

Table S1. Descriptive analysis of the sequence data

Gene	Number of alleles	Fragment analyzed (bp)	Polymorphic sites	Percent of variation sites (%)	Mean G+C content (mol%)
<i>purF</i>	5	757	118	15.58	60.42
<i>dnaJ1</i>	4	387	8	2.06	66.07
<i>rpoB</i>	8	860	35	4.07	64.49
<i>fusA</i>	6	644	37	5.74	60.02
<i>ileS</i>	10	542	37	6.82	61.95
<i>clpC</i>	8	524	21	4.01	62.58
<i>gyrB</i>	7	402	16	4.00	62.75
Total	48	4116	272	7.57	62.61

Table S2. Allelic profiles and frequencies of ST_s

ST	Allelic profile							Frequency ^a	%
	<i>dnaJ1</i>	<i>clpC</i>	<i>fusA</i>	<i>gyrB</i>	<i>ileS</i>	<i>purF</i>	<i>rpoB</i>		
1	5	1	4	1	9	4	5	4	4.88
2	2	1	4	1	3	1	5	1	1.22
3	5	3	1	2	3	7	1	5	6.1
4	5	3	1	2	3	2	1	2	2.44
5	2	1	3	5	6	2	1	3	3.66
6	5	1	3	1	8	8	1	3	3.66
7	1	1	1	1	8	8	1	1	1.22
8	5	1	2	1	2	2	2	1	1.22
9	5	1	1	6	2	4	1	5	6.1
10	5	1	2	1	2	7	1	3	3.66
11	5	1	5	2	6	3	1	1	1.22
12	5	1	5	1	6	7	1	1	1.22
13	5	1	2	5	6	2	1	1	1.22
14	5	4	5	2	3	3	1	1	1.22
15	2	1	2	1	4	7	6	3	3.66
16	5	1	1	3	3	4	5	1	1.22
17	5	1	1	1	10	2	4	3	3.66

18	5	1	1	1	9	4	4	1	1.22
19	5	1	1	3	10	4	5	1	1.22
20	5	1	1	3	9	4	7	1	1.22
21	5	1	1	1	10	2	1	1	1.22
22	5	1	1	2	5	2	1	8	9.76
23	5	3	1	2	5	2	1	2	2.44
24	5	1	1	3	5	4	5	1	1.22
25	5	1	3	5	2	4	1	9	10.98
26	5	1	6	5	2	4	1	2	2.44
27	5	1	3	5	2	4	4	1	1.22
28	5	1	8	4	7	6	3	2	2.44
29	5	1	3	5	7	4	1	1	1.22
30	3	2	8	4	7	6	3	3	3.66
31	1	1	1	1	1	1	1	5	6.1
32	1	1	1	1	1	5	1	1	1.22
33	4	1	1	6	1	1	1	1	1.22
34	5	1	7	5	2	4	1	2	2.44
35	5	1	1	5	2	4	1	1	1.22

^a Frequency indicates the number of individual strains generated the same STs.