

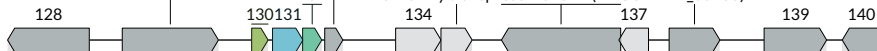
QacCGHJ-NCBIFAM QacCGHJ group quaternary ammonium compound efflux SMR transporter (NF000384.1) (AHCOELMN_00135)

Small, acid-soluble spore protein A (AHCOELMN_00133)

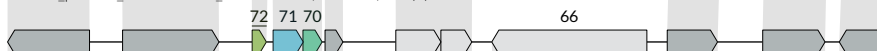
Tyrosine recombinase XerC (AHCOELMN_00129)

Pyrophosphatase PpaX (AHCOELMN_00138)

IS4 family transposase IS231B (AHCOELMN_00136)



IMGPR_plasmid_3300053491_054860: 123,279-133,952 (+)

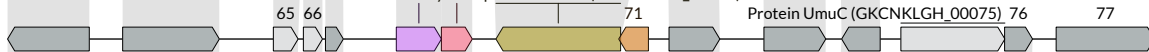


IMGPR_plasmid_3300053491_055309: 59,558-70,204 (-)

QacCGHJ-NCBIFAM QacCGHJ group quaternary ammonium compound efflux SMR transporter (NF000384.1) (GKCNKLGH_00069)

Nucleoid occlusion factor SlmA (GKCNKLGH_00068)

IS4 family transposase IS231B (GKCNKLGH_00070)



IMGPR_plasmid_3300053491_053186: 58,435-72,455 (+)

QacCGHJ-NCBIFAM QacCGHJ group quaternary ammonium compound efflux SMR transporter (NF000384.1) (DJLLHCGC_00186)

Nucleoid occlusion factor SlmA (DJLLHCGC_00187)

IS4 family transposase ISBth5 (EKP KIBON_00004)

Protein UmuC (GKCNKLGH_00075)

DNA polymerase IV (DJLLHCGC_00180)



IMGPR_plasmid_3300053491_055794: 180,419-194,405 (-)

IS4 family transposase ISBth5 (EKP KIBON_00004)



IMGPR_plasmid_3300053491_052356: 2,609-20,756 (+)



IMGPR_plasmid_3300053491_055310: 59,833-78,130 (-)



IMGPR_plasmid_3300053491_054859: 50,735-68,983 (+)

7000 nt

AMR (AMRFinderPlus)