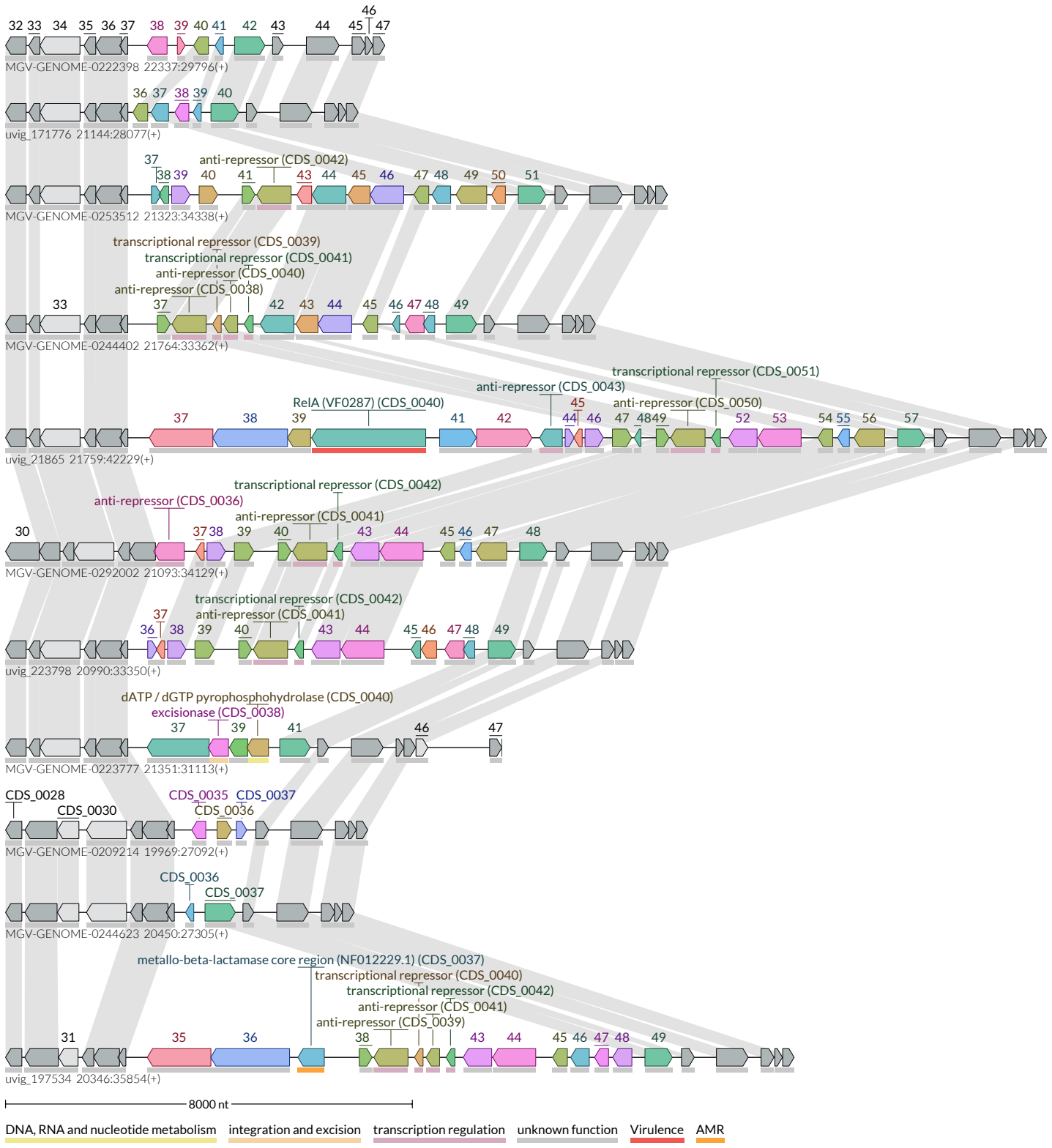
MGV-GENOME-0222398 22337:29796(+)
uvig_171776 21144:28077(+)
anti-repressor (CDS_0042)
MGV-GENOME-0253512 21323:34338(+)
transcriptional repressor (CDS_0039)
transcriptional repressor (CDS_0041)
anti-repressor (CDS_0040)
anti-repressor (CDS_0038)
MGV-GENOME-0244402 21764:33362(+)
transcriptional repressor (CDS_0051)
anti-repressor (CDS_0043)
RelA (VF0287) (CDS_0040)
anti-repressor (CDS_0050)
uvig_21865 21759:42229(+)
transcriptional repressor (CDS_0042)
anti-repressor (CDS_0036)
anti-repressor (CDS_0041)
MGV-GENOME-0292002 21093:34129(+)
transcriptional repressor (CDS_0042)
anti-repressor (CDS_0041)
uvig_223798 20990:33350(+)
dATP / dGTP pyrophosphohydrolase (CDS_0040)
excisionase (CDS_0038)
MGV-GENOME-0223777 21351:31113(+)
CDS_0028
CDS_0030
CDS_0035
CDS_0037
CDS_0036
MGV-GENOME-0209214 19969:27092(+)
CDS_0036
CDS_0037
MGV-GENOME-0244623 20450:27305(+)
metallo-beta-lactamase core region (NF012229.1) (CDS_0037)
transcriptional repressor (CDS_0040)
transcriptional repressor (CDS_0042)
anti-repressor (CDS_0041)
anti-repressor (CDS_0039)
uvig_197534 20346:35854(+)
8000 nt
DNA, RNA and nucleotide metabolism
integration and excision
transcription regulation
unknown function
Virulence
AMR