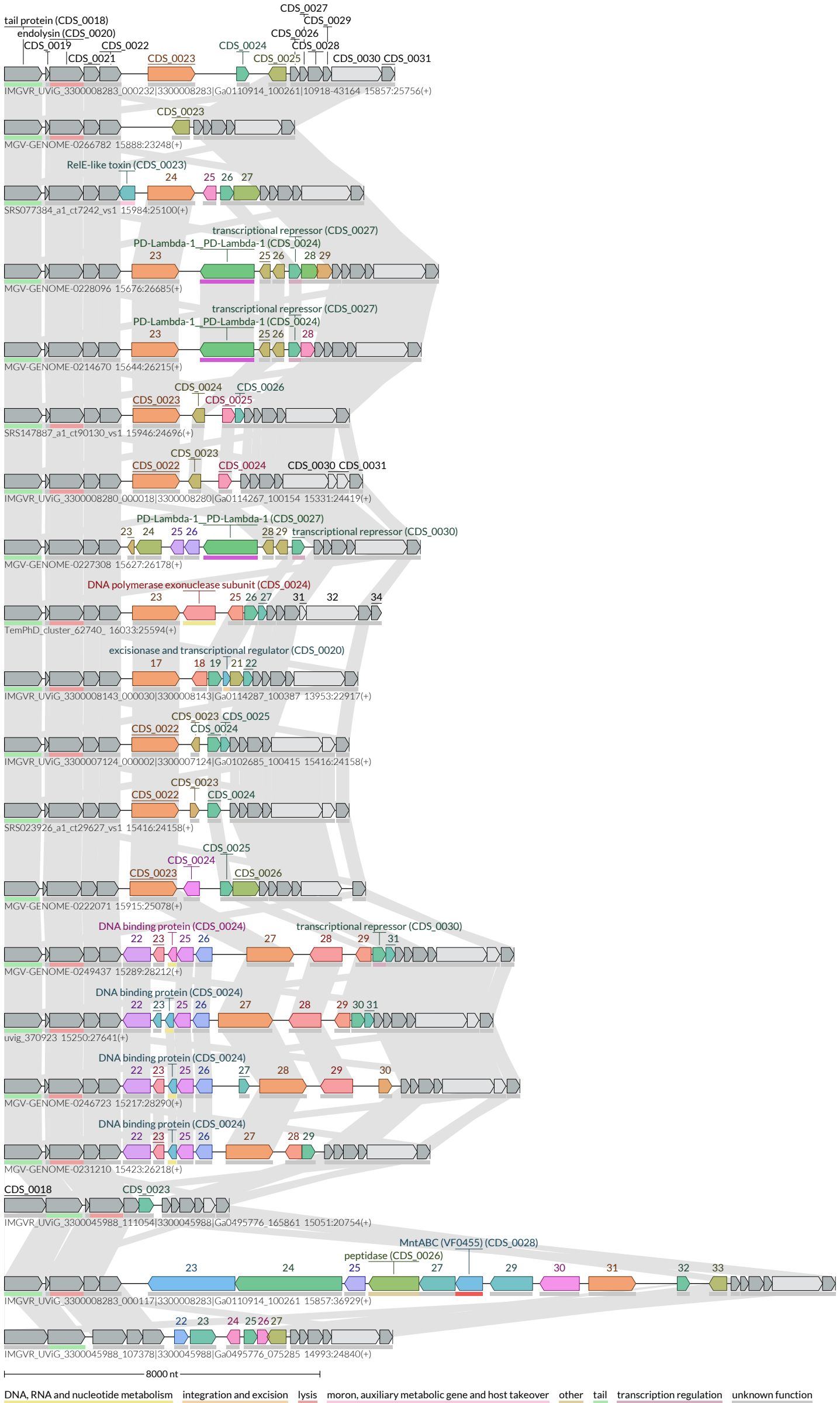
tail protein (CDS_0018)
endolysin (CDS_0020)
CDS_0019
CDS_0022
CDS_0021
CDS_0023
CDS_0024
CDS_0025
CDS_0026
CDS_0027
CDS_0028
CDS_0029
CDS_0030 CDS_0031
IMGVR_UViG_3300008283_000232|3300008283|Ga0110914_100261|10918-43164 15857:25756(+)
CDS_0023
MGV-GENOME-0266782 15888:23248(+)
RelE-like toxin (CDS_0023)
24
25 26 27
SRS077384_a1_ct7242_vs1 15984:25100(+)
transcriptional repressor (CDS_0027)
PD-Lambda-1_PD-Lambda-1 (CDS_0024)
23
25 26
28 29
MGV-GENOME-0228096 15676:26685(+)
transcriptional repressor (CDS_0027)
PD-Lambda-1_PD-Lambda-1 (CDS_0024)
23
25 26
28
MGV-GENOME-0214670 15644:26215(+)
CDS_0024
CDS_0026
CDS_0023
CDS_0025
SRS147887_a1_ct90130_vs1 15946:24696(+)
CDS_0023
CDS_0022
CDS_0024
CDS_0030 CDS_0031
IMGVR_UViG_3300008280_000018|3300008280|Ga0114267_100154 15331:24419(+)
PD-Lambda-1_PD-Lambda-1 (CDS_0027)
23 24
25 26
28 29
transcriptional repressor (CDS_0030)
MGV-GENOME-0227308 15627:26178(+)
DNA polymerase exonuclease subunit (CDS_0024)
23
25 26 27
31
32
34
TemPhD_cluster_62740_ 16033:25594(+)
excisionase and transcriptional regulator (CDS_0020)
17
18 19
21 22
IMGVR_UViG_3300008143_000030|3300008143|Ga0114287_100387 13953:22917(+)
CDS_0023 CDS_0025
CDS_0022
CDS_0024
IMGVR_UViG_3300007124_000002|3300007124|Ga0102685_100415 15416:24158(+)
CDS_0023
CDS_0022
CDS_0024
SRS023926_a1_ct29627_vs1 15416:24158(+)
CDS_0025
CDS_0024
CDS_0023
CDS_0026
MGV-GENOME-0222071 15915:25078(+)
DNA binding protein (CDS_0024)
transcriptional repressor (CDS_0030)
22 23 25 26
27
28
29 31
MGV-GENOME-0249437 15289:28212(+)
DNA binding protein (CDS_0024)
22 23 25 26
27
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29 30 31
uvig_370923 15250:27641(+)
DNA binding protein (CDS_0024)
22 23 25 26
27
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29
30
MGV-GENOME-0246723 15217:28290(+)
DNA binding protein (CDS_0024)
22 23 25 26
27
28 29
MGV-GENOME-0231210 15423:26218(+)
CDS_0018
CDS_0023
IMGVR_UViG_3300045988_111054|3300045988|Ga0495776_165861 15051:20754(+)
MntABC (VF0455) (CDS_0028)
peptidase (CDS_0026)
23
24
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33
IMGVR_UViG_3300008283_000117|3300008283|Ga0110914_100261 15857:36929(+)
22 23
24 25 26 27
IMGVR_UViG_3300045988_107378|3300045988|Ga0495776_075285 14993:24840(+)
8000 nt
DNA, RNA and nucleotide metabolism
integration and excision
lysis
moron, auxiliary metabolic gene and host takeover
other
tail
transcription regulation
unknown function
Defence
Virulence