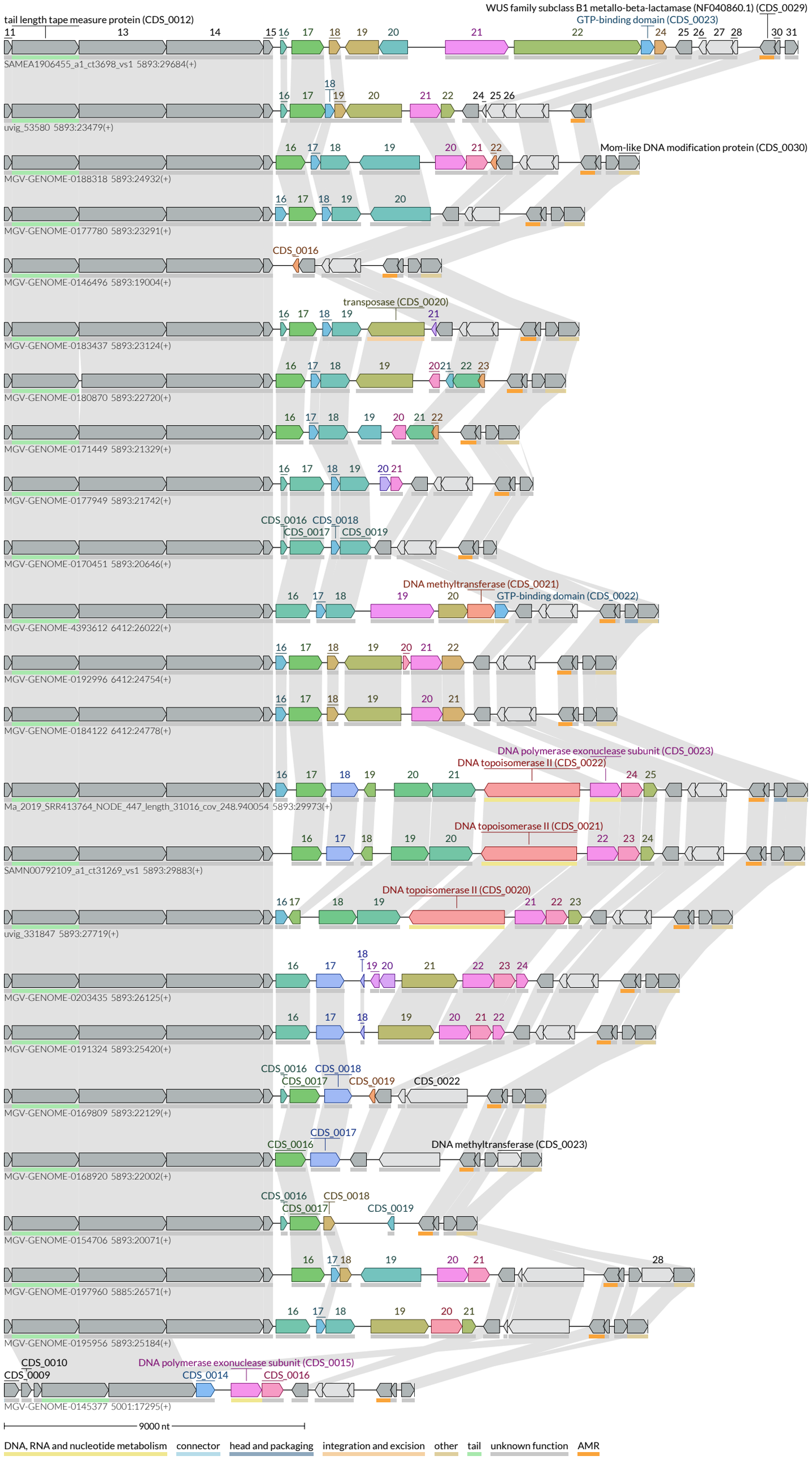
tail length tape measure protein (CDS_0012)
WUS family subclass B1 metallo-beta-lactamase (NF040860.1) (CDS_0029)
GTP-binding domain (CDS_0023)
SAMEA1906455_a1_ct3698_vs1 5893:29684(+)
uvig_53580 5893:23479(+)
Mom-like DNA modification protein (CDS_0030)
MGV-GENOME-0188318 5893:24932(+)
MGV-GENOME-0177780 5893:23291(+)
CDS_0016
MGV-GENOME-0146496 5893:19004(+)
transposase (CDS_0020)
MGV-GENOME-0183437 5893:23124(+)
MGV-GENOME-0180870 5893:22720(+)
MGV-GENOME-0171449 5893:21329(+)
MGV-GENOME-0177949 5893:21742(+)
CDS_0016 CDS_0018
CDS_0017 CDS_0019
MGV-GENOME-0170451 5893:20646(+)
DNA methyltransferase (CDS_0021)
GTP-binding domain (CDS_0022)
MGV-GENOME-4393612 6412:26022(+)
MGV-GENOME-0192996 6412:24754(+)
MGV-GENOME-0184122 6412:24778(+)
DNA polymerase exonuclease subunit (CDS_0023)
DNA topoisomerase II (CDS_0022)
Ma_2019_SRR413764_NODE_447_length_31016_cov_248.940054 5893:29973(+)
DNA topoisomerase II (CDS_0021)
SAMN00792109_a1_ct31269_vs1 5893:29883(+)
DNA topoisomerase II (CDS_0020)
uvig_331847 5893:27719(+)
MGV-GENOME-0203435 5893:26125(+)
MGV-GENOME-0191324 5893:25420(+)
CDS_0016 CDS_0018
CDS_0017 CDS_0019 CDS_0022
MGV-GENOME-0169809 5893:22129(+)
CDS_0017
CDS_0016
DNA methyltransferase (CDS_0023)
MGV-GENOME-0168920 5893:22002(+)
CDS_0016 CDS_0018
CDS_0017 CDS_0019
MGV-GENOME-0154706 5893:20071(+)
MGV-GENOME-0197960 5885:26571(+)
MGV-GENOME-0195956 5893:25184(+)
CDS_0010
CDS_0009
DNA polymerase exonuclease subunit (CDS_0015)
CDS_0014 CDS_0016
MGV-GENOME-0145377 5001:17295(+)
9000 nt
DNA, RNA and nucleotide metabolism
connector
head and packaging
integration and excision
other
tail
unknown function
AMR