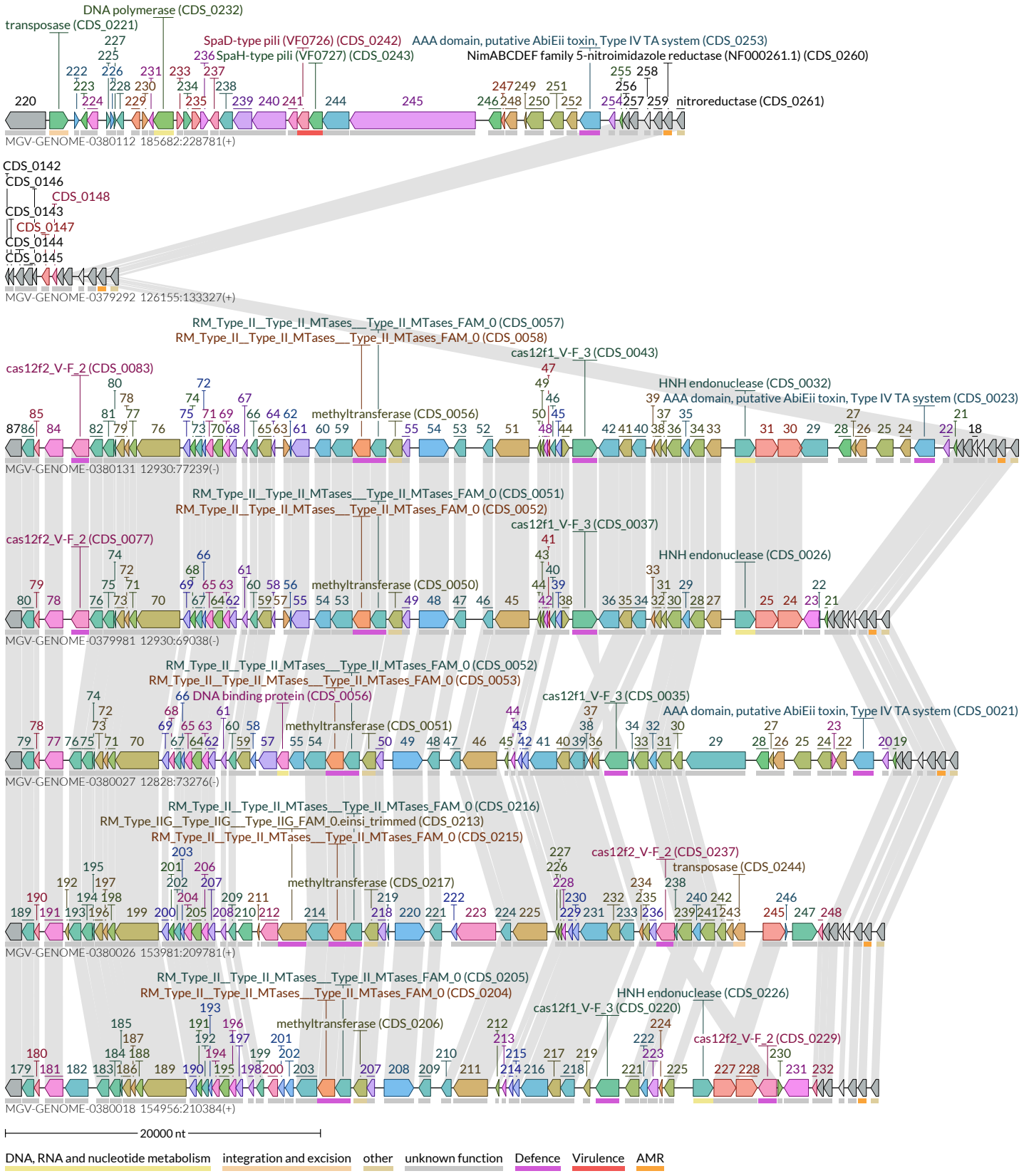
DNA polymerase (CDS_0232)
transposase (CDS_0221)
SpaD-type pili (VF0726) (CDS_0242)
SpaH-type pili (VF0727) (CDS_0243)
AAA domain, putative AbiEii toxin, Type IV TA system (CDS_0253)
NimABCDEF family 5-nitroimidazole reductase (NF000261.1) (CDS_0260)
nitroreductase (CDS_0261)
MGV-GENOME-0380112 185682:228781(+)
CDS_0142
CDS_0146
CDS_0148
CDS_0143
CDS_0147
CDS_0144
CDS_0145
MGV-GENOME-0379292 126155:133327(+)
RM_Type_II__Type_II_MTases__Type_II_MTases_FAM_0 (CDS_0057)
RM_Type_II__Type_II_MTases__Type_II_MTases_FAM_0 (CDS_0058)
cas12f1_V-F_3 (CDS_0043)
cas12f2_V-F_2 (CDS_0083)
HNH endonuclease (CDS_0032)
AAA domain, putative AbiEii toxin, Type IV TA system (CDS_0023)
methyltransferase (CDS_0056)
MGV-GENOME-0380131 12930:77239(-)
RM_Type_II__Type_II_MTases__Type_II_MTases_FAM_0 (CDS_0051)
RM_Type_II__Type_II_MTases__Type_II_MTases_FAM_0 (CDS_0052)
cas12f1_V-F_3 (CDS_0037)
cas12f2_V-F_2 (CDS_0077)
HNH endonuclease (CDS_0026)
methyltransferase (CDS_0050)
MGV-GENOME-0379981 12930:69038(-)
RM_Type_II__Type_II_MTases__Type_II_MTases_FAM_0 (CDS_0052)
RM_Type_II__Type_II_MTases__Type_II_MTases_FAM_0 (CDS_0053)
DNA binding protein (CDS_0056)
cas12f1_V-F_3 (CDS_0035)
AAA domain, putative AbiEii toxin, Type IV TA system (CDS_0021)
methyltransferase (CDS_0051)
MGV-GENOME-0380027 12828:73276(-)
RM_Type_II__Type_II_MTases__Type_II_MTases_FAM_0 (CDS_0216)
RM_Type_IIG__Type_IIG__Type_IIG_FAM_0.einsi_trimmed (CDS_0213)
RM_Type_II__Type_II_MTases__Type_II_MTases_FAM_0 (CDS_0215)
cas12f2_V-F_2 (CDS_0237)
transposase (CDS_0244)
methyltransferase (CDS_0217)
MGV-GENOME-0380026 153981:209781(+)
RM_Type_II__Type_II_MTases__Type_II_MTases_FAM_0 (CDS_0205)
RM_Type_II__Type_II_MTases__Type_II_MTases_FAM_0 (CDS_0204)
HNH endonuclease (CDS_0226)
cas12f1_V-F_3 (CDS_0220)
methyltransferase (CDS_0206)
cas12f2_V-F_2 (CDS_0229)
MGV-GENOME-0380018 154956:210384(+)
20000 nt
DNA, RNA and nucleotide metabolism
integration and excision
other
unknown function
Defence
Virulence
AMR