

Figure S1 Sequence comparison of pNDM_SCLZS30 with other GR59 plasmids. pNDM_SCLZS30 was used as a reference, and its gene annotations are indicated by black arrows in the outer circle. *parAB* and *repB* are highlighted in pink. Gaps indicate regions that are missing in the respective plasmid when compared to the reference plasmid.

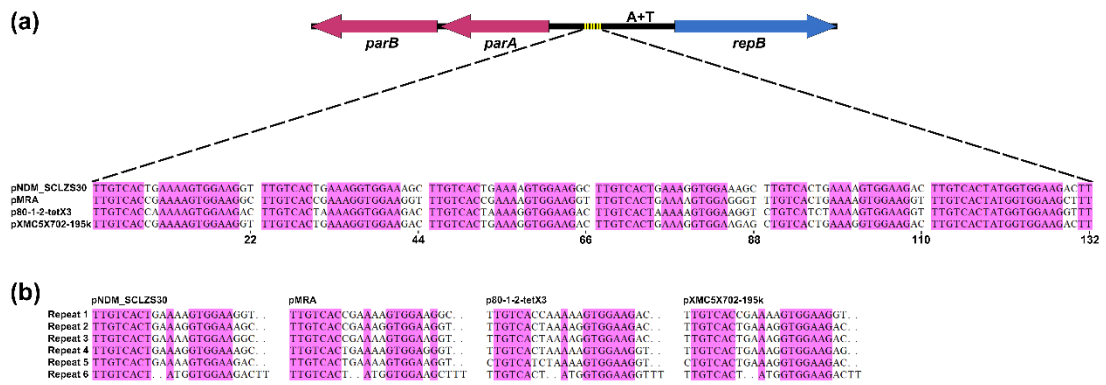


Figure S2 Characterization of the conserved region of GR59 plasmids. (a) Comparison of the different repeats between plasmids. *parAB* and *repB* are indicated by red and blue arrows. Yellow vertical bars indicate tandem repeats, and ‘A+T’ represents the AT-rich region. (b) Comparison of the repeats in different plasmids. Identical nucleotide sequences are indicated by pink shading.