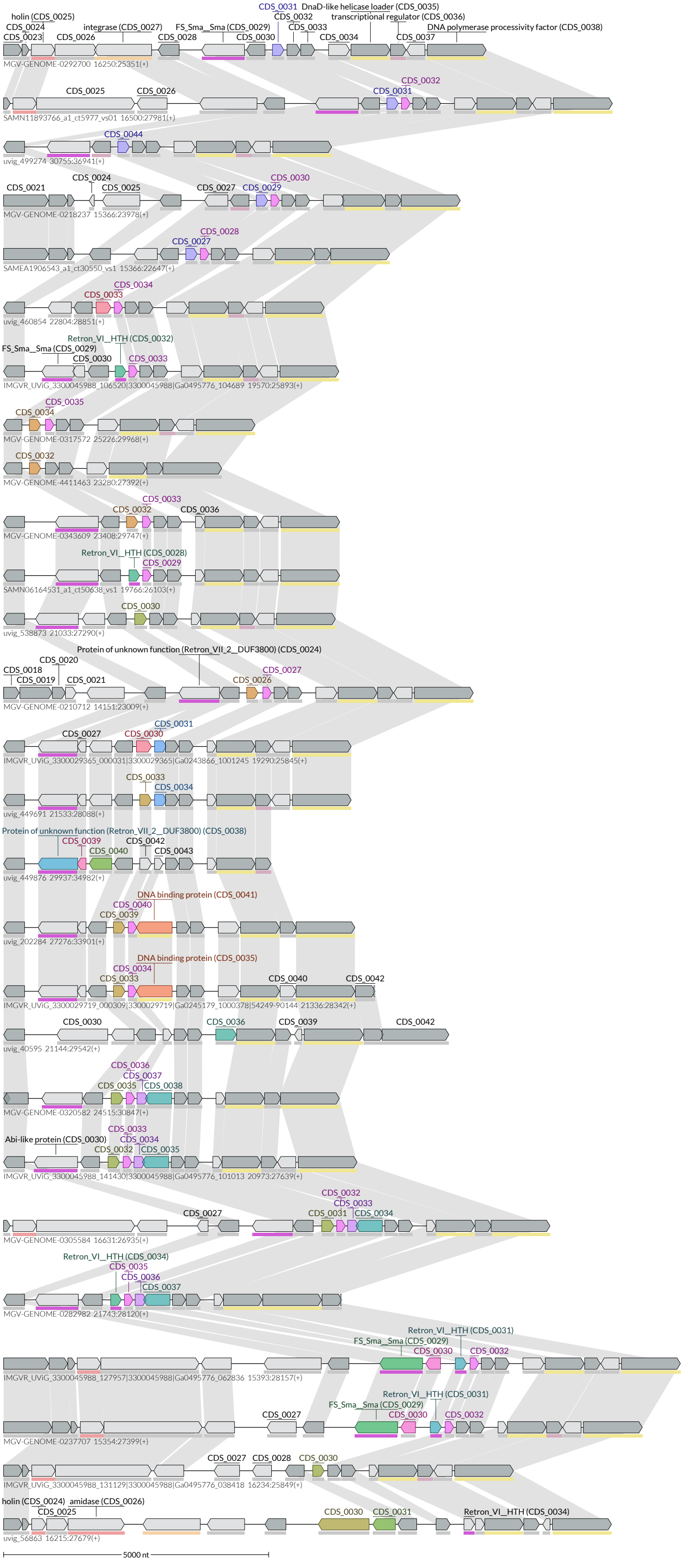
holin (CDS_0025)
CDS_0024
CDS_0023
integrase (CDS_0027)
CDS_0026
FS_Sma__Sma (CDS_0029)
CDS_0028
CDS_0030
CDS_0031
DnaD-like helicase loader (CDS_0035)
CDS_0032
transcriptional regulator (CDS_0036)
CDS_0033
CDS_0034
DNA polymerase processivity factor (CDS_0038)
CDS_0037
MGV-GENOME-0292700 16250:25351(+)
CDS_0025
CDS_0026
CDS_0032
CDS_0031
SAMN11893766_a1_ct5977_vs01 16500:27981(+)
CDS_0044
uvig_499274 30755:36941(+)
CDS_0024
CDS_0030
CDS_0021
CDS_0025
CDS_0027
CDS_0029
MGV-GENOME-0218237 15366:23978(+)
CDS_0028
CDS_0027
SAMEA1906543_a1_ct30550_vs1 15366:22647(+)
CDS_0034
CDS_0033
uvig_460854 22804:28851(+)
Retron_VI__HTH (CDS_0032)
FS_Sma__Sma (CDS_0029)
CDS_0030
CDS_0033
IMGVR_UViG_3300045988_106520|3300045988|Ga0495776_104689 19570:25893(+)
CDS_0035
CDS_0034
MGV-GENOME-0317572 25226:29968(+)
CDS_0032
MGV-GENOME-4411463 23280:27392(+)
CDS_0033
CDS_0032
CDS_0036
MGV-GENOME-0343609 23408:29747(+)
Retron_VI__HTH (CDS_0028)
CDS_0029
SAMN06164531_a1_ct50638_vs1 19766:26103(+)
CDS_0030
uvig_538873 21033:27290(+)
Protein of unknown function (Retron_VII_2__DUF3800) (CDS_0024)
CDS_0020
CDS_0018
CDS_0019
CDS_0021
CDS_0027
CDS_0026
MGV-GENOME-0210712 14151:23009(+)
CDS_0031
CDS_0027
CDS_0030
IMGVR_UViG_3300029365_000031|3300029365|Ga0243866_1001245 19290:25845(+)
CDS_0033
CDS_0034
uvig_449691 21533:28088(+)
Protein of unknown function (Retron_VII_2__DUF3800) (CDS_0038)
CDS_0039
CDS_0042
CDS_0040
CDS_0043
uvig_449876 29937:34982(+)
DNA binding protein (CDS_0041)
CDS_0040
CDS_0039
uvig_202284 27276:33901(+)
DNA binding protein (CDS_0035)
CDS_0034
CDS_0033
CDS_0040
CDS_0042
IMGVR_UViG_3300029719_000309|3300029719|Ga0245179_1000378|54249-90144 21336:28342(+)
CDS_0030
CDS_0036
CDS_0039
CDS_0042
uvig_40595 21144:29542(+)
CDS_0036
CDS_0037
CDS_0035
CDS_0038
MGV-GENOME-0320582 24515:30847(+)
CDS_0033
Abi-like protein (CDS_0030)
CDS_0034
CDS_0032
CDS_0035
IMGVR_UViG_3300045988_141430|3300045988|Ga0495776_101013 20973:27639(+)
CDS_0032
CDS_0033
CDS_0027
CDS_0031
CDS_0034
MGV-GENOME-0305584 16631:26935(+)
Retron_VI__HTH (CDS_0034)
CDS_0035
CDS_0036
CDS_0037
MGV-GENOME-0282982 21743:28120(+)
Retron_VI__HTH (CDS_0031)
FS_Sma__Sma (CDS_0029)
CDS_0030
CDS_0032
IMGVR_UViG_3300045988_127957|3300045988|Ga0495776_062836 15393:28157(+)
Retron_VI__HTH (CDS_0031)
FS_Sma__Sma (CDS_0029)
CDS_0027
CDS_0030
CDS_0032
MGV-GENOME-0237707 15354:27399(+)
CDS_0027
CDS_0028
CDS_0030
IMGVR_UViG_3300045988_131129|3300045988|Ga0495776_038418 16234:25849(+)
holin (CDS_0024)
amidase (CDS_0026)
CDS_0025
CDS_0030
CDS_0031
Retron_VI__HTH (CDS_0034)
uvig_56863 16215:27679(+)
5000 nt
DNA, RNA and nucleotide metabolism
integration and excision
lysis
transcription regulation
unknown function
Defence