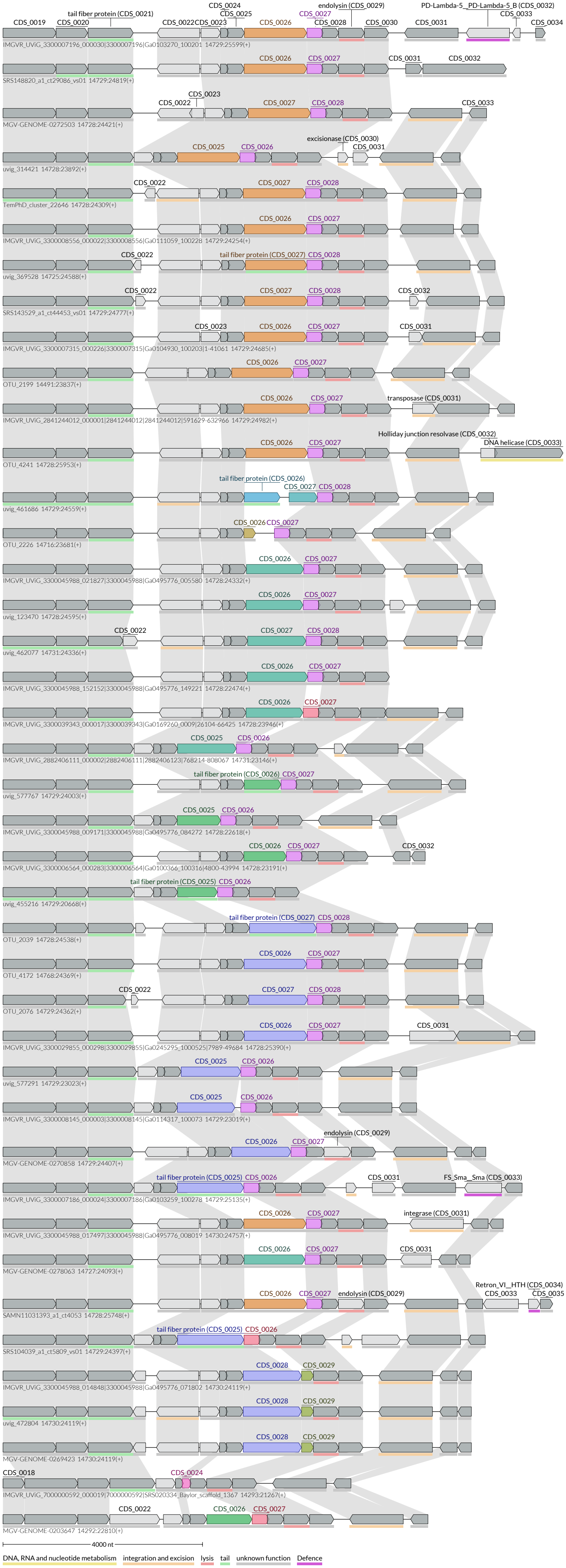
CDS_0019
tail fiber protein (CDS_0021)
CDS_0020
CDS_0022
CDS_0024
CDS_0023
CDS_0025
CDS_0026
CDS_0027
endolysin (CDS_0029)
CDS_0028
CDS_0030
CDS_0031
PD-Lambda-5__PD-Lambda-5_B (CDS_0032)
CDS_0033
CDS_0034
IMGVR_UViG_3300007196_000030|3300007196|Ga0103270_100201 14729:25599(+)
SRS148820_a1_ct29086_vs01 14729:24819(+)
MGV-GENOME-0272503 14728:24421(+)
excisionase (CDS_0030)
uvig_314421 14728:23892(+)
TemPhD_cluster_22646 14728:24309(+)
IMGVR_UViG_3300008556_000022|3300008556|Ga0111059_100228 14729:24254(+)
tail fiber protein (CDS_0027)
uvig_369528 14725:24588(+)
SRS143529_a1_ct44453_vs01 14729:24777(+)
IMGVR_UViG_3300007315_000226|3300007315|Ga0104930_100203|1-41061 14729:24685(+)
OTU_2199 14491:23837(+)
transposase (CDS_0031)
IMGVR_UViG_2841244012_000001|2841244012|2841244012|591629-632966 14729:24982(+)
Holliday junction resolvase (CDS_0032)
DNA helicase (CDS_0033)
OTU_4241 14728:25953(+)
tail fiber protein (CDS_0026)
uvig_461686 14729:24559(+)
OTU_2226 14716:23681(+)
IMGVR_UViG_3300045988_021827|3300045988|Ga0495776_005580 14728:24332(+)
uvig_123470 14728:24595(+)
uvig_462077 14731:24336(+)
IMGVR_UViG_3300045988_152152|3300045988|Ga0495776_149221 14728:22474(+)
IMGVR_UViG_3300039343_000017|3300039343|Ga0169260_0009|26104-66425 14728:23946(+)
IMGVR_UViG_2882406111_000002|2882406111|2882406123|768214-808067 14731:23146(+)
tail fiber protein (CDS_0026) CDS_0027
uvig_577767 14729:24003(+)
IMGVR_UViG_3300045988_009171|3300045988|Ga0495776_084272 14728:22618(+)
IMGVR_UViG_3300006564_000283|3300006564|Ga0100366_100316|4800-43994 14728:23191(+)
tail fiber protein (CDS_0025) CDS_0026
uvig_455216 14729:20668(+)
tail fiber protein (CDS_0027) CDS_0028
OTU_2039 14728:24538(+)
OTU_4172 14768:24369(+)
OTU_2076 14729:24362(+)
IMGVR_UViG_3300029855_000298|3300029855|Ga0245295_1000525|7989-49684 14728:25390(+)
uvig_577291 14729:23023(+)
IMGVR_UViG_3300008145_000003|3300008145|Ga0114317_100073 14729:23019(+)
MGV-GENOME-0270858 14729:24407(+)
FS_Sma__Sma (CDS_0033)
IMGVR_UViG_3300007186_000024|3300007186|Ga0103259_100278 14729:25135(+)
integrase (CDS_0031)
IMGVR_UViG_3300045988_017497|3300045988|Ga0495776_008019 14730:24757(+)
MGV-GENOME-0278063 14727:24093(+)
Retron_VI__HTH (CDS_0034)
CDS_0035
SAMN11031393_a1_ct4053 14728:25748(+)
SRS104039_a1_ct5809_vs01 14729:24397(+)
CDS_0029
IMGVR_UViG_3300045988_014848|3300045988|Ga0495776_071802 14730:24119(+)
uvig_472804 14730:24119(+)
MGV-GENOME-0269423 14730:24119(+)
IMGVR_UViG_7000000592_000019|7000000592|SRS020334_Baylor_scaffold_1367 14293:21267(+)
MGV-GENOME-0203647 14292:22810(+)
4000 nt
DNA, RNA and nucleotide metabolism
integration and excision
lysis
tail
unknown function
Defence