

S1 Table. Oligonucleotide sequences of primers used in this study.

| Gene | Primer sequences | PCR product size (bp) | Reference |
|--------------|--------------------------------|-----------------------|---------------|
| <i>fosA</i> | F- ATCTGTGGGTCTGCCTGTCGT | 271 | Lu et al. [1] |
| | R- ATGCCCGCATAGGGCTTCT | | |
| <i>fosA3</i> | F- CGGAGCCTATCTCTCCTGTG | 219 | this study |
| | R- CCGTCAGGGTCGAGAAAATA | | |
| <i>fosA5</i> | F- GCGCCGAGCGTGGCGTTT | 283 | this study |
| | R- GCCATCGGGATCGAGGAA | | |
| <i>fosB</i> | F- AGGTGAGACCTCGGCCTATT | 302 | this study |
| | R- GAGGTTTAGCCTCTTTATAATAACTCA | | |
| <i>fosC2</i> | F- GGGCATATCTGAGCTTGGAG | 212 | this study |
| | R- CAATTTATGGCCGTCAGGAT | | |
| <i>fosX</i> | F- GTTGCGTTTAAGGCAGGAAG | 425 | this study |
| | R- GGCTCCATTTGTTGGACAGT | | |

Reference

1. Lu PL, Hsieh YJ, Lin JE, Huang JW, Yang TY, Lin L, Tseng SP (2016) Characterisation of fosfomycin resistance mechanisms and molecular epidemiology in extended-spectrum beta-lactamase-producing *Klebsiella pneumoniae* isolates. Int J Antimicrob Agents 48 (5):564-568. doi:10.1016/j.ijantimicag.2016.08.013 PMID: 27765412