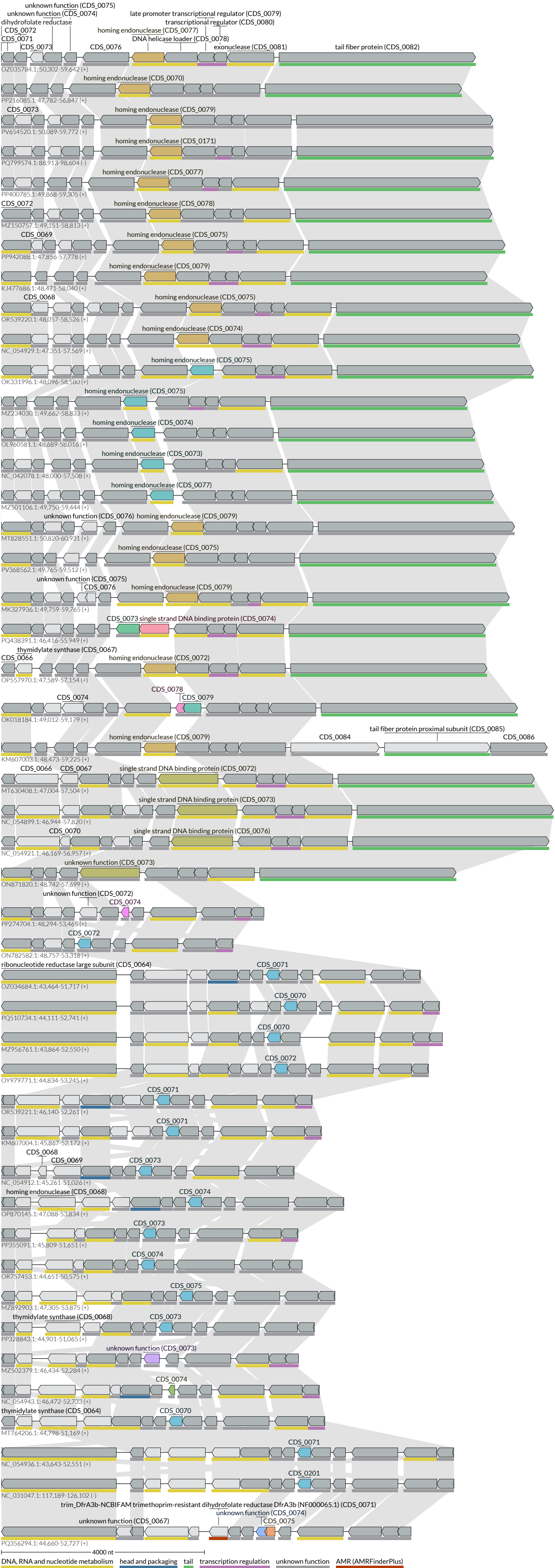
unknown function (CDS_0075)
unknown function (CDS_0074)
late promoter transcriptional regulator (CDS_0079)
dihydrofolate reductase
transcriptional regulator (CDS_0080)
CDS_0072
homing endonuclease (CDS_0077)
CDS_0071
DNA helicase loader (CDS_0078)
CDS_0073
CDS_0076
exonuclease (CDS_0081)
tail fiber protein (CDS_0082)
OZ035784.1: 50,302-59,642 (+)
homing endonuclease (CDS_0070)
PP216085.1: 47,782-56,847 (+)
CDS_0073
homing endonuclease (CDS_0079)
PV654520.1: 50,089-59,772 (+)
homing endonuclease (CDS_0171)
PQ799574.1: 88,913-98,604 (-)
homing endonuclease (CDS_0077)
PP400785.1: 49,868-59,305 (+)
CDS_0072
homing endonuclease (CDS_0078)
MZ150757.1: 49,151-58,813 (+)
CDS_0069
homing endonuclease (CDS_0075)
PP942088.1: 47,856-57,778 (+)
homing endonuclease (CDS_0079)
KJ477686.1: 48,471-58,040 (+)
CDS_0068
homing endonuclease (CDS_0075)
OR539220.1: 48,057-58,526 (+)
homing endonuclease (CDS_0074)
NC_054929.1: 47,351-57,569 (+)
homing endonuclease (CDS_0075)
OK331996.1: 48,096-58,580 (+)
homing endonuclease (CDS_0075)
MZ234030.1: 49,662-58,833 (+)
homing endonuclease (CDS_0074)
OL960581.1: 48,689-58,016 (+)
homing endonuclease (CDS_0073)
NC_042078.1: 48,000-57,508 (+)
homing endonuclease (CDS_0077)
MZ501106.1: 49,750-59,444 (+)
unknown function (CDS_0076)
homing endonuclease (CDS_0079)
MT828551.1: 50,820-60,931 (+)
homing endonuclease (CDS_0075)
PV368562.1: 49,765-59,512 (+)
unknown function (CDS_0075)
CDS_0076
homing endonuclease (CDS_0079)
MK327936.1: 49,759-59,765 (+)
CDS_0073 single strand DNA binding protein (CDS_0074)
PQ438391.1: 46,416-55,949 (+)
thymidylate synthase (CDS_0067)
CDS_0066
homing endonuclease (CDS_0072)
OP557970.1: 47,589-57,154 (+)
CDS_0078
CDS_0074
CDS_0079
OK018184.1: 49,012-59,179 (+)
tail fiber protein proximal subunit (CDS_0085)
homing endonuclease (CDS_0079)
CDS_0084
CDS_0086
KM607003.1: 48,473-59,225 (+)
CDS_0066
CDS_0067
single strand DNA binding protein (CDS_0072)
MT630408.1: 47,004-57,504 (+)
single strand DNA binding protein (CDS_0073)
NC_054899.1: 46,944-57,820 (+)
CDS_0070
single strand DNA binding protein (CDS_0076)
NC_054921.1: 46,169-56,957 (+)
unknown function (CDS_0073)
ON871820.1: 48,742-57,699 (+)
unknown function (CDS_0072)
CDS_0074
PP274704.1: 48,294-53,465 (+)
CDS_0072
ON782582.1: 48,757-53,318 (+)
ribonucleotide reductase large subunit (CDS_0064)
CDS_0071
OZ034684.1: 43,464-51,717 (+)
CDS_0070
PQ510734.1: 44,111-52,741 (+)
CDS_0070
MZ956761.1: 43,864-52,550 (+)
CDS_0072
OY979771.1: 44,834-53,245 (+)
CDS_0071
OR539221.1: 46,140-52,261 (+)
CDS_0071
KM607004.1: 45,867-52,172 (+)
CDS_0068
CDS_0069
CDS_0073
NC_054912.1: 45,261-51,026 (+)
homing endonuclease (CDS_0068)
CDS_0074
OP870145.1: 47,088-53,834 (+)
CDS_0073
PP355091.1: 45,809-51,651 (+)
CDS_0074
OR757453.1: 44,651-50,575 (+)
CDS_0075
MZ892903.1: 47,305-53,875 (+)
thymidylate synthase (CDS_0068)
CDS_0073
PP328843.1: 44,901-51,065 (+)
unknown function (CDS_0073)
MZ502379.1: 46,434-52,284 (+)
CDS_0074
NC_054943.1: 46,472-52,733 (+)
thymidylate synthase (CDS_0064)
CDS_0070
MT764206.1: 44,798-51,169 (+)
CDS_0071
NC_054936.1: 43,643-52,551 (+)
CDS_0201
NC_031047.1: 117,189-126,102 (-)
trim_DfrA3b-NCBIFAM trimethoprim-resistant dihydrofolate reductase DfrA3b (NF000065.1) (CDS_0071)
unknown function (CDS_0074)
unknown function (CDS_0067)
CDS_0075
PQ356294.1: 44,660-52,727 (+)
4000 nt
DNA, RNA and nucleotide metabolism
head and packaging
tail
transcription regulation
unknown function
AMR (AMRFinderPlus)