

**Table S2. Primers used in this study.**

<b>Primer</b>	<b>Sequence (5' – 3')<sup>a</sup></b>
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

<sup>a</sup>Restriction enzyme recognition sequences are underlined; FP= forward primer; RP= reverse primer.