

Table S1: Allelic combinations of sequence types (ST) and their Genbank accession numbers

Strains	Sequence Types	Combinations of alleles						
<i>E. coli</i>		<i>adk</i>	<i>fumC</i>	<i>gurB</i>	<i>icd</i>	<i>mdh</i>	<i>purA</i>	<i>recA</i>
	ST648	92	4	87	96	70	58	2
	Genbank accession number	MZ346383.1	MW519917.1	MF381225.1	MF381234.1	MH561342.1	MZ346152.1	KM072943.1
	ST8149	64	11	5	8	693	8	35
	Genbank accession number	LC187169.1	MW557688.1	MW519952.1	MW557701.1	MN315061.1	KM072796.1	AB648888.1
	ST2914	34	36	207	87	67	16	4
	Genbank accession number	KF914279.1	MH561275.1	MG194495.1	LC187109.1	MG204771.1	LC187111.1	MH561375.1
	ST155	6	4	14	16	24	8	14
	Genbank accession number	MW519923.1	MW519917.1	MN529017.1	MW557695.1	MN315063.1	KM072796.1	MN315096.1
	ST23	6	4	12	1	20	13	7
Genbank accession number	MW519923.1	MW519917.1	MN315005.1	MW519943.1	MN529105.1	MN315064.1	OU342919.1	
<i>K. pneumoniae</i>		<i>gapA</i>	<i>infB</i>	<i>mdh</i>	<i>pgi</i>	<i>phoE</i>	<i>rpoB</i>	<i>tonB</i>
	ST2010	17	55	39	20	243	18	156
	Genbank accession number	HG933781.1	LC049012.1	AM051118.1	LC049121.1	KT270588.1	MF345933.1	FJ483747.1
	ST551	3	1	1	1	9	4	135
	Genbank accession number	LT615348.1	AJ227992.1	AJ890392.1	AJ890402.1	AJ890421.1	MT882043.1	AJ890491.1

