

S1 Table. Accession numbers of the sequences used in this study.

Strains	GenBank Accession numbers	
	16S rRNA	<i>rpoB</i> ^c
JZ1	KX359581 ^b	KY013617
JZ2	KU252651 ^a	KU298439
JZ3	KU252652 ^a	KU298440
JZ4	KX359591 ^b	KY013618
JZ5	KX359592 ^b	-
JZ6	KU252653 ^a	KU870633
JZ10	KX359583 ^b	KX965686
JZ11	KX359582 ^b	KX965687
JZ13	KU252654 ^a	KU870634
JZ14	KU252655 ^a	KU870629
JZ15	KU252656 ^a	KU298442
JZ16 ^T	KR534194 ^a	KU298441
JZ19	KX359590 ^b	-
JZ20	KU252657 ^a	KU870630
JZ21	KX359584 ^b	KX965688
JZ22	KX359589 ^b	KX965689
JZ23	KX359586 ^b	KX965690
JZ25	KU252658 ^a	KU870631
JZ26	KX359585 ^b	KX965691
JZ27	KU252659 ^a	KU870632
JZ28	KX359588 ^b	KX965692
JZ29	KX359587 ^b	KX965693
JZ32	KX359593 ^b	KX965694
JZ34	KX359594 ^b	KX965695
JZ36	KX359595 ^b	KX965696
N1	KU252662 ^a	KU298443
FF1	KU252600 ^a	KX965697
FF3	KU252661 ^a	KX965698

^a Complete 16S rRNA gene sequences (1,400 – 1,500 bp).

^b Partial 16S rRNA gene sequences (600 – 800 bp).

^c Partial *rpoB* gene sequences (300 – 400 bp).