

**Table S1. Primers used in this study**

Primer <sup>a</sup>	Sequence (5'-3') <sup>b</sup>	Purpose
2A*-F	<u>GGATCC</u> AAATAGTGCCTCTGGGCAAA	Amplification primers of p3M-2A*
2A*-R	<u>GGATCC</u> ACGGGCGAGTTAGTTAGCAA	
2B*-F	<u>GGATCC</u> GTATGAATCAATTTTAGCTAGAG	Amplification primers of p3M-2B*
2B*-R	<u>GGATCC</u> AAATCCCTTTTATGTAAACGG	
q3M-16S-F	TCGCCTAGGTGAGCCTTTAC	Primers of reference gene 16S for qRT-PCR in P3M and corresponding strains
q3M-16S-R	CGAGCGGTAACAGGAGAAAG	
qDH-16S-F	CGGACGGGTGAGTAATGTCT	Primers of reference gene 16S for qRT-PCR in DH5a and corresponding strains
qDH-16S-R	TCAGACCAGCTAGGGATCGT	
q- <i>qnrD</i> -F	GCTCAAGGAGCTGATTTTCG	Primers of <i>qnrD</i> for qRT-PCR
q- <i>qnrD</i> -R	ACGATGAAAACCTCCCCACAA	
YZ-2A-F	GGTTGCATCGCATTTTACA	Validation primers of p3M-2A and p3M-2A*
YZ-2A-R	AGCTTTCGACACCTCCAAGA	
YZ-2B-F	CAGTTGCCAGCTATGTTGA	Validation primers of p3M-2B and p3M-2B*
YZ-2B-R	TCTGTCACTGCATCCAAACC	

<sup>a</sup> F, forward primer; R, reverse primer.

<sup>b</sup> Restriction sites are underlined.