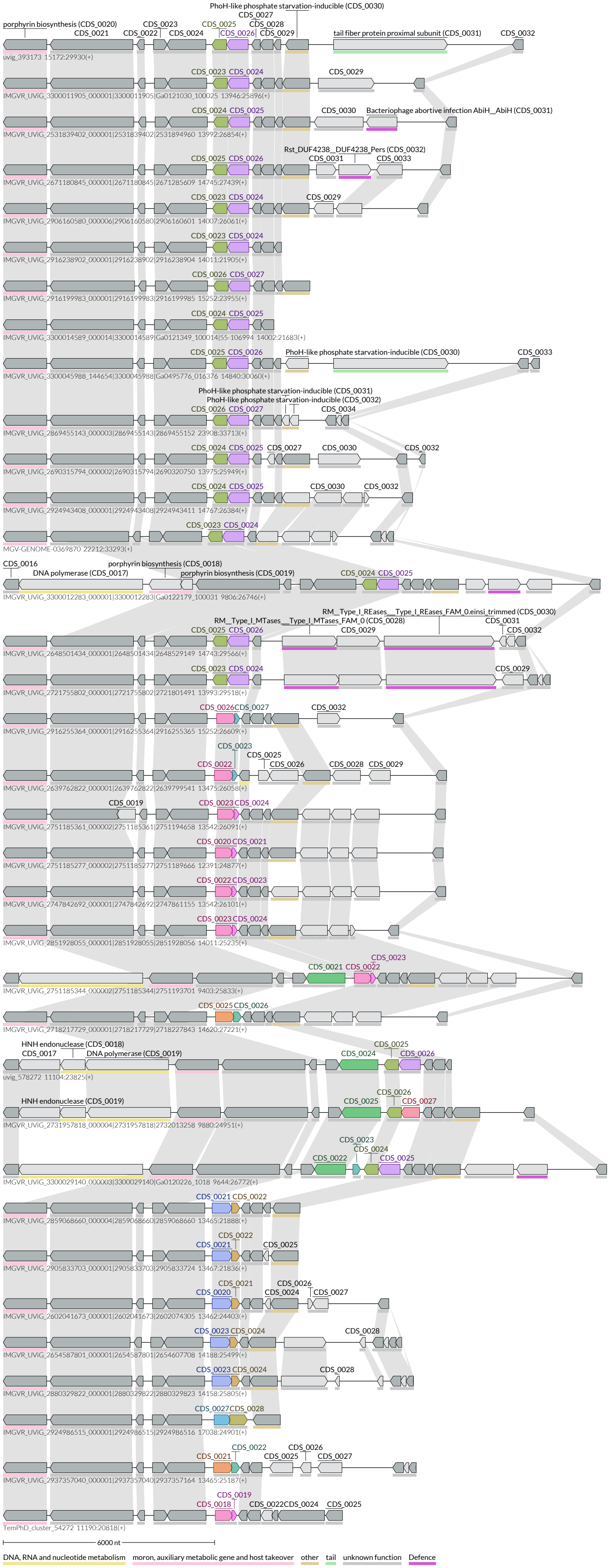
PhoH-like phosphate starvation-inducible (CDS_0030)
CDS_0027
porphyrin biosynthesis (CDS_0020)
CDS_0023
CDS_0025
CDS_0028
CDS_0021
CDS_0022
CDS_0024
CDS_0026
CDS_0029
tail fiber protein proximal subunit (CDS_0031)
CDS_0032
uvig_393173 15172:29930(+)
CDS_0023 CDS_0024
CDS_0029
IMGVR_UViG_3300011905_000001|3300011905|Ga0121030_100025 13946:25896(+)
CDS_0024 CDS_0025
CDS_0030
Bacteriophage abortive infection AbiH__AbiH (CDS_0031)
IMGVR_UViG_2531839402_000001|2531839402|2531894960 13992:26854(+)
Rst_DUF4238__DUF4238_Pers (CDS_0032)
CDS_0025 CDS_0026
CDS_0031
CDS_0033
IMGVR_UViG_2671180845_000001|2671180845|2671285609 14745:27439(+)
CDS_0023 CDS_0024
CDS_0029
IMGVR_UViG_2906160580_000006|2906160580|2906160601 14007:26061(+)
CDS_0023 CDS_0024
IMGVR_UViG_2916238902_000001|2916238902|2916238904 14011:21905(+)
CDS_0026 CDS_0027
IMGVR_UViG_2916199983_000001|2916199983|2916199985 15252:23955(+)
CDS_0024 CDS_0025
IMGVR_UViG_3300014589_000014|3300014589|Ga0121349_100014|55-106994 14002:21683(+)
CDS_0025 CDS_0026
PhoH-like phosphate starvation-inducible (CDS_0030)
CDS_0033
IMGVR_UViG_3300045988_144654|3300045988|Ga0495776_016376 14840:30060(+)
PhoH-like phosphate starvation-inducible (CDS_0031)
PhoH-like phosphate starvation-inducible (CDS_0032)
CDS_0026 CDS_0027
CDS_0034
IMGVR_UViG_2869455143_000003|2869455143|2869455152 23908:33713(+)
CDS_0024 CDS_0025
CDS_0027
CDS_0030
CDS_0032
IMGVR_UViG_2690315794_000002|2690315794|2690320750 13975:25949(+)
CDS_0024 CDS_0025
CDS_0030
CDS_0032
IMGVR_UViG_2924943408_000001|2924943408|2924943411 14767:26384(+)
CDS_0023 CDS_0024
MGV-GENOME-0369870 22212:33293(+)
CDS_0016
DNA polymerase (CDS_0017)
porphyrin biosynthesis (CDS_0018)
porphyrin biosynthesis (CDS_0019)
CDS_0024 CDS_0025
IMGVR_UViG_3300012283_000001|3300012283|Ga0122179_100031 9806:26746(+)
RM__Type_I_REases__Type_I_REases_FAM_0.einsi_trimmed (CDS_0030)
RM__Type_I_MTases__Type_I_MTases_FAM_0 (CDS_0028)
CDS_0031
CDS_0025 CDS_0026
CDS_0029
CDS_0032
IMGVR_UViG_2648501434_000001|2648501434|2648529149 14743:29566(+)
CDS_0023 CDS_0024
CDS_0029
IMGVR_UViG_2721755802_000001|2721755802|2721801491 13993:29518(+)
CDS_0026CDS_0027
CDS_0032
IMGVR_UViG_2916255364_000001|2916255364|2916255365 15252:26609(+)
CDS_0023
CDS_0025
CDS_0022
CDS_0026
CDS_0028
CDS_0029
IMGVR_UViG_2639762822_000001|2639762822|2639799541 13475:26058(+)
CDS_0019
CDS_0023 CDS_0024
IMGVR_UViG_2751185361_000002|2751185361|2751194658 13542:26091(+)
CDS_0020 CDS_0021
IMGVR_UViG_2751185277_000002|2751185277|2751189666 12391:24877(+)
CDS_0022 CDS_0023
IMGVR_UViG_2747842692_000001|2747842692|2747861155 13542:26101(+)
CDS_0023 CDS_0024
IMGVR_UViG_2851928055_000001|2851928055|2851928056 14011:25235(+)
CDS_0023
CDS_0021 CDS_0022
IMGVR_UViG_2751185344_000002|2751185344|2751193701 9403:25833(+)
CDS_0025 CDS_0026
IMGVR_UViG_2718217729_000001|2718217729|2718227843 14620:27221(+)
HNH endonuclease (CDS_0018)
CDS_0017
DNA polymerase (CDS_0019)
CDS_0025
CDS_0024
CDS_0026
uvig_578272 11104:23825(+)
CDS_0026
HNH endonuclease (CDS_0019)
CDS_0025
CDS_0027
IMGVR_UViG_2731957818_000004|2731957818|2732013258 9880:24951(+)
CDS_0023
CDS_0024
CDS_0022
CDS_0025
IMGVR_UViG_3300029140_000003|3300029140|Ga0120226_1018 9644:26772(+)
CDS_0021 CDS_0022
IMGVR_UViG_2859068660_000004|2859068660|2859068660 13465:21888(+)
CDS_0022
CDS_0021
CDS_0025
IMGVR_UViG_2905833703_000001|2905833703|2905833724 13467:21836(+)
CDS_0021
CDS_0026
CDS_0020
CDS_0024
CDS_0027
IMGVR_UViG_2602041673_000001|2602041673|2602074305 13462:24403(+)
CDS_0023 CDS_0024
CDS_0028
IMGVR_UViG_2654587801_000001|2654587801|2654607708 14188:25499(+)
CDS_0023 CDS_0024
CDS_0028
IMGVR_UViG_2880329822_000001|2880329822|2880329823 14158:25805(+)
CDS_0027CDS_0028
IMGVR_UViG_2924986515_000001|2924986515|2924986516 17038:24901(+)
CDS_0022
CDS_0026
CDS_0021
CDS_0025
CDS_0027
IMGVR_UViG_2937357040_000001|2937357040|2937357164 13465:25187(+)
CDS_0019
CDS_0018
CDS_0022CDS_0024
CDS_0025
TemPhD_cluster_54272 11190:20818(+)
6000 nt
DNA, RNA and nucleotide metabolism
moron, auxiliary metabolic gene and host takeover
other
tail
unknown function
Defence