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Table S2. Primers used in this study.

Primer	Sequence (5' – 3')^a
pCI372 FP	CGGGAAGCTAGAGTAAGTAG
pCI372 RP	CCTCTCGGTTATGAGTTAG
<i>stlA</i> FP	AAA <u>ACTGCAG</u> TTCTGGCAGCAGTGATTTG
<i>stlA</i> RP	GCT <u>CTAGACGGTC</u> GAGCAAGGTAATAGG
<i>stlA</i> -J FP	TGCTCTTCCGAAGCAGTCAG
<i>stlA</i> -J RP	AGCATATCGAAGACGGCCAG
<i>stlA</i> -OUT FP	CTGCTCTGTTGATGGGGTTT
<i>stlA</i> -OUT RP	CGGGCAACTACAAGGATGAT
<i>stlA</i> -IN FP	TATGGGAGGGGCTACTACGG
<i>stlA</i> -IN RP	ACCCAGTTGCCAAGCATATC
EZ-Tn FP-1	GCCAACGACTACGCAC TAGCCAAC
EZ-Tn RP-1	GAGCCAATATGCGAGAACACCCGAGAA

^aRestriction enzyme recognition sequences are underlined; FP= forward primer; RP= reverse primer.