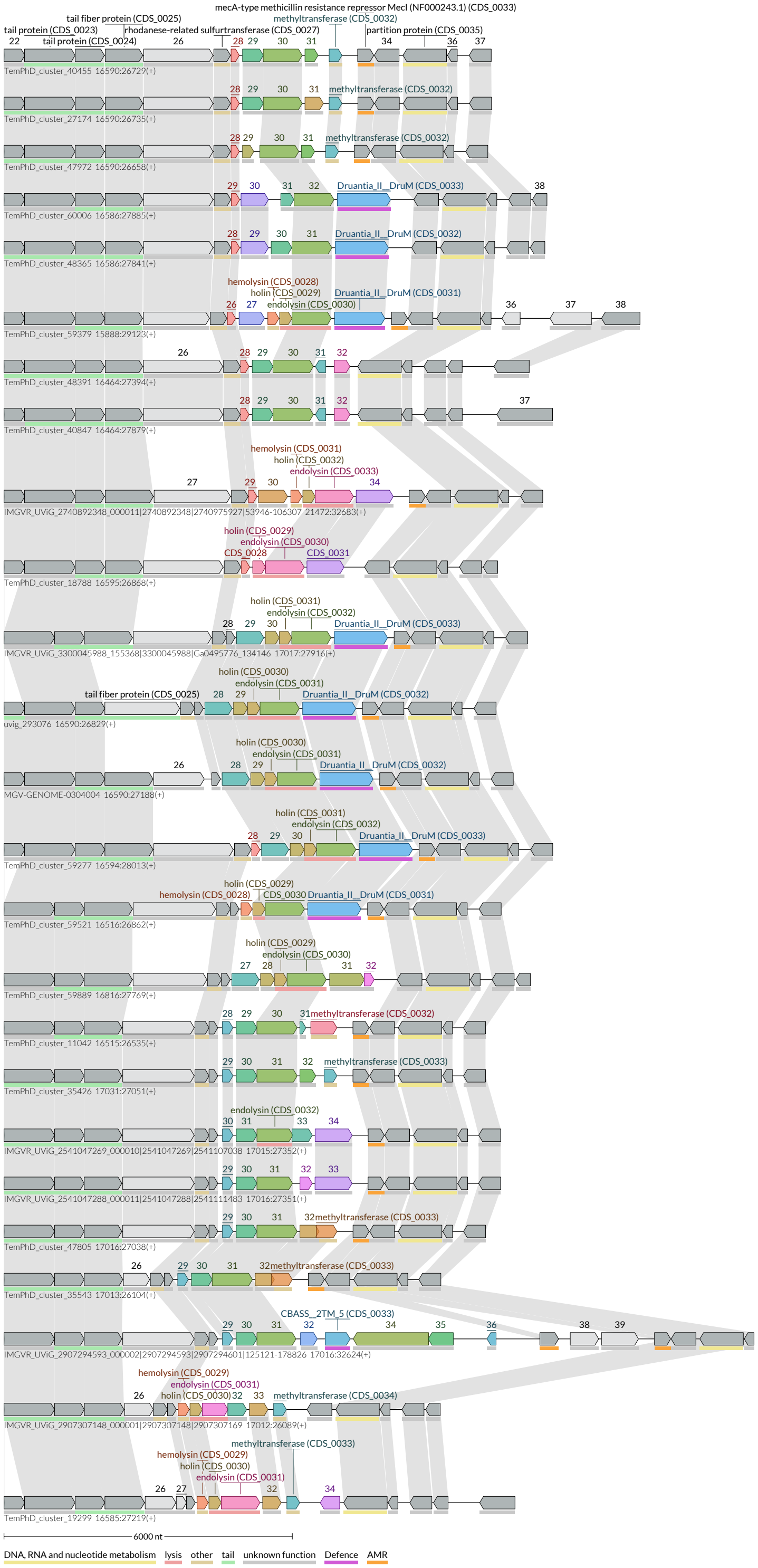
mecA-type methicillin resistance repressor MecI (NF000243.1) (CDS_0033)
tail fiber protein (CDS_0025)
methyltransferase (CDS_0032)
tail protein (CDS_0023)
rhodanese-related sulfurtransferase (CDS_0027)
partition protein (CDS_0035)
tail protein (CDS_0024)
TemPhD_cluster_40455 16590:26729(+)
TemPhD_cluster_27174 16590:26735(+)
TemPhD_cluster_47972 16590:26658(+)
Druantia_II_DruM (CDS_0033)
TemPhD_cluster_60006 16586:27885(+)
Druantia_II_DruM (CDS_0032)
TemPhD_cluster_48365 16586:27841(+)
hemolysin (CDS_0028)
holin (CDS_0029)
Druantia_II_DruM (CDS_0031)
endolysin (CDS_0030)
TemPhD_cluster_59379 15888:29123(+)
TemPhD_cluster_48391 16464:27394(+)
TemPhD_cluster_40847 16464:27879(+)
hemolysin (CDS_0031)
holin (CDS_0032)
endolysin (CDS_0033)
IMGVR_UViG_2740892348_000011|2740892348|2740975927|53946-106307 21472:32683(+)
holin (CDS_0029)
endolysin (CDS_0030)
CDS_0028
CDS_0031
TemPhD_cluster_18788 16595:26868(+)
holin (CDS_0031)
endolysin (CDS_0032)
Druantia_II_DruM (CDS_0033)
IMGVR_UViG_3300045988_155368|3300045988|Ga0495776_134146 17017:27916(+)
holin (CDS_0030)
endolysin (CDS_0031)
tail fiber protein (CDS_0025)
Druantia_II_DruM (CDS_0032)
uvig_293076 16590:26829(+)
holin (CDS_0030)
endolysin (CDS_0031)
Druantia_II_DruM (CDS_0032)
MGV-GENOME-0304004 16590:27188(+)
holin (CDS_0031)
endolysin (CDS_0032)
Druantia_II_DruM (CDS_0033)
TemPhD_cluster_59277 16594:28013(+)
holin (CDS_0029)
hemolysin (CDS_0028)
CDS_0030
Druantia_II_DruM (CDS_0031)
TemPhD_cluster_59521 16516:26862(+)
holin (CDS_0029)
endolysin (CDS_0030)
TemPhD_cluster_59889 16816:27769(+)
methyltransferase (CDS_0032)
TemPhD_cluster_11042 16515:26535(+)
methyltransferase (CDS_0033)
TemPhD_cluster_35426 17031:27051(+)
endolysin (CDS_0032)
IMGVR_UViG_2541047269_000010|2541047269|2541107038 17015:27352(+)
IMGVR_UViG_2541047288_000011|2541047288|2541111483 17016:27351(+)
TemPhD_cluster_47805 17016:27038(+)
TemPhD_cluster_35543 17013:26104(+)
CBASS_2TM_5 (CDS_0033)
IMGVR_UViG_2907294593_000002|2907294593|2907294601|125121-178826 17016:32624(+)
hemolysin (CDS_0029)
endolysin (CDS_0031)
holin (CDS_0030)
methyltransferase (CDS_0034)
IMGVR_UViG_2907307148_000001|2907307148|2907307169 17012:26089(+)
methyltransferase (CDS_0033)
hemolysin (CDS_0029)
holin (CDS_0030)
endolysin (CDS_0031)
TemPhD_cluster_19299 16585:27219(+)
6000 nt
DNA, RNA and nucleotide metabolism
lysis
other
tail
unknown function
Defence
AMR