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| Authors                     | Culligan, Eamonn P.;Sleator, Roy D.;Marchesi, Julian R.;Hill, Colin                                                                                                                                                                                                                                                                                                       |
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**Table S1. Oligonucleotide primers**

| <b>Primer</b>              | <b>Sequence (5' – 3')<sup>a</sup></b>   |
|----------------------------|-----------------------------------------|
| pCI372 FP                  | CGGGAAGCTAGAGTAAGTAG                    |
| pCI372 RP                  | CCTCTCGGTTATGAGTTAG                     |
| pBAD FP                    | ATGCCATAGCATTTTTATC                     |
| pBAD RP                    | GATTTAATCTGTATCAGG                      |
| <i>brpA<sub>S</sub></i> FP | AAA <u>ACTGCAG</u> ACCCAACACGATGCCATATT |
| <i>brpA<sub>S</sub></i> RP | GCT <u>CTAG</u> AACAGGGTGCGGTGATACA     |
| <i>brpAatfA</i> FP         | AAA <u>ACTGCAG</u> TAGCGGCTGGATCGGTAGTA |
| <i>brpAatfA</i> RP         | GCT <u>CTAG</u> AACCCAACACGATGCCATATT   |
| <i>brpA<sub>L</sub></i> FP | AAA <u>ACTGCAG</u> GCCGAATATCAACCCAACAC |
| <i>brpA<sub>L</sub></i> RP | GCT <u>CTAG</u> AAGGTATTTGTGCCTTGTGCT   |
| EZTn-FP-1                  | GCCAACGACTACGCACTAGCCAAC                |
| EZTn-RP-1                  | GAGCCAATATGCGAGAACACCCGAGAA             |

<sup>a</sup>Restriction enzyme cut-sites are underlined (*Pst*I, CTGCAG; *Xba*I, TCTAGA)