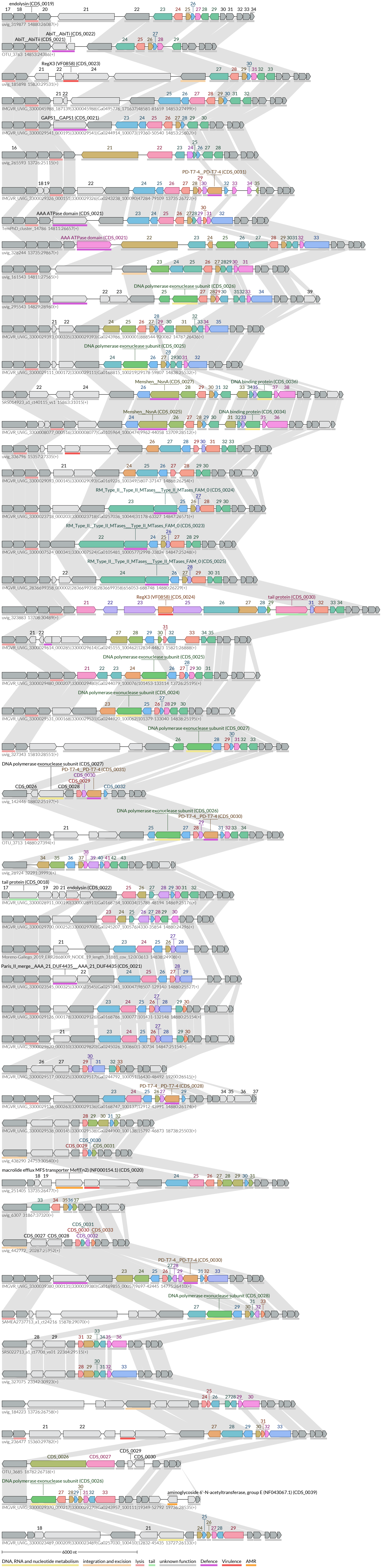
endolysin (CDS_0019)
uvig_319877 14880:26087(+)
AbiT__AbiTi (CDS_0022)
AbiT__AbiTii (CDS_0021)
OTU_3763 14853:24386(+)
RegX3 (VF0858) (CDS_0023)
uvig_185898 15830:29531(+)
IMGVR_UViG_3300045988_187139|3300045988|Ga0495776_171637|48581-81659 14853:27499(+)
GAPS1__GAPS1 (CDS_0021)
IMGVR_UViG_3300029541_000195|3300029541|Ga0244914_100073|19360-50540 14853:25602(+)
uvig_265593 13726:25115(+)
PD-T7-4__PD-T7-4 (CDS_0031)
IMGVR_UViG_3300029326_000151|3300029326|Ga0243238_100090|47284-79109 13735:26722(+)
AAA ATPase domain (CDS_0021)
TemPhD_cluster_14786 14811:26657(+)
AAA ATPase domain (CDS_0021)
uvig_328244 13735:29867(+)
uvig_161543 14811:27565(+)
DNA polymerase exonuclease subunit (CDS_0026)
uvig_295543 14829:28960(+)
IMGVR_UViG_3300029393_000335|3300029393|Ga0243986_1000001|888544-920082 14787:26436(+)
DNA polymerase exonuclease subunit (CDS_0025)
IMGVR_UViG_3300029111_000172|3300029111|Ga0168815_100219|29178-59807 14838:25532(+)
Menshen__NsnA (CDS_0027)
DNA binding protein (CDS_0036)
SRS014923_a1_ct40115_vs1 15863:31015(+)
Menshen__NsnA (CDS_0025)
DNA binding protein (CDS_0034)
IMGVR_UViG_3300008077_000516|3300008077|Ga0105964_1000474|9962-44058 13709:28512(+)
uvig_336796 15357:27335(+)
IMGVR_UViG_3300029093_000145|3300029093|Ga0169226_100349|5807-37147 14888:26254(+)
RM_Type_II__Type_II_MTases___Type_II_MTases_FAM_0 (CDS_0024)
IMGVR_UViG_3300023718_000203|3300023718|Ga0257036_10044|31178-63327 14847:26571(+)
RM_Type_II__Type_II_MTases___Type_II_MTases_FAM_0 (CDS_0023)
IMGVR_UViG_3300007524_000341|3300007524|Ga0105481_1000577|2998-33824 14847:25248(+)
RM_Type_II__Type_II_MTases___Type_II_MTases_FAM_0 (CDS_0025)
IMGVR_UViG_2836699358_000002|2836699358|2836699358|656053-688748 14880:26229(+)
RegX3 (VF0858) (CDS_0024)
tail protein (CDS_0030)
uvig_323883 13708:30489(+)
IMGVR_UViG_3300029614_000285|3300029614|Ga0245155_100462|12834-44823 15821:26888(+)
DNA polymerase exonuclease subunit (CDS_0025)
IMGVR_UViG_3300029480_000207|3300029480|Ga0244079_100076|101453-133114 13726:25195(+)
DNA polymerase exonuclease subunit (CDS_0024)
IMGVR_UViG_3300029531_000168|3300029531|Ga0244920_100062|101379-133040 14838:25195(+)
DNA polymerase exonuclease subunit (CDS_0027)
uvig_327343 15810:28551(+)
DNA polymerase exonuclease subunit (CDS_0027)
PD-T7-4__PD-T7-4 (CDS_0031)
CDS_0030
CDS_0029
CDS_0026
CDS_0028
CDS_0032
uvig_142446 18802:25197(+)
DNA polymerase exonuclease subunit (CDS_0026)
PD-T7-4__PD-T7-4 (CDS_0030)
OTU_3713 14880:27394(+)
uvig_28924 32291:39993(+)
tail protein (CDS_0018)
endolysin (CDS_0022)
IMGVR_UViG_3300028911_000190|3300028911|Ga0168754_100034|15788-48194 14869:25176(+)
IMGVR_UViG_3300029700_000252|3300029700|Ga0245207_100576|4330-35854 14880:24296(+)
Moreno-Gallego_2019_ERR2868009_NODE_19_length_31881_cov_12.003613 14838:24908(+)
Paris_II_merge__AAA_21_DUF4435__AAA_21_DUF4435 (CDS_0021)
IMGVR_UViG_3300023545_000325|3300023545|Ga0257041_100047|98507-129140 14880:25527(+)
IMGVR_UViG_3300029126_000178|3300029126|Ga0168786_100077|101431-132148 14880:25154(+)
IMGVR_UViG_3300029820_000310|3300029820|Ga0245026_100860|1-30734 14847:25154(+)
IMGVR_UViG_3300029517_000225|3300029517|Ga0244792_100051|16430-48492 19200:26515(+)
PD-T7-4__PD-T7-4 (CDS_0028)
IMGVR_UViG_3300029136_000263|3300029136|Ga0168747_100137|12912-43991 14880:26174(+)
IMGVR_UViG_3300029538_000145|3300029538|Ga0244900_100138|15792-46873 18738:25503(+)
CDS_0030
CDS_0029
CDS_0031
uvig_438290 24753:30540(+)
macrolide efflux MFS transporter Mef(En2) (NF000154.1) (CDS_0020)
uvig_251405 13735:26477(+)
uvig_6307 31867:37320(+)
CDS_0031
CDS_0030
CDS_0033
CDS_0027
CDS_0028
CDS_0032
uvig_442772_ 20287:25952(+)
PD-T7-4__PD-T7-4 (CDS_0030)
IMGVR_UViG_3300039380_000131|3300039380|Ga0169855_00057|9697-42445 14775:26410(+)
DNA polymerase exonuclease subunit (CDS_0028)
SAMEA2737713_a1_ct24216 15878:29070(+)
SRS022713_a1_ct7708_vs01 22384:29515(+)
uvig_327075 23342:30923(+)
uvig_184223 13726:26758(+)
uvig_236477 15360:29782(+)
CDS_0029
CDS_0026
CDS_0027
CDS_0030
OTU_3685 18782:26718(+)
DNA polymerase exonuclease subunit (CDS_0026)
aminoglycoside 6'-N-acetyltransferase, group E (NF043067.1) (CDS_0039)
IMGVR_UViG_3300029370_000217|3300029370|Ga0243957_100111|19349-52792 19736:28535(+)
IMGVR_UViG_3300023489_000209|3300023489|Ga0257030_100410|12832-45435 13727:26133(+)
6000 nt
DNA, RNA and nucleotide metabolism
integration and excision
lysis
tail
unknown function
Defence
Virulence
AMR