

Table S3. Polymerase chain reaction (PCR) detection¹ of six iron-acquisition genes in *Klebsiella pneumoniae* isolates tested in this study.

Isolate	Farm	<i>iucA</i>	<i>entB</i>	<i>fepA</i>	<i>ybtS</i>	<i>psn</i>	<i>kfu</i>	Number of the genes detected by PCR
#1	A	-	-	-	-	-	-	0
#2	A	-	-	-	-	-	-	0
#3	A	-	-	-	-	-	-	0
#4	A	-	-	-	-	-	-	0
#5	A	-	-	+	-	-	-	1
#6	A	-	+	+	-	-	+	3
#7	A	-	+	+	+	+	-	4
#8	A	-	+	+	+	+	+	5
#9	B	-	+	-	-	-	-	1
#10	B	-	-	+	-	-	+	2
#11	B	-	+	+	-	-	-	2
#12	B	-	+	+	-	-	-	2
#13	C	-	-	+	-	-	-	1
#14	C	-	+	+	-	-	-	2
#15	C	-	+	+	-	-	-	2
#16	C	-	+	+	-	-	+	3

#17	D	-	+	-	-	+	-	2
#18	D	-	+	+	-	-	-	2
#19	D	-	+	+	-	-	-	2
#20	E	-	-	+	-	-	-	1
#21	E	-	+	+	-	-	-	2
#22	E	-	+	+	-	-	+	3
#23	F	-	-	-	-	-	-	0
#24	F	-	+	+	-	-	+	3
#25	G	-	-	+	-	-	+	2
#26	G	-	+	+	-	-	-	2
#27	H	-	-	-	-	-	-	0
#28	I	-	-	-	-	+	-	1
#29	J	-	+	-	-	-	-	1
#30	K	-	+	-	-	-	+	2
#31	L	-	+	+	-	-	-	2
#32	M	-	+	+	-	-	-	2
#33	N	-	+	-	-	+	+	3

#34	O	-	+	+	-	-	+	3
#35	P	-	+	+	-	+	+	4
#36	Q	-	+	+	+	+	+	5
#37	R	-	+	+	+	+	+	5
Proportion ² (%)	0	67.6	67.6	10.8	21.6	35.1	2.6 (1.7) ³	

¹+ and -, Each iron-acquisition gene is detected or not detected by the PCR test, respectively.

²Proportion of the isolates with PCR-detection of each iron-acquisition gene, out of 37 strains tested in this study.

³Mean (standard deviation) of PCR-detection numbers of iron-acquisition genes in 37 isolates.