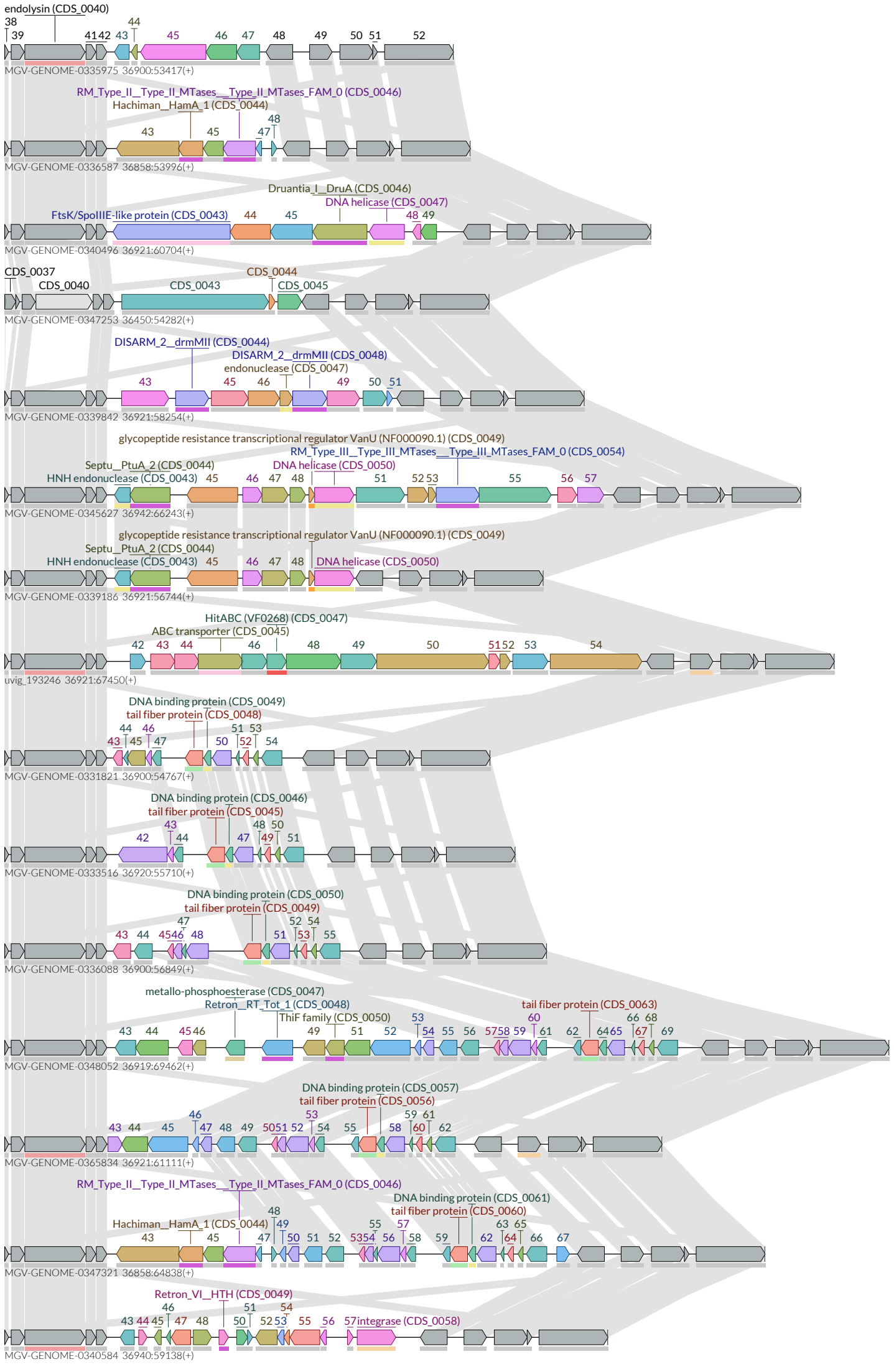
endolysin (CDS_0040)
MGV-GENOME-0335975 36900:53417(+)
RM_Type_II__Type_II_MTases__Type_II_MTases_FAM_0 (CDS_0046)
Hachiman__HamA_1 (CDS_0044)
MGV-GENOME-0336587 36858:53996(+)
Druantia_I__DruA (CDS_0046)
DNA helicase (CDS_0047)
FtsK/SpoIIIE-like protein (CDS_0043)
MGV-GENOME-0340496 36921:60704(+)
CDS_0037
CDS_0040
CDS_0043
CDS_0044
CDS_0045
MGV-GENOME-0347253 36450:54282(+)
DISARM_2__drmMII (CDS_0044)
DISARM_2__drmMII (CDS_0048)
endonuclease (CDS_0047)
MGV-GENOME-0339842 36921:58254(+)
glycopeptide resistance transcriptional regulator VanU (NF000090.1) (CDS_0049)
RM_Type_III__Type_III_MTases__Type_III_MTases_FAM_0 (CDS_0054)
Septu__PtuA_2 (CDS_0044)
DNA helicase (CDS_0050)
HNH endonuclease (CDS_0043)
MGV-GENOME-0345627 36942:66243(+)
glycopeptide resistance transcriptional regulator VanU (NF000090.1) (CDS_0049)
Septu__PtuA_2 (CDS_0044)
HNH endonuclease (CDS_0043)
DNA helicase (CDS_0050)
MGV-GENOME-0339186 36921:56744(+)
HitABC (VF0268) (CDS_0047)
ABC transporter (CDS_0045)
uvig_193246 36921:67450(+)
DNA binding protein (CDS_0049)
tail fiber protein (CDS_0048)
MGV-GENOME-0331821 36900:54767(+)
DNA binding protein (CDS_0046)
tail fiber protein (CDS_0045)
MGV-GENOME-0333516 36920:55710(+)
DNA binding protein (CDS_0050)
tail fiber protein (CDS_0049)
MGV-GENOME-0336088 36900:56849(+)
metallo-phosphoesterase (CDS_0047)
Retron__RT_Tot_1 (CDS_0048)
ThiF family (CDS_0050)
tail fiber protein (CDS_0063)
MGV-GENOME-0348052 36919:69462(+)
DNA binding protein (CDS_0057)
tail fiber protein (CDS_0056)
MGV-GENOME-0365834 36921:61111(+)
RM_Type_II__Type_II_MTases__Type_II_MTases_FAM_0 (CDS_0046)
DNA binding protein (CDS_0061)
tail fiber protein (CDS_0060)
Hachiman__HamA_1 (CDS_0044)
MGV-GENOME-0347321 36858:64838(+)
Retron_VI__HTH (CDS_0049)
integrase (CDS_0058)
MGV-GENOME-0340584 36940:59138(+)
10000 nt
DNA, RNA and nucleotide metabolism
integration and excision
lysis
moron, auxiliary metabolic gene and host takeover
other
tail
unknown function
Defence
Virulence
AMR