



FIG S1 Depth of reads mapped on pE044_IMP6. DNA fragments extracted from bands with sizes of (A) 50 kbp, (B) 100 kbp, (C) 150 kbp, and (D) 200 kbp in S1-PFGE were sequenced by Illumina MiSeq. The read depth against pE044_IMP6 is indicated. The three blue shades represent minimum, average and maximum coverage values for the aggregated mapped reads.