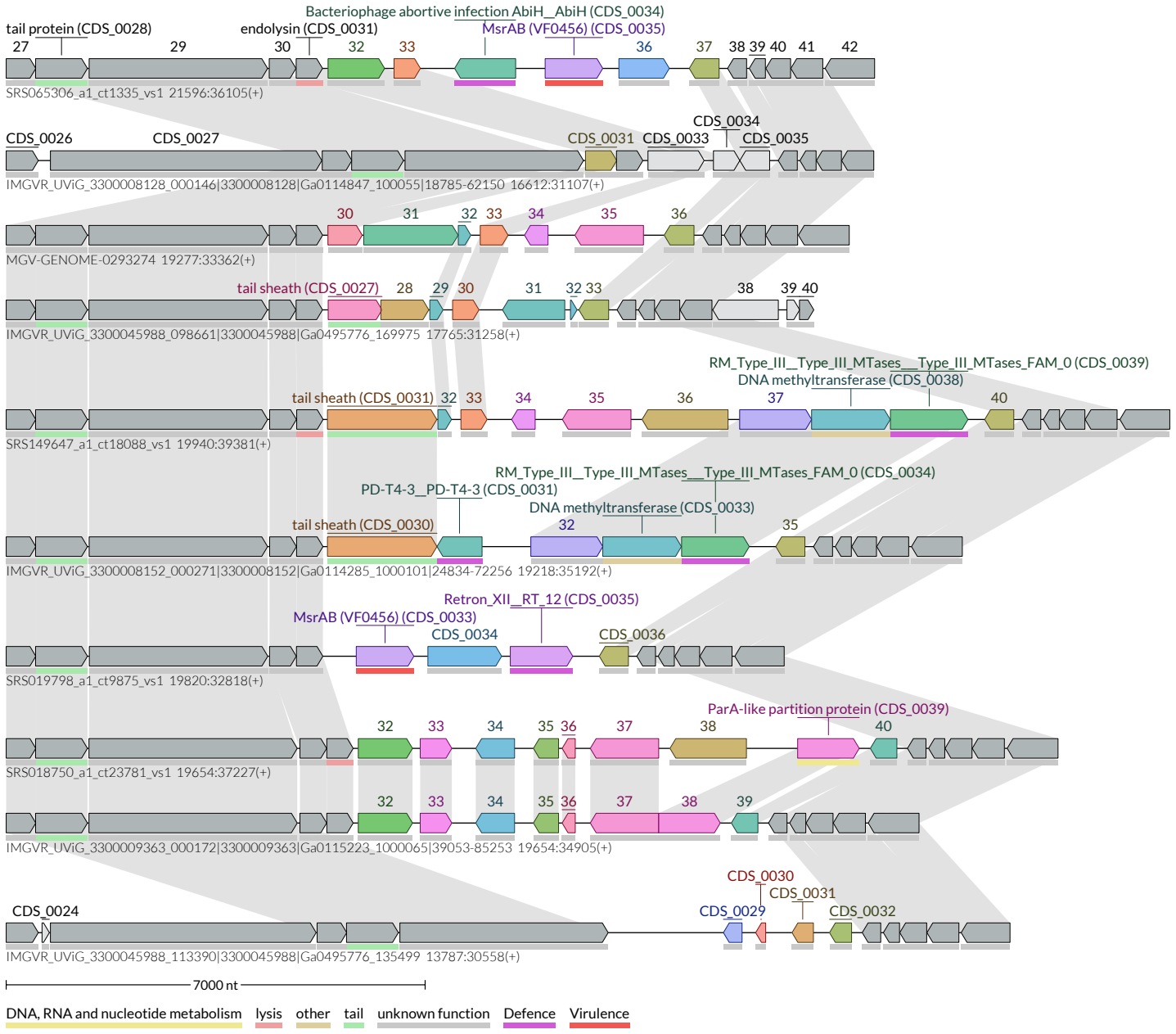
Bacteriophage abortive infection AbiH__AbiH (CDS_0034)
tail protein (CDS_0028)
endolysin (CDS_0031)
MsrAB (VF0456) (CDS_0035)
27 29 30 32 33 36 37 38 39 40 41 42
SRS065306_a1_ct1335_vs1 21596:36105(+)
CDS_0034
CDS_0026 CDS_0027 CDS_0031 CDS_0033 CDS_0035
IMGVR_UViG_3300008128_000146|3300008128|Ga0114847_100055|18785-62150 16612:31107(+)
30 31 32 33 34 35 36
MGV-GENOME-0293274 19277:33362(+)
tail sheath (CDS_0027)
28 29 30 31 32 33 38 39 40
IMGVR_UViG_3300045988_098661|3300045988|Ga0495776_169975 17765:31258(+)
RM_Type_III__Type_III_MTases___Type_III_MTases_FAM_0 (CDS_0039)
DNA methyltransferase (CDS_0038)
tail sheath (CDS_0031)
32 33 34 35 36 37 40
SRS149647_a1_ct18088_vs1 19940:39381(+)
RM_Type_III__Type_III_MTases___Type_III_MTases_FAM_0 (CDS_0034)
PD-T4-3__PD-T4-3 (CDS_0031)
DNA methyltransferase (CDS_0033)
tail sheath (CDS_0030)
32 35
IMGVR_UViG_3300008152_000271|3300008152|Ga0114285_1000101|24834-72256 19218:35192(+)
Retron_XII__RT_12 (CDS_0035)
MsrAB (VF0456) (CDS_0033)
CDS_0034 CDS_0036
SRS019798_a1_ct9875_vs1 19820:32818(+)
ParA-like partition protein (CDS_0039)
32 33 34 35 36 37 38 40
SRS018750_a1_ct23781_vs1 19654:37227(+)
32 33 34 35 36 37 38 39
IMGVR_UViG_3300009363_000172|3300009363|Ga0115223_1000065|39053-85253 19654:34905(+)
CDS_0030
CDS_0031
CDS_0024 CDS_0029 CDS_0032
IMGVR_UViG_3300045988_113390|3300045988|Ga0495776_135499 13787:30558(+)
7000 nt
DNA, RNA and nucleotide metabolism
lysis
other
tail
unknown function
Defence
Virulence