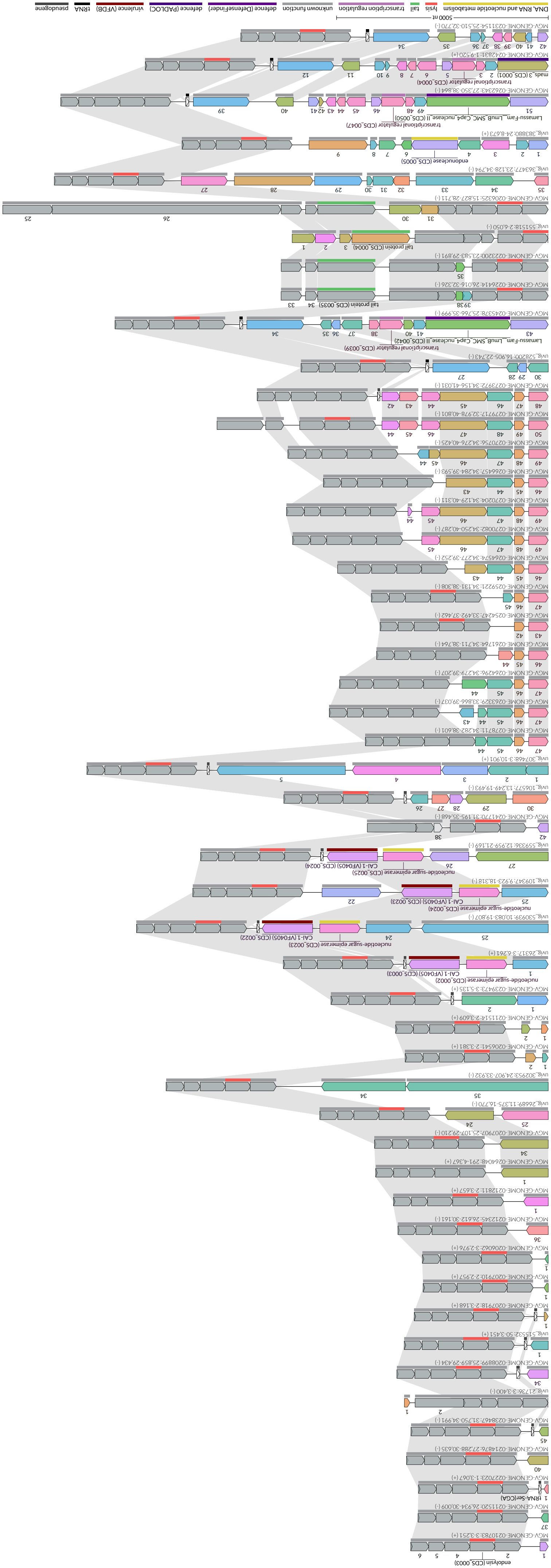
endolysin (CDS_0003)
MGV-GENOME-021078 3: 3-3,251 (+)
MGV-GENOME-021152 0: 26,934-30,009 (-)
MGV-GENOME-022702 3: 1-3,067 (+)
MGV-GENOME-021487 6: 27,288-30,635 (-)
MGV-GENOME-023846 7: 31,750-34,991 (-)
uvig_21736: 3-3,400 (-)
MGV-GENOME-020899 9: 25,859-29,434 (-)
uvig_515532: 50-3,451 (+)
MGV-GENOME-020791 8: 2-3,168 (+)
MGV-GENOME-020791 0: 2-2,957 (+)
MGV-GENOME-020606 2: 3-2,976 (+)
MGV-GENOME-021234 5: 26,612-30,161 (-)
MGV-GENOME-021281 1: 2-3,657 (+)
MGV-GENOME-026404 8: 291-4,367 (+)
MGV-GENOME-020790 7: 25,107-29,210 (-)
uvig_26689: 11,375-16,770 (-)
uvig_302953: 24,907-33,932 (-)
MGV-GENOME-020654 1: 2-3,381 (+)
MGV-GENOME-021151 4: 2-3,609 (+)
MGV-GENOME-023947 3: 3-5,135 (+)
nucleotide-sugar epimerase (CDS_0002)
CAI-1 (VF0405) (CDS_0003)
uvig_26317: 3-6,261 (+)
nucleotide-sugar epimerase (CDS_0023)
CAI-1 (VF0405) (CDS_0022)
uvig_530939: 10,083-19,807 (-)
nucleotide-sugar epimerase (CDS_0024)
CAI-1 (VF0405) (CDS_0023)
uvig_109347: 9,923-18,318 (-)
nucleotide-sugar epimerase (CDS_0025)
CAI-1 (VF0405) (CDS_0024)
uvig_559336: 12,959-21,169 (-)
MGV-GENOME-024177 0: 31,195-35,468 (-)
uvig_106577: 13,249-19,493 (-)
5000 bp
DNA, RNA and nucleotide metabolism
lysis
tail
transcription regulation
unknown function
defence (DefenseFinder)
defence (PADLOC)
virulence (VFDB)
tRNA
pseudogene