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Table S2. Primers used in this study.

Primer	Sequence (5' – 3')^a
pCI372 FP	CGGGAAGCTAGAGTAAGTAG
pCI372 RP	CCTCTCGGTTATGAGTTAG
<i>stI</i> A FP	AAA <u>ACTGCAGTTCTGGCAGCAGT</u> GATTTTG
<i>stI</i> A RP	G <u>CTCTAGACGGTCGAGCAAGGTA</u> ATAGG
<i>stI</i> A-J FP	TGCTCTTCCGAAGCAGTCAG
<i>stI</i> A-J RP	AGCATATCGAAGACGGCCAG
<i>stI</i> A-OUT FP	CTGCTCTGTTGATGGGGTTT
<i>stI</i> A-OUT RP	CGGGCAACTACAAGGATGAT
<i>stI</i> A-IN FP	TATGGGAGGGGCTACTACGG
<i>stI</i> A-IN RP	ACCCAGTTGCCAAGCATATC
EZ-Tn FP-1	GCCAACGACTACGCACTAGCCAAC
EZ-Tn RP-1	GAGCCAATATGCGAGAACACCCGAGAA

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