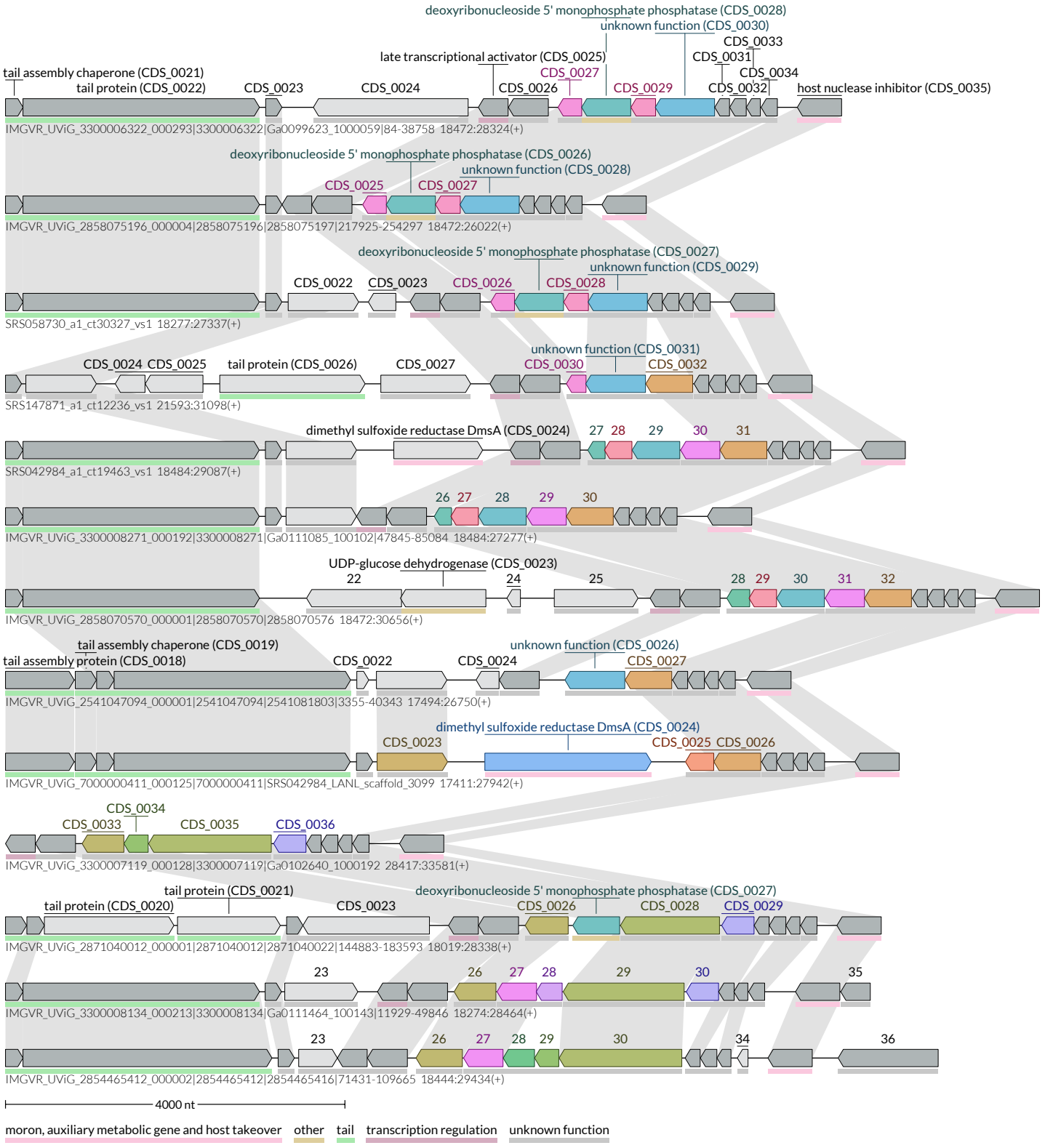
deoxyribonucleoside 5' monophosphate phosphatase (CDS_0028)
unknown function (CDS_0030)
CDS_0033
late transcriptional activator (CDS_0025)
CDS_0031
CDS_0027
CDS_0034
tail assembly chaperone (CDS_0021)
tail protein (CDS_0022)
CDS_0023
CDS_0024
CDS_0026
CDS_0029
CDS_0032
host nuclease inhibitor (CDS_0035)
IMGVR_UViG_3300006322_000293|3300006322|Ga0099623_1000059|84-38758 18472:28324(+)
deoxyribonucleoside 5' monophosphate phosphatase (CDS_0026)
unknown function (CDS_0028)
CDS_0025
CDS_0027
IMGVR_UViG_2858075196_000004|2858075196|2858075197|217925-254297 18472:26022(+)
deoxyribonucleoside 5' monophosphate phosphatase (CDS_0027)
unknown function (CDS_0029)
CDS_0022
CDS_0023
CDS_0026
CDS_0028
SRS058730_a1_ct30327_vs1 18277:27337(+)
unknown function (CDS_0031)
CDS_0024
CDS_0025
tail protein (CDS_0026)
CDS_0027
CDS_0030
CDS_0032
SRS147871_a1_ct12236_vs1 21593:31098(+)
dimethyl sulfoxide reductase DmsA (CDS_0024)
27
28
29
30
31
SRS042984_a1_ct19463_vs1 18484:29087(+)
26
27
28
29
30
IMGVR_UViG_3300008271_000192|3300008271|Ga0111085_100102|47845-85084 18484:27277(+)
UDP-glucose dehydrogenase (CDS_0023)
22
24
25
28
29
30
31
32
IMGVR_UViG_2858070570_000001|2858070570|2858070576 18472:30656(+)
tail assembly chaperone (CDS_0019)
unknown function (CDS_0026)
tail assembly protein (CDS_0018)
CDS_0022
CDS_0024
CDS_0027
IMGVR_UViG_2541047094_000001|2541047094|2541081803|3355-40343 17494:26750(+)
dimethyl sulfoxide reductase DmsA (CDS_0024)
CDS_0023
CDS_0025
CDS_0026
IMGVR_UViG_7000000411_000125|7000000411|SRS042984_LANL_scaffold_3099 17411:27942(+)
CDS_0034
CDS_0033
CDS_0035
CDS_0036
IMGVR_UViG_3300007119_000128|3300007119|Ga0102640_1000192 28417:33581(+)
tail protein (CDS_0021)
deoxyribonucleoside 5' monophosphate phosphatase (CDS_0027)
tail protein (CDS_0020)
CDS_0023
CDS_0026
CDS_0028
CDS_0029
IMGVR_UViG_2871040012_000001|2871040012|2871040022|144883-183593 18019:28338(+)
23
26
27
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35
IMGVR_UViG_3300008134_000213|3300008134|Ga0111464_100143|11929-49846 18274:28464(+)
23
26
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29
30
34
36
IMGVR_UViG_2854465412_000002|2854465412|2854465416|71431-109665 18444:29434(+)
4000 nt
moron, auxiliary metabolic gene and host takeover
other
tail
transcription regulation
unknown function