Table S1 The primers used in this study.

Primer	Sequences (5'-3')
PF1	ATGGCTAAGCGGCTGGTCCTCTTTGCGGC
PR1	CTATCACTGGCTAGCCGACAATGCGTCGC
PF2	GAAGGTGAGGCCTCTGGGCAACTACA
PR2	CTATCACTGGCTGGCCGACAATGCGTCG
PF3	GACAAGGGCAACAAGGGCAC*T*GCCCAACGT
PR3	AGCAGCCAGGACAAGGGCAACAAGGGCAC*T*

The positions of single-base substitution mutation were shown in bold italics.