

Table S1. Oligonucleotide primers used in this study

locus	PCR / gene	forward-primer	sequence	reverse-primer	sequence	fragment length
HPI-MLST PCRs						
	<i>intB</i>	intB.for	5'-CCAGTCAGAGGAGCCAATTTT-3'	intB.rev	5'-CCGGGAGAAATTAGCGAGCAT-3'	1,7 kb
	<i>ybtQ</i>	ybtQ.for	5'-GCGGCAGGGCGGGTCAAATGA-3'	ybtQ.rev	5'-TGCGACGGGAAGCGGTGCTAT-3'	650 bp
	<i>ybtA</i>	ybtA.for	5'-GCCATAGCTAAAATAACCTTCCT-3'	ybtA.rev	5'-CCGGTAACAGCGAATCCTGAGAT-3'	1,5 kb
	<i>irp2</i>	irp2.for	5'-CTCCGCAGAACAGGTAGCCGA-3'	irp2.rev	5'-GCCAGATAATGTCCGCCGAGT-3'	637 bp
	<i>irp1</i>	irp1.for	5'-CCATCTACAGGCGGCTACGTC-3'	irp1.rev	5'-GTCTCCCGGCGGCATACACCA-3'	654 bp
	<i>fyuA</i>	fyuA.for	5'-CTACGACATGCCGACAATGCC-3'	fyuA.rev	5'-TGCTTCCCGCGCCATAACGTG-3'	670 bp
„upstream HPI“						
in HPI-positive strains	1	P1	5'-GGAAGTCGGGAGATAAGTCATTGATAAAGTGGC-3'	P8	5'-CCAATTTTCCCGATGCGTAG-3'	1,3 kb
in HPI-negative strains	2	P1	5'-GGAAGTCGGGAGATAAGTCATTGATAAAGTGGC-3'	P2	5'-GGCCGATTCCAACGCTCCAAG-3'	1,2 kb
“downstream HPI”						
in HPI-negative strains	3	P5	5'-GCGGCAGCACAAAGGTGTGGTAAA-3'	P6	5'-CTAAGCCGTTGTTAATCTGCGTA-3'	600 bp
	4	P10	5'-ACCAACCACGACGGCAATTTTAA-3'	P11	5'-AACCGCCTTGTCGAGAGTAAAG-3'	600 bp
	5	P12	5'-ACTTTGATTGCCGATGCTGGTAC-3'	P9	5'-CATGGTGACCTCGGCTACTCCCT-3'	650 bp
“downstream HPI”						
in HPI-positive strains	6	P7	5'-TGGCAACATCCTACTATCGGTC-3'	P4	5'-GACGTTCCGGCACATAGGTCA-3'	
	7	P3	5'-CACGTGGATGGGCCTCTTCTCAG-3'	P9	5'-CATGGTGACCTCGGCTACTCCCT-3'	1 kb